

# Persistence of Influenza A (H1N1) Virus on Stainless Steel Surfaces

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## ABSTRACT

As annual influenza epidemics continue to cause significant morbidity and economic burden, an understanding of viral persistence and transmission is critical for public health officials and health care workers to better protect patients and their family members from infection. The infectivity and persistence of two influenza A (H1N1) virus strains (A/New Caledonia/20/1999 and A/Brisbane/59/2007) on stainless steel (SS) surfaces were evaluated using three different surface matrices (2% fetal bovine serum, 5 mg/ml mucin, and viral medium) under various absolute humidity conditions ( $4.1 \times 10^5$  mPa,  $6.5 \times 10^5$  mPa,  $7.1 \times 10^5$  mPa,  $11.4 \times 10^5$  mPa,  $11.2 \times 10^5$  mPa, and  $17.9 \times 10^5$  mPa) for up to 7 days. Influenza A virus was deposited onto SS coupons ( $7.07 \text{ cm}^2$ ) and recovered by agitation and sonication in viral medium. Viral persistence was quantified using a tissue culture-based enzyme-linked immunosorbent assay (ELISA) to determine the median (50%) tissue culture infective dose (TCID<sub>50</sub>) of infectious virus per coupon. Overall, both strains of influenza A virus remained infectious on SS coupons, with an approximate 2 log<sub>10</sub> loss over 7 days. Factors that influenced viral persistence included absolute humidity, strain-absolute humidity interaction, and time ( $P \leq 0.01$ ). Further studies on the transfer of influenza A virus from fomites by hand and the impact of inanimate surface contamination on transmission should be performed, as this study demonstrates prolonged persistence on nonporous surfaces.

## IMPORTANCE

This study tested the ability of two influenza A (H1N1) virus strains to persist and remain infectious on stainless steel surfaces under various environmental conditions. It demonstrated that influenza A (H1N1) viruses can persist and remain infectious on stainless steel surfaces for 7 days. Additional studies should be conducted to assess the role played by contaminated surfaces in the transmission of influenza A virus.

Influenza virus continues to be a pathogen of significant interest, as the World Health Organization estimates that annual influenza epidemics cause approximately 5 million cases of serious illness and over 250,000 deaths per year (1). In the United States alone, it has been estimated that there are approximately 226,000 hospitalizations and 3,000 to 48,000 deaths during an influenza virus season, leading to a total economic burden that can reach \$87.1 billion (2–4). Influenza A virus is an enveloped RNA virus which has been shown to be transmissible primarily by droplet, with less evidence of contact and airborne transmission being available (5, 6). Previous research has shown that fomites and surface contamination caused by large respiratory droplets may play a significant role in transmission (6–8).

Current Centers for Disease Control and Prevention health care infection control guidelines for seasonal influenza recommend performing hand hygiene before and after patient contact, after contact with contaminated surfaces or materials, and before personal protective equipment is put on and removed. Additional strategies include disinfection of potentially contaminated non-critical surfaces, including frequently touched surfaces that are in close proximity to the patient (e.g., hospital bed rails, overbed tables), with U.S. Environmental Protection Agency-registered hospital disinfectants (9, 10). However, lapses in infection control, most notably, lapses in hand hygiene implementation (11–13), may facilitate the transmission of influenza virus. Boone and Gerba (14) have shown that 50% of fomites found in homes and day care centers had influenza virus present during an active influenza season. Influenza A virus has also been shown to be transferred from stainless steel (SS) countertops to hands for up to 24 h

after surface contamination (15); however, indirect transmission through contact with these surfaces has not been demonstrated.

Regardless of the precise role of fomites and surface contamination in transmission, it is important to have a clear understanding of what factors influence the persistence of pathogens on surfaces and how this persistence impacts transmission from contaminated environmental surfaces. Previous studies of persistence on environmental surfaces have found that various strains of influenza A (H1N1) virus remained infectious on SS surfaces for approximately 24 h but no longer than 72 h (15–19). In this study, we evaluated the persistence and infectivity of two influenza A (H1N1) virus strains, A/New Caledonia/20/1999 (A/NC-H1N1) and A/Brisbane/59/2007 (A/Br-H1N1), in three different substrate matrices (viral medium, 2% fetal bovine serum [FBS], and 5 mg/ml mucin) on SS surfaces stored for various times up to 7 days under a range of absolute humidity conditions.

