



*NLRP1* (NM\_033004.4) - cDNA + Protein - 2026-03-01

GTTGACTAGG CGCTGTTCTT GCTGGCTGGT GCCCCAGGGC CTGGAGAGGT -503  
CTGAAGAAAC CTGGGAGCCA GCAGCCCAGG GCTCCACTCT GGGTTCTGAA -453  
AGCCCATTCC CTGCTCTGCG GCTCCTCCCA CCCCACCTCT TCTCAGCCTT -403  
GCAGCTCAAG GGTTGATCTC AGGAGTCCAG GACCCAGGAG AGGGAAGAAT -353  
CTGAGGAACA CAGAACAGTG AGCGTTGCCC ACACCCCATC TCCCGTCACC -303  
ACATCTCCCC TCACCCTCAC CCTCCCTGCC TGGCCCTGGA CCCCATCCCA -253  
GGACCTCCCT ATCAGCTGAC TTCTTCCAGT GTCTTGCAGG CCCCTCTGGG -203  
CTCCTCCCTC CCCTGGCTTT TCCTACCACT CCCCCTCTAT CGGCGTCTAT -153  
CTGTAGGTGC CCTGGGATTT ATAAAAGTGG GTTCCGAATG CTGAATAAGA -103  
GACGGTAAGA GCCAAGGCAA AGGACAGCAC TGTTCCTCTGC CTGCCTGATA -53  
CCCTCACCAC CTGGGAACAT CCCCAGACA CCCTCTTAAC TCCGGGACAG -3  
AGATGGCTGG CCGAGCCTGG GGCCGCTGG CCTGTTACTT GGAGTTCCTG 48 A2V G4R  
MetAlaG1 yGlyAlaTrp GlyArgLeuA laCysTyrLe uGluPheLeu 16  
  
AAGAAGGAGG AGCTGAAGGA GTTCCAGCTT CTGCTCGCCA ATAAAGCGCA 98 E20del K22 L26del F24L  
LysLysGluG luLeuLysG1 uPheGlnLeu LeuLeuAlaA snLysAlaHi 33  
  
CTCCAGGAGC TCTTCGGGTG AGACACCCGC TCAGCCAGAG AAGACGAGTG 148  
sSerArgSer SerSerGlyG luThrProAl aGlnProGlu LysThrSerG 50  
  
GCATGGAGGT GGCTTCGTAC CTGGTGCTC AGTATGGGGA GCAGCGGGCC 198 A54T A59P A66V  
lyMetGluVa lAlaSerTyr LeuValAlaG lnTyrGlyG1 uGlnArgAla 66

TGGGACCTAG CCCTCCATAC CTGGGAGCAG ATGGGGCTGA GGTCACTGTG 248 M77T  
TrpAspLeuA laLeuHisTh rTrpGluGln MetGlyLeuA rgSerLeuCy 83

CGCCCAAGCC CAGGAAGGGG CAGGCCACTC TCCCTCATT C CCTACAGCC 298 G91D  
sAlaGlnAla GlnGluGlyA laGlyHisSe rProSerPhe ProTyrSerP 100

CAAGTGAACC CCACCTGGGG TCTCCCAGCC AACCCACCTC CACCGCAGTG 348 G106R  
roSerGluPr oHisLeuGly SerProSerG lnProThrSe rThrAlaVal 116

CTAATGCCCT GGATCCATGA ATTGCCGGCG GGGTGCACCC AGGGCTCAGA 398 P119L P125L  
LeuMetProT rpIleHisGl uLeuProAla GlyCysThrG lnGlySerGl 133

GAGAAGGGTT TTGAAGACAGC TGCCTGACAC ATCTGGACGC CGCTGGAGAG 448 L137F R138\*  
uArgArgVal LeuArgGlnL euProAspTh rSerGlyArg ArgTrpArgG 150

AAATCTCTGC CTCACTCCTC TACCAAGCTC TTCCAAGCTC CCCAGACCAT 498  
luIleSerAl aSerLeuLeu TyrGlnAlaL euProSerSe rProAspHis 166

GAGTCTCCAA GCCAGGAGTC ACCCAACGCC CCCACATCCA CAGCAGTGCT 548  
GluSerProS erGlnGluSe rProAsnAla ProThrSerT hrAlaValLe 183

