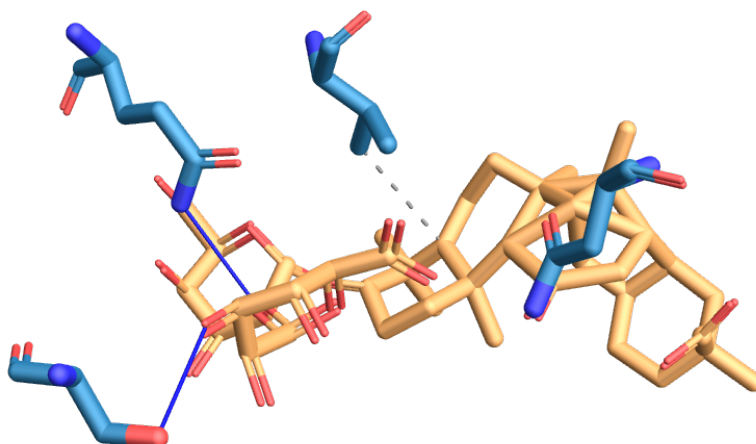


## 6lu7\_pdb-Glycyrrhizic

### Docking Scores

No.	Affinity	rmsd l.b	rmsd u.b
1	-6.1	0	0
2	-6	2.141	5.255
3	-5.7	3.41	9.15
4	-5.6	2.172	3.141
5	-5.4	2.275	3.934
6	-5.3	2.291	3.4
7	-5	3.279	10.79
8	-5	3.524	6.357
9	-4.9	2.379	4.326

### Visualisation



### Interactions

Hydrophobic Interactions Found

Hydrophobic Interactions Found

Hydrogen Bonds Found

No Water Bridges Found

No Salt Bridges Found

No Pi Stacks Found

No Pi Cation Interactions Found

No Halogen Bonds Found

No Metal Complexes Found

## Hydrophobic Interactions

No.	Res.	AA	Dist	Ligand Atom	Proton Atom
1	3	VAL	3.66	2421	2388

## Hydrogen Bonds

No.	Res.	AA	Dist H-A	Dist D-A	Don Angle	Protisdon?	Sidechain?	D. Atom	A. Atom
1	46	SER	2.73	3.34	121.63	False	True	2475[O3]	346[O3]
2	142	ASN	2.48	3.22	132.48	True	True	1103[Nam]	2461[O2]
3	189	GLN	2.38	3.33	161.89	True	True	1464[Nam]	2470[O3]

## Figures