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## MATERIALS AND METHODS

**Cell culture and viral propagation.** Madin-Darby canine kidney (MDCK) epithelial cells (CCL-3; ATCC, Manassas, VA) that had been passaged fewer than 90 times were maintained on a medium containing Dulbecco's modified Eagle medium (DMEM; Gibco, Grand Island, NY), 2% (vol/vol) FBS (Atlanta Biologicals, Lawrenceville, GA), and 2% (vol/vol) penicillin-streptomycin (stock concentration, 10,000 units/ml penicillin G sodium and 10,000 µg/ml streptomycin sulfate; Life Technologies, Carlsbad, CA). Cells were incubated at 37°C in 5% CO<sub>2</sub> for 24 to 48 h until they were 90% confluent. MDCK cells were rinsed with 1× phosphate-buffered saline (PBS), and the virus was added to obtain a multiplicity of infection of 0.01. The two influenza A (H1N1) virus strains used in this experiment, A/Brisbane/59/2007 (A/Br-H1N1) and A/New Caledonia/20/1999 (A/NC-H1N1), were obtained from the Lovelace Respiratory Research Institute (Albuquerque, NM). These strains were chosen because they were seasonal circulating strains and were included in the WHO yearly influenza vaccine recommendations (20). The cells were incubated at 37°C in 5% CO<sub>2</sub> until an approximately 75% cytopathic effect (CPE) was observed. The virus was harvested, cell debris was removed by centrifugation, and aliquots of the supernatant were stored at -80°C. The starting concentration for A/NC-H1N1 was approximately  $1.2 \times 10^4$  50% tissue culture infective doses (TCID<sub>50</sub>)/ml, and that for A/Br-H1N1 was  $6.95 \times 10^5$  TCID<sub>50</sub>/ml.

**Inoculation of stainless steel coupons and environmental conditions.** This study evaluated the persistence of the influenza A (H1N1) virus in various substrates: viral medium (DMEM, 7.5% bovine serum albumin [BSA; Life Technologies, Carlsbad, CA], 2% penicillin-streptomycin, HEPES buffer [Gibco, Grand Island, NY], and tosylsulfonyl phenylalanyl chloromethyl ketone [TPCK]-treated trypsin [Thermo Scientific, Rockford, IL]), 2% FBS, and 5 mg/ml mucin (MP Biomedicals, Solon, OH). SS sheets (24 gauge; T-304; Stewart Stainless Supply, Suwanee, GA) were punched into 7.07-cm<sup>2</sup> coupons. SS was used as a surrogate for nonporous surfaces because it is a material commonly found in hospitals (e.g., safety grip bars, poles holding intravenous solution) and representative of household surfaces (e.g., doorknobs). Prior to inoculation, the SS coupons were washed with a low-residue cleaner, rinsed thoroughly in reverse osmosis water, soaked in 70% ethanol for 1 h, and then left to air dry. The SS coupons were placed in glass petri dishes and autoclaved at 121°C for 20 min and then an additional 20 min for drying. The three sterile SS coupons were transferred to 6-well plates that had been UV sterilized in a biosafety cabinet (BSC). Equal volumes of virus suspension were combined with the undiluted sample matrix to achieve the desired 2% FBS and 5-mg/ml mucin concentrations. Virus mixture aliquots of 100 µl were added to each SS coupon and spread with a cell spreader, making the concentration for A/NC-H1N1 equal to  $1.2 \times 10^3$  TCID<sub>50</sub>/coupon and that for A/Br-H1N1 equal to  $6.9 \times 10^4$  TCID<sub>50</sub>/coupon. The SS coupons were allowed to air dry for 1 h at room temperature in the BSC (sash closed, no blower). The 6-well plates were transferred to a temperature- and humidity-controlled environmental chamber (Caron, Marietta, OH) under the following conditions for up to 7 days:  $4.1 \times 10^5$  mPa (18°C and 20% relative humidity [RH]),  $6.5 \times 10^5$  mPa (25°C and 20% RH),  $7.1 \times 10^5$  mPa (18°C and 35% RH),  $11.4 \times 10^5$  mPa (25°C and 35% RH),  $11.2 \times 10^5$  mPa (18°C and 55% RH), and  $17.9 \times 10^5$  mPa (25°C and 55% RH). Each condition was tested once, with each time point tested in triplicate.

**Sample processing.** Inoculated SS coupons were processed at 0 h ( $T_0$ ), 24 h ( $T_{24}$ ), 48 h ( $T_{48}$ ), 96 h ( $T_{96}$ ; 4 days), and 168 h ( $T_{168}$ ; 7 days). For sample processing, 1.9 ml of viral medium was added to the 6-well plate and, as described by Donlan et al., alternately agitated by sonication and vortexing three times for 30 s each time (21). The medium was collected and dispensed into two separate tubes. The samples were stored at -80°C until analysis.

**ELISA.** The tissue cultured-based enzyme-linked immunosorbent assay (ELISA) method, which measures the ability of the virus particles to infect and replicate, has been fully described by Coulliette et al. (22).

