

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q80VC9	CAMP3_MOUSE	Mus musculus	Calmodulin-regulated spectrin-associated protein 3	26.425892	T1197;S1209	T184;S193;S334;S347;S351;S368;S373;S382;S548;S555;S561;S683;S767;T797;S812;S881;S1077	22645316;36852467	MVEAAPAGSGPLRRTFLVPEIKSLDQ YDFSRAKAAASLAWVLRAAFGGAEH VPELWEPFYTDQYAEHVKPPVTR LLLSAELYCRAWRQALPQLEPSPSPS ALLALLARRGTVPSLPEHPVREADLK HQPILMGAHLAVIDALMVAFSFEWT KTLPGPLALSSLEHKLLFWVDTTVR RLQEKTEQEAQAQRASPAAPLDGASP AQPSIRYRKDRAIARRAPCFPNVTTL QDLASGAALAATIHCYCPQLLRLEEV CLKDPMSVADSLYNLQLVQDFCASH LPRGCPLSLEDLLYVPPPLKVNLVVL LAEMYMCFEVLKPDFVQAKDLPDG HAVSPRNTETVPSQNNSSGSSSPVFN FRHPLLSPGGPQSPLRGSTGSLKSSP SMHMEALGKAWNRQLSRPLSQAV SFSTPFGLDSDVDVVMGDPVLLRSV SSDSLGP RPVSTSSRN SAQPAPESG DLPTIEEALQIIHSAEPRLLPDGAADG SFYLHSPEGLSKPPLSPYPPEGASKP LSDRLNKAPIYISHPENPSKSSPCST GEILKPPPPSEGS PKAVASSPAANNS EVKMTSFAERKKQLVKAESAESGLGS PTSTPVAPEALSSEMSELGARLEEKR RAIEAQKRRIE AIFAKHRQLGKSAF LQVQPREAAGEAE EAEELGSVPGGE RPAGEGQGEPSLRHKS VTFSPDLGP VPPEGLGDYNRAVSKLSAALSSLQR DMQRLTDQQRLLAPPEAPGPAPPP AAWVIPGPATGPKAASPS PARRAPAA RRSPGPGSPTPRSPKHARPAELKLA PLTRVLT PPHD VDSLPHLRKFSPSQV PVQTRSSILLSEGT PPEEPTTKPALIEI PLASLGEPAADEEGDGSPPGAEDSL EEEASSEGEPRSGLGFFYKDEDKPE DEMAQKRASLLERQRRVEEARRR KQWQEA EKEQKREEAARLAQEAPG LAFTTPVVASAAPVATLAPTTRAMAP AEEVGP RR GDFTRLEYERRAQLKL MDDLDKVLRPRASGTGGPGRGRR ATRPRSGCCDSALARSPARGLLGS RLSKVYSQSTLSLSTVANEAPNNLG VKRPTSRAPSPSGLMSPSRLPGSRE RDWENGSNASSPASVPEYTGPRLYK EPSAKSNKFIHINALSHCCLAGKVN EPQKNRILEEIEKSKANHFLILFRDS SCQFRALYTLSGETEELSRLAGYGPR TVT PAMVEGIYKYNSDRKRFTQIPAK TMSMSVDAFTIQGHLWQSKKPTTPK KGGGTPK