

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasm meml
M9PI47	M9PI47_DROME	Drosophila melanogaster	NaN	30.60235	T114;S131;T144;T177;T336;T344;T424;T425;T474;T740;T741;S836;T844	NaN	33925313;40245499	MSAPSEGGAGGGPVGGGSTATAPSA HPAVAAKITHVKSDGGGGISLACTPIL SGGTIKVASANQVQOTSLGGTPIALN AGQATTAASIRAIGAGGQSTQVRVV MPMIKQLENVATGRTQITAIPAAPA RSGANASITVTRPVTQATYLPASVT ATQMGVGVGVGQRLVTPLRATSSA TVTPTAPTIVLSSTGGFVRGATASVSR SGVALTSQPSSAVISSSNSGAWMQS QGQVQLIRTIHQPRQRIITTSAGVSN ASVVTTTPVQAPTALSVSAGOPLATQ QATGPGGGPQAYVATVLPQRHQHAT LVYSSNVSAPNASQNPQQQFNPRFA VATPTGGVTTTTPGGTIVTTPRQVRPI PLGKSFPTAKLNTTISISIRAPSIQNLN TSVTASPTAVSVSGGATVAPGRGPGA GGTALTATNLPTTRIIQLQQPATGTT QQIIGSGARLAGNVMLQPFLLMSTTA AAKMGRPPVTMTAKVQPSLTITQLN PIGKLSAGGGGGQTMGPQGVASIQ AVPVNVSASVVNSSSVTGATSAAAG GATVPLTISGRGAGVPGGNILTGT LTPIKNASGITLGKMMMAPASSAPG ADGSSTSGATVTFQRTWNPVVASQ SQLMKIDEKGSAAITYYESMPASAST GVLSTTTTTVTQSTQAAQQLVSSAGG SLSVSSLPFASHATAGAGSGGGGSO ATFTVLPAGAGGTARTTIGHQQLIIPA GPNSAPPQHMVPLHTSVKVTGAG NPQQATASGALVSSFMRKRDAEGSP IRAAKNLGPILLSMASNSGQNPMP PAPGTTFNIPVSVTVPSVSGSLTV EALAKKERERVTHGVASSAASVPAT VGPVNSTRNSRADSPASSDGSTTVS ANSSPGVDQQLQDRIGDQPATGGAG GRDNSTHFNPINENLGHSSLGARSG SSAFVDVQAPTQQQQPQVPHLVG STQMQQGTRMNGTGNSSSSSYDCS SRKKPRRSTNDSQHSTQSQASLSLP PPSSEPTPPAGASYMOKHNNGLLV TAATPGVAPNSAPGDANHRNSTPAV AGNKENANPVDFVIRRRNCALLNT YKPTHKLANNHFHRYTDVKPREERR ATVIDLANQPNVQKINGWKIYHLR SQMEDLNESEVFSLGRLETMLQDL EKDKEKHSEIERVTELLKGNIRSKII TDGINEAQNLMKIFEHKKPHVSNIIIT RCASKRNFKKREKI	NaN	NaN	NaN	NaN	NaN	NaN	NaN	

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