

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
G3JX32	G3JX32_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MAQPQLLQIKLSRFDAQPWGFRLOG GTDFAQPLLQKVNAGSLSEQAGLQ PGDAVVKINDVDVFNLRHKDAQDIV VRSGNNFVITVQRGGSTWRPHVTPT GNVPOPNSPYLQTVTKTSLAHKQOD SQHIGCGYNNAARPFSSNGGDGQV SIVNKQYNTVPVGIYSDEIAETLSAQ EVLGGVVLGVNFKKNEKEYQGRSE VLKFLREEETGQSTPEPHSPANFYW TQSHAIGGNERRTPLHHQHQQPQQ DERIGVPLQSNLTAPEATHRPSLVA PKDNEEQARQDQEQPDPRIVLPIC PGLQGPEYKAEMEAALATDQDG RPRPLAASGHPACQLCGVGIVGVFV RIKDKNLHVECFKCATCGTSLKNQG YYNFNNKLYCDIHAKQAAIINNPPTG TEGYVPVPIKNTKLSASTISSALNS HGYGGHSNGYSNGNSTPAPAPVASS QATATVATVAPSAATAATAATPQAA TATDSPAATASSDNMSAYVADEPS SIYQISAESVALAPPPQPPTAGGG DQPFYVTLTGNVIRSVQAPKGAC PSYKVNQGYARPFGAAPKSPVSYPP QQQQSPRPAPGGQNPYATLPRSNV GQQGRNVRYQQQQQQQQYNNQQ KQYRNSYPMGNSNYSTPSQSPYITS NTNNYSSNSYNNNNSYNNNNN VYRAPGSASAPAPVPSAAPTATAPF KAPIVKSVIANAVNAAPAPAVFP PDLSDLNLNSNVDNSPGAGGKSAG AFGATSAPKRGRGILNKAAGPVRIP LCNSCNVQIRGPFITLALGRIWCPDHF ICVNGNCRRLQDIGFVEEKDLYC EYCFEKYLAPTSKCGKIKGDCLNA IGKHFHPECFCTCGQCGKIFGNRPF LEDGNAYCEADWNELFITTKCFACG FPVEAGDRWVEALNHNHYSQCFNC TFCKQNLLEGQSFYKGRPFCKNH AR	None	None	None	None	None	None	None	None	