

Supplemental Data 3: Oligonucleotides Used in this Study

| Name | Sequence (5' to 3') | Notes |
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| MembProt n87 PS F* | AAATACAAATAAACAACTAAGACAG | Membrane protein protospacer targeting nucleotide 87 |
| MembProt n150 PS F* | AAATAAAATTCTACTTATGTACTG | Membrane protein protospacer targeting nucleotide 150 |
| Hypo n57 PS F* | AAATTTCAAAGTGAATGAAGTGTG | Hypothetical protein protospacer targeting nucleotide 57 |
| Hypo n270 PS F* | AAATAATCTTCTAACACGTGAAATCG | Hypothetical protein protospacer targeting nucleotide 270 |
| <i>mrvR</i> n103 PS F* | AAATTAGAAGGCAATTGCATTCCG | <i>mrvR</i> protospacer targeting nucleotide 103 |
| <i>mrvR</i> n128 PS F* | AAATATTAAGTTGCCCTGCTAATTG | <i>mrvR</i> protospacer targeting nucleotide 128 |
| <i>deoR</i> n316 PS F* | AAATAATTCTCTAACGGTTGTTCCG | <i>deoR</i> protospacer targeting nucleotide 316 |
| <i>deoR</i> n326 PS F* | AAATTTGCGTAGCTAACCTCTAG | <i>deoR</i> protospacer targeting nucleotide 326 |
| ABC n103 PS F* | AAATCTTAATTATCTATCCCTACG | ABC transporter protospacer targeting nucleotide 103 |
| ABC n158 PS F* | AAATAAATTGGGTAAATACTGTGTG | ABC transporter protospacer targeting nucleotide 158 |
| 3015b NT PS F* | AGACCGCGTACTACGCCTATTGGT | Non-targeting (sham) protospacer used in control knockdown strain |
| MembProt qRT-PCR F | TCATCGGAAGTGTCACTCCTT | Membrane protein qRT-PCR primer |
| MembProt qRT-PCR R | CCGGATGTTCTAGCTTAGACT | Membrane protein qRT-PCR primer |
| Hypo qRT-PCR F | AAGACCTCATAAGCGTCGTAAG | Hypothetical protein qRT-PCR primer |
| Hypo qRT-PCR R | GAGCATTGGCATTGACTGG | Hypothetical protein qRT-PCR primer |
| <i>mrvR</i> qRT-PCR F | GAAGTGGATTAGAATTCTGTAGAG | <i>mrvR</i> qRT-PCR primer |
| <i>mrvR</i> qRT-PCR R | TGGTACCAACGATCAGCAATAA | <i>mrvR</i> qRT-PCR primer |
| <i>deoR</i> qRT-PCR F | GGTAGTAACCAATAGTCTTCTGTT | <i>deoR</i> qRT-PCR primer |
| <i>deoR</i> qRT-PCR R | GCACCAGTAATTCTCTGTACTCT | <i>deoR</i> qRT-PCR primer |
| ABC qRT-PCR F | TCACTCGTTCTACAGTATCT | ABC protein qRT-PCR primer |
| ABC qRT-PCR R | AATTCCCTCCAATTGCCACTAAA | ABC protein qRT-PCR primer |

