

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8K4G5	ABLM1_MOUSE	Mus musculus	Actin-binding LIM protein 1	40.481242	S496;S499	S216;S411;T417;T440;S466;S470;S475;T477;S479;T483;S496;S499;S502;S582;S671;S723;S738;S760;S789	19458039;22517741;35822049;21540332;22645316;29187734	MPSLLGLKCLGKLCSSSEIGKVPSPER ASLRNSHRLLIEDLSVPETPDPAHR RRGTVIHLVLYSAGCGPPELRFSSY DPSVAHPQDPHHSSEKPIVHCHKCG EPCKGEVLRVQTKHFHFKCFTCKVC GCDLAQGGFFIKNGDYLDYQRM YGTRCHGCGEFVEGEVVTALGKTYH PNCFACTICKRPFPPGDRVTFNGRD CLCQLCAQPMSSSPKEASCSSNCAG CGRDIKNGQALLALDKQWHLGCFK CKSCGKVLGTGEYISKDGSPLYCEKDYQ GLFGVKCEACHQFITGKVL EAGDKH YHPSCARCSRCNQMFTEGEEMYLQ GSTVWHPDCKQSTKTEEKL RPPNIP RSSSDFFYPKSLIRRTGRSPALQLLS PPCLTNSNKNPRQPTRTSSESIYSRP GSSIPGSPGHTIYAKVDNEILDYKDL AAIPKVKAIYDIERPDLITYEPFYTSGY EDKQERQSLGESPRTLSPTPSAEGY QDVRDRMIHRSTSQGSINSPVYSRH SYTPTRSRSPOHFHRPELLSPGVHR WSPLRTSSFSTHSDSRPNPPFRHH FLPHVKGNPSSGRNSPLPYRPDSR PLTPTYAQAPKHFHVPDQGINIYRKP PIYKQHAALAAQSKASEDIKFSKFPA AQAPDPNEIPKIETDHWPGPPSLAAV GTDPRRRSSGREDEEELLRRRQLQ EEQLMKLN SGLGQLILKEEMEKESR ERASLASRYDSPLHSASHAPSSKTSS LPGYGKNGLHRPVSTDF AQYNSYGD ISGGVRDYQTL PDGHMPAVRMDRG VSMPNMLEPKIFPYEMLMVTNRGR NKILRDVDRTRLERHLAPEVFWEIF GMSIQEFDKLP LWRNDM KKKAKL F

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