



AP1M2 (NM_005498.5) - cDNA - 2026-06-15

GCTTCCGCAG GAAGAAGGAA GCGGCGCCGC CATCGCCTCC CGGCGCTCCC -32
TCCCCGACTC CTAAGTCCTT CGGCCGCCAC **CATG**TCCGCC TCGGCTGTCT 19
TCATTCCTGGA CGTTAAGGGC AAGCCATTGA TCAGCCGCAA CTACAAGGGC 69
GATGTGGCCA TGAGCAAGAT TGAGCACTTC ATGCCTTTGC TGGTACAGCG 119
GGAGGAGGAA GGCGCCCTGG CCCCCTGTCT GAGCCACGGC CAGGTCCACT 169
TCCTATGGAT CAAACACAGC AACCTCTACT TGGTGGCCAC CACATCGAAG 219
AATGCCAATG CCTCCCTGGT GTACTCCTTC CTGTATAAGA CAATAGAGGT 269
ATTCTGCGAA TACTTCAAGG AGCTGGAGGA GGAGAGCATC CGGGACAACT 319
TTGTCATCGT CTACGAGTTG CTGGACGAGC TCATGGACTT TGGCTTCCCG 369
CAGACCACCG ACAGCAAGAT CCTGCAGGAG TACATCACTC AGCAGAGCAA 419
CAAGCTGGAG ACGGGCAAGT CACGGGTGCC ACCCACTGTC ACCAACGCTG 469
TGTCCGGCG CTCCGAGGGT ATCAAGTATA AGAAGAACGA GGTCCTCATT 519
GATGTCATAG AGTCTGTCAA CCTGCTGGTC AATGCCAACG GCAGCGTCCT 569
TCTGAGCGAA ATCGTCGGTA CCATCAAGCT CAAGGTGTTT CTGTCAGGAA 619
TGCCAGAGCT GCGGCTGGGC CTCAATGACC GCGTGCTCTT CGAGCTCACT 669
GGCCGCAGCA AGAACAAATC AGTAGAGCTG GAGGATGTAA AATTCCACCA 719
GTGCGTGCGG CTCTCTCGCT TTGACAACGA CCGCACCATC TCCTTCATCC 769
CGCCTGATGG TGACTTTGAG CTCATGTCAT ACCGCCTCAG CACCCAGGTC 819
AAGCCACTGA TCTGGATTGA GTCTGTCTATT GAGAAGTTCT CCCACAGCCG 869
CGTGGAGATC ATGGTCAAGG CCAAGGGGCA GTTTAAGAAA CAGTCAGTGG 919
CCAACGGTGT GGAGATATCT GTGCCTGTAC CCAGCGATGC CGACTCCCCC 969
AGATTCAAGA CCAGTGTGGG CAGCGCCAAG TATGTGCCGG AGAGAAACGT 1019

CGTGATTTGG AGTATTAAGT CTTTCCCGGG GGGCAAGGAG TACTTGATGC 1069
GAGCCCACTT TGGCCTCCCC AGTGTGGAAA AGGAAGAGGT GGAGGGCCGG 1119
CCCCCATCG GGGTCAAGTT TGAGATCCCC TACTTCACCG TCTCTGGGAT 1169
CCAGGTCCGA TACATGAAGA TCATTGAGAA AAGTGGTTAC CAGGCCCTGC 1219
CCTGGGTTTCG CTACATCACC CAGAGTGGCG ATTACCAACT TCGTACCAGC 1269
TAGAAGGGAG AAGAGATGGG GGCTTGAACA CGGGGCTTCC TTACAGCCCC *47
GGATGCAGAT TTTAGAGGGA GGCAGGTGC GGGCTGTGTG TGTCTGTGTG *97
AGGGCAGGTC CTGGACTTGG CAGTTTCTTG CTCCAGCAC CCGCCCCTTC *147
CTCACCTCTT CTTTATTCCA TAGGCTGGGA GAGAACTCT CTCTGCTTCC *197
CTCGCCCTTG GAGCTTTCCC CATCCCCCTG ATTTTATATG AAGAAATAGA *247
AGAGGGGCTT GAAGTCCTCC TCGCGAGTGC CTTCTTGCAA TTACCTGCCT *297
TAGCGGGTGT TCGGGTCCC TCCTTCACAG CCGCTGAGCC CAGAGGTCCC *347
GCTGCCCCCT CCTCTGAATT TTAGGATGTC ATTA AAAAGA TGAATCTA

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