



CDC42 (NM_001791.4) - cDNA + Protein - 2026-03-05

GCAGTGCTGC CAACGCCCCG GTGGAGAAGC TGAGGTCATC ATCAGATTTG -35
AAATATTTAA AGTGGATACA AAAC TATTT C AGCAATG CAG ACAATTAAGT 16
MetGln ThrIleLysC 6

GTGTTGTTGT GGGCGATGGT GCTGTTGGTA AAACATGTCT CCTGATATCC 66 I21T
ysValValVa lGlyAspGly AlaValGlyL ysThrCysLe uLeuIleSer 22

TACACAACAA ACAAATTTCC ATCGGAATAT GTACC GACTG TTTT GACAA 116 T23C P34Q
TyrThrThrA snLysPhePr oSerGluTyr ValProThrV alPheAspAs 39

CTATGCAGTC ACAGTTATGA TTGGTGGAGA ACCATATACT CTTGGACTTT 166
nTyrAlaVal ThrValMetI leGlyGlyGl uProTyrThr LeuGlyLeuP 56

TTGATACTGC AGGGCAAGAG GATTATGACA GATTAC GACC GCTGAGTTAT 216 Y64C R66G R68Q
heAspThrAl aGlyGlnGlu AspTyrAspA rgLeuArgPr oLeuSerTyr 72

CCACAAACAG ATGTATTTCT AGTCTGTTTT TCAGTGGTCT CTCCATCTTC 266 C81Y C81E S83P
ProGlnThrA spValPheLe uValCysPhe SerValValS erProSerSe 89

ATTTGAAAAC GTGAAAGAAA AGTGGGTGCC TGAGATAACT CACCACTGTC 316
rPheGluAsn ValLysGluL ysTrpValPr oGluIleThr HisHisCysP 106

CAAAGACTCC TTTCTTGCTT GTTGGGACTC AAATTGATCT CAGAGATGAC 366

roLysThrPr oPheLeuLeu ValGlyThrG lnIleAspLe uArgAspAsp 122

CCCTCTACTA TTGAGAAACT TGCCAAGAAC AAACAGAAGC CTATCACTCC 416

ProSerThrI leGluLysLe uAlaLysAsn LysGlnLysP roIleThrPr 139

AGAGACTGCT GAAAAGCTGG CCCGTGACCT GAAGGCTGTC AAGTATGTGG 466

oGluThrAla GluLysLeuA laArgAspLe uLysAlaVal LysTyrValG 156

AGTGTTC~~TG~~C A~~CT~~TACACAG AAAGGCCTAA AGAATGTATT TGAC~~G~~AAGCA 516 [A159V](#) [E171K](#)

luCysSerAl aLeuThrGln LysGlyLeuL ysAsnValPh eAspGluAla 172

ATATTGGCTG CCCTGGAGCC TCCAGAACCG AAGAAGAGCC GCAGGT~~G~~TGT 566 [R186C](#) [C188Y](#)

IleLeuAlaA laLeuGluPr oProGluPro LysLysSerA rgArgCysVa 189

GCTGCTA~~TGA~~ ACATCTCTCC AGAGCCCTTT CTGCACAGCT GGTGTCGGCA *40 [*192C*24](#)

