

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
X2JIM3	X2JIM3_DROME	Drosophila melanogaster	NaN	25.496522	NaN	NaN	33925313	MMEQSVNRNQTMSKTTNRRRTAG GIEAPSIANASATSTASASALANASTA ASASAAASASSSSSSVSSATCSATIVQ AKQRPPPLKTATTTVTTLVSSSEGG QPIPPSAHATGAAGGGSGGANASG SYRGATLTTPTPRIELSRASSSSH HEEDSRESSPENVFQAKRSSPHIF KFDDHQSYLQQHQKSDASSEVAAL LCISGRTSRISVGSQGSVAVRLSAIS GVSRSPSPHRMLETFCGPKPVEL AGSGGASNASTAASAHSQADGMP GAGTVAANAALLEQLLLARKHDPTQ AVLAEGIKVLPAPTGSTATTSSRSRS RQTSTLANGERNEKEPKVKQTIGAQ RRSTKSPGQMVVGKTASGTEYIRINL RPDHMYSKGIASNERVVEAPNQLA PVYSKPASLSLQGAPIEHRHTPTAS PKPGRRLTSRSPATGAGGGPGGS VSRKSSFSSLFRLGKSDSPDSPRETAR SRSKSKERPATATGGLGGGGSQNV TPSKQKSVLAIFKPKGRGSSKSSS PIESSVEPVPPPPPPAPGSGSTNRSR TPVGSRPGSRLRYDEPVEGVIHPL HTPPEEREARTLLQGIQLTHAAQA SMDPSPLPTHRAPPASAPVAPGTQI TATQLAAAAALRPVAQRRVAQRPD LDAIQSLPSQEDPPTAPGEEQILQOP EPPNWSLEVQRHSSQDSQETTAESV GSVVSARAANLAQQREQAAMQRLP SVESTQSACEPRLVHTVATVNAPNP DSEAAAKERRLLFATRLGSGSQEQ LFTSQFSIKTESQSSQLSEQVEHSG SETTEGLQRQGTVIRRMETVPPPPPP RGVCSSEEEQGGVMSGVVMRAKR TKSRPSSSEDEAAPTAAAAADVRHS RYLENFDVRRKLHENLTAEVGRSG EMPGGQIAIPKPKRQSVPEPRRET SYSSGTVPTTPTPTQATPVGQAKOP PSSATTPTSKGTPVTTTPADAVQAST SVTGPPGTSFNEPINTTRDQPSTSY KEPPTGTSREPVTTSIREHPSTSHHD SLTSVSYPPPERRHSQSTDEHEPVES SESERDSMAGSSSVTAVPAGGEA GRRHHNFPNRVNCVIEHESTGLVSQ ESFEDELPIPTTLPEERAQGVSIIPM RERANMELKTCPVDRPRSTTPLNPS HLEEYCTGIMTPQEAIPGGYQETDG VGGVTTGIAGGGVAGTPVRGEKLRIS LPRKESIGKERIQNSKSPRRVSNSTSG KSWFEEAEGLRANNNLERKDSAK QLLPVVTPPPAAPSTLEPKPKASTQ RKLSGHWIDFENIPEKRKPAKRITTL PKETLTSSVPATATSSSSACGSGHR SGSGAGHHHHHHHHHHQATKTNP DGGDKLHYNYVKPEDCQCECHAEAE RGESSQATAGAGDTSTETGTGTGTG SESLASVELLQQGEDMLPLEPDTS AEGRIYGDEEQAPLMRHSGKGVNQ PRSQHEEFSSPRNRKFPDRK	None	None	None	None	None	None	None		

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