

**Selective hydride generation- cryotrapping- ICP-MS for arsenic speciation analysis at picogram levels: analysis of river and sea water reference materials and human bladder epithelial cells**

Tomáš Matoušek, Jenna M. Currier, Nikola Trojánská, R. Jesse Saunders, María C. Ishida, Carmen González-Horta, Stanislav Musil, Zoltán Mester, Miroslav Stýblo, Jiří Dědina

**Supplementary information:**

Table S1:  
ISIS program for HG-CT-ICP-MS System 1

Step	Duration, s	Scan	Pump 1, rps	Pump 2, rps	Valve 1	Aux1 (heating power)	Aux 2 (heating indicator)
1	<P1>	-	0	<P17>	LOAD	OFF	OFF
2	<P2>	-	<P12>	<P15>	LOAD	OFF	OFF
3	<P3>	-	<P13>	0	INJECT	OFF	OFF
4	<P4>	-	<P13>	<P16>	INJECT	OFF	OFF
5	<P5>	-	0	0	INJECT	OFF	OFF
6	12	-	0	0	INJECT	OFF	ON
7	<P6>	-	0	0	INJECT	ON	ON
8	<P7>	Start	0	0	INJECT	ON	ON
9	<P8>	-	0	0	INJECT	OFF	OFF
10	<P9>	-	0	0	INJECT	ON	ON
11	<P10>	-	0	<P17>	INJECT	ON	ON
12	<P11>	-	<P14>	<P17>	INJECT	OFF	OFF
13	1	-	0	0	LOAD	OFF	OFF

Parameters	Description	Unit	Default value
<P1>	Time Pre-cool	s	30
<P2>	Time Prefill	s	5
<P3>	Time HG, P2 OFF	s	90*
<P4>	Time HG, P2 ON	s	0*
<P5>	Time Delay	s	80
<P6>	Time Acq. delay	s	5
<P7>	Time Acquisition 1	s	37**
<P8>	Time Acq. 2 (Heat off)	s	7
<P9>	Time Acquisition 3	s	22***
<P10>	Time U-Tube cleaning	s	35
<P11>	Time HG cleaning	s	50
<P12>	P1 Rate Prefill	rps	0.25
<P13>	P1 Rate HG	rps	0.25
<P14>	P1 Rate Post-run	rps	0
<P15>	P2 Rate Prefill	rps	0.5
<P16>	P2 Rate HG	rps	0
<P17>	P2 Rate Post-run	rps	0.65

\* for 500 µl loop

\*\* Set to the approximate MAs peak maximum retention time.

\*\*\* (64s - (<P6> + <P7>)), to keep total heating time constant.

Table S2: FIAS program for HG-CT-ICP-MS Systems 2 and 3

Step	Time, s	Pump 1	Pump 2	Inj. Vent	Remote 5 (heating)	
P	30	0	100	Fill	-	Precooling of U tube
1	25*	40	100	Fill	-	Prefill of injection loop
2	60*	40	0	Inject	-	HG step
3	90	0	0	Inject	-	Liquid nitrogen Dewar is removed 7 sec before the end of Step 3
4	42**	0	0	Fill	ON	READ step; hydrides released from U-tube
5	7	0	0	Inject	-	Heating break to improve DMA <sub>s</sub> - TMA <sub>s</sub> <sup>V</sup> O resolution
6	22***	0	0	Fill	ON	
7	35	0	120	Inject	ON	U-tube drying
8	3	0	0	Fill	-	

\*for 500 µl loop

\*\* set to the approximate MAs peak retention time.

\*\*\* (64sec- Step4), to keep total heating time (steps 4 to 7) constant.



Table S3. Results of comparative BEC sample analyses.

Sample code	Cells per 300 ul	System 2 (UNC) pg As per 300 ul aliquot ± 95% confidence interval						System 1 (IAC) pg As per 300ul aliquot ± 95% confidence interval					
		iAs	MAs	DMAs	iAs	MAs	DMAs	iAs	MAs	DMAs	iAs	MAs	DMAs
904	710500	676 ± 21	180 ± 17	21 ± 8	811 ± 26	203 ± 3	24 ± 1	362 ± 22	45 ± 1	10 ± 1	72 ± 9	24 ± 2	188 ± 9
994	198000	345 ± 23	46 ± 11	11 ± 8	89 ± 9	10 ± 1	20 ± 1	1242 ± 76	26 ± 2	11 ± 1	138 ± 6	45 ± 2	44 ± 3
995	198000	60 ± 14	25 ± 17	192 ± 8	249 ± 6	103 ± 3	24 ± 1	645 ± 24	94 ± 2	172 ± 7	216 ± 13	95 ± 22	7 ± 1
1105	283200	60 ± 31	9 ± 28	17 ± 29	670 ± 9	149 ± 3	38 ± 2	216 ± 13	94 ± 2	172 ± 7	253 ± 19	103 ± 3	24 ± 1
1107	201600	1220 ± 80	29 ± 28	15 ± 29	231 ± 9	99 ± 22	7 ± 1	264600	39 ± 6	11 ± 1	139 ± 13	45 ± 2	44 ± 3
1167	88200	139 ± 13	40 ± 12	32 ± 11	39 ± 6	10 ± 1	20 ± 1	168300	10 ± 1	11 ± 1	127 ± 25	103 ± 3	24 ± 1
1168	514500	253 ± 19	127 ± 25	21 ± 8	1242 ± 76	26 ± 2	11 ± 1	167 ± 25	149 ± 3	38 ± 2	645 ± 24	149 ± 3	38 ± 2
1171	737450	645 ± 24	167 ± 25	40 ± 8	138 ± 6	45 ± 2	44 ± 3	176 ± 11	99 ± 22	7 ± 1	216 ± 13	95 ± 22	7 ± 1
1183	168300	216 ± 13	91 ± 11	176 ± 11	231 ± 9	94 ± 2	172 ± 7	12 ± 20	99 ± 22	7 ± 1	139 ± 13	45 ± 2	44 ± 3
1196	563500	92 ± 16	12 ± 20	5 ± 10	39 ± 6	10 ± 1	11 ± 1	10 ± 20	39 ± 6	11 ± 1	127 ± 25	103 ± 3	24 ± 1
1199	264600	56 ± 16	10 ± 20	9 ± 10	1242 ± 76	26 ± 2	11 ± 1	167 ± 25	149 ± 3	38 ± 2	645 ± 24	149 ± 3	38 ± 2

Fig. S1. Stability of the HG-CT-ICP-MS signals of a mixed standard  $200 \text{ pg mL}^{-1}$  iAs<sup>V</sup> and  $100 \text{ pg mL}^{-1}$  MAs<sup>V</sup> and DMA<sup>V</sup>, after cys treatment; B- blank. \* Injection error.

