

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q920R0	ALS2_MOUSE	Mus musculus	Alsin	20.273723	NaN	S459;S460;S477;S486;T504;S1329	22645316;34678516	MDSKKKSSTEAEGSKERGLVHVWQ AGSFSLTPERLPGWGGKTVLQAALG VRHGVLLTEDGEVYSFGTLPWKSES AEICPSSPLLESALVGHVHVITVATGS FHSGAVTESGVVYMWGENAAGQCA VANQQYVPEPSPVSISDSETSPSLAV RILQLACGEEHTLALSLSREIWAWG TGCQLGLITTTFPVTKPQKVEHLAGR VVLQVACGAFHSLALVQCLPPQDLK PVPERCNQCSQLLITMTDKEDHVIIS DSHCCPLGVTLSESQAEKHASPAPS PHPEALDEOGEVFENTVVEAELNM GSSQTTSGSAISTQQNVVGTAEVSSA RTAPSYPDTHAVTAYLQKLSEHSMR ENHEPGEKPPQVQPLVEEAVPDLHS PPTTSTSALNSLVVSCASAVGVRVAA TYEAGALSLKKVMNFYSTAPCETAA QSGSASTGPESLKDLREEQVKQESL QGKKSSSLMDIREEELEGGSRRLSL PGLLSQVSPRLLRKAARVKTRTVVLT PTYSGEADALLPSLRTEVWTWGK GK EGQLGHGDVLPRLQPLCVKCLDGKE VIHLEAGGSHSLALTAKSQVYSWGS NTFGQLGHSEFPTTVPRLSKVSEN GVWSVAAGQDYSFLVDTEDFQPGL YYSGRQDRAEGDTLPENPSGKTPV LLSCSKLGYISRVTAGKDSYLALVDK NIMGYIASLHELASTERRFYSKLSEI KSQILRPLLSLENLGTVTTVQLLQEV ASRFSKLCYLIGQH GASLSSYLQGM KEASSLVIMKHSSLFLDSYTEYCTSV SNFLVMGGFQLLAKPAIDFLNKNQE LLQDLSEVNDENTQLMEILNMLFFL PIRRLHNYAKVLLKLATCFEVTSP EY QKLQDSSSCYESLALHLGKKRKEAE YTLFWKTFPGKMTDSL RKPERRLL CESSNRALSLQHAGRFVSNWFILFN DALVHAQFSTHHVFPLATLWAEPLS EEAGSVNGLKITTP EEQFTLISSTPQ EKT KWLRAISQAVDQALRGTSDFPL YGGGSSVQRQEPPISRSAKYTFYKDT RLKDATYDGRWLSGKPHGRGV LKW PDGKMYSGMFRNGLEDGYGEYRIP NKALNKEDHYVGHWKEGKMCGQG VYSYASGEVFEGCFQDNMRHGHGL LRSGKLTSSSPSMFIGQWVMDKKAG YGVFDDITRGEKYMGMWQDDVCQG NGVVVTQFGLYYEGNFHLNKMMG NGVLLSEDDTIYEGEFSDDWTL SGK

