

4th Edition

Handbook of ICP-QQQ Applications using the Agilent 8800 and 8900

Primer



Foreword

Agilent Technologies launched its 8800 Triple Quadrupole ICP-MS (ICP-QQQ) at the 2012 Winter Conference on Plasma Spectrochemistry in Tucson, Arizona, USA. At that time, ICP-MS had already been around for almost three decades and was widely praised for its low limits of detection. In fact, it was considered as the technique-par-excellence for multi-element (ultra-)trace analysis in a wide variety of fields. However, spectral interferences were still causing concern in some applications. Significant progress had been made in providing ICP-MS users with adequate tools to cope with spectral overlaps compared to the early commercial instruments introduced in 1983. By using a double-focusing sector-field mass spectrometer instead of a quadrupole filter for mass analysis, many spectral interferences can be resolved, but this approach requires expensive instrumentation. Quadrupole-based instruments could be equipped with a multipole-based collision/reaction cell (CRC), which alleviated spectral interferences to a significant extent, for instance, by using a non-reactive collision gas such as helium to slow down polyatomic interfering ions to a larger extent than the atomic analyte ions, such that the former could be selectively discriminated against on the basis of their lower kinetic energy. The analytical community first saw Agilent's 8800 ICP-QQQ instrumentation as an improved version of a guadrupole-based ICP-MS equipped with a CRC. But the unique applications being performed using Agilent ICP-QQQ instruments installed in hundreds of laboratories across industry, research and academia clearly demonstrates that it is much more than that.

In Agilent's ICP-QQQ, an octopole CRC is preceded by an additional quadrupole, enabling double mass selection, i.e. before the ions enter the CRC and afterwards. When the first quadrupole is used as an ion guide only, the ICP-QQQ system can be used as a "traditional" quadrupole-based ICP-MS instrument. This mode could be useful for carrying out routine analysis not significantly challenged by spectral interferences. When operated in tandem or MS/MS mode, however, the double mass selection only allows the analyte ion and the interfering ion(s) with the same mass-to-charge ratio to enter the CRC; all ions with a different mass-to-charge ratio are removed at this stage. Consequently, the control over the processes in the cell is greatly improved as the reaction of other (e.g. matrix) ions with the cell gas no longer hinders the desired reaction process. In case of a mass-shift reaction—i.e. chemical conversion of the analyte ion into a reaction product ion that can be measured interference-free at another mass-to-charge ratio—the absence of other ions at the new "location" of the product ion in the mass spectrum is guaranteed. As a result, interesting but challenging elements, such as S and P in biochemical applications, As and Se in environmental and food applications, or Si in nanoparticle applications can be easily assayed, interference-free

Profiting from the analytical advantages offered by MS/MS functionality, some ICP-QQQ users have demonstrated a larger degree of creativity by using very reactive gases such as $\rm NH_3$ or $\rm CH_3F$ in the CRC and monitoring reaction product ions at much higher mass-to-charge ratios than could be adequately exploited previously. Although this might initially sound complicated, the ICP-QQQ's software offers tools like product ion scanning, precursor ion scanning and differential mass scanning that provide the user with a clear insight into the reactions proceeding in the cell and allow the product ion that will provide the best, often unprecedented limits of detection to be easily identified. This level of freedom and ease of use leads to a situation in which every type of spectral overlap — whether caused by a polyatomic ion, doubly charged ion or isobaric nuclide — can be successfully overcome. Moreover, ICP-QQQ users have also been charmed by the additional advantages provided by this type of instrumentation, such as the unparalleled abundance sensitivity, which is an added benefit of double mass selection.

In 2016, the 8800 ICP-QQQ was replaced by the Agilent 8900 ICP-QQQ series. While maintaining the performance to resolve spectral interferences, this second generation ICP-QQQ instrument provides enhanced sensitivity and a faster detector system with a 100 μ s minimum dwell time. The latter feature is of specific importance in single-nanoparticle analysis, a rapidly emerging type of application, and in handling fast transient signals, such as those generated via laser ablation systems equipped with ultra-fast ablation cells.

In my opinion, ICP-QQQ has not only fulfilled its initial promises, but has greatly surpassed the anticipations of the diverse community of ICP-MS users.

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Introduction to Agilent Triple Quadrupole ICP-MS

The Agilent 8800 ICP-QQQ signaled a major advance in ICP-MS technology and redefined performance for trace element analysis. Compared to existing quadrupole ICP-MS instruments, the 8800 ICP-QQQ offered significant analytical benefits across many applications in both industry and academic research laboratories. Building on this success, the Agilent 8900 ICP-QQQ was launched four years later, in 2016.

Configuration of Agilent's ICP-QQQ instrumentation

According to IUPAC (term 538 from the 2013 Recommendations), a triple quadrupole mass spectrometer is a "Tandem mass spectrometer comprising two transmission quadrupole mass spectrometers in series, with a (non-selecting) RF-only quadrupole (or other multipole) between them to act as a collision cell."

The cell containing the ion guide—the Octopole Reaction System (ORS) in the case of Agilent ICP-QQQ—can be pressurized with a collision or reaction gas to allow the selective attenuation of potential interfering ions.

In MS/MS operation, where both quadrupoles are operated as unit mass filters, ions at the target analyte mass are selected by the first quadrupole (Q1) and passed to the ORS cell, where the analyte ions are separated from overlapping interfering ions. The resulting product ions that emerge from the cell are then filtered by the second quadrupole (Q2) before being passed to the detector. This configuration releases the full potential of reaction cell gas methods to resolve spectroscopic interferences including isobaric and doubly-charged interferences, as well as polyatomic ion overlaps.

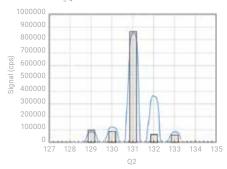
As a result, ICP-QQQ can determine a wider range of analytes at much lower concentrations with greater reliability and higher confidence.

Agilent introduced the world's first triple quadrupole ICP-MS (ICP-QQQ) in 2012.



Figure 1. Cutaway diagram of the Agilent 8900 ICP-QQQ.

Ti + (NH)(NH₂)₄ spectrum; Δ m = 2.1 amu of Q1



Ti + (NH)(NH₂)₄ spectrum; $\Delta m = 0.7$ amu of Q1

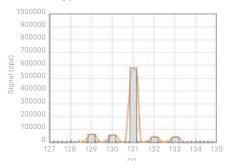


Figure 2. Ti + (NH)(NH₃)₄ product ion spectrum of five Ti isotopes obtained with NH₃ cell gas mode: ⁴⁶Ti (8.25%), ⁴⁷Ti (7.44%), ⁴⁸Ti (73.72), ⁴⁹Ti (5.41%), and ⁵⁰Ti (4.29%). A Ti standard solution was analyzed using ICP-QQQ with NH₃ cell gas. Top: Q1 was set at Δ m = 2.1 amu. Bottom: spectrum was acquired with Q1 set at Δ m = 0.7 amu.

ICP-MS/MS technology

The tandem MS configuration of Agilent ICP-QQQ instruments – with two fully functioning mass filters, one either side of the ORS cell – allows unprecedented control over the ions that enter the collision/reaction cell (CRC). Q1 rejects all non-target masses/elements, ensuring more consistent reaction processes in the CRC, even when the sample composition varies. Precise ion control by the first quadrupole, Q1, is crucial to the accurate analytical performance in MS/MS mode. For reliable, consistent control of reaction processes, Q1 must allow only ions at the target mass-to-charge ratio (*m/z*) to pass into the cell. Inefficient mass filtering would cause results to be compromised by interferences arising from non-target ions entering the cell. To ensure the most effective mass filtering and the best quality data, Agilent ICP-QQQ instruments use the same hyperbolic quadrupole mass spectrometer for Q1 and Q2. Both quadrupoles are placed in the high vacuum region to ensure optimum mass filtering. This arrangement allows both quadrupoles to operate at unit mass resolution and low abundance sensitivity without a significant loss of sensitivity.

The impact of varying Q1 mass filter performance is illustrated in the comparison of product ion spectra for Ti–NH $_3$ cluster ions shown in Figure 2. The product ion spectrum (top) was obtained using Q1 settings that provide compromised mass resolution, with Q1 passing all masses in a 2.1 amu window. This wider Q1 mass window allowed non-target ions to enter the cell, so the reaction processes and product ions formed were not under control. In this case, the overlapping ions were derived from different NH $_3$ clusters formed from the other Ti isotopes. For example, when 49 Ti is the target analyte but Q1 fails to exclude 48 Ti from the cell, 48 Ti(NH $_2$)(NH $_3$) $_4$ + forms in the cell and overlaps 49 Ti(NH)(NH $_3$) $_4$ + at m/z 132. In contrast, the Ti isotopic pattern (bottom) was obtained with Q1 set to operate with mass resolution of 0.7 amu; i.e. as a true mass filter providing genuine MS/MS operation. The Ti-NH $_3$ product ions fit the expected Ti isotopic pattern perfectly, confirming that each Ti isotope entered the cell in isolation to react with the

 ${
m NH_3}$ cell gas, i.e. ${
m ^{48}Ti^+} + {
m NH_3} \rightarrow {
m ^{48}Ti}({
m NH})({
m NH_3})_4^+$. The results show that if the mass resolution of Q1 is greater than 0.7 amu, precise analysis of the specific target analyte ion/isotope is impossible. Without true MS/MS operation (both quadrupoles operating with unit mass resolution), analytical results acquired for any analyte could be compromised because unexpected reaction product ions can be formed and cause overlap on the target product ion.

Second-generation triple quadrupole ICP-MS: the Agilent 8900 ICP-QQQ

Building on the success of the 8800, the 8900 ICP-QQQ provides performance and productivity improvements to address a wider range of applications:

- Double the sensitivity of the 8800: users of the 8900 can achieve lower detection limits or improve matrix robustness by diluting samples, without degrading detection capability. Note that the sensitivity of the Agilent 8900 Semiconductor configuration ICP-QQQ now exceeds 1Gcps/ppm.
- Axial Acceleration on the 8900 Advanced Applications and Semiconductor configurations controls the energy of ions in the cell. This increases the sensitivity in reaction cell mode and reduces potential product ion overlaps due to slow moving ions.
- Lower contribution from instrumental background: the 8900 ICP-QQQ is
 designed and manufactured to control background signals arising from the
 instrument itself. This attention to detail allows users to achieve even lower
 BECs than the 8800. The DL specification of the 8900 Advanced Applications
 and Semiconductor configurations for sulfur and silicon is < 50 ppt.
- 0.1 ms dwell time: the 8900 ICP-QQQ uses a new fast detector with fast time resolved analysis (TRA) capability suitable for the accurate analysis of single nanoparticles (sNPs). High speed is combined with effective interference removal and specialized software to process the signals and reveal the particle size and size distribution.

With true triple quadrupole performance, the advanced features and the robustness of the 8900 ICP-QQQ make it the world's most powerful and flexible multi-element analyzer. Agilent's Triple Quadrupole ICP-MS instruments will continue to open up new possibilities for analysts, especially for the most challenging applications.

Semiconductor

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Ultratrace Measurement of Calcium in Ultrapure Water

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Keywords

semiconductor, process chemicals, ultra pure water, UPW, calcium, method of standard additions, hydrogen on-mass

Introduction

In the semiconductor industry, the control of metal impurities in the process chemicals used in the manufacture of semiconductor devices is critical to achieve the required product performance and yield. As device performance is continually increasing, the required impurity control becomes ever more stringent. For example, metal content of the ultra-pure water (UPW) used in the manufacturing process must be at the sub-ppt level. ICP-MS is the standard technique used for the trace metals analysis of semiconductor chemicals and devices. The most common instrument and measurement technique used in the semiconductor industry is single quadrupole ICP-MS (ICP-QMS) with cool plasma. The cool plasma technique [1], developed in the mid 1990's, enables the quantification of key contaminant elements at the single ppt level. Collision and reaction cell ICP-QMS, developed from 2000 onwards, enabled the direct analysis of more complex semiconductor matrices, but did not improve on the DLs or BECs of cool plasma for low-matrix samples. To achieve measurement at the sub-ppt level, reduction of the BEC is required. As outlined in this paper, the Agilent 8800 ICP-QQQ provides new reaction cell technology that enables a significant reduction in the BEC that can be achieved for Ca, to 100 ppq.

Table 1. Cool plasma operating conditions.

Parameter	Unit	Tuning value
RF	w	600
Sampling depth	mm	18
Carrier gas flow	L/min	0.7
Make-up gas flow	L/min	1.0
Spray chamber temp.	°C	2

Experimental

Instrumentation: Agilent 8800 #200.

Plasma conditions: For the ultra-trace measurement of Ca, cool plasma operating conditions were used (Table 1). The sample was self-aspirated at a carrier gas flow rate of 0.7 L/min.

Reagents and sample preparation: A Ca standard was prepared in UPW acidified with 0.1% high purity HNO_3 . This was used to make 50 ppt and 100 ppt additions to a UPW blank acidified with 0.1% high purity HNO_3 .

Results and Discussion

Ultra-low BEC for Ca using MS/MS mode

Figure 1 shows the BECs obtained for Ca, measured at its major isotope of $^{40}\mathrm{Ca}$, using the method of standard additions (MSA) under three different operating conditions on the 8800 ICP-QQC: Single Quad mode with no cell gas, MS/MS mode with no cell gas, and finally MS/MS mode with a $\mathrm{H_2}$ cell gas flow of 1 mL/min. The Single Quad mode uses operating conditions with Q1 acting as an ion guide, to emulate the Agilent 7700 ICP-QMS. The obtained BEC of 6.8 ppt is similar to that routinely achieved with the Agilent 7700 operated in cool plasma mode.

Using MS/MS mode (without cell gas) improved the Ca BEC to 1.4 ppt. MS/MS mode with $\rm H_2$ at 1 mL/min in the cell further improved the BEC down to 0.041 ppt (41 ppq). The obtained MSA plot is shown in Figure 2. The Agilent 8800 ICP-QQQ in MS/MS mode with $\rm H_2$ cell gas achieved a BEC for Ca in UPW two orders of magnitude lower than the BEC obtained using conventional ICP-QMS.

Figure 3 shows the spectrum obtained for UPW using cool plasma conditions in Single Quad mode with no cell gas. As can be seen, Ar^+ (m/z 40) is suppressed under the lower temperature plasma conditions, but two intense background peaks are observed at m/z = 19 and 30. These are (H_2O) H^+ and NO^+ respectively. In Single Quad mode, all ions formed in the plasma, including these two intense ions, pass through to the cell. Even with no gas added to the cell, a reaction occurs in the cell which causes a new interfering ion at m/z = 40. The likely reaction occurring in the cell is: $NO^+ + Ar \rightarrow Ar^+ + NO$ (charge transfer reaction), which increases the BEC for Ca by several ppt. Although the ionization potential (IP) of NO (IP = 9.26 eV) is lower than that of Ar (IP = 15.7 eV), a metastable ion, NO^+ , exists close to the ionization potential of Ar [2]. So it is reasonable to assume that the charge transfer reaction shown occurs in the cell.

With MS/MS mode on the 8800 ICP-QQQ, Q1 rejects all non-target ions such as NO $^+$ and (H $_2$ O)H $^+$, preventing unwanted reactions from occurring in the cell, which lowers the Ca BEC. The addition of H $_2$ in the cell also removes any residual 40 Ar $^+$ that is formed even under cool plasma conditions.

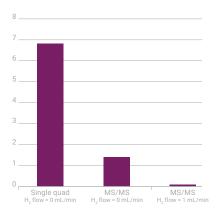


Figure 1. BECs for Ca obtained using Single Quad mode with no cell gas [6.8 ppt], MS/MS mode with no cell gas [1.4 ppt], and MS/MS mode with an $\rm H_2$ cell gas flow of 1 mL/min [0.041 ppt].

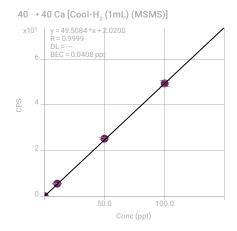
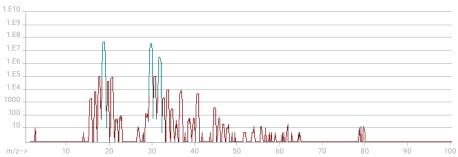


Figure 2. MSA calibration plot for Ca using MS/MS mode with H₂ flow of 1 mL/min.

[1] Spectrum No. 1 [187.530 sec]DIW_COOL.D/Tune#1[CPS][Log]



 $\textbf{Figure 3.} \ Spectrum \ of \ UPW \ acquired \ using \ cool \ plasma \ conditions \ in \ Single \ Quad \ mode \ with \ no \ gas \ mode.$

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Determination of Ti, V and Cr in 9.8% Sulfuric Acid

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Keywords

semiconductor, process chemicals, ultra pure water, UPW, calcium, method of standard additions, hydrogen on-mass

Introduction

High purity $\rm H_2SO_4$ is frequently used in the manufacturing of semiconductor devices, in processes such as the removal of organic substances from the surface of silicon wafers. The required metallic impurity level is lower than 100 ppt in the concentrated (usually 98%) acid. ICP-MS is the technique of choice for the measurement of trace metal impurities in semiconductor process chemicals. There are, however, some limitations for the measurement of elements such as Ti, V and Cr in $\rm H_2SO_4$. Because of its high viscosity of 27 cP, it is not possible to introduce $\rm H_2SO_4$ directly into the ICP without dilution. A 10 times dilution in UPW is normally applied, thus the BEC of the calibration curve must be lower than 10 ppt in the 9.8% $\rm H_2SO_4$ solution measured. In addition, spectral interferences from $\rm SO^+$, $\rm S_2^+$ and $\rm ArS^+$ originating from $\rm H_2SO_4$ make it difficult to determine elements such as Ti and Cr at low concentration even by quadrupole ICP-MS (ICP-QMS) equipped with collision/reaction cell (CRC). As outlined in this report, the Agilent 8800 ICP-QQQ with MS/MS mode allows the successful determination of the most problematic elements including Ti, V and Cr in $\rm H_2SO_4$.

Experimental

Instrumentation: Agilent 8800 #200. Operating parameters are given in Table 1.

Reagents and sample preparation: Highly purified H_2SO_4 , TAMAPURE-AA-100 (98% H_2SO_4) was purchased from Tama Chemicals Co., Ltd. (Kanagawa, Japan). 5 g of H_2SO_4 was diluted by a factor of 10 in a chilled PFA bottle.

Table 1. ICP-QQQ operating conditions.

		0 ₂ MS/MS ¹⁾		NH ₃ MS/MS ²⁾
RF power	W		1600	
Sampling depth	mm		8	
Carrier gas flow rate	L/min		0.8	
Make-up gas flow rate	L/min		0.41	
Octopole bias V	V		-20	
KED	V		-20	
Не	mL/min	3		1
0,	mL/min	0.4		0
NH ₃	mL/min	0		3

^{1) 100%} O₂ (purity 99.995%)

Results and Discussion

Of the potential polyatomic interferences formed from the H_2SO_4 matrix, the SO+ ion is very stable and difficult to eliminate because its dissociation energy is as high as 5.44 eV. In addition, its ionization potential is 10.3 eV, which is almost the same as that of S, 10.36 eV. The spectral interferences caused by SO+ and SOH+ overlap with 48 Ti ($^{32}S^{16}O$), ^{51}V ($^{33}S^{18}O$, $^{34}S^{16}OH$ and $^{32}S^{18}OH$) and ^{52}Cr ($^{34}S^{18}O$). Quadrupole ICP-MS operating in He

^{2) 10%} NH₃ balanced with 90% He (purity 99.995%)

collision mode provides BECs of 60 ppt for 47 Ti (the BEC for the preferred isotope 48 Ti is much higher), 3 ppt for V and 8 ppt for Cr in 9.8% H_2SO_4 . The BEC of Ti, in particular, is not acceptable for producers and users of semiconductor grade H_2SO_4 .

Appropriate reaction gases to remove SO $^+$ successfully in ICP-QMS are difficult to find. NH $_3$ can reduce SO $^+$ by two orders of magnitude but the background signal remains too high for this application. Additionally, cluster ions of NH $_3$ such as N $_m$ H $_n$ produced by the reaction between Ar $^+$ and the NH $_3$ cell gas lead to new reaction product ion interferences that increase the background at m/z 51, for example.

The 8800 ICP-QQQ operating in MS/MS mass-shift mode with NH $_3$ or O $_2$ reaction gas provides reliable and consistent measurement of Ti as 48 Ti 14 NH(14 NH $_3$) $_3$ (Figure 1) and Cr as 52 Cr 16 O in H $_2$ SO $_4$. Furthermore, in MS/MS mode, the Ar+ ion is removed by Q1, preventing it from reacting with NH $_3$ to form new product ion interferences in the cell. This reduces the background at m/z 51 improving the BEC for V, as shown in Figure 2. The final BECs obtained by ICP-QQQ in 9.8% high purity H $_2$ SO $_4$ are summarized in Table 2.

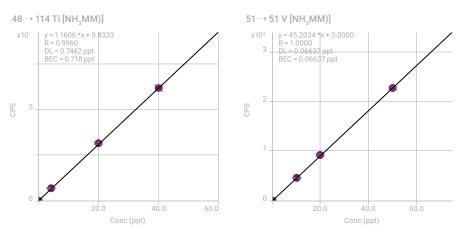


Figure 1. Calibration curve of Ti in 9.8% H₂SO₄.

Figure 2. Calibration curve of V in $9.8\% H_2SO_4$.

Table 2. BECs of Ti, V and Cr in 10x diluted 98% H₂SO₄, measured by ICP-QQQ.

Element	Ti	V	С
Mode (cell gas)	MS/MS (NH ₃)	MS/MS (NH ₃)	MS/MS (O ₂)
Measured ion	⁴⁸ Ti ¹⁴ NH(¹⁴ NH ₃) ₃ +	51 V +	⁵² Cr ¹⁶ O ⁺
Mass pair	Q1 = 48, Q2 = 114	Q1 = Q2 = 51	Q1 = 52, Q2 = 68
BEC - ppt	0.72	0.07	3.70

Conclusions

ICP-QQQ operating in MS/MS mode provides a reliable means for manufacturers of high purity $\rm H_2SO_4$ to guarantee all metallic impurity concentrations at less than 100 ppt in the concentrated acid.

More Information

Determination of challenging elements in ultrapure semiconductor grade sulfuric acid by Triple Quadrupole ICP-MS, Agilent publication, <u>5991-2819EN</u>.

Direct Determination of V, Cr, Ge and As in High-Purity 20% Hydrochloric Acid

Author

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Keywords

semiconductor, RCA Standard Clean, silicon wafer, hydrochloric acid, vanadium, chromium, germanium, arsenic, ammonia on-mass, ammonia mass-shift, oxygen mass-shift

Introduction

Since the 1970s, the RCA Standard Clean (SC) method has been used extensively in many countries for cleaning silicon wafer surfaces. SC-2 refers to a mixture of HCl and H₂O₂ that is used to remove ionic and metallic contaminants from the surface of silicon wafers. Because cleaning solutions are in direct contact with semiconductor devices, ultra high purity is required for these solutions. The SEMI standard Tier-D protocol for HCl defines the contaminant level to be <10 ppt for each element. Some elements have been very difficult to determine at ppt level by quadrupole ICP-MS (ICP-QMS) due to significant spectral interferences arising from the CI matrix, even when analyzed by ICP-MS equipped with a collision/ reaction cell (CRC). Consequently, some methods for the analysis of high purity HCl by ICP-MS have recommended sample pre-treatment steps to remove the chloride matrix, which can lead to analyte loss and sample contamination. In this study, ICP-QQQ was used to analyze undiluted HCl directly. Using MS/MS mode with mass-shift to remove polyatomic ions, the most problematic elements, such as V, Cr, Ge and As could be determined in HCl at single-figure ppt detection limits.

Experimental

Instrumentation: Agilent 8800 #200. Operating parameters are given in Table 1.

Reagents: 20% TAMAPURE-AA-100 HCl (metallic impurities are guaranteed to be below 100 ppt) was purchased from Tama Chemicals Co., Ltd. (Kanagawa, Japan). The undiluted HCl was introduced directly into the ICP-QQQ.

Table 1. ICP-QQQ operating conditions.

		O ₂ MS/MS ¹⁾		NH ₃ MS/MS ²⁾
RF power	w		1600	
Sampling depth	mm		8	
Carrier gas flow rate	L/min		0.8	
Make-up gas flow rate	L/min		0.41	
Octopole bias V	V		-20	
KED	V		-20	
He	mL/min	3		1
02	mL/min	0.4		0
NH ₃	mL/min	0		3

^{1) 100%} O₂ (purity 99.995%)

^{2) 10%} NH₃ balanced with 90% He (purity 99.995%)

Results and Discussion

Determination of BECs of V, Cr, Ge, and As in high purity HCl

ICP-QMS with a CRC using He collision mode can successfully eliminate some polyatomic ions such as ArCl [1], and the use of NH $_3$ as a reaction gas also works to remove the ClO $^+$ ion for the determination of V. However, ICP-QMS has some serious limitations when highly reactive cell gases (such as NH $_3$) are used in the CRC. Principal among these limitations is the fact that all ions enter the CRC, so predicted reaction pathways can be disrupted and new reaction product ion overlaps can be formed if the analyte levels in the sample change. ICP-QQQ with MS/MS removes this limitation, as the first quadrupole mass filter (Q1) allows precise selection of the ions that are allowed to enter the cell. This ensures that reaction processes and product ions are strictly controlled, dramatically improving detectability of the analyte ions shown in Table 2.

Table 2. Spectral interferences arising from the CI matrix on some key elements.

Polyatomic interference	m/z	Analyte ion
CIO+	51, 53	51 V +
CIOH+	52, 54	⁵² Cr+, (⁵⁴ Fe+)*
CICI+	70, 72, 74	⁷⁰ Ge ⁺ , ⁷² Ge ⁺ , ⁷⁴ Ge ⁺
ArCI+	75, 77	⁷⁵ As+, (⁷⁷ Se+)*

^{*}Alternative isotopes can be chosen to avoid spectral interferences on Fe and Se.

The MS/MS acquisition mode using O_2 or NH_3 as the reaction gas enables the determination of trace ^{51}V (measured directly as V+ using NH_3 cell gas), Cr as $^{52}Cr^{16}O^+$ (using O_2), Ge as $^{74}Ge^{14}NH_2^+$ (using NH_3) and As as $^{75}As^{16}O^+$ (using O_2). In the case of As, the $^{91}Zr^+$ ion is removed by Q1 (which is set to the As+ precursor ion mass of m/z 75), so the potential overlap from Zr on the AsO+ product ion at m/z 91 is also removed. The complete cut-off of cluster ions by Q1 also eliminates the possibility that $^{14}NH_2^{35}Cl$ is created in the cell, so the potential new product ion interference on ^{51}V is avoided. Representative calibration curves for V and Ge are shown in Figure 1. BECs and DLs determined by the ICP-QQQ for V, Cr, Ge, and As are given in Table 3.

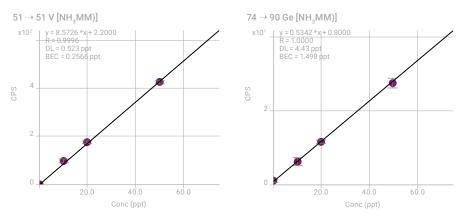


Figure 1. Calibration curves of V (NH2 on-mass mode) and Ge (NH2 mass-shift mode) in 20% HCl.

Table 3. BECs and DLs for V. Cr. Ge and As in 20% HCl.

Element	Ti	Cr	Ge	As
Mode (cell gas)	MS/MS (NH ₃)	MS/MS (O ₂)	MS/MS (NH ₃)	MS/MS (O ₂)
Measured ion	⁵¹ V +	⁵² Cr ¹⁶ O ⁺	⁷⁴ Ge ¹⁴ NH ₂ ⁺	⁵² Cr ¹⁶ O ⁺
Mass pair	Q1 = Q2 = 51	Q1 = 52, Q2 = 68	Q1 = 74, Q2 = 90	Q1 = 75, Q2 = 91
BEC - ppt	0.3	8.0	1.5	19.7
DL - ppt	0.5	1.1	4.4	3.4

Investigation of arsenic contamination

As the BEC for arsenic in high purity HCl was relatively high (Table 3), the signal count at m/z 91 was investigated further. The signals of the mass-pairs 75/75, 77/77, 75/91 and 77/91 were measured by ICP-QQQ with MS/MS, the mass pair number represents the set mass of Q1 followed by the set mass of Q2, so an MS/MS mode acquisition of mass pair 75/91 represents a mass-shift mode with Q1 = 75 and Q2 = 91, for example. The four mass pairs were measured in HCl blanks from three different lots, and the results are shown in Table 4. The following observations were made:

- 1. The ratio of the signal of 75/75 to 77/77 is around four, which is close to the ratio of the abundance of 35 Cl to 37 Cl, i.e. 3.13.
- 2. The ratio of the signal of 75/91 to 77/93 is 200-1000, which is far in excess of the ratio of 35 Cl to 37 Cl.
- 3. While the signals of 75/75 and 77/77 are similar for the three HCl blanks, those of 75/91 and 77/93 vary.

Table 4. Comparison of background counts (cps) in 3 different lots of 20% HCI*.

Mass pair	75->75	77->77	75->91	77->93
Sample A	509.3	133.5	584.4	2.5
Sample B	508.4	126.0	1172.6	1.9
Sample C	612.7	130.0	3175.6	2.6

^{*}All the samples were obtained from the new bottles of high purity HCl.

Finding #1 suggests that the remaining signal on 75/75 and 77/77 was mostly from ArCl*. This is a reasonable assumption since ArCl* doesn't react with $\rm O_2$ very efficiently so most ArCl* remains at the original masses of 75 and 77. Finding #2 suggests that the signal of 75/91 is not due to ArCl*. Assuming that all counts of 77/93 arise from $^{40}\rm Ar^{37}Cl$, the contribution of $^{40}\rm Ar^{35}Cl$ to the signal of 75/91 in the HCl blank is estimated to be just 7-8 cps, which is two orders of magnitude lower than the signal that is actually observed. Observation #3, together with #1 and #2, suggests the high count obtained for 75/91 in HCl is due to As impurity in the acid.

Reference

1. Direct analysis of trace metallic impurities in high purity hydrochloric acid by Agilent 7700s ICP-MS, Agilent application note, <u>5990-7354EN</u>.

Silicon Wafer Analysis by ICP-QQQ: Determination of Phosphorus and Titanium in a High Silicon Matrix

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Keywords

semiconductor, silicon wafer, phosphorus, titanium, Vapor Phase Decomposition, VPD, oxygen massshift

Introduction

The semiconductor industry first used ICP-MS for trace element analysis in the early 1980s. Nowadays the technique is widely used for control of trace impurities in materials and chemicals, particularly by silicon device manufacturers. The major challenge for quadrupole ICP-MS (ICP-QMS) is the presence of spectroscopic interferences on key contaminant elements, although performance has been gradually improved through developments such as cool plasma and collision/ reaction cells (CRC), and improved performance has also been provided by high resolution ICP-MS. Consequently metallic impurity control of silicon wafers can be successfully monitored by ICP-MS in the case of low silicon samples such as Vapor Phase Decomposition (VPD) of native silicon wafers. However, difficulties of Si-based spectral interferences, particularly on P and Ti, still affect the analysis of samples that contain high concentrations of Si, such as VPD samples of thermally oxidized wafers and samples relating to bulk silicon wafers. These interferences cannot be reduced adequately by ICP-QMS and have required HR-ICP-MS. In this paper, we evaluate triple quadrupole ICP-MS with MS/MS technology for the determination of ultratrace P and Ti in a high Si matrix.

Experimental

Instrumentation: Agilent 8800 #200 with an inert sample introduction kit including a low flow nebulizer (PFA-20) and a Pt/Ni skimmer cone. The actual sample uptake rate was 36 μ L/min. The sample was self-aspirated from an Agilent I-AS autosampler.

Plasma conditions: Robust tuning conditions were applied as summarized in Table 1.

Ion lens tune: Extract 1 = 0 V was used and other lens voltages were optimized using Auto tune.

Sample preparation: Silicon wafer samples were dissolved in TAMAPURE HF/HNO₃ and the final Si concentration was adjusted to 2000 ppm.

Table 1. Robust tuning conditions.

		O ₂ MS/MS		H ₂ MS/MS
RF power	W		1600	
Sampling depth	mm		8	
Carrier gas flow rate	L/min		0.6	
Make-up gas flow rate	L/min		0.6	
He	mL/min	3		0
02	mL/min	0.4		0
H ₂	mL/min	0		10

Results and Discussion

Phosphorus is monoisotopic at m/z 31, and suffers an interference from 30 SiH. While P+ can be detected as PO+ under cool plasma conditions, it is difficult to maintain cool plasma when the matrix concentration is high. Si sample solutions always contain HF, so Si will form SiF (IP: 7.54 eV) that also interferes with Ti. Table 2 shows the Si-based spectral interferences on P and Ti. Using the 8800 ICP-QQQ operating in MS/MS mode with O_2 mass-shift, P and Ti can be determined as their oxide ions, avoiding the Si-based interferences.

Table 2. Spectral interferences of Si on P and Ti.

Polyatomic interference	m/z	Analyte ion
³⁰ SiH⁺	31	³¹ P +
³⁰ Si ¹⁶ O ⁺	46	⁴⁶ Ti ⁺
²⁸ Si ¹⁹ F+, ³⁰ Si ¹⁶ OH+	47	⁴⁷ Ti+, ³¹ P ¹⁶ O+
²⁹ Si ¹⁹ F ⁺ , ³⁰ Si ¹⁸ O ⁺	48	⁴⁸ Ti ⁺
³⁰ Si ¹⁹ F ⁺	49	⁴⁹ Ti ⁺

For Ti analysis, Q1 is set to m/z 48, and so will transmit $^{48}\text{Ti}^+$ and any other interfering ions at mass 48, such as $^{29}\text{Si}^{19}\text{F}^+$ and $^{30}\text{Si}^{18}\text{O}^+$. But only ^{48}Ti reacts with oxygen in the CRC, producing the product ion $^{48}\text{Ti}^{16}\text{O}^+$, which is transmitted by setting Q2 to m/z 64. NH $_3$ can be used as an alternative reaction gas, as it produces $^{48}\text{Ti}^{14}\text{NH}^+$ that can be detected at m/z 63.

 $^{31}\text{P}^+$ reacts readily with O_2 to form $^{31}\text{P}^{16}\text{O}^+$. The selection of ions at m/z 31 by Q1 eliminates the spectral interference of $^{28}\text{Si}^{19}\text{F}$. However, ^{30}SiH passes through Q1 and reacts with O_2 to create $^{30}\text{Si}^{16}\text{OH}$. In order to determine P in a high Si matrix, H_2 mass-shift is a preferred option, despite the relatively low efficiency of production of PH_3^+ or PH_4^+ ions. The MSA calibration curves for P and Ti in a matrix of 2000 ppm Si are shown in Figure 1. The calculated BECs are summarized in Table 3. A long term stability test was carried out by analyzing a spiked sample repeatedly over five hours (Figure 2).

Table 3. BECs of P and Ti in 2000 ppm Si.

Element	Р	Ti
Mode (cell gas)	MS/MS (H ₂)	MS/MS (O ₂)
Measured ion	³¹ PH ₄ ⁺	⁴⁸ Ti ¹⁶ O ⁺
Mass pair	Q1 = 31, Q2 = 35	Q1 = 48, Q2 = 64
BEC - ppt	227	13

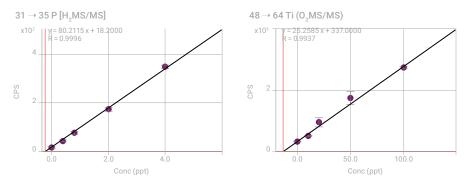


Figure 1. MSA curves of P and Ti in 2000 ppm Si.

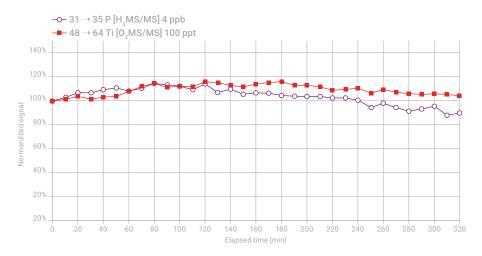


Figure 2. Five-hours test of P and Ti spiked in 2000 ppm Si.

Conclusions

The MS/MS mass-shift mode of the ICP-QQQ is effective for the determination of P, Ti and other trace elements in high purity silicon matrices, providing effective removal of the potential Si-based polyatomic interferences.

More Information

Improvement of ICP-MS detectability of phosphorus and titanium in high purity silicon samples using the Agilent 8800 Triple Quadrupole ICP-MS, Agilent publication <u>5991-2466EN</u>.

Analysis of Sulfur, Phosphorus, Silicon and Chlorine in N-methyl-2-pyrrolidone

Author

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Keywords

N-methyl-2-pyrrolidone, NMP, semiconductor, process chemicals, sulfur, phosphorus, silicon, chlorine, method of standard additions, oxygen mass-shift

Introduction

N-Methyl-2-Pyrrolidone (NMP), chemical formula: C_sH_9NO , is a stable, water-soluble organic solvent that is widely used in the pharmaceutical, petrochemical, polymer science and especially semiconductor industries. Electronic grade NMP is used by semiconductor manufacturers as a wafer cleaner and photo resist stripper and as such the solvent comes into direct contact with wafer surfaces. This requires NMP with the lowest possible trace metal (and non-metal) contaminant levels. ICP-MS is the technique of choice for the measurement of trace metal impurities in semiconductor process chemicals. It is a challenge, however for ICP-MS to measure non-metallic impurities such as sulfur, phosphorus, silicon, and chlorine in NMP. The low ionization efficiency of these elements greatly reduces analyte signal, while the elevated background signal (measured as background equivalent concentration, BEC) due to N-, O-, and C-based polyatomic ions formed from the NMP matrix makes low-level analysis even more difficult (Table 1).

Table 1. ICP-QMS BECs obtained in no gas mode for selected analytes in NMP.

Element	m/z	lonization potential (eV)	Ionization ra	tio (%) BEC without cell (ppm)	Interference
Si	28	8.152	87.9	>100	¹⁴ N ₂ +, ¹² C ¹⁶ O+
Р	31	10.487	28.8	0.39	¹⁴ N ¹⁶ OH ⁺ , COH ₃ ⁺
S	32	10.360	11.5	9.5	¹⁶ O ₂ +, NOH ₂ +
CI	35	12.967	0.46	0.26	¹⁶ O ¹⁸ OH ⁺

Experimental

Instrumentation: Agilent 8800 #200 with narrow injector (id =1.5 mm) torch (G3280-80080) typically used for the analysis of organic solvents. A C-flow 200 PFA nebulizer (G3285-80000) was used in self-aspiration mode. An option gas flow of 20% $\rm O_2$ in Ar was added to the carrier gas to prevent carbon build up on the interface cones.

Plasma conditions: NMP analysis requires hotter plasma conditions than normal. This was achieved by reducing Make-up Gas (MUGS) by 0.2 L/min. Plasma tuning conditions are summarized in Table 2.

Table 2. Plasma conditions for NMP analysis.

Parameter	Unit	Tuning value
RF	W	1550
Sampling depth	mm	8.0
Carrier gas flow	L/min	0.50
Make-up gas flow	L/min	0.10
Option gas flow	L/min	0.12 (12% of full scale)
Spray chamber temperature	°C	0

CRC conditions: Table 3 summarizes the cell tuning parameters (gas flow rate and voltages) used.

Table 3. CRC operating conditions.

Parameter	Unit	O ₂ reaction cell	l	H ₂ reaction cell	l
Method	-	On-mass	Mass-shift	On-mass	Mass-shift
Cell gas	-	0,		H ₂	
Gas flow rate	mL/min	0.30		4.0	
Octopole bias	V	-14		-10	
KED	V	-5	-5	0	-5

Reagents and sample preparation: Electronic industry grade NMP was distilled at 120 °C and acidified by adding high purity HNO₃ to a concentration of 1% w/w.

Results and Discussion

NMP was analyzed directly using the method of standard additions (MSA). Three replicate measurements (ten replicates for the blank) were acquired for S, P, Si and Cl using an integration time of 1 s per isotope.

P and S measurement in NMP

The mass-shift method using O_2 worked well for P and S measurement in NMP. The reactions of P and S with O_2 are exothermic, indicated by the negative value for Δ H, as shown below; therefore P+ and S+ are efficiently converted to their oxide ions, PO+ and SO+. P and S can be measured as the product ions, avoiding the original spectroscopic interferences on their elemental masses, m/z 31 and m/z 32.

$$P^+ + O_2 \rightarrow PO^+ + O$$
 $\Delta Hr = -3.17 \text{ eV}$
 $S^+ + O_2 \rightarrow SO^+ + O$ $\Delta Hr = -0.34 \text{ eV}$

In MS/MS mode, Q1 rejects 36 ArC+ before it can enter the cell, preventing it from overlapping S0+. This allows ICP-QQQ to control the reaction chemistry pathways and reaction product ions, ensuring that the analyte product ion is measured free from overlap, regardless of the levels of other co-existing analyte (or matrix) elements. MS/MS mode with the $\rm O_2$ mass-shift method achieved BECs of 0.55 ppb and 5.5 ppb for P and S respectively in NMP. The low BECs and linear calibration plots achieved in MS/MS mode also prove that the matrix-based interferences do not react with $\rm O_2$, allowing the analytes to be separated from the interferences.

Si measurement in NMP

 $\rm H_2$ cell gas was applied to the measurement of Si in NMP. The reaction kinetics for Si and its major interferences with $\rm H_2$ cell gas are shown below. The reaction rate data suggests that Si does not react with $\rm H_2$ cell gas (endothermic reaction indicated by the positive value for $\rm \Delta H$), and so could be measured in NMP using the direct, on-mass method. While the reaction of Si* with $\rm H_2$ is endothermic, the reactions of the major interfering ions on Si at mass 28 ($\rm N_2^+$ and CO+) are exothermic, and these interferences are therefore neutralized or reacted away.

Si⁺ + H₂
$$\rightarrow$$
 SiH⁺ + H Δ Hr = 1.30 eV
N₂⁺ + H₂ \rightarrow HN₂⁺ + H Δ Hr = -0.60 eV
CO⁺ + H₂ \rightarrow COH⁺ + H Δ Hr = -1.63eV

The results obtained are shown in Figure 2 (top). The $\rm H_2$ on-mass method achieved a BEC of 15.8 ppb for Si in NMP.

Oxygen cell gas was also tested of the measurement of Si in NMP. As shown below, the reaction of Si $^+$ with O $_2$ to form SiO $^+$ is endothermic. However, collisional processes in the cell provide additional energy which promotes the reaction, enabling the O $_2$ mass-shift method to be applied.

$$Si^+ + O_2 \rightarrow SiO^+ + O$$
 $\Delta Hr = 0.11 \text{ eV}$

Unfortunately a major interference on Si at m/z 28 (CO⁺) also reacts with O₂, so the BEC achieved using the O₂ mass-shift method to measure Si as SiO⁺ (Q1 = 28, Q2 = 44) was not satisfactory. Fortunately, another Si reaction product ion (SiO₂⁺) also forms and this can be measured at m/z 60 (Q1 = 28, Q2 = 60) giving a BEC of 11.9 ppb for Si in NMP (Figure 2, bottom).

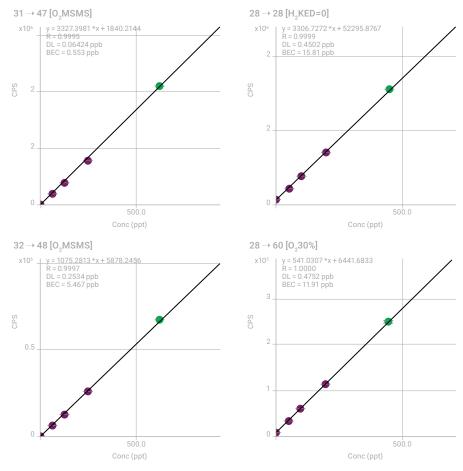


Figure 1. Calibration curve using MS/MS with O₂ mass-shift for P (top) and S (bottom) in NMP.

Figure 2. Calibration plots for Si in NMP. Top: H_2 on-mass method (Q1=Q2=28). Bottom: O_2 mass-shift method (Q1=28, Q2=60).

CI in NMP

Cl $^+$ reacts exothermically with H $_2$ to form HCl $^+$ as shown below. HCl $^+$ continues to react via a chain reaction to form H $_2$ Cl $^+$.

Cl⁺ + H₂ → HCl⁺ + H
$$\triangle$$
Hr = -0.17 eV
HCl⁺ + H₂ → H₂Cl⁺ + H \triangle Hr = -0.39 eV

Figure 3 (left) shows calibration plots obtained for CI in NMP using the $\rm H_2$ mass-shift method. The plot obtained using the $\rm O_2$ mass-shift method (Figure 3, below) is also shown for comparison. A slightly better BEC of 34.2 ppb was achieved with much higher sensitivity for CI in NMP using the $\rm H_2$ mass-shift method.

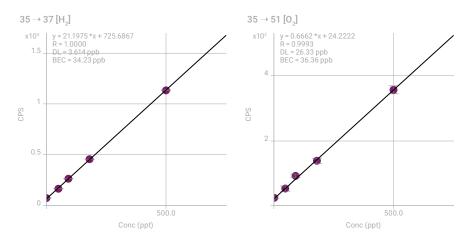


Figure 3. Calibration plots for Cl in NMP. Left: H_2 mass-shift method (Q1 = 35, Q2 = 37). Right: O_2 mass-shift method (Q1=35, Q2=51).

More Information

Trace level analysis of sulfur, phosphorus, silicon and chlorine in NMP using the Agilent 8800 Triple Quadrupole ICP-MS, 2013, Agilent publication, <u>5991-2303EN</u>.

Analysis of Silicon, Phosphorus and Sulfur in 20% Methanol

Authors

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Keywords

organic solvents, methanol, silicon, phosphorus, sulfur, hydrogen on-mass, oxygen mass-shift

Introduction

Analysis of organic solvents for trace metals presents a number of challenges to ICP-MS, many of which have been overcome to varying degrees on Agilent's 7700 Series quadrupole ICP-MS systems. However, even with these advances, several elements remain challenging in organic solvents, particularly silicon, phosphorus and sulfur. All three elements are subject to intense interferences from polyatomic ions based on carbon, nitrogen and oxygen, which are difficult to completely remove using conventional quadrupole ICP-MS (ICP-QMS). Examples include CO+, COH+, N_2 + and NO+ on silicon 28, 29 and 30; COH+, NOH+, N_2 H+, NO+ and CO+ on phosphorus 31 and O_2 +, NO+, NOH+ and NOH $_2$ + on sulfur 32 and 34. Additionally, phosphorus and sulfur have high first ionization potentials (IP) of 10.5 eV and 10.4 eV respectively, resulting in relatively poor sensitivity compared to more typical elements whose IPs are in the range of ~6 – 8 eV.

Experimental

Instrumentation: Agilent 8800 #200.

Plasma conditions and ion lens tune: RF power = 1550 W, Sampling depth = 8.0 mm and Carrier gas flow rate = 1.05 L/min were used with soft extraction tune, Extract 1 = 0 V and Extract 2 = -190 V.

Ultra pure methanol was spiked with silicon (Si), phosphorus (P) and sulfur (S) at 1, 5, 10 and 50 ppb and measured using the ICP-QQQ in several operational modes in order to evaluate the optimum conditions for the simultaneous analysis of all three analytes. Hydrogen and oxygen reaction gases were evaluated, with H_2 cell gas used in both Single Quad (SQ) and MS/MS modes. In addition, helium collision gas was investigated in both SQ and MS/MS mode to determine the effects of using MS/MS with a non-reactive cell gas.

The CRC conditions are outlined in Table 1, which includes the five analysis modes evaluated. Two Single Quad modes were tested, using both He and H_2 in the cell, to simulate the capability of a single quadrupole ICP-MS. In addition, three conditions using MS/MS mode were tested using H_2 , He and O_2 as cell gases.

Table 1. 8800 ICP-QQQ acquisition conditions tested, including five operational modes.

Parameter	Unit	He SQ	H ₂ SQ	H ₂ MS/MS	He MS/MS	O ₂ MS/MS
Acquisition mode		SQ	SQ	MS/MS	MS/MS	MS/MS
Cell gas		Не	H ₂	H ₂	He	02
Cell gas flow rate	mL/min	5.0	7.0	7.0	7.0	0.40
KED	V	5	0	0	5	-7

Results and Discussion

The BECs and DLs results are summarized in Tables 2–4, for silicon, phosphorus and sulfur respectively, for all 5 analysis modes tested. SQ and optimum MS/MS results are in bold type for comparison. A few mass-pairs were measured in each mode as shown. For example, Table 2 shows silicon monitored in MS/MS mode with $\rm O_2$ cell gas, using a mass-pair of Q1 = 28 and Q2 = 44. With Q1 set to m/z 28, only silicon 28 and any on-mass interferences are allowed to enter the ORS cell. The silicon 28 in the cell reacts with the oxygen cell gas to form SiO+, and Q2 is set to measure at Q1 + 16 (m/z = 44), ensuring that only the M + 16 O reaction transition is measured.

Table 2. DLs and BECs for silicon. Silicon was not measurable at the spiked concentrations in helium mode.

Mode	Mass or mass pair	BEC (ppb)	DL (ppb)	
H ₂ SQ	Q2=28	25.46	0.12	
H ₂ MS/MS	Q1=28, Q2=28	2.17	0.03	
O ₂ MS/MS	Q1=28, Q2=44	85.54	28.21	
O ₂ MS/MS	Q1=29, Q2=45	N/A	N/A	
O ₂ MS/MS	Q1=30, Q2=46	99.09	21.26	

Table 3. DLs and BECs for phosphorus. Phosphorus was not measurable at the spiked concentrations in $\rm H_2$ Single Quad mode.

Mode	Mass or mass pair	BEC (ppb)	DL (ppb)
He SQ	Q2=31	3.81	0.63
He MS/MS	Q1=31, Q2=31	2.99	0.72
H ₂ MS/MS	Q1=31, Q2=33	0.56	0.07
H ₂ MS/MS	Q1=31, Q2=34	0.58	0.67
O ₂ MS/MS	Q1=31, Q2=47	0.40	0.05
O ₂ MS/MS	Q1=31, Q2=63	0.41	0.02

Table 4. DLs and BECs for sulfur using MS/MS mode with O_2 cell gas. Sulfur was not measurable at the spiked concentrations in helium or hydrogen mode.

Mode (cell gas)	Mass or mass pair	BEC (ppb)	DL (ppb)
MS/MS (O ₂)	Q1 = 34, Q2 = 34	51.17	4.37
MS/MS (O ₂)	Q1 = 32, Q2 = 48	3.13	0.10
MS/MS (O ₂)	Q1 = 34, Q2 = 50	3.11	0.20

Sample calibration plots are displayed in Figure 1. They are displayed in pairs showing the results obtained using SQ mode with a typical cell gas (upper calibration), compared to MS/MS mode using the optimum conditions (lower calibration).

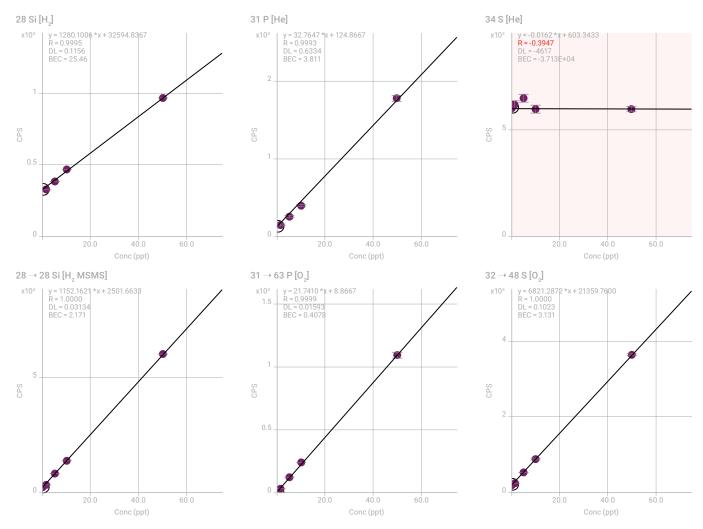


Figure 1. Calibration curves for Si, P and S showing SQ results (upper) compared with MS/MS results (lower). ³²S was not measurable at the spiked concentrations in methanol in SQ mode due to the intense ¹⁶O₂+ polyatomic interference.

Conclusions

It can be seen that in all cases the use of MS/MS mode significantly improves both the BEC and instrument detection limit when compared to Single Quad mode. The most notable improvement was for sulfur which cannot be measured at the spiked concentrations (1, 5, 10, 50 ppb) in SQ He mode due to the intense polyatomic background resulting from the methanol matrix. By contrast, on the 8800 ICP-QQQ using MS/MS mode with $\rm O_2$ mass-shift, S can be measured with a DL of 0.1 ppb.

More Information

Analysis of silicon, phosphorus and sulfur in 20% methanol using the Agilent 8800 Triple Quadrupole ICP-MS, Agilent publication, 5991-0320EN.

Ultratrace Measurement of Potassium and Other Elements in UPW Using ICP-QQQ in Cool Plasma/Reaction Mode

Authors

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Keywords

semiconductor, process chemicals, ultra pure water, UPW, potassium, cool plasma, ammonia on-mass

Introduction

The level of metal contaminants is strictly controlled in semicon device manufacturing processes, but the elements K, Ca and Fe are difficult to determine by ICP-MS due to argide interferences e.g. ArH+ on ³⁹K+, Ar+ on ⁴⁰Ca+ and ArO+ on ⁵⁶Fe+. Cool plasma employs a relatively low temperature plasma to remove the argide interferences allowing the analyst to measure these elements at trace levels. The low temperature plasma also reduces the background signal from any Easily Ionized Elements (EIEs) such as Li and Na that may deposit of on the interface of the ICP-MS. Even after the introduction of high concentration of EIEs, cool plasma ensures a low background level of these elements is maintained.

The Agilent 8800 ICP-QQQ provides improved cool plasma performance in combination with reaction cell technology, to enable a BEC of 30 ppq for K to be achieved in ultrapure water (UPW), and BECs at the ppq level for all the other elements studied: Li, Na, Mg, Al, Ca, Cr, Mn, Fe, Ni and Cu.

Table 1. Cool plasma operating conditions.

Parameter	Unit	Tuning value
RF	W	600
Carrier gas	L/min	0.7
Make-up gas	L/min	0.8
Sampling depth	(mm)	18
NH ₃ (10% in He) cell gas flow rate	mL/min	1

Experimental

Instrumentation: Agilent 8800 #200 (semiconductor configuration).

Plasma conditions: Cool plasma operating conditions (Table 1).

Reagents and sample preparation: The blanks and samples were acidified using high purity HNO_3 (TAMAPURE-AA-10, TAMA Chemicals Co. Ltd. Kanagawa, Japan). Standard solutions were prepared by serial dilution from a SPEX 331 mixed standard (SPEX CertiPrep, NJ, USA).

Cool plasma/NH3 reaction cell mode

Investigation of the signal at m/z 39 under cool plasma conditions indicated the presence of ³⁸ArH⁺ which decreases with lowering plasma temperature, indicating a reduction in the ionization of the polyatomic ion. However, there was also a contribution from a water cluster ion, $(H_3O^+)(H_2O)$, which is likely to form under low temperature plasma conditions. The combination of these two interferences means that there is no plasma temperature at which both interferences can be minimized (Figure 1). As the water cluster ion is known to react with deuterated ammonia (ND_3) via a fast proton transfer reaction [1], it was assumed that reaction with NH_3 would proceed at a similar rate, so this cell gas mode was investigated in order to remove the water cluster ion in cool plasma conditions.

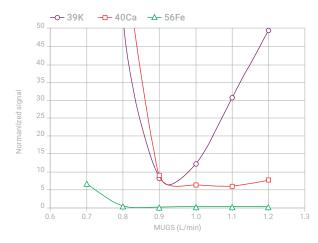


Figure 1. Investigation of the background signal under cool plasma conditions. BEC of K, Ca and Fe as a function of make-up gas (MUGS) flow rate.

Results and Discussion

Ultra-low BEC for K using MS/MS mode

Ammonia reaction gas mode under cool plasma conditions was used to determine K in UPW. The BEC of K was measured at 30 ppq. A comparative study carried out using a 7500cs quadrupole ICP-MS in cool plasma/NH $_3$ reaction mode achieved a BEC of 500 ppt for K [2]. It would be reasonable to attribute the improvement of BEC achieved with the 8800 to the MS/MS reaction capability of the ICP-QQQ. In conventional quadrupole ICP-MS, ions formed under cool plasma conditions enter the reaction cell and react with NH $_3$ or with impurity residues present in the cell to form product ions at m/z 39. MS/MS prevents any unwanted precursor ions from entering the cell, thus minimizing the production of undesired product ions.

Multielement analysis

The cool plasma/NH $_3$ reaction mode method was applied to the multielement analysis of UPW. As can be seen from the results in Table 2, a BEC < 150 ppq was achieved for all elements, including K, Ca and Fe.

Table 2. DL and BEC of elements in UPW.

Mass/Element	Sensitivity, cps/ppt	DL, ppt	BEC, ppt
7 Li	6.2	0.000	0.000
23 Na	94.0	0.014	0.035
24 Mg	44.0	0.010	0.005
27 AI	42.7	0.010	0.002
39 K	96.8	0.000	0.030
40 Ca	42.5	0.035	0.091
52 Cr	36.5	0.029	0.037
55 Mn	64.5	0.020	0.011
56 Fe	42.2	0.488	0.134
60 Ni	13.4	0.270	0.101
65 Cu	15.5	0.014	0.029

Conclusions

The Agilent 8800 ICP-QQQ was used to show the background signal at m/z 39 under cool plasma conditions was due to a water cluster ion, $H_3O(H_2O)^+$, which was removed using NH $_3$ cell gas. The ICP-QQQ BEC for ^{39}K was more than a factor of 10 lower than that achieved using a conventional quadrupole ICP-MS. This demonstrates the benefit of MS/MS mode for reaction gas methods: MS/MS mode prevents all non-target ions from entering the cell, and thereby eliminates the possibility of unwanted reactions from occurring.

References

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- 2. Junichi Takahashi et al., Use of collision reaction cell under cool plasma condition in ICP-MS, Asia Pacific Winter Plasma Conference 2008 (0-10)

More Information

Ultra trace measurement of potassium and other elements in ultrapure water using the Agilent 8800 ICP-QQQ in cool plasma reaction cell mode, Agilent publication 5991-5372EN.

GC-ICP-QQQ Achieves Sub-ppb Detection Limits for Hydride Gas Contaminants

Authors

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Emmett Soffey, Steve Wilbur and Chris Scanlon, Agilent Technologies Inc., USA

Keywords

semiconductor, petrochemical, phosphine, arsine, hydrogen sulfide, carbonyl sulfide, germane, silane, oxygen mass-shift, hydrogen on-mass

Introduction

Hydride gases, such as phosphine and arsine, are important contaminants in process chemicals used in both the petrochemical and semiconductor industries. The presence of phosphine, arsine, hydrogen sulfide, and carbonyl sulfide in polymer grade ethylene or propylene can have a deleterious effect on catalysts used in the production of polypropylene plastics. In the semiconductor industry, phosphine is used as a precursor for the deposition of group III-V compound semiconductors, and as a dopant in the manufacturing of semiconductor devices, such as diodes and transistors. The presence of unwanted hydride gas impurities can have a profound effect on the performance of the final device.

To date, measurement of these contaminants at ppb levels has been sufficient, but increasing competition within the industry and evolving performance criteria are pushing specifications ever lower. In addition, high purity gas manufacturers often require analytical detection limits 5-10 times lower than reported specifications. In anticipation of increasing industry demand for lower level detection, a new high sensitivity GC-ICP-QQQ method was developed for this application.

Experimental

Instrumentation: An Agilent 7890 GC was coupled to an Agilent 8800 #200 using the Agilent GC-ICP-MS interface.

Acquisition conditions: MS/MS mass-shift mode using oxygen as the cell gas for the measurement of Ge, As, P and S. MS/MS mode with hydrogen cell gas was used for the on-mass measurement of the primary isotope of Si at m/z 28.

 Table 1. Agilent 8800 ICP-QQQ operating conditions.

	O ₂ mode		H ₂ mode
RF power (W)		1350	
Sample depth (mm)		8.4	
Argon carrier (make-up) gas flow (L/min)		0.85	
Extract 1 (V)		-150	
Extract 2 (V)		-190	
Kinetic Energy Discrimination (V)	-4		0
Cell gas/flow (mL/min)	0.35		5.0

Reagents and sample preparation: Gas standards of silane, phosphine, germane, arsine (all balanced with H_2), and hydrogen sulfide and carbonyl sulfide (balanced with Ar) were supplied by Custom Gas Solutions at a nominal value of 10 ppmv. These standards were dynamically diluted in helium using a pressure/fixed restrictor based diluter supplied by Merlin MicroScience.

Results and Discussion

Low level phosphine analysis

The purpose of this experiment was to establish a detection limit for phosphine (PH $_3$) using GC-ICP-QQQ under ideal conditions. Q1 was set to m/z 31 (the precursor ion $^{31}P^{+}$) and Q2 was set to m/z 47 to measure the product ion $^{31}P^{16}O^{+}$. Since the eluting peaks are relatively narrow, with duration of no more than \sim 12 seconds, a maximum of 1 second was set for the total scan time. For the single element analysis of phosphine (measured as PO $^{+}$), an integration time of 1.0 second was used. A multi-point calibration curve was generated for PH $_3$ at concentrations of 8.2, 18.8 and 50.8 ppb. This covers the representative concentration range required for the measurement of this contaminant.

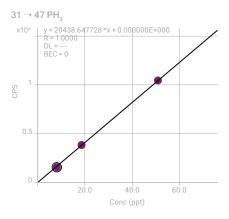


Figure 1. Phosphine calibration curve with an R value of 1.000 over the concentration range.

A low level phosphine standard (\sim 0.42 ppb) was also prepared, to allow the detection limit (DL) to be calculated. Two different methods for DL calculation were used:

- i. Two times the signal to noise (S/N) of the phosphine peak in the low level standard based on "Peak to Peak" noise method
- ii. The standard deviation of the concentrations measured in seven replicate analyses of the low level standard.

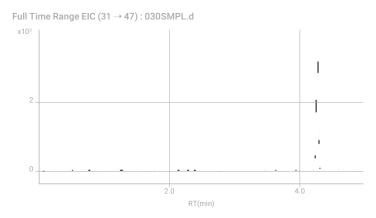


Figure 2. Chromatogram of 0.4 ppb PH3 standard. S/N: 96.9.

In the chromatogram shown in Figure 2, a S/N ratio of 96.9 was determined for the phosphine peak. Using the equation DL = $2 \times ((concentration of standard) / (S/N))$, a detection limit approximation of 8.67 ppt was calculated based on $2 \times ((0.42 \text{ ppb}) / (96.9))$. Using the standard deviation method, where multiple replicates of the low level standard were analyzed, the detection limit was 19 ppt.

Analysis of additional hydride gases

The GC-ICP-QQQ method was applied to the multielement analysis of germane, arsine and phosphine within a single analysis. Ge and As were measured as their $\rm O_2$ reaction product ions, GeO+ and AsO+, as was the case with P (PO+). Hydrogen sulfide (H₂S) and carbonyl sulfide (COS) were also analyzed using O₂ mass-shift mode, based on the ICP-QQQ measurement of sulfur as the $^{32}\rm S^{16}\rm O^{+}$ reaction product ion at m/z 48. For the analysis of silane, Si was measured directly (on-mass) at its major isotope $^{28}\rm Si$, using H₂ cell gas. The primary polyatomic interferences on $^{28}\rm Si^{+}$ are $^{12}\rm C^{16}\rm O^{+}$ and $^{14}\rm N_2^{+}$, due to the presence of CO₂, N₂ and O₂ in the argon supply and from air entrainment into the plasma. H₂ was selected as the reaction gas as both the CO+ and N₂+ interferences react readily with H₂ cell gas. Si+ remains unreactive and so can be measured, free from interferences, at its original mass.

Comparison of GC-ICP-QQQ and GC-ICP-MS DLs

For comparison purposes, H_2S , COS, PH_3 , GeH_4 , AsH_3 , and SiH_4 were analyzed by both GC-ICP-QQQ with the 8800 ICP-QQQ, and GC-ICP-MS using the same GC method with an Agilent 7900 conventional quadrupole ICP-MS. A summary of the detection limits (DLs) for both techniques is given in Table 1. For analytes where the background noise is very low (Ge-74, As-75), single digit ppt level detection limits are easily achieved using either GC-ICP-MS or GC-ICP-QQQ. However, for analytes that are prone to higher backgrounds (P-31 and S-32), significantly lower detection limits can be achieved by using MS/MS with O_2 cell gas and measuring the oxygen addition reaction product ions PO^+ and SO^+ in mass-shift mode. In addition, MS/MS mode with O_2 cell gas provides effective removal of background interferences at mass 28, allowing on-mass measurement of Si at its primary isotope.

Table 1. Detection limit comparison between GC-ICP-QQQ and GC-ICP-MS.

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MDL 2 X S/N 0.0013 MDL 2 X S/N 0.006	
SiH ₄ 28->28 (H ₂) 28 (H ₂)	
MDL 7 reps 0.14 MDL 7 reps 1.09	
MDL 2 X S/N 0.196 MDL 2 X S/N 1.18	

NA = not available

GC-ICP-QQQ sets benchmark detection limits

The significantly lower background and higher sensitivity of the Agilent 8800 ICP-QQQ resulted in a GC-ICP-QQQ method that shows a clear advantage for the determination of a range of contaminants in high purity gases at the low detection levels demanded by the industry. Compared to GC-ICP-MS with conventional quadrupole ICP-MS, GC-ICP-QQQ DLs for silane, phosphine, hydrogen sulfide, and carbonyl sulfide were lower by a factor of 5 to 10, with silane detection limits in the $\sim\!\!15$ ppt range.

More Information

Sub-ppb detection limits for hydride gas contaminants using GC-ICP-QQQ. Agilent Publication <u>5991-5849EN</u>.

Find out more about CONSCI at www.consci.com or contact William Geiger at bill@conscicorp.com

Ultralow Level Determination of Phosphorus, Sulfur, Silicon, and Chlorine

Author

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Keywords

sulfur, silicon, phosphorus, chlorine, ultralow level, UPW

Introduction

With the introduction of Agilent's second-generation ICP-QQQ instrument, the Agilent 8900 Triple Quadrupole ICP-MS, reaction cell operation with MS/MS mode has been further refined. This note describes the performance of the 8900 ICP-QQQ for the analysis of some of the most challenging elements for ICP-MS: phosphorus (P), sulfur (S), silicon (Si), and chlorine (Cl). As ICP-MS technology has developed, there has been a growing demand and expectation to measure these difficult elements, together with more conventional elements, in high purity chemicals and materials.

Experimental

Instrumentation: An Agilent 8900 Semiconductor configuration ICP-QQQ was used for all measurements. The sample introduction system comprised a PFA concentric nebulizer, a quartz spray chamber and torch, and platinum interface cones.

Tuning: Hot plasma conditions were used throughout this study. RF power = $1550 \, \text{W}$, sampling depth = $8 \, \text{mm}$, nebulizer gas flow rate = $0.70 \, \text{L/min}$, and make-up gas flow = $0.52 \, \text{L/min}$. Based on previous studies, oxygen (O_2) mass-shift mode was used for the analysis of P and S [1]; hydrogen (O_2) on-mass mode was used for Si; and Cl was determined using O_2 mass-shift mode [2]. Cell tuning conditions are summarized in Table 1.

Table 1. Cell tuning conditions.

Parameter	Unit	³¹ P	³² S	²⁸ Si	³⁵ Cl
Cell gas		02		H ₂	
Flow rate	mL/min	0.41		5	
Mass pair		(31047)	(32048)	(28□28)	(35037)
Octopole Bias	V	-3		-18	
KED	V	-8		0	
Axial Acceleration	V	1		0	

Results and Discussion

To prepare the ICP-QQQ for the analysis, a 1% HNO $_3$ solution was aspirated overnight to thoroughly clean the sample introduction system. Running the plasma for several hours would also help to remove any contaminants in the Ar gas flow line. P, S and Si were measured together, and Cl was analyzed in a separate batch since it benefited from an alkaline rinse between solutions. Figures 1 shows the calibration curves of the four elements in ultrapure water (UPW, supplied from Organo, Japan). The background equivalent concentrations (BEC) and detection limits (DL) are summarized in Table 2.

Table 2. BEC and DL of P, S, Si, and Cl in UPW.

		³¹ P	³² S	²⁸ Si	³⁵ Cl	
BEC	ppt	10.5	75.4	259	1830	
DL (n=10 x 1s integration)	ppt	3.3	5.5	14.7	280	

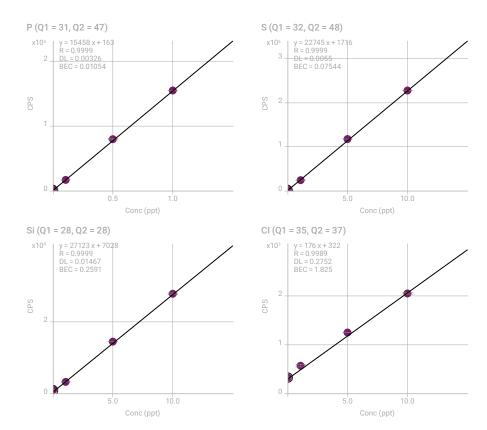


Figure 1. Calibration plots of P, S, Si, and Cl in UPW. All values in ug/L (ppb).

Conclusions

The Agilent 8900 ICP-QQQ operating in MS/MS mode with $\rm O_2$ and $\rm H_2$ cell gases successfully eliminated problematic spectral interferences on non-metallic impurities P, S, Si, and Cl in UPW. The results highlight the advanced performance of the second-generation ICP-QQQ for the analysis of challenging elements. The method delivered the lowest ever reported BECs for the four elements in UPW.

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More Information

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Determination of Ultratrace Elements in SEMI Grade 5 High Purity Hydrogen Peroxide

Author

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Keywords

SEMI, H₂O₂, hydrogen peroxide, semiconductor, high purity chemicals

Introduction

Hydrogen peroxide (H_2O_2) is one of the most important process chemicals used in semiconductor device manufacturing. As a strong oxidizer, it is used for cleaning silicon wafers, removing photoresists, and etching metallic copper on printed circuit boards.

Semiconductor Equipment and Materials International (SEMI) publishes standards regarding the specifications for semiconductor process chemicals including $\rm H_2O_2$ (SEMI C30-1110). SEMI Grade 5 is the highest purity level, with maximum contamination levels of 10 ppt for most trace elements. The industry therefore requires analytical methods capable of measuring the trace elements at single- or sub-ppt level background equivalent concentrations (BECs). SEMI C30-1110 also includes specifications for the maximum concentrations of sulfate and phosphate allowed in high purity $\rm H_2O_2$, with a limit of 30 ppb. This limit equates to elemental concentrations of sulfur and phosphorus of 10 ppb. These two contaminants are not currently measured by ICP-QMS. However, the recent development of triple quadrupole ICP-MS (ICP-QQQ) permits much lower limits of detection for S and P. It is now possible to monitor all SEMI elements using a single technique.

Experimental

Instrumentation: Agilent 8900 Semiconductor configuration ICP-QQQ.

Tuning: To achieve the lowest DLs, a multi-tune method was used. The tuning parameters are summarized in Table 1. For data acquisition, a 2 s integration time was used for all isotopes with three replicates (10 replicates for the blank to calculate the DLs).

Table 1. ICP-QQQ tuning conditions.

	Cool	Cool-NH ₃	Cool-NH ₃ (2)	No gas	H ₂	Не	0 ₂ (1)	0, (2)	
Scan mode	Single Q			MS/I	MS/MS				
RF power (W)		600			1500				
Nebulizer gas flow (L/min)				0.70	0.70				
Make-up gas flow (L/min.)		0.90				0.48			
Sampling depth (mm)		18.0				8.0			
Ex1 (V)	-15	50.0	-100.0	4.2	4.7	4.2	4.5	3.5	
Ex2 (V)	-18.0	-17.0	-12.0		-25	50.0		-120.0	
Omega (V)		-70.0		-140.0				-70.0	
Omega Bias (V)		2.0		10.0	8.0	-10.0	10.5	4.0	
Q1 Entrance (V)	-1	5.0			-50.0				
Cell gas	-	N	IH ₃	-	H ₂	He	C)2	
Cell gas flow (ml/min)		2.0	2.0		7.0	5.0	0.3	0.3	
Axial Acceleration (V)	0.0	1	.5		0.0		1.	.0	
KED (V)	15.0	-:	5.0	5.0	0.0	3.0	-7	.0	

Sample Preparation

TAMAPURE-AA-10 hydrogen peroxide (35%, Tama Chemicals, Japan) was used as the sample matrix. To stabilize the spiked elements, ultrapure nitric acid (TAMAPURE-AA-10) was added to the $\rm H_2O_2$ samples at one part of 70% HNO $_3$ to 1000.

Results and Discussion

Table 2 shows quantitative results and detection limits for the SEMI specification elements in high purity 35% $\rm H_2O_2$. Comparative quantitative results and DLs are also shown for the same elements in ultrapure water. Long-term stability was evaluated by measuring a $\rm H_2O_2$ sample spiked at 10 ppt for most elements and 100 ppt for sulfur. Calibration curves were generated at the beginning of the sequence. The spiked samples were then run as unknown samples for a total analysis period of 3 h 40 min. The RSDs of the 13 results are shown in Table 2 (Stability RSD %).

Table 2. ICP-QQQ tuning conditions.

					Н	Hydrogen peroxide		Ultrapure water	
Element (Q1	Q2	Scan mode	Tune	Conc.	DL	Stability	Conc.	DL
					(ppt)	(ppt)	RSD (%)	(ppt)	(ppt)
Li		7	single quad	Cool	< DL	0.003	4.7	< DL	0.004
В	11	11	MS/MS	No gas	7.7	0.69	8.1	4.6	0.57
Na		23	single quad	Cool	0.39	0.031	3.3	0.5	0.069
Mg		24	single quad	Cool	0.017	0.017	4.1	< DL	0.012
Al		27	single quad	Cool	0.39	0.071	2.9	0.11	0.11
Р :	31	47	MS/MS	0, (1)	4.2	0.89	3.3	3.4	0.91
s :	32	48	MS/MS	0, (1)	190	5.1	7.8	41	3.8
к :	39	39	MS/MS	cool+NH ₃ (2)	0.21	0.11	2.2	0.2	0.088
Ca	40	40	MS/MS	cool+NH ₃ (2)	< DL	0.23	1.9	< DL	0.10
Ti 4	48	64	MS/MS	0, (2)	0.097	0.045	2.6	< DL	0.028
V :	51	67	MS/MS	O ₂ (2)	0.067	0.027	2.6	< DL	0.023
Cr :	52	52	MS/MS	cool+NH ₃ (1)	0.13	0.075	3.5	< DL	0.031
Mn !	55	55	MS/MS	cool+NH ₃ (1)	< DL	0.012	2.7	< DL	0.004
Fe :	56	56	MS/MS	cool+NH ₃ (1)	0.13	0.074	3.3	< DL	0.027
Ni (60	60	MS/MS	cool+NH ₃ (1)	0.16	0.14	3.7	< DL	0.030
Cu	63	63	MS/MS	cool+NH ₃ (1)	< DL	0.048	5.0	0.19	0.18
Zn (64	64	MS/MS	Не	0.22	0.14	4.5	0.35	0.17
As	75	91	MS/MS	0, (2)	< DL	0.087	3.5	< DL	0.081
Cd	114	114	MS/MS	No gas	< DL	0.02	2.3	< DL	0.017
Sn	118	118	MS/MS	No gas	0.088	0.063	2.0	< DL	0.037
Sb	121	121	MS/MS	H ₂	< DL	0.015	1.6	< DL	0.022
Ва	138	138	MS/MS	H ₂	0.061	0.033	1.2	< DL	0.004
Pb :	208	208	MS/MS	No gas	0.081	0.053	1.0	0.056	0.035

Conclusions

All the elements specified in SEMI C30-1110 were measured at sub-ppt to ppt levels in high purity 35% hydrogen peroxide using the Agilent 8900 ICP-QQQ. For almost all elements, sub ppt quantitative results were obtained, with the remaining elements having single-ppt detection limits (except Si, 25 ppt). Reproducibility between 1.0 – 8.1 % RSD was obtained at the 10 ppt level (100 ppt for S) for the spiked analytes in a high purity 35% hydrogen peroxide sample analysis sequence that lasted 3 hours 40 minutes. This performance demonstrates the suitability of the Agilent 8900 Semiconductor configuration ICP-QQQ instrument for the routine analysis of the highest-purity semiconductor reagents and process chemicals.

More Information

Determination of ultra trace elements in high purity hydrogen peroxide with Agilent 8900 ICP-QQQ, Agilent publication, <u>5991-7701EN</u>.

Automated Ultratrace Element Analysis of Isopropyl Alcohol with the Agilent 8900 ICP-QQQ

Authors

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Riro Kobayashi IAS Inc, Japan Online calibration using the IAS Automated Standard Addition System (ASAS)

Introduction

Contamination control is critical in the semiconductor industry. Inorganic impurities are of particular concern, as they affect the electrical properties of the insulating and conducting layers from which semiconductor devices are made. Trace element contamination during wafer fabrication can therefore reduce the manufacturing yield and operational reliability of semiconductor devices. To minimize contamination, process chemicals must be monitored for ultratrace (ng/L; ppt) levels of elemental impurities.

Isopropyl alcohol (IPA) is an important organic solvent used in semiconductor manufacturing to remove organic and metallic residues and impurities from the surface of silicon wafers. Since IPA comes into direct contact with the wafer surface, the concentration of trace metals present in the solvent should be extremely low. For high purity Grade 4 IPA, SEMI standard C41-0705 specifies a maximum contaminant level of 100 ppt for each element (1). Delivering accurate analysis at these low concentrations requires a highly sensitive analytical instrument, together with a suitable clean laboratory environment, and advanced sample handling skills. Modern ICP-MS systems include predefined settings and auto-optimization routines to simplify operation. But the sample preparation, sample processing, and calibration steps still require a highly skilled analyst. Automating these steps would simplify the method, reducing the level of skill required for analysts to reliably perform the analysis.

Agilent ICP-MS systems can be integrated with various automated sample introduction systems, depending on a laboratory's requirements. Systems are available that automate a range of sample handling steps such as dilution, acidification, spiking, and calibration. One of the simplest, easiest-to-use and most cost-effective systems for automating semiconductor sample handling is the Automated Standard Addition System (ASAS) from IAS Inc. The ASAS can automatically add online spikes to generate a method of standard additions (MSA) or external calibration curve. In addition to simplifying the analysis, the automated sample introduction system decreases manual sample handling, reducing errors, and lowering the potential for sample contamination.

In this study, trace element impurities in IPA were quantified by online MSA using an IAS ASAS (Tokyo, Japan) and Agilent 8900 Triple Quadrupole ICP-MS (ICP-QQQ). The method allows the accurate and reliable quantification of ultratrace level impurities in IPA without requiring a highly skilled analyst.

Experimental

Reagents and samples

High purity IPA was prepared for analysis by distilling electronic-grade IPA in the lab. The IPA samples were introduced into the ICP-QQQ undiluted, to minimize the risk of contamination and to achieve the lowest possible detection limits (DLs).

A 1 μ g/L (ppb) mixed multi-element standard solution was used to create the MSA calibration spikes. The working standard solution was prepared by diluting a 10 ppm mixed multi-element standard (SPEX CertiPrep, Metuchen, NJ, US) with distilled IPA. To stabilize the spiked elements, nitric acid (68% ultrapure HNO₃) was added to the IPA samples at a final acid concentration of 1%.

The 1 ppb working standard solution was placed into a clean sample bottle and connected to the standard line of the ASAS. All MSA calibration (spike) solutions required for the analysis were automatically prepared and added online by the ASAS. Spike concentrations of 0, 5, 10, 20, and 50 ppt were added to the IPA sample. The sample preparation and analysis steps were performed in a Class 10,000 clean-room.

Instrumentation

An Agilent 8900 Semiconductor configuration ICP-QQQ instrument was used. The instrument was fitted with a glass concentric nebulizer (G1820-65138) self-aspirating with PFA sample tubing (G1820-65478; 0.3 mm id, 1.6 mm od).

A Peltier-cooled quartz spray chamber, quartz torch, platinum-tipped sampling and skimmer cones, and s-lens were used.

When organic solvents are analyzed, carbon in the sample aerosol can be deposited on the sampling cone, causing instability and signal drift. To prevent carbon deposition during the analysis of solvents such as IPA, oxygen is added to the carrier gas to oxidize the carbon in the plasma. Volatile organic solvents also cause a very high solvent vapor pressure in the spray chamber, leading to plasma instability. To reduce solvent vapor pressure and ensure reliable plasma ignition and operation, the spray chamber temperature is reduced to below zero degrees. In this work, the spray chamber was cooled to -5 °C using the Peltier device that is standard on all Agilent ICP-MS systems. For stable operation of the plasma, a torch with a narrow (1.5 mm) injector replaced the standard 2.5 mm injector torch.

Combining cool plasma with collision/reaction cell (CRC) operation has been shown to be a powerful mode for interference removal in ICP-MS (2). These conditions can also be used for the analysis of organic solvent samples, but such samples require more plasma energy to decompose the organic matrix. The analyst must balance reducing the plasma temperature enough to control the ionization of interfering species, while maintaining sufficient energy to decompose the matrix. With Agilent ICP-MS systems, the ShieldTorch System provides effective reduction of the plasma potential, so ionization of polyatomic ions is minimized, even at higher forward power. "Cool plasma" conditions provide better robustness and matrix tolerance on Agilent ICP-MS systems than on systems that do not have such effective control of plasma potential.

Furthermore, all Agilent ICP-MS systems have two separate gas controls contributing to the total "carrier" gas flow passing through the central, injector tube. The nebulizer gas flow (the flow that passes through the nebulizer and aspirates the sample) is adjusted to give optimum sample aspiration. The make-up gas flow is then optimized to control the total carrier gas flow that transports the sample aerosol through the central channel of the plasma. This total carrier flow, combined with the plasma power and sampling depth, determines the "coolness" of the plasma conditions.

In advanced semiconductor applications, the key requirement is to deliver the absolute lowest possible detection limits for each analyte. Laboratories measuring ultratrace levels of contaminant metals often use a multitune method, where several tuning steps are applied sequentially during the measurement of each solution. This approach allows the tuning conditions to be optimized for the removal of different types of interferences, while maintaining sensitivity for each analyte. In this work, several reaction cell gases (He, $\rm H_2$, $\rm O_2$, and $\rm NH_3$) were used for the analytes being measured.

Instrument tuning conditions are shown in Table 1 and other acquisition parameters are shown in Table 2.

Table 1. Agilent 8900 ICP-QQQ operating conditions.

	H ₂ (cool plasma*)	NH ₃ (cool plasma*)	O ₂ He	H ₂	He	No gas		
Scan type			MS/M	s				
RF power (W)			1500	1				
Sampling depth (mm)	18.0							
Nebulizer gas flow rate (L/ min)	0.70							
20% O ₂ Ar balance gas flow rate (L/min)	0.30 (30%)**							
Spray chamber temp (°C)			-5.0					
Make-up gas flow rate (L/min)	0.80 0.70 0.50							
Extract 1 (V)	-100							
Extract 2 (V)	-10.0							
Omega bias (V)	-70.0							
Omega lens (V)			4.0					
Q1 entrance (V)			-50.0	1				
He cell gas flow rate (mL/min)	-	1.0	12.0	-	5.0	-		
H ₂ cell gas flow rate (mL/min)	5.0	-	-	10.0	-	-		
NH ₃ cell gas flow rate (mL/ min)***	-	2 (20%)**	-	-	-	-		
O ₂ cell gas flow rate (mL/min)	-	-	0.075 (5%)**	-	-	-		
OctP bias (V)	-18	-5	-3	-30	-20	-10		
Axial acceleration (V)	1.0 0							
Energy discrimination (V)	0		-10			3		

Table 2. Acquisition parameters.

Parameter	Setting
Q2 peak pattern	1 point
Replicates	3 (spiked samples) 10 (unspiked sample)
Sweeps/replicate	10
Integration time (s)	1 (all elements except phosphorus) 10 (phosphorus)

^{*} Optimum cool plasma conditions were achieved by adjusting the make-up gas flow while maintaining a high forward power setting.

^{**} Values in parentheses are % of the maximum flow of the gas controller, as displayed in the tuning pane of ICP-MS MassHunter.

^{*** 10%} NH₃ balanced with 90% He.

Automated Standard Addition System (ASAS)

The IAS ASAS is an automated online sample processing device. It uses a precise, microflow syringe pump to add specific volumes of spike solutions or diluent to the sample flow as it passes to the ICP-MS nebulizer. The small footprint of the ASAS allows it to be easily positioned between the autosampler and ICP-MS, as shown in Figure 1. This arrangement is beneficial in the small workspace typically available in semiconductor clean-rooms. Once connected in line, the ASAS can be used to automatically generate a calibration curve using either external standards or MSA.

MSA calibrations have the advantage of exact matrix matching, since the calibration is created in the actual sample matrix. However, conventional manual MSA spiking is often regarded as complicated and time-consuming. With automatic spike additions using ASAS, the complexity is eliminated. Also, the Agilent ICP-MS MassHunter software allows an MSA calibration in one sample to be automatically converted to an external calibration. This function allows other samples of the same type to be run against the MSA calibration, without requiring the subsequent samples to be spiked individually. With these two improvements, MSA can be as fast and easy to run as conventional external calibration.

Spike recoveries are typically carried out as a routine performance check during semiconductor chemical analysis. This can be automated to simplify and speed up the analysis. The ASAS microvolume syringe pump adds the spike to the continuously flowing sample stream, so the risk of sample contamination and errors is minimized.

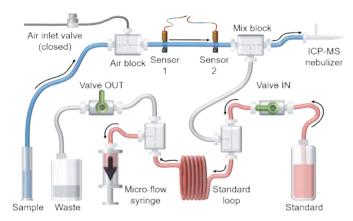


Figure 1. An Agilent ICP-MS fitted with an IAS ASAS automated standard addition system and the Agilent I-AS integrated autosampler.

ASAS operation consists of the following four steps:

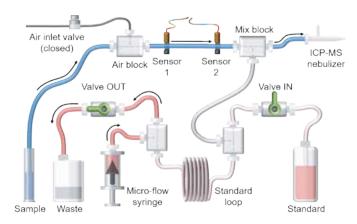
Step 1: Loop is filled with the standard solution.

"Valve IN" in the standard line opens and the syringe pump activates. This loads the standard solution along a dedicated uptake line from the standard bottle to the loop.



Step 2: Excess standard solution is pumped to waste.

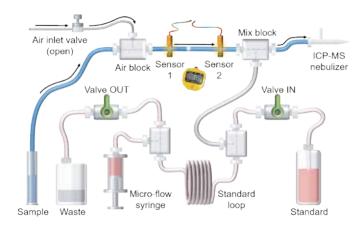
"Valve OUT" in the standard line opens, "Valve IN" closes, and the syringe pump discharges, pumping the remaining standard solution to the waste bottle—bypassing the loop.



Step 3: Sample flow rate is measured to allow calculation of MSA spike volumes.