lLeuLeuSto p

TCATACTAAA AGCAATGTTT AAATCAAAC~~T~~ AAAGATTAAA AATTAAAAAT *90

CGTTTTT~~TG~~CA ATAATGACAA ATGCCCTGCA CCTACCCACA TGCACTCGTG *140

TGAGACAAGG CCCATAGGTA TGGCCCCCCC CTTCCCCCTC CCAGTACTAG *190

TTAATTTTGA GTAATTGTAT TGTCAGAAAA GTGATTAGTA CTATTTTTTTT *240

TTGTTGTTTC AAAAAAAAAA TTTTTGTGTG TGTGTGTTTT TTTTTTTTTT *290

TTTTTTGTTG TTTAAAAGCA AGGCATGCTT GTGGATGACT CTGTAACAGA *340

CTAATTGGAA TTGTTGAAGC TGCTCCCTGG TTCCACTCTG GAGAGTAATC *390

TGGGACATCT TAGTGTTTTG TTTTGTTTTT TTCCCTCCTC TTTTTTTTGG *440

GGGGGAGTGT GTGTGGGGTT TGTTTTTTAG TCTTGTTTTT TTAATTCATT *490

AACCAGTGGT TAGCCCTTAA GGGGAGGAGG ACGGATTGAT TCCACATTCC *540

ACTTCCTAGA TCTAGTTT TAG AAAACATGTT CCCCATCTGG TGCTCTTAGG *590
AAGGAGTATA GTAAATGCCT CATTTAATAA CATACTCCTT TTTGAAAGTT *640
GCCTTTTCTC TCCACCCTTG AGTAGATCCA GTATTTGATG AAACTCATGA *690
AAGTGGGTGG AGCCCATCTT GCCCCTCCTC TTTTCTAGGA CGCACTATAT *740
GTGACTGTGA CTTTCAAGGA CATTGTTTTG CCATTTGCTG ATTTTTTTGG *790
GAAGTTAATT TCTAACTTCT TTCACTGATA AATGAAGAAA AGTATTGCAC *840
CTTTGAAATG CACCAAATGA ATTGAGTTTG TAATTAATAA AATTTTTTTC *890
CCTTTCAGTC ATTGTCTTAT ATGCTTAGCA TAGATTTGCA GCTCAGTAGT *940
ATATGGTGTT CCTAGAATGC AGCTGAAGAC CTGTTATGTA GAGGAAATAC *990
GAGGGGTGGT GCTAGAAGAC AGACATCTGT GGAATGATTC ACATCCTCTC *1040
AAGTTAGGAG GATGGAGGCC TGCTTCATTA AGAAGCTGGG GGTAGGGTGG *1090
GGGTGGGGAG AACACTTAAC AACATGGGGA CCAGTCAGGG GAATCCCCTT *1140
ATTTCTGTTT TGCATATGAG GAACCCTAGA GCAGCCAGGT GAGGCTCTCT *1190
AGTTTAATAA AAATCATGGA AAGACTCTTA ATGCAGACTC TTCTTAAGTG *1240
TTAATAGGGA TTTTTTCAGC TTATTTTGGT TGCAGTTTCC AATTTTTAAA *1290
AATGTTGAGG TAATCTTTCC CACCTTCCCA AACCTAATTC TTGTAGATGC *1340
ATTAGTGTTG AACCAATGCT TTCTCATGTC TCAATCTTT GTATATGCAT *1390
TCTTTTCAGA TGTATTAAC AAACAAAAAC CCTTCACAAG CCAGCCTGAG *1440
GGTTGTTATT TTCCCTCCGT CTCTTACTTT TCAGATATTG AGGAGTGGGA *1490
ATTTGACTCT TGATAACATC AATTTCTAAC AAACTTTGGG ATAAAAATTT *1540
AAAGCTTTAT TATTTTTATT TTCTGGCATT TTAGTTTGAG ATAATATTTA *1590
GCCCTCTATA TGTTCAGGTT TGTGCTTCTT CCTCTAAGTT TGACCCCTCC *1640
CTTAAATACC AAGTAGGTCC ATAATAATTC TTAGTGAAGC AGTGTTCAGG *1690
AAGCTCACTG CTCATGGTGG AAGGAAATGT CAGTGTACCT TTAATCCATA *1740
GATCATTGAA AAGCAGCTCA TTTTCCCCCA AGTTTTCTGC ATCAGTGGTT *1790
CTCCAAC TGG AGTGCAGCTT GAACAATGTG ATTTTTAAGT TTTCCAGGCA *1840
TTTTCTAAAAT GTAGCCAAGT TTGGGAATCA CCACACCTAG TTGGTTATCT *1890
CAAAC TACTA CTATCAGAAT ACAGGTTCTG TGCTGCGAAT CTGAATATGG *1940

