

# Infevers - CARD14 (NM\_024110.4) - cDNA + Protein - 2022-05-22

```

GCTCTTCCTT CTGCCAGCT CCGTCCCACC CAGCAGCCCG CAGAGAAAGG -119
AGGCAGCTGG CACCACACTG GGCTTTGGAG ACACTGCGGG GACTGTGGAC -69
CCCACCCTGC TGCACGGAGC TCCTGCAAAA GCAAACCTGA GAACCTTGGG -19
TCCTCCCAGC GCCCAGCCAT GGGGGAAGT TGCCGCAGGG ACTCCGCACT 32
                Me tGlyGluLeu CysArgArgA spSerAlaLe 11

CACGGCACTG GACGAGGAGA CACTGTGGGA GATGATGGAG AGCCACCGCC 82
uThrAlaLeu AspGluGluT hrLeuTrpGl uMetMetGlu SerHisArgH 28

ACAGGATCGT ACGCTGCATC TGCCCCAGCC GCCTCACCCC CTACCTGCGC 132
isArgIleVa lArgCysIle CysProSerA rgLeuThrPr oTyrLeuArg 44

CAGGCCAAGG TGCTGTCCA GCTGGACGAG GAGGAGGTGC TGCACAGCCC 182  C50Y
GlnAlaLysV alLeuCysGl nLeuAspGlu GluGluValL euHisSerPr 61

CCGGCTCACC AACAGCGCCA TGCGGGCCGG GCACTTGCTG GATTGCTGA 232  R62Q R69W R69Q
oArgLeuThr AsnSerAlaM etArgAlaGl yHisLeuLeu AspLeuLeuL 78

AGACTCGAGG GAAGAACGGG GCCATCGCCT TCCTGGAGAG CCTGAAGTTC 282  K78N K93Q
ysThrArgGl yLysAsnGly AlaIleAlaP heLeuGluSe rLeuLysPhe 94

CACAACCCTG ACGTCTACAC CCTGGTCACC GGGCTGCAGC CTGATGTTGA 332  V110A
HisAsnProA spValTyrTh rLeuValThr GlyLeuGlnP roAspValAs 111

CTTCAGTAAC TTTAGCGGTC TCATGGAGAC ATCCAAGCTG ACCGAGTGCC 382  G117S M119V M119R M119T M119K T121I L124P C127S
pPheSerAsn PheSerGlyL euMetGluTh rSerLysLeu ThrGluCysL 128

TGGCTGGGGC CATCGGCAGC CTGCAGGAGG AGCTGAAACCA GGAAAAGGGG 432  E138del E138K E138A
euAlaGlyAl aIleGlySer LeuGlnGluG luLeuAsnGl nGluLysGly 144

CAGAAGGAGG TGCTGCTGCG GCGGTGCCAG CAGCTGCAGG AGCACCTGGG 482  L149R L150R R151W R151Q C153S L156P Q157P
GlnLysGluV alLeuLeuAr gArgCysGln GlnLeuGlnG luHisLeuGl 161

CCTGGCCGAG ACCCGTGCCG AGGGCCTGCA CCAGCTGGAG GCTGACCACA 532  R166H D176H
yLeuAlaGlu ThrArgAlaG luGlyLeuHi sGlnLeuGlu AlaAspHisS 178

GCCGCATGAA GCGTGAGGTT AGCGCACACT TCCATGAGGT GCTGAGGCTG 582  R179H R182C

```

erArgMetLy sArgGluVal SerAlaHisP heHisGluVa lLeuArgLeu 194  
  
AAGGACGAGA TGCTCAGCCT CTCGCTGCAC TATAGCAATG CGCTGCAGGA 632 [E197K](#) [S200N](#) [L209P](#)  
LysAspGluM etLeuSerLe uSerLeuHis TyrSerAsnA laLeuGlnGl 211  
  
GAAGGAGCTG GCCGCCTCAC GCTGCCGAG CCTGCAGGAG GAGCTGTATC 682 [A216T](#) [R218C](#)  
uLysGluLeu AlaAlaSerA rgCysArgSe rLeuGlnGlu GluLeuTyrL 228  
  
TACTGAAGCA GGAGCTGCAG CGAGCCAACA TGGTTTCCTC CTGTGAGCTG 732  
euLeuLysGl nGluLeuGln ArgAlaAsnM etValSerSe rCysGluLeu 244  
  
GAATTGCAAG AGCAGTCCCT GAGGACAGCC AGCGACCAGG AGTCCGGGGA 782 [A254T](#)  
GluLeuGlnG luGlnSerLe uArgThrAla SerAspGlnG luSerGlyAs 261  
  
