

Infervers - RELA (NM_021975.4) - cDNA + Protein - 2023-02-09

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ATTTCCGCCT CTGGCGAATG GCTCGTCTGT AGTGCACGCC GCGGGCCCAG -35
CTGCGACCCC GGCCCCGCCC CCGGGACCCC GGCCATGGAC GAACTGTTCC 16
                MetAsp GluLeuPheP 6

CCCTCATCTT CCCGGCAGAG CCAGCCCAGG CCTCTGGCCC CTATGTGGAG 66
roLeuIlePh eProAlaGlu ProAlaGlnA laSerGlyPr oTyrValGlu 22

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IleIleGluG lnProLysGl nArgGlyMet ArgPheArgT yrLysCysGl 39

GGGGCGCTCC GCGGGCAGCA TCCCAGGCGA GAGGAGCACA GATACCACCA 166
uGlyArgSer AlaGlySerI leProGlyGl uArgSerThr AspThrThrL 56

AGACCCACCC CACCATCAAG ATCAATGGCT ACACAGGACC AGGGACAGTG 216
ysThrHisPr oThrIleLys IleAsnGlyT yrThrGlyPr oGlyThrVal 72

CGCATCTCCC TGGTCACCAA GGACCCTCCT CACCGGCCTC ACCCCCACGA 266
ArgIleSerL euValThrLy sAspProPro HisArgProH isProHisGl 89

GCTTGTAGGA AAGGACTGCC GGGATGGCTT CTATGAGGCT GAGCTCTGCC 316
uLeuValGly LysAspCysA rgAspGlyPh eTyrGluAla GluLeuCysP 106

CGGACCGCTG CATCCACAGT TTCCAGAACC TGGGAATCCA GTGTGTGAAG 366
roAspArgCy sIleHisSer PheGlnAsnL euGlyIleGl nCysValLys 122

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LysArgAspL euGluGlnAl aIleSerGln ArgIleGlnT hrAsnAsnAs 139

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nProPheGln ValProIleG luGluGlnAr gGlyAspTyr AspLeuAsnA 156

CTGTGCGGCT CTGCTTCCAG GTGACAGTGC GGGACCCATC AGGCAGGCCC 516
laValArgLe uCysPheGln ValThrValA rgAspProSe rGlyArgPro 172

CTCCGCCTGC CGCCTGTCTT TTCTCATCCC ATCTTTGACA ATCGTGCCCC 566
LeuArgLeuP roProValLe uSerHisPro IlePheAspA snArgAlaPr 189

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CAACACTGCC GAGCTCAAGA TCTGCCGAGT GAACCGAAAC TCTGGCAGCT 616
 oAsnThrAla GluLeuLysI leCysArgVa lAsnArgAsn SerGlySerC 206

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 ysLeuGlyGl yAspGluIle PheLeuLeuC ysAspLysVa lGlnLysGlu 222

GACATTGAGG TGTATTTTAC GGGACCAGGC TGGGAGGCC GAGGCTCCTT 716
 AspIleGluV alTyrPheTh rGlyProGly TrpGluAlaA rgGlySerPh 239

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 yrGluThrPh eLysSerIle MetLysLysS erProPheSe rGlyProThr 322

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 aSerValPro LysProAlaP roGlnProTy rProPheThr SerSerLeuS 356

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 nAlaProAla ProAlaProA laProAlaMe tValSerAla LeuAlaGlnA 406

CCCCAGCCCC TGTCCCAGTC CTAGCCCCAG GCCCTCCTCA GGCTGTGGCC 1266 [V410Sfs*4](#)
 laProAlaPr oValProVal LeuAlaProG lyProProGl nAlaValAla 422

CCACCTGCCC CCAAGCCCAC CCAGGCTGGG GAAGGAACGC TGTCAGAGGC 1316
 ProProAlaP roLysProTh rGlnAlaGly GluGlyThrL euSerGluAl 439

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 aLeuLeuGln LeuGlnPheA spAspGluAs pLeuGlyAla LeuLeuGlyA 456

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 snSerThrAs pProAlaVal PheThrAspL euAlaSerVa lAspAsnSer 472

GAGTTTCAGC AGCTGCTGAA CCAGGGCATA CCTGTGGCCC CCCACACAAC 1466 [H487Tfs*7](#)
 GluPheGlnG lnLeuLeuAs nGlnGlyIle ProValAlaP roHisThrTh 489

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 LeuProAsnG lyLeuLeuSe rGlyAspGlu AspPheSerS erIleAlaAs 539

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ATCTGTTGCC AAGCTGG

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