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| <i>recA</i> qRT-PCR F | GTGGGATTGCTGCCTTATTG | For qRT-PCR normalization |
| <i>recA</i> qRT-PCR R | CTGAGTCAGGTTGAGACAAGAG | For qRT-PCR normalization |
| CRISPRi qPCR R | CAGCTGAATAACGGACTGGATAT | Reverse qPCR primer used for CRISPRi competition experiment |
| 3015b qPCR F | GCGCCTTATCCGGTAACATAC | qPCR primer binds the 3015b plasmid backbone independent of the protospacer; for qPCR normalization |
| 3015b qPCR R | CTACATACCTCGCTTGCTAAC | qPCR primer binds the 3015b plasmid backbone independent of the protospacer; for qPCR normalization |
| <i>mrvR</i> comp GA F | ACAGCTATGACATGATTACGGATAAAGACCTCC TTAATGTC | For Gibson assembly cloning into pBSU101 double-digested with XbaI and EcoRI |
| <i>mrvR</i> comp GA R | TGCATGCCTGCAGGTCGACTCTTAATGAGAGGT AATGGAC | For Gibson assembly into pBSU101 double-digested with XbaI and EcoRI |
| <i>dmrVR</i> mut GA F | ATTGGGTACCGGGCCCC | For Gibson assembly of the <i>mrvR</i> allelic exchange knockout cassette into pMBSacB digested with Xhol and NotI |
| <i>dmrVR</i> mut GA R | TGGAGCTCCACCGCGGTGGCAAAAGTAGCAA AACAAATGATTGCTATTACCTTAGATTGAGATCG | For Gibson assembly of the <i>mrvR</i> allelic exchange knockout cassette into pMBSacB digested with Xhol and NotI |
| 1084 Phage R1 F | ACACCGAAAGACCAAGCGAT | For PCR confirmation of prophage region 1 presence |
| 1084 Phage R1 R | TTACGACGGCGGTATGTACG | For PCR confirmation of prophage region 1 presence |
| 1084 Phage qPCR R1 F | AGGGCAAAGACTCGGTAAATG | Prophage region 1 qRT-PCR primer |
| 1084 Phage qPCR R1 R | CATTCTTATGGGTCGGGAATT | Prophage region 1 qRT-PCR primer |
| 1084 Phage R2 F | GGTGAATGTCGTCGTTGTCT | For PCR confirmation of prophage region 2 presence |
| 1084 Phage R2 R | GCCACACGTTGTTCTCC | For PCR confirmation of prophage region 2 presence |
| 1084 Phage qPCR R2 F | CTAGATGCCTGTCCAACTTCTT | Prophage region 2 qRT-PCR primer |

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| 1084 Phage qPCR R2 R | ACTCTATGTCCATCATTCCCTTATC | Prophage region 2 qRT-PCR primer |
| 1084 Phage R3 F | GCTGGTGGCTGAAATGCA | For PCR confirmation of prophage region 3 presence |
| 1084 Phage R3 R | CATCATCAGGTACGCGACGA | For PCR confirmation of prophage region 3 presence |
| Erm pFfluc GA F | CACTCTTAAGGTATTAAAGATACCCCAAGAAG | For Gibson assembly replacement of the kanamycin resistance gene in the luciferase expression plasmid pLZ12Km2-P23R:TA:fFfluc with the erythromycin resistance gene (<i>erm</i>) |
| Erm pFfluc GA R | CTTTAACTACAAGCTTTTAGAC | For Gibson assembly replacement of the kanamycin resistance gene in the luciferase expression plasmid pLZ12Km2-P23R:TA:fFfluc with the erythromycin resistance gene (<i>erm</i>) |
| Luc1 pFfluc GA F | TAAAAAGCTTGAGTTAAAGGCGACTCATGAA TTATTTC | For Gibson assembly replacement of the kanamycin resistance gene in the luciferase expression plasmid pLZ12Km2-P23R:TA:fFfluc with the erythromycin resistance gene (<i>erm</i>) |
| Luc1 pFfluc GA R | AGTAACGATTGACCGCCAATACTCATGAA CGGCTTG | For Gibson assembly replacement of the kanamycin resistance gene in the luciferase expression plasmid pLZ12Km2-P23R:TA:fFfluc with the erythromycin resistance gene (<i>erm</i>) |
| Luc2 pFfluc GA F | GCGGTCAATGAATCGTTAC | For Gibson assembly replacement of the kanamycin resistance gene in the luciferase expression plasmid pLZ12Km2-P23R:TA:fFfluc with the erythromycin resistance gene (<i>erm</i>) |
| Luc2 pFfluc GA R | CTTTAAATACCTAACAGAGTGTGTTGATAGTG | For Gibson assembly replacement of the kanamycin resistance gene in the luciferase expression plasmid pLZ12Km2-P23R:TA:fFfluc with the erythromycin resistance gene (<i>erm</i>) |
| Tn-seq Amp F | CAAGCAGAAGACGGCATACGAGGTTGAAGTC TGATCTTCGG | For transposon-genome junction amplification in Tn-seq. See reference 41 |

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| Tn-seq Amp R | AATGATAACGGCGACCACCGAGATCACACTTT CCCTACACGACGCTTCC | For transposon-genome junction amplification in Tn-seq. See reference 41 |
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*Forward protospacer oligonucleotide; also used as forward qPCR primer in CRISPRi competition experiment