Briefly, viral samples were placed in a 96-well plate and diluted in series 1:3 in 100 µl of viral medium. MDCK cells were removed from the flask by trypsin-EDTA (Gibco, Grand Island, NY), centrifuged, and resuspended in viral medium, and the suspension was added to each well. The plate was incubated at 37°C in 5% CO<sub>2</sub> for 17 to 19 h. The plates were rinsed with 1× PBS, fixed with 80% acetone-1× PBS, and allowed to air dry in the biosafety cabinet. The primary antibody, a mouse anti-influenza A virus monoclonal antibody (Millipore, Temecula, CA) which binds to the nucleoprotein epitope, was diluted 1:1,000 in 1× PBS-Tween 20-1% BSA, the diluted antibody was added to each well, and the plates was incubated for 1 h and washed with 1× PBS-Tween 20 (PBST). The secondary antibody, peroxidase-labeled affinity-purified goat anti-mouse IgG antibody (KPL, Gaithersburg, MD), was diluted 1:1,000 in 1× PBS-Tween 20-1% BSA, the diluted antibody was added to each well, and the plate was incubated for 1 h and washed with 1× PBST. To develop the substrate, a solution containing phosphate-citrate buffer with sodium perborate (Sigma-Aldrich, St. Louis, MO), *o*-phenylenediamine dihydrochloride tablets (10 mg; Sigma-Aldrich, St. Louis, MO), and hydrogen peroxide (Sigma-Aldrich, St. Louis, MO) was added to each well, the plate was incubated at room temperature, and then the reaction was completed with sulfuric acid (Sigma-Aldrich, St. Louis, MO). The 96-well plates were read at an absorbance λ of 490 nm in a Synergy II plate reader (BioTek Instruments, Winooski, VT) using the Gen5 program (versions 1.11 and 2.00).

**Data analysis and statistics.** The absorbance data were exported to Microsoft Excel software (version 14; Redmond, WA). The number of TCID<sub>50</sub> per coupon was determined using the standard Reed and Muench method (23). The limit of detection for the ELISA was  $1.44 \times 10^1$  to  $3.40 \times 10^5$  TCID<sub>50</sub>/ml. The number of TCID<sub>50</sub> per coupon recovery results for each time point were log<sub>10</sub> transformed, and the mean log<sub>10</sub> recovery and change in log<sub>10</sub> recovery relative to the number of TCID<sub>50</sub> per coupon at  $T_0$  were calculated. SPSS software (version 19; Somers, NY) was used to generate box plots and descriptive statistics. SAS software (version 9.2; Cary, NC) was used to conduct linear modeling to assess the independent importance and best parameterization of each factor (i.e., sample matrix, absolute humidity [AH], and time). A generalized estimating equations (GEE) approach based on the best linear model was used to account for the correlation of the mean log<sub>10</sub> differences due to the clustering of replicates over time, with the intercept representing the mean log<sub>10</sub> change for the referent strata (24). The significance level for all tests performed was a *P* value of 0.01. It is important to note that the data for several groups could be combined for a more robust statistical analysis, as group maximum likelihood estimation (MLE) analyses showed that there was no significant difference (*P* > 0.01) between the levels at times of 24 and 48 h (identified as 24 to 48 h) and among the following AH groups:  $6.5 \times 10^5$  mPa,  $7.1 \times 10^5$  mPa,  $11.4 \times 10^5$  mPa, and  $11.2 \times 10^5$  mPa (identified as  $11.4 \times 10^5$  mPa).

## RESULTS

The mean number of TCID<sub>50</sub> of A/NC-H1N1 recovered per coupon was  $1.68 \times 10^2$  (range,  $1.22 \times 10^1$  to  $6.87 \times 10^2$ ) across all AHs and times, which is equivalent to a 13.9% recovery relative to the mean number of TCID<sub>50</sub> per coupon in the inoculum at  $T_0$ . The mean number of TCID<sub>50</sub> of A/Br-H1N1 recovered per coupon was  $1.41 \times 10^3$  (range,  $2.77 \times 10^2$  to  $4.89 \times 10^3$ ), which is equivalent to a 2.04% recovery relative to the mean number of TCID<sub>50</sub> per coupon in the inoculum at  $T_0$ . The recovery of both influenza A virus strains was compared using an MLE (not shown), which determined that the mean log<sub>10</sub> change in recovery across all AHs and times was -0.44 (standard deviation [SD], 0.85) for A/NC-H1N1 and -1.30 (SD, 0.87) for A/Br-H1N1. Due to the significant (*P* < 0.001) difference in recovery between the two influenza A virus strains, where a greater change in log<sub>10</sub> recovery was experienced for A/Br-H1N1 than for A/NC-H1N1, the