TGAGGAGCTG AACCGCCTGA AGGAGGAGAA TGAGAAACTG CGCTCGCTGA 832 [R266C](#)  
pGluGluLeu AsnArgLeuL ysGluGluAs nGluLysLeu ArgSerLeuT 278  
  
CTTTCAGCCT GCGGAGAAG GACATTCTGG AGCAGAGCCT GGACGAGGCG 882  
hrPheSerLe uAlaGluLys AspIleLeuG luGlnSerLe uAspGluAla 294  
  
CGGGGAGGC GACAGGAGCT GGTGGAGCGC ATCCACTCGC TCGGGAGCG 932 [R298\\*Stop](#) [R311W](#)  
ArgGlySerA rgGlnGluLe uValGluArg IleHisSerL euArgGluAr 311  
  
GGCCGTGGCT GCCGAGAGGC AGCGAGAGCA GTACTGGGAA GAGAAGGAAC 982 [R319Q](#)  
gAlaValAla AlaGluArgG lnArgGluGl nTyrTrpGlu GluLysGluG 328  
  
AGACCCTGCT GCAGTTCCAG AAGAGTAAGA TGGCCTGCCA ACTCTACAGG 1032 [M338V](#)  
lnThrLeuLe uGlnPheGln LysSerLysM etAlaCysGl nLeuTyrArg 344  
  
GAGAAGGTGA ATGCGCTGCA GGCCCAGGTG TCGGAGCTGC AGAAGGAGCG 1082 [L350P](#) [V354M](#)  
GluLysValA snAlaLeuGl nAlaGlnVal CysGluLeuG lnLysGluAr 361  
  
AGACCAGGCG TACTCCGCGA GGGACAGTGC TCAGAGGGAG ATTTCCGAGA 1132 [A364V](#)  
gAspGlnAla TyrSerAlaA rgAspSerAl aGlnArgGlu IleSerGlnS 378  
  
GCCTGGTGGGA GAAGGACTCC CTCCGAGGC AGGTGTTTCA GCTGACGGAC 1182  
erLeuValGl uLysAspSer LeuArgArgG lnValPheGl uLeuThrAsp 394  
  
CAGGTCTGCG AGCTGCGCAC ACAGCTTCGC CAGCTGCAGG CAGAGCCTCC 1232

GlnValCysG luLeuArgTh rGlnLeuArg GlnLeuGlnA laGluProPr 411  
 GGGTGTGCTC AAGCAGGAAG CCAGGACCAG GGAGCCCTGT CCACGGGAGA 1282 [T420A](#) [E422K](#)  
 oGlyValLeu LysGlnGluA laArgThrAr gGluProCys ProArgGluL 428  
 AGCAGCGGCT GGTGCGGATG CATGCCATCT GCCCAGAGA CGACAGCGAC 1332 [R430W](#)  
 ysGlnArgLe uValArgMet HisAlaIleC ysProArgAs pAspSerAsp 444  
 TGCAGCCTCG TCAGCTCCAC AGAGTCTCAG CTCTTGTCGG ACCTGAGTGC 1382  
 CysSerLeuV alSerSerTh rGluSerGln LeuLeuSerA spLeuSerAl 461  
 CACGTCCAGC CGCGAGCTGG TGGACAGCTT CCGCTCCAGC AGCCCCGCGC 1432  
 aThrSerSer ArgGluLeuV alAspSerPh eArgSerSer SerProAlaP 478  
 CCCCCAGCCA GCAGTCCCTG TACAAGCGGG TGGCCGAGGA CTTCGGGGAA 1482 [P479R](#)  
 roProSerGl nGlnSerLeu TyrLysArgV alAlaGluAs pPheGlyGlu 494  
 GAACCCCTGGT CTTTCAGCAG CTGCCTGGAG ATCCCGGAGG GAGACCCGGG 1532 [c.1488del](#) [P506L](#) [A512Sfs\\*6](#)  
 GluProTrpS erPheSerSe rCysLeuGlu IleProGluG lyAspProGl 511  
 AGCCCTGCCG GGAGCTAAGG CAGGCGACCC ACACCTGGAT TATGAGCTCC 1582 [A512T](#)  
 yAlaLeuPro GlyAlaLysA laGlyAspPr oHisLeuAsp TyrGluLeuL 528  
 TAGACACGGC AGACCTTCCG CAGCTGGAAA GCAGCCTGCA GCCAGTCTCC 1632 [L528P](#)  
 euAspThrAl aAspLeuPro GlnLeuGluS erSerLeuGl nProValSer 544  
 CCTGGAAGGC TTGATGTCTC GGAGAGCGGC GTCCTCATGC GGCGGAGGCC 1682 [R547S](#) [V555I](#)  
 ProGlyArgL euAspValSe rGluSerGly ValLeuMetA rgArgArgPr 561  
 AGCCCGCAGG ATCCTGAGCC AGGTCACCAT GCTGGCGTTC CAGGGGGATG 1732  
 oAlaArgArg IleLeuSerG lnValThrMe tLeuAlaPhe GlnGlyAspA 578  
 CATTGCTGGA GCAGATCAGC GTCATCGGCG GGAACCTCAC GGGCATCTTC 1782 [T591M](#)  
 laLeuLeuGl uGlnIleSer ValIleGlyG lyAsnLeuTh rGlyIlePhe 594  
 ATCCACCGGG TCACCCCGGG CTGGCGGGC GACCAGATGG CCTTGCGCC 1832 [P600L](#) [S602L](#) [A603V](#) [R610H](#)  
 IleHisArgV alThrProGl ySerAlaAla AspGlnMetA laLeuArgPr 611  
 GGGCACCCAG ATTGTGATGG TTGATTACGA AGCCTCAGAG CCCTTGTTCA 1882

