

**Supporting Information to**  
**Accurate high throughput quantification of selenium in biological samples-**  
**the potential of combining isotope dilution ICP-tandem mass spectrometry**  
**with flow injection**

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**Table S1.** Precision for each Se isotope ratio is determined from a CRM blend by four consecutive measurements in H<sub>2</sub> and O<sub>2</sub> -ICP-MS/MS mode. CRM blends were prepared and digested as described in the experimental section.

Precision of isotope ratios from CRM blend					
Gas	H <sub>2</sub>		O <sub>2</sub>		
CRM blend	<sup>78</sup> Se/ <sup>77</sup> Se	<sup>80</sup> Se/ <sup>77</sup> Se	<sup>78</sup> Se/ <sup>77</sup> Se	<sup>80</sup> Se/ <sup>77</sup> Se	<sup>82</sup> Se/ <sup>77</sup> Se
Average (n=4)	0.26	0.59	0.24	0.54	0.10
STDEV	0.003	0.006	0.003	0.002	0.002
RSD [%]	1.13	1.11	1.11	0.35	1.96
Mass Bias [%]	3.87	12.6	2.64	13.9	17.6

**Figure S1. (1)** Isotope dilution equation employed for calculating concentration of selenium in direct infusion. The concentration of the spike ( $C_{spk}$ ) was calculated by reverse IDMS, whereas ( $W_{spk}$ ) and ( $W_{std}$ ) respectively refer to the weights of spike and standard.  $R$  refers to the ratio reference-to-enriched isotope into the blend. ( $h_y$ ) refers to the abundance of the enriched isotope while ( $h_x$ ) is the IUPAC correspondent abundance. **(2)** Equation employed for calculating the analyte mass flow ( $M_x(t)$ ). **(3)** The mass flow of the spike ( $M_y(t)$ ) was assessed by multiplying  $C_{spk}$  with the make-up flow rate ( $f_{sp}$ ) **(4)** Selenium concentration was calculated by dividing absolute amounts ( $M_{xj}$ ) as integrated between  $t1$  and  $t2$ , through the injection volume ( $IV$ ). **(5)** ( $K$ ) Mass bias correction was calculated in external calibration: a  $100 \mu\text{g L}^{-1}$  Se-standard was measured 7 times consecutively and the true ratio ( $R_{true}$ ) was divided by the observed ratio ( $R_{obs}$ ).

(1)

$$C = \frac{C_{spk} * W_{spk}}{W_{std}} * \frac{R_y - R_b * K}{K * R_b - R_x} * \frac{h_{iy}}{h_{ix}}$$

(2)

$$M_x(t) = M_y(t) * \frac{R_y - R_b * K}{K * R_b - R_x} * \frac{h_{iy}}{h_{ix}}$$

(3)

$$M_y(t) = f_{sp} * C_{spk}$$

(4)

$$C = \frac{\int_{t1}^{t2} M_{xj}}{IV}$$

(5)

$$K = \frac{R_{true}}{R_{obs}}$$

**Table S2.** Table reporting value/unit and respective standard uncertainties for each factor included in the TCU. Quantification parameters as well as variables are reported in model equation (1).

H <sub>2</sub> mode							
Quantification parameter	Variables	<sup>78</sup> Se		<sup>80</sup> Se		<sup>82</sup> Se	
		Value/unit	Standard uncertainty	Value/unit	Standard uncertainty	Value/unit	Standard uncertainty
Weight of spike	W <sub>spk</sub>	198.2 mg	0.002	198.2 mg	0.002	198.2 mg	0.002
Weight of CRM/serum	W <sub>crm</sub>	182.65 mg	0.002	182.65 mg	0.002	182.65 mg	0.002
Concentration of isotopically enriched <sup>77</sup> Se "spike"	C <sub>spk</sub>	0.67 μmol L <sup>-1</sup> (n=5)	0.026	0.68 μmol L <sup>-1</sup> (n=5)	0.024	0.71 μmol L <sup>-1</sup> (n=5)	0.041
Isotopic abundance of the reference isotope in the spike	A <sub>spk</sub>	0.3 % (certificate)	0.020	0.2 % (certificate)	0.020	0.2 % (certificate)	0.020
Isotopic abundance of the enriched isotope in the spike	B <sub>spk</sub>	99.2 % (certificate )	0.020	99.2 % (certificate)	0.020	99.2 % (certificate)	0.020
Isorope ratio in the test blend	R <sub>b</sub>	0.219 (n=5)	0.003	0.500 (n=5)	0.006	0.097 (n=5)	0.002
Mass bias	K	1.08 (n=7)	0.005	0.986 (n=7)	0.005	0.935 (n=7)	0.006

  

O <sub>2</sub> mode							
Quantification parameter	Variables	<sup>78</sup> Se		<sup>80</sup> Se		<sup>82</sup> Se	
		Value/unit	Standard uncertainty	Value/unit	Standard uncertainty	Value/unit	Standard uncertainty
Weight of spike	W <sub>spk</sub>	200.04 mg	0.002	200.04 mg	0.002	200.04 mg	0.002
Weight of CRM/serum	W <sub>crm</sub>	191.93 mg	0.002	191.93 mg	0.002	191.93 mg	0.002
Concentration of isotopically enriched <sup>77</sup> Se "spike"	C <sub>spk</sub>	0.77 μmol L <sup>-1</sup> (n=5)	0.039	0.79 μmol L <sup>-1</sup> (n=5)	0.018	0.85 μmol L <sup>-1</sup> (n=5)	0.074
Isotopic abundance of the reference isotope in the spike	A <sub>spk</sub>	0.3 % (certificate)	0.020	0.2 % (certificate)	0.020	0.2 % (certificate)	0.020
Isotopic abundance of the enriched isotope in the spike	B <sub>spk</sub>	99.2 % (certificate )	0.020	99.2 % (certificate)	0.020	99.2 % (certificate)	0.020
Isorope ratio in the test blend	R <sub>b</sub>	0.204 (n=5)	0.003	0.491 (n=5)	0.002	0.085 (n=5)	0.002
Mass bias	K	1.05 (n=7)	0.004	0.913 (n=7)	0.003	0.85 (n=7)	0.003

**Figure S2.** Uncertainty budget “pies”: each pie illustrates the contribution brought to respectively  $^{78}\text{Se}$ ,  $^{80}\text{Se}$ ,  $^{82}\text{Se}$  in either gas mode as given by I- Weight of spike. II- Weight of CRM. III- Concentration of selenium in spike. IV- Isotopic abundance of reference isotope in spike. V- Isotopic abundance of enriched isotope in spike. VI- Blend isotope ratio. VII- Mass bias.

