

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q80X90	FLNB_MOUSE	Mus musculus	Filamin-B	25.527009	S2103	T216;T519;S730;S886;S932;S983;S1028;T1307;S1316;S1433;S1505;S1602;S2083;S2113;S2369;S2465;S2478;S2481;S2492	24788674;34887587;22645316;34418053;25153642	MPVTEKDLAEDAPWKKIQNTFTRWCNEHLKCVNKRIGNLQTDLSDGLRLIALLEVLVSQKRMHHKHQRPTFRQMKLENVSVALEFLDHESIKLVSIDSKAIVDGNLKLILGLVWTLILHYSISMVWEDEGEDDDAKKQTPKQRLLGWIQNKIPYLPITNFNQNWQDGKALGALVDSAPGLCPDWESWDPRKPDNAREAMQQADDWLGVPQVITPEEIIHPDVDEHSVMTYLSQFPKAKLKPGAPLKPCLNPKKARAYGRGIEPTGNMVKQPAKFTVDTISAGQGDVMVFVEDPEGNKEEARVTPDSDKNKTYVVEYLPKV TGLHKVIVLFAQGHISKSPFEVNVDKAQGDASKVTAKGPGLETTGNIANKPTYFDIYTAGAGVGDIGIEVEDPQGKNSVELLEDVRGNQVYRCVYKPVQPGPHVVKVSFAGDAIPKSPFGVQIGEACNPNACRASGRGLQPKGVRIRETADFKVDTKAAGSGELGVTVKGPKGLEELVKQKGFLDGVYSFEYYPSTPGKYSVAVTWGGHHIPKSPFEVQVQVPEAGMOKVRAWGPGLHGGIVGRSADFFVESIGSEVGTGFAIEGPSQAKIEYDDQNDGSCDVKYWPKEPGEYAVHIMCDEEDIKDSPYMAFIHPATGDYNPDLVQAYGPGLEKSGCTINNPAEFIVDPKDAGSA PLKILAQDGEQPIDIQMKSMDGT YACSYTPLKAIKHTIAVVWGGVNIPHSPYRVNIGQGSHQPQKVKVFGPGVER SGLKANEPHFTVDCTEAGEGDVSVGIKCDARVLSDDDEEDVDFDIIHNAN DTFTVKYVPPAPGRYTIKVLFAEQEIP ASPFRVKVDPSHDASKVKAEGPGLS KAGVENGKPTHFTVHTKGAGKAPLN VQFSSPLPGEAVKDLDIIDNYDYSHT VKYTPTQQGNMQVLVITYGGDPIPKS PFTVGVAAPLDLSKIKINGLENRVEV GKDQEFADTNGAGGQKLDVTILS PSRKVVPCLVAPVAGRECSTAKFIPR EEGLFAVDVTYDGHVPVPGSPYTVEA SLPPDPTKVKAHGPGLEGGLVGKPA EFTIDTKGAGTGGLGLTVEGPCEAKI ECSDNGDGTCSVSYLPTKPGYFVNI LFEEVHIPGSPFKADIEPFDPKVV ASGPGLHGVGEPGILCVDCSEAG PGTLGLEAVSDSGAKAEVSIQNNKD GTYAVTYVPLTAGMYTLTMKYGGEL VPHFFAWVKVEPAIDTSGIKAFGPGI EGKDVFREATTDFTVDSRPLTQVGG DHIKAQITNPSGASTEFCVKNADG TYQVEYTPFEKGFHVVEVYDDVPIP NSPFKVAVTEGCQPSRVHAQGPGLK

EAFTNKS NVFTV VTRGAGIGGLGITV
EGPSESKINCRDNKDGSCSAEYIPFA
PGDYDVNITYGGVHIPGSPFRVPSKD
VVDPSKVKIAGPGLSSCVRACIPQSF
TVDSSKAGLAPLEVRVLGPRGLVEPV
NVVDNGDGTHTVTYTPSQEGPYIVS
VKYADEEIPRSPFKVKVLPYDASKV
TASGPGLSAYGVPASLPVEFAIDARD
AGEGLLAVQITDQEGKQRATVHDN
KDGTAVTYIPDKTGRYMIGVTYGGD
NIPLSPYRIRATQTGDASKCLATGPGI
APT VKTGEEVGFVVD AKTAGKGKVT
CVILTPDGTEAEADVIENEDGTYDIF
YTAAKPGTYVIYVRFGGVDIPNSPFT
VMATDGEVTAMEEAPVNACPPGFR
PWVTEEAYVPVSDMNGLGFKPFDLV
IPFAVRKGEITGTVHMPSGKKATPEI
VDNKDGTVTVRYAPTEVGLHEMHIK
YRGSHIPESPLQFYVNPNSGVSAY
GPGLVYGVANKTATFTIVTEDAGEG
GLDLAIEGPSKAEISCIDNKDGTCTV
TYLPTLPGDYSILVKYNDKHIPGSPFT
AKITDDNRRCSQVKLGSAADFLLDIS
ETDLSTLTASIKAPSGRDEPCLLKRL
PNNHIGISFIPREVGEHLVSIKKNGN
HVANSPVSIMVVQSEIGDARRAKVY
GQGLEGRTFEMSDFIVDTRDAGYG
GISLAVEGPSKVDIQTEDLEDGTCKV
SYFPTVPGVYIVSTKFADEHVPGSPF
TVKISGEGRVRESITRTSRAPAVATV
GSICDLNLKIPEINSSDMSAHVTSFS
GHVTEAEIVPMGKNSHCVRFPQE
MGVHTVSVKYRGQHVTGSPFQFTV
GPLGEGGAHKVRAGGPGLERGEAGI
PAEFSIWTREAGAGGLSIAVEGPSKA
EITFDDHKNGSCGVSYIAQEPGNYE
VSIKFNDEHIPDSPYLVPVIAPSDDAR
CLTVLSLQESGLKVNQPASFAIRLNG
AKGKIDAKVHSPSGAVEECHVSELE
PDKYAVRFIPHENGIHTIDVKFNESH
VVGSPFKVRVGEPPQAGNPALVSAY
GAGLETGTTGIQSEFFINTTQAGPGT
LSVTIEGPSKVKMDCQEIPEGYKVM
YTPMAPGNYLIGVKYGGPNHISRSPF
KAKVTGQRLVSPGSANETSSILVESV
TRSSTETCYS AIPKSSSDASKVTSKG
AGLSKAFVVGQSSFLVDCSKAGSNM
LLIGVHGPTTPCEEVSMKHVKGQQY
NVTYVVKERGDYVLAVKWGEEHIPG
SPFHVTVP