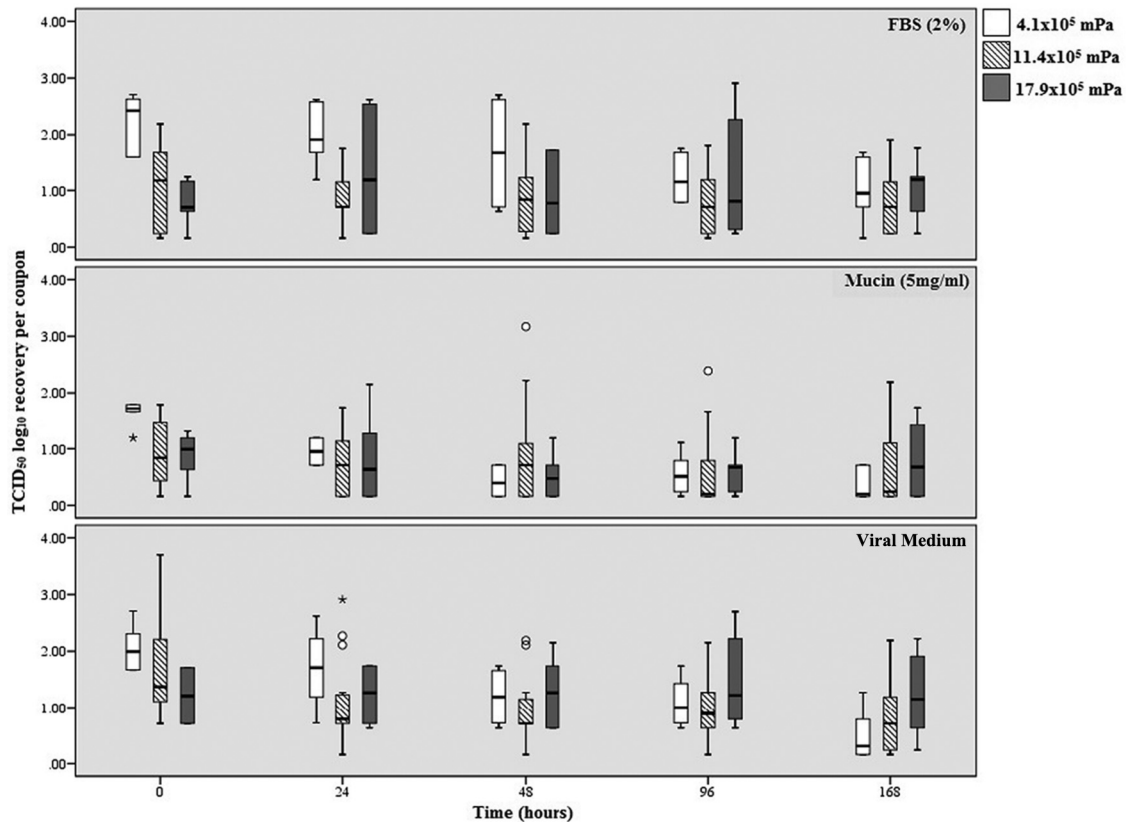


FIG 1 The  $\log_{10}$  TCID<sub>50</sub> recovery of the A/NC-H1N1 strain per coupon over time for the three different matrices and absolute humidities. The middle bar within each box represents the median; the top and bottom of the bar represent the 75th and 25th quartiles, respectively; and the error bars represent the 95% confidence intervals. Open circles represent outliers, and asterisks represent extreme outliers (three times the range of variation).

remainder of the data and statistical analyses were performed by consideration of the two strains independently.

The median  $\log_{10}$  recovery of infectious A/NC-H1N1 is presented in Fig. 1 for each matrix (FBS [2%], mucin [5 mg/ml], viral medium), time point ( $T_0$ ,  $T_{24}$ ,  $T_{48}$ ,  $T_{96}$ ,  $T_{168}$ ), and AH ( $4.1 \times 10^5$  mPa,  $11.4 \times 10^5$  mPa,  $17.9 \times 10^5$  mPa). The median  $\log_{10}$  range of recovery for virus held at an AH of  $4.1 \times 10^5$  mPa was 1.01 to 2.23 for virus held in FBS, 0.36 to 1.65 for virus held in mucin, and 0.50 to 2.06 for virus held in viral medium. The median  $\log_{10}$  range of recovery for virus held at an AH of  $11.4 \times 10^5$  mPa was 0.79 to 1.08 for virus held in FBS, 0.62 to 0.95 for virus held in mucin, and 0.90 to 1.69 for virus held in viral medium. The median  $\log_{10}$  range of recovery for virus held at an AH of  $17.9 \times 10^5$  mPa was 0.77 to 1.22 for virus held in FBS, 0.81 to 1.21 for virus held in mucin, and 0.88 to 1.04 for virus held in viral medium. The mean  $\log_{10}$  change in recovery for infectious A/NC-H1N1 compared to the mean number of TCID<sub>50</sub> of virus in the inoculum at  $T_0$  for each of the parameters is shown in Table 1. The median  $\log_{10}$  recovery of influenza A virus at  $T_0$  was higher when it was held at an AH of  $4.1 \times 10^5$  mPa than when it was held at AHs of  $11.4 \times 10^5$  mPa and  $17.9 \times 10^5$  mPa for all three substrates. Virus held at an AH of  $17.9 \times 10^5$  mPa was the most stable across all matrices and times, whereas virus held at an AH of  $4.1 \times 10^5$  mPa experienced the greatest loss in  $\log_{10}$  recovery across all three matrices.

