

Impact of Prior Infection on Protection and Transmission of SARS-CoV-2 in Golden Hamsters

4 Cheng Zhang^{1, 2&}, Zhendong Guo^{1&}, Nan Li^{1&}, Huan Cui^{1, 3}, Keyin Meng¹, Lina Liu¹, Li Zhao¹,
5 Shanshan Zhang¹, Chengfeng Qin⁴, Juxiang Liu², Yuwei Gao^{1#}, Chunmao Zhang^{1#}

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⁷ ¹ Military Veterinary Research Institute, Changchun, China

⁸ ²College of Veterinary Medicine, Hebei Agricultural University, Baoding, China

9 ³College of Veterinary Medicine, Jilin University, Changchun, China

⁴ Beijing Institute of Microbiology and Epidemiology, Beijing, China

11 & Authors contributed equally to this work.

12 [#] Correspondence author:

13 Chunmao Zhang: jk704715@sina.com

14 Yuwei Gao: gaoyuwei@gmail.com

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16 **Abstract**

17 The severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) has caused over 100 million
18 confirmed human infections, and 2 million more deaths globally since its emergence in the end of
19 2019. Several studies have shown that prior infection provided protective immunity against
20 SARS-CoV-2 in non-human primate models. However, the effect of prior infection on blocking
21 SARS-CoV-2 transmission is not clear. Here, we evaluated the impact of prior infection on
22 protection and transmission of the SARS-CoV-2 virus in golden hamsters. Our results showed that
23 prior infection provided protective immunity against SARS-CoV-2 re-challenge, but it was not
24 sterilizing immunity. The transmission experiment results showed that SARS-CoV-2 was efficiently
25 transmitted from naive hamsters to prior infected hamsters by direct contact and airborne route,
26 but not by indirect contact. Further, the virus was efficiently transmitted from prior infected
27 hamsters to naive hamsters by direct contact, but not by airborne route and indirect contact.
28 Surprisingly, the virus can be transmitted between prior infected hamsters by direct contact during
29 a short period of early infection. Taken together, our study demonstrated that prior infected
30 hamsters with good immunity can still be naturally re-infected, and the virus can be transmitted
31 between prior infected hamsters and the naive through different transmission routes, implying the
32 potential possibility of human re-infection and the risk of virus transmission between prior
33 infected population and the healthy. Our study will help to calculate the herd immunity threshold
34 more accurately, make more reasonable public health decisions, formulate an optimized
35 population vaccination program, as well as aid the implementation of appropriate public health
36 and social measures to control COVID-19.

37 Key words: SARS-CoV-2, prior infection, re-infection, direct contact, airborne transmission

38 **The main text**

39 As of Jan 29, 2021, more than 100 million confirmed human infections and over 2 million deaths
40 have been caused by the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), the
41 causative agent of coronavirus infectious disease 2019 (COVID-19) pandemic, with devastating
42 impact on lives and economy. At present, multiple vaccine candidates are in phase 3 clinical trials
43 and several of them have been approved for emergency use authorization with conditions¹⁻⁸.
44 Previous studies have shown that prior infection or vaccination provided protective immunity
45 against SARS-CoV-2 in different animal models⁹⁻¹⁷, but most are not sterilizing immunity. The prior
46 infected or vaccinated animals still shed large quantities of virus in their upper respiratory
47 tracts^{9,12,14,16}. Besides protection from diseases, reducing or blocking SARS-CoV-2 transmission
48 between humans is crucial for COVID-19 pandemic control. However, the impact of prior
49 infection on blocking SARS-CoV-2 transmission is not clear. As a small animal model, golden
50 hamsters have been used for studying pathogenesis^{18,19}, transmission ability²⁰ of SARS-CoV-2 and
51 for evaluating potential vaccines^{21,22} and antiviral drugs²³. Here, we evaluated the impact of prior
52 infection on protection and transmission of SARS-CoV-2 in golden hamsters. Our results showed
53 that prior infection provided good protective immunity against SARS-CoV-2 re-challenge, but the
54 prior infected hamsters can still be re-infected. Moreover, the virus was efficiently transmitted
55 from naive hamsters to prior infected hamsters by direct contact and airborne route, but not by
56 indirect contact. The virus was also transmitted from prior infected hamsters to the naive and prior
57 infected hamsters by direct contact, but not by airborne route. Our findings will help governments
58 and public health agencies to make more reasonable public health decisions as well as aid the
59 implementation of appropriate public health and social measures to control COVID-19.

