	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	A0A0B4K6T1	A0A0B4K6T1_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MSTQTRSGGGGGGGHARNQKKSNA SNSGGGGTGHHDGVSHAAAAGKKG GQDASKTDKPEKAQPKATTEQLRIA QITNSTTEDPQINEKVLLLLTMTQRS EEEVCCALNECDYDLEAAANFLIEE LPQGAFAKYEKKRKNKAANNTADGA AGDGDWADGNGNADRREKSRNRS SNRGGTRGSSDSRGWRGRETRENE RNQRESREPWSGQNAGQDRGDDRA NDNYRGQRNGGGRSGPGGGGRGG GFVSRSGRGGGRMGGRTGGPRGDR GSGGPGGAYGSGRGGNANEDHHEV ELWDNTIAQNAEKQQQAHDDAWG DWNNEEYEGSLKDSKVFTTSNLATQ SAANVVSGTGASVTAVPAAAGTEISA PPGLEHQLVQQGSHLEESSSSGPAA VTPPATLSGSATTPLLQYSAAVSNPP PQLQSQGTQSGAGTGASAAAGGGAG STPSSFVSASPDTFSSAASAAATLVH QAQKQQQLQQQTTPIKPSATLSVEQ SQYFNSLASQGVSPGSVPVQSAPAG YAQNPVAAYSQTSTSVGVSQYPNTY ANVFASGTAAGAGTAEQSQQQPQIR RARVKLPPPSKIPASAVEMPGDNAL NNIGYLDVQFGALDFGTDDGFEPLP EKVGSGFSIDGQQQQQQQPDDYQSKS QQQQQVTLAAGLQSSQISDALNAAG YTSRSTSQQQQGVSSAVNATIDQLT KSDPYGQTGGSGNAYQNAYQSSGAS KTASGFPTTAPGGYSSSTYANVQSSV ANSYQQQGYGSYQPSSYQQQAGSGA QSGTGAVSGGGGTATQNIPVGGSSS QNSTSGNASSAYLTSGYSTPQSAYQ SSQSVYGNTGLSNSSGFSGSASNAS SQYANFSASAKLKDATTASSAAHYD SVSTSSGVSSNSGSTGNGGVVSGQT GANQAAVSNNNSVSGSSSVSNVTA GVASGNVAGVGGGVSQSGVSSGVG VPGGSASSVGVNVNNNSSSASSVGA ATVAQTATGTTAAVLASLTNKNTSSS NSSGSGGSAATTTGNASGQGAGAST GGVGSSSGAGGAGSGGGSGSGLVPT NIQMQGYYDLNYPPASLGAGRDNLG SVTYSAMNDGRFARTDNNSSPVGN VSSTMSQQAGSSAPMLNVPYAYFYG GNVMPGSFQYGTPAIYPQIPAANTAS GQQFPKPSYSAGYGSTSYDTLSQTTQ DYSKGGYSSSVNQQSKTQTVSNQSQ AGTGSDLTSSMYGKGHVALNKVNSY EKQSFHSGTPPPFNMPNTQTAGGTS AQPYGMYLPMPAAGHHNMIHQPIH QDSNSAGQRQQSTSQSKSAGKQGY SPSYWAGQN	None	None	None	None	None	None	None	None	None