

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion
Q9H4G0	E41L1_HUMAN	Homo sapiens	Band 4.1-like protein 1	35.126883	S523;T697;T699;T775;T778;T793;S794;T799;T802	T30;S75;T79;T343;S378;S430;S437;S461;S466;T475;S510;S540;S541;S544;S546;T550;S564;S578;T580;S639;S648;S650;S667;S672;S678;S685;T686;S722;S784;S870	29351928;30379171;39534244;28657654;31492838;35132862;29237092;30059200;35254053;32574038	MTTETGPDSEVKKAQEEAPQQPEAA AAVTTPTVTPAGHGHPEANSNEKHPS QQDTRPAEQSLDMEEKDYSEADGLS ERTTPSKAQKSPQKIAKKYKSAICRV TLLDASEYECEVEKHHGRGQVLFDLV CEHLNLLLEKDYFGLTFCDADSQKN WLDPSKEIKKQIRSSPWNFAFTVKFY PPDPAQLTEDI TRY YLCLQLRADIITG RLPCSFVTHALLGSYAVQAE LGDYD AEEHVGNYSELRFAPNQTRELEER IMELHKTYRGMTPGEAEIHFL ENAK KLSMYGVDLHHAKDSEGIDIMLGVC ANGLLIYRDRLRINRF AWP KILKISYK RSNFYKIRPGEYE QFESTIGFKLPN HRS AKRLWKVCIEHHTFFRLVSPEP PPKGFLVMGSKFRYSGRTOAQTRQA SALIDRPAPFFERSSSKRYTMSRSLD GAEFSRPASVSENHDAGPDGDKRD EDGESGGQRSEAE EGEV RPTKIKE LKPEQETTPRHKQEF LDKPEDVLLK HQASINELKRTLKEPNSKLIHRDRD WERERRLPSSPASPSPKGTPEKANE RAGLREGSEEKVKPPRPRAPESDTG DEDQDQERD TVFLKDNHLAIERKCS SITVSSTSSLEAEVDFTVIGDYHGSAF EDFSRSLPELDRDKSDSDTEGLLFS RDLNKGAPSQDDESGGIEDSPDRGA CSTPDMPOFEPVKTETMTVSSLAIR KKIEPEAVLQTRVSA MDNTQQVDGS ASVGREFIATTPSITTETISTTMENSL KSGKGAAMIPGPQIVATEIRSLSP II GKDVLTSTYGATAETLSTSTTTHVTK TVKGGFSETRIEKRIITGDEDVDQD QALALAIKEAKLQHPDMLVTKAVVY RETDPSPEERDKKPQES	True	False	4.651	2.425	0.868