

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9Z2H5	E41L1_MOUSE	Mus musculus	Band 4.1-like protein 1	33.918701	S721;T791;S792;T793;T800;T803	T30;S75;T79;T343;S378;S430;S437;S461;S466;T475;S510;S540;S541;S544;S546;T550;S564;S578;S639;S648;S650;S665;S666;S671;S677;S684;T685;S721;S782;S868	22645316;22517741;36852467;33300544;35822049;30059200;34678516	MTTETGPDSEVKKAQEETPQQPEAA AAVTTTPVTPAGHSHPETNSNEKHLT QQDTRPAEQSLDMDKDYSEADGL SERITTPSKAQKSPQKIAKKFKSAICR VTLLDASEYECEVEKHGRGQVLFDL VCEHLNLLKDYFGLTFCDADSQKN WLDPSKEIKKQIRSSPWNFAFTVKFY PPDPAQLTEDITRYYLCLQLRADIITG RLPCSFVTHALLGSYAVQAEFGDYD AEEHVGNYVSELRFAPNQTRELEER IMELHKTYRGMTPGEAEIHFLNAK KLSMYGVDLHHAKDSEGIDIMLGVC ANGLLIYRDRLRINRFAWPKILKISYK RSNFYIKIRPGEYEQFESTIGFKLPN HRSKRLWKVCIEHHTFFRLVSPPEP PPKGFLVMGSKFRYSGRTQAQTRQA SALIDRPAPFFERSSSKRYTMSRSLD GAEFSRPASVSENHDAGPEGDKRED DAESGRRSEAEEGEVRTPTKIKELK PEQETTPRHKQEFLDKPEDVLLKHQ ASINELKRTLKEPNSKLIHRDRDWD RERRLPSSPASPSKGTPEKASERAG LREGSEEKVKPPRPRAPESDMGDED QDQERDAVFLKDNHLAIERKCSSITV SSTSSLEAEVDFTVIGDYHGGAFEDF SRSLPELDRDKSDSETGLVFAQDL KGPSSQEDES GGLEDSPDRGACSTP EMPQFESVKAETMTVSSLAIRKKIEP EAMLQSRVSAADSTQVDGGTPMVK DFMTTPPCITTETITTTMENSLSKSGK GAAAMIPGPQTVATEIRSLSPIIGKDV LTSTYGATAETLSTTTTHVTKTVKG GFSETRIEKRIITGDEDVDQDQALAL AIKEAKLQHPDMLVTKAVVYRETDP SPEERDKKPQES

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