	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	B8A429	B8A429_DROME	Drosophila melanogaster	NaN	28.06426	T535;T985	NaN	40245499	MSSSVFTEAPSARPAAAEADSNLVV VYSKSGISFDERAIGNIMEEKSISTIS VVRITSPTPSMVDQEEDERLEELAN FMETATDSELDSDNGSQSGGDNAS EMDEHDEENHNTGDDNNSDSTET EADAPRTRKRVGRKPKAIKKRKRRK PKERPQIVGLRVEQFYADNTADMVV KNALKLAGVSVYKRTAETDALMEVI RNDHNYTPFTSPEQLKNHKRDEKM AMEMQAQSRKIIMQAPGSVKLINAK KRVQAIPLSPVQVKVQRLPVKPHQV QQAAQLQPKVQLIRKPIHSPLPRQKP EIIANNSVNARLRPTRAPVKVIAPPV QEYIEDDDDAQSSDSAEEENADKAY DSEEMSEESDEFQESDNDRDSDMD FDMRHSNRRNPNKRKRVRKVVRQT PSKPMKPLPPPPTTPQHFPELKKRK EVLEPKAPQIGQIIQMPATKSVPAVF ALGRGLPSTSFLKIRPTHQSLPVKRP PLPTTPANSNVRRMPAVQLGRIVNS PQEVKEIIINKNMASPKGVFTNLNTL LGEDNNAATKSSPVPLRHRAVMSPS TPRSSYNQQMSPIVVPVPAQAHTPS KGFMPIGVDTAQSHKLPAQIVIETHQ SSSELAAENDKQLDLINSIVQDELLK STLVEQPVVNADEDIPKLVKMLEST AADLDPPPVSLPTNIFPAPEMQGVG NAAAADPNNIMDTANEDEITADFLQ HVVGLIEEDKQFEAEVVKQVLASTEP GTLDAIVSMPTSIEPVDVPQVQAHTN LLPNASLTEPAQSMTSLPIACSTPSR SVAASTPPTSAKVVRGYGRVIYLPPIE APTTRAKRRAQFPSAPGMAATSSSD AGNLSFGESSLDASINQPLNTSSLS NDSQPGSGPKRPNPREPSMARRSTA PRRSKKLDASQNNDPDASESQEDD DDPNKLWCICRQPHNNRFMICCDL CEDWFHGTCVGVTKAMGTDMENK GIDWKCPKCVKRQEERSQPRITDML VTRPTTQPEQRPSETKVLTTTAEIVQ VAAPSAPRRTLPVVLTVASSPMRIPM AKAAKKFPTGAISHQQQQQLNFIRL GPSPGKRISETLCVVCKRPASTSSVY CGEECIRKYAQSAIQAHAATKGPLPQ NAGAQSLLNNSFDAKKNKKKDLFE DVLRQADTVSKVERVCNKG	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None