oGlyThrGln IleValMetV alAspTyrGl uAlaSerGlu ProLeuPheL 628

AGGCAGTCCT GGAGGACACG ACCCTGGAGG AGGCCGTGGG GCTTCTCAGG 1932 A639G  
ysAlaValLe uGluAspThr ThrLeuGluG luAlaValGl yLeuLeuArg 644

AGGGTGGACG GCTTCTGCTG CCTGTCTGTG AAGGTCAACA CGGACGGTTA 1982 G648S  
ArgValAspG lyPheCysCy sLeuSerVal LysValAsnT hrAspGlyTy 661

TAAGAGGCTA CTCCAGGACC TGGAGGCCAA AGTGGCGACC TCGGGGGACT 2032  
rLysArgLeu LeuGlnAspL euGluAlaLy sValAlaThr SerGlyAspS 678

CATTCTACAT CGGGTCAAC CTGGCCATGG AGGGCAGGGC CAAAGGGGAG 2082 R682W  
erPheTyrIl eArgValAsn LeuAlaMetG luGlyArgAl aLysGlyGlu 694

CTGCAGGTGC ATTGCAACGA GGTCTCTGCAC GTCACCGACA CCATGTTCCA 2132  
LeuGlnValH isCysAsnGl uValLeuHis ValThrAspT hrMetPheGl 711

GGGCTGCGGC TGCTGGCATG CCCACCGCGT GAACTCTTAC ACCATGAAGG 2182 Y724\*Stop  
nGlyCysGly CysTrpHisA laHisArgVa lAsnSerTyr ThrMetLysA 728

ATACTGCCGC GCACGGCACC ATCCCCAACT ACTCCAGGGC TCAGCAGCAG 2232  
spThrAlaAl aHisGlyThr IleProAsnT yrSerArgAl aGlnGlnGln 744

CTCATAGCCC TCATCCAGGA CATGACTCAG CAGTGCACCG TGACCCGCAA 2282  
LeuIleAlaL euIleGlnAs pMetThrGln GlnCysThrV alThrArgLy 761

GCCATCTTCT GGGGGACCAC AGAAGCTGGT CCGCATCGTC AGTATGGACA 2332  
sProSerSer GlyGlyProG lnLysLeuVa lArgIleVal SerMetAspL 778

AAGCCAAGGC CAGCCCTCTG CGTTTGTCTT TTGACAGGGG CCAGTTGGAC 2382  
ysAlaLysAl aSerProLeu ArgLeuSerP heAspArgGl yGlnLeuAsp 794

CCCAGCAGGA TGGAGGGCTC CAGCACGTGC TTCTGGGCCG AGAGCTGCCT 2432  
ProSerArgM etGluGlySe rSerThrCys PheTrpAlaG luSerCysLe 811

CACCCTGGTG CCCTATACCC TGGTGCGGCC CCATCGACCC GCCCGGCCCC 2482 R820W R826W  
uThrLeuVal ProTyrThrL euValArgPr oHisArgPro AlaArgProA 828

