

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9QUH6	SYGP1_RAT	Rattus norvegicus	Ras/Rap GTPase-activating protein SynGAP	28.104173	NaN	T34;T39;S117;S371;S379;S385;S449;S466;S752;S766;S780;S823;S825;T828;S836;S840;S842;S876;S892;S895;S898;S985;S1114;S1118;S1121;S1165;S1204	17496889;22645316;19322778	MSRSRASIHRGSPAMSYAPFRDVR GPPMHRTQYVHSPYDRPGWNPRFC IISGNQLLMLDEDEIHPLLRDRRSE SSRNKLLRRTVSVPEGRPHGEHEY HLGRSRRKSVPGGKQYSMEAAPAAP FRPSQGFSLRRLKSSIKRTKSQPKLD RTSSFRQILPRFRSADHDRARLMQS FKESHSHESLLSPSSAAEALELNLD EDSIIKPVHSSILGQEFCEVTTSSGT KCFACRSAAERDKWIENLQRAVKPN KDNSRRVDNVLKLWIEARELPPKK RYYCELCLDDMLYARTTSKPRSASG DTVFWGEHFEFNLPVAVRALRLHLY RDSDKKRKKDKAGYVGLVTPVATL AGRFTEQWYPVTLPTGSGGSGGM GSGGGGGSGGGSGGKGGGCPAVR LKARYQTMSILPMELYKEFAEYVTN HYRMLCAVLEPALNVKGKEEVASAL VHILQSTGKAKDFLSDMAMSEVDRF MEREHLIFRENTLATKAIEEYMRLIG QKYLKDAIGEFIRALYESEENCEVDP IKCTASSLAEHQANLRMCCELALCK VVNSHCVFPRELKEVFASWRLRCAE RGREDIADRLISASLFLRFLCPAIMSP SLFGLMQEYPDEQTSRTLTLIAKVIQ NLANFSKFTSKEDFLGFMNEFLELE WGSMQQFLYEISNLDTLTNSSSFEG YIDLGRELSTLHALLWEVLPQLSKEA LLKLGPLPRLSDISTALRNPNIQRQ PSRQSERARSQPMVLRGPSAEMQG YMMRDLNSSIDLQSFMARGLNSSM DMARLPSPTKEKPPPPPPGGGKDLF YVSRPPLARSSPAYCTSSSDITEPEQ KMLSVNKSVMMLDLQGDGPGGRLN SSSVSNLAAVGDLHSSQASLTAAL GLRPAPAGRLSQSGSSITAAGMRL SQMGVTTDGVPAQQLRIPLSFQNPL FHMAADGPGPPAGHGGSSGHGPPS SHHHHHHHHHHRGGEPPGDTFAP FHGYKSEDLSTGVPKPPAASILHSH SYSDEFGPSGTDFTRRQLSLQDNLQ HMLSPPQITIGPQRPA PSGGGGSG GGSGGGGGGQPPPLQRGKSQQLTV SAAQKPRSSGNLLQSPESYGPARP RQQLSKEGSIGSGGSGGGGGGGL KPSITKQHSQTPSTLNPTMPASERTV AWVSNMPLSADIESAHIEREEYKL KEYSKSMDESRLDRVKEYEEIHS KERLHMSNRKLEEYERRLLSQEEQT SKILMQYQARLEQSEKRLRQQQVEK DSQIKSIIGRLMLVEEELRRDHPAMA

