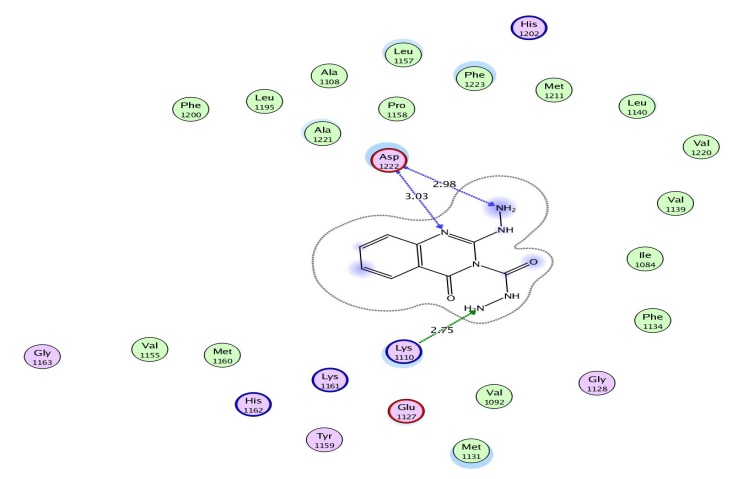
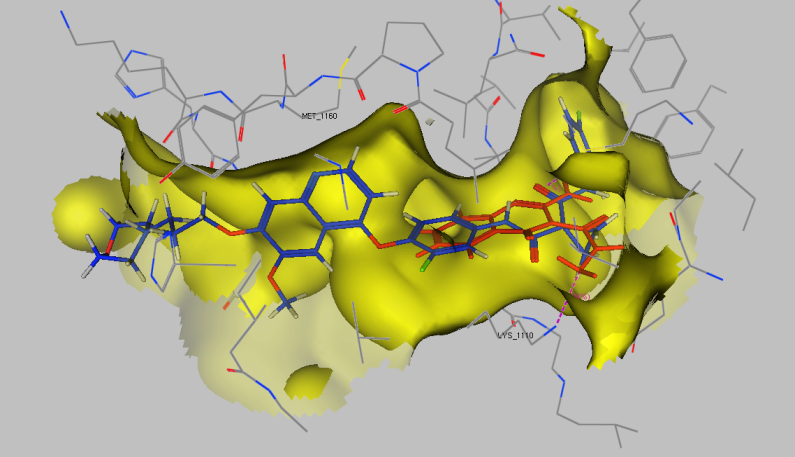


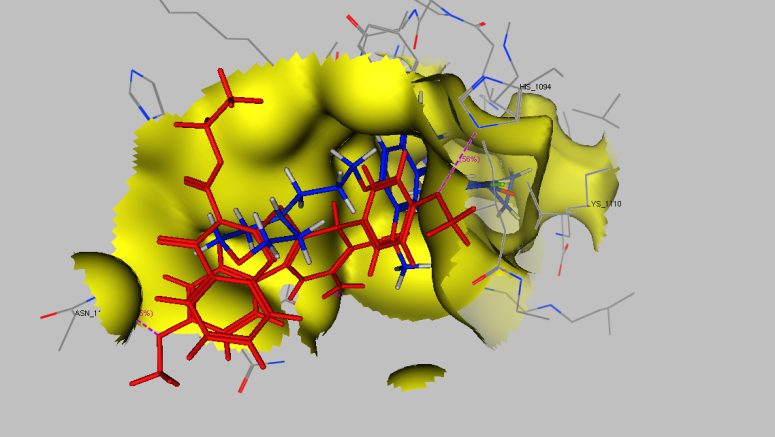
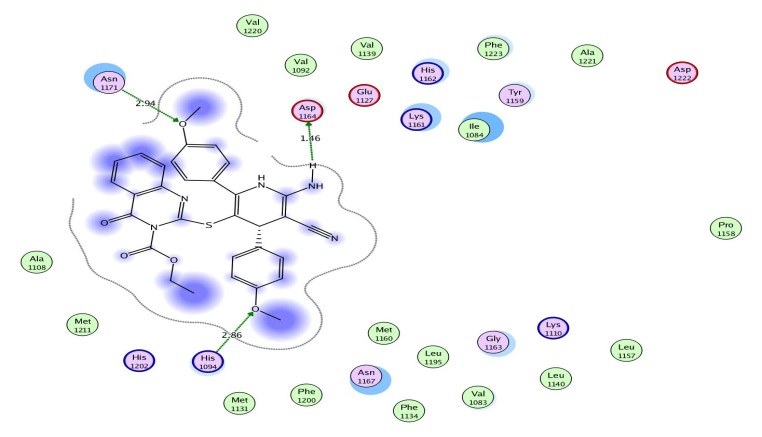
1. (b)