60 **Prior infection protects hamsters against SARS-CoV-2 re-challenge**

61 We evaluated protective immunity of prior infection against SARS-CoV-2 re-challenge. Hamsters
62 were divided into high dose infected group (HD) and low dose infected group (LD), and
63 intranasally inoculated with 10^5 TCID₅₀ or 10^3 TCID₅₀ of the virus respectively (Supplementary
64 Table S1). At 21 days post infection (dpi), Hamsters in HD and LD were inoculated with 10^6
65 TCID₅₀ of the virus. As the infected control (IC), another six naive hamsters were inoculated with
66 10^6 TCID₅₀ of the virus. At 2 and 4 dpi, nasal washes and the supernatants of the homogenized
67 nasal turbinates and lungs were collected for virus titration in Vero-E6 cells and RNA
68 quantification using real-time qPCR. The glutaraldehyde-fixed nasal turbinates were prepared for
69 histological examination.

70 Serum was collected from hamsters in HD and LD at 21 dpi. The results of virus neutralization
71 assay revealed that all hamsters inoculated with the virus had a much higher neutralizing antibody
72 titer, and the neutralizing antibody titer in HD was slightly higher than that in LD (Figure S1),
73 indicating that prior infection elicited effective immunity in hamsters. For nasal washes, viral load
74 in IC peaked at 2 dpi, with a titer of $10^{5.17}$ TCID₅₀/mL, and was significantly much higher than that
75 in HD and LD (Figure 1A). At 2 and 4 dpi, RNA copies in IC were slightly higher than that in HD
76 and LD with significant difference (Figure 1D). For nasal turbinate, high levels of viral load were
77 observed in IC at 2 and 4 dpi, with a titer of $10^{5.75}$ TCID₅₀/mL and $10^{3.75}$ TCID₅₀/mL respectively,
78 about 3500-fold and 180-fold higher than that in HD and LD (Figure 1B). Viral load in HD and
79 LD decreased to below the detection limit of TCID₅₀ assay at 4 dpi. RNA copies in IC at 2 dpi,
80 $10^{10.45}$ copies/mL, were 80-fold and 25-fold higher than that in HD and LD respectively and RNA
81 copies in IC at 4 dpi, $10^{9.83}$ copies/mL, were about 4000-fold and 10000-fold higher than that in

82 HD and LD (Figure 1E). Compared with at 2 dpi, RNA copies at 4 dpi in HD and LD decreased
83 about 200-fold and 1600-fold respectively (Figure 1E). Viral RNA assays were further confirmed
84 by the sgmRNA assays. The sgmRNA copies in IC at 2 and 4 dpi, $10^{7.1}$ copies/mL and
85 $10^{6.4}$ copies/mL, were about 80-fold and 5000-fold higher than that in HD and LD, averagely
86 $10^{5.2}$ copies/mL and $10^{2.7}$ copies/mL respectively (Figure 1G). Compared with at 2 dpi, the
87 sgmRNA copies in HD and LD at 4 dpi decreased about 250-fold and 600-fold respectively
88 (Figure 1G). For lungs, viral load at 2 and 4 dpi in HD and LD was under the detection limit,
89 which was significantly lower than that in IC, $10^{5.33}$ TCID₅₀/mL and $10^{4.17}$ TCID₅₀/mL respectively
90 (Figure 1C). RNA copies at 2 and 4 dpi in IC, $10^{10.2}$ copies/mL and $10^{9.5}$ copies/mL, was about
91 13000-fold and 40000-fold higher than that in HD and LD, about $10^{6.1}$ copies/mL and
92 $10^{4.9}$ copies/mL (Figure 1F). Similar to the trend of viral RNA copies, the sgmRNA in IG at 2 and
93 4 dpi, $10^{6.6}$ copies/mL and $10^{6.0}$ copies/mL, was about 16000-fold and 8000-fold higher than that in
94 HD and LD, averagely $10^{2.4}$ copies/mL and $10^{2.1}$ copies/mL respectively (Figure 1H). Additionally,
95 we examined the presence of SARS-CoV-2 virus in nasal tissues using transmission electron
96 microscopy. Several coronavirus-like particles were observed in intracellular compartments of
97 nasal tissues of hamsters that were re-challenged with SARS-CoV-2 (Figure S2).
98 The substantially reduced viral titers, RNA and sgmRNA copies in nasal washes, nasal turbinates
99 and lungs showed that prior infection provided good protective immunity against SARS-CoV-2.
100 However, a moderate level of live virus was still detected in nasal washes and nasal turbinates,
101 despite with a relatively short shedding period, and considerable sgmRNA copies were also
102 detected in nasal turbinates at 2 dpi. The results of the detected live virus and the considerable
103 sgmRNA copies, in combined with observation of coronavirus-like particles in intracellular

