

## Robustness of the ferret model for influenza risk assessment studies: a cross-laboratory exercise

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## 1    Abstract

2    Ferrets represent the preferred animal model for assessing the transmission potential of newly  
3    emerged zoonotic influenza viruses. However, heterogeneity among established experimental  
4    protocols and facilities across different laboratories may lead to variable results, complicating  
5    interpretation of transmission experimental data. Between 2018-2020, a global exercise was  
6    conducted by 11 participating laboratories to assess the range of variation in ferret transmission  
7    experiments using two common stock H1N1 influenza viruses that possess different transmission  
8    characteristics in ferrets. Inoculation route, dose, and volume were standardized, and all  
9    participating laboratories followed the same experimental conditions for respiratory droplet  
10   transmission, including a strict 1:1 donor:contact ratio. Additional host and environmental  
11   parameters likely to affect influenza transmission kinetics were monitored throughout. Overall  
12   transmission outcomes for both viruses across 11 laboratories were concordant, suggesting the  
13   robustness of the ferret model for zoonotic influenza risk assessment. To attain high confidence in  
14   identifying zoonotic influenza viruses with moderate-to-high or low transmissibility, our analyses  
15   support that as few as three but as many as five laboratories, respectively, would need to  
16   independently perform viral transmission experiments with concordant results. This exercise  
17   facilitates the development of a more homogenous protocol for ferret transmission experiments  
18   that are employed for the purposes of risk assessment.

19 **Introduction**

20 Pandemic influenza viruses may arise through interspecies transmission events of animal  
21 influenza viruses. Assessing the human-to-human transmission potential of animal influenza  
22 viruses that cause spillover infections in humans is essential for pandemic risk assessment. Ferrets  
23 have been used as a surrogate model for investigating the transmission potential of influenza  
24 viruses in humans, as they are naturally susceptible to infection with human and zoonotic influenza  
25 viruses, exhibit clinical signs during infection which closely resemble those of humans, and  
26 support influenza virus transmission via similar modes as humans. In particular, the respiratory  
27 droplet transmissibility of a specific influenza strain among ferrets often correlates with its  
28 transmission potential in humans (1). As such, ferrets are commonly used for assessing the  
29 pandemic potential of newly emerged zoonotic influenza viruses, and data from these experiments  
30 inform formal risk assessment rubrics established by the WHO and CDC (2, 3).

31 The transmission potential of influenza viruses is determined by multiple viral, host, and  
32 environmental parameters. As the ferret model becomes commonly employed in laboratories  
33 worldwide, there is an underappreciated heterogeneity among established experimental protocols  
34 and facility setups across different laboratories, which may lead to variable results between  
35 transmission experiments performed (4). Some of these variables, such as the dose, volume, and  
36 route of inoculation and animal age, have been confirmed to affect the kinetics of virus infection,  
37 replication, and transmission in the ferret model (5-7). However, the impact of other parameters,  
38 such as virus propagation procedures, caging designs, airflow directionality and number of air  
39 exchanges, and environmental conditions such as relative humidity, is largely unknown.  
40 Consequently, interpretation of results from ferret transmission experiments can represent a  
41 challenge when comparing data generated from multiple laboratories, even when the same virus

42 strain or subtype is being investigated (8). Considering the statistical limitations on small sample  
43 sizes in ferret experiments, and high potential for strain-specific variability, investigators often  
44 assess the pandemic potential of emerging virus subtypes as an aggregate of multiple viruses tested  
45 (9-11). As many public health efforts require cross-laboratory risk assessment studies for newly  
46 emerged zoonotic influenza viruses (12) and antiviral efficacy studies aiming to block influenza  
47 transmission between ferrets (13), a greater understanding of variability in transmission results  
48 obtained between independent groups is critical.

49 To assess the variability of ferret transmission results across laboratories under established  
50 protocols, we performed a global exercise using two common stock influenza viruses that possess  
51 different transmission characteristics in ferrets. Eleven independent laboratories inoculated ferrets  
52 with these stock viruses under uniform conditions; parameters known to affect influenza  
53 transmission kinetics were controlled in the experimental protocols while other potential  
54 parameters were carefully monitored and recorded, both prior to and during the transmission  
55 experiments. All aggregated data from these experiments were blinded and analyzed by an  
56 independent statistician. To inform future risk assessment activities, the confidence of drawing  
57 conclusions on virus transmissibility with concordant or discordant outcomes from multiple  
58 laboratories was also investigated. By assessing the range of variation present among ferret  
59 transmission experiments performed under established experimental protocols, this global exercise  
60 provides helpful guidance for data interpretation when cross-laboratory results are to be compared.  
61 The relatively concordant transmission results across 11 laboratories suggest that the ferret model  
62 is highly robust for influenza pandemic risk assessment studies under the semi-standardized  
63 conditions employed here. Furthermore, analyses investigating the role of host and environmental  
64 parameters as they contribute to virus transmission kinetics and outcomes is valuable for both

65 current risk assessment activities and for evaluation of countermeasures to block influenza  
66 transmission.

67

68 **Results**

69 **Transmissibility of human A(H1N1)pdm09 virus.** To evaluate potential heterogeneity in the  
70 transmission results between 11 laboratories, we first compared the transmissibility of a cell-grown  
71 isolate of the A(H1N1)pdm09 virus A/California/7/2009 (Cal/09), representative of early 2009  
72 pandemic isolates and anticipated to exhibit moderate to high respiratory droplet transmissibility  
73 (14-17). Transmissibility was evaluated with 4 donor:contact pairs at a 1:1 ratio in each laboratory.  
74 Transmission to exposed respiratory droplet contact ferrets was defined by detection of infectious  
75 virus or seroconversion to the homologous virus in post-exposure sera. Following establishment  
76 of contact with donor ferrets 24 hours post-inoculation, detection of infectious virus and  
77 seroconversion in contacts was observed in 10/11 and 11/11 laboratories, respectively, with the  
78 reported transmission frequency ranging from 50-100% (Table 1). One out of 11 laboratories  
79 determined viral loads in nasal swabs and throat swabs (Group F, with throat swab viral loads used  
80 for subsequent analysis), while the other laboratories determined viral loads in nasal washes.  
81 Employing both virological and serological results, by Fisher's exact test of homogeneity, there  
82 was no significant difference in the transmission outcomes across labs with this virus ( $p=0.797$ ).  
83 Collectively, infectious virus was detected from the nasal wash or throat swabs of 72.7% (32/44)  
84 exposed contacts, and seroconversion of contact ferrets to homologous virus was detected from  
85 79.5% (35/44) of exposed contacts.

