

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P02468	LAMC1_MOUSE	Mus musculus	Laminin subunit gamma-1	18.754982	T1216;T1380	S1147;S1491	29187734;36852467;33300544;36288343;30059200	MTGGGRAALALQPRGRLWPLLAVLA AVAGCVRAAMDECADEGGRPQRCM PEFVNAAFNVTVATNTTCGTPPEEY CVQTGVTGVTKSCHLCDAGQQHLQ HGA AFLTDYNNQADTTWWQSQTM LAGVQYPNSINLTLHLGKAFDITYVR LKFHTSRPESFAIYKRTREDGPWIPY QYYSGSCENTYSKANRGFIRTGGDE QQALCTDEFSDISPLTGGNVAFSTLE GRPSAYNFDNSPVLQEWVTATDIRV TLNRLNTFGDEVFNPKVLKSYYYAI SDFAVGGRCCKNGHASECVKNEFD KLMCNCKHNTYGVDCEKCLPFFND RPWRRATAESASESLPCDCNGRSQE CYFDPELYRSTGHGGHCTNCRDNT DGAKCERCENFFRLGNTEACSPC HCSPVGSLSLSTQCDYGRCSCKPGVM GDKCDRCQPGFHSLEAGCRPCSC DLRGSTDECNVETGRCVCKDNVEG FNCERCKPGFFNLESSNPKGCTPCF CFGHSSVCTNAVGYVYDISSTFQID EDGWRVEQRDGSEASLEWSSDRQD IAVISDSYFPRYFIAPVKFLGNQVLSY GQNLSFSFRVDRRDTRLSAEDLVLE GAGLRVSVPLIAQNSYPSETTVKYI FRLHEATDYPWRPALSPFEFQKLLN NLTSIKIRGTYSERTAGYLDVTLQS ARPGPGVPATWVESCTCPVGYGGQF CETCLPGYRRETPSLGPYSPCVLCTC NGHSETCDPETGVCD CRDNTAGPH CEKCS DGYGDSTLGTSSDCQPCPC PGGSSCAIVPKTKEVVCTHCPTGTAG KRCELCD DGYFGDPLGSNGPVRLCR PCQCNDNIDPNAVGN CNRLTGECL KCIYNTAGFYCDRCKEGFFGNPLAP NPADKCKACACNPYGT VQQSSCN PVTGQCQCLPHVSGRDCGTCDPGY NLQSGQCERCDC HALGSTNGQCD IRTGQCECQPGITGQH CERCETNHF GFGPEGCKPCDCHHEGSLSLQCKD DGRCECREGFVGNRC DQCEENYFY NRSWPGCQEC PACYRLVKDKAAEH RVKLQELES LIANLGTGDDMVT DQA FEDRLKEAEREVTDLLREAEVKDV DQNLMDRLQRVNSSLHSQISRLQNI RNTIEETGILAEARSRVESTEQLIEI ASRELEKAKMAANVSITQPESTGE PNNMTLLAEEARLAERHKQEADDI VRVAKTANETS AEAYNLLLRLTAGE