104 compartments of nasal cells, powerfully proved that the virus can replicate in prior infected
105 hamsters, especially in nasal turbinates, undoubtedly indicating that prior infected hamsters can be
106 re-infected by the virus, even with a higher neutralizing antibody titer.

107 **Impact of prior infection on SARS-CoV-2 transmission in hamsters**

108 The SARS-CoV-2 virus was transmitted between hamsters via multiple routes, including direct
109 contact, indirect contact and airborne transmission. Here we systematically evaluated the impact
110 of prior infection on SARS-CoV-2 transmission between prior infected hamsters and the naive
111 hamsters.

112 **Transmission of SARS-CoV-2 from naive hamsters to prior infected hamsters**

113 For the potential transmission of the virus from naive hamsters to prior infected hamsters by direct
114 contact, three naive donor hamsters were intranasally inoculated with 10^6 TCID₅₀ of the virus.
115 After 24 hours' inoculation, the three donors were transferred to a direct contact transmission cage
116 (supplementary Figure S3A) and co-housed with another three prior infected hamsters that were
117 inoculated with 10^5 TCID₅₀ or 10^3 TCID₅₀ of the virus 21 days ago. Nasal washes were collected
118 every other day from the donors and the contacts for 8 days. For donors, the infectious viral load
119 in nasal washes peaked at 2 dpi and then declined rapidly, while viral RNA copies was relatively
120 stable during the first six infection days, and then substantially declined at 8 dpi (Figure 2A). At 1
121 days post exposure (dpe), live SARS-CoV-2 virus was detected in nasal washes of two prior
122 infected contact hamsters, and one with a very low viral titer. At 3 dpe, live virus was detected in
123 all three prior infected contact hamsters (Figure 2A). The viral titers in the contacts were much
124 lower than that in the donor hamsters. Viral RNA copies in two prior infected contact hamsters
125 were significantly improved at 3 dpe (Figure 2A). The experiment results showed that the virus

