	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	A8WHG6	A8WHG6_DROME	Drosophila melanogaster	NaN	28.06426	T156;T162	NaN	40245499	MASKAIWLTILRFMGDLPDVVSSPTL HVFDNENLMSDLASLLNTSDSYKPR LFVRQSQRRIPKPLASGEKEAQEFYQ HWLNVPTSHLEKIHFIIGHGIIKNSL RDEILAQICKQLYLNPSRSSYSRGWL LLSLCLSCFPPSKEFEPHLRSFMKQ GTAQLQATPSLQRLERTLVNGPRCQ PPSLFELHAIRGRHPLRLDIHLMDG QQRRLQVDAASTAREAVNQLCQGM GLTDTFGFGLVMSLNGKLMPLGAG QEHVLDAISECEQRQLDAPWKLYIR KEMFATWYDPSMDPKATQLIYKQIL NGLKCGEYRCRSEKDIAMVCALACF VEYGPGEILRLKPSEITAFVPSDLLAP GERAIENWSRLIAATYEKSSYVKEEQ NDLLLEAQKRAKEDICLFAHLSWPM RHSRLFEVVRKEGPKLQSDELMLGI NSAGLFLIDETEQVLASCCFSEVLKV HVESDDKLHVMTFQHVNFVLQCSS AQDANEVINYMLDNLRQRSSYGVAL DPVVEGDLEDCLVLNPGDLIEFEAG VTGAQLMAGNAQDCYRGCVNGQW GQFLAGNVRVLATLTKPSEKLQDILR EGRFQEPPKPTPRANYSRRRQHNIS QLAESHFREPLDSDKAPLSKFSPEPL KAPLLKAVVKVPPLFQQALVMHHHI LKYMGDIARSNLPVNTDLIFQPALQ HPLLCDELYCQLMKQLSDNPSSESE KRGWDLLYLATGLVAPSVLVMRELII LLRMRADALADACLKRLKRSLAQGQ RKKAPHLIEVEGIQQRCLHIYHKIYFP DDTVEAFEIESHTRGAELIADIAQRL ELKSPVGYSIFLKTGDRVYAMPEEEF VFDFITQLIYWLRQQRTIRSISDGQY QLHFMRKLWLNNHPGEDLNGDMI FSYPQELHKYLKGYYPIDCEQASRLA ILVYSADHDVSLQRLPEVLTRLIPED LIPLQTVAEWRQQILPKVHRDHLTE DHAKILFLQELSHFACFGSTFFVVKQ QNDDALPETLLIAINSTGFHMLDPTT KEILRSYEYSQLGIWSSGKNHFHIRF GNMIGASKLLCSTTQGYKMDDLLAS YVKYFNEHE	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None