GGCCTGTGCT CCTCGTGCCC AGGGCGGTTG GGAAGATCCT GAGCGAGAAA 2532

rgProValLe uLeuValPro ArgAlaValG lyLysIleLe uSerGluLys 844

CTGTGCCTCC TCCAAGGGTT TAAGAAGTGC CTGGCAGAGT ACTTGAGCCA 2582 L847F

LeuCysLeuL euGlnGlyPh eLysLysCys LeuAlaGluT yrLeuSerGl 861

GGAGGAGTAT GAGGCCTGGA GCCAGAGAGG GGACATCATC CAGGAGGGAG 2632

nGluGluTyr GluAlaTrpS erGlnArgGl yAspIleIle GlnGluGlyG 878

AGGTGTCCGG GGGCCGCTGC TGGGTGACCC GCCATGCTGT GGAGTCCCTC 2682 R883H

luValSerGl yGlyArgCys TrpValThrA rgHisAlaVa lGluSerLeu 894

ATGGAAAAGA ACACCCATGC CCTCCTGGAC GTCCAGCTGG ACAGTGTCTG 2732

MetGluLysA snThrHisAl aLeuLeuAsp ValGlnLeuA spSerValCy 911

CACCCTGCAC AGGATGGACA TCTTCCCAT CGTCATCCAC GTCTCTGTCA 2782

sThrLeuHis ArgMetAspI lePheProIl eValIleHis ValSerValA 928

ACGAGAAGAT GGCAAAGAAG CTCAAGAAGG GCCTACAGCG GTTGGGCACC 2832 c.2789\_2791del

snGluLysMe tAlaLysLys LeuLysLysG lyLeuGlnAr gLeuGlyThr 944

TCAGAGGAGC AGCTCCTGGA GGCTGCGAGG CAGGAGGAGG GAGACCTGGA 2882 E957DEL

SerGluGluG lnLeuLeuGl uAlaAlaArg GlnGluGluG lyAspLeuAs 961

CCGGGCGCCC TGTCTATACA GCAGCCTGGC TCCTGACGGC TGGAGCGACC 2932 D973E

pArgAlaPro CysLeuTyrS erSerLeuAl aProAspGly TrpSerAspL 978

TGGACGGCCT GCTCAGCTGT GTCCGCCAGG CCATCGCCGA CGAGCAGAAG 2982

euAspGlyLe uLeuSerCys ValArgGlnA laIleAlaAs pGluGlnLys 994

AAGGTGGTGT GGACGGAGCA GAGCCCCGA TGATGCACCG TGCCCCCTTC \*17

LysValValT rpThrGluGl nSerProArg Stop

CGGGACTGTG GGGGCTTCTG TGTGCCTGTT AATGCAGTCC TGTTCTCTCAG \*67 \*18C>T \*19G>A

CCCAGGCCCT CTTGGCACAG CTGTGGGCTC CTTGGCACAT GAGGCCGGCT \*117

CTCCCCACTG GCTGGGGTCT AACCTTGAAC CCTCACCACG TGCAGGTCAC \*167

ACACAGTGAA GCCACTTGTA ACTGCACACT TTTCTGTGGA AACATCTTCA \*217

CCCTTTACCA GGCTTGGCAT GGTCTGAACT GGAAACCCTG AGAATGTTTC \*267

TGCAGTGGGA CAGGAGGGAC GTCTTCCCAT GCCTTCCCTA GAACCGGAGG \*317

CCCCGGACTT CTCTGGAAAA CCGCCTGTCT GCAGGCCCGA TTCAAATCTA \*367

TGGGGGCTGC ACTTCCCTTT TACATTTTGA TGTGTCAAAG GCTTTTGGAG \*417  
TGACCAAAAG CACAGAGGCA GCGGGTGGGG CGCCTGGGTG GTCCCAAGG \*467  
TCGCTGCCAC CCTTGCCCGG GGCAGAGGCA GAAGCCCACA TATGCTGTGA \*517  
CGCTGGCCAC CTTTTCTCAG CTTCTGAGGC TGCGATGCCT CAGGAACTCC \*567  
AGTTTACAGA GACCAGTGTG TTTACTTGTA AATAAAGCCT CTGCGTGGTG \*617  
GAGACGGTAC TTTTCAGTGGG TCTGTGCCCC GTGGCCCCTG TGCCTGTTTG \*667  
GTGGGGGTGT CCCAGAGAAG CCTGGCACCA GTACCCCCGT ACAAGGCCCA \*717  
GCGGACTCTG CCTTCCCCTG ACCTGGCTTT GCACCCCAGC CCTTCTTGGG \*767  
CCAAACATCT TTAATCCACC TTCAGGGCTC GGGGAGGACC CAGGTCCGCC \*817  
AGCACCTGGC CTTGCCCTG CCTCCTGGGG CTGTTGCAGA CTGAATGTCA \*867  
TTTTGACAGC AGTGTCCAAG AATCAGGAAG CTGTTCTAGA ATTCAGGTTG \*917  
GTATCATCAT AAATGAGTTC AGAAAAAGAA CTTCTGTATA TTTTACTAAA \*967  
ATAAAAAGCT TTTACAATA

**Infevers - CARD14 (NM\_024110.4) - cDNA + Protein - 2022-05-22**