

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q91Z67	SRGP2_MOUSE	Mus musculus	SLIT-ROBO Rho GTPase-activating protein 2	27.379393	S990	S206;S427;S500;S691;S695;S724;S795;S916;S930;S990;S994;S1013;S1027	22517741;28528544;36852467;33300544	MTSPAKFKKDKKEIIAEYDTQVKEIRA QLTEQMKCLDQQCELRVQLLQDLQ DFFRKKAEIEMDYSRNLEKLAERFL AKTRSTKDQQFKKDQNVLSPVNCW NLLLNQVKRESRDHTTSLDIYLNII PRFVQVSEDSGRLFKKSKEVGGQQLQ DDLKVLNELYSVMKTYHMYNADS ISAQSKLKEAEKQEEKQIGKSVKQED RQTPRSPDSTANVRIEEKHVRSSV KKIEKMKEKRQAKYTENKLKAIKAR NEYLLALEATNASVFKYIHDLSDIID QCCDLGYHASLNRLRFLSAELNL EQSKHEGLDAIENAVENLDATSDKQ RLMEMYNNVFCPPMKFEFQPHMG DMASQLCAQQPVQSELVQRCQQLQ SRLSTLKIENEEVKKTMEATLQTIQD IVTVEDFDVSDCFQYSNSMESVKST VSETFMSKPSIAKRRANQOETEQFY FTKMKEYLEGRNLITKLQAKHDLQ KTLGESQRTDCSLARRSSTVRKQDS SQAIPLVVEESCIRFISRHLQHEGIFR VSGSQVEVNDIKNAFERGEDPLAGD QNDHDMDSIAGVLKLYFRGLEHPLF PKDIFHDLIACVTMDNLQERAVHIR KVLVLVLPKPTLIIMRYLFAFLNHLSQ FSEENMMDPYNLAICFGPSLMSVPE GHDQVSCQAHVNELIKTIIQHENIF PNPRELEGPIYSRGGSMEDYCDSTH GETTSAEDSTQDVTAEHHTSDDECE PIEAIKFDYVGRTARELSFKKGASL LLYQRASDDWWEGRHNGIDGLIPH QYIVVQDTEGTVVERSSPKSEIEVMS EPPEEKVTARTGASCPGGHVADIYL ANINKQRKRPESGSIRKAFRSDSHG LGSSLTDSSSLGVGASCRPSSQPIMS QNLPKGPDKCSISGHGSLNSISRH SSLKNRMDSPQIRKTATAGRSKSFN NHRPMDPEVIAQDIEATMNSALNEL QELERQSSAKHTPDVVLDTLEPLKT SPVVAPTSEPSSPLHTQLLKDPEPAF QRSASTAGDIACAFRPVKSVMKMAAP VKPPATRPKPTVFPKTNATSPGVNSS ASPQATDKSCTV