

M (1027 bp)

Nature: cRNA

Source: DQ487329.1 Influenza A virus (A/Moscow/10/99(H3N2)) segment 7

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AGCAAAAGCAGGTAGATATTGAAAGATGAGCCTTCTAACCGAGGTCGAAACGTATGTTCTCTCTATCGTT
CCATCAGGCCCCCTCAAAGCCGAAATCGCGCAGAGACTTGAAGATGTCTTTGCTGGGAAAAACACAGATC
TTGAGGCTCTCATGGAATGGCTAAAGACAAGACCAATCCTGTCACCTCTGACTAAGGGGATTTTGGGGTT
TGTGTTTCACGCTCACCGTGCCAGTGAGCGAGGACTGCAGCGTAGACGCTTTGTCCAAAATGCCCTCAAT
GGGAATGGGGATCCAAATAACATGGACAAAGCAGTTAAACTGTATAGAAAACTTAAGAGGGGAGATAACAT
TCCATGGGGCCAAAGAAATAGCACTCAGTTATTCTGCTGGTGCACCTTGCCAGTTGCATGGGCCCTCATATA
CAATAGGATGGGGGCTGTAACCACTGAAGTGGCATTGTCCTGGTATGTGCAACATGTGAACAGATTGCT
GACTCCCAGCACAGGTCTCATAGGCAAATGGTGGCAACAACCAATCCATTAATAAGACATGAGAACAGAA
TGGTTTTTGGCCAGCACTACAGCTAAGGCTATGGAGCAAATGGCTGGATCAAGTGAGCAGGCAGCGGAGGC
CATGGAGATTGCTAGTCAGGCCAGGCAAATGGTGCAGGCAATGAGAGCCGTTGGGACTCATCCTAGCTCC
AGTACTGGTCTAAGAGATGATCTTCTTGAAAATTTGCAGACCTATCAGAAACGAATGGGGGTGCAGATGC
AACGATTCAAGTGACCCGCTTGTTGTTGCCGCGAATATCATTGGGATCTTGCACTTGATATTGTGGATTCT
TTGATCGTCTTTTTTTTCAAATGCATCTATCGACTCTTCAAACACGGCCTTAAAAAGAGGGCCTTCTACGGA
AGGAGTACCTGAGTCTATGAGGGAAGAATATCGAAAGGARCAGCAGAATGCTGTGGATGCTGACGACAGT
CATTTTGTTCAGCATAGAGTTGGAGTAAAAAACTACCTTGTCTTCTACT
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M1 protein

Source: ABE73098.1 (252 aa)

Sequence: 26- 784

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MSLLTEVETYVLSIVPSGPLKAEIAQRLEDVFAGKNTDLEALMEWLKTRPILSPLTKGILGFVFTLTVPSP
ERGLQRRRFVQNALNGNGDPNNMDKAVKLYRKLKREITFHGAKEIALSYSAGALASCMGLIYNRMGAVTT
EVAFGLVCATCEQIADSQHRSHRQMVATTNPLIRHENRMVLASTAKAMEQMAGSSEQAAEAMEIASQAR
QMVQAMRAVGTHPSSSTGLRDDLLENLQTYQKRMGVQMQRFK
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M2 protein

Source: ABE73099.1 (97 aa)

Sequence: 26- 1007

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MSLLTEVETPIRNEWGCRCNDSSDPLVVAANIIGILHLILWILDRLFFKCIYRLFKHGLKRGPSSTEGVPE
SMREEYRKEQQNAVDADDSHFVSIELE
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M (1027 bp)

