	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	B3DNL9	B3DNL9_DROME	Drosophila melanogaster	NaN	28.06426	T382	NaN	40245499	LVIANENMAANETNIFQPLYKQDEH ANILRQQIIDWAREQRKRLEQMNLD HLLIRLKDKLKHREEKLKVSIIKQVN AKSFEDKHMKLTPNYWKALKLTKQ ALDKIEQKTTLAGSRHPNKRSVDQP NEEPNFDEVTVDTLVVHGAVKADHI NGVDTRNPVYDSVTANKVYVSAKYT EPAGKERKLLLDELIVKDLQLDGRL NGLKWADLFDLTLKRSNKDVQFIKA AADFDHLESDAAQVNNNEVNDRPL GLLIPVDGGDFIVQQDVQFSQPIQVN RLLINQRLNHIHVDRQRFDVLTKQA NHTQVIEGSMRFENVRVMEPITIAG QIMGAELRAMSPAKVTHKSLQLNG DFVVDGDVTIGRLLQMKDFVDESTQ KSAAEALTRGLRLDQPLEDVNLNFV EPLIANNTELSFINTQDLQNLVKLNV GEVQLVQASKIFPQNVEIRDGFGEIK WLNGIDTERLPEIMLTKSGNQTISSS IQLQGVELNHVNSSQILLNDLGVED YLQTGQDQKSNGTLYIDNMDVREM SVNDLHLNGHLFGQTMSSIYEHGS KSLDSWHLPPDFNGTIHAQNLWLS GHINKVKVVQLEQQLQQLAGNVKYV GDFTFEHDVNISSLSFENSLNGIEA NKFGSCWLETDRDQTFTAPQKLASL DSQQGLWLRGQLNNHTLQDLENRS YRLNSPEHLQSVRFANPIVLQSELQL DRLNGLRVPEDVIYHNGGAYISAPVS VDGDLEALKLSNITNLNGYPLEALEI YLSGQNVRTFVAENVKFGGEPSYQL LNGHNLGNLLDQVWLDNEQIELREI KLESGQFEGLLEFQGPINGHQVDHI THNYFSRTRSNLLKTPLSFQHDVTF AQPPVAKYVKLRRGSSGVLMEGTSS NFSLDFDDFVTHSLKTAGSHAISGY WKLPEAFISGNLNDIYINNLNLAND VVRMPLQNDSINLPLIVEALKTVQG GTINRLHTTPESQVADVPVAKWIHG AAYLYGNHSISGITHLNNVNLYNDL TVNGSVNGVPWQADKLILMDNEQD IPGSLLVANSLPEQKRVMSNNVKNL WVDFVNDLPVNELLVSKAENRPNL HVESQLIFTQPLTVGHYEAGNGNGL LDKYYKKKRGVESNDIHDWQQLQE NVAAIKNRLAKPPKVLEGFAVLQKLP HKASHMKIISVDNEDVLAIWEQMPS EVGTTYTWQPDIHLFGRSSSTALTQL NKTNLNPINEEAQNPQKLQFRSYGF QCRALQEENQIVIRCLDEQNNSREA ILPRRAVKQLLPVADSDDMTILLRTP RAVELWKWSNGNYSLSSRLIDGQVE QMALSQGGAGIESGYVALMASSPAA EIRIYSFGSPDTSSFQLDQVLELEDS QQSRVMRFMHLPESDDLLLCVSNV LPQQPLRIYQHRGAAGFQQILGDSAL PKAHALEVLYLPSHKLRLLSMATEE SVYLLQPQFTTL	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None