LACC1 (NM_153218.4) - cDNA - 2026-11-22

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AATTTCGTTG AGGCTCAGGA CGAGGGACGG GGCGGCAAAT TGTCACAGAG -201
GTCGCCGTAG ACCGGTGTGC AACCGAAGGT GAAGAGCGGG AAGGCGAGGA -151
CCGAATCGCC TCTGGTGTGC GTTGTCCGAC AGGAAGTCCC GAATGGGCTG -101
TGAGCCGCCG GTGGAGGTGC CCGGGCAGGT GCAGAGCCAC AGCCTGCGCC -51
CCGTGAGCCG AGGCGGGTGA TTTATTTGGC ATAAAAGTAT TCTTTCAAGG -1
ATGGCAGAAG CTGTTTTGAT TGATCTTTTT GGTTTGAAAT TGAACTCTCA 50    c.3G>A
AAAAAACTGC CATCAGACAT TACTGAAGAC TTTGAATGCT GTCCAATACC 100   c.56_57insA
ACCATGCTGC CAAGGCCAAG TTTCTCTGTA TAATGTGTTG CAGTAACATC 150   C43fs
AGCTATGAAA GGGATGGAGA ACAAGATAAT TGTGAAATAG AAACAAGCAA 200
TGGATTATCA GCTCTCTTGG AAGAATTTGA GATTGTTAGC TGTCCCAGCA 250
TGGCTGCCAC TTTGTATACC ATTAAACAGA AAATTGATGA AAAAAATCTG 300
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TTTTATTGAT CAACTCTTCA CTGATGTTTA CAATTTTGAA TTTGAAGATT 400   c.372del
TGCAAGTGAC TTTTAGGGGA GGGCTTTTTA AACAGTCCAT TGAAATAAAC 450
GTAATCACAG CTCAAGAACT AAGAGGAATT CAGAATGAAA TAGAAACATT 500
TTTGAGAAGT CTGCCAGCAC TGAGAGGAAA ATTAACTATT ATCACTTCTT 550
CTTTGATCCC AGATATTTTC ATACATGGAT TTACTACAAG AACAGGTGGG 600   T195I
ATATCTTATA TACCAACTCT TAGCTCATTC AATCTCTTCA GTAGTTCCAA 650
ACGGAGAGAT CCCAAGGTAG TGGTTCAAGA AAATCTGCGT AGGTTGGCGA 700
ATGCTGCAGG ATTTAATGTG GAGAAATTTT ACCGAATAAA GACTCATCAT 750
TCCAATGACA TCTGGATTAT GGGAAGAAAG GAGCCTGACT CTTATGATGG 800   I254V
AATAACCACA AATCAGAGAG GAGTCACAAT AGCAGCTCTT GGTGCAGACT 850   T276fs A278P C284R
GTATACCGAT AGTTTTTGCA GATCCAGTCA AAAAAGCATG TGGGGTTGCT 900
CACGCTGGTT GGAAAGGTAC TTTGTTGGGT GTTGCTATGG CTACAGTGAA 950
TGCTATGATA GCAGAATATG GCTGCAGTTT GGAAGACATT GTTGTTGTAC 1000 I330del
TTGGACCTTC AGTAGGACCT TGCTGTTTTA CTCTTCCAAG GGAATCAGCA 1050 p.Glu348Ter
GAGGCATTTC ATAATCTTCA TCCTGCATGT GTACAACTAT TTGATTCACC 1100 S366*
AAATCCCTGT ATCGACATCC GTAAAGCCAC AAGGATTCTT CTAGAACAGG 1150 C370Y
GAGGAATTCT TCCACAGAAT ATTCAGGACC AGAACCAAGA TCTCAACCTC 1200
TGTACATCTT GCCATCCTGA CAAGTTTTTC TCCCATGTCC GAGATGGCCT 1250 R414X
TAATTTTGGT ACACAGATTG GCTTCATATC AATTAAAGAA TGAGATACTT *7
GACTGGATTT TTGTATAACT GCTTCCTGCC TCCTTCCAAA CTGACTGCAA *57
GAGAGAAATT TAGCTGTTTG ATTTACTTAA AACCAAATGG ATTACAATGG *107
ATAATTCATC TTTTGGGTAT ATTTTTACTA TTATTCAAAG CCAAATGATT *157
TTCATTTAAT TGTAATAATA ACTGACAAAA ATCAGTATGT TGTAGCTAAT *207
ATGTTTTATG CATGAGAATT ATTCTTAAAG TTTGTTCTCC CTGTTTATTA *257
CACAGATCAG GAATAGATTT GTTCAGTTCA GTATTTATTG GATACCCTCT *307
ATTGGTCAGG CATTGTGTTA AGCATATGTG AATCAAAATG AACACAACTT *357
TTTCCTTTGA GTCTGATACA GTGAAGGAGA TAAACACTTC TACAACTTAA *407
ATTTAATTTT AATAGCAGTA GAAGAGAACA TAAGGAATAG AGGTTAATTT *457
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TACCCAGAAG CAGGATAGAG AAAATATTAC AGAGAAAATC ACATATCACA *507