To minimize the potential for contamination from peristaltic pump tubing, high-purity samples are usually introduced using self aspiration. This means that the flow rate varies, depending on the sample viscosity and tubing length. To allow the MSA spike additions to be calculated accurately, the ASAS system first measures the sample flow rate, as follows:

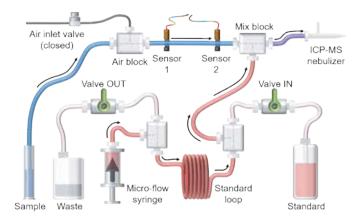
- When the autosampler probe moves to the sample vial, it triggers measurement of the sample uptake rate.
- An air bubble is introduced via the "Air inlet valve".
- Optical fiber sensors measure the elapsed time between the air bubble passing Sensor 1 and Sensor 2. The elapsed time is inversely proportional to the sample flow rate, allowing the actual solution flow rate to be calculated.



Step 4: MSA spikes are added automatically.

The syringe pump delivers the standard solution from the loop into the sample line via the "Mix Block". The standard solution flow rate needed to give each of the required MSA spike levels is calculated automatically, as explained later in the worked example.

The MSA spike solution is mixed with the sample online and the spiked sample is passed to the ICP-MS nebulizer.



Worked example: Addition of a 50 ppt MSA spike to a sample flowing at 200 µL/min.

- A 1 ppb spike standard is prepared and placed in a sample bottle connected to the Standard line of the ASAS.
- The microflow syringe loads the spike standard into the ASAS loop via the dedicated uptake line.
- The autosampler moves to the next sample vial.
- The ASAS measures the sample uptake line flow rate (as previously outlined in Step 3). In this example, we will use a nominal flow rate of 200 µL/min.
- Based on the measured sample flow rate, the ASAS software calculates the flow rate of the 1 ppb stock standard required to give a 50 ppt spike concentration. In this case, to achieve a spike level of 50 ppt in the sample flow of 200 μ L/min, the standard would need to be introduced at a flow rate of 10.0 μ L/min (20x dilution of the 1 ppb standard).
- The standard in the loop is added to the sample via the "Mix Block" at the calculated flow rate. The mixed, spiked sample then passes to the ICP-MS for analysis.

The ASAS can accurately add the standard solution at any flow rate between 0.10 and 99.99 μ L/min. However, to avoid over diluting the sample, the recommended standard flow rate is between 1.00 and 10.00 μ L/min. This assumes a typical nebulizer self-aspiration flow rate of 200 μ L/min.

The volume of the ASAS microflow syringe is about 800 μ L, and the loop volume is 700 μ L. When the volume of the standard solution remaining in the syringe falls below a set value, the syringe is automatically refilled. This happens after the current set of standard additions has completed.

If the total volume of standard solution required for the spike additions exceeds the loop volume, the syringe automatically refills the loop to ensure continuous operation.

Integrating the ASAS with the Agilent 8900 ICP-QQQ offers the following advantages for ultratrace elemental analysis of semiconductor samples:

- Compatibility with the Agilent I-AS autosampler and self-aspirating nebulizers
- Compact, easy-to-use, online system
- Automated creation of MSA or external calibrations
- Automated spike addition for spike recovery studies
- The ASAS can also be installed as part of the IAS Continuous Chemical Samples Inspection (CSI) system. This system provides online monitoring of multiple streams, baths, and containers of semiconductor process chemicals.

Results and Discussion

DLs and BECs

In total, 47 elements—including all 22 elements specified in SEMI standard C41-0705—were measured using the 8900 ICP-QQQ. The instrument was operated in multiple tune modes, which were switched automatically during a single visit to each sample vial. Data for each of the modes was combined automatically into a single report for each sample. Detection Limits (DLs) and Background Equivalent Concentrations (BECs) in undiluted IPA are given in Table 3. The DLs were calculated from 3 x the standard deviation (SD) of 10 replicate measurements of the blank (unspiked) IPA sample. The DLs and BECs for all SEMI required elements (shown in bold) were all well below the grade 4 requirements of 100 ppt; many were below 0.1 ppt. These results illustrate how the 8900 ICP-QQQ provides performance that ensures compliance with higher chemical purities that will be required for semiconductor manufacturing in the future. DLs and BECs for Hf and Re could not be calculated, as the measured background signal was zero counts per second in all replicates of the blank IPA. The BEC for Cu reported using the normal, preferred isotope of Cu-63 was unexpectedly high, at 6.4 ppt. This result was compared to the BEC measured using the secondary isotope, 65Cu, and the two measured concentrations were in agreement. This suggests the high BEC observed using 63Cu was due to trace Cu contamination in the IPA sample rather than any interference on ⁶³Cu.

The ASAS was used to perform an automatic spike recovery test. Ten separate IPA solutions were spiked at 20 ppt and measured against an external calibration that was created automatically by converting the MSA calibration. The spike recovery accuracy and repeatability (%RSD) results are also shown in Table 3. Excellent spike recoveries of between 91–108% were achieved for all elements at the 20 ppt level,

and RSDs (n=10) were between 1.6 and 8.9%. The results show the excellent reproducibility of the ASAS spike additions, as well as the good stability of the 8900 ICP-QQQ when aspirating organic solvents. This demonstrates the suitability of the ASAS-ICP-QQQ method for the routine analysis of ppt-level contaminant elements in IPA.

Table 3. DLs, BECs, and spike recoveries in IPA. Analytes shown in bold are SEMI grade 4 elements.

Analyte	Q1	Q2	Tune mode	DL (ng/L)	BEC (ng/L)	20 ng/L Recovery (%)	20 ng/L n=10 RSD (%)	SEMI stand- ard C41-0705 Grade 4 (ng/L)
Li	7	7	*H ₂	0.010	0.040	99	2.4	< 100
Ве	9	9	No gas	0.023	0.005	99	2.4	
В	11	11	No gas	1.2	12	96	8.0	< 100
Na	23	23	*NH ₃	0.060	0.97	109	5.7	< 100
Mg	24	24	*NH ₃	0.020	0.082	102	2.7	< 100
Al	27	27	*NH ₃	0.042	0.16	100	2.8	< 100
Р	31	47	O ₂ He	2.6	43	99	7.9	<16,000*
K	39	39	*NH ₃	0.64	1.1	107	4.9	< 100
Ca	40	40	*NH ₃	0.19	0.62	108	4.7	< 100
Ti	48	64	O ₂ He	0.23	1.3	99	2.4	< 100
٧	51	67	O ₂ He	0.020	0.030	99	2.3	< 100
Cr	52	52	*NH ₃	0.16	0.48	92	1.7	< 100
Mn	55	55	*NH ₃	0.030	0.030	102	2.4	< 100
Fe	56	56	*NH ₃	0.16	0.72	101	2.5	< 100
Со	59	59	He	0.020	0.020	99	2.1	
Ni	60	60	He	0.43	0.80	101	2.0	<100
Cu	63	63	O ₂ He	0.38	6.4	97	2.3	<100
Zn	64	64	He	0.71	0.72	98	6.9	<100
Ga	71	71	O ₂ He	0.013	0.005	100	2.8	
Ge	74	74	He	0.30	0.070	96	8.1	
As	75	91	O ₂ He	0.41	0.26	108	2.7	<100
Rb	85	85	H ₂	0.17	0.59	101	2.4	
Sr	88	88	O ₂ He	0.005	0.002	98	2.4	
Zr	90	90	O ₂ He	0.030	0.020	99	2.7	
Nb	93	93	H ₂	0.14	0.41	102	4.0	
Мо	98	130	O ₂ He	0.17	0.11	103	4.1	
Ru	101	101	He	0.080	0.03	99	2.9	
Rh	103	103	O ₂ He	0.070	0.18	99	2.1	
Pd	105	105	O ₂ He	0.070	0.040	100	2.5	
Ag	107	107	O ₂ He	0.014	0.006	97	2.6	
Cd	111	111	O ₂ He	0.035	0.004	98	4.0	<100
In	115	115	O ₂ He	0.012	0.008	99	1.8	
Sn	118	118	O ₂ He	0.058	0.034	100	4.9	<100
Sb	121	121	O ₂ He	0.056	0.009	103	2.3	<100
Те	125	125	O ₂ He	0.78	0.29	97	8.9	
Cs	133	133	*H ₂	0.060	0.022	96	4.2	
Ва	138	138	O ₂ He	0.009	0.004	99	2.4	<100

Analyte	Q1	Q2	Tune mode	DL (ng/L)	BEC (ng/L)	20 ng/L Recovery (%)	20 ng/L n=10 RSD (%)	SEMI stand- ard C41-0705 Grade 4 (ng/L)
Hf	178	178	He	0.000	0.000	105	5.1	
W	182	214	O ₂ He	0.21	0.049	97	5.7	
Re	185	185	O ₂ He	0.000	0.000	96	3.0	
Ir	193	193	No gas	0.060	0.006	101	6.7	
Pt	195	195	O ₂ He	0.51	0.45	100	3.0	
TI	205	205	O ₂ He	0.018	0.008	99	2.1	
Pb	208	208	O ₂ He	0.047	0.042	100	2.7	<100
Bi	209	209	O ₂ He	0.021	0.004	98	1.6	
Th	232	248	O ₂ He	0.11	0.022	97	4.6	
U	238	254	O ₂ He	0.18	0.048	91	6.6	

^{*} High-power cool plasma conditions: the temperature of the plasma was adjusted by changing the make-up gas flow rate.

Resolving polyatomic interferences on Mg, Al, and Cr

In this work, high-power cool plasma conditions were combined with reaction cell gases to provide the most effective control of intense background and matrix-based interferences. Cool plasma conditions were obtained by adjusting the make-up gas flow, while maintaining plasma energy with normal "hot plasma" RF power of 1500 W. These plasma conditions ensured sufficient plasma robustness to allow long-term analysis of the organic matrix, while providing effective control of carbon-based interferences on analytes such as Mg, Al, and Cr (Table 4).

Table 4. Main interferences arising from organic solvent matrix.

Analyte	Interferences	DL (ng/L)	BEC (ng/L)
²⁴ Mg	¹² C ₂ ⁺	0.020	0.082
²⁷ Al	¹² C ¹⁵ N+, ¹³ C ¹⁴ N+, ¹² C ¹⁴ N ¹ H+	0.042	0.16
³¹ P	¹⁵ N ¹⁶ O+, ¹⁴ N ¹⁷ O+, ¹³ C ¹⁸ O+	2.6	43
⁵² Cr	⁴⁰ Ar ¹² C ⁺	0.16	0.48

The major isotope of magnesium, 24 Mg $^{+}$, suffers an intense polyatomic interference from 12 C $_2^{+}$ in organic samples. Cool plasma conditions can suppress the ionization of C $_2$, and CRC mode can also be employed successfully to resolve the interference. In this work, the lowest DLs for Mg were achieved using a combination of high-power cool plasma conditions and on-mass measurement in MS/MS mode with NH $_3$ cell gas. The calibration curve for 24 Mg shows that the 12 C $_2^{+}$ interference was removed successfully, achieving a BEC less than 0.1 ng/L (ppt), and a detection limit of 0.020 ppt (Figure 2).

^{** &}lt; 16,000 ppt is the concentration limit for elemental P that is equivalent to the SEMI specified limit of 50 ppb (50,000 ppt) for PO_4 .

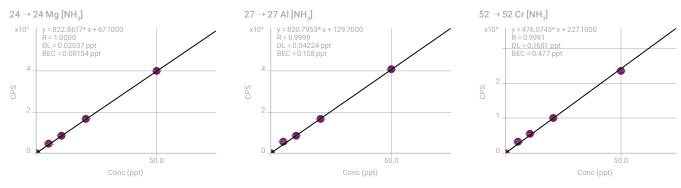


Figure 2. MSA calibration curves for ²⁴Mg, ²⁷Al, and ⁵²Cr.

The same approach was effective for the determination of other elements that suffer carbon-based polyatomic interference in organic solvents, such as 27 Al and 52 Cr. The calibrations shown in Figure 2 demonstrate that the interferences from 12 C 15 N $^+$, 13 C 14 N $^+$, 12 C 14 NH $^+$ on 27 Al $^+$ and 40 Ar 12 C $^+$ on 52 Cr $^+$ were minimized using high-power cool plasma conditions and NH $_3$ cell gas. These conditions gave BECs and DLs of 0.16 and 0.042 ppt for Al, and 0.48 and 0.16 ppt for Cr, respectively (Table 4).

P determination

SEMI Standard C41-0705 specifies the maximum concentration of phosphate allowed in high purity IPA, with a limit of 50 μ g/L (ppb) or 50,000 ppt. This limit equates to an elemental concentration of about 16,000 ppt for phosphorus (P).

The analysis of P in IPA needs an ORS cell mode that can resolve P from the normal plasma background polyatomic ions formed from N and O - ¹⁴N¹⁶O¹H⁺, ¹⁵N¹⁶O⁺, and ¹⁴N¹⁷O⁺. In addition, potential carbon-based interferences that overlap P⁺ at m/z 31 must also be resolved. MS/MS mass shift operation with oxygen reaction cell gas has been shown to be suitable for the analysis of P. Using this method, the P⁺ ions react with O₂ cell gas to form a reaction product ion PO⁺ at m/z 47, mass-shifted away from the original on-mass interferences.

A previous study (3) demonstrated that a relatively high octopole bias (-3 V) with a mix of cell gases comprising 0.075 mL/min oxygen plus 12 mL/min helium could be used successfully for low-level analysis of P. The relative cell gas flow rates mean that the density of helium atoms in the cell is 160 (12/0.075) times greater than that of oxygen. Most of the ions entering the cell will therefore collide multiple times with helium atoms before they collide (and react) with an oxygen atom. Helium works as a buffer gas, reducing the kinetic energy of the ions before they react with the O_2 cell gas. This low collision energy should reduce the in-cell formation of certain unwanted reaction product ions, for example, $^{13}C^{18}O^+ + O_2$ II $^{13}C^{18}O^{16}O^+ + O$. Suppressing these reactions reduces the formation of interfering product ions that could overlap the analyte product ion $^{31}P^{16}O^+$ at m/z 47. Using these mixed cell gas conditions, a minimum BEC for P of 27 ppt in IPA was reported (3).

The measurement conditions for P described in reference 3 were also used in this study. The calibration curve in Figure 3 shows good linearity from 5 to 50 ppt for P. The BEC was 43 ng/L (ppt) and the DL was 2.6 ppt (Table 3). Given the much higher typical contaminant levels for P compared to the other trace metals, it would be reasonable to calibrate P at a higher concentration level than the other elements. This modification could easily be applied to the ASAS methodology

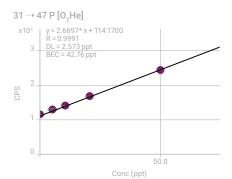


Figure 3. MSA calibration curve for ³¹P.

described here, if a higher P concentration was added to the working stock standard solution. The ASAS would then prepare and inject the online MSA spikes from the mixed standard, including the higher P spike levels. The upper limit for P defined in SEMI Grade 4 purity chemicals is also much higher than for the other trace elements. The relatively high BEC of 43 ppt is still several orders of magnitude lower than the 16,000 ppt concentration limit specified for P.

Conclusions

By automating the processes of sample preparation and standard spiking, the IAS ASAS automated-MSA system simplifies the elemental analysis of semiconductor process chemicals using the Agilent 8900 ICP-QQQ. The multi-element standard is prepared and connected to the ASAS, and the samples are loaded into the I-AS autosampler. The ASAS system then automatically performs all required steps, including online MSA spike additions and introduction of the sample to the ICP-QQQ.

Eliminating manual sample handling steps during ultratrace analysis lowers the risk of contamination. Limiting the handling of reagents and samples also reduces the likelihood of errors arising during the experimental procedure. Automating calibration and spike addition leads to increased consistency and higher confidence in the quality of the results.

The Agilent 8900 ICP-QQQ was operated using optimized plasma conditions and MS/MS mode to measure 47 elements in IPA. BECs at sub-ppt to ppt levels were acquired for all analytes – including all the elements specified in SEMI C41-0705. The results easily meet the current SEMI grade 4 specifications for all elements, including P, in IPA.

The excellent spike recovery and repeatability results for all elements at the 20 ppt level show the suitability of the automated ASAS method for the routine analysis of semiconductor process chemicals. The long-term robustness of this method is enhanced by using high-power cool plasma conditions. These conditions provide superior matrix decomposition and improved analyte ionization in the presence of the organic solvent matrix.

References

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Automated Analysis of Semiconductor Grade Hydrogen Peroxide and DI Water using ICP-QQQ

Authors

Kazuhiro Sakai Agilent Technologies, Japan Austin Schultz Elemental Scientific, USA Online MSA calibration using prepFAST S automated sample introduction and Agilent 8900 ICP-QQQ

Introduction

Maximizing product yield and performance of semiconductor devices requires manufacturers to address the potential for contamination at every stage of the production process. Contamination from particles, metals, and organic residues can affect the electrical properties of the semiconductor, reducing the quality and reliability of the final product. For example, following each photolithography step during wafer processing, the organic photoresist mask must be completely removed from the silicon wafer surface. A mixture of sulfuric acid (H_2SO_4) and hydrogen peroxide (H_2O_2) known as a sulfuric/peroxide mix (SPM) is used for this cleaning procedure. SPM is also used for degreasing the wafer surface. H_2O_2 is also used in the RCA Standard Clean steps (SC-1 and SC-2) used to clean silicon wafers, and for etching metallic copper on printed circuit boards.

Ultrapure water (UPW) is used throughout the wafer fabrication process. As well as working as a rinse solution between processing steps, UPW is also the diluent for many process chemistries such as SC-1 and SC-2 solutions. As these chemicals are in frequent and prolonged contact with the wafer surface, minimizing metal impurities is essential to prevent wafer surface contamination.

Semiconductor Equipment and Materials International (SEMI) publishes standards for semiconductor process chemicals. The standard for H_2O_2 is SEMI C30-1110 – Specifications for Hydrogen Peroxide (1). SEMI Grade 5 is the highest purity level, with maximum contamination levels of 10 ppt for most trace elements.

Quadrupole ICP-MS (ICP-QMS) is the standard technique used to monitor trace element contaminants in the semiconductor industry. However, the drive for ever smaller device architectures and higher yields requires an increasing number of contaminant elements to be monitored at lower concentrations.

In addition to trace elements, SEMI Standard C30-1110 specifies the maximum concentration of sulfate and phosphate allowed in high purity $\rm H_2O_{2'}$ with a limit of 30,000 ppt. This limit equates to an elemental concentration of sulfur (S) and phosphorus (P) of 10,000 ppt. Due to the relatively high detection limits achievable with conventional single quadruple ICP-MS, these two elements are not currently measured using ICP-MS.

Triple quadrupole ICP-MS (ICP-QQQ) provides much lower limits of detection for S and P (among many other elements). Uniquely, the technique offers the potential for the sulfate and phosphate analysis to be combined with the other trace metals. The adoption of ICP-QQQ therefore enables all SEMI specified elements to be monitored using a single technique (2, 3).

Contamination control

Ultratrace analysis at the pg/g (ppt) or fg/g (ppq) level is susceptible to contamination from the lab environment, reagents, or errors arising from manual tasks, such as pipetting. To deliver consistently accurate results at these ultratrace concentrations, a skilled and experienced analyst is typically required.

One approach to simplifying the analysis for less expert analysts is to use an automated sample introduction system. These systems automate typical sample handling steps such as dilution, acidification, and spiking. They can also automatically generate a calibration curve using either external standards or Method of Standard Additions (MSA).

In this study, an automated procedure was developed to quantify ultratrace elemental impurities in de-ionized (DI) water and $\rm H_2O_2$ using an Agilent 8900 ICP-QQQ fitted with an ESI prepFAST S automated sample introduction system. The prepFAST S automates sample preparation and calibration, saving time and minimizing the risk of sample-contamination from manual sample handling operations.

Experimental

Reagents and samples

TAMAPURE-AA-10 hydrogen peroxide (35%, Tama Chemicals, Japan) and ultrapure DI water (Milli-Q water, Molsheim, France) were used as the samples.

Standard stock solution for MSA: a 1000 ppt mixed multi-element standard solution was prepared by diluting a 10 ppm mixed multi-element standard solution (SPEX CertiPrep, NJ, US) with 1% HNO $_3$.

Nitric acid for sample acidification: a 10% nitric acid solution was prepared by diluting 68% ultrapure $\rm HNO_3$ (TAMAPURE-AA-10) with DI water. $\rm HNO_3$ was automatically added the $\rm H_2O_2$ samples, giving a final concentration of 0.5% $\rm HNO_3$ to stabilize the spiked elements. UPW samples are often also acidified to ensure trace element stability (see reference 2). However, in this work, the DI water was analyzed unacidified, without the addition of a $\rm HNO_3$ spike, providing results that can be compared with the earlier work.

The standard stock and ${\rm HNO_3}$ spike solutions were loaded on the prepFAST S. All solutions run in the analysis were automatically prepared from these stock solutions by the prepFAST S system. The prepFAST S method used DI (Milli-Q) water as the carrier solution, at a flow rate of 100 μ L/min.

All preparation and analysis steps were performed in a Class 10,000 clean room.

Instrumentation

A standard Agilent 8900 semiconductor configuration ICP-QQQ instrument was equipped with a PFA concentric nebulizer that is included with the prepFAST S automated sample introduction system. The semiconductor configuration ICP-QQQ is fitted with a Peltier cooled quartz spray chamber, quartz torch (2.5 mm id), platinum-tipped sampling and skimmer cones, and s-lens.

The 8900 ICP-QQQ was connected to the ESI prepFAST S automated sample introduction system. The prepFAST S is a specialized, semiconductor version of the standard ESI prepFAST. The S version has a high purity, low-contamination, inert sample path and features an automated MSA spike addition mode. ICP-QQQ

instrument operating conditions are given in Table 1. Tuning: To achieve the lowest DLs, a multi-tune method was used. The tuning parameters are summarized in Table 1. For data acquisition, a 2 s integration time was used for all isotopes with three replicates (10 replicates for the blank to calculate the DLs).

Table 1. ICP-QQQ operating conditions

	Cool-no gas	Cool-NH ₃ (1)	Cool-NH ₃ (2)	No gas	H ₂	Не	O ₂ (1)	0, (2)
Acquisition mode	Single Q	MS/	MS/MS					
RF power (W)		600				1500		
Carrier gas (L/min)				0.70			,	
Make-up gas (L/min.)	0.90					0.48		
Sampling depth (mm)	18.0					8.0		
Ex1 (V)	-150		-100.0	4.2	4.7	4.2	4.5	3.5
Ex2 (V)	-18.0 -17.0 -1				-25	0.0		-120.0
Omega bias (V)		-70.0			-140.0			
Omega lens (V)		2.0		10.0	8.0	10.0	10.5	4.0
Q1 entrance (V)	-15	5.0		-50.0				
NH ₃ flow (mL/min)*	-	3.0 (3	0%)**	-	-	-	-	-
He flow (mL/min)	-	1.	.0	-	-	5.0	-	-
H ₂ flow (mL/min)	-	-	-	-	7.0	-	-	-
O ₂ flow (mL/min)			-	-	-	-	4.5 (4	5%)**
Axial acceleration (V)	0.0 1.5			0.0 1				.0
Energy discrimination (V)	15.0	-5	.0	5.0	0.0	3.0	-7	.0

^{*10%} $\mathrm{NH_3}$ balanced with 90% He

The most advanced semiconductor manufacturing facilities require the lowest possible levels of contamination, so they require analytical techniques that can deliver the lowest possible detection limits (DLs). This requirement is critical in the analysis of trace contaminants in process chemicals such as UPW and $\rm H_2O_2$, which are used at multiple stages of the wafer fabrication process. UPW and $\rm H_2O_2$ also come into direct contact with the wafer surface.

The 8900 ICP-QQQ satisfies this requirement by offering the flexibility to optimize the measurement parameters (plasma conditions, quadrupole scan mode, cell gas type, and flow rate) to give the highest sensitivity and lowest background for each analyte.

In this work, several reaction cell gases (He, H_2 , O_2 , and NH_3) were used in the collision/reaction cell (CRC) of the 8900, as appropriate for the large number of analytes being measured. Since DI water and H_2O_2 are low-matrix samples, cool plasma conditions were also applied for the elements where this mode provides the lowest background equivalent concentrations (BECs).

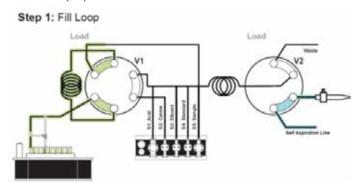
The tuning steps were applied sequentially during the measurement of each solution. This approach allows the tuning conditions to be optimized for the removal of different types of interferences, while maintaining maximum sensitivity for the analytes. Q1 and Q2 settings are shown in Table 2 along with DLs, BECs, and quantification results.

^{*} Values in parentheses are % of the maximum flow of the gas controller, as displayed in the tuning pane of ICP-MS MassHunter software

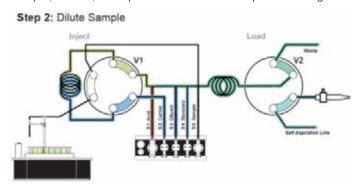
ESI prepFAST S operation

The prepFAST S automated sample introduction system combines an autosampler with a system of ultrapure valves (S1 -5), and a set of high precision syringe pumps. Undiluted chemicals can be placed on the autosampler and the system will perform the actions—such as dilution, acidification, and spiking—required to prepare the sample for introduction to the ICP-MS or ICP-QQQ. The operation of the prepFAST S is outlined in the four schematics shown in Figure 1.

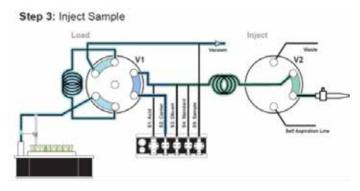
1. Loading of sample: Syringe S5 loads a precise amount of sample to the loop of valve 1 (V1).



2. Sample dilution and sample spiking: Syringes S1, S2, S3, and S4 mix the acid, sample, diluent, and spike solution into a loop connecting V1 and V2.



3. Sample injection: The prepared sample is introduced into the ICP-QQQ via the carrier solution pumped by S2. S2 provides a precise flow rate regardless of sample type. The V1 loop is washed simultaneously.



4. Valve wash: UPW or acidified UPW is used to clean the lines between V1 and V2.

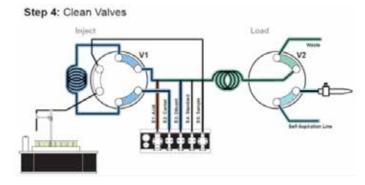


Figure 1. ESI prep*FAST* S system schematic, illustrating four distinct steps: sample loading during spray chamber rinse, sample preparation, injection, and cleaning.

The prepFAST S removes the need for analyst intervention in the analysis of semiconductor grade chemicals, reducing the risk of sample contamination. The integrated system offers the following advantages for the ultratrace elemental analysis of semiconductor samples:

- 1. Automated dilution of samples
- 2. Automated creation of external or MSA calibrations
- 3. Automated acidification of samples
- 4. Injection of samples at a precise flow rate
- 5. High speed rinsing of the ICP-MS sample introduction system

Results and Discussion

Figures 2 and 3 show calibration curves for Na, K, Si, P, and S in DI water and Ca, Zn, and As in $\rm H_2O_{2^1}$ respectively. All elements were measured using the MSA calibration prepared automatically using the prepFAST S. These elements are difficult to analyze at low levels due to raised backgrounds. The analytes Si, P, and S are not commonly measured with conventional single quadrupole ICP-MS, due to the presence of intense polyatomic interferences. However, the controlled reaction chemistry of the 8900 ICP-QQQ operating in MS/MS mode gives far superior control of background interferences. MS/MS mode allows these elements to be calibrated and quantified at ppt concentrations.

Good linearity at the ppt level was observed for all elements measured in both sample matrices, although Si, P, and S had relatively high BECs of 85, 10, and 118 ppt, respectively. These elements are typically present at higher levels than the trace metals, as they are more difficult to control in the lab environment and in reagents. They are also less critical contaminants, as reflected in the higher levels for P and S (of 30 ppb for phosphate and sulfate) permitted in high purity H_2O_2 . However, despite the higher BECs, the calibration curves for Si, P, and S were still linear over the calibration range from 10 to 50 ppt. The same calibration levels were used for all analytes, as the mixed stock standard contained all elements at the same concentration.

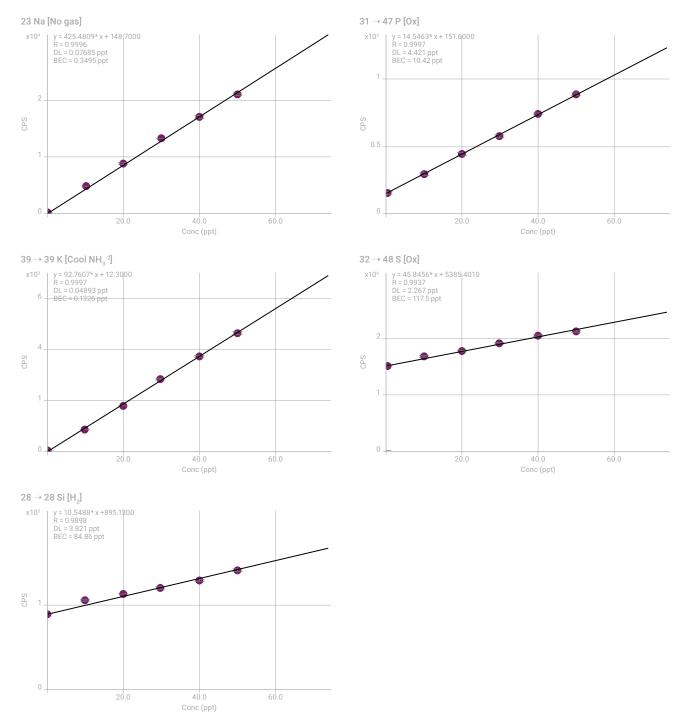


Figure 2. Calibration plots for Na, K, Si, P, and S in DI water. All values in ng/L (ppt).

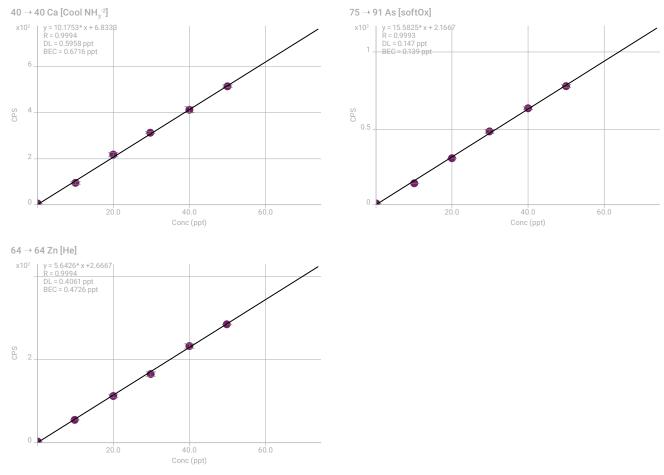


Figure 3. Calibration plots for Ca, Zn, and As in H₂O₂. All values in ng/L (ppt).

DLs and quantitative results

Forty-nine elements in total, including all the elements listed in SEMI C30-1110, were measured by MSA in DI water and $\rm H_2O_2$, using the 8900 multi-tune mode method. Data was acquired in an automated sequence of cool plasma, no gas, and gas modes, during a single visit to the sample vial. Data for each of the modes was combined automatically into a single report for each sample.

Quantitative results, DLs, and BECs for all analytes—including all the SEMI specified elements—are shown in Table 2. Detection limits were calculated as 3-sigma of 10 replicate measurements of the blank DI water or H₂O₂ sample.

DLs <1 ng/L (ppt) were obtained for 46 elements in DI water. The DLs for the remaining 3 elements, Si, P, and S, were at the single-ppt level. Measured concentrations of all elements apart from B, Si, P, and S were <1 ng/L or <DL, confirming the purity of the sample. This analytical performance easily meets the requirements for monitoing UPW in semiconductor manufacturing.

In $\rm H_2O_2$, DLs <1 ng/L were obtained for 45 elements. The DLs for B, P, and S, were at the single-ppt level, while the DL for Si was 26 ppt. All elements were measured at <1 ng/L or <DL apart from B (22 ppt), Na (1.1 ppt), Si (500 ppt), P (9.4 ppt), and S (220 ppt) in 35% $\rm H_2O_2$. Only B and Si exceed the 10 ppt maximum limit in the SEMI specifications, and of these, only B is a SEMI specified element. P and S were quantified well below the 10,000 ppt SEMI specified limit in $\rm H_2O_2$.

Table 2. Quantification of trace elements in DI water and 35% H2O2. SEMI specification elements are in bold.

	Q1	Q2	Scan type	Tune mode		DI Water			H ₂ O ₂	
					DL (ng/L)	BEC (ng/L)	Conc (ng/L)	DL (ng/L)	BEC (ng/L)	Conc (ng/L)
Li		7	SQ	Cool no gas	0.003	0.001	<dl< td=""><td>0.025</td><td>0.022</td><td><dl< td=""></dl<></td></dl<>	0.025	0.022	<dl< td=""></dl<>
Ве	9	9	MS/MS	No gas	0.096	0.040	<dl< td=""><td>0.089</td><td>0.017</td><td><dl< td=""></dl<></td></dl<>	0.089	0.017	<dl< td=""></dl<>
В	11	11	MS/MS	No gas	0.52	1.7	1.7	1.9	22	22
Na	-	23	SQ	Cool no gas	0.077	0.35	0.35	0.11	1.1	1.1
Mg		24	SQ	Cool no gas	0.015	0.009	<dl< td=""><td>0.040</td><td>0.053</td><td>0.053</td></dl<>	0.040	0.053	0.053
Al		27	SQ	Cool no gas	0.040	0.028	<dl< td=""><td>0.22</td><td>0.63</td><td>0.63</td></dl<>	0.22	0.63	0.63
Si	28	28	MS/MS	H ₂	3.8	85	85	26	500	500
P	31	47	MS/MS	0 ₂	4.4	10	10	2.6	9.4	9.4
s	32	48	MS/MS	0 ₂	2.3	120	120	7.5	220	220
		39			0.049					
K	39		MS/MS	Cool NH ₃ (2)		0.13	0.13	0.19	0.45	0.45
Ca	40	40	MS/MS	Cool NH ₃ (2)	0.082	0.044	<dl< td=""><td>0.60</td><td>0.67</td><td>0.67</td></dl<>	0.60	0.67	0.67
Ti	48	64	MS/MS	0 ₂ (2)	0.042	0.021	<dl< td=""><td>0.24</td><td>0.21</td><td><dl< td=""></dl<></td></dl<>	0.24	0.21	<dl< td=""></dl<>
v	51	67	MS/MS	0, (2)	0.021	0.026	0.026	0.058	0.068	0.068
Cr	52	52	MS/MS	Cool NH ₃ (1)	0.085	0.047	<dl< td=""><td>0.24</td><td>0.69</td><td>0.69</td></dl<>	0.24	0.69	0.69
Mn	55	55	MS/MS	Cool NH ₃ (1)	0.010	0.010	0.010	0.039	0.020	<dl< td=""></dl<>
Fe	56	56	MS/MS	Cool NH ₃ (1)	0.070	0.076	0.076	0.29	0.17	<dl< td=""></dl<>
Со	59	59	MS/MS	Cool NH ₃ (1)	0.017	0.002	<dl< td=""><td>0.025</td><td>0.005</td><td><dl< td=""></dl<></td></dl<>	0.025	0.005	<dl< td=""></dl<>
Ni	60	60	MS/MS	Cool NH ₃ (1)	0.080	0.016	<dl< td=""><td>0.24</td><td>0.18</td><td><dl< td=""></dl<></td></dl<>	0.24	0.18	<dl< td=""></dl<>
Cu	63	63	MS/MS	Cool NH ₃ (1)	0.12	0.11	<dl< td=""><td>0.17</td><td>0.12</td><td><dl< td=""></dl<></td></dl<>	0.17	0.12	<dl< td=""></dl<>
Zn	64	64	MS/MS	He	0.063	0.28	0.28	0.41	0.47	0.47
Ga	<u> </u>	71	SQ	Cool no gas	0.011	0.001	<dl< td=""><td>0.032</td><td>0.031</td><td><dl< td=""></dl<></td></dl<>	0.032	0.031	<dl< td=""></dl<>
Ge	74	74	MS/MS	He	0.36	0.32	<dl< td=""><td>0.27</td><td>0.20</td><td><dl< td=""></dl<></td></dl<>	0.27	0.20	<dl< td=""></dl<>
As	75	91	MS/MS	0, (2)	0.072	0.035	<dl< td=""><td>0.15</td><td>0.14</td><td><dl< td=""></dl<></td></dl<>	0.15	0.14	<dl< td=""></dl<>
Se	78	78	MS/MS	H ₂	0.20	0.14	<dl< td=""><td>0.40</td><td>0.13</td><td><dl< td=""></dl<></td></dl<>	0.40	0.13	<dl< td=""></dl<>
Rb		85	SQ	Cool no gas	0.031	0.015	<dl< td=""><td>0.052</td><td>0.035</td><td><dl< td=""></dl<></td></dl<>	0.052	0.035	<dl< td=""></dl<>
Sr	88	88	MS/MS	He	0.024	0.002	<dl< td=""><td>0.000*</td><td>0.000*</td><td>0.000*</td></dl<>	0.000*	0.000*	0.000*
Nb	93	93	MS/MS	He	0.018	0.010	<dl< td=""><td>0.030</td><td>0.029</td><td><dl< td=""></dl<></td></dl<>	0.030	0.029	<dl< td=""></dl<>
Мо	98	98	MS/MS	He	0.093	0.045	<dl< td=""><td>0.065</td><td>0.063</td><td><dl< td=""></dl<></td></dl<>	0.065	0.063	<dl< td=""></dl<>
Ru	101	101	MS/MS	He	0.077	0.058	<dl< td=""><td>0.075</td><td>0.014</td><td><dl< td=""></dl<></td></dl<>	0.075	0.014	<dl< td=""></dl<>
Rh	103	103	MS/MS	0, (2)	0.057	0.10	0.10	0.018	0.097	0.097
Pd	105	105	MS/MS	No gas	0.078	0.12	0.12	0.055	0.090	0.090
Ag	107	107	MS/MS	No gas	0.099	0.14	0.14	0.031	0.016	<dl< td=""></dl<>
Cd	114	114	MS/MS	No gas	0.045	0.021	<dl< td=""><td>0.047</td><td>0.009</td><td><dl< td=""></dl<></td></dl<>	0.047	0.009	<dl< td=""></dl<>
In	115	115	MS/MS	No gas	0.009	0.003	<dl< td=""><td>0.022</td><td>0.019</td><td><dl< td=""></dl<></td></dl<>	0.022	0.019	<dl< td=""></dl<>
Sn	118	118	MS/MS	No gas	0.038	0.059	0.059	0.20	0.17	<dl< td=""></dl<>
Sb	121	121	MS/MS	H_2	0.029	0.032	0.032	0.028	0.005	<dl< td=""></dl<>
Те	125	125	MS/MS	No gas	0.18	0.043	<dl< td=""><td>0.000*</td><td>0.000*</td><td>0.000*</td></dl<>	0.000*	0.000*	0.000*
Cs		133	MS/MS	Cool no gas	0.074	0.020	<dl< td=""><td>0.088</td><td>0.059</td><td><dl< td=""></dl<></td></dl<>	0.088	0.059	<dl< td=""></dl<>
Ba	138	138	MS/MS	H ₂	0.023	0.014	<dl< td=""><td>0.039</td><td>0.018</td><td><dl< td=""></dl<></td></dl<>	0.039	0.018	<dl< td=""></dl<>
Та	181	181	MS/MS	No gas	0.024	0.041	0.041	0.12	0.28	0.28
W	182	182	MS/MS	No gas	0.037	0.009	<dl< td=""><td>0.044</td><td>0.044</td><td>0.044</td></dl<>	0.044	0.044	0.044
Re	185	185	MS/MS	No gas	0.040	0.037	<dl< td=""><td>0.062</td><td>0.056</td><td><dl< td=""></dl<></td></dl<>	0.062	0.056	<dl< td=""></dl<>
lr	193	193	MS/MS	H ₂	0.023	0.016	<dl< td=""><td>0.040</td><td>0.027</td><td><dl< td=""></dl<></td></dl<>	0.040	0.027	<dl< td=""></dl<>
Pt	195	195	MS/MS	No gas	0.28	0.33	0.33	0.088	0.39	0.39
Au	197	197	MS/MS	No gas	0.051	0.048	<dl< td=""><td>0.22</td><td>0.15</td><td><dl< td=""></dl<></td></dl<>	0.22	0.15	<dl< td=""></dl<>
TI	205	205	MS/MS	No gas	0.036	0.082	0.082	0.015	0.010	<dl< td=""></dl<>
Pb	208	208	SQ	No gas	0.042	0.066	0.066	0.056	0.035	<dl< td=""></dl<>
Bi	209	209	MS/MS	No gas	0.034	0.048	0.048	0.027	0.054	0.054
U	238	238	MS/MS	No gas	0.004	0.001	<dl< td=""><td>0.012</td><td>0.008</td><td><dl< td=""></dl<></td></dl<>	0.012	0.008	<dl< td=""></dl<>

Conclusions

By combining superior detection limits with a high degree of automation, the Agilent 8900 ICP-QQQ fitted with ESI's prepFAST S automated sample introduction system provides unmatched performance. The method also simplifies the elemental analysis of semiconductor process chemicals.

User handling of the samples is limited to loading the multielement stock standards, acid used for spiking, and samples into the prepFAST S automated sample introduction system. All subsequent steps, including introduction of the sample to the ICP-QQQ, are performed automatically by the prepFAST S. Benefits of the method include:

- Autodilution of samples
- Auto-acidification of samples
- Auto-creation of MSA calibrations
- Injection of samples at a precise flow rate
- High-speed rinsing of the ICP-MS sample introduction system.

A complete analysis of the two samples, meausred using separate, automated MSA calibrations, was achieved in less than 30 minutes.

Automating the sample handling steps speeds up the analytical procedure, while also making the overall analysis easier for the analyst to perform. Eliminating manual tasks such as sample dilution and spiking lowers the risk of contamination during ultratrace analysis. Limiting the handling of reagents and samples also reduces the likelihood of errors arising during the experimental procedure, leading to an increased confidence in the data quality.

All the elements specified in SEMI C30-1110, including P and S, were measured at sub-ppt to ppt levels in DI water and high purity 35% $\rm H_2O_2$. The results easily meet the current SEMI Grade 5 specifications for $\rm H_2O_2$.

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Direct Analysis of Trace Metal Impurities in High Purity Nitric Acid Using ICP-QQQ

Authors

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Introduction

The manufacturing yield of semiconductor devices has always been susceptible to contamination from trace metals. As the industry continues to progress towards devices with smaller features and a higher density of integration, susceptibility to contamination in microfabrication processing presents an increasing challenge. Controlling contamination at these small scales requires ever-higher purity of process chemicals and manufacturing conditions.

The semiconductor device fabrication industry uses well-established cleaning procedures to remove organic and metallic residues and impurities from the surface of silicon wafers. The purity of reagents used during manufacturing processes and the air quality in the fabrication plant are important considerations.

Nitric acid (HNO_3) plays an important role in the fabrication of semiconductor devices so needs to be of ultrahigh purity. A mix of nitric and hydrofluoric acid is used to etch single crystal silicon and polycrystalline silicon. HNO_3 is also combined with phosphoric acid and acetic acid for wet etching of aluminum. As a reagent, HNO_3 is used in the preparation of other semiconductor materials.

SEMI standard C35-0708 Tier-B protocol for HNO_3 (69.0–70.0%) specifies contaminant levels of <1 μ g/L (ppb) for several elements [1]. The concentration of industrial grade HNO_3 is usually 60–68%, depending on the method of production.

In this study, undiluted ${\rm HNO_3}$ was analyzed directly by triple quadrupole ICP-MS (ICP-QQQ). This approach simplified sample preparation and avoided the potential introduction of contaminants during dilution.

Experimental

Samples and standards

Two samples of HNO₃ were used in this study:

- Sample 1: 68 % HNO₃ (high purity-grade)
- Sample 2: 61 % HNO₂ (electronic-grade lower purity)

No further sample preparation was necessary as all samples were introduced directly into the ICP-QQQ.

Calibration and quantification were done using the method of standard additions (MSA). Standard solutions were prepared by spiking a multi-element standard solution (SPEX CertiPrep, NJ, US) into each HNO₃ sample to give spike levels of 5, 10, 20, 30 and 40 ppt. The density of the nitric acid solution varies with the concentration of the acid, which affects the sample transport, nebulization and droplet evaporation processes in the ICP-MS sample introduction. Therefore, for the most accurate analysis, the acid grade (concentration) used for the spiked MSA calibration solutions should be approximately matched to the acid concentration of the samples. ICP-MS MassHunter allows an MSA calibration to be converted to an external calibration to determine contaminant levels in other nitric acid samples with similar acid concentration. The solutions were prepared just before analysis. All preparation and analyses were performed in a Class 10,000 clean room.

Instrumentation

An Agilent 8900 Semiconductor configuration ICP-QQQ instrument was used in this study. The instrument is fitted as standard with a PFA-100 nebulizer, Peltier-cooled quartz spray chamber, quartz torch, platinum-tipped sampling and skimmer cones, and s-lens. The nebulizer was operated in self-aspiration mode to minimize the potential for sample contamination from contact with the peristaltic pump tubing. If large numbers of undiluted HNO $_3$ samples are run routinely, it is recommended that the large (18 mm) insert Pt cone is fitted. Long-term corrosion of internal ICP-MS components can be minimized by fitting the dry pump option and ball-type interface valve kit.

In advanced semiconductor applications, the key requirement is to deliver the absolute lowest possible detection limits (DLs) for each analyte. To achieve this goal, laboratories measuring ultratrace levels of contaminants can use a multitune method, where several tuning steps are applied sequentially during the measurement of each solution. This approach allows the tuning conditions to be optimized for the removal of different types of interferences, while maintaining sensitivity for each analyte. In this work, several reaction cell gases (He, H_{2} , O_{2} , and NH_{3}) and both hot and cool plasma conditions were used as appropriate for the large number of analytes being measured. Tuning conditions are shown in Table 1 and other acquisition parameters are shown in Table 2.

Table 1. ICP-QQQ operating conditions.

	Cool-NH ₂	No gas	Н,	He	0,	0, -soft		
	3		2		2	2		
Acquisition mode			MS/	MS				
RF power (W)	600		1500					
Sampling depth (mm)	18.0			8.0				
Nebulizer gas (L/min)		0.70						
Make-up gas (L/min)	0.78	0.36						
Ex1 (V)	-150	4.2	4.7	4.2	4.5	3.5		
Ex2 (V)	-17.0	-250.0 -120.0						
Omega bias (V)	-70.0		-14	0.0		-70.0		
Omega lens (V)	2.0	10.0	8.0	10.0	10.5	4.0		
Q1 entrance (V)	-15.0			-50.0				
He flow (mL/min)	1.0	-	-	5.0	-	-		
H ₂ flow (mL/min)	-	-	7.0	-	-	-		
*NH ₃ flow (mL/min)	2.0 (20%)**	-	-	-	-	-		
O ₂ flow (mL/min)	-	4.5 (30%)**						
Axial acceleration (V)	1.5	0.0 1.0						
Energy discrimination (V)	-5.0	5.0	0.0	3.0	-7	.0		

^{*10%} NH $_{\rm 3}$ balanced with 90% He

^{**} Values in parentheses are % of the maximum flow of the gas controller, as displayed in the tuning pane of ICP-MS MassHunter

Table 2. Acquisition parameters.

Parameter	Setting
Q2 peak pattern	1 point
Replicates	3 (spiked samples) 10 (unspiked solution for DL measurement)
Sweeps/replicate	10
Integration time	2 s for all isotopes

Results and Discussion

DLs and BECs

In total, 49 elements were measured using the 8900 ICP-QQQ operating in multiple tune modes, switched automatically during a single visit to each sample vial. Data for each of the modes was combined automatically into a single report for each sample. DLs and Background Equivalent Concentrations (BECs) in undiluted 68% $\rm HNO_3$ (Sample 1) are given in Table 3. The stability test results are discussed in the "long-term stability" section of the report.

Table 3. DLs and BECs in high purity 68% HNO₃.

Element	Tune	Q1	Q2	DL (ng/L)	BEC (ng/L)	30 ppt Recovery %	Stability test RSD %
Ве	No gas	9	9	0.12	0.071	92	3.5
В	No gas	11	11	0.43	3.5	94	6.3
Na	Cool-NH ₃	23	23	0.53	2.3	93	3.1
Mg	Cool-NH ₃	24	24	0.085	0.049	93	2.0
Al	Cool-NH ₃	27	27	0.10	0.16	93	3.6
Р	0,	31	47	8.1	83	95	_**
S	0,	32	48	2.6	65	93	_**
K	Cool-NH ₃	39	39	0.38	0.73	93	2.9
Ca	Cool-NH ₃	40	40	0.54	0.38	93	1.2
Sc	0,	45	61	0.007	0.013	93	0.5
Ti	O ₂ -soft	48	64	0.039	0.081	93	3.3
V	O ₂ -soft	51	67	0.041	0.17	93	1.5
Cr	Cool-NH ₃	52	52	0.42	0.25	93	3.0
Mn	Cool-NH ₃	55	55	0.084	0.014	93	2.5
Fe	Cool-NH ₃	56	56	0.75	1.1	92	4.7
Со	Cool-NH ₃	59	59	0.21	0.075	93	4.3
Ni	O ₂ -soft	60	60	0.067	0.38	93	2.0
Cu	Cool-NH ₃	63	63	0.12	0.50	94	3.8
Zn	He	64	64	0.52	0.46	93	2.9
Ga	Cool-NH ₃	71	71	0 cps	0 cps	92	2.5
Ge	H ₂	74	74	0.060	0.10	93	1.4
As	O ₂ -soft	75	91	0.082	0.081	93	1.8
Se	H ₂	78	78	0.78	0.41	93	5.5
Rb	Cool-NH ₃	85	85	0.089	0.030	93	3.0
Sr	He	88	88	0.014	0.012	93	0.8

Element	Tune	Q1	Q2	DL (ng/L)	BEC (ng/L)	30 ppt Recovery %	Stability test RSD %
Zr	0,	90	106	0.22	1.0	93	0.4
Nb	He	93	93	0.012	0.014	93	0.8
Мо	He	98	98	0.088	0.10	93	1.0
Ru	He	101	101	0.032	0.034	93	1.2
Pd	No gas	105	105	0.066	0.14	92	1.0
Ag	No gas	107	107	0.029	0.025	93	0.9
Cd	No gas	114	114	0.058	0.046	92	1.4
In	No gas	115	115	0.004	0.004	93	0.6
Sn	No gas	118	118	0.099	0.35	93	0.9
Sb	H ₂	121	121	0.056	0.028	93	1.6
Те	H ₂	125	125	0.57	0.45	93	5.2
Cs	Cool-NH ₃	133	133	0 cps	0 cps	93	2.4
Ba	H ₂	138	138	0.014	0.010	93	0.4
Hf	No gas	178	178	0.014	0.005	93	0.9
Та	He	181	181	0.052	0.065	93	0.5
W	No gas	182	182	0.030	0.022	93	0.7
Ir	No gas	193	193	0.016	0.011	93	0.9
Au	No gas	197	197	0.049	0.068	93	1.7
TI	No gas	205	205	0.090	0.46*	93	0.6
Pb	No gas	208	208	0.060	0.21	93	0.7
Bi	No gas	209	209	0.018	0.025	93	0.4
Th	No gas	232	232	0.004	0.003	93	0.8
U	No gas	238	238	0.025	0.013	93	0.6

DLs were calculated as 3-sigma of 10 replicate measurements of a blank HNO3 sample (cps refers counts per second).

Table 4 shows quantitative data for all SEMI specification elements [1] in high purity 68% HNO $_3$ and electronic-grade 61% HNO $_3$ determined by MSA. For the greatest accuracy, the two different concentration grades of nitric acid measured in this study were calibrated using separate MSA calibrations. However, if additional samples of similar grade (acid concentration) are measured, the MSA calibration can be easily and automatically converted to an external calibration plot. External calibration allows subsequent samples to be measured without requiring MSA spike additions into each additional sample.

Good linearity was obtained for all SEMI target elements, as shown in the representative calibration curves for B, Na, Al, K, Ca, As, and Pb (Figure 1).

Normally, the concentration in each sample is obtained by multiplying the quantitative value by the dilution factor (usually about 10 times for nitric acid). However, in this study, the quantitative value equals the sample concentration in the original sample, as the acids were measured undiluted. The results given in Table 4 show that all 49 elements studied can be analyzed at significantly lower levels than the <1 ppb maximum limit specified for HNO_3 in SEMI standard C35-0708 Tier-B [1].

^{*}The BEC of TI was higher than expected, most likely due to residual signal from the ICP-MS tuning solution.

^{**}P and S concentration in the mixed spike (30 ppt) was too low for reliable quantification above the blank (83 ppt and 65 ppt, respectively).

Table 3. Quantitative results for SEMI specification elements [1] in high purity 68% HNO $_3$ and electronic-grade 61% HNO $_3$.

Element	High-purity grade 68% HNO ₃ , ng/L	Electronic grade 61% HNO ₃ , ng /L	SEMI C35-0708 Tier-B max limit, ng/L
Li	<0.061	0.19	<1000
В	3.5	270	<1000
Na	2.3	130	<1000
Mg	<0.085	11	<1000
Al	0.16	93	<1000
К	0.73	6.5	<1000
Ca	<0.54	50	<1000
Ti	0.081	1.1	<1000
V	0.17	0.24	<1000
Cr	<0.42	70	<1000
Mn	<0.084	3.4	<1000
Fe	1.1	270	<1000
Ni	0.38	28	<1000
Cu	0.50	0.99	<1000
Zn	<0.52	3.8	<1000
As	<0.082	0.25	<1000
Cd	<0.058	0.80	<1000
Sn	0.35	13	<1000
Sb	<0.056	0.11	<1000
Ba	<0.014	0.43	<1000
Pb	0.21	0.31	<1000

Measured values shown as "<" indicate that the measured concentration was below the detection limit.

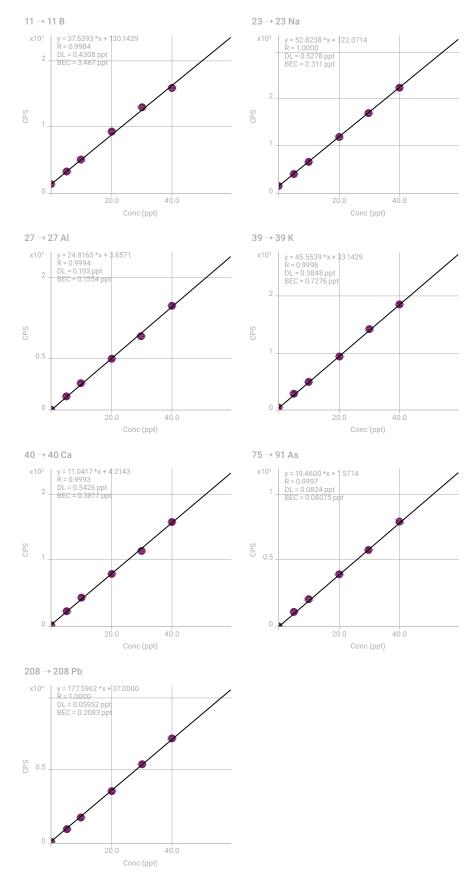


Figure 1. Calibration curves for several SEMI specification elements in high purity 68% HNO₃.

Long-term stability

Long-term stability was evaluated by measuring a 68% HNO $_3$ sample spiked at 30 ppt for all elements. Calibration curves were generated at the beginning of the sequence. The spiked samples were then run as unknown samples for a total analysis period of 6.5 hours. The RSDs of the 21 analysis results are shown in Table 3 (stability test RSD (%)). Good stability was maintained throughout the run, with RSDs between 0.4 and 5.5%. S and P gave less reliable long-term results due to the low concentration of the spike (30 ppt) measured above the relatively high concentration (83 ppt for P; 65 ppt for S) in the unspiked sample.

Conclusions

The Agilent 8900 ICP-QQQ operating in MS/MS mode provides the sensitivity, low backgrounds, and unmatched control of interferences required for the analysis of ultratrace elements in high purity nitric acid.

Forty-nine elements were measured at sub-ppt to ppt levels in undiluted high purity 68% HNO $_3$. Calibrations were linear for all elements between 0-40 ppt. SEMI-specified elements were quantified at the single-figure ppt or sub-ppt level in high purity 68% HNO $_3$. The reproducibility results for 30 ppt spikes in high purity undiluted 68% HNO $_3$ were between 0.4-5.5% RSD for all elements except P and S, in a sequence lasting 6.5 hours.

The results demonstrate the suitability of the Agilent 8900 Semiconductor configuration ICP-QQQ for the routine analysis of the highest-purity semiconductor-grade reagents and process chemicals.

References

1. SEMI C35-0708, Specifications and guidelines for nitric acid (2008).

More Information

For more information on Agilent ICP-MS products and services, visit our website at www.agilent.com/chem/icpms

When analyzing 61-68 % HNO $_3$ on a routine basis, it is recommended to use the following options:

- G3280-67056 Pt sampling cone (18 mm insert)
- G4915A Upgrade to dry pump
- G3666-67030 Interface valve kit ball type valve

Analysis of Trace Metal Impurities in High Purity Hydrochloric Acid Using ICP-QQQ

Authors

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Introduction

Hydrochloric acid (HCI) is a component of the standard RCA cleaning process used to remove organic and metallic residues and impurities from the surface of silicon wafers used in semiconductor manufacturing. The cleaning steps are performed before high temperature processing steps such as oxidation and chemical vapor deposition (CVD). RCA Standard Clean 2 (SC-2) removes ionic contaminants from the wafer surface. SC-2 follows SC-1, which removes organic residues and particles. SC-2 consists of HCl combined with hydrogen peroxide (H_2O_2) and de-ionized water (DIW). Since the cleaning solutions are in direct contact with the silicon wafer surface, ultrahigh purity reagents are required for these solutions.

SEMI standard C27-0708 Tier-C protocol for HCl specifies a maximum contaminant level of 100 ppt for each element (HCl 37.0 - 38.0 %) [1]. The concentration of industrial grade HCl is usually 20 or 35%, depending on the method of production. The Cl matrix leads to the formation of several polyatomic ions, which cause significant spectral interferences on some key elements. For example, H₂³⁷Cl⁺ on ³⁹K⁺, ³⁵Cl¹⁶O⁺ on ⁵¹V⁺, ³⁵Cl¹⁶OH⁺ on ⁵²Cr⁺, ³⁷Cl¹⁶O⁺ on ⁵³Cr⁺, ³⁵Cl³⁷Cl⁺ on ⁷²Ge⁺, ³⁷Cl₂⁺ on ⁷⁴Ge⁺, and ⁴⁰Ar³⁵Cl⁺ on ⁷⁵As⁺. As a result of these polyatomic interferences, it has been difficult to determine these elements at the required levels using conventional single quadrupole ICP-MS (ICP-QMS). Even ICP-QMS instruments fitted with a collision/reaction cell (CRC) or bandpass filter can only offer limited reduction of the spectral interferences arising from the Cl matrix. Consequently, some methods for the analysis of high purity HCl by ICP-QMS have recommended sample pretreatment steps to remove the chloride matrix, which can lead to analyte loss and/or sample contamination.

In this study, triple quadrupole ICP-MS (ICP-QQQ) was used to analyze 50 elements in HCl, using MS/MS mode to resolve the polyatomic interferences. All analytes, including the most problematic elements such as K, V, Cr, Ge, and As, could be determined directly in the undiluted HCl with single digit ppt detection limits.

Experimental

Instrumentation

An Agilent 8900 Semiconductor configuration ICP-QQQ was used in this study. The instrument was fitted with a PFA-100 nebulizer, Peltier-cooled quartz spray chamber, quartz torch, platinum-tipped sampling and skimmer cones and s-lens.

The nebulizer was operated in self-aspiration mode to minimize the potential for sample contamination from the peristaltic pump tubing. In advanced semiconductor applications, the key requirement is to deliver the absolute lowest possible detection limits (DLs) for each analyte. To achieve this goal, laboratories measuring ultratrace levels of contaminants can use a multi-tune method, where several tuning steps are applied sequentially during the measurement of each solution. This approach allows the tuning conditions to be optimized for the removal of different types of interferences, while maintaining maximum sensitivity for each analyte. In this work, several reaction cell gases (H₂, O₂, and NH₃) were

used as appropriate for the large number of analytes being measured. He was used as a buffer gas in the $\mathrm{NH_3}$ reaction gas modes. Tuning conditions are shown in Table 1 and other acquisition parameters are shown in Table 2.

 Table 1. ICP-QQQ operating conditions.

	Cool	Cool-NH ₃	No gas	H ₂	0,	NH ₃	O ₂ -soft	
Acquisition mode				MS/MS		ı		
RF power (W)	600			15	00			
Sampling depth (mm)	18.0			18 8	•			
Nebulizer gas (L/min)				0.70				
Make-up gas (L/min)	0.90			0	48			
Extract 1 (V)	-1	50	4.2	4.7	4.	5	3.5	
Extract 2 (V)	-18.0	-17.0		-25	0.0 -		-120.0	
Omega bias (V)	-70	0.0		-14	0.0		-70.0	
Omega lens (V)	2	.0	10.0	8.0	10.5 4.0			
Q1 entrance (V)	-1	5.0			-50.0			
He flow (mL/min)	-	1.0	-	-	-	1.0	-	
H ₂ flow (mL/min)	-	-	-	7.0	-	-	-	
NH ₃ flow (mL/min)		2.0 (20%)	-	-	-	2.0 (20%)	-	
O ₂ flow (mL/min)	-	-	-	-	0.45 (30%)		0.45 (30%)	
Axial acceleration (V)	0.0	1.5	0.	.0	1.0	0.2	1.0	
Energy discrimination (V)	15.0	-5.0	5.0	0.0		-7.0		

 Table 2. Acquisition parameters.

Parameter	Setting
Q2 peak pattern	1 point
Replicates	3 (spiked samples) 10 (unspiked solution)
Sweeps/replicate	10
Integration time	2 s for all isotopes

Samples and standards

The samples of HCl used in this study included:

- Sample 1: 20% HCl (high purity grade).
- Sample 2: 36% HCl (non-high purity grade).
- Sample 3: 20% HCl (34% high purity grade diluted to 20% with DIW).

No further sample preparation was necessary as all samples were introduced directly into the ICP-QQQ. To run undiluted HCl routinely, it is recommended that the large (18 mm) insert Pt cone is fitted. Long-term corrosion of internal ICP-MS components can be minimized by fitting the dry pump option.

Calibration and quantification were done using the method of standard additions (MSA). Standard solutions were prepared by spiking a multi-element standard solution (SPEX CertiPrep, NJ, US) into each HCl sample type to give spike levels of 10, 20, 30, and 40 ppt. The MSA calibrations were then automatically converted to external calibrations in the ICP-MS MassHunter data analysis table. This conversion allows other samples of the same type (HCl concentration) to be quantified without requiring separate MSA spike additions into each sample. All solutions were prepared just before analysis.

All preparation and analysis was performed in a Class 10,000 clean room.

Results and Discussion

DLs and BECs

In total, 50 elements including all SEMI specification analytes were measured using the 8900 ICP-QQQ operating in multiple tune modes. Data for each mode was combined automatically into a single report for each sample. Detection limits (DLs) and background equivalent concentrations (BECs) in 20% HCl are given in Table 3.

Table 3. DLs and BECs in high purity 20% HCI*.

Element	Cell gas mode	Q1 mass	Q2 mass	DL ng/L	BEC ng/L
Li	Cool-NH ₃	7	7	0.032	0.016
Ве	No gas	9	9	0.022	0.021
В	No gas	11	11	0.55	4.1
Na	Cool-NH ₃	23	23	0.064	0.15
Mg	Cool-NH ₃	24	24	0.077	0.056
Al	Cool-NH ₃	27	27	0.20	0.19
P	O ₂ -soft	31	47	1.1	2.6
К	Cool-NH ₃	39	39	0.087	0.17
Ca	Cool-NH ₃	40	40	0.44	0.68
Sc	O ₂ -soft	45	61	0.014	0.012
Ti	O ₂ -soft	48	64	0.051	0.074
V	NH ₃	51	51	0.11	0.19
Cr	Cool-NH ₃	52	52	0.18	0.12
Mn	Cool-NH ₃	55	55	0.016	0.006
Fe	Cool-NH ₃	56	56	0.24	0.27
Со	Cool-NH ₃	59	59	0.10	0.038
Ni	Cool-NH ₃	60	60	0.66	0.26
Cu	Cool-NH ₃	63	63	0.10	0.12
Zn	NH ₃	66	66	0.14	0.097
Ga	NH ₃	71	71	0.015	0.026
Ge	NH ₃	74	107	0.90	3.0
Ge	NH ₃	74	107	0.32	0.77
As	0,	75	91	1.4	48
As	02	75	91	0.73	6.2
Se	H ₂	78	78	0.44	0.52

^{*} Shaded rows for Ge and As indicate results measured in Sample 3, due to suspected contamination for these elements in Sample 1.

Element	Cell gas mode	Q1 mass	Q2 mass	DL ng/L	BEC ng/L
Rb	Cool-NH ₃	85	85	0.041	0.013
Sr	NH ₃	88	88	0.003	0.001
Υ	O ₂ -soft	90	106	0.010	0.006
Zr	O ₂ -soft	93	125	0.012	0.004
Nb	02	93	125	0.004	0.005
Мо	He	98	98	0.13	0.57
Ru	He	101	101	0.016	0.003
Pd	He	105	105	0.010	0.001
Ag	He	107	107	0.032	0.014
Cd	He	114	114	0.090	0.10
In	He	115	115	0.035	0.021
Sn	He	118	118	0.57	3.3
Sb	He	121	121	0.66	1.5
Те	H ₂	125	125	0.37	0.31
Cs	NH ₃	133	133	0.008	0.019
Ba	NH ₃	138	138	0.005	0.005
Hf	No gas	178	178	0.005	0.004
Та	He	181	181	0.013	0.010
W	No gas	182	182	0.039	0.062
Re	No gas	185	185	0.12	0.50
Ir	No gas	193	193	0.017	0.012
Au	He	197	197	0.027	0.022
TI	No gas	205	205	0.007	0.004
Pb	H ₂	208	208	0.028	0.023
Bi	No gas	209	209	0.024	0.030
Th	No gas	232	232	0.017	0.021
U	No gas	238	238	0.009	0.005

Quantitative results

Table 4 shows quantitative data for all SEMI specification elements in high purity 20% HCl and non-high purity 36% HCl determined by MSA. The results show that the 8900 ICP-QQQ can measure contaminants in HCl at a much lower level than the 100 ppt maximum limit specified in the SEMI specifications. It is important to note that the concentration specified by SEMI is for 37–38% HCl while the data presented here is for 20 and 36% HCl. Even taking this difference into account, the 8900 ICP-QQQ is clearly able to measure contaminants at levels far lower than current industry requirements for high-purity HCl.

Table 4. Quantitative results for SEMI specification elements in high purity 20% HCl (Sample 1) and non-high purity 36% HCl (Sample 2).

Element	Cell gas mode	Q1	Q2	Sample 1 20% HCl, ng/L	Sample 2 36% HCl, ng/L	DL, ng/L
Li	Cool-NH ₃	7	7	<dl< td=""><td><dl< td=""><td>0.032</td></dl<></td></dl<>	<dl< td=""><td>0.032</td></dl<>	0.032
В	No gas	11	11	4.1	15	0.55
Na	Cool-NH ₃	23	23	0.15	6.4	0.064
Mg	Cool-NH ₃	24	24	<dl< td=""><td>6.5</td><td>0.077</td></dl<>	6.5	0.077
Al	Cool-NH ₃	27	27	<dl< td=""><td>23</td><td>0.20</td></dl<>	23	0.20
К	Cool-NH ₃	39	39	0.17	1.5	0.087
Ca	Cool-NH ₃	40	40	0.68	13	0.44
Ti	O ₂ -soft	48	64	0.074	1.4	0.051
V	NH ₃	51	51	0.19	4.6	0.11
Cr	Cool-NH ₃	52	52	<dl< td=""><td>0.55</td><td>0.18</td></dl<>	0.55	0.18
Mn	Cool-NH ₃	55	55	<dl< td=""><td>0.071</td><td>0.016</td></dl<>	0.071	0.016
Fe	Cool-NH ₃	56	56	0.27	7.6	0.24
Ni	Cool-NH ₃	60	60	<dl< td=""><td><dl< td=""><td>0.66</td></dl<></td></dl<>	<dl< td=""><td>0.66</td></dl<>	0.66
Cu	Cool-NH ₃	63	63	0.12	0.57	0.10
Zn	NH ₃	66	66	<dl< td=""><td>1.1</td><td>0.14</td></dl<>	1.1	0.14
As	0,	75	91	48	39	0.73*
Cd	He	114	114	0.10	0.34	0.090
Sn	He	118	118	3.3	2.3	0.57
Sb	He	121	121	1.5	0.95	0.66
Ва	NH ₃	138	138	0.005	<dl< td=""><td>0.005</td></dl<>	0.005
Pb	H ₂	208	208	0.023	0.13	0.028

^{*}DL for As measured in Sample 3, due to suspected contamination for this element in Sample 1.

Cr and K determination

Cool plasma is a proven technique used to remove plasma-based interferences. Although it has been largely superseded by CRC methodology, cool plasma remains the most effective analytical mode for some elements in certain matrices. Combining cool plasma with CRC technology has been shown to be a powerful mode for interference removal [2]. Because the major isotope of chromium (52 Cr $^+$) suffers an interference from 35 Cl 16 OH $^+$ in high purity HCl, Cr was determined using cool plasma with ammonia cell gas. The calibration curve for 52 Cr shows that 35 Cl 16 OH $^+$ interference was removed successfully, allowing a BEC of 0.12 ng/L (ppt) to be achieved, with a detection limit of 0.18 ppt (Figure 1). The DL and BEC displayed in the ICP-MS MassHunter calibration plots are based on the 10 replicates of the unspiked high-purity 20% HCl sample.

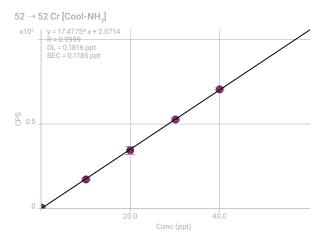


Figure 1. 52 Cr calibration curve obtained using cool plasma and NH_3 cell gas, showing low BEC and good linearity.

The same approach is effective for the determination of other interfered elements such as K. Figure 2 shows that the interference from $H^{237}Cl^+$ on $^{39}K^+$ was suppressed using cool plasma and NH_3 cell gas, giving a BEC and DL for K of 0.17 ppt and 0.09 ppt, respectively.

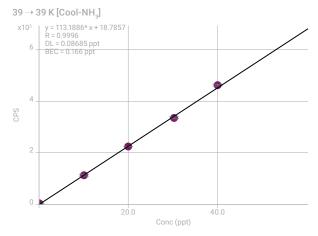


Figure 2. ³⁹K calibration curve obtained using cool plasma and NH₃ cell gas.

V and Ge determination

ICP-QMS fitted with a CRC operating in helium collision mode can successfully eliminate many polyatomic ions using He collision cell gas and kinetic energy discrimination (KED) [3]. However, ICP-QMS has some serious limitations when highly reactive cell gases, such as NH₂, are used in the CRC.

ICP-QMS has no mass selection step before the cell, so all ions enter the CRC. It is likely, therefore, that new reaction product ions will form in the CRC that may overlap the target analyte mass of interest. Bandpass ICP-QMS instruments, where all ions within a certain mass range (usually about 10 u) of the target analyte can enter the cell and react, have similar limitations to traditional ICP-QMS in terms of controlling reaction chemistry with highly reactive cell gases.

ICP-QQQ with MS/MS removes this limitation, as the first quadrupole mass filter (Q1), which is located before the CRC, allows precise selection of the specific mass of ions that are allowed to enter the cell. This extra mass selection step ensures that reaction processes in the cell are controlled, which removes the

potential for non-target product ion overlaps and dramatically improves the detectability of the analyte ions.

MS/MS acquisition mode using NH $_3$ as the reaction cell gas was used for the trace determination of V and Ge. The CIO $^+$ interference on 51 V was removed using NH 3 on-mass mode. Potentially, 14 NH 235 Cl $^+$ could form in the cell and interfere with V at m/z 51. However, the unit mass resolution of Q1 on the 8900 ICP-QQQ ensures that only ions at m/z 51 can enter the cell. All other matrix and analyte ions, e.g. 35 Cl $^+$, are prevented from entering the cell and cannot, therefore, contribute to the signal at the analyte mass. This simple approach avoids the formation of any new product ion interferences on 51 V.

The CICl $^+$ interference on 74 Ge was avoided by measuring a Ge-ammonia cluster ion, 74 Ge[14 NH $_2$ (14 NH $_3$)] $^+$, in mass-shift mode at mass 107. Q1 (set to m/z 74 to allow the 74 Ge $^+$ precursor ions to enter the cell) rejects all non-target masses, including 107 Ag $^+$, which would otherwise overlap the Ge-NH $_3$ product ion mass. Q1 (in contrast to a bandpass filter) also rejects all other nearby analyte ions, 70 Zn $^+$, 71 Ga $^+$, 73 Ga $^+$, 75 As $^+$, 78 Se $^+$, etc., preventing them from forming potentially overlapping ammonia clusters at the target product ion mass.

Representative calibration curves for V and Ge are shown in Figure 3, again illustrating the low BEC (0.19 ppt for V and 0.77 ppt for Ge) and DL (0.11 ppt for V and 0.32 ppt for Ge) achieved with the 8900 with NH₂ cell gas in MS/MS mode.

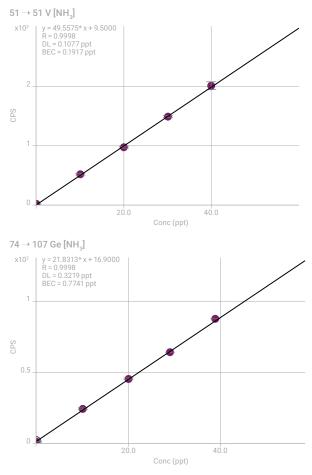


Figure 3. 51 V and 74 Ge calibration curve obtained using NH $_{3}$ cell gas.

Determination of As

Arsenic has a single isotope at m/z 75 that suffers an interference from the polyatomic ion 40 Ar 35 Cl $^+$. Since ArCl $^+$ readily forms in a chloride matrix, the polyatomic interference compromises the determination of As at ultratrace levels in concentrated HCl using ICP-QMS. Oxygen can be used as the cell gas to avoid this overlap, with As being measured as the AsO $^+$ product ion at m/z 91. However, with ICP-QMS, the AsO $^+$ product ion at mass 91 suffers an interference from 91 Zr $^+$. Helium collision mode in the Agilent ORS cell can reduce ArCl $^+$ effectively, allowing a BEC of less than 20 ppt to be achieved by ICP-QMS [3]. But, as semiconductor industry demands become more stringent, this sensitivity may not be sufficient for the lowest level of ultratrace analysis.

Using the 8900 ICP-QQQ with MS/MS, the $^{91}Zr^+$ ion is removed by Q1, which is set to the As+ precursor ion mass of 75. MS/MS mode allows O_2 cell gas to be used successfully, with As being measured as the AsO+ product ion at m/z 91 without overlap from $^{91}Zr^+$. A further benefit of O_2 cell gas is that measuring AsO+ provides more sensitivity that direct measurement of As+ in He mode.

A calibration curve for As in 20% HCl (Sample 3) is shown in Figure 4, demonstrating a BEC of 6.17 ppt and a DL of 0.73 ppt. While lower than the industry requirements for high-purity HCl, this BEC doesn't represent the best performance that can be achieved with the 8900 ICP-QQQ, so further investigation was done to identify the cause of the relatively high background.

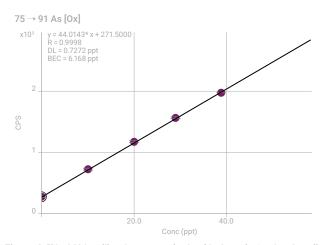


Figure 4. 75 As MSA calibration curve obtained in Sample 1 using O_2 cell gas.

Investigation of arsenic contamination

As the measured result for As was relatively high in high purity HCl Sample 1 (Table 4), the signal count at m/z 91 (mass of the product ion AsO+) was investigated further. In a high Cl matrix, the polyatomic ion 40 Ar 35 Cl+ forms in the plasma and during ion extraction. This polyatomic ion has the same nominal mass as the target 75 As+ precursor ion, so it passes through Q1 and enters the cell. While not thermodynamically favored, the ArCl+ might react with the O_2 cell gas to form ArClO+, which would therefore remain as an interference on AsO+ at m/z 91. This possibility can be checked by comparing the isotopic signature of the Cl-based product ions observed in the mass spectrum. Since chlorine has two isotopes, 35 and 37, the ratio of the natural abundances of these isotopes (75.78%: 24.22%) can be used to confirm whether a product ion is Cl-based.

The signals of the mass-pairs 75/91 and 77/93, representing the potential CI interferences 40 Ar 35 Cl 16 O+ and 40 Ar 37 Cl 16 O+ respectively, were measured by ICP-QQQ with MS/MS. A neutral gain scan spectrum (where Q1 and Q2 are scanned synchronously, with a fixed mass difference between them) was measured and the scan is presented in Figure 5. For this neutral gain scan, Q1 was scanned across the mass range from 74 to 78 u to pass any precursor ions to the CRC, and Q2 was scanned synchronously at Q1 + 16, monitoring any product ions formed by O-atom addition. The peak at mass-pair m/z 75/91 that caused the relatively high BEC for As in Sample 1 is clearly visible. However, if the signal at 75/91 was due to interference from 40 Ar 35 Cl 16 O+, there would also be a corresponding signal from 40 Ar 37 Cl 16 O+ at mass-pair 77/93. Since there was no signal observed at 77/93, we can conclude that the signal at m/z 75/91 is not due to any contribution from ArClO+, and the high reported concentration of As in Sample 1 is due to contamination.

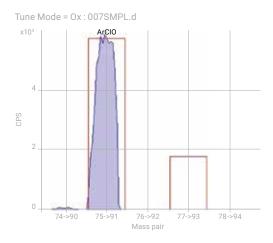


Figure 5. Neutral gain scan spectrum for 20% high purity HCl showing the theoretical isotope template for 40 Ar 35 Cl 16 O $^{+}$ and 40 Ar 37 Cl 16 O $^{+}$. Q1 was scanned from m/z 74 to 78, while Q2 was set to Q1 + 16.

Conclusions

The high performance of Agilent ICP-QQQ systems for the analysis of trace metallic impurities in concentrated HCl has been described previously [4]. Now, the Agilent 8900 Semiconductor configuration ICP-QQQ with flexible cell gas support, unique MS/MS capability, and unparalleled cool plasma performance, further improves the detection limits for the analysis of a wide range of trace metal contaminants in high purity acids. The advanced reaction cell methodology supported by the 8900 ICP-QQQ allows the SEMI elements, including those elements with potential matrix-based interferences such as K, V, Cr, Ge, and As, to be determined at lower concentrations in a chloride matrix than was previously possible.

References

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- 2. Junichi Takahashi and Katsuo Mizobuchi, Use of Collision Reaction Cell under Cool Plasma Conditions in ICP-MS, 2008 Asia Pacific Winter Conference on Plasma Spectroscopy
- 3. Junichi Takahashi, Direct analysis of trace metallic impurities in high purity hydrochloric acid by Agilent 7700s/7900 ICP-MS, Agilent publication, 2017, 5990-7354EN

More Information

When analyzing 20–36% HCl on a routine basis, it is recommended to use the following options:

- G3280-67056 Pt sampling cone (18 mm insert)
- G4915A Upgrade to dry pump
- G3666-67030 Interface valve kit ball type valve

Since hydrochloric acid is corrosive, avoid placing open sample bottles near the instrument.

Ultra-low level determination of phosphorus, sulfur, silicon and chlorine using the Agilent 8900 ICP-QQQ

Author

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Introduction

Quadrupole ICP-MS (ICP-QMS) is one of the most sensitive and versatile analytical tools used in inorganic analysis. With sensitivity approaching 1,000 million counts per second/part per million (1 G cps/ppm) and background signals typically less than 1 cps, the latest instrumentation achieves detection limits (DL) in the ppq (pg/L) range for most of the elements in the periodic table. Detection limits tend to be lowest for elements at masses higher than 80 amu, while some lower mass elements are more difficult to measure at trace levels due to the presence of spectral overlaps from polyatomic interferences. ICP-QMS can utilize cool plasma and/or collision/reaction cell methods to address the problem of background interferences, with successful results in many applications.

More recently, the introduction of triple quadrupole ICP-MS (ICP-QQQ) has dramatically improved the reliability and performance of reaction cell methods by allowing a double mass filter (MS/MS) to be applied to control reaction chemistry in the cell. This now allows analysts to resolve interferences on a wide range of elements in a controlled and effective manner [1].

With the introduction of Agilent's second generation ICP-QQQ instrument, the Agilent 8900 Triple Quadrupole ICP-MS, reaction cell operation with MS/MS mode has been further refined. This note describes the performance of the 8900 ICP-QQQ for the analysis of some of the most challenging elements for ICP-MS: phosphorus (P), sulfur (S), silicon (Si), and chlorine (Cl). The first ionization potentials of these elements are relatively high, which reduces the degree of ionization and therefore the analyte signal. Furthermore, the background signals are elevated due to plasma-, solvent- and matrix-based polyatomic ions, making low-level analysis even more difficult. As ICP-MS technology has developed, there has been a growing demand and expectation to measure these difficult elements together with more conventional elements in high purity chemicals and materials. Details of the methods used to control the interferences on the four elements are presented, together with background equivalent concentrations (BECs) and detection limits (DLs) for P, S, Si and Cl in ultra-pure water (UPW), and P, S and Si in the highest grade hydrogen peroxide (H_2O_2).

Experimental

Instrumentation

An Agilent 8900 ICP-QQQ (#200, Semiconductor Configuration) was used for all measurements. The sample introduction system comprised a PFA concentric nebulizer, a quartz spray chamber and torch, and platinum interface cones. The 8900 #200 ICP-QQQ is fitted with a new argon gas flow control system specially designed to minimize sulfur/silicon contamination from the gas line components.

Normal, hot plasma conditions were used throughout. Extraction lens voltages were optimized for maximum sensitivity using an Agilent 1 ppb tuning solution containing Li, Y, Ce and Tl. Operating and tuning parameters are summarized in Table 1.

Table 1. Agilent 8900 ICP-QQQ operating parameters.

Parameter	Unit	Value
RF power	W	1500
Sampling depth	mm	8.0
Carrier gas flow rate	L/min	0.70
Make-up gas flow rate	L/min	0.52
Extraction lens 1	V	4.0
Extraction lens 2	V	-210
Omega lens bias	V	-80
Omega lens	V	8.0

Method and cell tuning

Based on previous studies, oxygen (O_2) mass-shift mode was used for the analysis of P and S [2], hydrogen (H_2) on-mass mode was used for Si, and Cl was determined using H_2 mass-shift mode [3]. The reaction processes used for removal of the primary interference on each analyte were as follows:

Sulfur by oxygen mass-shift mode

The intense polyatomic interference from $^{16}O_2^+$ on the primary isotope of S, $^{32}S^+$ at m/z 32, is avoided by shifting S⁺ away from the interfering O_2^+ ion, using an O-atom addition reaction. S⁺ reacts readily with O_2 cell gas to form the product ion SO⁺, which can be measured free of interference at M + 16 amu (m/z 48 for the primary $^{32}S^{16}O^+$ isotope product ion), as shown in the following equations:

$$^{32}S^+ + O_2 < cell gas > \rightarrow ^{32}S^{16}O^+ + O$$

 $^{16}O_2^+ + O_2 < cell gas > \rightarrow$ no reaction

Phosphorus by oxygen mass-shift mode

A similar mass-shift approach is used for the measurement of P as P0 $^{+}$. The native mass of P (m/z 31) suffers an intense background interference from $^{14}N^{16}O^{1}H^{+}$, $^{15}N^{16}O^{+}$, and $^{14}N^{17}O^{+}$. These background polyatomic ions are avoided by reacting P $^{+}$ with O $_{2}$ cell gas, shifting the P $^{+}$ away from the interfering ions, and measuring it as the PO $^{+}$ product ion at m/z 47:

$$^{31}P^+ + O_2$$
 \rightarrow $^{31}P^{16}O^+ + O$
NOH+/NO+ + O_2 \rightarrow no reaction

Silicon by hydrogen on-mass mode

The analysis of Si uses on-mass measurement with H_2 cell gas, as the primary interferences on the major Si isotope at m/z 28, $^{14}N_2$ + and $^{12}C^{16}O$ +, react readily with H_2 , while Si⁺ does not react. Thus the N_2 ⁺ and CO⁺ interferences can be removed, and ^{28}Si ⁺ can be measured free from the interferences at its original mass:

28
Si⁺ + H₂ → no reaction
 14 N₂⁺ + H₂ → N₂H⁺ + H
 12 C¹⁶O⁺ + H₂ → COH⁺ + H

Chlorine by hydrogen mass-shift mode

Cl is a difficult element to analyze at low concentrations using ICP-MS, because it is a common contaminant and is often present in reagents used in the laboratory environment. In addition, its first ionization potential of 12.967 eV is higher than that of any other commonly measured element, meaning that Cl is very poorly ionized, so the sensitivity for Cl $^+$ is extremely low. A further issue for low-level Cl analysis is the presence of a polyatomic interference from $^{16}O^{18}O^{1}H^+$ on the primary Cl isotope at m/z 35. The O_2H^+ overlap can be avoided by measuring Cl as a ClH_2^+ product ion, produced from sequential reaction with H_2 reaction gas:

$$^{35}\text{Cl}^+ + \text{H}_2 < \text{cell gas} > \rightarrow ^{35}\text{Cl}^1\text{H}^+ + \text{H}$$
Followed by $^{35}\text{Cl}^1\text{H}^+ + \text{H}_2 < \text{cell gas} > \rightarrow ^{35}\text{Cl}^1\text{H}_2^+ + \text{H}$
 $^{16}\text{O}^{18}\text{O}^1\text{H}^+ + \text{H}_2 < \text{cell gas} > \rightarrow \text{no reaction}$

In all of these methods, the Agilent 8900 ICP-QQQ was operated in MS/MS mode (where both Q1 and Q2 function as mass filters) ensuring that only the target ion or product ion was measured. MS/MS means that potentially overlapping ions are excluded from the collision/reaction cell, so the reaction chemistry is controlled and consistent, even if other matrix elements or analytes are present in the sample. For example, in the case of 32S16O+ product ion measured at *m/z* 48, the product ion mass could be overlapped by other ions, such as 48Ca+, 48Ti+, and 36Ar12C+, if these ions were not rejected by Q1. This is the main reason for the improved reaction mode performance of ICP-QQQ compared to ICP-QMS, as ICP-QMS has no mass filter step before the collision/reaction cell.

The ORS4 collision/reaction cell of the 8900 #200 instrument has the facility to utilize an axial acceleration voltage, which was found to be effective to increase sensitivity in the O2 mass-shift method used for the determination of P and S. Cell parameters were optimized separately for each mode while aspirating a 1 ppb standard solution of each of the elements. Cell tuning parameters are summarized in Table 2.

