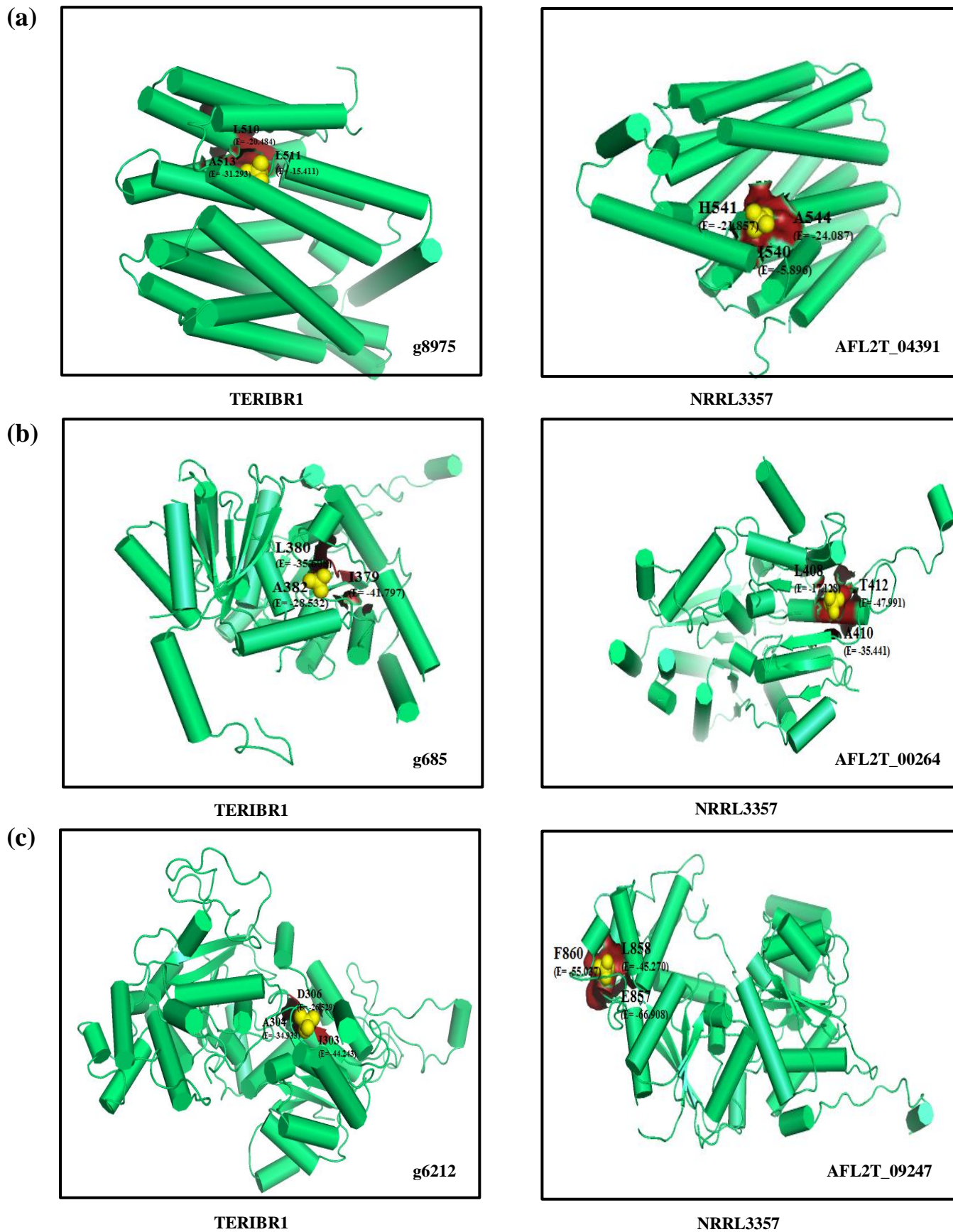
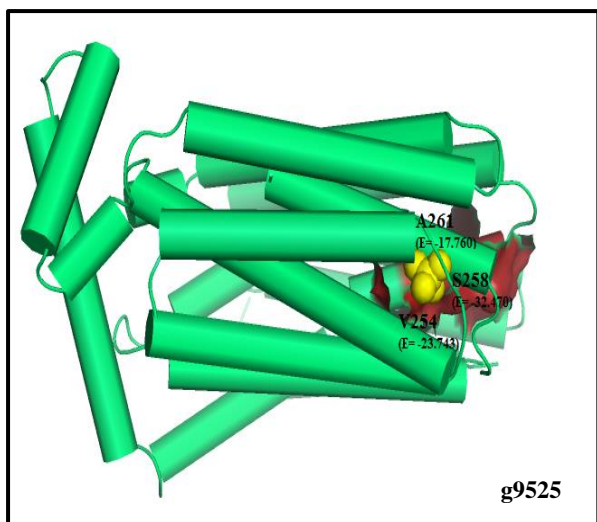


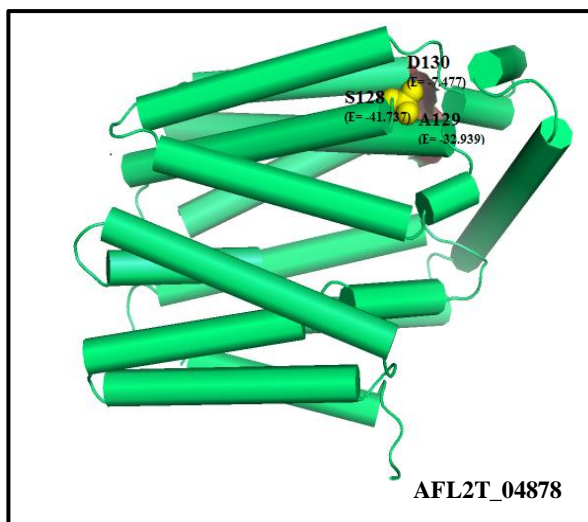
Figure S1. Protein-ligand interaction observed with four homologous pairs of protein of *A. flavus* strains TERIBR1 and NRRL3357



(d)



TERIBR1



NRRL3357

Chromate ion is depicted in yellow. Amino acid residues present in the proximity of ligand docking position are marked in black, binding sites predicted by 3DLigandSite are marked in fire brick and surfaced.