126 was efficiently transmitted from the naïve donors to the prior infected contacts. For airborne
127 transmission of the virus from naive hamsters to prior infected hamsters, three naive donor
128 hamsters were inoculated with 10^6 TCID₅₀ of the virus, and at 24 hours' inoculation, the donor
129 hamsters and another three prior infected recipient hamsters were transferred to the airborne
130 transmission cage, with two wire-mesh partition that prevented the direct and indirect contact
131 between animals and allowed the spread of the virus through air (supplementary Figure S3B). At 1
132 dpe, live SARS-CoV-2 virus was detected in nasal washes of two prior infected recipient hamsters
133 and one with a very low viral titer (Figure 2B). At 3 dpe, all three prior infected recipient hamsters
134 were infected by SARS-CoV-2 and viral titers in nasal washes were significantly improved, and at
135 5 dpe, the viral titer in one prior infected recipient hamster was still relatively high, about $10^{3.25}$
136 TCID₅₀/mL (Figure 2B). Viral RNA copies in nasal washes in the prior infected recipient hamsters
137 peaked at 3 dpe, and one hamster still had a higher viral RNA copies in nasal washes at 5 dpe
138 (Figure 2B). Therefore, the virus was efficiently transmitted from the naive to prior infected
139 hamsters by airborne route as well. For indirect contact transmission of the virus from the naive to
140 prior infected hamsters, three naive donor hamsters were inoculated with 10^6 TCID₅₀ of the virus,
141 and at 48 hours' inoculation, the donor hamsters were removed and transferred to a new cage, and
142 another three prior infected recipient hamsters were placed into the initial cage housing the donor
143 hamsters. During the whole experiment period, no live virus was detected in nasal washes of the
144 three prior infected recipient hamsters (Figure 2C). The virus was not transmitted from the naive
145 to prior infected hamsters by indirect contact. In summary, SARS-CoV-2 was efficiently
146 transmitted from the naive donors to prior infected hamsters by direct contact and airborne
147 transmission, but not by indirect contact.

148 **Transmission of SARS-CoV-2 from prior infected hamsters to naive hamsters**

149 For direct contact transmission of the virus from prior infected hamsters to naive hamsters, three

150 prior infected hamsters were as the donors, and another three naive hamsters were as the recipients

151 in direct contact transmission group. The donors were inoculated with 10^6 TCID₅₀ of the virus, and

152 at 24 hours' inoculation, the donors and the direct contacts were co-housed together in a new cage.

153 At 2 dpi, viral titers in nasal washes of two donor hamsters were very low, and another donor

154 hamster with a moderate titer of $10^{2.75}$ TCID₅₀/mL (Figure 3A). At 4 dpi, live virus was not

155 detected in nasal washes of two donor hamsters, and another one with a low viral titer. Viral RNA

156 copies in nasal washes of the donor hamsters were about 10^8 copies/mL at 2 dpi, and substantially

157 declined at 6 and 8 dpi (Figure 3A). Live virus was detected in nasal washes of all three contact

158 hamsters at 3 days post exposure (dpe), one with a very high titer of $10^{5.5}$ TCID₅₀/mL, and another

159 with a very low titer of $10^{0.75}$ TCID₅₀/mL (Figure 3A). Viral titers in nasal washes of all hamsters

160 were very high at 5 dpe. Viral RNA copies in nasal washes of the contact hamsters were improved

161 rapidly at 3 dpe and later held at a high level. The results showed that the virus was efficiently

162 transmitted from prior infected hamsters to naive hamsters by direct contact. For airborne

163 transmission of the virus from prior infected hamsters to naive hamsters, three prior infected donor

164 hamsters were inoculated with 10^6 TCID₅₀ of the virus, and another three naive hamsters were as

165 recipients. At 24 hours' inoculation, the prior infected donor hamsters and the naïve recipient

166 hamsters were transferred to an airborne transmission cage. Live virus was not detected at 2 dpi

167 and later in nasal washes of the prior infected donor hamsters (Figure 3B). No live virus was also

168 detected in nasal washes of the recipient hamsters in the airborne transmission group (Figure 3B).

169 The airborne transmission experiment was also performed similarly at two hours' inoculation. A

170 moderate level of virus titer was detected at 1 and 3 dpi in prior infected donor hamsters, and no
171 live virus was still detected in all recipient hamsters in the airborne transmission group during the
172 whole experiment period (Figure S4B). The results demonstrated that the virus was not
173 transmitted from prior infected donors to naive hamsters through airborne route. For indirect
174 contact transmission of the virus from prior infected hamsters to naive hamsters, three prior
175 infected donor hamsters were inoculated with 10^6 TCID₅₀ of the virus, and at 48 hours' inoculation,
176 the prior infected donor hamsters were removed and transferred to a new cage, and another three
177 naive recipient hamsters were placed into the initial cage housing the donors. No live virus was
178 detected in nasal washes of the recipient hamsters in the indirect contact transmission group
179 (Figure 3C). The virus was not transmitted from prior infected hamsters to the naive by indirect
180 contact. To sum up, SARS-CoV-2 can be transmitted from prior infected hamsters to naive
181 hamsters by direct contact, but not by airborne route and indirect contact.

