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2 **Main Manuscript for**

3 **Hedgehogs are the major amplifying hosts of severe fever with**  
4 **thrombocytopenia syndrome virus**

5

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31 **Classification:** Biological sciences, Microbiology

32

33 **Abstract**

34 Severe fever with thrombocytopenia syndrome virus (SFTSV) is a tick-borne bandavirus mainly  
35 transmitted by *Haemaphysalis longicornis* in East Asia, mostly in rural areas. To date, the  
36 amplifying host involved in the natural transmission of SFTSV remains unidentified. Our  
37 epidemiological field survey conducted in endemic areas in China showed that hedgehogs were  
38 widely distributed, had heavy tick infestations, and had high SFTSV seroprevalence and RNA  
39 prevalence. After experimental infection of *Erinaceus amurensis* and *Atelerix albiventris*  
40 hedgehogs with SFTSV, robust but transitory viremias were detected, which lasted for around  
41 nine to eleven days. The infected hedgehogs experienced light weight loss and histopathology of  
42 the spleen showed hemorrhagic necrosis and lymphopenia, with infected hedgehogs recovering  
43 after viral clearance. Remarkably, SFTSV transmission cycle between hedgehogs and  
44 nymph/adult *H. longicornis* was easily accomplished under laboratory condition with 100%  
45 efficiency. Furthermore, naïve *H. longicornis* ticks could be infected by SFTSV-positive ticks co-  
46 feeding on naïve hedgehogs, with transstadial transmission of SFTSV also confirmed. We also  
47 found that SFTSV viremia remained high in hedgehogs during hibernation, suggesting that this  
48 mechanism might contribute to the persistence of SFTSV from one year to the next. Of concern,  
49 we recently found evidence of the natural circulation of SFTSV in the urban area of Beijing City in  
50 China involving *H. longicornis* ticks and *E. amurensis* hedgehogs. Our study suggests that the  
51 hedgehogs are the major wildlife amplifying hosts of SFTSV and that urban outbreaks of SFTSV  
52 might occur in the future.

53

54 **Keywords:** SFTSV, hedgehog, *Haemaphysalis longicornis*, transmission, host

55

56 **Main Text**

57

58 **Introduction**

59

60 Severe fever with thrombocytopenia syndrome virus (SFTSV) is a new tick-borne bandavirus first  
61 identified in China in 2009(Yu et al., 2011), followed by Korea in 2011(Denic et al., 2011), Japan  
62 in 2014(Takahashi et al., 2014), Vietnam in 2019(Tran et al., 2019) and Pakistan in 2020(Zohaib  
63 et al., 2020). The symptoms of SFTS include fever, thrombocytopenia, leukocytopenia, and  
64 gastrointestinal disorders, with a case-fatality rate of between 2 and 30%(Liu, He, Huang, Wei, &  
65 Zhu, 2014; S. Liu et al., 2014; Yu et al., 2011). The earliest Chinese cases were reported in the  
66 Dabie mountain range, which is located at the intersection of Henan, Hubei, and Anhui provinces  
67 in central China. Besides the Dabie mountain range, Shandong, Liaoning, and Zhejiang provinces  
68 are the other main hot spots for SFTS in China(J. Sun et al., 2018). Within Zhejiang Province,  
69 Daishan County, an archipelago of islands located in the East China Sea, is one of the most  
70 endemic areas(Fu et al., 2016). The major industry in Daishan County is fishing and tourism,  
71 agriculture is relatively unimportant with only 4000 sheep and 150 cattle on the islands in 2019,  
72 as reported by the local government. As of 2020, SFTS cases have been reported in most other  
73 Chinese provinces(Lin et al., 2020; J. Sun et al., 2018; Zhu et al., 2019).