Table 2. Cell mode related tuning parameters.

Parameter	Unit	O ₂ mass-shift	H ₂ on-mass	H ₂ mass-shift			
Element		³¹ P, ³² S	²⁸ Si	³⁵ CI			
Mass pair	(Q1 → Q2)	(31 → 47), (32 → 48)	(28 → 28)	(35 → 37)			
Cell gas		02	H ₂				
Flow rate	mL/min	0.41	5.0				
OctpBias	V	-3	-18				
KED	V	-8	0				
Axial acceleration	V	1	0				
Cell exit	V	-90	-70				
Deflect	V	8	-6				
Plate bias	٧		-60				

Reagents

Standard solutions for P, S and Si were prepared from single element standards purchased from SPEX CertiPrep (NJ, USA), by serial dilution with UPW. The UPW was supplied from ORGANO Corp (Tokyo, Japan). The Cl standard was prepared from high purity HCl purchased from Wako Pure Chemicals Industries Ltd (Osaka, Japan). The highest purity grade $\rm H_2O_2$, TAMAPURE-AA-10, was purchased from TAMA Chemicals Co Ltd (Kanagawa, Japan). The calibration standard addition spikes were added directly to the undiluted $\rm H_2O_2$. A 1% TMAH alkaline rinse was used during the analysis of Cl to maximize the effectiveness of the washout between samples, and prevent any carryover. All pipette tips, vials and bottles were thoroughly cleaned using diluted high purity acids, and rinsed in UPW prior to use.

Results and Discussion

To prepare the ICP-QQQ for the analysis, a 1% $\mathrm{HNO_3}$ solution was aspirated overnight to thoroughly clean the sample introduction system. Running the plasma for several hours would also help to remove any contaminants in the Ar gas flow line. P, S and Si were measured together, and Cl was analyzed in a separate batch since it benefited from an alkaline rinse between solutions. Figures 1 and 2 show the calibration curves of the four elements in UPW and P, S and Si in $\mathrm{H_2O_2}$, respectively, measured using the method of standard addition (MSA). The background level of Cl present in the $\mathrm{H_2O_2}$ sample was too high to permit accurate analysis at the spike levels used.

Good linearity at low and sub-ppb levels was observed for all elements measured in both of the sample matrices. The DL for each element was calculated as 3 times the standard deviation of 10 replicates of the blank using an integration time of 1 sec for each element. The results are summarized in Table 3.

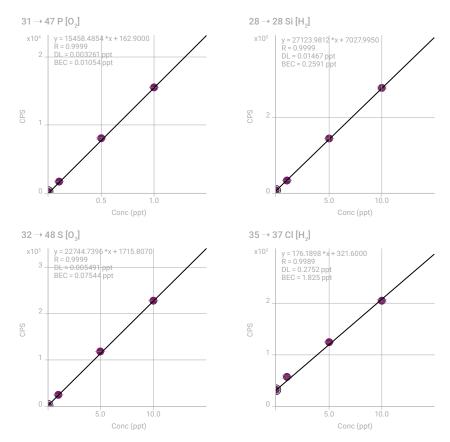


Figure 1. Calibration plots of P, S, Si and Cl in UPW. All values in ug/L (ppb).

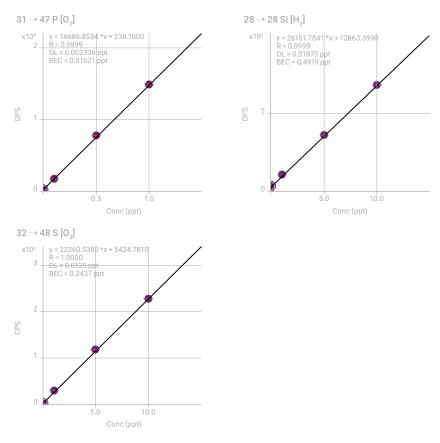


Figure 2. Calibration plots of P, S and Si in H_2O_2 .

Table 3. BEC and DL of P, S, Si and Cl in UPW and P, S and Si in the highest grade H₂O₂.

	P (ppt)		S (ppt)		Si (ppt)		Cl (ppt)	
Element	BEC	DL	BEC	DL	BEC	DL	BEC	DL
UPW	10.5	3.3	75.4	5.5	259	14.7	1.83	0.28
H ₂ O ₂	16.2	2.3	244	12.5	492	18.8		

Conclusions

The Agilent 8900 ICP-QQQ operating in MS/MS mode with $\rm O_2$ and $\rm H_2$ cell gases successfully eliminated problematic spectral interferences on non-metallic impurities P, S, Si and Cl in UPW and P, S and Si in $\rm H_2O_2$. The results highlight the advanced performance of the second generation ICP-QQQ for the analysis of challenging elements, by achieving the lowest ever reported BECs for the four elements in UPW.

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Arsenic Measurement in Cobalt Matrix using MS/ MS Mode with Oxygen Mass-Shift

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Keywords

arsenic, high purity metals, cobalt, zirconium, oxygen mass-shift

Introduction

Measuring the purity of materials such as high purity metals is of interest across advanced technology industries, to support the development of new materials and/or improve the performance of existing products. ICP-MS is widely used for determining elemental impurities in these materials due to its unique features: High sensitivity, low DLs, multi element analysis capability, wide dynamic range, fast analysis and minimal sample preparation requirements.

For many applications, the errors caused by spectral interferences in quadrupole ICP-MS have been adequately addressed by the introduction of CRC technology. However, the analysis of trace contaminants in high purity materials presents a particular challenge due to the high matrix levels and the need to determine impurities at the trace level. For example, the determination of As in Co is difficult for quadrupole ICP-MS due to the signal from CoO^+ that overlaps the only isotope of arsenic at m/z 75. Although only about

0.01% of the Co ions in the plasma are present as CoO^+ ions, the Co concentration in a 1000 ppm solution is 6 or 7 orders of magnitude higher than the trace levels of As that are of interest in this application. Consequently the CoO^+ interference is still very significant relative to the As $^+$ signal. This note describes the measurement of trace As in a 1000 ppm Co solution using an Agilent 8800 ICP-QQQ in MS/MS mass-shift mode, using oxygen as the reaction gas.

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/HMI-mid.

CRC conditions: O_2 gas at 0.3 mL/min, Octopole bias = -5 V, KED = -7 V.

Acquisition conditions: Three oxygen (O₂) mass-shift operational modes were compared:

- Single Quad mode A with low mass cut off at m/z < 59
- Single Quad mode B with low mass cut off at m/z = 59
- MS/MS mode with Q1 as a 1 amu mass filter, Q1 = 75 and Q2 = 91

Sample: SPEX CLCO2-2Y (SPEX CertiPrep Ltd., UK) was used as 1000 ppm Co solution.

Results and Discussion

BEC of As in 1000 ppm Co solution using O2 mass-shift method

From the equation and reaction enthalpy below, it can be seen that arsenic reacts readily with O_2 cell gas via an O-atom transfer reaction. This creates the reaction product ion AsO+ at m/z 91, moving the analyte away from the CoO+ interference on As+ at m/z 75.

$$As^{+} + O_{2} \rightarrow AsO^{+} + O \quad \Delta Hr = -1.21 \text{ eV}$$

The reaction enthalpy for CoO^+ with the O_2 cell gas is much less favorable, so the overlap from the CoO^+ polyatomic interference is successfully avoided. To evaluate the effectiveness of MS/MS mode for this application, the 8800 ICP-QQQ was operated in three acquisition modes: MS/MS mode and two "Single Quad" modes, in which Q1 functions as a bandpass filter rather than a unit (1 amu) mass filter. In Single Quad mode A, Q1 was set to allow most of the plasma-formed ions to enter the cell; in Single Quad mode B, Q1 was set with a low mass cutoff around m/z 59 to allow only ions with a mass greater than 59 to enter the cell (most $^{59}Co^+$ ions are rejected); and finally in MS/MS mode Q1 was set to allow only ions at m/z 75 to enter the cell (all $^{59}Co^+$ rejected).

The BECs for As obtained using the three acquisition modes are shown in Figure 1. MS/MS mode achieved the lowest BEC for As of 330 ppt in 1000 ppm Co. The BEC obtained by the Single Quad modes were orders of magnitude higher, which suggests the occurrence of the following undesired reactions in the cell and indicates the incomplete rejection of Co⁺ by Single Quad mode B:

$$^{59}\text{Co}^+ + \text{O}_2 \rightarrow ^{59}\text{CoO}^+ + \text{O}_3 + ^{59}\text{CoO}_2 + \text{O}_3 + ^{59}\text{CoO}_2 + \text{O}_3 + ^{59}\text{CoO}_2 + \text{O}_3 + ^{59}\text{CoO}_3 + ^{59}\text{CoO}_2 + ^{59}\text{CoO}_3 +$$

Note that this sequential reaction chemistry leads to a relatively intense signal for CoO_2^+ , because the number of precursor ions for the reaction (the Co^+ ions from the plasma) is so high (10000 times higher intensity than the CoO^+ signal in the plasma). Consequently, in Single Quad mode, the CoO^+ overlap cannot be successfully avoided by moving the As^+ to its AsO^+ product ion at m/z 91 using O_2 cell gas, because CoO_2^+ (also at m/z 91) is formed relatively easily when a large number of Co^+ ions are allowed to enter the cell.

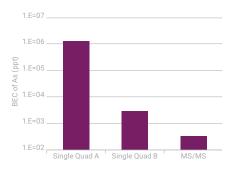


Figure 1. BEC of As in 1000 ppm Co solution with $\rm O_2$ mass-shift method using three acquisition modes (note log intensity scale).

AsO+ in the presence of zirconium

To successfully avoid interferences using the mass-shift method, the mass of the analyte product ion must itself be free from interference. For example, in this application the AsO+ product ion is measured at m/z 91 where it could be overlapped by an isotope of zirconium ($^{91}Zr^{+}$). The presence of Zr in a sample may therefore cause an error in the results for As measured as AsO+ using O_2 reaction mode on ICP-QMS. The potential effect of co-existing Zr on AsO+ measurement using ICP-QMS was investigated.

Figure 2 is a spectrum of 10 ppm Zr obtained using Single Quad mode A with O_2 mass-shift. Zr reacts with O_2 very efficiently (Δ Hr = -3.84) and is converted to ZrO $^+$. However not all the Zr $^+$ ions are converted to ZrO $^+$ so some Zr $^+$ remains, interfering with the measurement of AsO $^+$ at m/z 91. In contrast, in MS/MS mode the 91 Zr $^+$ ion is rejected by Q1, so the potential overlap on the AsO $^+$ product ion at m/z 91 is removed.

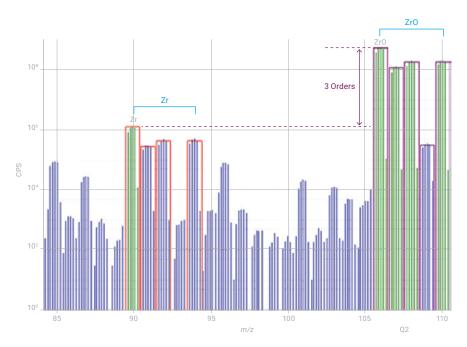


Figure 2. Spectrum of 10 ppm Zr obtained using Single Quad mode A with O_2 mass-shift-method

Conclusions

Trace levels of arsenic in a 1000 ppm cobalt matrix can be successfully measured (BEC of 330 ppt) using the 8800 ICP-QQQ operating in MS/MS mass-shift mode, with oxygen as the reaction gas. There are two main advantages of using MS/MS compared to ICP-QMS:

- 1. In MS/MS mode, Co^+ is prevented from entering the cell by Q1, which is set to m/z 75. In ICP-QMS, CoO_2^+ is formed in the cell via a chain reaction, and will interfere with AsO⁺ at m/z 91.
- 2. In MS/MS mode, the potential ${}^{91}Zr^+$ overlap on the AsO+ product ion at m/z 91 is eliminated, as ${}^{91}Zr^+$ ions (and all other ions apart from m/z 75) are rejected by Q1.

Determination of Sulfur, Phosphorus and Manganese in High Purity Iron

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Keywords

phosphorus, sulfur, manganese, iron, steel, JSS 001-6, JSS 003-6, abundance sensitivity, oxygen mass-shift

Introduction

ICP-MS is the analytical technique of choice for the analysis of trace elements in iron and steel. However, the sensitivity and interference removal performance of quadrupole ICP-MS (ICP-QMS) is not sufficient for the determination of difficult analytes such as phosphorus (P) and sulfur (S) at the low levels required. Furthermore, the determination of manganese (Mn) in an iron matrix is extremely challenging for ICP-QMS due to overlap (or tailing) from the very intense 54 Fe and 56 Fe peaks that occur either side of the single isotope of manganese at m/z 55.

The Agilent 8800 Triple Quadrupole ICP-MS (ICP-QQQ) provides more effective and reliable removal of polyatomic interferences, such as $^{14}\rm{N}^{16}\rm{OH}$ on $^{31}\rm{P}$ and $^{16}\rm{O_2}$ on $^{32}\rm{S}$, using controlled chemical reaction in the CRC. This note describes the performance of the 8800 ICP-QQQ operating in MS/MS mode, for the determination of the trace elements S, P and Mn in two high purity iron CRMs (JSS 001-6 and 003-6).

Experimental

Instrumentation: Agilent 8800 #100.

Plasma condition: Preset plasma/HMI-mid.

Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -175 V.

CRC and acquisition conditions:

- MS/MS O_2 mass-shift method to remove the $^{14}N^{16}OH^+$ and $^{16}O_2^+$ interferences on $^{31}P^+$ and $^{32}S^+$ respectively: O_2 gas at 0.3 mL/min, Octopole bias = -5 V and KED = -7 V.
- MS/MS He on-mass mode to measure ⁵⁵Mn+:
 He gas at 5.0 mL/min, Octopole bias = -18 V and KED = 4 V.

All other parameters were optimized by Autotune in the MassHunter software. Figure 1 shows the mechanism used on the 8800 ICP-QQQ to avoid the $^{14}N^{16}OH^+$ and $^{16}O_2^+$ interferences on $^{31}P^+$ and $^{32}S^+$ by mass-shift mode (Q1 \neq Q2) using O_2^- reaction gas.

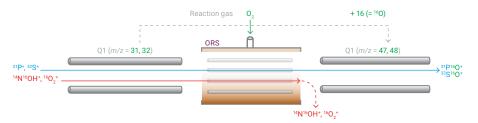


Figure 1. Mechanism of MS/MS mass-shift mode (Q2 = Q1+16) using O_2 reaction gas for the measurement of P as $^{31}P^{16}O^+$ and S as $^{32}S^{16}O^+$ at m/z 47 and 48 respectively.

Sample preparation: Two Steel CRMs, JSS-001 and JSS-003 were purchased from The Japan Iron and Steel Federation (Tokyo, Japan). 0.1 g of each Steel CRM was digested in a mixture of 1 mL HCl and 2 mL HNO $_3$ and diluted to 100 mL with UPW. No further chemical matrix separation, e.g., solvent extraction, ion exchange, etc. was applied. The digested CRM samples containing 0.1% (1000 ppm) Fe were analyzed directly on the ICP-QQQ using the robust plasma conditions provided by Agilent's HMI aerosol dilution system.

Results and Discussion

BEC and DL of P and S

The calibration plots shown in Figure 2 demonstrate that the 8800 ICP-QQQ with MS/MS mass-shift mode can successfully perform the trace level (single ppb) quantitation of P and S in 0.1% Fe solutions. The BEC and DL achieved for P were 0.14 ppb and 0.05 ppb respectively, and the BEC and DL for S were 6.45 ppb and 0.75 ppb respectively.

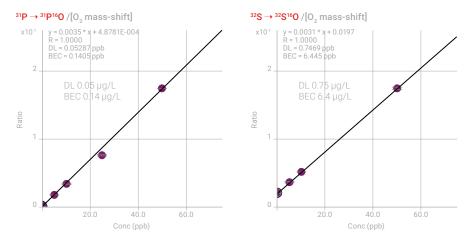


Figure 2. Calibration curve for P (left) and S (right) in 0.1% Fe matrix, obtained using \boldsymbol{O}_2 mass-shift mode under robust plasma conditions.

Trace Mn analysis in Fe matrix

The abundance sensitivity (AS, a measure of peak separation) of ICP-QQQ in MS/MS mode is the product of the Q1 AS x Q2 AS. This means the AS of the 8800 ICP-QQQ is theoretically about 2x that achievable on ICP-QMS, and the ICP-QQQ is therefore able to successfully separate the 55 Mn peak from the very intense overlaps from 54 Fe and 56 Fe in a high iron matrix. This is demonstrated in Figure 3 which shows the spectra of 10 ppb Mn in a 0.1% Fe matrix sample solution measured in Single Quad mode (left) and MS/MS mode on the ICP-QQQ (right). Helium was used as the cell gas in both cases to remove 54 FeH $^+$ and ArNH $^+$ interferences by KED.

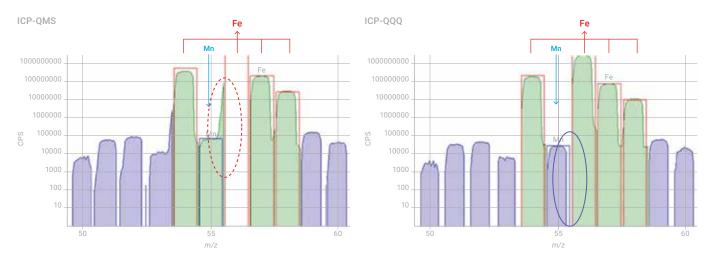


Figure 3. Spectra of 10 ppb Mn in a 0.1% Fe matrix sample solution obtained in Single Quad mode (left) and ICP-QQQ in MS/MS mode (right).

Determination of P, S and Mn in high purity iron CRMs

Trace elements including P, S and Mn were determined by ICP-QQQ in high purity iron CRMs: JSS 001-6 and 003-6, using $\rm O_2$ mass-shift mode (for P and S) and He mode (for Mn). As summarized in Table 1, excellent agreement was obtained between the measured (found) and certified values for all three elements, indicating the effective interference removal offered by the 8800 ICP-QQQ in MS/MS mode. Excellent spike recovery at the 50 ppb level was also confirmed with JSS 003.

Table 1. Analytical results for P, S and Mn in two high purity iron (
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				JSS 001-6			JSS 003-6			
Element	Q1	Q2	ORS	Certified value [mg/kg]	Uncertainty	Found [mg/kg]	Certified value [mg/kg]	Uncertainty	Found [mg/kg]	50 ppb spike recovery %
Р	31	47	0,	0.5*		0.458	3.5	0.7	3.170	103
s	32	48	0,	1.5	0.3	1.512	1.3	0.5	1.287	92
Mn	55	55	He	0.03*		0.036	3.2	0.2	3.432	101

Direct Measurement of Trace Rare Earth Elements in High Purity REE Oxides

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Keywords

Rare Earth Elements, REE, rare earth oxide, REO, samarium oxide, gadolinium oxide, oxygen mass-shift, ammonia on-mass

Introduction

The rare earth elements (REEs) are widely used in advanced technologies including high-power permanent magnets, lasers, phosphors used in fluorescent lamps, radar screens and plasma displays. REEs are also used in petroleum refining, automobile catalytic converters and batteries, and in high-technology glasses. It is clear from these examples that REEs play a key role in many types of materials used in high-technology industries. However, the presence of other REEs as contaminants in a purified single-element REE material often impacts the functionality of the final product, so impurities in the REE oxide raw material must be carefully controlled.

ICP-MS is the most commonly used atomic spectrometry technique for the measurement of trace REEs due to its simple REE spectra — particularly when compared to emission techniques. The measurement of mid- and high-mass REEs in a low-mass REE matrix is, however, very challenging for ICP-MS because REEs have among the highest metal-oxide (M-O) bond strengths of any element, and the oxide ions of the low mass REE overlap the preferred isotopes of the mid-mass and high-mass REEs. Table 1 shows the interferences observed in the analysis of trace REEs in high-purity samarium (Sm) oxide and gadolinium (Gd) oxide.

Separation of the trace REE analytes from the REE matrix can be performed utilizing a chelating resin, but this technique is time-consuming and customization is needed according to the analyte and matrix element. The direct analysis of trace REEs in a variety of high-purity REE matrices is therefore desired. In this work, an Agilent 8800 Triple Quadrupole ICP-MS was used for the direct analysis of trace REE in two high-purity REE materials: ${\rm Sm_2O_3}$ and ${\rm Gd_2O_3}$. Operating the ICP-QQQ in MS/MS mode effectively removes the challenging interferences, enabling the determination of REE impurities at trace levels in these two materials.

Table 1. Preferred isotope for ICP-MS analysis of each REE, and the potential interferences caused by Sm_aO_a and Gd_aO_a matrices.

Element	La	Се	Pr	Nd	Sm	Eu	Gd	Tb	Dy	Но	Er	Tm	Yb	Lu
Mass	139	140	141	146	147	153	157	159	163	165	166	169	172	175
Gd_2O_3							N/A	GdH⁺					GdO⁺	GdOH+
Sm ₂ O ₃					N/A	SmH⁺			Sm0+	Sm0+	Sm0+	Sm0H ⁺		

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/General purpose.

Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -180 V.

Acquisition parameters: Three cell modes were used with MS/MS acquisition: No gas, O_2 mass-shift, and NH $_3$ on-mass mode. In MS/MS O_2 mass-shift mode, the REEs were determined as their oxide ions. REE ions react efficiently with the O_2 cell gas and are converted to the oxide ion REE-O $^+$. For example, in the measurement of 153 Eu $^+$, Q1 is set to m/z 153 (153 Eu $^+$) and Q2 is set to m/z 169 (153 Eu 16 O $^+$). Cell tuning parameters are summarized in Table 2.

Table 2. CRC tuning parameters.

Cell mode	Unit	No gas	02	*NH ₃
Scan mode	MS/MS			
Cell gas		N/A	0,	NH ₃
Cell gas flow rate	mL/min	N/A	0.35	9.0
Octopole bias	٧	-8	-5	-18
KED	٧	5	-8	-8
Cell exit	٧	-80	-90	-110
Deflect lens	٧	20	10	-3
Plate	٧	-80	-90	-110

^{*10%} NH₃ balanced in Ar

Results and Discussion

Two high purity REE oxide materials $\mathrm{Gd_2O_3}$ (5N) and $\mathrm{Sm_2O_3}$ (4N8) were gently dissolved in semiconductor grade $\mathrm{HNO_3}$ and diluted to a concentration of 1 ppm (as the REE). The other (trace) REEs were measured in each matrix solution using the three cell modes. The results are given in Figure 1 and Figure 2. As expected, analysis of the 1 ppm Gd solution in no gas mode gave positive errors on some elements due to interferences from Gd polyatomic ions: $\mathrm{GdH^+}$ interferes with $^{159}\mathrm{Tb^+}$, $\mathrm{GdO^+}$ interferes with $^{172}\mathrm{Yb^+}$ and $\mathrm{GdOH^+}$ interferes with $^{175}\mathrm{Lu^+}$.

Preliminary studies showed that NH_3 cell gas reacts with many of the polyatomic ions that interfere with the REE. However, NH_3 also reacts quickly with some of the REE ions, leading to reduced sensitivity of < 1 cps/ppt [1], so this mode is only suitable for the measurement of the less reactive analytes: Pr, Eu, Dy, Ho, Er, Tm and Yb. For these elements, NH_3 on-mass mode gave excellent results, including for Yb in the Gd matrix, where the measured Yb background concentration was reduced by four orders of magnitude (Figure 1) indicating effective removal of the GdO+ overlap. Background signals for Dy, Ho, Er and Tm in the Sm matrix were also dramatically improved (Figure 2).

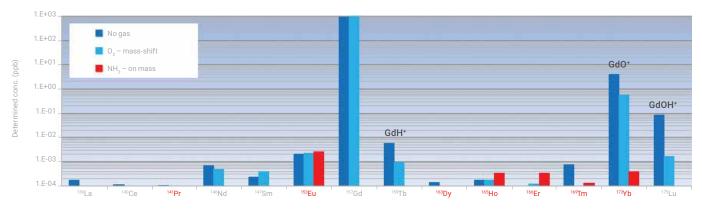


Figure 1. Measured concentration of REE impurities in 1 ppm Gd solution. Gd based interferences are observed on Tb, Yb and Lu. Only the elements in red were measured in NH, on-mass mode.

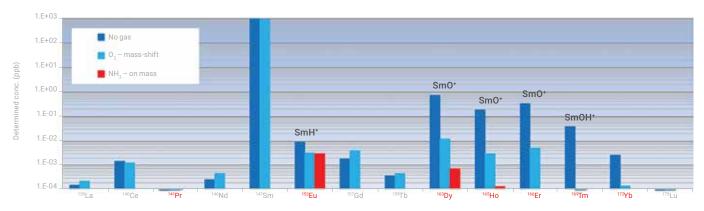


Figure 2. Measured concentration of REE impurities in 1 ppm Sm solution. Sm based interferences are observed on Eu, Dy, Ho, Er, Tm and Yb. Only the elements in red were measured in NH₃ on-mass mode.

For the REEs that react with NH_3 (La, Ce, Nd, Sm, Gd, Tb and Lu), O_2 mass-shift mode and measurement of the target analyte as its REE-O+ ion is the preferred approach. Most REEs are effectively converted to the oxide ion via reaction with O_2 cell gas [1], and this mode was applied to the measurement of Lu in the Gd matrix, avoiding the GdOH+ interference on the Lu+ isotope and giving a good improvement in the background signal. Compared to no gas mode, O_2 mass-shift mode also gave a good improvement in the background signals for Dy, Ho, Er, Tm and Yb in the Sm matrix, but for all these analytes the backgrounds in NH_3 mode were lower still.

Reference

 Direct measurement of trace rare earth elements (REEs) in high-purity REE oxide using the Agilent 8800 Triple Quadrupole ICP-MS with MS/MS mode, Agilent application note, 5991-0892EN.

The Benefits of Improved Abundance Sensitivity with MS/MS for Trace Elemental Analysis of High Purity Metals

Author

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Keywords

quadrupole, resolution, hyperbolic, abundance sensitivity, copper, high purity zinc

Introduction

The use of a quadrupole mass filter for the separation of compounds in mass spectrometry is well established. Initially used for organic mass spectrometry and residual gas analysis, the quadrupole spectrometer was adopted for the earliest ICP-MS systems, and has remained the default choice throughout the history of ICP-MS. However, the performance characteristics of the quadrupole mass filter do impose several limitations on quadrupole ICP-MS (ICP-QMS).

The resolution (R) of a mass filter (meaning its ability to separate adjacent masses) is defined as $M/\Delta M$, the mass of the target peak/the mass difference to nearest adjacent peak that can be distinguished (separated). However, for practical specifications, the resolution is often simply quoted as the width of the peak at a given peak height. The quadrupole mass filter used in an ICP-QMS instrument is typically operated with a nominal peak width of about 0.75 amu at 10% peak height, illustrated in Figure 1.

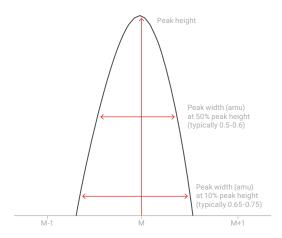


Figure 1. Illustration of resolution calculation for a mass spectrometer.

For two peaks within the normal signal range of the instrument, this allows the complete, baseline separation of masses 1 amu apart, within the elemental mass range from Li (7 amu) to U (238 amu) and beyond. Higher resolution of 0.4 amu peak width is possible by adjusting the quadrupole voltages, but the signal is reduced (less ion transmission) due to rejection of a higher proportion of the ions that are nominally "on-mass". Typically the signal loss at higher resolution is around 10-50%, depending on the design and operating characteristics of the quadrupole.

Both the efficiency of transmission of ions at the set-mass (i.e., the sensitivity) and the rejection of ions at other masses (i.e., the resolution of adjacent peaks) are affected by the shape of the field within the quadrupole, and the frequency of the alternating RF fields.

A hyperbolic field (generated by rods with a hyperbolic profile) alternating at high frequency gives more effective filtering of the ion beam than a lower frequency field generated by round quadrupole rods. The practical benefit of hyperbolic rods and high frequency RF voltage is therefore better ion transmission at higher resolution. Some of the many real-world applications where the combination of high sensitivity and good peak separation is required for adjacent low/high concentration elements measured by ICP-MS are shown in Table 1.

Low concentration	High signal	Example matrix
³¹ P	¹⁶ O ₂ , ³² S	Soil, plants, biological
⁵⁵ Mn	⁵⁶ Fe, ⁴⁰ Ar ¹⁶ O	Blood, iron and steel, soil
⁶³ Cu, ⁶⁵ Cu	⁶⁴ Zn, ⁶⁶ Zn	Metal refining
¹¹ B	¹² C	Soils, solvents, petrochem
¹³ C	¹⁴ N	Laser imaging of biological

Due to the ion transmission characteristics of a quadrupole, the peak that is generated from the ion signals at each mass forms a non-symmetric Gaussian distribution with a negative skew; i.e., the peak has a longer tail on the leading edge (low mass side) than the trailing edge (high mass side). These "tails" may extend significantly beyond the limits of the nominal 0.75 amu peak width, but since they are at intensities far below 10% of the peak height, they cannot be measured using the simple resolution figure quoted above. The contribution that a peak at mass M makes to its neighbors at M-1 amu and M+1 amu can be quantified, however, and this figure is referred to as the abundance sensitivity (AS) of the quadrupole, illustrated in Figure 2.

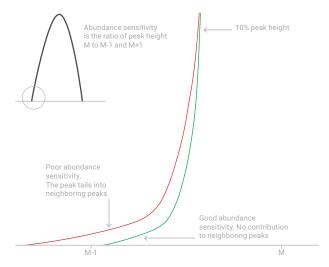


Figure 2. Illustration of abundance sensitivity calculation for a mass spectrometer.

For a good quadrupole mass spectrometer in ICP-QMS, the AS would typically be of the order of 10^{-7} , meaning that for an on-mass signal of 10^{7} counts, there is a contribution of one count at the adjacent mass (M+/-1 amu).

In applications where the trace analyte must be separated from a very intense matrix peak at the adjacent M+1 mass, such as the examples shown in Table 1, the matrix peak may be at an intensity greater than 10⁹ or 10¹⁰, and the AS of a quadrupole mass spectrometer is insufficient for accurate trace measurement of an adjacent overlapped analyte at low/sub ppb levels.

The problem of adjacent mass overlaps now has an elegant solution in the Agilent 8800 Triple Quadrupole ICP-MS (ICP-QQQ). The 8800 ICP-QQQ uses a tandem mass spectrometer configuration with two quadrupole mass filters (Q1 and Q2) separated by a collision/reaction cell. In MS/MS mode, both quadrupoles are operated as unit mass filters, so the overall AS of the instrument is the product of the Q1 AS x the Q2 AS. With two research-grade, high frequency, hyperbolic quadrupoles, each operating with AS of 10^{-7} , the combined AS of the 8800 ICP-QQQ is theoretically 10^{-14} , although this cannot be verified experimentally as the magnitude of the signal difference exceeds the dynamic range of the detector.

Experimental

Trace copper in high purity zinc

Major uses of Zn include galvanized coating to protect steel, die castings, and solder. Impurities in the metal cause Zn plating to lift, die casts to crack, or solder to 'de-wet', hence high purity zinc (>99.995 %) is a preferred commodity. Common impurities are Cu, Au and Sb, but may also include Cd, Al, Fe, Ag, Bi, As, In, Ni, P, and S.

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/Low matrix.

Ion lens tune: Auto tune was used for optimization.

CRC conditions: Helium cell gas at 4.8 mL/min with KED of 4 V.

Results and Discussion

Analysis was performed on high purity Zn, dissolved to give a 0.1% (1000 mg/L) Zn solution in a final acid concentration of 2% HNO $_3$. The sensitivity of the 8800 ICP-QQQ was reduced to bring the signal for the major Zn isotopes (64, 66) within the detector's upper limit of dynamic range ($\sim 10^{10}$ cps). The intense Zn signals were measured automatically in analog detector mode, while the Cu isotopes were measured in pulse mode. It can clearly be seen in Figure 3 that the intense Zn peaks at m/z 64 and 66 made no contribution to the signal at the two adjacent trace Cu isotopes at m/z 63 and 65. The Cu isotope ratio matched the theoretical abundances (63 Cu/ 65 Cu natural ratio of 69.17/30.83), at 1 µg/L concentration. If there was a contribution from an adjacent Zn mass then the isotope ratio would be biased.

From the 63 Cu calibration (Figure 4), the BEC and DL measured for Cu in the 0.1% Zn matrix were 1.7 ppb and 0.01 ppb respectively, indicating a low and stable background signal. The sensitivity of Cu was 7700 cps/ppb in 1000 μ g/L Zn, under the "de-tuned" conditions used to bring the Zn peaks within the detector range. This represents about a 2x reduction in the signal that would be obtained under normal tuning conditions for this type of matrix, if measurement of the matrix element peaks was not required.

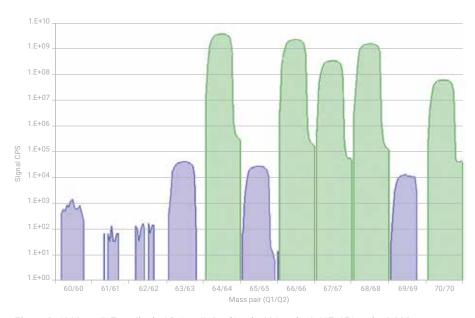


Figure 3. 1000 mg/L Zn spiked with 1 ug/L Cu. (1 ppb: 63/total = 0.667, 65/total = 0.333, no mass bias correction performed. If there was some contribution from Zn it would influence m/z 63 differently from m/z 65, because of the different abundance of the Zn isotopes).

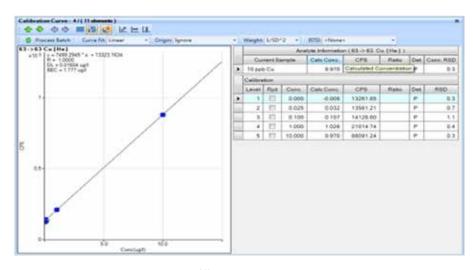


Figure 4. Standard addition calibration of 63 Cu in 1000 mg/L Zn

Ultra Trace Copper Analysis in a Semiconductor Grade Organometallic Titanium Complex

Authors

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Keywords

semiconductor, organometallic, copper, titanium, ammonia mass-shift

Introduction

Most quadrupole ICP-MS (ICP-QMS) instruments use CRC technology to resolve spectroscopic interferences. Helium collision mode is widely accepted due to its versatility and ease of use for multi-element analysis of complex and variable samples. While the performance achievable with He mode meets the requirements for most applications, there are some applications, for example impurity analysis of semiconductor materials, that require improved interference removal capability. For these applications, a reactive cell gas (reaction mode) may be used, but the use of highly reactive cell gases in quadrupole ICP-MS is prone to unexpected interferences and overlaps, especially when the matrix is complex, or other analytes are present at varying concentrations. The new Agilent 8800 Triple Quadrupole ICP-MS (ICP-QQQ) eliminates the variability associated with reactive cell gases in ICP-QMS, by using the first quadrupole (Q1) to control the ions that enter the CRC. This ensures that the reactions are predictable and the product ion spectrum is simple and consistent.

This report describes the measurement of trace Cu in a semiconductor grade organometallic Ti complex used in advanced semiconductor processing. It is a challenging application for quadrupole ICP-MS since both isotopes of copper, ⁶³Cu and ⁶⁵Cu, suffer interference from TiO and TiOH ions, and the use of reactive cell gases to avoid the overlap leads to a very complex product ion spectrum, particularly for organic samples. We demonstrate that the Agilent 8800 ICP-QQQ, operating in MS/MS mass-shift mode using ammonia as a reaction gas, was able to separate Cu⁺ from the Ti-based interferences and measure Cu at low ppt levels in a matrix of 500 ppm Ti. Results were also acquired using MS/MS He collision mode, for comparison.

Experimental

Instrumentation: Agilent 8800 #200 with narrow injector (id = 1.5 mm) torch (G3280-80080) used for organic solvent analysis. A low flow PFA nebulizer (G3285-80002) was used in self-aspiration mode. An option gas flow of 20% $\rm O_2$ balanced in Ar was added to the carrier gas via the standard option-gas line to prevent carbon build up on the interface cones.

Operating conditions: Table 1 summarizes plasma, ion lens and cell tuning conditions.

Acquisition conditions: MS/MS mode was used; cell gas was either NH_3 or He.

Sample and sample preparation: Semiconductor grade organometallic Ti complex (ADEKA Corp., Japan) was diluted with high purity IPA (Tokuyama Corp., Japan) to 500 ppm Ti solution. A spiked standard was prepared from the multielement standard, xstc-331, purchased from SPEX CertiPrep Ltd. (UK).

Table 1. Experimental conditions.

		Units	He collision cell mode	NH ₃ reaction cell mode
Cell conditions	Cell gas		Не	NH ₃ (10% NH ₃ in He)
	Cell gas flow rate	mL/min	8.0	6.5
	Octopole bias	٧	-18	-18
	KED	٧	4	-10
	Cell exit	٧	-100	-70
	Deflect	٧	-3	-12
	Plate	٧	-70	-60
Plasma conditions	RF	W	1600	
	SD	mm	12.0	
	Carrier gas	L/min	0.70	
	Make-up gas	L/min	0.20	
	Opt gas flow rate	L/min	0.20	
Ion lens	Extract 1	٧	-60	
	Extract 2	٧	-10	

Results and Discussion

He collision mode

The He cell gas flow rate was optimized for the lowest BEC of Cu in a 500 ppm Ti solution. As the BEC for 63 Cu was lower than the BEC for 65 Cu due to the higher abundance of the 63 isotope, and the more significant interference from TiO⁺ at m/z 65, Cu was determined on-mass at m/z 63. In MS/MS mode, this is achieved by the acquisition conditions: Q1 = 63; Q2 = 63 (63, 63).

Two solutions were analyzed: 500 ppm Ti solution and 500 ppm Ti + 1 ppb Cu spike. Figure 1 (left) shows the signal at m/z 63 obtained from the analysis of the two solutions, plotted as a function of He flow rate. The BEC calculated from these signals is also given in the figure. It shows that the lowest Cu BEC in He mode was 46 ppt, achieved at a flow rate of 8.0 mL/min He.

NH₃ reaction cell mode

Cu⁺ reacts efficiently with NH₃ to form NH₃ cluster ions with the general form Cu(NH₃)_n⁺. TiO⁺ does not follow the same reaction pathway as Cu⁺, so the Cu product ion can be measured free from Ti overlap. Based on a preliminary study, one of the intense product ions, Cu (NH₃)₂⁺, was selected to measure Cu separated from the original TiO⁺ interference. A mass pair of Q1 = 63, Q2 = 97 was used with NH₃ as the reaction gas. Figure 1 (right) shows the result. A BEC of 11 ppt for Cu in 500 ppm Ti solution was achieved in NH₃ mode (10% NH₃/He mixed gas), at a flow rate of 6.5 mL/min NH₃.

Conclusions

Table 2 summarizes the analytical performance achieved by the 8800 ICP-QQQ operating in MS/MS mode with He collision and NH $_{\rm 3}$ reaction gas. As can be seen, NH $_{\rm 3}$ reaction mode is more effective than He collision mode for the removal of the TiO+ interference on Cu. The BEC obtained for Cu in a Ti matrix by NH $_{\rm 3}$ reaction mode is four times lower than He mode, with seven times higher sensitivity.

Table 2. Summary of Cu measurement in Ti matrix.

	Flow rate (mL/min)	BEC (ppt) of Cu in 500 ppm Ti	Sensitivity (cps/ppb)
He collision mode	8.0	45.5	810
NH ₃ reaction mode	6.5	10.9	5900

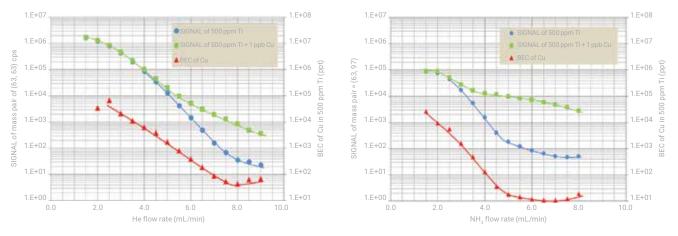


Figure 1. (Left) Cu signal (mass pair 63, 63) vs. He cell gas flow rate, for 500 ppm Ti matrix unspiked and with 1 ppb Cu spike, and calculated BEC. (Right) Cu signal (mass pair 63, 97) vs NH₃ cell gas flow rate, for 500 ppm Ti matrix unspiked and with 1 ppb Cu spike, and calculated BEC.

Removal of MH⁺ Interferences in Refined REE Material Analysis

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Keywords

Rare Earth Elements, REE, geochemistry, mining, material science, lanthanum, barium, cerium, method of standard additions, MSA, oxygen mass-shift

Introduction

The measurement of Rare Earth Elements (REEs) is of great importance in geochemistry, mining and material science. Manufacturers of high purity REE materials need to quantify metal impurities, including trace levels of the other REEs, in the refined, single element REE matrix. ICP-MS is the technique of choice for the measurement of REEs, but most of the REE isotopes suffer from interference by polyatomic species (predominantly hydride ions, MH+ and oxide ions, MO+) derived from other, lower-mass REE elements. While MH+ interferences are lower in intensity than MO+ interferences, they present a more challenging problem for REEs that have no isotope free from interference. For example ¹³⁹La+ is interfered by ¹³⁸BaH+ and ¹⁴⁰Ce+ by ¹³⁹LaH+. These interferences are too close in mass to be resolved by high-resolution (HR-)ICP-MS [1]. In this paper, we describe the removal of the MH+ interferences using an Agilent 8800 ICP-QQQ in MS/MS mass-shift mode, with oxygen as the reaction gas.

Experimental

Instrumentation: Agilent 8800 #100. The standard glass nebulizer was replaced with a C-flow nebulizer (G3285-80000) for optimal washout between the high matrix samples.

Plasma conditions: Preset plasma/General purpose.

Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -180 V. **CRC conditions:** O_2 gas at 0.3 mL/min, Octopole bias = -5 V, KED = -5 V. **Acquisition parameters:** MS/MS mode with O_2 mass-shift method.

Figure 1 illustrates the mechanism of MS/MS O_2 mass-shift mode used for measuring Ce in a La matrix sample. The major isotope of Ce at m/z 140 suffers an interference from 139 LaH $^+$. Q1 is set to m/z 140, allowing only the analyte ion 140 Ce $^+$ and any other ions at m/z 140 to pass through to the cell. All other ions not at m/z 140 are rejected. In the cell, Ce reacts with oxygen to form CeO $^+$ at m/z 156. Q2 is set to m/z 156, allowing CeO $^+$ to pass to the detector. Since 139 LaH $^+$ does not react with O_2 to form 139 LaOH $^+$, it remains as LaH $^+$ at m/z 140 and is rejected by Q2. The same principle is used for the separation of 139 La $^+$ from 138 BaH $^+$ in a Ba matrix.

Results and Discussion

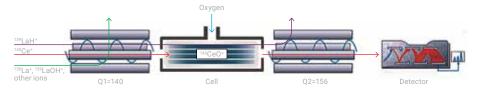


Figure 1. MS/MS mass-shift method with $\rm O_2$ reaction gas; used for the measurement of Ce, as CeO at m/z 156, in a La matrix.

Using the Method of Standard Addition (MSA), the BECs and DLs of La in a matrix of 50 ppm Ba, and Ce in a matrix of 50 ppm La were determined. Data was acquired using MS/MS mode with O_2 mass-shift, and also using Single Quad (SQ) mode with O_2 reaction gas to emulate conventional quadrupole ICP-MS (ICP-QMS) for comparison.

As shown in Figures 2A and 2B, SQ mode with $\rm O_2$ reaction gas suffers from interferences that prevent the measurement of La in the Ba matrix and Ce in the La matrix, respectively. In contrast, the calibration plots shown in Figures 2C and 2D demonstrate that MS/MS mode with $\rm O_2$ mass-shift can successfully remove the matrix overlaps to permit the trace quantitation of La in a Ba matrix and Ce in a La matrix. The BECs and DLs achieved were 8.5 ppt and 2.5 ppt respectively for La in a 50 ppm Ba solution, and 10.6 ppt and 0.8 ppt respectively for Ce in a 50 ppm La solution.

Investigation of unexpected product ion observed at m/z 156 in the 50 ppm

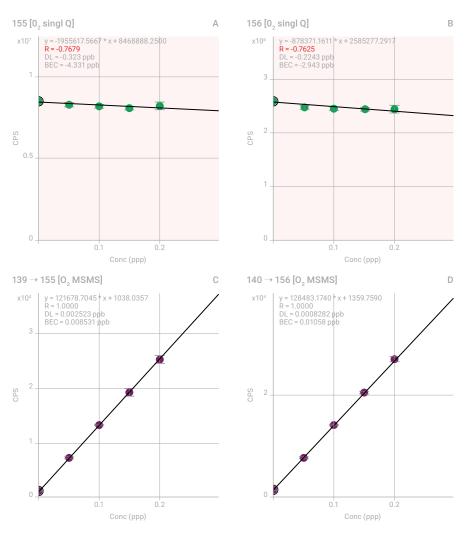


Figure 2. Top: Calibration plots up to 0.2 ppb for La in 50 ppm Ba matrix (A) and Ce in 50 ppm La matrix (B), acquired in SQ mode with oxygen reaction gas (emulating conventional quadrupole ICP-MS). Bottom: Calibration plots up to 0.2 ppb for La in 50 ppm Ba matrix (C) and Ce in 50 ppm La matrix (D) acquired in MS/MS mode with oxygen mass-shift.

La matrix

The background signals that contributed to the poor result obtained for Ce in the La matrix using SQ mode with $\rm O_2$ reaction gas (Figure 2B) were investigated by carrying out a precursor ion scan for product ion mass 156. The precursor ion scan capability of the 8800 ICP-QQQ provides a uniquely powerful approach to identifying the source of potential polyatomic and reaction product interferences. Oxygen cell gas was introduced into the cell and a precursor ion spectrum was obtained by scanning Q1 from 2 to 260 amu (Figure 3) with Q2 fixed at mass 156. From the spectrum, we can identify which precursor ions react with $\rm O_2$ to produce product ions at mass 156, overlapping $\rm ^{140}CeO^+$ in SQ mode.

Figure 3 shows the precursor ion scan spectrum for product ion mass 156 for the 50 ppm La matrix, with intense peaks at m/z 139 ($^{139}La^+$) and 156 ($^{139}La^{16}OH^+$). In SQ mode, as with conventional ICP-QMS, these ions all enter the cell, and with Q2 set to 156 amu, the $^{139}La^{16}OH^+$ polyatomic ions contribute to the signal measured at m/z 156 (^{140}Ce measured as analyte product ion $^{140}CeO^+$). These unwanted precursor ions cannot be rejected by a CRC operating as a bandpass filter in ICP-QMS, as they are too close in mass to the target analyte precursor ion. Only by using MS/MS mode on the 8800 ICP-QQQ, where Q1 operates as a unit mass filter, can non-target masses (like $^{139}La^{16}OH^+$ in this example) be prevented from entering the cell.

Reference

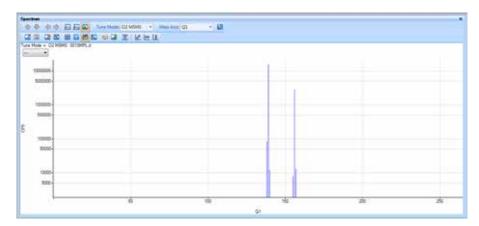


Figure 3. Precursor ion scan from 2-260 amu for product ion mass 156, in a 50 ppm La matrix. Six peaks are seen at m/z = 138, 139, 140, 155, 156 and 157, with the intense peaks at m/z 139 and m/z 156 being due to ¹³⁹La⁴ and ¹³⁹La⁶O¹H⁺ respectively.

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More Information

Removal of hydride ion interferences (MH+) on Rare Earth Elements using the Agilent 8800 Triple Quadrupole ICP-MS, Agilent publication, <u>5991-1481EN</u>.

Direct Analysis of Trace REEs in High Purity Nd₂O₃

Authors

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Keywords

Rare earth elements (REE), high purity metals, neodymium, neodymium (III) oxide, oxygen mass-shift, ammonia on-mass, ammonia mass-shift, geochemistry, mining, materials science

Introduction

Advanced technology products containing Rare Earth Elements (REEs) are increasing at a rapid rate. However, the presence of other REEs as contaminants in a purified single-element REE material may affect the functionality of the final product, so impurities in the REE oxide raw material must be carefully controlled.

The measurement of mid- and high-mass REEs in a low-mass REE matrix is challenging for ICP-MS because REEs have high metal-oxide (M-O) bond strengths, and the oxide ions of the low mass REEs overlap the preferred isotopes of the mid-mass and high-mass REEs. For example, in the analysis of trace REEs in high purity Nd_2O_3 , $^{145}Nd^{16}OH_2^+$ and $^{146}Nd^{16}OH^+$ overlap the preferred isotope of dysprosium ($^{163}Dy^+$), $^{143}Nd^{16}O^+$ overlaps the only isotope of terbium ($^{159}Tb^+$) and $^{148}Nd^{16}OH^+$ overlaps the sole isotope of holmium ($^{165}Ho^+$). While separation of the trace REEs from the REE matrix can be performed using a chelating resin, this technique is time-consuming and needs to be customized to the particular analyte and matrix under investigation. Clearly there is a requirement for a method capable of the direct analysis of trace REEs in a variety of high purity REE matrices.

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/HMI-L.

Acquisition parameters:

Five operational modes were evaluated:

- No gas
- Helium mode, 5 mL/min
- O₂ mass-shift, 0.3 mL/min
- NH₃ on-mass, 8 mL/min (NH₃ as 10% NH₃ in He)
- NH₃ mass-shift, 3 mL/min, (NH₃ as 10% NH₃ in He).

Sample and sample prep: High purity Nd_2O_3 (99.999%, purchased from the Baotou Research Institute of Rare Earths, China) was dissolved gently in semiconductor grade HNO_3 , and diluted to 500 ppm as Nd_2O_3 .

Results and Discussion

Thirteen trace REEs were measured in the $\mathrm{Nd}_2\mathrm{O}_3$ sample using the five different cell modes, and the results are summarized in Table 1. As expected, the BEC of low- and mid-mass REEs, such as La, Ce, Pr, Sm, Eu and Gd (Pr and Sm were present as impurities) were comparable in all modes, as these elements are free from interferences due to Nd. In contrast, the BECs for high-mass REEs in He mode were lower in He mode than in no gas mode, suggesting that high mass REEs suffered interferences from Nd-derived polyatomic ions.

Table 1. BECs of 13 REEs in 500 ppm Nd₂O₃. All units ug/kg (ppb).

Element	Isotope	No gas	He	O ₂ mass-shift	NH ₃ on-mass	NH ₃ mass-shift
La	139	0.143	0.127	0.143	-	-
Се	140	0.018	0.012	0.011	-	-
Pr	141	1.376	1.202	1.056	-	-
Sm	152	1.061	0.950	0.999	-	-
Eu	153	0.032	0.026	0.028	-	-
Gd	155	0.035	0.046	0.033	-	-
Tb	159	442.6	74.6	1.258	-	0.022
Dy	163	250.3	196	1.161	0.040	-
Но	165	20.43	16.2	0.101	0.004	-
Er	170	0.065	0.020	0.013	-	-
Tm	169	0.084	0.031	0.003	-	-
Yb	174	0.251	0.120	0.058	-	-
Lu	175	0.014	0.006	0.004	-	-

0, mass-shift mode

All 13 REEs react with O_2 efficiently to form REE-oxide ions, as shown below.

The MS/MS capability of the 8800 ICP-QQQ enables the removal of spectral interferences on each element using "mass-shift". For example in $\rm O_2$ mass-shift mode, all 13 REEs can be detected as REE-O+ ions at 16 amu higher than the original elemental mass (M+16). From Table 1, it can be seen that $\rm O_2$ reaction mode with mass-shift further reduced the BEC for Tb, Dy, Ho, Er, Tm, Yb, and Lu, compared to He mode.

While the improvement in O_2 mass-shift mode is significant for Tb, Dy and Ho that suffer intense interference from NdO+, the BECs of the other high-mass REEs such as Er, Tm, Yb and Lu were also improved in this mode, indicating that these elements also suffer interferences from Nd-based polyatomic ions: 150 NdOH $_3^+$ interferes with 169 Tm+, 142 NdO $_2^+$ (or 142 NdCO+) and 144 NdCN+ with 170 Er+, 142 NdO $_2^+$ with 174 Yb+, 143 NdO $_2^+$ and 144 NdONH+ (or 144 NdC $_2$ H+) with 175 Lu+. The contribution of the above mentioned interferences on Er, Tm, Yb and Lu are not overly significant. However, O_2 mass-shift mode was shown to be an effective approach for the removal of all polyatomic ion interferences, typically leading to a 5-10x lower BEC compared to no gas mode.

NH₃ on-mass mode for Dy and Ho

A previous study showed that NH_3 cell gas reacts with many of the polyatomic ions that interfere with the REEs. However, NH_3 also reacts quickly with some of the REE ions, leading to reduced sensitivity of < 1 cps/ppt for La, Ce, Nd, Sm, Gd, Tb and Lu. NH_3 on-mass mode is valuable for the determination of a limited number of REEs; Pr, Eu, Dy, Ho, Er, Tm and Yb [1]. The results in Table 1 show that NH_3 on-mass mode gave excellent results for Dy and Ho in the Nd_2O_3 matrix, with an improvement in BECs of 20x compared to O_2 mass-shift mode.

NH, mass-shift mode for Tb

For the REEs that react efficiently with NH $_3$ (La, Ce, Nd, Sm, Gd, Tb and Lu), NH $_3$ mass-shift mode can be used. In this study, NH $_3$ mass-shift mode was investigated for the determination of Tb, and the reaction product ion TbNH $^+$ (m/z 174) was found to give the lowest BEC. A BEC of 22 ppt for Tb in a 500 ppm Nd $_2$ O $_3$ solution was achieved, which is 50x lower than the result achieved in O $_2$ mass-shift mode, indicating the effective removal of the NdO $^+$ overlap.

Conclusions

The Agilent 8800 ICP-QQQ with MS/MS capability was used to successfully measure 13 REE impurities in a high-purity $\mathrm{Nd_2O_3}$ sample solution. Tandem MS with MS/MS mode is essential for accurate reaction mode analysis in a complex matrix. On conventional quadrupole ICP-MS, there is no additional quadrupole (Q1) to select which ions can enter the cell. As a result, all ions enter the cell so, when a reactive cell gas is used, a complex and variable population of reaction product ions is created, depending on the sample matrix and other analytes. With ICP-QQQ, the first quadrupole selects only the target mass to pass into the cell, so the reaction chemistry is controlled and consistent. With the combination of HMI and MS/MS reaction cell mode, the 8800 ICP-QQQ provided effective removal of the polyatomic interferences from the Nd matrix.

Reference

 Naoki Sugiyama and Glenn Woods, Direct measurement of trace rare earth elements (REEs) in high-purity REE oxide using the Agilent 8800 Triple Quadrupole ICP-MS with MS/MS mode, Agilent publication, 2012, 5991-0892EN.

More Information

Application note: Routine determination of trace rare earth elements in high purity Nd_2O_3 using the Agilent 8800 ICP-QQQ. Agilent publication <u>5991-5400EN</u>.

Determination of Trace Level Impurities of 49 Elements in High Purity Copper

Author

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Keywords

high purity metal, high purity copper, material science, mass-shift, m-lens, alkali elements

Introduction

Metals such as Cu, Al, Ta, W, and Hf are widely used in semiconductor devices. High purity metals are required to ensure reliable performance and a high production-yield. ICP-MS is often used for the quality-control of these metals. The application is not easy though due to the requirement for ultralow level impurity measurements in relatively high-matrix metal sample digests. The ultra-trace measurement of alkali elements in the presence of the high matrix is especially challenging. Cool plasma is accepted in the semiconductor industry as a reliable technique to remove argon-based interferences such as Ar+ and ArO+. The method enables the low-level analysis of Ca and Fe by ICP-MS. It can also be applied to the analysis of the alkali elements, providing lower background equivalent concentrations (BECs) than hot plasma conditions. However, the cooler, lower energy plasma has a poorer matrix tolerance, making it unsuitable for the analysis of high-matrix metal sample digests.

Experimental

Instrumentation: Agilent 8900 Semiconductor configuration ICP-QQQ fitted with an optional m-lens.

Tuning and method: Tuning parameters are summarized in Table 1.

- Internal standards (ISTD) Be, Sc and In were added to all solutions before analysis.
- Hot plasma conditions were optimized to give a 1% CeO⁺/Ce⁺ ratio.
- A single cell gas mode, $O_2 = 0.2$ (mL/min) + $H_2 = 7.0$ (mL/min), was used for all 49 analytes + 3 ISTD elements.
- Mass-shift mode was used for the analysis of P, S, As, Zr, Nb, Mo, Hf, Ta, W, Th, and U. On-mass mode was used for all other elements.

Sample preparation

All samples and standards were prepared in 5% nitric acid (HNO_3) using TAMAPURE AA-100 semiconductor grade HNO_3 (TAMA Chemicals Co. Ltd, Kanagawa, Japan).

A 0.1% Cu sample was prepared as follows. A piece of 9N high purity copper was cleaned in diluted HNO_3 then rinsed with ultrapure water (UPW). It was weighed (about 0.05 g) and dissolved in 5 mL of 50% HNO_3 (UPW: HNO_3 = 50:50). The solution was then brought up to 50 mL with UPW.

Table 1. Cool plasma operating conditions.

Parameter	Unit	Value	
RF power	W	1550	
Carrier gas flow rate	L/min	0.70	
Make-up gas flow rate	L/min	0.46	
Sampling depth	mm	8.0	
Extract 1	V	0.0	
Extract 2	V	-70.0	
Omega	V	8.0	
Omega bias	V	-60.0	
Cell gas and flow rate	mL/min	O ₂ = 0.2 and H ₂ = 7.0	
Octopole bias	V	-10.0	
KED	V	-10.0	
Axial Acceleration	V	2.0	

Results and Discussion

BECs in 5% HNO₃

Background equivalent concentrations (BECs) for each element in 5% HNO $_3$ were obtained from the calibration plots. The results are summarized in Figure 1. The plot shows that parts per trillion (ppt) level BECs were achieved for the alkali elements, Li, Na, and K, using hot plasma conditions. In addition, impressive BECs were achieved for the most challenging elements: S (84 ppt) and Si (231 ppt).

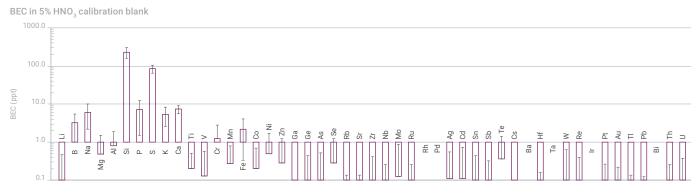


Figure 1. BECs of 49 elements in 5% HNO_3 blank. Error bar = 3σ DL.

Measured concentrations in 0.1% high purity copper

The method was used to determine the concentration of 49 elements in the 0.1 % high purity copper solution. The results are summarized in Figure 2. The observed matrix suppression for low mass elements like Li and B was 30%, and 10% or less for mid and high mass elements. Except for Si, S and Te, all elements were present at < 10 ppt concentration level. The $\rm O_2+H_2$ reaction cell gas removed significant spectral interferences caused by ArCu+ on Ru+, Rh+ and Pd+, allowing the determination of these elements at ultralow levels.



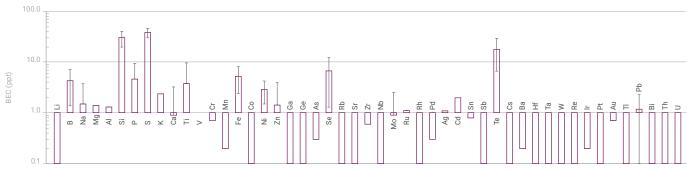


Figure 2. Measured concentrations of 49 elements in 0.1% high purity copper.

Conclusions

Using a single cell gas with MS/MS on-mass and mass-shift modes of the ICP-QQQ and hot plasma conditions, ppt level BECs were achieved for the alkali elements. Low-level BECs were also obtained for challenging elements, sulfur and silicon. Overall, 49 elements were determined at ultralow levels in 0.1% high purity Cu sample using a simple, single tune method.

More Information

Application note: Analysis of Ultratrace Impurities in High Purity Copper using the Agilent 8900 ICP-QQQ: Low-ppt determination of alkali metals in high matrix samples using the optional "m-lens", Agilent publication 5994-0383EN.

Direct Determination of Challenging Trace Rare Earth Elements in High Purity Lanthanide REE Oxides

Authors

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Keywords

Rare Earth Elements, REE, rare earth oxides, REO, oxygen mass-shift, ammonia on-mass, ammonia mass-shift

Introduction

ICP-MS is widely used for trace impurity analysis of high purity rare earth element (REE) oxide materials. But the analysis of trace REEs in high purity REE oxide materials remains challenging. Matrix-based polyatomic ions such as REEO⁺, REEOH⁺, and REEH⁺ cause severe spectral interferences on some REE elements. Trace REE analytes can be separated from the REE matrix using a chelating resin, but this technique is time-consuming, and customization is needed per the analyte and matrix element.

In this study, trace REEs in lanthanide oxide materials were determined using an Agilent 8900 ICP-QQQ with $\rm O_2$ and $\rm NH_3$ reaction cell gases. Since the analysis of $\rm La_2O_3$, $\rm Tm_2O_3$ and $\rm Lu_2O_3$ is relatively interference free, these matrices weren't included in the study.

Experimental

Instrumentation: An Agilent 8900 Advanced Applications configuration ICP-QQQ was used without any modification. For the analysis of 500 ppm REE matrix samples, 'general-purpose plasma' conditions were selected in the MassHunter software. The preset plasma function automatically sets all plasma-related parameters, simplifying instrument set-up.

Five cell modes were investigated: no gas, helium (He), oxygen (${\rm O_2}$), and ammonia (20% NH $_3$ in He). Tuning conditions are summarized in Table 1. In NH $_3$ mass shift mode, a pre-study was done using 'product ion scan' to identify the most abundant NH $_3$ cluster ion. The masses of the cluster ions used for the analysis are given in Tables 2 and 3, together with the analytical results.

Table 1. Cell gas mode-related tuning parameters.

Cell gas mode	No gas	Не	O ₂ mass-shift	NH ₃ on-mass	NH ₃ mass-shift
Scan Mode	Sing	jle Quad		MS/MS	
Octopole bias (V)	-8	-18	-3	-5	-5
Octopole RF (V)	140	180	180	180	180
KED (V)	+5	+4	-7	-7	-7
Axial Acceleration (V)	0	1	1.5	0.5	0.5
He (mL/min)		5		1	1
O ₂ (mL/min)			0.45		
NH ₃ (mL/min)				4.0 ~ 6.0	1.0 ~ 8.0

Results and Discussion

Ten REE oxide materials of the highest-grade purity (5N) including Ce_2O_3 , Pr_6O_{11} , Nd_2O_3 , Gd_2O_3 , Sm_2O_3 , Eu_2O_3 , Tb_4O_7 , Dy_2O_3 , Er_2O_3 , and Yb_2O_3 were dissolved in semiconductor grade HNO_3 and diluted to 500 ppm (as REE oxide). H_2O_2 was added during the dissolution of Ce_2O_3 and Tb_4O_7 . REEs were measured in each matrix solution using the five cell modes specified in Table 1. The results are given in Tables 2 and 3.

As expected, in no gas mode, the BECs for the REEs were relatively high due to spectral interferences. He collision cell mode was able to alleviate some of the interferences, but not all. Previous studies have shown that both $\rm O_2$ and $\rm NH_3$ are effective for the removal of polyatomic ions that interfere with the REEs [1, 2]. A drawback of $\rm NH_3$ mode has been low sensitivity. However, Axial Acceleration of cluster ions in the cell of the 8900 ICP-QQQ increases sensitivity. The results reported in Tables 2 and 3 show that the BECs for all REEs were dramatically improved using a reactive cell gas. The improvement factor data relates to the difference in BEC obtained in reaction mode compared to no gas mode.

Table 2. BECs of REE impurities in 500 ppm Ce, Pr, Nd, and Gd oxide solutions.

Sample				Ce ₂ O ₃		PrO		Nd_2O_3			$\mathrm{Gd_2O_3}$	
Analyte			Pr	Gd	Tb	Tb	Tb	Dy	Но	Tb	Yb	Lu
Isotope			141	160	159	159	159	163	165	159	172	175
Interference			¹⁴⁰ CeH ⁺	¹⁴² Ce ¹⁸ O+	¹⁴² Ce ¹⁶ OH ⁺	¹⁴¹ Pr ¹⁸ O+	¹⁴² NdOH+, ¹⁴³ NdO+	¹⁴⁵ Nd ¹⁸ O ⁺	¹⁴⁸ NdOH ⁺	158GdH+	¹⁵⁶ GdO+	¹⁵⁹ GdOH ⁺
BEC (ppb)	No gas		6.17	3.36	29.2	10.3	721	163	13.4	2.23	3420	75.0
	He		3.79	11.9	0.725	2.50	234	36.6	3.06	2.16	1200	66.4
	0,		0.064	0.030	9.76	0.001	1.95	0.804	0.070	0.106	284	0.444
	NH ₃	BEC			0.284	0.055	0.039	0.255	0.021		0.030	7.16
		mass pair			(159/174)	(159/244)	(159/174)	(163/163)	(165/165)		(172/172)	(175/260)
Improvement	t factor		x100	x100	x100	x10,000	x20,000	x1000	x1000	x20	x100,000	x200

Table 3. BECs of REE impurities in 500 ppm Sm, Eu, Tb, Dy, Er, and Yb oxide solutions.

Sample			Sm ₂ O ₃				Eu ₂ O ₃	Tb ₄ O ₇	Dy ₂ O ₃	Er ₂ O ₃	Yb ₂ O ₃
Analyte			Dy	Но	Er	Tm	Tm	Lu	Но	Tm	Lu
Isotope			162	165	167	169	169	175	165	169	175
Interference			¹⁴⁷ SmO ⁺	¹⁴⁸ SmOH ⁺ , ¹⁴⁹ SmO ⁺	150SmOH+	¹⁵² SmOH ⁺	¹⁴¹ EuO+	¹⁵⁹ TbO ⁺	¹⁶⁴ DyH ⁺	¹⁶⁸ ErH ⁺	¹⁷⁴ YbH ⁺
BEC (ppb)	No gas		0.408	185	44.9	39.0	64.8	3270	2.13	1.26	0.97
	Не		0.169	61.9	18.1	13.6	38.20	1670	1.28	1.57	1.38
	0,		0.083	0.158	0.916	0.240	2.73	26.1	0.057	0.025	0.195
	NH ₃	BEC	0.035	0.055	0.092	0.127	0.002	0.244	0.074		
		mass pair	(162/162)	(165/165)	(167/167)	(169/169)	(169/169)	(175/260)	(165/165)		
Improvement	factor		x10	x3000	x500	x200	x30,000	x10,000	x50	x50	x5

Conclusions

The Agilent 8900 ICP-QQQ method was used to measure REE impurities in high purity REE oxide materials. REE matrix-based hydride, oxide, and hydroxide polyatomic ion interferences were removed by operating the ICP-QQQ in MS/MS mode with $\rm O_2$ and $\rm NH_3$ reaction cell gases. The BECs were improved by one to four orders of magnitude using reactive cell gases compared to no gas mode. The method is suitable for the direct analysis of trace REEs in the presence of high concentration matrix-REEs.

References

- Direct measurement of trace rare earth elements (REEs) in high-purity REE oxide using the Agilent 8800 Triple Quadrupole ICP-MS with MS/MS mode, Agilent application note, 2012, 5991-0892EN
- 2. Routine determination of trace rare earth elements in high purity Nd_2O_3 using the Agilent 8800 ICP-QQQ, Agilent application note, 2015, 5991-5400EN

Analysis of 10 nm Gold Nanoparticles using the High Sensitivity of the Agilent 8900 ICP-QQQ

Authors

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Keywords

nanoparticles, single nanoparticle analysis, TRA, dwell time, gold nanoparticles

Introduction

The measurement of nanoparticles (NPs) is of public and scientific interest. More information is needed to understand the fate of NPs in the environment and the potential toxic effects once absorbed into the body. Gold (Au) NPs have a broad range of uses in medical, industrial, and technology applications. Au is a relatively easy element to measure by ICP-MS, as it is not affected by common spectral interferences. However, the detection of very small particles (<20 nm) remains challenging for ICP-MS, due to the low signal generated from such particles. The Agilent 8900 ICP-QQQ has a low background (<0.2 cps) and sensitivity up to Gcps/ppm, making it suited to small particle detection.

Experimental

Instrumentation: Agilent 8900 Advanced Applications configuration ICP-QQQ with 1-mm i.d. injector torch and standard sample introduction system.

Method: All aspects of method setup and data analysis were carried out using the fully integrated Single Nanoparticle Application Module option for ICP-MS MassHunter. The "Batch at a Glance" data table shown in Figure 1 summarizes the sample results for an entire batch. Detailed graphical results are displayed for each selected sample, allowing results to be viewed and compared, or method settings to be optimized if necessary. Reference [1] gives details of the particle size calculation used in the module.

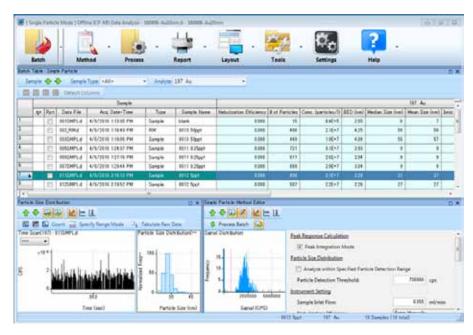


Figure 1. Data analysis view of the Single Nanoparticle Application software module.

Tuning conditions: For highest sensitivity, ¹⁹⁷Au was measured in single quad mode with no cell gas.

Plasma parameters: RF power =1550 W, sampling depth = 7.0 mm, and carrier gas flow rate = 0.78 L/min.

Data acquisition: TRA analysis with a dwell time of 0.1 ms. Data acquisition time was 60 s.

Reference materials and sample preparation:

Three Au NP reference materials (RMs) were used in the study: NIST 8011 with a nominal diameter of 10 nm (8.9 \pm 0.1 nm determined by Transmission Electron Microscopy (TEM)); NIST 8012 with a nominal diameter of 30 nm (27.6 \pm 2.1 nm determined by TEM); and NIST 8013 with a nominal diameter of 60 nm (56.0 \pm 0.5 nm determined by TEM). All final solutions containing Au nanoparticles or ionic Au standards were prepared in 0.01% L-cysteine for stabilization.

Results and Discussion

Analysis of Au NP samples

Solutions containing gold NPs with particle sizes of 10 nm, 30 nm, and 60 nm were prepared at concentrations of 0.25 ng/L, 5 ng/L and 50 ng/L, respectively. The solutions were measured using fast TRA acquisition. Figure 2 shows the measured raw signal event frequency and the calculated size distribution for a solution containing 10 nm particles. From the results, the practical detection limit of the particle diameter was estimated to be around 30,000 cps (equivalent to \sim 6.5 nm) and the background equivalent diameter (BED) was 3 nm.

The 30 nm and 60 nm particles were also measured, and the results are summarized in Table 1. The results for the median, mode, and mean particle sizes for all three standards agreed well with the reference sizes obtained by TEM.

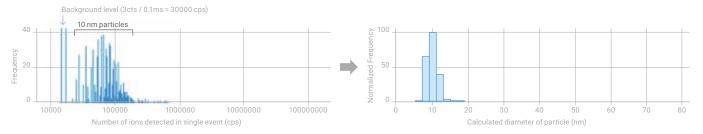


Figure 2. Raw signal event frequency (left) and calculated size distribution of 10 nm particles (right).

Table 1. Measured particle size for Au NPs in three NIST RMs.

Nominal size (nm)	, , , , , , , , , , , , , , , , , , , ,							
			Median		Mode		Mean	
			size (nm)	RSD (%)	size (nm)	RSD (%)	size (nm)	RSD (%)
10	8.9	± 0.1	9.0	3.3	10	0.0	9.2	3.3
30	27.6	± 2.1	26.9	0.3	28	0.0	27.0	0.3
60	56.0	± 0.5	56.1	0.3	56	1.8	57.2	0.4

Conclusions

The low background and high sensitivity of the Agilent 8900 ICP-QQQ make it suitable for single particle analysis of solutions containing the smallest-sized nanoparticles. The size and composition of gold NP solutions were characterized from 10 nm up to 60 nm, with good accuracy. The practical detection limit of the particle diameter was estimated to be 6.5 nm and the BED was 3 nm.

Reference

1. H. E. Pace, N. J. Rogers, C. Jarolimek, V.A. Coleman, C.P. Higgins, and J. F. Ranville, *Anal. Chem.*, **2011**, 83, 9361-936

More Information

Analysis of 10 nm gold nanoparticles using the high sensitivity of the Agilent 8900 ICP-QQQ, Agilent publication, <u>5991-6944EN</u>.

High Sensitivity Analysis of SiO₂ Nanoparticles using the Agilent 8900 ICP-QQQ

Authors

Michiko Yamanaka, Takayuki Itagaki, and Steve Wilbur, Agilent Technologies

Keywords

nanoparticles, single nanoparticle analysis, TRA, dwell time, silicon dioxide NPs

Introduction

ICP-MS is a well-established technique for measuring the elemental content of materials. With the recent development of Single Particle ICP-MS (spICP-MS) acquisition mode, ICP-MS can also be used to characterize the nanoparticle (NP) content of a sample.

Silicon dioxide (SiO_2) NPs are used for many applications including paints, coatings, adhesives, food additives, polishing micro-electronic devices etc. Given their wide spread use, there is a clear requirement for SiO_2 NPs to be monitored. Si measurement by ICP-MS is not easy since the major isotope of Si, ^{28}Si (92.23% abundance), is interfered by the background polyatomic ions CO^+ and N_2^+ . The interferences can be addressed using reaction chemistry in the collision/reaction cell of an ICP-MS. However, for controlled and consistent reaction processes, a tandem mass spectrometer instrument such as the Agilent 8900 Triple Quadrupole ICP-MS (ICP-QQQ) is required.

Experimental

Instrumentation: Agilent 8900 Advanced Applications configuration ICP-QQQ fitted with a 1 mm i.d. injector torch and standard sample introduction system.

Method: All aspects of the method setup and data analysis were carried out using the fully integrated Single Nanoparticle Application Module option of the ICP-MS MassHunter software. The "Batch at a Glance" data table summarizes the sample results for an entire batch. The detailed graphical results are displayed for each selected sample, allowing results to be viewed and compared, or method settings to be optimized if necessary. References [1 and 2] provide details of the particle size calculation used in the module.

Tuning conditions: H_2 on-mass mode was used to remove potential interferences by CO⁺ and N_2 ⁺ on ²⁸Si⁺. H_2 cell gas flow = 3 mL/min.

Plasma parameters: RF power =1550 W, sampling depth of 7.0 mm and carrier gas flow rate of 0.76 L/min.

Data acquisition: TRA analysis with dwell time of 0.1 ms.

Reference materials and sample preparation: SiO_2 NP reference materials (RMs) with nominal diameters of 50 nm, 60 nm, 100 nm, and 200 nm were bought from nanoComposix (San Diego, USA). They were diluted to a particle concentration of between 40 and 1000 ng/L with de-ionized (DI) water, and sonicated for 5 min to ensure sample homogeneity. A Si ionic standard of 5 μ g/L was prepared with DI water and used to measure the elemental response factor.

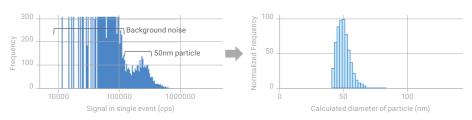


Figure 1. Raw signal event frequency (left) and calculated size distribution of 50 nm particles (right).

Results and Discussion

Analysis of SiO, NPs in UPW

The frequency distribution plots of the signals obtained from 50 nm $\mathrm{SiO_2}$ NPs are shown in Figure 1. The particle signals were clearly distinguished from the background (dissolved, ionic component). From the results, we can estimate that the practical detection limit for the particle diameter was below 50 nm and the background equivalent diameter (BED) was 22 nm.

The results for the size analysis of different ${
m SiO_2}$ NP solutions are summarized in Table 1. The results for median, mode, and mean particle sizes agreed well with the reference sizes obtained by TEM.

Table 1. Measured particle size for SiO, NPs in four RMs.

Nominal size (nm)	Particle size (n by TEM	m) Prepared particle concentration (ng/L)	Measured p	oarticle size (r	n = 10)
			Median	Mode	Mean
			size (nm)	size (nm)	size (nm)
50	46.3 ± 3.1	40	49	50	50
60	57.8 ± 3.5	40	61	62	62
100	97.0 ± 4.8	100	99	100	102
200	198.5 ± 10	1000	200	204	200

Analysis of SiO₂ NPs in a high-level carbon matrix

Real samples such as biological samples, food matrices, pharmaceutical ingredients, and organic solvents contain carbon matrices that give rise to a $^{12}\mathrm{C^{16}O^{+}}$ polyatomic ion interference on $^{28}\mathrm{Si^{+}}$. Figure 2 shows the particle size distribution for a mixed solution of the 100 and 200 nm $\mathrm{SiO_{2}}$ measured in a sample containing 1% ethanol. Despite the high concentration of carbon, the size distribution for each group of particle sizes was consistent with the results obtained by TEM. The 8900 ICP-QQQ in MS/MS mode with hydrogen cell gas was able to eliminate the $^{12}\mathrm{C^{16}O^{+}}$ interference effectively.

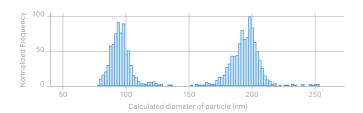


Figure 2. Size distribution result of 100 and 200 nm SiO_2 NPs in 1% ethanol.

Conclusions

 ${
m SiO}_2$ NPs can be determined and characterized successfully using the Agilent 8900 ICP-QQQ operating in MS/MS mode with H $_2$ cell gas. Even in the presence of a high level of carbon matrix. The Single Nanoparticle Application Module for ICP-MS MassHunter was used to calculate the particle sizes. The spICP-QQQ method provides fast analysis times, excellent detection limits for particle size and concentration, and accurate results for ${
m SiO}_2$ particles less than 100 nm.

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More Information

High sensitivity analysis of SiO_2 nanoparticles using the Agilent 8900 ICP-QQQ in MS/MS mode, Agilent publication <u>5991-6596EN</u>.

Analysis of Ultratrace Impurities in High Purity Copper using the Agilent 8900 ICP-QQQ

Author

Naoki Sugiyama Agilent Technologies, Japan Low-ppt determination of alkali metals in high matrix samples using the optional "m-lens"

Introduction

Metals such as copper (Cu), aluminum (Al), tantalum (Ta), tungsten (W), and hafnium (Hf) are essential for the manufacture of semiconductor devices. Metal sputtering targets are used to form conducting or insulating (dielectric) layers by thin film deposition using chemical vapor deposition (CVD) or physical vapor deposition (PVD). Conducting metals, originally Al but now typically Cu, are used as interconnects within wiring levels and as "vias" between layers. A complex, large-scale integrated circuit (IC) microprocessor chip may contain tens of layers of interconnect "wires" with a total length up to about 100 km (1, 2). To ensure high performance and high production-yield of the final devices, very high purity metals are required for these components.

Semiconductor manufacturers may require high-purity, electronic-grade metals at grades of 5N (5 9s – 99.999% purity) up to 9N (99.999999% purity) or above, depending on the proposed application. A 6N metal (99.9999% purity) contains a total of only 1 mg/kg (ppm) of the impurities of interest, so each individual impurity element would typically be certified as <0.01 or <0.005 ppm in the solid metal.

Determination of trace contaminants in high-purity metals is often performed using glow discharge mass spectrometry (GD-MS). However, GD-MS is expensive and requires the availability of solid metal calibration standards containing the trace elements of interest. GD-MS also has relatively slow speed of data acquisition leading to low sample throughput—around 10 minutes or more per sample—often longer when a cryo-cooled source is used. The fact that solid samples are analyzed also makes automation of sample changeover for unattended analysis more problematic than for liquid sample digests.

ICP-MS is widely used for quality-control of semiconductor materials, but some elements are difficult to measure at ultratrace levels in the presence of a high matrix. ICP-MS operating with a "cool" or reduced-energy plasma has been widely employed since the 1990s as a powerful mode for the analysis of high-purity chemicals and materials. Cool plasma suppresses the formation of intense argon-based interferences such as Ar+ and ArO+, allowing low-level analysis of 40Ca and 56Fe, respectively. Cool plasma conditions are also beneficial in the analysis of the alkali metal elements, providing lower background equivalent concentrations (BECs) than hot plasma conditions. A lower-temperature plasma causes less re-ionization of traces of easily-ionized elements (EIEs) from the cones and ion lens, giving lower background signals for these elements. Cool plasma is not universally applicable, though, as the lower power plasma has less energy, which reduces its ability to decompose the sample matrix. Poor tolerance of high matrix levels is especially problematic for the analysis of high matrix, high-purity samples such as electronic-grade metals.

This note describes a new approach to the measurement of ultratrace impurities in high-purity copper using triple quadrupole ICP-MS (ICP-QQQ). An optional ion lens (called the "m-lens") has been developed for the Agilent 8900 ICP-QQQ to allow ultra-low-level measurement of alkali metals under matrix tolerant, high-power plasma conditions. The m-lens has an optimized geometry that minimizes background signals from EIEs deposited on the ICP-MS interface components. matrix samples using the optional "m-lens".

Experimental

Sample preparation

All samples and standards were prepared in 5% semiconductor grade TAMAPURE AA-100 nitric acid (HNO_3) bought from Tama Chemicals Co. Ltd, Kanagawa, Japan. Solutions were prepared and analyzed in PFA vials, which were cleaned with diluted HCl and HNO_3 and then rinsed using ultrapure water (UPW) before use.

A 0.1% copper (Cu) solution was prepared for analysis. A sample of 9N high purity copper was cleaned in diluted ${\rm HNO_3}$, rinsed with UPW, weighed (about 0.05 g), and dissolved in 5 mL of 50% ${\rm HNO_3}$ (1:1 concentrated ${\rm HNO_3}$:UPW). The solution was brought up to volume (50 mL) with UPW, giving a total dilution of 1000x, and a matrix level of 0.1%. The 8900 ICP-QQQ can tolerate % levels of dissolved solids, but higher dilutions allow non-matrix-matched calibrations to be used. This removes the need for certified metal standards containing every element of interest. The exceptionally low detection limits of the 8900 ICP-QQQ (sub-ppt for most elements) enable ultratrace analysis even in higher sample dilutions.

The 1000x dilution factor simplifies conversion of the measured concentrations in ng/L (ppt) in the digest solution to the concentrations in μ g/kg (ppb) in the original solid.

Calibration standards for 49 elements were prepared from several mixed, multi-element stock standards (SPEX CertiPrep, NJ, USA). To minimize signal suppression due to physical sample transport and nebulization effects, the calibration standards were matrix matched to the ${\rm HNO_3}$ concentration (5%) of the Cu sample digest.

All samples and standards were spiked with a mix of three internal standard (ISTD) elements, Be, Sc, and In, at 5.0, 0.5, and 0.5 ppb, respectively. ISTDs were added to compensate for matrix differences between the standards (no Cu) and the 0.1% Cu solutions, and to correct for any long-term signal drift.

Instrumentation

An Agilent 8900 Semiconductor configuration ICP-QQQ was used for all measurements. The standard PFA nebulizer was used in self-aspiration mode, connected to the standard quartz spray chamber and quartz torch with 2.5 mm i.d. injector. The 8900 ICP-QQQ was fitted with the standard Pt-tipped sampling cone, optional m-lens (part number G3666-67500), and optional Pt-tipped, Ni-based skimmer cone for m-lens (part number G3666-67501). The skimmer cone for m-lens also requires a non-standard skimmer cone base (part number G3666-60401).

Tuning and method

Hot plasma conditions (1% CeO^+/Ce^+) were used to ensure good tolerance of the high concentration of Cu matrix. A single collision/reaction cell (CRC) tuning mode was used to measure all 49 analyte elements in the Cu samples. A cell gas mixture of oxygen (O_2) and hydrogen (O_2) was used to remove interferences using a combination of MS/MS on-mass and mass-shift modes. Operating conditions are summarized in Table 1, and acquisition parameters and given in Table 2.

Table 1. Agilent 8900 ICP-QQQ operating parameters.

Parameter	Setting
RF power (W)	1550
Sampling depth (mm)	8.0
Carrier gas flow rate (L/min)	0.70
Make-up gas flow rate (L/min)	0.46
Extract 1 (V)	0.0
Extract 2 (V)	-70
Omega bias (V)	-60
Omega lens (V)	8.0
Cell gas flow rate (mL/min)	O ₂ = 0.2; H ₂ = 7.0
Octopole bias (V)	-10
KED (V)	-10
Axial acceleration (V)	+2.0

 Table 2. Acquisition parameters.

Element	Q1/Q2	Main	Scan	Measured	Integration	ISTD
		interferences	method	ion	time (s)	
Li	7/7		On-mass	Li*	0.5	Ве
В	11/11		On-mass	B ⁺	2.0	Ве
Na	23/23		On-mass	Na⁺	0.5	Sc
Mg	24/24		On-mass	Mg⁺	0.5	Sc
Al	27/27		On-mass	Αl⁺	0.3	Sc
Si	28/28	N ₂ +, CO+	On-mass	Si⁺	0.5	Sc
Р	31/47	NOH⁺, Cu⁺⁺	Mass shift	PO ⁺	2.0	Ве
S	32/48	0 ₂ +, Cu++	Mass shift	SO⁺	2.0	Ве
К	39/39	ArH⁺	On-mass	K ⁺	0.5	Ве
Ca	40/40	Ar⁺	On-mass	Ca⁺	0.3	Sc
Ti	48/48	SO⁺	On-mass	Ti⁺	0.5	Sc
V	51/51	(CIO+)	On-mass	V ⁺	0.3	Sc
Cr	52/52	ArC⁺	On-mass	Cr+	0.3	Sc
Mn	55/55	ArNH⁺	On-mass	Mn⁺	0.3	Sc
Fe	56/56	ArO⁺	On-mass	Fe⁺	0.3	Sc
Со	59/59		On-mass	Co⁺	0.3	Sc
Ni	60/60		On-mass	Ni ⁺	0.5	Sc
Zn	68/68	ArNN⁺, CuHHH⁺	On-mass	Zn⁺	2.0	Sc
Ga	71/71		On-mass	Ga⁺	0.5	In

Element	Q1/Q2	Main interferences	Scan method	Measured ion	Integration time (s)	ISTD
Ge	72/72	ArAr+	On-mass	Ge⁺	0.5	ln
As	75/91	(ArCl+)	Mass shift	As0+	1.0	ln
Se	78/78	ArAr+	On-mass	Se⁺	3.0	In
Rb	85/85		On-mass	Rb⁺	0.3	In
Sr	88/88		On-mass	Sr+	0.5	In
Zr	90/106		Mass shift	ZrO+	0.5	In
Nb	93/125	CuNO⁺	Mass shift	Nb00+	0.3	In
Мо	95/127	CuOO⁺	Mass shift	MoOO⁺	0.5	In
Ru	99/99	ArCu⁺	On-mass	Ru⁺	0.5	In
Rh	103/103	ArCu⁺	On-mass	Rh⁺	0.3	In
Pd	105/105	ArCu⁺	On-mass	Pd⁺	0.5	ln
Ag	107/107		On-mass	Ag⁺	0.3	ln
Cd	111/111		On-mass	Cd⁺	1.0	ln
Sn	118/118		On-mass	Sn⁺	0.5	ln
Sb	121/121		On-mass	Sb⁺	0.5	ln
Те	125/125		On-mass	Te⁺	3.0	In
Cs	133/133		On-mass	Cs⁺	0.5	In
Ва	137/137		On-mass	Ba⁺	0.5	In
Hf	178/194		Mass shift	HfO⁺	0.5	In
Та	181/213		Mass shift	Ta00+	0.5	In
w	182/214		Mass shift	WOO+	0.5	In
Re	185/185		On-mass	Re⁺	0.5	In
ir	193/193		On-mass	lr ⁺	0.5	In
Pt	195/195		On-mass	Pt ⁺	0.5	In
Au	197/197		On-mass	Au⁺	0.5	In
TI	205/205		On-mass	TI⁺	0.3	In
Pb	208/208		On-mass	Pb⁺	0.3	In
Bi	209/209		On-mass	Bi ⁺	0.3	In
Th	232/248		Mass shift	ThO⁺	0.3	ln
U	238/270		Mass shift	U00⁺	0.3	In

Results and Discussion

BECs and DLs in 5% HNO $_{_3}$ blank

Background equivalent concentrations (BECs) in 5% HNO $_3$ were obtained from the calibration plots for each analyte. Calibration plots for three alkaline elements (Li, Na, and K) are shown in Figure 1. The BECs for the three elements were 0.1, 6.1, and 5.4 ppt, respectively, indicating very low background signals obtained using the m-lens. Calibration plots for Si, P, and S are also given in Figure 1. The BECs for these challenging elements were 231, 7.2, and 84 ppt, respectively. P and S have relatively high first ionization potentials (IPs) and so are poorly ionized under cool plasma conditions. Using hot plasma conditions in this work, these poorly ionized elements – together with others such as B, Zn, As, Cd, Ir, Pt, and Au – were all measured at low ppt levels.

