

Supplementary Table 5: Single nucleotide polymorphism matrix of *Enterococcus faecalis* ST4 isolates.

	MEZEF110	2684_P9_CL_A7	2539_90
MEZEF110		6581	6693
2539_90	6581		553
2684_P9_CL_A7	6693	553	

Table. Estimates of Evolutionary Divergence between Sequences

The number of base differences per sequence from between sequences are shown. This analysis involved 3 nucleotide sequences. All positions containing gaps and missing data were eliminated (complete deletion option). There were a total of 6912 positions in the final dataset. Evolutionary analyses were conducted in MEGA X [1]

1. Kumar S., Stecher G., Li M., Knyaz C., and Tamura K. (2018). MEGA X: Molecular Evolutionary Genetics Analysis across computing platforms. *Molecular Biology and Evolution* 35:1547-1549.

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