

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P22681	CBL_HUMAN	Homo sapiens	E3 ubiquitin-protein ligase CBL	32.901925	T245;S589;S601;S604;S605;S606;T610;S626	T371;S439;S452;S483;S619;S642;S668;S669;T674;T700;T731;T774;S900	3626323;30379171;33214551;30059200;31492838;28657654;32574038;29351928;31373491;28580691;29237092;34019948;30620550;23301498	MAGNVKKSSGAGGGSGSGGSGSGG LIGLMKDAFQPHHHHHHHHLSPHPP GTVDKKMVEKCKWKLMDKVVRLCQ NPKLALKNSPPYILDLLPDTYQHLRT ILSRYEGKMETLGENEYFRVFMENL MKKTKQTISLKFKEGKERMYEENSQP RRNLTKLSLIFSHMLAELKGIFPSGL FQGDTRITKADAAEFWRKAFGEKTI VPWKSFRQALHEVHPISSGLEAMAL KSTIDLTCNDYISVFEFDIFTRLFQP WSSLLRNWNSLAVTHPGYMAFLTY DEVKARLQKFIHKPGSYIFRLSCTRL GQWAIGYVTADGNILQTI PHNKPLF QALIDGFREGFYLFPPDGRNQNPDLT GLCEPTQDHIKVTQEYELYCEMG STFQLCKICAENDKDVKIEPCGHLM CTSCLTSWQESEGGQCPFCRCEIKG TEPIVDPFDPGSGSLLRQGAEGAP SPNYDDDDDERADDTL FMMKELAG AKVERPPSPFSMAPQASLPPVPPRLD LLPQRVCVPSSASALGTASKAASGSL HKDKPLVPPTLRDLPPPPPPDRPYS VGAESRPQRRPLCTPGDCPSRDKL PPVPSSRLGDSWLPRPIPKVPVSAPS SSDPWTGREL TNRHSLPFSLPSQME PRPDVPRLGSTFSLDTSMSMNSSPL VGPECDHPKIKPSSANAIYSLAARP LPVVKLPPGEQCEGEDTEYMT PSS RPLRPLDTSQSSRACDCDQID SCTY EAMYNIQSQAPSITESSTFGEGNLAA AHANTGPEESENEDDGYDVPKPPVP AVLARRTLSDISNASSFGWLSLDG DPTTNVTEGSQVPERPPKPPRRINS ERKAGSCQQSGPAASAATASPQLS SEIENLMSQGYSYQDIQKALVIAQN NIEMAKNILREFVSISSPAHVAT

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