# QUANTITATION OF INFLUENZA A VIRUS FROM NASAL AND LUNG TISSUE OF COTTON RATS USING REAL-TIME RT-PCR AND CULTURE

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# Background

The 2007/08 influenza season was the worst in three seasons in terms of overall morbidity and mortality (1), particularly due to antigenic drift in the current trivalent influenza vaccine. Additionally, an increase in the cumulative confirmed cases of high pathogenic H5N1 has raised concerns of a potential pandemic in humans. Since Jan 1, 2008 avian influenza has continued to circulate in Indonesia, Vietnam, and Egypt with 30 confirmed human cases and 23 fatalities (76% mortality) (2).

A highly sensitive and specific set of real-time RT-PCR (rRT-PCR) assays for point of care detection of influenza virus typing (influenza A or B) and subtyping (H3N2, H1N1, and H5N1) would facilitate patient care and decrease costly evaluations. In a potential pandemic, rapid detection of infected individuals would enhance intervention and patient cohorting, and could help prevent widespread dissemination of disease in the community.

The cotton rat (*Sigmodon hispidus*) is a useful small animal model for influenza pathogenesis (Figure 1) and several studies exploring influenza nasal and pulmonary infection, immunological response, and antiviral therapy have been performed (3-5). This study compared the sensitivity of 'gold standard' culture to the recently developed PrimeMix real-time RT-PCR System for the detection of influenza virus in primary infected and non-infected (sentinel) cotton rats.



Figure 1. *Sigmodon hispidus*, a commonly used cotton rat in biomedical research.

## Objective

To compare quantitative sensitivities rRT-PCR and culture for detecting influenza virus in lung and nasal tissue from primary infected and sentinel (non-infected) cotton rats.

### Methods

Cotton rats (primary infected) were inoculated intranasally with 10<sup>7</sup> TCID<sub>50</sub> influenza A (H3N2) virus and housed with non- infected (sentinel) rats. Nasal and lung tissue homogenates obtained on post infection day 1,4,10,21, and 28 were analyzed using quantitative culture and real-time with an exogenous control and calibrator sample.

The exogenous control is a distinct nucleic acid sequence (e.g., influenza B) of known concentration added to each sample and serves as a reference to control for variations from extraction and pipetting (6). The calibrator is an untreated control used for normalizing viral concentration (Figure 2).

RNA extractions from homogenized nose and lung samples were lysed and preserved in PrimeStore Solution. Extraction was performed using the RNaqueous Micro Kit (Ambion). Realtime RT-PCR assays for influenza A were designed, optimized, and evaluated according to Daum *et al.* (7) and adapted into PrimeMix, a ready-use simplified rRT-PCR blend.

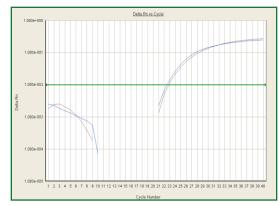




Figure 2. Real-Time RT-PCR Assays Detect Known Amounts of Calibrator and Exogenous Template and can Quantify Viral Unknowns in Nose and Lung Samples. The calibrator and exogenous control reaction represent 0.1 pg of template cRNA corresponding to approximately  $5.8 \times 10^5$  influenza copies. The calibrator and exogenous controls were used in subsequent real-time RT-PCR to quantify viral copy number from cotton rat nose /lung samples inflected with

