**Table S2: Comparative transcriptome analysis of *E. faecium* E1162 during mid-exponential growth in the presence (20 µg ml−1) and absence of ampicillin.**

|  |  |  |  |
| --- | --- | --- | --- |
| **LocusTag*a***  | **Accession code** | **Annotation** | **Expression ratio BHI+amp/BHI** |
| EfmE1162\_0046 | ZP\_06675891 | PyrR bifunctional protein | 1.9 |
| EfmE1162\_0047 | ZP\_06675892 | uracil permease | 2.0 |
| EfmE1162\_0048 | ZP\_06675893 | aspartate carbamoyltransferase | 1.9 |
| EfmE1162\_0049 | ZP\_06675894 | Dihydroorotase | 2.1 |
| EfmE1162\_0050 | ZP\_06675895 | carbamoyl-phosphate synthase, small subunit | 1.9 |
| EfmE1162\_0051 | ZP\_06675896 | carbamoyl-phosphate synthase, large subunit | 1.9 |
| EfmE1162\_0052 | ZP\_06675897 | dihydroorotate dehydrogenase electron transfer subunit | 2.0 |
| EfmE1162\_0053 | ZP\_06675898 | dihydroorotate dehydrogenase B, catalytic subunit | 1.9 |
| EfmE1162\_0054 | ZP\_06675899 | orotidine 5'-phosphate decarboxylase | 2.0 |
| EfmE1162\_0055 | ZP\_06675900 | orotate phosphoribosyltransferase | 2.0 |
| EfmE1162\_0229 | ZP\_06676074 | conserved hypothetical protein | 1.8 |
| EfmE1162\_0230 | ZP\_06676075 | accessory gene regulator protein B, putative | 1.7 |
| EfmE1162\_0231 | ZP\_06676076 | hypothetical protein | 1.8 |
| EfmE1162\_0443 | ZP\_06676288 | hypothetical protein | 1.7 |
| EfmE1162\_0683 | ZP\_06676528 | peptidoglycan-binding LysM | 0.5 |
| EfmE1162\_2616 | ZP\_06678315 | aggregation promoting protein | 0.4 |

*a* All genes exhibiting significantly different expression during mid-exponential growth (OD660 = 0.3) in BHI and BHI with 20 µg ml−1 ampicillin for all four probes per gene (Bayesian P-value <0.001) in microarray hybridizations are included.