

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O75044	SRGP2_HUMAN	Homo sapiens	SLIT-ROBO Rho GTPase-activating protein 2	7.726555	NaN	S206;S427;S500;S691;S695;S724;S795;S916;S930;S990;S994;S1013;S1027	30379171;23301498	MTSPAKFKKDKEIIAEYDTQVKEIRA QLTEQMKCLDQOCELRVQLLQDLQ DFFRKKAEIEMDYSRNLEKLAERFL AKTRSTKDQQFKKQNVLSPVNCW NLLLNQVKRESRDHTTSLSDIYLNII PRFVQVSEDSGRLFKKSKEVGGQLQ DDLKVLNELYSVMKTYHMYNADS ISAQSKLKEAEKQEEKQIGKSVKQED RQTPRSPDSTANVRIEEKHVRSSV KKIEKMKEKRQAKYTENKLKAIKAR NEYLLALEATNASVFKYYIHDLSDLI DQCCDLGYHASLNRLRTFLSAELN LEQSKHEGLDAIENAVENLDATSDK QRLMEMYNNVFCPPMKFEFQPHM GDMASQLCAQQPVQSELVQRCQQL QSRLSTLKIENEVVKKTMEATLQTIQ DIVTVEDFDVSDCFQYSNSMESVKS TVSETFMSKPSIAKRRANQQETEQF YFTKMKEYLEGRNLITKLQAKHDLL QKTLGESQRTDCSLARRSSTVRKQD SSQAIPLVVESCIRFISRHGLQHEGIF RVSGSQVEVNDIKNAFERGEDPLAG DQNDHDMDSIAGVLKLYFRGLEHPL FPKDIFHDLMACVTMDNLQERALHI RKVLLVLPKTTLIIMRYLFAFLNHL QFSEENMMDPYNLAICFGPSLMSVP EGHDQVSCQAHVNELIKTIIQHENI FPSPRELEGPVYSRGGSMEDYCDSP HGETTSVEDSTQDVTAEHHTSDDEC EPIEAIKFDYVGRtarelsfkkgas LLLYQRASDDWWEGRHNGIDGLIPH QYIVVQDTEdgvversspkseiEIVISE PPEEKVTARAGASCPSGGHVADIYLA NINKQRKRPESGSIRKTFRSDSHGLS SSLTDSSSPGVGASCRPSSQPIMSQS LPKEGPKCSISGHGSLNSISRHSSL KNRLDSPQIRKTATAGRSKSFNNHR PMDPEVIAQDIEATMNSALNELREL ERQSSVKHTPDVVLDTLEPLKTSPPV APTSEPSSPLHTQLLKDPEPAFQRSA STAGDIACAFRPVKSVMKMAAPVKPPA TRPKPTVFPKTNATSPGVNSSTSPQS TDKSCTV