GGGGAGCTGG GGATCCCCAC CTCAGCCCAG CCTAGCACCC AGAGAGCAGG 598  
uGlySerTrp GlySerProP roGlnProSe rLeuAlaPro ArgGluGlnG 200

AGGCTCCTGG GACCCAATGG CCTCTGGATG AAACGTCAGG AATTTACTAC 648  
luAlaProGl yThrGlnTrp ProLeuAspG luThrSerGl yIleTyrTyr 216

ACAGAAATCA GAGAAGAGAGA GAGAGAGAAA TCAGAGAAAAG GCAGGCCCCC 698 [E225Nfs\\*13](#)  
ThrGluIleA rgGluArgGl uArgGluLys SerGluLysG lyArgProPr 233

ATGGGCAGCG GTGGTAGGAA CGCCCCACA GGCGCACACC AGCCTACAGC 748 [A244V](#) [T246A](#)  
oTrpAlaAla ValValGlyT hrProProGl nAlaHisThr SerLeuGlnP 250

CCCACCACCA CCCATGGGAG CCTTCTGTGA GAGAGAGCCT CTGTTCCACA 798  
roHisHisHi sProTrpGlu ProSerValA rgGluSerLe uCysSerThr 266

TGGCCCTGGA AAAATGAGGA TTTTAACCAA AAATTCACAC AGCTGCTACT 848 [F274L](#)  
TrpProTrpL ysAsnGluAs pPheAsnGln LysPheThrG lnLeuLeuLe 283

TCTACAAAGA CCTCACCCC GAAGCCAAGA TCCCCTGGTC AAGAGAAGCT 898 [R290G](#)  
uLeuGlnArg ProHisProA rgSerGlnAs pProLeuVal LysArgSerT 300

GGCCTGATTA TGTGGAGGAG AATCGAGGAC ATTTAATTGA GATCAGAGAC 948  
rpProAspTy rValGluGlu AsnArgGlyH isLeuIleGl uIleArgAsp 316

TTATTTGGCC CAGGCCTGGA TACCCAAGAA CCTCGCATAG TCATACTGCA 998  
LeuPheGlyP roGlyLeuAs pThrGlnGlu ProArgIleV alIleLeuGl 333

GGGGGCTGCT GGAATTGGGA AGTCAACACT GGCCAGGCAG GTGAAGGAAG 1048 [S341L](#)  
nGlyAlaAla GlyIleGlyL ysSerThrLe uAlaArgGln ValLysGluA 350

CTGGGGGAG AGGCCAGCTG TATGGGGACC GCTTCCAGCA TGTCTTCTAC 1098 [W351G](#) [G352W](#)  
laTrpGlyAr gGlyGlnLeu TyrGlyAspA rgPheGlnHi sValPheTyr 366