The median  $\log_{10}$  recovery of A/Br-H1N1 is presented in Fig. 2 for each matrix (FBS [2%], mucin [5 mg/ml], viral medium), time point ( $T_0$ ,  $T_{24}$ ,  $T_{48}$ ,  $T_{96}$ ,  $T_{168}$ ), and AH ( $4.1 \times 10^5$  mPa,  $11.4 \times 10^5$

mPa,  $17.9 \times 10^5$  mPa). The median  $\log_{10}$  range of recovery for virus held at  $4.1 \times 10^5$  mPa was 1.53 to 2.83 for virus held in FBS, 1.60 to 2.23 for virus held in mucin, and 1.84 to 2.73 for virus held in viral medium. The median  $\log_{10}$  range of recovery for virus held

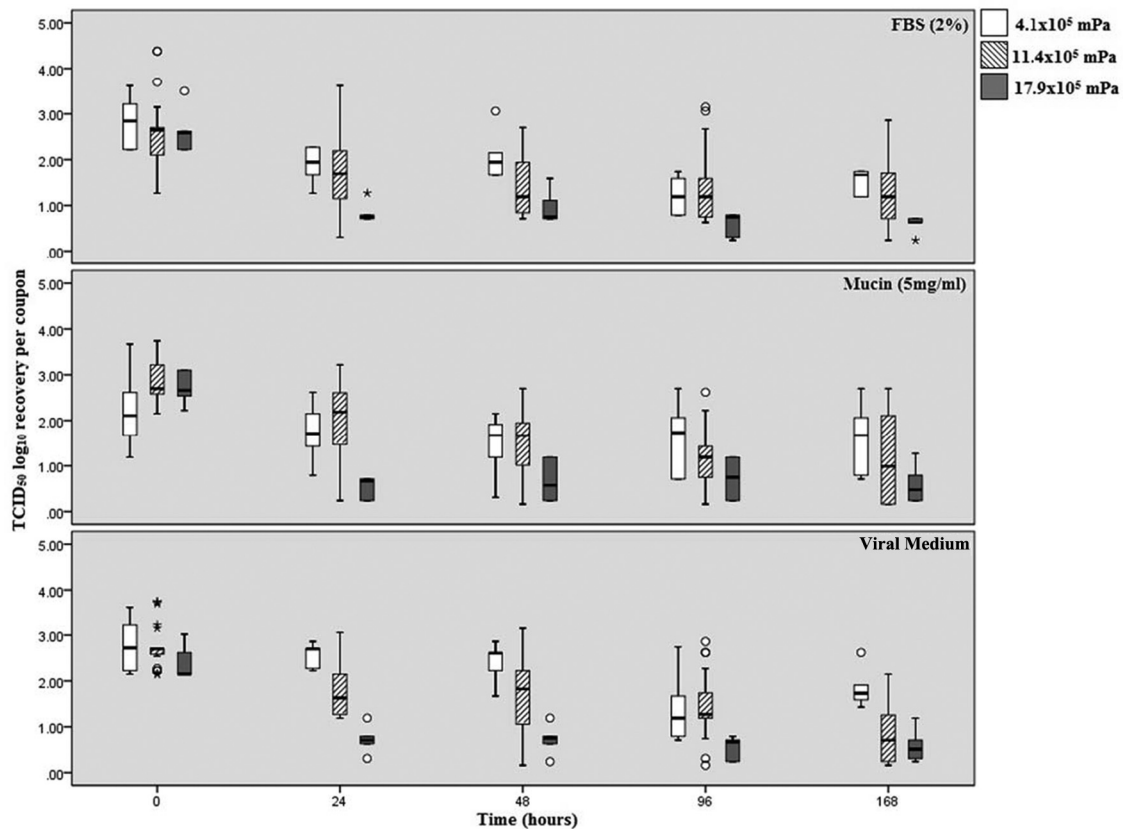
TABLE 1 Change in  $\log_{10}$  TCID<sub>50</sub> recovery of A/NC-H1N1 per SS coupon over time for each matrix and absolute humidity

A/NC-H1N1 sample matrix	Time	Mean change in $\log_{10}$ TCID <sub>50</sub> <sup>a</sup> (SD) at an absolute humidity of:		
		$4.1 \times 10^5$ mPa <sup>b</sup>	$11.4 \times 10^5$ mPa <sup>c</sup>	$17.9 \times 10^5$ mPa <sup>b</sup>
FBS	$T_{24}$	-0.25 (0.7)	-0.23 (0.6)	0.57 (1.2)
	$T_{48}$	-0.56 (0.7)	-0.18 (0.8)	0.14 (0.8)
	$T_{96}$	-1.01 (0.4)	-0.31 (0.8)	0.46 (1.3)
	$T_{168}$	-1.22 (0.5)	-0.32 (0.9)	0.28 (0.7)
Mucin	$T_{24}$	-0.70 (0.2)	-0.23 (0.7)	-0.04 (0.5)
	$T_{48}$	-1.23 (0.5)	-0.16 (0.9)	-0.35 (0.3)
	$T_{96}$	-1.10 (0.5)	-0.40 (0.8)	-0.27 (0.4)
	$T_{168}$	-1.29 (0.3)	-0.36 (0.8)	-0.07 (0.4)
Viral medium	$T_{24}$	-0.36 (0.3)	-0.66 (0.7)	0.01 (0.1)
	$T_{48}$	-0.87 (0.3)	-0.75 (0.8)	0.07 (0.2)
	$T_{96}$	-0.97 (0.1)	-0.77 (1.2)	0.25 (0.5)
	$T_{168}$	-1.56 (0.5)	-1.01 (1.1)	0.01 (0.4)

<sup>a</sup> Data are relative to the number of TCID<sub>50</sub> in the inoculum at  $T_0$ .