86

87

88 **Table 1. Summary of virus transmissibility results from all laboratories.**

Group	A(H1N1)pdm09 virus, A/California/7/2009			A(H1N1) avian influenza virus, A/ruddy turnstone/Delaware/300/20/2009		
	Viral load of inoculated donors (area under the curve) <sup>a</sup>	Transmission to aerosol contacts		Viral load of inoculated donors (area under the curve) <sup>a</sup>	Transmission to aerosol contacts	
		Virus detection <sup>b</sup>	Sero- conversion <sup>c</sup>		Virus detection <sup>b</sup>	Sero- conversion <sup>c</sup>
A	6.51±0.49	3/4	3/4	4.28±0.35	0/4	0/4
B	5.70±0.42	4/4	4/4	4.25±0.39	1/4	1/4
C	5.30±0.78	4/4	4/4	5.10±0.11	1/4	1/4
D	6.86±0.40	2/4	2/4	5.73±0.21	0/4	0/4
E	5.53±0.32	3/4	3/4	4.43±0.56	0/4	0/4
F	5.77±0.60	3/4	3/4	5.34±0.60	0/4	0/3
G	6.57±0.06	2/4	2/4	6.48±0.37	0/4	0/4
H	5.82±0.43	0/4	3/4	4.72±0.31	0/4	0/4
I	6.48±0.80	3/4	3/4	6.24±0.31	0/4	0/4
J	5.62±0.54	4/4	4/4	4.92±0.39	1/4	1/4
K	4.07±0.72	4/4	4/4	4.15±0.57	3/4	3/4

89 <sup>a</sup>Area under the curve (AUC) of viral titers from inoculated ferrets (following normalization of  
90 infectious units to TCID<sub>50</sub>), expressed as log<sub>10</sub> mean ± standard deviation. <sup>b</sup>Number of contact  
91 ferrets with infectious virus detected in respiratory specimens between 1-13 days post-contact  
92 (p.c.). <sup>c</sup>Number of contact ferrets that seroconverted to the exposed virus at the end of the study  
93 using hemagglutination inhibition assay.

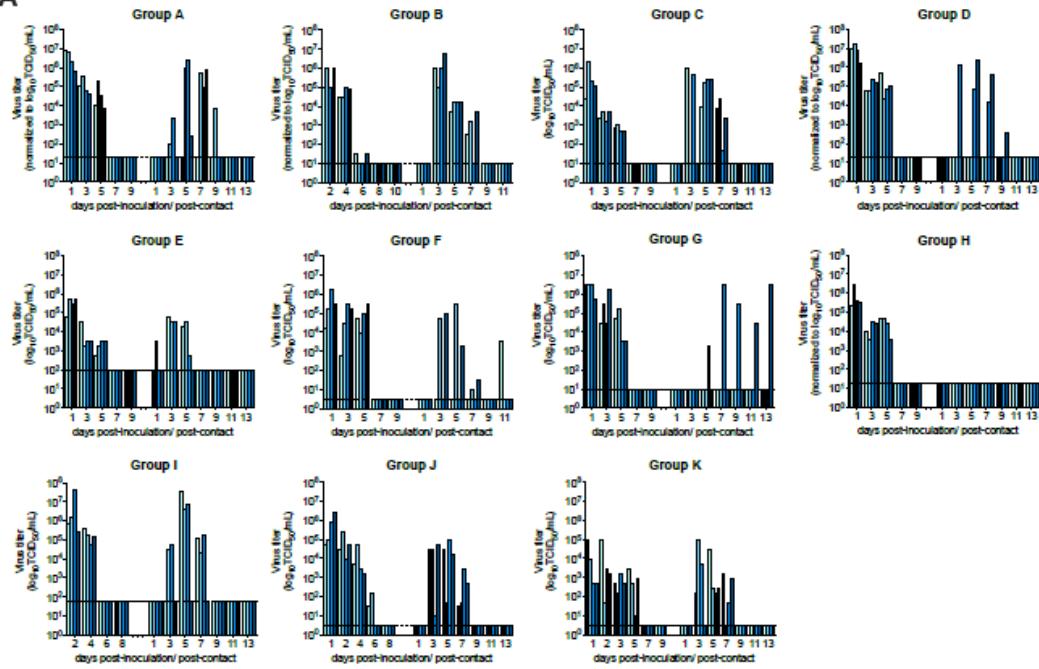
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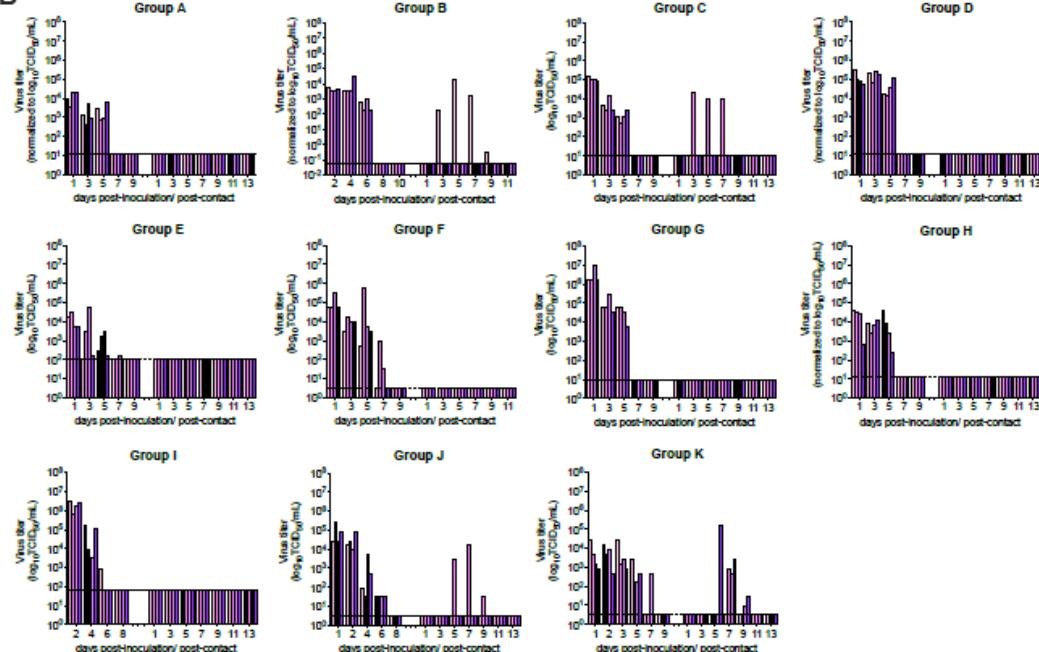
96 To allow comparison of the effect of viral load on transmissibility, viral titer units from  
97 nasal wash/throat swab samples (inclusive of TCID<sub>50</sub>, PFU, and EID<sub>50</sub> units, Supplemental Figures  
98 1-2) were normalized to TCID<sub>50</sub> units (Figure 1), employing strain-specific conversions prior to  
99 analyses (Supplemental Table 1). From the inoculated donor ferrets, the peak viral titers detected  
100 in the nasal washes or throat swabs were at 5.72 ± 0.95, mean ± SD log<sub>10</sub> TCID<sub>50</sub>/mL after  
101 normalization, with the peak titers detected from 95.5% (42/44) of donors on the first sampling  
102 time point on 1 or 2 days post-inoculation (dpi), followed by a decline of infectious titer over time  
103 (Figure 1A). Area under the curve (AUC) after normalization was calculated to approximate total  
viral load shed by the Cal/09-inoculated donors, with a mean ± SD log<sub>10</sub> AUC of 5.84 ± 0.89.

