

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q99NH0-2	ANR17_MOUSE	Mus musculus	Isoform 2 of Ankyrin repeat domain-containing protein 17	36.8604	NaN	S15;S42;S152;S799;S1453;S1631;S1692;S1696;S1705;S2038;S2040;S2041;S2043;S2055;S2063;S2373;S2401	27669760;22645316;22517741;29187734;36852467;35822049;30059200;34678516	MLETASKLLLSGTADGADLRTVDPE TQARLEALLEAAGIGKLSTADGKAFA DPEVLRRLTSSVSCALDEAAAALTR MRAESTANAGQSDNRSLAEACSEG DVNAVRKLLIEGRSVNEHTEEGESL LCLACSAGYYELAQVLLAMHANVED RGIKGDITPLMAAANGGHVKIVKLLL AHKADVNAQSSTGNTALTYACAGGY VDVVKVLLSESGASIEDHNENGHTPL MEAGSAGHVEVARLLENGAGINTH SNEFKESALTLACYKGHLEMVRFLL EAGADQEHKTDEMH TALMEACMD GHVEVARLLLD SGAQVNMPADSFES PLTLAACGGHVELAALLIERGASLEE VNDEGYTPLMEAAREGHEEMVALL LGQGANINAQTEETQETALTLACCG GFLEVADFLIKAGADIELGCSTPLME AAQEGHLELVKYLLAAGANVHATTA TGDTALTYACENGHTD VADVLLQAG ADLEHESEGGRTPLMKAARAGHVC TVQFLISKGANVNRRTTANNDHTVLS LACAGGHLAVVELLLAHGADPTHRL KDGSTMLIEAAKGGHTSVVCYLLDY PNNLLAAPPDVTQLTPPSHDLNRA PRVPVQALPMVPPQEPDKPPANLA ATLPVRSKAVSGRASAMSNTPTHSIA ASVSQPQTPTPSPIISPSAMLPYPAID IDAQTESNHDTALT LACAGGHEELV QTLLERGASIEHRDKKGFTPLILAAT AGHVG VVEILLDNGADIEAQSERTK DTPLSLACSGGRQEVVELLLARGAN KEHRNVSDYTPLSLAASGGYVNIKI LLNAGAEINSRTGSKLGISPLMLAA MNGHTAAVKLLLDMGSDINAQIETN RNTALTLACFQGRTEVVSLLLDRKA NVEHRAKTGLTPLMEAASGGYAEVG RVLLDKGADVNAPPVSSRD TALTIA ADKGHYKFCCELLIGKAHIDVRNKK GNTPLWLAANGGHLDVVQLLVQAT ADVDAADNRKITPLMAAFRKGHVKV VRYLVKEVNQFPSDSECMRYIATITD KEMLKKCHLCMESIVQAKDRQAAE ANKNASILLELDLEKLREESRRRLAL AAKREKRKEKRRKKKEEQRRKLEEI EAKNKENFELQAAQEKEKLVVEEP EVLTEPPSATTTTTIGISATWTTLGS HGKRNTTTTTSSKRKNRKNKITPE NVQIIFDDPLPISYSQPEKVN GESKS SSTSESGSDNMRISSCSDESSNSN SSRKSNNHASAVVTTT MASKKQPSV LVTFPKEERKSVSGKASIKLSETVNE GTSNSLSTCTKSGPSPLSSPNGKLTV ASPKRGPKREEGWKEVRR