

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|--|--------------|---------------|--|----------------------------|---|
| Q3UH68 | LIMC1_MOUSE | Mus musculus | LIM and calponin homology domains-containing protein 1 | 20.995369 | T506 | S169;S177;S192;S201;S204;S207;T215;S217;S226;S231;S233;S297;S303;T313;S377;T402;S471;T472;S518;S523;T529;T537;S603;S672;S719;T725;S751;S757;S760;T899;S973 | 22517741;36288343;35822049 | MACPALGLEVLQPLQPEPPPEPAFA EAQKWIEQVTGRSFGDKDFRTGLEN GILLCELLNAIKPGLVKKINRLPTPIA GLDNTILFLRGCKELGLKESQLFDPS DLQDTSNRVTVKLNLDYSRKLKKNLVV TIYWLGAANSASYGGTTLNLKEF EGLLAQMRKETDDIDSPKRSIRDSG YIDCWDSERSDSLSPRHGRDSDSFD SLDSFGSRSRQTPSPDVVLRGSSDG RGSDSESDLPHRKLDPVKKDDMSA RRTSHGEPKSAVFPNQYLPNKSNT AYVPAPLRKKKAEREFRKSWSTAT SPLGGERPFRYPRTVPVSDDAESTS MFDMRCEEEAAVLPHSRARQEQLQ LINNQLREEDDKWQDDLARWKSRR RSASQDLIKKEEERKKMEKLMGSED GTSERRKSIKTYREIVQEKERREREL HEAYKNARSQEEAEGILQQYIERFTI SEAVLERLEMPKILERSHSTEPNVSS FPNDPSPMKYLRQQLPPPFTATV ETTIARTSVPESIASAGTGPSKIIITPN TVPMLTPRPYSQPKNSQEVLKTFFKV DGKVMNGEMAPGDEEGKEKEGPA AAAPGPSLTKSQMFEGVATVHDSVP QVKQGSNSIEINIKKPNSAPQELTAA SEETESNGQEDEDGEERP GTGDLEP DSAEPQHFTTTVTRCSPTVALVEFSS NPQLKNEVPEQGQKPEDEMSGKV ELVLSQKVAKPKSPEPEATLTFPFLD KMPETNQLHLPNPSSQADSPSSEKS PGSTPFKFWAWDPEEERRRQEKWQ QEQRLLQERYQKEQDKLKEEWEK AQKEVEEEERRYYEEERKIIEDTVVP FTISSSADQLSTSLSVTEGSGTRNK MDLENC PDKENERRQKTPFQENDG DSSLKTREGGLPEEQSLTPSPSANPE ISVSKGIHQDPQLEAEAGAPHCCTN PQPAQDPPRNQQIPNPPTSTSEDVK PKTLALEKTINHQMESPGERRKSISG KKLCSSCGLTLGKGAAMIETLNLYF HIQCFRCGICKGQLGDAVSGTDVIRIR NGLLNCTDCYMRSRSAGQP TTL |