

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q867Z4	LOLA4_DROME	Drosophila melanogaster	Longitudinals lacking protein, isoforms F/I/K/T	25.496522	NaN	NaN	33925313	MDDDQFCLRWNNHQSTLISVFDL LLENETLVDCFLAAEGKFLKAHKVV LSACSPYFATLLQEYDKHPFILKD VKYQELRAMMDYMYRGEVNIQDQ LAALLKAAESLQIKGLSDNRTGGVA PKPESSGHHRRGGKLSGAYTLEQTKR ARLATGGAMDTSGDVSGSREGSSSP SRRRRKVVRRRSMENDAHDNSNSSV LQAAASNQSIQQGTGAGLAVSALVTT QLSSGPAAGTSSQASSTQQQPLTS TNVTKTESAKLTSSTAAPASGASAS AAVQQAHLHQQAQTSDAINTENV QAQSQGGAQGVQGDDEDIDEQSAV GGPNSATGPNPASASASAVHAGVVV KQLASVVDKSSSNHKKIKDINSVSS VGSEMVIKAEYDDDAHDENVEDL TLDEEDMTMEELDQTAGTSQGGEG SSQTYATWQHRSQDELGLMAQDA QQRDPQASKQDKGEQTEGAQDEFE LDDCLESNDIVITQNKDGFVLHVK KLGNTAAKLEENQAVAQQGQAAV TVTGPAGQPTPTITELLNAAAASHSE PKPTLTLTSTPIKLPSSCELINIKKI IPATTTIATHHPHTSSTIHPHHIIQH VSQEPHHEHHQHQHTIIEEVPQT SQQHHQQHHHQLQTVQPTHQV QSIHTAHPGQITINLVGLRNVQLADSK PIASRIRYRSGKIIGPTVQNLQIVETH EPIQHHEHLSDGTKEYEISEIDLNNP NASAAIISDLVKYAEIDDIELPDGTKI GIGFAPSEITEHMQTSGGETHITIE HEPQELQTVHQEQTQTHHHIAG QLQTHHIQTVVQSSSGQQHQDQQQ HHQHHSIELQDDDGVEITITPEELGM HDSKSYTILTRPMKEESEHDPSC MTYELSLSDSSLGPCDDPESRYVCR HCGKKYRWKSTLRRHENVECGGKE PCHPCPYCSYKAKQRGNLGVHVRK HHPEKPQLESKRGRKV	None	None	None	None	None	None	None		

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