74 *Haemaphysalis longicornis* (Asian long-horned tick) is the major vector for SFTSV and the  
75 dominant human-biting tick in the SFTSV endemic areas(Li et al., 2016; Yun et al., 2015; G.  
76 Zhang, Zheng, Tian, & Li, 2019). *H. longicornis* has both bisexual and parthenogenetic  
77 populations, with the parthenogenetic populations being widely distributed in China and strongly  
78 correlated with the distribution of SFTS cases (X. Zhang et al., 2022). *H. longicornis* ticks go  
79 through a three-stage life cycle (larva, nymph, and adult). At each stage, they feed on a wide  
80 range of wild and domestic animals including mammals, birds, companion animals and  
81 livestock(Zhao et al., 2020).

82 Extensive reports suggest that *H. longicornis* is the reservoir of SFTSV(Luo et al., 2015; S. W.  
83 Park et al., 2014; Zhuang et al., 2018). However, the transstadial transmission efficiencies of

84 SFTSV from egg pools to larvae pools, larval pools to nymph pools and nymph pools to adults  
85 were 80%, 92%, 40% or 100%, 100%, 50% under laboratory conditions according to two  
86 reports(Y. Y. Hu et al., 2020; Zhuang et al., 2018). Correspondingly, the SFTSV prevalence was  
87 extremely low in different developmental stages of host-seeking *H. longicornis* ticks collected  
88 from vegetation, ranging from 0.2% to 2.2%(Luo et al., 2015; S. W. Park et al., 2014; Wang et al.,  
89 2015). These findings suggest that ticks alone are not sufficient to maintain a reservoir of SFTSV  
90 in the natural environment, therefore one or more additional amplifying hosts are required.

91 Antibodies to SFTSV and viral RNA have been detected in a wide range of domestic animals,  
92 including goats, cattle, dogs, and pigs, and wild animals such as shrews, rodents, weasels, and  
93 hedgehogs. The highest seroprevalence was found in sheep (69.5%), cattle (60.4%), dogs  
94 (37.9%) and chickens (47.4%)(Chen et al., 2019; Huang et al., 2019; Niu et al., 2013). Given that  
95 most of the SFTS patients are farmers, who have frequent contacts with many of the domestic  
96 and wild animals listed above, this makes understanding the epidemiology of SFTSV both difficult  
97 and complex.

98 Hedgehogs belong to the family *Erinaceinae*, which includes twenty four genera and are widely  
99 distributed in the Eurasian continent and Africa(He et al., 2012). Some genera have even been  
100 introduced into countries with no indigenous hedgehogs, including Japan and New  
101 Zealand(Brockie, 1975; ISAAC, 2005). The Amur hedgehog *Erinaceus amurensis* is closely  
102 related to the European hedgehog, *Erinaceus europaeus*, but is slightly bigger and lighter in  
103 color. It is native to Amur Oblast and Primorye in Russia, the Korean Peninsula, and is common  
104 in northern and central China. The African pygmy hedgehog *Atelerix albiventris*, native to central  
105 and eastern Africa, has been introduced into many countries as pets, including China, where they  
106 are available in many suburban petshops(Brockie, 1975; ISAAC, 2005). Both the Amur hedgehog  
107 and the African pygmy hedgehog can become heavily infested by all kinds of ticks and are known  
108 to carry many zoonotic diseases, such as Tickborne encephalitis virus, Bhanja virus, and Tahyna  
109 virus(Dziemian, Sikora, Pilacinska, Michalik, & Zwolak, 2015; Jahfari et al., 2017; Riley & Chomel,  
110 2005). Hedgehogs are poikilothermal animals and hibernate during winter. During hibernation,  
111 their metabolism and immune system are suppressed (Bouma, Carey, & Kroese, 2010) which  
112 has led to the suspicion that hibernating hedgehogs contribute to the long-term persistence of  
113 these viruses(Simkova, 1966). A few previous studies reported that SFTSV antibodies and RNA  
114 were detected in *E. amurensis* in Shandong and Jiangsu Province. However, the prevalence of  
115 SFTV infection appeared low as compared to that in other animals such as goats, sheep, and  
116 cattle(Li et al., 2016; Y. Sun et al., 2017).

