



Figure S8. Predicted 3D structures of phenol hydroxylase protein component P2 aligned to the characterized PDB structure (1HQI)

(A) LSNL_2977 of *Acinetobacter lwoffii* NL1 and (B) LSNL_2977 of *A. lwoffii* NL115.
The PDB structure 1HQI is highlighted in yellow.