	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	H1UUH1	H1UUH1_DROME	Drosophila melanogaster	NaN	28.06426	S340;S341	NaN	40245499	METPTGSGRPSRMATPRLSERKRQL FGSPRSRLRQINDDEDDADVDSLGV LPLKTHVAANRKGRSLFAAVPGKSS SSANSSPETNKENKKTRGGVMTATA EQLPQLFTATMRLNSNSSSNSRNSS PRQTRVQRKRADSSMSSPTSSSEGT PSSRARNSIRRSPRTFSAQKDPDAFS SPESFQTRLSKVAAMLMKGQDSRS MLEKSKKKHNHSLKTTAQVHTTKP KKTSPAEESQSDDEKPSSSKNSRKN TEVRETRSSQIISPKTRNRRRPFTSA DINCKTLKAAAHLHENMRSYDEEKT AAVKLENSRSRSKSPVEVFKSNDDA VKRNTGNTNNKTAKSSEVATAKRPE SPGSSMKIDVEVPESDEEASNHKPQ KRQHPETSTPVAPSADADSGSPQSK MRKVTLSSSIPTMAFYSHSGEAVTK SRRRPSISKNSLKQPTKISPTSRPLLG INKGVHHKIRKRHGFANRLPATDMD NILNSLSNERLKNLITTKREERAKVE EVHQILRNAKDPIKMAKPLSVIEADD ANNNNNLPATAWQETSADFSDLSD VEDIDPIIEVEPIIPIIRHEPVQKSPTA EPADLSKRKFFKSGRRSSTCMEVRIT DNIRASVSQGKIELVQTIRRKPRQVR VKSATIFSAEQATVDAILKNLDDTVV DEIVEANPVVQATPIDAEETTMETES LPDIIEYAPEANDVEIDPFAEFRQRLP YQTDDPNVVEQQQILLEFLISNNICT EKNFEIFIANPDDYKDEANQIVDNLY MVVNSEEAAQLAQMETVENTAVAIA PKQDAPAVEEVQPKLFPIFTQRLQPV VQKSLRRRPDTSMRLLTAAGGSNQY QIDAGQKAFGARQCQQCGLVYTVHE PEEELLHREYHNSIHVLRFKGWIDE DIVSVYPEWASDGRIIRINERAPTARL DRLRDLIGVVDKELGYSSYIVPKIFVA FIAVRKQQIVGFCLVQPLSQAHRFIQ VDGTDYFSEESYPASCGVSRIWVSPL QRRSGIASKLLRVVQCHTVLGQEIAR ECIAFSTPTDDGRALARQFTGLDNFL TYDQ	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None