Nature: vRNA

Source: Illumina sequences from virus stocks_RPS2022

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NNNNNNNGCAGGTAGATATTGAAAGATGAGCCTTCTAACCGAGGTCGAAACGTATGTTCTCT
CTATCGTTCCATCAGGCCCCCTCAAAGCCGAAATCGCGCAGAGACTTGAAGATGTCTTTGCT
GGGAAAAACACAGATCTTGAGGCTCTCATGGAATGGCTAAAGACAAGACCAATCCTGTCACC
TCTGACTAAGGGGATTTTGGGGTTTGTGTTACACGCTCACCGTGCCCAGTGAGCGAGGACTGC
AGCGTAGACGCTTTGTCCAAAATGCCCTCAATGGGAATGGGGATCCAAATAACATGGACAAA
GCAGTTAAACTGTATAGAAAACCTTAAGAGGGAGATAACATTCCATGGGGCCAAAGAAATAGC
ACTCAGTTATTCTGCTGGTGCACCTTGCCAGTTGCATGGGCCTCATATACAATAGGATGGGGG
CTGTAACCACTGAAGTGGCATTGTCCTGGTATGTGCAACATGTGAACAGATTGCTGACTCC
CAGCACAGGTCTCATAGGCAAATGGTGGCAACAACCAATCCATTAATAAGACATGAGAACAG
AATGGTTTTGGCCAGCACTACAGCTAAGGCTATGGAGCAAATGGCTGGATCAAGTGAGCAGG
CAGCGGAGGCCATGGAGATTGCTAGTCAGGCCAGGCAAATGGTGCAGGCAATGAGAGCCGTT
GGGACTCATCCTAGCTCCAGTACTGGTCTAAGAGATGATCTTCTTGAAAATTTGCAGACCTA
TCAGAAACGAATGGGGGTGCAGATGCAACGATTCAAGTGACCCGCTTGTTGTTGCCGCGAAT
ATCATTGGGATCTTGCACTTGATATTGTGGATTCTTGATCGTCTTTTTTTTCAAATGCATCTA
TCGACTCTTCAAACACGGCCTTAAAAGAGGGCCTTCTACGGAAGGAGTACCTGAGTCTATGA
GGGAAGAATATCGAAAGGAACAGCAGAATGCTGTGGATGCTGACGACAGTCATTTTGTGTCAGC
ATAGAGTTGGAGTAAAAAACTACNNNNNNNNNNNN
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M (1027 bp)

Nature: cDNA_pHW2000

Source: GATC sequences from Maxiprep_RPS_2023

AGCAAAAGCAGGTAGATATTGAAAG**ATG**AGCCTTCTAACCGAGGTCGAAACGTATGTTCTCT
CTATCGTTCCATCAGGCCCCCTCAAAGCCGAAATCGCGCAGAGACTTGAAGATGTCTTTGCT
GGGAAAAACACAGATCTTGAGGCTCTCATGGAATGGCTAAAGACAAGACCAATCCTGTCACC
TCTGACTAAGGGGATTTTGGGGTTTGTGTTACGCTCACCGTGCCCAGTGAGCGAGGACTGC
AGCGTAGACGCTTTGTCCAAAATGCCCTCAATGGGAATGGGGATCCAAATAACATGGACAAA
GCAGTTAAACTGTATAGAAAACCTTAAGAGGGAGATAACATTCCATGGGGCCAAAGAAATAGC
ACTCAGTTATTCTGCTGGTGCACCTTGCCAGTTGCATGGGCCTCATATACAATAGGATGGGGG
CTGTAACCACTGAAGTGGCATTGTCCTGGTATGTGCAACATGTGAACAGATTGCTGACTCC
CAGCACAGGTCTCATAGGCAAATGGTGGCAACAACCAATCCATTAATAAGACATGAGAACAG
AATGGTTTTGGCCAGCACTACAGCTAAGGCTATGGAGCAAATGGCTGGATCAAGTGAGCAGG
CAGCGGAGGCCATGGAGATTGCTAGTCAGGCCAGGCAAATGGTGCAGGCAATGAGAGCCGTT
GGGACTCATCCTAGCTCCAGTACTGGTCTAAGAGATGATCTTCTTGAAAATTTGCAGACCTA
TCAGAAACGAATGGGGGTGCAGATGCAACGATTCAAGTGACCCGCTTGTTGTTGCCGCGAAT
ATCATTGGGATCTTGCACTTGATATTGTGGATTCTTGATCGTCTTTTTTTTCAAATGCATCTA
TCGACTCTTCAAACACGGCCTTAAAAGAGGGCCTTCTACGGAAGGAGTACCTGAGTCTATGA
GGGAAGAATATCGAAAGGAACAGCAGAATGCTGTGGATGC**TGA**CGACAGTCATTTTGTGTCAGC
ATAGAGTTGGAGTAAAAAACTACCTTGTTTCTACT