TTCAGCTGCA GAGAGCTGGC CCAGTCCAAG GTGGTGAGTC TCGCTGAGCT 1148

PheSerCysA rgGluLeuAl aGlnSerLys ValValSerL euAlaGluLe 383

CATCGGAAAA GATGGGACAG CCACTCCGGC TCCCATTAGA CAGATCCTGT 1198

uIleGlyLys AspGlyThrA laThrProAl aProIleArg GlnIleLeuS 400

CTAGGCCAGA GCGGCTGCTC TTCATCCTCG ATGGTGTAGA TGAGCCAGGA 1248 I408T D410H

erArgProGl uArgLeuLeu PheIleLeuA spGlyValAs pGluProGly 416

TGGGTCTTGC AGGAGCCGAG TTCTGAGCTC TGTCTGCACT GGAGCCAGCC 1298

TrpValLeuG lnGluProSe rSerGluLeu CysLeuHisT rpSerGlnPr 433

ACAGCCGGCG GATGCACTGC TGGGCAGTTT GCTGGGGAAA ACTATACTTC 1348 S442N

oGlnProAla AspAlaLeuL euGlySerLe uLeuGlyLys ThrIleLeuP 450

CCGAGGCATC CTTCCCTGATC ACGGCTCGGA CCACAGCTCT GCAGAACCTC 1398 R459G R459Q

roGluAlaSe rPheLeuIle ThrAlaArgT hrThrAlaLe uGlnAsnLeu 466

ATTCCTTCTT TGGAGCAGGC ACGTTGGGTA GAGGTCCTGG GTTTCTCTGA 1448 G480E

IleProSerL euGluGlnAl aArgTrpVal GluValLeuG lyPheSerGl 483

GTCCAGCAGG AAGGAATATT TCTACAGATA TTTCACAGAT GAAAGGCAAG 1498

uSerSerArg LysGluTyrP heTyrArgTy rPheThrAsp GluArgGlnA 500

CAATTAGAGC CTTTAGGTTG GTCAAATCAA ACAAAGAGCT CTGGGCCCTG 1548 K511E

laIleArgAl aPheArgLeu ValLysSerA snLysGluLe uTrpAlaLeu 516

TGTCTTGTGC CCTGGGTGTC CTGGCTGGCC TGCACTTGCC TGATGCAGCA 1598 C527Y

CysLeuValP roTrpValSe rTrpLeuAla CysThrCysL euMetGlnGl 533

GATGAAGCGG AAGGAAAAAC TCACACTGAC TTCCAAGACC ACCACAACCC 1648 [R536Q](#)  
nMetLysArg LysGluLysL euThrLeuTh rSerLysThr ThrThrThrL 550

TCTGTCTACA TTACCTTGCC CAGGCTCTCC AAGCTCAGCC ATTGGGACCC 1698 [H553R](#)  
euCysLeuHi sTyrLeuAla GlnAlaLeuG lnAlaGlnPr oLeuGlyPro 566

CAGCTCAGAG ACCTCTGCTC TCTGGCTGCT GAGGGCATCT GGCAAAAAAA 1748 [S573C](#) [G578V](#)  
GlnLeuArgA spLeuCysSe rLeuAlaAla GluGlyIleT rpGlnLysLy 583

GACCCTTTTT AGTCCAGATG ACCTCAGGAA GCATGGGTTA GATGGGGCCA 1798  
sThrLeuPhe SerProAspA spLeuArgLy sHisGlyLeu AspGlyAlaI 600

TCATCTCCAC CTTCTTGAAG ATGGGTATTC TTCAAGAGCA CCCCATCCCT 1848 [K606R](#)  
leIleSerTh rPheLeuLys MetGlyIleL euGlnGluHi sProIlePro 616

CTGAGCTACA GCTTCATTCA CCTCTGTTTC CAAGAGTTC TGCAGCAAT 1898 [F626S](#) [F629C](#)  
LeuSerTyrS erPheIleHi sLeuCysPhe GlnGluPheP heAlaAlaMe 633

GTCCATATGTC TTGGAGGATG AGAAGGGGAG AGGTAAACAT TCTAATTGCA 1948 [G644C](#)  
tSerTyrVal LeuGluAspG luLysGlyAr gGlyLysHis SerAsnCysI 650

TCATAGATTT GGAAAAGACG CTAGAAGCAT ATGGAATACA TGGCCTGTTT 1998 [F666L](#)  
leIleAspLe uGluLysThr LeuGluAlaT yrGlyIleHi sGlyLeuPhe 666

GGGGCATCAA CCACACGTTT CCTATTGGGC CTGTAAAGTG ATGAGGGGGA 2048 [T670I](#) [G676S](#) [S679N](#) [D680dup](#)  
GlyAlaSerT hrThrArgPh eLeuLeuGly LeuLeuSerA spGluGlyGl 683

GAGAGAGATG GAGAACATCT TTCACTGCCG GCTGTCTCAG GGGAGGAACC 2098 [M686V](#)  
uArgGluMet GluAsnIleP heHisCysAr gLeuSerGln GlyArgAsnL 700

TGATGCAGTG GGTCCCGTCC CTGCAGCTGC TGCTGCAGCC ACACTCTCTG 2148  
euMetGlnTr pValProSer LeuGlnLeuL euLeuGlnPr oHisSerLeu 716

GAGTCCCTCC ACTGCTTGTA CGAGACTCGG AACAAAACGT TCCTGACACA 2198 [E724K](#) [R726W](#)  
GluSerLeuH isCysLeuTy rGluThrArg AsnLysThrP heLeuThrGl 733

AGTGATGGCC CATTTCGAAG AAATGGGCAT GTGTGTAGAA ACAGACATGG 2248 [M735I](#)  
nValMetAla HisPheGluG luMetGlyMe tCysValGlu ThrAspMetG 750

