

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|------------|--------------|--|-----------------------|-------------------------------------|---|
| Q8VHG2 | AMOT_MOUSE | Mus musculus | Angiomotin | 32.735269 | S183;T196;S214;T275;T377;S697;S753;T755;S818 | S693 | 22645316;22517741;30059200;36852467 | MRSSDDQPSGGTTVLQRLQLQEQLRY GNPSENRSLLAIHQATGNSSPFST GSGNQGPQNDVLSQDHHQQQLVA HPARQEPQGQEIQSENGVMEKQLS PRMQNNEELPTYEEAKVQSQYFRG QQHASVGAIFYVTGVTNQKMRTEG RPSVQRLTPGKMHQDEGLRDLKQG HVRSLSERLMQMSLATSGVKAHPPV TSAPLSPQPNDLYKNATSSSEFYKA QGPPPSQHSKGMHEHRGPPPEYPFK GVPSQSVVCKSQEPGHFYSEHRLNQ PGRTEGQLMRYQHPPEYGAARATQ DISLSLSARN SQPHSPTSSSLTAGAS SLPLLQSPSTRLPQGHLVSNQGD HSAHLSRHQQHLLSSQSHQGDHYR HAQASLTSQQQPGEAYSAMPRAQ QSASYQPMPADPFAMVSRAQMVE ILSDENRNLRLQELDGCYEKVARLQK VETEIQRVSEAYENLVKSSSKREALE KAMRNKLEGEIRRMHDFNRDLRDR LETANKQLAEKEYEGSEDTRKTISQL FAKHKENQREKEKLEAELATARSTN EDQRRHIEIRDQALSNAQAKVVKLE EELKKKQVYVDKVEKMQQALVQLQ AACEKREQLEHRLRTRLERELESRLI QQRQGNSQPTNASEYNAAALMELL REKEERILALEADMTKWEQKYLEEN VMRHFALDAAATVAAQRDTTVISHS PNTSYDTALEARIQKEEEEILMANKR CLDMEGRIKTLHAQIIEKDAMIKVLQ QRSRKEPSKTEQLSSMRPAKSLMSI SNAGSGLLAHSSTLTGAPIMEEKR DKSWKGLGVLLGGDYRVEVPVSTP SPVPPSTPLLSAHSKTGSRDCSTQTE RGPESTKTAAVTPISAPMAGPVAAAA PAAAINATAATNTATAATNTTIMVAA APVAVAAVAPAAAAATPSPANAAAL AAAAAPATSVSAATSVSAANSISPAA PVAPAAVPPAAPVSPAAAVQIPAAA SLTPATVSPTAATAATAAVAAATTAIT AAAAAATTAIQVAPATSAPVSPASIP APATAQASAPTPTQASTPAPTEPPSP VPTPALVQTEGPANPGASSGPRRL STPNLMCNPKDPDAPAFHSSTLERK TPIQILQPEPDAEMVEYLI |