Table S3. Bacterial strains, phages, plasmids and primers used in this study.

Strains, phages, plasmids and primers	Characteristics and/or description	Reference /Source
Enterococcus faecalis		
OG1RF	Human oral isolate; Rf^R, Fa^R	(1)
Δpip$_{EF}$	OG1RF background with a deletion of OG1RF_10588 (PIP)	(2)
OG1RF_10820-Tn	*E. faecalis* OG1RF *OG1RF_10820* (LytR family response regulator) transposon mutant (Genomic position: 860054). Rf^R, Fa^R, Cm^R	(3)
cscK-Tn	*E. faecalis* OG1RF OG1RF_10951 (fructosekinase) transposon mutant (Genomic position: 993104). Rf^R, Fa^R, Cm^R	(3)
OG1RF_12241-Tn	*E. faecalis* OG1RF OG1RF_12241 (LysR family transcriptional regulator) transposon mutant (Genomic position: 2363777). Rf^R, Fa^R, Cm^R	(3)
OG1RF_11710-Tn	*E. faecalis* OG1RF OG1RF_11710 (O-antigen polymerase) transposon mutant (Genomic position: 1789261). Rf^R, Fa^R, Cm^R	(3)
OG1RF_11714-Tn	*E. faecalis* OG1RF OG1RF_11714 (group 2 glycosyl transferase) transposon mutant (Genomic position: 1793746). Rf^R, Fa^R, Cm^R	(3)
OG1RF_11715-Tn	*E. faecalis* OG1RF OG1RF_11715 (*epaOX*, glycosyltransferase) transposon mutant (Genomic position: 1794475). Rf^R, Fa^R, Cm^R	(3)
ΔepaOX	JD102; *E. faecalis* OG1RF *ΔepaOX* (Δ*OG1RF_11715,* glycosyltransferase) markerless deletion. Rf^R, Fa^R	(4)
OG1RF_12434-Tn	*E. faecalis* OG1RF OG1RF_12434 (*mutL*) locus transposon mutant (Genomic position: 2567339), Rf^R, Fa^R, Cm^R	(3)
OG1RF_12435-Tn	*E. faecalis* OG1RF OG1RF_12435 (*mutS*) locus transposon mutant (Genomic position: 2569574), Rf^R, Fa^R, Cm^R	(3)
Wild type - E	OG1RF strain carrying pAT28 empty vector	This study

Δ*pip*$_{EF}$ - E	*Δpip*$_{EF}$ strain carrying pAT28 empty vector	This study
OG1RF_10820-Tn - E	OG1RF_10820 transposon mutant carrying pAT28 empty vector. Rf^R, Fa^R, Cm^R,Sp^R	This study
cscK-Tn - E	OG1RF_10951 transposon mutant carrying pAT28 empty vector. Rf^R, Fa^R, Cm^R, Sp^R	This study
OG1RF_12241-Tn - E	OG1RF_12241 transposon mutant carrying pAT28 empty vector. Rf^R, Fa^R, Cm^R, Sp^R	This study
OG1RF_10820-Tn - C	OG1RF_10820 transposon mutant carrying pAT28 complementation vector. Rf^R, Fa^R, Cm^R,Sp^R	This study
cscK-Tn - C	OG1RF_10951 transposon mutant carrying pAT28 complementation vector. Rf^R, Fa^R, Cm^R, Sp^R	This study
OG1RF_12241-Tn - C	OG1RF_12241 transposon mutant carrying pAT28 complementation vector. Rf^R, Fa^R, Cm^R, Sp^R	This study
mutL-Tn - E	OG1RF_12434 (*mutL*) locus transposon mutant carrying pAT28 empty vector. Rf^R, Fa^R, Cm^R, Sp^R	This study
mutS-Tn - E	OG1RF_12435 (*mutS*) locus transposon mutant carrying pAT28 empty vector. Rf^R, Fa^R, Cm^R, Sp^R	This study
mutL-Tn - C	OG1RF_12434 (*mutL*) locus transposon mutant carrying pAT28 complementation vector. Rf^R, Fa^R, Cm^R, Sp^R	This study
mutS-Tn - C	OG1RF_12435 (*mutS*) locus transposon mutant carrying pAT28 complementation vector. Rf^R, Fa^R, Cm^R, Sp^R	This study
mutL-Tn-NPh_C1	*mutL*-Tn colony # 1 selected from O/N THB agar plate. Rf^R, Fa^R, Cm^R	This study
mutL-Tn-NPh_C2	*mutL*-Tn colony # 2 selected from O/N THB agar plate. Rf^R, Fa^R, Cm^R	This study
mutL-Tn-NPh_C3	*mutL*-Tn colony # 3 selected from O/N THB agar plate. Rf^R, Fa^R, Cm^R	This study
mutL-Tn-NPh_C4	*mutL*-Tn colony # 4 selected from O/N THB agar plate. Rf^R, Fa^R, Cm^R	This study
mutL-Tn-Ph_C1	*mutL*-Tn colony # 1 selected from O/N THB agar plate containing VPE25. Rf^R, Fa^R, Cm^R	This study
mutL-Tn-Ph_C2	*mutL*-Tn colony # 2 selected from O/N THB agar plate containing VPE25. Rf^R, Fa^R, Cm^R	This study

