

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
Q9Z1Z0	USO1_MOUSE	Mus musculus	General vesicular transport factor p115	32.695381	NaN	S50;S940	25153642;28528544;36288343	MNFLRGVMGGQSAGPQHTAEATIQ KLCDRVASSTLLDERRNAVRALKSL SKKYRLEVGIQAMEHLIHVLQTDRS DSEIIAYALDTLYNIISNDEEELEEN STRQSEDLGSGQFTEIFIKQPENVTLL LSLLEEFDFHVRWPGVRLTSLKQ LGPPVQQIILVSPMGVSRMLDLLADS REIIRNDGVLLQALTRSNNGAIQKIVA FENAFERLLDIITEEGNSDGGIVVED CLILLQNLKTNNSNQFFKEGYSYI QRMKPWFVEVGEENSGWSAQKVTNL HLMQLVLRVLPVPTNPPGATSSCQK AMFQCGLLQQLCTILMATGIPADILT ETINTVSEVIRGCQVNDYFASVNP SNPPRPAIVLLMSMVNERQPFVLR CAVLYCFQCFLYKNEKGQGEIVATLL PSTIDATGNSVSAGQLCGGLFSTDS LSNWCAAVALAHALQGNATQKEQL LRVQLATSIGNPPVSLLOQCTNILSQ GSKIQTRVGLLMLLCTWLSNCPAVT HFLHNSANVPFLTGQIAENLGEEEQ LVQGLCALLGISIYFNDSLENYTK EKLKQLIEKRIGKENFIEKLGFIKHE LYSRASQKPPNPPSPPEYMIFDHEF TKLVKELEGVITKAIYKSSEEDKKEE EVKKTLEQHDNIVTHYKNMIREQDL QLEELKQVSTLKCQNEQLQTAVTQ QASQIQQHKDQYNLLKVQLGKDNH HOGSHGDGAQVNGIQPEISRLREE IEELKSQALLQGQLAEKDSLIENLK SSQASGMSEQASATCPPRPDPEQVAE LKQELTALKSQLCSQSLEITRLQTEN CELLQRAETLAKSVPEGESEHVSA AKTTDVEGRLSALLQETKELKNEIKA LSEERTAIQKQLDSSNSTIAILQTEKD KLDLEVTDSKKEQDDLLVLLADQDQ KILSLKSKLDLGHVPEEEDESGDQ EDDDDEIDDGDKQDI	True	False	5.0	3.101	0.98	3.617	3.845	1.319	1.095