

M. putrefaciens



Me. florum L1



S. taiwanense CT-1



S. diminutum CUAS-1



S. culicicola AES-1



S. apis B31



S. sabaudiense Ar-1343



A. brassicae O502



A. palmae J233



A. laidlawii PG-8A



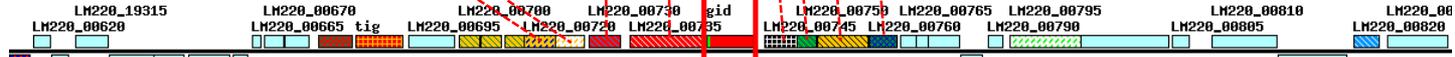
S. aureus N315



B. subtilis 168



L. monocytogenes J1-220



molluscates

non-molluscates

Figure S5. Genomic context of *trmFO* homologs. Analysis of synteny and genome context was performed using the MGD database. Homologous *trmFO* genes are framed and coloured in red. Other homologs are coloured the same and connected with red dotted lines.