

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q61033	LAP2A_MOUSE	Mus musculus	Lamina-associated polypeptide 2, isoforms alpha/zeta	44.030548	NaN	S59;S66;S67;T74;S79;S82;T153;S155;S158;T159;T163;S165;S167;S310;S349;S352;S368;S420;S422	21540332;34887587;22645316;24788674;25153642;28528544	MPEFLEDPSVLTKDKLSELVANNV TLPAGEQRKDVYVQLYLQHLTARNR PPLAAGANSKGGPPDFSSDEEREPTP VLGSGASVGRGRGAVGRKATKKTDK PRLEDKDDLVDVTELSNEELLQLVLR YGVNPGPIVGTTRKLYEKLLKLREQ GTESRSSTPLPTVSSSAENTRQNGS NDSDRYSDNDEGKKKEHKKVKSAR DCVPFSELASTPSGAFFQGISFPEIST RPPLGRTELQAARKVQTTKRDPPRE TCTDTALPGKGQTHKLAPGRSLFIPS ESSYDRCVEKSSPSSQREFAARLVLS AAASPLIRETTTTYSKDIVENICRGG KSRAQPLRAEEPVSQSVFSSERE VLOESERSQVISPLAQAIRDVVNSL LVQGGVGSPLPGTSDSVPTLDVENICK RLSQSSYQDSSELSPPRKVPRLSEKP ARGGDSGSCVAFQNTPGSEHRSSFA KSVVSHSLTTLGVEVSKPPPQHDKIE ASEPSFPLHESILKVVEEWEQQIDRQ LPSVACRYPVSSIEAARILSVPKVDDE ILGFISEATPRAATQASSTESC DKHL DLALCRSYEAAASALQIAAHTAFVAK SLQADISQAAQIINSDPSDAQALRIL NRTYDAASYLCDAADFVVRMSACA MGSSTMGRRYLWLKDCIKSPASKN KLTVPFPGGTLFGGEVHKVIKRG NKQ	True	False	2.133	5.0	1.312	1.298	0.767