	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membra
0	P60469	LIPA3_MOUSE	Mus musculus	Liprin-alpha-3	36.610221	S639	S17;S142;S207;S431;S508;S640;S645;S668;S683;T714;S737;T792;S794;S1123;S1164	34678516;35822049;22645316	MMCEVMPTISEDGRRGSALGPDEA GGELERLMVTMLTERERLLETLREA QDGLATAQLRLRELGHEKDSLQRQL SIALPQEFAALTKELNLCREQLLERE EEIAELKAERNNTRLLLEHLECLVSR HERSLRMTVVKRQAQSPGGVSSEVE VLKALKSLFEHHKALDEKVRERLRM ALERVAVLEEELELSNQEALNLRDQ LSRRRSGLEEPGKDGDGQTLANGLG PVGESNRRTAELEEALERQRAEVCQ LRERLAVLCRQMSQLEEELGTAHRE LGKAEEANSKLQRDLKEALAQRED MEERITTLEKRYLSAQREATSLHDA NDKLENELASKESLYRQSEEKSRQL AEWLDDAKQKLQQTLQKAETLPEIE AQLAQRVAALNKAEERHGNFEERL RQLEAQLEEKNQELQRARQREKMN DDHNKRLSETVDKLLSESNERLQLH LKERMGALEEKNSLSEEIANMKKLQ DELLLNKEQLLAEMERMQMEIDQL RGRPPSSYSRSLPGSALELRYSQAPT LPSGAPLDPYGAGSGRAGKRGRWSG AKDESSKDWDRSAPAGSIPPPFPGE LDGSDEEEAEGMFGAELLSPSGQAD VQTLAIMLQEQLEAINKEIKLIQEEK ETTEQRAEELESRVSSSGLDSLGRYR SSCSLPPSLTTSTLASPSPPSSGHSTP RLAPPSPAREGTDKTNHVSKEEAGV PRGEGPAVPGDTPPPTPRSARLERM AQALALQAGSPEDGAPPRGSESTPD SLHKAPKRKSIKSSIGRLFGKKEKGR MGPPGRESVSLAGTPSDETLATDPL GLAKLTGPGDKDRRNKRKHELLEEA CRQGLPFAAWDGPTVVSWLELWVG MPAWYVAACRANVKSGAIMANLSD TEIQREIGISNPLHRLKLRLAIQEMV SLTSPSAPASSRTPTGNVWMTHEE MESLTAATKPETKEISWEQILAYGD MNHEWVGNDWLPSLGLPQYRSYF MESLVDARMLDHLNKKELRGQLKM VDSFHRVSLHYGIMCLKRLNYDRKD LERRREESQTQIRDVMVWSNERVM GWVSGLGLKEFATNLTESGVHGALL ALDETFDYSDLALLLQIPTQNAQARQ LLEKEFSNLISLGTDRRLDEDSAKSF SRSPSWRKMFREKDLRGVTPDSAE MLPPNFRSAAAGALGSPGLPLRKLQ PEGQTSGSSRADGVSVRTYSC	None	None	None	None	None	None	None	None