



Supplementary Materials

Antibacterial Activity and Molecular Docking Studies of a Selected Series of Hydroxy-3-arylcoumarins

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Figure S1. Validation of docking protocol. Ligand superimposition: co-crystal and docked conformations.



Figure S2. Docked conformation of the investigated ligands in (PDB id: 1JIL) Staphylococcus aureus tyrosyl-tRNA synthetase proteinstructure.



Figure S3. Docked conformation of the investigated ligands in (PDB id: 2XCS) S. aureus Gyrase complex.