



*PLCG1* (NM\_002660.3) - cDNA - 2026-05-25

GTCTGCCGCC TCAGCCTCAG CCCCAACCTC AGCCGCCGCC GTTGCCTTG -50  
CTCCCGGGCG GTCCTGGCCT GTGCCGCCGC CGCCCCCAGC GTCGGAGCCA 1  
TGGCGGGCGC CGCGTCCCCT TGCGCCAACG GCTGCGGGCC CGGCGCGCCC 51  
TCGGACGCCG AGGTGCTGCA CCTCTGCCGC AGCCTCGAGG TGGGCACCGT 101  
CATGACTTTG TTCTACTCCA AGAAGTCGCA GCGACCCGAG CGGAAGACCT 151  
TCCAGGTCAA GCTGGAGACG CGCCAGATCA CGTGGAGCCG GGGCGCCGAC 201  
AAGATCGAGG GGGCCATTGA CATTTCGTAA ATTAAGGAGA TCCGCCCAGG 251  
GAAGACCTCA CGGGACTTTG ATCGCTATCA AGAGGACCCA GCTTTCGGGC 301  
CGGACCAGTC ACATTGCTTT GTCATTCTCT ATGGAATGGA ATTTCGCCTG 351  
AAAACGCTGA GCCTGCAAGC CACATCTGAG GATGAAGTGA ACATGTGGAT 401  
CAAGGGCTTA ACTTGGCTGA TGGAGGATAC ATTGCAGGCA CCCACACCCC 451  
TGCAGATTGA GAGGTGGCTC CGGAAGCAGT TTTACTCAGT GGATCGGAAT 501  
CGTGAGGATC GTATATCAGC CAAGGACCTG AAGAACATGC TGTCCCAGGT 551  
CAACTACCGG GTCCCAACA TGCGCTTCCT CCGAGAGCGG CTGACGGACC 601  
TGGAGCAGCG CAGCGGGGAC ATCACCTACG GGCAGTTTGC TCAGCTGTAC 651  
CGCAGCCTCA TGTACAGCGC CCAGAAGACG ATGGACCTCC CTTCTTGGGA 701  
AGCCAGTACT CTGAGGGCTG GGGAGCGGCC GGAGCTTTGC CGAGTGTCCC 751  
TTCTTGAGTT CCAGCAGTTC CTTCTTGA CT ACCAGGGGGA GCTGTGGGCT 801  
GTTGATCGCC TCCAGGTGCA GGAGTTCATG CTCAGCTTCC TCCGAGACCC 851  
CTTACGAGAG ATCGAGGAGC CATACTTCTT CCTGGATGAG TTTGTCACCT 901  
TCCTGTTCTC CAAAGAGAAC AGTGTGTGGA ACTCGCAGCT GGATGCAGTA 951  
TGCCCGGACA CCATGAACAA CCCTCTTTCC CACTACTGGA TCTCCTCCTC 1001

