

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q99P72	RTN4_MOUSE	Mus musculus	Reticulon-4	32.530015	T509;T834	S7;S16;S105;S145;S165;S167;S329;S344;T348;S426;T430;S489;S690;S727;S768;S832;T834;S857;S961	24788674;34678516;36288343;22645316;22517741;34418053;28528544	MEDIDQSSLVSSADSPRRPPPAFKY QFVTEPEDEEEDDEEEDDEDELE ELEVELRKPAAGLSAAPVPPAAAPLL DFSSDSVPPAPRGPLPAAPPTAPERQ PSWERSPAASAPSLPPAAAVLPSKLP EDDEPPARPPAPAGASPLAEPAPPS TPAAPKRRGSGSVDETLFALPAASEP VIPSSAEKIMDLKEQPGNTVSSGQE DFPSVLFETAASLPSLSPLSTVSFKE HGYLGNLSAVASTEGTIEETLNEASR ELPERATNPFVNRESAEFSVLEYSE MGSSFNKSPKGESAMLVENTKEEVI VRSKDKEDLVCSAALHNPQESPATL TKVVKEDGVM SPEKTMDFNEMKM SVVAPVREEYADFKPFEQAWEVKDT YEGSRDVLAAANMESKVDKCCFE DSLEQKGGHGDSESRNENASFPRT ELVKDGSRAYITCDSFSSATESTAN IFPVLEDHTSENKTDEKKIEERKAQII TEKTSPKTSNPFLVAIHDSEADYVTT DNLSKVTEAVVATMPEGLTPDLVQE ACESELNEATGTKIAYETKVDLVQTS EAIQESIYPTAQLCPSFEEAEATPSV LPDIVMEAPLNSLLPSTGASVAQPSA SPLEVPSVSYDGKLEPENPPPYEE AMSVALKTSDSKEEIKEPESFNAAA QEAEAPYISIACDLIKETKLSTEPSPE FSNYSEIAKFEKSVDPHCELVDDSSP ESEPVDLFSDDSIPEVPQTQEEAVM LMKESLTEVSETVTQHKHKERLSAS PQEVGKPYLESFQPNLHITKDAASN EIPTLTKKETISLQMEEFNTAIYSND DLLSSKEDKMKESETFSDSSPIIID EFPTFVSAKDDSPKEYTDLEVS NKSE IANVQSGANSLPCSELPCDLSFKNTY PKDEAHVSDEFKSRSSSVSKVPLLLP NVSALESQIEMGNIVKPKVLTKEAEE KLPSDTEKEDRSLTAVLSAELNKTSV VDLLYWRDIKKTGVVFGASLFLLSL TVFSIVSVTAYIALALLSVTISFRIYKG VIQAIQKSDEGHPFRAYLESEVAISE ELVQKYSNSALGHVNSTIKELRRLFL VDDLVDLKFVLMWVFTYVGFALFN GLTLLILALISLFSIPVIYERHQAQIDH YLGLANKSVKDAMAKIQAKIPGLKR KAE