

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6ZQ58	LARP1_MOUSE	Mus musculus	La-related protein 1	29.834462	T747	S68;S81;S138;S188;S193;T196;S198;S201;S299;S302;T351;S494;S498;T503;S525;S568;S604;S608;T626;T701;S743;T746;T747;S751;T754;T759;T762;T765;T774;S801;T822;S824;S828;S830;S838;T842;S845;S1034;S1065	28135057;25153642;22645316;36852467;34887587;28528544	MATQVEPLLPA GAPLLQAEEHGLAR KKPAPDAQAESGPGDGGGEPDGGV RRPRPACARPRGRDGAERESPRPPAA AEAPAGSDGEDGGRRDFVEAPPPKV NPWTKHAPPPAAVNGQPPPEPSAPA KVVRAAAPKPRKGSKVGDFGDVAVN WPTPGEIAHKSVPQPSHKPQARKL PPKKDMKEQEKGSDGSDSKESPKTKS DESGEEKNGDEDCQRGGQKKKGSK HKWVPLQIDMKPEVPREKLASRPTR PQEPRHTPAVRGEMKGSEPATYMPV SVAPPTPAWQPETKVEPAWHDQDE TSSVKSDGAGGARASFRGRGRGRGR GRGRGRGGTRTHFDYQFGYRKFDG TEGPRTHKYMNNITYYFDNVSSNEI YSMDQELLKDYIKRQIEYYFSVDNLE RDFFLRRKMDADGFLPITLIASFHRV QALTTDISLIFAALKDSKVVEMVEEK VRRREEPEKWPLPGPIVDYSQTDFS QLLNCPEFVPRQHYQKETESAPGSP RAVTPVPTKTEEVSNLKTLPKGLSAS LPDLDESIEWIEVKRPRPSPARPKP EEPRFSHTALPQQLPSSQQLMSKDQ DEQEELDFLFDEEMEOMDGRKNTF TAWSEEDSDYEIDDRDVNKILIVTQT PPYMRRHPGGDRTGNHTSRKMSA ELAKVINDGLFYEQDLWTEKFEPE YSQIKQEVENFKKVNMSREQFDL TPEPPVDPNQEVPPGPPRFQVPTD ALANKLFGAPEPSTIARSLPTTVPEP NYRNARTPRTPTPQLKDSSQTPRF YPVVKEGRTLDAKMPRKRKTRHSSN PPLESHVGWVMSREHRPRTASISS SPSEGTPAVGSYGCTPQSLPKFQHPS HELLKENGFTQHVVYHKYRRRCLNE RKRLGIGQSQEMNTLFRFWSFFLRD HFNKKMYEEFKQLALEDAKEGYRY GLECLFRYYSYGLEKKFRDLDFKDFQ EETVKDYEAGQLYGLEKFWAFLKYS KAKNLDIDPKLQEYLGKFRRLDFR VDPPMGEENHNRHPVAVAGGSSEG RKRCPSQSSSRPATGISQPPTTPTGQ ATREDAKWTSQHSDTLTLRK