104 **Figure 1. Transmission kinetics of A(H1N1) viruses in ferrets.** **A**, normalized viral loads of  
105 donors (left bars) and aerosol contact ferrets (right bars) after inoculation or exposure to  
106 A(H1N1)pdm09 virus Cal/09. **B**, normalized viral loads of donors (left bars) and aerosol contact  
107 ferrets (right bars) after inoculation or exposure to avian H1N1 virus ruddy turnstone/09. Nasal  
108 washes (all groups except Group F) or throat swabs (Group F) were sampled to determine  
109 infectious viral loads which were normalized to  $\log_{10}$  TCID<sub>50</sub>/mL. Each bar represents individual  
110 ferrets. Limit of detection is indicated with a dashed line.

**A**



**B**

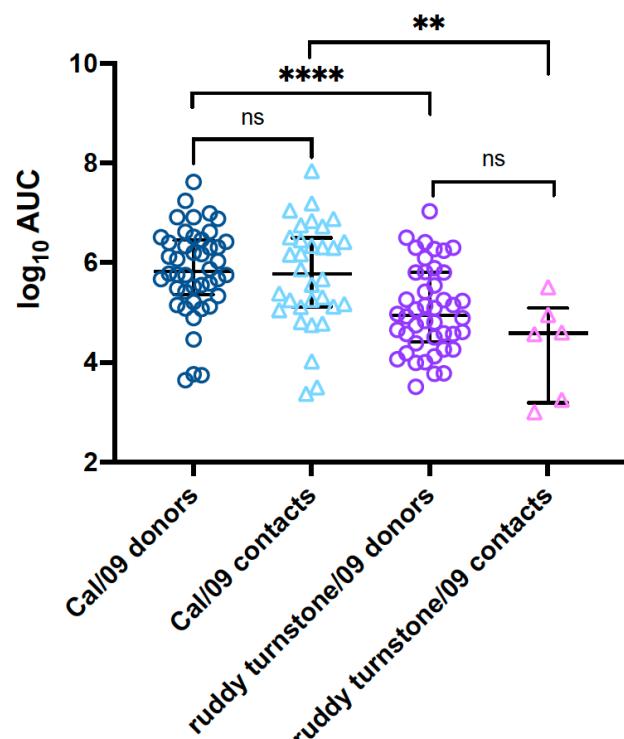


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112 Next, to evaluate the transmission efficiency, the serial interval (first detection of viral  
113 shedding in contacts post-exposure from specimens collected every-other-day) was calculated for  
114 each infected contact ferret. The serial interval was 1 day for 3.1% (1/32) of the Cal/09 infected  
115 contact ferrets, followed by 3 days for 68.8% (22/32), 5 days for 21.9% (7/32), and 11 days for  
116 6.3% (2/32), with a median serial interval of 3 days post-contact. Peak viral titers detected in the  
117 contact nasal washes or throat swabs were at  $5.41 \pm 1.06$  mean  $\pm$  SD  $\log_{10}$  TCID<sub>50</sub>/mL after  
118 normalization, with peak titers detected from 50% (16/32) and 34.4% (11/32) infected contacts on  
119 3 dpi and 5 dpi, respectively. Altogether, the AUC for Cal/09 infected contact ferrets was  $5.75 \pm$   
120 1.05, comparable to the Cal/09 virus-inoculated donors (Mann-Whitney test,  $p=0.6547$ ) (Figure  
121 2).

122

123 **Figure 2. Area under the curve of infectious viral loads detected from inoculated donors or**  
124 **infected contacts.** Data points represent AUC values from individual ferrets from which  
125 infectious virus was detected. \*\*,  $p < 0.01$ , \*\*\*\*,  $p < 0.0001$ , Mann-Whitney test.  
126



127

128

129 **Transmissibility of avian A(H1N1) influenza virus.** We further evaluated the range of  
130 heterogeneity present in transmission results when using the A(H1N1) avian influenza virus  
131 A/ruddy turnstone/Delaware/300/2009 (ruddy turnstone/09) (18, 19), which has been reported to  
132 transmit in ferrets via respiratory droplets under the experimental setting of donor: direct contact:  
133 respiratory droplet contact at a 1:1:1 ratio, but not at a 1:1 donor:respiratory droplet contact ratio  
134 (R Fouchier, unpublished data) (18, 19). Here, the experimental setup and conditions were  
135 identical to those assessing Cal/09 virus transmissibility including a donor: respiratory droplet  
136 contact 1:1 ratio with no direct contact ferret. Transmission of an egg-derived isolate of ruddy  
137 turnstone/09 virus to exposed respiratory droplet contacts was only observed in 4 out of the 11  
138 laboratories, with the transmission frequencies ranging from 25-75% across these four laboratories  
139 (Table 1). Compared to Cal/09 virus, there were greater differences in the ruddy turnstone/09 virus  
140 transmission outcomes across 11 laboratories, but the difference did not reach statistical  
141 significance by Fisher's exact test of homogeneity ( $p=0.068$ ). Viral shedding and seroconversion  
142 to ruddy turnstone/09 virus were detected from 6/43 exposed contact ferrets across all laboratories,  
143 resulting in a transmission efficiency of 14.0%, which was significantly lower compared to that of  
144 Cal/09 virus (72.3%, paired t-test,  $p < 0.001$ ).

145 From the inoculated donor ferrets, the peak viral titers detected in the nasal washes or throat  
146 swabs were at  $4.85 \pm 0.94$ , mean  $\pm$  SD  $\log_{10}$  TCID<sub>50</sub>/mL after normalization, which was  
147 significantly lower than those detected in the Cal/09 inoculated donors (Mann-Whitney test,  $p$   
148  $<0.0001$ ). Peak titers were detected from 88.6% (39/44) donors on the first sampling time point (1  
149 or 2 dpi) followed by a decline of infectious titer over time (Figure 1B). Compared with Cal/09  
150 virus inoculated donors, the mean  $\pm$  SD  $\log_{10}$  AUC of ruddy turnstone/09 virus-inoculated ferrets

151 was  $5.06 \pm 1.86$ , significantly lower than those inoculated with the Cal/09 virus (Mann-Whitney  
152 test,  $p < 0.0001$ ) (Figure 2). Overall, ruddy turnstone/09 virus-inoculated donor ferrets shed lower  
153 titers of infectious virus than the Cal/09 virus-inoculated donors.

