

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q4VCS5	AMOT_HUMAN	Homo sapiens	Angiomotin	24.567491	S182;T195;S213;T776	S714	34019948;29485866;29249144;30620550;29217192;35254053;35138101;32870666;32119511;28657654	MRNSEEQPSGGTTVLQRLQLQEQLRY GNPSENRSLLAIHQATGNGPPFPS GSGNPGPQSDVLSPODHHQQLVAH AARQEPQGQEIQSENLIMEKQLSPR MQNNEELPTYEEAKVQSQYFRGQQ HASVGAAFYVTGVTNQKMRTEGRPS VQRLNPGKMHQDEGLRDLKQGHVR SLSERLMQMSLATSQVKAHPPVTS PLSPPQPNDLYKNPTSSEFYKAQGP LPNQHSLKGMHRGPPPEYPFKGM PPQSVVCKPQEPGHFYSEHRLNQPG RTEGQLMRYQHPPPEYGAARPAQDIS LPLSARNSQPHSPTSSTLTSGGSLPLL QSPPSTRLSPARHPLVFNQGDHSAH LPRPQQHFPLPNQAHQGDHYRLSQP GLSQQQQQQQQHHHHHHHQQQ QQQQPQQQGEAYSAMPRAQPSSA SYQVPADPFAIVSRAQQMVEILSDE NRNLRQELEGCYEKVARLQKVETEI QRVSEAYENLVKSSSKREALEKAMR NKLEGEIRRMHDFNRDLRERLETA NKQLAEKEYEGSEDTRKTISQLFAK NKEQREKEKLEAELATARSTNEDQ RRHIEIRDQALSNAQAKVVKLEELK KKQVYVDKVEKMQQALVQLQAACE KREQLEHRLRTRLERELESRLRIQR QGNCQPTNVSEYNAALMELLREK EERILALEADMTKWEQKYLEENVM RHFALDAAATVAAQRDTTVISHSPN TSYDTALEARIQKEEEEILMANKRCL DMEGRIKTLHAQIIEKDAMIKVLQQR SRKEPSKTEQLSCMRPAKSLMSISN AGSGLLSHSSTLTGSPIMEEKRDDK SWKGSLLGILLGGDYRAEYVPSTPSV PPSTPLLSAHSKTGSRDCSTQTERGT ESNKTAAPVAPISVPAPVAAAAATAAIT ATAATITTTMVAAAPVAVAAAAAPAA AAAPSPATAAATAAVSPAAAGQIPA AASVASAAAVAPSAAAAAAVQVAPAA PAPVPAPALVPVPAPAAAQASAPAQT QAPTSAPAVAPTPAPTTPAVAQAEV PASPATGPGPHRLSIPSLTCNPKTD GPVFHSNTLERKTPIQILGQEPDAE MVEYLI