

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8BMJ2	SYLC_MOUSE	Mus musculus	Leucine--tRNA ligase, cytoplasmic	23.300923	NaN	S169;S722	28528544;22645316;34887587	MAGRKGTAKVDFLKEIEKEAQQKW EAEKVFEVSASRLEKQKQSSKGYF VTFPYPYMNRLHLGHTFSLSKCEF AVGYQRLKKGKSLFPFGLHCTGMPI KACADKLRKREIELYCPPDFPEEEEE EESSAKPGDIVVRDKAKGKSKAA AKAGSSKYQWDIMKSLGLSDDDIVK FSEAEHWLDYFPPLAVQDLKTIGLK VDWRRSFITTDVNPYYDSFVRWQFL TLRERNKIKFGKRYTIYSPKDGQPCM DHDROTGEGVGPQEYTLVKLVLEP YPSKLSGLKGKNIFLVAATLRPETMF GQTNCWVRPDMKYIGFETANGDIFI CTQRAARNMSYQGFTKHNGVVPVV KELMGEEILGASLSAPLTCYKVYVVL PMLTIKEDKGTGVVTSVPSDSPDDLA ALRDLKKKQALRTKFGIRDDMVLPF EPVPVLEIPGIGNLPAVTVCDELKIQS QNDREKLAEAKEKLYLRGFYDGVML VDGFKGQKIQHVKKTIQKNMIDAGD ALIYMEPEKQVMSRSADECVALCD QWYLDYGDENWKKQTFQCLKNME TFCEESRKNFEASLDWLQEHACSRT YGLGTRLPWDEQWLIESLSDSTIYM AFYTVAHLLQGGDLNGQAESPLGIR PQQMTKDVWDYVFFKDAPFPKTQIP KEKLDQLKQEFEFWYPVDLRASGKD LIPNHLSYYIYNHVAMWPEQSDKWP VSVRANGHLLLNSEKMSKSTGNFLT LSQAVDKFSADGMRLALADAGDTVE DANFVEAMADAGILRLYTWVEWVK EMLASCSSLRSGPADSFNDRVFASE MNAGIIKTDQNYEKMMFKEALKTG FFEFQAAKDKYRELATEGMHRELVF RFIEVQTILLTPFCPHLCEHIWTLLG KPD SIMHASWPVAGPVDESLIRSSQ YLMEVAHDLRLRLKNYMMPAKGKK TDKQPAQRPSHCTIYVAKNYPVWQH ITLTTLRSHFEANNGKLPDNKVIASE LGSLPELKKYMKKVMPFVAMIKEN MEKKGPRVLDLELEFDEQAVLMENI VYLTNSLELEHIEVKFASEAEDKVRE ECCPGKPLNVFRTEPGVPVSLVNPQ PSSGHFSTKIDIRQGDSCESIIRRLM KTDRGIKDLKVKLMRFDDPLLGPR RVPVLGREHSEKTLISENAV FHVLDL VSKKVHLTENGLRTDIGDTMVYLVH