

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
Q0KHY3	MESH_DROME	Drosophila melanogaster	Protein mesh	25.496522	NaN	S1249;T1251;S1254;T1295;S1297	33925313	MRFKLFVVCALACGFMAFVLANENI STEEFEDQINAVPEPVKVKPKKAQA EFVEVPKATPAPKKAEDNPKTPLA LQKPTLIPVVHVSSRGTTDDILTVAGL KPEKMFALIMPNDYLGEYDNDNS GYNWDPNPNVAPPTTYSTAAGGYTI TAARLAELRSNFMYWYFDKDMYGG RGDYQFDIHASMTQLHKNLNFQLP FYGFRFNYTRLSLNGYLEFSDPPEYL TYPLVFPKDWPAKRDPSPMGIFFSK CRVGRIYPSDIDQRTPGVYFRVERDL MGRTDRFGVEVRETRMWDIRQGVV GADTFIPKHVVIATWKNVSFAGGIDN SLYTTNTFQMVLTDEYTYIIFNYA VLNWLSHTEAGDITKGEVGPAYV GFNAGNGTQAYEYNPYSQNMVIRD LANRGGWANGFPGRHIFRVDEQILIG SCNKDIDAALLPLTFAPESGNMLGG QVVNITGPCFDPFAIRVTCHFDTESVL GTYVDRNRVICVQPYLKAEGYIRFQI SVGTQRKWRGKYFVETPAAATEKI FFTTDDVHKKNPAEIRITWNQYNLT SNANANVMISLWGYRETKIEPQLEYI DVIEASYSNSGSYVITPSNYINRNNI NRDMQFGFLQINLTQPDQYSGLAIS PVLWSRPIPLAWYMAPQWERQHGK RWARALCDNWIRADRFLRNFAADL PLCPCTLDQAVLDKGRFRPDRECDK DSNPSCLRHRAIHCVVSGTPVAQG AEQQCCYDRYGFLMLTYDQMWGSR PRRVHNLGKMPWNEASKVPSLSM WFHDMRPFYSCCYWQEEQAVGCET YRFERRPSQDCVAYQAPGIAGVFGD PHFVTFDGTAYTFNGLGEFVLARSV DESNKFEVQGRFEQLPVNYYGEVKA TQLTAVAMRGNTTTIEVRLRPLHA RWRYRLDVLADGRRVYFDRESLKFQ HFDGVTVYPTYLLNQSQVVVQFDA GIGVEVVENEGYMTGRVFLPWKFIN KTAGLFGNWSFNKLDLDFMLPNGQV AQLNLNDRSIHTNFGIKWMLTDRE VPGVGAALFKREFGRMSGYANATF QPNYVLDPADFLPANRSYDLERAAE LCGECMQCQYDYAMTLNRDLAHT KNYYDITVNMQAENAVRVVSCGVLQ TPRFGRKLSDFMPGAKVSFECNEG FILIGDQRRECLSNGLWNIPEYGYTE CLREVVYTRRVAFIAGIIFLVICPLML CIVCGVYRQKQLKEDPOWQMPM LPRSASSARNLRTLNYDDSDTDA STLKKRSYDKVYRTNEPLPGKQID FPAKWDLDENDEDTSSSEGEKPK QLGRRSLHSASGTELDQHGSPVDGD HPDEDDDFDEADVHTGTHPVGYH QPLSPEEADQLHRQQYSPTFSGLDS RTSGASSINYPQAATQNRFGGVPL PSNTQYSRPPSSVNTVPLRTAPTPL TQDTGLPSPLATSPTPSVQKSTEV	None	None	None	None	None	None	None		

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