Supplementary materials:

Complete Genome Sequence and Analysis of a ST573 Multidrug-Resistant Methicillin-Resistant *Staphylococcus aureus* SauR3 Clinical Isolate from Terengganu, Malaysia

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Please refer Excel for supplementary tables,
Table S1: List of genes encoding virulence factors detected in SauR3 genome.

Table S2: List of *Staphylococcus aureus* strains used to construct the phylogenetic tree depicted in Figure 1.