

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
014639	ABLM1_HUMAN	Homo sapiens	Actin-binding LIM protein 1	46.493258	T351;S352;S358;S362;S363;T371;T429;S450;T451;S452;S455;S458;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	28580691;35132862;35254053;29237092;20068230;30059200;35289036;37340703;29485866;30620550;31492838;29351928;28657654	MPAFLGLKCLGKLCSSSEKSKVTSSE RTSARGSNRKRLLIVEDRRVSGTSFTA HRRATITHLLYLCPKDYCPRGRVCN SVDPFVAHPQDPHHPSEKPVHCHK CGEPCCKGEVLRVQTKHFHIKFTCK VCGCDLAQGGFFIKNGEYLCTLDYQ RMYGTRCHGCGEFVEGEVVTALGK TYHPNCFACTICKRPFPPGDRVTFN GRDCLCQLCAQPMSSSKETTFSSN CAGCGRDIKNGQALLALDKQWHLG CFKCKSCGKVLGTGEYISKDGAPYCEK DYQGLFGVKCEACHQFITGKVLEAG DKHYHPSCARCSRCNQMFTEGEEM YLQGSTVWHPDCKQSTKTEEKL RPT RTSSESIYSRPGSSIPGSPGHTIYAKV DNEILDYKD LAAIPKVKAIYDIERPDL ITYEPFYTSGYDDKQERQSLGESPR LSPTPSAEGYQDVRDRMIHRSTSQC SINSPVYSRHSYPTTTSRSPQHFHRP GNEPSSGRNSPLPYRPDSRPLTPTYA QAPKHFHVPDQGGINIYRKPIYKQHA ALAAQSKSSEDIKFSKFPAAPDP SETPKIE TDHWPGPPFAVVGPD RRSSGREEDDEELLRRRQLQEEQL MKLNSGLGQLILKEEMEKSRRSS LLASRYDSPINSASHIPSSKTASLPGY GRNGLHRPVSTDFAQNSYGDVSG GVRDYQTLPDGHMPAMRMDRGVS MPNMLEPKIFPYEMLMVTNRGRNK ILREVDRTRLERHLAPEVFREIFGMS IQEFDRLPLWRRNDMKKKAKLF