AGCTCTTAGT GTGCACTTTC TGCATTAAAT TCAGCCGCCA CGTGAAGAAG 2298 [T755N](#) [V764L](#)  
luLeuLeuVa lCysThrPhe CysIleLysP heSerArgHi sValLysLys 766

CTTCAGCTGA TTGAGGGCAG GCAGCACAGA TCAACA TGGA GCCCCACCAT 2348 [W779R](#) [P781T](#)  
LeuGlnLeuI leGluGlyAr gGlnHisArg SerThrTrpS erProThrMe 783

GGTAGTCCTG [TTCAGGTGGG](#) TCCCAGTCAC AGATGCCTAT TGGCAGATTC 2398 [F787](#) [R843del](#) [V790I](#)  
tValValLeu PheArgTrpV alProValTh rAspAlaTyr TrpGlnIleL 800

TCTTCTCCGT CCTCAAGGTC ACCAGAAACC TGAAGGAGC T GGACCTAAGT 2448 [L813P](#)  
euPheSerVa lLeuLysVal ThrArgAsnL euLysGluLe uAspLeuSer 816

GGAAACTCGC TGAGCCACTC TGCAGTGAAG AGTCTTTGTA AGACCCTGAG 2498 [S821R](#) [K826N](#)  
GlyAsnSerL euSerHisSe rAlaValLys SerLeuCysL ysThrLeuAr 833

ACGCCCTCGC [TGCCCTCCTGG](#) AGACCCTGCG GTTGGCTGGC TGTGGCCTCA 2548 [C837Y](#) [R843Q](#)

gArgProArg CysLeuLeuG luThrLeuAr gLeuAlaGly CysGlyLeuT 850

CAGCTGAGGA CTGCAAGGAC CTTGCCTTTG GGCTGAGAGC CAACCAGACC 2598

hrAlaGluAs pCysLysAsp LeuAlaPheG lyLeuArgAl aAsnGlnThr 866

CTGACCGAGC TGGACCTGAG CTTCAATGTG CTCACGGATG CTGGAGCCAA 2648 [E869K](#)

LeuThrGluL euAspLeuSe rPheAsnVal LeuThrAspA laGlyAlaLy 883

ACACCTTTGC CAGAGACTGA GACAGCCGAG CTGCAAGCTA CAGCGACTGC 2698 [P892L](#)

sHisLeuCys GlnArgLeuA rgGlnProSe rCysLysLeu GlnArgLeuG 900

AGCTGGTCAG CTGTGGCCTC ACGTCTGACT GCTGCCAGGA CCTGGCCTCT 2748 [A915T](#)

lnLeuValSe rCysGlyLeu ThrSerAspC ysCysGlnAs pLeuAlaSer 916

GTGCTTAGTG CCAGCCCCAG CCTGAAGGAG CTAGACCTGC AGCAGAACAA 2798

ValLeuSerA laSerProSe rLeuLysGlu LeuAspLeuG lnGlnAsnAs 933

CCTGGATGAC GTTGGCGTGC GACTGCTCTG TGAGGGGCTC AGGCATCCTG 2848 [D935N](#) [R940\\*](#) [R947S](#)

nLeuAspAsp ValGlyValA rgLeuLeuCy sGluGlyLeu ArgHisProA 950

CCTGCAAACCT CATA CGCCTG GGGCTGGACC AGACAACTCT GAGTGATGAG 2898 [T961K](#)

laCysLysLe uIleArgLeu GlyLeuAspG lnThrThrLe uSerAspGlu 966

ATGAGGCAGG AACTGAGGGC CCTGGAGCAG GAGAAACCTC AGCTGCTCAT 2948 [M967K](#)

