

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P22682	CBL_MOUSE	Mus musculus	E3 ubiquitin-protein ligase CBL	15.818779	T511	T369;S437;S450;S481;S617;S640;S666;S667;T672;S692;T698;T737;T780;S907	36852467	MAGNVKKSSGAGGGGSGGSGAGGL IGLMKDAFQPHHHHHHLSPHPPCT VDKMMVEKCKWKLMDKVVRLCQNP KLALKNSPPYILDLLPDTYQHLRTVL SRYEGKMETLGENEYFRVFMENLM KTKQTISLFKEGKERMYEENSQPR RNLTKLSLIFSHMLAELKGIFPSGLF QGDTFRITKADAAEFWRKAFGEKTIV PWKSFRQALHEVHPISSGLEAMALK STIDLTCNDYISVFEFDIFTRLFQPW SLLRNWNSLAVTHPGYMAFLTYDE VKARLQKFIHKPGSYIFRLSCTRLGQ WAIGYVTADGNILQTIPHNKPLFQAL IDGFREGFYLFDPGRNQNPDLTGLC EPTPQDHIKVTQEYELYCEMGSTF QLCKICAENDKDVKIEPCGHLMCTS CLTSWQESEGQGCPCRCCEIKGTEPI VVDPDFPRGSGSLLRQGAEGAPSPN YDDDDDERADDSLFMMKELAGAKV ERPSSPFSMAPQASLPPVPPRLDLLQ QRAPVPASTSVLGTASKAASGSLHK DKPLPIPPTLRDLPPPPPPDRPYSVGA ETRPQRRPLPCTPGDCPSRDKLPPVP SSRPGDSWLSRPIKVPVATPNPGDP WNGRELTNRHSLPFSLPSQMEPRA DVPRLGSTFSLDTSMTMNSSPVAGP ESEHPKIKPSSANAIYSLAARPLPM PKLPPGEQGESEEDTEYMTPTSRPV GVQKPEPKRPLEATQSSRACDCDQQ IDSCTYEAMYNIQSQALSVAENSASG EGNLATAHTSTGPEESENEDDGYDV PKPPVPAVLARRTLSDISNASSSFGW LSLDGDPTNFNEGSQVPERPPKFPF RRINSERKASSYQQGGGATANPVAT APSPQLSSEIERLMSQGYSDIQKA LVIAHNNIEMAKNILREFVSISSPAH VAT