

## Supplemental Material

Bacteria Resistance to Metal Pollution in a Copper Tailing Dam Area in Northern China

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**TABLE S1 Target genes and primer sequences used in this study**

|      | gene name   | Function                                      | Primer 5'-3'<br>forward and reverse                                               | Product size (bp) | Annealing temperature (°C) | Reference             |
|------|-------------|-----------------------------------------------|-----------------------------------------------------------------------------------|-------------------|----------------------------|-----------------------|
| MRGs | <i>copA</i> | Cu translocating ATPase                       | F CGG TCT CTA CGA ATA CCG CTT CAA<br>R GAA ATA GCT CAT TGC CGA GGC GTT            | 300               | 55                         | Bouskill et al., 2007 |
| MRGs | <i>copB</i> | Cu binding protein                            | F TTC CTG CTC GAC CAG TTG GAA TAC<br>R GGT TGG TCA ACA GGA TGT CGT ACT            | 450               | 58                         | Bouskill et al., 2007 |
| MRGs | <i>pcoA</i> | multi-copper oxidase                          | F ATA CCA TCT TCG CAC AAT CCA<br>R GCT GTT ATC CAT GCC TCC C                      | 194               | 60                         | Lee et al., 2002      |
| MRGs | <i>pcoC</i> | Periplasmic protein                           | F TGT CGA TTT TAA ATA AAG CCA TTC TT<br>R TCA TAA TCA GGT CGT TCA TAA TAT TAC TTC | 400               | 60                         | Lee et al., 2002      |
| MRGs | <i>pcoD</i> | Periplasmic protein                           | F GAT ATC TGA CTC CCT GGC<br>R CCG TAA AAT CAA AGG GCT                            | 734               | 60                         | Lee et al., 2002      |
| MRGs | <i>czcA</i> | Co/Zn/Cd efflux protein                       | F GTT CAC CTT GCT CTT CGC CAT GTT<br>R ACA GGT TGC GGA TGA AGG AGA TCA            | 320               | 56                         | Bouskill et al., 2007 |
| MRGs | <i>czcC</i> | Outer part of a Co/Zn/Cd efflux protein (RND) | F AGC CGY CAG TAT CCG GAT CTG AC<br>R GTG GTC GCC GCC TGA TAG GT                  | 418               | 62                         | Roosa et al., 2014    |
| MRGs | <i>czcD</i> | Cation diffusion facilitator                  | F TCA TCG CCG GTG CGA TCA TCA T<br>R TGT CAT TCA CGA CAT GAA CC                   | 272               | 55                         | Roosa et al., 2014    |
| MRGs | <i>nccA</i> | Inner part of a Ni/Co efflux protein (RND)    | F TTY AGC CAG GTV ACS GTS ATY TT<br>R GCY GCR TCS GCR CGC ACC AGR TA              | 532               | 63                         | Roosa et al., 2014    |

