

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8VHX6	FLNC_MOUSE	Mus musculus	Filamin-C	31.717039	S2647	S5;S1162;S1339;S2043;S2234;S2237;T2239;S2587;S2618;S2621;S2633;S2715;S2719	24788674;22645316;37507081;25153642;36288343;27301331;34418053	MMNNSNYSDASGLGLVDEADEMPS TEKDLAEDAPWKKIQNTFTRWCN EHLKCVGKRLTDLQRDLSGLRLIA LLEVLSSQKRMFKFHRPNFRQMK LENVSVALEFLEREHIKLVSIDSKAIV DGNLKLILGLIWLTLIHYSISMPMWE DEDEDARKQTPKQRLLGWIQNKV PQLPITNFNRDWQDGKALGALVDN CAPGLCPDWEAWDPNQPVQNAREA MQQADDWLGVPQVIAPEEIVDPNVD EHSVMTYLSQFPKAKLKPGAPVRSK QLNPKKAIAYGPIEPQGNTVLQPAH FTVQTVDAGVGEVLVYIEDPEGHTE EAKVVPNNDKDRTYAVSYVPKVAGL HKVTVLFAGQNIERSPFEVNVGMAL GDANKVSARGPGLPVGNVANKPTY FDIYTAGAGTGDVAVVIVDPQGRRT VEVALEDKGDNTFRCTYRPVMEGP HTVHVAFAGAPITRSPFPVHVAEAC NPNACRASGRGLQPKGVRVKEVADF KVFTKGAGSGELKVTVKGPKGTEEP VKVREAGDGVFECEYYPVPGKYVV TITWGGYAI PRSPFEVQVSPEAGA QK VRAWGPGLETGQVGKSADFVVEAIG TEVGTGFGSIEGPSQAKIECDDKGDG SCDVRYWPTEPGEYAVHVICDDEDI RDSPFIAHIQPAPPDCFPDKVKAFGP GLEPTGCIVDRPAEFTIDARAAGKGD LKLYAQDADGCPIDIKVIPNGDGTFR CSYVPTKPIKHTIIVSWG GVNVPKSP FRVNVGEGSHPERVKVYGPVVEKTG LKANEPYFTVDCSEAGQGDV SIGIK CAPGVVGPVEADIDFDIHKNDNDTFT VKYTPPGAGHYTIMVLFANQEIPASP FHIKVDPSHDASKVKAEGPGLSRTG VEVGKPTHFTVLTGAGKAKLDVHF AGAAKGEAVRDFEIDNHDYSYTVKY TAVQQGNMAVTVTYGGDPVPKSPFV VNVAPPLDLSKVKVQGLNSKVAVGQ EQAFSVNTRGAGGQGLDVRMTSP SRRPIPCKLEPGGGAEAQAVRYMPP EEGPYKVDITYDGHPVPGSPFAVEGV LPPDPSKVCAYGPKLGGGLVGT PAFP SIDTKGAGTGGLGLTVEGPCEAKIEC QDNGDGSCAVSYLPTEPGEYTINILF AEAHIPGSPFKATI QPVFDPSKVRAS GGLER GKAGEAATFTVDCSEAGEA ELTIEILSDAGVKAEVLIQNNADGTY HITYSPAFPGTYTITIKYGGHPKFPPT RVHVQPAVDTS GIKVSGPGVEPHGV LREVTTEFTVDARSLTATGGNHVTA RVLNPSGAKTDTYVTDNGDGT YRVQ YTAYEEGVHLVEVLYDEVAVPKSPFR

VGVTEGCDPTRVRAFGPGLEGGLVN
KANRFTVETRGAGTGGLGLAIEGPS
EAKMSCKDNKDGSCCTVEYIPFTPGD
YDVNITFGGQPIPGSPFRVPVKDVVD
PGKVKCSGPGGLGTGVRARVPQTFTV
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VRDNGDGTHTVHYTPATDGPYTVAV
KYADQEVPRSPFKIKVLP SHDASKVR
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GQGLLTVQILDPEGKPKKANIRDNG
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PNSPFHVLACDPLPHVEEPAEMLQ
MRQPYAPLRPGTCPTHWATEEPVVP
VEPLESMLRPFNLVIPFTVQKGELTG
EVRMPSGKTARP NITDNKDG TITVR
YAPTEKGLHQMGIKYDGNHIPGSPL
QFYVDAINSRHVSAYGPGLSHGMVN
KPATFTIVTKDAGEGGLSLAVEGPSK
AEITCKDNKDG TCTVSYLPTAPGDYS
IIVRFDDKHIPGSPFTAKITGDDSMR
TSQLNVGTSTDVSLKITEGDLSQLTA
SIRAPSGNEEPCLLKRLPNRHIGISF
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ILVGPSEIGDASKVRVWGKGLSEGQT
FQVAEFIVDTRNAGYGGGLSIEGPS
KVDINCEMEDGTCKVTYCPTPEPGT
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MKESITRRRQAPSIATIGSTCDLNLKI
PGNWFQMVSAQERL TRTFTRSSHT
YTRTERTEISKTRGGETKREVRVEES
TQVGGDPFPAVFGDFLGRERLGSFG
SITRQQEGEASSQDMTAQVTSPSGK
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HTVTVKYRGQHVP GSPFQFTVGPLG
EGGAHKVRAGGTGLERGVAGVPAEF
SIWTREAGAGGLSIAVEGPSKAEIAF
EDRKDGSCGVS YVVQEPGDYEVS IK
FNDEHIPDSPFVVPV ASLSDDARRLT
VTSLQETGLKVNQPASFAVQLNGAR
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GLEGGTTGVSSEFIVNTQNAGSGAL
SVTIDGPSKVQLDCRECPEGHVVITYT
PMAPGNYLIAIKYGGPQHIVGSPFKA
KVTGPRLSGGHS LHETSTVLVETVT
KSSSSRGASYSSIPKFSSDASKVVTR
GPGLSQAFV GQKNSFTVDCSKAGTN
MMMVG VHGPKTPCEEVYVKHMG N
RVYNVITYTVKEKGDYILIVKWGDES V
PGSPFKVNVP