

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q8CHS8	VP37A_MOUSE	Mus musculus	Vacuolar protein sorting-associated protein 37A	37.053079	T171;S172;S174;T178;S180;S201;T208;T209	S18	40885482;40021952;36852467	MSWLFPLAKSASSAAGSPAGLTSL QQKQRLIESLRNSHSSIAEIQKDVE YRLPFTVNNLTININILLPPQFPQEK VISVYPPIRHHLMDSQGLYVTSPLVS NFTMHSDLGKIIQSLLEDFWKNPPV LAPTSTTFPYLYSNPGGMPPYPSQGF PFLPPYPPPEANRNITSLSVADTVSS STTSYTAAPVAPSFGLSSLPVPPT TESSASVNONQFGYKMPDIPDAFPE LSELSVSQLTDMNEQEEVLEQFLM LPQLKQIITDKEDLVKNIEELARKNL LLEHSLEGGKQTVLDKYELLQMK TFEKKMQRQHELSESCSASALQARL KVAAHEAEEESDNIAEDFLEGKTEID DFLNSFKEKRTICHCRRAKEEKLHQ VIAMHSQFHAPL	False	False	2.966	2.867	2.2	1.093	0.825