MO_M_Virus	NNNNNNNGCAGGTAGATATTTGAAAGATGAGCCTTCTAACCGAGGTCGAAACGTATGTTCT	60
MO_M_NCBI	AGCAAAAAGCAGGTAGATATTTGAAAGATGAGCCTTCTAACCGAGGTCGAAACGTATGTTCT	60
MO_M_pHW2000	AGCAAAAAGCAGGTAGATATTTGAAAGATGAGCCTTCTAACCGAGGTCGAAACGTATGTTCT	60

MO_M_Virus	CTCTATCGTTCCATCAGGCCCCCTCAAAGCCGAAATCGCGCAGAGACTTGAAGATGTCTT	120
MO_M_NCBI	CTCTATCGTTCCATCAGGCCCCCTCAAAGCCGAAATCGCGCAGAGACTTGAAGATGTCTT	120
MO_M_pHW2000	CTCTATCGTTCCATCAGGCCCCCTCAAAGCCGAAATCGCGCAGAGACTTGAAGATGTCTT	120

MO_M_Virus	TGCTGGGAAAAACACAGATCTTTGAGGCTCTCATGGAATGGCTAAAGACAAGACCAATCCT	180
MO_M_NCBI	TGCTGGGAAAAACACAGATCTTTGAGGCTCTCATGGAATGGCTAAAGACAAGACCAATCCT	180
MO_M_pHW2000	TGCTGGGAAAAACACAGATCTTTGAGGCTCTCATGGAATGGCTAAAGACAAGACCAATCCT	180

MO_M_Virus	GTCACCTCTGACTAAGGGGATTTTGGGGTTTGTGTTACGCTCACCGTGCCAGTGAGCG	240
MO_M_NCBI	GTCACCTCTGACTAAGGGGATTTTGGGGTTTGTGTTACGCTCACCGTGCCAGTGAGCG	240
MO_M_pHW2000	GTCACCTCTGACTAAGGGGATTTTGGGGTTTGTGTTACGCTCACCGTGCCAGTGAGCG	240

MO_M_Virus	AGGACTGCAGCGTAGACGCTTTGTCCAAAATGCCCTCAATGGGAATGGGGATCCAAATAA	300
MO_M_NCBI	AGGACTGCAGCGTAGACGCTTTGTCCAAAATGCCCTCAATGGGAATGGGGATCCAAATAA	300
MO_M_pHW2000	AGGACTGCAGCGTAGACGCTTTGTCCAAAATGCCCTCAATGGGAATGGGGATCCAAATAA	300

MO_M_Virus	CATGGACAAAGCAGTTAAACTGTATAGAAAACTTAAGAGGGAGATAACATTCCATGGGGC	360
MO_M_NCBI	CATGGACAAAGCAGTTAAACTGTATAGAAAACTTAAGAGGGAGATAACATTCCATGGGGC	360
MO_M_pHW2000	CATGGACAAAGCAGTTAAACTGTATAGAAAACTTAAGAGGGAGATAACATTCCATGGGGC	360

MO_M_Virus	CAAAGAAATAGCACTCAGTTATTCTGCTGGTGCACCTTGCCAGTTGCATGGGCCTCATATA	420
MO_M_NCBI	CAAAGAAATAGCACTCAGTTATTCTGCTGGTGCACCTTGCCAGTTGCATGGGCCTCATATA	420
MO_M_pHW2000	CAAAGAAATAGCACTCAGTTATTCTGCTGGTGCACCTTGCCAGTTGCATGGGCCTCATATA	420

MO_M_Virus	CAATAGGATGGGGCTGTAACCACTGAAGTGGCATTTGGCCTGGTATGTGCAACATGTGA	480
MO_M_NCBI	CAATAGGATGGGGCTGTAACCACTGAAGTGGCATTTGGCCTGGTATGTGCAACATGTGA	480
MO_M_pHW2000	CAATAGGATGGGGCTGTAACCACTGAAGTGGCATTTGGCCTGGTATGTGCAACATGTGA	480

MO_M_Virus	ACAGATTGCTGACTCCCAGCACAGGTCTCATAGGCAAATGGTGGCAACAACCAATCCATT	540
MO_M_NCBI	ACAGATTGCTGACTCCCAGCACAGGTCTCATAGGCAAATGGTGGCAACAACCAATCCATT	540
MO_M_pHW2000	ACAGATTGCTGACTCCCAGCACAGGTCTCATAGGCAAATGGTGGCAACAACCAATCCATT	540

