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3 **A novel MAs(III)-selective ArsR transcriptional repressor**

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9 **Supplemental Materials**

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Table S1: Oligonucleotide primers used in this study

Primer	Sequence	
pBAD- <i>SparsR</i> :		
SparsRF:	5'-CATG <u>CCATGGT</u> GAAACATAGCCGATATGAAT-3'	(<i>Nco</i> I site underlined)
SparsRR:	5'-GCGT <u>CGACCT</u> GGCCACTACAACATTGATCC-3'	(<i>Sal</i> I site underlined)
pACYC184- <i>ParsP-gfp</i> :		
ParsPF:	5'-CG <u>GGATCC</u> ATTGGCCTCATCTTATTAAG-3'	(<i>Bam</i> HI site underlined)
ParsPR:	5'-GCGT <u>CGACT</u> TTTTACCTCTGTGGTATTTATA-3'	(<i>Sal</i> I site underlined)
pBAD- <i>SpArsR_{AC}</i> :		
SparsRF:	5'-CATG <u>CCATGGT</u> GAAACATAGCCGATATGAAT-3'	(<i>Nco</i> I site underlined)
HSpArsR ₁₀₂ R:	5'-CCCA <u>AAGCTT</u> ACAACATTGATCCTGTAAAAAAT-3'	(<i>Hind</i> III site underlined)
HAfArsR ₉₇ F:	5'-CCCA <u>AAGCTT</u> ACCCGGGACTGTGCCCTATCCG-3,	(<i>Hind</i> III site underlined)
HAfArsR ₉₇ R:	5'-ATGATGATGATGATGATG <u>GTCGACCT</u> GGT-3'	(<i>Sal</i> I site underlined)
Cysteine residue mutagenesis		
C83S forward	5'-AAGGAAGGGTGTGTAC <u>AGCGT</u> GCCTGATTATGAG-3'	
C83S reverse	5'-CTCATAATCAGGCAC <u>GCT</u> GTACAACACCCTTCCTT-3'	
C101S forward	5'-GCATTTTTTACAGGATCAA <u>AGTT</u> GTAGTGGCCAGCATCA-3'	
C101S reverse	5'-TGATGCTGGCCACTACA <u>ACT</u> TTGATCCTGTAAAAAATGC-3'	
C102S forward	5'-CATTTTTTACAGGATCAATGT <u>AGT</u> AGTGGCCAGCATCATCATC-3'	
C102S reverse	5'-GATGATGATGCTGGCCACT <u>ACT</u> TACATTGATCCTGTAAAAAATG-3'	
Reverse transcription polymerase chain reaction (RT-PCR)		
ArsP forward	5'-GTGGGCATTCTACAAGCCTTTATCA-3'	579bp
ArsP reverse	5'-TAATGAACCGAGTAAAACACCCACA-3'	
ArsR forward	5'-AATGTAGAGAATGCAGCTAAGGTAT-3'	266bp
ArsR reverse	5'-TGATCCTGTAAAAAATGCACTAACC-3'	
ArsPR forward	5'-GATAAAGGCTTGTAGAATGCCCACT-3'	644bp
ArsPR reverse	5'-TGATCCTGTAAAAAATGCACTAACC-3'	

11 **Legends to the Supplemental Figures:**

12

13 **Supplemental Fig. S1. Multiple alignment of SpArsR orthologs.** Representative ArsR
14 orthologs (accession numbers in parentheses) are from: *A. ferrooxidans* (ACK80311),
15 *Ralstonia solanacearum* (NP_522690), *Caulobacter crescentus* (NP_420316),
16 *Mesorhizobium loti* (NP_103579), *Sinorhizobium meliloti* 1021 (NP_385183),
17 *Agrobacterium fabrum* (NP_354498), *Rhodopseudomonas palustris* (NP_947601),
18 *Desulfospira joergensenii* (WP_027362847), *Serratia liquefaciens* FK01 (GAK27386),
19 *Endozoicomonas elysicola* (WP_020582082), *Enterovibrio calviensis* (WP_017007765),
20 *Psychromonas ossibalaenae* (WP_026339319), *Glaciecola agarilytica* NO2
21 (GAC07582.1), *Photobacterium* sp. AK15 (ELR66560), *Moritella marina*
22 (WP_019439555) and *S. putrefaciens* (ADV53698). The multiple alignment was
23 calculated with CLUSTAL W.

