	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	C7LA93	C7LA93_DROME	Drosophila melanogaster	NaN	28.06426	T899;S911	NaN	40245499	MFPNQNNGAAPGQGPAVDGQQSLN YNGLPAQQQQQLAQSTKNVRKKPYV KITEQPAGKALRFRYECEGRSAGSIP GVNSTPENKTYPTIEIVGYKGRAVVV VSCVTKDTPYRPHPHNLVGKEGCKK GVCTLEINSETMRAVFSNLGIQCVK KKDIEAALKAREEIRVDPFKTGFSHR FQPSSIDLNSVRLCFQVFMESEQKG RFTSPLPPVVSEPIFDKKAMSDLVIC RLCSCSATVFGNTQIILLCEKVAKED ISVRFFEEKNGQSVWEAFGDFQHTD VHKQTAITFKTPRYHTLDITEPAKVFI QLRRPSDGVTSEALPFEYVPMDSGK HTFWNLHRHLKRKPDEDLFQQILRL DAKREVQPPTIEVIDLDTPKIDVQREI PSEMEFNHEESQQSEPALEQEQSV QQEQYTQEQSLQQEQYTQEQSLQQ EQYLQQLEQQQSFQLEEPMQQDQE LPAQQSFDQAIDHLPDHTSDHIPED MEAADAHAEAEAHRLRSEQEKEIDT IIDEKVRELEQLDLGQQLEPRPLTAN DKITEWMKSSEIEQQVHEPSPTAEA DVLDSALEISKADKTLDELLETVAEL DEIYTDFKVQRDTYKNTIQNELAGLQ GRAPLQVEDSFDDAATYTSLQIAFKN PVLIPMDDIMPPTPPMSQCAPEDAH QHYDPVEVNSQARKPETPMRPVPPV PPAILTIQYPPEEDKLPPLPPKRIRKQ DSNAENRSIEANTVQTKPSTGESPL NKRLPPAPKNPNFNTLPRQKKPGFF SKLFSRRKSKPDLAQGQENSSILDSK ANSREPSIGHFNMQDPMRASLRSS KSAAPFISNPAPAKSSPVKAKKPGSK LTKPVGRSVSSVSGKRPAYLNADVV HIPLKGDSVNSLPQQQRTEGYSQSS TISVGAGLDRRTASALQLADIPISEGG MELVAIADRQSLHNLVSSIEGHFNV QLDPNLDLTEAEHFALYTSIPPLAAA SEFDETSAYYAPVDAGEILTPDEVAK RLAAANGI	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None