|      | gene<br>name  | Function                   | Primer 5'-3'<br>forward and reverse                                    | Product<br>size<br>(bp) | Annealing<br>temperature<br>(°C) | Reference             |
|------|---------------|----------------------------|------------------------------------------------------------------------|-------------------------|----------------------------------|-----------------------|
| MRGs | <i>pbrT</i>   | Pb uptake protein          | F AGC GCG CCC AGG AGC GCA GCG TCT T<br>R GGC TCG AAG CCG TCG AGR TA    | 448                     | 55                               | Roosa et al., 2014    |
| MRGs | <i>chrB</i>   | Cr efflux protein          | F GTC GTT AGC TTG CCA ACA TC<br>R CGG AAA GCA AGA TGT CGA TCG          | 450                     | 57                               | Nie et al., 1990      |
| MRGs | <i>arsB</i>   | Arsenite efflux ATPase     | F GGT CTA TGC GCT GGA GCA ATT GAA<br>R TGC TGG GCA TGT TGT TCA TTA CCG | 500                     | 46                               | Bouskill et al., 2007 |
| MRGs | <i>arsC</i>   | Arsenate reductase protein | F GCA GCA TTC TTT CCG AAG CCA TGT<br>R TCG CAA ACG GTG ATG ACG ATG T   | 215                     | 46                               | Bouskill et al., 2007 |
| ARGs | <i>tetA</i>   | Tetracycline efflux        | F GCT ACA TCC TGC TTG CCT TC<br>R CAT AGA TCG CCG TGA AGA GG           | 210                     | 55                               | Ng et al., 2001       |
| ARGs | <i>tetC</i>   | Tetracycline efflux        | F CTT GAG AGC CTT CAA CCC AG<br>R ATG GTC GTC ATC TAC CTG CC           | 418                     | 55                               | Ng et al., 2001       |
| ARGs | <i>tetE</i>   | Tetracycline efflux        | F GTT ATT ACG GGA GTT TGT TGG<br>R AAT ACA ACA CCC ACA CTA CGC         | 278                     | 55                               | Ng et al., 2001       |
| ARGs | <i>tetG</i>   | Tetracycline efflux        | F GCT CGG TGG TAT CTC TGC TC<br>R AGC AAC AGA ATC GGG AAC AC           | 468                     | 55                               | Ng et al., 2001       |
| ARGs | <i>tetK</i>   | Tetracycline efflux        | F CGA AAA CAG ACT CGC CAA TC<br>R TCC ATA ATG AGG TGG GGC              | 169                     | 55                               | Ng et al., 2001       |
| ARGs | <i>tetL</i>   | Tetracycline efflux        | F TCG TTA GCG TGC TGT CAT TC<br>R GTA TCC CAC CAA TGT AGC CG           | 267                     | 55                               | Ng et al., 2001       |
| ARGs | <i>tetA/P</i> | Tetracycline efflux        | F CTT GGA TTG CGG AAG AAG AG<br>R ATA TGC CCA TTT AAC CAC GC           | 676                     | 55                               | Ng et al., 2001       |

|      | gene<br>name  | Function   | Primer 5'-3'<br>forward and reverse                                | Product<br>size<br>(bp) | Annealing<br>temperature<br>(°C) | Reference           |
|------|---------------|------------|--------------------------------------------------------------------|-------------------------|----------------------------------|---------------------|
| ARGs | <i>tetS</i>   | protection | F CAT AGA CAA GCC GTT GAC C<br>R ATG TTT TTG GAA CGC CAG AG        | 667                     | 55                               | Ng et al., 2001     |
| ARGs | <i>tetX</i>   | unkown     | F CAA TAA TTG GTG GTG GAC CC<br>R TTC TTA CCT TGG ACA TCC CG       | 468                     | 58                               | Ng et al., 2001     |
| ARGs | <i>tetM</i>   | protection | F ACA GAA AGC TTA TTA TAT AAC<br>R TGG CGT GTC TAT GAT GTT CAC     | 171                     | 45                               | Aminov et al., 2001 |
| ARGs | <i>tetO</i>   | protection | F ACG GAR AGT TTA TTG TAT ACC<br>R TGG CGT ATC TAT AAT GTT GAC     | 171                     | 45                               | Aminov et al., 2001 |
| ARGs | <i>tetQ</i>   | protection | F AGA ATC TGC TGT TTG CCA GTG<br>R CGG AGT GTC AAT GAT ATT GCA     | 169                     | 55                               | Aminov et al., 2001 |
| ARGs | <i>tetT</i>   | protection | F AAG GTT TAT TAT ATA AAA GTG<br>R AGG TGT ATC TAT GAT ATT TAC     | 169                     | 40                               | Aminov et al., 2001 |
| ARGs | <i>tetW</i>   | protection | F GAG AGC CTG CTA TAT GCC AGC<br>R GGG CGT ATC CAC AAT GTT AAC     | 168                     | 60                               | Aminov et al., 2001 |
| ARGs | <i>tetB/P</i> | protection | F AAA ACT TAT TAT ATT ATA GTG<br>R TGG AGT ATC AAT AAT ATT CAC     | 169                     | 40                               | Aminov et al., 2001 |
| ARGs | <i>suI</i>    | protection | F CGC ACC GGA AAC ATC GCT GCA C<br>R TGA AGT TCC GCC GCA AGG CTC G | 163                     | 56                               | Pei et al., 2006    |
| ARGs | <i>suII</i>   | protection | F TCC GGT GGA GGC CGG TAT ATG G<br>R CGG GAA TGC CAT CTG CCT TGA G | 191                     | 61                               | Pei et al., 2006    |
| ARGs | <i>suIII</i>  | protection | F TCC GTT CAG CGA ATT GGT GCA G<br>R TTC GTT CAC GCC TTA CAC CAG C | 128                     | 60                               | Pei et al., 2006    |