182 **Transmission of SARS-CoV-2 between prior infected hamsters**

183 We evaluated the potential transmission of SARS-CoV-2 between prior infected hamsters by direct
184 contact and airborne route. For direct contact transmission, four prior infected donor hamsters
185 were inoculated with 10^6 TCID₅₀ of the virus. At two hours' inoculation, another four prior
186 infected hamsters were co-housed together with those four donors in a new cage. A high level of
187 virus titer was detected in nasal washes of the donor hamsters at 1 and 3 dpi, averagely $10^{3.56}$
188 TCID₅₀/mL and $10^{2.25}$ TCID₅₀/mL, while viral RNA copies were maintained at 10^7 to
189 10^8 copies/mL. Since 5 dpi, no live virus was found in nasal washes of the donors. At 3 dpe, live
190 virus was detected in nasal washes in one contact hamster, with a titer of $10^{4.25}$ TCID₅₀/mL, and
191 viral RNA load in this hamster was also greatly improved to $10^{8.26}$ copies/mL (Figure 4A). At 5 dpe,

192 live virus was detected in nasal washes of another hamster in the contact transmission group, with
193 a low titer of $10^{0.75}$ TCID₅₀/mL, while viral RNA copies were improved to 10^6 copies/mL. At 7 dpe,
194 virus titer in nasal washes of this hamster was substantially improved to $10^{3.25}$ TCID₅₀/mL, and
195 viral RNA copies were further improved to $10^{8.46}$ copies/mL (Figure 4A). It seems that the virus
196 was first transmitted from the artificially inoculated hamsters to a prior infected contact hamster,
197 and then was sequentially transmitted to another prior infected contact hamster. The experiment
198 was also performed similarly at 24 hours' inoculation, four prior infected donors and four prior
199 infected contact hamsters were co-housed in a new cage at 24 hours' inoculation. No live virus
200 was detected in nasal washes of all contact hamsters during the experiment period (Figure S5A).
201 The experiment results showed that SARS-CoV-2 was transmitted between prior infected hamsters
202 by direct contact during a very short period of early infection. For airborne transmission, three
203 prior infected donor hamsters were inoculated with 10^6 TCID₅₀ of the virus and another three prior
204 infected hamsters were as the recipients in airborne transmission group. The three donor hamsters
205 and the three recipient hamsters were transferred to an airborne transmission cage at two hours' or
206 24 hours' inoculation, no live virus was detected in nasal washes of all recipient hamsters in the
207 two airborne transmission groups (Figure 4B & Figure S5B). The results showed that
208 SARS-CoV-2 was not transmitted between prior infected hamsters by airborne route. Taken
209 together, SARS-CoV-2 has limited transmission ability between prior infected hamsters by direct
210 contact during a short period of early infection, but without the ability to transmit by airborne
211 route.

