

Supporting Information for MS:

A new role for sulfur in arsenic cycling

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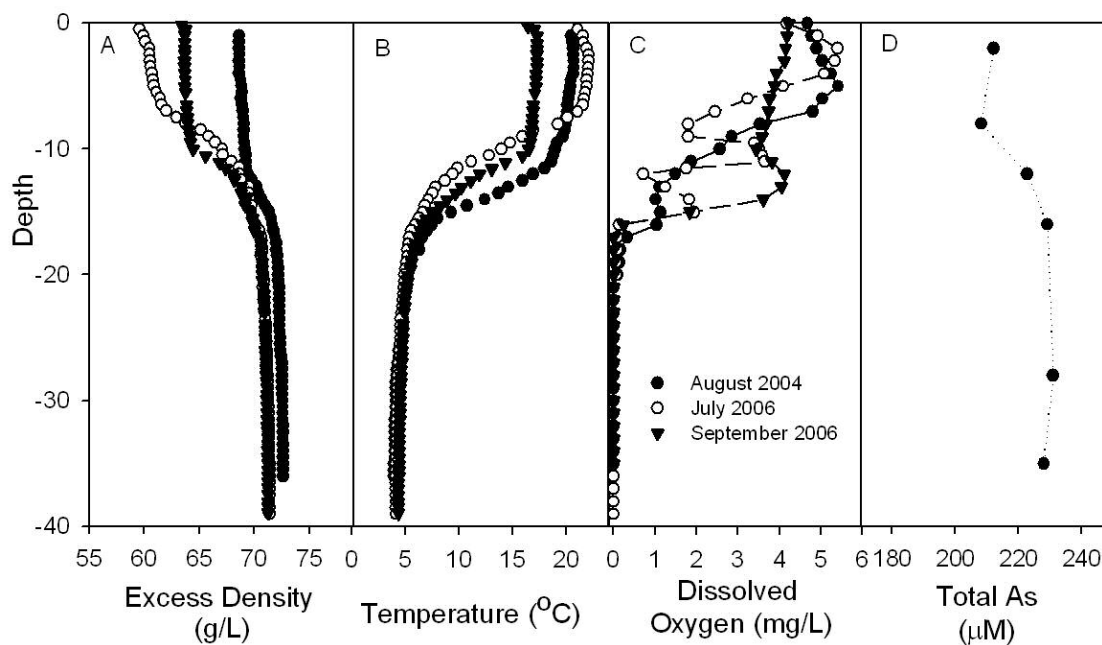
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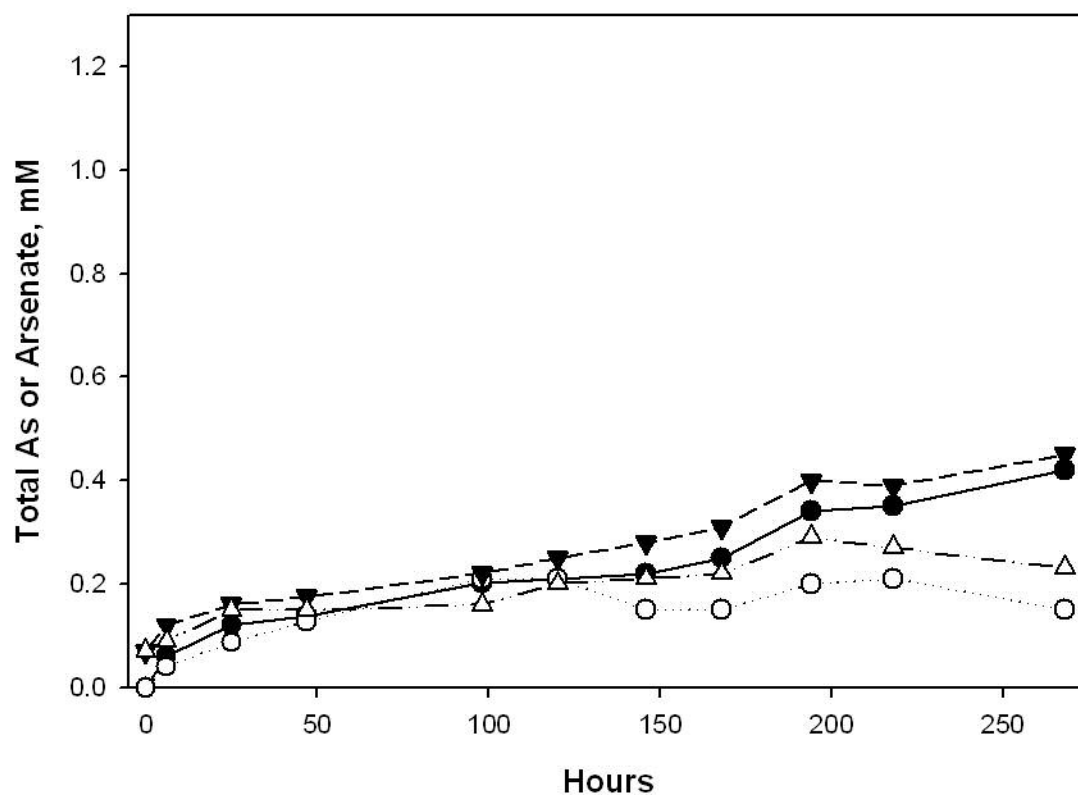
Number of pages: 10

