

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q62417	SRBS1_MOUSE	Mus musculus	Sorbin and SH3 domain-containing protein 1	44.332106	S593;S597;S1199;S1200	T51;S55;S58;S62;T179;S185;S204;S209;S254;S261;S270;T325;S345;S376;S407;T421;S432;T475;S969;T1189;T1193;T1198;S1201;S1209;T1238;S164;S346;S194;S376;S164;S346;S175;S357;S288;S470	26192747;30016717;22517741;36288343;35822049;22645316;16452088	MSSECDVGSSKAVVNGLASGNHGP DKDMDPTKICTGKGTVTLRASSSYR GTPSSSPVSPQESPKHESKSGLEPED PSADEWKLSSADTNGNAQPSPLAA KGYRSVHPSLSADKPKQSPLLNEVS SSHIEDTSQDFPPTSRPSSAYPSTTIV NPTIVLLQHNRPASERRAGEQDPV PTPAELTSPGRASERRAKDASRRVVR SAQDLSDVSTDEVGIPLRNTERSKD WYKTMFKQIHKLNRDDSDVHSPR YFSDDTKSPLSVPRSKSEMNYIEGE KVKRSATLPLPARSSSLKSSPERND WEPLDKKVDTRKYRAEPKSIYEQP GKSSVLTNEKMSRDISPEEIDLKNEP WYKFFSELEFGRPTNLEKDLSFCQA ELEADLEKVETVNKSPSANSPOSSA VSPTPDITSEPPGYIYSSNFHAVKRES DGTGGLASLENERQIYKSVLEGGDI PLQGLSGLKRPSSASTKVDRKGGN AHMISSSSVHSRTFHTSNALGPGCK HKKPLSAAKACISEILPSKFKPRLSAP SALLQEQKSVLLPSEKAQSCENLCV SLNDSKRGLPLRVGGSIENTLLMRSR RDYDSKSSSTMSLQEYGTSSRRPCP LSRKAGLHFMSFYRDMHQINRAGL SLGSISSSVRDLASHFERSLTLAR GELGASQEGSEHIPKHTVSSRITAFE QLIORSRSMPSLDFSGRLSKSPTPVL SRSGLTSARSAESLLESTKLRPREM DGMDSGGVYASPTCSNMADHALSF RSLVPSSEPLSICSELDHCSNVSND REGSGGSVHGDFPKHRLNKCKGTC PASYTRFTTIRKHEQQSSRQSDWRS DSRGDKNSLLRNIHLMSPFPRLKK PLQQHPRQPPPSDSESAPAGQKADL PCHDPQDQPHSAGKPQVPTRLSSRH TMARLSHNSEPLDRPAGLEDCTRA INNGNPVPYSDHGLDRNNNPQSEL AAAHGDESEPRHFIPADYLESTEEFI RRRHDDKEKLLADQRRLLKREQEEA DIAARRHTGVIPTHHQFITNERFGDL LNIDDTAKRKSGLMRPARAKFDK AOTLKELPLQKGDVVYIRQIDQNW YEGEHHGRVGFPRTYIELLPPAEKA QPRKLAPVQVLEYGEAIAKFNFNNGD TQVEMSRKGERITLLRQVDENWYE GRIPGTSRQGIFFITYVDVLRPLVKT PVDYIDLPSYSSPSRSATVSPQQPQA QRRRVTDRSQPSLDLCSYQALYSY VPQNDDELELRDGDIVDMEKCDL GWFVGTSSRRTRQFGTFPGNYVKPLY L