

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q7TSC1	PRC2A_MOUSE	Mus musculus	Protein PRC2A	19.954247	S1575;S1787;S1795	S18;S19;S30;S146;S166;S204;S342;S350;S363;S378;S381;S454;T609;S759;S761;S764;T807;S808;S931;T995;S1002;T1081;S1083;S1087;S1090;T1092;S1104;S1108;S1118;S1145;S1217;S1302;S1306;T1319;S1321;S1324;T1343;T1349;S1380;S1382;T1401;S1521;S1757;S2037;T2077;S2114	34678516;29187734;36852467	MSDRSGPTAKGKDGKKYSSLNLFDTYKGSLEIQKPAVAPRHGLQSLGKVAIARRMPPPPANLPSLKAENKGNDPNVSLVPKDGTGWASKQEQSDPKSSDASTAQPPESQLPASQTPASNQPKRPPTAPENTPESVPSGVKSWAQAVTHGAHGDGGRASNLLSRFSREFFTLQAAGDQDKAAKERESAEQSSGPGPSLRPQNSTTWRDGGGRGPDDLEGPDSKLGHHGDPRGGLQPSGPPQFPYRGMMPFFMYPPYLPFPPPYGPQGPYRYPTPDGPSRFPRVAGPRGSGPPMRLVEPVGRPSILKEDNLKEFDQLDQENDDGWAGAHEEVDYTEKLKFSDEEDGRDSDEEGAEGHKDSQSAAAEPEPDTGDKGTSPGSELPPPKTAWTENARPSETEPAPPTPKPPPPPHRGPVGNWGGPPGDYPDRGGPPCKPPAPEDEDEAWRQRRKQSSSEISLAVERRRRREEEERRMQEERRAACAEKLRLEKDFGAPDKRLKAEPAAAPPVTPAAPALPPVVPKEIPAAPALPPTPTTPEKEPEEPAQAPPVQAAPSPGVAPVPTLVSGGGCTANSNSSGSFEASVPEQLPSKEGPEPPEVPPPTTPAPKMEPKGDGVGSTRQPPSQGLGYPKYQKSLPPRFQRQQQEQLLKQQQQQQWQQQQGTAPPAPVPPSPPQPVTLGAVPAPQAPPPPKALYPGALGRPPMPPMNFDPRWMMIPPYVDPRLQGRPPLDFYPPGVHPSGLVPRERSDSGGSSSEPFERHAPPLLERGTPPVDPKLAWGDVFTTTPTDPRPLTSPLRQAADDEEEKSMRSETPPVPPPPPYLANYPGFPENGTPGPPISRFPLEESAPPGRPLPWPPGNDEAAKMQAPPPKKEPSKEEPPQLSGPEAGRKPARGGQGGPPPRRENRTETRWGPRPGSCRRGIPPEEPGVPPRRAGPIKKPPPVKVEELPPKSLEQGDETPKVPKPDALKTAKGKVGPKETPPGGNLSAPRLRRDYSYERVGPTSCRGRGRGEYFARGRGFRGTYGGRGRGARSREFRSYREFRGDDGRGGGSGGTNHPSAPRGRTASSTRSEGEYEEIPKRRRQRGSETGSETHESDLAPSDKEAPPPKEGVLGQVPLAPPQPGAPPSPAPARFSTARGGRVFTPRGVPSRRRGRGGGRPPPVCSGWSPPAKSLVKKPPTGPLPSKEPLKEKLISGPLSPMSRAGNMGVGMEDGERPRRRRHGRAQQQDKPPRFRLKQERENAARGADGKPPSLTAASTPGPEETLTAATVPPPPRRTAAKSPDLSNQNSDQANEWEWETASESSDFASERRGDKETPPAALMTSKAVGTPGANAGGAGPGISAMSRGDLSQLAKDLSKRSF

SSQRPGMDRQNRRTGKTGSGG
GSSGGGAGPGGRTGPGRGDKRSW
PSPKNRSRPPEERPPGLPLPPPPSS
SAVFRLDQVIHSNPAGIQQALQLSS
RQGNVTAPGGHPRPKGPPQAPQGS
SPRPTRYDPPRASSAISDPHFEEP
GPMVRGVGGTTPRDSAGVNPFPKRR
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SGPCSQRSSPDGGLKGS AEGPPKRP
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LPVVNFGSLPPAPPAPPPLSLLPVG
PALQPPNLAVRPPPAPAARVLPSPAR
PFAPSLGRAELHPVELKPFQDYRKLS
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