	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrac region
0	O75335	LIPA4_HUMAN	Homo sapiens	Liprin-alpha-4	26.157505	T313;S331;T609	S640;S681	28657654;39531497;30379171	MCEVMPTINEGDRLGPPHGADADA NFEQLMVNMLDEREKLLESLRESQ ETLAATQSRLQDAIHERDQLQRHLN SALPQEFATLTRELSMCREQLLERE EEISELKAERNNTRLLLEHLECLVSR HERSLRMTVVKRQAQSPSGVSSEVE VLKALKSLFEHHKALDEKVRERLRA ALERVTTLEEQLAGAHQQVSALQQG AGVRDGAAEEEGTVELGPKRLWKED TGRVEELQELLEKQNFELSQARERL VTLTTTVTELEEDLGTARRDLIKSEE LSSKHQRDLREALAQKEDMEERITT LEKRYLAAQREATSIHDLNDKLENE LANKESLHRQCEEKARHLQELLEVA EQKLQQTMRKAETLPEVEAELAQRI AALTKAEERHGNIEEHLRQLEGQLE EKNQELARTAVQVRQREKMNEDHN KRLSDTVDRLLSESNERLQLHLKER MAALEEKNTLIQELESSQRQIEEQH HHKGRLSEEIEKLRQEVDQLKGRGG PFVDGVHSRSHMGSAADVRFSLGTT THAPPGVHRRYSALREESAKLALPLT VTLRSPTWMRMSQGVCCNLEYHSS GTLCGSSGPLPVPEMIQEEKESTELR AEEIETRVTSGSMEALNLKQLRKRG SIPTSLTALSLASASPPLSGRSTPKLT SRSAAQDLDRMGVMTLPSDLRKHR RKLLSPVSREENREDKATIKCETSPP SSPRTLRLEKLGHPALSQEEGKSALE DQGSNPSSSNSSQDSLHKGAKRKGI KSSIGRLFGKKEKGRLIQLSRDGATG HVLLTDSEFSMQEPMVPAKLGTQAE KDRRLKKKHQLLEDARRKGMPFAQ WDGPTVVSWLELWVGMPAWYVAA CRANVKSGAIMSALSDTEIQREIGIS NALHRLKLRLAIQEMVSLTSPSAPPT SRTSSGNVWVTHEEMETLETSTKTD SEEGSWAQTLAYGDMNHEWIGNE WLPSLGLPQYRSYFMECLVDARML DHLTKKDLRVHLKMVDSFHRTSLQ YGIMCLKRLNYDRKELEKRREESQH EIKDVLVWTNDQVVHWVQSIGLRDY AGNLHESGVHGALLALDENFDHNT LALILQIPTQNTQARQVMEREFNNL LALGTDRKLDDGDDKVFRRAPSWR KRFRPREHHGRGGMLSASAETLPAG FRVSTLGTLQPPPAPPKKIMPEAHSH YLYGHMLSAFRD	True	False	4.616	3.65	0.994	0.73	False	1.093	1.057