154 In contrast to the transmission efficiency of Cal/09 virus with a median serial interval of 3  
155 days, for the ruddy turnstone/09 transmission experiments, the serial interval was 3 days, 5 days,  
156 or 7 days for 33.3% (2/6), 33.3% (2/6), and 33.3% (2/6) of the infected contact ferrets, respectively,  
157 with the median serial interval at 5 days. Peak viral loads ( $3.94 \pm 0.94$  mean  $\pm$  SD  $\log_{10}$   
158 TCID<sub>50</sub>/mL) detected from the six infected contact ferrets were lower when compared to the Cal/09  
159 infected contact ferrets (Mann-Whitney test,  $p=0.0022$ ). Peak titers were detected from 16.7%  
160 (1/6), 33.3% (2/6), and 50% (3/6) infected contacts on 3 dpi, 5 dpi, and 7 dpi, respectively.  
161 Furthermore, ruddy turnstone/09 virus-infected contact ferrets shed significantly less infectious  
162 virus ( $4.31 \pm 0.98$ , mean  $\pm$  SD  $\log_{10}$  AUC) when compared to those animals directly inoculated  
163 with Cal/09 virus (Mann-Whitney test,  $p=0.0033$ ) (Figure 2). Taken together, there was a longer  
164 serial interval and lower infectious virus shed by ruddy turnstone/09 virus-exposed contact ferrets  
165 when compared to those exposed to Cal/09 virus.

166  
167 **Contributing factors to ruddy turnstone/09 virus transmissibility.** By standardizing the source  
168 stock virus, dose and volume of inoculation, and donor-to-contact ratio, we show that while  
169 infrequent discordant results were documented, the transmission outcomes of Cal/09 and ruddy  
170 turnstone/09 viruses independently performed by 11 laboratories were in general concordant,  
171 despite variabilities in the laboratory settings that were not standardized in the experiments  
172 (Supplemental Tables 2-4). As the transmission outcomes for the highly transmissible Cal/09 virus  
173 were more concordant than the less transmissible ruddy turnstone/09 virus, we attempted to

174 examine if any variable, including those not standardized between laboratories, may have been  
175 associated with differences in ruddy turnstone/09 virus transmissibility results.

176 Univariable logistic regression was performed to first evaluate if donor viral shedding  
177 kinetics were linked to ruddy turnstone/09 virus transmission efficiency. However, examination  
178 of several parameters, including AUC ( $p=0.193$ ), peak viral titer ( $p=0.197$ ), and days to peak titer  
179 ( $p=0.473$ ), were not statistically associated with different transmission outcomes observed  
180 between laboratories (Supplemental Table 5), indicating that differences observed between  
181 laboratories were not attributable to virological measurements.

182 Numerous studies have indicated a role for environmental parameters in virus  
183 transmissibility (20, 21). Room temperature was generally consistent across all groups, with means  
184 of daily recordings within 3°C for all experiments performed (20.5-23.2°C, Supplemental Table  
185 4). In contrast, the relative humidity reported varied widely, with regard to both the range of daily  
186 readings reported during 14-day individual experiments (varying 1-60% between low and high  
187 readings) and the mean recordings over the entirety of each experiment (32.7% to 77.0%). Despite  
188 this variability, there was no statistically significant association between transmission of ruddy  
189 turnstone/09 virus and temperature, relative humidity, or absolute humidity (all  $p>0.3$ ,  
190 Supplemental Table 5).

191 Experimental cage setups varied widely between different groups, with extensive  
192 heterogeneity present with regard to cage dimensions, airflow directionality and air changes per  
193 hour, distance between cages, and other parameters (Supplemental Table 3). Groups employing  
194 caging with airflow directionality from inoculated to contact cages more frequently reported  
195 moderate to high transmissibility ( $\geq 50\%$ ) of both viruses compared with groups lacking this  
196 airflow directionality (6/6 vs 3/5 groups for Cal/09 virus, 3/6 vs 1/5 groups for ruddy turnstone/09

197 virus), however these findings did not reach statistical significance (both  $p>0.3$ , Supplemental  
198 Table 5). Other specific features of cage setups, including distance between inoculated and contact  
199 cages and air changes per hour (ACH) were also not statistically linked to the ruddy turnstone/09  
200 transmission outcomes (both  $p>0.4$ , Supplemental Table 5). Taken together, despite substantial  
201 heterogeneity in numerous non-standardized parameters in experimental setups employed between  
202 groups, no one feature was identified as modulating transmission outcomes to a significant degree.  
203

204 **Contributing factors to virus pathogenicity.** All ferrets inoculated with either Cal/09 or ruddy  
205 turnstone/09 were productively infected, however measurements of morbidity varied between  
206 groups for both viruses. Among Cal/09 virus-inoculated ferrets, mean maximum weight loss and  
207 peak rise in body temperature between groups ranged from <1.0-15.6% and 0.6-2.1°C,  
208 respectively (Supplemental Table 6, Supplemental Figure 3). Following ruddy turnstone/09 virus  
209 inoculation, infected ferrets generally exhibited greater mean maximum weight loss (up to 19.6%)  
210 and transient fevers (up to 3°C) (Supplemental Table 7, Supplemental Figure 4) compared to  
211 ferrets with Cal/09 virus infections; ruddy turnstone/09-inoculated ferrets reached humane  
212 experimental endpoints in 2/11 groups. The coefficient of variation between mean maximum  
213 weight loss reported between groups was generally similar (56% and 52% for Cal/09 and ruddy  
214 turnstone/09 viruses, respectively). No commonality with increased morbidity and ferret vendor,  
215 gender, or pre-inoculation body weight was identified. Furthermore, no association was found  
216 between morbidity and viral load (peak titer or AUC) or other environmental parameters, with the  
217 exception of room temperature (with higher mean room temperatures associated with greater mean  
218 weight loss) (Supplemental Table 8).

219

220 **Confidence in virus transmission results generated from multiple laboratories.** Collectively,  
221 the results from this exercise demonstrate a capacity for groups possessing differences in facilities  
222 designs and experimental protocols to report varying levels of relative transmissibility and  
223 pathogenicity following inoculation of ferrets with the same virus. To illustrate how confidence in  
224 risk assessments of virus transmissibility can increase as results from multiple groups are  
225 combined, we evaluated the hypothetical risk of a virus capable of moderate to high transmission  
226 (defined as  $p \geq 50\%$  transmission events per total pairs of ferrets as defined in Table 2) or non-  
227 transmissible (defined as  $p \leq 25\%$  transmission events). In these analyses, concordant results are  
228 defined as multiple groups identifying a virus exhibiting the same transmission capacity, and  
229 discordant results are defined as multiple groups identifying a virus with different transmission  
230 capacities, as defined above. By assuming concordant results across laboratories which permits  
231 pooling of all transmission outcomes, as few as three groups (12 pairs of ferrets) will yield a  
232 probability of over 80% to conclude moderate to high transmissibility when transmission was  
233 observed in at least half of all experiments, and a probability of over 85% to conclude low  
234 transmissibility when at most one transmission event was observed over all experiments.

