

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O15084	ANR28_HUMAN	Homo sapiens	Serine/threonine-protein phosphatase 6 regulatory ankyrin repeat subunit A	19.089993	T1009	S1007;S1011	37217939;28657654	MAFLKLRDQPSLVQAIENGDPDEVR ALIFKKEDVNFQDNEKRTPLHAAAY LGDAEIIELLILSGARVNAKDSKWL PLHRAVASCSEEAQVLLKHSADVN ARDKNWQTPLHIAAANKAVKCAEAL VPLLSNVNVSDRAGRTALHHAAFSG HGEMVLLLLSRGANINAFDCKDRRA IHWAAYMGHIEVVKLLVSHGAEVTC KDKKSYTPLHAAASSGMISVVKYLLD LGVDMNEPNAYGNTPLHVACYNGQ DVVVNELIDCGAIVNQKNEKGFTPL HFAAASTHGALCLELLVGNGADV MKS KDGKTPLHMTALHGRF SRSQTI IQSGAVIDCEDKNGNTPLHIAARYGH ELLINTLITSGADTAKRGIHGMFPLH LAALSGFSDCCRKLLSSGFDIDTPDD FGRTCLHAAAAGGNLECLNLLLNTG ADFNKDKFGRSPLHYAAANCNYQ CLFALVSGASVNDLDERGCTPLHY AATSDTDGKCLEYLLRNDANPGIRD KQGYNAVHYSAAAYGHRCLQLIASE TPLDVLMETSGTDMLSDSNRATIS PLHLAAYHGHQALEVLVQSLLDLD VRNSSGRTPDLAAAFKGHVECV DVL INQGASILVKDYILKRTPIHAAATNG HSECLRLLIGNAEPQNAVDIQDGNG QTPLMLSVLNGHTDCVYSLLNKGA NVD AKDKWGRTALHRGAVTGHEEC VDALLQH GAKCLLRDSRGRTPHLS AACGHIGVLGALLQSAASMDANPAT ADNHGYTALHWACYNGHETCVELL LEQEVFQKTEGNAFSP LHC AVINDN EGAAEMLIDTLGASIVNATDSKGRTP LHAAAFDTHVECLQLLLSHNAQVNS VDSTGKTPLMMAAENGQTN TVEML VSSASAELTLQDNSKNTALHLACSK GHETSALLILEKITDRNLINATNAAL QTPLHVAARNGLTMVVQELLGKGAS VLAVDENGYPALACAPNKDVADCL ALILATMMPVSSSSPLS SLTFNAINR YTNTSKTVSFEALPIMRNEPSSYCSF NNIGGEQEYLYTDVDELNDS DSEY