

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P48634	PRC2A_HUMAN	Homo sapiens	Protein PRC2A	17.889641	S1282;S1619;T1786;S1788;T1790;S1796;T2093	S18;S19;S30;S146;S166;S204;S342;S350;S363;S380;S383;S456;T610;S759;S761;S764;T807;S808;S908;S932;T997;S1004;T1083;S1085;S1089;S1092;T1094;S1106;S1110;S1120;S1147;S1219;S1306;S1310;T1323;S1325;S1328;T1347;T1353;S1384;S1386;S1525;S1778;S2036;T2076;S2082;S2113	23301498;30379171;35254053;28657654;20305658;33214551;32119511;34019948;30620550	MSDRSGPTAKGKDGGKYSSLNLFDTYKGSLEIQKPAVAPRHGLQSLGKVAIARRMPPANLPSLKAENKGNPNVSLVPKDGTGWASKQEQSDPKSSDASTAQPPESQPLPASQTPASNQPKRPPA APENTPLVPSGVKSWAQASVTHGAHGDGGRASSLLSRFSREEFPTLQAAGDQDKAAKERESAEQSSGPGPSLRPQ NSTTWRDGGGRGPDELEGPDSKLGHHGHDPRGGLQPSGPPQFPYRGMMPFFMYPPYLPFPPPYGPGQGPYRYPTPDGPSRFPRVAGPRGSGPPMRLVEPVGRPSILKEDNLKEFDQLDQENDDG WAGAHEEVVDYTEKLKFSDEEDGRDSDEEGAEGRHDSQSASGEERPPEADGKKGNPNSEPTPKTAWAETSRPPE TEPGPPAPKPLPPPHRGPAGNWP PPGDYPPDRGGPPCKPPAPEDEDEAWRQRRKQSSSEISLAVERARRRREEEERRMQEERRAACAEKLRLEKFGAPDKRLKAEPAPPAAPSTPAPPPAVPK ELPAPPAPPASAPTPETEPEEPAQA PPAQSTPTPGVAAAPTLVSGGGSTSS TSSGSFEASVPEQLPSKEGPEPPEE VPPPTTPVPKVEPKGDGIGPTRQPP SQGLGYPKYQKSLPPRFQRQQEQL LKQQQQHQWQQHQGSAPPTPVPP SPPQPVTLGAVPAPQAPPPPKALYP GALGRPPMPPMNFDPWRMMIPPY VDPRLQGRPPLDFYPPGVHPSGLV PRERSDSGSSSEPFDRHAPAMLRE RGTPVDPKLAWVGDVFTATPAEPR PLTSLRQAADDDKGMRETPPVPP PPPPYLASYPGFPENGAPGPPISRFP EEPGRPLPWPPGSDEVAKIQTTPPK KEPPKEETAQLTGPEAGRKPARGVG SGGQGP PPRRESRTETRWGPRPGS SRRGIPPEEPGAPRRRAGPIKKPPPT KVEELPPKPLEQGDETPKPPKPDPLK ITKGLGGPKETPPNGNLSPAPRLRR DYSYERVGPTSCRGRGRGEYFARGR GFRGTYGGRGRGARSREFRSYREFR GDDGRGGGTGGPNHPPAPRGRTAS ETRSEGSEYEEIPKRRRQRGSETGSE THESDLAPSDKEAFTPKEGTLTQVPL APPPPGAPSPAPARFTARGGRVFT RGVPSRRGRGGGRPPPQVCPGWSP AKSLAPKKPPTGPLPSPKEPLKEKLIP GPLSPVARGGSNGGSNVGMEDGER PRRRRHGRAQQQDKPPFRRLKQE RENAARGSEGKPSLTPASAPGPEE ALTTVTVAPAPRRAAAKSPDLSNQ SDQANEETWETASESSDFTSERRGDK EAPPVLLTPKAVGTPGGGGGGGAVP GISAMSRGDLSSQRAKDLKRSFSSQ RPGMERQNRPPGPGGKAGSSGSSS GGGGGGPGGRTGPRGDKRSWPSP KNRSRPPPEERPPGLPLPPPPSSSAV FRLDQVIHSNPAGIQQALQLSSRQ GSVTAPGGHPRHKPGLPQAPQGPSP RPPTRYEPQRVNSGLSSDPHFEEPG PMVRGVGGTPRDSAGVSPFPKRRR