235

236 **Table 2. Confidence in conclusions derived from pooled samples from multiple**  
237 **laboratories.**

No. labs <sup>a</sup>	No. transmission pairs tested	No. pairs with transmission <sup>b</sup>	Probability of moderate to high transmissibility <sup>c</sup>	No. pairs without transmission	Probability of low transmissibility <sup>d</sup>
1	4	4	99%	0	76%
		3	88%	1	37%
2	8	≥7	>99%	0	92%
		6	99%	1	70%
		5	92%	2	40%
		4	68%		
3	12	≥8	>99%	0	98%
		7	95%	1	87%
		6	82%	2	67%
				3	42%
4	16	≥10	>99%	0	99%
		9	97%	1	95%
		8	90%	2	84%
		7	73%	3	65%
				4	43%

238 <sup>a</sup>Number of laboratories providing results (assumes 4 pairs of donor:contact at 1:1 ratio per  
239 laboratory; transmission in each pair is an independent event). <sup>b</sup>Number of transmission events  
240 among total number of ferrets from all groups specified (defined as detection of infectious virus  
241 and seroconversion to homologous virus in contact ferrets). <sup>c</sup>Moderate to high transmissibility is  
242 defined as  $p \geq 50\%$  (eg.  $\geq 2$  infected out of 4 ferrets). <sup>d</sup>Low transmissibility is defined as  $p \leq 25\%$  (eg.  
243 0 or 1 infected out of 4 ferrets).

244

245       Alternatively, a voting system can be considered by first drawing a conclusion on  
246 transmissibility in each laboratory, with an overall conclusion drawn based on these ‘votes’ from  
247 multiple labs. When testing for moderate to high transmissibility, and assuming n=4 ferrets per  
248 laboratory, 3 laboratories are needed to conclude moderate to high transmissibility with confidence  
249 >90% if concordant results are obtained. In agreement with probabilities shown in Table 3, a  
250 greater number of laboratories contributing results are needed to demonstrate statistically  
251 significant results when testing for low transmissibility; to conclude low transmissibility with

252 >90% confidence, this would necessitate 5 contributing laboratories if concordant results are  
253 obtained. In this scenario, a greater number of contributing laboratories (or a greater number of  
254 donor:contact pairs per laboratory) would be required if the true transmission probability was  
255 higher for confirming low transmissibility, or when the true transmission probability was lower  
256 for confirming moderate to high transmissibility.

257

258 **Table 3. Confidence in conclusions derived from multiple laboratories considering a voting**  
259 **system.**

No. labs <sup>a</sup>	No. labs with concordant result <sup>b</sup>	Probability of moderate to high transmissibility <sup>c</sup>	Probability of low transmissibility <sup>d</sup>
1	1	76%	56%
2	2	87%	72%
	1 (discordant)	31%	18%
3	3	92%	81%
	2 (discordant)	48%	32%
4	4	95%	87%
	3 (discordant)	62%	45%
	2 (discordant)	23%	12%
5	5	97%	95%
	4 (discordant)	72%	56%
	3 (discordant)	35%	21%
6	6	98%	94%
	5 (discordant)	80%	65%
	4 (discordant)	47%	30%

260 <sup>a</sup>Number of laboratories providing votes on the transmissibility of the tested virus. Each  
261 laboratory will vote if the tested virus possess moderate to high transmissibility ( $p \geq 50\%$ , eg.  $\geq 2$   
262 infected out of 4 ferrets) or low transmissibility ( $p \leq 25\%$ , eg. 0 or 1 infected out of 4 ferrets)  
263 based on the experimental result. <sup>b</sup>Number of laboratories with concordant results on viral  
264 transmissibility, or the number of laboratories reaching concordant results when discordant  
265 results are included. <sup>c</sup>Moderate to high transmissibility is defined as  $p \geq 50\%$  (eg.  $\geq 2$  infected out  
266 of 4 ferrets). <sup>d</sup>Low transmissibility is defined as  $p \leq 25\%$  (eg. 0 or 1 infected out of 4 ferrets).

267

268 Despite generally consistent results between all groups in this exercise, discordant results  
269 are possible (Table 1), highlighting the need to better understand how to responsibly interpret and

270 account for these findings. As such, we also considered the scenario when discordant results  
271 between laboratories are recorded. To demonstrate moderate to high transmissibility, we found  
272 that 6 laboratories with 1 discordant result could still provide 80% confidence in the conclusion,  
273 while any discordant result significantly reduced confidence for concluding low transmissibility  
274 (Table 3). In both scenarios, if the results from different laboratories were more heterogeneous,  
275 the uncertainty around the conclusion from each lab increases and the overall confidence would  
276 decrease. This exercise is an illustration of the possible scenarios and confidence in drawing  
277 conclusions on transmissibility but would be affected by how moderate to high or low  
278 transmissibility were defined.

279

## 280 **Discussion**

281 The importance of the ferret model for influenza virus risk assessment studies cannot be  
282 understated (4, 22). Recent advances in molecular biology, aerobiology, genomics, and other areas  
283 highlight the ways the ferret model in general, and studies evaluating virus transmissibility by the  
284 airborne route specifically, continue to contribute towards our understanding of influenza viruses  
285 and the threat they pose to human health (23-25). However, as this model becomes more  
286 commonly employed in laboratories worldwide, there is a pressing need to capture the level of  
287 variability and heterogeneity intrinsic to this research. Cross-laboratory exercises have been  
288 employed in the past to evaluate the reproducibility of assays employed for influenza virus public  
289 health efforts (26), but no such exercise has been performed to date evaluating influenza virus  
290 transmissibility in the ferret. In this exercise, 11 laboratories across different continents  
291 independently evaluated the transmission potential of Cal/09 and ruddy turnstone/09 viruses with  
292 distinct transmission potential. With only a few experimental parameters (common virus stock,

293 standardized inoculation dose, route, volume, and the 1:1 donor:contact ratio) being controlled  
294 across the participating laboratories, we observed homogenous transmission outcomes (that is,  
295 outcomes did not differ statistically) across laboratories. Our results demonstrate the robustness of  
296 the ferret model in influenza risk assessment studies.