## Results

Table 1. Influenza detection in nose and lung tissue using real-time RT-PCR and

culture.						
		Nose		Lung		
Sample	Treatment	Culture (TCID₅o)	rRT-PCR (copy #)	Culture (TCID₅o)	rRT-PCR (copy #)	
71073	control	(ICID <sub>50</sub> )	(сору #)	(TCID <sub>50</sub> )	(copy #)	
71073	control	-	· · · · · · · · · · · · · · · · · · ·	- · · · · · · · · · · · · · · · · · · ·		
71075	control	-	·	- · · · · · · · · · · · · · · · · · · ·	: _	
71076	control	-	-	-	· _	
71077	Primary infected	300	5x10⁵	325	3x10 <sup>4</sup>	
71078	Primary infected	350	2x10 <sup>6</sup>	375	2x10 <sup>7</sup>	
71079	Non-infected sentinel	-	<del>.</del>	-	5x10 <sup>6</sup>	
71080	Non-infected sentinel	-	-	-	2x10 <sup>4</sup>	
71081	Primary infected	325	6x10⁵	325	124	
71082	Non-infected sentinel	-	<del>.</del>	-	9x10 <sup>6</sup>	DAY 1
71083	Non-infected sentinel	-		-	:	
71084	Primary infected	300	6x10⁵	425	6x10 <sup>6</sup>	
71085	Primary infected	325	6x10⁵	350	4x10 <sup>6</sup>	
71086	Primary infected	325	2x10⁵	375	3x10 <sup>6</sup>	7
71087	Primary infected	300	2x10 <sup>6</sup>	375	3x10 <sup>4</sup>	
71088	Primary infected	300	3x10 <sup>6</sup>	· · · · · · · · · · · · · · · · · · ·	9893	
71089	Primary infected	275	6x10 <sup>6</sup>	<del>.</del>	9x10 <sup>4</sup>	
	Non-infected sentinel		116		82	
71091 71092	Non-infected sentinel	275	 6x10⁵		5x10⁴	
71092	Primary infected Primary infected	275 300	1x10 <sup>6</sup>		1x10 <sup>4</sup>	
71093	Primary infected	250	;1X10 6x10⁵	- · · · · · · · · · · · · · · · · · · ·	:	DAY 4
71094	Non-infected sentinel	-	124	-		
	Non-infected sentinel	-	210	-		
	Non-infected sentinel	-	53	-	· -	
	Primary infected	425	2x10 <sup>6</sup>	-	5x10 <sup>5</sup>	
71099	Non-infected sentinel	-	582	-	-	
71100	Primary infected	375	1x10 <sup>9</sup>	-	4x10 <sup>5</sup>	
71101	Primary infected	400	7x10 <sup>6</sup>		8x10⁵	
71102	Primary infected	-	482	-	: :	
71103	Primary infected	· <del>-</del>	6x10⁴	<del>.</del>	193	
	Non-infected sentinel		489		i	
	Primary infected		7565	<del>.</del>	<u>.</u>	
	Non-infected sentinel			<del>.</del>	· · · · · · · · · · · · · · · · · · ·	
	Primary infected		4x10 <sup>4</sup>		174	
	Non-infected sentinel Non-infected sentinel		114		}	→ DAY 10
	Primary infected		2x10 <sup>7</sup>		<u> </u>	DAI 10
	Primary infected		4x10 <sup>6</sup>		<u>:</u>	
	Non-infected sentinel		-			
	Non-infected sentinel	-	-	-	-	
	Primary infected	-	4x10 <sup>4</sup>	-	17	
	Primary infected	-	9x10⁴	-	31	
	Primary infected	-		-	· · · · · · · · · · · · · · · · · · ·	
	Primary infected	-	145	-	-	
	Primary infected	-	112		-	
71119	Non-infected sentinel	-		-	29	
	Non-infected sentinel	-			-	
	Primary infected		582	<del>.</del>	76	►DAY 21
	Primary infected	· · · · · · · · · · · · · · · · · · ·	67	<del>.</del>	489	
	Primary infected	<del>.</del>	103	<del>.</del>	ļ	
	Non-infected sentinel		33	<del>.</del>	;	
	Non-infected sentinel		66	<del>.</del>		
	Primary infected		85	<del>.</del>	<u>:</u>	
	Non-infected sentinel			<del>.</del>	·····	
	Non-infected sentinel				500	
	Primary infected		9	· · · · · · · <del>·</del> · · · · · ·	569	
	Primary infected Primary infected	· · · · · · · · · · · · · · · · · · ·			······	
71131	Primary infected Primary infected	<del>.</del>	0		<del>.</del>	<b>→</b> DAY 28
	Non-infected sentinel	-	· · · · · · · · · · · · · · · · · · ·	- · · · · · · · · · · · · · · · · · · ·		
	Primary infected	-	10	-	· · · · · · · · · · · · · · · · · · ·	
	Non-infected sentinel	-	<del></del>	-	· · · · · · · · · · · · · · · · · · ·	
	Primary infected	-	,	-	: -	

#### Culture

❖All primary infected cotton rats (N=36) became ill and culture-positive influenza virus was detected from all nasal (6/6) and lung (6/6) samples on Day 1 post-infection.

VACCINES OF DIAGNOSTICS

- ♦On Day 4 primary infected nose samples (9/9) but no matched lung samples (0/9) were culture-positive, and thereafter (Day 10,21,28) all primary-infected lung and nasal samples were culture-negative.
- ❖All sentinel (non-infected) rats (Day 1-28) were culturenegative. Sentinel animals analyzed on Day 21 had increased antibody levels to influenza virus.

#### Real-time RT-PCR

❖rRT-PCR detected influenza in lung and nasal tissue from primary-infected and non-infected animals at times when culture was negative and at levels < 10 viral copies.

#### Conclusions

- The rRT-PCR method described here is rapid (<2 hours), more sensitive than traditional culture and could be valuable for point-of-care patient influenza detection.
- ➤ PrimeMix influenza assays could be valuable for early therapeutic intervention for high risk populations and to decrease spread of infection during a pandemic.
- Studies are currently underway to evaluate these diagnostic assays for point of care detection in children and families.

#### References

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