	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	extracellul region
0	G3X9J0	SI1L3_MOUSE	Mus musculus	Signal-induced proliferation-associated 1-like protein 3	25.161187	NaN	S94;S140;S394;S1358;T1381;S1538;S1541;S1614;S1617;S1672;T1694;T1698	34887587	MTTYRPLPNDGVDLAASCGARSTDI LPGPHPGDYTPMGFWAQNGSMPQP LGESPAATTTRPSPTTPAMPKMGVR ARVADWPPKRDALREQSNPSPSQDT DGVKTTKVAHSMRNLQNGQLPSST PASSGSRAFHRLSRRRSKDVEFQDG WPRSPGRAFLPLRHRSSSEITLSECD VEEPGDPRGTRHPGVLPLFREYGST SSIDVQGVPEQSFFDILNEFRSEQPE ARGSQNLRELLQVDPGALSGGSCGT KGDPRNGQPTKDSLQSLQPLKEKEK SRKKPVRGLGSGDTVDSSIFRKLRSS KPEGEVGRPLGETEESRSPPEASRP WVCQKSFAHFDVQSMLFDLNEAAA NRVSVAQRRNTTTGASAASAASAMV TLTASRAHSLGTLDPAFTSTEDLNCK ENLEQDLGDDNSNDLLLSCPHFRN EIGGERERNVSFSRASVGSPGGSSE AHMAEPTLSTHRTNASISVLEVPKE QQRTQSRPRQYSIEHVDLGARYYQD YFVGKEHANYFGVDEKLGPVAVSIK REKLEDHKDHGPQYQYRIIFRTRELI TLRGSILEDATPTATKHGTGRGLPLK DALEYVIPELNIHCLRLALNTPKVTE QLLKLDEQGLCRKHKVGILYCKAGQ SSEEEMYNNEEAGPAFEEFLDLLGD KVCLKGFTKYAAQLDVKTDSTGTHS LYTTYQDYEIMFHVSTLLPYTPNNRQ QLLRKRHIGNDIVTIIFQEPGALPFTP KNIRSHFQHVFIIVRVHNPCTENVCY SMAVTRSKDAPPFGPPIPNGTTFRKS DVFRDFLLAKVINAENAAHKSDKFH TMATRTRQEYLKDLAENCVSNTPID SSGKFNLISLTSKKKEKTKARAGAEQ HSAGAIAWRVAAQDYAQGSEIDCILG ISNEFVVLLDLRTKEVVFNCYCGDVI GWTPDSSTIKIFYGRGDHIFLQAAEG SVEDIRDIVQRLKVMTNGWETVDMT LRRNGLGQLGFHVKYDGTVAEVEDY GFAWQAGLRQGSRLVEICKVAVVTL SHDQMIDLLRTSVTVKVVIIPPFEDG TPRRGWPETYDMNASEPKTESETTT PGGRPPYRSNAPWQWSGPASHNSL PATKWTTPATPGHAQSLSRLPKQTP VVPFRESQPLHSKRPVSFPETPFAAS PAGADRVPPYRQPSGSFSTPGSATYA RYKPSPERYAAAPHPLLSFDPHFMH DGMSSGDSSSGGLTSQESTMERPK PEPLWHVPAQARLSAMTGSIGSKHP SRQDAAGKDSPNRHSKGEPQYSSHS SSNTLSSNASSSHSDDRWFDPLDPL EPEQDPFSKGGSSDSGIDTTLYTSSP SCMSLAKAPRPTKPHKPPGNIGLCG GGRESAGRPHPVDRRREVSPAPVVA GQNKGYRPKLYSSGSCTPPGLVGGS RDPPRQPSDMGSRAGYPTQVYKTAS AETPRPSQLSQCSPFQLSTSVPKSFF SKQPAHNKHSTGWKRTDEPPPRPLP FTDSKKQVDTNAKNVFGQPRLRASL RDLRSPRKNYKSTIEDDLKKLIVMD NLGPEQERDTGQSPQKSLQRTLSDE SLCSGRREPSFASPASLEPGLPSDVL FTSTCTFPSSTLPARRQHQHAHPPS GAPSTTPATGNGFPEKKSAISASELS LADGRDRPLRRLDPGMMPLPDTAA GLEWSSLVNAAKAYEVQRAVSLFSL NDPALSPEIPPAHSPVHSHLSLERGP QTPRATPTMSEESPLDLTGKVYQLE VMLKQLHTDLQKEKQDKVVLQSEV ASLRQNNQRLQEESQAASEQLRKFA ELFSREKKEL	None	None	None	None	None	None	None	None	None