

Appendix B

Supplementary Data

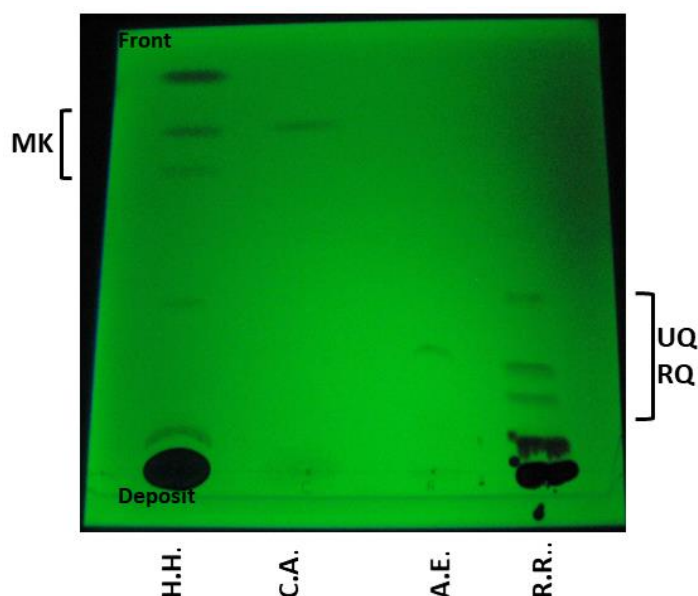


Figure B.1: Thin Layer Chromatography of quinone extracts from membranes from diverse bacteria. Quinones are extracted from membranes (Mb) by a mixing of 1.6 mL Mb/2mL chloroform/4 mL methanol. After a centrifugation of 20 min at 16000g, the lower phase is sampled. The solvent is evaporated under argon flux and in the absence of light. The sample is resuspended in ethanol for depositing on the TLC plate. *Halorhodospira halophila* (H.H), containing both MK and UQ, was grown under Na₂S/As(V) conditions. *Chrysiogenes arsenatis* (C.A.), containing only MK, was grown under As(V) conditions. *Alkalilimnicola ehrlichii I* (A.E.), containing only UQ, was grown under As(III) conditions. *Rhodospirillum rubrum* (R.R.), containing both UQ and RQ, is not an arsenicals-converting strain. To unambiguously identify the quinone from A.E as a high redox potential quinone, it has been studied by cyclic voltammetry determining its redox potential at $E_{\text{mph}7} = +65$ mV.

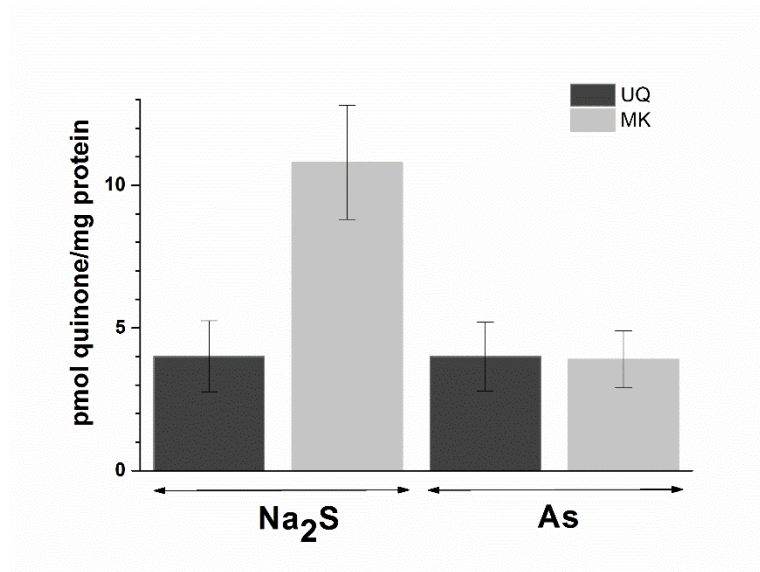


Figure B.2: Quantification of cellular UQ- and MK-content in lipid extracts from *H. halophila*. *H. halophila* has been grown anaerobically and photosynthetically on Na₂S or As^{III}, as exclusive electron donor. Error bars show standard deviations.

