| | UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochndrion | endoplasm reticulum |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | Q60520 | SIN3A_MOUSE | Mus musculus | Paired amphipathic helix protein Sin3a | 47.583305 | S28;S251;T266;S295 | S10;S277;T284;S833;S861;S941;S1090;S1113 | 22826440;36852467;23835259;34887587;21606357;22517741;34678516;29187734 | MKRRLDDQESPVYAAQQRRIPGSTE<br>AFSHQHRVLAPAPPVYEAVSETMQS<br>ATGIQYSVAPNYQVSAVPQSSGSHG<br>PAIAAVHSSHHHPTAVQPHGGQVVQ<br>SHAHPAPPVAPVQGQQQFQRLKVED<br>ALSYLDQVKLQFGSQPQVYNDFLDI<br>MKEFKSQSIDTPGVISRVSQLFKGHP<br>DLIMGFNTFLPPGYKIEVQTNDMVN<br>VTTPGQVHQIPTHGIQPQPQPPPQHP<br>SQPSSQSAPTPAQPAPQPTAAKVSKP<br>SQLQAHTPASQQTPPLPPYASPRSPP<br>VQPHTPVTISLGTAPSLQNNQPVEF<br>NHAINYVNKIKNRFQGQPDIYKAFLE<br>ILHTYQKEQRNAKEAGGNYTPALTE<br>QEVYAQVARLFKNQEDLLSEFGQFL<br>PDANSSVLLSKTTAEKVDSVRNDHG<br>GTVKKPQLNNKPQRPSQNGCQIRR<br>HSGTGATPPVKKKPKLMSLKESSMA<br>DASKHGVGTESLFFDKVRKALRSAE<br>AYENFLRCLVIFNQEVISRAELVQLV<br>SPFLGKFPELFNWFKNFLGYKESVH<br>LESFPKERATEGIAMEIDYASCKRLG<br>SSYRALPKSYQQPKCTGRTPLCKEVL<br>NDTWVSFPSWSEDSTFVSSKKTQYE<br>EHIYRCEDERFELDVVLETNLATIRV<br>LEAIQKKLSRLSAEEQAKFRLDNTLG<br>GTSEVIHRKALQRIYADKAADIIDGLR<br>KNPSIAVPIVLKRLKMKEEEWREAQ<br>RGFNKVWREQNEKYYLKSLDHQGI<br>NFKQNDTKVLRSKSLLNEIESIYDER<br>QEQATEENAGVPVGPHLSLAYEDKQ<br>ILEDAAALIIHHVKRQTGIQKEDKYKI<br>KQIMHHFIPDLLFAQRGDLSDVEEE<br>EEEEMDVDEATGAPKKHNGVGGSP<br>PKSKLLFSNTAAQKLRGMDEVYNLF<br>YVNNNWYIFMRLHQILCLRLLRICS<br>QAERQIEEENREREWEREVLGIKRD<br>KSDSPAIQLRLKEPMDVDVEDYYPA<br>FLDMVRSLLDGNIDSSQYEDSLREM<br>FTIHAYIAFTMDKLIQSIVRQLQHIVS<br>DEVCVQVTDLYLAENNNGATGGQL<br>NSQTSRSLLESAYQRKAEQLMSDEN<br>CFKLMFIQSQGQVQLTVELLDTEEE<br>NSDDPVEAERWSDYVERYMSSDTTS<br>PELREHLAQKPVFLPRNLRRIRKCQ<br>RGREQQEKEGKEGNSKKTMENVES<br>LDKLECRFKLNSYKMVYVIKSEDYM<br>YRRTALLRAHQSHERVSKRLHQRFQ<br>AWVDKWTKEHVPREMAAETSKWL<br>MGEGLEGLVPCTTTCDTETLHFVSI<br>NKYRVKYGTVFKAP | None | None | None | None | None | None |