

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q96ST3	SIN3A_HUMAN	Homo sapiens	Paired amphipathic helix protein Sin3a	26.236195	S251;S254;S263;T266;T284;T287;S289;T292;S295;S1231	S10;S277;T284;S832;S860;S940;S1089;S1112	29351928;34725712;34019948;28510447;32522876;22661428;35254053;35132862;12150998;30059200;23301498;35008409;32119511;33214551	MKRRLDDQESPVYAAQQRRIIPGSTE AFPHQHRVLPAPPVYEAVSETMQS ATGIQYSVTPSYQVSAMPQSSGSHG PAIAAVHSSHHHPTAVQPHGGQVVQ SHAHPAPPVAPVQGGQQFQRLKVED ALSYLDQVKLQFGSQPVYNDFLDI MKEFKSQSIDTPGVISRVSQLFKGHP DLIMGFNTFLPPGYKIEVQTNDMVN VTTPGQVHQIPTHGIQPPQPQQHP SQPSAQSAAPAPAQAPQPPPAKVSKEP SQLQAHTPASQQTPLPPYASPRSP VQPHTPVTISLGTAPSLQNNQPVF NHAINVYVNIKNRFFQGGQPDYKAFLE ILHTYQKEQRNAKEAGGNYTPALTE QEVYAQVARLFKNQEDLLSEFGQFL PDANSSVLLSKTTAEKVDSVRNDHG GTVKKPQLNKNKQRPQPSQNGCQIRR HPTGTTPPVKKKPKLLNLKDSMAD ASKHGGGTESLFFDKVRKALRSAEA YENFLRCLVIFNQEVISRAELVQLVS PFLGKFPPELFNWFKNFLGYKESVHL ETYPKERATEGIAMEIDYASCKRLGS SYRALPKSYQQPKCTGRTPLCKEVL NDTWVSFPWSVEDSTFVSSKKTQYE EHIYRCEDERFELDVVLETNLATIRV LEAIQKKLSRLSAEEQAKFRDNTLG GTSEVIHRKALQRIYADKAADIIDGLR KNPSIAVPIVLKRLKMKEEWEWREAQ RGFNKVVWREQNEKYLLKSLDHQGI NFKQNDTKVLRSKSLLNEIESIYDER QEATEENAGVPVPHLSLAYEDKQ ILEDAAALIIHHVKRQTGIQKEDKYKI KQIMHHFIPDLLFAQRGDLSDVEEE EEEEMDVDEATGAVKKHNGVGGSP PKSKLLFSNTAAQKLRGMDEVYNLF YVNNNWWYIFMRLHQILCLRLLRICS QAERQIEEENREREWEREVLGIKRD KSDSPAQLRLKEPMDVDVEDYYP FLDMVRSLLDGNIDSSQYEDSLREM FTIHAYIAFTMDKLIQSIVRQLQHIVS DEICVQVTDLYLAENNGATGGQLN TQNSRSLESTYQRKAEQLMSDENC FKLMFIQSQQVQLTIELLDTEEENS DDPVEAERWSDYVERYMNSDTTSP ELREHLAQKPVFLPRNLRRIRKQOR GREQOEKEGKEGNSKKTMENTVDSL DKLECRFKLNSYKMVYVIKSEYMY RRTALLRAHQSHERSKRLHQRFQA WVDKWTKEHVPREMAAETSKWLM GEGLEGLVPCTTTCDTETLHFVSINK YRVKYGTVFKAP