	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	O61380	O61380_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MQQAIPTLPTQSDIDKAMQPHSAQN MYVSTGNNNSGNTRSNPQSGGIFR GPPSTPNAPRGASGGATRHVHVQP MYSQPLHQNMVLQQYTQYNPRQQT FPASHLQYAPAPMPYYQYQYVPTLQ QQPPHTRSAVTVNTNVNVGNNLQP VQSGPNGPLPVPGASSSQIQLLTSTV QPGASTVMGVGGPGSTMGQVGVPP MVGVGVMTTSVQPQPVQVQPASRR RHQHRLQIIDPTTKKNILDDFDKTKS NTDNEFSDQVTSTNTPATVLSEGPIR IPQQESVGLNNLTSTSSQGSESRTN APYIPIETTPISRTDVGPTPIVSAMTD APSVEILPTPQRGRSKKIPIVSPKNVS ESIVAPITDETDDAGSTPISRATEESM PPNQTHPNLLISDSSKHKQAVSNSEI SKDAPTGLKEMHVAELESSVASENH PAGILDVNVNKSSQSLDFSADPEIDS IGAISPTSPNAVSPILHEVLDNTELSK KLENSTTERFKDSQSVEKPTHQELS LRNATDETEISAMALQDVNSLDNN QIEKTYPSKPKLNVDVSEDISSRESAI KSTSTKNTGVDVGLQSDSKPETTLN DKQDSTDLKVKVSAKISSIINYNEGQ WSPNNPSGKKQYDREQLLQLREVK ASRIQPEVKNVSILPQPNLMPSFIRN NNNNKRVQSMVGIIGNRSNESAGN YIGKQMSMSGVQSGGGRSSMKGMI HVNLSLNQDVKLSENENAWRPRVL NKSDGDSDAKSALEKDELVRRVRGI LNKLTPERFDTLVEEIIKLKIDTPDKV DEVIVLVFEKAIDEPNFSVSYARLCQ RLAAEVKVIDERMESETKSNSAHFR NALLDKTEQEFTQNVSQSTAKEKKL QPIVDKIKKCTDANEKAELEAFLEEE ERKIRRRSGGTVRFIGELFKISMLTG KIIYSCIDTLLNPHSEDMLECLCKLL TTVGAKFEKTPVNSKDPSRCYSLEKS ITRMQAIASKTDKDGARVSSRVRFM LQDVIDLRKNKWQTSRNEAPKTMG QIEKEAKNEQLSAQYFGTLSSTTPGG SQGGSGKRDDRGNSRYGESRSSSAY GGSHSQRGDNGNLRHQQQNNVGG NVSGGAGHSNGNNDENTWHVQTS KGSRSLAVDSNKLEGLSKLSDQNLE TKKMGGLTQFIWISSDTTRLSSAPTP TPSNPFAVLSSLIDKNSNERDRDRS GPRNKGSYNKGSMERDRYDRGMHS RTGSSQGSRENSSSRGGQQGRTLLS SSVQKSTSHSKYTQQAPPTRHTVKA QSSVGSSNVNTGPLYRGSEQQTSAT FSQTTRSVAPVAVFIEASETDLKLIKS VVSEIVDLSAASKEVTPGAVSCIKRV PEKLRCSFIYYILTDYLHLANVGKQY RRYLSIAVSQLIQQNYISADHLRLAY NEFTVYANDLIVDIPELWLYILQFAG PLIVKKILTISDLWNNNLKENSPSNV AKKFLKTYLIYCTQEVGPNFARNMW IKFNLKWSDFMPESEVADFIKFNRL EYVENESKSPVIDHRETPEKHVKNVI DHIEHLLKEGTTADCIIDYSNGNIMV VDKLFIRGLTETLSNFSIHYKDNSYK LESETFQKFCIPVLQRYIDSNEDHQL ECLYTLQLLVHGLEHPRGLLSELIGE LYDAFVIQKESLCKWRDSKDQSAGK GVAVKSLNPFFNSLLNDDAN	None	None	None	None	None	None	None	None	None