Acronym	Enzyme	strain name	phylum	A SU accession number	C type SU	Predicted Q pathway	biochemical Q analysis
SheAN	Arr	<i>Shewanella sp. ANA-3</i>	gamma Proteobacteria	AAQ01672.1	CymA	Ubi- Men	
ShePu	Arr	<i>Shewanella putrefaciens/Alivibrio salmonicida LFI1238</i>	gamma Proteobacteria	WP_014610142	CymA	Ubi- Men	
ShePi	Arr	<i>Shewanella piezotolerans WP3</i>	gamma Proteobacteria	ACJ28932	CymA	Ubi- Men	
SheW3	Arr	<i>Shewanella sp. W3-18-1</i>	gamma Proteobacteria	WP_011790217.1	CymA	Ubi- Men	
SheP1	Arr	<i>Shewanella sp. P1-14-1</i>	gamma Proteobacteria	WP_055024219.1	CymA	Ubi- Men	
SheWa	Arr	<i>Shewanella waksmanii</i>	gamma Proteobacteria	WP_028773911.1	CymA	Ubi- Men	
FerBa	Arr	<i>Ferrimonas balearica DSM 9799</i>	gamma Proteobacteria	WP_013345579.1	CymA	Ubi- Men	
FerFu	Arr	<i>Ferrimonas futtsuensis</i>	gamma Proteobacteria	WP_028109898.1	CymA	Ubi- Men	
FerKy	Arr	<i>Ferrimonas kyonanensis</i>	gamma Proteobacteria	WP_028113667.1	CymA	Ubi- Men	
FerSe	Arr	<i>Ferrimonas senticii</i>	gamma Proteobacteria	WP_028116765.1	CymA	Ubi- Men	
ThiTh	Arr	<i>Thioalkalivibrio paradoxus Arh 4</i>	gamma Proteobacteria	WP_006746746.1	ArrC	Ubi- Rqu Men	
AerHy	Arr	<i>Aeromonas hydrophila</i>	gamma Proteobacteria	WP_017788451.1	ArrC	Ubi- Men	
PhoGa	Arr	<i>Photobacterium ganghwense</i>	gamma Proteobacteria	WP_047886784.1	ArrC	Ubi- Men	
PsyAq	Arr	<i>Psychromonas aquimarina</i>	gamma Proteobacteria	WP_028865486.1	ArrC	Ubi- Men	
HalALS9	Arr	<i>Halomonas sp. ALS9</i>	gamma Proteobacteria	WP_064235365.1	No	Ubi- Men	
BurBa	Arr	<i>Burkholderiales bacterium 1_1_47</i>	beta Proteobacteria	ZP_07342827.1	ArrC	Men partial	
ParEx	Arr	<i>Parasutterella excrementihominis YIT 11859</i>	beta Proteobacteria	ZP_08322656.1	CymA	Ubi- Men	
SulHy	Arr	<i>Sulfuritalea hydrogenivorans sk43H</i>	beta Proteobacteria	BAO28251	ArrC	Ubi-	UQ
DesThAS	Arr	<i>Desulfonatrosipira thiodismutans ASO3-1</i>	delta Proteobacteria	ZP_07017810.1	ArrC	Mqn	
MLMS	Arr	<i>delta proteobacterium MLMS-1</i>	delta Proteobacteria	ZP_01288668.1	ArrC	Mqn	
GeoLo	Arr	<i>Geobacter lovleyi SZ</i>	delta Proteobacteria	WP_012469220.1	No	Mqn	MK
DesAl	Arr	<i>Desulfovibrio alkalitolerans DSM 16529</i>	delta Proteobacteria	EPR32287.1	ArrC	Mqn	
DesHal	Arr	<i>Desulfovermiculus halophilus</i>	delta Proteobacteria	WP_051564251.1	ArrC	Mqn	
DesBR	Arr	<i>Desulfatitalea sp. BRH_c12</i>	delta Proteobacteria	KJS28921	No	Mqn	
DesThio	Arr	<i>Desulfonatronum thioautotrophicum</i>	delta Proteobacteria	WP_045221492.1	No	Mqn	
DesTh	Arr	<i>Desulfonatronum thiodismutans</i>	delta Proteobacteria	WP_031387844	ArrC	Mqn	
DesLa	Arr	<i>Desulfonatronum lacustre</i>	delta Proteobacteria	WP_028572649.1	ArrC	Mqn	

