

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extra regio
Q15437	SC23B_HUMAN	Homo sapiens	Protein transport protein Sec23B	35.300052	S765	NaN	34229054;30397120;20305658;33214551;40307207;40914422;35008409;34019948	MATYLEFIQQNEERDGVRFSSWNVW PSSRLEATRMVVPLACLTLPLKERPD LPPVQYEPVLCSPRTCKAVLNPLCQV DYRAKLWACNFCFQRNQFPAYGGI SEVNQPAELMPQFSTIEYVIQGAQS PLIFLYVVDTCLEEDDLQALKEQLQM SLSLPPDALVGLITFGRMVQVHEL CEGISKSYVFRGTDLTAKQIQDML GLTKPAMPMPQARPAQOEHPFAS SRFLQPVHKIDMNLTDLLGELQRPD WPVTQGKRPLRSTGVALSIAVGLLEG TFPNTGARIMLFTGGPPTQPGMVV GDELKIPRSWHDIEKDNARFMKKA TKHYEMLANRTAANGHCIDIYACAL DOTGLEMKCCANLTGGYVMVGDS FNLSLQKQTFQRIFTKDFNGDFRMA FGATLDVKTRELKIAGAIGPCVSLN VKGPCVSENELGVGGTSQWKICGLD PTSTLGIYFEVNVQHNTPIPQGGGGA IQFVTHYQHSSTQRRIRVTIARNWA DVQSOLRHIEAADFQEAAMVLMARL GVFRAESEEGPDVLRWLDRLRLC QKFGQYNKEDPTSFRLSDSFSLYPQ FMFHLRRSPFLQVFNNSPDESSYR HHFARQDLTQSLIMIPILYSYFHHG PPEVLLDSSSILADRILLMDYFFQIV IYLGETIAQWRKAGYQDMPEYENFK HLLQAPLDDAQEILQARFPMRYINT EHGGSQARFLLSKVNPSQTHNLYA WGQETGAPILTDVSLQVFMHLLK LAVSSAC	True	True	4.537	2.562	1.184	5.0	2.117	1.578	1.356