

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A0A0B4LF93	A0A0B4LF93_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MMKRTRVDEVQFGRPVPTSGGV GVGVVGVVGGGPTSGGGGTATVGVN TTGVTIGTVVPSAHNATISGIGSIHHR ILTPQHGGQAQTIAYLPSTTPTATNLKT TTSIVDSTTAGGPGVAGAVGVGS AAGGGVVVSTGSGTQTLQYTSY SVASIQAGGTLKANTADGANTVQIH VTGGGAANNPASAQTVSSSSQTGTI RQRTISGTQTVATAVGNLATISQQQP VQQSPLGKAQTPPSSVVANSIPVGGT TPPQQSGNATPRLKVEDALSYLDQ VKYQYADQPQIYNNFLDIMKEFKSH CIDTPGVIERVSTLFGKHTELIYGFN MFLPPGYKIEIHS DALGCSVPVVSMP SPPGAPTSTGTVHMLTGNSSMSGAG HIAIKTTNAATLTPAAGAGAAAAAAA VAQIQSAGAVNLMTHGGASLTQTTI HALQQATPPQSQSPGGGHVHVSVT NSTAANAVVPGQPGISVAHNVPQN YSRDRERATITPTQVAGAAANVNA SASIVVGGPPTPNSLSESPHGGAGG GPGAGAAQHNLHHIQAHQSILLGE TGQQNQPVFEFNHAITYVKNKRNRFQ NQPAKYKKFLEILHDYQREQKVMKE GSLNQGKMLTEQEVYTVAKLFGQ DEDLLREFGQFLPDATNHQSGQYM SKSASVHNDHGKRPTATLGGGAHIT MSSASPAPSGSPLHLGATTLPOIDKS AHHAAIGNLSAVNTSVSIKTYNNNQ QQQNHVIGSGLNATRNDILFEKDYH AGLQQQAHORGAGVGGHHHLAGTA AGANIGRPGVGASVMVSYDKEHRN NHHVQKYVGHAPNQNLTHGHNAK KSPSYGIPSVIGSMPHISDNLDRSS PGISYATPPLPSGPHGQHNSGSATRR PGDDSLVGHYASGAPPKRPKPYCR DVSFSEASSKCTISDAFFDKVRKAL RSPVYDNFLRCLTLFNQEIIVSKTEL LGLVSPFLMKFPDLLRWFTDFLGPP SGQPAGGLIDGMPLAATQROGGSS NSSHDRGTSHQSAAEVVDVLDLSSC KRLGASYCALPQSTVPKCCSRTALC REVLNDKVVSPPTWASEDSTFVTSR KTQFEETIYRTEDEFELDLVIEVNS ATIRVLENLQKKMSRMSTEELSKFH LDDHLGGTSQTIHQRAIHRIYGDKSG EIITGMKKNPFVAVPIVKRLKVKKEE EWREAQKTFNKQWREQNEKYLLKS LDHQAINFKPNDMKALRSKSLFNEI ETLYDERHDQEDDAMEPFGPHLVLP YKDKTILDDAANLLIHHVKRQTGIQK QEKQIKQIIRQFVDPDLFFAPROPLS DDEKDAFPFLVDDNTKMDVDSPL GRTESSTRNAKSTPSESASPARSNAS TSSVTPAGIKKETDDSKATTGSFAPA SSATASSATPVDDATPSTSSAAAAAS AASSTVSGTEGPKKDDPLSSHKEE GAGSTSSGVATSPRQAQDTAGAGVD VEIKLEHPADFSNPKLLPPHAHGQR EDES YTLFFANNNWYFLRLHAILC DRLHVMYERARLLAIEEERCVRNRR ESTATALRLKPKPEIQVEDYYPFFLD MLKNVLDGNMDSNTFEDTMREMF GIYAYISFTLDKVVSNVAVRQLQYCVT ERAALDCVELFATEQRRGCTGGFCR DAHKTFDREMSYORKAESILNEENC FKVYIYKIDCRVTIELLDSEPEEVDKP AALKAQKFSKYVERLANPALGGGGN TGRSDSALGNDSVVDGSDIKTEADE DTAEA	None	None	None	None	None	None	None		