117 In China, the density of large wild animals is extremely low, especially in East China where SFTS  
118 is endemic. Instead, the most abundant wildlife in these areas are rodents and insectivores(Jiang,  
119 Liu, Wu, Jiang, & Zhou, 2017). However, the potential role of rodents in the transmission of  
120 SFTSV was refuted when it was shown that immunocompetent rodents cannot develop SFTSV  
121 viremia after artificial inoculation(Matsuno et al., 2017). In contrast, hedgehogs are the only small  
122 wild animals which consistently show high SFTSV seroprevalence, high density, as well as high  
123 *H. longicornis* infestation in the SFTS endemic areas(Li et al., 2014; Y. Sun et al., 2017). They  
124 are widely distributed both in the urban and wild ecosystem(Smith et al., 2010), which has led us  
125 to speculate that hedgehogs might play an important role in the natural circulation of SFTSV in  
126 China. To test this hypothesis, we first carried out an epidemiological survey to confirm the role of  
127 hedgehogs as potential wild amplifying hosts for SFTSV. Then a series of linked laboratory  
128 experiments were performed to investigate the susceptibility and tolerance of hedgehogs to  
129 SFTSV infection, and to establish the transmission of infection between hedgehogs and *H.*  
130 *longicornis* ticks at both the nymph and adult stages.

131

## 132 **Results**

133

### 134 **Field survey of hedgehogs in SFTS endemic areas**

135 To confirm the role of hedgehogs as potential wild amplifying hosts for SFTSV, we firstly  
136 performed an animal survey in Daishan County, an archipelago of islands in the East China Sea  
137 (Figure 1A). Daishan County is the worst affected area for SFTS in Zhejiang Province (Fu et al.,

138 2016) and between 2011 and 2019 one hundred and thirty-three SFTS cases were reported by  
139 Daishan CDC. SFTS cases were reported on all the major Daishan County islands including  
140 Daishan Island, Qushan Island and Changtu Island, with the exception of Xiushan Island, even  
141 though Xiushan Island has a similar landscape, vegetation, and population density as Daishan  
142 Island, Qushan Island and Changtu Island (Fig. 1B). Small mammal traps were set, and the  
143 following numbers of wildlife caught, 33 on Daishan Island and 75 on Xiushan Island. On Daishan  
144 Island, 28% of the captured small mammals were *E. amurensis* (Amur hedgehog), 18% were  
145 *Rattus. Norvegicus* (brown rat), 36% were *Sorex araneus* (common shrew), and 18% were  
146 *Apodemus agrarius* (striped field mouse). On Xiushan Island no hedgehogs were caught, 48% of  
147 the small mammals caught were *R. norvegicus*, 44% were *S. araneus* and 8% were *Rattus losea*  
148 (lesser ricefield rat) (Fig. 1C). Antibody testing showed that 3/9 (33%) of *E. amurensis* hedgehogs  
149 from Daishan Island were positive for SFTSV (Fig. 1D). Hedgehogs are abundant in the two  
150 villages in Daishan Island, with an estimated population density of greater than 80 individuals per  
151 square kilometer based on the results of the trapping study (Table 1). In addition, these nine  
152 trapped hedgehogs were all heavily infected by ticks with an average of 145 ticks per hedgehog,  
153 including *H. longicornis* (Table 2).

154

155 **Table 1.** The density of hedgehogs in rural and urban areas.

156

Site	Location	Density
Daao village*	Daishan County, Zhejiang Province	>80
Dongsha village*	Daishan County, Zhejiang Province	>90
Olympic Forest Park <sup>&amp;</sup>	Chaoyang District, Beijing	>60
Southeast Community <sup>&amp;</sup>	Haidian District, Beijing	>75

157

158 The density was calculated by the number of trapped hedgehogs divided by the area (Number of  
159 animals per square kilometer). \*Rural; <sup>&</sup>urban.

