

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
Q91T77	LAR4_DROME	Drosophila melanogaster	La-related protein Larp4B	25.496522	NaN	S1123;S1370;S1413	33925313	MEYFLGGYVMNGDALKIQPPVYTN VPVETAMLATNLFGATTQIAASPAAA AAAHPLITAAAAAAGAPAVAPP VAVAPVATASAAQAAVVPAAQQQQA HPHQAILAHTVVAHQQQQQQQQ TIQQLHQQQQQSPHPAQHLTYG HQPALSQATAGGSGTISGGAVGPSG VVVEGQDTNEYAIMNGALGQTQDG TVVCYTTDALNNTNLVALDPQPHQI VVPVQVPVQVPVQPLPANGSADPQ QGSNNAAGGEEPNIPLDKLQMLAT QLEYFYSRENLANDTYLLSQMDSQ YVPIYTVARFNLVRKLTNDINLITEVL RESPNVQVDDKGLRVRPNRKRCHIL REISNNTPLDDVKALFSNESCPRPIS CEFAANNSWYITFESDEDAQKAYKY LREEVKEFQGKPIMARIKPKSFINRI QAVPKNGYRLTSPPTANVFDPAGAG GATVSYAAAQPPRYLYTNGAAIAAPA SVQYSNPVLIPQNPQFYPLVAGPW PPGTVAATTAHQNFYELGGNIFTT NPLPPOPVTAGFAQAPPPTSQAMVT PKPQGGGRYNNHRGNPNNVGGGA NSGPRAESRSKPPRSTPSASHIQQG NIMPIQITAPIPGGMVSVVPTNIVQD PLQPAPLQQQQQVIQIQQQPSSSA QQQQAATVQMNNTRHYPIKSNWK GGMQKNLDKSYSGGVATHHHYTTQ VQALPAHSHHLQAQQQLGQGTQQQ HYQLASSAASAPQVTVTTAPAPQQ PGQHQHHLVVQQTQPQQQQQQQ VVLQQQVQVQELQEAQDGVESHSS HAMQQVNRSSNMSASSSSSLATSM SKEPLQWQNRPRRRRDEEGGINYS PGGRVGLYGSSPSNPHQQLMSSS TGSNVQSAGGTDGGGASHRGGGERQ SHYNTIHDVPPPQHRGNYKGNYN PHYHAQHSSMASGSHHHHHHNL SSQQQHHLLTTGTGQQGSGHHYHH HHHHNSIGSNVGNSSGLGVSSGGS GGGGSGGSGSNLSGGSNRERGIH HSSLGYQGHQHQHQQQQQQQQQ QQQQAAPVQPPQFDLEDAAFPPLP ATSATAPHTPQATGGASLHNSTTSS SSSTGLGQKQTLHQQQQAPQQHQ VLSSVESAGGDEQRSSNQGIETS NIHLANSSTANNWVENRLSDVVRT GGGGGGGKGRKARDNRHPQGQN QPHMAPNYQQHQRNPPVSPTPAL GAEYGIQIEGNNTKNVTQSSSNN NSIHVICKQTPNINASSKEEAAGAQQ QQQQQLDKSNKTEDEMPKQPSQQ RLVEGYVQQQLPLLAHNEYSAAAL AAGAGACSVVEGLTTSLAECSLGQ HKKPPSVAALHAKKDANLLEKGA HKSVAATSTSTENLSAAATASKLSYAQ VAQHKKAAASDSKETGSGGSGTTL SPTGSHKMDLVFGDPVVVAAVEKS GLATVSTEKRVAGGASPAALIGKPA GLPATNTTQGSAAVVSAAKEKDNRP NASVRPLSKEKQQQHYGSATEHNT NANILGASGNRRSRANNS	None	None	None	None	None	None	None		