GATACATTTT CTTTAGGTAG CAAGAAAGGT TTATTTACCT TTAGGACATT *1990
ATACTGATCT GCATGAGAAT CTTAGAAACT AAGAACCCTT AAAATCAGTT *2040
AAGGTTGGTA CACTGTGGCC TCATTTTGAT AGTTCCTGTC ATGGAGTCAT *2090
GTTTGGCATG ATTTGCATAC TGTGCTAGAA TTAAGCTGGG TTGGTTGAGA *2140
AGGGGTTAGG AAGAGGCAGA TGTAAGCTTT AACAGTTCTA ACCAAGAAAT *2190
GCATCTTATT TATTCCTTGT TGACTATACT AGGGAGAGAT ACCTGGATCA *2240
TGTGATACCC TGGAAAGCAG GGACATATGT ATCCTTAGTG TAGTCAGGGG *2290
CGTATAGGTC CTTGACTATT GCTATATCTT TGTCCATAAG CAATGCTTGA *2340
CATGATATGG CTCTAGAAGT AGTCATTGGA TGGGTATATT ATTTTATAAA *2390
TGAGTAAGTG AAATAAAGCA GGTCATACAG TTGTGTCAAC GAATTATCCG *2440
TGTATCAGAA GAACAAGGAT GATGGGGATT GATTGAAGCT AATGTTTTCT *2490
TAAATATTTT CAGATTGATT TCTGCATGTC AGAAGAACTT GGGTAGCAAG *2540
TGGATGCAGC TTAGGTTTTT CATTAAATA TTGTTTTCTA TTTGACGTAA *2590
CAAAC TTGCT TATAGTCGTG GCTTAGAAGG TGAGTGATAT TCTCCAGGGA *2640
GTATGATTTA GAGGCTGGAA AGGTTATTG AAAATAGTCC TTACAGGCTT *2690
AGTTTGTGTC AGAATTTATA GTATTTAATA CGATGTTAAT ATATTATTGC *2740
TATTTATAAG TGTTTTACTG AACATCCTAG AAATAGATTT TTTTTTTTTT *2790
TTTTTTTTGA GACGGAGTCT CGCTCTGTCG CCCAGGCTGG AGTGCAGTGG *2840
CGCGATCTCG GCTCACTGCA AGCTCCGCCT CCCGGGTTCA CGCCATTCTC *2890
CTGCC TCAGC CTCCCGAGTA GCTGGGACTA CCGGCGCCCG CTACCACGCC *2940
CGGCTAATTT TTTGTATTTT TAGTAGAGAC GGGGTTTCAC CGTGTTAGCC *2990
AGAATGGTCT CGATCTCCTG ACCTCGTGAT CCGCCCGCCT CGGCCTCCCA *3040
AAGTGCTGGG ATTACAGGCG TGAGCCACCG CGCCCGGCCA GAAATAGATT *3090
TTATAGTTAA AGCGATGTTT TTCAAGCTGG ACTTTATGCC ACAAGGGGAG *3140
TGTATTCTAA AGGGGAAGAA AAGGATTGTG GTTTAAAAGA ACAATTGTAA *3190
TATTTCTTC ATTCTGAGGT GTATTTTTCT CACATTTTCA TATTGTAATT *3240
AGGATACCTG TTAACATTGG TGTACATTTA ATGTAGTAGT ACTTTTTCTC *3290
CTTTTGCCCT GCAAGTTGTC ACTAAATTTG TGCTATCATA ATGTTTACCT *3340

