

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	20.655088	S990	S206;S427;S500;S691;S695;S724;S795;S916;S930;S990;S994;S1013;S1027	33300544;28528544;22517741	MTSPAKFKKDKEIIAEYDTQVKEIRA QLTEQMKCLDQQCELRVQLLQDLQ DFFRKKAEIEMDYSRNLEKLAERFL AKTRSTKDQQFKDQNVLSPVNCW NLLLNQVKRESRDHTTSLDIYLNII PRFVQVSEDSGRLFKKSKEVGGQLQ DDLKVLNELYSVMKTYHMYNADS ISAQSKLKEAEKQEEKQIGKSVKQED RQTPRSPDSTANVRIEEKHVRRSSV KKIEKMKEKRQAKYTENKLAIKAR NEYLLALEATNASVFKYYIHDLSDIID QCCDLGYHASLNRALRTFLSAELNL EQSKHEGLDAIENAVENLDATSDKQ RLMEMYNNVFCPPMKFEFQPHMG DMASQLCAQQPVQSELVQRCQQLQ SRLSTLKIENEVVKKTMEATLQTIQD IVTVEDFDVSDCFQYSNSMESVKST VSETFMKPSIAKRRANQQETEQFY FTKMKEYLEGRNLITKLQAKHDLQ KTLGESQRTDCSLARRSSTVRKQDS SQAIPLVVESCIRFISRHGLQHEGIFR VSGSQVEVNDIKNAFERGEDPLAGD QNDHDMDSIAGVLKLYFRGLEHPLF PKDIFHDLIACVTMDNLQERAVHIR KVLLVLPKPTLIIMRYLFAFLNHLSQ FSEENMMDPYNLAICFGPSLMSVPE GHDQVSCQAHVNELIKTIIHQHENIF PNPRELEGPIYSRGGSMEDYCDSTH GETTSAEDSTQDVTAEHHTSDDECE PIEAIKFDYVGRTARELSFKKGASL LLYQRASDDWWEGRHNGIDGLIPH QYIVVQDTEDEGVVERSSPKSEIEVMS EPPEEKVTARTGASCPSGGHVADIYL ANINKQRKRPESGSIRKAFRSDSHG LGSSLTDSSSLGVGASCRPSSQPIMS QNLKPEGPKCSISGHGSLNSISRH SSLKNRMDSPQIRKTATAGRSKSFN NHRPMDPEVIAQDIEATMNSALNEL QELERQSSAKHTPDVVLDLTLPLKT SPVVAPTSEPSSPLHTQLLKDPEPAF QRSASTAGDIACAFRPVKSVMKMAAP VKPPATRPKPTVFPKTNATSPGVNSS ASPQATDKSCTV