<sup>b</sup>  $n = 6$ .

<sup>c</sup>  $n = 24$ .



**FIG 2** The  $\log_{10}$  TCID<sub>50</sub> recovery of the A/Br-H1N1 strain per coupon over time for the three different matrices and absolute humidities. The middle bar within each box represents the median; the top and bottom of the bar represent the 75th and 25th quartiles, respectively; and the error bars represent the 95% confidence intervals. Open circles represent outliers, and asterisks represent extreme outliers (three times the range of variation).

at  $11.4 \times 10^5$  mPa was 2.29 to 2.61 for virus held in FBS, 1.19 to 2.84 for virus held in mucin, and 0.78 to 2.75 for virus held in viral medium. The median  $\log_{10}$  range of recovery for virus held at  $17.9 \times 10^5$  mPa was 0.59 to 2.71 for virus held in FBS, 0.58 to 2.37 for virus held in mucin, and 0.58 to 2.62 for virus held in viral medium. The mean  $\log_{10}$  change in recovery of A/Br-H1N1 compared to the mean number of TCID<sub>50</sub> of virus in the inoculum at  $T_0$  for each of the parameters is shown in Table 2. As for A/NC-H1N1,  $\log_{10}$  recovery was the most stable at  $17.9 \times 10^5$  mPa. In addition, influenza A virus held at an AH of  $11.4 \times 10^5$  mPa experienced the greatest change in  $\log_{10}$  recovery, with only samples in viral medium demonstrating the worst recovery, though samples in 2% FBS and 5 mg/ml mucin also demonstrated similarly low recoveries over time.

A GEE multivariate analysis was created using the cumulative  $\log_{10}$  change in recovery for each parameter when modeled concurrently, and the results are shown in Table 3. The model showed that AH, the AH-strain interaction, and time were factors that significantly impacted the overall change in  $\log_{10}$  recovery ( $P < 0.0001$ ) and that the virus strain did not impact persistence ( $P = 0.45$ ). An MLE (not shown) showed that the sample matrix did not have an impact on influenza A virus persistence and therefore was not included in the GEE analysis. When the AH was increased from  $4.1 \times 10^5$  mPa to  $17.9 \times 10^5$  mPa, there was a significant ( $P < 0.0001$ ) cumulative decrease in  $\log_{10}$  recovery ( $-1.74 \log_{10}$ ). Additionally, when the increase in AH occurred in combination with the A/NC-H1N1 strain, there was a significant ( $P < 0.0001$ )

cumulative  $\log_{10}$  increase (1.41  $\log_{10}$ ) compared to that for the A/Br-H1N1 strain. When time was modeled, samples had to be held for 7 days ( $T_{168}$ ) before a significant ( $P = 0.0013$ ) cumulative  $\log_{10}$  decrease in recovery ( $-1.05 \log_{10}$ ) was observed.

**TABLE 2** Change in  $\log_{10}$  TCID<sub>50</sub> recovery of A/Br-H1N1 per SS coupon over time for each matrix and absolute humidity

A/Br-H1N1 sample matrix	Time	Mean change in $\log_{10}$ TCID <sub>50</sub> <sup>a</sup> (SD) at an absolute humidity of:		
		$4.1 \times 10^5$ mPa <sup>b</sup>	$11.4 \times 10^5$ mPa <sup>c</sup>	$17.9 \times 10^5$ mPa <sup>b</sup>
FBS	$T_{24}$	-0.93 (0.3)	-0.80 (0.5)	-1.79 (0.5)
	$T_{48}$	-0.76 (0.5)	-1.16 (0.8)	-1.68 (0.6)
	$T_{96}$	-1.61 (0.2)	-1.25 (0.7)	-2.02 (0.5)
	$T_{168}$	-1.30 (0.7)	-1.32 (1.2)	-2.03 (0.6)
Mucin	$T_{24}$	-0.49 (0.5)	-0.84 (0.8)	-2.17 (0.4)
	$T_{48}$	-0.74 (1.3)	-1.33 (0.9)	-2.05 (0.2)
	$T_{96}$	-0.62 (1.4)	-1.69 (0.7)	-1.99 (0.3)
	$T_{168}$	-0.63 (1.4)	-1.65 (1.1)	-2.13 (0.4)
Viral medium	$T_{24}$	-0.20 (0.6)	-0.94 (0.5)	-1.65 (0.3)
	$T_{48}$	-0.34 (0.7)	-0.99 (0.9)	-1.65 (0.5)
	$T_{96}$	-1.39 (0.7)	-1.27 (0.8)	-1.82 (0.4)
	$T_{168}$	-0.94 (0.6)	-1.95 (0.6)	-1.80 (0.3)

<sup>a</sup> Data are relative to the number of TCID<sub>500</sub> in the inoculum at  $T_0$ .

<sup>b</sup>  $n = 6$ .

<sup>c</sup>  $n = 24$ .

