

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q5H9F3	BCORL_HUMAN	Homo sapiens	BCL-6 corepressor-like protein 1	12.136309	S127;S135;T641;T652;T824;S834;S855	S496;S599;S613;S1029;S1033;S1162;S1476	30379171;29237092;35254053;29351928;33214551;30059200;35289036;34019948;30620550	MISTAPLYSGVHNWTTSSDRIRMCGI NEERRAPLSDEESTTGDQCQHFQSGE FCVSSSFQKVELTAVGSGSNARGAD PDGSATEKLGHKSEDKPDDPQPKM DYAGNVAEAEGLLVPLSSPGDGLKL PASDSA EASNSRADCSWTPLNTQM SKQVDCSPAGVKALDSRQGVGEKNT FILATLGTGVPVEGTLPLVTTNFSPLP APICPPAPGASVPHSVPDFQVPLS VPAPVPHSGLVPVQVATSVPAPSPPL APVPALAPAPPSVPTLISDSNPLSVSA SVLVVPASAPPSPGVPLSAPAPAPL SVPVSAAPLALIQA PVPSPAPTLLVAP VPTPVLAMPASTPPAAPAPPSVPMMP TPTPSSGPPSTPTLIPAFAPTPVPAPT PAPIFTPAPTPMPAATPAAIPTSAPIP ASFSLSRVCFPAAQAPAMQKVPLSF QPGTVLTPSQPLVYIPPPSCGQPLSV ATLPTLGVSTLTL PVLPSYLQDRC LPGVLASPELRSYPYAFSVARPLTSD SKLVSLVNRPLPCTSPSGSTTTQPAP DGVPGPLADTSLVTASAKVLPTPQPL LPAPSGSSAPHPAKMPSGTEQQTE GTSVTFSPKSPQLEREMASPPPCS EMPLDLSSKSNRQKLPNQRKTPP MPVLT PVHTSSKALLSTVLSRSQRTT QAAGGNVTSCLGSTSSPFVIFPEIVR NGDPSTWVKNSTALISTIPGTIVGVA NPVPASLLLKNKDPNLGLNRDPRHLP KQEPISIIDQGEKGTGATCGKKGSQ AGAEGQPSTVKRYTPARIAPGLPGCQ TKELSLWKPTGPANIYPRCSVNGKP TSTQVLPVGWSPYHQASLLSIGISSA GQLTPSQGAPIRPTSVVSEFSGVPSL SSSEAVHGLPEGQPRPGGSFVPEQD PVTKNKTCRIAAPYEEQVNPVLLTL SPQTGTLALSVPQSGGDIRMNOGPE ESEHLCSDSTPKMEGPPQACGLKL AGDTKPKNQVLATYMSHELVLATPQ NLPKMPPELPLPHDSHPKELILDVVP SSRRGSSTERPQLGSQVDLGRVKME KVDGDVVFNLATCFRADGLPVAPQR GQAEVRAKAGQARVKQESVGVFACK NKWQPDDVTESLPPKKMKCGKEKD SEEQQLQPQAKAVRSSHRRPKCRKL PSDPQESTKKS PRGASDSGKEHNGV RGKHKHRKPTKPEQSPPGKRADSHE EGSLEKKAKSSFRDFIPVVLSTRTRS QSGSICSSFAGMADSDMGSQEVFPT EEEEVTPTPAKRRKVRKTQRDTQY RSHHAQDKSLLSQGRRHLWRAREM PWRTAARQMWDTNEEEEEEEEEEG LLKRKRKRKRQSRKYQTGEYLTEQE DEQRRKGRADLKARKQKTSSQSLE HRLRNRNLLLPNKVQGISDSPNGFL PNNLEEPALENSEKPSGKRKCKTK HMATVSEEAKGKGRWSQOKTRSPK SPTPVKPTPECTPSKSRASSEEASE SPTARQIPPEARRLIVNKNAGETLLQ RAARLGKDVVLYCLOKQDSEDVNH DNAGYTALHEACSRGWTDILNILLE HGANVNC SAQDGRFPVHDAVVNDN LETIWLLLSYGADPTLATYSQGTAMK

LASSDTMKRFLSDHLSDLQGRAEGD
PGVSWDFYSSSVLEEKDGFACDLLH
NPPGSSDQEGDDPMEEDDFMFELS
DKPLLPCYNLQVSVSRGPCNWFIFS
DVLKRLKSSRIFQARFPHFEITTMP
KAEFYRQVASSQLLTPAERPGLDD
RSPPGSSETVELVRYEPDLLRLLGSE
VEFQSCNS