

Methylated arsenic species in plants originate from soil microorganisms

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Supplementary Materials

Table S1 Recovery and speciation of the certified reference material NIST 1568a rice flour

Total iAs ($\mu\text{g g}^{-1}$ DW)	MMA(V) ($\mu\text{g g}^{-1}$ DW)	DMA(V) ($\mu\text{g g}^{-1}$ DW)	Sum As ($\mu\text{g g}^{-1}$ DW)	Recovery (%)	Reference
0.09 ± 0.003	0.01 ± 0.001	0.18 ± 0.007	0.28 ± 0.010	96	This study
0.11 ± 0.006	0.02 ± 0.004	0.14 ± 0.005	0.27 ± 0.010	94	Batista <i>et al.</i> (2011)

Data are means \pm SD.

For a full list of previous analyses of As speciation in NIST 1568a, see Batista *et al.* (2011).

Batista BL, Souza JMO, De Souza SS, Barbosa Jr F. 2011. Speciation of arsenic in rice and estimation of daily intake of different arsenic species by Brazilians through rice consumption. *Journal of Hazardous Materials* **191**: 342-348.

Table S2 Relative abundance of *arsM* sequences detected in a Bangladeshi paddy soil using GeoChip.

Genbank ID	Organism	Mean relative abundance
158320129	<i>Alkaliphilus oremlandii</i> OhILAs	0.039
255008414	<i>Bacteroides fragilis</i> 3_1_12	0.024
167761029	<i>Clostridium scindens</i> ATCC 35704	0.055
254430241	<i>Cyanobium</i> sp. PCC 7001	0.217
57233796	<i>Dehalococcoides ethenogenes</i> 195	0.335
89896894	<i>Desulfitobacterium hafniense</i> Y51	0.300
158520779	<i>Desulfococcus oleovorans</i> Hxd3	0.305
257797304	<i>Desulfohalobium retbaense</i> DSM 5692*	1.814
258405077	<i>Desulfohalobium retbaense</i> DSM 5692*	1.733
258515753	<i>Desulfotomaculum acetoxidans</i> DSM 771	0.341
219869487	<i>Desulfovibrio desulfuricans</i> subsp. <i>desulfuricans</i> str. ATCC 27774	0.669
254448130	<i>Gamma proteobacterium</i> HTCC5015	0.424
168699430	<i>Gemmata obscuriglobus</i> UQM 2246	0.303
78223987	<i>Geobacter metallireducens</i> GS-15	0.364
284015962	<i>Haloterrigena turkmenica</i> DSM 5511	0.148
21226763	<i>Methanosarcina mazei</i> Go1	0.094
167461246	<i>Paenibacillus larvae</i> subsp. <i>larvae</i> BRL-230010	0.013
253575807	<i>Paenibacillus</i> sp. oral taxon 786 str. D14*	0.096
251844850	<i>Paenibacillus</i> sp. oral taxon 786 str. D14*	0.097
149277880	<i>Pedobacter</i> sp. BAL39	0.150
147676775	<i>Pelotomaculum thermopropionicum</i> SI	0.634
211959444	<i>Pseudovibrio</i> sp. JE062	0.590
254510744	<i>Rhodobacteraceae bacterium</i> KLH11	0.108
283559495	<i>Rhodomicrobium vannielii</i> ATCC 17100	0.269
283843001	<i>Rhodopseudomonas palustris</i> DX-1	0.421
83815482	<i>Salinibacter ruber</i> DSM 13855	0.640
116623500	<i>Solibacter usitatus</i> Ellin6076	0.839
269838062	<i>Sphaerobacter thermophilus</i> DSM 20745	0.553
51893507	<i>Symbiobacterium thermophilum</i> IAM 14863	0.507
269792113	<i>Thermanaerovibrio acidaminovorans</i> DSM 6589	0.596
219994748	<i>Thioalkalivibrio</i> sp. HL-EbGR7	0.498
220933438	<i>Thioalkalivibrio</i> sp. HL-EbGR7	0.330
74317446	<i>Thiobacillus denitrificans</i> ATCC 25259	0.445
257456586	<i>Treponema vincentii</i> ATCC 35580	0.637
269306796	<i>Xylanimonas cellulosilytica</i> DSM 15894	0.610

* Duplicate organisms but different GenBank sequences.

