

Nucleotide sequence of the PA gene of influenza A/WSN/33(H1N1)

Takato Odagiri and Kiyotake Tobita

Department of Virology, Jichi Medical School, Minamikawachi-machi, Kawachi-gun, Tochigi-ken
329-04, Japan

Submitted January 9, 1990

EMBL accession no. X17336

The cloned DNA was cut with the appropriate restriction endonucleases and was subcloned into the M13 phage vectors mp18 and mp19 (1). The sequence was determined on both strands of the cDNA by the dideoxynucleotide chain termination method (2) using modified T7 polymerase sequenase. The PA gene of A/WSN contained 2233 nucleotides which can code for 716 amino acids and indicated a high homology (97%) to that of A/PR/8/34(H1N1) (3).

ACKNOWLEDGEMENT

We thank E.Kurokawa, N. Ogawa and K.Uchiyama for their technical assistances. This work was supported by the Naito Foundation, Japan.

REFERENCES

- Norlander, J., Kempe, T. and Messing, J. (1983) *Gene* **26**, 101-106.
- Sanger, F., Nicklen, S. and Coulson, A.R. (1977) *Proc. Natl. Acad. Sci. USA* **74**, 5463-5467.
- Fields, S. and Winter, G. (1982) *Cell* **28**, 303-313.

```

AGCGAAGAGCGUAGUUGUAAAUGGAAAGUUUUGGCGCAAUUGCUUCAAUCCGAGUAGUUGGAGCUU 72
M E D F V R Q C F N P M I V E L (16)
GCGGAAAAGGCAUUGAAGAGUAGGAGGACCGAAAUUGGAAACAAUUGGAGCAUUAUUGCACU 144
A E K A M K E Y G E D L K I E T N K F A A I C T (40)
CACUUGGAAGUGUCUUGUUAUUGAUAUUGUUAUUGUUAUUGUUAUUGUUAUUGUUAUUGUUAUUG 216
H L E V C F M Y S D F H F I D E Q G E S I V V E (64)
CUUGGGCAUCCAAUUGCAUUGAAGCACAGUUAUUGAAUUAUUGGAGGAGAGUUGCCAAUAGCCUGG 288
L G D P N A L L K H R F E I I E G R D R T I A W (88)
ACAGUAAUAAACAGUUAUUGCAACUACAGGGGUGAGAAACAAAGUUUUCACAGUUAUUGUUAUUGU 360
T V I N S I C N T T G A E K P K F L P D L Y D Y (112)
AAGAAGAAUAGUUAUUGGAAUUGGAAUUGGAAUUGGAAUUGGAAUUGGAAUUGGAAUUGGAAUUG 432
K K N R F I E I G V T R R E V H I Y Y L E K A N (136)
AAAUUUAUUGUAGAGACACAUCCACAUUUAUUGUUAUUGUUAUUGUUAUUGUUAUUGUUAUUG 504
K I K S E K T H I H I F S F T G E E M A T K A D (160)
UACACUCUGAAGAAAGCGAGGCUAGGAUCAAACAGGCUUUAUCCAUAGACAAAGAAUUGGCUAGC 576
Y T L D E E S R A R I K T R L F T I R Q E M A S (184)
AGAGGCCUCUGGAAUUCUUGCUGACUCCGAGAGGGCGAAGAGACAAUUGAAGAAUUGAAGAAUUG 648
R G L W D S F R Q S E R G E E T I E E R F E I T (208)
GGAACAAUGCGCAAGCUUGCCGACCAAGUCUCCGCAACUUCUCCAGCCUUGAAAUUUAUUGAGCCU 720
G T M R K L A D Q S L P P N F S S L E N F R A Y (232)
GUGGAGUUAUUGCAACGACCUAUGAGGGCAAGCUUUCUCAAUUGUUAUUGUUAUUGUUAUUGUUA 792
V D G F E P N G Y I E G K L S Q M S K E V N A R (256)
AUUGAACUUUUUUAUUGAAUUAUUGAAUUAUUGAAUUAUUGAAUUAUUGAAUUAUUGAAUUAUUG 864
I E P F L K S T P R P L R L P D G P P C S Q R S (280)
