

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
A0A0B4LIF3	A0A0B4LIF3_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MANVNRDSDTSLWLHNKLGTSNDS WINGSICSQLNKEVLRNKECFDDL QTQVKLKLKLLSFLHIPRRLVEEWKAE LEEVEVAGLDSELWVSMMLAETMKT FPATSSLNTEISDYEDTRPFIIDMVN DLRKLVTKHSDLGMLPLECQYLNKN ALISVVGQQPAPVKHFTLKRKPKSAQ LRTELLHKSADAQSSLKASAPTPL RSRGMPRKMTDTPPLKGIPSRMPTT GFRSATVPGNAQRPNLSRTPAGRK DGGIKLIEFTEQPLGYAAAKRKRREQ QLEEQKKQEQQQAAAAA TAAASNAGDSSPTDAATASGGGETPV TPTSANNSEIKMEPQQLNQSAGSL EEPQDDMSGKANMECDIETPEYAT ATLDFQAATSTSVADGQPGKPAVET KLKTPRTPKSAAKLNNNNNNNNNN SFNHTPKRIKQEIIEIKSEEHVPASIKL EKIETSPSTQRVSIQQQPPSLVQRT HLLIRSSPQKRQNNGATTSAGTTTTT VGNTTIKMEKLDIKPMIRATGVSPST SAGTTTTLLTPQQLRQAANPLANLP NNISVKITSAKAAAAAATSSGSQ TQSQQPOTLQVQAPQQSHPLLIN SSTPVILASSPSAQAALALPSSSTSA TTTTTIPSAIKTMPLSQLKTATNSG PVIISQTIQPAKRAQQAGTSSAAAA SQQQQQQSQQQLLHQIQQQTQQ QAQLPQQPQQQTQYILATPQQQPQ QIQQQQQQPILPTLTSFHSRPM PQTTLYQATTPSGSGQPTPKILLKTS GSSSVMTPLRQAQPQATAVVSSN PPPLVATSAVVSPGQTLNINQNVQL PNRPVTIQPASQAAQQQHMQAQLQ QQQPHPQHTIVANTTATQPKLSQV LMOPTAAVGTSGVSALNVSPSTGKN KTIILTQKGVILRNIGGDMYQQIPISN VGNMQGLGGTTLMTTATGPPSLVKT TPSSGVQLQQQSGKQILPTLIPTSS LGGQHVVIVQQQPTNVIGNSQQTII RPVMTNVGGGLTTLPOGLTLIQRPG QQPQLVQVQAAPGSTQRTIITQSN AAASQQPQQQQQILVQHKPAPTL QQLVTSSTSGGQGGQGNPNAGLP RTVQVQVQAQQQQPQQQATQQQ SQQAPQRRGLSLSNEHVHKAHEMF RKANRVS RPKALILGFMAGLREN RPNNENVLVIKLGETEKEVQQDNG HTALCLVESHIRLDYNTGEWKTQFN YRLQDQSAAS	None	None	None	None	None	None	None		

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