

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A0A0B4KFC5	A0A0B4KFC5_DROME	Drosophila melanogaster	NaN	27.156312	NaN	NaN	28604694	MSTQTRSGGGGGGHARNQKKSNA SNSGGGGTGHHDGVSHAAAAGKKG GQDASKTDKPEKAQPKATTEQLRIA QITNSTTEDPQINEKVLVLLMTQRS EEEVCCALNECDYDEAAAANFLIEE LPQGAFAYEKKRKNKAANTADGA AGDGDWADGNGNADRREKSRNRS SNRGGTRGSSDSRGWRGRETRENE RNQRESREPWSGQNAQDRGDDRA NDNYRQQRNGGGRSGPGGGRRGG GFVSRSGRGGRMGGRTGGPRGDR GSGGPGGAYGSGRGGNANEDHHEV ELWDNTIAQNAEKQQQAHDHDAWG DWNNEEYEGSLKDSKVFTTSLNATQ SAANVVSGTGASVTAVPAAAGTEISA PPGLEHQLVQQGSHLEESSSSGPAA VTPPATLGSATTPLLOYSAAVSNPP PQLQSQGTQSGAGTGASAAAAGGGAG STPSSFVSAASPDTFSSAASAAATLVH QAQKQQQLQQQTTPIKPSATLSVEQ SQYFNLSLQGVSPGSPVQSAAPAG YAQNPVAAYSQTSTSVGVSQYPNTY ANVFASGTAAGAGTAEQSQQPQIR RARVKLPPPSKIPASAVEMPGDNAL NNIGYLDVQFGALDFGTDDGFELPL EKVGSGFSDIDGQQQQPDDYQSKS QQQQQVTLAAGLQSSQISDALNAAG YTSRSTSQQQGVSSAVNATIDQLT KSDPYGQTGGSGNAYQNAQYSSGAS KTASGFPTTAPGGYSSSTYANVQSSV ANSYQQQGYGSYQPSYQQQAGSSA QSGTGAVSGGGTATQNIPIVGGSSS QNSTSVSTSSGVSSNSGSTGNGGVV SGQTGANQAAVSNNSVSGSSSVS NVTAGVASGNVAGVGGVVSQSGVSS GVGVPGGSASSVGVNNSNSASS VGAATVAQTATGTTAAVLASLTNKN TSSNSSSGGGSAATTTGNASGQGA GASTGGVSSSGAGGAGSGGGSGSG LVPTNIQMVSQYIQTGLPYYQQPVYS YEELQMMQQRVPHVQGYDLNYP ASLGAGRDNLGVSSTYSAMNDGRFA RTDNNSSPVGNVSTMSQQAGSSA PMLNVPYAYFYGGNVMPGSFYQYGP AIYPQIPAANTASGQQFPKPSYAGY GSTSYDTLSQTTQDYSKGGYSSVN QSKTQTVSNQSQAGTGSDLTSSMY GKGHVALNKVNSYEKQSFHSCTPPP FNMPNTQTAGGTSAPYGMYPMP AAGHHNMIHQPIHQDSNSAGQRQ STSQSKSAGKQGYSPSYWAGQNXLP ENTIWRPLQIARANDFYSLRNSST TTITTTTTTPIHHIYSKSSASNART LPITLGTNSKVTIAASNTTSDAS NTPAAIIVKTTTSAMPGIPGQLMLV KEQQQPQPQPPPPQQQHQALADKV GSEKSSPASENDYLIYDDKRSYFT	None	None	None	None	None	None	None		