	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	Q9VC19	Q9VC19_DROME	Drosophila melanogaster	NaN	28.06426	S93;S97	NaN	40245499	MESRTQMNQSNQQQQQQRNRLPS SPSPPLSVVSLTGGPNANTTLMGLE HDAVPSSQLGRNNNNGGGHHRSH SITNAATTSLANLLDGFTARVSTAVS VAGGTNMLMDRSSQSAASTPGSTH SIASLTSMGSLGTTANALYSSIGSAL NSITSPIQAAAAAATATTTEVVVAAAS VITNHLSSPTSGTPPHGLDVDEGDD ECDDCEEDEDDPVDNDGHIAAPTPN KSWAENKQVVQEIRKKMCNLSSMV PTNVQFRHLSDGRARCYFLGTPPQS WETTLLFADINLTQSEEQQLLVQRL EGIASDEWSPTMNAGSPTSSGHQPA FLFNSLPRPRLPWSPLLQQPIQSSGG SGGSGSASPYAREYQLLQERKRLST WGITSYELHKPSGKLVFPCFNDLYQ CLDTGYNSGLLFPTQLRTCPQWTAL DPQICPQNSDMIAYISDCDLFVTHTL SGHEKRLTYTSTGRHSYVDDALSAG VPSYVMQEEFSRYQGFWWQPHSND GIYRIVYEEVDESEVSVYTFPSSTAM HGRVDEYRFPRTGSPNAKSKLKLVQ FVLNEALQVSEIAIKDLPYSLLAVFS WLEYIVRVGWTPDAKYVWVQGLDR KQQRLDVILIPLDNFCESYSSQVSTP TDSIGDHSWRSLYSRTITPLQVIYTER SDSWINVHDMLHFLDLTETSVTFL WASEETGFRHLYLVTASLLLSQANG QPDPGSVGAQPSFVDLSALQPRILNK VALTSGEWEVLARNLWVDKANKLV YFVGLRDTPLEKHLYVVSLERPEHIR LLTEPGYSYLVEFDDQCKLMLLVYC NIQRLPSCKVMRVNQTCSNGGVNGI QISLVGYLHEGGKPEPQYCPQIFSPQ LPSGDIVYAMVFKPHNFELGVKYPT VLNVYGGPEVQTVNNTFKGKHQLR MHMLAAQGYCVICIDSRGSRHRGKR FESHIRGRMGQVELTDQVDALRSLS DQLGYIDMDRVAIHGWSYGGYLSLM GLVQYPKIFKVAIAGAPVTNWEYYDT GYTERYMDMPQNNEAGYSAGSVLE YVNSFPEEDKRLLLIHGLIDENVHFC HTSRLISALNKANKPYEVHLFPEER HSLRNLESNKNYETKLLSFLQNL	False	False	1.789	1.781	False	False	False	False	False