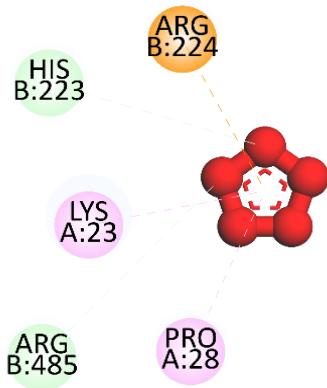
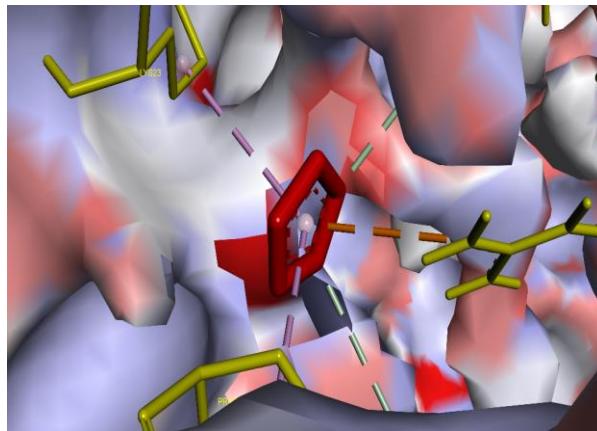
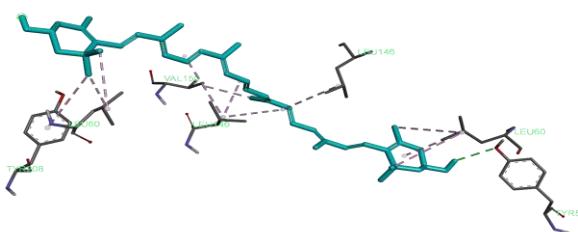


Molecular Docking Sample Projects



Interactions
Carbon Hydrogen Bond
Pi-Alkyl
Pi-Cation



The figures above illustrate the molecular interaction and docking poses of ACHE with Furan. Furan interacts moderately with the sidechain amino acid residues at the active site of ACHE. Two hydrogen bonds were detected on HIS 223 (3.6 Å) and ARG 485 (3.6 Å)