TABLE 3 GEE analysis of the recovery of influenza A (H1N1) virus on SS coupons<sup>a</sup>

Parameter	Cumulative log <sub>10</sub> change	Estimate (SE)	Confidence limit	P value (α) <sup>c</sup>
Intercept		-0.67 (0.11)	-0.88 to -0.46	<0.0001
AH				
4.1 × 10 <sup>5</sup> mPa <sup>b</sup>				
11.4 × 10 <sup>5</sup> mPa	-1.15	-0.44 (0.13)	-0.69 to 0.18	<0.0001
17.9 × 10 <sup>5</sup> mPa	-1.74	-1.07 (0.12)	-1.29 to -0.84	<0.0001
Strain				
A/NC-H1N1	-0.77	-0.10 (0.13)	-0.35 to 0.15	0.45
A/Br-H1N1 <sup>b</sup>				
Interactions				
4.1-A/NC-H1N1 <sup>b</sup>				
4.1-A/Br-H1N1 <sup>b</sup>				
11.4-A/NC-H1N1	0.24	0.92 (0.18)	0.57 to 1.27	<0.0001
11.4-A/Br-H1N1 <sup>b</sup>				
17.9-A/NC-H1N1	1.41	2.08 (0.17)	1.76 to 2.41	<0.0001
17.9-A/Br-H1N1 <sup>b</sup>				
Time point				
24-48 h <sup>b</sup>				
96 h	-0.93	-0.26 (0.10)	-0.46 to -0.06	0.01
168 h	-1.05	-0.38 (0.12)	-0.62 to -0.15	0.0013

<sup>a</sup> The GEE can be used to calculate the cumulative change in the log<sub>10</sub> TCID<sub>50</sub> by adding the estimate for an individual parameter to the intercept value. A further model estimation can be obtained by combining the intercept (-0.67) with parameters in a given scenario; for example, at 11.4 × 10<sup>5</sup> mPa, the combination of the AH estimate (-0.44) with the A/NC-H1N1 estimate (-0.10) for 168 h (-0.38) results in a -1.59 cumulative log<sub>10</sub> change in the TCID<sub>50</sub> of influenza A (H1N1) virus recovered from stainless steel.

<sup>b</sup> This group is the referent group and is reflective of the intercept.

<sup>c</sup> The significance limit was a P value of <0.01.

## DISCUSSION

Our study demonstrates that influenza A (H1N1) virus strains A/NC-H1N1 and A/Br-H1N1 persisted and remained infectious on SS coupons for 7 days ( $T_{168}$ ) under the various AH conditions. Overall, A/NC-H1N1 experienced a 1.5-log<sub>10</sub> decrease in recovery (Table 1) across the 7 days ( $T_{168}$ ), while A/Br-H1N1 experienced a 2-log<sub>10</sub> decrease in recovery (Table 2) across the 7 days ( $T_{168}$ ). The GEE model (Table 3) revealed that AH, the interaction between AH and the virus strain, and time were the factors that significantly affected the ability of influenza A virus to persist on SS. The sample matrix did not influence viral persistence.

The results from this study contradict those from previously

published research. For example, it has previously been demonstrated that there is an increased persistence of influenza A virus in the presence of mucus (25, 26). However, in our study, the sample matrix (e.g., viral medium, FBS, mucin) did not influence influenza A virus persistence, which was also demonstrated by Coulliette et al. (22). It is important to note that the source of mucus for all three studies was different and data are not available to explain how this may impact virus persistence. Additional differences in persistence between our study and previous studies, which are summarized in Table 4, may be attributed to differences in the influenza A virus strains, viral surface recovery methods, quantification methods, sample matrices, and environmental conditions used.

TABLE 4 Summary of published research and our study on the persistence of influenza A virus on SS surfaces

Influenza A virus strain	AH (mPa)	Temp (°C)	RH (%)	Detection methodology	Log <sub>10</sub> loss of recovery	Time to loss (h)	Sample size	Authors (reference)
A/Brazil/11/78-like	10.8 × 10 <sup>5</sup>	27.8-28.3	35-50	CPE <sup>a</sup>	3.5	48	NS <sup>b</sup>	Bean et al. (15)
A/PuertoRico/8/34	13.2 × 10 <sup>5</sup>	25.2	55	CPE	3	24	2	Sakaguchi et al. (19)
A/PuertoRico/8/34 and A/Cambridge/AH04/2009	3.8 × 10 <sup>5</sup>	17-21	23-24	Plaque assay, PCR	>4.2	24	NS	Greatorex et al. (16)
Novel H1N1 strain (untyped)	NS	NS	NS	Antigen detection assay	>5	0.5	2	Mukherjee et al. (17)
A/California/7/2009	NS	NS	NS	HA <sup>c</sup>	>5.5	48	4	Oxford et al. (18)
A/New Caledonia/20/1999 and A/Brisbane/59/2007	4.1 × 10 <sup>5</sup> , 11.4 × 10 <sup>5</sup> , and 17.9 × 10 <sup>5</sup> <sup>c</sup>	18-25	20-55	ELISA <sup>d</sup>	2	168	1	Perry et al. (this study)

<sup>a</sup> CPE, cytopathic effect assay.

