

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrac region
Q9D219	BCL9_MOUSE	Mus musculus	B-cell CLL/lymphoma 9 protein	24.992371	T315;S318	S104;S157;T172;T315;S318;S352;S686;S688;S906;S916	36852467	MHPSNPKVRRSSPSGNTQSSPKSKQE VMVRRPTVMSPSGNPQLDSKFSNQ GKPGGSASQSQSPSCDSKSGGHTPK ALPGPGGSMGLKNGAGNGAKGKGGK RERSISADSFQDRDPGTPNDDSDIKE CNSADHIKSQESQHTPHSMTPTSTAT APRSSTPSHGQTPAPEPIAQAQKTPAK VVYVFSTEMANKAAEAVALKGOVETI VSFHIQNISNSKSERSTAPLNTQIPT LRNDPKPLPQQPPAPANQDNSSQ NARLQPTPIQAPAPKPTAAPRPLDR ESPGVENKLIIPPVSGSGSSTPLPPDG TGPNSTPNRAVTPVVSQGSNSSSAD PKAPPPPPVSGGEPPTLGENPDGLS QEQLHRERSLQTLRDIQRMFLFDE KEFTAGQTGGPQONTGVLDGPQKKP DGPIQAMMSQSQSLGKGGPRTDV GAPFGPQGHRDVPFSPDEMVPNNM SSQSGPIGPDHLDHMTPEQIAWLK QQEFYEEKRRKQEQVVVQCSLQD MMVHQHGRGPRGVVGGPPPYQMAPG EGWAPGAEPFDPDGINISHSLPPRGM APHPNMPGSQMRLPGFAGMINSEM EGPNVNPASRPGLSGVSWPDDVPK IPDGRNFPPGGQVFSGPRGRERFPN PQGLSEEMFQQQLAEQLALPPGM SMEGIRPGMEMNRMIPGSQRHMEP GSNPIFPRIPVEGPLSPSRGDFPKGM PPQIGPRELEFGMVPGGMKGEVN LNVNMGSSSQMIPQKMREAGAGPE EMMKLRPGSSEMLPAQQKMVPLPF GEHPQQEYGVGPRPFLPMSQGPGS NSGLRNLREPIGPDORTNSRLSHMP PLPLNPSNPTSLSTAPPVORGLGRK PLDISVAGSQVHSPGINPLKSPTMH QVQSPMLGSPSGNLKSPQTPSQLAG MLAGPAAAASIKSPVVLGSAASPVH LKSPLPAPSPGWTSPPKPLQSPGI PPNHKAPLTMASAPMLGSVESGGPP PPTASQPAVNIPGSLPSTPYPMPP EPTLSQNPLSIMMSRMSKFAMPSST PLYHDAIKTVASSDDSPPARSPNLP SMNSMPGMGINTQNPRISSGPNPVV PMPTLSMGMGTQPLSHSNQMPSPN AMGSPIPPVPMGPMGLMSHNPIM GHGSQEPPMVPGGRMGFPQGFPPV QSPQVPPPHNGPTGGQGNFPGGI GFPGEPLGRPSNLPQSSADPALCK PGGPGAPDSFTVLGNSMPSVFTDPD LQEVIRPGATGIFEFDLSRIIPSEKPS QTLQYFPRGEVPRGRKQPGQGGPFS HMQGMMSDQAPRMGLALPMGGGP GPVGTDPDPLGTSPSPMGPHNPMRPP AFLQQGMMGPHHRMMSPAQSTVP GPATLMTNPAAAVGMIPGKDRGPAG LYTHPGPVGSPGMMMSMQGMMGP QQNIMIPPQMRPRGMAADVGMGGP SQGPGNPGNMMF	True	True	2.227	5.0	0.689	0.577	5.0	1.291	1.059