

S3 Table. Strains and plasmid list used in this study.

| Strain name | Parental background | Source | Description | Reference |
|------------------------------|---------------------|------------------|---|-------------|
| <i>Enterococcus faecalis</i> | | | | |
| DK11 | JH2-2 | Clinical | WT plasmidless strain derived from JH2 | 74 |
| DK10 | OG1RF | Oral | WT plasmid free, commensal; Rif ^r , Fus ^r | 77 |
| DK13 | OG1RF | -- | <i>ΔepaB</i> | Gilmore Lab |
| DK255 | OG1RF | -- | <i>ΔebpABC</i> | 26 |
| DK256 | OG1RF | -- | <i>ΔebpA</i> | 26 |
| DK29 | OG1RF | -- | pMV158-GFP, conjugation donor | 78 |
| DK30 | V583 | Blood | Clinical WT Van ^r | 75 |
| DK1 | VE14089 | Blood | Plasmidless strain derived from V583 | 83 |
| DK2 | VE14089 | -- | <i>ΔepaX</i> | 46 |
| DK3 | VE14089 | -- | <i>ΔepaX::epaX</i> ; Erm ^r | 46 |
| DK287 | VE14089 | -- | WT, pMV158-GFP, Tet ^r | This study |
| DK288 | VE14089 | -- | <i>ΔepaX</i> , pMV158-GFP, Tet ^r | This study |
| DK8 | MMH594 | Blood | Clinical WT Erm ^r , Gen ^r | 76 |
| DK250 | MMH594 | -- | WT, empty pAT28 vector; Sp ^r | This study |
| DK19 | MMH594 | -- | WT, pMV158-GFP, Tet ^r | This study |
| DK20 | MMH594 | -- | WT, pMV158-mCherry, Tet ^r | Gilmore lab |
| DK16 | MMH594 | -- | <i>glnA::TnM</i> ; Cm ^r | This study |
| DK251 | MMH594 | -- | <i>glnA::TnM</i> , empty pAT28 vector; Cm ^r , Sp ^r | This study |
| DK252 | MMH594 | -- | <i>glnA::TnM</i> , pAT28- <i>glnRA</i> ; Cm ^r , Sp ^r | This study |
| DK18 | MMH594 | -- | <i>glnA::TnM</i> ; pMV158-GFP; Cm ^r , Tet ^r | This study |
| DK14 | MMH594 | -- | <i>rpiA::TnM</i> ; Cm ^r ; | This study |
| DK253 | MMH594 | -- | <i>rpiA::TnM</i> , empty pAT28 vector; Cm ^r , Sp ^r | This study |
| DK15 | MMH594 | -- | <i>rpiA::TnM</i> , pAT28- <i>rpiA</i> ; Cm ^r , Sp ^r | This study |
| DK17 | MMH594 | -- | <i>rpiA::TnM</i> ; pMV158-GFP; Cm ^r , Tet ^r | This study |
| DK267 | MMH594 | -- | Δ EF2170 | This study |
| DK21 | 1230 | Clinical | WT | 73 |
| <i>Escherichia coli</i> | | | | |
| DK91 | DH5 α | -- | F ⁻ Φ 80 <i>lacZ</i> Δ M15 Δ (<i>lacZYA-argF</i>) U169 <i>recA1 endA1 hsdR17</i> (rk ⁻ , mk ⁺) <i>phoA supE44 λ thi⁻¹ gyrA96 relA1</i> | Stratagene |
| <i>Staphylococcus aureus</i> | | | | |
| DK43 | MN8 | Urogenital tract | WT | 32 |
| DK44 | MN8 | -- | <i>Δica</i> , PNAG(PIA) deficient strain | 32 |
| <i>Candida albicans</i> | | | | |
| DK80 | SC5314 | -- | Prototrophic clinical isolate | 80 |
| Plasmids | | | | |
| pMINIMAD | | | Thermosensitive plasmid; Amp ^r | 81 |
| pLT06 | | | Deletion construct used for amplification of Cm ^r cassette (<i>camR</i>) | 10 |
| pIC333 | | | Deletion construct used for amplification of Sp ^r cassette (<i>spcR</i>) | 79 |
| pJR02 | | | pMINIMAD vector with EF2170:: <i>spcR</i> deletion construct; Amp ^r | This study |
| pJR03 | | | Derived from pJR02 with Cm ^r from pLT06 vector; Amp ^r , Cm ^r | This study |
| pAT28 | | | Complementation vector | 82 |
| pJR01 | | | <i>glnRA</i> complementation vector | This study |
| pAH01 | | | <i>rpiA</i> complementation vector | This study |
| pMV158 | | | GFP cassette; Cm ^r , Tet ^r | 92 |

Amp^r, ampicillin resistant; Rif^r, rifampin resistant; Fus^r, fusidic acid resistant; Sp^r, spectinomycin resistant; Tet^r, tetracycline resistant; Cm^r, chloramphenicol resistant; Em^r, erythromycin resistant. *E. faecalis* MMH594 strains harboring pAT28 vector and derivatives required 750 μ g/mL spectinomycin. Transposon insertion strains required 10 μ g/mL chloramphenicol. Fluorescent reporter strains harboring pMV158 derivatives needed 15 μ g/mL tetracycline. *E. faecalis* harboring pLT06 derivatives required 15 μ g/mL chloramphenicol. pMINIMAD derivatives required 100 μ g/mL ampicillin. WT: Wild type.