



## **Influenza A(H1N1) M2 detailed pyrosequencing protocol for antiviral susceptibility testing**

**13 May 2009**

The WHO Collaborating Centre for influenza at CDC Atlanta, United States of America, has made available the M2 detailed pyrosequencing protocol, attached, for antiviral susceptibility testing of influenza A(H1N1).

This document updates and replaces the Protocol for antiviral susceptibility testing by pyrosequencing, published on 29 April 2009.



**Molecular Epidemiology Team**  
**Virus Surveillance and Diagnosis Branch**

*Centers for Disease Control and Prevention, NCIRD/CCID  
 Influenza Division*

SOP Title: Swine-H1N1 M2 detailed pyrosequencing protocol  
 Effective Date: May 11, 2009

**Pyrosequencing of M2 gene to detect resistance to adamantanes (amantadine and rimantadine)**

*RT-PCR primers (20 uM):*

sw-M2-F670, AGC TCC AGT GCT GGT CTG AAA G

sw-M2-R900-biot, GAC TCA GGC ACT CCT TCC GTA GAA

*Sequencing primer (100 uM):*

sw-M2-F747-seq, GCG ATT CAA GTG ATC C

**RT-PCR protocol:** using Invitrogen Superscript III One step RT-PCR with HiFi Taq system.

	Volume (ul)
<b>2X Reaction Mix</b>	<b>25</b>
<b>Superscript III HiFi Taq</b>	<b>1</b>
<b>Forward primer(20uM)</b>	<b>1</b>
<b>Reverse primer(20uM)</b>	<b>1</b>
<b>Rnase Inhibitor</b>	<b>0.5</b>
<b>Nuclease free water</b>	<b>16.5</b>
<b>Total/sampels</b>	<b>45</b>
<b>Template</b>	<b>5</b>
<b>Final volume</b>	<b>50</b>

**PCR Cycling Parameters:**

Temp	Length	
<b>50</b>	<b>30min</b>	
<b>94</b>	<b>2min</b>	
<b>94</b>	<b>15sec</b>	<b>45 cycles</b>
<b>55</b>	<b>30sec</b>	
<b>68</b>	<b>60 sec</b>	
<b>68</b>	<b>5min</b>	
<b>4</b>	<b>hold</b>	

**Pyrosequencing protocol:** using PyroGold Reagents (for 96 samples)

*1- Binding Reaction*

- 1- Binding Buffer: 4 ml
- 2- Dist H<sub>2</sub>O: 2 ml
- 3- Sepharose beads (vortex well): 300 ul
- 4- Mix well by vortexing
- 5- Dispense 60 ul of binding buffer to each well of a clean 96-RT-PCR plate (Use 12 channel)
- 6- Transfer 20 ul of RT-PCR product to appropriate wells containing binding buffer.
- 7- Cover with film tape
- 8- Shake plate for at least 10 min

*2- Annealing Reaction:*

- 1- Annealing buffer: 4.4 ml
- 2- Sequencing primer: 20 ul (sw-M2-F747-seq) at 100 uM
- 3- Vortex and add 40 ul annealing reaction to each well of a 96-well soft pyrosequencing plate.

*3- Dispensation order:* cyclic 60 (CATG/GT)

*4- Module used:* SQA (sequence analysis)

*5- Sequence to be analyzed:*

**M2 pyrosequencing target regions of seasonal H1N1 and H3N2 compared to that of the swine origin 2009 H1N1 viruses: (nucleotide and amino acid differences compared to seasonal H1N1 viruses are shown in bold font and underlined).**

	25	26	27	28	29	30	31	32	33	34	35	36	37	38		
	P	L	V	V	A	A	N	I	I	G	I	L	H	L		
<b>H3:</b>	AC	CCG	CTT	GTT	GTT	GCC	GCG	<b>AAT</b>	ATC	ATT	GGG	ATC	TTG	CAC	TTG	AT
	P	L	V	V	A	A	S	I	I	G	I	V	H	L		
<b>H1:</b>	AT	CCT	CTT	GTT	GTT	GCC	GCA	AGT	ATA	ATT	GGG	ATT	GTG	CAC	CTG	AT
	P	L	V	<u>I</u>	A	A	<b>N</b>	I	I	G	I	L	H	L		
<b>swH1:</b>	AT	CCT	<u>CTC</u>	<u>GTC</u>	<u>ATT</u>	<u>GCA</u>	GCA	<b>AAT</b>	ATC	ATT	GGG	ATC	TTG	CAC	CTG	AT

Established markers of resistance to M2 blockers (amantadine and rimantadine):

L26F (CTT/C to TTT)

V27A (GTT to GCT)

A30V/T (GCG to GTG), or GCN to GTN, GCG to ACG

S31N (AGT/C to AAT/C)

G34E (GGG to GAG)

**The 2009 swine-origin H1N1 viruses are adamantane resistant and carry the S31N change in M2 protein (AGT----AAT).**