

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
Q9W3G1	Q9W3G1_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MSAVYSPQPPQQQPQQQQLQQQQ QQPPPPQQQQQQQQQLQQPPNSA PNQQPPGSGVGGGGGGGIPGGGGI LSMGMQQQQRRPPMGVPGSPGSIIS GGGGGGVIGGGMPGGGMMVTP TSTPRGGANMQGGGGGVPOQVNS AQIQKMLDENCGLIQTIQDFQSMGK AQECMSYHVALHRNLVYLAQLADPA MNISQILPPPHILQTQAMQGGGQQT PPTGPHGMLGGPPQQQQPGQGPV PGQPGQPPQMGMQHGGDPQGP PVQMPPYGAQQQPQPHPLPPGAQ QQSQQQQQQQQQQQQQQQQQQQ QAAAAAAAAAAAAAAGQQQGPQ VSQAGPQQQQQQHPVYRNAQGGG QPAGQVPGQGGQPVQSVINPNAAP NQORPNNGLPLSGPQNPQQQQQP QPGGQPPNQQQQQQTGPGGPGP QPGAGGPGVPPQSPYRVSYQQQQQ QHSYYPGYPPQPTQYQPQGAYPYG PPTQGYGPPPPPPNAAQGGYHHGP AGAATGASGHGYQPNAGAGGGPPP GAYPPPGSQVPPVPGQQPPPGP PPPGQPPTGGQQPPPGPPQSQYGP PPPQNSAGGPPPMGYAGYPPNPGQY GQAGAGGGPPPSGYWPPPTSSAQ SPYQAYQQQQQAAAGGGAGAPP GSSYPGGPPTSGAAPPPPGAYSTT APSQTPPPQGGGGAGGGNNNPNGP NAQQSTPPPGGGAGGGAGPSGPGG AGQYAGPPQQPPQQPPGVVVS GVAPLFTQVQPTYSTPGNYNQPGA PPPNQQQQQQQQQQQQQTPPS AGGSAGGGGAPNAQQGQNPQPPN GATPPMPPNQYPAPGAPQGPYGGP PPPQAYGPPPGSAYPGHAYHPQ AGGYAQPPPTQGYQYRPAQAQMP PGAPQGGPPTGAYGYSSQPPQ	None	None	None	None	None	None	None		

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