MO_M_Virus	AATAAGACATGAGAACAGAATGGTTTTTGGCCAGCACTACAGCTAAGGCTATGGAGCAAAT	600
MO_M_NCBI	AATAAGACATGAGAACAGAATGGTTTTTGGCCAGCACTACAGCTAAGGCTATGGAGCAAAT	600
MO_M_pHW2000	AATAAGACATGAGAACAGAATGGTTTTTGGCCAGCACTACAGCTAAGGCTATGGAGCAAAT	600

MO_M_Virus	GGCTGGATCAAGTGAGCAGGCAGCGGAGGCCATGGAGATTGCTAGTCAGGCCAGGCAAAT	660
MO_M_NCBI	GGCTGGATCAAGTGAGCAGGCAGCGGAGGCCATGGAGATTGCTAGTCAGGCCAGGCAAAT	660
MO_M_pHW2000	GGCTGGATCAAGTGAGCAGGCAGCGGAGGCCATGGAGATTGCTAGTCAGGCCAGGCAAAT	660

MO_M_Virus	GGTGCAGGCAATGAGAGCCGTTGGGACTCATCCTAGCTCCAGTACTGGTCTAAGAGATGA	720
MO_M_NCBI	GGTGCAGGCAATGAGAGCCGTTGGGACTCATCCTAGCTCCAGTACTGGTCTAAGAGATGA	720
MO_M_pHW2000	GGTGCAGGCAATGAGAGCCGTTGGGACTCATCCTAGCTCCAGTACTGGTCTAAGAGATGA	720

MO_M_Virus	TCTTCTTGAAAATTTGCAGACCTATCAGAAACGAATGGGGGTGCAGATGCAACGATTCAA	780
MO_M_NCBI	TCTTCTTGAAAATTTGCAGACCTATCAGAAACGAATGGGGGTGCAGATGCAACGATTCAA	780
MO_M_pHW2000	TCTTCTTGAAAATTTGCAGACCTATCAGAAACGAATGGGGGTGCAGATGCAACGATTCAA	780

MO_M_Virus	GTGACCCGCTTGTTGTTGCCGCGAATATCATTTGGGATCTTGCACTTGATATTGTGGATTC	840
MO_M_NCBI	GTGACCCGCTTGTTGTTGCCGCGAATATCATTTGGGATCTTGCACTTGATATTGTGGATTC	840
MO_M_pHW2000	GTGACCCGCTTGTTGTTGCCGCGAATATCATTTGGGATCTTGCACTTGATATTGTGGATTC	840

MO_M_Virus	TTGATCGTCTTTTTTTTCAAATGCATCTATCGACTCTTCAAACACGGCCTTAAAAAGAGGGC	900
MO_M_NCBI	TTGATCGTCTTTTTTTTCAAATGCATCTATCGACTCTTCAAACACGGCCTTAAAAAGAGGGC	900
MO_M_pHW2000	TTGATCGTCTTTTTTTTCAAATGCATCTATCGACTCTTCAAACACGGCCTTAAAAAGAGGGC	900

MO_M_Virus	CTTCTACGGAAGGAGTACCTGAGTCTATGAGGGAAGAATATCGAAAGGAACAGCAGAAATG	960
MO_M_NCBI	CTTCTACGGAAGGAGTACCTGAGTCTATGAGGGAAGAATATCGAAAGGARACAGCAGAAATG	960
MO_M_pHW2000	CTTCTACGGAAGGAGTACCTGAGTCTATGAGGGAAGAATATCGAAAGGAACAGCAGAAATG	960

MO_M_Virus	CTGTGGATGCTGACGACAGTCATTTTGTGAGCATAGAGTTGGAGTAAAAAACTACNNNNN	1020
MO_M_NCBI	CTGTGGATGCTGACGACAGTCATTTTGTGAGCATAGAGTTGGAGTAAAAAACTACCTTGT	1020
MO_M_pHW2000	CTGTGGATGCTGACGACAGTCATTTTGTGAGCATAGAGTTGGAGTAAAAAACTACCTTGT	1020

MO_M_Virus	NNNNNNN	1027
MO_M_NCBI	TTCTACT	1027
MO_M_pHW2000	TTCTACT	1027