TCATAGAATG CTTAAATAGT GGTACTTCTT TGACTTGTTA GGTTTGGACT *3390
ATTTTCCTTT TCTTCTTCCA TTTGTGAGGT AAATTAGTGG TTATTTTGGT *3440
GCTGGGTGCT AAGAACTCG CAAAACAAGG GCCCTGCTCA AGAAGGATTG *3490
AAACTTAGTC CTTGCTGGGT ACAAATTTAG AAATTTGTTA TATAACTGTG *3540
GGGCAAGGGA CAACATTTCA CCAGGAAGTA GCAACTCCAT CATCTTTCAG *3590
GAAGGGCTGA TGGTTAGGGC ATCTCTGAAT GATTATTGGA GGGAAAGTGGC *3640
TTTATGTCAC ACTTGTTTTT TGTAAGTCTT ACTCTGTATC TACCCGACTT *3690
CAGATCATGG TGATTTAATT ACATGGCCCA ATCATGTTTT CTTTCCAGGG *3740
CTGCTTGAAT GCTTGAATAC TTTTTTTTTT GAGATGGAGT CTCGTTCTGT *3790
CGCCCAGGCT GGAGTGCAGT GGTGCAATCT CGGCTCATTG CAAGCTCCGC *3840
CTCCCAGGTT CACACCATT CACTGCCTCA GCCTCCCGAG TAGCCAGGAC *3890
TACAGGCATG TGCCACCATG CCTGGCTAAT GTTTTGTATT TTTAGTAGAG *3940
ATAGGGTTTC ACCATGTTAG CCAGGATGGT CTCGATCTCC TGACCTCGTG *3990
ATCCGCCCGC CTTGGCCTCC CAAAGTGCTA GGATTACAGG CGTGAGCCAC *4040
CGCGCCTGGC CAGAATGAAT ACTTTTAATT ACTACTATGA GTCTTTTTTC *4090
AAAGTGCCCT CACCTTAAAA AAAATTCCTT ATTTTATTCA CTTGCCAGAA *4140
TAATGTTGCA TTTGAACCAT GCCAACGGTA GCTTATTAAA ACAGCATTTA *4190
TTTTTCTCAT TCATGGGTCA GCTGAACCTG TCTCTAGCCT TCAGATTGGG *4240
CTCAGGTGGG CTTTGTGTTT GTCATTCTGG AGCCTAGTTC CAGAGAGCAG *4290
TGGCTGCTTG AGTGTGGACT TGCTTATGGT GGGCCAAAGA GGGGCAAGGG *4340
AAAGCGCATT TACACTTCT CGTATTCCAT TGCCAGAGA TGGCATATAG *4390
CCGATCCCAA GTCAGTGCTA TTCATTCAAT TTTTTAAAAA AATTTATTTA *4440
TTTATTTATT TATTTATTTT CAGACGGGGT CTTGCACTGT TGCCCCGGCT *4490
GGAGTGCAGT GGTGGCATCT CCGCTCACTG CAACCTCCAC CTCCCAGGTT *4540
CACATGATTC TCCTGCCTCA GCCTCCTGAG TAGCTGGGAT TACAGGTGTC *4590
TGCCACTGCG CCCGGCTAAT TTTTTGTATT TTTAAGGGAG ACTGGGTGTC *4640
ACTATGTTGG CCAGACAGAT CTCAAACCTC TGACCTCCTG ATCCGCCCTC *4690
CTCAGCTTCC CAAAGTGCTG GGATTATAGG CATGAGTCAC TGCGCCCGGC *4740

