	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	Q8SX64	Q8SX64_DROME	Drosophila melanogaster	NaN	28.06426	T627;T981;T990	NaN	40245499	MAGTATATPMEIDEFVNSSIISWLES CLPRAELLAGYTSLLDGHIIHSVWLQ IDPEPQNNPSELNDLNGKSLSIARAK NFECVVRNLKSFFEEELGQTILVLPD AFTLGHHPESKNGLEQMKTLLTLLL GAAVQCPNKELFIARIKELDLETQHA IVGLIKQVTDSHSLVLTEDSLERLTP QSMYTHILRLTKERDVMYLKWIDLA CVETEMTASDNLVECGQGVSVTRSP SNGTATSTPSSSSNSESNHLAVECA DLRSKNRKLRQELEEKSENLLELRE ELDDKKARFDKLRQESQEWFTEAKR ASAYRDEVDILRERAERADRLEVEV QKYREKLGDSDFYKSRVEELREDNR VLLESKEMLEEQLQRYRKRSEHAIS LESEIIKYKQKINDMALERDVDRSKL EELLEENSQLQLVARNLNSTMDLD KSFSENEDDCNSGDNSLSEQLTNN AQTRALKLELENRRLTAALEQLKES SFHESTSKMLELEKEKKKLSLKIEQ MQENINRLTQQNVELEGVFKNALE ENKKLQDAVDNRQKSYDRQSLERE ADRQKLSDAEQHVETLNKEKQRIQT LNESIQRRADDLERLAESKTKELEQY LEKSRQYELTKQKLYEIEARVSTYER ENASLLKEVSKLKEGSEQKSVQLDD SINRLDVQSKELQKLGKALEDSEQV HQKLVELEKQNQELASQRIIDQEMI STLRNDLVTGTLVTKKVRNNLEKLG LADEEPGELNVEHVVEKLVRNPETF KTVREIMLNVTREQLEEEEREGGVK SDMCVLCHRQEIFTVEKNIELAATPA PAPAQPSSQELRFEHKVRLSPARES AELTRIKDSNTQLQTENARLSVDVA ALGSQITSLNTQHVALQLANSQLAA EKDSLLKEIDSLQQEHKHALQDQVT LQCLHDQLSAEYESLNKDKEQLKAA VRDLRQELRDTREQQSALEQRIEEL TIQNSNMKTCSEDLSILRTEHSKLTD DFRNLFATSDRFKNEYKNIQEQYKM VRMEHSSLKLQNTELSGELNAKSD QVRCLQMEYSKVQQRCEMLIQNNA ELDSERKALMDNVSQLLSQYQELLA ISLEDKKHFHEEEKNYTERVHSLKR QKEKLEEKIMEHYKKSETTVHKKKP FASMLVRRVKKASSDLMNKVPSRN RRSWVDDARTNSQFVIGSESGGNES DNSNEEPLSIASDTHLLQRNVPLRQ SLQRDLLDNSIQRGGAVRSSLQAQK LLKRQT	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None