



National Institute for Public Health  
and the Environment  
*Ministry of Health, Welfare and Sport*

# **RIVM Laboratory Protocol Library**

## **Influenzavirus A and B PCR LightCycler – probe test A-Matrix-B-NS-H1pdm09-H3-H5-N1pdm09-N1avian-N2**

Updated November 2016

This protocol has been made available by:

**Pieter Overduin  
Dr. Adam Meijer  
National Institute for Public Health and the Environment,  
Diagnostic Laboratory for Infectious Diseases and Screening  
Department Virology  
PO Box 1  
3720 BA Bilthoven  
The Netherlands  
e-mail: [pieter.overduin@rivm.nl](mailto:pieter.overduin@rivm.nl), [adam.meijer@rivm.nl](mailto:adam.meijer@rivm.nl)**

The copyright of this protocol resides with the original author(s).  
The original author(s) are responsible for the correctness of the contents of this protocol.  
This protocol may be used by diagnostic laboratories in the Netherlands and the members of the European Influenza Surveillance Network (EISN) in the context of the Community Network of Reference Laboratories for Human Influenza in Europe.  
Distribution of this protocol outside the network and publication in any form are prohibited without obtaining explicit permission of the original author(s), unless the protocol has been published in a journal.

# 1. Reagents

## 1.1. RNA isolation

Automated MagNAPure LC Total Nucleic Acid Kit (Roche cat no. 03 038 505 01)

or,

Manual High Pure RNA isolation Kit (Roche cat no. 11 828 665 001)

**One-step protocol for: A-Matrix, B-NS, H3, H1pdm09 H5, N1pdm09, N1avian and N2**

## 1.2. Lightcycler PCR

PCR reactions are validated on the LightCycler 480 with:

- one-step: TaqMan® Fast Virus 1-Step Master Mix (LifeTechnologies cat no. 4444436)

## 1.3. Primersets \*\*

**Matrix Protein gene (specific for all influenzavirus A subtypes, H1-H16).**

INFAM Sense1	AAG ACC AAT CCT GTC ACC TCT GA*
INFAM Sense2	AAG ACC AAT CTT GTC ACC TCT GA*
INFAM Sense4	AAG ACC AAT TCT GTC ACC TCT GA*
INFAM a-Sense	CAA AGC GTC TAC GCT GCA GTC C
INFAM a-Sense2	TAA AGC GTC TAC GCT GCA GTC C

INFAM probe3 5'6 *Fam* TTT GTK TTC ACG CTC ACC GTG CC 3'*EDQI*

\* INFAM sense primer is a mixture of equal amounts of the three primer variants to a final concentration of 30 pmol/ul. The reason is the following distribution of ambiguities at positions 10 and 11 of the primer site in the matrix genome segment:

\* INFAM a-sense primer is a mixture of equal amounts of the two primer variants to a final concentration of 30 pmol/ul. The reason is the following distribution of ambiguities at position 1 in the matrix genome segment:

\*\* Adapted primers are summarised in the appendix below.

A(H1N1)pdm09	TCTT
A(H3N2) 2004-2016	TTCT
A(H1N1) seasonal	TCCT
A(H5N1) human viruses	TCCT

### NS-INF B

INFB-NS779F	GTC TTA ATG AAG GAC ATT CAA AGC C
INFB-NS886R	TAA AGT TCT TCC GTG ACC AGT CTA
INFB-848P	5' <i>Yak Yellow</i> GTC AAG AGC ACC GAT TAT CAC CAG AAG AG 3' <i>BHQ1</i>

### HA seasonal H3 subtype

H3-1541F2	CRAT GTR TAC AGG GAT GAA GCWTTA AAC A
H3-1600R	TAG GAT CCA ATC TTT GTA CCC TGA CTT
H3-1571P1	5' <i>Yak Yellow</i> AGC TCA ACT CCC TTG ATC TGG AAY CGG 3' <i>BHQ1</i>

### HA human H1pdm09 (H1v) subtype

H1v-1306F	TGG ACT TAC AAT GCC GAA CT
H1v-1423R	CAG CCG TTT CCA ATT TCC TT
H1v-1357P2	5' <i>Texas Red</i> GGAC TAT CAC GAT TCA AAT GTG AAG AAC T 3' <i>BHQ2</i>

