1	Engineering the soil bacterium <i>Pseudomonas putida</i> for arsenic removal
2	
3	Jian Chen <sup>1,2</sup> , Jie Qin <sup>1</sup> , Yong-Guan Zhu <sup>2,3</sup> , Víctor de Lorenzo <sup>4</sup> and Barry P. Rosen <sup>1*</sup>
4	<sup>1</sup> Department of Cellular Biology and Pharmacology, Florida International University, Herbert
5	Wertheim College of Medicine, Miami, FL 33199, USA
6	<sup>2</sup> State Key Lab of Regional and Urban Ecology, Research Center for Eco-environmental
7	Sciences, Chinese Academy of Sciences, Beijing 100085, China
8	<sup>3</sup> Key Lab of Urban Environment and Health, Institute of Urban Environment, Chinese
9	Academy of Sciences, Xiamen 361021, People's Republic of China
10	<sup>4</sup> Centro Nacional de Biotecnología-CSIC, Campus UAM-Cantoblanco, Madrid 28049, Spain
11	
12	*Correspondence: Barry P. Rosen, Florida International University Herbert Wertheim College
13	of Medicine, 11200 SW 8th Street, Miami, FL 33199 Tel: (+1) 305-348-0657, Fax: (+1)

14 305-348-0651, Email: brosen@fiu.edu

15 **Supplemental materials**:

16

Abbreviations: polymerase chain reaction, PCR; sodium dodecyl sulfate polyacrylamide
 gel electrophoresis (SDS-PAGE); high pressure liquid chromatography (HPLC); inductively
 coupled plasma mass spectroscopy (ICP-MS).

20

## 21 Supplemental Figure Legends:

Supplemental Fig. 1S. Methylation of As(III) by purified CrArsM. <u>A</u>: Transformation of 10 µM As(III) into soluble methylated species was determined by high pressure liquid chromatography (HPLC) coupled to inductively coupled mass spectroscopy (ICP-MS) using a C18 reverse phase column after incubation for various times with 1 µM purified CrArsM. <u>Curve 1</u>, standards. <u>Curve 2</u>: 0 time; <u>Curve 3</u>: 7 h; <u>Curve 4</u>: 14 h. <u>B</u>: Production of volatile species determined by anion exchange HPLC-ICP-MS analysis. <u>Curve 1</u>, 0 time; <u>Curve 2</u>: 7 h; <u>Curve 3</u>: 14 h.

29

Supplemental Fig. 2S. Construction of plasmid pBAM1-P<sub>km</sub>-CrArsM. The gene for CrArsM 30 with a C-terminal six-histidine tag was cloned from a *C* reinhardtii cDNA library using forward 31 primer arsM-F 5'-GGAATTCCATATGGTGGAGCCGGCTTCC ATCGCGGAGCTT-3' (Ndel 32 underlined) 5'-CCG<u>CTCGAG</u>TTAATG 33 site and reverse primer arsM-R 34 ATGATGATGATGATGGCAGCAGGCGCCGCCGGGG-3' (Xhol site underlined). The polymerase chain reaction (PCR) fragment was gel purified and cloned into vector plasmid 35 pET28a vector with the indicated restriction enzymes, generating plasmid pET28a-CrArsM. 36

37 The kanamycin promoter was cloned from pBAM1 vector using forward primer 5'-GAAGATCTTGTCTC TTATACACATCTGACG-3' (*Bgl*II site underlined) and reverse primer 38 5'-GGAATTCCATATGAA CACCCCT TGTATTAC-3' (*Nde*l site underlined). The T7 promoter 39 in pET28a-CrArsM was replaced with the kanamycin promoter, generating plasmid 40 41 pET28a-P<sub>km</sub>-CrArsM. Finally, a fragment contained the kanamycin promoter and arsM was 42 excised by digestion with Bg/II and XhoI and ligated into vector plasmid pBAMI vector that had been digested with BamHI and Sall, generating plasmid pBAM1-P<sub>km</sub>-CrArsM. 43 Functional elements of the plasmid include relevant restriction sites, transposase (tnpA), 44 45 origin of replication (R6K), the origin of transfer region (oriT), mosaic element O (ME-O), and mosaic element I (ME-I), as shown. 46

