

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q8IMH4	Q8IMH4_DROME	Drosophila melanogaster	NaN	30.60235	T14;T229	NaN	33925313;40245499	MVTQARDQLKSNQTEELKEVFEGS CKLIPKIQKECIKVAADDLPELVEA LASQMNPDQVCSVAGLCNSARIDEL YKNGIQAGLDGTVQNEDDSSEETEL AMQPQLSCGCNLLSRLMHSKFA ATDRDDMVETMLHMCGLSSFSDA CANIVLTYFNDIYDHVSKHLTTDAVC HVSGVCASRYHQHEEEKQPQEALV ALDAGDDIPCELCEQLVKHLRDVLV ANTTETEFKQVMEGFCKQSKGFKD ECLSIVDQYHVIYETLVSKLDANGA CCMIGICQKNSASSMKDVPIMPLLP VIEPAQVKITIEKLEKHEKQLGASE PKFSQQEILDMQLPIDHLMGAANPG ALVEGGELCTLCEYMLHFIQETLATP STDDEIKHTVENICAKLPSGVAGQCR NFVEMYGDAVIALLVQGLNPRDVCV LMQMCPKNLPKKEDVEVFNPPQAS DEQDPPTCPLCLFAVEQAQMKIRDN KSKDNIKKVLNGLCSHLPNEIKEEC VDFVNTYSNELIDMLITDFKPQEICV QLKLCPKTTYALWDLRISLEDDVDG EDKSSSEEIFNDIESLEELPPQLAF DPGFTAAPNCLICEELVKTLEKRMG KHPTRDSIKHILEESCDRMRKPMNT KCHKVIDKYGDKIADLLLKEMDPKLI CTELGMCILADLDDLEVDALKYDVI ALPRQDNKLSSEIKPPTCVLCEFIM TKLDADLKNKTEQDDIKRAIEAVCN RLPATVRKQCDTFVDGYASAVLKLLS DVPPKQVCQKQLCFSVAVTDEVLE CGVCHGVTQALLPFLREKKNVSEV TALQMTSVGCENLPKYYKICSEMIS YGSSEIKNLAKRPYIDQSHICAEIGKC FESEKSSLAFARISA	NaN	NaN	NaN	NaN	NaN	NaN	NaN		

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