

# Molecular Docking Results (Boltz-2)

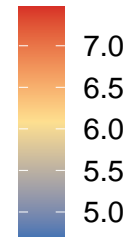
Predicted affinity between candidate ligands and hub proteins

pIC50 (higher is better)

Protein (Hub Genes)

AKT1  
MTOR  
PIK3CA  
ENO1  
SLC27A5

AKT1	NA	6.62	5.69	5.77	NA	6.06	6.04	NA	5.45	NA	5.77	5.14	5.79	5.26	NA
MTOR	6.74	7.38	6.56	6.31	6.39	6.12	5.42	5.28	5.61	5.82	5.53	6.16	5.41	5.17	4.70
PIK3CA	6.60	5.88	6.37	5.80	5.88	6.63	6.20	5.93	6.12	5.86	5.43	6.21	5.74	5.84	4.95
ENO1	6.59	6.00	7.46	7.13	6.44	6.22	6.42	6.59	6.01	5.80	5.91	5.92	5.48	5.37	5.03
SLC27A5	7.24	7.00	6.39	6.73	6.65	5.84	6.22	6.14	6.35	6.10	6.73	5.84	5.97	6.24	4.82



5,7-dimethoxyflavan  
Aeruginol  
Curdione  
Furanodiene  
(S)-(+)-2-Heptanol  
(-)-Isodene  
(S)-(+)-4-methyl-4-vinylcyclohexene  
Undecan-2-one  
(+)-alpha-thujene  
Nonan-2-one  
Dodeca-3,7-diene  
Curcumenone  
Curzerenone  
(-)-Sabinene  
Eucalyptol

Ligand (Active Compounds)