297 Risk assessment rubrics have thoroughly evaluated a wide scope of influenza A viruses,  
298 from viruses associated with poultry outbreaks in the absence of confirmed human infections, to  
299 viruses such as A(H5N1) and A(H7N9) influenza viruses that have caused substantial human  
300 disease and death (3, 27). As such, there is a need to evaluate heterogeneity of ferret transmission  
301 models employing viruses possessing a similar scope of transmissibility phenotypes. While the  
302 variability in transmission results for either the Cal/09 or ruddy turnstone/09 viruses tested in this  
303 study were not statistically significant, the range of results obtained, especially with the ruddy  
304 turnstone/09 virus, nonetheless illustrates a level of variability that can be present in transmission  
305 readouts of viruses exhibiting both low to high transmission efficiency (Table 1). This variability  
306 was present despite a high degree of standardization of virus stock, inoculation procedures, and  
307 uniformity of donor:contact ratio.

308 As shown in the Supplemental Methods and Supplemental Tables 1-6, this exercise  
309 captured the extensive heterogeneity in laboratory protocols and setups present between different  
310 groups. Documented variation was present in every parameter examined, inclusive of ferrets, cage  
311 setups, titration methods, and environmental conditions, among other features. Caging and airflow  
312 considerations were especially variable (Supplemental Table 2). It is impossible to standardize all  
313 contributing variables to these experiments, as institutional, animal welfare, and governmental  
314 guidelines and requirements vary worldwide, as do cost implications. That said, this exercise  
315 supports the capacity to harmonize results generated between disparate groups when a small

316 number of procedural parameters are fixed. Interestingly, the four groups that detected infectious  
317 virus in contact nasal wash specimens in ruddy turnstone/09 transmission experiments all found  
318 4/4 virus transmission in the Cal/09 experiment; transmission percentages between the two viruses  
319 were highly correlated between laboratories (Spearman correlation = 0.86,  $p < 0.001$ ).  
320 Furthermore, while directional airflow (OR=4) did not reach statistical significance, it is  
321 nonetheless of note that 3/4 laboratories for which ruddy turnstone/09 virus transmission was  
322 detected possessed directional airflow, versus 3/7 of the laboratories for which transmission with  
323 this virus was not detected; directional airflow from inoculated to contact animals was a feature in  
324 6/11 laboratories in this exercise (Supplemental Table 3). While our results did not conclusively  
325 identify any one experimental parameter statistically associated with enhanced transmissibility  
326 outcomes, it is possible that a confluence of parameters is nonetheless capable of creating a more  
327 permissive environment for virus transmission to occur.

328 To improve interpretation of results from this standardization exercise, we concurrently  
329 investigated the hypothetical confidence in concluding low transmissibility ( $\leq 25\%$  or  $\leq 1$  ferret  
330 infected out of 4 ferrets) or moderate to high transmissibility ( $\geq 50\%$  or  $\geq 2$  ferrets infected out of 4  
331 ferrets) from multiple contributing laboratories. These analyses assumed both a uniform prior  
332 distribution for the transmission probability for a novel pathogen, and independent transmission  
333 outcomes from the laboratories. We considered two scenarios: one scenario where strong  
334 homogeneity across laboratories could be assumed so the samples were pooled from multiple  
335 laboratories, and another scenario where each laboratory drew their own conclusion on  
336 transmissibility such that an overall conclusion was drawn as a voting system. As influenza viruses  
337 of notable public health importance are frequently assessed across multiple independent  
338 laboratories, these analyses provide a framework to rigorously interpret independently generated

339 findings, especially when discordant results between laboratories are reported. This is most critical  
340 in the event of a novel virus believed to possess moderate-to-high transmissibility; our analyses  
341 support that 4 independent laboratories with concordant results supporting an enhanced  
342 transmissibility phenotype yields a 95% probability of this finding, with additional independent  
343 groups or a greater number of total ferret donor:contact pairs necessary when discordant results  
344 are present.

345 Collectively, the findings of this exercise support the potential benefit of increased  
346 uniformity, or standardization, of some parameters when conducting risk assessment-specific  
347 activities on the same viruses. Specifically, the donor:contact ratio represents such a parameter.  
348 For a virus with moderate to high transmissibility, such as Cal/09 virus, modulation of this ratio  
349 (e.g., conducting experiments with a 2:1 donor:contact ratio, as is the case when transmission  
350 evaluations in a direct contact setting and via respiratory droplets employ a common donor) would  
351 not substantially alter conclusions drawn. However, for a virus with reduced transmissibility at a  
352 1:1 ratio, such as the ruddy turnstone/09 virus evaluated here, it is likely that an increased  
353 donor:contact ratio (eg., 2:1) may enhance transmissibility by increasing virus-laden aerosols  
354 exhaled from infected ferrets. Previous studies on ruddy turnstone/09 virus demonstrated airborne  
355 transmission potential when employing a donor: direct contact: aerosol contact at 1:1:1 ratio;  
356 efficient transmission by direct contact will subsequently affect the quantity and kinetics of virus-  
357 laden aerosols that mediate transmission by air (18, 19). There is a need to better understand how  
358 modulation of this ratio contributes to assessments of virus transmissibility. However, this does  
359 underscore the potential complications posed by harmonizing data generated for risk assessment  
360 purposes for which the donor:contact ratio diverges. With increased heterogeneity in results  
361 between labs, uncertainty around the conclusions increases, and there is a corresponding decrease

362 in confidence in the results (Table 3), showing the utility in increasing homogeneity across findings  
363 from different labs in order to reduce the total number of labs required to yield statistically  
364 meaningful results in this sort of analysis.

365 The emergence of SARS-CoV-2 further corroborates the pandemic potential of viruses of  
366 zoonotic origin. Early identification and risk assessments of novel viruses are essential for  
367 preventing the next pandemic. Continued optimization and refinement of risk assessment protocols  
368 will facilitate data interpretation in response to emerging pandemic threats. Collectively, a greater  
369 appreciation of this heterogeneity, and understanding of the scope of variability present in risk  
370 assessment settings, will permit more robust conclusions to be drawn from these efforts in the  
371 future.

372

### 373 **Materials and methods**

374 **Viruses.** The A(H1N1)pdm09 virus A/California/07/2009 (Cal/09) was propagated in MDCK  
375 cells (passage C3) at the US CDC as described previously (28). The low pathogenic avian influenza  
376 A(H1N1) virus A/ruddy turnstone/Delaware/300/2009 (ruddy turnstone/09) was propagated in  
377 eggs (passage E3) by St. Jude Children's Research Hospital as described previously (19). Stocks  
378 were fully sequenced and tested for exclusivity to rule out the presence of other influenza virus  
379 subtypes prior to distribution.