DesBRc16	Arr	<i>Desulfobulbaceae bacterium BRH_c16a</i>	delta Proteobacteria	KJS00427	No	Mqn	
DesWTL	Arr	<i>Desulfuromonas soudanensis</i>	delta Proteobacteria	WP_053552011.1	No	Mqn	
WolSu	Arr	<i>Wolinella succinogenes DSM 1740</i>	epsilon Proteobacteria	WP_011138677.1	ArrC	Mqn	
SulBa	Arr	<i>Sulfurospirillum barnesii SES-3</i>	epsilon Proteobacteria	WP_014770549	ArrC	Mqn	MK
SulMu	Arr	<i>Sulfurospirillum multivorans</i>	epsilon Proteobacteria	WP_025346199.1	ArrC	Mqn	
SulAr	Arr	<i>Sulfurospirillum arsenophilum</i>	epsilon Proteobacteria	WP_041956283.1	ArrC	Mqn	
AnaAr	Arr	<i>Anaerobacillus arseniciselenatis</i>	Firmicutes	WP_071313526.1	ArrC	Mqn	MK
DesHaDCB	Arr	<i>Desulfitobacterium hafniense DCB-2</i>	Firmicutes	WP_015943299.1	ArrC	Men	
AlkOr	Arr	<i>Alkaliphilus oremlandii OhILAs</i>	Firmicutes	WP_012158954.1	No	None Full	MK
BacSeMLS10	Arr	<i>Bacillus selenitireducens MLS10</i>	Firmicutes	WP_013173528.1	ArrC	Mqn	
HalSi	Arr	<i>Halarsenatibacter silvermanii</i>	Firmicutes	ACF74513	No	None	cultivation failed
NatTh	Arr	<i>Natranaerobius thermophilus JW/NM-WN-LF</i>	Firmicutes	WP_012447122.1	No	None Full	None
DesHaY51	Arr	<i>Desulfitobacterium hafniense Y51</i>	Firmicutes	BAE85920.1	ArrC	Men	
DesYo	Arr	<i>Desulfosporosinus youngiae DSM 17734</i>	Firmicutes	WP_007781366.1	ArrC	Mqn	
DesDi	Arr	<i>Desulfitobacterium dichloroeliminans LMG P-21439</i>	Firmicutes	WP_015261495.1	ArrC	None Full	MK imported
AlkMe	Arr	<i>Alkaliphilus metalliredigens QYMF</i>	Firmicutes	WP_012062249.1	No	None Full	
BacZY	Arr	<i>Bacillus sp. ZYK</i>	Firmicutes	WP_017755305.1	ArrC	Men	
BacMa	Arr	<i>Bacillus massiliosenegalensis</i>	Firmicutes	WP_019156627.1	ArrC	Men	
BacBo	Arr	<i>Bacillus boroniphilus JCM 21738/Bacillus sp. 17376</i>	Firmicutes	GAE44117 +GAE44118	ArrC	Men	
CloBR	Arr	<i>Clostridiaceae bacterium BRH_c20a</i>	Firmicutes	KJS19576	CymA	Men partial	
PepBR	Arr	<i>Peptococcaceae bacterium BRH_c23</i>	Firmicutes	KJS46169	ArrC	Mqn	
PepBIC	Arr	<i>Peptococcaceae bacterium BICA1-8</i>	Firmicutes	KJS88232	ArrC	Men partial	
GraBR	Arr	<i>Gracilibacter sp. BRH_c7a</i>	Firmicutes	KUO58299	ArrC	Men	
DesBRc19	Arr	<i>Desulfitibacter sp. BRH_c19</i>	Firmicutes	KUO49382	ArrC	Men partial	
BacSe	Arr	<i>Bacillus selenatarsenatis</i>	Firmicutes	WP_041964264.1	ArrC	Men	
TepZ9	Arr	<i>Tepidibacillus decaturensis</i>	Firmicutes	KXG44095	ArrC	Men	
SpjC	Arr	<i>Spirochaeta sp. JC202</i>	Spirochaetes	WP_052080012.1	No	Men partial	
DenAc	Arr	<i>Denitrovibrio acetiphilus DSM 12809</i>	Deferribacteria	WP_013011386.1	No	Mqn	
Geol21	Arr	<i>Geovibrio sp. L21-Ace-BES</i>	Deferribacteria	WP_022851114 .1	No	Mqn	