**Figure 2.** Interactions of XL880 (Foretinib) with the amino acid residues of the active site of c-Met 3D(a) and 2D(b)



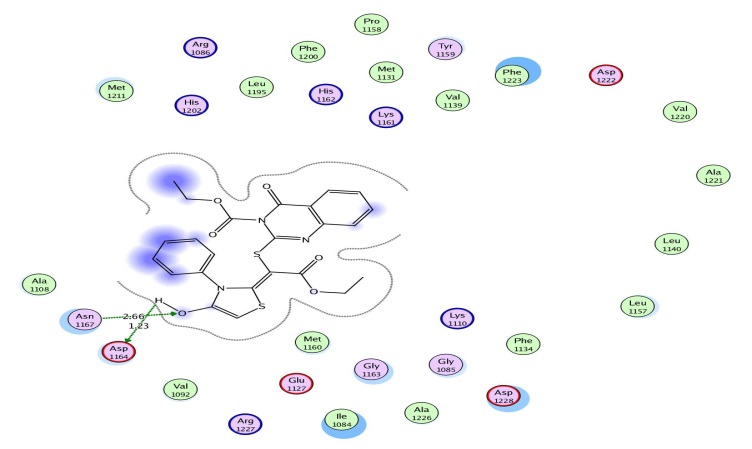
1. (b)

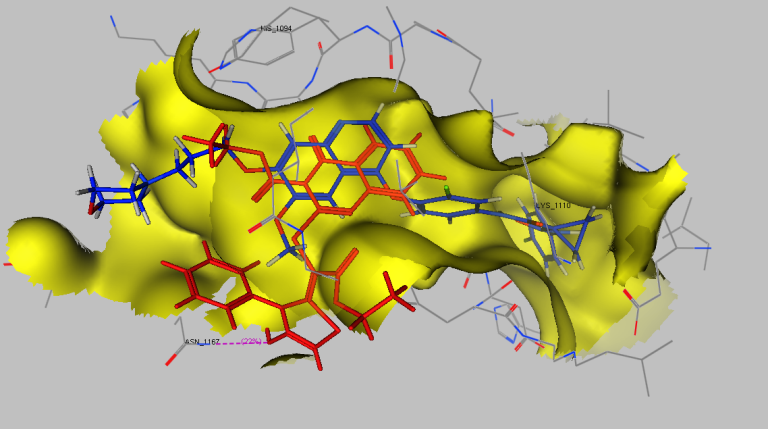
**Figure 3.** (a) The superposition of **Foretinib** (Blue) and compound **13** (red) docked in the binding site of c-Met, the dotted lines represent H-bonding interactions (b) 2D ligand interaction of **13** in binding site of c-Met.



1. (b)

**Figure 4.** (a) The superposition of **Foretinib** (Blue) and compound **21e** (red) docked in the binding site of c-Met, the dotted lines represent H-bonding interactions (b) 2D ligand interaction of **21e** in binding site of c-Met.





**Figure 5.** (a) The superposition of **Foretinib** (Blue) and compound **24a** (red) docked in the binding site of c-Met, the dotted lines represent H-bonding interactions (b) 2D ligand interaction of **24a** in binding site of c-Met.