# *AP1M2* (NM_005498.5) - cDNA + Protein - 2026-01-28

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TCCCCGACTC CTAAGTCCTT CGGCCGCCAC CATGTCCGCC TCGGCTGTCT 19
                                  MetSerAla SerAlaValP 7

TCATTCTGGA CGTTAAGGGC AAGCCATTGA TCAGCCGCAA CTACAAGGGC 69
heIleLeuAs pValLysGly LysProLeuI leSerArgAs nTyrLysGly 23

GATGTGGCCA TGAGCAAGAT TGAGCACTTC ATGCCTTTGC TGGTACAGCG 119
AspValAlaM etSerLysIl eGluHisPhe MetProLeuL euValGlnAr 40

GGAGGAGGAA GGCGCCCTGG CCCCGCTGCT GAGCCACGGC CAGGTCCACT 169
gGluGluGlu GlyAlaLeuA laProLeuLe uSerHisGly GlnValHisP 57

TCCTATGGAT CAAACACAGC AACCTCTACT TGGTGGCCAC CACATCGAAG 219
heLeuTrpIl eLysHisSer AsnLeuTyrL euValAlaTh rThrSerLys 73

AATGCCAATG CCTCCCTGGT GTACTCCTTC CTGTATAAGA CAATAGAGGT 269
AsnAlaAsnA laSerLeuVa lTyrSerPhe LeuTyrLysT hrIleGluVa 90

ATTCTGCGAA TACTTCAAGG AGCTGGAGGA GGAGAGCATC CGGGACAACT 319
lPheCysGlu TyrPheLysG luLeuGluGl uGluSerIle ArgAspAsnP 107
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heValIleVa lTyrGluLeu LeuAspGluL euMetAspPh eGlyPhePro 123

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GlnThrThrA spSerLysIl eLeuGlnGlu TyrIleThrG lnGlnSerAs 140

CAAGCTGGAG ACGGGCAAGT CACGGGTGCC ACCCACTGTC ACCAACGCTG 469
nLysLeuGlu ThrGlyLysS erArgValPr oProThrVal ThrAsnAlaV 157

TGTCCTGGCG CTCCGAGGGT ATCAAGTATA AGAAGAACGA GGTCTTCATT 519
alSerTrpAr gSerGluGly IleLysTyrL ysLysAsnGl uValPheIle 173

GATGTCATAG AGTCTGTCAA CCTGCTGGTC AATGCCAACG GCAGCGTCCT 569
AspValIleG luSerValAs nLeuLeuVal AsnAlaAsnG lySerValLe 190

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uLeuSerGlu IleValGlyT hrIleLysLe uLysValPhe LeuSerGlyM 207

TGCCAGAGCT GCGGCTGGGC CTCAATGACC GCGTGCTCTT CGAGCTCACT 669
etProGluLe uArgLeuGly LeuAsnAspA rgValLeuPh eGluLeuThr 223

GGCCGCAGCA AGAACAAATC AGTAGAGCTG GAGGATGTAA AATTCCACCA 719
GlyArgSerL ysAsnLysSe rValGluLeu GluAspValL ysPheHisGl 240

GTGCGTGCGG CTCTCTCGCT TTGACAACGA CCGCACCATC TCCTTCATCC 769
nCysValArg LeuSerArgP heAspAsnAs pArgThrIle SerPheIleP 257

CGCCTGATGG TGACTTTGAG CTCATGTCAT ACCGCCTCAG CACCCAGGTC 819
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roProAspGl yAspPheGlu LeuMetSerT yrArgLeuSe rThrGlnVal 273

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LysProLeuI leTrpIleGl uSerValIle GluLysPheS erHisSerAr 290

CGTGGAGATC ATGGTCAAGG CCAAGGGGCA GTTTAAGAAA CAGTCAGTGG 919
gValGluIle MetValLysA laLysGlyGl nPheLysLys GlnSerValA 307

CCAACGGTGT GGAGATATCT GTGCCTGTAC CCAGCGATGC CGACTCCCCC 969
laAsnGlyVa lGluIleSer ValProValP roSerAspAl aAspSerPro 323

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ArgPheLysT hrSerValGl ySerAlaLys TyrValProG luArgAsnVa 340

CGTGATTTGG AGTATTAAGT CTTTCCCGGG GGGCAAGGAG TACTTGATGC 1069
lValIleTrp SerIleLysS erPheProGl yGlyLysGlu TyrLeuMetA 357

GAGCCCACTT TGGCCTCCCC AGTGTGGAAA AGGAAGAGGT GGAGGGCCGG 1119
rgAlaHisPh eGlyLeuPro SerValGluL ysGluGluVa lGluGlyArg 373

CCCCCCATCG GGGTCAAGTT TGAGATCCCC TACTTCACCG TCTCTGGGAT 1169
ProProIleG lyValLysPh eGluIlePro TyrPheThrV alSerGlyIl 390

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eGlnValArg TyrMetLysI leIleGluLy sSerGlyTyr GlnAlaLeuP 407

CCTGGGTTCG CTACATCACC CAGAGTGGCG ATTACCAACT TCGTACCAGC 1269
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Stop

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CTCGCCCTTG GAGCTTTCCC CATCCCCCTG ATTTTATATG AAGAAATAGA *247

AGAGGGGCTT GAAGTCCTCC TCGCGAGTGC CTTCTTGCAA TTACCTGCCT *297

TAGCGGGTGT TGCGGGTCCC TCCTTCACAG CCGCTGAGCC CAGAGGTCCC *347

GCTGGCCCCT CCTCTGAATT TTAGGATGTC ATTAAAAAGA TGAATCTA

## *AP1M2* (NM_005498.5) - cDNA + Protein - 2026-01-28

nfevers