DesIn	Arr	<i>Desulfurispirillum indicum S5</i>	Chrysiogenetes	WP_013505023.1	No	Mqn	
ChrAr	Arr	<i>Chrysiogenes arsenatis DSM 11915</i>	Chrysiogenetes	WP_027389593.1	No	Mqn	MK
NatGr	Arr	<i>Natronobacterium gregoryi SP2</i>	Archaea; Euryarchaeota; Halobacteria	WP_005576997.1	ArrC	Men	
HalNi	Arr	<i>Halobiforma nitratireducens</i>	Archaea; Euryarchaeota; Halobacteria	WP_006671188.1	ArrC	Men partial	
cMetNi1	Arr	<i>Candidatus Methanoperedens nitroreducens</i>	Archaea; Euryarchaeota; Methanomicrobia	WP_048088693.1	ArrC	Mqn	
AlkEh	Arx	<i>Alkalilimnicola ehrlichii MLHE-1</i>	gamma Proteobacteria	WP_011627967.1	ArrC	Ubi	UQ
HalHa	Arx	<i>Halorhodospira halophila SL1</i>	gamma Proteobacteria	WP_011813170.1	ArrC	Ubi- Men	UQ/MK
EctPH	Arx	<i>Ectothiorhodospira sp. PHS-1</i>	gamma Proteobacteria	WP_008932021.1	ArrC	Ubi- Rqu Men	
ThiALM	Arx	<i>Thioalkalivibrio sp. ALMg11</i>	gamma Proteobacteria	WP_018950173.1	ArrC	Ubi-Rqu	
HalA3	Arx	<i>Halomonas sp. A3H3</i>	gamma Proteobacteria	WP_022524225.1	ArrC	Ubi- Men	
ThiNi	Arx	<i>Thioalkalivibrio nitratireducens DSM 14787</i>	gamma Proteobacteria	WP_015257777.1	ArrC	Ubi- Rqu Men	
ThiALJ	Arx	<i>Thioalkalivibrio sulfidiphilus</i>	gamma Proteobacteria	WP_018952916	ArrC	Ubi-Rqu	
HalCh	Arx	<i>Halomonas chromatireducens</i>	gamma Proteobacteria	AMD01645	ArrC	Ubi	
HalBo	Arx	<i>Halomonas boliviensis</i>	gamma Proteobacteria	WP_040480658.1	ArrC	Ubi- Men	
MagMa	Arx	<i>Magnetospirillum magnetotacticum MS-1</i>	alpha Proteobacteria	WP_041039509.1	ArrC	Ubi	
AzoCIB	Arx	<i>Azoarcus sp. CIB</i>	beta Proteobacteria	WP_050415005.1	ArrC	Ubi-Rqu	
RIF67	Arx	Rhodospirillales bacterium RIFCSPLOWO2_12_FULL_67_15	alpha Proteobacteria	OHC82914.1	ArrC	Ubi? ?	
HydWC	Arx	Hydrogenophilales bacterium CG18_big_fil_WC_8_21_14_2_50_58_12	Proteobacteria	PIQ11014.1	ArrC	Ubi- Rqu Men	
MuRBG	Arx	<i>Candidatus Muproteobacteria bacterium RBG_16_65_34</i>	Proteobacteria	OGI47522.1	ArrC	Ubi	
AzoTo	Arx	<i>Azoarcus tolulyticus</i>	beta Proteobacteria	WP_076602992.1	ArrC	Ubi	
Gal59	Arx	<i>Gallionellales bacterium RIFCSPLOWO2_02_FULL_59_110</i>	beta Proteobacteria	OGS99566.1	ArrC	Ubi	
RIF62	Arx	<i>Betaproteobacteria bacterium RIFCSPLOWO2_12_FULL_62_13</i>	beta Proteobacteria	OGA50751.1	ArrC	Ubi	
SulDe	Arx	<i>Sulfuricella denitrificans</i>	beta Proteobacteria	WP_009207744.1	ArrC	Ubi- Rqu Men	
ThioKS	Arx	<i>Thiocapsa sp. KS1</i>	gamma Proteobacteria	WP_093186008.1	ArrC	Ubi- Rqu Men	
EctBL	Arx	<i>Ectothiorhodospira sp. BSL-9</i>	gamma Proteobacteria	WP_063465591.1	ArrC	Ubi- Rqu Men	
MarCe	Arx	<i>Marinospirillum celere</i>	gamma Proteobacteria	WP_091965171.1	ArrC	Ubi- Men	