mutL-Tn-Ph_C3	*mutL*-Tn colony # 3 selected from O/N THB agar plate containing VPE25. Rf^R, Fa^R, Cm^R	This study
mutL-Tn-Ph_C4	*mutL*-Tn colony # 4 selected from O/N THB agar plate containing VPE25. Rf^R, Fa^R, Cm^R	This study
mutS-Tn-NPh_C1	*mutS*-Tn colony # 1 selected from O/N THB agar plate. Rf^R, Fa^R, Cm^R	This study
mutS-Tn-NPh_C2	*mutS*-Tn colony # 2 selected from O/N THB agar plate. Rf^R, Fa^R, Cm^R	This study
mutS-Tn-NPh_C3	*mutS*-Tn colony # 3 selected from O/N THB agar plate. Rf^R, Fa^R, Cm^R	This study
mutS-Tn-NPh_C4	*mutS*-Tn colony # 4 selected from O/N THB agar plate. Rf^R, Fa^R, Cm^R	This study
mutS-Tn-Ph_C1	*mutS*-Tn colony # 1 selected from O/N THB agar plate containing VPE25. Rf^R, Fa^R, Cm^R	This study
mutS-Tn-Ph_C2	*mutS*-Tn colony # 2 selected from O/N THB agar plate containing VPE25. Rf^R, Fa^R, Cm^R	This study
mutS-Tn-Ph_C3	*mutS*-Tn colony # 3 selected from O/N THB agar plate containing VPE25. Rf^R, Fa^R, Cm^R	This study
mutS-Tn-Ph_C4	*mutS*-Tn colony # 4 selected from O/N THB agar plate containing VPE25. Rf^R, Fa^R, Cm^R	This study
Escherichia coli		
TG1	*[F' traD36 proAB lacIqZ ΔM15] supE thi-1 Δ(lac-proAB) Δ(mcrBhsdSM)5(rK - mK -)*	Lucigen
Phage		
VPE25	Siphoviridae; Wastewater isolate	(2)
NPV1	Siphoviridae; Wastewater isolate	(5)
Plasmids		
pAT28	*Escherichia coli* and in Gram-positive bacteria shuttle vector; Sp^R	(6)
pAT28: OG1RF_10820	pAT28 expressing OG1RF_10820 from the native promoter. Cloned into SacI/BamHI site. Sp^R	This study
pAT28: *cscK*	pAT28 expressing *cscK* from the native promoter. Cloned into SacI/BamHI site. Sp^R	This study