212 **Impact of a lower dose inoculation on SARS-CoV-2 transmission**

213 We evaluated the impact of a lower dose inoculation on SARS-CoV-2 transmission between prior

214 infected hamsters and naïve hamsters by direct contact. For transmission of the virus from naïve
215 donors to prior infected contacts, four naïve donor hamsters were inoculated with 10^4 TCID₅₀ of
216 the virus, and at 24 hours' inoculation, the donors and another four prior infected hamsters were
217 co-housed together in a new cage. Similar to the high dose inoculation, viral load in nasal washes
218 was maintained at a higher level at 2 and 4 dpi as well, and then followed a rapid decline (Figure
219 5A). Viral RNA copies were maintained at about 10^9 copies/mL during the first four infection days,
220 and then slowly declined. Live virus was detected in two contact hamsters at 1 dpe, one with a
221 very low titer of $10^{0.75}$ TCID₅₀/mL. At 3 dpe, live virus was found in nasal washes of all four prior
222 infected hamsters, two of which was significantly improved than before (Figure 5A). Viral RNA
223 copies in nasal washes at 3 dpe were also significantly improved than before. Obviously,
224 SARS-CoV-2 was efficiently transmitted from naïve donor hamsters to prior infected hamsters by
225 direct contact. For transmission of the virus from prior infected hamsters to naïve hamsters, four
226 prior infected donor hamsters were inoculated with 10^4 TCID₅₀ of the virus. At 24 hours'
227 inoculation, the donor hamsters and another four naïve hamsters were co-housed together in a new
228 cage. At 2 dpi, virus load in nasal washes of the donor hamsters was at a moderate level, about
229 $10^{2.25}$ TCID₅₀/mL, and viral RNA copies were about $10^{7.8}$ copies/mL (Figure 5B). Live virus was
230 detected in nasal washes of three of the four naïve contacts at 3 dpe, with titers from $10^{3.25}$
231 TCID₅₀/mL to $10^{5.5}$ TCID₅₀/mL (Figure 5B). At 5 dpe, another contact hamster was also infected by
232 SARS-CoV-2, with a high viral titer of $10^{5.25}$ TCID₅₀/mL in nasal washes. Viral RNA copies were
233 rapidly improved to $10^{8.72}$ copies/mL at 3 dpe, and further improved to $10^{10.15}$ copies/mL at 5 dpe,
234 then slowly declined at 7 dpe. The results showed that SARS-CoV-2 was efficiently transmitted
235 from prior infected donor hamsters to the naïve contacts. In summary, with a lower inoculation,

236 SARS-CoV-2 was still efficiently transmitted between prior infected hamsters and naïve hamsters.

237 **Discussion**

238 Our results showed that prior infection of SARS-CoV-2 elicited a higher titer of neutralizing

239 antibodies in hamsters, and provided protective immunity against SARS-CoV-2 re-challenge.

240 However, it was not sterilizing immunity, and the virus still moderately replicated in nasal

241 turbinates of prior infected hamsters, indicating that prior infected hamsters can be artificially

242 re-infected after a short recovery period, even with a high level of neutralization antibodies. The

243 conclusion is consistent with recent reports showing that recovered COVID-19 patients were

244 re-infected in the presence of neutralizing antibodies^{24,25}. A large study of a recovered cohort of

245 175 COVID-19 patients revealed that 6% of COVID-19 patients did not show any antibody

246 response at all, and about 30% COVID-19 patients showed very low neutralizing antibodies²⁶.

247 Considering the gradual decay of neutralizing antibodies²⁷⁻³⁰ and a considerable population with

248 very low neutralizing antibodies, the re-infection of some recovered COVID-19 patients will be

249 unavoidable in the future. We also showed that prior infected hamsters can be naturally re-infected

250 by direct contact or airborne route. The results of transmission study showed that SARS-CoV-2

251 can be transmitted effectively from naïve hamsters to prior infected hamsters by direct contact and

252 airborne routes, but not by indirect contact. Additionally, SARS-CoV-2 can be transmitted

253 effectively from prior infected hamsters to naïve hamsters by direct contact, but not by airborne

254 route and indirect contact. Furthermore, SARS-CoV-2 can be transmitted between prior infected

255 hamsters by direct contact during a very short period of early infection, but the transmission

256 efficiency was limited. Taken together, prior infection substantially reduced the transmission

257 efficiency of SARS-CoV-2 from prior infected hamsters to the naïve or prior infected hamsters by