<sup>b</sup> NS, not stated.

<sup>c</sup> HA, hemagglutination assay.

<sup>d</sup> ELISA, enzyme-linked immunosorbent assay.

<sup>e</sup> See Materials and Methods for defined temperature and relative humidity ranges.

The aforementioned previous studies (Table 4) used different strains of influenza A viruses that represent virus strains from various regions of the world that were collected at different times over a period spanning decades. Influenza virus is known for rapidly mutating, whether it is due to natural mutation or host immune selective pressure (27). This mutation could contribute to the variability between virus strains, rendering it difficult to make a general conclusion regarding persistence. In our own study, a maximum likelihood estimate (not shown) that compared the cumulative change in  $\log_{10}$  recovery determined that a significant difference in persistence between the A/NC-H1N1 and A/Br-H1N1 strains ( $P \leq 0.0001$ ) existed. However, when the recoveries of both virus strains were simultaneously compared against all parameters in the GEE model (Table 3), there was no significant difference in recovery between the strains ( $P = 0.45$ ). The A/NC-H1N1 strain had a greater decrease in  $\log_{10}$  recovery when it was held at  $4.1 \times 10^5$  mPa ( $-1.56 \log_{10}$  [SD,  $0.5 \log_{10}$ ]), while A/Br-H1N1 had a greater decrease in  $\log_{10}$  recovery when it was held at  $17.9 \times 10^5$  mPa ( $-2.17 \log_{10}$  [SD,  $0.4 \log_{10}$ ]). The differences in persistence between virus strains that were seen may be attributed to slight variations within the fatty acid composition in the outer phospholipid membrane of the influenza A viruses. The lipid content of influenza viruses can range from 20 to 24% (28), and it has been observed that viruses with greater lipid contents are more stable at lower RHs (29). Studies that compared the lipid contents of influenza A and B virus strains, as well as those of two influenza A virus strains, where the lipid content of one was made incomplete, have demonstrated that these slight differences do exist (30, 31). Additionally, influenza A virus strain A/Puerto Rico/8/34 has been shown to resist desiccation on surfaces for a week or longer at ambient temperature (32). This may be in part due to the protective nature of the envelope and an ability to form clumps on surfaces (29).

Methods previously used to recover influenza virus from environmental surfaces included a premoistened swab (15–17), surface rinsing (18), or a combination of residual liquid and a premoistened swab (19). To date, no recommended, standardized, or validated guidelines that describe the best methodology for recovery of viruses from any type of environmental surface are available. Therefore, it is difficult to say if a method used in a study to recover influenza A virus from environmental surfaces is the most efficient. In addition, cell culture-based methods, such as the CPE assay (15, 19), the plaque assay (16), and the hemagglutination (HA) assay (18), as well as molecular methods, like PCR (7, 14, 16), were used in previous research to determine the persistence of influenza A virus. While CPE and plaque assays can extrapolate infectivity and viability, HA assays and PCR cannot, since surface proteins and RNA can be present even after the virus is no longer viable. Our study used a tissue culture-based ELISA, which helped improve the sensitivity and removed subjectivity from interpretation of the results, as it is a direct measure of viral presence and infectivity by immunofluorescence. An analysis and/or side-by-side evaluation of methods for the recovery and determination of infectivity of influenza virus was not the focus of this study, although such an analysis would be a significant contribution to this field.

It is important to note that our study evaluated the viral persistence of different influenza virus strains in various sample matrices and environmental conditions on a nonporous surface. These combined parameters have not been previously investi-

gated using a tissue culture-based ELISA. Although the two strains chosen for use in this study no longer circulate in humans, they act as a surrogate for the pandemic strain, which has been shown to have survival characteristics similar to those of the strains used in this study (22). Previous studies of virus persistence on environmental surfaces have found that various strains of influenza A (H1N1) virus remain infectious on stainless steel surfaces for approximately 24 h but no longer than 72 h (15–19). Inactivation studies found a minimal decrease in infectious influenza A (H1N1) virus particles on stainless steel surfaces at room temperature and 50 to 60% RH (33) but that the virus was rapidly inactivated under conditions of high heat (temperature range, 55 to 65°C) and humidity (RH range, 25 to 75%) (34). Initially, it was indicated that temperature and RH are key factors that influence viral persistence and transmission (35, 36) and that influenza virus has higher levels of persistence and increased rates of transmission at lower temperatures and RHs (29, 34–37). Recent studies have found that AH, which is an actual measure of the water vapor content of air, is better suited for determining the influence of environmental factors on viral persistence (34, 38, 39).

Overall, our study showed that two influenza A (H1N1) virus strains tested under a range of AH conditions that represent the different conditions of indoor and outdoor settings remained infectious on SS for at least 7 days, with less than a 2- $\log_{10}$  decrease in recovery, as measured by infectivity, being found. Further research is needed to determine the role of contaminated environmental surfaces on the transmission of influenza A virus so that improved infection control guidelines can be devised and implemented.

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