The BECs and 3σ DLs for all 49 elements in the 5% HNO $_3$ blank are shown in Figure 2. The BECs for most elements were below 1 ng/L (ppt) in solution. This value is equivalent to <1 µg/kg (ppb) relative to the solid Cu, taking the 1000x dilution into account. This sensitivity indicates that the 8900 ICP-QQQ method is suitable for ultratrace analysis of these impurity elements in high-purity Cu. Low ppt BECs were also achieved for alkali elements, Li, Na, and K, under the hot plasma conditions used. BECs at the 10s to 100s ppt-level were achieved for the most challenging elements: S (84 ppt) and Si (231 ppt).

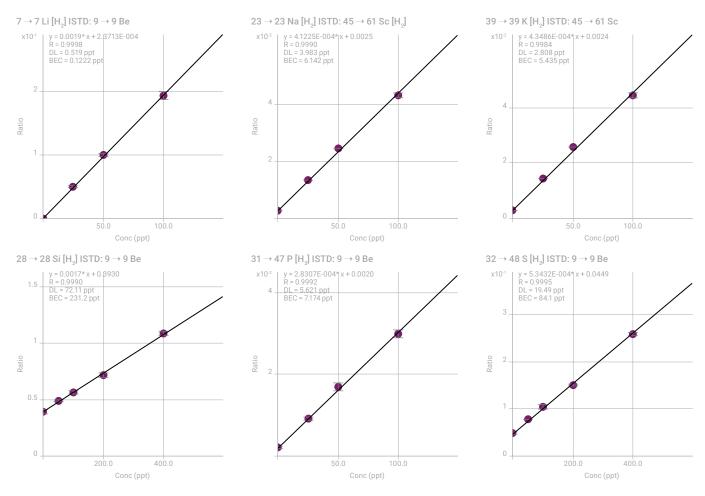


Figure 1. Calibrations for easily-ionized elements Li, Na, and K, and challenging elements Si, P, and S.

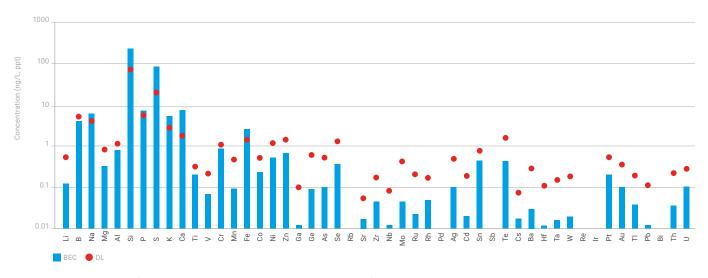


Figure 2. BECs and 3σ DLs for 49 elements in 5% HNO $_3$ blank. The BEC and DL for Rb, Pd, Sb, Re, Ir, and Bi could not be calculated, as the measured counts were zero in all replicates of the blank.

Determination of impurities in 0.1% 9N high purity copper

The 8900 ICP-QQQ method was used to determine the concentration of 49 elements in the 0.1% high purity copper solutions. ISTD correction was applied to correct for signal differences between the synthetic standards (with no Cu matrix) and the 0.1% Cu samples. Signal differences were all less than 30% between the non-matrix and Cu-matrix samples, demonstrating the robustness of the hot plasma conditions used.

All elements measured—apart from Si, S, and Te—were <10 ppt in the digest, as shown in Figure 3. Most elements were measured at 1 ppt or below, which is equivalent to <1 g/kg (ppb) in the solid metal. The mixed $\rm O_2$ + $\rm H_2$ reaction cell gas removed the significant spectral interferences caused by ArCu+ on Ru+, Rh+, and Pd+ (see Table 2). Removing the interferences allowed the determination of these elements at single- or sub-ppt levels (equivalent to single- or sub-ppb in the solid metal).

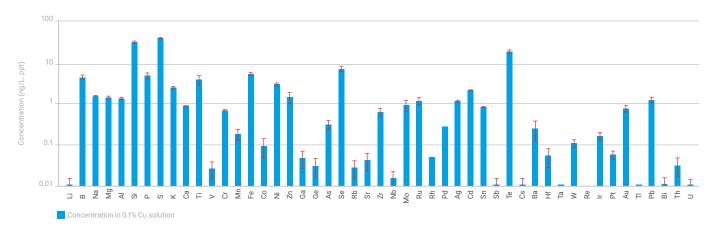


Figure 3. Measured concentrations of 49 elements in 0.1% 9N Cu sample (error bars = standard deviation for three samples). The values shown in ng/L (ppt) in solution) are equivalent to the values in μ g/kg (ppb) in the original solid metal. The concentration reported for Re was 0.000 ppt. SDs were zero for Rh, Pd, Ta, Re, and Tl.



Figure 4. Spike recovery test at 50 ppt (200 ppt for S, P, and Si) in 0.1% Cu solution. Most elements were within 90–110% recovery. The red lines indicate upper and lower limits of 85 to 115% recovery.

To validate the method, a spike recovery test was carried out for all 49 impurity elements. A 0.1% 9N copper blank solution was spiked at 50 ppt (200 ppt for Si, P, and S). The recoveries were within 84-112% for all 49 elements, with most being within 90-110%, as shown in Figure 4.

Conclusions

Ultratrace level impurities can be analyzed quickly and accurately in high purity copper metal digests using the Agilent 8900 ICP-QQQ. The optional m-lens ensures that the background signals for the alkali elements are minimized under hot plasma conditions. Using MS/MS mode with a mixed cell gas $(O_2 + H_2)$, the method delivered the following performance benefits:

- Low ppt level BECs were achieved for most impurities, including the alkali elements, using matrix-tolerant hot plasma conditions.
- Low-level BECs at the 10s to 100s ppt-level were obtained for sulfur and silicon—the most challenging elements to measure using ICP-MS.
- No matrix matching for Cu matrix was required, as ISTDs corrected for matrix differences between the standards (in 5% HNO3) and the samples (in 0.1% Cu).
- Using the fast and simple method with a single, mixed cell gas mode, a total of 49 elements were determined at ultralow levels in 0.1% high purity Cu sample.

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Single Nanoparticle Analysis of Asphaltene Solutions using ICP-QQQ

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Francisco Lopez-Linares, Laura Poirier, and Estrella Rogel Chevron Energy Technology Company, USA Agilent 8900 and ICP-MS MassHunter software module simplify spICP-MS analysis

Introduction

Single Particle ICP-MS (spICP-MS) is increasingly being used to characterize the nanoparticle (NP) content of samples dispersed in an aqueous media (1–5). In several industries—including oil refining, petrochemicals, and semiconductor manufacturing—there is also interest in determining NPs in hydrocarbon matrices. In this study, we report a new method using triple quadrupole ICP-MS (ICP-QQQ) for the multi-element characterization of NPs in the heavy asphaltene fraction of petroleum (6). The method can be used to differentiate between metals present in NPs and the dissolved metal content. It will therefore extend the understanding of the role and form of metals present in crude oils and petroleum-based products. The method also has wider applicability to the characterization of NP populations in other hydrocarbon-based matrices, such as NMP, PGMEA, butyl acetate, and other organic solvents used in the semiconductor industry.

In spICP-MS analysis, the ICP-MS uses a fast time resolved acquisition mode to measure the signal generated by each NP as it passes through the plasma. The high sensitivity and low background noise of ICP-MS enables the signals generated from individual NPs to be distinguished, and these key performance characteristics are greatly enhanced with ICP-QQQ. The superior control of interferences achieved using tandem MS (MS/MS) operation means that ICP-QQQ is especially suitable for some of the elements of most interest in NP analysis, such as Si, Ti, Fe, S, and others.

The intensity of the NP signal peak is proportional to the size of the particle and the concentration (mass fraction) of the analyte element within the particle. The frequency of the individual NP signals is directly proportional to the number of NPs in the sample, allowing calculation of the NP size distribution, particle number, particle concentration, and dissolved metal concentration, all from a single ICP-MS measurement. Nanoparticle method setup, acquisition, calibration, and data reporting are simplified using the optional Single Nanoparticle Application Module of Agilent ICP-MS MassHunter software.

In this study, the spICP-MS acquisition mode of the Agilent 8900 ICP-QQQ was used to identify and characterize trace elements in asphaltenes—a complex class of high molecular weight hydrocarbons found in heavy oil fractions and bitumens. Asphaltenes are defined by their solubility class. They are soluble in aromatics such as benzene or toluene, but insoluble in lighter paraffins, such as *n*-pentane or *n*-heptane. Asphaltenes, together with waxes and resins, are of interest in petrochemical processing as they can deposit in equipment and pipelines leading to production problems. Asphaltenes also contain a high proportion of the metals in crude oil, including elements such as V and Ni, which act as catalyst poisons, affecting the oil refining process.

Iron- and Mo-based NPs were identified in the asphaltene samples. In contrast, V and Ni were found to be present mainly as dissolved metals, likely metal porphyrins and other organometallic species. Data is provided on the concentration and size distribution of Fe and Mo NPs in the asphaltene samples, and the levels of dissolved metals is also presented. The results highlight the potential of spICP-MS for the routine characterization of metal NPs—as well as dissolved metals—in asphaltenes, crude oils, petroleum-derived materials, and other organic sample-types.

Experimental

Reagents and samples

Trace metal grade purity chemicals were used throughout (6).

Three separate samples of asphaltene were obtained from different sources:

- Asphaltene A—a heavy Mexican crude oil (14° American Petroleum Institute, API).
- Asphaltene B—an asphaltenic deposit recovered from a submersible pump.
- Asphaltene C—an oxidized asphalt obtained from a commercial plant that produces specialty asphalts.

Sample preparation

The asphaltenes were extracted from the sample matrix using n-heptane at a sample/solvent ratio of 1/20. The blended crude oil/heptane was heated to 80 °C. After one hour, the undissolved asphaltenes were recovered by filtering the mixture through a 0.8 μ membrane filter.

Calibration standard preparation for total metals

Calibration standard solutions for direct analysis were prepared from Conostan (Quebec, Canada) S-21+K oil-based multi-element organometallic standard. The diluent comprised trace metal grade purity o-xylene (Fisher Scientific), a matrix modifier (made from mineral oil; Fisher Scientific), and a dispersant (Chevron Oronite). Scandium and yttrium were used as internal standards, spiked at 0.1 and 5 mg/kg, respectively.

Multiple calibration standards ranging from 1 to $1000 \,\mu\text{g/kg}$ for each of the target elements were prepared by weight from the 10 mg/kg Conostan multi-element standard and o-xylene diluent. The diluent solution was used as the blank for calibration.

Nanoparticle reference materials and sample preparation

A 60 nm silver (Ag) NP reference material (nanoComposix) was used to calculate the nebulizer efficiency. The Ag NP reference material and the three asphaltene samples were diluted to a particle concentration of between 40 and 1000 ng/g with o-xylene (via propylene glycol methyl ether acetate, PGMEA). The solutions were sonicated for 5 min to ensure sample homogeneity. Elemental response factors were determined by measuring elemental standards for each target analyte $(10.0 \ \mu g/g)$ prepared with o-xylene.

Wet acid digestion for total metals

One to 5 g of each sample was heated on a hot plate at 100 °C for four hours with 1 to 2 mL of $\rm H_2SO_4$ (93–98 % w/w). The solution was then subjected to an ashing sequence, as described in reference 6. Six mL HCl (34 to 37 % w/w) and 2 mL HNO $_3$ (67 to 70 % w/w) were added, before further heating on a hot plate at 100 °C for 1 hour. Before analysis, Sc was added as an internal standard (to give a final Sc concentration of 5 mg/kg) and the solution was brought to a final volume of 25 mL with Milli-Q water.

Instrumentation

An Agilent 8900 Triple Quadrupole ICP-MS (ICP-QQQ) equipped with the standard glass concentric nebulizer and Peltier-cooled quartz spray chamber was used. For the analysis of samples prepared in organic solvent (spICP-MS analysis and total metals determinations in the diluted asphaltene samples), an optional "organics" quartz torch with a 1 mm ID injector was used in place of the standard quartz torch, which has a 2.5 mm injector.

For the organic sample analyses, oxygen (20% in Ar) was added to the injector gas stream after the spray chamber. $\rm O_2$ addition serves to decompose the carbon matrix thus avoiding carbon deposition on the interface cones. The more reactive plasma environment with $\rm O_2$ addition requires the use of the more chemically resistant optional platinum-tipped sampling and skimmer cones.

The high sensitivity of the ICP-QQQ enabled the samples for NP analysis to be diluted by a factor between 1:2100 and 1:2700 in o-xylene. Applying a high dilution factor minimizes the risk of colloidal particles forming an agglomerate after nebulization. The dilution ensures that the NPs are dispersed in the solution so that each NP passes through the plasma separately from any other NPs. As a result, the signal peaks measured are each generated by a single particle event and not from multiple, overlapping particle signals.

Signal intensities for each NP target element were acquired in fast Time Resolved Analysis (fast TRA) mode using a dwell time of 0.1 ms (100 μ s) per point, with no settling time between measurements. For Fe and Mo NPs, the signals were measured on-mass in MS/MS mode. For on-mass measurements, both quadrupoles (Q1 and Q2) were set to the target analyte ion mass of m/z 56 (for Fe) and m/z 95 (for Mo). Helium (He) cell gas was used in the 8900 ORS, to control the polyatomic interferences (mainly ArO on Fe at m/z 56). On-mass measurement with He cell gas was also used for the measurement of V and Ni (dissolved concentrations only—no NPs detected). ICP-QQQ operating conditions are given in Table 1.

Table 1. ICP-QQQ operating conditions

Parameter	Value
RF power (W)	1600
Sampling depth (mm)	10
Carrier gas (L/min)	0.35
Spray chamber temperature (°C)	-5
Option gas (L/min) (Ar 80%, O ₂ 20%)	0.35
Dwell time (ms)	0.1
He cell gas flow rate (mL/min)	5.0

Simplified workflow

The optional Single Nanoparticle Application Module of the ICP-MS MassHunter software was used for NP data acquisition and analysis. The spICP-MS Method Wizard guides the user through the process of nanoparticle method setup, data acquisition, data analysis, and presentation of the NP results data.

Nebulization efficiency

Nebulization efficiency is the ratio of the amount of analyte (aerosol) delivered to the plasma as a proportion of the amount of analyte (solution) entering the nebulizer. In this work, the nebulization efficiency was determined using the Ag NP reference material of known (60 nm) particle size. The reference material was first dispersed in PGMEA, and then further diluted in o-xylene. Nebulization efficiency, calculated from the certified size of Ag NP reference material, was found to be 0.065 or 6.5%.

Results and Discussion

Nanoparticle size distributions

Nanoparticles containing Fe and Mo were detected in the asphaltene samples using the sp-ICP-MS method. By contrast, the signals for V and Ni were continuous, rather than the discrete signal pulses caused by the presence of clusters or particles of these elements. This finding indicates that V and Ni were most likely in the form of dissolved metal complexes. The TRA signal charts for Fe in sample B (Figure 1) and Mo in sample A (Figure 2) show the signal-intensity as a function of time. In spICP-MS, the peak area for each particle signal "plume" can be used to calculate the particle mass and therefore size.

According to the literature, the Fe NPs are most likely to be present as iron oxides $(Fe_2O_3 \text{ or } Fe_3O_4)$ (7) and pyrrhotite (FeS) (8). The Mo NPs are most likely present as molybdenite (MoS_2) (9), which is readily formed from oil-soluble Mo complexes present in heavy fractions from crude oils (10, 11).

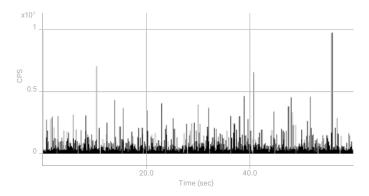


Figure 1. Asphaltene sample B: typical signals in counts per second (cps) for Fe (m/z 56) as a function of time.

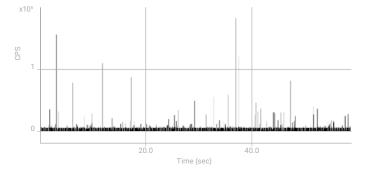


Figure 2. Asphaltene sample A: typical signals in counts per second (cps) for Mo (m/z 95) as a function of time.

The Fe and Mo NP size distribution plots for the three different asphaltene samples were calculated on the assumption that the Fe NPs were composed of ${\rm Fe_2O_3}$, and the Mo NPs were composed of ${\rm MoS_2}$. As shown in Figure 3, the average size of the Fe NPs varied among the samples. In contrast, the particle size distributions for Mo NPs are similar, with an average particle diameter in the range 70 to 80 nm (Figure 4).

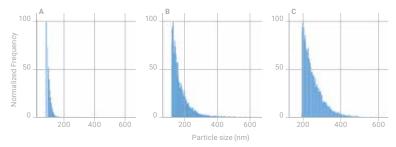


Figure 3. Comparison of size distributions for Fe NPs as Fe_2O_3 in the three asphaltene samples: A, B, and C. Modified with permission from J. Nelson et al., *Energy Fuels*, 2017, 31 (11), 11971–11976. © 2017 American Chemical Society.

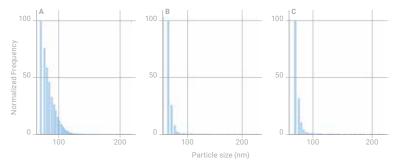


Figure 4. Comparison of size distributions for Mo NPs as MoS_2 in the three asphaltene samples: A, B, and C. Modified with permission from J. Nelson et al., *Energy Fuels*, 2017, 31 (11), 11971–11976. © 2017 American Chemical Society.

Concentration of the different forms of Fe and Mo

Uniquely, spICP-MS can distinguish between metal content that is contained in NPs (insoluble) and metal content that is dissolved in the sample matrix. The relative NP and soluble concentration data for Fe and Mo in the three asphaltene samples is given in Table 2. The data indicates that there was some variation in the distribution of metals among the asphaltene samples. In samples A and B, Fe was mostly present as NPs (76 and 91 wt. %, respectively), while in asphaltene sample C, less than half the Fe content was present as NPs. By contrast, Mo was almost all present as soluble forms (between 60 and 99 wt. %) in all three asphaltene samples, as shown in Table 2.

Table 2. Interference check results for ⁴⁸Ti in various matrices, with and without cell gas.

Asphaltene		Iron concentration, mg/kg					Molybdenum concentration, mg/kg				
samples	NPs	Soluble	Total, spICP-MS	Total, direct dilution	Total, wet acid digestion	NPs	Soluble	Total, spICP-MS	Total, direct dilution	Total, wet acid digestion	
Α	54.0 (76%)	17.0	71.0	39.5	68.0	3.48	5.33 (60%)	8.81	39.7	40.4	
В	173 (91%)	0.001	0.18	0.054	0.001	0.04	3.23 (99%)	3.27	0.78	0.52	
С	457 (47%)	508	965	420	750	0.07	6.33 (99%)	6.40	5.89	6.22	

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Total concentration of Fe and Mo

The total concentrations of Fe and Mo from the spICP-MS analysis (sum of the particle concentration and the dissolved concentration) were compared to the total metal concentrations measured by direct dilution and wet acid digestion. The results, which are given in Table 2, indicate that there was some variation between the three separate results for total concentrations of Fe and Mo. The spICP-MS and acid digestion approaches gave similar results for total Fe in all three samples, with the direct dilution results being consistently lower. This low bias for Fe present as particles in samples prepared and introduced using direct dilution has been reported in the literature (12). A study using Laser Ablation-ICP-MS (13) has also shown that large particles are not completely vaporized and ionized in the plasma. This finding could account for the low recovery observed for Fe in the direct analysis of the diluted samples. The difference compared to the spICP-MS total concentration may be due to the way the relatively large particles in these asphaltene samples are calibrated in spICP-MS vs the effect of incomplete dissociation and ionization of these larger particles measured by direct dilution.

The concentrations for Mo following direct dilution compare well with those obtained using wet acid digestion. For sample C, the total concentration found using spICP-MS also agrees well. But for the other two samples, the spICP-MS results do not tally with the total Mo concentrations found by the dilution and digestion methods. Total Mo by spICP-MS was found to be lower in Sample A and higher in Sample B, compared to total Mo determined by the other two approaches. Further studies are underway to investigate the discrepancies in the total concentration values for Mo calculated using the spICP-MS method compared with direct dilution and wet acid digestion.

Conclusions

Single particle-ICP-MS is becoming a widely used and well-established technique for the characterization of NPs in aqueous-based solutions. In this study, we show the potential for the spICP-MS methodology to be applied to complex hydrocarbon-based matrices of interest in petroleum refining and other industries.

The Agilent 8900 ICP-QQQ is especially suited to spICP-MS analysis because of its high sensitivity, low background, and unmatched control of spectral interferences. Setup and analysis for NP applications is facilitated by the optional Single Nanoparticle Application Module for ICP-MS MassHunter software.

Iron and molybdenum NPs were determined in three asphaltene samples from different sources associated with oil refining and petroleum-related product processing. No nickel or vanadium-containing NPs were detected in the heavy petroleum fractions suggesting that these elements are more likely to form dissolved organometallic complexes, such as porphyrins. This spICP-MS method is also able to differentiate between metal-containing NPs and dissolved metals.

Further work is in progress to expand the spICP-MS method for the routine characterization of metals in petroleum-derived samples, as well as in other hydrocarbon-based samples.

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Accurate Determination of TiO₂ Nanoparticles in Complex Matrices using the Agilent 8900 ICP-QQQ

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Introduction

Titanium dioxide (TiO_2) nanoparticles (NPs) are widely used in paints, food colorants, cosmetics, pharmaceuticals, and many other applications. Due to their high refractive index, TiO_2 NPs are common ingredients in sun protection products used to guard against UV exposure. However, the fate of NPs in the environment and the potential for toxic effects once absorbed into the body remain largely unknown. Many researchers have investigated different methodologies to measure TiO_2 NPs in cosmetic or food samples [1, 2, 3, 4].

 ${
m TiO}_2$ NPs have three principal levels of structure, beginning with nanoscale crystallites. These crystals fuse to form 'hard' nanoscale aggregates, which in turn associate to form microscale agglomerates [5]. When aqueous dispersions of ${
m TiO}_2$ NPs are prepared, the particle sizes observed are the aggregation or agglomeration sizes, which are usually different from (larger than) the primary (crystal) particle sizes [5, 6]. Typically, the primary size is measured by transmission electron microscopy (TEM) or X-ray diffraction (XRD), and the dispersion size is measured by laser diffraction spectrometry (LDS) or dynamic light scattering (DLS).

The relatively recent development of Single Particle ICP-MS (spICP-MS) now provides a powerful tool to characterize the NP content of dispersed samples. spICP-MS is used to measure the target element signals generated from individual NPs in the solution analyzed. This approach allows the simultaneous determination of the number, concentration, and size of particles present, as well as the dissolved element concentration.

In practice, however, there are some challenges for the measurement of ${\rm TiO}_2$ NPs using conventional single quadrupole ICP-MS (ICP-QMS). Many real samples may contain P, S, Ca, Si and C, and all these elements cause interferences that hinder the measurement of Ti. Also, the most abundant isotope of Ti, ⁴⁸Ti (73.7% abundance), suffers an isobaric interference from ⁴⁸Ca; therefore, the less interfered isotopes ⁴⁷Ti or ⁴⁹Ti are typically measured. However, the less abundant isotopes provide lower sensitivity, which limits the detection of smaller-sized ${\rm TiO}_2$ NPs by ICP-QMS.

The Agilent 8900 Triple Quadrupole ICP-MS (ICP-QQQ) can operate in MS/MS mode to resolve the spectral interferences on Ti, including the isobaric interference from ⁴⁸Ca on ⁴⁸Ti. The 8900 ICP-QQQ is a tandem mass spectrometer, meaning that it has an additional mass spectrometer with unit (1 u) resolution, positioned before the collision/reaction cell. This extra mass filter selects the ions that can enter the cell, providing control of the reaction chemistry when reactive cell gases are used. ICP-QQQ with MS/MS provides an elegant and effective approach for solving the most challenging spectral overlaps [7].

In this study, ${\rm TiO_2}$ NPs in sunscreen were measured in spICP-MS mode using the Agilent 8900 ICP-QQQ in MS/MS mode. The optional Single Nanoparticle Application Module software for ICP-MS MassHunter was used for method setup and data processing.

Current regulations

The methodologies used to evaluate the properties of nanomaterials are not yet considered to be finalized and approved, which may be impeding the introduction of specific regulations relating to NPs. In June 2014, the USA Food and Drug Administration (FDA) issued guidance on the safety assessment of nanomaterials in cosmetic products [8]. As part of the FDA, the Center for Drug Evaluation and Research (CDER) is examining the safety of titanium dioxide (and zinc oxide) nanomaterials for sunscreen use as part of an ongoing regulatory process to establish a final monograph for over-the-counter (OTC) sunscreen drug products [9].

Currently, the European Union Scientific Committee on Consumer Safety (SCCS) considers that it is safe to use ${\rm TiO_2}$ NPs as a UV filter at a concentration up to 25% in sunscreens. Manufacturers must respect this limit according to European legislation (annex VI list of UV filters) of the EU regulation on cosmetic products; regulation EC 1223/2009 [10]. The regulation was amended in 2016 to state that in the case of combined use of titanium dioxide and titanium dioxide (nano), the sum shall not exceed 25% [11].

In 2016, following a request from the European Commission to the European Food Safety Authority (EFSA), the Scientific Panel on Food Additives and Nutrient Sources added to Food (ANS) considered the safety of titanium dioxide (${\rm TiO}_2$, E 171) when used as a food additive [12]. The Panel will establish a health-based guidance value for acceptable daily intake (ADI) once more data is available on the reproductive toxicity of E 171.

Experimental

Reference materials and calibration solutions

The TiO2 standard reference material (SRM) NIST 1898 Titanium Dioxide (Maryland, US) was used. The SRM contains crystal or primary sized particles <50 nm, but the size of particles dispersed in the aqueous phase range from 71 to 112 nm due to nanoscale aggregation [5]. The SRM was diluted with de-ionized water to a particle concentration that was calculated to give 500 – 2000 particle counts per minute, and sonicated to ensure sample homogeneity. A 1 ppb Ti ionic standard prepared in 1 % nitric acid was used to measure the elemental response factor for Ti. A gold NP RM with a nominal particle size of 60 nm (NIST 8013 Gold Nanoparticles) was used to measure the nebulization efficiency of the ICP-QQQ.

Sunscreen samples

Sunscreen products were bought in a local store in Tokyo, Japan. The samples were diluted with de-ionized water plus 0.1~% Triton[™] X-100. The results obtained from an initial screening analysis using the spICP-MS method, showed the size-range of TiO_2 particles present in the different sunscreen samples varied. One of the samples contained particles <30 nm, while another product contained particles sized 30 to 200 nm. A sunscreen that contained TiO_2 NPs sized 30 to 100 nm was selected for further investigation. The selected sunscreen was

prepared in various diluent matrices: de-ionized water; tap water; and a "matrix mixture" containing 100 ppm of P and S, 50 ppm of Ca and Si, and 0.1 % of ethanol. The matrix mixture was used to check the impact of matrix-based interferences on the measurement of Ti.

Instrumentation

An Agilent 8900 Advanced Applications configuration ICP-QQQ was used throughout. The instrument was equipped with the standard glass concentric nebulizer and quartz spray chamber, optional quartz torch with 1.0 mm i.d. injector, and standard nickel sampling and skimmer cones. Samples were introduced directly into the ICP-QQQ using the standard peristaltic pump and 1.02 mm i.d. pump tubing. Analyses were performed in fast Time Resolved Analysis (fast TRA) mode, using a dwell time of 0.1 ms (100 μ s) per point, with no settling time between measurements. The major titanium isotope, ⁴⁸Ti, was measured in MS/MS mass-shift mode, using a mixed cell gas containing oxygen and hydrogen to resolve all the polyatomic and isobaric interferences. Q1 was set to m/z 48 (the mass of the precursor ⁴⁸Ti ion) and Q2 was set to m/z 64 (the mass of the target product ion ⁴⁸Ti¹⁶O). O_2 and O_2 and O_3 are detailed in Table 1. The operating conditions of the Agilent 8900 ICP-QQQ are detailed in Table 1.

Table 1. ICP-QQQ operating conditions.

Parameter	Value
RF power	1550 W
Sampling depth	8.0 mm
Carrier gas	0.70 L/min
Sample uptake rate	0.35 L/min
Spray chamber temp.	2 °C
Dwell time	0.1 ms
Settling time	None
Acquisition mode	MS/MS (Q1: m/z 48, Q2: m/z 64)
Oxygen flow rate	0.15 mL/min (10% of full scale)
Hydrogen flow rate	7.0 mL/min
Axial Acceleration	1.0 V
Octopole bias voltage	-6 V
Energy discrimination	-15 V

The Single Nanoparticle Application Module of the ICP-MS MassHunter software was used for method setup and data analysis. Sample results for an entire batch are summarized in the interactive 'Batch at a Glance' table. Detailed graphical results are displayed for selected samples, permitting visual confirmation and optimization of parameters if needed.

Results and Discussion

Optimization of cell gas conditions using ionic Ti solution

Before measurement of the NPs, cell gas conditions were investigated. Ti reacts readily with oxygen, so can be measured as TiO+ in oxygen mass shift mode. The first quadrupole (Q1) was set to pass only m/z 48, to allow 48Ti+ (and any on-mass interferences) to enter the cell. The second quadrupole (Q2), which is located after the collision/reaction cell, was set to m/z 64 to pass the target product ion (48 Ti 16 O $^{+}$) to the detector. Any potential native ion overlaps at m/z 64 (e.g. 64Zn and 64Ni) are rejected by Q1. Most of the primary interferences at m/z 48, such as ${}^{32}S^{16}O+$, ${}^{30}Si^{18}O^+$, ${}^{31}P^{16}OH^+$, ${}^{12}C^{18}O_2^+$, can be avoided by measuring Ti as TiO+ in oxygen cell gas mode. However, some of the ⁴⁸Ca ions also react with oxygen to form ⁴⁸CaO+, which interferes with the ⁴⁸TiO+ product ions at m/z 64. Adding hydrogen gas can eliminate the Ca interference by converting CaO+ to CaOH+. TiO+ does not react in the same way with H_a cell gas, so remains as the TiO+ product ion at m/z 64. Inter-isotope overlaps (such as 46 Ti 18 O and ⁴⁶Ca¹⁸O) can affect the ⁴⁸Ti¹⁶O measurement at *m/z* 64 when a single quadrupole or bandpass MS system is used. With MS/MS, however, these overlaps are avoided as the precursor ions (46Ti and 46Ca) are rejected by Q1 and so do not enter the cell to react.

Table 2 shows the quantitative results for Ti (measured as 48 Ti+ in no gas mode and 48 TiO+ in O $_2$ /H $_2$ mode) in various matrices. The quantitative results obtained in no gas mode show a large positive error due to the interferences on 48 Ti. In contrast, O $_2$ /H $_2$ cell gas mode effectively reduces the interferences including the potential CaO+ product ion overlap formed from 48 Ca. This method enables the TiO+ product ion from the most abundant isotope of Ti (mass 48; 73.7% relative abundance) to be measured, providing the sensitivity required for detection of small particles.

Table 2. Interference check results for ⁴⁸Ti in various matrices, with and without cell gas.

Cell gas mode	Sensitivity	Apparent concentration of Ti, measured as ⁴⁸ Ti+ or ⁴⁸ TiO+ (ppb)						
	(cps/ppb)	100 ppm P	100 ppm S	50 ppm Ca	50 ppm Si	0.1% ethanol	Matrix mixture*	
No gas	155,000	1.7	6.0	225	0.39	0.14	261	
O ₂ + H ₂	79,000	0.010	0.001	0.18	0.054	0.001	0.23	

^{*}Includes all the matrices (100 ppm of P and S, 50 ppm of Ca and Si, and 0.1% ethanol).

Measurement of a TiO, NP reference material

NIST 1898 ${\rm TiO_2}$ NP reference material was measured by ICP-QQQ in MS/MS mass-shift mode with ${\rm O_2/H_2}$ reaction gas. The time resolved signal chart for NIST 1898 (Figure 1) shows clear NP peaks with a wide variation in intensity (peak heights). In single particle ICP-MS, the peak height for each particle signal "plume" is representative of the particle mass (size). Figure 2 shows the signal frequency distribution for NIST 1898 (upper), and the calculated particle size distribution (lower). The mean size of 71 nm agrees well with the results by LDS (71 \pm 4 nm), X-Ray Disc Centrifugation (77 \pm 7 nm), and DLS (112 \pm 4 nm) according to the NIST certificate [5]. Note that DLS measures the hydrodynamic particle size, which includes the layer where the particle surface interacts with the solvent. Consequently DLS has been reported to give particle sizes that are larger than the value measured by other techniques [13].

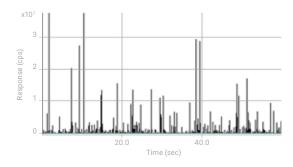


Figure 1. Time resolved signal for NIST 1898 TiO_2 NP reference material. The blue line represents a baseline automatically set by the MassHunter software function.

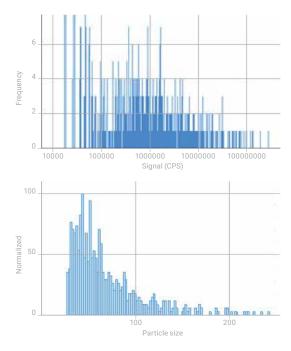


Figure 2. Signal frequency distribution (upper) and particle size distribution (lower) for NIST 1898 TiO, NP reference material.

Analysis of TiO, NPs in sunscreens

 ${
m TiO_2}$ NPs were measured in a commercial sunscreen prepared (dispersed) in several different solutions and the results are presented in Figure 3. Figure 3-A shows the ${
m TiO_2}$ signal distribution and Figure 3-B shows the particle size distribution for the sunscreen dispersed in de-ionized water (plus Triton X-100). The mean particle size of ${
m TiO_2}$ was calculated as 77 nm. The same sunscreen was dispersed in tap water (Figures 3-C and 3-D), and a synthetic matrix mixture comprising 100 ppm of P and S, 50 ppm of Ca and Si, 0.1 % of ethanol (Figures 3-E and 3-F). These results show signal distributions that are almost the same as the ones obtained for the sunscreen dispersed in de-ionized water. The mean particle sizes (79 nm for tap water and 84 nm for the matrix mixture) are similar. The particle size detection limit (the threshold between the baseline noise and particle signals) was about 30 nm for the dispersed sunscreens in all the matrices. The synthetic high matrix (Figure 3-E and 3-F) did not affect the size-DL or the accuracy of the particle size measurement.

The results show that ${\rm TiO_2}$ NPs <100 nm diameter can easily be measured using the MS/MS capability of the 8900 ICP-QQQ, even in a high concentration matrix.

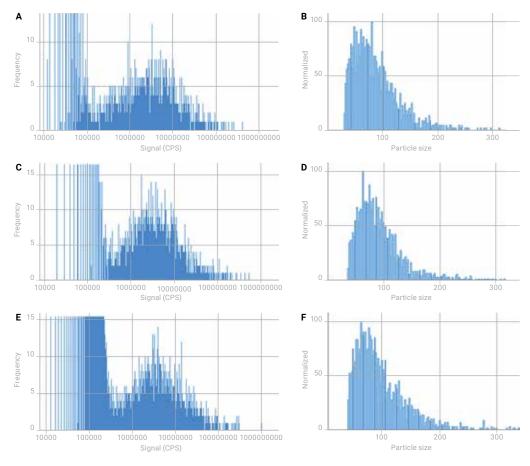


Figure 3. TiO₂ NP measurement of commercial sunscreen using ICP-QQQ. Signal distribution A) and particle distribution B) of sunscreen dispersed in de-ionized water. Signal distribution C) and particle distribution D) of sunscreen dispersed in tap water. Signal distribution E) and particle distribution F) of sunscreen dispersed in the matrix mixture.

Conclusions

The Agilent 8900 ICP-QQQ operating in MS/MS mode with $\rm O_2/H_2$ cell gas was used for the successful determination and characterization of $\rm TiO_2$ nanoparticles in various sample matrices. MS/MS mass-shift mode effectively resolved the polyatomic and isobaric ions that interfere with the measurement of Ti at its most abundant isotope. This unique MS/MS capability provided a particle size detection limit of \sim 30 nm.

Overall, the method delivered low background signals and excellent sensitivity, even in the presence of a high level of P, S, Ca, Si, and C matrix.

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Foods

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Accurate and Sensitive Analysis of Arsenic and Selenium in Foods using ICP-QQQ to Remove Doubly-Charged REE Interferences

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Keywords

arsenic, selenium, rare earth elements, environmental, agricultural, human health, NIST 1547 Peach Leaves, NIST 1515 Apple Leaves, oxygen mass-shift

Introduction

Concern about the impact on public health from potentially toxic elements and compounds present in everyday foodstuffs has led to new legislative guidance [1, 2]. The inorganic forms of arsenic (As) are known to be toxic and carcinogenic to humans, and food and drinks are a potential source of exposure [3]. Selenium (Se) is an essential micro-nutrient that can be deficient in the diet as Se-poor soils yield Se-poor food crops. Accurate quantification of Se in food is necessary to assess nutrient status.

As and Se can be difficult to quantify accurately at trace levels by conventional quadrupole ICP-MS (ICP-QMS), as all the analytically useful isotopes can suffer from multiple spectral interferences, as shown in Table 1. Potential interferences on As and Se include the doubly-charged ions of the Rare Earth Elements (REE++) and matrix and plasma-born polyatomic ions. The quadrupole mass spectrometer separates ions based on mass to charge ratio (*m/z*), so the REE++ ions appear at half their true mass, overlapping the singly-charged analyte ions of As and Se. Typically the REE content in food samples is low, but crops grown in REE-enriched soils may take up higher concentrations of these elements [4, 5] leading to false positive results for As and Se. In this study, we evaluated the capability of the Agilent 8800 ICP-QQQ in MS/MS reaction mode to remove interferences, including REE++, on As and Se.

Table 1. Spectroscopic interferences on As and Se isotopes.

As and Se isotope			Interference			
Element	Mass	Abundance %	Doubly charged	Matrix	Dimer	
As	75	100	¹⁵⁰ Sm ⁺⁺ , ¹⁵⁰ Nd ⁺⁺	⁴⁰ Ar ³⁵ Cl ⁺ , ⁴⁰ Ca ³⁵ Cl ⁺		
Se	77	7.63	¹⁵⁴ Sm ⁺⁺ ,	⁴⁰ Ar ³⁵ Cl ⁺ , ⁴⁰ Ca ³⁵ Cl ⁺		
	78	23.77	¹⁵⁶ Gd ⁺⁺ , ¹⁵⁶ Dy ⁺⁺	⁴¹ K ³⁷ Cl ⁺	³⁸ Ar ⁴⁰ Ar ⁺ , ³⁹ K ³⁹ K ⁺	
	80	49.61	¹⁶⁰ Gd ⁺⁺ , ¹⁶⁰ Dy ⁺⁺ ,	³² S ₂ ¹⁶ O ⁺ , ³² S ¹⁶ O ₃ ⁺ , ⁴⁰ Ca ⁴⁰ Ar ⁺ , ⁴⁵ Sc ³⁵ Cl ⁺	⁴⁰ Ar ⁴⁰ Ar⁺, ⁴⁰ Ca ⁴⁰ Ca⁺	
	82	8.73	¹⁶⁴ Dy ⁺⁺ , ¹⁶⁴ Er ⁺⁺	⁴⁵ Sc ³⁷ Cl ⁺		

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/General purpose.

Acquisition parameters: MS/MS mass-shift mode using O_2/H_2 at a gas flow of 0.6 mL/min and 1.0 mL/min respectively. As was measured as the reaction product ion AsO+ at m/z 91, and Se was measured as SeO+ at m/z 96.

Reagents: Two National Institute of Standards and Technology (NIST) standard reference materials (SRMs), NIST 1547 Peach Leaves and NIST 1515 Apple Leaves, were studied. These SRMs contain low $\mu g/kg$ levels of As and Se in the presence of mg/kg levels of REEs.

Sample prep: All samples were acid digested using a closed vessel microwave digestion system. The SRMs were prepared in triplicate. First, 0.25 g sample was digested in 2.5 mL of 9:1 HNO $_3$:HCl acid mix, and the digest was then diluted to a final weight of 25 g with ultra-pure water. 5% butanol was added to the internal standard mixture to equalize the organic plasma load between samples and standards. NIST 1547 and 1515 contain low μ g/kg concentrations of As and Se (Table 2) and high concentrations of REEs. Reference (non-certified) values for Nd, Sm and Gd are 7, 1, and 1 mg/kg in NIST 1547, and 17, 3, and 3 mg/kg in NIST 1515, respectively.

Results and Discussion

Both SRM digests were analyzed using the 8800 ICP-QQQ with $\rm O_2/H_2$ as the reaction gas (Table 2). A previous study showed that the presence of $\rm H_2$ in the cell further improved Se detection capability [6]. The measured values for As and Se in NIST 1547 and 1515 were well within the certified range for both SRMs, demonstrating the successful elimination of the REE++ interferences in $\rm O_2/H_2$ MS/MS mode on the 8800. These results were obtained without the need for correction equations (i.e. uncorrected).

Table 2. Analysis of As and Se in NIST 1547 and 1515 in He mode and H_2 mode using ICP-QMS (both uncorrected and corrected data is given) and by ICP-QQQ in MS/MS mode. All concentrations are in mg/kg, and are averages of 3 replicate sample digests expressed as mean \pm standard deviation.

		ICP-QMS He mode		ICP-QMS ${ m H_2}$ mode		ICP-QQQ 0 ₂ / H ₂ mass-shift
SRM	Certified	Uncorrected	Corrected	Uncorrected	Corrected	Uncorrected
As (mg/kg)						
NIST 1547	0.060±0.018	0.170±0.016	0.068±0.003*	0.113±0.004	0.079±0.004*	0.065±0.002*
NIST 1515	0.038±0.007	0.250±0.016	0.026±0.021*	0.126±0.005	0.047±0.004*	0.032±0.002*
Se (mg/kg)						
NIST 1547	0.120±0.009	0.394±0.04	0.113±0.02*	0.119±0.009*	0.119±0.009*	0.127±0.006*
NIST 1515	0.050±0.009	0.808±0.04	0.013±0.04*	0.050±0.003*	0.050±0.003*	0.047±0.006*

^{*95%} confidence interval overlaps with the certified range

Results obtained using ICP-QMS are included for comparison purposes. "Corrected" refers to the use of correction equations. ICP-QMS operating in helium mode is suitable for the analysis of As and Se in general routine sample types that might contain a small amount of REEs, and $\rm H_2$ mode has been shown to be effective at reducing doubly charged species. Compared to these conventional methods, ICP-QQQ had 10-fold lower detection limits, which makes it particularly suited to low level determination of As and Se in complex sample matrices.

ICP-QMS can also use O_2 mass-shift mode but, with ICP-QMS, all matrix and analyte ions enter the cell, so the analyte reaction product ions measured (75 As 16 O $^+$ at m/z 91, and 78 Se 16 O $^+$ at m/z 94), could suffer overlap from existing analyte or matrix ions at the product ion mass (e.g. $^{91/94}$ Zr $^+$ and 94 Mo $^+$). To confirm that the ICP-QQQ MS/MS method can be applied to samples that contain high concentrations of Zr and Mo, an aliquot of NIST 1547 was spiked with 1 mg/L (1000 ppm) Zr and Mo, and the results are shown in Table 3. The measured values for As (as AsO $^+$) and Se (as SeO $^+$) in the spiked sample are the same as for the unspiked samples, demonstrating that MS/MS mode is effective at rejecting existing overlapping ions present at the mass of the cell-formed analyte product ions. This capability is unique to the tandem mass spectrometer configuration of the 8800 ICP-QQQ, where the ions that enter the cell are controlled by an additional mass filter, Q1, positioned in front of the collision/reaction cell.

Table 3. ICP-QQQ measured results for As and Se in NIST 1547 unspiked and spiked with 1 mg/L Zr and Mo. No correction equations were applied.

ICP-QQQ O ₂ /H ₂ mass-shift				
SRM NIST 1547	Certified	Unspiked (n=3)	Spiked with 1 mg/L Zr & Mo (n=1)	
As (mg/kg)	0.060±0.018	0.065±0.002	0.063	
Se (mg/kg)	0.120±0.009	0.127±0.006	0.13	

Conclusions

The Agilent 8800 ICP-QQQ with MS/MS capability has been shown to be the optimum method to successfully measure trace levels of As and Se in the presence of high concentration of REEs in NIST 1547 Peach Leaves and NIST 1515 Apple Leaves. All REE doubly-charged and matrix-based polyatomic interferences that affect As and Se measurement at m/z 75 and m/z 78 are avoided using O_2/H_2 cell gas and MS/MS mass-shift mode. Arsenic is shifted to its product ion AsO+ which is measured at m/z 91, and Se is shifted to SeO+, measured at m/z 94. Importantly, MS/MS mode also eliminates potential ion overlaps at m/z 91 and m/z 94 from O_2/H_2 and O_2/H_2 and O_3/H_2 and O_3

More Information

For a full account of this application see publication: Advantages of reaction cell ICP-MS on doubly charged interferences for arsenic and selenium analysis in foods, Brian P. Jackson, Amir Liba and Jenny Nelson, *J. Anal. At. Spectrom.*, **2015**, Advance Article. DOI: 10.1039/C4JA00310A.

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High Throughput Determination of Inorganic Arsenic in Rice using Hydride Generation-ICP-QQQ

Authors

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Keywords

arsenic, arsenite, arsenate, IMEP-7 rice, NIST 1568a Rice Flour, oxygen mass-shift

Introduction

The concentration of potentially toxic chemicals such as arsenic in rice is closely monitored to ensure food safety. However, the toxicity of arsenic depends on the chemical form or "species" of the element that is present rather than total concentration. Inorganic arsenic (iAs) species, arsenite (As(III)) and arsenate (As(V)), are known to be carcinogenic and highly toxic, whereas the common organoarsenic species monomethylarsonic acid (MMA) and dimethylarsinic acid (DMA) are less toxic [1].

Rice is an important food source for a large percentage of the world's population, but it does contain relatively high concentrations of iAs due to the uptake of As from the soil and water in which the rice plants are grown. Available As from soils can be both naturally occurring and due to anthropogenic sources such as As-based pesticides that were widely used until the 1970s. Clearly there is an urgent food-safety requirement for a simple and quick analytical method to screen large numbers of rice and other food samples for iAs. In this study, a fast and sensitive method using hydride generation (HG) with ICP-QQQ is described for the separation and detection of iAs in commercial rice samples.

Experimental

Instrumentation: A Hydride Generation (HG) accessory for Agilent's ICP-MS Integrated Sample Introduction System (ISIS) was used with an Agilent 8800 #100 ICP-QQQ. When treated with NaBH $_4$ under acidic conditions, iAs is very efficiently converted into volatile arsine (AsH $_3$), whereas organically bound As compounds are not converted, or form only less volatile arsine species such as dimethylarsine (CH $_3$) $_2$ AsH, which has a boiling point of 35°C. Adding high concentrations of HCI further reduces the production of the less volatile arsines, and iAs is almost entirely converted to arsine, enabling the measurement of iAs without species separation using chromatography.

Plasma conditions and ion lens tune: RF power = 1550 W, Sampling depth = 8.0 mm and Carrier gas flow rate = 0.93 L/min were used with soft extraction tune, Extract 1 = 0.5 V and Extract 2 = -170 V.

Acquisition parameters: MS/MS mass-shift mode using O_2 cell gas at a flow rate of 2.0 mL/min. As was measured as the reaction product ion AsO+ at m/z 91.

Samples: Samples included 31 different rice products purchased from local stores. Sub-samples (30g) of the commercially sourced rice samples were ground to a fine homogeneous powder using a coffee grinder. Two rice reference materials IMEP-107 rice (Institute for Reference Materials and Measurements Geel, Belgium) and rice SRM NIST 1568a Rice Flour were used as quality control for iAs concentration measurements.

Sample prep: 0.15g of each rice sample was digested in 1 mL concentrated HNO_3 and 2 mL H_2O_2 (30 % w/w) using open vessel digestion in a CEM Mars microwave system. All samples were diluted to a final volume of 30 mL using deionized water.

Table 1. Cool plasma operating conditions.

Sample flow rate (mL/min)	0.5
HCl flow rate (mL/min)	2.5
NaBH4 flow rate (mL/min)	0.5
Reaction coil volume (mL)	0.23
Ar flow rate (for HG) (L/min)	0.3
Ar flow rate (for nebulization of IS) (L/min)	0.85-0.95

Results and Discussion

The speciation results for iAs in IMEP-7 rice and NIST 1568a Rice Flour obtained using HG-ICP-QQQ were in good agreement with the values obtained using HPLC-ICP-QQQ and with the reported values (Table 2).

Table 2. iAs results obtained using HG-ICP-QQQ and HPLC-ICP-QQQ.

Inorganic As			
	HG-ICP-QQQ (μg/kg)	HPLC-ICP-QQQ (µg/kg)	Reported value (µg/kg)
IMEP-7 rice	100 ± 11 (n=15)	110 ± 12 (n=15)	107 ± 14 [2]
NIST 1568a	94 ± 8 (n=3)	105 ± 4 (n=3)	94 ± 12

A summary of the values for iAs, DMA and total As determined in the commercial rice samples is given in Table 3. The dominant arsenic species found in rice are iAs and dimethylarsinic acid (DMA), with only trace amounts of methylarsonic acid (MA). The method uses HCl (5 M) and NaBH $_4$ for the selective generation of arsines where iAs and DMA are converted almost exclusively to AsH $_3$, with only minor (2-4%) conversion of DMA to dimethylarsine. MA forms methylarsine at approximately 40% efficiency with the method; however, since MA is generally absent from rice – or only present in trace amounts – this should not affect the quantification of iAs.

Table 3. Speciation results of iAs in 31 rice products determined by HG-ICP-QQQ and HPLC-ICP-QQQ. Results are also given for DMA and MMA, and the total As concentration determined by ICP-QQQ. All data \pm SD, with n=3 for speciation and n=2 or 3 for total As.

Rice product					
	HG iAs (μg/kg)	HPLC iAs (μg/kg)	HPLC DMA (μg/kg)	HPLC MMA (μg/kg)	Total As (μg/kg)
Arborio Risotto	113 ± 13	120 ± 18	63 ± 7	<l0q< td=""><td>236 ± 15</td></l0q<>	236 ± 15
Organic ArbRis	109 ± 12	119 ± 13	60 ± 8	<lod< td=""><td>150 ± 7</td></lod<>	150 ± 7
Basmati, 1	41 ± 4	53 ± 7	8 ± 1	<lod< td=""><td>100 ± 12</td></lod<>	100 ± 12
Basmati, 2	76 ± 6	88 ± 6	28 ± 4	<lod< td=""><td>91 ± 8</td></lod<>	91 ± 8
Basmati (white)	72 ± 11	69 ± 9	24 ± 1	<lod< td=""><td>240 ± 5</td></lod<>	240 ± 5
Organic Basmati (white)	95 ± 3	104 ± 3	21 ± 2	<lod< td=""><td>117 ± 13</td></lod<>	117 ± 13
Brown Rice	127 ± 6	137 ± 5	35 ± 2	<lod< td=""><td>205 ± 2</td></lod<>	205 ± 2
Japanese Rice	101 ± 5	99 ± 5	123 ± 1	<l0q< td=""><td>252 ± 10</td></l0q<>	252 ± 10
Long Grain (white)	89 ± 2	85 ± 1	16 ± 1	<lod< td=""><td>121 ± 11</td></lod<>	121 ± 11
Long Grain Rice, 1	103 ± 2	94 ± 1	218 ± 9	<l0q< td=""><td>392 ± 23</td></l0q<>	392 ± 23
Long Grain Rice, 2	40 ± 2	52 ± 10	39 ± 3	<l0q< td=""><td>111 ± 8</td></l0q<>	111 ± 8
Long Grain white	47 ± 2	61 ± 4	19 ± 4	<lod< td=""><td>102 ± 9</td></lod<>	102 ± 9
Organic Long Grain (brown)	111 ± 7	131 ± 14	54 ± 7	<l0q< td=""><td>207 ± 15</td></l0q<>	207 ± 15
Organic (white)	65 ± 4	65 ± 2	11 ± 1	<lod< td=""><td>92 ± 4</td></lod<>	92 ± 4
Paella, 1	60 ± 5	65 ± 2	38 ± 1	1.2 ± 0.1	136 ± 1
Paella, 2	66 ± 4	70 ± 3	17 ± 1	<lod< td=""><td>121 ± 6</td></lod<>	121 ± 6
Spanish Paella	67 ± 2	67 ± 3	13 ± 1	<lod< td=""><td>109 ± 7</td></lod<>	109 ± 7
Pudding Rice	124 ± 9	125 ± 11	44 ± 5	<lod< td=""><td>202 ± 4</td></lod<>	202 ± 4
Rice Flour	40 ± 1	46 ± 5	19 ± 2	<lod< td=""><td>102 ± 6</td></lod<>	102 ± 6
Carnaroli Risotto Rice	81 ± 2	82 ± 4	84 ± 2	<lod< td=""><td>210 ± 15</td></lod<>	210 ± 15
Risotto Rice	97 ± 11	114 ± 10	72 ± 9	<l0q< td=""><td>221 ± 17</td></l0q<>	221 ± 17
FLG Thai (white)	88 ± 3	102 ± 3	52 ± 5	<lod< td=""><td>197 ± 9</td></lod<>	197 ± 9
Thai Jasmine	61 ± 4	64 ± 3	49 ± 5	<lod< td=""><td>143 ± 3</td></lod<>	143 ± 3
Thai Jasmine (white)	62 ± 4	62 ± 3	49 ± 2	<lod< td=""><td>171 ± 5</td></lod<>	171 ± 5
Vietnamese Rice Paper	21 ± 2	28 ± 1	<l0q< td=""><td><lod< td=""><td>58 ± 10</td></lod<></td></l0q<>	<lod< td=""><td>58 ± 10</td></lod<>	58 ± 10
White Rice	71 ± 5	76 ± 5	14 ± 4	<l0q< td=""><td>124 ± 1</td></l0q<>	124 ± 1
Whole Grain	133 ± 2	127 ± 2	151 ± 12	7.2 ± 0.3	370 ± 19

LOQ HG-ICP-QQQ: 5 μg/kg, HPLC-ICP-QQQ: 1.1 μg/kg

Conclusions

Inorganic arsenic (iAs) was quantified at low ppb levels in extracts of 31 rice samples using an Agilent Hydride Generator/ISIS coupled to an Agilent 8800 ICP-QQQ. Results obtained using HG-ICP-QQQ were in good agreement with HPLC-ICP-QQQ values across a wide linear range, with comparable limits of detection.

Following a simple sample preparation using microwave extraction, quick separation of iAs and DMA by HG-ICP-QQQ was performed online. A previous study has shown that HG-ICP-QQQ requires only 4 minutes total run time per sample (5 replicate measurements) compared to speciation with HPLC which commonly takes between 5 and 10 minutes for each sample replicate [3]. Data handling for the HG method is also straightforward as no peak-integration is necessary.

The new HG-ICP-MS method offers fast analysis time, high throughput, and simple, reliable operation. This makes it ideally suited to screening large numbers of food samples to meet the increasing demand for the routine determination of iAs in food, especially rice-based products.

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More Information

For a full account of this application see publication: Ásta H. Pétursdóttir *et al.*, Hydride generation ICP-MS as a simple method for determination of inorganic arsenic in rice for routine biomonitoring, *Anal. Methods*, **2014**,6, 5392-5396.

Determination of Pesticides using Phosphorus and Sulfur Detection by GC-ICP-QQQ

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Keywords

pesticides, GC-ICP-MS, GC-ICP-MS/ MS, GC-ICP-QQQ, sulfur, phosphorus

Introduction

The determination of pesticide residues in food products is important. Most pesticide residue laboratories use some variation of the QuEChERS (Quick, Easy, Cheap, Effective, Rugged, and Safe) extraction methods. Typically, the extracted material is analyzed using GC/MS/MS the thermally stable, less polar pesticides, or LC-MS/MS for the less volatile and/or more polar ones. A more recently developed alternative technique involves coupling GC to triple quadrupole ICP-MS (GC-ICP-QQQ). Pesticides can be quantified using GC-ICP-QQQ by measuring the heteroatoms P and S (also Cl and Br) contained in most pesticides. GC-ICP-QQQ offers good selectivity, specificity, and sensitivity that can be greater than the established methods.

Experimental

Instrumentation: An Agilent 7890 GC was coupled to an Agilent 8800 #100 ICP-QQQ using an Agilent GC-ICP-MS interface (G3158D).

GC: Two Agilent columns were used in series. The first column was a 5 m length cut from a 20 m x 0.18 mm x 0.18 μ m film thickness DB-35ms Ultra Inert (UI) capillary column (p/n 121-3822UI). This column was installed between the inlet and one end of the purged union. It was back flushed shortly before the run had ended to prevent high boiling point contaminants from entering the second column. The second column was a 15 m x 0.25 mm i.d. 0.25 μ m film thickness DB-5MS UI capillary column (p/n 19091S-431UI). The column was installed between the other end of the purged union and the ICP-QQQ transfer line connection inside the GC oven. Sample injections of 1 μ L volume were made under splitless conditions with the inlet held at 280 °C. GC operating parameters are detailed in a previous study [1].

ICP-QQQ: O_2 mass-shift method was applied to detect P and S. The O_2 flow rate was 0.2 mL/min. P and S were detected as PO⁺ and SO⁺, respectively.

Samples and preparation: Three standard pesticide mixes were obtained from Ultra Scientific (Kingstown, RI, USA) and Agilent Technologies (p/n 5190–0468). The standard solutions were diluted with high purity grade acetonitrile to form intermediate solutions. These solutions were then used to prepare calibration standard solutions following serial dilutions in acetonitrile.

Results and Discussion

Figure 1 shows overlaid chromatograms for P and S in a mixed pesticide standard. The pesticides that contain more than one hetero-element can be identified easily.

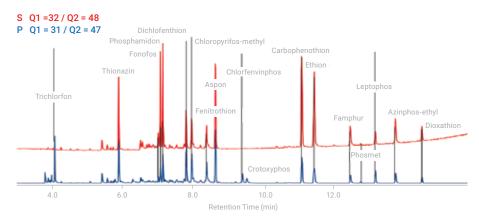


Figure 1. Chromatograms showing the heteroatom traces for P and S in the mixed pesticide standard, with identified pesticide compounds. Reprinted with permission from J. Agric. Food Chem., 2015, 63, 4478–4483. Copyright 2015

American Chemical Society.

Table 1 summarizes the retention time and compound detection limits (DLs) of the pesticides. DLs for pesticides using current GC/MS/MS instrumentation typically vary from about 0.1 to 10 μ g/L depending on the pesticide and instrument used. The data in Table 1 suggests that GC-ICP-QQQ offers similar or slightly lower DLs than GC/MS/MS for the determination of organophosphorus pesticides. For S-containing pesticides, detection limits are similar to, or slightly higher than DLs achieved by GC/MS/MS. All of the pesticides listed in Table 1, which were detected via their P content, were quantified well below the 10 μ g/L limit of quantitation (LOQ) required by most food safety laboratories.

Table 1. Detection limits as the compound for pesticides.

Pesticide	RT (min)	Compound	d DL (µg/L)
		Р	S
Trichlorfon	4.103	0.178	
Thionazin	5.926	0.221	11.93
Terbufos	7.071	0.718	9.708
Fonofos	7.185	0.455	7.917
Phosphamidon	7.299	0.923	
Dichlofenthion	7.858	0.362	15.8
Chloropyrifos-methyl	7.973	0.613	24.18
Fenitrothion	8.44	0.907	19.52
Aspon	8.705	0.200	9.912
Chlorfenvinphos	9.486	2.020	
Crotoxyphos	9.541	3.338	
Carbophenothion	11.158	0.583	9.585
Ethion	11.527	0.707	11.51
Famphur	12.547	2.206	20.61
Phosmet	12.851	3.829	
Leptophos	13.263	1.125	18.35
Azinphos-ethyl	13.827	1.812	21.33
Dioxathion	14.587	1.392	7.84

Conclusions

The GC-ICP-QQQ method is suitable for the selective and sensitive detection of organophosphorus and organosulfur pesticides by measurement of their heteroatoms. Due to the significantly lower background of the Agilent 8800 ICP-QQQ, GC-ICP-QQQ provides good sensitivity performance for the determination of organophosphorus pesticides compared to GC/MS/MS.

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More Information

Determination of pesticides in foods using phosphorus and sulfur detection by GC-ICP-QQQ, Agilent publication, <u>5991-6260EN</u>.

Benefits of the Agilent 8900 ICP-QQQ with MS/MS operation for routine food analysis

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Introduction

Growing awareness of and concern about the issue of food safety is reflected in the tightening of regulations governing toxic elements and compounds in food. Many toxic elements such as As, Hg, Cd, Pb etc. are routinely monitored to ensure food safety, while minerals that are beneficial/essential to human health such as Se, Na, Mg, K, Ca, etc., are also measured. As a fast, high throughput, multi-element technique, with a wide dynamic range and high sensitivity, ICP-MS is increasingly used for routine food analysis. Recent improvements in matrix tolerance with Agilent's High Matrix Introduction (HMI/UHMI) technology is a further benefit for the application as food matrices are varied and can be complex. UHMI uses aerosol dilution to reduce the sample matrix load on the plasma, allowing matrix levels up to several percent total dissolved solids (TDS) to be analyzed routinely. This is much higher than the limit of 0.2% (2000 ppm) which has traditionally applied to samples intended for ICP-MS analysis.

Control of polyatomic ion interferences in quadrupole ICP-MS has also improved significantly with the development of collision/reaction cells (CRCs), which use kinetic energy discrimination (KED) to attenuate polyatomic ions in helium (He) collision mode. Agilent's octopole-based CRC, the ORS⁴, is routinely used to suppress a wide range of matrix-based polyatomic ion interferences under one set of cell conditions [1]. Hence, reliable and accurate quantification of all required elements at regulated levels in a variety of sample matrices is now possible using conventional quadrupole ICP-MS (ICP-QMS).

However, some food-analysis applications require greater sensitivity for specific elements, while some complex sample matrices may cause spectral interferences that remain a challenge for ICP-QMS. For example, doubly charged ions of some rare earth elements (REEs) appear at the same mass as key analytes, hindering accurate low-level measurement of arsenic (As) and selenium (Se) in some sample types [2, 3].

Improved interference removal with ICP-QQQ

The Agilent 8900 Triple Quadrupole ICP-MS (ICP-QQQ) has a unique tandem MS configuration, comprising two scanning quadrupole mass analyzers either side of an octopole-based ORS⁴ collision reaction cell. As a result, the 8900 ICP-QQQ is able to utilize reactive cell gases and ion/molecule reaction chemistry in combination with MS/MS mode to resolve difficult spectral interferences [4]. The superior interference removal offered by reaction chemistry with MS/MS led to the previous generation Agilent 8800 ICP-QQQ being widely accepted in industry and research labs in fields such as semiconductor device and high purity chemical/material manufacturing, life-science, geoscience, radionuclides and many others [5-8]. MS/MS mode is also beneficial for the analysis of certain elements which are subject to problematic interferences in routine applications, such as the analysis of food samples, soils, waste water and groundwater. Since the matrix tolerance and robustness of the Agilent 8900 ICP-QQQ is

comparable to Agilent's market-leading single quadrupole ICP-MS systems, the 8900 ICP-QQQ can be used to analyze these high-matrix samples routinely.

Solving problems associated with As and Se analysis

Arsenic is a well-known toxic element, while Se is an essential element that can be toxic in excess. Consequently, many countries regulate the permitted concentrations of As and Se in food, animal feed, drinking water, surface water and soils. However, As and Se can suffer spectral interferences from polyatomic ions including ArCl+, CaCl+, ArAr+, $\rm S_2O^+, SO_3^+, GeH^+, and BrH^+.$ These interferences can be reduced using ICP-QMS operating in helium (He) cell mode, allowing the accurate and precise measurement of As and Se at the concentration levels required to meet typical regulatory demands.

However, He mode is not effective against doubly-charged ion overlaps. The lanthanides or rare earth elements (REE) can form doubly charged ions (REE++) which overlap As and Se. These doubly-charged overlaps can be avoided using mass-shift mode with O_2 as the reaction cell gas. In this mode, the analytes are measured as reaction product ions $^{75}\text{As}^{16}\text{O}^+$ and $^{78}\text{Se}^{16}\text{O}^+$, mass-shifted to m/z 91 and 94 respectively, where they are free from the original REE++ overlaps. This reaction chemistry can be used in the CRC of an ICP-QMS, but existing ions from the plasma may overlap the newly-formed product ions; e.g. $^{91}\text{Zr}^+$ on $^{75}\text{As}^{16}\text{O}^+$, and $^{94}\text{Mo}^+$ on $^{78}\text{Se}^{16}\text{O}^+$. To ensure controlled and consistent reaction chemistry, MS/MS mode on an ICP-QQQ is required, where the first quadrupole (Q1) operates as a mass filter set to the appropriate As+ or Se+ precursor ion mass. Q1 rejects all other masses, thereby removing the existing Zr+ and Mo+ ions and preventing them from overlapping the new analyte product ions.

Typically, the REE content of food and other natural samples is low, but crops grown in REE-enriched soils can take up high concentrations of these elements. The use of MS/MS mode with $\rm O_2$ reaction cell gas avoids the potential risk of reporting incorrect results for As and Se in the case of an unexpectedly high level of REEs.

In this study, the Agilent 8900 ICP-QQQ was evaluated as a routine tool for the analysis of 30 elements, including As and Se, in food sample digests.

Experimental

Certified Reference Materials (CRMs)

Five food CRMs purchased from National Institute of Standards and Technology (NIST) and High-Purity Standards Inc. (Charleston, SC, USA) were analyzed in this study. The CRMs used were NIST 1567b Wheat Flour, NIST 1568b Rice Flour, NIST 1515 Apple Leaves, NIST 1573a Tomato Leaves and High Purity Standards Mixed Food Diet Solution.

Sample preparation

Due to the requirement to measure several volatile elements, including Hg, closed vessel microwave digestion using a Milestone ETHOS 1 Advanced Microwave Digestion System was used to digest the food CRMs. Sample weights of approximately 1.0 g for each of the flour CRMs (NIST 1567b, NIST 1568b) and 0.5 g for each of the other sample types (NIST 1515, NIST 1573a) were accurately weighed into closed microwave vessels. 6 mL of HNO $_3$ and 1 mL of HCl (electronics (EL) grade acids, Kanto Chemicals) were added to the microwave

vessels. After 15 minutes held at room temperature, microwave heating was applied, using the heating program shown in Table 1. All CRMs were completely dissolved, resulting in clear solutions which were diluted to a final volume of 100 mL with ultrapure water (Merck, Darmstadt, Germany).

Table 1. Microwave digestion heating programs for four CRM food samples.

Power (W)	Temp (°C)	Ramp (min)	Hold (min)
500	70	2	3
1000	140	5	5
100	200	5	15
	Ventilation		30

It is well known that carbon present in the sample solution enhances the ICP-MS signal of some elements, notably As, Se and P, although the precise mechanism of the enhancement is not clearly understood [9, 10]. With the high digestion temperature used in this work (200 °C), the carbon matrix was effectively decomposed during digestion. If any residual carbon did remain its effect could be mitigated by adding 2% butan-1-ol online with the internal standard solution.

Instrumentation

An Agilent 8900 ICP-QQQ (Standard configuration) with the standard sample introduction system consisting of a glass concentric nebulizer, quartz spray chamber, and Ni interface cones was used. UHMI technology is included on the 8900 ICP-QQQ Standard configuration, allowing matrices as high as 25% NaCl solution to be analyzed [11]. The plasma conditions were selected according to the sample type and expected matrix level using the "Preset plasma" function of the MassHunter software.

Acquisition conditions

For the multi-element analysis of the food samples, a multi-tune method was used so all elements could be acquired in the optimum cell gas mode. Multi-tune permits samples to be automatically analyzed using the optimum tune and cell conditions for each analyte element. He mode was used for all elements except P, S, As and Se which were determined in mass-shift mode using $\rm O_2$ cell gas. The method was based on an appropriate preset method for food samples, which was modified to include $\rm O_2$ cell gas mode. Preset plasma condition "UHMI-4" was selected, where the number 4 represents the approximate aerosol dilution factor. The UHMI setting automatically applies the predefined and calibrated parameters for RF power, sampling depth, carrier gas flow rate and dilution gas flow rate, giving precise and reproducible plasma conditions for the target sample types. The lens voltages were auto-tuned for maximum sensitivity. Table 2 summarizes the instrument operating parameters.

Table 2. Agilent 8900 ICP-QQQ operating conditions.

Parameter		Setting
Cell mode	He mode	O ₂ mode
Scan type	Single Quad	MS/MS
Plasma conditions	ι	JHMI-4
RF power (W)		1600
Sampling depth (mm)		10
Carrier gas flow rate (L/min)		0.77
Dilution gas flow rate (L/min)		0.15
Extract 1 (V)		0
Extract 2 (V)		-250
Omega Bias (V)		-140
Omega lens (V)		8.8
Cell gas flow (mL/min)	5.5	0.3 (20% of full scale)
KED (V)	5	-7

Bold parameters are predefined by selecting preset plasma condition UHMI-4.

Calibration standards and internal standards

Calibration standards were prepared from an Agilent multi-element environmental calibration standard (p/n 5183-4688) which contains 1000 ppm each of Fe, K, Ca, Na, Mg and 10 ppm each of Ag, Al, As, Ba, Be, Cd, Co, Cr, Cu, Mn, Mo, Ni, Pb, Sb, Se, Th, Tl, U, V, and Zn. Standards for B, Rb, Sr, Sn and Hg were prepared from 1000 ppm Atomic-Absorption grade single element standards from Kanto Chemicals (Tokyo, Japan). S and P were prepared from 10,000 ppm Spex single element standards (SPEX CertiPrep, NJ, USA). The internal standard (ISTD) solution was prepared from an Agilent internal standard stock solution for ICP-MS systems (p/n 5188-6525) containing 6-Li, Sc, Ge, Rh, In, Tb, Lu, and Bi. Ir was added from an Atomic-Absorption grade single element standard purchased from Kanto Chemicals. The ISTD was added to the sample using the standard online ISTD kit.

Calibration standards were prepared in 6% HNO $_3$ and 1% HCl to match the acid content of the sample solutions. The ISTDs were prepared in 1% HNO $_3$ and 0.5% HCl. The calibration ranges were as follows: major elements: 0-100 ppm, trace elements: 0-500 ppb, B: 0-200 ppb, Hg: 0-1 ppb and Sn: 0-2 ppb.

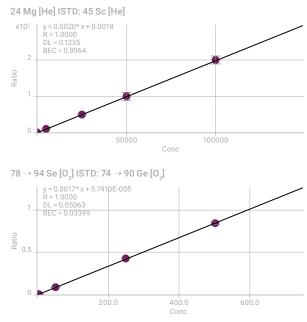


Figure 1. Representative calibration curves for a major element (Mg) and a trace element (Se).

Sequence of calibrants, samples, and QC solutions

The sequence consisted of an initial multi-level calibration, covering the typical range for the target analytes, followed by a QC block containing an Initial Calibration Blank (ICB) check and Initial Calibration Verification (ICV) solution. After calibration and initial QC check, twelve sample blocks were analyzed per the flow chart shown in Figure 2; each block consisted of 2 preparation blanks and 10 samples (2 each of Wheat Flour, Rice Flour, Apple Leaves, Tomato Leaves and Mixed Food Diet). A Periodic Block consisting of Continuing Calibration Blank (CCB) and Continuing Calibration Verification (CCV) samples was automatically inserted into the sequence after each sample block.

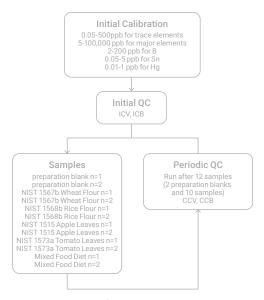


Figure 2. Sequence of calibrants, samples, and QC solutions analyzed in a single 15-hour sequence. Sample Block was repeated continuously with automatic insertion of Periodic QC Block after each Sample Block.

The total number of analyses of calibration standards, QC samples and food digest samples was 183 over \sim 15 hours. The sample-to-sample run time was about 5 minutes, which included a 10 s probe rinse and 60 s sample introduction system rinse at 0.3 rps peristaltic pump rate. Table 3 shows the detection limits (DL) obtained using this method.

Table 3. Method detection limits.

Element	Scan Mode	Q1	Q2	DI (ppb)
В	Single Quad		11	0.3653
Na	Single Quad		23	0.1945
Mg	Single Quad		24	0.1235
Al	Single Quad		27	0.1847
P	MS/MS	31	47	0.0919
S	MS/MS	32	48	0.4367
K	Single Quad		39	7.0656
Ca	Single Quad		44	8.7579
V	Single Quad		51	0.0079
Cr	Single Quad		52	0.0880
Mn	Single Quad		55	0.0099
Fe	Single Quad		56	0.1595
Со	Single Quad		59	0.0009
Ni	Single Quad		60	0.0484
Cu	Single Quad		63	0.0102
Zn	Single Quad		66	0.0308
As	Single Quad		75	0.0044
As	MS/MS	75	91	0.0040
Se	Single Quad		78	0.3158
Se	MS/MS	78	94	0.0506
Rb	Single Quad		85	0.0115
Sr	Single Quad		88	0.0006
Мо	Single Quad		95	0.0090
Ag	Single Quad		107	0.0063
Cd	Single Quad		111	0.0018
Sn	Single Quad		118	0.0074
Sb	Single Quad		121	0.0026
Ва	Single Quad		138	0.0008
Hg	Single Quad		202	0.0005
TI	Single Quad		205	0.0104
Pb	Single Quad		208	0.0016
Th	Single Quad		232	0.0018
U	Single Quad		238	0.0009
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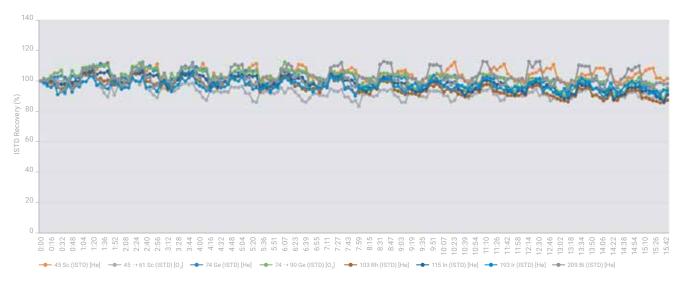


Figure 3. ISTD signal stability for the sequence of 183 samples analyzed over 15 hours.

Results and Discussion

ISTD and CCV stability

Figure 3 shows the ISTD signal stability for the sequence of 183 samples analyzed over 15 hours. The ISTD recoveries for all samples were well within ±20 % of the value in the initial calibration standard. These ISTD recoveries are comparable to the results obtained routinely using ICP-QMS, demonstrating the equivalent robustness of the 8900 ICP-QQQ.

The midpoint of the calibration standards was used as the CCV solution. CCV recovery over the 15-hour analysis was stable and within $\pm 10\%$ for all elements, as shown in Figure 4, again demonstrating that the 8900 ICP-QQQ has the high matrix tolerance required for routine food digest analysis.

CRM recovery results

The accuracy of the method was evaluated by analyzing the five food CRMs as unknown samples. Each CRM was measured 24 times in the batch. The mean concentration and relative standard deviation (%RSD) were calculated for each element and compared to the certified value, as shown in Tables 4 to 8. Using the preferred measurement mode, the results for all elements were in good agreement with the certified and reference values. Results are shown for both He mode and $\rm O_2$ mass-shift mode for As and Se, to compare the results for samples where a sample might contain an unexpected high level of REEs. NIST 1515 Apple Leaves CRM contains low $\rm \mu g/kg$ concentrations of As and Se (Table 6) and high concentrations of REEs. Reference (non-certified) values for Nd, Sm, and Gd are 17, 3, and 3 mg/kg, respectively. In the case of Apple Leaves and, to a lesser extent, Tomato Leaves, more accurate recovery was obtained for As and Se using $\rm O_2$ mass shift mode, illustrating the potential error that can be caused by the relatively high level of REE in these two reference materials.

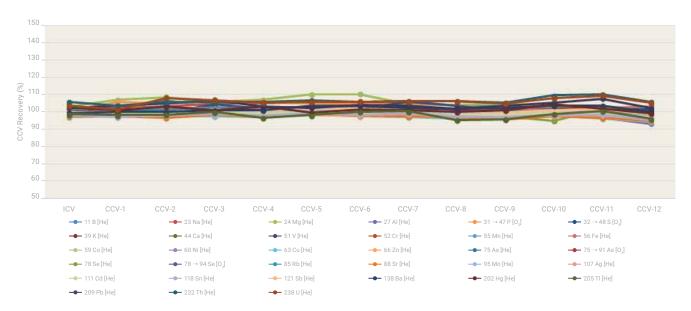


Figure 4. CCV recovery for all elements over the 15-hour analysis.

Table 4. Results for NIST 1567b Wheat Flour, n = 24.

Element	Measured Solution Concentration (μg/L)	RSD (%)		nple ng/kg)	Cert	Recovery (%)			
23 Na	65.2	2.3	6.50	±	0.15	6.71	±	0.21	97
24 Mg	3842	1.6	383	±	6	398	±	12	96
27 Al	39	2.8	3.9	±	0.1	4.4	±	1.2	88
31 -> 47 P	12936	2.0	1291	±	26	1333	±	36	97
32 -> 48 S	15496	2.2	1546	±	34	1645	±	25	94
39 K	12700	2.3	1267	±	29	1325	±	20	96
44 Ca	1871	1.8	186.7	±	3.4	191.4	±	3.3	98
51 V	0.10	8.1	0.010	±	0.001		0.01*		100
55 Mn	86	1.7	8.54	±	0.14	9.00	±	0.78	95
56 Fe	142	1.6	14.20	±	0.22	14.11	±	0.33	101
63 Cu	19	1.6	1.94	±	0.03	2.03	±	0.14	96
66 Zn	112	1.9	11.17	±	0.21	11.61	±	0.26	96
75 As	0.047	16.5	0.0046	±	0.001	0.0048	±	0.0003	97
75 -> 91 As	0.049	19.4	0.0049	±	0.001	0.0048	±	0.0003	101
78 Se	11.5	4.2	1.15	±	0.05	1.14	±	0.10	101
78 -> 94 Se	11.8	1.9	1.17	±	0.02	1.14	±	0.10	103
85 Rb	6.54	1.8	0.652	±	0.012	0.671	±	0.012	97
95 Mo	4.60	2.1	0.459	±	0.009	0.464	±	0.034	99
111 Cd	0.239	5.7	0.0238	±	0.0014	0.0254	±	0.0009	94
118 Sn	0.0355	12.8	0.0035	±	0.0005		0.003*		118
202 Hg	0.0066	11.3	0.0007	±	0.0001		0.0005*		131
208 Pb	0.0937	4.4	0.0094	±	0.0004	0.0104	±	0.0024	90

^{*} Reference value

Table 5. Results for NIST 1568b Rice Flour, n = 24.

Element	Measured Solution Concentration (μg/L)	RSD (%)	RSD (%) Calculated Sample Concentration (mg/kg)			Certif	Recovery (%)		
23 Na	65.6	3.2	6.54	±	0.28	6.74	±	0.19	97
24 Mg	5454	1.5	543	±	8	559	±	10	97
27 Al	40.3	3.3	4.01	±	0.13	4.21	±	0.34	95
31 -> 47 P	15162	2.8	1510	±	43	1530	±	40	99
32 -> 48 S	11369	2.5	1133	±	28	1200	±	10	94
39 K	12371	2.0	1233	±	24	1282	±	11	96
44 Ca	1158	2.1	115.3	±	2.5	118.4	±	3.1	97
51 V	182.3	1.0	18.2	±	0.2	19.2	±	1.8	95
55 Mn	75.4	1.0	7.51	±	0.08	7.42	±	0.44	101
56 Fe	0.173	1.7	0.0173	±	0.0003	0.0177	±	0.0005*	98
63 Cu	22.7	1.0	2.26	±	0.02	2.35	±	0.16	96
66 Zn	191.7	1.4	19.10	±	0.26	19.42	±	0.26	98
75 As	2.97	1.4	0.296	±	0.004	0.285	±	0.014	104
75 -> 91 As	3.01	1.7	0.300	±	0.005	0.285	±	0.014	105
78 Se	3.4	8.9	0.341	±	0.030	0.365	±	0.029	93
78 -> 94 Se	3.5	3.8	0.352	±	0.013	0.365	±	0.029	96
85 Rb	61.1	1.1	6.088	±	0.069	6.198	±	0.026	98
95 Mo	13.96	1.2	1.391	±	0.017	1.451	±	0.048	96
111 Cd	0.201	4.9	0.0201	±	0.0010	0.0224	±	0.0013	90
118 Sn	0.060	7.4	0.0060	±	0.0004	0.005	±	0.001*	121
202 Hg	0.0529	2.1	0.0053	±	0.0001	0.0059	±	0.0004	89
208 Pb	0.068	3.0	0.0068	±	0.0002	0.008	±	0.003*	85

^{*} Reference value

Table 6. Results for NIST 1515 Apple Leaves, n = 24.

Element	Measured Solution Concentration (μg/L)	RSD (%) Calculated Sample Concentration (mg/kg)				Certi	Recovery (%)		
11 B	141	2.9	28	±	0.8	27	±	2	104
23 Na	196	1.6	39.1	±	0.6	24.4	±	1.2	160*1
24 Mg	14083	1.3	2812	±	36	2710	±	80	104
27 AI	1458	1.6	291	±	5	286	±	9	102
31 -> 47 P	8088	2.2	1615	±	35		1590*		102
32 -> 48 S	9211	1.4	1839	±	26		1800*		102
39 K	80429	2.2	16057	±	361	16100	±	200	100
44 Ca	74060	1.2	14786	±	172	15260	±	1500	97
51 V	1.20	2.8	0.24	±	0.01	0.26	±	0.03	92
52 Cr	1.3	1.4	0.25	±	0.00	0.3*			85
55 Mn	265	1.0	53	±	1	54	±	3	98
56 Fe	379	0.8	76	±	1		80*		95
59 Co	0.44	1.5	0.088	±	0.001		0.09*		98
60 Ni	4.4	1.7	0.88	±	0.02	0.91	±	0.12	97
63 Cu	28.2	1.0	5.62	±	0.06	5.64	±	0.24	100
66 Zn	60.3	0.9	12.0	±	0.1	12.5	±	0.3	96
75 As	2.0	1.2	0.395	±	0.005	0.038	±	0.007	1040
75 -> 91 As	0.2	3.7	0.036	±	0.001	0.038	±	0.007	94
78 Se	13.43	5.8	2.7	±	0.2	0.050	±	0.009	5364
78 -> 94 Se	0.271	13.8	0.054	±	0.008	0.050	±	0.009	108
85 Rb	46.3	0.9	9.2	±	0.1		9*		103
88 Sr	123.0	1.0	25	±	0	25	±	2	98

Element	Measured Solution Concentration (μg/L)	RSD (%)		culated Sar entration (n	•	Certified Concentration (mg/kg)			Recovery (%)
95 Mo	0.44	5.3	0.088	±	0.005	0.094	±	0.013	94
111 Cd	0.06	7.0	0.013	±	0.001		0.014*		91
121 Sb	0.06	4.6	0.011	±	0.001		0.013*		85
138 Ba	245	1.9	49	±	1	49	±	2	100
202 Hg	0.21	2.0	0.041	±	0.001	0.044	±	0.004	93
208 Pb	2.3	1.3	0.452	±	0.006	0.470	±	0.024	96
232 Th	0.14	2.2	0.028	±	0.001		0.03*		93
238 U	0.034	3.7	0.0068	±	0.0003		0.006*		113

^{*}Reference value.

Bold values for As and Se were obtained in single quad mode with He cell gas. The accurate results obtained using MS/MS mode with O_2 mass-shift are shown in the lines below.

Table 7. Results for NIST 1573a Tomato Leaves, n = 24.

