	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	A1Z9L2	A1Z9L2_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MEHSYMRDTPVRSDVHNGLTPARN ILVRHPPQCPSCHTYPPQHEELSEIQ QAHVPPYDEVVAKEAMNECARIAKY VKNNNSDEQDWVARVNKFNWTPM QEMVFEKVCSILDQDQLARLANDKR QHEPIHRRISSDKSASRFRKLLASVA WESRITQWIHALLMEHLSPSYMASY LEILQTLKTKLPTLVDKMLFGRPLNN SQELLAPVMKKRWEPNIMPKSRQLT HNAIMVVLPNMPTTGTVSDRMQKW YQALSTITQVVQISLPNTNNRIGNQN LDHVAETIVSITRVKINELRTENPSR GIILVGFNAGAALALQVALSESVACV VCMGFAYNTIRGPRGTPDDRMLDIK APILFVIGQNSARTSQEEMEGLRER MQSESSLVVVGSADDALRVPKSKRR IEGVTQSMVDYMVVEEIFEFVNRTL SNPPGPRMPTCLMHQQGYQRLPKQ SPHILADGNANKAVQQMRKRKMDG GLDDSMGQPSKTKFVPHNDAQDYG DAQTTAGGSASLPKVITPGVLGKQLP PVNLAAGTKIKMIPSSQFVQIKSLPS QSKLINYTLNKPSVGTSSTANIGSVV KTLPCSSVSGQQIITLKTPSGQTQQF ATAATSTGAAGSSTSTGTGQKKYTVF KNANGMTMLHLTKSVPTTGSSSGS ANMDLSNIIDMPIVFADNEGNIPEH QQADKDIKPAPIRSANKSPLIISQKIIK EANKPSGMVTSGSITAVSKPGNIVLN KGLQQLLNSSPGGTLATQNKVVYLN RSTIKPMGNVAPGTHRVPIRIVSTPN TGGAVTTTASSSPIMVDPATGSKVAK TVNVQTIKPTGSVLAQGTIRQAGNVG TKSYLQMINTAPGASKTVAGDGKGS VRNIYFKSATGLKQLPVQMLGNRIP GGAVISSTASASSGSPGLRRVVNIGP VSKVTATSTTSSSIAATLTHNKM	None	None	None	None	None	None	None	None	None