24

Fig. S1

Acidithiobacillus 1 -----MEPLQDPAQIVARLEALASPVRLIEIFRLLVVEQEPTGLV
 Ralstonia 1 -----MEETDAIRSLAALAHSLRLRVFRMLVVGAPAGLT
 Caulobacter 1 -----MLSALSHEGRLAIFRLLVQAGPAGVA
 Mesorhizobium 1 -----MDKATALQALAALGQETRLVFRLLVRAGAKGVP
 Sinorhizobium 1 -----MDTVDIIAFAALAQPTRLDAFRFLVKHEPDGLP
 Agrobacterium 1 -----MDNIGAIAAISALAQTTTLETFRLLVQHEPEGIP
 Rhodopseudomonas 1 -----MDNDSAIASIGALAQGTTRLDVFRLLVRHEPDGLA
 Desulfospira 1 -----MENETAAKKLAELGHTTTRLSIFRYLVKVGQCAS
 Serratia 1 -----MELNEVANAKKELGHPTRLSIYKQLVRAGHGGLP
 Endozoicomonas 1 -----MNIDDYAKTLKELGHPVRLSIYKRLVKAGRSVIP
 Enterovibrio 1 -----MELDAVAKAKKELGHPTRLTIFRRLVRAGHNGIA
 Psychromonas 1 -----MELEVIAKAKKELGHPTRLAIFKRLVKSGFQOGIA
 Glacielcola 1 -----MDIENTAKAKKELGHPTRLNIIYRSVVRAGYQOGIA
 Photobacterium 1 -----MDIEIVAKAKKELGHPTRLAIFKSVVRAGYQOGIA
 Moritella 1 -----MDIDIIAKAKKELGHPTRLTIFKSIIRAGYQOGIA
 Shewanella 1 -----MNIADMNVADMNVENAAKVLKELGHPTRIALFRLLVKGGYTQVA

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Acidithiobacillus 39 SCDIAEHLGQPHNGISFHLKNIQHAGLVTVQREGRYQRYRAAMPVVRALVAYLTENCCHG
 Ralstonia 35 PCATAEQLDVPGATLSFHLKELMNAALVTQERDGRHLIYRAAFDHMNGLLGFLTENCQOG
 Caulobacter 27 AGETARALNVLPNLSLSANLNVLSHSGLIASRRDGRSIIYTYADYDAMTGLLGFLMEDCCAG
 Mesorhizobium 35 AGETATRLSTVQNTMSAHLKILGQAGLVHAERDGRSIRYADMTGFRDLLAYLMEDCCNG
 Sinorhizobium 35 AGETARLINVPHNTMSTHTAQLQRAGLVTTTRRQSRSSIIYRADLDGLRHVVSFILKDCAG
 Agrobacterium 35 AGETARLTHVPQNTMSAHLATLARAGLVKSERQSRSSIIYRADLEGLRALTLFLLKDCGG
 Rhodopseudomonas 35 AGETARQLDVPQNTMSAHLGILARAGIVRSEHSRSIIYRADLDGLRALTLFLVKDCAG
 Desulfospira 35 VGGIQQKLGIPGSTLSHHLRRLVAAGLTKQVRESRVLYCIPQFDVLDRLLEFLKSECCTE
 Serratia 35 VGGIQQKRLDIPNSTLSHHLAALLSVGLVHQQREGRTLFCPLQYDVLNGIVAFLEIECCAE
 Endozoicomonas 35 VGGIQQKELDIPGSTLSHHLAALLSVGLVQQIRDGRLLYCIPQFQVFDQLLDFLRDQCCVD
 Enterovibrio 35 VGGVQDDLSIPGSTLSHHLSSLMSAGLITQRREGRVLYCVPQFEQLVAVLAFLODECCVD
 Psychromonas 35 VGGVQEEAIPGSTLSHHLISGLVSAAGLVTQRREGRVLYCVAEYKKLESVITFLQDECCAD
 Glacielcola 35 VGGIQQKLEIPGSTLSHHLISGLLSAGLITQRRESRTLFCVAEYDKLQSVIDFLQEECCAD
 Photobacterium 35 VGGIQQKLGIPGSTLSHHLISGLLSAGLISQRREGRRTLFCVAEYDKLQAVIGFLQDECCLD
 Moritella 35 VGGIQQQLAIPGSTLSHHLISGLLSAGLISQRREGRTLFCVAEYDKLKAVIGFLQDECCID
 Shewanella 45 VGGIQQEALQIPGSTLSHHLISALMSAGIISQRREGRVLYCVPDYELQGLVHFLQDQCCSG