380 **Animal and experimental variability.** Groups obtained ferrets from multiple vendors and  
381 independent breeders from North America, Europe and Asia, and animals varied in their age,  
382 gender, health status, and other parameters (Supplemental Table 1). There was substantial  
383 differences between laboratories in the specific caging employed for transmission experiments,  
384 distance between cages, airflow directionality between cages, and air changes per hour

385 (Supplemental Table 2). Anesthesia protocols, sample collection methods, and decontamination  
386 procedures to prevent cross-contamination between contact and donor animals varied between  
387 groups and are reported in Supplemental Methods. All experiments were performed under country-  
388 specific legal guidelines and approved institutional-specific animal protocols as specified in the  
389 Supplemental Methods.

390 **Standardized procedures.** All laboratories received common stock viruses prepared by CDC and  
391 St. Jude Children's Research Hospital with the shipping temperature recorded. Stock viruses were  
392 diluted to  $10^6$  plaque forming units (PFU) in 500 $\mu$ l PBS based on predetermined viral titers, and  
393 donor ferrets were inoculated intranasally under in-house protocols for anesthesia (Supplemental  
394 Methods). On day 1 post-inoculation, one respiratory droplet contact ferret was introduced and  
395 exposed to each donor by housing in an adjacent cage, employing a strict 1:1 donor:contact ratio,  
396 with 4 transmission pairs tested for each virus. Ferret temperatures, weights, and nasal  
397 washes/swabs were collected every 24-48 hours. Daily room temperature and relative humidity  
398 readings were collected and are reported in Supplemental Table 3 employing pre-validated  
399 thermohygrometers with comparable readings (Testo Inc., 608-H1). Sera was collected at the end  
400 of each experiment for determination of seroconversion to homologous virus by hemagglutinin  
401 inhibition assay using established in-house serology protocols.

402 **Sample titration and normalization.** Infectious virus titers were determined by plaque assay,  
403 50% tissue culture infectious dose (TCID<sub>50</sub>) assay, or 50% egg infectious doses (EID<sub>50</sub>) assay at  
404 each laboratory with varying limits of detection (Supplemental Table 4). To facilitate subsequent  
405 statistical assessments across laboratories, reported titers from each laboratory were normalized to  
406 TCID<sub>50</sub>/mL for each virus based on PFU, TCID<sub>50</sub>, and EID<sub>50</sub> values pre-determined by a single  
407 laboratory to minimize titration methodology-specific variation.

408 **Data blinding and analyses.** Data blinding, aggregation and all statistical analyses were  
409 performed by an independent statistician. Transmission outcomes were compared across  
410 laboratories by each virus, using Fisher's exact test of homogeneity. Viral load between viruses  
411 were compared by testing difference in area under the curve (AUC) using t-test. Factors associated  
412 with transmissibility and morbidity were assessed by using logistic regression and linear regression  
413 models. We also investigated the confidence in concluding low transmissibility ( $\leq 25\%$ , or  $\leq 1$  ferret  
414 infected out of 4 ferrets) or moderate to high transmissibility ( $\geq 50\%$  or  $\geq 2$  ferrets infected out of 4  
415 ferrets) from multiple contributing laboratories. We assumed a uniform prior distribution for the  
416 transmission probability for a novel pathogen was assumed, and independent transmission  
417 outcomes from the laboratories. The confidence of drawing conclusion on transmissibility with  
418 concordant or discordant outcomes from the laboratories is presented. We considered a scenario  
419 where strong homogeneity across laboratory can be assumed so the samples were pooled from  
420 multiple laboratories, and another scenario that each laboratory draw their own conclusion on  
421 transmissibility and the overall conclusion was drawn as voting system. All analyses were  
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423

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428 Registry.

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## 432 References

- 433 1. Buhnerkempe MG, Gostic K, Park M, Ahsan P, Belser JA, Lloyd-Smith JO. 2015.  
434 Mapping influenza transmission in the ferret model to transmission in humans. *Elife* 4.
- 435 2. Cox NJ, Trock SC, Burke SA. 2014. Pandemic preparedness and the Influenza Risk  
436 Assessment Tool (IRAT). *Curr Top Microbiol Immunol* 385:119-36.
- 437 3. WHO. 2016. Tool for Influenza Pandemic Risk Assessment (TIPRA).
- 438 4. Belser JA, Barclay W, Barr I, Fouchier RAM, Matsuyama R, Nishiura H, Peiris M,  
439 Russell CJ, Subbarao K, Zhu H, Yen HL. 2018. Ferrets as Models for Influenza Virus  
440 Transmission Studies and Pandemic Risk Assessments. *Emerg Infect Dis* 24:965-971.
- 441 5. Moore IN, Lamirande EW, Paskel M, Donahue D, Kenney H, Qin J, Subbarao K. 2014.  
442 Severity of clinical disease and pathology in ferrets experimentally infected with  
443 influenza viruses is influenced by inoculum volume. *J Virol* 88:13879-91.
- 444 6. Gustin KM, Belser JA, Wadford DA, Pearce MB, Katz JM, Tumpey TM, Maines TR.  
445 2011. Influenza virus aerosol exposure and analytical system for ferrets. *Proc Natl Acad  
446 Sci U S A* 108:8432-7.
- 447 7. Bodewes R, Kreijtz JH, van Amerongen G, Fouchier RA, Osterhaus AD, Rimmelzwaan  
448 GF, Kuiken T. 2011. Pathogenesis of Influenza A/H5N1 virus infection in ferrets differs  
449 between intranasal and intratracheal routes of inoculation. *Am J Pathol* 179:30-6.
- 450 8. Belser JA, Eckert AM, Tumpey TM, Maines TR. 2016. Complexities in Ferret Influenza  
451 Virus Pathogenesis and Transmission Models. *Microbiol Mol Biol Rev* 80:733-44.
- 452 9. Belser JA, Maines TR, Katz JM, Tumpey TM. 2013. Considerations regarding  
453 appropriate sample size for conducting ferret transmission experiments. *Future Microbiol*  
454 8:961-5.
- 455 10. Nishiura H, Yen HL, Cowling BJ. 2013. Sample size considerations for one-to-one  
456 animal transmission studies of the influenza A viruses. *PLoS One* 8:e55358.
- 457 11. Linster M, van Boheemen S, de Graaf M, Schrauwen EJA, Lexmond P, Manz B,  
458 Bestebroer TM, Baumann J, van Riel D, Rimmelzwaan GF, Osterhaus A, Matrosovich  
459 M, Fouchier RAM, Herfst S. 2014. Identification, characterization, and natural selection  
460 of mutations driving airborne transmission of A/H5N1 virus. *Cell* 157:329-339.
- 461 12. Martinez-Sobrido L, Blanco-Lobo P, Rodriguez L, Fitzgerald T, Zhang H, Nguyen P,  
462 Anderson CS, Holden-Wiltse J, Bandyopadhyay S, Nogales A, DeDiego ML, Wasik BR,  
463 Miller BL, Henry C, Wilson PC, Sangster MY, Treanor JJ, Topham DJ, Byrd-Leotis L,  
464 Steinhauer DA, Cummings RD, Luczo JM, Tompkins SM, Sakamoto K, Jones CA, Steel  
465 J, Lowen AC, Danzy S, Tao H, Fink AL, Klein SL, Wohlgemuth N, Fenstermacher KJ,  
466 El Najjar F, Pekosz A, Sauer L, Lewis MK, Shaw-Saliba K, Rothman RE, Liu ZY, Chen  
467 KF, Parrish CR, Voorhees IEH, Kawaoka Y, Neumann G, Chiba S, Fan S, Hatta M,  
468 Kong H, Zhong G, et al. 2020. Characterizing Emerging Canine H3 Influenza Viruses.  
469 *PLoS Pathog* 16:e1008409.