pAT28: OG1RF_12241	pAT28 expressing OG1RF_12241 from the native promoter. Cloned into SacI/BamHI site. Sp^R	This study
pAT28: *muL*	pAT28 expressing *mutL* from the native promoter. Cloned into SacI/XbaI site. Sp^R	This study
pAT28: *mutS*	pAT28 expressing *mutS* from the native promoter. Cloned into SacI/EcoRI site. Sp^R	This study
Primers		
lytR-F	NNNNNNGAGCTCACAGTTATTTTTTTTGTTTGGTGTTAC; Forward primer to generate pAT28:*OG1RF_10820*; SacI site	This study
lytR-R	NNNNNNGGATCCGAATCTATGTTATAACTAACATCATTG; Reverse primer to generate pAT28:*OG1RF_10820*; BamHI site	This study
cscK-F	NNNNNNGAGCTCTCATTGTTAGATAACCGTGAATGCTC; Forward primer to generate pAT28:*cscK*; SacI site	This study
cscK-R	NNNNNNGGATCCTCATTCTTAGCCATTTCCTTCAAAT; Reverse primer to generate pAT28:*cscK*; BamHI site	This study
lysR-F	NNNNNNGAGCTCTCGCCTCCTTTTGCTAAATTTATCG; Forward primer to generate pAT28:*OG1RF_12241*; SacI site	This study
lysR-R	NNNNNNGGATCCAGTATTCAACGACCTATTATTCAG; Reverse primer to generate pAT28:*OG1RF_12241*; BamHI site	This study
mutS-F	NNNNNNGAGCTCCCTCCTAACCGGTTCGGTTCCTAATTAGA; Forward primer to generate pAT28:*mutS*; SacI site	This study
mutS-R	NNNNNNGAATTCGAAGAAAAGTTAATCTCGGATCATTAAGAGG; Reverse primer to generate pAT28:*mutS*; EcoRI site	This study
mutL-F	NNNNNNGAGCTCAATGATGACGCCTCGTTTCTTTTTAATGGG; Forward primer to generate pAT28:*mutL*; SacI site	This study
mutL-R	NNNNNNTCTAGAAGGAACCGAACCGGTTAGGAGGTCGTGAAATGG; Reverse primer to generate pAT28:*mutL*; XbaI site	This study

RT-*fsrA* -F	GCAGGATTTGAGGTTGCTAAAG; qPCR forward primer for *fsrA*	This study
RT-*fsrA* -R	GAGCTGAAACCATATATTTGTAGGAAG; qPCR reverse primer for *fsrA*	This study
RT-*fsrBD* -F	TCTTCTGTGAGCTTACCGTTTATT; qPCR forward primer for *fsrBD*	This study
RT-*fsrBD* -R	CAGGTTCAATTGCTGTTCCTTG; qPCR reverse primer for *fsrBD*	This study
RT-*fsrC* -F	TAACGAAGAGCTAGCGATGTTT; qPCR forward primer for *fsrC*	This study
RT-*fsrC* -R	GGTTGGTGCAATCGTTTCTTC; qPCR reverse primer for *fsrC*	This study
RT-*gelE* -F	CGCCATCACTAGCGACATTA; qPCR forward primer for *gelE*	This study
RT-*gelE* -R	GGCATCCCTCGATCATCAATAC; qPCR reverse primer for *gelE*	This study
RT-*sprE* -F	AGAAGACAAGAAGTGGCAGATAC; qPCR forward primer for spr*E*	This study
RT-*sprE* -R	CAACAACAAAGCCTGTTCCTAAA; qPCR reverse primer for spr*E*	This study
RT-10296 -F	GCTTCACCAGGCTCCATTATAG; qPCR forward primer for OG1RF_10296	This study
RT-10296 -F	TGACTGCGGATACTGCAAAG; qPCR reverse primer for OG1RF_10296	This study
RT-10297 -F	TGTGTTGAGCCCGCTTATATC; qPCR forward primer for OG1RF_10297	This study
RT-10297 -F	TGGCTTCTTCTGTTCCATGAG; qPCR reverse primer for OG1RF_10297	This study
RT-10875 -F	GACTGTATTACGCTTCGCTAAAG; qPCR forward primer for OG1RF_10875	This study
RT-10875 -R	CCACATTGAACTGCCCATAAA; qPCR reverse primer for OG1RF_10875	This study
RT-10876 -F	TGTGGGCTAAGCTAATGAATTT; qPCR forward primer for OG1RF_10876	This study

