

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q14315	FLNC_HUMAN	Homo sapiens	Filamin-C	14.578219	S2490	S5;S1161;S1338;S2042;S2233;S2236;T2238;S2586;S2617;S2620;S2623;S2632;S2714;S2718	23301498;30379171;16408927;29351928;33214551;34725712;34229054;34019948	MMNSNGYS DAGLGLGDETDEMPST EKDLAEDAPWKKIQNTTFTRWCNE HLKCVGKRLTDLQRDLS DGLRLIAL EVL SQKRM YRKFHPRPNFRQMKLE NVSVALEFLEREHIKLV SIDSKAIVD GNLKLILGLIWTLILHYSISMPMWED EDDEDARKQTPKORLLGWIQNKVP QLPITNFNRDWQDGKALGALVDNC APGLCPDWEAWDPNQPVENAREAM QQADDWLGVPQVIAPEEIVDPNVDE HSVMTYLSQFPKAKLKPGAPVRSKQ LNP KKA IAYGPGIEPQGNTVLQPAHF TVQTV DAGVGEVLVYIEDPEGHTEE AKVVPNNDKDRTYAVSYV PKVAGLH KVTVLFAGQNIERSPFEVNVGMALG DANKVSARGPGLPVGNVANKPTYF DIYTAGAGTGDVAVVIVDPQGRRTV EVALEDKGDSTFRCTYRPAMEGPHT VHVAFAGAPITRSPFPVHVSEACNP NACRASGRGLQPKGVRVKEVADFKV FTKGAGSGELKVTVKGPKGTEEPVK VREAGDGVFECEYYPVVPK YVVTIT WGGYAI PRSPFEVQVSPEAGVQKVR AWGPGLETQV GKSADFVVEAIGTE VGT L GFSIEGPSQAKIECDDKGDGSC DVR YWPTEPGEYAVHVICDDEDIRD SPFIAHILPAPPDCFPDKVKAFGPGL EPTGCIVDKPAEFTIDARAAGKGLK LYAQDADGCPIDIKVIPNGDGTFRCS YVPTKPIKHTIIISWGGVNVPKSPFRV NVGEGSHPERVKVYGPVEKTGLKA NEPTYFTVDCSEAGQGDSIGIKCAP GVVGP AEADIDFDIKN DNDTFTVKY TPPGAGRYTIMVLFANQEIPASPFHI KVDPSHDASKVKAEGPGLNRTGVEV GKPTHFTVLT KGAGKAKLDVQFAGT AKGEVVRDFEIIDNHDYSYTVKYTAV QQGNMAVTVYGGDPV PKSPFVVN VAPPLDLSKIKVQGLNSKVAVGQEQ AFSVNTRGAGGQGLDVRMTSPSR RPIPCKLEPGGAEAQAVRYMPPEE GPYKVDITYDGHPVPGSPFAVEGVL PDPSKVCAYGPGLKGGLVGT PAFPSI DTKGAGTGGLTVEGPCEAKIECQ DNGDGSCAVSYLPTEPGEYTINILFA EAHIPGSPFKATIRPVFDPSKVRASG PGLERGVGEAATFTVDCSEAGEAE LTIEILSDAGVKA EVL IHNADGTYH ITYSPA FPGTYTITIKYGGHPVPKFPT RVHVQPAVDTS GVKVSGPGVEPHGV LREVTTEFTVDARSLTATGGNHVTA RVLNPSGAKTDTYVTDNGDGT YRVQ YTAYEEGVHLVEVLYDEVAVPKSPFR

VGVTEGCDPTRVRAFGPGLEGGLVN
KANRFTVETRGAGTGGLGLAIEGPS
EAKMSCKDNKDGSCCTVEYIPFTPGD
YDVNITFGGRPIPGSPFRVPVKDVVD
PGKVKCSGPGLGAGVRARVPQTFTV
DCSQAGRAPLQVAVLGPTGVAEPVE
VRDNGDGTHTVHYTPATDGPYTVAV
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GGLLTQILDPEGKPKKANIRDNG
DGTYYTVSYLPDMSGRYTITIKYGGDE
IPYSPFRIHALPTGDASKCLVTVSIGG
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DGTFDIYYTAPEPGKYVITIRFGGEHI
PNSPFHVLACDPLPHEEEEPSEVPQL
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TSQLNVGTSTDVSLKITESDLSQLTA
SIRAPSGNEEPCLLKRLPNRHIGISF
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ILVGPSEIGDASKVRVWGKGLSEGH
TFQVAEFIVDTRNAGYGGLGLSIEGP
SKVDINCEMEDGTCKVYCPTEPG
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ARGVIDARVHTPSGAVEECYVSELDS
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GLEGGTTGVSSEFIVNTLNAGSGALS
VTIDGPSKVQLDCRECPGHHVVTYTP
MAPGNLYLAIKYGGPQHIVGSPFKAK
VTGPRLSGHSLHETSTVLVETVTKS
SSSRGSSYSSIPKFSSDASKVVTRGP
GLSQAFVVGQNSFTVDCSKAGTNM
MMVGVHGPKTPCEEVYVKHMGNR
VYNVTYTVKEKGDYILIVKWGDESVP
GSPFKVKVP