|      | gene<br>name               | Function   | Primer 5'-3'<br>forward and reverse                                  | Product<br>size<br>(bp) | Annealing<br>temperature<br>(°C) | Reference               |
|------|----------------------------|------------|----------------------------------------------------------------------|-------------------------|----------------------------------|-------------------------|
| ARGs | <i>qnrA</i>                | unkown     | F TCA GCA AGA GGA TTT CTC A<br>R GGC AGC ACT ATG ACT CCC A           | 516                     | 50                               | Kehrenberg et al., 2006 |
| ARGs | <i>qnrB</i>                | unkown     | F TCG GCT GTC AGT TCT ATG ATC G<br>R TCC ATG AGC AAC GAT GCC T       | 469                     | 54                               | Kehrenberg et al., 2006 |
| ARGs | <i>qnrS</i>                | unkown     | F TGA TCT CAC CTT CAC CGC TTG<br>R GAA TCA GTT CTT GCT GCC AGG       | 566                     | 58                               | Kehrenberg et al., 2006 |
| ARGs | <i>ereA</i>                | deactivate | F AAC ACC CTG AAC CCA AGG GAC G<br>R CTT CAC ATC CGG ATT CGC TCG     | 420                     | 52                               | Sutcliffe et al., 1996  |
| ARGs | <i>ereB</i>                | deactivate | F AGA AAT GGA GGT TCA TAC TTA CCA<br>R CAT ATA ATC ATC ACC AAT GGC A | 546                     | 52                               | Sutcliffe et al., 1996  |
| ARGs | <i>mphA</i>                | deactivate | F AAC TGT ACG CAC TTG C<br>R GGT ACT CTT CGT TAC C                   | 837                     | 52                               | Sutcliffe et al., 1996  |
| ARGs | <i>bla<sub>CTX-M</sub></i> | deactivate | F ATG TGC AGY ACC AGT AAR GT<br>R TGG GTR AAR TAR GTS ACC AGA        | 593                     | 50                               | Shahid, 2010            |
| ARGs | <i>bla<sub>TEM</sub></i>   | deactivate | F KAC AAT AAC CCT GRT AAA TGC<br>R AGT ATA TAT GAG TAA ACT TGG       | 936                     | 58                               | Shahid, 2010            |
| ARGs | <i>bla<sub>SHV</sub></i>   | deactivate | F TTT ATC GGC CYT CAC TCA AGG<br>R GCT GCG GGC CGG ATA ACG           | 930                     | 58                               | Shahid, 2010            |
| ARGs | <i>bla<sub>ampC</sub></i>  | deactivate | F CCC CGC TTA TAG AGC AAC AA<br>R TCA ATG GTC GAC TTC ACA CC         | 634                     | 58                               | Shahid, 2010            |
| MGE  | <i>intI1</i>               | integrase  | F CTG GAT TTC GAT CAC GGC ACG<br>R ACA TGC GTG TAA ATC ATC GTC G     | 473                     | 60                               | Hardwick et al., 2010   |

|     | gene<br>name | Function    | Primer 5'-3'<br>forward and reverse                          | Product<br>size<br>(bp) | Annealing<br>temperature<br>(°C) | Reference        |
|-----|--------------|-------------|--------------------------------------------------------------|-------------------------|----------------------------------|------------------|
| MGE | <i>tnpA</i>  | transposase | F CCG ATC ACG GAA AGC TCA AG<br>R CCG ATC ACG GAA AGC TCA AG | 101                     | 60                               | Han et al., 2016 |

MRGs: metal resistance genes. ARGs: antibiotic resistance genes. MGE: mobile gene elements.

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