F12 (NM_000505.3) - cDNA + Protein - 2024-12-06

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CTATTGATCT GGACTCCTGG ATAGGCAGCT GGACCAACGG ACGGATGCCA 1
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TGAGGGCTCT GCTGCTCCTG GGGTTCCTGC TGGTGAGCTT GGAGTCAACA 51
etArgAlaLe uLeuLeuLeu GlyPheLeuL euValSerLe uGluSerThr 17

CTTTCGATTC CACCTTGGGA AGCCCCCAAG GAGCATAAGT ACAAAGCTGA 101
LeuSerIleP roProTrpGl uAlaProLys GluHisLysT yrLysAlaGl 34

AGAGCACACA GTCGTTCTCA CTGTCACCGG GGAGCCCTGC CACTTCCCCT 151
uGluHisThr ValValLeuT hrValThrGl yGluProCys HisPheProP 51

TCCAGTACCA CCGGCAGCTG TACCACAAAT GTACCCACAA GGGCCGGCCA 201
heGlnTyrHi sArgGlnLeu TyrHisLysC ysThrHisLy sGlyArgPro 67

GGCCCTCAGC CCTGGTGTGC TACCACCCCC AACTTTGATC AGGACCAGCG 251
GlyProGlnP roTrpCysAl aThrThrPro AsnPheAspG lnAspGlnAr 84

ATGGGGATAC TGTTTGGAGC CCAAGAAAGT GAAAGACCAC TGCAGCAAAC 301
gTrpGlyTyr CysLeuGluP roLysLysVa lLysAspHis CysSerLysH 101

ACAGCCCCTG CCAGAAAGGA GGGACCTGTG TGAACATGCC AAGCGGCCCC 351
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isSerProCy sGlnLysGly GlyThrCysV alAsnMetPr oSerGlyPro 117

CACTGTCTCT GTCCACAACA CCTCACTGGA AACCACTGCC AGAAAGAGAA 401
HisCysLeuC ysProGlnHi sLeuThrGly AsnHisCysG lnLysGluLy 134

GTGCTTTGAG CCTCAGCTTC TCCGGTTTTT CCACAAGAAT GAGATATGGT 451
sCysPheGlu ProGlnLeuL euArgPhePh eHisLysAsn GluIleTrpT 151

ATAGAACTGA GCAAGCAGCT GTGGCCAGAT GCCAGTGCAA GGGTCCTGAT 501
yrArgThrGl uGlnAlaAla ValAlaArgC ysGlnCysLy sGlyProAsp 167

GCCCACTGCC AGCGGCTGGC CAGCCAGGCC TGCCGCACCA ACCCGTGCCT 551
AlaHisCysG lnArgLeuAl aSerGlnAla CysArgThrA snProCysLe 184

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uHisGlyGly ArgCysLeuG luValGluGl yHisArgLeu CysHisCysP 201

CGGTGGGCTA CACCGGAGCC TTCTGCGACG TGGACACCAA GGCAAGCTGC 651
roValGlyTy rThrGlyAla PheCysAspV alAspThrLy sAlaSerCys 217

TATGATGGCC GCGGGCTCAG CTACCGCGGC CTGGCCAGGA CCACGCTCTC 701
TyrAspGlyA rgGlyLeuSe rTyrArgGly LeuAlaArgT hrThrLeuSe 234

GGGTGCGCCC TGTCAGCCGT GGGCCTCGGA GGCCACCTAC CGGAACGTGA 751
rGlyAlaPro CysGlnProT rpAlaSerGl uAlaThrTyr ArgAsnValT 251

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gLeuSerTrp GluTyrCysA spLeuAlaGl nCysGlnThr ProThrGlnA 301

CGGCGCCTCC GACCCCGGTG TCCCCTAGGC TTCATGTCCC ACTCATGCCC 951
laAlaProPr oThrProVal SerProArgL euHisValPr oLeuMetPro 317

GCGCAGCCGG CACCGCCGAA GCCTCAGCCC ACGACCCGGA CCCCGCCTCA 1001
AlaGlnProA laProProLy sProGlnPro ThrThrArgT hrProProGl 334

GTCCCAGACC CCGGGAGCCT TGCCGGCGAA GCGGGAGCAG CCGCCTTCCC 1051
nSerGlnThr ProGlyAlaL euProAlaLy sArgGluGln ProProSerL 351

TGACCAGGAA CGGCCCACTG AGCTGCGGGC AGCGGCTCCG CAAGAGTCTG 1101
euThrArgAs nGlyProLeu SerCysGlyG lnArgLeuAr gLysSerLeu 367

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SerSerMetT hrArgValVa lGlyGlyLeu ValAlaLeuA rgGlyAlaHi 384

CCCCTACATC GCCGCGCTGT ACTGGGGCCA CAGTTTCTGC GCCGGCAGCC 1201
sProTyrIle AlaAlaLeuT yrTrpGlyHi sSerPheCys AlaGlySerL 401

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euIleAlaPr oCysTrpVal LeuThrAlaA laHisCysLe uGlnAspArg 417
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CAGCTGTGAG CCGTGCCAGA CGTTGGCCGT GCGCTCCTAC CGCTTGCACG 1351
sSerCysGlu ProCysGlnT hrLeuAlaVa lArgSerTyr ArgLeuHisG 451

AGGCCTTCTC GCCCGTCAGC TACCAGCACG ACCTGGCTCT GTTGCGCCTT 1401
luAlaPheSe rProValSer TyrGlnHisA spLeuAlaLe uLeuArgLeu 467

CAGGAGGATG CGGACGGCAG CTGCGCGCTC CTGTCGCCTT ACGTTCAGCC 1451
GlnGluAspA laAspGlySe rCysAlaLeu LeuSerProT yrValGlnPr 484

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oValCysLeu ProSerGlyA laAlaArgPr oSerGluThr ThrLeuCysG 501

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lnValAlaGl yTrpGlyHis GlnPheGluG lyAlaGluGl uTyrAlaSer 517

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PheLeuGlnG luAlaGlnVa lProPheLeu SerLeuGluA rgCysSerAl 534

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aProAspVal HisGlySerS erIleLeuPr oGlyMetLeu CysAlaGlyP 551

TCCTCGAGGG CGGCACCGAT GCGTGCCAGG GTGATTCCGG AGGCCCGCTG 1701
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GTGTGTGAGG ACCAAGCTGC AGAGCGCCGG CTCACCCTGC AAGGCATCAT 1751
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ValCysGluA spGlnAlaAl aGluArgArg LeuThrLeuG lnGlyIleIl 584

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eSerTrpGly SerGlyCysG lyAspArgAs nLysProGly ValTyrThrA 601

ATGTGGCCTA CTACCTGGCC TGGATCCGGG AGCACACCGT TTCCTGATTG *3
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CTCAGGGACT CATCTTTCCC TCCTTGGTGA TTCCGCAGTG AGAGAGTGGC *53
TGGGGCATGG AAGGCAAGAT TGTGTCCCAT TCCCCCAGTG CGGCCAGCTC *103
CGCGCCAGGA TGGCGCAGGA ACTCAATAAA GTGCTTTGAA AATGCTGAGA *153
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