Number of tables: 1

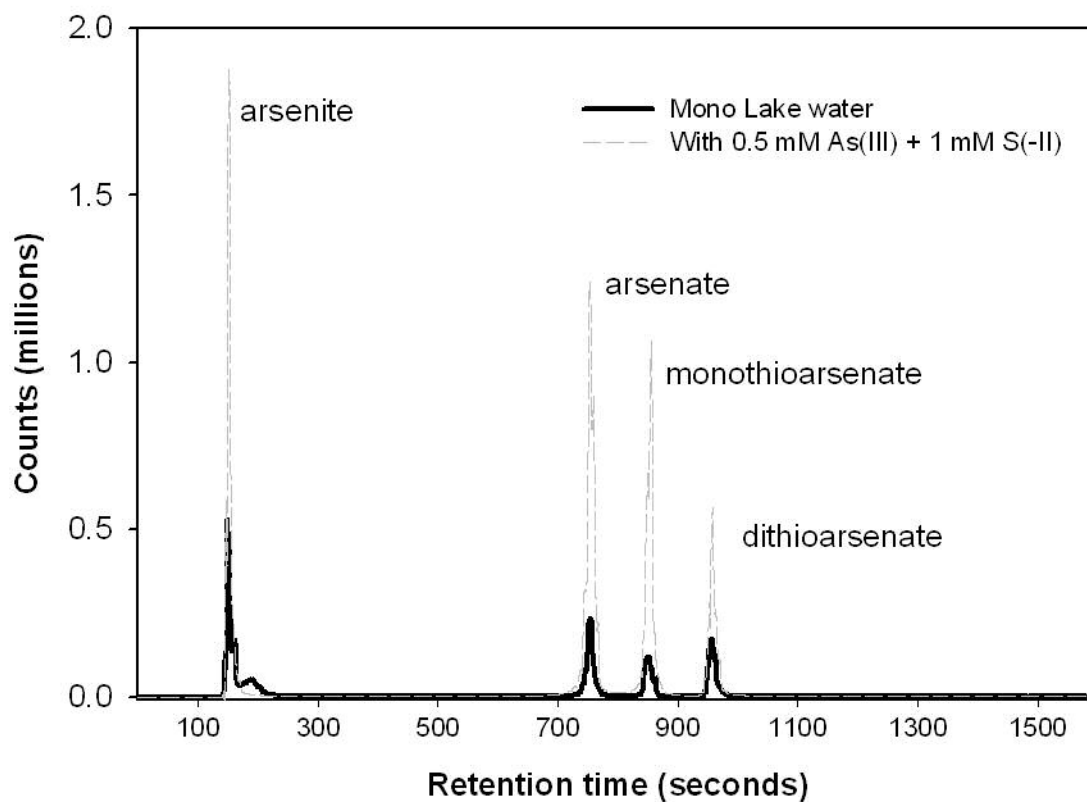
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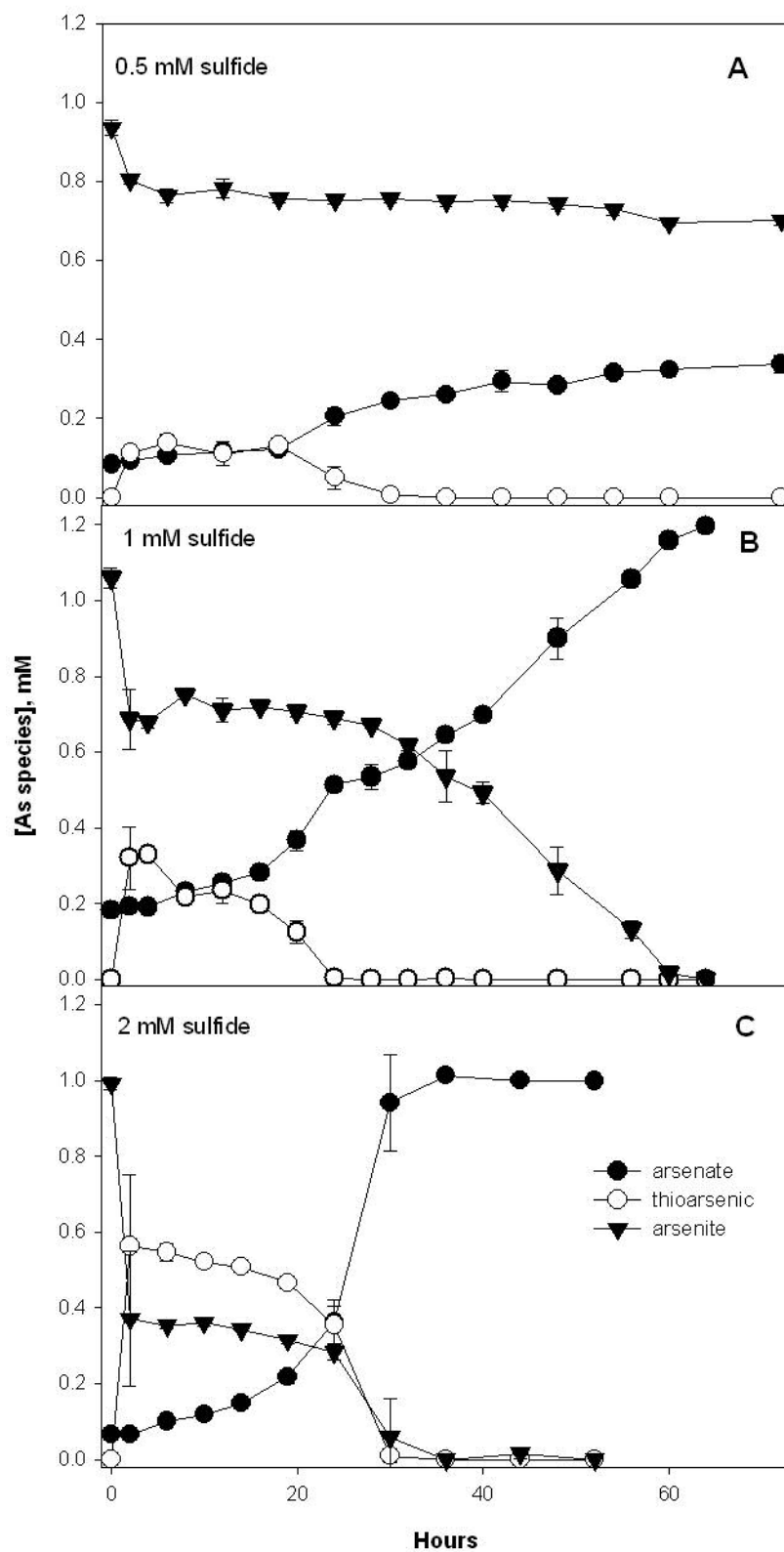
Supporting Figure S1. CTD measurements (excess density and temperature, A and B) and dissolved oxygen concentrations (C) from CTD casts taken in August 2004 (●), July 2006 (○), and September 2006 (▼). A set of samples from August 2004 was analyzed for total arsenic (●, D).

