

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A2ASQ1	AGRIN_MOUSE	Mus musculus	Agtrin	9.761602	NaN	S569;S571	33300544	MPPLPLEHRPRQPGASVLVRYFMI PCNICLILLATSTLGFVLLFLSNYKP GIHFTAAPSMPPDVCRGMLCGFGAV CEPSVEDPGRASCVCCKNVCPAMVA PVCSDASTYSNECELQRAQCNOQ RRIRLLRQGPCSRDPCANVTCSFG STCVPSADGQTASCLCPTTCFGAPD GTVCGSDGVDYPSECQLLRHACANQ EHIFKKFDGPCDPCQGSMSDLNHIC RVNPRTRHPPELLRPENCPAQHTPI CGDDGVTYENDCVMSRIGAARGLL QKVRSGQCQTRDQCPETCQFNVC LSRRGRPHCSCDRVTCGAYRPVCA QDGHTYDNCWRQQAECRQQQTIP PKHQGPCDQTPSPCRGAQCAFGATC TVKNGKAVCECQRVCSSGGYDPVCGS DGVTYGSVCELESMACTLGREIRVA RRGPCDRCGQCRFGSLCEVETGRCV CPSECVESAQPVCSDGHTYASECE LHVHACTHQISLYVASAGHCQTCGE TVCTFGAVCSAGQVCPRCEHPPPG PVCSDGVTYLSACELREAACQQQV QIEEARAGPCEPAECSSGSSGSGED NACEQELCRQHGGVWDEDESDGPC VCDFSCQSVLKSPVCSDGVTYSTE CHLKKARCEARQELYVAAQGACRGP TLAPLLPMASPHCAQTPYGCCQDNV TAAQGVGLAGCPSTCHCNPHGSYSG TCDPVTGQCSCRPGVGGRLRCDRCEP GFWNFRGIVTDGHSGCTPCSCDPRG AVRDDCEQMTGLCSCRPGVAGPKC GQCPDGQALGHLGCEADPTTPVTCV EMHCEFGASCVEEAGFAQCVCPTLT CPEANSTKVCSDGVTYGNQCQLKT IACRQRDLISIQSLGPCRESVAPGVSP TSASMTTPRHILSRTLASPHSSLPLS PSTTAHDWPTPLPTSPQTVVGTTPRST AATPSDVAFLATAIFRESGSTNGSGD EELSGDEEASGGGSGGLEPPVGSVV VTHGPPIERASCYNSPLGCCSDGKTP SLDSEGSNCPATKAFQGVLELEGVE GQELFYTPEMADPKSELFGETARSIE STLDDLFRNSDVKKDFWSIRLRELG PGKLVRAIVDVHFDPTTAFQAPDVG QALLQQIQVSRPWALAVRRPLREHV RFLDFDWFPTFFTGAATGTAAVAT ARATTVSRLSASSVTPRVPSYTSRP VGRTTAPLTTTRPPTTTASIDRPTP GPQRPPKSCDSQPCLHGGTCQDLDS GKGFSCSCTAGRAGTVCEKVQLPSV

PAFKGHSFLAFPTLRAYHTLRLALEF  
RALETEGLLLYNGNARGKDFLALAL  
LDGHVQFRFDTGSGPAVLTSVPVE  
PGRWHRLELSRHRWRQGTLSVDGEA  
PVVGESPSGTDGLNLDTKLYVGGLP  
EEQVATVLDRTSVGIGLKGCIIRMLDI  
NNQLELSDWQRAVVQSSGVGECG  
DHPCSPNPCHGGALCQALEAGVFLC  
QCPPGRFGPTCADEKNPCQPNPCH  
GSAPCHVLSRGGAKCACPLGRSGSF  
CETVLENAGSRPFLADFNGFSYLEL  
KGLHTFERDLGEKMALEMVFLARG  
PSGLLLYNGQKTDGKGFVSLALHN  
RHLEFRYDLGKGAAIIRSKEPIALGT  
WVRVFLERNRKGALQVGDGPRVL  
GESPKSRKVPHTMLNLKEPLYVGGGA  
PDFSKLARGAAVASGFDGAIQLVSLR  
GHQLLTQEHVLRADVAPFAGHPCT  
QAVDNPCLNGGSCIPREATYECLCP  
GGFSGLHCEKGIVEKSVGDLLETAF  
DGRTYIEYLNAVTESELTNEIPAPETL  
DSRALFSEKALQSNHFELSLRTEAT  
QGLVLWIGKVGERADYMALAIVDGH  
LQLSYDLGSQPVLRSTVKVNTNRW  
LRVRAHREHREGSLQVGNEAPVTGS  
SPLGATQLDTDGALWLGGLOKLPVG  
QALPKAYGTGFVGCRLRDVVVGHRL  
HLEDAVTKPELRPCPTL