LACC1 (NM_001128303.2) - cDNA + Protein - 2025-04-02

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TTCCCGCCGC CCGGCTTTCT AACCGGGCCC CTAGTTCCCC GCCCCGTGCC -492
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CGGGGCTGCG GGATGCCGAC TCCGCGGACC GCCCAGACCC GGAACTGCTG -242
AGGCAGCAGC GGGCTCGCGG CGCTTGGCTC ATCCCGGGAT TCCCCAGCTC -192
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TACCTCCCTC CCGCCTCCCG GCAGCTGGCA CGAGGGAACC TGGCCGTCAG -92
GTTTCCCCTG GGATCCTGGG ACGGTATCAG GCGGGGAATC TGTGCGGCCG -42
CGGCGAGGTG ATTTATTTGG CATAAAAGTA TTCTTTCAAG GATGGCAGAA 9     c.3G>A
                                              MetAlaGlu 3

GCTGTTTTGA TTGATCTTTT TGGTTTGAAA TTGAACTCTC AAAAAAACTG 59    c.56_57insA
AlaValLeuI leAspLeuPh eGlyLeuLys LeuAsnSerG lnLysAsnCy 20

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sHisGlnThr LeuLeuLysT hrLeuAsnAl aValGlnTyr HisHisAlaA 37

CCAAGGCCAA GTTTCTCTGT ATAATGTGTT GCAGTAACAT CAGCTATGAA 159   C43fs
laLysAlaLy sPheLeuCys IleMetCysC ysSerAsnIl eSerTyrGlu 53
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rAlaLeuLeu GluGluPheG luIleValSe rCysProSer MetAlaAlaT 87

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LysValIleV alProArgHi sArgLysThr LeuMetLysA laPheIleAs 120

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pGlnLeuPhe ThrAspValT yrAsnPheGl uPheGluAsp LeuGlnValT 137

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hrPheArgGl yGlyLeuPhe LysGlnSerI leGluIleAs nValIleThr 153

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AlaGlnGluL euArgGlyIl eGlnAsnGlu IleGluThrP heLeuArgSe 170

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rLeuProAla LeuArgGlyL ysLeuThrIl eIleThrSer SerLeuIleP 187

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TGCCATCCTG ACAAGTTTTT CTCCCATGTC CGAGATGGCC TTAATTTTGG 1259 R414X
CysHisProA spLysPhePh eSerHisVal ArgAspGlyL euAsnPheGl 420

TACACAGATT GGCTTCATAT CAATTAAAGA ATGAGATACT TGACTGGATT *16
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GCAGGATAGA GAAAATATTA CAGAGAAAAT CACATATCAC ATGGGCTCGA *516
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ATAAGTAAGG GTCTCAAATG CAATGTCAAA GAGCTTGCAG TTTATTTTCC *716
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LACC1 (NM_001128303.2) - cDNA + Protein - 2025-04-02