

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
A0A0B4K171	A0A0B4K171_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MEHHEDNAQLDNYLQNRRLAESLQI CGGAGEHNPHLADATGGNGCAPGI APSKSDEVGDGEDEEWKYIHEVRQS EKLQQEKPLTKETGNGFGPGRDSD NQVHGNGAAAVFNLYEEDVEVIKN DGDFTNSNTTSTDEVVARQAQEP NQLPEQLQQQQIESQGVHEDPRQ EDEDEHSSVATTYGTSSLSENNSSP LDQEEVVMVAQTVGQEQLVDFDNK ENSYFVKNLEENHSQLNPNVAFVFP GVGSQSSSPLAAEDPLPGVQPRPFL PGGTLDDLVAESPRKEFARINMDGI AVPDEREFDIEADMRRPHELEQESDT FGAGHLEMQLLNGIGTADQAALRDV LDHGPETSDMELPLDQVPNDADI MKQSIYAEHNSSIEDILNSVQPLPIQ TCDDKELIHVEEKEHVSKSPSTEEL QFQSDFPNNQESHTLFNNTQDPM QASFYLEHTSQKAQEGCQEQMQLP AECSDIFADQSLLLDTSAPQLSSEAD SPVAKLELESQQAGIVDITPSPLSSTA EKHLVEDTKELVEEYTLDPESHFFG VVSSQAPLQNDENAVFESVSGYET QNFDEISSPPEGINPFAQPFTPAHLV IEQANTMMEDVGGMPIPASEDFAIC DKVANEIPLSSASKEKLLPDTTDEQL LTSALEEKLRVAPPEESVSTAADGQS ISQFDEYVIASNKPLEDILEPEKDVEV AKSLSEKTSVATVAGGAVVGATKTH SATKAGSTAASAKSKTETLVMKKTTA SSTS VYGANKSAAPRPSTARLGIKST SIATKTSSTSLTGNPRKSLSSNVGS TVKPPTKLSGTRPATAPVSKVTLGAK TITNKPTASGTASDNVTRITLRPLVS TNARRPATSGTGSVASSTARRPVTN AKGSAPGSAASTKVRPAATMTAPVK PKVLSPRSTISSITTVRKVPSTSTPSF STRSPNKQQSNGLGKNTSSTTTSTA TATITKSFTARSAPKFTHSASLTYNN GSTRRLVPGSSSTTTSSLRKSSP LKAAPGKAASKPLTPQSKDGTAKSSP AVLKARNSTLMGGEGVAPSNECVPT PNGQINAEVVKLGKGLAEVVKIIE EQLHEQDVPAHNAEVPLLDLF	None	None	None	None	None	None	None		