CTATTCATTC ATTCTTACAG TGAATTAGTA AATTCAACAA ATAGGTTTTT *4790
TTAAAGCAAT TAAGGCAC TG AGGATATACA GAGAGTAAAA TAGATGTCTC *4840
CTGACTTCTA CTTGGAATTT TCTTTCCCTT ACAGTTTTGA TAGCTGCTTT *4890
CTCTTGAATA ACTATGTCAT CTCTTTCCAC TTTGCATTGC TCCCAAGAGA *4940
GTAATGTAAA AGTTTTATCT GTGCATGCAT ATTCATGACA CATTCAATTTG *4990
TCATTATTTA TCAAAAACCT GTTCCCTTCC CTGGCACTGG AGACATTGGA *5040
GAATGACAGT CTCAAGCAGT GCAGAGATGT AAGCAGGCAG AGGTAAGGAC *5090
TTTGGTCTCA ATAGCCTACA GGGTGCCTCA TCCACTCACT GTATTCTGTG *5140
TGCCGTGTGCC GCGGTGCAGG TGGTCAGTTG GAACCCCTGA TGGGTATCTG *5190
GGCCACTGCC CAGGCCACAC AGAGCCTTTG TGCAAATTAG AAAAGGCCCT *5240
TCCAAAGATA CTTTATGAAC GACTTGTAAT AATTTAGTGA TTTTATTTTC *5290
CAAAGCAAGG CATTGGTTG CTACCTGCTA GAACTGTTAC AGTAAATACA *5340
CAGATCTATT TTGTGAGCAA ATAGTACTGC CTTGGATGTG GTACCCTTGT *5390
GCAGTGCACA ACTTGGACAG TTCATACACA ATGTTGTATT GAGATGATTG *5440
TTATCTACA TTGTATTTAT TTAGATAGTG GGATTGAAAA TAGGCTAAGA *5490
TGGAGAACTG GGTGGGGAAC AATTTGTAAT CCTTTTTAGT TCATATGTCA *5540
CAATCACAGA ATTGAGACTA GCTAGTTTAA ATCCCTGCAT CTTGCACATG *5590
GTGAGGAAAG GTGAGGACAT TTAGATGACC TTTGCCTGTC ACCCACACAGC *5640
TGTATGGTAG CAGAGCTGGG CGTGAAACCT GCATTCCTGG CCTCTCCTTT *5690
TGTGGTGCTT ACAGTGCTGT GTTAGGTCCA GTTTTACCTC TTAAATCTGT *5740
ATTATTACTT TTTTTTCGAG ACCAGATCTT ACTCTGTCGC CCAGGCTGGA *5790
GTGCAGTGGC GTGATCTCGG CTTGCTGCAA TCTCTGCAAT TCTCATGCCT *5840
CAGCCTCCCA AGTAGCTGGG ACTACAGGCG TGAGCCACCG TGCCCTGGCTA *5890
ATTTTTGTAT TTTTAGTAGA GATGGGGTTT CACCATGTTG GCCAGGCTGG *5940
TCTTGAACTC CTGACCTCAG GTGATCCGCC CGTTGCGGCC TCCCTAAGTG *5990
CTGGGATTAT AGGCGTGAGC CACTGCGCCC GGCCTGAAAT GTGTATTCTT *6040
AAGTCCGCCT TTCAAAGTTG AGTACTGCAT TCTTGCGGCT CTACCTAAGT *6090
TTCTCCTAGA ATCTCCCTGC ACTTCTTGTT TTGCTGCCAT CAGGTAATTT *6140

