	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	B7YZL8	B7YZL8_DROME	Drosophila melanogaster	NaN	28.06426	T770;S772	NaN	40245499	MDSLAIFIFLWAVLALFSVWLIKLLY HKYLNKDKSASAANSRQTSVAPTSG SPTSVAGKTEKRLSEPRDLLATKSKV EGLDLSKPLAGASGGRGRSSASPLN TAGAAAGGPRRRVVRQSSTGPENRK KRYVPPPSNVVGPETSSVTWTSQVF RWLYSDLVIVNELLMSWVIAINDTL RKSVEEHGVAVEVVRVLPDSPAPGL NNIFCNCDENNPADMLITFDCDAM PVLQVKTFRQKSGKVETSHYKVTVS RFRARMAIPMNYNSLKGEMRVEGY PDVRIAMNSVGAIKAMDQDEQQLQT VISDILTTALRDTIYPVDFSIYSTCPRA EVEPLDLPVIYPVHYDSLAHNMEQH MGGVGLRDSQHMVSGRRLLVKIVK GDGIRDAQNPYVVIEMDEPAQKNQT GTQRGGKPFWDEHFLFELSPQSAEI LFEVYDHPVIASDPPKFLGLGLVGID ELAVGPASTQLLQLQPRPYETQPVSG AITVDFVFIEGAEIPAGARPQRLKEAL RLSTPAINEHIRNGADLADAAVRALQ DGALSSSGSGGQPSKSTLIIHSVQRN SSSPNAFKVELNRDGQIEVTDTATEL DQAVAQAFERAANEAQTELELELAK EEKSSQLGEPLIDSGNGTVNEESTA EFGQPNAASSPNGSSYHNNYSLNG NGNSNGAGSGGYNSLPRNGGAQQL AQQSGLLEGHDVVDDRGRSKKRNF FGTLKKRLSRSKTRTLSADQPNNNS HKSLSATNSNTTTATGFPRTATGTL NGDSSRSLSVDRATLSKSNSLGPRM GIGHSITDHSRRSSISESSAISGFSSA SNKTYVHEASTLVLETIENGIKRHFI VPLAIAQRPRWRRKGTKLHIYNDHT FIAKHLSGSGLQCSICMKSIPRRPGK QGYECRDCQLISHKQCHIRAPQACP NPTVLSMELTSFPVLTERDINLVN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None