

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q60520	SIN3A_MOUSE	Mus musculus	Paired amphipathic helix protein Sin3a	32.932374	S251	S10;S277;T284;S833;S861;S941;S1090;S1113	29187734;34887587;21606357;34678516;22826440;22517741;23835259	MKRRLDDQESPVYAAQRRIPGSTE AFSHQHRVLPAPPVYEAVSETMQS ATGIQYSVAPNYQVSAVPQSSGSHG PAIAAVHSSHHHPTAVQPHGGQVVQ SHAHPAPPVAPVQGGQQFQRLKVED ALSYLDQVKLQFGSQPVYNDFLDI MKEFKSQSIDTPGVISRVSQLFKGHP DLIMGFNTFLPPGYKIEVQTNDMVN VTTPGQVHQIPTHGIQPQPQPPQHP SQPSSQSAPTPAQAPQPTAAKVS KP SQLQAHTPASQQTPLPPYASPRSP VQPHTPVTISLGTAPSLQNNQPVEF NHAINYVNKIKNRFQGPDIYKAFLE ILHTYQKEQRNAKEAGGNYTPALTE QEVYAQVARLFKNQEDLLSEFGQFL PDANSSVLLSKTTAEKVDSVRNDHG GTVKKPQLNKNPQRPSQNGCQIRR HSGTGATPPVKKKPKLMSLKESMA DASKHGVGTESLFFDKVRKALRSAE AYENFLRCLVIFNQEVISRAELVQLV SPFLGKPELFWFKNFLGYKESVH LESFPKERATEGIAMEIDYASCKRLG SSYRALPKSYQQPKCTGRTPLCKEVL NDTWVSFPSWSEDSTFVSSKKTQYE EHIYRCEDERFELDVVLETNLATIRV LEAIQKLSRLSAEEQAKFRDNTLG GTSEVIHRKALQRIYADKAADIIDGLR KNPSIAVPIVLKRLKMKEEWREAQ RGFNKVVREQNEKYLLKSLDHQGI NFKQNDTKVLRSKSLLNEIESIYDER QEATEENAGVPVGPVPHLSLAYEDKQ ILEDAAALIIHHVKRQTGIQKEDKYKI KQIMHHFIPDLLFAQRGDLSDVEEE EEEEMDVDEATGAPKKHNGVGGSP PKSKLLFSNTAAQKLRGMDEVYNLF YVNNNWIYFMRHQILCLRLLRICS QAERQIEEENREREREVLEVLGKRD KSDSPAQLRLKEPMDVDVEDYYP FLDMVRSLLDGNIDSSQYEDSLREM FTIHAYIAFTMDKLIQSIVRQLQHIVS DEVCVQVTDLYLAENNGATGGQL NSQTSRSLLESAYQRKAEQLMSDEN CFKLMFIQSQQGQVQLTVELLDTTEE NSDDPVEAERWSYVERYMSSDTTS PELREHLAQKPVFLPRNLRRIRKCCQ RGREQEKEGKEGNSKKTMENTVES LDKLECRFKLNSYKMYVVIKSEYDM YRRTALLRAHQSHERVSKRLHQRFQ AWVDKWTKEHVPREMAAETSKWL MGEGLEGLVPCTTTCDTETLHFVSI NKYRVKYGTVFKAP