	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparat
0	Q9UPQ0-3	LIMC1_HUMAN	Homo sapiens	Isoform 3 of LIM and calponin homology domains-containing protein 1	25.756906	S437 (Q9UPQ0-3)	S169;S177;S192;S201;S204;S207;T215;S217;S226;S231;S233;S297;S303;T313;S377;T402;S471;T472;S516;S518;S523;T529;T537;S601;S670;S681;S718;T724;S750;S756;S875;T900;S966;S973	30379171;31492838;28657654;28411811;31373491;35254053;29485866;38253038;29237092;37217939;28510447	MRKDTDDIESPKRSIRDSGYIDCWD SERSDSLSPPRHGRDDSFDSLDSFG SRSRQTPSPDVVLRGSSDGRGSDSE SDLPHRKLPDVKKDDMSARRTSHG EPKSAVPFNQYLPNKSNQTAYVPAP LRKKKAEREEYRKSWSTATSPLGGE RPFSFPETIEEEGSEVGSAGEDNPAG QMNPGWKPSDGGCELPDGSGKEHP SSDGAVVAPAPKSEEKDAAEIQKRER LEQAGIKVMPAAQRFASQKQLSEEK EAIRDIVLRKENSFLTHQHGNDSEA EGEVVCRLPDLEKDDFAARRARMN QTKPMVPLNQLLYGPYPKKGAEKSD GSKQLSKGISKKRSLEYKRNQGHTE EVKLIVTCNMRAQESEPVEGGLRKV PDLHKDDLAQQRIQGSLAPHREPPS FITLSNITEADLETWERLKVSEKARD GDVQHICASEPSPEIKAETAIRDDFA NRKARASKKASSPRQKFVHFGPVTE LDQQKWKRLSIGKAGPREDEEEVIC HGSKIQMDSVSPVSAATSSLKGHQI FNRQNDCRTMNCGRGDYCRRASW LAPVPESQEEWVCSLGECPRGTEEV TSKQLPQDGKEETESAPRDSERLSK AERSEDSSQPLVCPLASECEASGTEE KLEKMTAPAWSGSGLKGQRKLDDS RKDDMMARRTGMSLRHTGSNPNQ FLPVPFAKQQDVEESSKGLPMKDQR YGPRTPVSDDAESTSMFDMRCEEEA AVQPHSRARQEQLQLINNQLREEDD KWQDDLARWKSRRRSVSQDLIKKE EERKKMEKLLAGEDGTSERRKSIKT YREIVQEKERRERELHEAYKNARSQ EEAEGILQQYIERFTISEAVLERLEM PKILERSHSTEPNLSSFLNDPNPMK YLRQQSLPPPKFTATVETTIARASVL DTSMSAGSGSPSKTVTPKAVPMLTP KPYSQPKNSQDVLKTFKVDGKVSVN GETVHREEEKERECPTVAPAHSLTK SQMFEGVARVHGSPLELKQDNGSIE INIKKPNSVPQELAATTEKTEPNSQE DKNDGGKSRKGNIELASSEPQHFTT TVTRCSPTVAFVEFPSSPQLKNDVSE EKDQKKPENEMSGKVELVLSQKVV KPKSPEPEATLTFPFLDKMPEANQL HLPNLNSQVDSPSSEKSPVMTPFKF WAWDPEEEERRRQEKWQQEQERLL QERYQKEQDKLKEEWEKAQKEVEE EERRYYEEERKIIEDTVVPFTVSSSSA DQLSTSSSMTEGSGTMNKIDLGNC QDEKQDRRWKKSFQGDDSDLLLKT RESDRLEEKGSLTEGALAHSGNPVS KGVHEDHQLDTEAGAPHCGTNPQL AQDPSQNQQTSNPTHSSEDVKPKTL PLDKSINHQIESPSERRKKSPREHFQ AGPFSPCSPTPPGQSPNRSISGKKLC SSCGLPLGKGAAMIIETLNLYFHIQC FRCGICKGQLGDAVSGTDVRIRNGL LNCNDCYMRSRSAGQPTTL	None	None	None	None	None	None	None