RT-10876 -R	AGTTCTCCTGAATCTATCCAAGTA; qPCR reverse primer for OG1RF_10876	This study
RT-*eutB* -F	CAATGGAAGCGCGTTGTTATG; qPCR forward primer for *eutB*	This study
RT-*eutB* -R	AGCGCGGATGACTTGTTT; qPCR reverse primer for *eutB*	This study
RT-*eutC* -F	CAGCTGCTATCGAAGCGAATA; qPCR forward primer for *eutC*	This study
RT-*eutC* -R	TTGCTGGCACACGACAATA; qPCR reverse primer for *eutC*	This study
RT-*eutH*-F	TAGCTGGCGCTTTCTGTTTAG; qPCR forward primer for *eutH*	This study
RT-*eutH* -R	AGCCGCTACTTCATTCATTCC; qPCR reverse primer for *eutH*	This study
RT-*epaA* -F	GCAGGAGGACTTGCAATCTA; qPCR forward primer for *epaA*	This study
RT-*epaA* -R	ACCATTCCACCAGCCAAA; qPCR reverse primer for *epaA*	This study
RT-*epaE* -F	AGGGATCAAACCGTCAGAAC; qPCR forward primer for *epaA*	This study
RT-*epaE* -R	GGCCCATTACCTCAACAGAA; qPCR reverse primer for *epaA*	This study
RT- *epaOX* -F	CCACGTTAACTCGTGCAGTA; qPCR forward primer for *epaOX*	This study
RT- *epaOX* -R	CCACTGTCATCAGAAGAACCA; qPCR reverse primer for *epaOX*	This study
RT- 11714 -F	GGAAGCATTACCGCTTCTAAGT; qPCR forward primer for OG1RF_11714	This study
RT- 11714 -R	ATGCAAGGTCGCTATATTCTGG; qPCR reverse primer for OG1RF_11714	This study
RT- 11710 -F	GTGTATGGTTTGTCTTAGCGTTAG; qPCR forward primer for OG1RF_11710	This study
RT- 11710 -R	TCGGCCTTCATAGAAACCTATAC; qPCR reverse primer for OG1RF_11710	This study

RT- 11100 *-F*	AAGGGCAAGCATTTCAAGCG; qPCR forward primer for OG1RF_11100	This study
RT- 11100 *-R*	TCTTGACGGTCACGTTCTGC; qPCR reverse primer for OG1RF_11100	This study
RT- 11101 *-F*	CCAATGGCTTGGCAACTGAC; qPCR forward primer for OG1RF_11101	This study
RT- 11101 *-R*	GCGAACGAACGTGCATTTTG; qPCR reverse primer for OG1RF_11101	This study
RT- 11104 *-F*	GGGAATGGCACCCTGAAAGA; qPCR forward primer for OG1RF_11104	This study
RT- 11104 *-R*	CTTCGCGCTTGGCTTTTTGA; qPCR reverse primer for OG1RF_11104	This study
RT- 11105 *-F*	TTGGAAAGGTGGCGGAATAG; qPCR forward primer for OG1RF_11105	This study
RT- 11105 *-R*	TCTGCTTTGATACTGGCTAAGG; qPCR reverse primer for OG1RF_11105	This study
RT- 11109 *-F*	GCTTTGGAGAACGCTGAACG; qPCR forward primer for OG1RF_11109	This study
RT- 11109 *-R*	TTTTGACAGTCTTGCGCTCG; qPCR reverse primer for OG1RF_11109	This study
RT- 11115 *-F*	CTCAACCGGATCGTGCTTATT; qPCR forward primer for OG1RF_11115	This study
RT- 11115 *-R*	CCTTGGTAGCGAATGGATCATAG; qPCR reverse primer for OG1RF_11115	This study
RT-16S -F	CGCTTCTTTCCTCCCGAGT; qPCR forward primer 16S rRNA gene	This study
RT-16S -F	GCCATGCGGCATAAACTG; qPCR reverse primer 16S rRNA gene	This study

Cm^R - chloramphenicol resistant; Rf^R - rifampicin resistance; Fa^R - fusidic acid resistance; SpR - spectinomycin resistance; Restriction enzyme sites are underlined

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