### HA avian H5 subtype

H5-HA-960F ATA TgT gAA ATC AAA CAR ATT AgT CC  
H5-HA-1100R gTR gAT TCT TTg TCT gCA gCg TA  
H5-HA-1050P 5'6Fam TAC CCA TAC CAA CCA TCT ACC ATT CCC T3'BBQI

### NA seasonal N2 subtype (specific for human NA)

N2-278F2 GTG GCA TTA CAG GAT TTG CAC  
N2-376R YTG TCC AAG GGC AAA TTG ATA AC  
N2-327P 5'Yak-Yellow GAC ATC TGG GTG ACA AGA GAA CCT TAT-3'BHQI

### NA human N1pdm09 (N1v) subtype

N1v-1142F1 GAC TGG GAC TGA CAA TAA ATT CTC AA  
N1v-1244R1 CAG CCC TGT TAG TTC TGG ATG.C  
N1v-1193P1 5'6 Fam AGT GGT CAG GGT ATA GCG GGA GT 3'BHQI

### NA avian N1 subtype (specific for H5 HPAI NA)

N1-1069F1\* CCA AAT GGG TGG ACT GRA A  
N1-1069F2\* CCA AAT GGG TGG gCT GgA a  
N1-1079F\* GGA CTG RAA CGG ACA GTA  
N1-1207R AGA ARC AAG GTC TTA TGC ART CTA  
N1-1150P1\*\* 5'Yak Yellow GGG AGT TTT GTY CAG CAT CCA GAA 3'BHQI  
N1-1150P2\*\* 5'Yak Yellow GGG AGT TTT GTC CAG CAT CCR GAA 3'BHQI

\* mix the N1 forward (F) primers in equimolar amounts

\*\* mix the N1 probes (P) in equimolar amounts

## 2. Procedure

1. Prepare appropriate dilutions of positive controls
2. Isolate RNA using the High pure RNA isolation Kit or MagNApure

### Mixes

Influenza A Matrix /B NS	[end-conc.]	µl
PCR grade water		
Taqman FV MM (4x)		5
INFAM Sense mix 1-3	0.75 µM	
INFAM a-Sense	0.75 µM	
INFAM probe 2 (Fam)	0.25 µM	
INFB-NS779F	0.75 µM	
INFB-NS886R	0.75 µM	
INFB-848P	0.25 µM	
Total volume		15

H1pdm09-H3 Human	[end-conc.]	µl
PCR grade water		
Taqman FV MM (4x)		5
H1-sw-1306F1	0.75 µM	
H1-sw-1423R	0.75 µM	
H1 sw-1357p2	0.25 µM	
H3-1541F1	1.5 µM	
H3-1600R	0.75 µM	
H3-1571P1	0.75 µM	
Total volume		15

<b>H5</b>	[end-conc]	µl
PCR grade water		
Taqman FV MM (4x)		5
H5-HA-960F	0.75 µM	
H5-HA-1100R	0.75 µM	
H5-HA-1050P	0.3 µM	
Total volume		15

<b>N1pdm09-N2</b>	[end-conc]	µl
PCR grade water		
Taqman FV MM (4x))		5
N1pdm09-1142F	0.75 µM	
N1pdm09-1244R	0.75 µM	
N1pdm09-1193P	0.25 µM	
N2-278F	1 µM	
N2-376R	0.63 µM	
N2-327P	0.25 µM	
Total volume		20

<b>NA-Avian H5-HPI</b>	[end-conc]	µl
PCR grade water		
Taqman FV MM (4x))		5
H5-NA-1069F1	0.25 µM	
H5-NA-1069F2	0.25 µM	
H5-NA-1079F	0.25 µM	
H5-NA-1207R	0.75 µM	
H5-Na-1150P1	0.15 µM	
H5-NA-1150P2	0.15 µM	
Total volume		20

