

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
014639	ABLM1_HUMAN	Homo sapiens	Actin-binding LIM protein 1	36.848719	S352;S358;S362;S363;T371;T429;S450;S452;S455;S462;S465;S471;S651;S677;S682	S216;S367;T373;T396;S422;S426;S431;T433;S435;T439;S452;S455;S458;S498;S587;S640;S655;S677;S706	20068230;30059200;31492838;28657654;29351928;29485866;28580691;29237092;30620550	MPAFLGLKCLGKLCSSSEKSKVTSSE RTSARGSNRKRLLIVEDRRVSGTSFTA HRRATITHLLYLCPKDYCPGRVCN SVDPFVAHPQDPHHPSEKPVHCHK CGEPCCKGEVLRVQTKHFHIKFTCK VCGCDLAQGGFFIKNGEYLCTLDYQ RMYGTRCHGCGEFVEGEVVTALGK TYHPNCFACTICKRPFPPGDRVTFN GRDCLCQLCAQPMSSSKETTFSSN CAGCGRDIKNGQALLALDKQWHLG CFKCKSCGKVLGTGEYISKDGAPYCEK DYQGLFGVKCEACHQFITGKVLEAG DKHYHPSCARCSRCNQMFTEGEEM YLQGSTVWHPDCKQSTKTEEKL RPT RTSSESIYSRPGSSIPGSPGHTIYAKV DNEILDYKD LAAIPKVKAIYDIERPDL ITYEPFYTSGYDDKQERQSLGESPR LSPTPSAEGYQDVRDRMIHRSTSQC SINSPVYSRHSYPTTTSRSPQHFHRP GNEPSSGRNSPLPYRPSRPLTPTYA QAPKHFHVPDQGINIYRKPIYKQHA ALAAQSKSSEDIKFSKFPAAPDP SETPKIE TDHWPGPPSFAVVGPD RRSSGREEDDEELLRRRQLQEEQL MKLNSGLGQLILKEEMEKSRRSS LLASRYDSPINSASHIPSSKTASLPGY GRNGLHRPVSTDF AQYNSYGDVSG GVRDYQTL PDGHMPAMRMDRGVS MPNMLEPKIFPYEMLMVTNRGRNK ILREVDRTRLERHLAPEV FREIFGMS IQEFDRLPLWRRNDM KKKAKLF