GCACAACACG TACCTGACCG GGGACCAGTT CTCCAGTGAG TCCTCCTTGG 1051  
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TGCTGGGACG GCCCGGATGG GATGCCAGTT ATTTACCATG GGCACACCCT 1151  
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CCTTTGTGGC CTCAGAGTAC CCAGTCATCC TGTCCATTGA GGACCACTGC 1251  
AGCATTGCC AGCAGAGAAA CATGGCCCAA TACTTCAAGA AGGTGCTGGG 1301  
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GAGGGCAGTG CCTACGAGGA GGTGCC TACA TCCATGATGT ACTCTGAGAA 1451  
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GGAACGGGAA AGTCCAGCAC TGCCGTATCC ACTCCCGGCA AGATGCTGGG 1851  
ACCCCAAGT TCTTCTTGAC AGACAACCTC GTCTTTGACT CCCTCTATGA 1901  
CCTCATCACG CACTACCAGC AGGTGCCCT GCGCTGTAAT GAGTTTGAGA 1951  
TGCGACTTTC AGAGCCTGTC CCACAGACCA ACGCCACGA GAGCAAAGAG 2001  
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CAGCAAGAGG GCCAGACAGT GATGCTAGGG AACTCGGAGT TCGACAGCCT 2201  
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GGAAACCAAG GCTGAGAAAT ACGTGAACAA GGCCAAAGGC AAGAAGTTCC 3001  
TTCAGTACAA TCGACTGCAG CTCTCCCGCA TCTACCCCAA GGGCCAGCGA 3051  
CTGGATTCCCT CCAACTACGA TCCTTTGCCC ATGTGGATCT GTGGCAGTCA 3101 [S1021F](#)  
GCTTGTGGCC CTCAACTTCC AGACCCCTGA CAAGCCTATG CAGATGAACC 3151  
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ACCATGCGGG ATGAGGCCTT CGACCCCTTT GACAAGAGCA GCCTCCGCGG 3251  
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AGCAGGTGCT GTGCGCCTTG TAGAATGCCG CGAACTGGGT TCTTTGGAAG \*75  
CAGCCCCCTG TGGCGGCCCTT CCGGGTCTCG CAGCCTGAAG CCTGGATTCC \*125  
AGCAGTGAAT GCTAGACAGA AACCAAGCCA TTAATGAGAT GTTATTACTG \*175  
TTTTGGGCCT CCATGCCCCA GCTCTGGATG AAGGCAAAAA CTGTACTGTG \*225  
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CCATCTTGTG GGGCCAGGAC CATGGCCGAA GCCCCTTGA GAGAGAGGCT \*325  
GCCTCAGCCA GTGGCACAGG AGACTCCAAG GAGCTACTGA CATTCCTAAG \*375  
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TGTCCTGTAG CTTTAAACC ACAGCTGGGC AGGGTGAGAA GCTAGATGCC \*475  
CCTGCAGTTT GGCCCTGGAG CCAGGGCAGA GGAATGTAGG GCCTGCATGG \*525  
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TACTCCTCCT GCTCCTCACC CCTGTAGGA AACCTTGGAG AGGAGAGTGG \*725  
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CTGTCAGGAA TGGGGGGCGG GCCAGGGTGG GAAGAGAAGA AATAGCAGAG \*825  
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AAGGTGCCTC ATAGCATAGC CAGCATTCAG CACACACAAA CCTACTGCCC \*1125  
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TCTGACTGCC AAATAAATCA TCCTCATGTC CTTTTTCCTT TGACTTGTAT \*1225  
GCTCTTTCGG GGGCTCAGGA AAGCCTGTTG CATGGGACAT GCTCACTAGA \*1275  
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GAAATTTCTC AGGTTAAGCA GCAAGAGCTG TAACCCCTCC TCTGGGCTAA \*1375  
CAGGAGTTGT GGGTCCACTC TCTCCTGCCC ACCCTCTGAG GGTGTGTCTG \*1425  
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CCCATCCACG TTGGTTAGGA CGTCCTTGGC GTGTTTGTTA GTGTTGACCC \*2025  
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AAAAGAGCAG GCAGGGCCGG GAGAGGGAAG TCAGTGGTAC CAGAAGGTAG \*2625  
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TCTGGCTGAC AGACCAGGA CAAGTAGACT GGGTTCAAAG TGACAGACCT \*2725

TTCAC<sup>1</sup>TTTCA ACAGC<sup>2</sup>TTTGG CTCAGCAGAC ATGTACACAT ACAAAGTAGA \*2775  
GCCTACAAGG TCAGGGGCAT TCTGCCCCCG CCACAGGACT AAAGACTGCC \*2825  
CTGCGGGAAG ATGGCAGGAG CAGTTTCTGA CCTCAGTTGA GTATCTGTGG \*2875  
CCATGAGCAG AAAAGGCAGG GGTCTGCCTC CTGACCAAGC ACATCTTGAA \*2925  
CATCACCTGA GAGCTTGAAC ATCACTAAGG ACCTAGACAC TCACTTGTCT \*2975  
TTCAACTTGA GCCCATCACT CACCATGTGA GTTCTGTTGA GGGTGGGTAG \*3025  
AAAGCAAATA GGTTCAGGTT ATCCCACCAG ACTAACTCGG TGAATGAAAG \*3075  
GATCATGCCT TCTTCACATT TTAATTAAAT GGATCAAGCA CA

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