	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golg appa
0	Q920R0	ALS2_MOUSE	Mus musculus	Alsin	46.837075	S377;S388;S473;S483	S459;S460;S477;S486;T504;S1329	22645316;40885482;40021952;34678516;40997131	MDSKKKSSTEAEGSKERGLVHVWQ AGSFSLTPERLPGWGGKTVLQAALG VRHGVLLTEDGEVYSFGTLPWKSES AEICPSSPLLESALVGHHVITVATGS FHSGAVTESGVVYMWGENAAGQCA VANQQYVPEPSPVSISDSETSPSLAV RILQLACGEEHTLALSLSREIWAWG TGCQLGLITTTFPVTKPQKVEHLAGR VVLQVACGAFHSLALVQCLPPQDLK PVPERCNQCSQLLITMTDKEDHVIIS DSHCCPLGVTLSESQAEKHASPAPS PHPEALDEQGEVFENTVVEAELNM GSSQTTSGSAISTQQNVVGTAEVSSA RTAPSYPDTHAVTAYLQKLSEHSMR ENHEPGEKPPQVQPLVEEAVPDLHS PPTTSTSALNSLVVSCASAVGVRVAA TYEAGALSLKKVMNFYSTAPCETAA QSGSASTGPESLKDLREEQVKQESL QGKKSSSLMDIREEELEGGSRRLSL PGLLSQVSPRLLRKAARVKTRTVVLT PTYSGEADALLPSLRTEVWTWGKGK EGQLGHGDVLPRLQPLCVKCLDGKE VIHLEAGGSHSLALTAKSQVYSWGS NTFGQLGHSEFPTTVPRLSKVSSEN GVWSVAAGQDYSLFLVDTEDFQPGL YYSGRQDRAEGDTLPENPSGTKTPV LLSCSKLGYISRVTAGKDSYLALVDK NIMGYIASLHELASTERRFYSKLSEI KSQILRPLLSLENLGTVTTVQLLQEV ASRFSKLCYLIGQHGASLSSYLQGM KEASSLVIMKHSSLFLDSYTEYCTSV SNFLVMGGFQLLAKPAIDFLNKNQE LLQDLSEVNDENTQLMEILNMLFFL PIRRLHNYAKVLLKLATCFEVTSPEY QKLQDSSSCYESLALHLGKKRKEAE YTLSFWKTFPGKMTDSLRKPERRLL CESSNRALSLQHAGRFSVNWFILFN DALVHAQFSTHHVFPLATLWAEPLS EEAGSVNGLKITTPEEQFTLISSTPQ EKTKWLRAISQAVDQALRGTSDFPL YGGGSSVQRQEPPISRSAKYTFYKDT RLKDATYDGRWLSGKPHGRGVLKW PDGKMYSGMFRNGLEDGYGEYRIP NKALNKEDHYVGHWKEGKMCGQG VYSYASGEVFEGCFQDNMRHGHGL LRSGKLTSSSPSMFIGQWVMDKKAG YGVFDDITRGEKYMGMWQDDVCQG NGVVVTQFGLYYEGNFHLNKMMG NGVLLSEDDTIYEGEFSDDWTLSGK GTLTMPHGDYIEGYFSGEWGSGIKIT GTYFKPSLYESDKDKPKAFRKLGNL AVAADEKWRAVFEECWRQLGCESP GQGEVWKAWDNIAVALTTNRRQHK DSPEILSRSQTQTLESLEYIPQHIGAF SVEKYDDIKKYLIKACDTPLHPLGRL VETLVAVYRMTYVGVGANRRLLQEA VKEIKSYLKRIFQLVRFLFPELPEEGS TIPLSAPLPTGRRSFCTGKSDSRSES PEPGYVVTSSGLLLPVLLPRLYPPLF MLYALDNDREEDIYWECVLRLNKQP DIALLGFLGVQKKFWPATLSILGESK KVLSTTKDACFASAVECLQQISTTFT PSDKLKVIQQTFEEISQSVLASLQED FLWSMDDLFPVFLYVVLRARIRNLG SEVHLIEDLMDPFLQHGEQGIMFTT LKACYFQIQREKLN	True	False	4.661	4.733	1.954	1.666	1.71