	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum
0	Q96PV0	SYGP1_HUMAN	Homo sapiens	Ras/Rap GTPase-activating protein SynGAP	32.965235	S809;T822;S840;S858;S859;S873;T878;S892;S898;T900;T912;T913;S983;S1005;S1079;S1087;T1134;T1147;S1159;T1310	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	35637289;28657654	MSRSRASIHRGSIPAMSYAPFRDVR GPSMHRTQYVHSPYDRPGWNPRFC IISGNQLLMLDEDEIHPLLIRDRRSE SSRNKLLRRTVSVPVEGRPHGEHEY HLGRSRRKSVPGGKQYSMEGAPAAP FRPSQGFLSRRLKSSIKRTKSQPKLD RTSSFRQILPRFRSADHDRARLMQS FKESHSHESLLSPSSAAEALELNLD EDSIIKPVHSSILGQEFCFEVTTSSGT KCFACRSAAERDKWIENLQRAVKPN KDNSRRVDNVLKLWIIEARELPPKK RYYCELCLDDMLYARTTSKPRSASG DTVFWGEHFEFNNLPAVRALRLHLY RDSDKKRKKDKAGYVGLVTVPVATL AGRHFTEQWYPVTLPTGSGGSGGM GSGGGGGSGGGSGGKGKGGCPAVR LKARYQTMSILPMELYKEFAEYVTN HYRMLCAVLEPALNVKGKEEVASAL VHILQSTGKAKDFLSDMAMSEVDRF MEREHLIFRENTLATKAIEEYMRLIG QKYLKDAIGEFIRALYESEENCEVDP IKCTASSLAEHQANLRMCCELALCK VVNSHCVFPRELKEVFASWRLRCAE RGREDIADRLISASLFLRFLCPAIMSP SLFGLMQEYPDEQTSRTLTLIAKVIQ NLANFSKFTSKEDFLGFMNEFLELE WGSMQQFLYEISNLDTLTNSSSFEG YIDLGRELSTLHALLWEVLPQLSKEA LLKLGPLPRLLNDISTALRNPNIQRQ PSRQSERPRPQPVVLRGPSAEMQGY MMRDLNSSIDLQSFMARGLNSSMD MARLPSPTKEKPPPPPPGGGKDLFY VSRPPLARSSPAYCTSSSDITEPEQK MLSVNKSVSMLDLQGDGPGGRLNS SSVSNLAAVGDLLHSSQASLTAALG LRPAPAGRLSQGSGSSITAAGMRLS QMGVTTDGVPAQQLRIPLSFQNPLF HMAADGPGPPGGHGGGGGHGPPSS HHHHHHHHHHRGGEPPGDTFAPF HGYSKSEDLSSGVPKPPAASILHSHS YSDEFGPSGTDFTRRQLSLQDNLQH MLSPPQITIGPQRPAPSGPGGGSGGG SGGGGGGQPPPLQRGKSQQLTVSAA QKPRPSSGNLLQSPEPSYGPARPRQ QSLSKEGSIGGSGGSGGGGGGGLKP SITKQHSQTPSTLNPTMPASERTVA WVSNMPHLSADIESAHIEREEYKLK EYSKSMDESRLDRVKEYEEEIHSLK ERLHMSNRKLEEYERRLLSQEEQTS KILMQYQARLEQSEKRLRQQQAEKD SQIKSIIGRLMLVEEELRRDHPAMAE PLPEPKKRLLDAQERQLPPLGPTNPR VTLAPPWNGLAPPAPPPPPRLQITEN GEFRNTADH	None	None	None	None	None	None