

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0 P24539	AT5F1_HUMAN	Homo sapiens	ATP synthase F(0) complex subunit B1, mitochondrial	24.097408	NaN	NaN	30444036;28510447	MLSRVLSAAATAAPSLKNAFLGP GVLQATRTFHTGQPHLVPVPLPEY GGKVRYGLIPEEFFQFLYPKTGVTGP YVLGTGLILYALSKEIYVISAETFTALS VLGVMVYGIKKYGPVADFADKLE QKLAQLEEAKQASIQHIQNAIDTEKS QQALVQKRHYLFDVQRNNIAMALE VTYRERLYRVYKEVKNRLDYHISVQ NMMRRKEQEHMINWVEKHVVQSIS TQEKETIAKCIADLKLAKKAQAQPV VM	True	False	2.5	4.517	5.0	1.323	1.02	1.405	1.289