

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O75369	FLNB_HUMAN	Homo sapiens	Filamin-B	12.430888	T401;S1422;T1593;S1602;S2478	T216;T519;S730;S886;S932;S983;S1028;T1307;S1316;S1433;S1505;S1602;S2083;S2107;S2113;S2369;S2465;S2478;S2481;S2492;S1474	32119511;29351928;34725712;34019948;35254053;30397120;35083852;23301498;30444036;33214551;35008409;30379171;26853435;34229054	MPVTEKDLAEDAPWKKIQNTFTFRWCNEHLKCVNKRIGNLQTDLS DGLRLIALLEVLVSQKRMRYKYHQRP TFRQMQLENVSVALEFLDRESIKLVSIDS KAIVDGNLKLILGLVWTLILHYSISM PVWEDEGDDDAKKQTPKQRL LGWIQNKIPYLPITNFNQNWQDGKALGAL VDSCAPGLCPDWESWDPQKPVDNA REAMQQADDWLGVPQVITPEEIIHP DVDEHSMVTYLSQFPKAKLKPGAPL KPCLNPKKARAYGRGIEPTGNMVKQ PAKFTVDTISAGQGDMVFVEDPEG NKEEAQVTPDSDKNKTY SVEYLPKV TGLHKVTVLFAGQHISKSPFEVSVDK AQGDASKVTAKGPGLEAVGNIANKP TYFDIYTAGAGVGDIGVEVEDPQGKN TVELLVEDKGNQVYRCVYKPMQPGP HVVKIFFAGDTIPKSPFVQVGEACN PNACRASGRGLQPKGV RIRETTDFK VDTKAAGSGELGVTMKGPKGLEELV KQKDFLDGVYAFEYYPSTPGRYSIAIT WGGHHIPKSPFEVQVGEAGMQKV RAWPGPLHGGIVGRSADFVVESIGS EVGSLGFAIEGPSQAKIEYNDQNDG SCDVKYWPKEPGEYAVHIMCDDEDI KDSPYMAFIHPATGGYNPDLV RAYGPGLEKSGCIVNNLAEFTVDPKDAGK APLKIFAQDGEQQRIDIQMKNRMDG TYACSYTPVKAIKHTIAVWGGVNIP HSPYRVNIGQGSHPQKV VFGPGVE RSGLKANEPHFTVDCTEAGEGDVS VGIKCDARV LSEDEEDVDFDIIHNAN DTFTVKYVPPAAGRYTIKVL FASQEIP ASPFRVKVDP SHDASKVKAEGPGLS KAGVENKPTHTFTVYTKGAGKAPLN VQFN SPLPGDAVKDLDIIDNYDYSHT VKYTPTQQGNMQVLV TYGGDPIPKS PFTVGVAAPLDLSKIKLNGLENRVEV GKDQEFTVDTRGAGGQKLDVTILS PSRKVVPCLVTPVTGRENSTAKFIPR EEGLYAVDV TYDGHVPVPGSPYV EA SLPPDPSKVKAHGPGLGGLVGKPA EFTIDTKGAGTGGLGLTVEGPCEAKI ECSDNNGDGTCSVSYLPTKPG EYFVNI LFEEVHIPGSPFKADIEMPFDP SKVV ASGPGL EHGKVG EAGLLSVDCSEAG PGALGLEAVSDSGTKA EVSIQNNKD GTYAVTYVPLTAGMYT LTMKYGGEL VPHFPARVKVEPAVDTSRIK VFGPGI EGKDVFREATTDFTVDSRPLTQVGG DHIKAHIANPSGASTE CFVTDNADG TYQVEYTPFEKGLHVVEV TYDDVPIP NSPFKVAVTEGCQPSRVQAQGPGLK EAFTNKNPVFTVVTRGAGIGGLGITV EGPSESKINCRDNKDGSCSAEYIPFA PGDYDVNITYGGAHIPGSPFRVPVKD VVDPSKVKIAGPGLSGV RARVLQSF TVDSSKAGLAPLEVRVLGPRGLVEPV NVVDNGDGTHVTYTPSQEGPYMV SVKYADEEIPRSPFKVKVLP TYDASK

VTASGPGGLSSYGVPASLPVDFDAIDAR  
DAGEGLLAVQITDQEGKPKRAIVHD  
NKDGTAVTYIPDKTGRYMIGVTYGG  
DDIPLSPYRIRATQTDGASKCLATGP  
GIASVTKTGEEVGFVVDAKTAGKKG  
VTCTVLTPDGTEAEADVIENEDGTYD  
IFYTAAKPGTYVIYVRFGGVDIPNSPF  
TVMATDGEVTAVEEAPVNACPPGFR  
PWVTEEAYVPVSDMNLGFKPFDLV  
IPFAVRKGEITGEVHMPSGKTATPEI  
VDNKDGTVTTRYAPTEVGLHEMHIK  
YMGSHIPESPLQFYVNYPNSSGSVSA  
YGPGLVYGVANKTATFTTIVTEDAGEG  
GLDLAIEGPSKAEISCIDNKDGTCTV  
TYLPTLPGDYSILVKYNDKHIPGSPFT  
AKITDDSRRC SQVKLGSAADFLDIS  
ETDLSSLTASIKAPSGRDEPCLLRL  
PNNHIGISFIPREVGEHLVSIKKNGN  
HVANSPVSIMVVQSEIGDARRAKVY  
GRGLSEGRTFEMSDFIVDTRDAGYG  
GISLAVEGPSKVDIQTEDLEDGTCKV  
SYFPTVPGVYIVSTKFADEHVPGSPF  
TVKISGEGRVKESITRTRAPSVATV  
GSICDLNLKIPEINSSDMSAHVTSPS  
GRVTEAEIVPMGKNSHCVRFPQE  
MGVHTVSVKYRGQHVGTGSPFQFTV  
GPLGEGGAHKVRAGGPGLERGEAG  
VPAEFSIWTREAGAGGLSIAVEGPSK  
AEITFDDHKNGSCGVSYIAQEPGNY  
EVSIFFNDEHIPESPYLVPVIAPSDDA  
RRLTVMSLQESGLKVNQPASFAIRL  
NGAKGKIDAKVHSPSGAVEECHVSE  
LEPDKYAVRFIPHENGVTIDVKFN  
GSHVVGSPFKVRVGEPPGQAGNPALV  
SAYGTGLEGGTTGIQSEFFINTTRAG  
PGTLSVTIEGPSKVKMDCQETPEGY  
KVMYTPMAPGNYLISVKYGGPNHIV  
GSPFKAKVTGQRLVSPGSANETSSIL  
VESVTRSSTETCYS AIPKASSDASKV  
TSKGAGLSKAFV GQKSSFLVDCSKA  
GSNMLLIGVHGPTTPCEEVSMKHV  
GNQQYNVTYVVKERGDYVLAVKWG  
EEHIPGSPFHVTVP