MetArgGlnG luLeuArgAl aLeuGluGln GluLysProG lnLeuLeuIl 983

CTTCAGCAGA CGGAAACCAA GTGTGATGAC CCCTACTGAG GGCCTGGATA 2998

ePheSerArg ArgLysProS erValMetTh rProThrGlu GlyLeuAspT 1000

CGGGAGAGAT GAGTAATAGC ACATCCTCAC TCAAGCGGCA GAGACTCGGA 3048

hrGlyGluMe tSerAsnSer ThrSerSerL euLysArgGl nArgLeuGly 1016

TCAGAGAGGG CGGCTTCCCA TGTTGCTCAG GCTAATCTCA AACTCCTGGA 3098

SerGluArgA laAlaSerHi sValAlaGln AlaAsnLeuL ysLeuLeuAs 1033

CGTGAGCAAG ATCTTCCCAA TTGCTGAGAT TGCAGAGGAA AGCTCCCCAG 3148

pValSerLys ilePheProI leAlaGluIl eAlaGluGlu SerSerProG 1050

AGGTAGTACC GGTGGAAGTC TTGTGCGTGC CTTCTCCTGC CTCTCAAGGG 3198

luValValPr oValGluLeu LeuCysValP roSerProAl aSerGlnGly 1066

GACCTGCATA CGAAGCCTTT GGGACTGAC GATGACTTCT GGGCCCCAC 3248

AspLeuHisT hrLysProLe uGlyThrAsp AspAspPheT rpGlyProTh 1083

GGGGCCTGTG GCTACTGAGG TAGTTGACAA AGAAAAGAAC TTGTACCGAG 3298

rGlyProVal AlaThrGluV alValAspLy sGluLysAsn LeuTyrArgV 1100

TTCACTTCCC TGTAGCTGGC TCCTACCGCT GGCCCAACAC GGGTCTCTGC 3348 [F1102L](#) [G1106D](#) [R1109C](#) [R1109H](#)

alHisPhePr oValAlaGly SerTyrArgT rpProAsnTh rGlyLeuCys 1116

TTTGTGATGA GAGAAGCGGT GACCGTTGAG ATTGAATTCT GTGTGTGGGA 3398

PheValMetA rgGluAlaVa lThrValGlu ileGluPheC ysValTrpAs 1133

CCAGTTCCTG GGTGAGATCA ACCCACAGCA CAGCTGGATG GTGGCAGGGC 3448 [Q1134H](#)

pGlnPheLeu GlyGluIleA snProGlnHi sSerTrpMet ValAlaGlyP 1150

CTCTGCTGGA CATCAAGGCT GAGCCTGGAG CTGTGGAAGC TGTGCACCTC 3498

roLeuLeuAs pIleLysAla GluProGlyA laValGluAl aValHisLeu 1166

CCTCACTTTG TGGCTCTCCA AGGGGGCCAT GTGGACACAT CCCTGTTCCA 3548 [A1171P](#) [G1175R](#)

ProHisPheV alAlaLeuGl nGlyGlyHis ValAspThrs erLeuPheGl 1183

AATGCCCCAC TTAAAGAGG AGGGGATGCT CCTGGAGAAG CCAGCCAGGG 3598 [E1190K](#)

nMetAlaHis PheLysGluG luGlyMetLe uLeuGluLys ProAlaArgV 1200

TGGAGCTGCA TCACATAGTT CTGGAAAACC CCAGCTTCTC CCCCTTGGGA 3648 [P1214R](#) [P1214L](#)

alGluLeuHi sHisIleVal LeuGluAsnP roSerPheSe rProLeuGly 1216

GTCCCTCCTGA AAATGATCCA TAATGCCCTG CGCTTCATTC CCGTCACCTC 3698

ValLeuLeuL ysMetIleHi sAsnAlaLeu ArgPheIleP roValThrSe 1233

TGTGGTGTGG CTTTACCACC GCGTCCATCC TGAGGAAGTC ACCTTCCACC 3748 [R1240C](#) [V1241I](#)

rValValLeu LeuTyrHisA rgValHisPr oGluGluVal ThrPheHisL 1250

TCTACCTGAT CCAAGTGAC TGCTCCATTC GGAAGGCCAT AGATGATCTA 3798 [K1261Rfs\\*3](#) [I1263T](#)

euTyrLeuIl eProSerAsp CysSerIleA rgLysAlaIl eAspAspLeu 1266

GAAATGAAAT TCCAGTTTGT GCGAATCCAC AAGCCACCCC CGCTGACCCC 3848

GluMetLysP heGlnPheVa lArgIleHis LysProProP roLeuThrPr 1283

ACTTTATATG GGCTGTCGTT ACACTGTGTC TGGGTCTGGT TCAGGGATGC 3898 [L1284I](#) [Y1290del](#)

oLeuTyrMet GlyCysArgT yrThrValSe rGlySerGly SerGlyMetL 1300

TGGAAATACT CCCCAAGGAA CTGGAGCTCT GCTATCGAAG CCCTGGAGAA 3948 [E1308Q](#)

euGluIleLe uProLysGlu LeuGluLeuC ysTyrArgSe rProGlyGlu 1316

GACCAGCTGT TCTCGGAGTT CTACGTTGGC CACTTGGGAT CAGGGATCAG 3998  
AspGlnLeuP heSerGluPh eTyrValGly HisLeuGlyS erGlyIleAr 1333

