

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9EP71	RAI14_MOUSE	Mus musculus	Ankycorbin	21.713034	NaN	S11;T249;S281;S286;S293;T295;T297;S300;S304;S318;S327;S329;S340;S341;S358;S513;S516;S667;S694;S914	22645316;24788674;28528544	MKSLKAKFRKSDTNEWNKNDRLRLLQAVENGDAEKVASLLGKKGASATKH DSEGKTAFLHAAAKGHVECLKVMVT HGVDVTAQDSSGHSALHVAAKNGH PECIRKLLQYKSPAENIDNSGKTALH YAAAQGCLOAVQLLCEHKSPINLKD LDGNIPLLVAVQNGHSEACHFLLDH GADVNSRDKNGRTALMLACETGSS NTVDALIKKGADLSLVDSLGHNALH YSKLSENAGIQNLLLSKISQDADLKT PTKPKQHDQVSKISSERSGTPKRRK APPPPISPTQLSDVSSPRISITSTPLSG KESVFFAEAPFKAIEISSIQENKDRLS DSTAGADSLDISSEADQQDLLVLLQ AKVASLTLHNKELQDKLQAKSPKDK EAEADLSFQSFHSTQTDLAPSPGKA SDIPSSDAKSSPPVEHPAGTSTTDND VIIRQLQDSLHDLQKRLESSEAEKKQ LQDELQSQRDTLCLNNTSEISNGS DLSQKLKETQSKYEEAMKEVLSVQK QMKLGLLSQESADGYSHLREAPADE DIDTLKQDLQKAVEESARNKERVRE LETKLAEKEQAEATKPPAEACEELRS SYCSVIENMNKEKAFLEKYQQAQE EIMKLDKTLKSQMPQEAPDDSGDM KEAMNRMIDELNKQVSELSQLYREA QAELEDYRKRKSLDAAEYIHKAEH ERLMHVSNLSRAKSEEALSEMKSQ YSKVLNELTQLKQLVDAHKENSVSIT EHLQVITTLRRTAKEMEEKISALTGH LANKEAEVAKLEKQLAEKAAVSDA MVPKSSYEKLQASLESEVNALATKL KESVREREKAHSEVAQVRSEVSQAR REKDNIQTLLKAKEQEVTAQVQKFQ RAQEELAGMRRRCSETSSKLEEDKDE KINEMTREVVLKKEALNSLSQLSYST SSSKRQSQQLDLLQQQVKQLQNQL AECKKHHQEVISVYRMHLLYAVQGO MDEDVQKVLKQILTMCKNQSQKK