

UniprotKB ID	Entry name	organism	full name	oglcnaacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O43166	SI1L1_HUMAN	Homo sapiens	Signal-induced proliferation-associated 1-like protein 1	13.195361	S59;S1114;T1135;S1178;T1408;S1424	S162;S187;S193;S208;S255;S288;S1078;S1087;S1116;S1127;S1149;S1170;S1181;S1255;S1270;S1326;T1330;S1349;S1366;S1390;S1391;S1412;S1431;S1433;S1528;S1549;T1551;S1554;S1565;S1568;S1585;S1588;S1603;S1645;S1647;S1650;S1708;S1711;S1728;S1729;S1734	30379171;31492838;35254053;28657654;35132862	MTSLKRSQTERPLATDRASVVGTDG TPKVHTDDDFYMRFRSQNSGLGSSV MAPVGPPESEGGSHHITSTPGVPMG VRARIADWPPRKENIKESRSSSQEIE TSSCLDSLSSKSSPVSQSSVLSNSN DSAMLKSIQNTLKNKTRPSENMS RFLMPEAYPSSPRKALRRIRQRSNS DITISELDVDSFDECISPTYKTGPSLH REYGSTSSIDKQGTSGESFFDLLKGY KDDKSDRGPTPTKLSDFLITGGGKG SGFSLDVIDGPISQRENLRLFKEREK PLKRRSKSETGDSIFRKLRLNAKGE LGKSSDLEDNRSEDSVRPWTCPKCF AHYDVQSILFDLNEAIMNRHNVIKR RNTTTGASAAVASLVSGPLSHSASF SSPMGSTEDLNSKGSLSMDQGDDK SNELVMSCPYPFRNEIGGEGGERKISLS KNSGSGFSGCESASFESTLSSHCTN AGVAVLEVPKENLVLHLDRVKRYIVE HVDLGAYYRKFYQKEHWNFYGA DENLGPVAVSIRREKPDENKENGSP YNYRIIFRTSELMTLRGSVLEDAIPST AKHSTARGLPLKEVLEHVPELVNQ CLRLAFNTPKVTEQLMKLDEQGLNY QOKVIMYCKAGQSTEEEMYNNE AGPAFEFLQLLGERVRLKGFYK AQLDTKTSTGTHSLYTTYKDYEIF HVSTMLPYTPNKKOQLLRKRHIGN DIVTIVFQEPGAQPFSPKNIRSHFQ VFVIVRVHNPSCDSVCYSVAVTRSRD VPSFGPPIPKGVTFPKSNVFRDFLLA KVINAENA AHKSEKFRAMATRTRQE YKDLAEKNVTNTPIDPSGKFPFISL ASKKKEKSKPYPGAELSSMGAIWWA VRAEDYNKAMELDCLLGISNEFIVLI EQETKSVVFNCSRDVIGWTSTDTS LKIFYERGECVSVGSFINIEEIKEIVK RLQFVSKGCESEVMTLRRNLGLQL GFHVNYEGIVADVEPYGYAWQAGLR QGSRLVEICKVAVATLSHEQMIDLLR TSVTVKVVIIPPHDDCTPRRSCSEY RMPVMEYKMNIEGVSVEYFKFPFRNN NKWQRNASKGPHSPQVPSQVQSPM TSRLNAGKGDGKMPPERAANIPRS ISSDGRPLERLSPGSDIYVTVSSMA LARSQRNSPNSLSSSDTGSVGGT YRQKSMPEGFGVSRSPASIDRQNT QSDIGSGKSTPSWQSEDSIADQM AYSYRGPQDFNSFVLEQHEYTEPTC HLPVSKVLPARESPSGRLMRQDP VVHLSPNKQGHSDSHYSSSSSNTL SSNASSAHSEKWDGDRTESELN SYNYLQGTADSGIDTTSYGPSHGST ASLGAATSSPRSGPGEKVAPLWHS SSEVISMADRTLETESHGLDRKTESS LSLDIHSKQAGSTPLTRENSTFSIN DAASHTSTMSSRHSASPVVFTSARS SPKEELHPAAPSQ LAPSFSSSSSSS GPRSFYPRQATSKYLIGWKKPEGTI NSVGFMDTRKRHQSDGNEIAHTRL RASTRDLRASP KPTSSTIEEDLKKLI DLESPTPESQKSFKHALSSPQSPFP STPTSRRALHRTLSDESIYNSQREHF

FTSRASLLDQALPNDVLFSSSTYPSLP  
KSLPLRRPSYTLGMKSLHGFSASD  
SSLTDIQETRRQMPDPGLMPLPDT  
AADLDWSNLVDAAKAYEVQRASFFA  
ASDENHRPLSAASNSDQLEDQALAQ  
MKPYSSSKDSSPTLASKVDQLEGML  
KMLREDLKKEKEDKAHLQAEVQHL  
REDNLRQLQEESQNASDKLKKFTEW  
VFNTIDMS