	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	Q8T9B4	Q8T9B4_DROME	Drosophila melanogaster	NaN	28.06426	T485	NaN	40245499	MAETLVSIKRVQKYMQSDETNVMD MSVDLTEDFQGSNQETVHADGDEE RDEAEDKLLGPPIATVNENAKLSEA GISISGLMAKWDVNSPDYSLNGVNL RVQPGTMLGIVGRTGSGKSSLIQAIL GELPAESGEIKVNGSMSYASQEPWL FSGTVRQNILFGQPMDRRRYAKVVK KCALERDFELLPFKDKTIVGERGASL SGGQKARISLARAVYRETSIYLLDDP LSAVDTHVARHLFEQCMRGYLRERI VILATHQLQFLQHADQIVIMDKGRV SAVGTYESLRESGLDFASMLADPER DEQSEERSRSRSGSYTHSHSDQRRN SEQSLLSMADSCMDDLEAEQANNQ ERQEAGQIGLRLYSKYFKAGGGFFA FFVMMGFCVLSQGLASLGDYFLSY WVTKKGNVAYRADNNDTTRSEELE PRLSTWLRDIGLSVDAEMLDTYIFTV ITVLTILVTVARSFLFFNLAMKASIRL HNSMFRGITRAAMYFFNTNPSGRIL NRFSKDMGQVDEILPAVMMDVIQIF LALAGIVIVIAVVNPLFLIPTVVLGIIFY QLRTFYLKTSRDVKRMEAITRSPVYS HLAASLTGLSTIRAFGAQRVLEAEFD NYQDMHSSAFYMFISTSRAFGYWL DCFCVIYIAIITLSFFIFPPANGGDVGL AITQAMGMTGMVQWGMRQSAELE NTMTAVERVVEYEDIEPEGALEAPA DKKPPKSWPEQGKIVFDELSLRYTP DPKSENVLKSLSFVIKPKEKVGIVGR TGAGKSSLINALFRLSYNDGSVLIDK RDTSEMGLHDLRSKISIIPQEPVLFS GTMRYNLDPFDEYSDDKLWRSLEE VKLKEVVADLPSGLQSKITEGGTNFS VGQRQLVCLARAILRENRILVMDEA TANVDPQTDGLIQTTIRNKFKECTVL TIAHRLHTIMDSDKVLVMDAGRAVE FGTPYELLTLADSKVFHGMVKQTGH ATYESLLKIAQKAFENRQNHSLSS	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None