	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	Q7KV35	Q7KV35_DROME	Drosophila melanogaster	NaN	28.06426	T163	NaN	40245499	MNFSKKLLGDKNLVRLDSDATRTFC PQVSEGRLTLENVIFDRGGSVTTKKP RKRPTPKPVIRNFNNLSVPEEVDVD NLQGQRKGVAVNSNNDEMMISSSD SSDSSDDYSSFGDDIFTPGPETSDTS DGDSSCEDELKIPDFKSSATSKDEKL IPSSKWNFTLTKDIIPPGEGKKSHIG ASLPEPVNRNFNNKSVLDLQGQRN QGSRFGAQGVAVNSNKDEMMISSS DSSDSSDDYSSFGDDIFTPGPETSDT SDGDSSCEDELKIPDFKSSATSKDKK MIPSSKRNFTLTKDIIPPGEGKKSHI GASLPEPVNRNSNNKSVLDLLGQR NQGVNSNKDELTILSSDTSDTSKEE KMIPSSKRNFFLTKDIIPPGEGKKSH IGASLPEPVNRNSNSKSVLDLLGQR NQGVNSNKDELTILSSDTSDTSKEE KMIPSSKRNFFLTKDNIPPGEGKKS HIGASLPEPVDRTSNNKSVLDLQGQ RNQGSKDEKMIPSSKRNFTLTKDIIP PGEGKKSHIGASLPEPLNRNSNNKS VLDLQGQRDQGSKDEKMIPSSKRNF TLTKDIIPPGEGKKSHIGASLPEPVNR NSNNKSVLDLQGQRDQGSKDEKMI PSSKRNFTLTKDIVPPGEGKKSHIGA SLPEPVNRNSNNKSVLDLQGQRNQ GSKDEKMIPSSKRNFTLTKDIIPPGE GKKSHIGASLPEPVNRNSNNKSVLD LQGQRNQGSKDEKMIPSSKRNFTLT KDIVPPVEGKKSHIGASLPEPVNRNS NSKSVLDLLGQRNQGVNSNKDELTI LSSDTSDTSKEEKMIPSSKRNFFLTK DNIPPGEGKKSHIGASLPEPVDRTSN NKSVLDLQGQRNQGSKDEKMIPSS KRNFTLTKDIIPPGEGKKSHIGASLP EPVNRNSNSKSVLDLLGQRNQGVN SNKDELTILSSDTSDTSKEEKMIPSS KRNFFLTKDNIPPGEGKKSHIGASLP EPVNRNSNNKSVLDLQGQRNQGSK DEKMIPSSKRNFTLTKDIIPPGEGKK SPFGATLPEPVGRNSNNKSVLDLLG QRNQGVNSNKDELTILSSDTSDTSK EEKMIPSSKRNFFLTKDNIPPGEGKK SHIGASLPEPVNRNSNNKSVLDLQG QRNQGSKDEKMIPSSKRNFTLTKDII PPGEGKKSPFGATLPEPVGRNSNNK SVLDLLGQRNQGVNSNKDELTILSS DTSDTSKEEKMIPSSKRNFFLTKDII PPGEGKKSHIGASLPEPVNRNSNNK SVLDLQGQRNQGSKDEKMIPSSKR NFTLTKDIIPPGEEKKSHIGASLPEPV NRNSNNKSVLEEVDVDNLLSQRNQ GNRFGAQRLVVNSNKDELIILGSDS SDTSGDANLCKDEIKIPSSKRKLTLM NDMVSPSKRAHTEAVGPTTPKPWAP VLADKKDNDYPYGGKEWARKFLEK KKTKAEVANPTLAEVSPQEDESKVV KELLETQEQLDMVEEMERKEMEKE KIDEWRKTSKQVLEKNKKEEKLNLQ KPKMAKGKVQKKKSKDRQCGKTLKI AKGKLEALKTAKDTMAAKVLEKQMI MKKKMAQKMSEIEALERHKKYVEIK EKMVMGNGKRNRSAPYRYLKK	False	False	1.375	2.265	0.521	False	0.406	False	False