

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
F6SEU4	SYGP1_MOUSE	Mus musculus	Ras/Rap GTPase-activating protein SynGAP	22.646418	NaN	T34;T39;S117;S371;S379;S385;S449;S466;S752;S766;S780;S823;S825;T828;S836;S840;S842;S876;S892;S895;S898;S985;S1111;S1115;S1118;S1162;S1201	16452088;33300544	MSRSRASIHRSIPAMSYAPFRDVR GPPMHRTOYVHSPYDRPGWNPRFC IISGNQLLMLDEDEIHPLLLRDRSE SSRNKLLRRTVSVGPVEGRPHGEHEY HLGRSRRKSVPGKQYSMEAAPAAP FRPSQGFLSRRLKSSIKRRTKSQPKLD RTSSFRQILPRFRSADHDRARLMQS FKESHSHESLLSPSSAAEALELNLD EDSIIKPVHSSILGQEFCEVTTSSGT KCFACRSAAERDKWIENLQRAVKPN KDNSRRVDNVLKLWIIARELPPKK RYYCELCLDDMLYARTTSKPRSASG DTVFWGEHFEFNNLPAVRALRLHLY RDSDKKRKKDKAGYVGLVTPVATL AGRHFTEQWYPVTLPTGSGGSGGM GSGGGGGSGGGSGGKGGGCPAVR LKARYQTMSILPMELYKEFAEYVTN HYRMLCAVLEPALNVKGKEEVASAL VHILQSTGKAKDFLSDMAMSEVDRF MEREHLIFRENTLATKAIEEYMRLIG QKYLKDAIGEFIRALYESEENCEVDP IKCTASSLAEHQANLRMCCELALCK VVNSHCVFRELKEVFASWRLRCAE RGREDIADRLISASFLRFLCPAIMSP SLFGLMQEYPDEQTSRTLTLIAKVIQ NLANFSKFTSKEDFLGFMNEFLELE WGSMQQFLYEISNLDLTLNSSSFEG YIDLGRELSTLHALLWEVLPQLSKEA LLKLGPLPRLNDISTALRNPNIQRQ PSRQSERTRSQPMVLRGPSAEMQG YMMRDLNSSIDLQSFMARGLNSSM DMARLPSPPTKEPPPPPPGGGKDLF YVSRPPLARSSPAYCTSSSDITEPEQ KMLSVNKSVSMLDLQGDGPGGRLN SSSVSNLAAVGDLLHSSQASLTAAL GLRPAPAGRLSQSGSSITAAGMRL SQMGVTTDGVPAQQLRIPLSFQNPL FHMAADGPGPPAGHGGSSGHGPPS SHHHHHHHHHHRGGEPPGDTFAP FHGYKSEDLSSGVPKPPAASILHSH SYSDEFGPSGTDFTRRQLSLQDSLQ HMLSPPQITIGPQRPAAPSGGGGSG GGSGGGQPPPLQRGKSQQLTVSAAQ KPRPSSGNLLQSPEPSYGPAPRQ SLSKEGSIGSGGGGGGGGGLKPSI TKQHSQTPSTLNPTMPASERTVAWV SNMPHLSADIESAHIEREEYKLKEYS KSMDESRLDRVKEYEEEEIHSKERL HMSNRKLEEYERRLLSQEEQTSKIL

