	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	Q9VWA7	Q9VWA7_DROME	Drosophila melanogaster	NaN	28.06426	T561	NaN	40245499	MRKNSKEMAANRRRKKTIGMLTSG LKSVVKWKIIVSSIMLLLLSGTSHVL AVRKGRLMSPSTFTALSGDLHVQIQ FNGNEMSSQAEEEHTIGGGPSNEE PLSSSDGLKSRNNTIMSTLVISKIIDE VESPLEKGLTPGPEVTNTSQVVLHR KRKEVVQNIPLFPDPRFNVTQVTVP CQTFLIGGRYEMQVVSNLKSVNRSE VTTTLVPAQDERLHQTLDVRWPSAE MIVSPVRLKTYPKHPVEVTLRFPEVN CNQSGSLALPEFWLELIYCGKDRSC NRNITPTSVLFAEQIRGFPKNKSLHL GCELFGLAGNYAVQLRPMIPAQNVP TTRRLLSVDWSDEFVFNVYARSIFPC DPHTGIGVLYEYPGCILEQGDRVRLY ARLRADVASLKPPTSLHYVSEQRVVK SQHSLYFACEHFSEKYVEYCFVYVS QAISGAVADVRMDCVPTLPVSDSDT GGWGPWSEWTPCSTNCLGGTRNRY RFCDSPPPRYGAKFCEGPSVETQKC GKSIADTWDCIYETSGTVITKENSTE VSQEIGPGCRCGCIVHLGSSKPKRIIA GATQSCPGRSFWLIQVDGEESISLAL SFLRLPCATQYIKVRDGPSLSSTLLV ELNGGGLILKGAGVPVVVDSTGGQL LVEFSAGDGPAGNATQEASPACTGG FMANVQQRLATRSSNSSITLVAATR LSGKTRSVVHKSMSRLRFTLVHLSA MIFASIIIIISSLLGAQYVVRYRKYHLA VARRQDEGSRLHTPRASLSSLQGPP SRAMSTTTLLSEVIYMVKMRPKHHL RHSILRESVDAENLTKETEFKESDSQ GMLAKCAEIKELGSSASMVTLRNER SSPARSLDTGSIIGSPSSAEAELAAVH SRSESKDSPSDEPIEPPSCKDSARDS ESIISSGMSSLCNSPPMNSKRLSKYS DRYDAVTLKLLSSRLDESLQDASSLL SVTESFRLGRLRTRSVSSSLTNGCYS PAASVVSTATIRTTSNPKESKEKQNR RKLLARPGSEFSLGNQEELEMDYYD YNVINAGSAPGSYLGMDPAYLIWIPP FEEVAEREEDDKTPEPEPDPERDER QPLYEEIRMPKYGHYLSPASNTESSK STTADNTPSSEERSPPDSGRPSKATS PCDGTERGGRSMPKQPTPFMSRKQ RRQSKQQMQSSLSLSAGSLDHEQR RSSKVPIEIAAVHQKVETELEPNDSM TGSYVEKETAVEKSSGDLQEFLSLD DIQFADESGSDYEAVNLKKVPKSEL QLSQDDARSKILRRKEPKSKEVAV	False	True	False	False	False	0.656	0.469	5.0	1.387