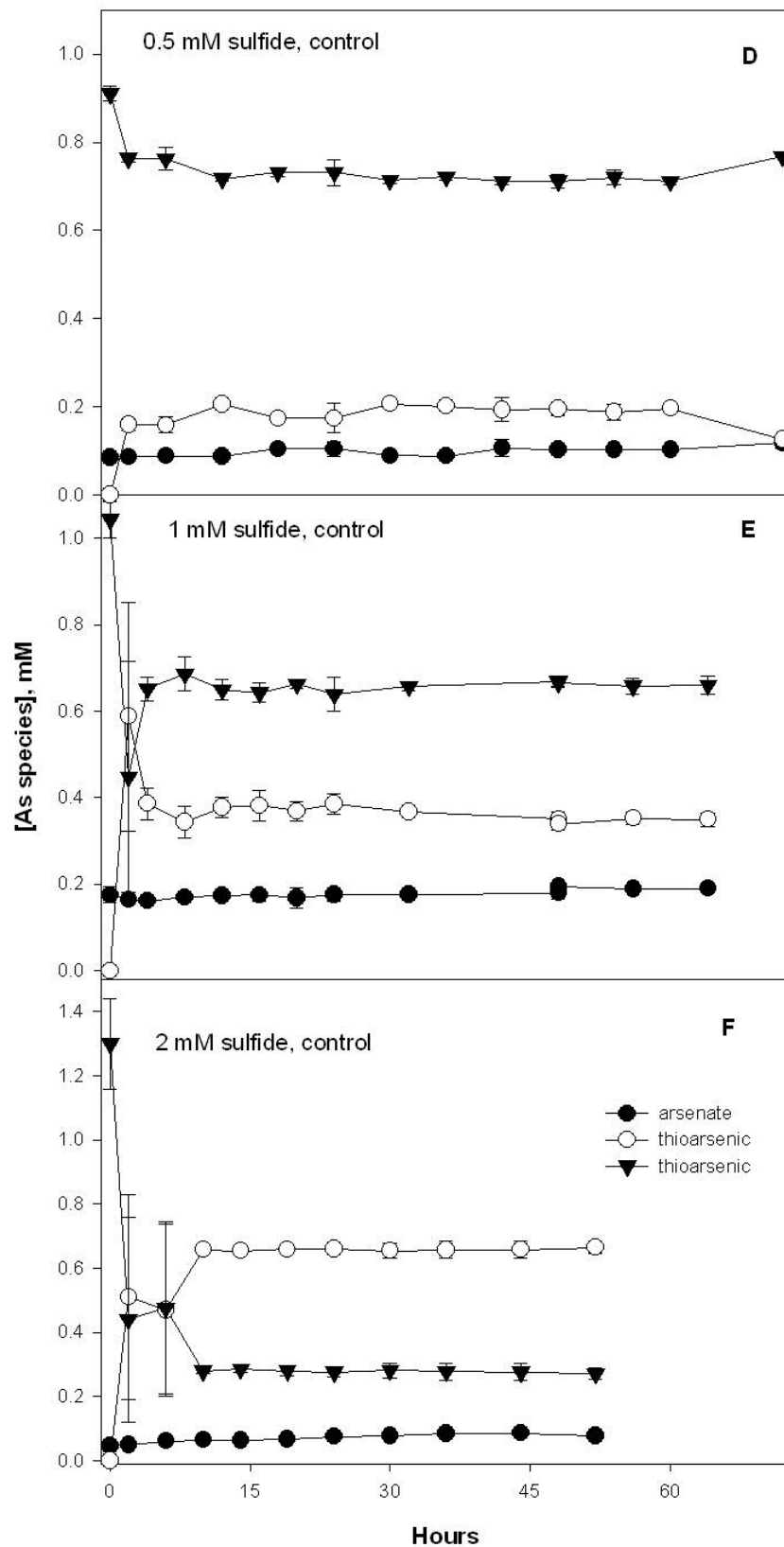


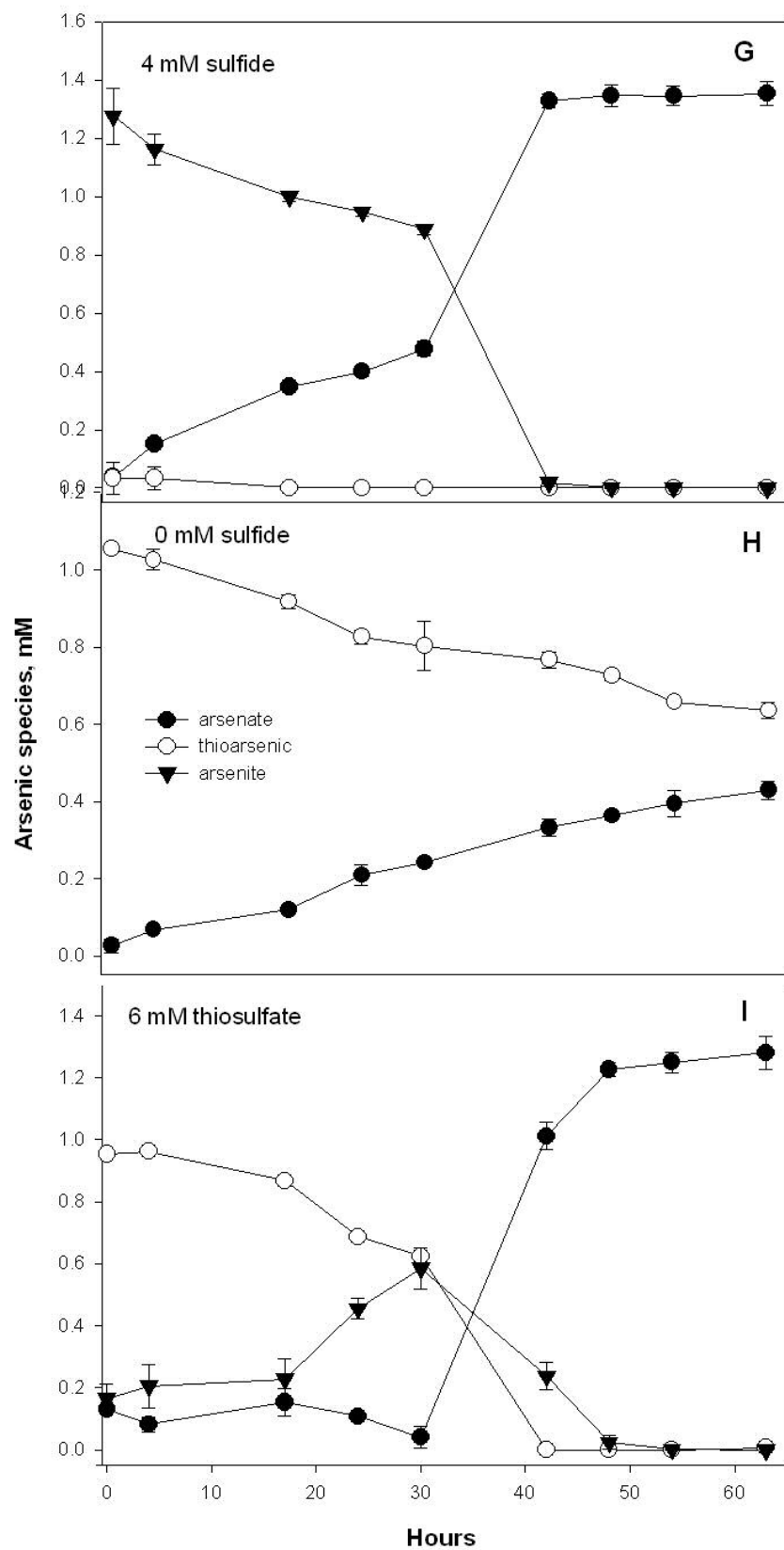
Supporting Figure S2. Comparison of arsenate production in Mono Lake water (MLW) or artificial Mono Lake water (AMLW) medium controls amended with 2 mM sulfide and 1 mM arsenite with or without formalin. Symbols: MLW, filtered (Δ); MLW with formalin (\blacktriangledown); AMLW, filtered (\circ); AMLW with formalin (\bullet).