47

48 Supplemental Fig. 3S. Purification of CrArsM from *E. coli*. A culture of *E. coli* BL21(DE3) pET28a-CrArsM was induced overnight with 0.3 mM isopropyl β-D-1-thiogalactopyranoside 49 and lysed by a single passage through a French pressure cell. Following high speed 50 centrifugation, the cytoplasmic solution was chromatographed on a Ni-NTA column, and the 51 fractions analyzed by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) 52 53 stained with Coomassie Blue. Lane 1: Standard protein markers. Lane 2-8: Successive column fractions eluted with an imidazole gradient. Lane 9, Cells of E. coli BL21(DE3) 54 pET28a-CrArsM without induction. Lane 10, Cells induced with 0.3 mM isopropyl 55 β-D-1-thiogalactopyranoside. 56

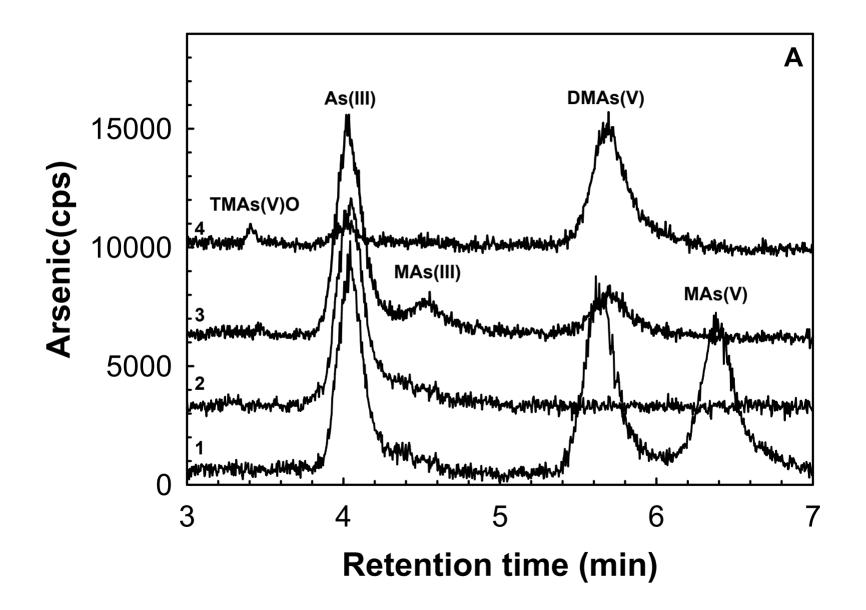
57

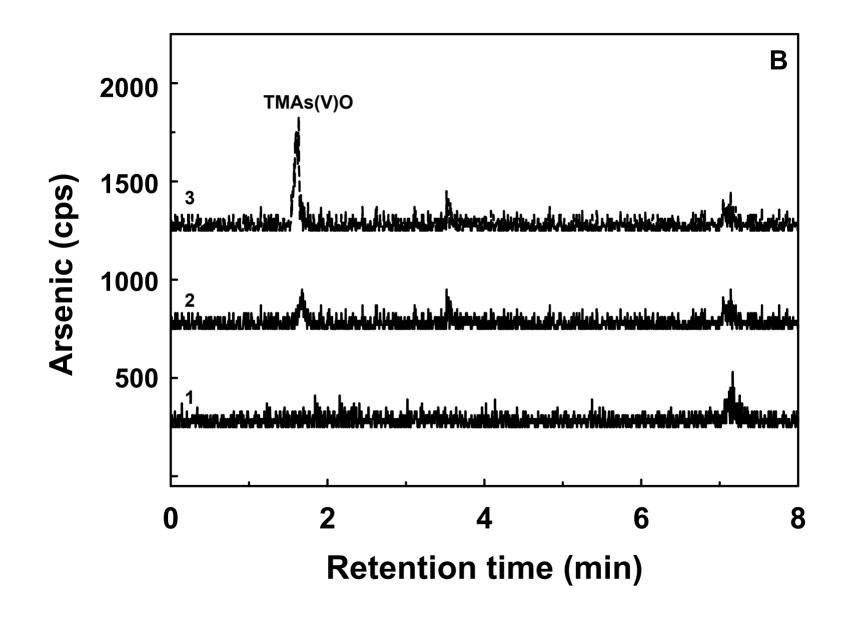
58 Supplemental Fig. 4S. Immunoblot detection of CrArsM. Cultures of both E. coli

