Table 2: Docking of all molecules against Transthyretin through PyRx

S.No	Name Of Molecule	Binding Affinity	Mode	RMSD Upper Bound	RMSD Lower Bound
1.	7-desacetyl-7-benzoylazadiradione	-10.6	0	0.0	0.0
2.	Lict_17_hydroxyazadiradiaze	-8.4	0	0.0	0.0
3.	Gedunin	-8.9	0	0.0	0.0
4.	n-hexacosonal	-4.9	0	0.0	0.0
5.	Nimbin	-8.0	0	0.0	0.0
6.	Nimolide	-7.8	0	0.0	0.0
7.	Nimbolinin	-8.8	0	0.0	0.0
8.	Polyphenolic flavonoids	-10.6	0	0.0	0.0
9.	Quercetin	-8.1	0	0.0	0.0
10.	Salannin	-9.2	0	0.0	0.0