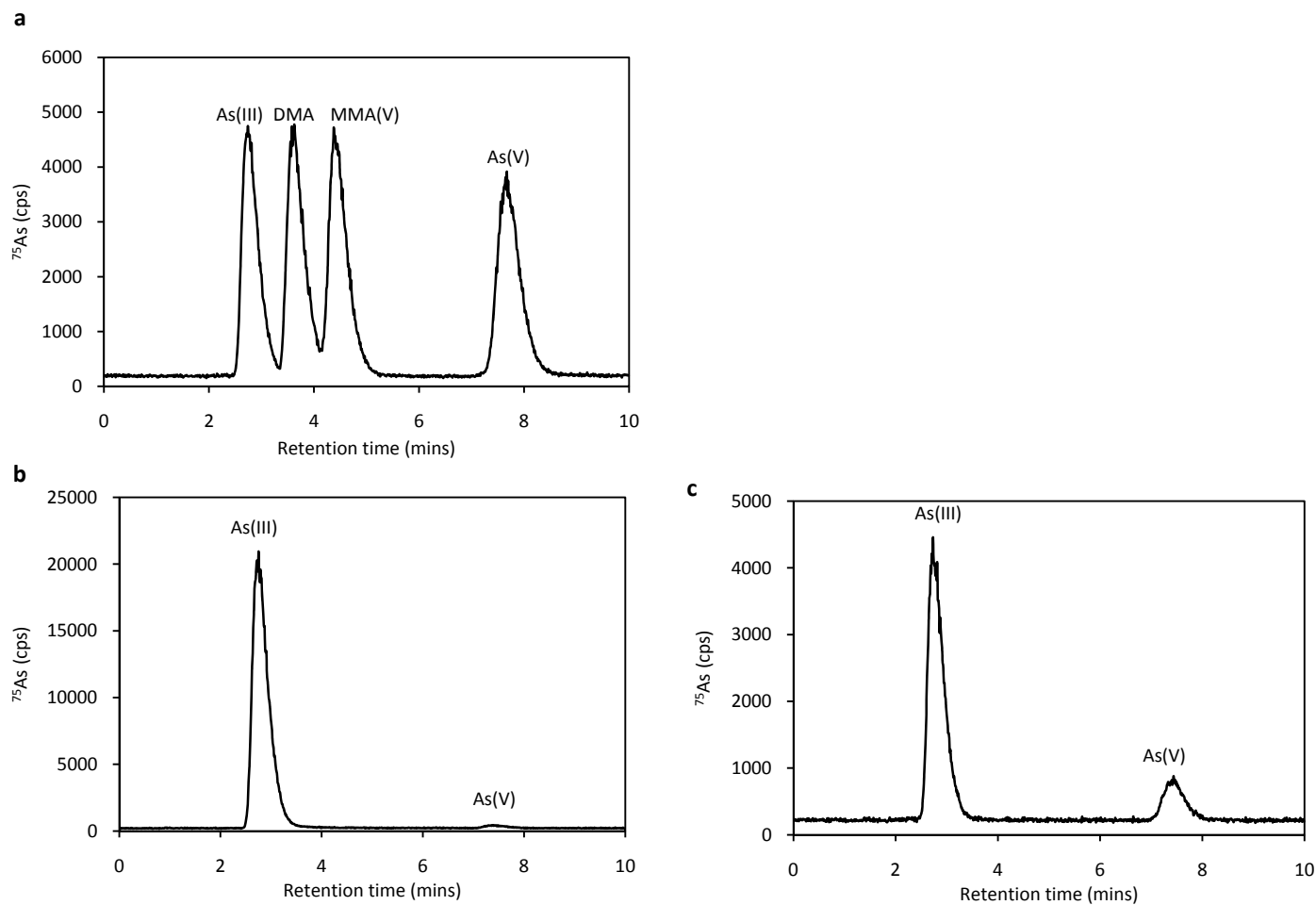


Fig. S1 Lack of methylated As species in axenically-grown plant extracts. **a**, Chromatograms of standards containing $50 \mu\text{g As L}^{-1}$ arsenite, DMA(V), MMA(V) and arsenate by HPLC-ICP-MS. Representative chromatograms of As speciation in extracts of rice (cv. Nipponbare) roots (**b**) and shoots (**c**) exposed to $10 \mu\text{M}$ arsenate. cps = counts per second.

