	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	H9XVM9	H9XVM9_DROME	Drosophila melanogaster	NaN	28.06426	T247;T250	NaN	40245499	MDSTLNIENVNDPTSIASDLSAENT KADLVSLNEPKTYEPTTIAPDLKMEN TKNSIPSPIKPNVNDQTSSASDLTAE NTKADHDSLNKPKDFNNQILNIISDI DINIKAQEKITQLKEQELKLIQKQND LANEIHKQQILAKQLSAQNQLKQNE CQSGELDLHSQYQNERGGPTSTLYQ EKNIASKQIPVVHNNMNCTVEKKRG YRPSDLIDPSKRQYLLSLSKKLWQEK KESANVSKTVDLRKIFTPATDAAEIL PKNRKLYASSAFYSPTLHPTVEDQVE LARRISHSLSDISNQTSKGQSMYVN RKKRSDKWVHEGGSQGNDAINPFK ENSETKSTMEVAKLEKIPLKLIMNP NGKVRDYNSLKDLINVEAGLLSPDN CAELITALQLHQGRGAELFAKRRRK ADNWVVDESHSGTQYHPSGIPDFQ QYQQRPVLSPNILPAYSDAGKHRVQ LNIHQNQLIEKYSKPGLQVVQSPWK AALQTGSASSAFLEDTKSFSPPALTPI PSSQAGSKDWTDANEPMPHGSSRR NTNAVASSPRSVIVPSNPQRDLAYTP CVAQGWGGRSVELPKDSFQSNESQ SSWQTEPHIGRADDSNPSFNNYIFG ERLTSNFAFDVQNRLHSLENFQKYF LEYQRLELEILRNRESSRVLQHDAYE SLLDFTPQNEFKVISGEDLKVNCDL DEKVNVRELIQSFEKQNTTDLRHDA APIEGLYVPKEISLSSYAAPSQQCEIH SKHFERPYEDTFPRTFPRNNYSANN NFPGNDGIGEFPLSKPTGLTTGTLH QQPYSPTSYKKTVPGLEKVENFARYE QPDRKRLTSGSPQYLNALQNSQVRN ASPIPFGTSLNEQTSWPPRSVLLQGG SPSPLPNHVQTFNKCAKGWGSIPVK QQSYSSQYNLPGNLPYSDF	False	False	0.938	3.938	False	False	False	False	False