Comparative oxidation state specific analysis of arsenic by high-performance liquid chromatography-inductively coupled plasma-mass spectrometry and hydride generationcryotrapping-atomic absorption spectrometry

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## SUPPLEMENTARY DATA

**Table S1:** The slopes of calibration curves generated by HG-CT-AAS (10-<sup>3</sup>s/ng) and HPLC-ICP-MS (cps\*s/pg) for As standards

	HG-CT-AAS <sup>a</sup>								
Solvents	iAs	II+V	MAs <sup>III+V</sup>		DMAs <sup>III+V</sup>				
DIW	81	.4	808		788				
	HPLC-ICP-MS <sup>b</sup>								
	iAs <sup>III</sup>	iAs <sup>v</sup>	MAs <sup>III</sup>	MAs <sup>V</sup>	DMAs <sup>III</sup>	DMAs <sup>V</sup>			
DIW	1503	1461	1152	1543	632	1425			
<b>Reaction Mixture</b>	1550	707	1145	1643	811	1486			

<sup>a</sup> Standards were prepared in DIW and were reduced with 2% cysteine prior to HG-CT-AAS analysis.

<sup>b</sup> Standards were prepared in DIW or in the *in vitro* reaction mixture containing Tris-HCl buffer, SAM and TCEP, but not AS3MT.

**Table S2:** Limits of detection (LOD) for the analysis of As species by HPLC-ICP-MS and HG-CT-AAS

	$LOD (pg mL^{-1})$									
Method	iAs <sup>III</sup>	<sup>I</sup> iAs <sup>V</sup>	iAs <sup>III+V</sup>	MAs <sup>III</sup>	MAs <sup>V</sup>	MAs <sup>III+V</sup>	DMAs <sup>III</sup>	DMAs <sup>V</sup>	DMAs <sup>III+V</sup>	
HPLC-ICP-MS	<sup>a</sup> 32	295		31	57		88	98		
HG-CT-AAS <sup>b</sup>	36	54	40	32	57	46	24	44	38	

<sup>a</sup>LODs were calculated as (3\*SD)/(slope\*0.02) for 10 injections of DIW (20 µl each).

<sup>b</sup>LODs were calculated as (3\*SD)/(slope\*0.5) for 10 injections of DIW or 2% cysteine in DIW (0.5 ml each).



Figure S1: The HPLC-ICP-MS profiles of As standards in DIW, including AsC, AsB, TMAO (A, 2,000 pg each) and iAs<sup>III</sup> (B, 280 pg). Detailed operating conditions are described in Table 1.



**Figure S2:** Comparison of the HPLC-ICP-MS profiles for As standards prepared in solutions containing the components of the *in vitro* methylation mixture: 0.5  $\mu$ M iAs<sup>III</sup> (A) and 0.5  $\mu$ M MAs<sup>III</sup> (B) in the 100 mM TRIS-HCl buffer (pH 7.4) in absence (—) and presence (---) of AS3MT (60  $\mu$ g/mL). 0.5  $\mu$ M DMAs<sup>III</sup> (C) in a mixture with TRIS-HCl buffer (—), 1 mM TCEP (—), AS3MT (60  $\mu$ g/mL; ---) or 1 mM SAM (---).