258 airborne route, but had limited impact on lowering the transmission by direct contact. In contrast
259 with SARS-CoV-2, seasonal influenza A virus transmission between ferrets can be substantially
260 reduced or blocked by natural infection or vaccination with live attenuated viruses^{31,32}. The
261 underlying mechanism behind the difference is not clear. Given the facts of re-infection, effective
262 transmission between the prior infected hamsters and the naive, and waning immunity of the
263 recovered COVID-19 patients^{28,29}, it would be much more difficult to achieve herd immunity by
264 natural infection or vaccination. A much higher vaccination coverage rate may be needed. At
265 present, many governments and public health agencies are considering introducing immunity
266 passport to help with recovery of social community and economy activities³³, but evidence
267 supporting this proposal is not enough. Reducing or blocking SARS-CoV-2 transmission is critical
268 for COVID-19 pandemic control. A roaring increase in confirmed infections and hospitalized
269 patients may lead to the collapse of the health care system, resulting in more deaths, social panic
270 and even economic paralysis. How does vaccination with COVID-19 vaccines impact
271 SARS-CoV-2 transmission in humans? There is still no clear-cut answer. Recent studies
272 demonstrated that intranasal immunization with an Ad vector vaccine provided near complete
273 sterilizing immunity to SARS-CoV-2 in mice³⁴, which might block the virus transmission in
274 humans. Further studies to evaluate the blocking efficiency of different vaccines vaccinated by
275 different routes, in particular, intranasal vaccination, on SARS-CoV-2 transmission in humans and
276 animal models are urgently needed. Our work will help to determine the herd immunity threshold
277 more accurately, make more reasonable public health decisions, as well as aid the implementation
278 of appropriate public health and social measures to control COVID-19.

279

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285 **Author Contributions**

286 CMZ, YWG conceived and designed the project, CMZ, CZ, ZDG, NL, HC, LNL, LZ, KYM and
287 SSZ performed the experiments. CMZ, CZ, ZDG and NL analyzed the data. CMZ drafted the
288 manuscript, YWG, CFQ, ZDG and JXL revised the manuscript critically.

289 **Declaration of interests**

290 All authors declared no competing interests.

291 **Data sharing**

292 Data will be made available on request, directed to corresponding author CMZ.

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389

390 **Figure legends**

391 **Figure 1** Viral load and histological examination in prior infected hamsters intranasally inoculated
392 with SARS-CoV-2. Sixteen hamsters were randomly divided into HD and LD groups and
393 inoculated with 10^5 TCID₅₀ or 10^3 TCID₅₀ of the SARS-CoV-2 virus respectively. At 21 dpi,
394 hamsters in HD and LD were re-challenged with 10^6 TCID₅₀ of the SARS-CoV-2 virus. At 2 and 4
395 dpi, nasal washes, nasal turbinate and lungs were collected from hamsters for viral titration, RNA
396 quantification and transmission electron microscopy examination. (A to C) Viral titers
397 (\log_{10} TCID₅₀/mL) detected in nasal washes (A), nasal turbinates (B) and lungs (C) of prior
398 infected hamsters challenged with the SARS-CoV-2 virus. (D to F) Viral RNA copies (\log_{10} RNA
399 copies/mL) detected in nasal washes (D), nasal turbinates (E) and lungs (F) of prior infected
400 hamsters challenged with SARS-CoV-2. (G and H) Viral sgmRNA copies (\log_{10} sgmRNA
401 copies/mL) detected in nasal turbinates (G) and lungs (H) of prior infected hamsters re-challenged
402 with SARS-CoV-2. One-way analysis of variance (ANOVA) and Tukey's multiple comparisons
403 test were used to analyze the statistical differences of viral titers, RNA copies and sgmRNA copies
404 in nasal washes, nasal turbinates and lungs between different experimental groups ($p > 0.05$, not
405 significant, [ns]; $p < 0.05$, * ; $p < 0.01$, **; $p < 0.001$, ***).

406 **Figure 2** Transmission of SARS-CoV-2 from naïve hamsters to prior infected hamsters. (A)
407 Infectious viral load (\log_{10} TCID₅₀ shown in bars) and viral RNA copies (\log_{10} RNA copies/mL,
408 shown in dots with matched color) detected in nasal washes of the naïve donor hamsters
409 inoculated with 10^6 TCID₅₀ of SARS-CoV-2 and prior infected contact hamsters in the
410 transmission group, which were previously inoculated with 10^5 TCID₅₀ or 10^3 TCID₅₀ of
411 SARS-CoV-2 twenty-one days ago. At 24 hours' inoculation, the donor hamsters and the prior