Element	Measured Solution Concentration (μg/L)	RSD (%)		culated Sar entration (n	•	Certi	ified Concentr (mg/kg)	ation	Recovery (%)
11 B	167	1.9	33.3	±	0.6	33.3	±	0.7	10
23 Na	613	2.5	122	±	3	136	±	4	90
24 Mg	57311	2.0	11412	±	225		12000*		95
27 AI	2573	2.4	512	±	12	598	±	12	86
31 -> 47 P	10928	2.7	2176	±	59	2160	±	40	101
32 -> 48 S	48387	1.4	9635	±	131		9600*		100
39 K	134250	2.2	26732	±	591	27000	±	500	99
44 Ca	243939	1.4	48574	±	671	50500	±	900	96
51 V	4.0	2.2	0.792	±	0.017	0.835	±	0.010	95
52 Cr	9.3	1.6	1.85	±	0.03	1.99	±	0.06	93
55 Mn	1236.5	1.5	246	±	4	246	±	8	100
56 Fe	1843.3	1.7	367	±	6	368	±	7	100
59 Co	2.8	1.4	0.55	±	0.01	0.57	±	0.02	96
60 Ni	7.9	1.9	1.56	±	0.03	1.59	±	0.07	98
63 Cu	23.7	1.5	4.71	±	0.07	4.70	±	0.14	100
66 Zn	149.4	1.5	29.8	±	0.5	30.9	±	0.7	96
75 As	0.7	2.3	0.141	±	0.003	0.112	±	0.004	126
75 -> 91 As	0.6	1.7	0.112	±	0.002	0.112	±	0.004	100
78 Se	1.03	15.6	0.205	±	0.032	0.054	±	0.003	380
78 -> 94 Se	0.31	11.2	0.061	±	0.007	0.054	±	0.003	113
85 Rb	69.7	1.2	13.88	±	0.16	14.89	±	0.27	93
88 Sr	421.0	1.3	84	±	1		85*		99
95 Mo	2.1	2.8	0.42	±	0.01		0.46*		91
107 Ag	0.09	9.1	0.018	±	0.002		0.017*		104
111 Cd	7.4	1.4	1.47	±	0.02	1.52	±	0.04	97
121 Sb	0.28	3.4	0.055	±	0.002	0.063	±	0.006	88
138 Ba	302.8	2.1	60.3	±	1.3		63*		96
202 Hg	0.15	2.4	0.030	±	0.001	0.034	±	0.004	88
232 Th	0.52	2.1	0.104	±	0.002		0.12*		87
238 U	0.14	2.3	0.029	±	0.001		0.035*		81

^{*} Reference value

Bold values for As and Se were obtained in single quad mode with He cell gas. The accurate results obtained using MS/MS mode with O2 mass-shift are shown in the lines below.

^{*1} The measured Na result was high compared to the reference value; the same result was obtained from a repeated analysis of the same solution, so a spike recovery test was performed for confirmation. The spike recovery result was good (recovery: 99%), suggesting that the original sample had suffered Na contamination.

Table 8. Results for the High Purity Standard Mixed Food Diet Solution, n = 24.

Element	Measured Solution Concentration (μg/L)	RSD (%)		culated Sar entration (n	•	Cert	ified Concentr (mg/kg)	ation	Recovery (%)
23 Na	15808	2.9	61.8	±	1.8	60.0	±	0.6	105
24 Mg	3300	2.3	12.9	±	0.3	12.0	±	0.1	108
27 Al	26	4.5	0.100	±	0.005	0.100	±	0.002	100
31 -> 47 P	15543	3.3	60.8	±	2.0	60.0	±	0.6	101
39 K	41898	2.2	164	±	4	160	±	2	102
44 Ca	9800	2.7	38.3	±	1.0	40.0	±	0.4	96
52 Cr	0.55	10.4	0.0021	±	0.0002		0.002*		107
55 Mn	49.2	1.7	0.192	±	0.003	0.200	±	0.004	96
56 Fe	204.5	1.8	0.80	±	0.01	0.80	±	0.01	100
59 Co	0.2	2.4	0.0008	±	0.0000		0.0008*		98
60 Ni	5.1	2.5	0.020	±	0.001	0.020	±	0.001	99
63 Cu	15.3	1.7	0.060	±	0.001	0.060	±	0.006	100
66 Zn	74.5	2.0	0.29	±	0.01	0.30	±	0.01	97
75 As	5.1	2.0	0.020	±	0.000	0.020	±	0.001	99
75 -> 91 As	5.2	2.6	0.020	±	0.001	0.020	±	0.001	102
78 Se	1.26	14.8	0.0049	±	0.0007		0.005*		99
78 -> 94 Se	1.31	6.6	0.0051	±	0.0003		0.005*		102
95 Mo	1.5	3.1	0.0059	±	0.0002		0.006*		98
111 Cd	2.0	2.1	0.0078	±	0.0002	0.0080	±	0.0008	98

^{*} Reference value

Bold values for As and Se were obtained in single quad mode with He cell gas. The accurate results obtained using MS/MS mode with O₂ mass-shift are shown in the lines below.

Conclusions

The Agilent 8900 Standard configuration ICP-QQQ with UHMI offers the robustness and matrix tolerance required for the routine analysis of the widest range of trace and major elements in high matrix samples, such as food digest samples. Doubly-charged REE interferences that can affect the accurate measurement of arsenic and selenium at trace levels were avoided using $\rm O_2$ cell gas with MS/MS mass-shift mode. Most other elements were measured in He mode; a field-proven method that is widely used to remove common matrix-based polyatomic interferences in complex and variable matrices.

While not all food products, soils and sediments contain significant concentrations of REEs, the use of ICP-QQQ with MS/MS improves the accuracy and confidence in the results for As and Se measured in food and environmental samples that often contain complex, variable, high TDS matrices.

Method development was greatly simplified with the use of Pre-set Methods and auto tuning, which ensures reproducible performance irrespective of operator experience.

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Speciated Arsenic Analysis in Wine Using HPLC-ICP-QQQ

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Validation of an extended FDA Elemental Analysis
Manual method

Introduction

In 2013, the US Food and Drug Administration (FDA) released Elemental Analysis Manual (EAM) Method §4.10. The method describes the Determination of Four Arsenic Species in Fruit Juice using High-Performance Liquid Chromatography-Inductively Coupled Plasma-Mass Spectrometry [1]. To extend the method to include wine, a multi-laboratory validation (MLV) of the method was carried out with three US-based laboratories sharing their data [2]. The data shown in this application note is supplementary to the published data. In addition to the paper, this note includes long term stability of the method, and extended quantitative analysis of five commercially available wines. The method required separation and analysis of all target species. This approach differs from another Agilent application note, which focused on the development of a fast method for inorganic arsenic (iAs) [3].

The US Environmental Protection Agency (EPA) set a maximum threshold of total As in drinking water of 10 μ g/kg [4]. There is no equivalent US regulation for As in wine. Studies have shown that As in wine can be the result of an accumulation of As in the grapes from the environment [5] or introduced during the wine making process [6].

Regulations in Canada (Vintners Quality Alliance VQA, Ontario) and Europe (International Organisation of Vine and Wine, OIV) specify limits for total As of $100~\mu g/L$ and $200~\mu g/L$, respectively [7, 8]. However, the toxicity of As is determined by its chemical form. Because the inorganic forms of As (iAs) are the most carcinogenic, the FDA has established an action limit for iAs in apple juice of $10~\mu g/kg$ in 2013 [9]. FDA EAM Method §4.10 details a relatively simple and robust method for the determination of As species in fruit juice using HPLC-ICP-MS [1]. The method describes a procedure to determine iAs (the sum of arsenite, As(III), and arsenate, As(V)); dimethylarsinic acid (DMA); and monomethylarsonic acid (MMA). The method also states that a solution containing arsenobetaine (AB) and As(III) is analyzed to demonstrate adequate separation between unretained arsenic-containing species and As(III).

Due to recent media attention on As levels in wine, and the lack of published research on As speciation in wine, extension of EAM §4.10 to include wine is a logical next step.

In this study, EAM §4.10 was modified for the determination of the main organic arsenic species (DMA and MMA) and the more toxic inorganic forms (As(V) and As(III)) in wine using HPLC coupled to a triple quadrupole ICP-MS (ICP-QQQ). The ICP-QQQ was utilized to provide the highest possible sensitivity of all the instruments available in the lab at UC Davis. ICP-QQQ also provides superior resolution of potential spectral interferences, but the potential Cl-based interferences on ⁷⁵As are resolved chromatographically, so QQQ with MS/MS is not essential. This application could also be done on a single quadrupole ICP-MS such as the Agilent 7800 or 7900.

Experimental

Reagents

Arsenite (As(III)) and arsenate (As(V)) were bought as 1000 mg/L standard solutions from Spex Certiprep (Metuchen, NJ, USA). Monomethylarsonic acid (MMA, 98.5% purity) and dimethylarsinic acid (DMA, 98.9% purity) were bought from Chem Service (West Chester, PA, USA). Arsenobetaine (AB, purum p.a., ≥95.0%) was bought from Fluka Analytical (Morris Plains, NJ, USA).

Samples and sample preparation

Five commercially available wine samples were bought from a local store in Davis, California. The wines were selected to represent the main types (and styles) of wine: red (Cabernet Sauvignon), white (Sauvignon blanc), rosé (Zinfandel), sparkling (sparkling white) and fortified (Port-style). To investigate the range of ethanol content that could be analyzed using the method, the alcohol concentrations of the wines selected ranged from 9.5–20% (v/v).

The sample preparation and analysis details were carried out according to the EAM §4.10 method. Each wine sample was diluted five times with de-ionized water and then filtered separately using syringe-filtration (0.45 μ m PVDF membrane).

Per EAM §4.10, calibration curves were prepared at nominal concentrations of 0.4, 0.5, 1, 5, 10, 20, 40 μ g/kg for the four arsenic species: As(III), DMA, MMA, and As(V). However, for this method, a fifth, low-level calibration point was also prepared at 0.1 μ g/kg. NIST 1643e Trace Elements in Water standard reference material (SRM), used to assess recovery and stability, was prepared using a 15-fold dilution. All calibration standards and the SRM were prepared in a 3% ethanol solution to approximately match the level of alcohol (carbon matrix) in the diluted wine samples. In addition to the effect that a change in sample viscosity has on sample transport and nebulization, the level of carbon also affects (increases) the degree of ionization of some elements in the ICP, including arsenic. Therefore, sample preparation for carbon-containing matrices should ensure a reasonably consistent level of carbon across all samples and standards, to avoid errors due to variable carbon enhancement in different sample solutions.

Instrumentation

An Agilent 1260 Infinity LC comprising a binary pump, autosampler, and vacuum degasser was coupled to an Agilent 8800 Triple Quadrupole ICP-MS (ICP-QQQ). HPLC and ICP-QQQ parameters are shown in Table 1.

Table 1. HPLC-ICP-QQQ hardware system and operating conditions.

LC conditions	Value
Column	Hamilton PRP-X100 anion exchange (4.1 x 250 mm) column with a matching Hamilton PRP-X100 guard column
Mobile phase	Mobile phase, aqueous 10 mM ammonium phosphate dibasic, 1% ethanol, pH 8.25 (±0.05)
Flow rate (mL/min)	1.0
Temperature	Ambient
Injection volume (µL)	100
Column compartment time table for introduction of ISTD	0.1 min, column position 1, 1.0 min; switch to column position 2, 2.0 min; switch back to column position 1
ICP-QQQ parameters	Value
RF power (W)	1550
Carrier gas flow (L/min)	1.0
Spray chamber temperature (°C)	2
Sample depth (mm)	8.5
Peristaltic pump speed (rps)	0.3 (~1.2 mL/min)
Scan mode	MS/MS
Helium cell gas flow (mL/min)	~2.0

Results and Discussion

Method blanks (3% ethanol) spiked with low levels of As(III), DMA, MMA, and As(V) were prepared and analyzed for the determination of the detection limits.

Figure 1 shows overlaid chromatograms obtained for the mixed As species standards, demonstrating excellent peak separation of the As species of interest. The calibration curves in Figure 2 show a linear response for each As species across the concentration range from 0.1 to 40 μ g/kg.

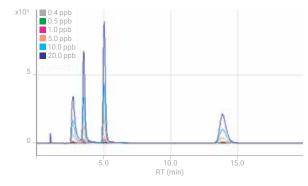


Figure 1. Overlaid chromatograms of As species standards at nominal concentrations of 0.4, 0.5, 1, 5, 10, 20 μ g/kg showing good peak separation. The 40 μ g/kg standard is not shown, to allow the lower concentration levels to be seen.

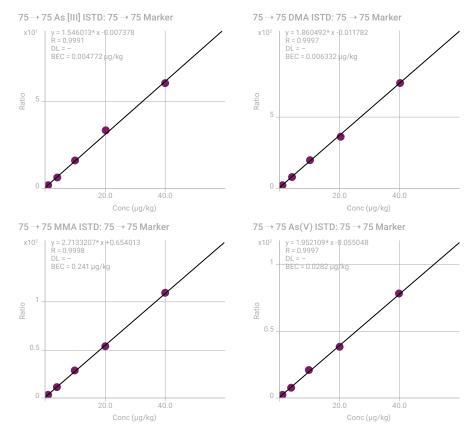


Figure 2. Calibration graphs for As(III), DMA, MMA, and As(V).

The limits of detection (LOD) for the As species in wine were calculated as described in the FDA's Elemental Analysis Manual Section 3.2 [1]. The limits of quantification (LOQ) for each species were calculated as LOQ = Dilution Factor (DF) x 30 x σ . The LOQs for As(III) and As(V) were 1.18 and 1.35 μ g/kg, respectively. The LOQ for total inorganic arsenic (calculated from the SD of the sum of the integrated peak areas for As(III) and As(V) in each repeat of the low standard) was 2.53 μ g/kg. The LODs and LOQs determined for the species DMA, MMA, and total iAs (sum of As(III) and As(V)) using the optimized method are given in Table 2. Results are reported for iAs since the current regulations only specify iAs, and not the individual species As(III) and As(V).

Table 2. LODs and LOQs for DMA, MMA, and iAs.

	LOD, µg/kg	LOQ, μg/kg
DMA	0.17	1.3
MMA	0.15	1.2
iAs	0.17	1.4

The iAs LOQ is well within the FDA's 10 μ g/L level of concern for iAs in juice samples. The sensitivity of the method is therefore sufficient to determine iAs in solution following a five-fold dilution of the samples.

Quantitative results

The five wines included in the MLV were analyzed in the lab at UC Davis using LC-ICP-QQQ and the results are shown in Table 3. The average percent recovery of the sum of the species compared to the total As present in the samples (determined using direct analysis without HPLC separation) was calculated using the mass balance approach. The percent recovery for all samples was between 91–107%. The results were found to be in good agreement with the results obtained from the other laboratories taking part in the MLV study [2].

Table 3. Quantitative results for the five wines analyzed at UC Davis as part of the MLV study. Average ± 1σ, n=3 for the individual species

Wine sample	% Ethanol (v/v)	DMA μg/kg	MMA μg/kg	iAs μg/kg	Sum of species µg/kg	Total As μg/kg	Mass balance %
Red (Cabernet)	9.5	0.81 ± 0.1*	<lod< td=""><td>14.4 ± 1.0</td><td>15.2 ± 1.1</td><td>15.3 ± 1.2</td><td>99</td></lod<>	14.4 ± 1.0	15.2 ± 1.1	15.3 ± 1.2	99
White (Chardonnay)	13	0.74 ± 0.04*	<lod< td=""><td>10.7 ± 0.2</td><td>11.4 ± 0.2</td><td>11.1 ± 0.8</td><td>103</td></lod<>	10.7 ± 0.2	11.4 ± 0.2	11.1 ± 0.8	103
Rosé (Zinfandel)	12	0.75 ± 0.1*	<lod< td=""><td>9.2 ± 0.4</td><td>9.9 ± 0.4</td><td>9.3 ± 1.1</td><td>107</td></lod<>	9.2 ± 0.4	9.9 ± 0.4	9.3 ± 1.1	107
Sparkling wine	20	1.7 ± 0.1	<lod< td=""><td>2.1 ± 0.3</td><td>3.8 ± 0.3</td><td>3.6 ± 0.3</td><td>105</td></lod<>	2.1 ± 0.3	3.8 ± 0.3	3.6 ± 0.3	105
Port-style wine	14.5	0.45 ± 0.01*	<lod< td=""><td>1.5 ± 0.3</td><td>2.0 ± 0.3</td><td>2.2 ± 0.1</td><td>91</td></lod<>	1.5 ± 0.3	2.0 ± 0.3	2.2 ± 0.1	91

^{*} Value between LOD and LOQ

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Long-term stability

To test the stability of the ICP-QQQ over an extended sampling period of 96 hours (four days), the wine samples were measured repeatedly in a continuous sequence. Two quality control (QC) samples—a 2-ppb mixed As species standard solution and NIST 1643e spiked with 3% ethanol— were analyzed after every 10 wine samples. The instrument was not recalibrated during the continuous analytical run. The plots shown in Figures 3 and 4 show exceptional stability was achieved over the course of the validation stability test.

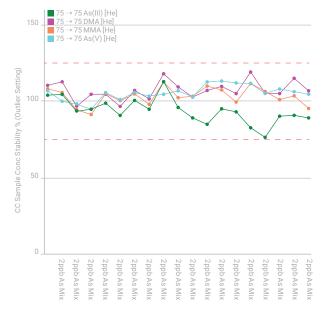


Figure 3. Stability plot of the 2-ppb mixed As species standard solution, analyzed over 96 hours (four days).

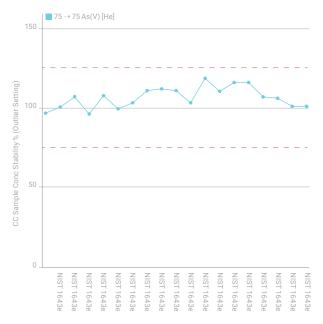


Figure 4. Stability plot of As in NIST 1643e spiked with ethanol and analyzed over four days.

Results of additional market basket wine analysis

In addition to the five wines used in the MLV study, an extra 60 wines were analyzed as part of the method validation [2]. In this study, a selection of previously untested wines (S1 to S5) were analyzed. The results shown in Table 4 are consistent with the published data from the reference paper [2]. Most of the As was in the more toxic, inorganic forms. While four of the five wine samples contained levels of total As higher than the EPA drinking water limit of 10 μ g/L, the levels in all five wines were below the 100 and 200 μ g/kg limits for total As in wine set in Canada and Europe, respectively. However, the measured concentrations for iAs in four out of five of the wines exceeded the FDA's action limit of 10 μ g/kg for iAs in apple juice.

Table 4. Quantitative results ($\mu g/kg$) for As species in five commercially available wines measured by LC-ICP-QQQ.

Wine Sample	iAs	DMA	ММА	Sum of Species
S1	17.13 ± 0.22	0.83 ± 0.03	<lod< td=""><td>17.96 ± 0.13</td></lod<>	17.96 ± 0.13
\$2	7.49 ± 0.15	0.30 ± 0.06	0.77 ± 0.32	8.56 ± 0.17
S3	14.63 ± 0.40	0.80 ± 0.08	<lod< td=""><td>15.43 ± 0.24</td></lod<>	15.43 ± 0.24
S4	25.03 ± 0.89	0.69 ± 0.26	0.47 ± 0.12	26.19 ± 0.42
S 5	23.45 ± 1.12	0.32 ± 0.05	<lod< td=""><td>23.77 ± 0.59</td></lod<>	23.77 ± 0.59

Spike recovery test

Table 5 shows the spike recoveries for the MLV samples fortified at levels of approximately 5, 10, and 30 μ g/kg for DMA, MMA, and iAs (the iAs spike concentration was the sum of As(III) and As(V) each spiked at 50% of the levels shown). The average recoveries of DMA, MMA, and iAs measured using LC-ICP-QQQ were 99, 92, and 104%, respectively. All the recoveries are within the FDA's EAM acceptability criteria of 100 \pm 20% for iAs, DMA, and MMA [1].

Table 5. Average spike recovery results for duplicate analyses of five samples spiked at 5, 10, and 30 μg/kg with DMA, MMA, and iAs. n=30.

	DMA	ММА	iAs
Average spike recovery, %	99	92	104
Recovery range	93 – 107	72 – 119	97 – 114

Conclusions

The As speciation results obtained using an Agilent 1260 Infinity LC coupled to an Agilent 8800 ICP-QQQ were used as part of an MLV to validate the extension of Elemental Analysis Manual Method §4.10 to include wine. The method was optimized for the analysis of four arsenic species including the toxicologically relevant inorganic forms, As(III) and As(V).

In addition to the data published as part of the MLV, five more wines were analyzed. The total As levels of the five wines were between 8.56 and 26.19 μ g/L. These levels are below the Canadian and European regulatory limits for total As in wine of 100 and 200 μ g/kg, respectively. The average percentage of As found in the form of iAs in the five wine samples was 95%.

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More Information

For a full account of this study, see Courtney K. Tanabe, Helene Hopfer, Susan E. Ebeler, Jenny Nelson, Sean D. Conklin, Kevin M. Kubachka, and Robert A. Wilson, Matrix Extension and Multilaboratory Validation of Arsenic Speciation Method EAM §4.10 to Include Wine, *J. Agric. Food Chem.*, **2017**, 65 (20), pp 4193–4199, DOI: 10.1021/acs.jafc.7b00855

Multielement Analysis and Selenium Speciation in Cattle and Fish Feed using LC-ICP-QQQ

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Validation of an extended FDA Elemental Analysis Manual method

Introduction

In the USA, animal feed is subject to regulation under the Federal Food, Drug, and Cosmetic Act (FFDCA), which defines food as "articles used for food or drink for man or other animals". The US Food and Drug Administration's (FDA) Center for Veterinary Medicine (CVM) is responsible for regulations relating to the safety of feed intended for animals (including but not limited to horses, cattle, swine, poultry, and fish), under the Animal Feed Safety System (AFSS). The regulations control many aspects of the production, storage, and labeling of animal feed, and the permitted levels of additives and contaminants, such as potentially toxic heavy metals.

Selenium (Se) has been approved by the FDA as a supplement for animal feed since the 1970s. The FDA's current method for Se quantification uses ICP-MS with helium collision/reaction cell (CRC) for control of interferences [1], but the sensitivity of this method is affected when high levels of interferences are present.

Selenium is an essential trace nutrient for vertebrates, and is involved in several vital metabolic processes. The recommended human dietary intake is approximately 55 μ g Se per day, which most people acquire through the consumption of plant-based foods, such as cereals. However, Se in the soil is not evenly distributed geographically, so dietary supplementation is commonplace in some parts of the world. This approach requires caution though, as Se is toxic in excess, with a tolerable upper intake level of about 200 μ g/day depending on gender and age [2].

Animal feeds including cattle feed and fish feed are often supplemented with selenium. Supplementation may be in the form of sodium selenite/selenate, which is approved by the European Food Safety Authority (EFSA) as a food additive for all animal species [3]. Selenized yeast is also commonly used as an additive; if properly produced, the fortified yeast should contain primarily selenomethionine (SeMet). It is noted that the US regulations for supplemented selenium applicable for cattle feed and other livestock do not apply to fish feed.

Total and speciation analysis of selenium

Determination of total Se concentration is commonly carried out as part of a multielement analysis using ICP-MS. More recently, Se analysis has benefited from the lower detection limits and greater freedom from spectral interferences provided by triple quadrupole ICP-MS (ICP-QQQ) [4, 5]. However, the toxicity of Se depends on the chemical form or species in which the Se occurs, so total elemental concentrations do not provide a complete picture of the element's potential toxicity. As a result, separation and detection of the individual species (speciation) is required. The major Se species that occur naturally in the types of crop plant used to produce animal feed include two inorganic species, selenite

(Se(IV)) and selenate (Se(VI)), and some selenoamino acids: selenocystine (SeCys₂), selenomethionine (SeMet), and Se-methyl selenocysteine (MeSeCys). Selenoamino acids are considered to be less toxic than the inorganic forms, with Se(IV) being the most toxic species [6]. In addition to the naturally occurring selenium species, animal feeds may contain various selenium compounds added during production to raise the level of total selenium in the animals' diet.

Se speciation analysis typically uses the well-established analytical method of HPLC (to separate the various Se-containing species) coupled to ICP-MS (to identify and quantify the individual Se species). HPLC-ICP-MS has been widely employed for analyzing various sample types [7] but there are few studies on selenium speciation in animal feed [8].

In this study, we developed an extraction and analytical method for the measurement of total Se using ICP-QQQ, and for Se speciation using LC-ICP-QQQ. The method provides low background and high sensitivity enabling low detection limits for total Se and Se compounds to be achieved. We then applied the speciation method to evaluate the selenium content and species distribution in cattle feed and fish feed.

Experimental

Samples and sample preparation

Two commercial cattle feeds and four commercial aquaculture feeds were bought from local stores.

Total Se (and multi-element) analysis of feeds

Cattle and fish feed samples were weighed to approximately 100 mg dry mass and microwave digested in a 1:1 mix of trace metal grade $\mathrm{HNO_3}$ and distilled de-ionized (DDI) $\mathrm{H_2O}$. A solution containing various internal standards was added before digestion, giving an internal standard concentration of 5 ng/g in the final diluted solutions as analyzed. The microwave program consisted of a first step at 300 W with a 10 min ramp to 95 °C and a second step at 300 W including a 10 min ramp to 200 °C, followed by a 20 min hold time. After cooling, 1 mL of 30% $\mathrm{H_2O_2}$ was added and a second digestion was performed using the same heating program. The sample digests were diluted with DDI water to give a final acid concentration of approximately 2% $\mathrm{HNO_3}$. Two certified reference materials (CRMs) NIST 1547 Peach Leaves (NIST, Gaithersburg, MD USA) and SELM-1 Selenium Enriched Yeast (National Research Council of Canada) were prepared as quality control samples.

Enzymatic extraction procedure for Se speciation in feeds

An extraction solution containing 50 mM ammonium phosphate monobasic was prepared and the pH was adjusted to 7.5. Feed samples were weighed to approximately 200 mg with 3 mL of the extraction solution [9]. Samples were sonicated using QSonica sonication probe. The sonication program consisted of a 2 second pulse, followed by a 3 second rest at 60% amplitude for 2 minutes. Following sonication, approximately 20 mg of protease type XIV (from Streptomyces griseus, Sigma-Aldrich) and 10 mg of lipase (from Candida rugose, Sigma-Aldrich) dissolved in buffer solution were added to each sample and placed on a hot block for 12 hours at 37 °C. After 12 hours, the samples were sonicated again at 30% amplitude for 30 seconds and placed on the hot block for a further 12 hours.

The "enzyme extract" was filtered with a 0.45 m Spin-X Centrifuge Tube Filter (Costar, USA). The resulting sample was then diluted 1:1 with mobile phase 1 ready for analysis by reversed phase ion pairing LC-ICP-QQQ.

Instrumentation

An Agilent 8800 Triple Quadrupole ICP-MS (ICP-QQQ) equipped with an Agilent SPS4 autosampler was used for analysis of total selenium and other elements in cattle and fish feed samples. Instrument operating conditions are given in Table 1.

Table 1. ICP-QQQ operating conditions.

Parameter	No gas	He	O ₂	
Spray chamber temp (°C)				
RF power (W)	15	1600		
Sampling depth (mm)	8.	8		
Carrier gas (L/min)	1.00			
Make up gas (mL/min)	0.1	0.15		
Cell gas (mL/min)	0.0	3.4	0.5 (30%*)	

^{*} Indicates % of full scale flow rate, as displayed in the ICP-MS MassHunter Tune screen

Agilent ICP-MS MassHunter software was used for the setup and operation of the ICP-QQQ for total Se and multielement data analysis. ICP-MS MassHunter with the optional Chromatographic Analysis module was used for combined instrument control and sequencing of the LC-ICP-QQQ Se-speciation study.

For the speciation studies in cattle and fish feed, an Agilent 1100 Series HPLC was coupled to the ICP-QQQ. Chromatographic separations were performed using an Agilent ZORBAX Extend column (80 Å C18, 4.6 x 250 mm, 5 μm). Details of the HPLC method used for Se speciation analysis of cattle and fish feed are given in Table 2.

The six Se species of interest, Se(IV), Se(VI), selenocystine (SeCys₂), selenomethionine (SeMet), methyl selenocysteine (MeSeCys), and selenomethionine-Se-oxide (SeOMet), were calibrated using mixed standard solutions containing each of the Se species at levels from 1 to 50 ng/g.

In common with any ion paring method, column equilibration is crucial to ensuring long-term reproducibility when using this LC-ICP-QQQ method. Equilibration is important after cleaning or when the column has been stored for a long time. To prevent deterioration of the column, 2 mM TBAH was added to the 65% acetonitrile storage solution. Following storage or cleaning, the column was equilibrated for 20 mins with $3x75~\mu L$ injections of 25 mM TBAH dissolved in the mobile phase.

Table 2.HPLC method used for the analysis of cattle and fish feed sample extracts.

Method	Salt gradient; reverse-phase ion-pairing					
Injection volume		25 μL				
Mobile phase 1	5 mM ammonium acetate, 2 mM ammonium phosphate, 2 mM TBAH, 2% MeOH, pH 6.5					
Mobile phase 1	15 mM ammonium acetate, 5 mM ammonium phosphate, 2 mM TBAH, 2% MeOH, pH 6.5					
Method	Minute	% Mobile phase 2	Flow rate (mL/min)			
	5	0	1			
	10	100	1			
	16	100	1			
	20	0	1			
	21	0	1.5			
	40	0	1.5			
	0	1				

Interference removal

Routine determination of total selenium concentrations or analysis of Se species using LC-ICP-MS does not necessarily require the use of ICP-QQQ. Conventional quadrupole ICP-MS (ICP-QMS) fitted with a CRC is able to resolve the ⁴⁰Ar³⁸Ar⁺ dimer interference on ⁷⁸Se sufficiently well to give acceptable results for Se speciation analysis in many sample types [7]. However, doubly-charged ion interferences such as ^{156/160}Gd⁺⁺ and ^{156/160}Dy⁺⁺ on ^{78/80}Se can lead to positive bias in samples containing relatively high levels of the rare earth elements (REEs) [5]. In these sample types, ICP-QQQ is able to completely resolve the doubly charged REE interferences along with other spectral interferences, giving lower detection limits and better accuracy than ICP-QMS for Se (and As).

Results and Discussion

Multielement analysis

The multielement analysis results including total Se content of the feed and CRM samples are summarized in Table 3. The measured value for total Se in Se-yeast SELM-1 CRM was in good agreement with the certified value at 94% recovery. The results validate the sample preparation method and the accuracy of the ICP-QQQ results. Accurate recovery of Se in NIST 1547 Peach Leaves was also obtained (102%, relative to the original 1991 certified value). This indicates the effective control of interferences including doubly charged rare earth elements, as NIST Peach Leaves contains up to 10 mg/kg (ppm) of the light rare earth elements. Table 3 also includes instrument detection limits (IDLs) demonstrating low ng/L (ppt) detection limits for most analytes.

Table 3. Total selenium dry weight concentration (mg/kg) determined in cattle feed, fish feed, and CRMs analyzed by ICP-QQQ.

Element	Tune	Q1 🛭 Q2 Set Mass	Cattle Feed 1	Cattle Feed 2	Fish Feed 1	Fish Feed 2
Mg	He	24->24	2,278 ± 49	2,762 ± 26	1,873 ± 37	2,419 ± 17
К	He	39->39	8,771 ± 77	9,556 ± 60	7,064 ± 129	10,080 ± 196
V	He	51->51	1.37 ± 0.05	0.28 ± 0.01	0.30 ± 0.03	0.36 ± 0.02
Cr	He	52->52	1.79 ± 0.07	0.90 ± 0.03	1.16 ± 0.06	0.74 ± 0.06
Fe	He	56->56	392 ± 21	166 ± 25	432 ± 10	255 ± 3
Co	He	59->59	0.66 ± 0.02	1.3 ± 0.1	0.127 ± 0.001	0.67 ± 0.04
Cu	He	63->63	31.1 ± 0.8	26 ± 2	8.68 ± 0.08	66 ± 2
As	02	75->91	0.21 ± 0.02	0.11 ± 0.01	1.14 ± 0.09	0.30 ± 0.05
Se	02	78->94	0.86 ± 0.04	0.69 ± 0.03	1.07 ± 0.08	0.98 ± 0.05
Sr	He	88->88	11 ± 2	11.4 ± 0.5	49 ± 1	16.4 ± 0.6
Мо	He	95->95	1.35 ± 0.03	2.17 ± 0.01	0.77 ± 0.01	1.52 ± 0.02
Cd	He	111->111	0.094 ± 0.004	0.072 ± 0.003	0.40 ± 0.02	0.049 ± 0.002
Pb	No gas	208->208	0.24 ± 0.03	0.12 ± 0.01	0.38 ± 0.06	0.23 ± 0.09

Element	Tune	Fish Feed 3	Fish Feed 4	NIST 1547 ^a	ELM-1ª	IDL, ppb
Mg	He	2,152 ± 29	1,586 ± 56	4,406 ± 72 (98)		0.2116
К	He	11,298 ± 110	7,520 ± 66	22,167 ± 364 (91)		7.16
V	He	0.42 ± 0.01	1.14 ± 0.08	0.341 ± 0.006 (93)		0.0123
Cr	He	2.37 ± 0.01	1.10 ± 0.05	1.067 ± 0.009 (107b)		0.0044
Fe	He	204 ± 15	642 ± 50	225 ± 3 (102)		0.1027
Со	He	0.146 ± 0.002	0.20 ± 0.01	0.068 ± 0.002 (97b)		0.0005
Cu	He	16.2 ± 0.3	10.42 ± 0.08	3.8 ± 0.2 (101)		0.027
As	02	0.171 ± 0.009	0.81 ± 0.03	0.08 ± 0.02 (133c)		0.0035
Se	02	0.55 ± 0.02	1.05 ± 0.04	0.122 ± 0.003 (102c)	1911 ± 97 (94)	0.0031
Sr	He	40 ± 2	32 ± 2	62 ± 1 (117)	-	0.018
Мо	He	1.67 ± 0.03	1.11 ± 0.03	0.063 ± 0.006 (104)		0.002
Cd	He	0.074 ± 0.004	0.056± 0.006	0.028 ± 0.001 (107)		0.0039
Pb	No gas	0.25 ± 0.04	0.41 ± 0.03	0.82 ± 0.03 (94)		0.1946

a. Values enclosed in parenthesis are recoveries of the certified value of reference material.

In the United States, the Association of American Feed Control Officials (AAFCO) 2011 Guidelines [10] and the US FDA's 21 CFR Part 573, Section 573.920 (Selenium) [11] state animal feeds intended for chickens, turkeys, swine, beef cattle, dairy cattle (and in the AAFCO Guidelines, bison, sheep, goats, llamas, alpacas, and horses) may contain selenium yeast at a level not to exceed 0.3 ppm (mg/kg) of selenium based on the complete feed [10]. Furthermore, the level of inorganic species should not exceed 2% of the total Se content in the final yeast product. Our results show that the two cattle feeds contained Se significantly above the 0.3 mg/kg limit, at 0.86 and 0.69 mg/kg Se. Similar results were found in the four fish feeds tested, which contained concentrations between 0.55 and 1.07 mg/kg Se.

b. Recovery determined relative to a non-certified, information value.

c. Recoveries for As and Se are calculated relative to the original certified values (1991 revision). These certified values have subsequently been removed from the certificate (2017 revision) so may not be reliable.

All feeds contained at least twice the maximum Se concentration of 0.3 ppm (mg/kg) in selenium supplemented feeds. The feeds were likely supplemented by the addition of "antioxidants" including Se-yeast to increase the selenium content. To further investigate the Se content of the feeds, speciation analysis was performed to separate and quantify the individual Se species present in the feed samples.

Selenium speciation analysis

Selenium speciation analysis was performed using LC coupled to ICP-QQQ. Se was measured using the oxygen cell gas tune mode as for Se in the multielement analysis. The selenium species concentrations were calibrated using mixed standard solutions containing each of the Se species at levels from 1 to 50 ng/g. The integrated peak areas for each species were plotted versus the standard concentrations to generate calibration curves covering the required calibration range.

The chromatogram shown in Figure 1 was obtained from the analysis of a mixed standard containing each Se species at 25 ng/g. The chromatogram demonstrates good sensitivity and peak separation for all species. Peak identities were confirmed by retention time (RT) matching and/or the use of standard spikes added to the extracts.

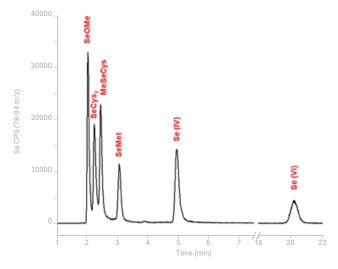


Figure 1. LC-ICP-QQQ chromatogram of standard containing six selenium species at 25 ng/g.

Se species in cattle feed samples

The two cattle feed samples were analyzed using LC-ICP-QQQ. The overlaid chromatograms in Figure 2 show that both samples contained primarily SeMet, while cattle feed 2 also contained significant levels of Se(VI). Other species were present at trace levels. The source of SeMet was likely to be selenized yeast, which is often added intentionally to enrich the feeds. However, natural sources, such as grains and soybeans, are common additives that have been found to accumulate SeMet when supplied with inorganic Se sources [12–14]. Plants naturally uptake Se from soils, and inorganic Se species tend to be more mobile, which leads to increased plant uptake compared with organic forms. Depending on soil conditions, either Se (IV) or Se (VI) may be the major Se source for plants [15]. When Se (IV) is the main source of selenium, it gets metabolized to organic Se compounds; while Se (VI) uptake generally results in higher accumulation

without transformation [12, 14]. It can be concluded that the Se (VI) found in the cattle feeds for this study likely originated from the addition of high-selenium plant additives often used in feed production.

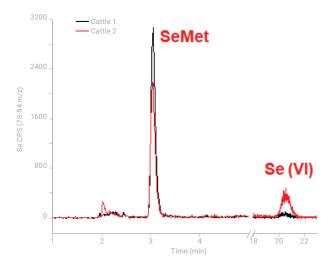


Figure 2. LC-ICP-QQQ speciation analysis of two cattle feed samples.

Se species in fish feed samples

Figure 3 shows chromatograms for the four fish feed samples. As with the cattle feeds, a variety of Se species were observed in these samples. SeMet was the primary species in all feeds, but inorganic forms such as Se(IV) and Se(VI) were also present. The quantitative results in Table 4 show that Se(IV) was considerably higher in fish feed 1 compared to the other samples, while Se(VI) was higher in the cattle feeds than the fish feeds.

Selenium supplementation has been shown to improve growth and antioxidant status for fish reared in the crowded conditions that are typical of mass production methods [16]. Previous aquaculture studies have shown supplementation with inorganic forms of Se, mainly Se(IV), leads to inferior bioavailability compared to SeMet or selenoyeast [17, 18]. Due to greater accumulation of Se in fillets and whole body, many studies currently use organic Se for aquaculture and supplementation research [19, 20].

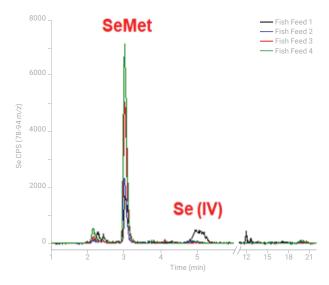


Figure 3. LC-ICP-QQQ speciation analysis of four fish feed samples.

Table 4. Enzymatic extraction and quantification of Se species for cattle and fish feeds using LC-ICP-QQQ.

Sample	Extraction total	Extraction	Quantification of known species (µg/kg, ppb)						
	(μg/kg, ppb)	efficiency (%)* -	SeOMet	SeOMet	MeSeCys	SeMet	Se (IV)	Se (VI)	
Cattle feed 1	386 ± 3	45 ± 1	2 ± 1	44 ± 41	5 ± 1	112 ± 8	0 ± 0	72 ± 3	
Cattle feed 2	346 ± 14	50 ± 2	6 ± 3	4 ± 2	3 ± 1	97 ± 3	24 ± 27	128 ± 15	
Fish feed 1	648 ± 32	61 ± 3	5 ± 2	19 ± 3	12 ± 1	79 ± 11	61 ± 8	23 ± 3	
Fish feed 2	484 ± 90	49 ± 9	2 ± 1	14 ± 6	9 ± 1	117 ± 11	10 ± 3	22 ± 4	
Fish feed 3	363 ± 103	66 ± 19	4 ± 2	17 ± 3	8 ± 2	222 ± 8	3 ± 5	30 ± 7	
Fish feed 4	710 ± 43	68 ± 4	2 ± 1	27 ± 1	14 ± 1	293 ± 39	14 ± 3	31 ± 1	

^{*} Compared to total Se concentration (shown in Table 3).

Conclusions

Total concentrations of several elements, including selenium, were determined in cattle and fish feed sample-extracts, using the Agilent 8800 ICP-QQQ. In all samples, the concentration of Se was above the maximum of 0.3 ppm (μ g/kg) Se recommended in the AAFCO and FDA guidelines for selenium supplemented feeds.

Reversed phase ion pairing LC-ICP-QQQ was then used successfully to separate and detect the selenium species at low mg/kg levels in the feed samples (low μ g/L in the solutions analyzed). The method provided valuable information on the Se species present in the feeds. SeMet was found to be the predominant species, although the toxicologically relevant inorganic forms of Se (Se(IV) and Se(VI)) were also found to be present in most of the samples.

More Information

For a full account of part of this application, see A. F. Oliveira, J. Landero, K. Kubachka, A. R. A. Nogueira, M. A. Zanetti and J. Caruso, Development and application of a selenium speciation method in cattle feed and beef samples using HPLC-ICP-MS: evaluating the selenium metabolic process in cattle. *J. Anal. At. Spectrom.*, **2016**, 31, 1034. DOI: 10.1039/c5ja00330j

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Sulfur isotope fractionation analysis in mineral waters using an Agilent 8900 ICP-QQQ

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Introduction

Stable isotope geochemistry is a branch of geology that investigates the age of natural materials, their origin and the processes they have undergone since formation [1]. Stable isotope analysis is also used in biogeochemical studies to monitor element cycling in ecosystems [2] and to identify geographical/regional differences for food provenance and archaeology. Of the elements of interest in stable isotope studies (hydrogen, carbon, nitrogen, oxygen and sulfur), only sulfur is accessible using aqueous solution analysis by ICP-MS, and even sulfur is difficult to measure by conventional quadrupole ICP-MS.

The relative abundance of the two major stable isotopes of sulfur, 32 S (94.99% abundance) and 34 S (4.25%), varies significantly in nature, so the 34 S/ 32 S ratio can be used to characterize a sample. In sulfur stable isotope analysis, the variation in the 34 S/ 32 S isotope ratio is calcuated and reported as a deviation or delta (δ) in 34 S abundance relative to a standard material, the troilite (iron sulfide) mineral from the Canyon Diablo meteorite, referred to as δ VCDT (Vienna Canyon Diablo Troilite). Natural variations in 34 S abundance, expressed in parts per thousand or "per mil" (%), can be of the order of -50% to +40% (and occasionally much greater), due to redox reaction. Examples of some values for natural S isotope fractionation are given in Table 1 [3].

Table 1. Sulfur isotope distribution in nature.

Source	δ ³⁴ S (‰) relative to VCDT
Igneous rocks	0
Sedimentary rocks	-40 to +40
Seawater SO ₄	+21
Atmospheric SO ₄	-30 to +30
Surface water/groundwater SO ₄	-22 to +135
Soil (organic sulfur)	-30 to +30
Vegetation (organic sulfur)	-34 to +32
Animals (organic sulfur)	-10 to +22
Fossil fuels (organic sulfur)	-11 to +28

Sulfur isotope ratio (IR) analysis has been mostly done by gas phase isotope ratio mass spectrometry (IRMS) but recent developments in ICP-MS technology have vastly improved its ability to measure sulfur accurately at low levels. In this work, we investigated the performance of a new, high senistivity ICP-MS instrument, the Agilent 8900 Triple Quadrupole ICP-MS (ICP-QQQ), for low level S IR analysis of mineral waters. Spectral interferences on S arising from O_2^+ can be removed by operating the ICP-QQQ in MS/MS mode, allowing both sulfur isotopes to be measured and potentially offering a faster and simpler S isotope analysis technique.

Experimental

Sulfur IR analysis method: mass-bias correction, matrix effects and background control

For accurate and precise IR analysis by ICP-MS, the instrumental mass bias must be corrected, and the effect of the sample matrix must be controlled.

As is typical for isotope ratio analysis by ICP-MS, instrumental mass-bias was corrected using sample-standard bracketing. A standard solution of known S isotope composition was measured before and after each sample, and the sample IR was corrected by the average IR of the two standard measurements. A 0.5 ppm solution of IAEA-S-1 was used as the mass bias correction standard [4].

The sample matrix can also affect the relative transmission of different mass ions in ICP-MS, and consequently the mass bias and the measured IR. To overcome this effect, a chelation technique can be used to remove the sample matrix before analysis [5]. Alternatively, the variation in sample matrix composition can be reduced by diluting all samples and standards in a consistent matrix. In this work, the mass bias standard and samples were diluted using a solution which contained 50 ppm calcium (Ca) and 100 ppm sodium chloride (NaCl). Use of this diluent reduced the matrix variation that could otherwise have caused fluctuations in the mass bias. The S concentration in the matrix blank was around 0.7 ppb which was low enough not to affect the accuracy of the IR analysis.

Sample dilution in a consistent matrix avoided the necessity for time consuming matrix removal. The matrix dilution approach was made possible by the high sensitivity and low S background of the 8900 ICP-QQQ. Sulfur is ubiquitous in laboratory consumables, supplies, and many of the materials used in instrument components, typically leading to a high elemental background signal. To minimize the contribution from the ICP-MS hardware, key components of the argon gas flow path of the 8900 #100 (Advanced Applications configuration) ICP-QQQ have been replaced using more inert materials. This has successfully reduced the background signal for S (and Si), allowing a detection limit specification of < 50 ppt for S, Si (and P) to be quoted¹. In a recent study, S was measured with a sensitivity of 10⁴ cps/ppb using the 8900 #100 ICP-QQQ to achieve a background equivalent concentration (BEC) of less than 100 ppt S in ultrapure water [6].

Instrumentation

An Agilent 8900 ICP-QQQ (#100, Advanced Applications configuration) equipped with the standard Ni cones and x-lens was used. The standard glass concentric nebulizer was replaced with a PFA nebulizer, run using self-aspiration for better signal precision.

The two most abundant isotopes of S, 34 S and 32 S, were measured using the Agilent 8900 #100 in MS/MS mass-shift mode with O_2 cell gas [6]. The polyatomic interference from $^{16}O_2^{+}$ on the primary isotope of S, 32 S+ at m/z 32, and from $^{16}O^{18}O^{+}$ on the minor 34 S isotope at m/z 34 were avoided by shifting S+ to a new mass. S+ reacts readily with O_2 cell gas to form the product ion SO+, while the O_2^{+} interference does not react in the same way with the O_2^{-} cell gas. Consequently, the SO+ product ions can be measured free of interference at M + 16 amu (m/z 48 for the primary 32 S 16 O+ isotope product ion and m/z 50 for 34 S 16 O+). Tuning conditions and method parameters are summarized in Table 2.

1. This specification is verified on every Agilent 8900 Advanced Applications and Semiconductor configuration instrument during factory testing

Table 2. Agilent 8900 ICP-QQQ tuning and method conditions

	Tuning parameter	Value
Plasma	RF power (W)	1550
	Sampling depth (mm)	8.0
	Nebulizer gas flow rate (L/min)	0.90
	Make up gas flow rate (L/min)	0.30
Lens	Extract 1 (V)	-80
	Extract 2 (V)	-150
	Omega (V)	10.0
	Omega bias (V)	-120
Cell	Octp Bias (V)	-5.0
	Axial Acceleration (V)	2.0
	KED (V)	-8.0
	Cell gas	Oxygen
	Cell gas flow rate (mL/min)	0.45

	Method parameter	Value
Data Acquisition	Integration time (s)	1 and 5 for 32S and 34S
	Number of sweeps	1000
	Number of replicates	10
Rinse	1% HNO ₃ rinse (s)	20
	50 ppm Ca/100 ppm NaCl rinse (s)	30
Peripump	Uptake time (s)	30
	Stabilization time (s)	30

The high sensitivity and low background of 32 S and 34 S can be clearly seen in Figure 1, which shows a spectrum of sulfur obtained using the MS/MS method.

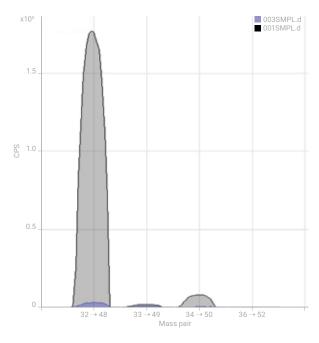


Figure 1. MS/MS spectrum of 10 ppb sulfur solution (grey) and blank (blue).

Sample and sample preparation

Sulfur isotope certified reference materials (CRMs) IAEA-S-1 ($^{34}\delta_{\text{VCDT}}$ = -0.3%) and IAEA-S-2 ($^{34}\delta_{\text{VCDT}}$ = +22.7%) were purchased from National Institute of Standards and Technology (NIST), Gaithersburg, MD, USA. Each CRM was gently dissolved in diluted nitric acid and diluted to the appropriate concentration. A matrix blank was prepared with 50 ppm Ca (SPEX Certiprep, US) and 100 ppm NaCl (Wako Pure Chemical Industries Ltd, Japan) in 1% nitric acid (Tamapure 100: Tama Chemicals Co. Ltd, Japan). This solution was also used to dilute the standards and samples.

The seawater and mineral water samples were diluted between 10 and 2000 times to give a S concentration in the range 0.2 to 0.8 ppm. Concentration matching contributed to accurate isotope ratio analysis because, at these levels, ³²S is measured in analogue mode and ³⁴S in pulse counting mode. Concentration matching also removes any potential errors caused by detector dead time (the instrument default dead time was 36.3 ns for mass 32).

Results and Discussion

Synthetic sample analysis

Sulfur isotope CRMs IAEA-S-1 and IAEA-S-2 and two mixes of the two CRMs were prepared to give four samples with theoretical $^{34}\delta_{\text{VCDT}}$ values of -0.3, 5.4, 11.2 and 22.6. Each blend was prepared at a S concentration of 0.5 ppm. The S IRs were measured six times (standard corrected as described previously).

Figure 2 shows the raw IR data for the IAEA-S-1 mass bias standard and the corrected IR data for the CRM blend samples. The average $^{34}\delta$ values and errors (two times the standard deviation) were determined for the four CRM mixes, and these measured values were plotted against the theoretical values for each mixed standard. The results can be seen in Figure 3, demonstrating the good linearity obtained.

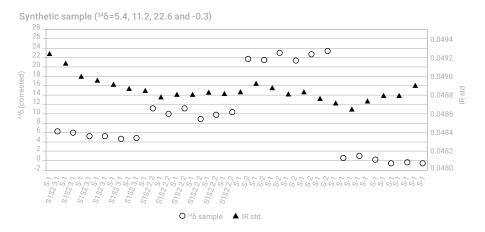


Figure 2. Raw IR analysis of bracketing mass bias standard IAEA-S-1 (triangles) and corrected S IRs of six separate measurements of each of four isotope CRM blends (points).



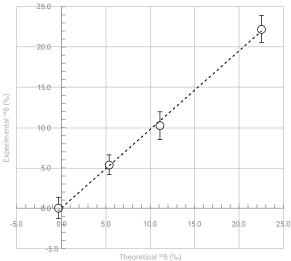


Figure 3. Average of Sulfur IR analysis of the four IAEA CRM blends.

Water sample analysis

Three different brands of mineral water were purchased at a local store in Tokyo, Japan. The mineral water samples were prepared for analysis, together with samples of; JSAC 0301, a Japanese river water CRM (from the Japan Society for Analytical Chemistry); a spring water collected from the IKAHO hot spring in the north of Japan; a NASS 5 seawater CRM (National Research Council, Canada); and Tamapure-AA 100 high purity sulfuric acid (Tama Chemicals Co., Ltd.).

Before the IR measurements were undertaken, the sulfur concentration of each sampe was checked to determine the appropriate dilution factor. The dilution factors applied to the samples are given in Table 3. Each sample was measured 10 times and the average and the standard deviation were calculated. Figure 4 shows the average IR and the error (as two times standard deviation) of the IR.

Table 3. Dilution factors

Sample	Dilution factor
Mineral water A	10
Mineral water B	10
Mineral water C	1000
JSAC 0301: Japanese river water CRM	10
IKAHO hot spring water	1000
NASS 5	2000
High purity sulfuric acid	50000

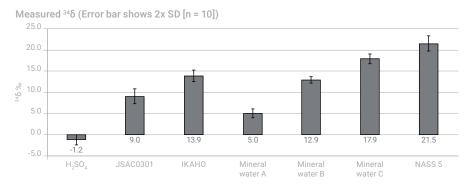


Figure 4. Measured sulfur IR for sulfuric acid, river water CRM, spring water, 3 commercial mineral waters (brands A, B and C) and seawater CRM.

The results show a clear difference in the S IRs for all of the samples, including between the 3 brands of mineral water. The $^{34}\delta$ value of +21.5% determined in the seawater reference material agrees well with the global average oceanic seawater value of +21% for seawater sulfate (see Table 1 and reference 3).

The new, fast ICP-QQQ method for sulfur isotope analysis could be useful in identifying the natural characteristics of a water-source, monitoring seasonal and biogeochemical variations, and also for determining the impact of man-made sources of sulfur on the environment.

Conclusions

The Agilent 8900 Advanced Applications configuration ICP-QQQ is ideally suited to ³⁴S/³²S isotope ratio analysis, which can provide valuable information for sample characterization in natural systems or to monitor anthropogenic impact. The 8900 ICP-QQQ provides a low background and high sensitivity for sulfur, which enabled a method to be developed that simply required the sample to be diluted with the matrix blank before analysis. Sample/standard bracketing was used to correct for any instrumental mass-bias or drift.

By operating the 8900 ICP-QQQ in MS/MS mode with O_2 cell gas, problematic spectral interferences due to O_2^+ overlaps on $^{32}S^+$ and $^{34}S^+$ were successfully avoided. The S IR analysis method was applied to various samples including three mineral waters, a river CRM, a seawater CRM, a hot spring water, and high purity sulfuric acid. The precision of the IRs achieved was excellent at 1-1.5 % (as two times the standard deviation).

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Fast Analysis of Arsenic Species in Wines using LC-ICP-QQQ

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Introduction

Arsenic (As) occurs naturally in the environment but human activity has contributed to the levels found in some locations. Man-made sources of As include industrial processes such as mining, smelting and power generation, as well as agricultural pesticides and timber preservatives [1]. Once contamination has occurred, As persists in the environment for decades. For example, the widespread use of As-containing agrochemicals ceased in the 1970s, but lead and calcium arsenate levels remain high in some soils. As can be absorbed from soil and water into crops. In the case of wine, the As content can also be affected by the wine making processes.

Arsenic exists in multiple forms in foods and beverages and not all forms have the same toxicity. The inorganic forms of As (iAs), comprising As(III) (arsenite) and As(V) (arsenate), are the most toxic, and are categorized as class 1 carcinogens. In contrast, arsenobetaine (AB), the most abundant form of As in fresh seafood, is essentially non-toxic to humans. Due to the high variability in the toxicity of the different species of As, and the potential health threat of iAs, it is important to determine the levels of the individual species in foodstuffs – and not just the total As concentration. The US Food and Drug Administration (FDA) has established an action limit for iAs in apple juice of $10 \, \mu g/kg$ (ppb) [2] but there are currently no regulations in the US controlling the As content of wine. Canada (Vintners Quality Alliance VQA, Ontario) and Europe (International Organisation of Vine and Wine, OIV) have set maximum acceptable limits for total As in wine of $100 \, \text{and} \, 200 \, \mu g/L$ (ppb), respectively [3, 4].

Arsenic contamination of food is of great public interest. There is a clear demand for rapid and reliable screening methods to accurately determine the levels of iAs in food and drink to support existing and future regulations. One of the most useful and reliable approaches uses high performance liquid chromatography (HPLC) to separate the species, which are then quantified by inductively coupled plasma mass spectrometry (ICP-MS) [5].

The methodology described here is based on a previous As speciation method developed by Jackson, who coupled HPLC to a triple quadrupole ICP-MS (ICP-QQQ) [6]. HPLC-ICP-QQQ was also used in this study. However, instead of analyzing the iAs species separately, As(III) was intentionally oxidized to As(V) with hydrogen peroxide before analysis [7, 8]. By converting As(III) and analyzing all inorganic species as As(V), this method was able to separate monomethylarsonic acid (MMA) and dimethylarsinic acid (DMA) from iAs (as As(V)) in less than 2 minutes. The analysis time is 10 times faster than the current FDA methods used for the speciation of As [9].

In this work, oxygen reaction gas was used in the collision/reaction cell (CRC) of the ICP-QQQ to resolve the spectral interferences on ⁷⁵As, while maintaining excellent sensitivity. Results are presented that demonstrate the accuracy and reproducibility of the new method. The method was further validated using a wine matrix that was analyzed by two participating laboratories.

Experimental

Standards

The As(III) and As(V) standards were purchased from Spex Certiprep (Christiansburg, VA; Metuchen, NJ, USA). The MMA and DMA standards were purchased from Chem Service (West Chester, PA, USA). An AB standard was also purchased from Chem Service to be used as a flow injection marker (internal standard) for post-column injection. Calibration standards were prepared at 0.1, 0.5, 1.0, 5.0, 10 and 20 µg/L (ppb) for each of DMA, MMA, and total iAs (sum of As(III) and As(V)).

Samples

Five different California wines were used for the validation (V) study. Each wine represented one of the five main styles of wine: red, white, rosé, sparkling, dessert. Five additional California wines were analyzed for a commercial market basket (MB) study. Details of the wine style, cultivar, growing region, vintage, and alcohol content for all samples are given in Table 1.

Table 1. Wine style, cultivar, regional origin, vintage, and alcohol content of the wine samples for the validation and commercial market basket studies.

Sample	Style	Cultivar	Region	Vintage	Alcohol (%v/v)
V-1	Rosé	Zinfandel	Napa and Lodi	NA	9.5
V-2	White	Sauvignon blanc	Oakville/Napa County	2013	13.0
V-3	Sparkling	Sparkling white blend	County	NA	12.0
V-4	Dessert	Petite Sirah Port-style	Clarksburg/Yolo County	2012	20.0
V-5	Red	Cabernet Sauvignon	Monterey County	2013	14.5
MB-1	Red	Cabernet Sauvignon	North Coast	2009	13.5
MB-2	Red	Pinot noir	Appellation Central Coast	2004	13.8
MB-3	White	Chardonnay	Santa Barbara County	2013	13.5
MB-4	Rosé	Zinfandel	Napa and Sonoma	2013	10.5
MB-5	White	Chardonnay	Central Coast	2013	13.5

Sample preparation

 $\rm H_2O_2$ was added to all samples at a 1:1 ratio to oxidize As(III) to As(V). Each sample was further diluted with de-ionized water to give a total dilution factor of 5 or 6 (there were no differences in results between the two dilution factors). Each sample was then passed through a 0.45 μ m syringe filter to remove any particulates. Samples V-1, V-4, V-5 were spiked in duplicate with all As species at three concentration levels: 5, 10, and 30 μ g/kg.

Instrumentation

An Agilent 1260 HPLC fitted with a Hamilton PRP-X100 5 μ m 50 x 2.1 mm column was coupled to an Agilent 8800 Triple Quadrupole ICP-MS (ICP-QQQ). The mobile phase was 40 mM ammonium carbonate ((NH₄)₂CO₃, trace metal grade 99.999% from Sigma Aldrich) with 3% v/v methanol (Optima LC/MS grade, Fisher Chemical) adjusted to a pH of 9.0 with ammonium hydroxide (Optima Grade, Fisher Scientific). The ICP-QQQ was equipped with a standard sample introduction system comprising a quartz torch with 2.5 mm i.d. injector, a quartz spray chamber, glass concentric nebulizer, and nickel-tipped interface cones. Peak integration was carried out according to FDA EAM §4.10 and 4.11.15 [9]. The instrument operating conditions are summarized in Table 2.

Table 2. HPLC-ICP-QQQ operating conditions.

Forward power Sampling depth Spray chamber temp. Carrier gas Make-up gas Extract 1 Octopole bias Energy discrimination Cell gas (O ₂) flow rate Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	
Spray chamber temp. Carrier gas Make-up gas Extract 1 Octopole bias Energy discrimination Cell gas (O₂) flow rate Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	1550 W
Carrier gas Make-up gas Extract 1 Octopole bias Energy discrimination Cell gas (O₂) flow rate Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	8.0 mm
Make-up gas Extract 1 Detopole bias Energy discrimination Cell gas (O₂) flow rate Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	2 °C
Extract 1 Octopole bias Energy discrimination Cell gas (O₂) flow rate Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	0.95 L/min
Octopole bias Energy discrimination Cell gas (O₂) flow rate Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	0.20 L/min
Energy discrimination Cell gas (0₂) flow rate Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	0 V
Cell gas (O₂) flow rate Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	-5.0 V
Scan mode Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	-7 V
Q1/Q2 mass HPLC Mobile phase flow rate njection volume Sample temperature	0.31 mL/min
HPLC Mobile phase flow rate njection volume Sample temperature	MS/MS
Mobile phase flow rate njection volume Sample temperature	75/91 u
njection volume Sample temperature	
Sample temperature	0.5 mL/min
· · ·	5 μL
	4 °C
STD injection volume	5 μL

Results and Discussion

Development of a fast method

For this study, the focus of the method development was to reduce the analysis time per sample. In the development of this method, we followed Jackson's use of a small injection volume, short ion-exchange column, oxygen cell gas, and a high mobile phase linear velocity [6].

Figure 1 shows overlaid chromatograms for a representative calibration set of 0.5, 1.0, 5.0, and 20 μ g/kg standards. All As species are clearly separated in less than two minutes. Simply by oxidizing As(III) to As(V) and analyzing all iAs in the form of As(V), the analysis time was reduced significantly compared to the current FDA regulatory method [9].

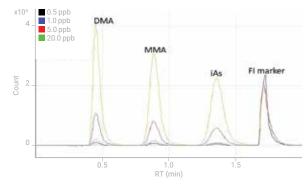


Figure 1. Overlay of the 0.5, 1.0, 5.0, and $20.0 \mu g/kg$ calibration standards. An AB internal standard (flow injection marker; fourth peak) was added post column via an external switching valve.

Linear calibrations

The calibration curves for DMA, MMA, and iAs show good linearity (Figure 2). All As concentrations in the wine samples were within the linear range except iAs, which was measured at a maximum concentration of 150% of the highest calibration standard.

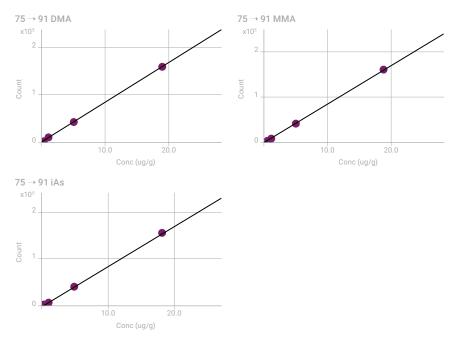


Figure 2. Calibration curves for DMA, MMA, and total iAs (sum of converted As(III) and As(V)).

Detection limits

The limits of detection (LOD) and limits of quantitation (LOQ) given in Table 3 are based on repeated measurements of the $0.05 \,\mu g/kg$ (ppb) mixed standard, n=15.

Table 3 I ODe	(3 sigma), LOQs	(30 ciama)	and actimated	wine I OO
i able 3. LODS	(3 Sigilia), LUQS	(SU Sigilla)	, anu estimateu	WITE LOQ.

	LOD, μg/kg	LOQ, μg/kg	Estimated wine LOQ, (6 x dilution) µg/kg
DMA	0.018	0.175	1.1
MMA	0.026	0.258	1.5
iAs	0.022	0.221	1.3

Spike recoveries

Samples V-1, V-4, V-5 were spiked in duplicate with each species (DMA, MMA, and total iAs as As(V)) at 5, 10, and 30 μ g/kg. The averaged recoveries for all As species at the three different fortification levels were 100 \pm 3% (Table 4).

Table 4. Percent recovery (mean and range) for three spiking levels of DMA, MMA and iAs in wines V-1, V-4 and V-5.

	DMA	MMA	iAs
Average, %	102	97	99
Range, %	97 – 107	91 – 102	95 – 103

Quantitative results

All 10 wines were analyzed using the new HPLC-ICP-QQQ method. Table 5 lists the measured concentrations for DMA and iAs. All MMA values were below the calculated LOD (0.026 μ g/kg) and could not be quantified. The measured concentrations using the new method were compared to the values obtained using the FDA EAM §4.10 extension method [10]. The agreement between the measurements was mostly within $\pm 10\%$. iAs represented the majority of As in all wines, while only one wine sample (MB-3) contained DMA levels significantly above the LOQ of 1.1 μ g/kg. A chromatogram of V-1 is shown in Figure 3.

Overall, the concentration of iAs ranged from 1.7 \pm 0.3 to 32.9 \pm 0.8 µg/kg (the latter being above the FDA's action limit for iAs in apple juice of 10 µg/kg). The sum of all As species (Table 5) ranged from a low of 2.2 \pm 0.3 µg/kg to a high of 32.9 \pm 0.8 µg/kg, which is under the Canadian limit of 100 µg/L and OIV limit of 200 µg/L.

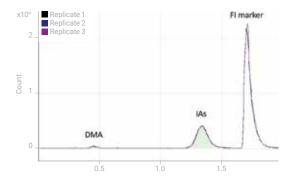


Figure 3. Chromatogram showing the overlay of the three replicates of wine sample V-1.

Table 5. Results from the fast and fit-for-purpose analysis method (measured at two different labs) compared to the FDA EAM §4.10 extension results for the five validation (V) and five market basket (MB) wines. % Recovery (shown in parentheses) calculated as "Measured" divided by "EAM §4.10" and "Sum of Species" divided by "Total".

Sample	DM	DMA (µg/kg) iiAs (µg/kg) Total		iiAs (μg/kg)		As (μg/kg)
	EAM §4.10	Measured	EAM §4.10	Measured	Total	Sum of Species
V-1	0.81 ± 0.1*	0.72 ± 0.04 (89%)	14.4 ± 1.0	16.0 ± 0.5 (111%)	16.5 ± 0.02	16.7 ± 0.5 (101%)
V-2	0.74 ± 0.04*	0.72 ± 0.06 (98%)	10.7 ± 0.2	11.4 ± 0.4 (107%)	12.6 ± 0.16	12.1 ± 0.3 (96%)
V-3	0.75 ± 0.1*	0.83 ± 0.04 (111%)	9.2 ± 0.4	9.5 ± 0.6 (103%)	10.4 ± 0.11	10.3 ± 0.5 (99%)
V-4	1.70 ± 0.1	1.86 ± 0.06 (109%)	2.1 ± 0.3	2.3 ± 0.4 (109%)	4.5 ± 0.01	4.1 ± 0.4 (92%)
V-5	0.45 ± 0.01*	0.47 ± 0.04 (105%)	1.5 ± 0.3	1.7 ± 0.3 (113%)	2.4 ± 0.03	2.2 ± 0.3 (90%)
MB-1	<lod< td=""><td><lod< td=""><td>30.2 ± 1.3</td><td>32.9 ± 0.8 (109%)</td><td>34.4 ± 0.4</td><td>32.9 ± 0.8 (96%)</td></lod<></td></lod<>	<lod< td=""><td>30.2 ± 1.3</td><td>32.9 ± 0.8 (109%)</td><td>34.4 ± 0.4</td><td>32.9 ± 0.8 (96%)</td></lod<>	30.2 ± 1.3	32.9 ± 0.8 (109%)	34.4 ± 0.4	32.9 ± 0.8 (96%)
MB-2	0.33 ± 0.04*	<lod< td=""><td>7.57 ± 0.49</td><td>9.1 ± 0.4 (120%)</td><td>9.1 ± 0.3</td><td>9.1 ± 0.4 (100%)</td></lod<>	7.57 ± 0.49	9.1 ± 0.4 (120%)	9.1 ± 0.3	9.1 ± 0.4 (100%)
MB-3	0.71 ± 0.08*	1.1 ± 0.0 (155%)	24.64 ± 0.40	27.6 ± 0.7 (112%)	28.9 ± 0.9	28.6 ± 0.7 (99%)
MB-4	1.16 ± 0.09*	1.0 ± 0.1 (86%)	26.3 ± 0.89	27.5 ± 0.9 (105%)	27.9 ± 0.9	28.5 ± 0.9 (102%)
MB-5	<lod< td=""><td><lod< td=""><td>3.5 ± 0.25</td><td>4.5 ± 0.1 (129%)</td><td>4.7 ± 0.1</td><td>4.5 ± 0.1 (96%)</td></lod<></td></lod<>	<lod< td=""><td>3.5 ± 0.25</td><td>4.5 ± 0.1 (129%)</td><td>4.7 ± 0.1</td><td>4.5 ± 0.1 (96%)</td></lod<>	3.5 ± 0.25	4.5 ± 0.1 (129%)	4.7 ± 0.1	4.5 ± 0.1 (96%)

Average \pm 1 σ , n=3. *Indicates value between LOD (0.17 μ g/kg) and LOQ (1.3 μ g/kg) for EAM §4.10 method. Refer to Table 3 for Measured LODs and LOQs

Conclusions

This note describes a simple, robust, and fast HPLC-ICP-QQQ method to measure the sum of the most toxic inorganic As species (As(III) and As(V)) and two organic As species in under two minutes. By oxidizing As(III) to As(V) with $\rm H_2O_2$ during sample preparation, total iAs can be determined as As(V), leading to a much faster separation of the species of interest in wine samples. The narrow bore column and 0.5 mL/min flow rate provided excellent sensitivity which allowed low volume injections to be used. Compared to the current FDA method for the determination of As in wines, sample run times were 10x faster with improved limits of detection and quantification.

In this study, total As concentrations ranged from 2.2 to 32.9 μ g/kg, which is well below the limit defined in regulations set in Ontario, Canada (100 μ g/kg) and the maximum level established by the International Organisation of Vine and Wine in Europe (200 μ g/kg). However, iAs was the predominant species present in the wines, and five of the wines tested contained iAs at concentrations that exceeded 10 μ g/kg, which is the FDA's action limit for iAs in apple juice.

The results obtained using the new fast and fit-for-purpose method were in good agreement with data obtained using the FDA's EAM §4.10.

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More Information

For a full account of this study, see Patrick J. Gray, Courtney K. Tanabe, Susan E. Ebeler, and Jenny Nelson, A fast and fit-forpurpose arsenic speciation method for wine and rice, *J. Anal. At. Spectrom.*, **2017**, 32, 1031–1034; DOI: 10.1039/C7JA00041C

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Fast Analysis of Arsenic Species in Infant Rice Cereals using LC-ICP-QQQ

Authors

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Introduction

Arsenic contamination of food can be harmful to human health. To assess the risk, several speciation methods have been developed to separate the toxic inorganic forms of As (iAs)—a class 1 carcinogen—from less toxic or non-toxic forms.

In a previous study (1), a speciation method specified in US FDA EAM: Section 4.11 (2) was used to separate four arsenic species in 31 baby rice cereals. The arsenic species included the inorganic forms; As(III) (arsenite) and As(V) (arsenate), and two organic forms; monomethylarsonic acid (MMA), and dimethylarsinic acid (DMA). The four species were separated using isocratic anion-exchange HPLC, and ICP-MS was used to detect arsenic-containing chromatographic peaks.

This study aimed to develop a rapid and reliable screening method for inorganic arsenic (iAs) analysis, to assist the food industry in meeting existing and future regulations. As shown in Table 1, the FDA has proposed an action limit of 100 ppb for iAs in infant rice cereals. This limit is in line with the European Union's limit for rice used for the production of food for infants and young children.

Table 1. Example regulatory maximum concentrations governing iAs in rice and rice-based products.

Regulating body	Action or Maximum Concentration for iAs (ppb)	Rice Type or Rice Product
US FDA (3)	100 (proposed)	Infant rice cereals
Codex Alimentarius	200	Polished (white) rice
Commission (4, 5)	350	Husked (brown) rice
European Union (6)	200	Polished rice
	250	Parboiled and husked rice
	300	Rice waffles, wafers, crackers, and cakes
	100	Rice used for the production of food for infants and young children
China (7)	150	Rice grains

The methodology described in this application note is based on a previous method developed by Jackson (8), where As species were determined using HPLC coupled to a triple quadrupole ICP-MS (ICP-QQQ). HPLC-ICP-QQQ was also used in this study, but instead of analyzing the iAs species separately, As(III) was intentionally oxidized to As(V) with hydrogen peroxide before analysis (9, 10). By converting As(III) and analyzing all inorganic species as As(V), this method was able to separate MMA and DMA from iAs (as As(V)) in less than 2 minutes. The analysis time is 10 times faster than the current FDA methods used for As speciation (2). The same fast HPLC-ICP-QQQ approach has also been applied to As speciation in wine (11).

Oxygen was used as a reaction gas in the collision/reaction cell (CRC) of the ICP-QQQ to resolve the Cl-based spectral interferences on As-75 for total As measurements. For the speciation measurements, the potential Cl-based interferences are resolved chromatographically, so ICP-QQQ with MS/MS is not essential. While this analysis could be done on a single quadrupole ICP-MS such as the Agilent 7800 or 7900 ICP-MS, ICP-QQQ offers higher sensitivity and lower detection limits where both As speciation and total As analysis is required. Results are presented that demonstrate the accuracy and reproducibility of the new method. The method was further validated by analyzing four rice standard and certified reference materials.

Experimental

Standards

The As(III) and As(V) standards were bought from Spex Certiprep (Metuchen, NJ, USA). The MMA and DMA standards were bought from Chem Service (West Chester, PA, USA). An arsenobetaine (AB) standard was also purchased from Chem Service to be used as a flow injection marker (internal standard) for post-column injection. Calibration standards were prepared at 0.1, 0.5, 1.0, 5.0, 10, and 20 μ g/L (ppb) for each of DMA, MMA, and total iAs (sum of As(III) and As(V)).

Standard/certified reference materials

Four SRM/CRMs were used as quality control materials for the As speciation measurements and total As measurements (without HPLC separation). The SRM used was the National Institute of Standards and Technology (NIST) 1568a Rice Flour. The three CRMs were the National Metrology Institute of Japan (NMIJ) 7503a White Rice Flour, the NMIJ 7532a Brown Rice Flour, and the Joint Research Centre (JRC) ERM-BC211 - Arsenic in Rice.

Samples and sample preparation

Six baby rice cereals were purchased from a local store in Berkeley, CA, USA. Each cereal was produced by a different manufacturer.

Arsenic was extracted from the rice matrix according to FDA method EAM 4.11 (2). Infant rice cereal (1 g) was weighed into a centrifuge tube and 10 mL of 0.28 mol/L $\rm HNO_3$ was added. The capped tube was placed in a preheated block digestion system at 95 °C for 90 minutes. The mixture was then diluted with 6.6 mL $\rm H_2O$, centrifuged, and filtered. Equal 0.5 mL portions of rice extract, $\rm H_2O_2$, and mobile phase were pipetted into a 2 mL plastic HPLC vial as the test solution. Each sample was prepared in duplicate.

Instrumentation

An Agilent 1260 HPLC fitted with a Hamilton PRP-X100 5 μ m 50 x 2.1 mm column was coupled to an Agilent 8800* Triple Quadrupole ICP-MS (ICP-QQQ). The mobile phase was 40 mM ammonium carbonate ((NH₄)₂CO₃, trace metal grade 99.999%, Sigma Aldrich) with 3% v/v methanol (Optima LC/MS grade, Fisher Chemical) adjusted to a pH of 9.0 with ammonium hydroxide (Optima Grade, Fisher Scientific). The ICP-QQQ was equipped with a standard sample introduction system comprising a glass concentric nebulizer, quartz spray chamber, quartz torch with 2.5 mm i.d. injector, and nickel-tipped interface cones. Peak integration was carried out according to FDA EAM §4.10 (12) and 4.11.15 (2). The instrument operating conditions are summarized in Table 2.

Table 2. HPLC-ICP-QQQ operating conditions.

ICP-QQQ	
Forward power	1550 W
Sampling depth	8.0 mm
Spray chamber temp.	2 °C
Carrier gas	0.95 L/min
Make-up gas	0.20 L/min
Extract 1	0 V
Octopole bias	-5.0 V
Energy discrimination	-7 V
O ₂ cell gas flow rate	0.31 mL/min
Scan mode	MS/MS
Q1/Q2 mass	75/91 u
HPLC	
Mobile phase flow	0.5 mL/min
Injection volume	5 μL
Sample temperature	4 °C
ISTD injection volume	5 μL

Results and Discussion

Development of a fast method

The focus of the method was to reduce the analysis time per sample compared to the current FDA method for As speciation. In common with Jackson's method (8), a small injection volume, short ion-exchange column, high mobile phase linear velocity, and oxygen cell gas mode were used.

Figure 1 shows overlaid chromatograms for a representative calibration set of 0.5, 1.0, 5.0, and 20 μ g/kg standards. All As species are baseline separated in less than two minutes. Simply by oxidizing As(III) to As(V) and analyzing all iAs in the form of As(V), the analysis time was reduced significantly compared to approximately 20 minutes for the current FDA regulatory method (2).

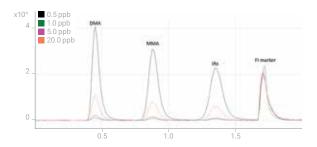


Figure 1. Overlay of the 0.5, 1.0, 5.0, and 20.0 μ g/kg As species calibration standards. An AB internal standard (flow injection marker; fourth peak) was added post column via an external switching valve.

Linear calibrations

The calibration curves for DMA, MMA, and iAs showed good linearity (Figure 2). All As concentrations in the rice samples were within the linear range except iAs, which was measured at a maximum concentration of 150% of the highest calibration standard.

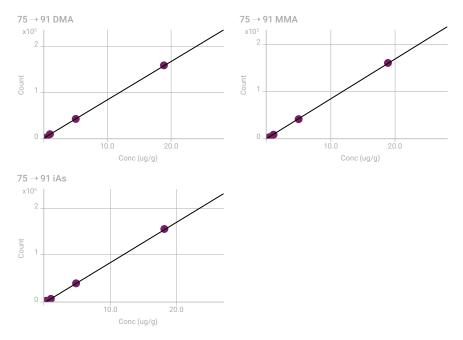


Figure 2. Calibration curves for DMA, MMA, and total iAs (sum of converted As(III) and As(V)).

Detection limits

The limits of detection (LOD) and limits of quantitation (LOQ) given in Table 3 are based on repeated measurements of the $0.05\,\mu\text{g/kg}$ (ppb) mixed standard, n=15.

Table 3. LOD (3 sigma)	100(30 ciams) and	d actimated LOO in rice	a for each Ac chacine

	LOD, μg/kg	LOQ, μg/kg	Estimated wine LOQ, (50 x dilution) μg/kg
DMA	0.018	0.175	8.8
MMA	0.026	0.258	12.9
iAs	0.022	0.221	11.0

Reproduced by permission of The Royal Society of Chemistry. P. J. Gray et al, J. Anal. At. Spectrom., 2017, 32, 1031 Analyis of rice RMs

Arsenic species were determined in four rice reference materials using the new HPLC-ICP-QQQ method. The total As concentration in each sample was also determined by direct ICP-QQQ analysis (no HPLC separation). Table 4 lists the reference and measured concentrations for DMA, MMA, iAs, and total As. Only one of the reference materials—NIST 1568b—had a reference value for MMA. The HPLC-ICP-QQQ measured concentrations were compared to the reference values, where available. Species recoveries ranged from 93 to 123% of their certified values when concentrations were above the LOQ. The recoveries for total As were also acceptable, ranging from 92 to 112%.

Table 4. Quantitative results for As species and total As in rice reference materials.

	DMA (mg/kg)		ММА (MMA (mg/kg)		iAs (mg/kg)		Total As (mg/kg)	
Rice RM	Reference	Measured	Reference	Measured	Reference	Measured	Reference	Measured	
NIST 1568b	180 ± 12	195 ± 4 (109%)	11.6 ± 3.5	14.9 ± 0.9 (128%)	92 ± 10	105 ± 1 (114%)	285 ± 14	315 ± 3 (110%)	
NMIJ 7503a	13.3 ± 0.9	15.4 ± 0.1 (116%)	None reported	<lod< td=""><td>84.1 ± 3a</td><td>79 ± 4 (94%)</td><td>98 ± 7</td><td>94 ± 4 (96%)</td></lod<>	84.1 ± 3a	79 ± 4 (94%)	98 ± 7	94 ± 4 (96%)	
NMIJ 7532a	18.6 ± 0.8	18.7 ± 1.3 (101%)	None reported	2.2 ± 1.9	298 ± 8	277 ± 12 (93%)	320 ± 10	297 ± 12 (93%)	
ERM BC-211	119 ± 13	146 ± 3 (123%)	None reported	19.9 ± 0.6	124 ± 11	124 ± 2 (100%)	260 ±13	290 ± 5 (112%)	

a. NMIJ 7503a iAs uncertainty estimated as the square root of the sum of squares of the AsIII and AsV uncertainties. Reproduced with permission of The Royal Society of Chemistry. P. J. Gray et al, J. Anal. At. Spectrom., 2017, 32, 1031.

To check the quality of the data, z-scores were also calculated. Z-scores are the number of standard deviations from the mean, with values between -3 and +3 being sufficient for regulatory purposes. The percent recovery for DMA in ERM BC211 RM was biased high, but the z-score was 2.1. The recovery for MMA in NIST 1568b RM was 128% but the reference concentration was below the method's LOQ. The z-score was 0.94.

Quantitative results in infant rice products

Six baby rice cereal samples were measured in duplicate using the HPLC-ICP-QQQ speciation method. Table 5 lists the measured concentrations for DMA and iAs; MMA was only present above the LOQ (0.026 ppb) in two of the rice samples (E and F). There was no significant difference between the two duplicates run for each cereal sample, showing the reproducibility of the method.

The concentration of iAs in four of the six rice samples was below the US FDA's proposed action limit and the EU's maximum limit of 100 ppb for iAs in infant rice cereals. Samples C and D exceeded the regulatory limit.

Table 5. Quantitative results in $\mu g/kg$ (ppb) for As species in six infant rice market basket samples measured in duplicate.

Sample Name	DMA	ММА	iAs	Proposed US FDA limit of 100 ppb for iAs
Baby rice cereal A_1	11.4	N/D	63.3	Door
Baby rice cereal A_2	11.2	N/D	62.3	— Pass
Baby rice cereal B_1	12.5	N/D	53.6	D
Baby rice cereal B_2	14.9	N/D	56.4	— Pass
Baby rice cereal C_1	33.9	N/D	106.4	F-9
Baby rice cereal C_2	36.0	N/D	113.5	— Fail
Baby rice cereal D_1	15.4	N/D	102.6	F-9
Baby rice cereal D_2	15.1	N/D	103.6	— Fail
Baby rice cereal E_1	41.9	2.2	87.9	
Baby rice cereal E_2	39.0	2.3	89.4	— Pass
Baby rice cereal F_1	46.4	8.7	89.4	D
Baby rice cereal F_2	46.7	9.0	90.4	— Pass

N/D = Not detected

Conclusions

A fast and fit-for-purpose HPLC-ICP-QQQ method is described for the measurement of inorganic As and two organic As species in baby rice cereal. A full speciation analysis can be completed in under two minutes.

- By oxidizing As(III) to As(V) with H2O2 during sample preparation, total iAs was determined as As(V).
- The narrow bore column and 0.5 mL/min HPLC flow rate provided excellent sensitivity, which allowed low volume injections to be used.
- Sample run times were 10x faster than the current FDA 4.11 method for the determination of As in rice.
- The HPLC-ICP-QQQ method delivered improved sensitivity, limits of detection and limits of quantification compared to the FDA 4.11 method.

The reproducibility of the method was demonstrated by the good agreement between the quantitative results for duplicate measurements of six rice cereal samples. The results showed that two of the samples contained iAs above 100 ppb.

This method provides valuable information for the safety of rice and rice-based infant cereals, as well as allowing food producers to meet regulatory requirements.

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More Information

For a full account of this study, see Patrick J. Gray, Courtney K. Tanabe, Susan E. Ebeler, and Jenny Nelson, A fast and fit-for-purpose arsenic speciation method for wine and rice, *J. Anal. At. Spectrom.*, **2017**, 32, 1031–1034; DOI: 10.1039/C7JA00041C

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Accurate Analysis of Trace Mercury in Cosmetics using the Agilent 8900 ICP-QQQ

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Effective removal of tungsten-based interferences on five Hg isotopes using MS/MS

Introduction

Many mercury (Hg) compounds are toxic, causing symptoms ranging from skin irritation, headaches, and tremors, through to nervous system damage, renal failure, and heart disease (1). Because Hg compounds are easily absorbed through the skin, their use in cosmetics is controlled. For example, the US Food and Drug Administration (FDA) does not allow Hg in cosmetics, except under specific conditions where there are no other safe and effective preservatives available (2). Increasingly, however, Hg has been found in cosmetic products such as skin creams, soap, and lotions sold as "anti-aging" or "skin lightening".

Mercury is a challenging element to determine at low levels by ICP-MS. It has a high first ionization potential (10.44 eV), so is relatively poorly ionized in the plasma, leading to low sensitivity. Also, Hg has seven naturally occurring isotopes, each with relatively low % abundance, further reducing sensitivity. Many forms of Hg are also volatile, and the element's chemistry means that it can be difficult to stabilize in solution. To address these issues, analysts must control the acid mix used for sample preservation and rinse solutions, to avoid problems with poor linearity and long washout times. Despite these difficulties, ICP-MS can still be used successfully to perform trace-level analysis of Hg, if appropriate sample stabilization—for example with the addition of HCI—is used.

Trace-level mercury analysis is even more difficult in samples—including some cosmetics—that contain a high concentration of tungsten (W). The W matrix forms polyatomic ions WO+ and WOH+ that overlap all the Hg isotopes, making Hg measurement even more challenging. For example, the most abundant Hg isotopes, ²⁰⁰Hg and ²⁰²Hg, suffer interferences from ¹⁸⁴W¹⁶O+ and ¹⁸⁶W¹⁶O+, respectively. Collision/reaction cells (CRCs) are used successfully to control many common polyatomic interferences in conventional single quadrupole ICP-MS (ICP-QMS). However, even with CRC operation, ICP-QMS cannot reduce the WO+ and WOH+ interferences sufficiently to allow the accurate determination of Hg at trace levels in samples that contain a high level of W.

The superior interference removal capability of triple quadrupole ICP-MS (ICP-QQQ) was investigated for this application. ICP-QQQ has dramatically improved the performance of reaction cell methods by using two mass-selection steps (MS/MS), one before and one after the CRC. In MS/MS, reaction chemistry in the CRC is controlled and consistent because only the target analyte mass enters the CRC. This capability offers a much more predictable and reliable approach to resolving interferences on a wide range of elements, particularly in complex and variable samples (3–5).

In this study, an Agilent 8900 ICP-QQQ was used for the measurement of Hg in a tungsten-rich cosmetic sample.

Experimental

Standards and samples

Mercury standards were prepared in 0.5 % high purity hydrochloric acid (TAMA-Pure-AA-100, Kanagawa, Japan).

A tungsten-rich cosmetic toning lotion was bought from a local store in Shanghai. The liquid sample was weighed to the nearest 0.100 g, and then diluted 100-fold with de-ionized water acidified with 0.5 % HCl to ensure Hg stability. The sample was shaken for a couple of minutes to ensure it was fully homogenized. The concentration of W in the original cosmetic sample was about 4000 mg/kg (ppm), as determined by ICP-QQQ in a diluted sample. Therefore, the W matrix in the sample as analyzed was about 40 mg/L (ppm) after the 100x dilution.

Instrumentation

An Agilent 8900 Standard configuration ICP-QQQ was used. The instrument was fitted with the standard sample introduction system comprising a glass concentric nebulizer, quartz double-pass spray chamber, quartz torch with 2.5 mm id injector, and Ni interface cones. The ICP-QQQ was operated in no gas mode, with He cell gas, and with $\rm O_2$ cell gas in both single quad (SQ) and MS/MS modes. The main operating conditions are shown in Table 1.

Table 1. 8900 ICP-QQQ operating conditions.

