

Docking scores of Fenebrutinib binding to predicted targets		
Uniprot ID	Protein name	PDB ID
Q96B26	Exosome complex component RRP43	2NN6_C
Q5JRX3	Presequence protease, mitochondrial	4L3T_A
Q99720	Sigma non-opioid intracellular receptor 1	5HK1_A
Q5VT66	Mitochondrial amidoxime-reducing component 1	6FW2_A
P29122	Proprotein convertase subtilisin/kexin type 6	AF-P29122-F1 (157-62)
Q96K12	Fatty acyl-CoA reductase 2	AF-Q96K12-F1 (1-478)
O94973	AP-2 complex subunit alpha-2	AF-O94973-F1 (3-622)
Docking scores of NMS-P715 binding to predicted targets		
Uniprot ID	Protein name	PDB ID
Q9UN86	Ras GTPase-activating protein-binding protein 2	5DRV_A
P67870	Casein kinase II subunit beta	1QF8_A
Q96B26	Exosome complex component RRP43	2NN6_C
P62877	E3 ubiquitin-protein ligase RBX1	2HYE_D
P61962	DDB1- and CUL4-associated factor 7	AF-P61962-F1 (9-341)
Q9NXH9	tRNA (guanine(26)-N(2))-dimethyltransferase	AF-Q9NXH9-F1 (53-5)
Q9NQT4	Exosome complex component RRP46	2NN6_D

Table 1: Docking scores for Fenebrutinib and NMS-P715