ThiDe	Arx	<i>Thioalkalivibrio denitrificans</i>	gamma Proteobacteria	WP_077279016.1	ArrC	Ubi-Rqu	Men
ThiTh	Arx	<i>Thioalkalivibrio thiocyanodenitrificans</i>	gamma Proteobacteria	WP_018231946.1	ArrC	Ubi-Rqu	Men
HalBC	Arx	<i>Halomonas</i> sp. BC04	gamma Proteobacteria	WP_043513786.1	ArrC	Ubi-	Men partial
ThiAR	Arx	<i>Thioalkalivibrio</i> sp. ARh3	gamma Proteobacteria	WP_018864019.1	ArrC	Ubi-Rqu	
NitLa	Arx	<i>Nitrincola lacisaponensis</i>	gamma Proteobacteria	WP_036549261.1	ArrC	Ubi	
DelGW	Arx	<i>Deltaproteobacteria bacterium</i> GWC2_42_11	delta Proteobacteria	OGP31207.1	ArrC	Ubi	
cMetNiBLZ1	Unk	<i>Candidatus Methanoperedens nitroreducens</i>	Archaea; Euryarchaeota; Methanomicobia	KPQ43588	ArrC	Mqn	
cMetNi2	Unk	<i>Candidatus Methanoperedens</i> spBLZ1	Archaea; Euryarchaeota; Methanomicobia	WP_048088677	ArrC	Mqn	
HR01	Unk	archaeon HR1	Archaea	GBC69680.1	No	none Full	
HR02	Unk	archaeon HR2	Archaea	GBC70990.1	No	none Full	

Table B.1: Arr/Arx sequences used for reconstruction of the phylogenetic trees shown in Figure 1 and Figure A.2 and analysis of the quinone content of the related strains.

Strain	<i>arr</i> or <i>arx</i> genes	Quinone content	As ^{III} conversion around pH7	As ^{III} conversion around pH9	As ^V conversion around pH7	A ^V conversion around pH 9
<i>A. ehrlichii</i>	<i>arx</i>	UQ	NO	YES, with growth	NO	NO
<i>D. hafniense</i>	<i>arr</i>	MK	NO	NO	YES, with growth	NO
<i>S. hydrogenivorans</i>	<i>arr</i>	UQ	NO	n.d.	YES, with growth	n.d.
with gramicidin			n.d.	n.d.	NO	n.d.
<i>H. halophila</i>	<i>arx</i>	MK and UQ	NO	YES, with growth	YES, no growth	NO
with gramicidin			n.d.	n.d.	NO	n.d.

Table B.2: Summary of the results of growth- and arsenical-conversion-experiments performed on the four strains tested, in relation to their quinone- and Arr- or Arx-enzyme-content. n.d. indicates that the experiment has not been performed.