	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	Q9VWL7	Q9VWL7_DROME	Drosophila melanogaster	NaN	27.156312	NaN	NaN	28604694	MPQQAKDINDNVGMEAHIYENLPG LTSTPKGHLNAELRKYFPREAIYENL CRGCGYGVFAAQGRYCHFCQCIVSG GVVSTPPPPPPPPLSTISGESSNIYEN ICDLCRGIYSDNERCRCQRSETPAKA PAPSTPTTPPTPPTKVKSRSLLGYSKS SAATLTRLFGSLKQLNRSSSLQRRIF PRTPTTGSSSAGSLEIIHNVDEVFRT QETFDLQRICELKLEQQQLKGSQSD QHIYGRVREPEESQSHKRERKRPRV SRIRLLQDESTSGSSSLYLNESICQW MTSLRRQVHHYDEGPWQQPKSVPA RDPSSSAEESTVTLRRSTEESAKCAD EQAKLVKDFKRKLQQKRAHNVEPE REQRRCAITANFVNDYLSALGEEKE AVQHQQVAAIEEHVAVVAIAIQPQVS VSLLRLMQSLALSASLHVSVRGFLT AYDKSFSLWLWHQQQQQRQQPPIN GYRRRRARDSVVGVVGDWRRRLPLF AFSFAGGNDRSVASASVSVSPAISES VDVCAAAASRRRANRSTRRRGTAIV REEGVCGGESSTSSNNSKKNNNNN SNNNSGALMAAPKPNGEAERPSTN SNNNTINCSSNNNNDNISNNNNN HHTVDNESGINNNKNNDGNANCN VNEQQQQQQQQPITQNCVIKVRRPA PKVPVRNPNTSLSNPRQIEEEQTKP NGGKKDNLRGNSFAAENSSPEEQE EESIYQPIWQFKTIEPVDQEIIEALPV EGEDEEDEDEEDDDVDDEEEDLGD YDESLPSDADPDVMALQAAAVFAAG VRHKLSTSTLTKRSKMMMPTPLPSP TASMDHGAWETDMEFMYSRESKEE ADETLSLGSTVTNSTATLGSISSTGS SVTGQPKDTPKDLPADQQPSVVVHP ILRNICIFYSQKDPKKYSAIFYDYHLS LVHQRFMTRIRRPAPPPPAQLATSN TTSLAPRSDDSGCSCEAEDSYNHQH NPTPAIPKPLKEEWERAVLAAGAVA GKRGVAPRTGRKLRVRSNARLLLTD SVLAWRETLQDVNCCEDEEDMIVPV TDILRAQQIQEDNEVQQTQQSTILQR YKSVSIGNLLDLPGEDDKKSIKNRM RMKFAVNSNVFRINRSPKSEKKSRS RRGQVNSLYLDEQKYPLFAAPLSAL ELNMTDHPNVPRFVVDVCAYIEQPE CIEQDGLYRASGNKVLVDELRKKLT HVYDPRWLKTDDIHTLTSLHKQFFR ELTSPLITQEAYERLGRSLNDDAAIE RMSLAFDDMPEPNRSTLRFLIRHLT RVAAASASNRMPSTNLAIVWGPCLL SANQIQLDIGRMNMLAKVLIENYDR IFHPDNERLVC	None	None	None	None	None	None	None	None	None