AAAUUUCUGCUGAUGGAAUUGCUUUAUUAUUGAAUUAUUGAAUUAUUGAAUUAUUGAAUUAUUG 936
K F L L M D A L K L S I E D P S H E G E G I P L (304)
UUGAUGCAUCAAUUGCAUUGAAGCAUUCUUGGAAUGGAAUUGGAAUUGGAAUUGGAAUUGGAAU 1008
Y D A I K C M R T F F G W K E P N V V K P H E K (328)
GGAUUAUUAUUAUUGUUAUUGUUAUUGUUAUUGUUAUUGUUAUUGUUAUUGUUAUUGUUAUUG 1080
G I N P N Y L L S W K Q V L A E L Q D I E N E E G (352)
AAAUUUCAAAGGCUAAUUAUUGAAGAAUUAUUGAAGAAUUAUUGAAGAAUUAUUGAAGAAUUAUUG 1152
K I P R T K N M K K T S Q L K W A L G E N M A P I (376)
GAAAAGGAGACUUGGAGUUAUUGAAGAAUUAUUGAAGAAUUAUUGAAGAAUUAUUGAAGAAUUAUUG 1224
E K V D F D D C K D V G D L K Q Y D S D E P E L (400)
AGGUCGCUUGCAUUGGAAUUGCAUUGAAGCAUUGAAGCAUUGAAGCAUUGAAGCAUUGAAGCAUUG 1296
R S L A S W I Q N E F N K A C E L T D S S W I E (424)
CUUGAUGAUGGAGAAUUGGAGGCUUCAAUUGAAGCAUUGAAGCAUUGAAGCAUUGAAGCAUUGAAG 1368
L D E I G E D A A P I E H I A S M R R N Y F T A (448)
GAGGUGUCUUAUUGCAGAGCCACAGAAUUAUUGAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1440
E V S H C R A T E Y I M K G V Y I N T A L L N A (472)
UCCUGGAGCAUUGGAAUUGUUAUUGCAUUAUUGCAUUAUUGCAUUAUUGCAUUAUUGCAUUAUUG 1512
S C A A M D D F Q L I P M I S K C R T K E G R R (496)
AAGACAAUUGUACGGUUAUUGCAUUAUUGAAGAAUUGAAGAAUUGAAGAAUUGAAGAAUUGAAGAA 1584
K T N L Y G F I I K G R S H L R N D T D V V N F (520)
GUGAGCAUUGGAGUUAUUGCUCACUGACCAAGCAUUGAAGCAUUGAAGCAUUGAAGCAUUGAAGCA 1656
V S M E F S L T D P R L E P H K W E K Y C V L E (544)
GUAGGAGAUUGCUUAAGAAUGCCAUAGGCGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUG 1728
V G D M L L R S A I G H V S R P M F L Y V R T N (568)
GAAACUCAAUUAUUAUUGAAUUGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1800
G T S K I K M K W G M E M R R C L L Q S L Q Q I (592)
GAGAGUUAUUGAAGCUGAGUCCUUGUUAUUGAAGGAAAGCAUUGAAGCAUUGAAGCAUUGAAGCA 1872
E S M I E A E S S V K E K D M T K E F F E N K S (616)
GAAACUUGGCCUUGGAGAGUCCCAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1944
E T W P V G E S P K G V E E G S I G K V C R T L (640)
UUGGCAAGGUGUUAUUGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGG 2016
L A K S V F N S L Y A S P Q L E G F S A E S R K (664)
CUGCUUUAUUGCUUAGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGG 2088
I L L I V Q A L R D N L E P G T F D L G G L Y E (688)
GCAUUGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGG 2160
A I E E C L I N D P W V L L N A S W F N S F L T (712)
CAUGCAUUGGAGAGUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUGGCAUUG 2232
H A L R * (716)
U (2233)

```

Nucleotide sequence of the PA gene of A/WSN/33 written in the mRNA sense. The first 12 nucleotides were assumed to be the same as primer used in the cDNA synthesis.