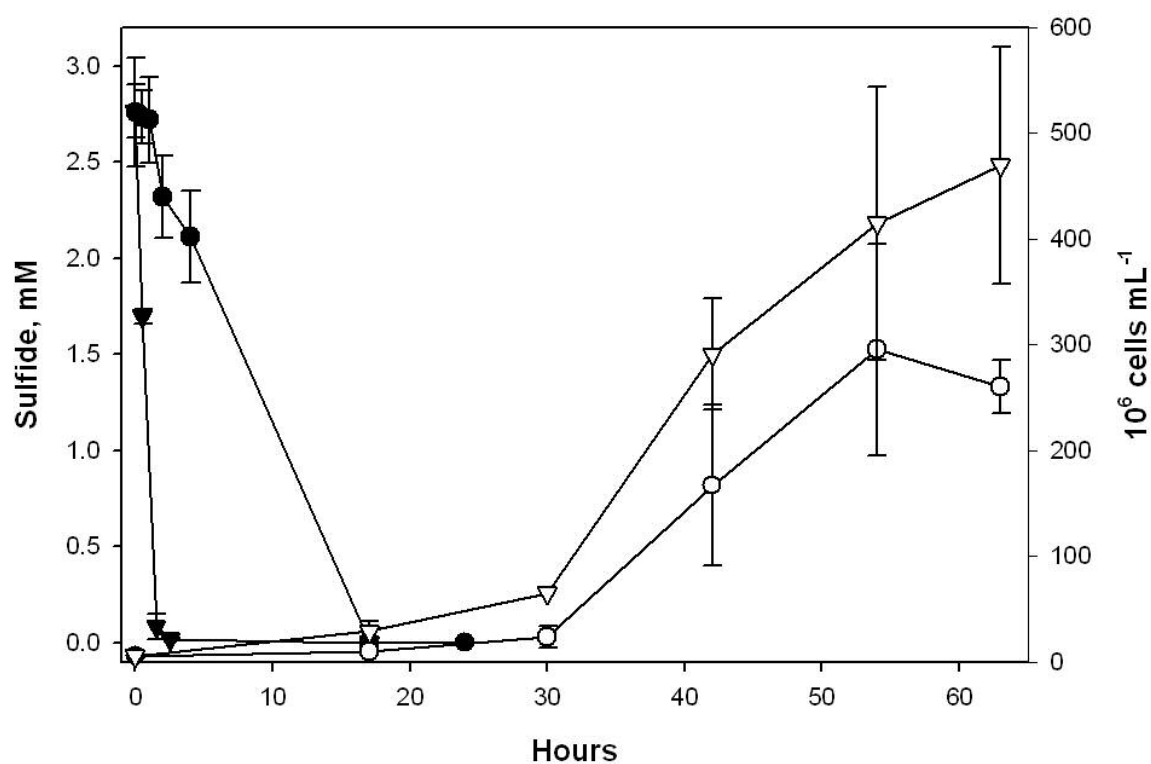
Supporting Figure S3. IC-ICP-MS chromatograms of an anoxic Mono Lake water sample (35m, September 2006; —) and the first sample from the second simulated mixing experiment (---).







Supporting Figure S4. Arsenic speciation in enrichment culture experiments. Arsenate (●) and arsenite (▼) were measured by HPLC and thioarsenic (○) was determined by difference. Artificial Mono Lake water was amended with 1 mM arsenite and A) 0.5 mM sulfide and live inoculum, B) 1 mM sulfide and live inoculum, C) 2 mM sulfide and live inoculum, D) 0.5 mM sulfide (sterile control), E) 1 mM sulfide (sterile control), F) 2 mM sulfide (sterile control), G) 4 mM sulfide, H) no sulfide, or I) 6 mM thiosulfate.



Supporting Figure S5. Sulfide was depleted rapidly (<24 hours) in the 4 mM sulfide + 1 mM arsenite treatment (As+S, ▼) and in the 4 mM sulfide only treatment (S, ●). Cell density in the As+S (Δ) and S (○) treatments increased significantly after sulfide was not detected, likely due to oxidation of polysulfides (S₅²⁻), thiosulfate, or sulfite.

Experiments were conducted in artificial Mono Lake water with a consortium of sulfoxidizing bacteria.

Supporting Table S1. Description of clones from an arsenite and sulfide enrichment culture.

Sequence (Accession number)	Closest sequence match (% similarity)	Closest cultured relative (% similarity)	Number of clones represented by sequence
MLAsHS-clone506-1 (EF643362)	Uncultured gamma proteobacterium clone ML615J-17 (99%)	<i>Thioalkalivibrio jannaschii</i> (98%)	5
MLAsHS-clone506-2 (EF643363)	Uncultured gamma proteobacterium clone ML615J-17 (99%)	<i>Thioalkalivibrio versutus</i> (97%)	2
MLAsHS-clone506-3 (EF643364)	<i>Idiomarina</i> sp. G- R2A15 (99%)	<i>Idiomarina salinae</i> strain ISL-52 (96%)	1
MLAsHS-clone506-8 (EF643369)	Uncultured gamma proteobacterium clone ML623J-18 (96%)	<i>Thioalkalivibrio nitratis</i> (95%)	1
MLAsHS-clone506-10 (EF643371)	<i>Arhodomonas</i> sp. EL- 201 (98%)	<i>Saccharospirillum impatiens</i> (98%)	1