Parameter	No gas	He	02	02	
Acquisition mode		Single Quad		MS/MS	
RF power (W)		155	50		
Sampling depth (mm)	8.0				
Carrier gas flow rate (L/min)	0.8				
Make-up gas flow rate (L/min)		0.	4		
Spray chamber temp. (°C)	2				
He cell gas flow rate (mL/min)	- 5.0 -				
O ₂ cell gas flow rate (mL/min)	- 0.9				

Results and Discussion

As the most abundant isotope, 202 Hg is selected as the preferred isotope for ICP-MS measurements. However, some analysts select 201Hg instead (or as well), as the 201 isotope has proportionally lower W-based interference. The Hg calibration was prepared in a matrix of dilute (0.5 to 1.0%)

HCl to ensure that the Hg remained stable in solution as a Cl-complex. The calibration plots for ²⁰¹Hg and ²⁰²Hg are shown in Figure 1. The figures of merit—linearity, detection limit (DL), and background equivalent concentration (BEC)—taken from the ²⁰²Hg calibration are presented in Table 2.

Table 2. DL, BEC, and R value of the calibration curve of ²⁰²Hg in dilute HCl determined in four different cell modes.

	8900 calibration performance figures of merit for ²⁰² Hg			
	No gas	He	O ₂ Single Quad	O ₂ MS/MS
R	0.997	0.999	0.999	0.999
DL (µg/L)	0.002	0.001	0.002	0.002
BEC (μg/L)	0.011	0.008	0.003	0.003

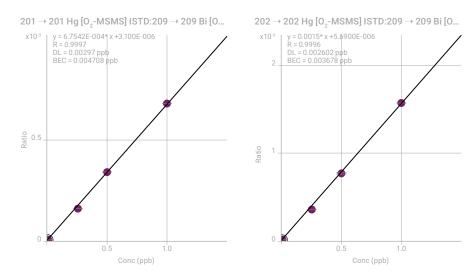


Figure 1. Calibration plots for ²⁰¹Hg and ²⁰²Hg, demonstrating good sensitivity and linearity due to effective stabilization of Hg by the addition of HCl.

ICP-MS/MS reaction mechanism used to resolve W-based interferences on Hg

The general reaction mechanism using MS/MS mode with $\rm O_2$ cell gas to resolve the WO+ and WOH+ interferences on Hg is shown in Figure 2. Since 200 Hg suffers the most serious polyatomic ion overlaps, the reaction mechanism is illustrated using the example of the 184 W 16 O+ overlap on 200 Hg+. Q1 is set to m/z 200, so 200 Hg+ and WO+ ions at m/z 200 pass through Q1 and enter the CRC. WO+ reacts with the $\rm O_2$ cell gas to form WO $_2$ + and WO $_3$ +, shifting to higher masses. The 200 Hg+ ions do not react with the $\rm O_2$ cell gas and so remain at m/z 200. By setting Q2 to m/z 200, 200 Hg+ ions pass to the detector free of interference. The same reaction mechanism is effective at resolving the WOH+ interferences, as WOH+ also reacts with the $\rm O_2$ cell gas to form higher-order product ions.

Multiple isotope analysis study

To investigate the effectiveness of interference removal in the different cell gas modes, a 1 μ g/L (ppb) Hg spike was added to the diluted W-rich cosmetic lotion sample. The five most abundant Hg isotopes were measured in the four different cell gas modes, and the isotopic ratios calculated. Comparing the measured isotope ratios with the theoretical natural ratios gives an excellent indication of the effectiveness of the interference-removal on each isotope. This capability is important for many ICP-MS applications, where the results calculated from a second isotope can be used to confirm the concentration reported using the primary or preferred isotope. Performing "confirmatory measurements" is recommended or required in several regulated methods across the environmental, food, and pharmaceutical industries. This approach is analogous to the use of "qualifier ions" in organic mass spectrometry.

The isotope ratios for several Hg isotope pairs measured in the W-rich cosmetic sample using the four cell gas modes are presented in Table 3. These results show that MS/MS mode with O_2 cell gas gives measured Hg ratios that are virtually identical to the theoretical natural ratios. O_2 in MS/MS mode is much more effective than the other modes for the removal of the tungsten oxide and hydroxide polyatomic interferences. The effective removal of the WOH+ overlap on 201 Hg is illustrated by the accurate ratios obtained for the 200 Hg/ 201 Hg ratio in MS/MS mode using O_2 cell gas.

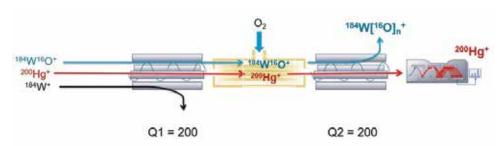


Figure 2. Reaction mechanism in MS/MS mode with O_2 cell gas for removal of WO $^+$ overlap to allow on-mass detection of Hg in a tungsten matrix. The same mechanism is also effective at resolving WOH $^+$ overlaps.

Table 3. Hg isotope ratios in tungsten-rich sample measured in different cell modes.

Hg ratio	Natural value	8900 measured results			
		No gas	He	O ₂ Single Quad	O ₂ MS/MS
198/199	0.591	1.738	1.769	1.435	0.598
198/200	0.432	0.831	0.823	0.739	0.43
200/201	1.75	49.4	61.4	7.75	1.76
201/202	0.441	0.022	0.017	0.126	0.445

As a further illustration of the ability of MS/MS to resolve interferences on multiple Hg isotopes, a scan spectrum comparison was made using on-mass measurement (Q1 = Q2). The mass range of the Hg isotopes was acquired for a simple Hg standard (1 μ g/L) and a solution containing the same concentration of Hg spiked into a high W matrix (10 mg/L). The overlaid spectra are shown in Figure 3, together with the template indicating the natural abundance of the Hg isotopes. The spectra show that the measured isotopic abundances match the natural Hg isotope pattern in both samples. This confirms the ability of MS/MS mode with O_2 cell gas to remove the W-based overlaps caused by the high W matrix in the second sample.

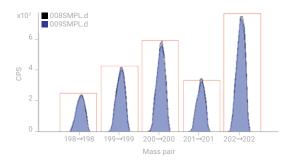


Figure 3. Hg isotopes with W matrix (blue shading) and without W matrix (gray shading), confirming accurate Hg isotope abundances and effective removal of W-based interferences on all Hg isotopes by ICP-QQQ with MS/MS.

Hg in tungsten-rich samples

Hg was measured in a tungsten-rich cosmetic sample using four different cell gas modes. The five most abundant isotopes of Hg (198, 199, 200, 201, and 202) were used for quantitation, giving five independently calibrated results for total Hg. The results for Hg in the original cosmetic sample, corrected for the 100 times dilution, are shown in Table 4.

Table 4. Apparent Hg concentration (μ g/kg) in tungsten-rich cosmetic sample, quantitated independently using five isotopes. The data shows errors due to the contribution from WO+ and WOH+ overlaps in no gas, He, and O₂ (SQ) cell modes.

	8900 measured results					
Cell mode	198	199	200	201	202	
No gas	267000	872000	126400	4770	98700	
He	173200	57700	90700	2460	66500	
O ₂ Single Quad	772	249	399	2.8	288	
O ₂ MS/MS	2.1	1.8	2.4	1.5	1.7	

Matrix-based interferences affect different isotopes of an analyte to different degrees, so giving different errors in the quantitative results calculated from each isotope.

Comparing the elemental concentrations calculated from the different isotopes of an element can therefore be used to identify whether the reported concentrations were affected by interferences. The reported Hg concentrations in MS/MS mode with O_2 (~2 μ g/kg in the original sample, or 0.02 μ g/L in the 100x diluted solution) are much lower than the results reported using the other cell modes. Also, the good agreement between the results obtained for the five isotopes in O2 (MS/MS) mode shows that this mode can simultaneously remove polyatomic interferences from all five Hg isotopes. These results contrast with the other modes, where incomplete removal of interferences from most of the isotopes led to erroneously high values and large differences between the results calculated using the different isotopes. The reported concentration of 2.8 µg/kg obtained for ²⁰¹Hg using O₂ cell gas in single-quad mode shows that the WOH+ interference could be reduced reasonably effectively. However, the other isotopes gave variable results in O₂(SQ) mode, so the ²⁰¹Hg result could not be verified by comparing it with a second, qualifier isotope. The data in Table 4 shows that, even when a suitable reaction gas is identified, MS/MS is essential for full control of the reaction chemistry.

Spike recovery test

A spike recovery test was carried out to further evaluate the interference removal capability and matrix tolerance of the method. Since ²⁰⁰Hg suffers the most serious polyatomic ion overlaps, it was selected as the target mass for the spike recovery test.

A 30 ppt spike of Hg was added to the diluted cosmetic lotion sample. The spike recovery in O_2 (MS/MS) mode was 104%, confirming the interference removal capability and matrix tolerance of the method (Table 5).

Table 5. 30 ppt Hg spike recovery results in different cell gas modes.

	Mode	Sample (µg/L)	Spike recovery (%)
200 Hg	No gas SQ	1364	6635
200 Hg	He SQ	906	879
200 Hg	O ₂ SQ	3.99	216
200 🛮 200 Hg	O ₂ MS/MS	0.024	104

Conclusions

The Agilent 8900 ICP-QQQ operating in MS/MS mode with $\rm O_2$ cell gas is highly effective for the removal of tungsten oxide/hydroxide polyatomic interferences on the five major Hg isotopes.

- Hg was measured accurately and consistently at trace levels in the presence of W using an MS/MS on-mass method with O₂ reaction cell gas.
- Compared to conventional single quadrupole ICP-MS, ICP-MS/MS reduced interferences by more than two orders of magnitude.
- The ICP-MS/MS method easily meets the requirements of trace level Hg analysis in tungsten-rich cosmetic samples.

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The Accurate Measurement of Selenium in Reference Materials using Online Isotope Dilution

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Keywords

selenium, environmental, agricultural, human health, online isotope dilution analysis, OIDA, oxygen mass-shift

Introduction

Selenium (Se) is an important element in environmental and agricultural studies and in human health, as it is an essential trace nutrient but is toxic in excess. The role of certain chemical forms of Se is the subject of significant research into cancer prevention. ICP-MS is the analytical method of choice for both total and speciated Se measurements, but Se is a difficult element to quantify accurately at trace levels by ICP-MS for several reasons:

- The signal for Se is low, since it is poorly ionized in the plasma due to its high Ionization Potential (IP) of 9.75 eV.
- Because Se is poorly ionized, it suffers signal suppression in high matrix samples — an issue that is further compounded by the lack of a suitable internal standard element with a similar mass and IP.
- All the analytically useful Se isotopes suffer from multiple spectral interferences, as summarized in Table 1.
- The resolution required to separate all of the spectral interferences is beyond the capabilities of sector-type high resolution (HR-)ICP-MS.

The Agilent 8800 ICP-QQQ in MS/MS mode has a unique ability to remove the complex spectral interferences from all the Se isotopes shown in Table 1, allowing the use of Isotope Dilution (ID) analysis, which requires at least two interference-free isotopes. ID is the most accurate quantification technique as it is based on direct measurement of isotopic abundances in each sample, rather than a relative measurement of analyte response compared to a standard. As a result, it offers better traceability and improved correction of non-spectroscopic interferences encountered in high matrix sample analysis. This note describes the application of the Agilent 8800 ICP-QQQ using ID for the accurate quantification of Se in a range of certified reference materials (CRMs).

Table 1. Spectral interferences on Se isotopes.

	Se isotope				Interferen	ce		
Mass	Abundance %	Isobaric	Argide	Oxides	Hydride	Chloride	Doubly charged	Dimer
77	7.63		³⁹ K ³⁸ Ar+	⁶¹ Ni ¹⁶ O+, ⁵⁹ Co ¹⁸ O+	⁷⁶ GeH⁺, ⁷⁶ SeH⁺	⁴⁰ Ar ³⁷ Cl ⁺ , ⁴⁰ Ca ³⁷ Cl ⁺	¹⁵⁴ Sm ⁺⁺ , ¹⁵⁴ Gd ⁺⁺	
78	23.77	⁷⁸ Kr ⁺	⁴⁰ Ca ³⁸ Ar ⁺	⁶² Ni ¹⁶ O+	⁷⁷ SeH⁺	⁴¹ K ³⁷ Cl ⁺	¹⁵⁶ Gd ⁺⁺ , ¹⁵⁶ Dy ⁺⁺	³⁸ Ar ⁴⁰ Ar ⁺ , ³⁹ K ³⁹ K ⁺
80	49.61	⁸⁰ Kr ⁺	⁴⁰ Ca ⁴⁰ Ar ⁺	64Ni ¹⁶ O+, 64Zn ¹⁶ O+, ³² S ₂ ¹⁶ O+, ³² S ¹⁶ O ₃ +	⁷⁹ BrH ⁺	⁴⁵ Sc ³⁵ Cl ⁺	¹⁶⁰ Gd ⁺⁺ , ¹⁶⁰ Dy ⁺⁺ ,	⁴⁰ Ar ⁴⁰ Ar ⁺ , ⁴⁰ Ca ⁴⁰ Ca ⁺
82	8.73	⁸² Kr ⁺	⁴² Ca ⁴⁰ Ar ⁺	⁶⁶ Zn ¹⁶ O+	⁸¹ BrH ⁺	⁴⁵ Sc ³⁷ Cl ⁺	¹⁶⁴ Dy ⁺⁺ , ¹⁶⁴ Er ⁺⁺	

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/General purpose. **Ion lens tune:** Soft extraction tune: Extract 1 = 0 V,

Extract 2 = -180 V.

CRC conditions: O_2 gas at 0.4 mL/min plus H_2 gas at 2.0 mL/min, Octopole bias = -18 V and KED = -6 V.

Acquisition parameters: MS/MS O_2 mass-shift method. The reaction of Se⁺ with O_2 to form SeO⁺ is endothermic (Δ Hr = 0.71 eV), but the reaction is efficiently promoted using high collision energy using a low octopole bias voltage setting [1]. Preliminary studies have shown that low BEC for Se isotopes can be achieved via the addition of a small amount of H₂ in MS/MS O_2 mass-shift method.

Method: Online isotope dilution analysis (OIDA) [2] was used. OIDA is a useful development of traditional isotope dilution, as it removes the time consuming step of spiking enriched-isotope standards into each individual sample. A ⁸²Se enriched standard purchased from Oak Ridge National Laboratory (USA) was prepared at the appropriate concentration and added via the standard online ISTD mixing kit to the samples. Product ions derived from the ¹⁶O-atom addition transition were measured for the three most analytically useful isotopes of Se. On the 8800 ICP-QQQ, this is simply achieved by defining the acquisition method with Q1/Q2 settings: Q1=78/Q2=94, Q1=80/Q2=96 and Q1=82/Q2=98 for the Se isotopes at *m/z* 78, 80 and 82 respectively.

It should be noted that the use of MS/MS (where Q1 acts as a 1 amu mass filter) is essential for this measurement, as it ensures that only one Se isotope enters the cell for any given mass pair measurement, and only the ¹⁶O atom addition is measured because the mass difference between Q1 and Q2 is 16 amu. This ensures that there is no overlap due to the precursor ions from different Se⁺ isotopes giving SeO⁺ product ions at the same mass, such as the ⁸⁰Se¹⁸O⁺ product ion overlap on ⁸²Se¹⁶O⁺, both of which appear at *m/z* 98. ICP-QQQ in MS/MS mode thereby removes one of the critical limitations of reaction chemistry with ICP-QMS, where all the sample ions enter the cell together so no specific reaction transition can be defined. Each Se isotope mass pair was measured with an integration time of 1 s and three replicates.

Sample preparation: The CRMs were microwave digested using a Milestone ETHOS closed vessel microwave digestion system (Milestone, Sorisole, Italy) and following the manufacturer's recommended procedures. The final dilution factor of the samples varied from 250 to 500x.

Results and Discussion

Study of cell gases for spectroscopic interference removal

Figure 1 shows the result of a preliminary study of the effects of the choice of cell gas on interference removal. The findings of the study showed that $\rm O_2/H_2$ mass-shift (Figure 1) enables the measurement of ⁷⁸Se, ⁸⁰Se and ⁸²Se relatively free from interferences in the range of synthetic matrices tested.

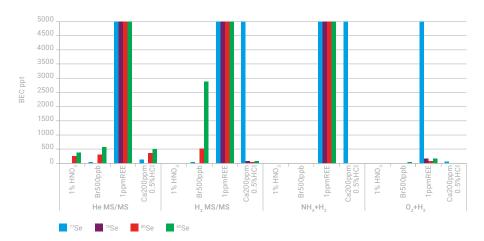


Figure 1. Preliminary study of the effectiveness of different cell gases for interference removal. Four synthetic matrices likely to give rise to interferences on the Se isotopes were measured using each of the 4 different cell gas modes.

Measurement of Se in CRMs

The concentration of Se was determined in 12 different CRMs using the OIDA method. The CRMs were obtained from NIST (Gaithersburg MD, USA), GSJ Geochemical Reference Samples (Tokyo, Japan), Japan Society for Analytical Chemistry (Tokyo, Japan), and National Institute of Metrology (Beijing, China). The matrices included environmental waters (NIST 1643e and JASC 0302-3 River Water), rock (JB-3 basalt), sedimentary rock (JSI-1 and NIST 1646a Estuarine Sediment), soil (JSAC0411 Volcanic Ash Soil), biological samples (NIST 1566a Oyster Tissue, NCSZC 81002 Human Hair, NIST 2976 Mussel Tissue), and plant materials (NIST 1575a Pine Needles, NIST 1515 Apple Leaves, NIST1573a Tomato Leaves).

Figure 2 shows the Se results for each CRM expressed as % recovery relative to the certified value. The measured results for Se were in good agreement with the CRM values (90%-112%), using two Se isotope pairs: 78/82 and 80/82. This demonstrates the effectiveness of the Agilent 8800 ICP-QQQ in MS/MS mode for the removal of multiple interferences on ⁷⁸Se, ⁸⁰Se and ⁸²Se.

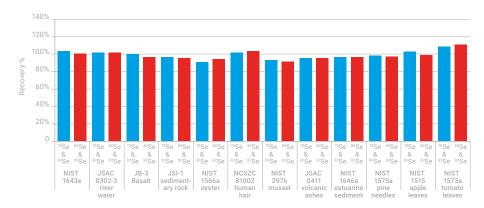


Figure 2. Result of Se quantification using OIDA in various CRMs.

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On-line isotope dilution analysis with the 7700 Series ICP-MS: Analysis of trace elements in high matrix samples, Giuseppe Centineo, Jose Angel Rodriguez Castrillon and Esther Munoz Agudo, 2011, Agilent application note, 5990-9171EN

More Information

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Accurate Sulfur Quantification in Organic Solvents using Isotope Dilution Mass Spectrometry

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Based upon the published work "Accurate determination of S in organic matrices using isotope dilution ICP-MS/MS" *J. Anal., At. Spectrom.* 2012 DOI: 10.1039/c2ja30265a by:

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Keywords

sulfur, ID-MS, biodiesel, environmental, ethanol, NIST SRM 2773, oxygen mass-shift

Introduction

Accurate measurement of sulfur in aqueous and organic media is relatively difficult for ICP-MS due to intense spectral interferences from polyatomic ions formed mainly from oxygen and nitrogen. Sulfur is an important element in environmental terms as it forms ${\rm SO}_{\rm x}$ when combusted, contributing to acid rain and photochemical smog. It is also a catalyst poison for some industrial processes and its accurate measurement can be critical.

Experimental

A quadrupole ICP-MS (ICP-QMS) with a collision/reaction cell set up for $\rm O_2$ mass-shift reaction chemistry can be used to avoid the $^{16}\rm O_2^+$ overlap on $^{32}\rm S^+$ by converting the S+ to SO+ reaction product ions that are then measured at a new mass (m/z 48) that is free from the $\rm O_2^+$ overlap. However, in practice, this approach has been of relatively limited use, as ICP-QMS has no way to reject existing ions at the mass of the new analyte product ions, so not all of the interferences are eliminated, particularly when complex or variable matrices are investigated. There has also been some limited success reported by using Xe as a reaction gas to attenuate the $\rm O_2^-$ -based interference particularly on the $^{34}\rm S$ isotope. Neither of these approaches reduces the backgrounds significantly enough to allow reliable trace level measurement of S, and they do not necessarily preserve the S isotopic abundances. In this investigation, ethanol was used as an example organic solvent and the Agilent 8800 ICP-QQQ was used to determine S by ID-MS in a biodiesel reference material to assess the measurement accuracy of MS/MS mode with $\rm O_2$ mass-shift for S determination.

Instrumentation: Agilent 8800 #100 with Micromist nebulizer (free aspiration). For organic solvent analysis, a narrow injector torch with id 1.0 mm (G3280-80005) and Pt cones were used. 20% $\rm O_2$ balanced in Ar was introduced via an option gas flow line to prevent carbon build up.

Plasma conditions: Plasma conditions were optimized manually. (RF power = 1450 W, Carrier gas flow rate = 0.98 L/min, Option gas flow rate = 0.75 L/min and spray chamber temp. = -5°C).

CRC conditions: O_2 gas at 0.4 mL/min, Octopole bias = -9 V, KED = -8 V.

Sample: Biodiesel certified reference material NIST SRM 2773.

Results and Discussion

When using mass-shift mode for sulfur (or any element) it is important to eliminate any potential interferences at the target mass of the reaction product ion, as well as on the primary element mass (the precursor ion). If the target mass suffers from interferences then the measurement would still be compromised. For sulfur, the corresponding isotopes are shifted as follows using M + 16 amu mass-shift:

32S I SO at 48 amu

33S I SO at 49 amu

34S I SO at 50 amu

Unfortunately, the SO+ product ion masses (m/z 48, 49 and 50) can suffer from multiple interferences including Ca+, Cr+, V+, Ti+, ArC+ and CCl+ in natural samples. Furthermore the 33 S and 34 S isotopes can suffer from overlaps due to other combinations of SO+ product ions, as well as pre-existing ions at the target mass. For example, the 34 S 16 O+ product ion formed at m/z 50 is overlapped by 32 S 18 O+ and 33 S 17 O+, as well as 50 Cr+, 50 V+, 50 Ti+, 38 Ar 12 C+, and 13 C 37 Cl+. When operating the 8800 ICP-QQQ in MS/MS mass-shift mode, these overlaps are eliminated and the sulfur isotope pattern is preserved. Figure 1 provides a graphical representation of the ICP-QQQ setup and the method of interference elimination.

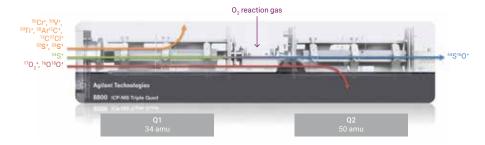


Figure 1. Mechanism of MS/MS mass-shift for sulfur isotope analysis. The mass difference between Q1 and Q2 is fixed at 16 amu, so only the + 16 O-atom transition is observed – the other oxygen isotope transitions are eliminated so the original sulfur isotopic pattern is preserved.

This method would not be useful if the reaction were not quantitative, so to check for linearity, a blank ethanol sample was spiked with sulfur – see Figure 2. Despite the wide variation in absolute sensitivity for the different S isotopes, the BEC was the same for all three isotopes, indicating that the background is due to sulfur in the ethanol.

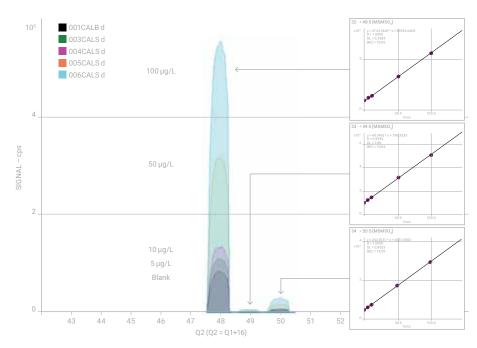


Figure 2. Ethanol with 0, 5, 10, 50 and 100 µg/L sulfur spikes and corresponding calibration curves.

An isotope dilution (ID) method was used to evaluate the accuracy of the 8800 ICP-QQQ MS/MS method, using a biodiesel certified reference material (NIST SRM 2773) and an enriched ³⁴S spike. The biodiesel sample was simply diluted into the ethanol solvent and the appropriate spike added. Reproducibility was tested by analyzing three separate samples of the CRM. The results are presented in Table 1. Repeat measurements were within the expected recovery limits for the material.

Table 1. Isotope dilution analysis of S in diluted biodiesel reference material NIST 2773.

Sample	S conc.(µg/g)
SRM 2773 - Certified	7.39 ± 0.39
SRM 2773 - measured 1	7.234
SRM 2773 - measured 2	7.227
SRM 2773 - measured 3	7.231
Average (measured)	7.231
Standard Deviation	0.003
95% confidence interval	7.231 ± 0.015

Conclusions

Until the introduction of ICP-QQQ with MS/MS capability, it was impossible to obtain reliable results for reaction chemistry methods combined with an ID approach, using a quadrupole-based ICP-MS. The novel QQQ configuration of the 8800 ICP-QQQ enables operation in MS/MS mode, which ensures precise control over the reaction chemistry in the cell. This allows the unique isotopic information of the analyte to be retained, while removing the interferences that could affect both precursor and product ions of the target analyte.

Removal of REE⁺⁺ Interference on Arsenic and Selenium

Authors

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Keywords

Rare Earth Elements, REE, arsenic, selenium, environmental, food, CRMs, oxygen mass-shift

Introduction

Trace analysis of arsenic (As) and selenium (Se) in environmental and food samples is of a great interest, since both elements can be toxic even at quite low levels. It is difficult to quantify As and Se accurately at trace levels in some matrices by quadrupole ICP-MS as all the analytically useful isotopes can suffer from multiple spectral interferences, as summarized in Table 1. This application investigates ICP-QQQ in MS/MS reaction mode to remove interferences on As and Se, with an emphasis on the removal of the doubly-charged ions arising from Rare Earth Elements (REE++). While the concentration of REEs in environmental and food samples is usually low, some plants will accumulate REEs from the soil, and a high concentration will lead to false positive results for As and Se.

Table 1. Spectroscopic interferences on As and Se isotopes.

	As and Se	isotope		Interference		
Element	Mass	Abundance %	Doubly charged	Matrix	Dimer	
As	75	100	¹⁵⁰ Sm ⁺⁺ , ¹⁵⁰ Nd ⁺⁺	⁴⁰ Ar ³⁷ Cl ⁺ , ⁴⁰ Ca ³⁷ Cl ⁺		
Se	77	7.63	¹⁵⁴ Sm ⁺⁺ , ¹⁵⁴ Gd ⁺⁺	⁴⁰ Ar ³⁷ Cl ⁺ , ⁴⁰ Ca ³⁷ Cl ⁺		
	78	23.77	¹⁵⁶ Gd ⁺⁺ , ¹⁵⁶ Dy ⁺⁺	⁴¹ K ³⁷ CI ⁺	³⁸ Ar ⁴⁰ Ar ⁺ , ³⁹ K ³⁹ K ⁺	
	80	49.61	¹⁶⁰ Gd ⁺⁺ , ¹⁶⁰ Gd ⁺⁺ ,	⁴⁵ Sc ³⁵ Cl ⁺	⁴⁰ Ar ⁴⁰ Ar ⁺ , ⁴⁰ Ca ⁴⁰ Ca ⁺	
	82	8.73	¹⁶⁴ Dy ⁺⁺ , ¹⁶⁴ Er ⁺⁺	⁴⁵ Sc ³⁷ Cl ⁺		

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/Low matrix.

Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -180 V.

CRC conditions: O_2 gas flow rate of 0.2 mL/min, Octopole bias = -8 V and KED =

-6 V.

Acquisition parameters: MS/MS O_2 mass-shift method to measure As⁺ (as AsO⁺) and Se⁺ (as SeO⁺), as illustrated in Figure 1. Unlike conventional quadrupole ICP-MS, the 8800 ICP-QQQ mass-shift method can be applied to complex matrix samples that may contain Zr and/or Mo. The MS/MS configuration prevents undesired ions such as 91 Zr⁺ and 94 Mo⁺ from overlapping the MO⁺ product ions, as they are rejected by Q1.

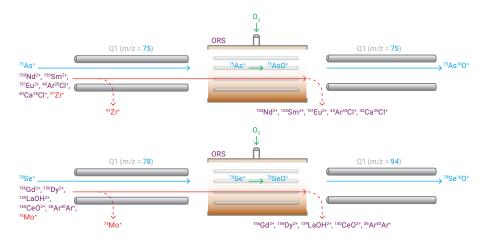


Figure 1. ICP-QQQ MS/MS 0, mass-shift method for measuring 75As (top) and 78Se (bottom)

Samples and sample preparation: SPEX XSTC-1 (a mixture of 10 ppm each of Ce, Dy, Er, Eu, Gd, Ho, La, Lu, Nd, Pr, Sm, Sc, Tb, Tm, Yb and Y) purchased from SPEX CertiPrep Ltd. (UK) was used. Four certified reference materials (CRMs): NIST 1515 Apple Leaves, NIST 1573a Tomato Leaves, NIST 1575a Pine Needles and NMIJ 7531a Brown Rice, were used for the method validation. It should be noted that NIST 1515 contains 3 mg/kg Sm and Gd, and 0.2 mg/kg Eu. NIST 1573a contains 0.19 mg/kg Sm, 0.17 mg/kg Gd, 5% Ca and 2.7% K, a combination of matrix elements that might be expected to cause severe interferences on As and Se. All CRMs were microwave-digested in HNO $_3$ and H $_2$ O $_2$, diluted and analyzed.

Results and Discussion

Effectiveness of O₂ mass-shift method for removing REE** interferences

To investigate the effectiveness of interference removal modes on the 8800 ICP-QQQ, As and Se were measured in a mixed REE solution containing 1 ppm each of Ce, Dy, Er, Eu, Gd, Ho, La, Lu, Nd, Pr, Sm, Sc, Tb, Tm, Yb and Y. Three different 8800 ICP-QQQ cell modes were used:

- Single Quad (SQ); no gas
- Single Quad (SQ); reaction mode using hydrogen (H₂) cell gas
- MS/MS; reaction mode using O₂ cell gas with + 16 amu mass-shift

[&]quot;Single Quad" represents the performance of conventional ICP-QMS while MS/MS mode is unique to ICP-QQQ.

Figure 2 shows the BECs of As and Se in each of the measurement modes. The results in Figure 2 illustrate the excellent interference removal performance of the O₂ mass-shift method for the detection of As and Se in a matrix containing REEs.

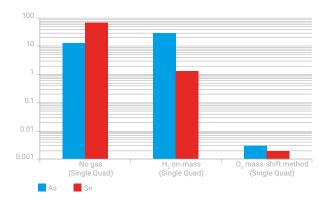


Figure 2. BEC of As and Se in 1ppm REE mixed solution with three measurement modes: no gas, H_2 on-mass and O_2 mass-shift mode.

Figure 3 shows the product ion scan spectra obtained using O_2 mass-shift mode for a solution containing 1 ppm REEs without (left) and with (right) a 1 ppb As spike. As illustrated in the schematic, Q1 was fixed at m/z = 75 and Q2 was scanned across the selected mass range to monitor all existing and cell-formed ions derived from precursor ions at m/z 75. Figure 3 (left) shows the product ions from m/z 75 in the blank REE matrix; the signal at Q2 m/z = 75 (mass of As) is due to REE⁺⁺. The absence of a signal at m/z = 91 (the mass of AsO⁺) in the blank REE matrix, indicates that the REEs do not react with O_2 in the cell to give rise to product ions (such as REEO₂⁺⁺) that overlap AsO⁺ at m/z 91. Consequently, As can be successfully measured as AsO⁺ at m/z = 91 as shown in Figure 3 (right).

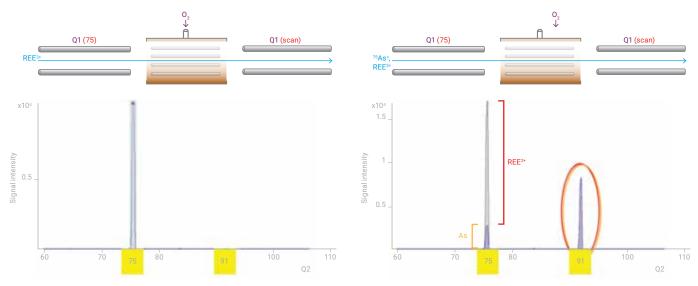


Figure 3. Product ion scan spectrum of O₂ mass-shift method. (Left) 1 ppm mixed-REE solution and (right) 1 ppm mixed-REE plus 1 ppb As spike.

Method validation with CRMs

The ICP-QQQ method was applied to the measurement of As and Se in four CRMs. Table 2 summarizes the results. The measured concentrations of As and Se in the CRMs were all in good agreement with the certified values.

Table 2. Results of the determination of As and Se in four CRMs using MS/MS $\rm O_2$ mass-shift mode on the 8800 ICP-QQQ.

	As	(as AsO+ at m/z	91)	Se (as SeO+ at m/z 94)			
	Certified mg/kg	Found average mg/kg	Recovery %	Certified mg/kg	Found average mg/kg	Recovery %	
NIST1515 Apple Leaves	0.038±0.007	0.037	97	0.050±0.009	0.050	100	
NIST1575a Pine Needles	0.039±0.002	0.038	97	0.099±0.004	0.099	100	
NIST1573a Tomato Leaves	0.112±0.004	0.113	101	0.054±0.003	0.058	107	
NMIJ 7531a Brown Rice	0.280±0.009	0.258	92	NA	0.032	NA	

Removal of Molybdenum Oxide Interference on Cadmium

Authors

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Keywords

cadmium, molybdenum oxide, environmental, food, CRMs, hydrogen on-mass

Introduction

Cadmium (Cd) is a well-known toxic element along with As, Hg and Pb. The maximum contamination level of these elements in food, pharmaceuticals, drinking water, wastewater and other matrices is strictly controlled under national and international legislation. Out of the eight natural isotopes of Cd, only ¹¹¹Cd is free from direct overlap by an atomic isobar (an isotope of a different element at the same mass as the Cd isotope), and even ¹¹¹Cd is potentially subject to spectroscopic interference by ⁹⁵MoO+. Fortunately, the concentration of Mo is low in most samples, and quadrupole ICP-MS (ICP-QMS) operating in helium collision mode can remove the interference, allowing the accurate measurement of Cd. However, there are some cases where the Mo concentration is high and a better interference removal technique is required in order to accurately determine Cd. This paper describes the application of MS/MS H₂ reaction mode on the Agilent 8800 ICP-QQQ for the determination of trace Cd in the presence of a high concentration of Mo.

Experimental

Instrumentation: Agilent 8800 #100. Indium (In) was introduced as the internal standard using the on-line ISTD kit.

Plasma conditions and ion lens tune: RF power = 1550 W; sampling depth = 8.0 mm; carrier gas = 1.01 L/min; make-up gas/dilution gas = 0.0 L/min; Soft extraction tune: Extract 1 = 0 V, Extract 2 = -165 V, Omega bias = -100 V, Omega = 11.4 V.

CRC conditions: H_2 flow rate 9.0 mL/min, Octopole bias = -22 V, KED = +5 V.

Acquisition parameters: MS/MS H_2 on-mass method i.e. ¹¹¹Cd was measured at m/z 111 using quadrupole settings of (Q1 = 111, Q2 = 111).

Results and Discussion

Optimization of H₂ flow rate

Figure 1 (left) shows the signal at m/z 111 for a 10 ppm Mo solution and a 10 ppm Mo + 1 ppm Cd solution, plotted as a function of H_2 flow rate. Figure 1 (right) shows the calculated BEC of Cd in the presence of 10 ppm Mo. The optimum cell gas flow rate of 9.0 mL/min was used for subsequent experiments.

In order to test the effectiveness of MS/MS mode with H_2 cell gas in comparison to no gas mode, a spike recovery test of 1 ppb Cd in a series of Mo matrix solutions ranging from 0.1 to 100 ppm was conducted. Figure 2 summarizes the results. In no gas mode, the error in quantification of the 1 ppb Cd spike dramatically increases with the concentration of Mo; in contrast, H_2 reaction mode delivers a consistent and accurate result for Cd even in the presence of 100 ppm Mo.

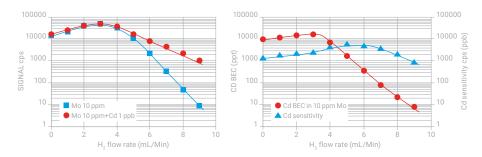


Figure 1. (Left): Signal for m/z 111 with 10 ppm Mo and 10 ppm Mo + 1 ppb Cd, plotted as a function of H_2 flow rate. (Right): Estimated Cd BEC in the presence of 10 ppm Mo as a function of H_2 flow rate.

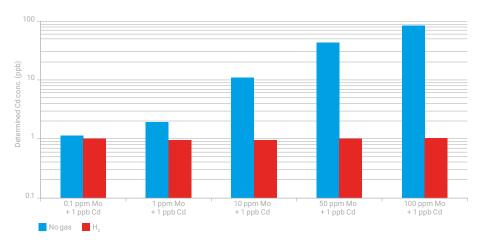


Figure 2. 1 ppb Cd spike recovery in a series of Mo matrix solutions using no gas mode and MS/MS ${\rm H_2}$ mode.

Method validation: Analysis of Cd in CRMs

The concentration of Cd was determined in four different CRMs: NIST 1515 Apple Leaves, NIST 1573a Tomato Leaves, NIST 1575a Pine Needles and NMIJ 7531a Brown Rice Flour (National Metrology Institute of Japan). Each sample was microwave digested following the manufacturer's recommended procedures, then diluted and analyzed by ICP-QQQ; the final dilution factor was around 100–200. For each CRM, the digested sample was analyzed using the developed method. A second sample of each CRM was prepared and analyzed after the addition of a 10 ppm Mo spike. As summarized in Table 1, good recoveries were obtained for all four references materials both for the unspiked samples and the duplicates with the high added Mo concentration, demonstrating the validity of the method for real sample analysis.

Table 1. Measurement of Cd in four CRMs using the 8800 ICP-QQQ in MS/MS mode with $\rm H_2$ reaction gas.

CRMs		Without Mo	addition	With 10 ppm Mo addition			
	Certified mg/kg	Determined mg/kg	Recovery %	Determined mg/kg	Recovery %		
NIST1515 Apple Leaves	0.014	0.013	93	0.016	100		
NIST1573a Tomato Leaves	1.52	1.496	98	1.475	100		
NIST1575a Pine Needles	0.223	0.220	99	0.224	107		
NMIJ 7531a Brown Rice Flour	0.308	0.298	97	0.293	NA		

Feasibility Study of Fluorine Detection by ICP-QQQ

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Keywords

fluorine-containing polyatomic ions, barium, oxygen on-mass, ammonia mass-shift

Introduction

Fluorine (19 F) cannot be directly detected by conventional quadrupole ICP-MS (ICP-QMS) because of severe water-derived interferences at m/z 19 from 1 H $_{3}^{16}$ O $^{+}$ and 1 H 18 O $^{+}$, and extremely low sensitivity due to the fact that it is very difficult to convert fluorine atoms to the positive ions that are measured in ICP-MS. The interference problem can be resolved by high resolution ICP-MS, but the sensitivity issue remains a challenge because almost no F atoms are ionized in an argon plasma due to F having an ionization potential (17.423 eV) that is higher than that of Ar (15.760 eV).

However, fluorine-containing polyatomic ions (XF $^+$) can be formed in the plasma and they may be used to determine fluorine. Candidate ions are those with a high bond-dissociation energy for the X $^+$ -F bond and low ionization potential of X or XF. Since oxygen is present in the plasma (from the water matrix or from air entrainment), the formation of XO $^+$ or XO often competes against that of XF $^+$. Therefore, a low bond-dissociation energy for X $^+$ -O and X-O bonds (low affinity of X $^+$ and X for O) is also desirable for the efficient formation of XF $^+$. Barium was selected as "X" for this feasibility study, based on its thermochemical properties (Table 1).

Table 1. Gas phase thermochemical properties of elements having an affinity for fluorine*.

Element X	D ₀ (X+-F)	IP (X)	D ₀ (X-F)	IP (XF)	D ₀ (X+-O)	D ₀ (X-O)
С	7.77	11.27	5.60	9.11	8.35	11.15
Al	3.16	5.99	6.99	9.73	1.81	5.31
Si	7.01	8.15	5.69	7.54	4.99	11.49
Ва	6.39	5.21	5.98	4.70	5.60	5.80
La	6.83	5.61	6.86	5.56	8.73	8.50
Eu	6.05	5.67	5.59	5.90	4.00	5.90

*Unit: eV. D0(A-F) is the bond-dissociation energy for A-F bond (affinity of A for F) and IP(B) is the ionization potential of B.

Experimental

Instrumentation: Agilent 8800 #200 with a Micromist nebulizer.

Plasma conditions and ion lens tune: RF power = 1500 W; Sampling depth = 8 mm; Carrier gas flow rate = 1.00 L/min; sample uptake rate 0.33 mL/min; 100 ppm Ba uptake rate = 0.03 mL/min; Make-up gas flow rate = 0.32 L/min; Extract 1 = -150 V. Extract 2 = -4 V.

CRC conditions: O_2 gas at 1 mL/min (100%), Octopole bias = -60 V, Energy discrimination = -10 V in O_2 mode; 10% NH₃/90% He flow rate 8.5 mL/min (85%), Octopole bias = -20 V, Energy discrimination = -10 V in NH₃ mode.

Acquisition parameters: MS/MS O_2 on-mass and MS/MS NH_3 mass-shift. Integration time per mass for BaF and BaF(NH_3)₃ = 1 sec; integration time per mass for BaF(NH_3)₄ = 10 sec.

In order to produce BaF⁺ in the plasma, Ba solution was mixed online with fluorine standards per a fixed mixing ratio of 1:10. The mixing occurred just before the nebulizer. BaF⁺ was efficiently formed under general plasma conditions with the BaO⁺/Ba⁺ ratio at about 11%. Under hotter plasma conditions, the formation of BaF⁺ decreases because it tends to break apart. Under cooler plasma conditions, the formation of BaF⁺ also decreases because of the formation of BaO⁺ or, possibly, BaO. The signal intensity of BaF⁺ was proportional to the concentration of Ba, which was fixed at about 10 ppm (after mixing).

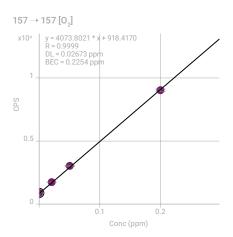
Interference removal using MS/MS mode

 $^{138}\text{Ba}^{19}\text{F}^+$ (*m/z*=157) suffers an interference from $^{138}\text{Ba}^{18}\text{O}^1\text{H}^+$. O_2 and NH $_3$ were tested as reaction gases to reduce the interference. It was found that O_2 reacts with BaOH+ more efficiently than it reacts with BaF+ in high energy reaction mode (octopole bias < -50 V). Therefore, using MS/MS mode, a mass pair (Q1 \square Q2) = (157 \square 157) was selected to detect BaF+ in O $_2$ mode. With Q1 set to 157 amu, $^{138}\text{Ba}^+$ was prevented from entering the cell and forming new interferences through unwanted reactions.

NH $_3$ was found to react with BaF+ at a high NH $_3$ flow rate to form BaF(NH $_3$) $_n$ +, where n = 2, 3, 4. The most abundant complex ion was BaF(NH $_3$) $_3$ + at m/z = 208, but BaF(NH $_3$) $_4$ + at m/z = 225 was preferable in terms of signal to background ratio or BEC. Mass pairs (Q1 \square Q2) = (157 \square 208) and (157 \square 225) were selected in NH $_3$ mode.

Results and Discussion

Figures 1 and 2 show calibration curves up to 2 mg/L (ppm) for fluorine in deionized water. The lowest detection limit (27 ppb) was obtained in $\rm O_2$ mode. The lowest BEC (87 ppb) was obtained by measuring BaF(NH $_3$) $_4$ $^+$ in NH $_3$ mode. Table 2 shows the BEC and DL results for F obtained from this study in comparison with the literature values.





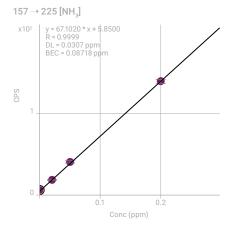


Figure 2. Calibration curve for F measured as $BaF(NH_3)_4^+$ in NH_3 mode.

Table 2. Analytical performance for fluorine detection by ICP-MS.

Analyte ion	Sensitivity [cps/ppm]	BEC [ppm]	DL [ppm]	Technique, reference
F-	60,000	NA	0.11	Negative ion mode ICP-MS, Appl. Spectrosc, 42, 425 (1988)
F ⁺	3,000	NA	0.023	He-ICP-MS, Japan analyst 52(4), 275-278, 2003
Al+ (AlF ²⁺ complex)	NA	0.0033	0.0001	IC-ICP-MS (indirect determination), Analyst. 1999 Jan;124(1):27-31
F+	26	2.05	5.07	HR-ICP-MS, J. Anal. At. Spectrom, 18, 1443, 2003
BaF ⁺	4,073	0.23	0.027	ICP-QQQ, O ₂ mode, this work
BaF(NH ₃) ₃ ⁺	929	0.17	0.043	ICP-QQQ, NH ₃ mode, this work
BaF(NH ₃) ₄ ⁺	67	0.087	0.031	ICP-QQQ, NH ₃ mode, this work

Conclusions

Based on this preliminary study, it is clear that the controlled reaction chemistry that is possible with MS/MS mode on the 8800 ICP-QQQ can provide a novel approach to the measurement of F by ICP-MS. In addition to demonstrating detection limits that are comparable with published data measured using conventional quadrupole ICP-MS or high-resolution ICP-MS, the 8800 ICP-QQQ also allows unprecedented flexibility to monitor specific reaction transitions, making it invaluable for method development.

ICP-QQQ with Oxygen Reaction Mode for Accurate Trace-Level Arsenic Analysis in Complex Samples

Authors

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Keywords

arsenic, zirconium, doubly-charged ion interferences, oxygen mass-shift

Introduction

Arsenic (As), with its high first ionization potential and single isotope at mass 75, is one of the most difficult elements to measure accurately by ICP-MS, particularly in complex matrices. The polyatomic interferences from ArCl⁺ and CaCl⁺ that overlap As⁺ at mass 75 can be removed effectively using quadrupole ICP-MS (ICP-QMS) in helium collision mode, but collision mode cannot resolve the doubly-charged ion interferences from 150 Nd⁺⁺ and 150 Sm⁺⁺. A quadrupole mass spectrometer separates ions based on their mass to charge ratio (m/z), so doubly-charged ions appear at half their true mass; 150 Nd⁺⁺ and 150 Sm⁺⁺ therefore give an apparent overlap on As at mass 75.

Oxygen reaction mode ($\mathrm{O_2}$ mode) offers a solution to these doubly-charged ion overlaps, since As can be converted to a reaction product ion $^{75}\mathrm{As^{16}O^{+}}$, measured at m/z 91, where it is separated from the doubly charged Nd and Sm, which do not form such product ions. However, the new mass of the AsO+ product ion is also overlapped by an isotope of zirconium ($^{91}\mathrm{Zr^{+}}$). The presence of Zr in a sample may therefore cause an error in the results for As measured as AsO+ using $\mathrm{O_2}$ reaction mode on ICP-QMS.

ICP-QQQ solves this problem, as MS/MS mode allows all masses apart from m/z 75 (including the ${}^{91}Zr^+$ ions) to be rejected by the first quadrupole (Q1), ensuring that the AsO+ product ions can be measured free from overlap. ICP-QQQ with MS/MS therefore allows the accurate determination of As in complex samples that contain any combination of Cl, Ca, Nd, Sm and Zr.

Experimental

Reagents and sample preparation: All of the sample matrices used for this work were prepared using single-element stock solutions (Spex CertiPrep, Claritas grade). The acid matrix and elemental standard concentrations are shown in the caption for each spectrum and are representative of the acid matrix (dilute HNO₂/HCl) and matrix levels commonly found in ICP-MS samples.

The sample matrices investigated were:

- Dilute nitric acid (1% HNO_s)
- Dilute hydrochloric acid (5% HCl)
- Calcium (100 ppm)
- Neodymium and samarium (1 ppm each element)
- Zirconium (0.5 ppm)

Instrumentation: Agilent 8800 #100.

Plasma conditions and ion lens tune: Preset plasma/General purpose, Soft extraction tune: Extract 1 = 0 V, Extract 2 = -170 V.

Acquisition conditions: Four operational modes were used, to investigate the different interference removal performance provided by the different cell modes:

- Single Quad (SQ); no gas
- Single Quad (SQ); collision mode (using helium (He) cell gas at a flow rate of 4 mL/min)
- Single Quad (SQ); reaction mode (using oxygen (O₂) cell gas at a flow rate of 0.2 mL/min).
- MS/MS; reaction mode (using O₂ cell gas at a flow rate of 0.2 mL/min)

KED bias voltage was +5 V in no gas and He mode, and -8 V in O₂ mode.

The three "Single Quad" modes represent the performance available on conventional ICP-QMS operating in collision or reaction mode. MS/MS mode is unique to the tandem mass spectrometer configuration of the 8800 ICP-QQQ.

Results and Discussion

Figures 1a, 1b and 1c illustrate how Single Quad mode with He cell gas is effective at removing the common ArCl $^+$ and CaCl $^+$ polyatomic interferences on As $^+$ at m/z 75, but is ineffective against the Nd $^{++}$ /Sm $^{++}$ interferences.

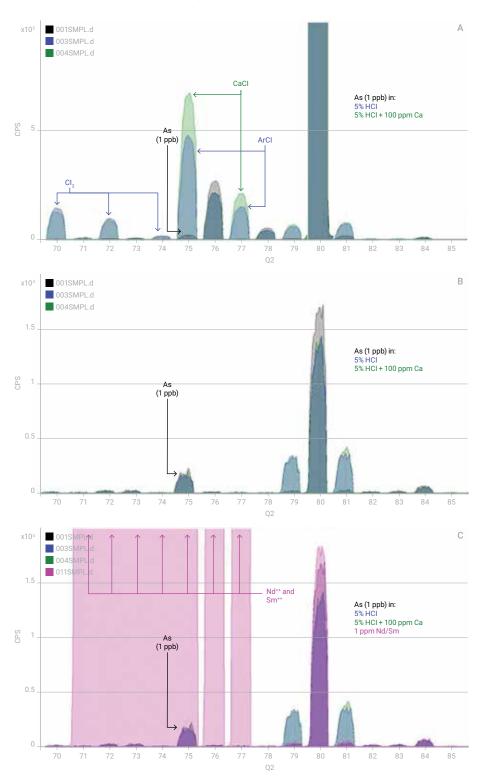


Figure 1. a) As* (m/z 75) in no gas mode, showing polyatomic interferences from ArCl* and CaCl*; **b)** ArCl* and CaCl* polyatomics are removed in He collision mode; **c)** He collision mode fails to remove Nd** and Sm** interferences at m/z 75.

Figures 2a and 2b show how Single Quad mode with $\rm O_2$ reaction gas successfully avoids the doubly-charged Nd and Sm interferences by mass-shifting the As to the new AsO+ product ion mass at m/z 91; but $\rm O_2$ reaction mode on ICP-QMS cannot remove the $\rm ^{91}Zr^+$ overlap on the AsO+ product ion.

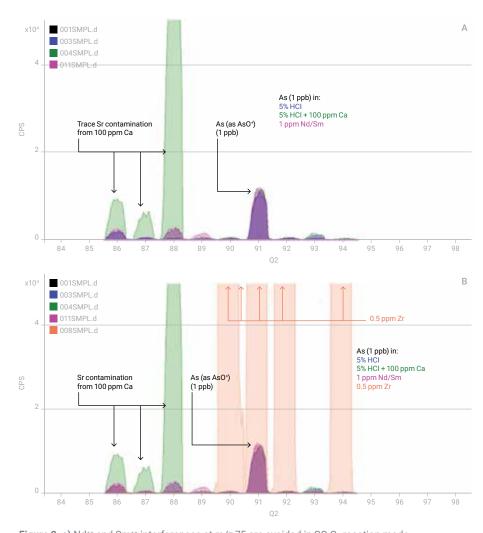


Figure 2. a) Nd⁺⁺ and Sm⁺⁺ interferences at m/z 75 are avoided in SQ O₂ reaction mode, by measuring As as the AsO⁺ product ion at m/z 91; **b)** SQ O₂ reaction mode fails to remove $^{91}\text{Zr}^+$ overlap on the AsO⁺ product ion.

Figure 3 shows that the 8800 ICP-QQQ in MS/MS mode with O_2 reaction gas provides reliable and consistent measurement of As (as AsO+) in all matrices. All the original polyatomic and doubly-charged interferences at m/z 75 are avoided by mass-shifting the As to m/z 91; and in MS/MS mode the $^{91}Zr^+$ ion is removed by Q1, so the potential overlap on the AsO+ product ion at m/z 91 is also removed.

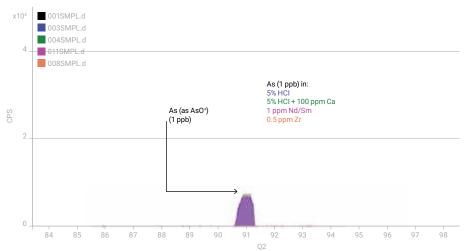


Figure 3. MS/MS mass-shift with 0_2 reaction mode provides consistent, interference-free measurement of As as AsO in all the matrices.

Conclusions

With the combination of $\rm O_2$ reaction mode and MS/MS operation, the 8800 ICP-QQQ provides a reliable approach to the accurate measurement of As in complex samples. All the polyatomic and doubly-charged interferences that affect As measurement at its native mass (m/z 75) are avoided by using $\rm O_2$ mode to mass-shift the As to its AsO⁺ product ion, measured at m/z 91. Furthermore, uniquely to the 8800 ICP-QQQ, MS/MS mode also eliminates potential native ion overlaps at m/z 91, as they are rejected by Q1 that is set to m/z 75 when measuring As.

Avoidance of Spectral Overlaps on Reaction Product Ions with O₂ Cell Gas: Comparison of Quadrupole ICP-MS and ICP-QQQ

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Keywords

titanium, reaction chemistry, oxygen mass-shift

Introduction

The 8800 ICP-QQQ opens up many new analytical possibilities and novel methodologies for interference removal based on reaction chemistry. The major benefit provided by the 8800 ICP-QQQ is its unique tandem mass spectrometer configuration, which permits operation in MS/MS mode. In MS/MS, the first quadrupole (Q1) operates as a 1 amu mass filter, providing precise selection of the ions that can enter the reaction cell, and therefore control of the reaction processes that occur. This level of reaction process control is fundamentally different to the operation of conventional quadrupole ICP-MS (ICP-QMS) when using these same reaction chemistries, as ICP-QMS has no way to reject ions before they enter the cell, and so cannot select which ions are involved in the reactions.

This difference is apparent in many reaction chemistries, including both on-mass measurements (where the interfering ions are reactive and are moved away from the analyte ions, which are then measured at the natural mass), and mass-shift methods (where the analyte ions are reactive and are moved to a new product ion mass that is free from the original overlap). Overlaps on analyte product ions commonly occur in ICP-QMS and can give severe errors in results, especially in cases where the sample matrix or co-existing analyte levels vary from sample to sample.

In this note, we compare the performance of ICP-QMS (the 8800 ICP-QQQ operated in Single Quad mode with Q1 as a bandpass filter) and ICP-QQQ (the 8800 ICP-QQQ operated in MS/MS mode) for the measurement of titanium (Ti) as TiO+ product ions, using oxygen reaction mode (O₂ mode).

The native ion overlaps that could affect the measurement of TiO+ product ions with oxygen reaction gas are shown in Table 1. It should be noted that these native ion overlaps cannot be rejected by the cell bandpass settings of a conventional quadrupole ICP-MS, because they occur at the same mass as the analyte product ion being measured.

Table 1. Potential native ion overlaps on TiO⁺ product ions in O₂ reaction mode.

Precursor ion	(Q1) Product ion (Q2)	ı	Potential overlaps from	other analytes
Ti	TiO	Ni	Cu	Zn
46	62	⁶² Ni	-	-
47	63	-	⁶³ Cu	-
48	64	-	-	⁶⁴ Zn
49	65	-	⁶⁵ Cu	-
50	66	-	-	⁶⁶ Zn

Experimental

For the spectral comparison, scan data were collected for the mass range from m/z 60 to 69, covering the TiO⁺ product ions formed from Ti in O₂ reaction mode.

Instrumentation: Agilent 8800 #100.

Plasma conditions and ion lens tune: Preset plasma/General purpose,

Soft extraction tune: Extract 1 = 0 V, Extract 2 = -180 V.

CRC conditions: Cell gas = 0, gas at 0.3 mL/min, Octopole bias = -5 V, KED = -7 V.

Acquisition parameters: Scan range = m/z 60 to 69; points per peak = 20; integration time per mass = 1 sec.

integration time per mass

Results and Discussion

The comparative results for TiO $^+$ measured in Single Quad (SQ) mode and MS/MS mode are shown in the overlaid spectra in Figures 1 and 2. In both cases, the TiO $^+$ ions at mass 62, 63, 64, 65 and 66 (from the 5 isotopes of Ti at 46, 47, 48, 49 and 50, respectively) are shown, measured using the same O $_2$ reaction mode conditions for both modes. The four solutions measured for the overlaid spectra are:

- 1 ppb Ti in 1% HNO₃
- 1 ppb Ti + 10 ppb Ni in 1% HNO₃
- 1 ppb Ti + 10 ppb Cu in 1% HNO₃
- 1 ppb Ti + 10 ppb Zn in 1% HNO₃

The overlaid spectra in Single Quad mode, shown in Figure 1, show that the peaks for the five TiO $^+$ isotopes match the theoretical isotopic template in the 1 ppb Ti sample. However, in the other samples containing the elements Ni, Cu and Zn, all of the TiO $^+$ product ions suffer significant overlap from the native Ni (m/z 62), Cu (m/z 63 and 65) and Zn (m/z 64 and 66) ions. Unexpected or variable levels of these common elements would lead to an error in the reported results for Ti measured as TiO $^+$ using quadrupole ICP-MS in O $_2$ reaction mode.

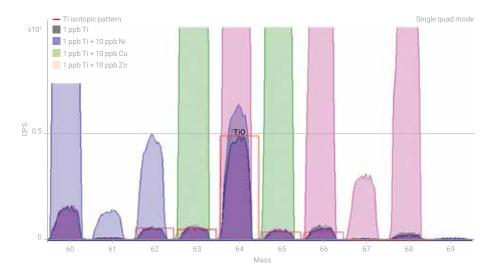


Figure 1. Overlaid spectra for TiO^+ product ions in variable samples measured using SQ mode (see text for sample composition).

In contrast, the overlaid spectra for MS/MS mode, shown in Figure 2, demonstrate consistent measurement of all five TiO+ product ions in all four solutions. The presence of the other elements Ni, Cu and Zn has no impact on the TiO+ peaks and all five TiO+ product ion isotopes could be used to give reliable results for Ti in these variable samples. This illustrates how MS/MS mode on the 8800 ICP-QQQ can simplify method development, because consistent cell conditions, acquisition parameters and isotope selection can be used for a range of variable sample types. A further benefit is that interferences are removed from all isotopes under the same cell conditions, so secondary (or qualifier) isotopes become available for data confirmation or isotope analysis.

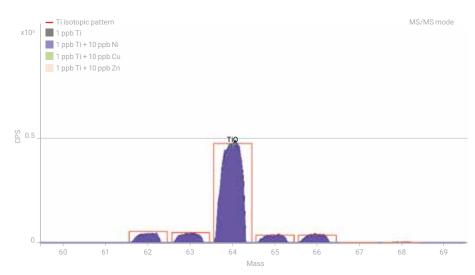


Figure 2. Overlaid spectra for TiO* product ions in variable samples measured using MS/MS mode (see text for sample composition).

Conclusions

The comparative spectra presented in this note illustrate the improved accuracy and consistency delivered by ICP-QQQ operating in MS/MS mode, compared to a conventional quadrupole ICP-MS using a reaction cell with bandpass filter. By rejecting non-target native ions that would occur at the same mass as analyte product ions, potential interferences can be eliminated by MS/MS. This allows simpler, more consistent method development, as well as improving accuracy for interfered elements in complex and variable samples.

Removal of Complex Spectral Interferences on Noble Metal Isotopes

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Keywords

Platinum Group Elements, gold, silver, ore exploration, geochemical, environmental, catalytic converter, ammonia

Introduction

The precise determination of the noble metals, comprising the Platinum Group Elements (PGEs: Ru, Rh, Pd, Os, Ir and Pt), Au and Ag, is of great interest in areas such as ore exploration and geochemical studies, and these metals are increasingly used for industrial applications including advanced materials and alloys, medical devices, and catalysts for pharmaceutical manufacturing. Environmental monitoring is also required as some of these elements are used in automobile catalytic converters. ICP-MS is widely used for these applications due to its high sensitivity and multi-element capability. However, the analysis is challenging because the metal concentrations are often low and they are subject to severe spectral overlaps.

Table 1 summarizes the interferences and abundance (%) of each isotope of the elements (the isotopes highlighted in yellow represent the recommended isotope for determination by ICP-MS). Several methods have been developed to resolve the interferences, such as mathematical correction, matrix removal and high-resolution magnetic sector (HR-)ICP-MS. However the mass resolution required to separate some of the interferences is beyond the capability of current commercial HR-ICP-MS. For example separation of $^{103}\text{Rh}^+$ from $^{87}\text{Sr}^{16}\text{O}^+$, $^{105}\text{Pd}^+$ from $^{89}\text{Y}^{16}\text{O}^+$, and $^{109}\text{Ag}^+$ from $^{93}\text{Nb}^{16}\text{O}^+$ requires mass resolution (M/ Δ M) of 102900, 27600 and 31500, respectively; commercial HR-ICP-MS instruments are limited to a maximum resolution of 10,000. To remove the multiple, complex interferences on noble elements, the Agilent 8800 ICP-QQQ was used in MS/MS mode, using ammonia as the reaction gas.

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/Low matrix.

Ion lens tune: Soft extraction tune: Extract 1 = -3 V, Extract 2 = -200 V.

CRC conditions: NH₃ (10% NH₃ in He) was used as CRC gas in MS/MS mode.

Following a preliminary optimization study, three different NH_3 gas flow rates (low (L), medium (M) and high (H)) were used. Cell conditions are given in Table 2. No gas mode was also applied for comparison purposes.

Table 1. Abundance (%) of each isotope of noble metals and the interference on each isotope.

m		96	97	98	99	100	101	102	103	104	105	106	107	108	109	110
Element	Ru	5.54		1.87	12.76	12.6	17.06	31.55		8.62						
	Rh							_	100							
	Pd							1.02		11.14	22.33	27.33		26.46		11.7
	Ag												51.84		48.16	
Interference	Atomic	Zr, Nb, Mo		Мо		Мо						Cd		Cd		Cd
	МН			МоН	МоН		МоН									
	мо, мон	SeO, BrOH		SeO		Sr0	RbO	Sr0	SrO, RbO	Sr0	YO, SrOH	YOH, ZrO	ZrO	ZrO, MoO	NbO	ZrO, MoC
	Argide			NiAr		NiAr	NiAr	NiAr	CuAr	ZnAr	CuAr	ZnAr	ZnAr	ZnAr		
	Others			CuCl	ZnCl	CuCl	ZnCl	CuCl, ZnCl	ZnCl, Pb ⁺⁺	ZnCl	ZnCl					
m		184	185	186	187	188	189	190	191	192	193	194	195	196	197	198
Element	Os	0.02		1.59	1.96	13.24	16.15	26.26		40.78						
	lr								37.3		62.7					
	Pt							0.014		0.782		32.97	33.83	25.24		7.16
	Au														100	
Interference	Atomic	W		w	Re									Hg		Hg
	МН				WH											
	мо, мон			YbO	YbO	YbO	YbO	YbO	LuO	YbO, LuD, HfO	HfO	HfO	HfO	HfO	TaO, HfOH	Wo, TaOl
	Argide	NdAr	NdAr	NdAr	SmAr	SmAr, NdAr	SmAr	SmAr, NdAr	EuAr	SmAr	EuAr	SmAr, GdAr	GdAr	GdAr	GdAr	GdA

Table 2. CRC conditions.

	No gas	NH ₃ -L	NH ₃ -M	NH ₃ -H		
Cell gas	na	NH ₃	NH ₃	NH ₃		
Gas flow rate (mL/min)	na	2.0	3.0	5.0		
Octopole bias (V)	-8	-5	-10	-12		
KED (V)	+5		-8			
Cell exit (V)		-9	0			
Deflect lens (V)	20	10	6	2		
Plate lens (V)	-110					

Method

The BECs of the noble metals were determined in a series of synthetic-matrix samples, using an external calibration method. Indium (In) internal standard (ISTD) was mixed online with the sample via the standard ISTD mixing T-connector. An integration time of 1 s per isotope was used with 3 replicates (7 replicates for blank).

Samples and sample preparation

Standards and matrix samples were prepared from single element stock solutions purchased from Kanto Chemical Co., Inc. (Saitama, Japan) and a REE mixture standard, XSTC-1 purchased from Spex certiPrep. All solutions were diluted into a final acid mix of 1% HNO $_3$ and 3% HCl.

Results

Matrix interference study

Tables 3 and 4 summarize the results of the spectral interference study obtained by analyzing individual synthetic matrix blank solutions. Table 3 shows the observed interferences, expressed as BEC (ppb), in each matrix blank measured using no gas mode. As expected from Table 1, the synthetic matrices caused significantly elevated BECs (>> 1 ppb) on all the primary and secondary isotopes of all the analytes except for Ru; Rh suffered a relatively minor increase in BEC of ~0.5 ppb in the 10 ppm Pb/1 ppm Hg matrix.

Table 4 shows the results obtained using $\mathrm{NH_3}$ reaction mode. The optimum gas flow rate for $\mathrm{NH_3}$ for each element was investigated and three gas flow rates (Low: 2.0, Medium: 3.0, and High: 5.0 mL/min) were used. The best isotope and method is highlighted in bold in the Table. It can clearly be seen that $\mathrm{NH_3}$ reaction mode effectively removes the interferences on all the analytes, giving BECs of << 0.1 ppb for the preferred isotope/cell mode in all the matrices. The mechanism for the removal of each interference using the MS/MS capability of the 8800 ICP-QQQ is as follows:

- Ru: slight interferences from Zn and Mo were resolved using on-mass method with NH₃-M.
- **Rh:** Pb⁺⁺ interference was resolved using on-mass method with NH₃-M.
- Pd: significant interferences from SrOH⁺ and YO⁺ were seen on ¹⁰⁵Pd, the only isotope free from atomic isobar. On-mass method with NH₃-H removed the interferences.
- Ag: significant ZrO+ interference on both ¹⁰⁷Ag and ¹⁰⁹Ag was resolved using on-mass method with NH₂-H.

- **Os:** YbO+ interference was observed on both 188 Os+ and 189 Os+. Since Os+ sensitivity in NH $_3$ mode is low, but Os+ forms a product ion of OsNH+, NH $_3$ -L with mass-shift gave the best result.
- **Ir:** LuO $^+$ and HfO $^+$ interfere with 191 Ir $^+$ and 193 Ir $^+$ respectively. NH $_3$ -M with mass-shift method worked for 191 Ir $^+$ as Ir $^+$ forms a product ion of IrNH $^+$.
- Pt: ¹⁹⁵Pt⁺ suffers a significant interference from HfO⁺. While the overlap is less significant on ¹⁹⁸Pt⁺, ¹⁹⁸Pt⁺ suffers an atomic isobar interference from ¹⁹⁸Hg⁺. However Hg⁺ is effectively neutralized by NH₃ so ¹⁹⁸Pt⁺ can be measured free from interference.
- **Au:** significant interferences by TaO⁺ and HfOH⁺ are resolved by mass-shift method with NH_3 -M. Au⁺ forms a product ion of Au(NH_3)₂⁺.

Table 3. Summary of spectral interferences in no gas mode, showing analyte BECs (ppb) in each matrix blank. Matrix overlaps that made a significant contribution to the analyte BECs are indicated in red (BEC > 10 ppb) and orange (BEC > 1 ppb).

	Du		Dh	Pd	۸۵		00	
I A	Ru	101	Rh		Ag	100	0s	100
Isotope	99	101	103	105	107	109	188	189
NH ₃ flow rate mL/min	NA							
Method	on-mass							
Mass pair	99-99	101-101	103-103	105-105	107-107	109-109	188-188	189-189
10 ppm Cu Zn	0.058	0.041	0.138	0.328	0.064	0.061	0.000	0.000
10 ppm Sr Rb	0.000	0.034	0.150	4.39	0.005	0.001	0.000	0.000
10 ppm Ni	0.007	0.019	0.000	0.022	0.012	0.016	0.000	0.000
10 ppm Mo	0.059	0.018	0.000	0.004	0.000	0.018	0.000	0.000
10 ppm Pb, 1 ppm Hg	0.000	0.000	0.472	0.002	0.033	0.034	0.000	0.000
10 ppm Zr Nb	0.000	0.000	0.000	0.022	21.9	1.59	0.000	0.000
10 ppm REE	0.004	0.000	0.009	165	0.147	0.005	2.78	2.99
10 ppm Ta	0.008	0.000	0.000	0.004	0.003	0.000	0.000	0.000
10 ppm Hf	0.000	0.000	0.000	0.004	0.312	0.026	0.000	0.000
10 ppm W	0.000	0.000	0.000	0.003	0.001	0.001	0.000	0.000
	lr				Pt			Au
Isotope	191		193		195		198	197
NH ₃ flow rate mL/min	NA							
Method	on-mass		on-mass		on-mass		on-mass	on-mass
Mass pair	191-191		193-193		195-195		198-198	197-197
10 ppm Cu Zn	0.003		0.002		0.000		0.279	0.001
10 ppm Sr Rb	0.002		0.000		0.001		0.310	0.004
10 ppm Ni	0.009		0.004		0.002		0.444	0.011
10 ppm Mo	0.000		0.000		0.000		0.295	0.000
10 ppm Pb, 1 ppm Hg	0.002		0.002		0.000		1293	0.000
10 ppm Zr Nb	0.002		0.775		1.98		3.17	0.417
10 ppm REE	123		0.712		0.788		2.17	0.138
10 ppm Ta	0.000		0.000		0.244		114	284
10 ppm Hf	0.071		28.1		70.9		2.34	14.1
10 ppm W	0.001		0.000	_	0.000		19.6	0.002

Table 4. Summary of spectral interferences in MS/MS NH₃ reaction cell mode, showing analyte BECs (ppb) in each matrix blank. Matrix overlaps that made a significant contribution to the analyte BECs are indicated in red (> 10 ppb) and orange (> 1 ppb).

t	Ru		Rh	Pd	Ag		Os			
Isotope	99	101	103	105	107	109	188		189	
NH ₃ flow rate mL/min	3.0		3.0	5.0	5.0		2.0			
Method	on-mass	on-mass	on-mass	on-mass	on-mass	on-mass	on-mass	mass-shift	on-mass	mass-shift
Mass pair	99-99	101-101	103-103	105-105	107-107	109-109	188-188	188-203	189-189	189-204
10 ppm Cu Zn	0.000	0.000	0.000	0.001	0.061	0.057	0.000	0.000	0.001	0.002
10 ppm Sr Rb	0.000	0.005	0.016	0.033	0.000	0.000	0.000	0.000	0.002	0.000
10 ppm Ni	0.000	0.000	0.000	0.000	0.010	0.009	0.000	0.000	0.000	0.000
10 ppm Mo	0.005	0.001	0.000	0.000	0.001	0.000	0.000	0.000	0.001	0.000
10 ppm Pb, 1 ppm Hg	0.000	0.000	0.000	0.001	0.033	0.035	0.000	0.000	0.001	0.000
10 ppm Zr Nb	0.000	0.000	0.000	0.000	0.003	0.000	0.000	0.000	0.002	0.001
10 ppm REE	0.000	0.000	0.000	0.014	0.004	0.004	2.79	0.003	5.85	0.010
10 ppm Ta	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.001	0.003	0.000
10 ppm Hf	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.001	0.058	0.000
10 ppm W	0.000	0.000	0.000	0.001	0.001	0.001	0.000	0.000	0.000	0.000
	Ir				Pt				Au	
Isotope	lr 191		193		Pt 195		198		Au 197	
Isotope NH ₃ flow rate mL/min			193				198			
·	191	mass-shift		mass-shift	195 5.0	mass-shift	3.0	mass-shift	197 3.0	mass-shift
NH ₃ flow rate mL/min	191 3.0	mass-shift		mass-shift	195 5.0	mass-shift	3.0	mass-shift	197 3.0	mass-shift 197-231
NH ₃ flow rate mL/min	191 3.0 on-mass		on-mass		195 5.0 on-mass	<u> </u>	3.0 on-mass		197 3.0 on-mass	
NH ₃ flow rate mL/min Method Mass pair	191 3.0 on-mass 191-191	191-206	on-mass 193-193	193-208	195 5.0 on-mass 195-195	195-229	3.0 on-mass 198-198	198-232	197 3.0 on-mass 197-197	197-231
NH ₃ flow rate mL/min Method Mass pair 10 ppm Cu Zn	191 3.0 on-mass 191-191 0.004	191-206 0.000	on-mass 193-193 0.003	193-208 0.004	195 5.0 on-mass 195-195 0.003	195-229 0.000	3.0 on-mass 198-198 0.002	198-232 0.000	197 3.0 on-mass 197-197 0.000	197-231 0.000
NH ₃ flow rate mL/min Method Mass pair 10 ppm Cu Zn 10 ppm Sr Rb	191 3.0 on-mass 191-191 0.004 0.002	191-206 0.000 0.000	on-mass 193-193 0.003 0.001	193-208 0.004 0.001	195 5.0 on-mass 195-195 0.003 0.000	195-229 0.000 0.000	3.0 on-mass 198-198 0.002 0.000	198-232 0.000 0.000	197 3.0 on-mass 197-197 0.000	197-231 0.000 0.000
NH ₃ flow rate mL/min Method Mass pair 10 ppm Cu Zn 10 ppm Sr Rb 10 ppm Ni	191 3.0 on-mass 191-191 0.004 0.002	191-206 0.000 0.000 0.000	on-mass 193-193 0.003 0.001	193-208 0.004 0.001 0.004	195 5.0 on-mass 195-195 0.003 0.000	195-229 0.000 0.000 0.000	3.0 on-mass 198-198 0.002 0.000	198-232 0.000 0.000 0.002	197 3.0 on-mass 197-197 0.000 0.000	197-231 0.000 0.000 0.000
NH ₃ flow rate mL/min Method Mass pair 10 ppm Cu Zn 10 ppm Sr Rb 10 ppm Ni 10 ppm Mo	191 3.0 on-mass 191-191 0.004 0.002 0.004 0.000	191-206 0.000 0.000 0.000 0.000	on-mass 193-193 0.003 0.001 0.001	193-208 0.004 0.001 0.004 0.000	195 5.0 on-mass 195-195 0.003 0.000 0.000 0.000	195-229 0.000 0.000 0.000 0.000	3.0 on-mass 198-198 0.002 0.000 0.000	198-232 0.000 0.000 0.002 0.000	197 3.0 on-mass 197-197 0.000 0.000 0.000 0.000	197-231 0.000 0.000 0.000 0.000
NH ₃ flow rate mL/min Method Mass pair 10 ppm Cu Zn 10 ppm Sr Rb 10 ppm Ni 10 ppm Mo 10 ppm Pb, 1 ppm Hg	191 3.0 on-mass 191-191 0.004 0.002 0.004 0.000 0.000	191-206 0.000 0.000 0.000 0.000 0.001	on-mass 193-193 0.003 0.001 0.001 0.000	193-208 0.004 0.001 0.004 0.000 0.000	195 5.0 on-mass 195-195 0.003 0.000 0.000 0.000	195-229 0.000 0.000 0.000 0.000 0.000	3.0 on-mass 198-198 0.002 0.000 0.000 0.000 0.003	198-232 0.000 0.000 0.002 0.000 0.001	197 3.0 on-mass 197-197 0.000 0.000 0.000 0.000 0.000	197-231 0.000 0.000 0.000 0.000 0.000
NH ₃ flow rate mL/min Method Mass pair 10 ppm Cu Zn 10 ppm Sr Rb 10 ppm Ni 10 ppm Mo 10 ppm Pb, 1 ppm Hg 10 ppm Zr Nb	191 3.0 on-mass 191-191 0.004 0.002 0.004 0.000 0.002 0.017	191-206 0.000 0.000 0.000 0.000 0.001 0.000	on-mass 193-193 0.003 0.001 0.001 0.000 0.000	193-208 0.004 0.001 0.004 0.000 0.000 0.066	195 5.0 on-mass 195-195 0.003 0.000 0.000 0.000 0.000 0.001	195-229 0.000 0.000 0.000 0.000 0.000 0.009	3.0 on-mass 198-198 0.002 0.000 0.000 0.003 0.005	198-232 0.000 0.000 0.002 0.000 0.001	197 3.0 on-mass 197-197 0.000 0.000 0.000 0.000 0.000 0.000	197-231 0.000 0.000 0.000 0.000 0.000 0.000
NH ₃ flow rate mL/min Method Mass pair 10 ppm Cu Zn 10 ppm Sr Rb 10 ppm Ni 10 ppm Mo 10 ppm Pb, 1 ppm Hg 10 ppm Zr Nb 10 ppm REE	191 3.0 on-mass 191-191 0.004 0.002 0.004 0.000 0.002 0.017	191-206 0.000 0.000 0.000 0.000 0.001 0.000 0.019	on-mass 193-193 0.003 0.001 0.001 0.000 0.000 0.679 1.56	193-208 0.004 0.001 0.004 0.000 0.000 0.066 0.019	195 5.0 on-mass 195-195 0.003 0.000 0.000 0.000 0.000 0.001 0.001	195-229 0.000 0.000 0.000 0.000 0.000 0.009 0.002	3.0 on-mass 198-198 0.002 0.000 0.000 0.003 0.005 0.003 0.000	198-232 0.000 0.000 0.002 0.000 0.001 0.001 0.000	197 3.0 on-mass 197-197 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000	197-231 0.000 0.000 0.000 0.000 0.000 0.000 0.000

Analysis of complex synthetic matrix sample using optimized NH₃ reaction mode

A complex synthetic matrix sample containing 10 ppm each of Cu, Zn, Sr, Rb, Ni, Mo, Pb, Zr, Nb, REEs, Ta, Hf, W and 1 ppm Hg was prepared, and this matrix was spiked with 1 ppb each of Ru, Rh, Pd, Ag, Os, Ir, Pt and Au as analytes. The concentration of the noble metals was determined in two modes: No gas mode and $\mathrm{NH_3}$ reaction cell mode, and the spike recovery results are displayed in Figure 1 for each mode. The results demonstrate that MS/MS mode with $\mathrm{NH_3}$ reaction cell gas successfully removes multiple interferences on all the noble metals, providing accurate results for these analytes even in a complex and challenging matrix.

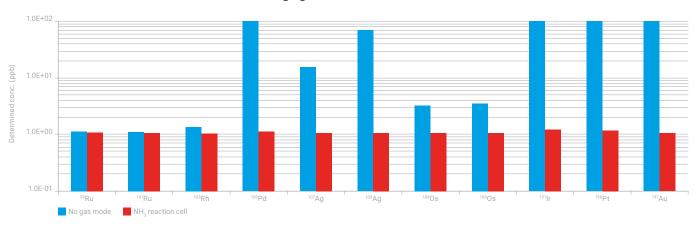


Figure 1. Result of synthetic matrix sample test. 1 ppb noble metals were measured in a multi-matrix sample containing 10 ppm of each Cu, Zn, Sr, Rb, Ni, Mo, Pb, Zr, Nb, REEs, Ta, Hf, W and 1 ppm Hg.

Routine Soil Analysis using the Agilent 8800 ICP-QQQ

Author

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Keywords

soil, sediment, routine analysis, As, Se, HMI, matrix tolerance, robustness

Introduction

Quadrupole ICP-MS is widely used in inorganic testing laboratories, due to its high sensitivity, low detection limits, wide dynamic range, and high speed multi-element analysis. The technique is well suited to the analysis of elemental contaminants present in soil and sediment samples. Helium (He) collision cell technology can be used successfully to remove many common matrix-based polyatomic interferences. He mode is less effective for the removal of interferences caused by doubly charged ions though. For example, interferences on arsenic (As) and selenium (Se) by doubly charged ions of rare earth elements (REEs). Typically, the REE content of environmental samples is low. However, all interferences, including the doubly charged ions of REEs on As and Se, can be removed using oxygen mass-shift mode of Agilent's ICP-QQQ. This approach provides a high level of confidence in the analysis of unknown samples. Agilent's ICP-QQQ instruments also offer the same robustness and matrix tolerance of Agilent's single-quadrupole ICP-MS systems.

This study demonstrates the robustness of the Agilent 8800 ICP-QQQ for routine soil analysis.

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/HMI-4. The ICP-MS MassHunter software automatically sets robust plasma tuning conditions that are suitable for soil/sediment analysis.

Method: the method was based on a preset method for soil (EPA 6020). It was modified to include $\rm O_2$ mass-shift mode for sulfur (S), As, and Se. All other elements were measured in He mode. After the calibration standards and initial QC samples had been analyzed, 13 sample blocks were analyzed. Each block consisted of 10 samples (two each of Soil A, Soil B, Estuarine Sediment, River Sediment A, River Sediment B). A Periodic Block consisting of Continuing Calibration Blank (CCB) and Continuing Calibration Verification (CCV) samples was automatically inserted into the sequence after each set of 10 samples.

Samples: Five soil and sediment CRMs bought from High-Purity Standards Inc. (Charleston, SC, USA) were analyzed in this study. These included CRM River Sediment A, CRM River Sediment B, CRM Estuarine Sediment, CRM Soil A, and CRM Soil B.

Results and Discussion

The total number of analyses of calibration standards, QC samples, and soil samples was 177 over ~12 hours. The internal standard (ISTD) stability plot, shown in Figure 1, met EPA 6020 requirements of between 70 and 120% of the value of the initial calibration standard.

The accuracy of the method was evaluated by analyzing the soil and sediment CRMs as unknown samples. Each CRM was measured 26 times in the batch. The mean concentrations and relative standard deviations (%RSD) were calculated and compared to the certified value, as shown in Table 1. The mean concentration for all elements was in good agreement with the certified value, with most RSDs well below 5% over the 12-hour analysis.

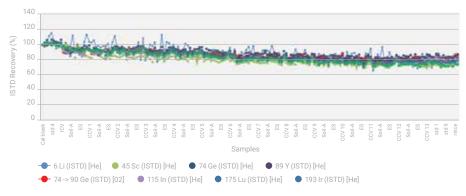


Figure 1. ISTD signal stability plot over 12 hours.

Conclusions

The Agilent 8800 ICP-QQQ with HMI offers the robustness and matrix tolerance required for the routine analysis of the widest range of trace and major elements in high matrix samples, such as soil and sediments. Doubly charged REE interferences that can affect arsenic and selenium measurement at trace levels were avoided using MS/MS mass-shift mode with $\rm O_2$ cell gas. Most other elements were measured in He mode, proven to remove common matrix-based polyatomic interferences in complex and variable matrices. Not all soils, sediments, and food products contain significant concentrations of REEs. However, the presence of REEs in samples that are analyzed using single quadrupole ICP-MS can lead to false positive results for As and Se. The use of the ICP-QQQ with MS/MS improves confidence in the results for these two important elements. Furthermore, method development is simplified with the use of preset methods and autotuning, ensuring reproducible performance from day-to-day and irrespective of operator experience.

More Information

Routine soil analysis using an Agilent 8800 ICP-QQQ, Agilent publication <u>5991-6409EN</u>.

 Table 1. Mean recovery % of three soil/sediment CRMs.

Element	Integration	MDL	River Sed	ment A		Estuarine	e Sediment A		Soil A		
	time (s)	(ppb)	Mean conc. (ppb)	RSD (%)	Mean recovery (%)	Mean conc. (ppb)	RSD (%)	Mean recovery (%)	Mean conc. (ppb)	RSD (%)	Mean recovery (%)
9Be	3	0.06	< MDL			2.1	5.5	106	< MDL		
²³ Na	0.1	0.98	5191	2.6	104	20862	2.4	104	7292	2.6	104
²⁴ Mg	0.1	0.73	7292	2.6	104	10553	2.7	106	7341	2.6	105
²⁷ Al	0.1	1.00	25862	2.4	103	70884	2.7	101	51034	2.5	102
³¹ P	1	3.20	< MDL			520	2.4	104	1042	2.0	104
³² S	1	4.10	< MDL			< MDL			< MDL		
³⁹ K	0.1	7.50	15623	2.2	104	15568	2.7	104	20678	2.3	103
⁴⁴ Ca	0.1	2.70	28860	2.1	96	7760	3.2	97	33670	1.8	96
⁵¹ V	0.3	0.021	26	2.6	105	103	2.9	103	10.4	3.4	104
52Cr	0.3	0.04	29792	2.7	99	83	3.0	104	< MDL		
55Mn	0.3	0.062	809	2.2	101	399	2.9	100	10.9	3.0	109
⁵⁶ Fe	0.1	0.45	120085	2.7	100	35335	3.3	101	20215	2.2	101
⁵⁹ Co	0.3	0.017	11	2.9	106	10.8	2.8	108	0.33	3.1	
⁶⁰ Ni	0.3	0.049	52	2.8	103	30.7	3.2	102	30.2	2.6	101
⁶³ Cu	0.3	0.021	102	2.9	102	20.2	3.1	101	30.2	2.4	101
⁶⁶ Zn	0.3	0.063	1499	2.5	100	151	2.9	101	101	2.3	101
⁷⁵ As	1	0.024	60	3.6	100	10.5	3.6	105	20.4	3.0	102
⁷⁸ Se	3	0.049	2.0	3.6	101	4.9	3.0	99	1.0	6.2	99
⁹⁵ Mo	0.3	0.022	0.19	10.5		< MDL			< MDL		
¹⁰⁷ Ag	0.3	0.015	0.15	9.0		0.015	16.4		0.038	17.3	
¹¹¹ Cd	3	0.012	10.3	2.0	103	0.11	4.5		0.37	2.9	125
¹²¹ Sb	0.3	0.011	50.8	2.1	102	0.58	4.4		3.2	3.5	106
¹³⁵ Ba	0.3	0.055	50.9	2.1	102	1.5	5.4		513	2.6	103
²⁰¹ Hg	1	0.003	< MDL			< MDL			0.018		
²⁰⁵ TI	0.3	0.008	0.97	2.0	97	< MDL			< MDL		
²⁰⁸ Pb	0.3	0.009	719	2.1	103	30.7	2.6	102	41	2.4	101
²³² Th	0.3	0.007	2.1	3.1	106	10.4	2.5	104	10	2.2	103
²³⁸ U	0.3	0.09	1.0	2.4	104	< MDL			1.0	2.5	102

HPLC-ICP-MS/MS: Fluorine Speciation Analysis

Authors

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Keywords

BaF*, fluorine speciation analysis, fluoride, fluoroacetate, trifluoroacetate, HPLC-ICP-QQQ

Introduction

Fluorine is often used in the form of organofluorine compounds in applications such as pharmaceuticals, agrochemicals, and materials. This usage has resulted in the accumulation of large quantities of unknown organofluorine compounds in the environment [1, 2]. Fluorine is a difficult element to determine by ICP-MS. Its high ionization potential (17.423 eV) results in a low yield of F⁺ ions in the plasma, leading to low sensitivity.