412 infected contact hamsters were co-housed together in a new cage. (B) Viral titers and viral RNA
413 copies detected in nasal washes of the naïve donor hamsters inoculated with SARS-CoV-2 and
414 those prior infected hamsters in airborne transmission group. At 24 hours' inoculation, the donor
415 hamsters and the prior infected hamsters were transferred to an airborne transmission cage. (C)
416 Viral titers and viral RNA copies detected in nasal washes of the naïve donor hamsters inoculated
417 with SARS-CoV-2 and the prior infected hamsters in the indirect contact transmission group. At
418 48 hours' inoculation, the donor hamsters were removed and transferred to a new cage and prior
419 infected hamsters were placed into the cage housing the naïve hamsters before. Nasal washes were
420 collected from all hamsters in different experiment groups every other day for virus titration and
421 RNA quantification.

422 **Figure 3** Transmission of SARS-CoV-2 from prior infected hamsters to naive hamsters. (A)
423 Infectious viral load (\log_{10} TCID₅₀ shown in bars) and viral RNA copies (\log_{10} RNA copies/mL,
424 shown in dots with matched color) detected in nasal washes of prior infected hamsters inoculated
425 with 10^6 TCID₅₀ of SARS-CoV-2 and the naïve contact hamsters. At 24 hours' inoculation, the
426 prior infected donor hamsters and the naïve contact hamsters were co-housed together in a new
427 cage. (B) Viral titers and viral RNA copies detected in nasal washes of the prior infected hamsters
428 inoculated with SARS-CoV-2 and naive hamsters in airborne transmission group. At 24 hours'
429 inoculation, the prior infected donor hamsters and naive hamsters were transferred to an airborne
430 transmission cage. (C) Viral titers and viral RNA copies detected in nasal washes of prior infected
431 hamsters inoculated with SARS-CoV-2 and naive hamsters in indirect contact transmission group.
432 At 48 hours' inoculation, the donor hamsters were removed and transferred to a new cage and the
433 naive hamsters were placed into the cage housing the prior infected donor hamsters before. Nasal

434 washes were collected from all hamsters every other day for viral titration and RNA
435 quantification.

436 **Figure 4** Transmission of SARS-CoV-2 between prior infected hamsters. A. Infectious viral load
437 (\log_{10} TCID₅₀ shown in bars) and viral RNA copies (\log_{10} RNA copies/mL, shown in dots with
438 matched color) detected in nasal washes of prior infected hamsters inoculated with 10^6 TCID₅₀ of
439 SARS-CoV-2 and the prior infected contact hamsters. At two hours' inoculation, the donor
440 hamsters and the contact hamsters were co-housed together in a new cage. (B) Viral titers and viral
441 RNA copies detected in nasal washes of the prior infected donor hamsters inoculated with
442 SARS-CoV-2 and the prior infected hamsters in airborne transmission group. At two hours'
443 inoculation, the donor hamsters and other hamsters were transferred to the airborne transmission
444 cage. Nasal washes were collected from all hamsters every other day for viral titration and RNA
445 quantification.

446 **Figure 5** Impact of a lower dose infection on SARS-CoV-2 transmission between naïve hamsters
447 and prior infected hamsters. (A) Infectious viral load (\log_{10} TCID₅₀ shown in bars) and viral RNA
448 copies (\log_{10} RNA copies/mL, shown in dots with matched color) detected in nasal washes of the
449 naïve hamsters inoculated with 10^4 TCID₅₀ of SARS-CoV-2 and the prior infected contact
450 hamsters. At 24 hours' inoculation, the naïve donor hamsters and the prior infected contact
451 hamsters were co-housed together in a new cage. (B) Viral titers and viral RNA copies detected in
452 nasal washes of the prior infected donor hamsters inoculated with 10^4 TCID₅₀ of SARS-CoV-2 and
453 naïve contact hamsters. At 24 hours' inoculation, the prior infected donor hamsters and naïve
454 contact hamsters were co-housed together in a new cage. Nasal washes were collected from all
455 hamsters for viral titration and RNA quantification.