GCTGCAAGTG AAAGACAAGA AAGATGAGAC TCTGGTGTGG GAGGCCTTGG 4048  
gLeuGlnVal LysAspLysL ysAspGluTh rLeuValTrp GluAlaLeuV 1350

TGAAACCAGG AGATCTCATG CCTGCAACTA CTCTGATCCC TCCAGCCCCG 4098  
alLysProGl yAspLeuMet ProAlaThrT hrLeuIlePr oProAlaArg 1366

ATAGCCGTAC CTTACCTCT GGATGCCCCG CAGTTGCTGC ACTTTGTGGA 4148  
IleAlaValP roSerProLe uAspAlaPro GlnLeuLeuH isPheValAs 1383

CCAGTATCGA GAGCAGCTGA TAGCCCGAGT GACATCGGIG GAGGTTGTCT 4198 [Q1384P](#) [R1392Q](#) [V1396A](#)  
pGlnTyrArg GluGlnLeuI leAlaArgVa lThrSerVal GluValValL 1400

TGGACAAACT GCATGGACAG GTGCTGAGCC AGGAGCAGTA CGAGAGGGTG 4248  
euAspLysLe uHisGlyGln ValLeuSerG lnGluGlnTy rGluArgVal 1416

CTGGCTGAGA ACACGAGGCC CAGCCAGATG CGGAAGCTGT TCAGCTTGAG 4298 [N1420K](#) [S1424C](#) [S1431I](#)  
LeuAlaGluA snThrArgPr oSerGlnMet ArgLysLeuP heSerLeuSe 1433

CCAGTCCTGG GACCGGAAGT GCAAAGATGG ACTCTACCAA GCCCTGAAGG 4348  
rGlnSerTrp AspArgLysC ysLysAspGl yLeuTyrGln AlaLeuLysG 1450

AGACCCATCC TCACCTCATT ATGGAAGTCT GGGAGAAGGG CAGCAAAAAG 4398  
luThrHisPr oHisLeuIle MetGluLeuT rpGluLysGl ySerLysLys 1466

GGACTCCTGC CACTCAGCAG CTGAAGTATC AACACCAGCC CTTGACCCTT \*26 G1467R

GlyLeuLeuP roLeuSerSe rStop

GAGTCCTGGC TTTGGCTGAC CCTTCTTTGG GTCTCAGTTT CTTTCTCTGC \*76  
AAACAAGTTG CCATCTGGTT TGCCTTCCAG CACTAAAGTA ATGGAACCTT \*126  
GATGATGCCT TTGCTGGGCA TTATGTGTCC ATGCCAGGGA TGCCACAGGG \*176  
GGCCCCAGTC CAGGTGGCCT AACAGCATCT CAGGGAATGT CCATCTGGAG \*226  
CTGGCAAGAC CCCTGCAGAC CTCATAGAGC CTCATCTGGT GGCCACAGCA \*276  
GCCAAGCCTA GAGCCCTCCG GATCCCATCC AGGCGCAAAG AGGAATAGGA \*326  
GGGACATGGA ACCATTTGCC TCTGGCTGTG TCACAGGGTG AGCCCCAAAA \*376  
TTGGGGTTCA GCGTGGGAGG CCACGTGGAT TCTTGGCTTT GTACAGGAAG \*426  
ATCTACAAGA GCAAGCCAAC AGAGTAAAGT GGAAGGAAGT TTATTCAGAA \*476  
AATAAAGGAG TATCACAGCT CTTTTAGAAT TTGTCTAGCA GGCTTTCCAG \*526  
TTTTTACCAG AAAACCCCTA TAAATTAATA ATTTTTTACT TAAATTTAAG \*576  
AATTAATAAAA ATACAAAAAA GAAAAAATGA AAATAAAGGA ATAAGAAGTT \*626  
ACCTACTCCA

NLRP1 (NM\_033004.4) - cDNA + Protein - 2026-03-01