Fluorine can be determined, however, by mixing barium and fluorine solutions and measuring the polyatomic ion BaF⁺ by triple quadrupole ICP-MS (ICP-QQQ) [3]. Because 138 Ba is the most abundant isotope, the highest sensitivity would be achieved for 138 Ba 19 F⁺ at m/z 157. Mechanisms for the formation of BaF⁺ are shown in equations 1 and 2.

- (1) 138 Ba⁺ + 19 F⁰ 138 Ba¹⁹F⁺
- (2) $^{138}\text{Ba}^{++} + ^{19}\text{F}^{-} ^{138}\text{Ba}^{19}\text{F}^{+}$

While this approach resolves the low ionization yield issue for F, the formation of potential interfering ions at m/z 157 from ¹³⁸Ba¹⁸O¹H⁺, ¹³⁸Ba¹⁶O¹H₃⁺, and ¹³⁸Ba¹⁷O₂H⁺ also need to be considered. These interferences can be reduced by operating the ICP-QQQ in MS/MS mode, using oxygen as the reaction gas. This approach was used for development of an online HPLC-ICP-QQQ speciation method for the determination of F.

Experimental

Instrumentation: Agilent 8800 ICP-QQQ with Micromist nebulizer and s-lens.

Operating conditions: Table 1 summarizes the plasma, ion lens, and cell tuning conditions.

Acquisition parameters: MS/MS mode with O_2 on-mass. Integration time per m/z for BaF+ = 1 sec.

HPLC system: Agilent 1290 with Metrosep A Supp 5 (150 mm x 4.0 mm) separation column and Metrosep RP Guard/3.5 column. Buffer = 3.2 mM sodium carbonate and 1.0 mM sodium bicarbonate (pH 10); flow rate = isocratic 0.7 mL/min of 70% buffer solution; sample injection = 100 μ L.

A transfer capillary was used to connect the chromatographic column to the nebulizer of the ICP-QQQ system via a T-pin, which allowed the mixing of Ba with F solution. The parameters were optimized in a previous study [4].

Table 1. ICP-QQQ operating conditions.

Parameter	Unit	Value
Plasma		
RF power	W	1500
Sampling depth	mm	8.0
Carrier gas flow rate	L/min	1.00
Make-up gas flow rate	L/min	0.36
Lenses		
Extract 1	V	-150.0
Extract 2	V	5.0
Deflect	V	-48.0
Cell		
Oxygen flow rate	mL/min	0.75
Octopole Bias	V	-60.0
Octopole RF	V	200
Energy discrimination	V	-10.0
Wait time offset	msec	2
Sample uptake rate	mL/min	0.33
32 mg/L Ba uptake rate	mL/min	0.22

Results and Discussion

Figures 1 and 2 show the chromatograms and calibration curves for fluorine speciation analysis of three different fluorine compounds: fluoride, fluoroacetate (FAA), and trifluoroacetate (TFA). All compounds were baseline separated within 10 minutes. The sensitivity of F is similar for each sample, indicating the method is compound independent and fluorine specific.

The limits of detection (LOD) of the HPLC-ICP-QQQ method were 0.012 mg/L, 0.073 mg/L, and 0.12 mg/L for fluoride, FAA, and TFA, respectively. Table 2 shows the LOD results for F from this study compared to data reported in the literature.

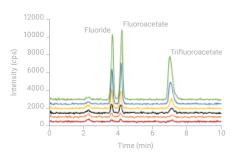


Figure 1. HPLC-ICP-QQQ chromatogram of fluoride, fluoroacetate, and trifluoroacetate at different F concentrations: 0.1, 0.25, 0.5, 1.0, 2.5, and 5.0 mg/L (as indicated by red to green colored lines).

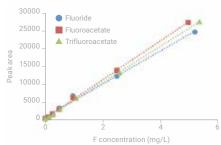


Figure 2. HPLC-ICP-QQQ calibration curves of fluorine compounds: fluoride, fluoroacetate, and trifluoroacetate.

Table 2. Limit of detection (LOD) of F analysis obtained by different methods.

Method	Analyte ion	LOD (mg/L)	Reference
IC-ICP-MS (indirect determination)	Al ⁺ (AlF ⁺ complex)	0.0001	Analyst, 1999, 124, 27-31
HR-ICP-MS	F ⁺	5.1	J. Anal. At. Spectrom., 2003, 18, 1443-1451
ETV-ICP-MS	F ⁺	3.2	J. Anal. At. Spectrom., 2001, 16, 539-541
ICP-MS/MS	BaF+	0.027	Agilent, 2015, 5991-2802EN
		0.043	J. Anal., At. Spectrom., 2017, 32, 942-950
HPLC-ICP-MS/MS	BaF+ for fluoride	0.012	This work
	BaF+ for fluoroacetate	0.073	This work
	BaF+ for trifluoroacetate	0.12	This work

Conclusions

For the first time, coupling an HPLC directly to an ICP-QQQ enabled the speciation analysis of fluorine-containing compounds through the formation of the polyatomic ion BaF⁺ [4]. The method was not only able to detect fluorine specifically but also has a comparable low LOD, which opens up possibilities for future non-targeted fluorine speciation analysis in environmental samples.

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Rapid Analysis of Radium-226 in Water Samples by ICP-QQQ

Authors

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Introduction

Radium-226 is a radionuclide that occurs naturally as part of the uranium-238 decay series. ²²⁶Ra decays with a half-life of 1,600 years to radon-222 with the emission of alpha and gamma radiation. The element is known for its historical use in the luminescent paint used in clocks, watches, and other instruments. These uses led to severe health problems for the so-called Radium Girls who painted the watch and clock dials. ²²⁶Ra has a long half-life compared to the other Ra isotopes, and is considered a significant contributor to occupational radiological dose with regards to industrial sources of naturally occurring radioactive materials (NORM).

²²⁶Ra occurs naturally in waters through interaction with uranium-bearing minerals [1]. It is also present as a result of waste from the industrial exploitation of mineral resources (including uranium mining and processing sites, and produced waters following hydraulic fracturing). Radium waste producers are required to comply with stringent limits when discharging to watercourses. Analytical methods must therefore be capable of detecting ²²⁶Ra at values ranging from 0.01 Bq/L to 1 Bq/L (equivalent to 0.3 – 30 pg/L (ppq) or 0.0003 – 0.03 ppt) [2,3,4].

²²⁶Ra analysis is typically performed by alpha spectrometry, which requires time-consuming and labor-intensive separation before measurement, followed by count times of several days per sample to reach the target detection limits.

This study outlines a new method developed by the National Physical Laboratory (NPL) Nuclear Metrology Group for the rapid analysis of ²²⁶Ra in water samples. The new method uses a preconcentration step prior to measurement of ²²⁶Ra using triple quadrupole ICP-MS (ICP-QQQ)[5]. The procedural time is significantly reduced compared to decay counting techniques, and ²²⁶Ra is measurable at concentrations required to meet the regulatory detection limits.

Experimental

Sample preparation

Radium-226 calibration standards were prepared from an in-house standard solution in a dedicated facility used for the preparation of aqueous radioactive sources for decay counting or mass spectrometry measurement. The calibration standards were diluted in 2% (v/v) HNO_3 .

Groundwater samples were also investigated to assess the impact of a more complex sample matrix. Samples were evaporated to dryness and redissolved in 2% (v/v) HNO $_3$. The solutions were then spiked with 226 Ra over a concentration range of 0.03-30 ppt to represent the concentrations expected following preconcentration.

High volume water samples (1 L) were spiked over the same concentration range as the groundwater samples to represent samples close to, and higher than, the regulatory discharge limits. Samples were acidified to pH 2 and passed through a chromatographic column to trap ²²⁶Ra [6]. The ²²⁶Ra was then eluted, evaporated to

incipient dryness and then made up in 5 mL 2% HNO $_3$, representing a concentration factor of ~200. Unspiked water samples were run through the same preconcentration procedure, and then measured to establish the elemental composition and confirm no contribution of polyatomic interferences to the background at m/z = 226. Matrix matched calibration standards were prepared by spiking water samples following preconcentration, which also enabled the recovery to be calculated ($\geq 70\%$ over the concentration range studied).

Instrumentation

An Agilent 8800 Triple Quadrupole ICP-MS (ICP-QQQ) was used throughout. The standard sample introduction system was used, comprising a quartz torch with 2.5 mm i.d. injector, a quartz spray chamber, glass concentric nebulizer, and nickel-tipped interface cones. The instrument operating conditions are summarized in Table 1.

Table 1. ICP-QQQ operating conditions; low matrix tuning is appropriate for samples where most of the matrix has been removed during analyte preconcentration.

Parameter	Setting	1		
Scan mode	Single Quad			
Plasma conditions	Low matrix (optimized for high sensitivity)	НМІ		
RF power (W)	1550			
Carrier gas (L/min)	1.07	0.60		
Dilution gas (L/min)	0	0.35		
Extract 1	0.0			
Extract 2	-200.0			
Omega Bias (V)	-100.0			
Omega lens (V)	13.6			
Octopole bias (V)	-8.0			
He cell gas (mL/min)	0 – 1.0	0.5 - 1.0		

Results and Discussion

Sensitivity of ICP-QQQ for ²²⁶Ra

The half-life of 226 Ra is relatively short with regards to ICP-MS measurements (1 Bq/kg is equivalent to 27.3 ppq, compared to long-lived 238 U (half-life 4.5×109 years), where 1 Bq/kg is equivalent to 8.0×107 ppq). In practice, this means that calibration should be performed using standards prepared for the radioisotope of interest, rather than calibrating using a long-lived or stable isotope as an analog. The instrument detection limits (IDLs) for several operating conditions (Q1 modes and cell gas flows), were calculated from a calibration curve prepared by spiking 2% (v/v) HNO $_3$ with 226 Ra at concentrations of 0.01–30 ppt (Table 2) .

Table 2. Limit of detection for different instrument conditions and cell gas flow rates.

Instrument mode	Single Quad			MS/MS		
He flow rate (mL/min)	0.0	0.5	1.0	0.0	0.5	1.0
Limit of detection (ppt)	0.08	0.10	0.02	0.04	0.04	0.07

The IDLs in Table 2 are close to the higher end of the regulatory limits quoted (0.03 ppt), and orders of magnitude higher than the lowest values (0.3 ppq). Measurement of ²²⁶Ra at environmentally relevant levels therefore requires an effective preconcentration step prior to ICP-QQQ analysis, to rival the detection limits of traditional alpha spectrometry measurement.

Interference removal by ICP-QQQ

Multiple potential interferences from polyatomic ions including $^{88}Sr^{138}Ba^+$, $^{87}Sr^{139}La^+$, $^{86}Sr^{140}Ce^+$, $^{208}Pb^{18}O^+$, $^{186}W^{40}Ar^+$, and $^{97}Mo^{129}Xe^+$ can potentially affect ICP-MS measurement of ^{226}Ra . Multiple separation stages prior to sample introduction are often required to remove the interferences. As an alternative approach, helium (He) collision mode was investigated for the removal of polyatomic interferences, initially by introducing up to 100 ppm Sr + Ba, Sr + La, Ce, W, and Pb standards. The background at m/z = 226 was 0 cps in single quad mode when using 0.5-1.0 mL/min He cell gas, confirming the ability of He mode to attenuate all the polyatomic ions. Given that the on-mass polyatomic interferences are formed during sample introduction and not in the collision/reaction cell (CRC), MS/MS was not required, so the instrument was operated in single quad mode throughout.

Groundwater samples from different locations in North West England were then analyzed to determine the impact of a more complex sample matrix on instrument performance. The samples were spiked with 226 Ra, and measured at varying He gas flow rates together with unspiked samples and blank solutions. Bismuth-209 was used as an internal standard. The impact of matrix suppression was overcome using robust plasma conditions and aerosol dilution with the High Matrix Introduction (HMI) system of the 8800. HMI allows higher matrix levels to be analyzed directly without requiring chemical separation prior to measurement, further reducing the total procedural time. The reduction in sensitivity when operating with 0.5 mL/min He cell gas was offset by the lower background, giving comparable or improved background equivalent concentrations (BECs) at m/z = 226 compared to no gas mode (Table 3). The sensitivity at 0.5 mL/min He cell gas is illustrated in the calibration plot shown in Figure 1.

Table 3. BECs of ^{226}Ra using no gas and He gas mode.

He flow rate (mL/min)	BEC (ppt)			
	Sample 1	Sample 2	Sample 3	
0	0.015	0.017	0.0085	
0.5	0.0083	0.0089	0.0092	
1.0	0.011	0.0092	0.013	

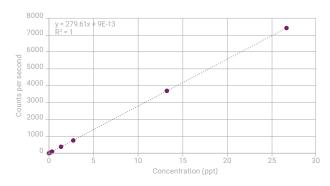


Figure 1. Calibration plot for ²²⁶Ra standards in single quad mode with 0.5 mL/min He.

Measurement of high volume water samples

In water samples, ²²⁶Ra was detected down to 0.03 ppt (1 Bq/L), which is equivalent to 5 mBq/L in the original sample, assuming a preconcentration factor of 200. The RSD was <10% at concentrations above 1.4 ppt (50 Bq/L), equivalent to 250 mBq/L in the original sample. The results demonstrate that ICP-QQQ combined with preconcentration from high volume water samples is capable of measuring ²²⁶Ra at concentrations relevant to regulatory discharge limits. Improved accuracy at the lower limits is potentially achievable through higher preconcentration factors.

Conclusions.

A method is presented that demonstrates the capabilities of ICP-QQQ for the measurement of the naturally occurring radionuclide ²²⁶Ra. The use of He collision gas effectively removes potential polyatomic interferences, while operating with HMI reduces the impact of matrix suppression. When combined with preconcentration using chromatographic separation techniques, the detection limits achievable are applicable to the regulatory limits for water. The measurement time of several minutes per sample represents a significant improvement compared to several days using traditional alpha spectrometry. The increase in sample throughput is potentially beneficial for routine monitoring of water supplies, as well as routine environmental monitoring at nuclear and industrial sites.

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Analysis of Radioactive Iodine-129 Using MS/MS with Oxygen Reaction Mode

Authors

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Keywords

radionuclide, iodine, ¹²⁹I, environmental, nuclear, xenon, NIST 3231 Level I and II, abundance sensitivity, oxygen on-mass

Introduction

lodine-129 is a long-lived radionuclide (half-life of 15.7 My) which has been released into the environment as a result of human activities such as nuclear weapons testing, accidents at nuclear power plants and especially by emissions from spent nuclear fuel reprocessing plants. The determination of iodine-129 in environmental samples is very difficult by ICP-MS due to the element's relatively low sensitivity, the very low concentrations at which 1291 must be determined, relative to potentially high levels of ¹²⁷I, the high background caused by ¹²⁹Xe impurities in the argon plasma gas, and possible polyatomic interference from ¹²⁷IH₂⁺. Iodine analysis is further complicated by the fact that it is rapidly volatilized from samples prepared using the acid digestions that are normal for ICP-MS analysis, so an alternative, alkaline sample solubilization and stabilization strategy is required. The isobaric interference from ¹²⁹Xe⁺ can be significantly reduced using ICP-QMS with an Octopole Reaction Cell operated in O₂ reaction mode, resulting in a measured ratio for 129 I/127 I of 10-7 in NIST 3231 SRM Level I (1). However, the problem of potential overlap due to tailing from ¹²⁷I and ¹²⁷IH remains, as the relative abundance of the ¹²⁹I to ¹²⁷I will typically exceed 10⁻⁷, which is of the same order as the abundance sensitivity (ability to separate adjacent peaks) of quadrupole ICP-MS (ICP-QMS). In order to overcome these challenges, ICP-QQQ operating in MS/MS mode with O2 reaction gas was applied to determine ultratrace levels of iodine-129 in aqueous samples.

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/Low matrix.

Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -190 V.

CRC conditions: O₂ gas at 0.8 mL/min,

Octopole bias = -18 V and KED = -1.5V. MS/MS O_2 on-mass mode was applied to measure iodine-127 and iodine-129 (Q1 = Q2 = 127 for iodine-127; Q1 = Q2 = 129 for iodine-129).

Reference materials and calibration standards: Calibration standards were prepared by diluting ¹²⁹I isotopic standards NIST SRM 3231 Level I and II (NIST, Gaithersburg MD, USA) with 0.5% TMAH in deionized water. The Level I Certified Value for ¹²⁹I/¹²⁷I = $0.981 \times 10^{-6} \pm 0.012 \times 10^{-6}$, Level II = $0.982 \times 10^{-8} \pm 0.012 \times 10^{-8}$. These reference materials were used to check the calibration linearity of the iodine isotopes and to validate the isotopic ratio of iodine-129 and iodine-127.

Results and Discussion

Optimization of oxygen cell gas flow

The oxygen gas flow rate was optimized by varying the O_2 flow over the full range of the mass flow controller (0–1.12 mL/min), while monitoring the ^{127,129}I signal and blank intensity, as shown in Figure 1. As the flow rate of O_2 increases, the background signal (due to ¹²⁹Xe) at m/z = 129 decreases rapidly, and the iodine signal remains high, dramatically improving the DL for ¹²⁹I.

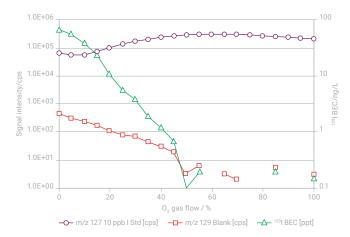


Figure 1. Profile of 127 I+, 129 Xe+ signals and estimated 129 I BEC. For the BEC calculation, the sensitivity of 129 I was assumed to be the same as 127 I. Scale of 0, flow: 100% = 1.12 mL/min.

Abundance sensitivity

Scan spectra over the mass range 127 to 129, covering both 127 I and 129 I, were acquired for the two SRMs, NIST 3231 Level I and II, using the Agilent 8800 ICP-QQQ in MS/MS on-mass mode with O_2 reaction gas. The overlaid spectra are shown in Figure 2. Excellent abundance sensitivity can be seen, with the sides of the intense (>109 cps) 127 I peak reaching baseline with no tailing of 127 I+ or 127 IH+ on 129 I+.

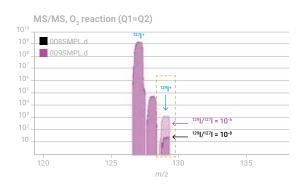


Figure 2. lodine spectra showing both 127 I and 129 I acquired using MS/ MS on-mass mode with O_2 cell gas. 127 IH+ remains to some extent while 127 IH $_2$ + is completely removed, as noted later.

Calibration curves for 127I and 129I

In order to check the linearity of both iodine isotopes, different concentration solutions of NIST 3231 SRM Level I were prepared in 0.5% TMAH and analyzed as calibration standards, as shown in Figure 3. The BECs for ^{127}I and ^{129}I were 2.9 $\mu g/L$ and 0.04 ng/L respectively, and the detection limits (3σ, n=10) were 0.26 $\mu g/L$ for ^{127}I and 0.07 ng/L for ^{129}I .

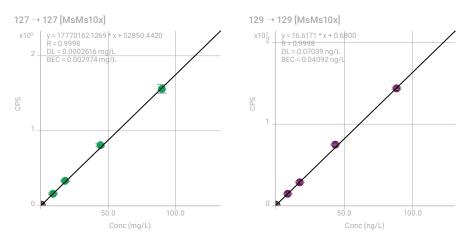


Figure 3. Calibration curve for iodine-127 (top) and iodine-129 (bottom) obtained from multiple dilutions of NIST 3231 SRM.

Analysis of NIST 3231 SRM Level I and Level II

The $^{129}\text{I}/^{127}\text{I}$ ratio in 10x diluted NIST 3231 SRM Levels I ($^{129}\text{I}/^{127}\text{I}$ = 0.981 x 10-6) and II ($^{129}\text{I}/^{127}\text{I}$ = 0.982 x 10-8) was measured using ICP-QQQ in MS/MS on-mass mode with O $_2$ cell gas. The results are summarized in Table 1. After subtracting the ^{129}I blank, the measured $^{129}\text{I}/^{127}\text{I}$ ratio of NIST 3231 SRM Levels I and II corresponded well with the certified values of 0.981x10-6 and 0.982x10-8 respectively. The good agreement with the certified ratio indicates that the potential interference of $^{127}\text{IH}_2^+$ on $^{129}\text{I}^+$ is completely removed by O $_2$ reaction with MS/MS mode.

Table 1. Analytical results for NIST 3231 Level I and Lev	/el II.
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Sample name	Dilution factor	Q1=Q2=127 CPS	Q1=Q2=129 CPS	¹²⁹ / ¹²⁷	¹²⁹ I/ ¹²⁷ I (average n = 5)	RSD (%)
NIST 3231 10 ⁻⁶ (¹²⁹ I/ ¹²⁷ I = 0.981 x 10 ⁻⁶)	10	594,277,896	585.6	0.971 x 10 ⁻⁶	 _ 0.981 x 10 ⁻⁶ 	0.8
		592,633,576	597.4	0.994 x 10 ⁻⁶		
		590,000,723	586.5	0.980 x 10 ⁻⁶		
		593,387,443	588.5	0.978 x 10 ⁻⁶		
		592,834,056	588.9	0.979 x 10 ⁻⁶		
NIST 3231 10 ⁻⁸ (¹²⁹ I/ ¹²⁷ I = 0.982 x 10 ⁻⁸)	10	608,737,949	15.1	1.12 x 10⁻8	 1.02 x 10 ⁻⁸ 	7.2
		608,536,242	14.8	1.07 x 10 ⁻⁸		
		602,626,536	14.2	0.979 x 10 ⁻⁸		
		603,091,763	13.9	0.929 x 10 ⁻⁸		
		603,250,003	14.5	1.03 x 10 ⁻⁸	_	
NIST Blank	10	600,444,851	8.3	_	_	_

Reference

1. The ultratrace determination of iodine 129 in aqueous samples using the 7700x ICP-MS with oxygen reaction mode, Agilent application note, 5990-8171EN.

More Information

The ultratrace determination of iodine 129 using the Agilent 8800 Triple Quadrupole ICP MS in MS/MS mode, Agilent publication, <u>5991-0321EN</u>.

Feasibility Study on the Analysis of Radioisotopes: Sr-90 and Cs-137

Authors

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Keywords

radioisotopes, radioactive, environmental, nuclear, strontium, ⁹⁰Sr, zirconium, cesium, ¹³⁷Cs, barium, abundance sensitivity, oxygen and hydrogen on-mass, nitrous oxide on-mass

Introduction

ICP-MS can be an effective analytical tool for the analysis of long half-life radioisotopes due to its high sensitivity, speed of analysis, low sample consumption, and ease of sample preparation. The challenge for ICP-MS analysis of radioisotopes arises from interferences; not only by polyatomic ions but also atomic isobar ions that cannot be separated even by high-resolution (HR-) ICP-MS.

Trace analysis of the radionuclide 90 Sr (half-life = 28.74 years) in environmental samples is of great interest. 90 Sr is a main fission product that may be present in the environment following accidental releases from nuclear power plants. Geiger-Muller (GM) detectors or Liquid Scintillation Counters (LSC) are used to measure 90 Sr, though both techniques require complex chemical separation prior to analysis, or long integration times. ICP-MS is also used to measure 90 Sr, especially when a quick turn-around time is desired. However detection limits of quadrupole ICP-MS are compromised by a spectral overlap from 90 Zr; in common with all direct isobaric interferences, the 90 Zr overlap is too close in mass to the 90 Sr to be resolved using sector field HR-ICP-MS, which is limited to a maximum resolution (M/ Δ M) of 10,000. This note describes a method for measuring trace 90 Sr in the presence of 90 Zr using ICP-QQQ in MS/MS reaction mode. Since it isn't possible to obtain 90 Sr, a natural isotope of strontium (88 Sr) was used to estimate the DL for 90 Sr. A similar approach was applied to 137 Cs (half-life = 30.0 years).

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/Low matrix.

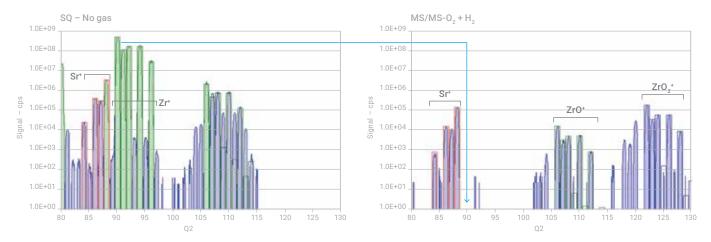
Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -190 V.

CRC and acquisition conditions: The following conditions were used for the analysis of ⁹⁰Sr and ¹³⁷Cs:

- For 90 Sr: MS/MS on-mass mode (Q1 = Q2 = 90) with O_2 + H_2 cell gas: 1 mL/min of O_2 and 10 mL/min of H_2 , Octopole bias = -5 V and KED = -13 V.
- For 137 Cs: MS/MS on-mass mode (Q1 = Q2 = 137) with N $_2$ O cell gas: 7 mL/min of N $_2$ O (10% N $_2$ O balanced in He, introduced via the 3rd cell gas flow line), Octopole bias = -5 V and KED = -13 V.

Radioactive Sr-90 (0, + H, on-mass mode)

Figure 1 shows spectra of a solution containing Sr and Zr (natural isotopes) acquired on the 8800 ICP-QQQ operated in Single Quad mode (Q1 operated as an ion guide to emulate conventional quadrupole ICP-MS) with no cell gas (left), and in MS/MS mode with O_2 + H_2 cell gas (right). As can be seen in the left hand spectrum, the overlap of 90 Zr+ on 90 Sr+ precludes the low-level determination of 90 Sr by conventional quadrupole ICP-MS. The spectrum on the right indicates that 90 Sr+ could be measured on-mass at m/z = 90 free from interference by 90 Zr+, since Zr+ reacts readily with the O_2 + H_2 gas to form ZrO+ and ZrO₂+. The signal-tonoise ratio for 90 Sr was improved by six orders of magnitude using MS/MS O_2 + H_2 reaction cell mode.



 $\textbf{Figure 1.} \ \ \text{Mass spectra of a solution containing 20 ppb Sr + 5 ppm Zr: (left) SQ no gas mode and (right) MS/MSO_2 + H_2 reaction mode.$

Figure 2 is a spectrum of 100 ppm Sr acquired using MS/MS on-mass mode with $O_2 + H_2$ reaction gas. The excellent abundance sensitivity (peak separation) of MS/MS mode can be confirmed. The peak sides reach the baseline with no tailing from the intense peak of the natural isotope of ⁸⁸Sr⁺. In addition, no ⁸⁸SrHH⁺ at m/z = 90 is formed in cell, even in a solution containing 100 ppm natural Sr.

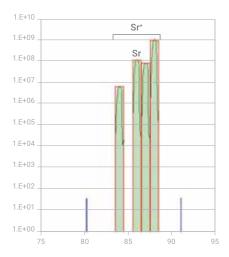


Figure 2. Spectrum of 100 ppm Sr solution acquired using MS/MS 0₂+ H₂ on-mass mode.

Radioactive Cs-137 (N₂O on-mass mode)

Figure 3 shows spectra of a solution containing Cs and Ba (natural isotopes) acquired on the 8800 ICP-QQQ operated in Single Quad mode with no gas mode (left), and in MS/MS mode with N_2 0 cell gas (right). As can be seen in the left hand spectrum, the 137 Ba+ overlap on 137 Cs+ is a problem in conventional quadrupole ICP-MS. As with 90 Sr, the right hand spectrum shows that 137 Cs+ could be measured on mass at m/z = 137, free from the 137 Ba+ interference. Ba+ reacts readily with N_2 0 to form BaO+ and BaOH+ while a part of the Cs+ analyte ion signal remains at its original mass (as shown by the substantial peak for 133 Cs in the right-hand spectrum).

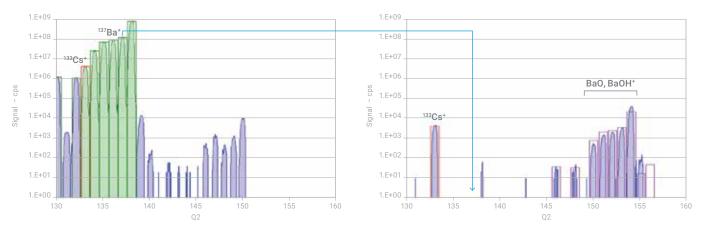


Figure 3. Mass spectra of a solution containing 20 ppb Cs + 5 ppm Ba: (left) SQ no gas and (right) MS/MS N₂O reaction mode.

Estimated BEC and DL for Sr-90 and Cs-137

The BEC and DL for two radioisotopes, 90 Sr and 137 Cs, were estimated from these spectra as summarized in Table 1. This feasibility study demonstrates the potential of ICP-QQQ for the measurement of radioisotopes such as 90 Sr and 137 Cs.

Table 1. Estimated BEC and DL for 90Sr and 137Cs.

Radioisotope	BEC (ng/L)	DL (ng/L)
⁹⁰ Sr	0.08	0.23
¹³⁷ Cs	2.9	15

Determination of Trace ²³⁶U as UOO+ using ICP-QQQ Oxygen Mass-shift Method

Authors

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Keywords

uranium, uranium-236, O_2 cell gas, mass-shift, extended mass range

Introduction

Uranium-236 is a long-lived radionuclide that is created from the naturally occurring trace isotope ²³⁵U (0.72% abundance) by thermal neutron capture. This process leads to a natural abundance of ²³⁶U in the range from 10⁻¹⁴ to 10⁻¹³ relative to the major ²³⁸U isotope (²³⁶U/²³⁸U). ²³⁶U is also created during the process of uranium enrichment for nuclear fuel or weapons. The ²³⁶U/²³⁸U ratio is increased up to 10⁻³ in spent nuclear fuel, with background levels in the environment at around 10⁻⁷ to 10⁻⁸ as a result of global fallout. The ²³⁶U/²³⁸U isotope ratio can therefore be used as a sensitive method to trace the accidental release of enriched uranium fuel, spent fuel, and nuclear waste.

The challenges for ICP-MS for this application are the interference on 236 U+ by the hydride ion 235 UH+, and the contribution at m/z 236 from tailing of the 235 U+ and 238 U+ peaks. The hydride overlap and peak tailing are more problematic in samples that have been enriched, as these samples contain a higher proportion of 235 U. Uranium was measured via its dioxide ion, UO_2^+ , due to the efficient conversion (almost 100%) of U+ to UO_2^+ with O_2 cell gas.

Experimental

Instrumentation: Agilent 8900 Advanced Applications configuration ICP-QQQ with PFA nebulizer (p/n G3139-65100).

Plasma tuning: RF power = 1550 W, sampling depth = 8.0 mm, nebulizer gas flow rate = 0.80 L/min, make-up gas flow rate = 0.30 L/min, and peristaltic pump = 0.1 rps.

Cell tuning: Octopole bias = 0 V, KED = -10 V, O_2 cell gas flow = 0 to 35% of full scale (0 to 0.53 mL/min).

Sample preparation: Uranium solutions were prepared at suitable concentrations by diluting SPEX multi element standard XSTC-331 (SPEX CertiPrep, Metuchen, NJ, USA) with de-ionized water. All samples, blank, and rinse solutions were spiked with high purity TAMAPURE 100 HNO $_3$ (Tama Kagaku, Saitama, Japan) to a concentration of 1%.

UO+ and UOO+ formation as a function of O2 cell gas flow rate

The rate of formation of UO+ and UO2+ was studied as a function of O2 cell gas flow rate. A solution containing 10 ppb uranium (1000x dilution of XSTC-331) was introduced into the ICP-QQQ. The signals of 238 U+, 238 U16O+, and 238 U16O16O+ were measured via three mass pairs (Q1 \rightarrow Q2) = (238 \rightarrow 238), (238 \rightarrow 254), and (238 \rightarrow 270), and plotted against the O2 cell gas flow rate. The octopole bias (Octp Bias) voltage was optimized to give the maximum UO2+ signal (0 V). Figure 1 shows that UO+ formation reaches a maximum at an O2 flow rate of 5% of full scale (equivalent to 0.074 mL/min as O2). Above 0.075 mL/min flow rate, the formation of UO+ decreased, while the formation of UO2+ increased, reaching a maximum at an O2 flow of 22% of full scale (0.33 mL/min). This indicates the conversion of UO+ to UO2+ via a chain reaction. The 8900 ICP-QQQ was optimized for highest sensitivity for the UO2+ product ion.

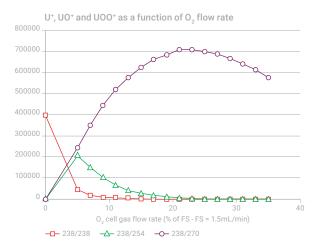


Figure 1. U* (238 \Rightarrow 238), U0* (238 \Rightarrow 254), and U0 $_2$ * (238 \Rightarrow 270) as a function of O $_2$ cell gas flow rate.

Effect of product ion selection on hydride ion formation rate

The hydride ratio was measured at the optimal O_2 flow rate for U+ and each of the U-oxide product ions: $^{238}UH^+/^{238}U^+$, $^{238}UOH^+/^{238}UO^+$, and $^{238}UO_2H^+/^{238}UO_2^+$.

A sample containing 50 ppb U (200x diluted XSTC- 331) was introduced for the measurement of the hydride formation ratio. Ten replicate measurements were made, with integration times of 1 s and 10 s for the analyte and hydride ions respectively. The results are summarized in Table 1. The data shows that measuring UO $^+$ decreases the hydride ratio by a factor of \sim 20, while measuring UO $^+$ leads to more than a three orders of magnitude improvement, reducing the hydride ratio to 10^{-8} .

Uranium detection limit

The detection limit (DL) of U was estimated using the UO_2^+ method. A blank solution was introduced and the signal of the mass pairs (236 \rightarrow 268 and 238 \rightarrow 270) corresponding to $^{236}U^+\rightarrow^{236}UO_2^+$ and $^{238}U^+\rightarrow^{238}UO_2^+$ were measured using an integration time of 10 s. The results in Table 2 are based on 10 replicate measurements. The DL for ^{236}U was calculated from the concentration equivalent to three times the standard deviation of the background, using the sensitivity of $^{238}UO_2^+$ given in Table 1 and the background for mass pair 236 \rightarrow 268 in Table 2. The DL for uranium-236 was calculated to be 0.50 ppq (fg/g).

Table 1. UH+/U+ ratios obtained by measuring uranium as U+, UO+, and UO₂+.

	O ₂ cell gas flow rate (%)	l	J+ analysis		UF	UH⁺ analysis			
	now rate (%)	Mass pair for U ⁺	Counts	RSD	Mass pair for UH ⁺	Counts	RSD	_	
		Q1/Q2	cps	%	Q1/Q2	cps	%	_	
as U⁺	0	238/238	24168974	2.8	239/239	1578.5	0.6	6.53E-05	
as UO⁺	5	238/254	14152816	4.2	239/255	48.9	4.3	3.46E-06	
as U00⁺	22	238/270	40527770	2.0	239/271	2.3	20.8	5.68E-08	

Table 2. Uranium background noise.

236/268		238/270	238/270			
Counts	RSD	Counts	RSD			
cps	%	cps	%			
0.15	90.3	0.18	51.1			

Conclusions

The Agilent 8900 ICP-QQQ operating in MS/MS mode with O_2 cell gas is suitable for the measurement of U via its reaction product ion UO_2^+ . This approach was successful in reducing the contribution from the hydride ion (i.e. 235 UH overlap on 236 U). The formation of 235 UH was decreased by three orders of magnitude compared to direct, on-mass measurement of U+. MS/MS mode with O_2 cell gas gave a UO_2 H+/ UO_2 + ratio in the 10^{-8} range, without the use of a desolvation system. The results suggest that the approach could be successful in reducing the interference of 235 UH+ on 236 U+, even in samples containing enriched U.

More Information

Using ICP-QQQ for $\rm UO_2^+$ product ion measurement to reduce uranium hydride ion interference and enable trace $^{236}\rm U$ isotopic analysis, Agilent publication $\underline{\rm 5991}$ - $\underline{\rm 6553EN}$.

Measurement of Neptunium in the Presence of Uranium: Benefits of Low Abundance Sensitivity and Oxygen Reaction Mode

Author

Glenn Woods Agilent Technologies, UK

Keywords

neptunium, radiochemistry, abundance sensitivity, oxygen reaction mode

Introduction

Neptunium is present in the environment at ultratrace levels due to natural neutron capture, nuclear bomb testing, and as a decay product of ²⁴¹Am. ²⁴¹Am is used in ionizing smoke detectors, radiography, and a neutron source, among other uses. By far the greatest quantity of Np is formed during energy production within uranium fission reactors. The predominant isotope formed is ²³⁷Np, with approximately 50 metric tonnes per annum being produced in nuclear waste. As the half-life of ²³⁷Np is ~2.14 billion years, ²³⁷Np in existence today is solely from the previously mentioned processes rather than remaining from the formation of the earth. However, the relatively long half-life ensures its persistence. Np will readily form aqueous solutions (more so than any other actinide element). It also attaches to particles and colloids rather than getting trapped in humic media (such as soil and peat). These properties mean that Np is fairly mobile once in the environment. Its high affinity for calcium-rich media causes it to concentrate within concrete and bone etc.

Trace and ultratrace measurement of ²³⁷Np is hindered by the presence of uranium within the sample. The biggest potential interference comes from peak broadening of the adjacent ²³⁸U isotope. This Abundance Sensitivity (AS) interference is difficult to overcome. AS depends on the fundamental design of the spectrometer – such as the mass separation process (e.g. quadrupole or magnets), vacuum system, and electronics. Furthermore, minor but important polyatomic interferences from the hydrides of lighter U isotopes; ²³⁶U¹H, ²³⁵U¹H₂, ²³⁵U²H, ²³⁴U¹H²H hinder the measurement of ²³⁷Np. Regardless of the interference source, its affect will vary depending on the concentration of uranium (and its isotope ratio), potentially causing false and variable measurements.

Experimental

Instrumentation: An Agilent 8900 Advanced Applications configuration ICP-QQQ was used. The instrument version features Axial Acceleration across the ORS⁴ collision/reaction cell that gives a higher product ion yield when using reaction chemistry.

Tuning: Np was measured under two sets of MS/MS conditions: on-mass (using no gas) and mass-shift (using O_2 reaction gas). In the latter mode, ²³⁷Np is shifted away from the UH_x interferences allowing Np to be measured as the product ion NpO₂+, free from interference, at m/z 269.

Calibration: Np was spiked into a 10 mg/L (ppm) U matrix to produce a set of calibration standards at 0.0, 0.19, 0.95, 1.9, 19.0, 95.0 ng/L (ppt).

The Single Quad mass scan in Figure 1 shows the problem associated with AS when the U concentration is relatively high. As can be seen, the 238 U peak overlaps the 237 Np peak, impeding the trace level measurement of Np. Conversely when operating the ICP-QQQ in MS/MS mode, the peak overlap on Np is eliminated. This improvement is due to two separate mass separations taking place, improving the AS from $\sim\!10^{-7}$ to $<<\!10^{-10}$. The background is significantly reduced under MS/MS mode but not eliminated. Uranium can form various hydride interferences that are not related to (or removed by) AS. However, reaction chemistry can be used to remove interference-based background levels.

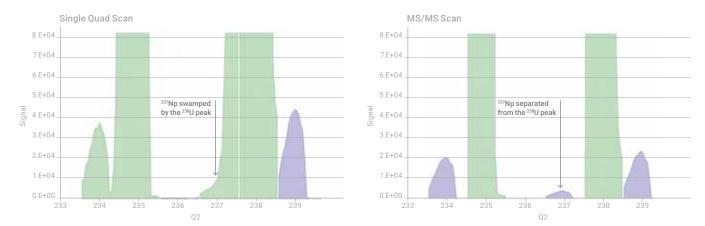


Figure 1. Spectrum of ²³⁷Np in presence of 10 ppm U. Left: Single Quad scan and Right: MS/MS mode. MS/MS mode eliminates the peak tail on the low mass side of the intense ²³⁸U peak.

To check the reaction efficiency of oxygen as a cell gas for this study, a spiked U matrix was measured under MS/MS mass-shift mode with $\rm O_2$ reaction gas. The Np spike was 1000x lower than the previous scans at 950 ppq (0.95 ng/L). Figure 2 shows the mass scan of the NpO $_2$ (and UO $_2$) product ions. It is worth noting that during quantitative analysis (rather than scanning, as shown in Figure 2), all the U isotopes would be eliminated by Q1, which would be set to m/z 237. The conversion efficiency of Np to NpO $_2$ was found to be 99%. Only 1% of total Np signal converted to NpO.

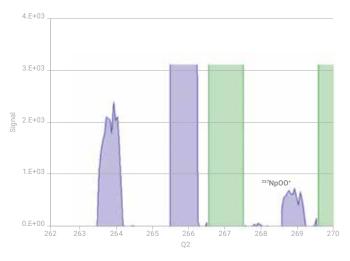


Figure 2. 950 ppt Np in 10 ppm U measured in MS/MS mass-shift mode with 0, cell gas.

Figure 3 shows the calibration graphs for Np in a 10 ppm U matrix generated in no gas mode (left) and oxygen reaction gas mode (right). Identical solutions were analyzed in both cases. The improvement in BEC and DL can be clearly seen in oxygen mass-shift mode. The DL and BEC under no gas conditions were 1.9 ppq and 2.4 ppq. Using $\rm O_2$ mode and measuring Np as NpO $_2$ improved the DL to 0.56 ppq and the BEC to 0.32 ppq (pg/L).

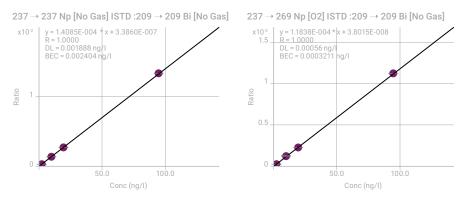


Figure 3. Np calibration in a 10 ppm U matrix. Left: no gas mode. Right: O_2 reaction gas mode – showing a 7.5x reduction in BEC. All UH-based interferences were avoided by measuring 237 N as 237 Np $^{16}O_2$.

Conclusions

The Agilent 8900 ICP-QQQ offers unrivalled abundance sensitivity performance allowing an ultratrace element to be measured in the proximity of a major matrix isotope. For the determination of Np, the removal of uranium-based interferences is essential, as U is present within the environment at significantly higher concentrations than Np. The unique MS/MS capability of the 8900 ICP-QQQ removes peak overlaps and uranium hydride-based interferences.

Lead Isotope Analysis: Removal of ²⁰⁴Hg Isobaric Interference on ²⁰⁴Pb using ICP-QQQ MS/MS Reaction Cell

Author

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Keywords

lead, isotope, ratio, geochronology, dating, mercury, artifacts, precious metals, food, ammonia, on-mass

Introduction

Lead isotope ratio analysis is important as it is used for Pb-Pb dating in geochronology, and to trace the origin of artifacts, precious metals and even foodstuffs. The natural isotopic pattern of lead varies more than any other element in the periodic table, because three of its isotopes are formed from the radioactive decay of uranium ($^{235}\text{U} \rightarrow ^{207}\text{Pb}$; $^{238}\text{U} \rightarrow ^{206}\text{Pb}$) and thorium ($^{232}\text{Th} \rightarrow ^{208}\text{Pb}$). The Pb isotopic pattern can therefore vary depending upon the geology of the rocks and minerals from which the lead was extracted, and the age of the material. In geochronology, the constant rate of U/Th decay allows the Pb/Pb, U/Pb and Th/Pb ratios to be used to date the age of rocks using a so-called geological clock.

When Pb ratios are measured, it is often necessary to correct for the lead naturally present in the sample, and the only non-radiogenic isotope of Pb (204Pb; natural or common lead), is used for this purpose. For Pb-Pb dating, 204Pb is the reference isotope against which the radiogenic isotopes are compared (206Pb/204Pb; 207Pb/204Pb). Unfortunately 204Pb is directly overlapped by an isotope of Hg (204Hg), which makes accurate measurement of 204Pb impossible by ICP-MS. Mass resolution of 204Pb from 204Hg is far beyond the capability of any commercial high-resolution (HR-) ICP-MS system, and until recently there has been no reliable chemical means to remove the Hg interference, so mathematical correction has been employed, which introduces error. Mercury does however undergo a gas-phase charge-transfer reaction with ammonia gas (NH₃), a reaction that can be utilized in the collision/reaction cell of a suitably equipped ICP-MS as follows:

 $Hg^{+} + NH_{3} \rightarrow Hg^{0} + "NH_{3}^{+"}$

This reaction offers the potential to remove the ²⁰⁴Hg interference from ²⁰⁴Pb, and could be applied to either solution or laser-based ICP-MS analysis.

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/General purpose.

Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -170 V.

CRC conditions: NH₃ gas (10% in He) at 1.7 mL/min, Octopole bias = -8 V,

KED = -8 V.

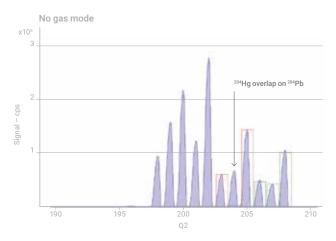
Acquisition parameters: Three acquisition modes were compared:

- No gas: No reaction cell gas; Single Quad (SQ) mode with Q1 operating as an ion guide.
- NH₃ bandpass: Ammonia reaction gas; SQ mode with Q1 operating as a bandpass filter.
- NH₃ MS/MS: Ammonia reaction gas; MS/MS mode with Q1 operating as a mass filter at unit mass resolution.

Results and Discussion

Removal of ²⁰⁴Hg⁺ interference on ²⁰⁴Pb⁺

A preliminary study showed that Pb is almost unreactive with NH $_3$ cell gas (<0.5% loss of Pb signal) indicating that on-mass sensitivity for Pb should be maintained. On-mass measurement of Pb in NH $_3$ cell gas mode was therefore investigated in the presence of Hg at 10 ppb. Figure 1 displays the spectra obtained in no gas (left) and NH $_3$ cell gas (right) modes. The 204 Hg interference on 204 Pb can be clearly seen in the no gas spectrum, while it has been completely removed under NH $_3$ reaction mode with MS/MS. A perfect isotopic pattern match was confirmed for Pb in NH $_3$ mode.



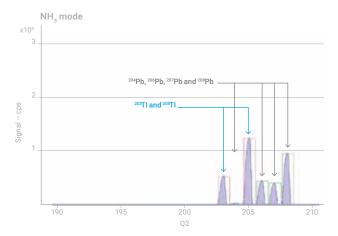


Figure 1. Standard solution (1 ppb each of Pb and Tl) spiked with 10 ppb Hg without cell gas (left) and with NH₃ (right) using MS/MS mode. Note the ²⁰⁴Hg interference on ²⁰⁴Pb in no gas mode.

Effectiveness of MS/MS

The $\mathrm{NH_3}$ reaction that removes the $^{204}\mathrm{Hg}$ interference would also work in the reaction cell of a single quadrupole ICP-MS (ICP-QMS), but ammonia is a highly reactive gas and can produce many adduct cluster ions, for example from Rare Earth Elements (REEs), see Table 1. The complex matrix composition of many natural samples means that the results obtained with $\mathrm{NH_3}$ cell gas in ICP-QMS are often extremely unreliable. With the 8800 ICP-QQQ, MS/MS mode allows all the co-existing matrix elements to be rejected by Q1, so only the target ions ($^{204}\mathrm{Pb}$ and $^{204}\mathrm{Hg}$) enter the CRC. The $\mathrm{NH_3}$ reactions are therefore controlled and consistent, and no overlapping reaction product ions are formed from other elements in the sample.

Table 1. Some possible Rare Earth Element cluster ions that can form in the CRC of an ICP-QMS when using $\mathrm{NH_3}$ reaction gas – the list is by no means exhaustive.

Mass	Potential Cluster Ions of REE
204	Eu(NH ₃) ₃ ; Yb(NH ₃) ₂ ; Ce(NH ₃) ₄
205	Yb(NH ₃) ₂ ; Gd(NH ₃) ₃
206	Yb(NH ₃) ₂ ; Lu(NH ₂) ₂ ; La(NH ₃) ₄ ; Ce(NH ₃) ₄ ; Gd(NH ₃) ₃
207	La(NH ₃) ₄ ; Yb(NH ₃) ₂ ; Gd(NH ₃) ₃
208	$Ce(NH_3)_4$; $GdNH(NH_3)_2$; $TbNH(NH_3)_2$; $Yb(NH_3)_2$; $Gd(NH_3)_3$

To check the formation of cluster ions, the ICP-QQQ was operated with $\mathrm{NH_3}$ cell gas; "Single Quad bandpass" and MS/MS modes were compared for the measurement of a 50 ppb REE mix. Figures 2a and 2b display the spectra obtained using bandpass and MS/MS conditions, respectively.

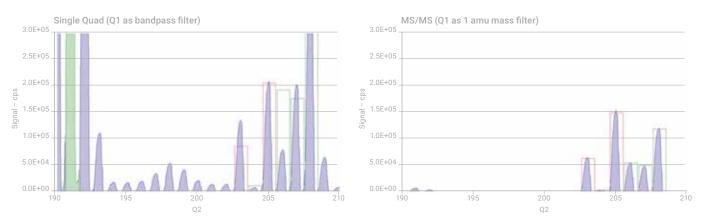


Figure 2. Cluster ion formation for 50 ppb REE standard in ammonia mode. Figure 2a (left): REE cluster ion formation using ammonia cell gas in bandpass mode; REE's are allowed into the cell if Q1 is operated as a bandpass filter. The REE cluster ions can be seen at all masses including those for Hg, Tl, Pb and Bi. Figure 2b (right): The identical sample under the same ammonia conditions but this time with Q1 operated at unit mass resolution (MS/MS mode). The REE's are removed from the ion beam before they can enter the cell and form reaction by-products.

²⁰⁴Pb/²⁰⁸Pb isotope ratio analysis in presence of Hg

To check the effectiveness of the 204 Hg removal, the 204 Pb/ 208 Pb ratio was measured in a 1 ppb lead solution spiked with increasing Hg concentration. Table 2 displays the measured Pb ratio results (without any mass bias correction), showing that the Pb isotope ratio remained constant, regardless of the Hg content.

Table 1. Uncorrected isotopic ratios measured in 1 ppb Pb solutions containing mercury at varying concentrations. The lead isotopic ratio 204/208 is not influenced by the presence of Hg.

	²⁰⁴ Pb	²⁰⁸ Pb	IR (204/208)
Sample		CPS	
Theoretical	NA	NA	0.02671
Pb	3518.5	136124.8	0.02585
Pb Hg 5 ppb	3510.0	139585.9	0.02515
Pb Hg 10 ppb	3439.2	132796.4	0.02590
Pb Hg 20 ppb	3464.8	134417.7	0.02578

Conclusions

With the successful removal of the 204 Hg interference on the natural 204 Pb isotope, ICP-QQQ displays great promise for Pb/Pb and U/Pb dating and for other applications where accurate measurement of 204 Pb is required.

Fractionation of Sulfur Isotope Ratio Analysis in Environmental Waters

Authors

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Keywords

sulfur, sulfur isotope ratio, isotope ratio analysis, fractionation, mass-shift

Introduction

Sulfur isotope ratio (IR) data is a useful indicator in geochemical and biochemical studies [1]. In stable sulfur isotope analysis, the variation in the $^{34}\text{S}/^{32}\text{S}$ IR is calculated and reported as a deviation or delta (δ) in ^{34}S abundance relative to a standard material, the troilite (iron sulfide) mineral from the Canyon Diablo meteorite. This standard is referred to as δVCDT (Vienna Canyon Diablo Troilite). Natural variations in ^{34}S abundance, expressed in parts per thousand or "per mil" (%), can be of the order of -50% to +40% (and occasionally much greater), due to redox reaction [2]. In this study, triple quadrupole ICP-MS (ICP-QQQ) was investigated as a fast and simple technique for S IR analysis. ICP-QQQ is a tandem ICP-MS that can resolve spectral interferences using reaction cell technologies. Using the method described in this paper, ICP-QQQ can measure S at a low concentration (background equivalent concentration < 0.2 ppb in UPW) with high sensitivity (^{32}S > 10000 cps/ppb).

Experimental

Instrumentation: Agilent 8900 Advanced Applications configuration ICP-QQQ with PFA nebulizer. Self-aspiration mode was used for better precision.

Tuning: O₂ mass-shift method. Tuning conditions are summarized in Table 1.

Method: the following procedures were used for the accurate determination of sulfur IRs:

- Matrix matching: all samples were diluted by the matrix blank, which contained 50 ppm Ca and 100 ppm NaCl in 1% HNO₃.
- Concentration matching: each sample was diluted by the matrix blank to ~0.5 ppm S concentration. This dilution was done to remove any errors caused by signal count differences. For example, NASS 5 was diluted 2000 times and mineral water A was diluted 10 times.
- Mass bias correction: to correct mass bias (including mass-bias drift), sample-standard bracketing was applied. IR of 0.5 ppm IAEA-S-1 [3] was measured before and after the IR analysis of each sample. The average of the IRs for the standard was used to correct the mass-bias and the drift.

 Table 1. ICP-QQQ tuning and method parameters.

Tuning parameter	Unit	Value
RF power	W	1550
Sampling depth	mm	8.0
Nebulizer gas flow rate	L/min	0.90
Make-up gas flow rate	L/min	0.30
Extract 1	٧	-80
Extract 2	V	-150
Omega	V	10.0
Omega bias	V	-120
Octp Bias	v	-5.0
Axial Acceleration	v	2.0
KED	v	-8.0
Cell gas		Oxygen
Cell gas flow rate	mL/min	0.45
Method parameter	Unit	Value
Integration time	s	1 and 5 for 32S and 34S
Number of sweeps	-	1000
Number of replicates	-	10
1% HNO ₃ rinse	s	20
50 ppm/100 ppm NaCl rinse	s	30
Load time	s	30
Stabilization time	s	30
	RF power Sampling depth Nebulizer gas flow rate Make-up gas flow rate Extract 1 Extract 2 Omega Omega bias Octp Bias Axial Acceleration KED Cell gas Cell gas flow rate Method parameter Integration time Number of sweeps Number of replicates 1% HNO ₃ rinse 50 ppm/100 ppm NaCl rinse Load time	RF power Sampling depth Mebulizer gas flow rate L/min L/min L/min Extract 1 Extract 2 Omega V Omega V Ottp Bias V Axial Acceleration KED Cell gas Cell gas flow rate Method parameter Unit Integration time Number of sweeps Number of replicates 1% HNO ₃ rinse S Load time W Mmm Memm W Mmm Mmm Mmm Mmm

Figure 1 shows a spectrum of three sulfur isotopes in a blank and 10 ppb S standard measured by ICP-QQQ in $\rm O_2$ mass-shift mode. The two spectra show the low BEC of sulfur in the blank, which allows accurate S IR analysis.

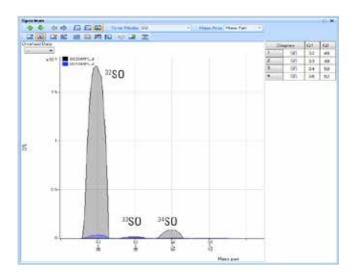


Figure 1. Spectra of S isotopes measured in O_2 mass-shift mode. The blank is indicated in blue and the 10 ppb S standard in grey. The spectra show that the BEC of the blank is < 200 ppt.

Synthetic samples were prepared and analyzed. Two standards, IAEA S-1 ($^{34}\delta$ = -0.3%) and IAEA S-2 ($^{34}\delta$ = +22.6%), were mixed to make four synthetic samples with a theoretical S IR of $^{34}\delta$ = -0.3, 5.4, 11.2 and 22.6 %. Each sample was measured six times, and the average IR and precision (as two times the standard deviation) were calculated. As shown by the linearity of Figure 2, the measured $^{34}\delta$ values were in excellent agreement with the theoretical values.

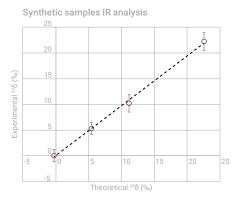


Figure 2. Sulfur IR of four synthetic samples.

The developed method was used to analyze seven samples: sulfuric acid (Tamapure AA-100); a Japan river water CRM, JSAC0301; a hot spring water, IKAHO; three mineral waters A, B, and C; and a seawater SRM, NASS-5. The concentration of S was first determined in each sample. The samples were then diluted with the matrix blank (50 ppm Ca + 100 ppm NaCl) to \sim 0.5 ppm of S. The IR of each sample was measured 10 times to determine the average value and precision (as two times the standard deviation). The results given in Figure 3 show \pm 1.2 to \pm 1.7 % error.

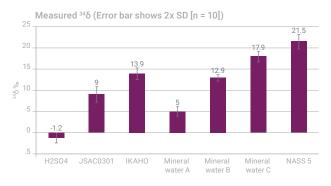


Figure 3. Measured δ^{34} S of seven samples.

Conclusions

The Agilent 8900 Advanced Applications configuration ICP-QQQ is ideally suited to 34 S/ 32 S isotope ratio analysis. The analysis can provide valuable information for sample characterization in natural systems or to monitor anthropogenic impact. The 8900 ICP-QQQ provides a low background and high sensitivity for sulfur, which enabled a method to be developed that required the sample to be diluted with the matrix blank before analysis. The precision of the IRs achieved was excellent at 1.2–1.7 % (as two times the standard deviation).

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More Information

Sulfur isotope fractionation analysis in mineral waters using an Agilent 8900 ICP-QQQ, Agilent publication <u>5991-7285EN</u>.

Direct Strontium Isotopic Analysis of Solid Samples by LA-ICP-MS/MS

Authors

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Martín Resano, University of Zaragoza, Spain

Keywords

isotopic analysis, LA-ICP-MS/MS, LA-ICP-QQQ, strontium, geology

Introduction

Strontium has four stable isotopes: 84 Sr (0.56%), 86 Sr (9.86%), 87 Sr (7.0%), and 88 Sr (82.58%). 87 Sr is either formed during nucleosynthesis with other stable Sr isotopes or via beta decay from 87 Rb (half-life of 4.88 x 10 10 years): 87 Rb [] 87 Sr + β^- + \bar{v} . Consequently, a high 87 Sr/ 86 Sr ratio is observed in rocks that are geologically old or which contain a high concentration of Rb (high Rb/Sr ratio). The 87 Sr/ 86 Sr ratio has been widely studied and reported in geological studies [1].

Measuring the ⁸⁷Sr/⁸⁶Sr ratio using mass spectrometry techniques is challenging because of the isobaric overlap of the signals from ⁸⁷Rb and ⁸⁷Sr. Chemical separation can be used to isolate Sr from Rb before analysis by ICP-MS. However, a simpler method uses triple quadrupole ICP-MS (ICP-QQQ) and chemical reaction in the CRC with a reactive gas. In this study, Laser Ablation coupled to ICP-QQQ (LA-ICP-QQQ) in MS/MS mode with CH₃F/He reaction gas was used to resolve the ⁸⁷Rb interference on ⁸⁷Sr. This approach allowed the direct Sr isotopic analysis of solid samples [2].

Experimental

A preliminary study showed that better precision was obtained using wet plasma conditions. The experimental setup shown in Figure 1 was used throughout. De-ionized water was continuously aspirated using a standard nebulizer. The sample aerosol that was generated by the LA system was carried by helium gas. Before being delivered to the plasma, the dry aerosol was combined with the liquid aerosol in the spray chamber, which was chilled to 2 °C.

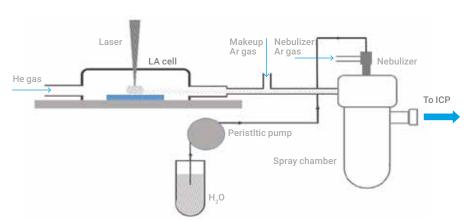


Figure 1. Schematic diagram of LA-ICP-QQQ using wet plasma conditions.

Instrumentation: An Analyte G2 193 nm ArF*excimer-based LA-unit (Teledyne CETAC Technologies, USA) equipped with a HELEX 2 ablation cell was coupled to an Agilent 8800 #100. The ICP-QQQ was fitted with a standard sample introduction system.

Method: Tuning conditions and method parameters are given in Table 1.

Reaction cell method: The CH $_3$ F/He (1:9) cell gas was introduced via the ICP-QQQ's fourth cell gas mass flow channel (0-1 mL/min as O $_2$). Rb $^+$ ions do not react with CH $_3$ F, whereas Sr $^+$ reacts with CH $_3$ F to form SrF $^+$. Thus 86 SrF $^+$, and 88 SrF $^+$ can be measured as the corresponding 86 SrF $^+$, 87 SrF $^+$ and 88 SrF $^+$ reaction product ions, free from interference.

Mass bias correction: The instrumental mass bias was corrected for using a double correction approach: internal correction assuming a constant ⁸⁸Sr/⁸⁶Sr isotope ratio (Russell's law, given below), followed by external correction in a sample-standard bracketing (SSB) approach using NIST 612 glass SRM.

$$\begin{split} &R^{87\text{Sr/86Sr}}_{\text{ sample, corrected}} = R^{87\text{Sr/86Sr}}_{\text{ sample, measured}} \times \left(m^{87\text{Sr}}/m^{86\text{Sr}}\right)^{\text{f}} \\ &f = In\left[R^{88\text{Sr/86Sr}}_{\text{ true}} / R^{88\text{Sr/86Sr}}_{\text{ measured}}\right] / In\left[m^{88\text{Sr}}/m^{86\text{Sr}}\right] \end{split}$$

Samples: Seven geological reference materials (RMs) were analyzed for their Sr isotopic composition. The RMs were selected to cover a wide range of matrix composition, Sr concentration, and Rb/Sr elemental ratio, as summarized in Table 2.

Table 1. LA-ICP-QQQ tuning conditions.

Laser Ablation		
Energy density	J/cm²	3.54
Repetition rate	Hz	40
Scan speed	μm/s	15
Beam size	μm	20-85
He carrier gas flow	L/min	0.42

ICP-QQQ		
RF power	W	1550
Sampling depth	mm	3.5
Nebulizer gas flow	L/min	1.0
Make-up gas flow	L/min	0.33
CH ₃ F/He cell gas flow	mL/min	0.90
Dwell time per acquisition point	ms	300
Acquisition time per replicate	s	60
Number of replicates		12
Total analysis time per sample	min	15.55

Removal of ⁸⁷Rb overlap using MS/MS mass-shift mode with CH₃F/He cell gas

Seven RMs were selected to cover a wide range of Rb/Sr ratios. The 87 Sr/ 86 Sr ratio was measured in each RM. For comparison purposes, the analysis was done using a no gas on-mass method and the CH $_3$ F/He mass-shift method. Figure 2 shows the measured 87 Sr/ 86 Sr and 88 Sr/ 86 Sr ratios obtained with the two methods as a function of the Rb/Sr ratio.

With both methods, a constant 88 Sr/ 86 Sr ratio was obtained regardless of the sample type. However, the measured 87 Sr/ 86 Sr ratio increased in no gas mode, indicating an interference from 87 Rb on 87 Sr. In contrast, the 87 Sr/ 86 Sr ratio measured in CH $_{3}$ F/He mode remained constant, regardless of the Rb/Sr ratio, showing that the method was effective at removing the 87 Rb isobaric overlap on 87 Sr.

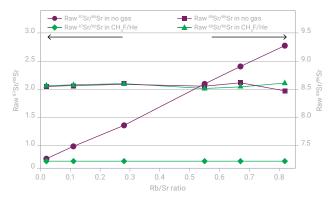


Figure 2. ⁸⁷Sr/⁸⁶Sr and ⁸⁸Sr/⁸⁶Sr isotope ratios measured using LA-ICP-QQQ in no gas and CH₃F/He cell gas modes. *Reproduced from J. Anal. At. Spectrom.*, 2016, 31, 464–472 with permission from the Royal Society of Chemistry.

Determination of 87Sr/86Sr ratio in seven RMs

The method was used to determine the ⁸⁷Sr/⁸⁶Sr ratio in seven RMs. The results are summarized in Table 2. After mass bias correction, excellent agreement was obtained between the measured ⁸⁷Sr/⁹⁶Sr ratios and the recommended reference values, even in samples with a high Rb content.

Table 2. 87Sr/86Sr isotope ratio results in seven reference materials.

Reference material	Type Rb/Sr ratio	Chem	hemical composition of the reference materials (%)					⁸⁷ Sr/ ⁸⁶ Sr rati	0						
			Al ₂ O ₃	CaO	FeO	K ₂ O	MgO	MnO	Na ₂ O	SiO ₂	Expe	imental	Recom	mended	Error (%)
USGS BHVO-2G	Basalt	0.02	13.6	11.4	11.3	0.51	7.13	0.17	2.4	49.3	0.70351	±0.00034	0.703469	±0.000007	0.006
USGS NKT-1G	Nephelinite	0.03	10.5	13.4	12.2	1.27	14.2	0.24	3.85	38.9	0.70363	±0.00017	0.703509	±0.000019	0.017
USGS TB-1G	Basalt	0.11	17.12	6.7	8.67	4.52	3.51	0.18	3.56	54.29	0.70576	±0.00030	0.705580	±0.000023	0.026
USGS GSD-1G	Basalt	0.55	13.4	7.2	13.3	3	3.6		3.6	53.2	0.70924	±0.00029	0.709416	±0.000050	-0.025
USGS BCR-2G	Basalt	0.14	13.4	7.06	12.4	1.74	3.56	0.19	3.23	54.4	0.70486	±0.00038	0.705003	±0.000004	-0.020
MPI-DING T1-G	Diorite	0.28	17.1	7.1	6.44	1.96	3.75	0.127	3.13	58.6	0.70990	±0.00035	0.710093	±0.000017	-0.027
MPI-DING ATHO-G	Rhyolite	0.67	12.2	1.7	3.27	2.64	0.103	0.106	3.75	75.6	0.70310	±0.00026	0.703271	±0.000015	-0.024

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Conclusions

LA-ICP-QQQ with wet plasma conditions can be used for the direct determination of the $^{87}\text{Sr}/^{86}\text{Sr}$ isotope ratio in geological samples. The isobaric interference from ^{87}Rb on ^{87}Sr was overcome using MS/MS mass-shift mode with CH $_3\text{F}/\text{He}$ cell gas. The Sr+ ions react in the CRC to form SrF+ reaction product ions, while Rb+ ions do not react. $^{87}\text{Sr}/^{86}\text{Sr}$ ratios were accurately determined in seven reference materials, regardless of the matrix composition, Sr concentration, and Rb/Sr elemental ratio.

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Resolution of ¹⁷⁶Yb and ¹⁷⁶Lu interferences on ¹⁷⁶Hf to enable accurate ¹⁷⁶Hf/¹⁷⁷Hf isotope ratio analysis using ICP-QQQ with MS/MS

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Keywords

Hafnium, Hf, geology, dating studies, rock dating, isotopic abundance

Introduction

Hafnium ¹⁷⁶Hf to ¹⁷⁷Hf isotope ratio analysis can provide insight into the different geological events and processes that a mineral underwent during its formation/ metamorphosis; ¹⁷⁶Hf/¹⁷⁷Hf ratios are also used for geochronology dating studies. Isotope geochronology is a dating technique in which the age of a rock or mineral is derived from differences in the abundance of two isotopes of an element. Changes in isotopic abundance may be caused by isotopic (mass) fractionation, or by radioactive decay; in each case, the ratio acts as a geological clock, allowing the time that the mineral was formed to be estimated. Hf has lower mobility than lead (Pb) in metamict minerals such as zircon, xenotime, euxenite etc., so Hf isotope ratios can offer an alternative to Pb/Pb or Pb/U ratios for dating these minerals.

Of the Hf isotopes of interest, ¹⁷⁷Hf is free from direct isobaric overlap from any other element and does not typically suffer from polyatomic interference from other co-existing elements. However, the second Hf isotope used in the isotope ratio calculation, ¹⁷⁶Hf, suffers isobaric overlap from ¹⁷⁶Lu and ¹⁷⁶Yb, as shown in Figure 1. In order to obtain accurate Hf ratios, it is therefore necessary to separate the ¹⁷⁶Hf signal from the overlapping Lu and Yb signals.

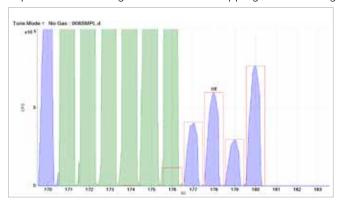


Figure 1. Hf (red peak template) in the presence of Lu and Yb matrix. The poor template fit for 176 Hf (highlighted in blue outline) is due to the contribution from 176 Lu and 176 Yb to the signal at m/z 176.

The mass resolution required to separate 176 Hf from the Lu and Yb isobaric interferences (M/ Δ M of \sim 140,000 for 176 Lu and >150,000 for 176 Yb) is far beyond the capability of commercial High Resolution Sector Field ICP-MS (SF-ICP-MS), so sample preparation (chemical separation) is required prior to analysis. In cases where chemical separation cannot be performed, for example in-situ measurement by Laser Ablation (LA), the Lu/Yb overlaps mean that accurate Hf isotope ratio analysis is not possible, or must rely on mathematical corrections (and the errors they can introduce).

An alternative direct approach is "chemical" resolution within a collision/reaction cell (CRC), using specific gas phase ion-molecule reaction(s) that will either:

- React with the interfering ion to neutralize it or move it to a new mass.
- React with the analyte to create a new product ion at a different, noninterfered mass.

In this study, the second approach, known as "mass-shift", was used. Hf reacts efficiently with ammonia cell gas to form Hf-ammonia cluster ions, while Lu and Yb are relatively unreactive. However, ammonia will react with the other Hf isotopes and other co-existing ions present in a typical sample matrix. These other ions also form ammonia-adduct ions, creating new interferences that vary depending on the matrix composition. These ammonia-adduct ions would interfere with the original Hf isotope pattern, making Hf isotope analysis unreliable, so control over the reaction process is essential.

The solution to this problem is to use a tandem mass spectrometer, which has an additional mass filter before the CRC. This extra mass filter prevents all ions apart from the target mass from entering the CRC, so the reaction chemistry is precisely controlled and unwanted side-reactions are avoided. This double mass filter approach is only possible with a tandem MS (or MS/MS) configuration, which provides unprecedented control of the ion/molecule reaction chemistry used in CRC-ICP-MS methods.

The Agilent 8800 and 8900 Triple Quadrupole ICP-MS (ICP-QQQ) instruments have an additional quadrupole mass filter (Q1), positioned in front of the CRC, with the capability of operating at unit mass resolution (MS/MS mode). In MS/MS operation, only a single mass-to-charge ratio (m/z) is transmitted through Q1, so the other Hf isotopes and any co-existing elements are rejected before they can enter the CRC. Unwanted side-reactions and potentially overlapping product ions are therefore eliminated. This method was used to measure Hf isotope ratios in a variety of samples containing Lu, Yb and mixed rare earth elements (REE). For this proof of concept, all work was performed using solution sample introduction, which allowed a greater flexibility to test interference removal. However, the same cell gas and MS/MS method can also be applied successfully to sample analysis using laser ablation (LA-ICP-QQQ).

Experimental

Instrumentation

The Agilent 8800* ICP-QQQ was configured with an SPS 4 autosampler and the standard sample introduction system consisting of a Micromist nebulizer (free aspiration), quartz spray chamber & torch, and Ni interface cones. Table 1 shows the key instrument parameters used for the analysis.

Table 1. Instrument parameters

Parameter	Value
RF power	1550 W
Sampling depth	7.0 mm
Nebulizer gas	1.15 L/min
Spray chamber temp	2 °C
Ammonia (10% in He) cell gas	22% of full scale (~2.2 mL/min)
Octopole bias	-6.0 V
Energy discrimination	-8.0 V

Samples and sample preparation

Due to the reactivity of ammonia, its use as a cell gas leads to a complex population of product ions, even in a simple sample matrix. However, selection of the most appropriate adduct ion is relatively simple with ICP-QQQ, by performing a Product Ion Scan. Unique to the MS/MS mode of operation, a Product Ion Scan uses a fixed mass setting for Q1, combined with a Q2 scan across the selected mass range. To identify useful ^{176}Hf - ammonia product ions, Q1 was fixed to mass (*m/z*) 176 amu, and Q2 was scanned across the mass range from *m/z* 170 to *m/z* 260, while aspirating a solution of 5 µg/L Hf. The resulting mass spectrum can be seen in Figure 2. Initially, the reaction product ion spectra may appear complex, but it should be noted that the use of a fixed mass setting for Q1 means that all these ammonia product ions are derived from the ^{176}Hf isotope. The most abundant ammonia adduct ion was Hf(NH)(NH₂)(NH₃)₃+, which occurs at M + 82 amu (*m/z* 258 for the ^{176}Hf isotope); this adduct was selected as the preferred mass transition.

It should be noted that the Hf adduct ion used is sensitive to CRC conditions, particularly the acceleration voltage applied from the Octopole Bias. This parameter was optimized to a lower value than is typically used, in order to favor the preferred transition and maximize the yield of the desired product ion. The cell gas flow rate was then re-optimized using the ICP-MS MassHunter autotune routines to further improve the product ion signal.

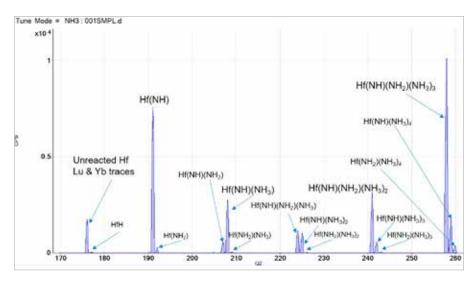


Figure 2. Product Ion Scan from m/z 170 to m/z 260 showing ammonia adduct reaction product ions formed from 176 Hf precursor ion (Q1 set to m/z 176)

A graphical representation of the 176 Hf transition can be seen in Figure 3; this schematic illustrates how Q1 (set to m/z 176) eliminates all ions apart from those at m/z 176, and Q2 (set to m/z 258) eliminates the unreacted 176 Lu and 176 Yb isotopes. The same mass transition is used for the other isotope of interest, 177 Hf, using Q1 and Q2 settings of m/z 177 and m/z 259, respectively.

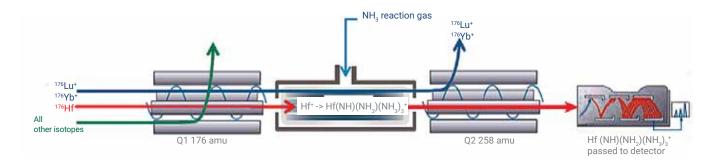


Figure 3. Schematic of the 176 Hf isotope reaction with ammonia cell gas in MS/MS mode.

To check that the Hf isotope pattern was maintained, a Neutral Gain Scan was performed where both Q1 and Q2 were scanned together, with a fixed mass difference of +82 amu applied to Q2. Figure 4 shows the resultant spectrum, confirming that the Hf-ammonia product ions match the overlay of the theoretical Hf isotopic abundances. Note that the higher mass Hf isotopes (178/179/180Hf) are not of interest in the isotopic analysis and were not measured. This spectrum demonstrates the unique benefit of MS/MS mode, which ensures that interisotope overlaps cannot occur, as only one Hf isotope mass is present in the cell at any given time.

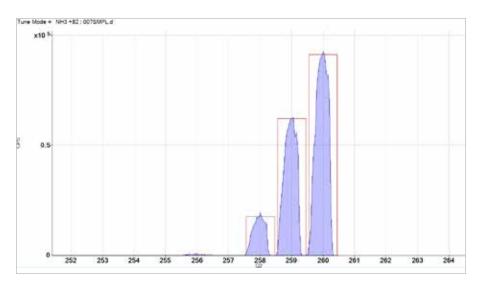


Figure 4. Neutral Gain Scan of the Hf isotopes as $Hf(NH)(NH_2)(NH_3)_3^*$ clusters; theoretical Hf isotopic abundances are shown in red, confirming that the isotope ratios are preserved in the product ion spectrum

To simulate real-world sample analysis, several potential sources of interference were introduced to assess whether bias or new interferences were created.

The test solutions included:

- Hf standard (5 ppb) also used for Mass Bias Calibration
- 100 ppb Yb & 5 ppb Hf
- 100 ppb Lu & 5 ppb Hf
- 100 ppb Yb + Lu & 5 ppb Hf
- Mixed 100 ppb "REE1" standard and 5 ppb Hf
- Mineral² sample with 100 ppb "REE1" and 5 ppb Hf
- 1. Agilent Standard 8500-6944 containing La, Ce, Pr, Nd, Sm, Eu, Gd, Tb, Dy, Ho, Er, Tm, Yb, Lu, plus Sc, Y, Th.
- 2. Natural mineral sample containing approximately 500 ppm Ca, 120 ppm Mg, 15 ppm Na, 5 ppm K, 1500 ppm SO,

Hf isotope ratio measurement

In order to provide comparative performance data, the ICP-QQQ was set to measure ^{176/177}Hf isotope ratios in three separate acquisition modes:

- No cell gas, "Single Quad" mode
 "Base" ICP-MS data not utilizing any mechanism to reduce isobaric overlaps
- $-\ \ {\rm NH_3}$ reaction gas, Single Quad Bandpass mode Non-MS/MS operation, allowing a limited mass range "window" into the CRC
- NH_3 reaction gas, MS/MS mode Q1 operating as a mass filter with unit mass resolution, allowing only a single m/z into the CRC

Table 2 displays the Hf isotope ratio (IR) data for each of the test solutions in each instrument mode. It can be seen that there was a large positive deviation from the expected ratio (i.e. the ¹⁷⁶Hf signal was high relative to its theoretical abundance) in both of the Single Quad modes of operation (no gas mode and

ammonia mode with bandpass filtering). This indicates that "Single Quad" operation did not resolve the Yb and Lu isobars at m/z 176, or stop the formation of new reaction product ion interferences.

By contrast, MS/MS mode with $\mathrm{NH_3}$ cell gas gave consistent, accurate Hf IR data in all the sample matrices.

To visualize and further investigate the potential overlaps that could have caused the poor Hf isotope ratio performance in Single Quad mode, a mass scan of the mineral sample was performed using Single Quadrupole Bandpass mode with $\mathrm{NH_3}$ reaction gas. The spectrum can be observed in Figure 5. The measured Hf isotopic pattern (far right of the spectrum) does not match the theoretical abundance template, showing that the Hf isotopes suffer overlap from new cell-formed cluster ions, due to the lack of control over the reaction processes. In a complex sample matrix, numerous cell-formed interferences are created, precluding the accurate analysis of many target product ions.

Table 2. ^{176/177}Hf isotope ratio (IR) data measured in samples containing various sources of interferences, using three different ICP-QQQ operating modes. The "deviation" is the error in the measured ratio relative to the true ratio of 0.282796.

	No gas Si	ngle Quad	NH3 Single Q	uad bandpass	NH3	MS/MS
Sample	IR	Deviation	IR	Deviation	IR	Deviation
Hf 5 ppb	0.27981	0.989	0.28252	0.999	0.28196	0.997
Hf 5 ppb, Yb 100 ppb	15.25251	53.935	0.30461	1.077	0.28370	1.003
Hf 5 ppb, Lu 100 ppb	3.18739	11.271	1.06062	3.750	0.28051	0.992
Hf 5 ppb, Yb, Lu 100 ppb	18.51262	65.463	1.06267	3.758	0.28099	0.994
Hf 5 ppb, REE mix 100 ppb	15.26995	53.996	0.64603	2.284	0.28139	0.995
Hf 5 ppb, Mineral REE mix 100 ppb	16.16150	57.149	0.63479	2.245	0.28230	0.998

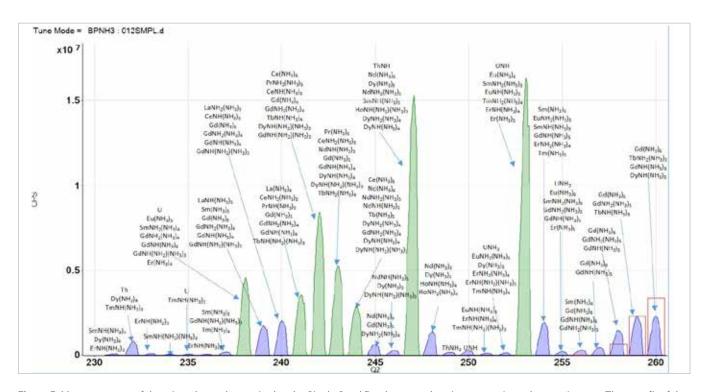


Figure 5. Mass spectrum of the mineral sample acquired under Single Quad Bandpass mode using ammonia as the reaction gas. The poor fit of the measured Hf isotopic pattern (far right) illustrates the interferences that occur in Single Quad mode. Some examples of cell-formed ammonia cluster ions are shown.

Many matrix elements and other analytes can react with ammonia to produce higher order reaction products, so MS/MS mode is essential to remove these precursor ions before they enter the cell and form new interferences.

The 8800 and 8900 ICP-QQQ use an additional quadrupole mass filter, operating at unit mass resolution and positioned before the CRC, to control which ions enter the reaction cell. This ensures unprecedented levels of control over the reaction processes that occur within the cell. MS/MS mode can quickly switch between on-mass measurement and off-mass measurement within a single acquisition, supporting multi-element analysis in each gas mode. Figure 6 shows the Hf isotopes measured using off-mass mode (Q2=Q1 + 82 amu) and the other masses measured on-mass (Q1=Q2). The small, residual peaks for unreacted Th and U can be seen, along with ThO and UO. Most of the Th and U would have reacted with ammonia cell gas, forming adduct species that are not measured in MS/MS on-mass mode. Any undesired side reactions are eliminated before they can proceed, so the underlying analyte isotope ratios are preserved in the product ion spectrum.

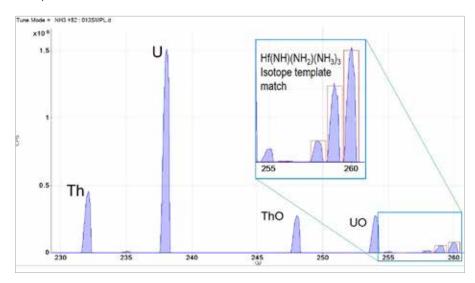


Figure 6. Mass spectrum of the mineral sample measured using $\mathrm{NH_3}$ mass-shift mode for Hf and on-mass mode for all other isotopes. The Hf isotopic pattern illustrates that all of the interferences that occurred in Single Quad mode (shown in Figure 5) have been resolved.

Conclusions

The MS/MS capabilities of Agilent's ICP-QQQ measured Hf isotope ratios with excellent accuracy—even in samples containing high levels of co-existing and potentially interfering matrix elements.

The isobaric overlaps from 176 Lu and 176 Yb on 176 Hf were eliminated using NH $_3$ as the reaction gas. The reaction chemistry was controlled in the cell by operating the first quadrupole mass filter at unit mass resolution set to $\it m/z$ 176. This excluded all ions apart from those at $\it m/z$ 176 (176 Lu, 176 Yb and 176 Hf). Since only Hf reacts readily with NH $_3$, 176 Hf was free to be measured via its most appropriate cluster ion at $\it m/z$ 258, effectively avoiding the isobaric overlaps from Lu and Yb. Together with the corresponding ammonia cluster ion formed from the 177 Hf isotope, this method allowed accurate Hf isotope analysis to be performed in a range of complex synthetic sample matrices.

In summary:

Chemical resolution using a reaction gas offers a powerful alternative to mass resolution, allowing access to isobars beyond the maximum resolution available with commercial High Resolution SF-ICP-MS.

Control over the reaction processes is essential to avoid new, unexpected interferences forming from the sample matrix and other coexisting elements and isotopes.

MS/MS technology affords unprecedented control over the reaction processes, greatly simplifying methodology regardless of the process or sample matrix.

Crucially, MS/MS operation allows access to higher order reaction product (cluster) ions, while still preserving the analyte's original isotopic information.

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Manganese Analysis in Whole Blood: Expanding the Analytical Capabilities of ICP-MS

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Keywords

manganese, whole blood, iron, abundance sensitivity, helium MS/MS

Introduction

Analysis of clinical samples is challenging due to their complex matrices. While ICP-MS is an immensely powerful multi-element analytical technique, it does suffer from some well-documented spectral interferences. Achieving low detection limits is limited by background signal from low level impurities and the presence of polyatomic interferences, which require the use of CRC technology for their removal. Although the use of CRC-ICP-MS has alleviated many of these analytical challenges, some spectral interferences remain problematic for quadrupole ICP-MS (ICP-QMS). One such interference is the signal overlap on ⁵⁵Mn due to peak tailing from both ⁵⁴Fe and ⁵⁶Fe. Whole blood contains an average of 500 ppm of Fe, and with the level of Mn in whole blood being roughly 10 ppb, analytical results for Mn tend to bias high due to the significant signal tailing and overlap from the adjacent Fe peaks. In this work, we use the superior abundance sensitivity of the 8800 ICP-QQQ to remove any signal overlap from Fe on Mn in whole blood.

Experimental

Instrumentation: Agilent 8800 #100.

Plasma conditions and ion lens tune: Preset plasma/General purpose with soft extraction tune: Extract 1 = 0 V.

Method: Samples were analyzed using the 8800 ICP-QQQ in both Single Quad (SQ) mode and MS/MS mode. In this study, the mass range of interest (from *m/z* 50 to 60) was scanned at twenty points per peak in both no gas and helium (He) modes. For the analysis of Mn in blood, MS/MS mode with on-mass measurement (Q2 set to the same mass as Q1) was used, with helium cell gas (typical flow of 4.3 mL/min) to remove polyatomic ion interferences such as FeH⁺ and ArOH⁺.

Sample preparation: A 5 ppb solution of Mn was prepared from a stock of 1000 ppm Mn and either analyzed separately or spiked into "base" whole blood (low level Mn). Whole blood was diluted using an alkali matrix containing ammonium hydroxide, EDTA, Triton X-100, and butanol.

Abundance sensitivity

The abundance sensitivity (AS) of a mass spectrometer is the contribution that the signal at mass M makes to the signals at the adjacent masses (M \pm 1), expressed as a ratio (M-1/M on the low-mass side and M+1/M on the high-mass side). Simply put, AS is the measure of the "peak tailing" to adjacent masses, which will contribute to a false positive signal, such as that seen on ⁵⁵Mn (present at trace levels) from the large contribution from ⁵⁴Fe and ⁵⁶Fe (which exists at very high concentration) in whole blood. The abundance sensitivity of the best quadrupole ICP-MS systems is of the order of 10^{-7} .

Abundance sensitivity study in SQ and MS/MS mode

SQ and MS/MS spectra for a 500 ppm Fe solution acquired in no gas mode are shown in Figure 1. The spectrum on the right illustrates the superior peak-to-peak resolution of the 8800 ICP-QQQ operated in MS/MS mode. Although no interference removal for polyatomic ions was employed, the elimination of the contribution to mass 55 from adjacent peaks is clearly evident in MS/MS mode. The "flat-top" peak shapes are the result of the logarithmic scale.

Abundance sensitivity plays an important role when samples contain a large concentration of Fe. Figure 1 looks at the contribution of "peak tailing" on ⁵⁵Mn due to high levels of Fe. The high concentration of Fe together with the ArN+ and ArO+ contribution in no gas mode resulted in the signals at 54 and 56 being over the range of the detector, and so they were automatically skipped. However, the signal contribution from ⁵⁶Fe on mass 55 is clearly visible in the SQ mode (indicated by the red box) while it is absent in the MS/MS mode.