TGGGCTCGAA AGATGTAGAG GTTTTTGACA AATGAAGAAC AACCATAACA *557

GGTAGAGGGA ACACCATGAA CCAGGGCATG AAACTGAAAG TGCATAACAT *607

ATTCTAGAGA GAGAAGGGTG TGGGCATGAG TTAGGGCTGG AAAAACAGGT *657

TGGAAACAGA TAAGTAAGGG TCTCAAATGC AATGTCAAAG AGCTTGCAGT *707

TTATTTTCCA GGCAATGAGT AGGCAGCCAA AAAAAAAAAA GTAAGGATGT *757

TTTTTTTTTT TTTCCCATGG CATCATATTT AAGAGGATGG ATTTAAATTG *807

TGTGAGACCA AAGCATAGAG ACTAGATAAG AGGCGATCAA AATATTTCAA *857

AAAGAAATAA TGAAGATCCA ATGAAGGAAG TGGAAATTAA AATAGGGAAG *907

AGAGTAGATG GATTAGAGAG ACATTTAAGA GATGGAATCA ATAGATCCTG *957

TTACTAGATA ATGGAAGTAA GAGGTGAGGA AGAGTGGAAA AGTCATTAAT *1007

GACTCTAAGA TTTCTGCTTG GCTGCTTACC AAGATTGGCA ACAAAGGGAG *1057

GGAGAAGGTT TGGAAAAAGA GAGAAGGATA ATGAGTTTGA CTTTACATAG *1107

AATGAAGGGC ATCCAGATAG AAATCTTTGG TTAATAATTA GAAATATAGA *1157

CCTAGAAATT AGGAGGAAAC CTGAGACAGA GACAAATATT TCAAAGCTTA *1207

CAATACAGAG ATGATACCTG ATTCTATTGG AGCAGGTTTG ATCATCTAGG *1257

CAGAAATTAG GATGAGAAAA AAGGAGATCC AATAATACAA CCTTATAGTC *1307

ACAGAAGTAA GAAAAAAAGG GTAGTTGTTT TGAAGAAGCC AGGATAGGTG *1357

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TCAAGGAAGG GACTTTTTTT GGATGAGCAG TTTTGAGTGT GTTTGTCAGT *1607

TAAAGAGAGG AATTAGGTTA GTTTTCATTT GGGAAAAATT GTATATATAT *1657

TTAATGTAAG TTATCACATT GCATCTTAAA AATATTCTTA TTTAATACAT *1707

ATATTTCCTA CATGTATATG TGGTAGCATG ATAGCAAATA ACATTTGTTT *1757

GGTATTTCCA AAGGACTTTC ATGTACATTG CCTCATTTTA CCTTTACAGC *1807

TACTCTGAAA TACACAGGCA TTATCCCTTT TATTCAGCTG AGAAAACTGA *1857

GCTTCATTGA GGTGGAGGTC AAAAATCACA AAATTTGTGA TGAATTAAGA *1907

TTTGAACATA TGTTTTGTGA CTCCAGTTTT CCTTTCAGAT TTTAAAATTA *1957

ATTAAAGGGA TCTTCATTAT ACTTTTATTG TTAACTTTTT GTTAACATAA *2007

TTTATTCATA CATTCAGTGA AAATTTTGTT GAGGTACTGG GACAGGTTAA *2057

AAAATACAGT TGTAGCCCTC AGGATATTTA ATATCCAGTG AAAAGTGACA *2107

GTCAGTAAAC CAACAATCTC AATACTTTGA TATATGTTGT GAGGTTGTGA *2157

TAACCGATTC TTGTTTAGTT TAATTCTATA TCTCCCTTAG ACCAGTGTTA *2207

AATTTAAATA AAACACCTCA TTTTTTCCAA TTCAGGGAAG GCACTAAACA *2257

TAAAGCATAG GATAGAAATG TTGAACTCAT CCAAAATATT ATTTTGTTTA *2307

ATGAAAATGA TGAAGATTAA GGAATACTTC CATGTATTGA GTAAGGTTGA *2357

TAATTTTCTA ATTCTTCACT GTGCATTATT TTGTTTGAAG TTGGTAAATT *2407

TGGAGTATCC TGCAGACACA TTTTGCTTTA TGTACTACAA CATTCTACAA *2457

CCAAATAAAA ATTATTTTGA TTATCTGA