

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9P1Y5	CAMP3_HUMAN	Homo sapiens	Calmodulin-regulated spectrin-associated protein 3	10.887383	T338;T565;S566	T184;S193;S334;S341;S347;S351;S368;S373;S382;S547;S554;S560;S685;S769;T799;S814;S1074	30379171;35132862;35254053;28657654	MVEAAPPGPGPLRRTFLVPEIKSLDQ YDFSRAKAAAASLAWVLRAAFGGAEH VPPELWEPFYTDQYAEHVKPPVTR LLLSAELYCRAWRQALPQLETPPNP SALLALLARRGTVPALPERPVREADL RHQPILMGAHLAVIDALMAAFAFEW TKTLPGLALTSLEHKLLFWVDTTVR RLQEKTEQEAQASPAAPADGAAP AQPSIRYRKDRVARRAPCFPTVTSL QDLASGAALAATIHICYCPQLLRLEEV CLKDPMSVADSLYNLQLVQDFCASR LPRGCPLSLEDLLYVPPPLKVNLVVM LAELFMCFEVLKPDFVQVKDLPDGH AASPRGTEASPPQNNSSGSSSPVFTF RHPLLSSGGPQSPLRGSTGSLKSSPS MSHMEALGKAWNRLSRPLSQAVS FSTPFGLDSDVDVVMGDPVLLRSVS SDSLGPPRPAPARTPTQPPPEPGDLP TIEEALQIIHSAEPRLLPDGAADGSFY LHSPGPKPSLASPYLPEGTSKPLS DRPTKAPVYMPHPETPSKPSCLVG EASKPPAPSEGSPKAVASSPAATNSE VKMTSFAERKKQLVKAEEAGAGSP TSTPAPPEALSSEMSELARLEEKRR AIEAQKRRIEAIFAKHRQRLGKSAFL QVQPREASGEAEAEAEADSGPVPG GERPAGEGQGEPTSRPKAVTFSPDL GPVPHEGLGEYNRAVSKLSAALSSL QRDMQRLTDQQRLLAPPEAPGSAP PPAAWVIPGPTTGPKAASPPARRVP ATTRSPGPGPSQSPRSPKHTRPAELR LAPLTRVLTTPHDVDSLPHLRKFSPS QVPVQTRSSILLAETPPEPAARPG LIEIPLGSLADPAAEDEGDGSPAGAE DSLEEEASSEGEPRVGLGFFYKDED KPEDEMAQKRASLLERQORRAEEA RRRKQWQVEVEQRREEAARLAQE EAPGPAPLVSAPMATPAPAARAPAE EEVGPRKGFTRQEYERRAQLKLM DDLKVLPRPRAAGSGGPGRGGRRAT RPRSGCCDDALARSARGLLGSRL SKIYSQSTLSLSTVANEAHNNLGVK RPTSRAPSPGLMSPSRLPGSRERD WENGSNASSPASVPEYTGPRLYKEP SAKSNKFIIHNALSHCCLAGKVN QKNRILEEIEKSKANHFLILFRDSSC QFRALYTLSTGTEELSRLAGYGPRTV TPAMVEGIYKYNSDRKRFTQIPAKT MSMSVDAFTIQGHLWQGGKPTTPK KGGGTPK