

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 Q01405	SC23A_MOUSE	Mus musculus	Protein transport protein Sec23A	25.295579	NaN	T308	37453647;33300544	MTTYLEFIQQNEERDGVRFWSNVW PSSRLEATRMVVPVAALFTPLKERPD LPPIQYEPVLCSTRITCRAVLNPLCQV DYRAKLWACNFCYQRNQFPPTYAGI SELNQPAELLPQFSSIEYVLRGPQM PLIFLYVVDTCIEDEDLQALKESMQ MSLSLLPPTALVGLITFGRMVQVHE LGCEGISKSYVFRGTDLSAKQLQE MLGLSKVPVTQATRGPVQVQPPPSN RFLQPVQKIDMNLTDLLGELQRPW PVPQGRPLRSSGVALSIAVGLLECT FPNTGARIMMFIGGPATQGGPMVVG DELKTPIRSWHDIEKDNKYVKKGT KHFEALANRAATTGHVIDIYACALDQ TGLEEMKCCPNLTGGYVMVMDGSFN TSLFKQTFQRVFTKDIHGQFKMGFG GTLEIKTSREIKISGAIGPCVSLNSKG PCVSENEIGTGGTCQWKICGLSPTTT LAIYFEVNVNQHNAIPQGGRGAVQF VTQYQHSSGQRRIRVTTIARNWADA QTQIQNIAASFDOEAAAAILMARLAIY RAETEEGPDVLRWLDRQLIRLQCKF GEYHKDDPNSFRFSETFSLYPQFMF HLRRSPFLQVFNNSPDESSYRHHF MRQDLTQSLIMIPIYAYFSFGPPE PVLLDSSSILADRILLMDTFFQILIYH GETIAQWRKSGYQDMPEYENFRHL LQAPVDDAQEILHSRFPMPRYIDTE HGGSQARFLLSKVNPSQTHNNMYA WGQESGAPILTDVSLQVFMHLKK LAVSSAA	True	True	4.555	2.68	1.424	4.516	5.0	1.934	1.844