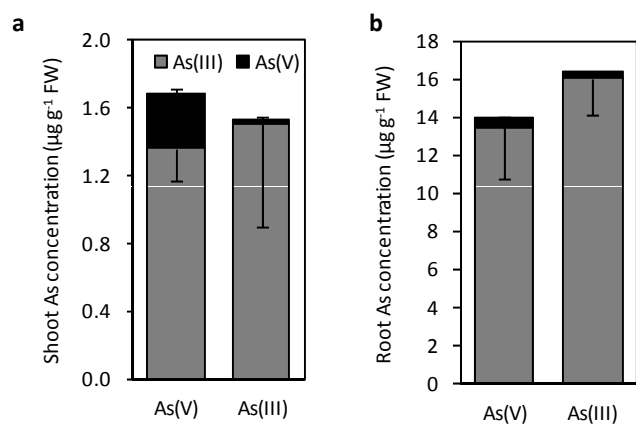


Fig. S2 Arsenic speciation in rice exposed to arsenate or arsenite in axenic culture. Arsenic speciation in rice (cv. Nipponbare) shoots (**a**) and roots (**b**) after exposure to 10 µM arsenate or arsenite. Error bars represent SE ($n = 5$).

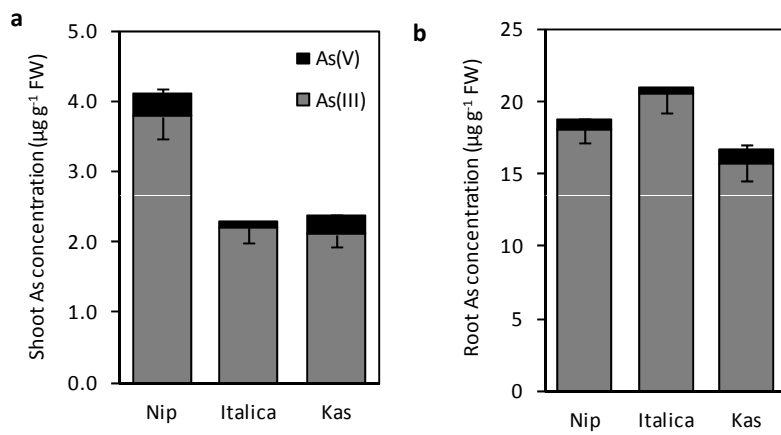


Fig. S3 Arsenic speciation in different rice cultivars exposed to arsenate in axenic culture. Arsenic speciation in shoots (**a**) and roots (**b**) of 3 different rice cultivars; Nip = Nipponbare (japonica), Italica = Italica carolina (japonica) and Kas = Kasalath (indica), after exposure to 10 μM arsenate. Error bars represent SE ($n=5$).

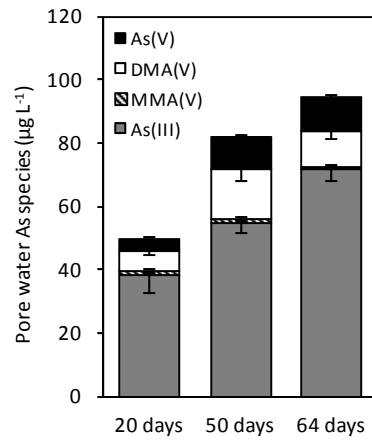


Fig. S4 Arsenic speciation in pore water of incubated UK arable soil. Pore water was collected at 20, 50 and 64 days after flooding. Error bars represent SE ($n=5$).

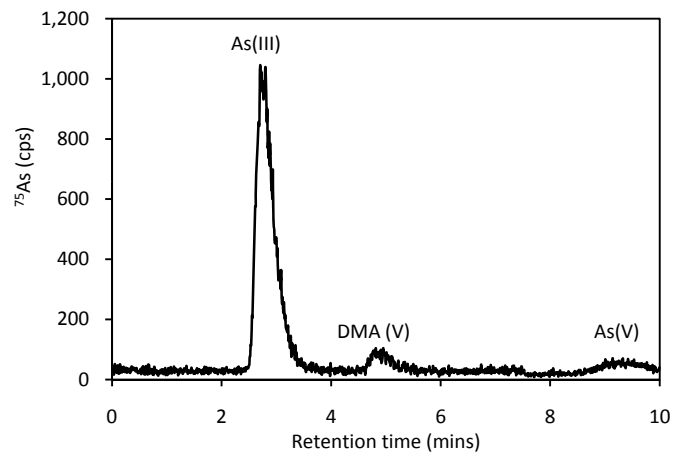


Fig. S5 Detection of methylated As in nutrient solution amended with arsenite. Representative chromatogram of As speciation of nutrient solution spiked with 10 μM arsenite at the end of the 7 day exposure period.