GTGCATTAGA ACTAGCTTGG AACAGAATGG GTAAACTTCT GTGAAGTTAG *6190
ACACTTGCCT CTTAGAGGCA TATCCAGTGA GTCTTTAAAG CCCTACGTTA *6240
TAGAAAAATG TGAAGTATGT CTAATAAAGG AGTACAGTGG AAGGTTTAAA *6290
AGCACATTGC TGATTAGTGG GGTCAGGATG GGAGGAGAGC CCTGATTTTA *6340
ATGCAGTGTC ATAGTCTCAT AGGTGATTTT CCCAAAAGCA AATAAATGCT *6390
CGATGGATTT TGGGTCCACC ATGCATCTCT GAAGTAGAAG TGACAGATTT *6440
TCAAGGCTAA GGTAAGTATA GCATTTTCTC AACCCATTATA CAATAGAGCA *6490
CAGGATCATA ACTGTTCTGT TTGCTCTGAT GCGTCCCGCA GAGCAGTGCT *6540
TTTTCTCGGT GTGAACGTG GATCACTTGT CTCAGTCATC AGGATTCCTG *6590
GACCCTGTAG GATTTACTCT CAATCTTTAG GGGTAGAGAC TAGGAGCCTA *6640
CATTTTGAAC AAGCATCTTT TTATTTTGAT GAAGTTTGCA TTATTACTCC *6690
TAACAACGTG CCCTTTGGCA GGTAGCTTCC ACTTTTCTGG CCTCATTATC *6740
TGTA AAAATTG CATACTTAGT TATCATTGAG GTCTCAAATG CAGTTCGGGT *6790
TCCTGGTTGG TGAACAATTA GATGGGAAGG CTGAAAAAAA AAATACTCCT *6840
GAGATTC TGA ATCTGTAAGT ATGCAATGGG ACCTGAGAAT CAATCTAATT *6890
TCTAAAAGAT TCCCAGATAG AACTTTGTTG AGCTATGTTT AGGGACCAGT *6940
GCACACCTTG GTGATAGAAC CTTTTTGAAA CTTTGAAAAA GTTAAAAGGG *6990
TTCTGTGTAA TGTTACCTAA GAAGCTTATT TTTAAAAAAG GAAAGCAGTG *7040
CTTAGGGCCA TATTGTTTTG AAATAGGATC TGAGCACCAT TTAAATTTTT *7090
ATAAATTATT TTTTAGAGGC ATCACTTCTT TAAAGCCCAT CAGATTAATG *7140
TGCCATTGGT TTGAAAACCA TGACATGAGA TGATTTCTTA AGGTTTTATT *7190
TGTGTCAGAC ATGAATTTTA GAGTATTTCT GGGCTATAGG TCATTACAGC *7240
TTTACTTATT TATTTTTTGG AGACAGGGTC TCACTCTGTC ACCCAGGCTG *7290
GAGTGCAGTG TCGTGGTCTT GGGTCACTGC ATCCTTGACC TCCCAGCTC *7340
AAGCGATCCT CCCGCCTCAG CCTCCCGAGT AGAAGGTACT ACAGGTGTGT *7390
GACACCACGC CTGGCTCATT TTTGCATTTT GTGTAGAGAG GGGGTTTTGC *7440
CATTTTACCC AGGCTGGTCT CGAACTCCTG AGCTGATCCT CCCACCTCGG *7490
CCTCCAACG TGCTGGGATT ACAGGCATTA GCCACTGTGC CCAGCTGGGT *7540

ATGGATTTCT GAGTGGAGGG AATGTGAACA AAGGAGGGCC CAGGCATGTT *7590
TGCAGAGTGG CTGGGGGCCG TGCTTCACCA CGGCATAGGG ACGGGTGCAG *7640
TAGGAAGAAA GGTAGCCTCA AGGCCCCCAT AAATAAATTG GTGCATTAAG *7690
GAAGGAATAT TAATCACATC CAGGAGGGCT CTGCCCTAAC TGCTGAGAAT *7740
CATAAAAGAA ATGGGGACAT ATTCTGGCGA AATAACAGCT TTGTTAATTG *7790
AGCTCTTACT GTGTGCCTTT GCACTGTGTT AAGCACATGG TGTATCACA *7840
CTTGCTACTC ACAACAGCTC TGAGGTTGAT AGTACAGGTA TCTTCTTTTT *7890
ATGGAGCAGG ACTCTGAGAC ATCATAATTG CTTGGTATAT GCAGCCAGAG *7940
TTGGAAATCC TATCTGTCTG ATTTTGGAAA CTGTGTGCTT ACCATTACAT *7990
ATCTTATGTG GAAATAAAAT GTATATGATT AATTACAAAG CTGAAAATCG *8040
TAGTTGAAGC ATCATATAGT GCACAGGGCA AGCAAAGCAG CATAAAAGGC *8090
TTCATCTGTA AGGGTAGGTT TGTTGGGTAG GAGCTTTTTG GGGGAGGTGA *8140
CTTTAGGATC AGTTGTAGCA TGAGAGAATA TGACACCTTC TTCCTTTAGA *8190
GGATCTAAAG GTGTAGCTTT AGAGGATTGT AGGGTGCCTT TGCCCATTGT *8240
GTCATTTGGT CCTTCTTAAA ATTTGAGTGT TTATTCTCTA CATGGCACTC *8290
TTTTAAGCCT TTGTGTGAAT TAACTCAGTC TTCTCAGCAA TCCCCATTT *8340
TTTCATATGT GGAAACTGAG GCATGTTATG ATACAAAGCT TGTCCAAGAA *8390
TTCAGAGCTA GCTTGTCCAA GAATTCAGAG CTAGTAAGTG AGGGAAGTGG *8440
GATTTGAATT TAAGCAGTCT GTCTCTAGAG TCTGCTCAGC TGGCCTTTTT *8490
TTCTTTCTTC TTCTTCTTCT TCTTTTTTTTT TTTTTTTGTA TAATAGGCCT *8540
GTGAGTTGGG AAGAGCAGGG TTTGTTTGGT GAGAAAGTTG CACCTTAGTG *8590
AATACCCAAG GTCCTGAGGA GGCTTGGAGC CAGCTCTGCA ACCTGGCTCC *8640
CTTCTTTTCT TAGCTTGACC CAAGCACAGG GAGATGACTG GGGCTCTGCC *8690
TTCATCTTCT CTGCAAACGG TCAGGGATAC TCATGCAGTT TTTGAGTTTA *8740
AGTAGTAAAC TCCTAAACTT CAAGTCATGT TTGTATTAGT GATTTCACTT *8790
GACCTCAGTA CAACTCTGTA AAAGAGGCAG GGATGGATGA TGTCATTTGT *8840
CAAATGGTAA AACTGAGGCG AAACAACCTG GCATTGTTGG TGTGTGCGA *8890
AGTACATGCA TAGGCTTTGG ATTTGAGTTT GGCTCTGCCA CTAACCTGAT *8940

