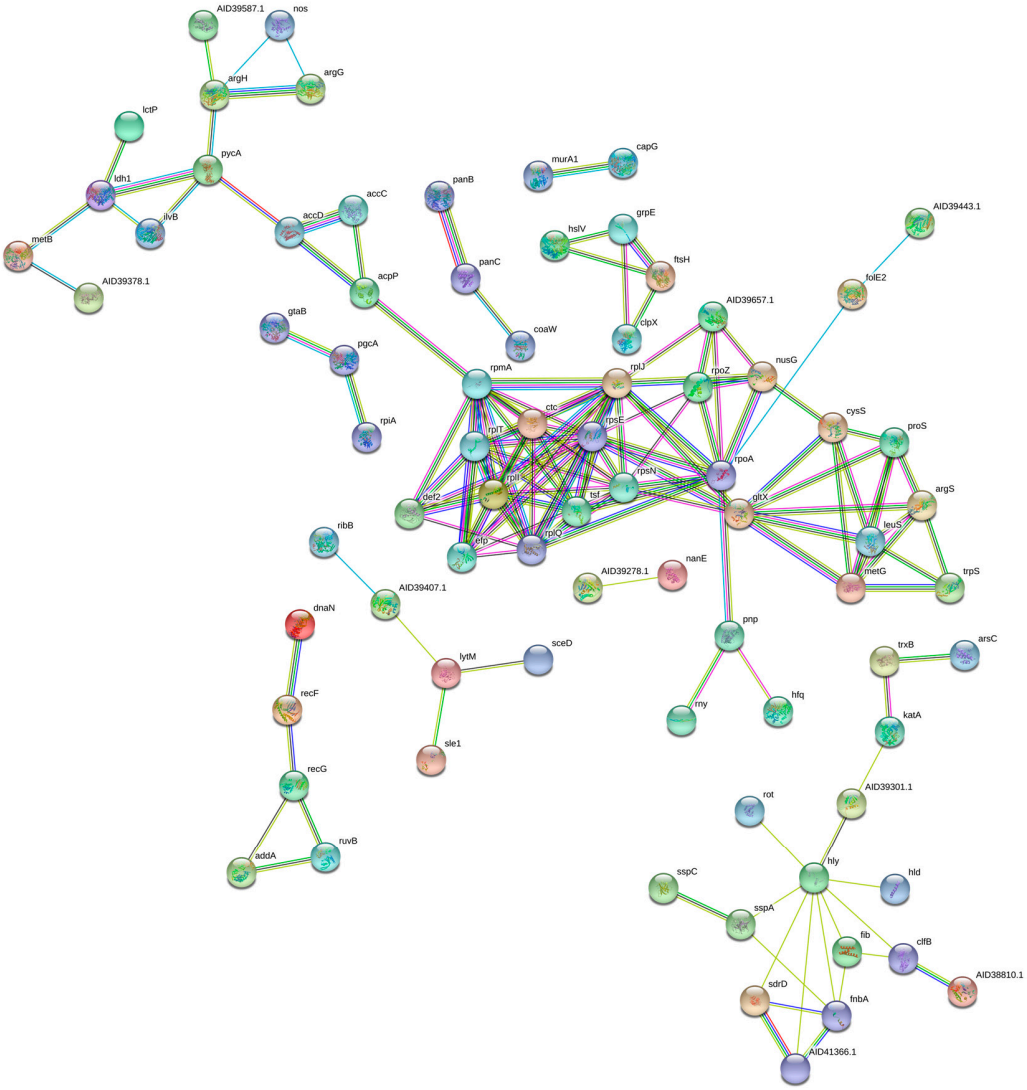

Figure S1. The network of DREPs of *S. aureus* biofilm was analysed by STRING software (v.10.0). High-confidence interactions (≥0.7) and omitted nodes that are not connected in the network in STRING software were selected for this study.