Mix per reaction with RNA

PCR grade water  
Taqman FV MM (4x)  
Primer/probe mix

7 µl  
5µl  
3µl  
\_\_\_\_\_  
+  
15 µl mix volume

RNA

5 µl  
\_\_\_\_\_  
+  
20 µl final volume

**LC 480 One-step PCR temperature protocol:**

PCR Program	Segment number	Temperature target (°C)	Hold Time	Ramp rate (°C/sec.)	Acquisition mode
Reverse transcription	1	50	15 min.	4.4	None
Reverse transcription	2	95	2 min.	4.4	None
Amplification (cycles:50)	1	94	10 sec.	4.4	None
	2	60	30sec.	2.2	Single
Cooling	1	40	30	4.4	None

## Exotic avian Influenza's H7N9 en H9

### HA1 avian H7 subtype\*

AI-H7N9-F1 cgt gca agC ttCctg aga gg  
 AI-H7N9-R2 gGc ctt ccc atc cat ttt ca  
 AI-H7N9-TM1 5'6 Fam' aac ccg cta tAg cac caa ata ggc ctc 3'BHQ1

*\*Adapted from Slomka et al. Influenza Other Respir Viruses 2009 Jul.3(4) 151-64 Nucleotides in Capital adapted*

### HA2 avian H7 subtype\*

H7-LH6FGGC CAG TAT TAG AAA TAA CAC CTA TGA  
 H7-RH4 GCC CCG AAG CTA AAC CAA AGT AT  
 H7-pro11 5'6 Fam CCG CTG CTT AGT TTG ACT GGG TCA ATC T 3'BHQ1

### NA avian (H7) N9 subtype

N9-1210F TCT TTC ATG GAC TAT TGG GC  
 N9-1334R GAA CAC ATC GAT ACT AKA CTA TTG C  
 N9-1271P 5'6 Fam'TAC GTG GAA GAC CCA AGG ARG ATA AAG 3'BHQ1

### HA avian H9 subtype

H9-FOR ATG GGG TTT GCT GCC  
 H9-REV TTA TAT ACA RAT GTT GCA YCT G  
 H9-PROBE 5'6 Fam'TTC TGG GCC ATG TCC AAT GG 3'BHQ1

H7-HA1	[end-conc]	µl
PCR grade water		
Taqman FV MM (4x)		5
AI-H7N9-F1	0.25 µM	
AI-H7N9-R2	0.25 µM	
AI-H7N9-TM1	0.125 µM	
Total volume		15

H7-HA2	[end-conc]	µl
PCR grade water		
Taqman FV MM (4x)		5
H7-LH6F	0.25 µM	
H7-RH4	0.25 µM	
H7-pro11	0.125 µM	
Total volume		15

PCR Program <b>H9</b>	Segment number	Temperature target (°C)	Hold Time	Ramp rate (°C/sec.)	Acquisition mode
Reverse transcription	1	50	15 min.	4.4	None
Reverse transcription	2	95	2 min.	4.4	None
Amplification (cycles:50)	1	94	10 sec.	4.4	None
	2	60	30sec.	2.2	Single
Cooling	1	40	30	4.4	None

<b>N9</b>	[end-conc]	µl
PCR grade water		
Taqman FV MM (4x)		5
N9-1210F	0.25 µM	
N9-1334R	0.25 µM	
N9-1271	0.125 µM	
Total volume		15

<b>H9</b>	[end-conc]	µl
PCR grade water		
Taqman FV MM (4x)		5
H9-FOR	0.5 µM	
H9-REV	0.5 µM	
H9 probe	0.25 µM	
Total volume		15

### Appendix November 2016.

Based on *in silico analysis* of Influenza sequences from the 2015-2016 season, the following primers and probes are adapted to emerged mutations.

All other primers and probes are not changed

INFAM-sense4	AAG ACC AAT TCT GTC ACC TYT GA
INFAM A-sense2	TAA AGC GTC TAC GCT GCA GTC C
INFAM-probe3	TTT GTK TTC ACG CTC ACC GTGCC
H3-1541F2	CRAT GTR TAC AGG GAT GAA GCW TTA AAC A
N2-278F2	GTG GCA TTA CAG GAT TTGCAC
H1-sw-1357P2	GGACTAT CAC GAT TCA AAT GTGAAG AACT
N1v-1142F1	GACTGGGA CTGACA ATA AAT TCT CAA
N1v-1244R1	CAGCCCT GTT AGT TCT GGA TGC
N1v-1193P1	AGT GGT CAG GGT ATA GCGGGA GT