	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	Q9VI72	Q9VI72_DROME	Drosophila melanogaster	NaN	27.156312	NaN	NaN	28604694	MQEAGESPVGMPLSQEAIAQRLAAL SRWQDEQKRLLQERQSNHRVLLGL EQRNMYKMLGLLHQETESHENSVL EEWDEEHSIQMPRLAADPDEHEPEI SMADPQPAKPKRPFLRRGEGLKQRF KINPDQLRLENLPRYKFANAHPQFR TPQMKKGILKKQVKSPPNPAPPPAP KPSAQLDDLNEQRFQQMLIGKNAC SSTPDLKSSYASSTTASSTSPRVRFV EQKSRAGQTTHADDEASSEASPMTG VCWAKVLDTQSIKPAQIQRRSAQIRV EDDSNVSIFELLEQKATEGNIDMNS SCIRTFMARKDQRRRVADDSDHIVV TQQVRQMRLQPQVDQMLVQELQEG DEEDTEPSSDQTLTMTPIQVGNTKV RVRFSDSNDTHEYSDATSLNDGSNI QLFEQFKSALFQALEQKKKSSPSKE SDEPITKDLQEKANLVRTRLEELETE IATFKEQNAQLLRLRQQHELEKAKC TQDHMEAMERVHDEKIQAEIYLHD ERMKIEEERRKFEQQMRLQKSNAN SKEKKEIAALKQEVEGLQLQLKQKE QAHVSAQARLRAQLRASEKEQRNYR DEIELLRKENKRLEQELVKIGRENN SKMLQEINRNIARLAPKVLPSATMS DILDENGRRTQSLDGTAGKPAKQRE AQNRRQSSGSAQVRSRSRSLSRNKK TPYALDESYASSSSVESEEVQPPVTA PKAATPPPAANSSSDFKREITNADGS KDIWYPNGNLKKISADGMNLRMLY FNKDIKETNIREGTVKYYYAETNTW HTSYLDGLEILEFPNGQTEHRRKDG TVEIHFPNNSIKIVDPSDTEKLEEWR YADGTHLVQLRNGDKILNLPNGQKE IHTKLNKRREYPDGTVKLVYPDGSQ ETRYSNGRVRLKDKDGKLIMDTDYA KY	None	None	None	None	None	None	None	None	None