CTATGTCTTC AGGCAAGTAA CTTAAATGCT CTGAGTGTTG CATTCTCTGT *8990
GAAGTGGAAA TAGCTGTACT TTGCCCAAGA GTTGTGAGAG TGACATGAAT *9040
TGCTATATGT GAGGTTCTTA GCATAGTGCC TGGCAAGTAG AAGGTACTCA *9090
TCTGTGAATG GGGTAGATAG CAGCTGTCTT CATCACTGTC ATCATTGTTG *9140
GCTCACATCA AATCAGGACC AGAAAACCCCT GTTGACTTCT GGGTCTTAAA *9190
CTGCTGTATC AGTGAGGGGC TTCCTCCTCA TACTCATTGC CAGGAGTAGA *9240
GTAAGGCTTT TAATTTTCATC TCTAACCAAC TCTGATAGGG CTGGAAATGT *9290
GTATGGAGTT CCAGACCCAC CATGGGGCCC CAGTGCTAGC TTCTTAGCCC *9340
TCCTTTGTAA GGTTCGCACT AGTTTGATGG GAGAGGGATT TTTTTTGTCT *9390
TGATTCTGAA GCTCTTAAAG AAATGCTCCT TTTTCTCCAT AACTTGAGGA *9440
TTTCACAATT GTATTGGAAT TATTGGTGT T AAGGAGTTAG GTCCATCTCT *9490
TGGCAGCTTG TGAAGGGTGG GCAGGCACCT GTGAATTCTT CACTGGCTTC *9540
TTGTAAGAAT GACTTTACTG GAGGGTGCTG AGTTTGTGAT TATCTCCTCT *9590
GCTGCTAGAA AACTCCGTGA ACCCTGGTAC ATATAGCGTG ATAAATCAAG *9640
TCCTGTTTTT GCTCCTAACT TGCATTATGG TGTGATAAC TATATCTTCA *9690
TTTCTCCCAT GGTAGTAATA ACACGTGTGG AAAGAGCTCT CAGTTGGAAG *9740
TTGAAGATCC AGGTTCTAGT TGAGGCACCA GAGTTTCCTT GGGCAAGTTG *9790
CCATACCTTT TTGGGCCTTG GTTTCCTCAT CTCAATAAAA TGAGTTTCTG *9840
TTC

CDC42 (NM_001791.4) - cDNA + Protein - 2026-03-05

