

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9VG78	Q9VG78_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MSSSVFTEAPSARPAAAEADSNLVV VYSKSGISFDERAINIMEEKSISTIS VVRITSPTPSMVDQEEDELEELAN FMETATDSELDSDNGSQSGGDNAS EMDEHDEENHNTGDDNNSDSTET EADAPRTRKRVGRKPKAIKRRKRRK PKERPQIVGLRVEQFYADNTADMVV KNALKLAGVSVYKRTAETDALMEVI RNDHNYTPFTSPEQLKNHHRDEKM AMEMQAQSRKIIHQAPGSVKLINAK KRVQAIPLSPVQVKVQRLPVKPHQV QQAALQPKVQLIRKPIHSPLPRQKP EIIANNSVNARLRPTRAPVKVIAPPV QEYIEDDDDAQSSDAEEENADKAY DSEEMSEESDEFQESDNRDSDMD FDMRHSNRRRNPNKRKRVRKVVQR PSKPMKPLPPPTTPQHFPCLKKRRK EVLEPKAFQIGQIIMPATKSVPAVF ALGRGLPSTSFLLKIRPTHQSLPVKRP PLPTTPANSNVRMPAVQLGRVNS PQEVKEIIIINKNMA SPKGVFTNLNLT LGEDNNAATKSSPVPLRHRVMSPS TPRSSYNQOMSPVIVVPAQAHTPS KGFMPIGVDTAQSHKLPQIVIETHQ SSSELAENDKQLDLINSIVQDELLK STLVEQPVVNADEDIPKLVKMLEST AADLDPVPSLPTNIFPAPEMQGVG NAAAADPNIMDTANEDEITADFLO HVVGLEEDKQFEAEVVKQVLASTEP GTLDAIVSMPTSIEPVDVPAHTNLL PNASLTEPAQSM TSLPIACSTPSRSV AASTPPTS AKVVRGYGRVIYLPPIEAP TTRAKRRAQFPSAPGMAATSSSDAG NLSFGESLDASINQPLNTSSLSNDS QPGSGPKRPNPREPSMARRSTAPRR SKKLDASQNNDDASESQEDDDDP NKLWCICRQPHNNRFMICCDLCE WFHGTCTVGVTKAMGDMENKID WKCPKCVKROEERSQPRITMLVTR PTTQPEQRPSKVLTTAEIVQVAA PSAPRRTL PVVLT VASSPMRIPMAKP AKKFPTGAISHQQQQLNFIRLGPSP GKRISETLCVCKRPASTSSVYCGEE CIRKYAQSAIQAHAATKGPLQONAGA QSLNNSFDKKNKKDLFEDVLRQ ADTVSKVERINVFERKSGRVTGHM APSAHQFRKWLQENPSFEVLPSGT QSADA EKRLKGAPEAATSTSEPAVL GVAKKPEGPAKLSHPQNTTVQASH QLGISSVRPLAKDKEKTTPTVQAPT PNRIAAGKPEPVIRIGIRRLREQLLA RIKEAQAAEENSGQPTTQWPVLE DQFVKNVELEMFN SFGRDVGAKYK AKYRSLMFMNIKDRKNRTLFEKICAK QVEPRQLVRMTPEQLASQELAKWR EEENRHQLDMIKKSELDLLACAQNY VVKTHKGEEVIATKVDVTLPEENVSE ASSMADTKQTSLVSEPSTTLERSTS REKAGSSKEKRHKSHKHHHRKRSR SRNSRGRSVDKRHRQHNEGGEQ PGSGEREHRSREKQSRERDEIVPSP LPKKRDENDQSPVPKKTAEKKTEAS AYNLVDQILESEKTVEQAANLQKPK PSPKPLPTLPSLKAPEMDNYSRYL HGLTSSSLWSGLK MIDLADFEIVM YPVQGNCHQLGNLMPQMDVIGRI TRVNVWEYIKLKKSPTEVVIVNIF PASPSETYKFDLFFEYLD SRQRLGVL GVDSDQIRDFYIFPLGSGDKLPPALQ TAEVPPFYEEAQRPN TLLGIIVRCLS KRPADAVPSTPSVPTPV PALSSKVAK RSRSSISYPAQSSPKRKISTHSTSSKD DEFDIDAIKAPIAKLQKTAPKVVQP SVPNDADEPYS PGGSDDELVPSAPQ RPNDLERQVNEINKQIAAQMEIAG	None	None	None	None	None	None	None		

