

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
R9PY26	R9PY26_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MVAAPPSAGASTSGAAATAQTASNK KEKHQRSKQADKEKDKKEEKQAS SGKRKKEKISVEKIDTGFVVGIGDK LKVNYHEKKSPSSHGSTYEAKVIEIS VQRGVPMYLVHYTGWNNRYDEWVP RERIAENLTGSKQKTRTISTSSANS GSGGGGGGGGGGGGGGGGGGSL VQGSQPPGVSDKQPGKDGCSKMSP SSGNSTGPGAPSLSGSLGGASSTPSL LSTVVKTPPTGAKRGRGRSDSMPP RSTTPSSVVAHSGRTKSPAASQPQL QQQMKKRPTRVVPGTTTTPRRVSDAS MASESDSDSEPVRRPKRQSAKDKP QAGKAQPPGKGRASSASTAPAAH PSDDSEDEEEEEPSAARAASSKQQ QQQASSLRGSRAGGNRAMSSGAAS AKGRDYDLSEIRSELKGFQPKLLTNA ASNEERKDLAKKEPSDEPALQDIKK EPKLESSAKSSSTELSETESYADE SQSSDYRKQLKSGGAGKKEPTASPS KLHHEPVTKRELAVKEEPLKIEPKTE PKEEETKSKPFLSGADIKPTALIAPAR FGNTSAAQAASSGSSSTTAKYTSVI VEKPLTIGGKKSVEQHVPKKAELLKK QSGGAGGTAASSAASQESKKFAEP VASLKVEMPAACSPSSSSSSSFCFS TGSAVSSSATRSLPDMSKLEISSGT VPGATPGAAALQPSNVAQVSSSGAA KESKYSSSGGAAAGSGISMRLKLLS DVYEFKDTPEPFEFKRIKSPMASVGGT VAAAATAAGAMAAGAASVITMVP TPGASGSGVAASASTAPVVVGSAR KQALKASAIQHILEHQSPAAGRERG GYGGMTSSISLLTAPKLRKRGSPKLE AALCMEKAKTYKLDKDQVQVPEQ KTQQLIQPTLGPVESYGAGSPEAMS NLSTPTTSSGHAQLSASTYSQLTPH HATPFDALRKSPFNLNITALNEELA QTVQETTRALTDALQPPSTVPPTVS ASGSTPTAVTPVITAMPTAGTAGLSC PSTPPTGANPVLASPKLSTPPQAINA SKQPPASHIVGSPFIETRNVFELSTS NEGSYSSGESKDNKFEKLENVKIL LAGAVGPFDL DAGSYEQKTSIADKV LKAISQKKEEVENNKGKPPVEPAAA AKTPGREDVVPSTAKLDCSSAIKL DTLKLLEPLKIQTGPLLGELYRPGP ATSSPETKSILESSLPAKNSELSETIQ KLECAIQQRKTPVGGALSASSTAGT PPTAHTPNSTATAGAGFDESMDST DSEQRLVIEDVIAEEHTTTTTTGEQK SPGSQEEGQNTATMATAVTAETE GTSSSTPTPPVAPVKLELGAKAMPG IQAPIPKPAEGSSFAGKLTGVVQPP PLAKLQQQEELGQAQQGSKSSSFGI PIVLEIPASVVVATPALMVASAATV MRHSPGSTPNSQAPVLISPKTFLTAP KVGEVGGLLLKAYTGVSGAVGAAPA GSVIASAGGGGPTVISNFPQQQQQ ASVLQASSEMPANYVLDMAEAPS SSAPVVARPFVMAVKGELFNVVAP AASSPLISDPHNESINLLCEETIPGSP APNYGGTDQLPTAVNSALAVGITPL PPDTPPPVPVAAIQQQQHQQLQSGQ QQQAAKTGEVNPSPGNSPDSASQD ESGEEAKKNAGLEHEDSTGLGALNK RKRNRKPLPMSAQTAAVAALQSL GKRRRQVSGMRNQKTTTTAAGSDT DDNSDNIAPNAGQQQQQPRQSARQ HMMPQSNAAQQQQQLLQQQQTQ LPQGIQSTNSSGVRPCPNFLVELDP ALSSDECITILRKQIQDLRKAYNTIKG ELAVIDRRRKLRRREREKKQQQLQ SQQQGKICA	None	None	None	None	None	None	None		