TLR7 (NM_016562.4) - cDNA + Protein - 2026-04-04

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CTGCTCTCTT CAACCAGACC TCTACATTCC ATTTTGGAAG AAGACTAAAA -1
ATGGTGTTTC CAATGTGGAC ACTGAAGAGA CAAATTCTTA TCCTTTTTAA 50
MetValPheP roMetTrpTh rLeuLysArg GlnIleLeuI leLeuPheAs 17

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nIleIleLeu IleSerLysL euLeuGlyAl aArgTrpPhe ProLysThrL 34

TGCCCTGTGA TGTCACTCTG GATGTTCCAA AGAACCATGT GATCGTGGAC 150
euProCysAs pValThrLeu AspValProL ysAsnHisVa lIleValAsp 50

TGCACAGACA AGCATTTGAC AGAAATTCCT GGAGGTATTC CCACGAACAC 200
CysThrAspL ysHisLeuTh rGluIlePro GlyGlyIleP roThrAsnTh 67

CACGAACCTC ACCCTCACCA TTAACCACAT ACCAGACATC TCCCCAGCGT 250
rThrAsnLeu ThrLeuThrI leAsnHisIl eProAspIle SerProAlaS 84

CCTTTCACAG ACTGGACCAT CTGGTAGAGA TCGATTTCAG ATGCAACTGT 300
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ValProIleP roLeuGlySe rLysAsnAsn MetCysIleL ysArgLeuGl 117

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nIleLysPro ArgSerPheS erGlyLeuTh rTyrLeuLys SerLeuTyrL 134

TGGATGGAAA CCAGCTACTA GAGATACCGC AGGGCCTCCC GCCTAGCTTA 450
euAspGlyAs nGlnLeuLeu GluIleProG lnGlyLeuPr oProSerLeu 150

CAGCTTCTCA GCCTTGAGGC CAACAACATC TTTTCCATCA GAAAAGAGAA 500
GlnLeuLeuS erLeuGluAl aAsnAsnIle PheSerIleA rgLysGluAs 167

TCTAACAGAA CTGGCCAACA TAGAAATACT CTACCTGGGC CAAAACTGTT 550
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AGCCGTCCCT ACTGTTTTGC CATCTACTTT AACAGAACTA TATCTCTACA 700
rAlaValPro ThrValLeuP roSerThrLe uThrGluLeu TyrLeuTyrA 234

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snAsnMetIl eAlaLysIle GlnGluAspA spPheAsnAs nLeuAsnGln 250

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PheGluLeuG lnValTyrAr gAlaSerMet AsnLeuSerG lnAlaPheSe 367

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rSerLeuLys SerLeuLysI leLeuArgIl eArgGlyTyr ValPheLysG 384

AGTTGAAAAG CTTTAACCTC TCGCCATTAC ATAATCTTCA AAATCTTGAA 1200

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ValLeuAspL euGlyThrAs nPheIleLys IleAlaAsnL euSerMetPh 417

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SerValGluS erTyrGluPr oGlnValLeu GluGlnLeuH isTyrPheAr 467

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LeuMetMetA snAspAsnAs pIleSerSer SerThrSerA rgThrMetGl 617

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uSerGluSer LeuArgThrL euGluPheAr gGlyAsnHis LeuAspValL 634

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CTA

TLR7 (NM_016562.4) - cDNA + Protein - 2026-04-04