

Docking scores of Fenebrutinib binding to predicted targets			
Uniprot ID	Protein name	PDB ID	Docking score (kcal/mol)
Q96B26	Exosome complex component RRP43	2NN6_C	-7.9
Q5JRX3	Presequence protease, mitochondrial	4L3T_A	-10.8
Q99720	Sigma non-opioid intracellular receptor 1	5HK1_A	-9.6
Q5VT66	Mitochondrial amidoxime-reducing component 1	6FW2_A	-10.4
P29122	Proprotein convertase subtilisin/kexin type 6	AF-P29122-F1 (157-622)	-8.5
Q96K12	Fatty acyl-CoA reductase 2	AF-Q96K12-F1 (1-478)	-10.1
O94973	AP-2 complex subunit alpha-2	AF-O94973-F1 (3-622)	-8.6
Docking scores of NMS-P715 binding to predicted targets			
Uniprot ID	Protein name	PDB ID	Docking score (kcal/mol)
Q9UN86	Ras GTPase-activating protein-binding protein 2	5DRV_A	-9.5
P67870	Casein kinase II subunit beta	1QF8_A	-8.6
Q96B26	Exosome complex component RRP43	2NN6_C	-9.3
P62877	E3 ubiquitin-protein ligase RBX1	2HYE_D	-7.9
P61962	DDB1- and CUL4-associated factor 7	AF-P61962-F1 (9-341)	-8.7
Q9NXH9	tRNA (guanine(26)-N(2))-dimethyltransferase	AF-Q9NXH9-F1 (53-556)	-9.0
Q9NQT4	Exosome complex component RRP46	2NN6_D	-8.6

Table 1: Docking scores for Fenebrutinib and NMS-P715