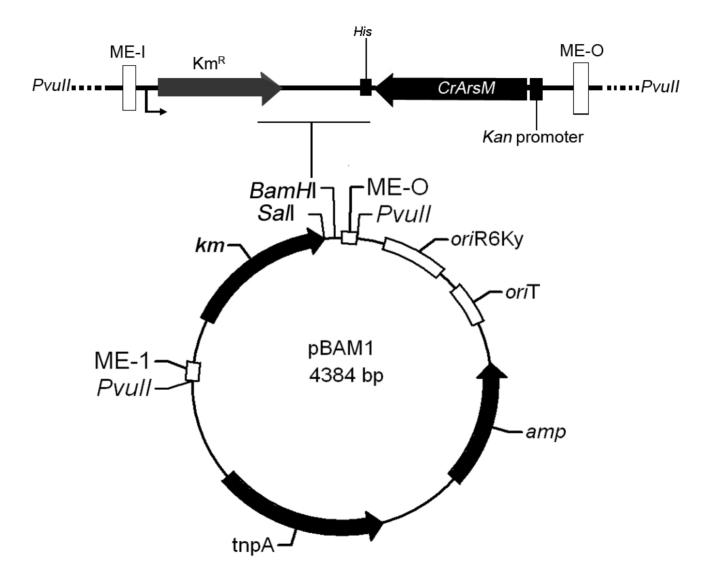
3

BL21(DE3) pET28a-CrArsM induced with 0.3 mM isopropyl β-D-1-thiogalactopyranoside and 59 P. putida pBAM1-Pkm-CrArsM were grown for 6 h in LB medium, centrifuged and suspended 60 61 in 1/10 the original volume of SDS sample gel buffer and boiled for 5 m. Portions (5 µl) were separated by SDS-PAGE on a 12% polyacrylamide gel, transferred to a nitrocellulose 62 membrane and electrophoresed at 25 V overnight. After transfer, the membrane was 63 blocked with 5% nonfat dried milk in physiological saline buffer for 4 hrs with shaking. The 64 membrane was incubated with the primary His-tag monoclonal antibody (anti-mouse cat. No. 65 70796-3 diluted 1:1000 (Novagen-EMD Millipore) in 5% blocking buffer for 1 hrs at room 66 67 temperature. The membrane was washed four times with the same buffer containing 0.05% Tween 20 for 25 min for each time. The membrane was then incubated with secondary 68 antibody (peroxidase-coupled anti-mouse IgG, Sigma Chemical Co.) diluted 1:5000 for 1 hrs 69 70 at room temperature and washed with the same buffer 4 times for 25 min each time. The proteins were detected using a Western Lightning<sup>™</sup> Ultra chemiluminescent substrate kit 71 (ECL Kit, 2 ml/membrane). In a separate tube, the solutions were mixed in a 1:1 ratio and 72 poured onto the membrane for 1 min before exposure. 73

Lane 1, Standard protein markers. Lane 2, vector plasmid pET28a in BL21(DE3), Lane 3,
vector plasmid pBAM1 in *P. putida*. Lane 4, ars*M* in plasmid pET28a expressed in BL21(DE3)
with IPTG. Lane 5, ars*M* chromosomally expressed in *P. putida*.







-



