	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q9W451	Q9W451_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MISPSVCAALFSVLVLNSYIAQVQSE LSAAEVAGDPSAAQLLPSSCRNGLN GSSSGPEIRLPFVDMMVRHEKPLFV LREPEEDLLLLLHGHEIFQLHVATG ERRLVPLGYLAGQVSGRRALGAEVV AWREFLLLAIVLDDSMEVYQLPKEL LLSEDPATQVSFEPLQEFSLPGAFLQ LHLLKPSGDQVLLLVASNHTRSHSK CRTFEWLDTYFNPLEEITLPAIRVLQ VVGRQPLYIIFGRSLKGSQSKLVLTIY ELDRNSRILQQRQALTVQAQSVQAV RFHGRNCLIACTSSASASCIFFRMVD GQFVVYRKHARKDLSFRRLSATKKG HLLIGARANGEVIVFSSTRLDCYSGF QVAGEADPSGLLIHKNLHNETFLLL AYRRSSSDLIRTVQLGGVEEISNQVG AAPDEDLSVVQLHRHEFEESINALR GLLLRRRSSTENLRNIVDILKRKSAI HLDTPLHLVQQGQIGRLQLVDEHLR TPTQLKQRLEELRQRFKAIQTRRTTR SFGTGNENDEPETLKARRLRVGNLI YKGQLISGHILDNSKPPMLTIRNSPL QTKKLRVNQLVTPQDFRNSSGAQIA PEETSGSECVVRHLRVEQINGVQWN DFLDSLFLRSRDTSIEGRLVMQSRV RVNSLKTPLLNGLVVDRLFNLRRAQ VISSNIYMSAFFAPRLEAQQVNGLDF AHDVLHRGSNDTWIKTPVRINQMSV SGDIFVGNTSKWQPQPRQLDDPME QRLQQYYTGRVTIRGSLTVTNVLRDT ANSLIVLGNQSLARKDIQSTYLLDQT PQNLTNLTFGHARVSVPSMTTSYVD GHPVRDHLLSGGQQNQHTHPNNTL HIIFMNATILGDVICRDYSSRLAEIAK DAVSHGDAVNITGHKHFQAPLTVQA LQANEINGVPVSEIVVKSNTTQNFTG TKTFARVVVANDINVHDHLNVSHLN GLPLQQLLGHDISLQRLELTETPHLE NLHFRRLNGLPFDELLSKISEDGEQ PLLLHKQLLIEGNVRFDKAVQVQNI NGIQWDEYLDRLVRSDSNAEIRGKK TFMADVQINDALQSPHINNLDLSSL LDNTLLRMTPQQISGAYTFDRMMAT NVDVNRINGVPKEEFIDTRKDLELK GDLYLKQLNINGSLKCPFVNEAVLN DLKQRVEQVQQLPWRNLIVTGDAL WDGEEQSQSQLDYLRQHAVRRVGN QTIAGHVLLKEPYLKDIQSKHSLPIDL NFTQLAEDALLRRSPTNETQVITAPQ ELQGLVRANTLHLEQDAHFELVNGI DIRRFNASLYRLSSGEAIAADLHFHQ APKIGKLQLEYPEVNGAPIQSIYQQG KGQSWPPVTFKNLFVEQDLNLKTV NNMNLDYFLQHRIPLKGDALEVFGS LTFEHLQLSGKPLLRSINGIQLDNLV FRHSNRTQSITGAKTFRGGVEFTGP GHVMNLNGRELSESFRGSIFRNKDY NIDSLVLDKALFPGGLVLAGLQSQIP NRMRATEVQSNPLEEFQELLALGN QTSPRRLLYLDHDTESLESTWVKGP FRGSSELQVSLPLGYSAPCQRRTLQA KLLIHERRVLLANSSVSNQLMRLHS GDIRVKVQNHCHRPARRLRSRIRISC RNETHTLGMRQSVEAMELLEMGE MVLLLLGTEEEVRVLRLKRSNCSLT DWQSVLPADGRLMKLVRMGVESRQ DLILTSGMSNHRPVLAIHYRDPLNH NFKMLQLIQGSYDLAEMQGNQLLV SCLGCRHIAIFKRYSHSAEILFEPIQQ LSFQERIQQLTPFKVGNEQYLLVVTQ PEEEHFYLFSYNEVGGWQQRTFGIK KQDQWAWSLVKSGQMLEAAETPILL LCGKKRECSLVKALLG	None	None	None	None	None	None	None	None	None