470 13. Lee LYY, Zhou J, Frise R, Goldhill DH, Koszalka P, Mifsud EJ, Baba K, Noda T, Ando  
471 Y, Sato K, Yuki AI, Shishido T, Uehara T, Wildum S, Zwanziger E, Collinson N,  
472 Kuhlbusch K, Clinch B, Hurt AC, Barclay WS. 2020. Baloxavir treatment of ferrets  
473 infected with influenza A(H1N1)pdm09 virus reduces onward transmission. PLoS Pathog  
474 16:e1008395.

475 14. Pulit-Penaloza JA, Jones J, Sun X, Jang Y, Thor S, Belser JA, Zanders N, Creager HM,  
476 Ridenour C, Wang L, Stark TJ, Garten R, Chen LM, Barnes J, Tumpey TM, Wentworth  
477 DE, Maines TR, Davis CT. 2018. Antigenically Diverse Swine Origin H1N1 Variant  
478 Influenza Viruses Exhibit Differential Ferret Pathogenesis and Transmission Phenotypes.  
479 J Virol 92.

480 15. Baz M, Boonnak K, Paskel M, Santos C, Powell T, Townsend A, Subbarao K. 2015.  
481 Nonreplicating influenza A virus vaccines confer broad protection against lethal  
482 challenge. MBio 6:e01487-15.

483 16. Lakdawala SS, Lamirande EW, Sugitan AL, Jr., Wang W, Santos CP, Vogel L,  
484 Matsuoka Y, Lindsley WG, Jin H, Subbarao K. 2011. Eurasian-origin gene segments  
485 contribute to the transmissibility, aerosol release, and morphology of the 2009 pandemic  
486 H1N1 influenza virus. PLoS Pathog 7:e1002443.

487 17. Munster VJ, de Wit E, van den Brand JM, Herfst S, Schrauwen EJ, Bestebroer TM, van  
488 de Vijver D, Boucher CA, Koopmans M, Rimmelzwaan GF, Kuiken T, Osterhaus AD,  
489 Fouchier RA. 2009. Pathogenesis and transmission of swine-origin 2009 A(H1N1)  
490 influenza virus in ferrets. Science 325:481-3.

491 18. Kocer ZA, Krauss S, Stallknecht DE, Rehg JE, Webster RG. 2012. The potential of avian  
492 H1N1 influenza A viruses to replicate and cause disease in mammalian models. PLoS  
493 One 7:e41609.

494 19. Zanin M, Wong SS, Barman S, Kaewborisuth C, Vogel P, Rubrum A, Darnell D,  
495 Marinova-Petkova A, Krauss S, Webby RJ, Webster RG. 2017. Molecular basis of  
496 mammalian transmissibility of avian H1N1 influenza viruses and their pandemic  
497 potential. Proc Natl Acad Sci U S A 114:11217-11222.

498 20. Marr LC, Tang JW, Van Mullekom J, Lakdawala SS. 2019. Mechanistic insights into the  
499 effect of humidity on airborne influenza virus survival, transmission and incidence. J R  
500 Soc Interface 16:20180298.

501 21. Gustin KM, Belser JA, Veguilla V, Zeng H, Katz JM, Tumpey TM, Maines TR. 2015.  
502 Environmental Conditions Affect Exhalation of H3N2 Seasonal and Variant Influenza  
503 Viruses and Respiratory Droplet Transmission in Ferrets. PLoS One 10:e0125874.

504 22. Albrecht RA, Liu WC, Sant AJ, Tompkins SM, Pekosz A, Meliopoulos V, Cherry S,  
505 Thomas PG, Schultz-Cherry S. 2018. Moving Forward: Recent Developments for the  
506 Ferret Biomedical Research Model. mBio 9.

507 23. Karlsson EA, Meliopoulos VA, Savage C, Livingston B, Mehle A, Schultz-Cherry S.  
508 2015. Visualizing real-time influenza virus infection, transmission and protection in  
509 ferrets. Nat Commun 6:6378.

510 24. Yen HL, Liang CH, Wu CY, Forrest HL, Ferguson A, Choy KT, Jones J, Wong DD,  
511 Cheung PP, Hsu CH, Li OT, Yuen KM, Chan RW, Poon LL, Chan MC, Nicholls JM,  
512 Krauss S, Wong CH, Guan Y, Webster RG, Webby RJ, Peiris M. 2011. Hemagglutinin-  
513 neuraminidase balance confers respiratory-droplet transmissibility of the pandemic H1N1  
514 influenza virus in ferrets. *Proc Natl Acad Sci U S A* 108:14264-9.

515 25. Varble A, Albrecht RA, Backes S, Crumiller M, Bouvier NM, Sachs D, Garcia-Sastre A,  
516 tenOever BR. 2014. Influenza A virus transmission bottlenecks are defined by infection  
517 route and recipient host. *Cell Host Microbe* 16:691-700.

518 26. Stephenson I, Heath A, Major D, Newman RW, Hoschler K, Junzi W, Katz JM, Weir JP,  
519 Zambon MC, Wood JM. 2009. Reproducibility of serologic assays for influenza virus A  
520 (H5N1). *Emerg Infect Dis* 15:1252-9.

521 27. CDC. 2021. Summary of Influenza Risk Assessment Tool (IRAT) Results.  
<https://www.cdc.gov/flu/pandemic-resources/monitoring/irat-virus-summaries.htm>.  
522 Accessed Feb 10 2021.

523 28. Maines TR, Jayaraman A, Belser JA, Wadford DA, Pappas C, Zeng H, Gustin KM,  
524 Pearce MB, Viswanathan K, Shriver ZH, Raman R, Cox NJ, Sasisekharan R, Katz JM,  
525 Tumpey TM. 2009. Transmission and pathogenesis of swine-origin 2009 A(H1N1)  
526 influenza viruses in ferrets and mice. *Science* 325:484-7.

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