	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	Q6NN63	Q6NN63_DROME	Drosophila melanogaster	NaN	28.06426	T923;T925	NaN	40245499	MSATRAAEARAEANPENERDPPAGE ESIDRQDEKEKLYAEVLHTIANTVGA PAPGGQFAHYKDEMYLHAQRAFGV NPDRHYRLLHASAEDKPKIIVLSAVV IEADGLEAKDANGFSDPYCMLGIQP DENGGPPVSPLPLPPTPRALSADMS GGFDNSATDSPPHRKHSFRLSFKRR EAGRRERRDSLGGPVPAKLIKATSIK PHTLTPKWNEKFKFDIDDINTDTFH LDIWDHDDESCVMDAVSRLNEVRG VRGLGRFFKQVCQSARQGSQDDFL GSVNIPLSEIPSTGLDAWFKLEARSH RSTVQGRIRLKLWLSTREDRNTVSE ENHEQQIHKVEQLQRVFMQHEVTT HEPSWTWCGDLPGPALTILHQLAVQ SDLSDLHCAMARYVAAAKLNRSTPL DPKFLHRLLSDVEKQWNQPNQEAL SRDLEQWLADAMNGFVERSLNQIR RHRDIFPALNPPSLIRLEFLLRCLGL LGSMRAFRAVCPFNKGVRGEVVNA LRKGSVMWGQTVMRESQSMPNPLS NFVTTLTADIQLGQTYYHALFDNTN GIQYFSIIYKQYDAMVGEEVYSRMAA GQVPGVRMNLSQYAVLEGDAEPVD TKPFELFLALQEFCQLKRHLSAQPQ PPEKPLALSNSHEWFIPTLERWMM VSKAKALQRIRAAIRMDAICEGERIV RYSSSSVDTASTLLNIKEFWKVLNW PDEDTAIMMESQLIDVVCSAAMHYC DLIHTALADSGYYEQQGPFRCSDDM CVTVNNVEYVRRTLAEFRSDEQPLE ESADNLLESSLTHMENRCERILSKL APLMQMSLQKAMFHLAWSPDSLPA NQAIVPLLEYLDCHLAALNSALLTKN FNRSLRFIWKTVLGEMSRQMETGD ETDKPTHFHKRLYEALQLLIDFFHAE GQGLLLECLHTEDFWRIEQRLQFHK TDTDRLIDMFYMQRLREQLMAVMP SPYGSLAVRAYFNHDSLCVEVLHAR DVVPLDPNGFSDPFVVIELLPRRVFL HCMEQQTNVHKRTLNPVFDECFEF SVTLEQCLTEGAMICFTVMDHDVLT ANDFGGEAYLALGNIPGVADYSTSV DNFHGLKQIELPLMEQKDKCNPILQ ILEVRVNDKQAQDFVRKQKSRFIN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None