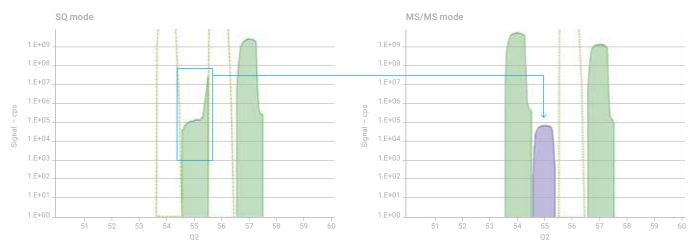


Figure 1. Comparison of no gas mode spectra for 500 ppm Fe solution, measured in SQ mode (left) and MS/MS mode (right). The signal colored blue was obtained in pulse counting while the green signal was obtained in analog mode. The dotted lines indicate over-range peaks (automatically skipped to protect the EM detector).

Figure 2 shows three spectra obtained in MS/MS mode with He cell gas. When He cell mode is used for interference removal, precise and accurate analysis is easily achieved. In He MS/MS mode, all interferences (arising from signal overlap from tailing of adjacent peaks and polyatomic ions isobaric interferences) are removed, yielding unbiased analysis and accurate results.



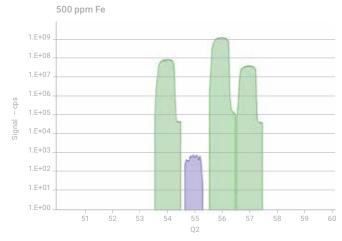




Figure 2. He MS/MS mode spectra: (top to bottm) 5 ppb Mn, 500 ppm Fe, and 5 ppb Mn + 500 ppm Fe.

Figure 3 is an overlay of three spectra measured using He MS/MS mode; 1) Blank, 2) 10x whole blood, and 3) 500 ppt Mn spike in 10x whole blood. Table 1 summarizes the results of 10x diluted whole blood analysis and 500 ppt Mn spike recovery test. As shown, very low blank levels were achieved, ⁵⁵Mn was clearly resolved in the spectrum and good spike recoveries were obtained.

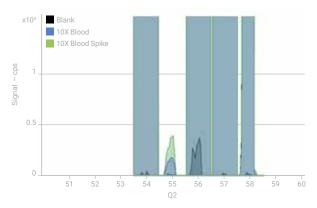


Figure 3. Spectra of three samples: blank, 10x diluted blood and 10x diluted blood spiked with 500 ppt Mn.

Table 1. 10x diluted whole blood analysis results for Mn.

	Blood sample	Blood sample + 500 ppt Mn	Spike recovery
	μg/L		%
Sample A	0.413	0.983	114
Sample B	0.432	0.924	98

Conclusions

Quadrupole ICP-MS has been almost universally accepted for low level analysis of trace analytes in complex matrices. However, many challenging interferences remain unresolved, especially when trace analytes must be measured close to matrix element peaks in complex samples. The Agilent 8800 ICP-QQQ with MS/MS capability has abundance sensitivity better than 10⁻¹⁰, which enables the analysis of trace analytes (such as Mn) in the presence of a high concentration of adjacent elements (such as Fe).

For Research Use Only. Not for use in diagnostic procedures.

Measurement of Titanium in Clinical Samples: Possible Application to Monitoring Patients with Joint Replacements

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Keywords

titanium, biological, serum, urine, joint-replacement, Seronorm, ammonia mass-shift

Introduction

Although titanium (Ti) has little or no direct biological role, it is widely used in dental, artificial/replacement joints and surgical reconstruction applications due to its high strength, light weight and the fact it is biocompatible. It is also used extensively as a pigment and abrasive polishing agent (as TiO₂) and is often found in foods and toothpaste due to its inert nature (this form is usually passed unaltered in faecal matter and is not normally transported or expressed through body fluids).

More recent applications with certain types of metal-on-metal (rather than ceramic or polymer based) joint replacements can lead to the release of wear metal particles or ions within the body of the patient. These can become highly concentrated in the synovial fluid (lubricating fluid of the joint), pass into the bloodstream and be expressed through urine. Unusually "high" concentrations of Ti can indicate a premature failure of a Ti-based joint and such failure can lead to infection or constant pain for the patient. It is therefore important to reliably determine the concentration of Ti within biological fluids at normal endogenous levels in order to obtain a basal concentration. An increase from this concentration could indicate an imminent failure of the joint.

Experimental

The determination of Ti in biological matrices is challenging for conventional ICP-MS, due to its low natural concentration and the presence of spectral interferences on all the Ti isotopes e.g. sulfur (as SO), P (as PO) and Ca. It is possible to use reaction chemistry with NH $_3$ cell gas in the CRC of a quadrupole ICP-MS (ICP-QMS), to mass-shift the Ti $^+$ to a higher mass product ion, leaving the interfering species behind. However, the use of highly reactive cell gases in ICP-QMS is prone to severe errors, as there is no way to control the ions that enter the CRC. This means that the reaction chemistry and the product ions created can change dramatically, with even slight differences in sample matrix or co-existing analyte concentrations. For this application, the 8800 ICP-QQQ was used to provide controlled reaction chemistry with ammonia as the reaction gas and measuring Ti as the TiNH $_2$ (NH $_3$) $_4$ $_1$ cluster ion at the M + 84 amu transition.

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/General purpose.

Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -170 V.

CRC conditions: NH₂ gas (10% in He) at 1.7 mL/min, Octopole bias = -8 V,

KED = -8 V.

Samples and sample preparation: Certified reference materials of human serum and urine were purchased from Seronorm (Norway). They were prepared in duplicate by 10x dilution into a basic diluent consisting of NH₄OH (0.5%), H₄-EDTA (0.01%), BuOH (2%) & Triton X-100 (0.01%) in ultrapure water. No further matrix matching was applied for the standards.

Selection of product ion for Ti measurement

In order to select the most appropriate Ti cluster ions in NH_3 mode, a product ion scan was performed for the ⁴⁸Ti isotope by introducing a 10 ppb Ti solution (Figure 1). Q1 was set to m/z 48, allowing only ions at the mass of the target precursor ion to enter the cell; Q2 was scanned over a selected mass range to measure all the product ions formed in the cell by NH_3 reactions with ⁴⁸Ti. Based upon this scan, the two most abundant cluster ions (Q1 + 84 amu [TiNH₂(NH₃)₄] and Q1 + 102 amu [Ti(NH₃)₆]) were selected for further study. For each of the two reaction transitions identified above, neutral gain scans (where Q1 and Q2 are scanned synchronously, with a set mass difference between them (Q2 = Q1 + 84 and Q2 = Q1 + 102 in this case)) were performed. These scans are shown in Figure 2 confirming the correct natural isotopic abundances for the different Ti isotopes. Without MS/MS capability, it would be impossible to preserve the isotopic information for this element due to the relatively complex nature of the Ti-ammonia adducts. The instrument cell conditions were optimized using simple HNO₃ acidified Ti standards and applied to the analysis of the CRMs.

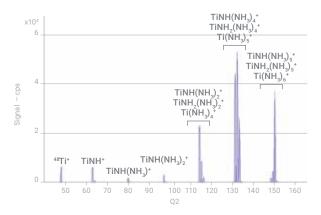


Figure 1. Product ion scan for ⁴⁸Ti+ in NH₃ mode.

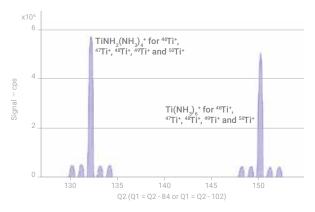


Figure 2. Neutral gain scan for two Ti \Box Ti cluster ion transitions: For TiNH₂(NH₃)₄+ cluster ions, Q2 = Q1+ 84 amu, and for Ti(NH₃)₆+ cluster ions, Q2 = Q1+102 amu. The preservation of the natural Ti isotope abundance pattern (⁴⁶Ti+, ⁴⁷Ti+, ⁴⁸Ti+, ⁴⁹Ti+ and ⁵⁰Ti+) can be seen, confirming that MS/MS mode provides complete control over the complex Ti-NH₂ reaction chemistry.

Table 1 displays the results for both serum and urine sample types measured against a single calibration. The 8800 ICP-QQQ was operated also under no gas and He mode to provide comparative data, and three Ti isotopes were monitored for the same cluster ion transition, to give confirmation of the results.

Table 1. Urine and serum sample recovery (μg/L) for Ti in Seronorm CRM using TiNH₂(NH₂)₄ cluster.

Sample Name	Target	⁴⁷ Ti [No gas]	⁴⁷ Ti [He]	47 -> 131 Ti [NH ₃]	48 -> 132 Ti [NH ₃]	49 -> 133 Ti [NH ₃]
Urine blank	4.6 (2.2-7.0)	1989.79	41.44	2.80	2.79	2.92
Urine blank	4.6 (2.2-7.0)	2004.91	44.30	3.50	2.93	3.33
Urine trace elements		1789.92	51.41	14.81	15.27	14.42
Urine trace elements		1749.13	52.58	14.99	15.49	15.50
Serum L1	1.28 (0.86-1.80)	144.18	3.79	1.21	1.15	1.14
Serum L1	1.28 (0.86-1.80)	128.97	2.95	1.27	1.18	1.09
Serum L2		100.16	3.95	1.76	1.92	1.61
Serum L2		95.65	3.02	1.82	1.64	1.76

Conclusions

Titanium was only certified in two of the four materials measured but the 8800 ICP-QQQ data were all comfortably within the measured ranges when operating under ammonia MS/MS mode, in contrast to no gas and He mode data. Importantly, the three Ti isotopes measured under ammonia MS/MS mode all gave equivalent data; this could indicate applicability of the method to the use of isotope-based analysis such as isotope dilution (ID) or isotope tracer analysis. The use of ammonia combined with MS/MS greatly simplifies the analysis of Ti in biological media for several isotopes. Furthermore, because MS/MS mode provides control over the reaction chemistry, no special attention needs to be paid to specific matrix matching regardless of the fluid investigated.

Simultaneous Quantitation of Peptides and Phosphopeptides by capLC-ICP-QQQ

Authors

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Keywords

proteins, peptides, phosphorus, phosphopeptides, sulfur, S-containing peptides, heteroatom, isotope ratio, pharmaceutical, clinical, drugs, metabolites, environmental, pesticides, nanotechnology, nanoparticles, oxygen mass-shift

Introduction

LC-MS/MS is used for the quantification of target proteins in pharma/biopharma and clinical research. The approach generally relies on the use of synthetic, isotopically-labeled forms of each target protein and peptide, which are used as internal standards for the specific quantitation of the corresponding target compound. In contrast, the high temperature plasma ionization source used in ICP-MS ensures that elemental response is practically independent of the original form of the compound, which enables non-species-specific (or compoundindependent) quantitation of compounds by measuring the signal for an element contained in the target compound. In this way, different proteins and peptides containing the heteroatoms S and P can be quantified using a single S- or P-containing compound as a generic standard. Unfortunately, with conventional quadrupole ICP-MS, the DLs for P and S are compromised by their high ionization potential and by multiple polyatomic interferences. The Agilent 8800 ICP-QQQ can effectively remove those interferences using reaction cell chemistry combined with the unique MS/MS mode, achieving excellent DLs for P and S even in organic solvents. This paper demonstrates the advantage of ICP-QQQ for the determination of proteins and peptides by measurement of P and S heteroatoms.

Experimental

Instrumentation: An Agilent 8800 Triple Quadrupole ICP-MS was used with an Agilent 1260 Series low flow capillary LC system. The standard 2.5 mm internal diameter (id) injector torch was replaced with the narrow injector, 1.5 mm id torch (G3280-80080) used for the analysis of volatile organic solvents. The exit of the LC column was interfaced to the ICP-MS via an Agilent capillary LC interface kit (G3680A) featuring a total consumption nebulizer and micro-volume spray chamber. $\rm O_2$ gas (20% $\rm O_2$ in Ar) was supplied to the plasma as an option gas at 0.08 L/min to prevent carbon build-up on the interface cones. Agilent ICP-MS MassHunter chromatographic software was used for integrated control of the LC-ICP-MS system and for data analysis.

CRC conditions: O_2 cell gas flow rate at 0.35 mL/min, Octopole bias = -18 V and KED = -6 V.

Acquisition conditions: MS/MS $\rm O_2$ mass-shift method was applied for P and S measurement as shown in Figure 1.

LC conditions: An Agilent Zorbax SB C18 (5 μ m, 150 x 0.3 mm) reverse phase column was used with a flow rate of 5 μ L/min. Mobile phases of water (A) and acetonitrile (B) were used for a gradient elution with the following profile: 0-3 min: 1% B; 3-35 min: 1-60% B linear. Both mobile phases contained 0.1% formic acid and 10 ppb Ge as ISTD and for tuning. The injection volume was 2 μ L.

Reagents: Bis-4-nitro-phenyl phosphate (BNPP, 99% purity) and methionine (≥ 99% purity) (Sigma- Aldrich, Steinheim, Germany) were used as calibration standards for phosphopeptides and S-containing peptides respectively. Amino acid sequences of the phosphopeptides were LRRA-pS-LG and KRS-pY-EEHIP, and the S-containing peptides were A-C-TPER-M-AE and VP-M-LK. All peptides were purchased from AnaSpec (Fremont, CA, USA) with purity ≥95%.

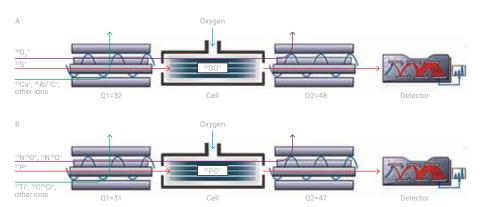


Figure 1. 8800 ICP-QQQ MS/MS operation in mass-shift mode to remove interferences on S (a) and P (b).

Results and Discussion

Calibration and DL

Calibration standards containing 25, 50, 100 and 200 ng/mL of both P and S (BNPP and methionine, respectively) were injected and measured. Excellent linearity and RSD of <4% was obtained (see Figure 2).

The chromatogram for the 50 ng/mL standard was used for signal to noise (S/N) and DL calculation. The DL achieved was 0.10 ng/mL for P and 0.18 ng/mL for S. As the injection volume was 2 μ L, the DLs in absolute weight were calculated to be 6.6 fmol and 11 fmol for P and S, respectively.

Measurement of phosphopeptide and S-containing peptides

Finally, a sample containing a mixture of phosphopeptides and S-containing peptides was analyzed. The sample was also spiked with the standards methionine and BNPP for non-species-specific calibration. The chromatogram shown in Figure 3 illustrates the excellent peak shape and S/N obtained, demonstrating the exciting potential of ICP-QQQ for quantitative protein and peptide analysis using measurement of P- and S-heteroatoms.

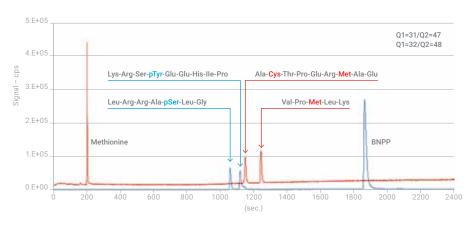


Figure 2. Chromatogram of phosphopeptides and S-containing peptides. Sample: 45 ng/mL of two phosphopeptides and two S-containing peptides, and 105 ng/mL of BNPP and methionine (conc. as P or S).

More Information

Simultaneous quantitation of peptides and phosphopeptides by capLC-ICP-MS using the Agilent 8800 Triple Quadrupole ICP-MS, Agilent publication, $\underline{5991}$ - $\underline{1461EN}$.

Analysis of Selenoproteins in Rat Serum using HPLC-ICP-QQQ

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Keywords

selenium, selenoprotein P, Sel P, glutathione peroxidase, eGPx, GPx-3, serum, biological, rat, mouse, hamster, guinea pig, speciation, mass-shift method, oxygen reaction mode

Introduction

Selenium (Se) is an essential micronutrient in animals and is present in several of the key proteins found in plasma. Two selenoproteins which contain Se as selenocysteine (SeCys) in their primary structures, extracellular glutathione peroxidase (eGPx, GPx-3) and selenoprotein P (Sel P), have been detected in animal plasma. Other Se-containing proteins which have Se incorporated into their peptide sequence as selenomethionine (SeMet), are also detected because animals are unable to discriminate SeMet from methionine (Met). The most abundant Se-containing protein in human plasma is albumin. However, some studies have indicated that no or little Se-containing albumin is detected in the blood plasma of experimental animals compared to human plasma [1-3]. This can be explained by the fact that humans ingest Se mainly as SeMet, whereas the major Se species in the feeds given to experimental animals is inorganic Se, such as selenite and selenate.

The three most abundant Se isotopes, 80 Se (49.6%), 78 Se (23.8%), and 76 Se (9.36%), suffer from interference by several polyatomic ions originating from the Ar plasma, namely, 40 Ar 40 Ar $^{+}$, 40 Ar 38 Ar $^{+}$, and 38 Ar 38 Ar $^{+}$, respectively. 77 Se is also subject to interference by 40 Ar 37 Cl $^{+}$ when chloride is present in the sample matrix, as is the case with biological samples. Sample matrix components such as S, Ca and K may also contribute polyatomic overlaps on isotopes of Se, for example 39 K 37 Cl $^{+}$ on 76 Se $^{+}$, 32 S $_{2}$ 16 O $^{+}$, 32 S $_{1}$ 60 O $_{3}$, and 40 Ca $_{2}$ $^{+}$ on 80 Se $^{+}$, and $^{79/81}$ BrH $^{+}$ on $^{80/82}$ Se $^{+}$.

ICP-QQQ can operate with oxygen cell gas and mass-shift mode, using 0-atom addition to move the analyte ions away from the interference for detection at M+16 amu. For example, $^{78}\mathrm{Se^+}$ is measured as $^{78}\mathrm{Se^{16}0^+}$ at 94 amu; $^{80}\mathrm{Se^+}$ is measured at 96 amu; and $^{82}\mathrm{Se^+}$ is measured at 98 amu. The aim of this study is to evaluate the performance of ICP-QQQ for the speciation of Se in rat serum.

Experimental

Instrumentation: Agilent 8800 #100 was used with an HPLC system.

CRC conditions: O₂ cell gas at a flow rate of 0.30 mL/min.

Acquisition conditions: MS/MS $\rm O_2$ mass-shift method: Se signals were monitored as SeO+ at m/z 94, 96, and 98

LC conditions: A multi-mode gel filtration column, Shodex Asahipak GS-520HQ (7.5 i.d. x 300 mm, with a guard column, 7.5 i.d. x 75 mm, Showa Denko, Tokyo, Japan), was used. A 200 μ L aliquot of serum sample was injected onto the column and then eluted with 50 mmol/L Tris-HCl, pH 7.4, at a flow rate of 0.6 mL/min. The eluate emerging from the column was introduced directly into the nebulizer of the ICP-QQQ.

Reagents: The instrument was tuned using an inorganic Se standard. Tris(hydroxymethyl) aminomethane (TRIZMA base and TRIZMA HCl) were purchased from Sigma (St. Louis, MO, USA).

Results and Discussion

Elution profiles of Se in rat serum

Blood was collected from the experimental rats after one week; the blood was separated by centrifugation, and the serum samples were stored at -30 °C prior to analysis by LC-ICP-QQQ. Two well-separated Se peaks were detected at retention times of 11.7 and 14.3 min (Figure 1). The former and latter peaks were assignable to eGPx and Sel P, respectively, per a previous study [4]. It was reported that albumin was eluted at the retention time of 15.0-16.0 min on this column [5]. However, we did not detect a Se peak at a retention time of 15.0-16.0 min, suggesting that SeMet was not incorporated into albumin in place of Met.

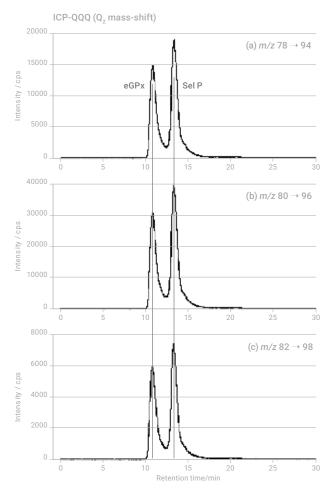


Figure 1. Elution profiles of Se in rat serum. A 200- μ L aliquot of a rat serum sample was injected into a GS-520HQ column and the eluate was monitored by ICP-QQQ (a-c) at m/z 94 (a), 96 (b), and 98 (c).

Conclusions

Two major selenoproteins, eGPx and Sel P, in rat serum were well separated on an HPLC column. ICP-QQQ was a more accurate detector for the speciation of serum selenoproteins than conventional quadrupole ICP-MS, because the ICP-QQQ analysis was completely free of interferences originating from the Ar plasma source and any matrix elements.

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Absolute Quantification of Intact Proteins without Specific Standards by capLC-ICP-QQQ

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Keywords

proteins, sulfur, isotope dilution analysis, absolute quantification, proteomics, venomics

Introduction

ICP-MS has become a feasible alternative to molecular MS-based proteomic workflows, which commonly require specific (mostly isotopically enriched) standards for each target protein. The introduction of triple quadrupole ICP-MS (ICP-QQQ) has boosted the applicability of ICP-MS-based methods for quantitative proteomic studies. This development is due to the sensitive interference-free detection of heteroelements, such as sulfur (S), and the possibility of generic quantification without the need for specific standards. This paper demonstrates the capacity of capillary reverse phase (RP)-LC-ICP-QQQ for the absolute quantification of intact protein standards using post-column isotope dilution (ID) analysis. The method was used for the individual quantification of up to 27 different proteins present in a relatively complex sample of snake venom.

Experimental

Instrumentation: Capillary LC separation was performed using an Agilent 1200 series HPLC equipped with a BIOshell A400 (3.4 μ m, 150 mm x 0.3 mm) reverse-phase column (Sigma-Aldrich, Germany). The exit of the column was coupled to the Agilent 8800 ICP-QQQ using an Agilent Capillary LC interface kit (G3680A). The kit comprised a total consumption nebulizer with single pass spray chamber. Enriched ³⁴S was added post-column. Sulfur isotopes were measured using an oxygen mass-shift MS/MS method and the S content of the proteins was determined by ID. BOC-L-methionine was used as the internal standard to correct for any injection errors.

Reagents: Pure standard BOC-L-methionine (Sigma, Germany) was used as a standard for quantification. Cytochrome C, bovine serum albumin (BSA), β -casein, transferrin (Sigma-Aldrich, Germany), and intact mAb (Waters, USA) were used as protein standards in the recovery study. The venom sample analyzed in the study was extracted from a Tanzanian *Naja mossambica* (Valence, France).

Results and Discussion

Protein standards analysis

Absolute quantification of proteins was achieved through the measurement of sulfur with post-column ID. The individual protein samples (BSA, cytochrome C, and transferrin) were spiked with a non-species-specific sulfur-containing standard, BOC-L-methionine, in this case. Quantitative (mass purity) data obtained for cytochrome C (94 \pm 2%), BSA (96 \pm 2%), and transferrin (94 \pm 2%) were in excellent agreement with the theoretical values supplied by the manufacturer, \geq 95%, \geq 98% and \geq 95%, respectively.

Quantification of 27 proteins in snake venom

The method was applied to the analysis and quantification of the proteins present in a snake venom sample. The sample was spiked with BOC-L-methionine. capLC-ICP-QQQ was then used to monitor the sulfur present in each protein and to quantify the concentration via ID analysis. Figure 1 shows the chromatogram (22-39 min) for the snake venom sample.

Parallel capLC-ESI-MS analysis was used to identify the proteins. By identifying the proteins, it was possible to know the S-to-protein stoichiometry of each peak. This information was then used to translate sulfur mass into individual protein quantities, in μ mol protein per gram of venom sample. The quantified results are summarized in Figure 2.

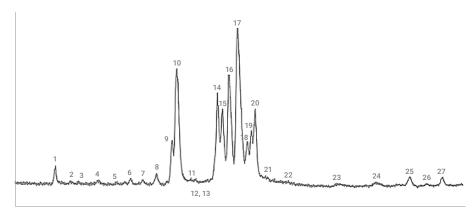


Figure 1. Chromatogram (22–39 min) of snake venom sample (S detection). Reprinted with permission from Anal. Chem., 2016, 88 (19), 9699–97. Copyright 2016 American Chemical Society.

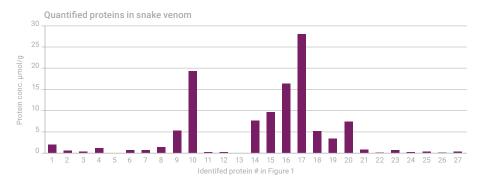


Figure 2. Quantified proteins in snake venom.

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Conclusions

The capLC-ICP-QQQ method is suitable for the absolute quantification of intact proteins without the need for specific standards. If quantitative chromatographic recoveries can be assured, it is even possible to quantify nonpure protein samples using this method. The potential of the methodology for the quantification of intact proteins present in relatively complex samples was demonstrated by analyzing 27 proteins in snake venom.

More Information

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Measurement of Selenium in Clinical Samples in the Presence of Gadolinium-Based Magnetic Resonance Imaging Contrasting Agents

Authors

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Keywords

selenium, enzyme, blood, serum, urine, MRI contrasting agents, gadolinium, molybdenum, zirconium, neutral gain scan, oxygen mass-shift

Introduction

Selenium is an important micronutrient for human, mammalian, bacterial, and plant life and is contained within several co-factors and enzyme systems. It is monitored in blood, serum and urine as part of human health, and a deficiency can indicate an illness (particularly if the levels change suddenly), such as cancer, diabetes and tuberculosis (TB).

For cancer patients, determining the location of the tumour often requires the use of Magnetic Resonance Imaging (MRI). However, for some soft tissues such as the brain, a "contrasting agent" is needed to effectively show the location of the tumour or problem area. There are several contrasting agents which are salts or chelates of gadolinium (III) (Gd(III)), trade names are given in brackets:

Gadodiamide (Omniscan), Gadobenate (MultiHance), Gadopentetate (Magnevist), Gadoteridol (ProHance), Gadofosveset (Ablavar, formerly Vasovist), Gadoversetamide (OptiMARK), Gadoxetate (Eovist), Gadobutrol (Gadavist)

Unfortunately Gd has a relatively low second ionization potential (12.09 eV) meaning it can form Gd $^{++}$ ions in the plasma. These Gd $^{++}$ ions appear at half their original mass (as a quadrupole measures ions based on their mass to charge ratio or m/z) and form interferences on all of the main analytical isotopes of Se. This is complicated to a greater extent as Gd has several odd-mass isotopes which form Gd $^{++}$ interferences at half-mass (e.g., 155 Gd $^{++}$ would appear at m/z 77.5). This makes the spectrum in the mass region of the Se isotopes quite complex when Gd is present in the sample. In a typical patient's sample, the Gd concentration can vary between zero to several thousand parts per billion (μ g/L). Because of the variability from patient-to-patient (which is also time-dependant on a sample-to-sample basis due to the contrasting agent's half-life in the body) a simple mathematical correction cannot always be made or a constant "background" be assumed.

Experimental

In order to remove the Gd-based interference, Se $^+$ can be reacted with oxygen cell gas in the collision/reaction cell to produce SeO $^+$ as a product ion. The Se-O reaction is slightly endothermic (Δ Hr = 0.71 eV) which means that the reaction yield for SeO $^+$ would be relatively low. However the bias voltage on the ORS can be adjusted to increase the ion energy improving reaction yield significantly over a more "thermalized" approach. These conditions are referred to as high ORS bias conditions.

Instrumentation: Agilent 8800 #100.

Plasma conditions: Preset plasma/General purpose.

Ion lens tune: Soft extraction tune: Extract 1 = 0 V, Extract 2 = -170 V. **CRC conditions:** O_2 gas at 0.3 mL/min, Octopole bias = -15 V, KED = -8 V.

Results and Discussion

O₂ mass-shift method

Using O_2 mass-shift, the analyte is measured at M +16 amu (e.g. 78 Se $^+$ is measured as 78 Se 16 O $^+$ at 94 amu). With conventional quadrupole ICP-MS, any 94 Mo or 94 Zr present in the sample would interfere with the measurement at this mass. However, with MS/MS mode, 94 Mo or 94 Zr are removed by Q1 as it is set to the mass of the Se $^+$ precursor ion at 78 amu, and 156 Gd $^{++}$ is eliminated as Q2 is set to the SeO $^+$ product ion mass of 94 amu. Even if Gd did form GdO $^{++}$ this would also be eliminated by Q2 as the apparent mass (m/z) of 156 Gd 16 O $^{++}$ is 172/2 (86 amu). Figure 1 is a graphical representation of the MS/MS setup.

To check for efficient conversion of Se $^+$ to SeO $^+$, a neutral gain scan covering the mass range of all the SeO $^+$ product ions was performed for a 5 ppb Se solution. Figure 2 displays the isotope pattern of the + 16 O-atom transitions for all the Se isotopes, showing a perfect match with the theoretical isotopic fit.

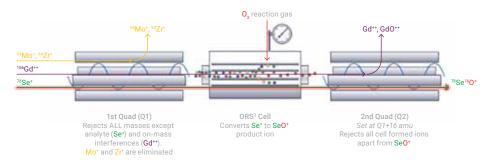


Figure 1. Representation of ICP-QQQ setup with Q1 set to 78 amu and Q2 set to 94 amu. Mo and Zr-based interferences are eliminated by Q1 and Gd^{++} is eliminated by Q2 allowing the measurement of ^{78}Se as $^{78}Se^{16}O^{+}$.

Se measurement in human serum

Instrument cell conditions were optimized using a Se standard in a simple HNO $_3$ matrix. A pooled human serum sample was prepared by 10x dilution into a basic diluent consisting of NH $_4$ OH (0.5%), H $_4$ -EDTA (0.01%), BuOH (2%) & Triton X-100 (0.01%) in ultrapure water. The sample was prepared unspiked and also spiked with Gd equivalent to 250, 500 and 1000 µg/L in the original sample, and analyzed using the 8800 ICP-QQQ in no gas and O $_2$ mass-shift modes of operation for comparison. The data is summarized in Table 1. The results show that, under no gas conditions, the apparent Se concentration is influenced by the variable Gd⁺⁺ interference. Recovery based upon the original unspiked sample demonstrates an over-recovery of almost 130% for the no gas data when Gd is at a concentration of 1000 µg/L. In contrast, the Se data measured with MS/MS mass-shift mode remains essentially constant at all levels of Gd matrix. This would indicate that the O $_2$ mass-shift reaction is independent of the Gd concentration and is highly applicable to this relatively difficult and important application.

Table 1. Serum sample data and recovery for Se with variable Gd concentration. Recovery is calculated based on determined Se concentration in unspiked serum sample. All data is dilution corrected.

	No gas mode		0, 1	mass-shift
	Conc. ppb Recovery %		Conc. ppb	Recovery %
Serum	93.64	NA	91.42	NA
Serum with 250 μg/L Gd	99.97	106.7	91.38	100.0
Serum with 500 μg/L Gd	112.1	120.0	91.70	100.3
Serum with 1000 μg/L Gd	121.1	129.3	91.78	100.4

Quantitative Analysis of Active Pharmaceutical Ingredients using Heteroatoms as Elemental Labels

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Keywords

protein, heteroatom, API, monoclonal antibody, mAb, sulfonamide, sulfur, zoledronic acid hydrate, phosphorus, clonidine hydrochloride, chlorine, mass-shift, oxygen reaction mode, hydrogen reaction mode

Introduction

Organic molecules and proteins can be detected and quantified indirectly using ICP-MS to measure a heteroatom "tag" element contained within the targeted compound. For example a large number of Active Pharmaceutical Ingredients (API) contain sulfur (S), phosphorus (P) or halogens. Unfortunately, S, P and the halogens have high first ionization potentials so they are poorly ionized in the ICP-MS plasma, leading to low sensitivity. S, P and chlorine (CI) are also difficult to measure by conventional quadrupole ICP-MS (ICP-QMS) due to intense spectral interferences. As a result, accurate analysis of S, P, and the halogens at the analytical ranges that are relevant to pharmaceutical molecules is nearly impossible to achieve by ICP-QMS. However, ICP-QQQ operating in MS/MS reaction cell mode can be applied to resolve these spectral interferences, allowing the quantification of S, P and CI at far lower levels (biologically relevant concentrations) than was previously possible by ICP-QMS.

In this study, five APIs and a monoclonal antibody (mAb) were analyzed using ICP-QQQ. The targeted compounds included small (m = 250-320 Da) and large (m = 146 kDa for the mAb) molecules.

Experimental

Instrumentation: An Agilent 8800 ICP-QQQ #100 was coupled to an Agilent 1260 Infinity Bio-inert HPLC system with quaternary pump (G5611A) and autosampler (G5667A). An HPLC flow rate of 0.4 mL/min and an injection volume of 20 μ L were applied throughout the study.

CRC conditions: O_2 at 0.3 mL/min. H_2 flow at 3.0 mL/min. Octopole bias = -4 V and KED = -8 V.

Acquisition conditions: MS/MS O_2 mass-shift method for S and P measurement and H_2 mass-shift method for CI measurement.

LC conditions: Two types of columns were used: an Agilent ZORBAX plus C18, 2.1 x 100 mm, 3.5 μ m (Agilent # 959793-902) was used for the analysis of the small molecules, and an Agilent Bio SEC-3 300 Å, 4.6 x 150 mm, 3 μ m (Agilent # 5190-2514) was employed for the mAb analysis.

Reagents: Sulfamethizole, sulfamethazine, sulfamethoxazole, zoledronic acid hydrate and clonidine hydrochloride were purchased from Sigma Aldrich (St. Louis, MO, US). The monoclonal antibody (IgG2a) was obtained from Agilent Technologies (Agilent #200473).

Results and Discussion

Sulfur-containing APIs

Three sulfonamide APIs, sulfamethizole, sulfamethazine and sulfamethoxazole, were dissolved separately in methanol or methanol/water. Each sample was filtered, diluted with the LC mobile phase of 13% acetonitrile with 0.1% formic acid, and injected into the HPLC using an isocratic separation. The resulting overlaid chromatograms are shown in Figure 1. The method detection limit (MDL) for the compound sulfamethizole was calculated to be 23 nM (6.3 ppb as the compound and 1.5 ppb as S).

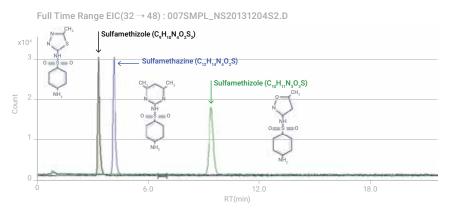


Figure 1. Overlaid chromatograms of three sulfur-containing APIs. The concentration of S in all three APIs injected is 100 ppb.

Antibodies are glycoproteins that contain about 1% sulfur and are therefore excellent targets for quantification via sulfur determination by ICP-QQQ. A mAb (IgG2a) obtained from Agilent was diluted with UPW and injected into the HPLC. An isocratic mobile phase of 50 mM phosphate buffer adjusted to pH 7.0 was used. Figure 2 shows the overlaid chromatograms obtained for two different concentrations of IgG2a. The MDL was calculated to be 14 nM (40 ng) as the compound.

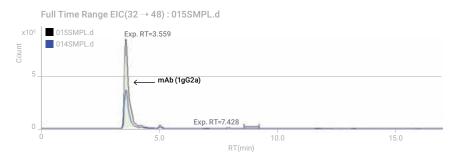


Figure 2. Overlaid chromatograms of 0.5 mg/mL and 1.0 mg/mL mAb (IgG2a) obtained by HPLC-ICP-QQQ.

Phosphorus-containing API

ZOMETA® is a commercial drug that is used for the treatment of hypercalcemia (high levels of blood calcium). It contains zoledronic acid monohydrate $(C_5H_{10}N_2O_7P_2\cdot H_2O)$, which acts as an inhibitor of osteoclastic bone resorption and an inducer of osteoclast apoptosis. A 5 mL vial of commercially supplied ZOMETA® (containing 4.264 mg of the API) was prepared by diluting the drug

2000-fold with the LC mobile phase to give a final API concentration of 426.4 μ g/L. The isocratic mobile phase consisted of a 70:30 mixture of A: 6 mM tetra-butyl-ammonium bromide and 5 mM acetic acid adjusted to pH 6.5 with NH₃(aq), and B: 95% MeOH. A calibration curve was prepared using zoledronic acid monohydrate standards, and the API in the sample was quantified based on the response for P compared to the external calibration. The concentration of the API in the sample was determined to be 433 ng/mL, which is a recovery of 102%. The MDL for the drug compound was calculated to be 25 nM (144 pg; 7.2 ppb as compound and 1.5 ppb as P).

Chlorine-containing API

Catapres® is a commercial drug that is used for the treatment of hypertension. It contains clonidine hydrochloride ($C_9H_9Cl_2N_3\cdot HCl$) which acts in the brain to suppress secretion of noradrenaline, lowering blood pressure. A tablet of Catapres® (containing 75 µg of the API) was dissolved in 50 mL water and sonicated for 60 minutes. The solution was then filtered and analyzed by HPLC-ICP-QQQ. A calibration curve was prepared by analyzing clonidine hydrochloride standards, and the API in the sample was quantified by external calibration. The isocratic HPLC method used a mobile phase consisting of 20% acetonitrile with 0.1% formic acid adjusted to pH 7.0 by NH $_3$ (aq). The calibration curve for CI measured as 35 CIH $_2$ + at m/z 37 and the chromatogram of clonidine hydrochloride measured in the Catapres® sample are presented in Figure 3. The concentration of the API in the sample was determined to be 1444 ppb, which is a recovery of 96%. The MDL of the compound was 146 nM (780 pg; 39 ppb as compound and 15 ppb as CI).

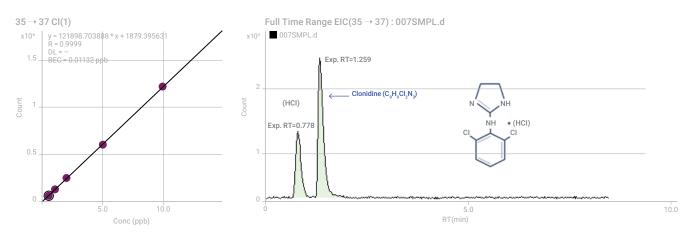


Figure 3. (Top) Calibration curve for CI (measured as 35 CIH $_{2}^{+}$) in clonidine hydrochloride (C_{9} H $_{9}$ CI $_{2}$ N $_{3}$ ·HCI) standards. (Bottom) Chromatogram of clonidine hydrochloride in Catapres® sample.

Conclusions

The advanced capability of the Agilent 8800 ICP-QQQ operating in MS/MS mode has been successfully applied to the analysis of APIs and mAb, based on the measurement of the heteroatoms S, P and CI — an analysis that is normally carried out using molecular-MS techniques. These preliminary studies are presented here in order to demonstrate the potential use of HPLC-ICP-QQQ in drug development and post manufacturing QA/QC control.

Fast and Accurate Absolute-quantification of Proteins and Antibodies using ID-ICP-QQQ

Authors

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Keywords

proteins, antibody, absolutequantification, isotope dilution, ICP-MS/MS, ICP-QQQ

Introduction

Triple quadrupole ICP-MS (ICP-QQQ) dramatically improves the efficiency and reliability of removing spectral interferences on a wider number of elements than conventional ICP-MS. Challenging elements such as sulfur (S), which suffer intense spectral overlaps, can be analyzed at low levels by ICP-QQQ. Furthermore, the effective removal of spectral overlaps allows access to multiple isotopes of elements, enabling quantification of metalloproteins and peptides using isotope dilution mass spectrometry (IDMS) analysis. IDMS is an absolute quantification technique that eliminates the requirement for compound-specific calibration standards. It allows accurate quantification without the need for a reference standard, which is a major benefit of ID-ICP-MS/MS for life science research, where many compounds are unknown.

In this study, we evaluated an Agilent 8800 ICP-QQQ and isotope dilution analysis (ID-ICP-QQQ) of sulfur, for the quantification of NIST Bovine Serum Albumin (BSA) 927e standard reference material (SRM) [1] and a monoclonal antibody, trastuzumab.

Experimental

Instrumentation: An Agilent 8800 #100 was used.

Acquisition conditions: two sulfur isotopes, 32 S and 34 S, were measured in MS/MS mass-shift mode with oxygen (O_2) reaction gas.

Plasma conditions: RF power =1550 W, nebulizer gas flow rate = 0.25 L/min, and dilution gas flow rate of 0.85 L/min.

Double isotope dilution method: in a simple ID method, a sample containing an unknown amount of S, which is primarily composed of the major ³²S isotope (94.93% abundance), is spiked with a known amount of a certified enriched isotopic standard solution containing ³⁴S. An aliquot of the resulting solution is analyzed, and the ratio of ³⁴S to ³²S is measured. From the measured ratio and the known amount of ³⁴S, it is possible to calculate the amount of ³²S and therefore the total S concentration (based on natural isotopic abundances) in the original sample.

However, as the 34 SO $_{4}^{2-}$ spiking solution used in this study was prepared by oxidation of a powder of 34 S sulfur, the exact concentration of 34 S present in the spike was unknown. Therefore, a high accuracy technique known as double IDMS was employed in this study per Equation 1. The concentration of 34 S in the H_{2}^{34} SO $_{4}$ solution was determined by reverse IDMS. A National Institute of Standards and Technology (NIST) certified solution of SO $_{47}$ with a natural sulfur isotopic abundance, was used as the reference standard.

$$w_x = w_z \cdot \frac{m_y \cdot m_z}{m_x \cdot m_{y\prime}} \cdot \frac{R_y - R_{xy}}{R_{xy} - R_x} \cdot \frac{R_{zy} - R_z}{R_y - R_{zy}}$$

- · x refers to the sample
- y and y' refer to the 34SO₄ spiking solution
- z refers to the NIST SO₄ standard solution
- w_x is the sulfur mass fraction (µg/g) in the sample
- $\mathbf{w_z}$ is the sulfur mass fraction ($\mu g/g$) in the NIST $\mathrm{SO_4}$ standard solution
- m_i is the mass of sample, standard, or spiking solution
- \bullet R $_{i}$ is the $^{34}\text{S}/^{32}\text{S}$ ratio measured by ICP-QQQ in the unspiked and spiked solutions
- R_v is the ³⁴S/³²S ratio measured in the sample solution
- R, is the 34S/32S ratio measured in the spiking solution
- R₇ is the ³⁴S/³²S ratio measured in the SO₄ standard solution
- \bullet $m_{_{\! X}}$ is spiked with my and ratio $R_{_{\! X\! Y\! }}$ is measured
- m_z is spiked with my' and ratio R_{zv} is measured

Equation 1. The double IDMS equation used in this study.

Samples: sample solutions were prepared for microwave digestion. First, an amount of sample (BSA standard and trastuzumab solution) estimated to contain approximately 50 μ g sulfur was weighed into a disposable glass tube. 50 μ g of 34 S (as $\rm H_2^{34}SO_4$) was added, followed by 2 mL of 69% HNO $_3$, 0.5 mL of 37% HCl, and 1 mL of 30 % $\rm H_2O_2$. Once the microwave digestion program had finished, the digest was transferred and diluted to 50 mL with $\rm H_2O$. The concentration of S in solution was about 1 ppm.

A standard was also prepared for the double IDMS method. 50 μ g of a 1000 mg/L sulfur ICP-MS standard (natural isotopic abundance) was weighed into another glass tube. The above procedure was then carried out.

Results and Discussion

Six samples of NIST BSA 927e were quantified using the ID-ICP-MS/MS method. The average recovery to the certified value (67.38 \pm 1.38 g/L as S) was 101.26 % and the RSD of six analyses was 0.22%.

Matrix effects were also investigated. The same amount of BSA was spiked with different formulation ingredients. The solutions were then digested and analyzed. The results in Figure 1 show good recoveries were obtained for S in all matrix solutions.

Trastuzumab is a monoclonal antibody (mAb) that interferes with the human epidermal growth factor receptor 2 (HER2/neu receptor). The mAb in solution was quantified using the developed method. The average recovery to expected value (21 mg/mL as S) was 97.8% and the RSD of three analyses was 0.02%.

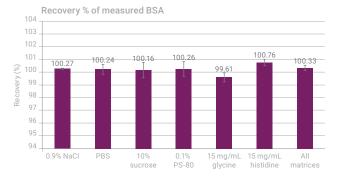


Figure 1. Recovery % (average of n = 3) of measured BSA in various matrix solutions. The error bars show the standard deviation of the three analyses.

Conclusions

The Agilent 8800 ICP-QQQ with MS/MS mode provides high analytical sensitivity and effective interference reduction for the determination of multiple sulfur isotopes. This capability allows the accurate analysis of biological molecules that contain sulfur, using isotope dilution analysis. The ID-ICP-QQQ method is suitable for the accurate and precise quantitative analysis of biological molecules, such as pure proteins and antibodies, without the need for compound-specific calibration standards.

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More Information

Fast and accurate absolute-quantification of proteins and antibodies using Isotope Dilution-Triple Quadrupole ICP-MS, Agilent publication <u>5991-6118EN</u>.

Determination of Diclofenac and Its Related Compounds using RP-HPLCICP- QQQ

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Compound structure-independent quantification of drugs

Introduction

Quantitative drug metabolite profiling is an important application in the pharmaceutical industry. Researchers involved in drug development require an analytical technique with a response that is independent of compound structure. This compound-independent response enables accurate quantification of the drug and its metabolites, without requiring compound-specific calibration. Currently, radiolabeling techniques followed by HPLC separation and radiodetection are used for this application, but a simpler, quicker, and safer alternative approach is desirable.

The very high temperature plasma ion source and elemental ion-based measurement of ICP-MS enables compound structure-independent quantification, so individual standards for the metabolites of the (candidate) drug are not required. ICP-MS also links seamlessly with chromatography systems for speciation studies, for example HPLC.

HPLC-ICP-MS is used in a wide range of applications, including speciation studies of metals and metalloids, such as arsenic, mercury, selenium, chromium, and antimony [1]. However, many drugs contain nonmetal heteroatoms such as phosphorus, sulfur, chlorine, fluorine, or bromine, rather than metals and metalloids. The determination of these nonmetals is difficult by conventional single quadrupole ICP-MS, due to poor ionization, spectral overlaps, high backgrounds, or a combination of these factors. Except for F, these "difficult" elements can be measured accurately at low levels by triple quadrupole ICP-MS (ICP-QQQ) operating in MS/MS mode with a reactive cell gas. Reversed phase (RP) HPLC coupled to ICP-QQQ can introduce further analytical challenges due to the changing composition of the mobile phase gradient. In this study, it was necessary to compensate for the effect of gradient elution on the instrumental response during the RP-HPLC-ICP-QQQ analysis [2].

This note describes the quantitative determination of the drug diclofenac and its related compounds in human plasma. Diclofenac is a prescription non-steroidal anti-inflammatory drug (NSAID) that is used to alleviate mild to moderate pain, fever, and inflammation. Compounds were quantified based on measurement of the CI heteroatom using RP-HPLC-ICP-QQQ.

Chlorine is not a typical analyte for ICP-MS, due to its poor ionization, high background signal, and the presence of intense spectral overlaps. The element's very high first ionization potential of 12.967 eV means that Cl atoms are only converted to positive ions (Cl*) with an efficiency of about 0.13 % in an argon plasma operating at a nominal temperature of 7000 K. Chlorine is also a common contaminant in the laboratory, either from handling of sample containers, sample preparation equipment, or instrument hardware. Also, HCl acid is commonly used for stabilization of many elements, and chlorine tablets are often used as a

biocidal treatment in deionized water systems, leading to high background. Finally, both isotopes of CI (35 CI and 37 CI) suffer from polyatomic interference from polyatomic ions including O_2 H $^+$, SH $^+$, and ArH $^+$.

Experimental

Samples

Diclofenac sodium (99.9% purity) and 4'-hydroxydiclofenac (99.0% purity) were bought from Sigma-Aldrich (St. Louis, MO, USA). Mixed working solutions containing diclofenac sodium and 4'-hydroxydiclofenac were used for method development, external standard calibration, and method validation. Full details are given in Reference 2. Diclofenac was synthetically degraded to generate degradation products covering a broad hydrophobicity range [2]. The synthetically degraded diclofenac samples were used as part of the mass balance study.

Human blood plasma was collected from healthy individuals, pooled, and stored at -20 °C until analysis. Sample preparation details are given in Reference 2.

Instrumentation

An 8800* Triple Quadrupole ICP-MS (ICP-QQQ) was used for all measurements; the instrument was fitted with a PFA nebulizer and platinum cones. The spray chamber was set to a temperature of -1 °C and a plasma torch with a 1.0 mm internal diameter injector was used. These changes helped ensure plasma stability with the high vapor pressure from the volatile organic solvent-based mobile phase (B). Oxygen (20% $\rm O_2$ in Ar) was added to the carrier gas flow at 0.20 L/min to prevent the build-up of carbon on the interface.

To address the spectral overlaps on CI, the major isotope 35 CI (75.78% abundance) was measured by ICP-QQQ in MS/MS mode using a mass-shift method with H $_2$ cell gas. In this mode, ICP-QQQ avoids the interferences on 35 CI by measuring the product ion 35 CIH $_2$ $^+$ at m/z 37 [3]. ICP-QQQ operating conditions and parameters are given in Table 1.

 $\textbf{Table 1.} \ \mathsf{ICP}\text{-}\mathsf{QQQ} \ \mathsf{operating} \ \mathsf{conditions} \ \mathsf{and} \ \mathsf{acquisition} \ \mathsf{parameters}.$

Parameter	Value	
RF power (W)	1570	
Ar carrier gas flow rate (L/min)	0.30	
Optional gas (20% O ₂ in Ar) mass flow controller setting	20% (0.2 L/min)	
Spray chamber temp (°C)	-1	
H ₂ cell gas flow rate (mL/min)	3.5	
Monitored transitions/masses, Q1 🏻 Q2 (m/z)	35 (CI+) [] 37 (CIH ₂ +)	
Data collection mode	TRA	
Integration time (s)	0.4 for m/z 37	

The ICP-QQQ was coupled to an Agilent 1260 Infinity HPLC System equipped with an Agilent 1260 Infinity Vacuum Degasser, an Agilent 1260 Infinity Binary Pump, an Agilent 1260 Infinity Autosampler, an Agilent 1290 Infinity Thermostatted Column Compartment, and an Agilent 1290 Infinity Series 2-position/10-port Microvalve. Column details and operating conditions are given in Table 2.

^{*} The 8800 ICP-MS has been superceded by the 8900 model.

To compensate for the increased sensitivity for Cl caused by the changing level of acetonitrile during the gradient elution, a mathematical correction was applied to the Cl response (measured as $^{35}\text{ClH}_2^{\,+}$). The correction was based on the measured variation of the Cl response with increasing acetonitrile concentration, as shown in Figure 1.

The chromatographic peaks for the drug and metabolite compounds were identified by retention times (RT), and each peak area was then integrated. The organic solvent concentration at the RT of each peak was calculated from the LC gradient program. The appropriate response factor for each peak was then determined from the organic solvent concentration and the response curve. Finally, the corrected CI concentration of each peak was calculated based on the peak area and the corresponding response factor.

Table 2. HPLC operating conditions.

	Online preconcentration
Analytical column	1570
Eluent A	0.1% (v/v) formic acid in MQ water
Eluent B	0.1% (v/v) formic acid in acetonitrile
Gradient	-1
Flow rate (mL/min)	1.0
Sample temp (°C)	5
Column temp (°C)	22-23 (room temp)
Injection volume (µL)	50

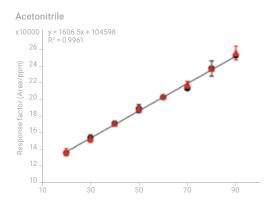


Figure 1. Measured response curve demonstrating the effect of the organic solvent (acetonitrile) content of the mobile phase on the CI response of ICP-QQQ for both inorganic CI and diclofenac-CI (95% confidence intervals, n=3). The response factor was found to be independent of the chemical form, as expected.

Method Development and Method Validation

Selectivity

Compound selectivity—the ability of a technique to distinguish an individual compound from other (often related) compounds—was confirmed by comparing the chromatograms shown in Figure 2. The chromatograms include (i) a blank, (ii) a mixture of 4'-hydroxydiclofenac and diclofenac, each at a concentration equivalent to 1 mg/L (ppm) Cl (iii) synthetically degraded diclofenac at a concentration equivalent to 10 mg/L Cl, and (iv) synthetically degraded diclofenac (at 10 mg/L Cl) spiked with 4'-hydroxydiclofenac (at 1 mg/L Cl).

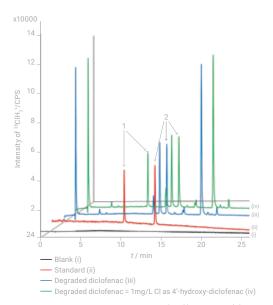


Figure 2. Chlorine chromatograms for (i) a blank, (ii) 4'-hydroxydiclofenac plus diclofenac, each at a concentration equivalent to 1 mg/L CI, (iii) synthetically degraded diclofenac at 10 mg/L CI, and (iv) synthetically degraded diclofenac at 10 mg/L CI, spiked with 4'-hydroxydiclofenac at 1 mg/L CI. Peak 1: 4'-hydroxydiclofenac; peak 2: diclofenac; other peaks: degradation products with unknown chemical structure.

Accuracy and precision

Accuracy and precision were investigated by spiking blank human plasma with 4'-hydroxydiclofenac and diclofenac at three concentration levels (three replicates at each level). The recovery was determined for both compounds. To assess the precision of the method, both intraday and interday precision were studied. As summarized in Table 3, excellent results were obtained with recoveries between 90-100% and RSDs below 4%.

Table 3. Accuracy and precision of the results for 4'-hydroxydiclofenac and diclofenac spiked into human plasma matrix.

Cl conc (mg/L)	Recovery (%)		Intraday precision (RSD%)		Interday precision (RSD%)	
	4'-hydroxy-diclofenac	Diclofenac	4'-hydroxy-diclofenac	Diclofenac	4'-hydroxy-diclofenac	Diclofenac
0.5	92.4	97.5	2.7	3.2	3.1	2.5
1.0	95.0	97.2	1.8	1.8	1.9	2.3
3.0	91.9	91.8	0.2	0.3	0.3	0.6

Linearity and Limit of Quantitation

The linearity of the method was tested by injecting diclofenac standard solutions at concentrations equivalent to between 0.05 and 5.0 mg/L Cl. Excellent linearity was achieved with an $R^2 > 0.99$ and with the origin included in the 95% confidence interval of the intercept.

The limit of quantification (LOQ) was determined according to the signal-to-noise (S/N) method described in the ICH Q2(R1) guidelines (Part II, section 7.2). The LOQ for diclofenac was a compound concentration equivalent to 0.05 mg/L Cl.

Mass balance study

The mass balance study was performed using a blank solution and a human plasma matrix. Each solution was spiked with synthetically degraded diclofenac at a level equivalent to a nominal total CI concentration of 10 mg/L (the actual total spike amounts are shown in Table 4). The total CI content of all the compounds in the spiked samples was measured and the concentration and recovery results are given in Table 4.

The recovery for the total CI content was excellent, both in the absence and presence of the human plasma matrix (92 and 93 %, respectively). This matrix-independent response was further confirmed by comparing the results separately for each degradation product peak with and without the plasma matrix. The relative percent differences (RPDs) observed were mostly less than 5%. Slightly higher differences—up to 12% RPD—were observed for compounds that were present at levels close to the LOQ of 0.05 mg/L. It can be concluded that the human plasma matrix does not introduce any bias to the results obtained using this method.

Table 4. Comparison of mass balance studies with and without human plasma sample matrix.

RT/min	v/v % of acetonitrile*	Cl conc without plasma (mg/L)	Cl conc with plasma (mg/L)	Relative difference (%)**
1.4	30.0	2.10	2.10	0.0
4.4	38.3	0.10	0.10	-2.3
11.1	57.1	0.43	0.44	3.3
11.8	59.2	1.29	1.30	1.5
12.7	61.6	1.25	1.27	1.0
14.7	67.2	0.07	0.07	-9.3
14.8	67.6	0.15	0.14	-5.6
17.0	73.6	2.89	2.94	1.5
18.9	79.0	0.15	0.16	0.2
Measured total CI content (mg/L)		8.49	8.57	
True spike amount (mg/L)		9.21	9.21	
Recovery (%)		92.2	93.1	

^{*}The eluent composition in which the compound with the indicated retention time is eluted.

^{**}Results in spiked plasma, relative to the values obtained without human plasma matrix.

Enhancing Sensitivity for CI

Online sample preconcentration

To improve the sensitivity of the method and enable metabolite profiling of low-dose Cl-based drugs, a simple sample preconcentration procedure was used. The drug-related compounds present in human plasma were trapped on a trapping column (Waters XBridge BEH C18 4.6x20 mm; 3.5 μ m) before analytical separation and ICP-QQQ detection. No additional sample pretreatment was required. The injection volume was increased to 1500 μ L to load the preconcentration column. More details can be found in Reference 2. Using preconcentration, the LOQ for diclofenac was equivalent to 0.002 mg/L Cl - a 25-fold improvement. Human plasma blanks were spiked with 4'-hydroxydiclofenac and diclofenac at three concentration levels between 0.005 and 0.05 mg/L Cl (5 to 50 μ g/L, ppb). Excellent recoveries between 94 and 98% were obtained for both compounds at all concentration levels, as shown in Table 5.

Table 5. Recoveries obtained for 4'-hydroxydiclofenac and diclofenac in the presence of human plasma matrix, when using a simple sample preconcentration procedure.

Cl conc (µg/L, ppb)	Recovery (%)		
	4'-hydroxydiclofenac	Diclofenac	
5	95.7	96.7	
30	97.8	95.7	
50	97.4	93.9	

Conclusions

A reversed phase HPLC-ICP-QQQ method has been successfully used for the compound-independent quantitative determination of diclofenac and its related compounds. Based on the measurement of the CI heteroatom, the new HPLC-ICP-QQQ approach is quicker, simpler, and safer than the traditional radiolabeling HPLC technique.

Since CI has a high first ionization potential and is poorly ionized in the ICP plasma, ICP-MS sensitivity is usually low. This was overcome using a simple online sample preconcentration procedure. The drug-related compounds from a larger injection volume of human plasma were trapped on the preconcentration column, leading to a 25-fold improvement in the LOQ of CI. This step broadens the application to metabolite profiling of low-dose pharmaceutical drugs containing CI at sub mg/L levels.

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- 3. Naoki Sugiyama, Trace level analysis of sulfur, phosphorus, silicon and chlorine in NMP using the Agilent 8800 Triple Quadrupole ICP-MS, Agilent publication, 2013, 5991-2303EN

More Information

For a full account of this application, see Balazs Klencsar, Lieve Balcaen, Filip Cuyckens, Frederic Lynen, Frank Vanhaecke, Development and validation of a novel quantification approach for gradient elution reversed phase high-performance liquid chromatography coupled to tandem ICP-mass spectrometry (RP-HPLC-ICP-MS/MS) and its application to diclofenac and its related compounds, *Analytica Chimica Acta* 974, **2017**, 43–53, doi.org/10.1016/j.aca.2017.04.030.

Absolute Quantification of Proteins in Snake Venom

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Introduction

Venoms are complex biological fluids that contain unique mixtures of peptides and proteins. Identifying and quantifying the composition of venoms is of increasing scientific interest. It is especially important to characterize the toxins responsible for the severe biological effects of the venom on humans. Better understanding of the clinical symptoms of envenoming would help with the development of effective therapies. Also, venoms are being investigated as potential sources of new compounds in drug development.

In this study, an absolute quantification method suitable for the direct quantification of intact proteins was applied to the analysis of the venom of the Mozambique spitting cobra (*Naja mossambica*). This cobra is one of the most dangerous snakes in Africa. The cobra's venom is mainly toxic to cells (cytotoxic), causing swelling of the bite wound that may evolve into tissue necrosis and gangrene [1, 2]. The cytotoxic components of the cobra venom have been identified mainly as members of the three-finger toxin (3FTx) and phospholipase A2 (PLA2) protein families [3, 4].

The methodology was based on capillary liquid chromatography (capLC) coupled to a triple quadrupole ICP-MS (ICP-QQQ). Absolute protein quantification was achieved by measuring the sulfur heteroelement in the proteins, calibrated using online isotope dilution analysis (IDA). ICP-QQQ uses MS/MS mode to control reaction chemistry in the collision/reaction cell (CRC), giving consistent removal of spectral interferences using reactive cell gases [5]. Efficient removal of spectral overlaps using MS/MS allows access to multiple isotopes of biologically important elements such as iron, sulfur, and selenium. ICP-QQQ enables the quantification of metalloproteins and peptides using IDA, without the need for compound-specific calibration standards [6, 7].

Most proteins (> 95%) contain sulfur from methionine and cysteine residues [8, 9], but sulfur determination is difficult by conventional single-quadrupole ICP-MS. The challenges are due to the element's high ionization potential (10.4 eV) that leads to low sensitivity, and the occurrence of spectral interferences from multiple polyatomic ions that overlap all isotopes of sulfur. ICP-QQQ provides low backgrounds and high sensitivity, and removes multiple matrix interferences using MS/MS, freeing up the three most abundant S isotopes for accurate measurement using IDA.

In this study, capLC-ICP-QQQ was used for the quantitative analysis of intact proteins, isolated, and present in simple mixtures. The method was also applied to the analysis and quantification of the major toxins comprising the venom proteome of the Mozambique spitting cobra.

Experimental

Reagents and samples

Methionine and BOC-Met-OH (Sigma-Aldrich, Germany) were used as standards. Bovine serum albumin (BSA), transferrin, β -casein, and cytochrome C (Sigma-Aldrich, Germany); and intact monoclonal antibody (mAb) Mass Check Standard (Waters, USA) were used as protein standards in the recovery study. Other reagents included sulfur (1000 mg/L S) ICP standard (Merck KGaA, Germany); solid isotopically enriched 34 S (Isoflex USA); and sodium hydroxide (VWR Chemicals Belgium). Lyophilized *Naja mossambica* venom was obtained from the specialist venom supplier Latoxan S.A.S., France. The venom was collected from a snake from Tanzania, and was stored at -20 °C before use. All solutions were prepared in Milli-Q water (ChemLabor Millipore system, with 0.22 μ m filter, Millipak - Millipore). Mobile phase B was prepared in acetonitrile (ACN) Optima® LC/MS (Fischer Scientific, USA). Formic acid (FA) was bought from Merck KGaA (Germany).

Instrumentation

Capillary LC separation was performed using an Agilent 1200 Series HPLC system fitted with a BIOShellTM A400 C4, 3.4 μ m, 150 mm x 0.3 mm reversed-phase column (Sigma-Aldrich, Germany) and autosampler. Chromatographic column and post-column connections comprised Agilent PEEK-coated fused silica capillaries 200 mm x 100 μ m id (ICP and syringe connection) and 50 μ m (column connection), and a 0.03" (0.8 mm) Agilent zero-dead volume T-connector. Post-column flow was provided by a syringe pump system kdScientific (Holliston, MA, USA). The column was heated using a Spark Holland oven (Mistral, The Netherlands) to improve chromatographic efficiency.

Sulfur isotope measurements were carried out using an Agilent 8800 Triple Quadrupole ICP-MS (ICP-QQQ). The capLC column was connected to the ICP-QQQ using the Agilent capillary LC interface kit (Agilent product number G3680A). The kit consists of a total consumption nebulizer with single pass spray chamber. Enriched 34S was added to the LC eluate solution post-column. Sulfur isotopes were measured by ICP-QQQ using an oxygen mass-shift MS/MS method and the S content of the proteins was determined by IDA [10]. BOCMet-OH was used as the internal standard (IS) to correct for any injection errors. capLC-ICP-QQQ operating conditions are given in Table 1.

Table 1. capLC-ICP-QQQ operating conditions.

ICP-QQQ				
RF power (W)	15	50		
Sampling depth (mm)	8.	.0		
Carrier gas flow rate (L/min)	0.	85		
Make-up gas flow rate (L/min)	0.0	00		
O ₂ cell gas flow rate (mL/min)	0.	16		
Data acquisition ion pairs, Q1 (S ⁺) » Q2 (SO ⁺) mass (m/z)	·-·	□ 48 □ 50		
capLC				
Chromatographic flow rate (µL/min)	4.5;	3.5*		
Mobile phase A	H ₂ 0/0	H ₂ O/0.2% FA		
Mobile phase B	AcN/0	AcN/0.2% FA		
Temperature (°C)	8	80		
Chromatographic gradient	Time range (Min)	% mobile phase B		
BSA and Intact mAb standards	0	2		
	2	2		
	16	60		
	18	90		
Naja mossambica sample	0	1.5		
	8	1.5		
	10	10		
	40	30		
	47	90		

^{*} Conditions for measurement of venom sample

Results and Discussion

Absolute protein quantification

Absolute quantification of proteins was achieved through the measurement of sulfur by capHPLC-ICP-QQQ, with postcolumn IDA. Two individual protein samples (BSA and mAb) were spiked with BOC-Met-OH IS. Quantitative ID (mass purity) results for BSA, 95 \pm 5% (n=3), compared well with the sample purity data provided by the manufacturer (\geq 98 %). The ID mass purity result for Intact mAb was 77 \pm 4% (n=3).

To validate the method further, digests of the two protein standards were quantified using external calibration of sulfur. The results obtained by external calibration were 96 \pm 1% for BSA and 79 \pm 2% for Intact mAb. These results are in good agreement with the ID mass purity results for BSA and Intact mAb shown above.

Quantitative analysis of snake venom proteome

The method was applied to the analysis and quantification of the proteins present in a snake venom sample. Before analysis, chromatographic recovery was calculated by measuring the sulfur mass of a sample eluting from the chromatographic column and comparing this to the sulfur mass recorded directly by flow injection (FI). Chromatographic recovery for a series of protein standards (cytochrome C, β -casein, transferrin, BSA, Intact mAb) was higher than 98%. For the *Naja mossambica* protein content, it was 99 ± 1% (n=3). The total sulfur mass content corresponded almost exactly to the sum of the sulfur contained in the different venom protein peaks detected via their sulfur heteroelement content (Figure 1). The excellent recovery obtained for the complex sample shows that quantitative protein recoveries for the chromatographic column are species (individual protein) independent.

Parallel capLC-ESI-MS analysis was used to identify the proteins from their molecular weight, according to database information (Table 2). By identifying the proteins, it was possible to know the S-to-protein stoichiometry of each peak. This information was then used to translate sulfur mass into individual protein quantities, in µmol protein per gram of venom sample. The quantified results are summarized in Table 2.

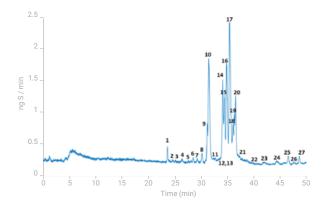


Figure 1. capLC-ICP-QQQ mass flow chromatogram of *Naja mossambica* venom. All the venom protein species eluted between 20 and 50 min (S detection). The 27 sulfur-containing peaks are numbered—see Table 2 for more information.

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Table 2. Matching of the masses of protein peaks from *Naja mossambica* venom to known protein families. Closest available protein species, estimated exact mass, and calculated concentration are listed. Uncertainty corresponds to one standard deviation (n=3).

Peak	Family	Closest homolog	Mol wt (Da)	µmol protein/g venom sample
1	3FTx	-	7064.2	1.99 ± 0.06
2	3FTx	~P29179	7417.4	0.471 ± 0.066
3	3FTx	~P29179	7451.6	0.325 ± 0.040
4	3FTx	~P01420	6892.4	1.10 ± 0.13
5	3FTx	~Q9W6W6	7786.4	< 0.1
6	3FTx	~P01452	7277.3	0.680 ± 0.050
7	3FTx	~P01452	7306.3	0.668 ± 0.057
8	3FTx	-	7246.2	1.35 ± 0.10
9	3FTx	P25517	6832.4	5.09 ± 0.28
10	3FTx	P01452	6704.3	19.0 ± 0.8
11	3FTx	-	6686.3	0.183 ± 0.035
12	3FTx	-	6829.3	0.220 ± 0.039
13	3FTx	-	6687.3	< 0.1
14	PLA2	P00604	13280.9	7.76 ± 0.32
15	3FTx	P01470	6882.4	9.54 ± 0.27
16	3FTx	P25517	6813.3	16.2 ± 0.4
17	3FTx	P01467	6814.3	27.8 ± 0.8
18	3FTx	~P01469	7046.4	5.13 ± 0.26
19	PLA2	P00604	13237.8	3.40 ± 0.14
20	PLA2	P00002	13196.6	7.35 ± 0.33
21	PLA2	-	13179.7	0.805 ± 0.049
22	Minor	-	42000	0.102 ± 0.009
23	Endonuclease	-	30000	0.619 ± 0.050
24	SVMP	Q10749	46700	0.165 ± 0.009
25	SVMP	Q10750	46700	0.264 ± 0.013
26	SVMP	Q10751	46700	0.097 ± 0.006
27	SVMP	Q10752	46700	0.257 ± 0.006

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Conclusions

The capLC-ICP-QQQ method uses measurement of sulfur by IDA to enable the absolute quantification of intact proteins without the need for protein-specific standards.

Agilent ICP-QQQ instrumentation is especially suited for IDA analysis of S as it uses MS/MS to remove multiple spectral interferences from several S isotopes using an oxygen massshift method. By adding enriched ³⁴S post-column and spiking each sample with a generic S-containing internal standard, multiple S isotopes can be measured as SO+ product ions. MS/MS ensures that each S isotope mass enters the CRC in isolation, so no interfering product ions can be formed from the other isotopes of S. This approach allows the sulfur content of the proteins to be determined by IDA.

The method was applied successfully to the quantification and mass purity confirmation of protein standards. If quantitative chromatographic recoveries can be assured, it is even possible to quantify nonpure protein samples using this method. The potential of the methodology for the quantification of intact proteins present in relatively complex samples was demonstrated by analyzing 27 proteins in snake venom.

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More Information

The full results of the study were published in: Francisco Calderón-Celis et al., Elemental Mass Spectrometry for Absolute Intact Protein Quantification without Protein-Specific Standards: Application to Snake Venomics, *Anal. Chem.*, **2016**, 88 (19), 9699–9706.

Glossary

Α

Acquisition conditions

Parameters including: Peak profile, mass, integration time,

scan number and replicate.

ADME Acronym of absorption, distribution, metabolite, and excretion

studies

Ammonia, NH₃ A reaction gas used in the collision/reaction cell. NH₃ is a

very reactive gas, which is used both in on-mass methods and mass-shift methods to remove/avoid interferences.

amu Atomic mass unit. An obsolete, non-SI unit that is still in

common use in its abbreviated form "amu", meaning the same thing as "unified atomic mass unit" (u) or dalton (Da). All are used to indicate the atomic mass of ions, atoms or

molecules, based on the carbon 12 standard.

API, Active Pharmaceutical Ingredient An API is a compound in a drug which has remedy effects

on the target disorder.

AS, abundance sensitivity

The measure of an analyzer's ability to separate adjacent peaks differing greatly in intensity. Agilent ICP-QQQ with MS/MS operation delivers unmatched peak separation (abundance sensitivity <10⁻¹⁰), as the resolution performance is the product of the abundance sensitivity of the two

quadrupoles.

ASX-520 Autosampler suitable for medium to high sample throughput

applications, with rack configurations providing up to 360 vial positions (up to 720 with the extended rack XLR-8 version).

Axial Acceleration

A function of the ion guide to accelerate/decelerate ions along

the axis of the ion guide.

B

Bandpass Mode of operation of a multipole ion guide, where both a

low-mass cut-off and high-mass cut-off are applied, rejecting ions below and above a certain m/z. A bandpass filter passes a "window" of masses (typically covering a 20-30 m/z range) through the ion guide, and is therefore distinct from a mass filter, which is capable of unit mass resolution (single m/z

mass selection).

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Batch

The acquisition batch contains all the information required for a sample analysis or experiment, including peripump program, tuning conditions, acquisition parameters, sample list and data analysis (calibration) parameters. The data analysis (DA) batch contains the measured results for the batch of samples, and includes the calibration plots, internal standard signals and QC flags.

Equivalent Concentration

BEC, Background The magnitude of a signal in a blank, expressed as a concentration.

BED, Background Equivalent Diameter

BED is the diameter equivalent to the background noise in single particle analysis.

C

Charge transfer

A reaction mechanism that relies on the exchange of charge between ions and cell gas molecules, e.g., $Ar^+ + NH_a \rightarrow Ar + NH_a^+$

Collision mode

A cell mechanism to remove interferences either by collisional dissociation or by kinetic energy discrimination (KED). With KED, ions entering the collision/reaction cell collide with the cell gas (such as helium). Since polyatomic ions have a larger ionic cross-section than mono-atomic analyte ions at the same mass, the polyatomic ions undergo more collisions than the analyte ions, and so lose more energy. By the cell exit, the lower energy ions (the polyatomics) can be separated from the higher energy (analyte) ions by applying a bias voltage "step". This is knows as kinetic energy discrimination (KED).

Cool plasma

A technique used to reduce interferences. Under low temperature plasma (cool plasma) conditions, the formation of interferences such as Ar+, ArO+ and ArH+ is suppressed, allowing the detection of Ca⁺, Fe⁺ and K⁺ at the trace level. Typical RF power for cool plasma is 600-900 W.

CRGS, Carrier gas

Carrier gas is an Ar gas supply flowing through the nebulizer to convert a liquid sample into a fine aerosol. It is a tuning parameter of the plasma.

CRC, Collision/ **Reaction Cell**

Device used to remove interferences from the ion beam, using settings such as cell gas, cell gas flow rate, octopole bias voltage, KED bias and deflection lens.

D

DL

Acronym of Detection Limit. Also called LOD (limit of detection). It is the concentration that is equivalent to 3 times the standard deviation (SD) of the background signal.

Desolvation system

A device to remove solvent from the aerosol generated by the nebulizer.

Dynamic range or analytical working range

The range of linearity of an analytical instrument. Agilent ICP-QQQ instruments are fitted with an advanced, dual-mode, discrete dynode electron multiplier (DDEM) that provides a full nine orders dynamic range under standard operating conditions.

DIGS, Dilution Gas

Argon gas flow added to the carrier gas via a dilution gas port located between the torch and the spray chamber. A dilution gas is used for Aerosol Dilution with HMI or UHMI. The gas supply used for the DiGS can also be switched automatically to add the gas flow to the spray chamber instead (known as make-up gas or MUGS). It is a tuning parameter of the plasma.

Dwell time

The period of time that the analytical instument accumlates

the signal.

Е

Enthalpy of reaction, ΔHr Amount of energy (heat) absorbed or released by a reaction. When ΔHr is positive ($\Delta Hr > 0$), the reaction is endothermic, meaning energy is required (absorbed) for the reaction to occur. When ΔHr is negative ($\Delta Hr < 0$), the reaction is exothermic, meaning energy is released by the reaction, which is spontaneous.

G

GC Interface kit

Agilent's GC-ICP-MS interface features a fully heated inert transfer line and separately heated inert torch injector that provides reliable separation of volatile compounds.

Н

Introduction

HMI, High Matrix HMI Aerosol Dilution technology is standard on Agilent ICP-QQQ, extending the TDS range to % level, while eliminating the added cost, time and potential errors of conventional

liquid dilution.

UHMI, Ultra High Matrix Introduction

Agilent's second generation aerosol dilution system, which allows the direct analysis of 25% NaCl solutions.

Hard extraction

A tuning condition when a negative voltage is applied to the extraction lens. Hard extraction provides higher sensitivity at lower plasma temperature than soft extraction. Cool plasma conditions require hard extraction.

mode

Helium mode, He See collision mode.

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HR-ICP-MS, high-resolution ICP-MS Also known as magnetic sector, sector field or double focusing. Magnetic sector based ICP-MS instruments are capable of resolution (M/ Δ M) of up to 10,000 and are able to resolve most polyatomic species from analytes at the same nominal mass.

Ī

I-AS, Integrated Auto sampler Integrated, covered auto sampler with pumped rinse station; ideal for ultra-trace analysis and small sample volumes (as low as 0.5 mL). Flexible rack configurations offer a maximum capacity of 89 vials, plus 3 rinse vials.

ICP

Inductively coupled plasma, generated by applying a high-power radio frequency (RF) field to a flow of argon gas. The plasma is a high temperature ion source, up to 10,000 K maximum and around 7,500 K in the central channel.

ICP-MS

Inductively coupled plasma mass spectrometer or spectrometry.

ICP-QQQ

Abbreviation for triple quadrupole ICP-MS.

IDA, ID, ID-MS

Isotope Dilution Analysis or Isotope Dilution Mass
Spectrometry is a highly accurate method to quantify
elements based on the change in isotope ratio that results
from the spiking of an unknown sample with a spike enriched
in one isotope of the target analyte. Because each sample
result is based on the measurement of the change in ratio in
that sample, rather than relative to a response in a separate
calibration standard, IDMS results are also directly traceable
to certified standards, which reduces uncertainty.

Inert Sample Introduction kit

O-ring-free and manufactured from PFA for the lowest contamination levels. Demountable torch with Pt or sapphire injector options. HF resistant, and suitable for high-purity reagents.

Interferences - spectral

Direct overlap from a different element with an isotope at the same nominal mass (isobar), or overlap from a polyatomic ion, or doubly-charged ions resulting from the loss of two electrons instead of just one. Because the quadrupole separates ions based on m/z (mass over charge ratio), a doubly-charged ion (M^{2+}) will appear at mass M/2.

Ion guide

Operation of an ion lens where no mass rejection is performed. Applies to simple electrostatic ion lenses, and also to multipole ion guides operated with no low- or high-mass cut-off.

IP, Ionization Potential

The first ionization potential of the element is the energy required to remove one electron from a neutral atom and is specific for each element. Most elements are largely converted (>90%) to singly-charged ions in an argon plasma. Elements with a low second IP will also form some doubly-charged ions.

IR, Isotope Ratio

Ratio of abundance of two isotopes of an element.

Isobar

Refers to isotopes of different elements that appear at the same nominal mass. These overlaps occur when atoms of two different elements (i.e. different number of protons in the nucleus, so different atomic number) each have an isotope with the same atomic weight (same total number of protons plus neutrons in the nucleus, e.g. ²⁰⁴Pb and ²⁰⁴Hg).

Isobaric interferences

Overlaps that occur at the same mass (see isobar). These overlaps/interferences can be resolved by reaction chemistry (e.g. $\rm NH_3$ is used to separate Pb from the Hg overlap), but cannot be separated by high-resolution ICP-MS; separation of 204 Pb from 204 Hg would require a resolution of around 500,000 (50x higher than can be achieved by any commercial high-resolution ICP-MS).

Isotope

A specific form (atomic weight) of an element. Many elements have atoms with different atomic weights, such as Pb 204, 206, 207 and 208; these are called isotopes. The different isotopes of Pb all have 82 protons in the nucleus (Pb has atomic number 82) but a different number of neutrons, so the atomic weight is different for each isotope.

ISTD, internal standard

Internal standards are commonly used in ICP-MS, particularly where samples vary in composition from the calibration standards. Changes in sample transport, nebulization efficiency and signal intensity (long-term drift) would all lead to errors, which may be corrected if an ISTD element with similar behavior is used as a reference.

K

KED, Kinetic Energy Discrimination KED is used to discriminate the analyte ion of interest from interfering ion(s) by the difference of kinetic energy. Refer to collision mode. KED is also used as a tuning parameter of CRC conditions: KED = (Q2 bias voltage) - (octopole bias voltage).

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LA, Laser Ablation

Method used for the direct analysis of solid samples using a laser to vaporize the sample before introduction to the plasma.

LC speciation kits

Sample introduction kits to facilitate LC coupling and provide turn-key methods for common speciation applications. A Capillary-LC connection kit is also available.

M

MUGS. Make-up gas

Make-up gas refers to Ar gas applied to the spray chamber to increase/adjust total injector gas flow rate. It is a tuning parameter of the plasma.

MS/MS mode

Acquisition mode unique to Agilent ICP-QQQ. MS/MS mode operates Q1 as a unit (1 amu window) mass filter and Q2 is also set to the single mass of the target ion or reaction product ion.

Mass balance

Balance between the amount of a substance introduced into the system and excreted from the system.

Mass pair

MS/MS mode requires a mass setting for Q1 and Q2. The selected mass settings for Q1 and Q2 are known as the mass pair. For example when As is measured in O₂ mode, Q1 is set to the precursor ion (As $^+$) at m/z 75 and Q2 is set to the product ion (AsO+) at m/z 91. 75 --> 91 is the mass pair for As in O₂ mode.

Mass filter

Generic term for any mass analyzer cabable of unit mass resolution. Note that the ion guide used in the CRC of some quadrupole mass spectrometers appears physically similar to a quadrupole mass filter. However, these ion guides cannot provide unit mass resolution because of the ion scattering effect at the higher pressures present in the CRC.

Mass spectrum

See spectrum.

Mass-shift method

A method where the analyte ion is reactive and is moved to a new mass free from the original interference. Sometimes referred to as "indirect" measurement, e.g. Se+ reacts with O₂ in the cell and is converted to SeO+. It can then be detected free from the original interference of ArAr+.

software

MH, MassHunter Software package that provides comprehensive instrument control for the Agilent ICP-QQQ and accessories, and integrated data processing.

m-lens

An optional lens of the Agilent 8900. It provides a low BEC for alkaline elements like K and Na under hot plasma conditions. Monoclonal antibody, mAb

Antibody produced by identical antibody-forming cell, which binds to a certain antigen.

MSA, Method of Standard Additions (also known as StdAdd) A calibration solution is spiked at multiple levels directly into the unknown sample, giving a calibration of response against added concentration. MSA eliminates matrix effects by calibrating in the sample matrix.

Ν

Nanoparticle, NP

Sub-microscale particles with a size range from 1 to 100 nm $\,$

in diameter.

Neutral Gain Scan Q1 and Q2 scan together, a fixed mass-shift apart. For example Q2 scans at Q1 + 16 amu for O-atom addition

reactions.

0

O₂, oxygen A reaction gas used with the Agilent ICP-QQQ. A number of

elements can be measured in mass-shift method using ${\rm O_{2^{\prime}}}$ e.g. Se⁺ can be measured as SeO⁺ using ${\rm O_{2}}$ cell gas. ${\rm O_{2}}$ is also added to the plasma carrier gas to decompose the carbon

matrix when organic solvents are analyzed.

Octopole bias (OctP Bias)

A CRC parameter. It is the bias voltage applied to the octopole ion guide, which determines the collision energy of analyte

ions with cell gas molecules.

OIDA, on-line isotope dilution analysis

A very powerful and useful development of traditional isotope dilution, using on-line addition of the isotope spike. Removes the time consuming step of spiking enriched-

isotope standards into each separate sample.

On-mass method A method where reactive interferences are removed to allow

an unreactive analyte to be measured at its original mass. Sometimes referred to as "direct" measurement, e.g., the interference of GdO+ on Yb+ can be removed by the reaction

of GdO+ with NH₃.

Organics kit Contains the sample introduction parts needed to run volatile

organic solvents. Includes organics torch, solvent-resistant

drain kit and uptake tubing.

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ORS

Agilent's CRC design is known as the Octopole Reaction System. It is a temperature-controlled collision/reaction cell with octopole ion guide and four cell gas lines as standard on the Agilent ICP-QQQ. Provides maximum flexibility in collision and reaction modes, and uses a small internal volume cell to ensure rapid cell gas switching and high ion transmission.

ORS4

Fourth generation Octopole Reaction System.

Oxygen atom abstraction or oxygen atom transfer

Reaction mechanism associated with the use of oxygen in the collision reaction cell.

P

Preset method

Preset methods are provided in the ICP-MS MH software. These built-in methods cover a range of predefined operating conditions to suit different applications. Using a preset method, a user can create a new batch with minimum or no customization.

Preset plasma

Preset plasma conditions are a function of ICP-MS MH software. The software provides several predefined plasma conditions that users can select according to the application. This greatly simplifies system optimization by automatically tuning and calibrating the plasma parameters, rather than the user having to set a number of individual plasma tuning parameters. There are three preset plasma conditions that can be selected depending on the sample matrix: Low matrix, general purpose and HMI/UHMI.

Polyatomic, polyatomic ion

A molecular ion (an ion composed of more than one atom) that arises in the plasma or during ion extraction, and can appear at the same nominal mass as analyte ions. Polyatomics are usually interferences (such as ArO+).

Precursor ion scan

Q1 scans a user set mass range, while Q2 is set to a single fixed mass, measuring all the reaction product ions at that mass, formed from the different ions entering the cell as Q1 scans the mass range.

Product ion scan Q1 is set to a fixed precursor ion mass, while Q2 scans a user set mass range measuring all reaction product ions formed from that single precursor ion.

Q

Quadrupole bias (QP Bias or Qpole Bias) Bias voltage applied to the Q2 rods. Used in conjunction with the Octopole bias to provide a bias voltage "step" at the cell exit, usually to reject unwanted low energy ions from the

ion beam.

Quantitation or quantification

Quantitative results are produced by comparing signal intensities of elements in the sample to those generated by

calibration standards.

Q1 First quadrupole in the configuration of the Agilent ICP-QQQ.

Q1 is positioned in front of the ORS, to control the ions that

are passed to the cell and enable MS/MS operation.

Q2 Second quadrupole in the configuration of the Agilent

ICP-QQQ. Q2 filters the ions that emerge from the cell exit,

passing only the target analyte ions to the detector.

R

Rare Earth Elements, REEs Comprise 17 elements: Sc, Y, La, Ce, Pr, Nd, (Pm), Sm, Eu, Gd,

Tb, Dy, Ho, Er, Tm, Yb, and Lu.

Resolution The ability of a mass filter to separate adjacent masses.

Defined as M/ Δ M; the mass of the target peak/the mass difference to the nearest adjacent peak that can be separated. Sometimes also quoted as the width of the peak at a given

peak height (e.g. 0.75 amu at 10% peak height).

S

SEMI Semiconductor Equipment and Materials International

standards are international standards for materials, chemicals and manufacturing devices used in

microelectronics industries.

Single particle (sp) analysis

In this handbook, spICP-MS analysis refers to particle size measurement using the signal generated from a single

particle.

Single Quadrupole MS, ICP-OMS Conventional ICP-MS containing a single quadrupole

mass filter.

Single Quad mode, SQ mode

Q1 operates as a wide band mass filter. SQ mode emulates

conventional quadrupole ICP-MS.

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Speciation measurement

Combination of chromatographic techniques with ICP-MS as a detector to determine the chemical form of elements in the sample.

Spectrum (mass spectrum)

After separation by the final mass filter (Q2), the ions are detected by an electron multiplier. The detector electronics count and store the total signal for each mass (m/z), creating a mass spectrum. The spectrum that is produced provides a simple and accurate qualitative representation of the sample. The magnitude of each peak is directly proportional to the concentration of an element in a sample.

System

STS, ShieldTorch A technique to eliminate capacitive coupling between the RF coil and plasma, keeping the plasma potential low and energy distribution of ions narrow. The technique is crucial for cool plasma and collision mode.

Т

TDS, total dissolved solids

The total summed concentration of all non-volatile, dissolved inorganic and organic substances in a liquid. The nominal matrix tolerance of ICP-MS instruments is 0.2 % TDS. On Agilent ICP-MS system, this can be extended to approximately 3% TDS with HDMI, and up to approximately 25% with UHDMI.

ICP-MS

Triple quadrupole ICP-MS with a tandem MS configuration, featuring a quadrupole mass filter (Q1) in front of the collision/reaction cell (CRC), which is followed by a second quadrupole mass filter (Q2).

U

UPW

Ultra Pure Water, purified by ion exchange to remove trace contaminants. Used for preparation of standards and for sample dilution for ultra-trace analysis

٧

Venomics

The study of venoms via genomic, proteomic, and transciptomic approaches.

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