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Acidithiobacillus 99 -TRDCALSGET----RSPSVQEGNQ-----
 Ralstonia 95 EA--CPEFSAA----GCKC-----
 Caulobacter 87 APQICAPLSAI----VSGGMACGAACA-----
 Mesorhizobium 95 APELCQPILQA----VTCKC-----
 Sinorhizobium 95 HPDVCAPLVAD----LTPCCSPMDRPARSHRSEGGGMDGEAADRKN
 Agrobacterium 95 ATELCAPLIAE----LTPCCQAEAL-----
 Rhodopseudomonas 95 APELCAPLLAE----LTPCC-----
 Desulfospira 95 AKCKM-----
 Serratia 95 QQPHC-----
 Endozoicomonas 95 SGVSSF-----
 Enterovibrio 95 ESSAS-----
 Psychromonas 95 MEDIG-----
 Glacielcola 95 EKT-----
 Photobacterium 95 EQASDDM-----
 Moritella 95 EVEFSATASNK-----
 Shewanella 105 Q-----

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27 **Supplemental Fig. S2. Response of the bacterial biosensor with SpArsR to As(III)**

28 **and MAs(III). A)** Diagrammatic representation of the construction of the bacterial

29 biosensor with SpArsR. In pBAD-*SparsR*, *SparsR* is under control of the *ara* promoter. In

30 pACYC184-*ParsP-gfp*, the *gfp* reporter is under control of the *arsP* promoter. **B)**

31 Conditions for constitutive, repressed or derepressed *gfp* expression. In cells of *E. coli*

32 AW3110 with both plasmids, *SparsR* is not expressed in the absence of arabinose, and

33 *gfp* expression is constitutive, producing cellular fluorescence. In the presence of

34 arabinose, *SparsR* is expressed, and *gfp* is repressed, so the cells are not fluorescent.

35 In the presence of both arabinose and arsenical inducer, *gfp* expression is derepressed,

36 and the cells are fluorescent. **C)** The bacterial biosensor with *SparsR* responds only to

37 MAs(III). Expression of the *gfp* reporter gene was assayed as described under

38 Experimental Procedures. Cells of *E. coli* strain AW3110(DE3) bearing wild type *SparsR*

39 were grown without arabinose, 0.2% arabinose, or 0.2% arabinose and arsenicals at the

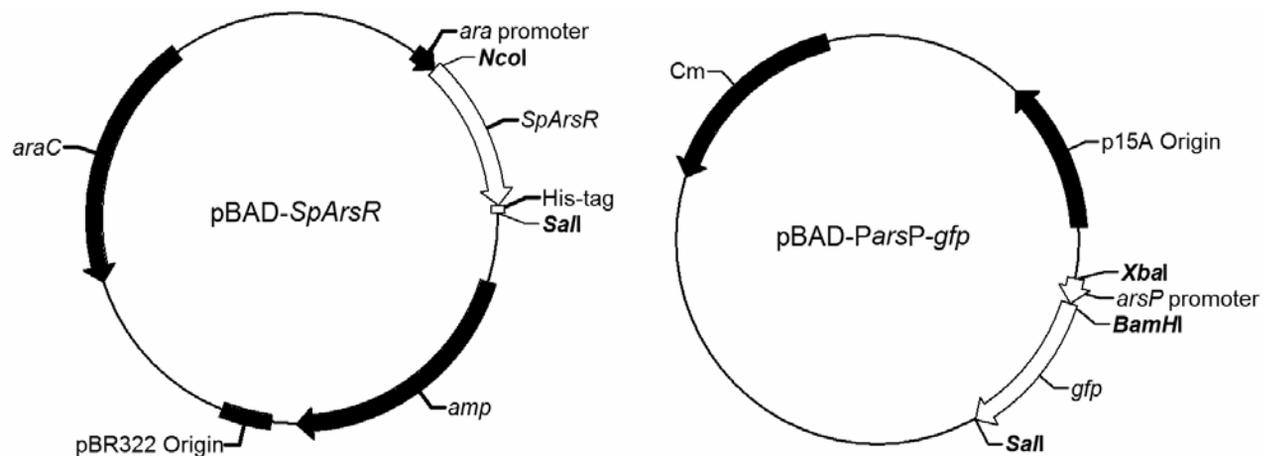
40 indicated concentrations. Fluorescence intensities were quantified by spectrofluorometry.

41 The data are the mean \pm SE (n = 3).

42

Fig. S2

A



B

Condition	arabinose	As	<i>arsR</i>	<i>gfp</i>	fluorescence
Constitutive	-	-	repressed	expressed	yes
Repression	+	-	expressed	repressed	no
Derepression	+	+	induced	expressed	yes

C