160

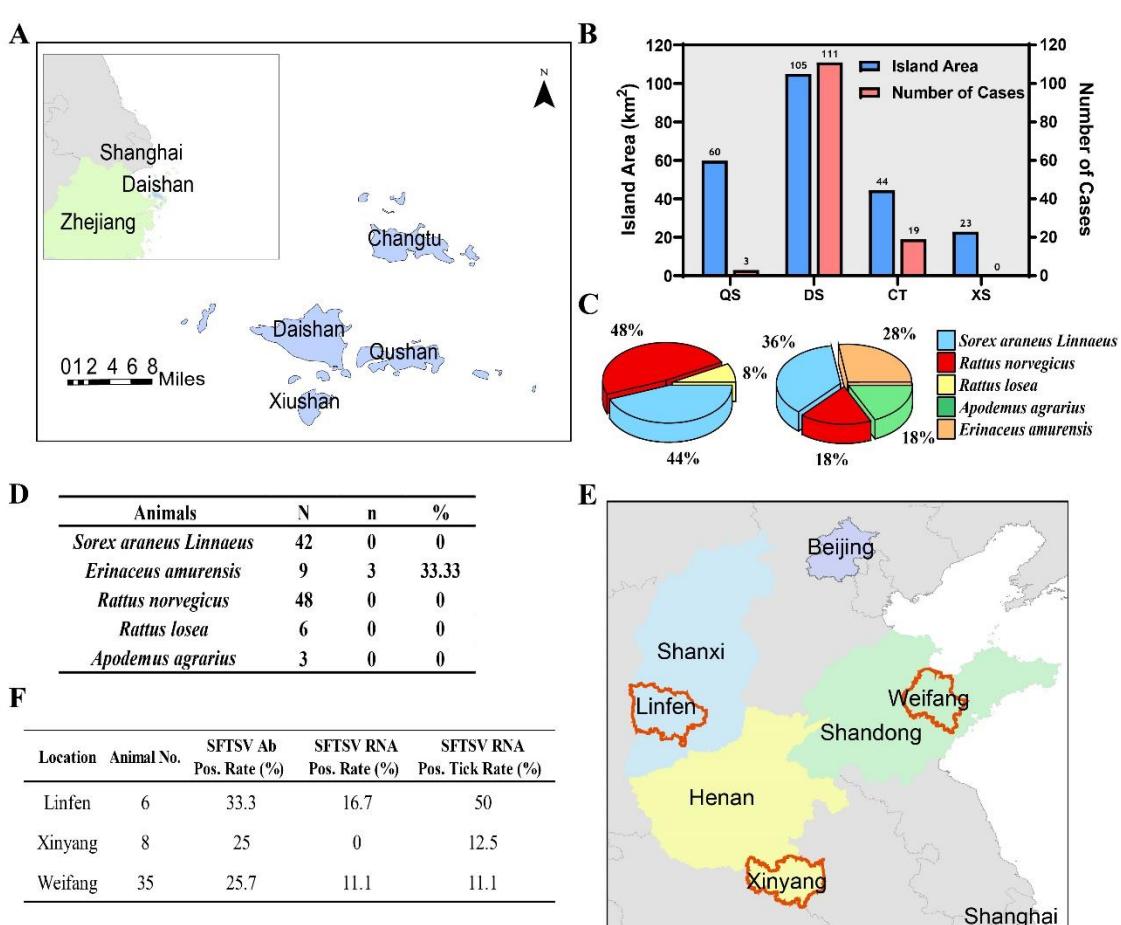
161 **Table 2.** Average number of ticks collected from wild mammals captured in Daishan County.

162

Animal	Number
<i>Sorex araneus Linnaeus</i>	1.5
<i>Rattus norvegicus</i>	1
<i>Rattus losea</i>	0
<i>Apodemus agrarius</i>	0
<i>Erinaceus amurensis</i>	145

163

164 Additional *E. amurensis* hedgehog serum samples were collected from trapping studies  
 165 conducted in other SFTS endemic areas, including Weifang City of Shandong Province, Linfen  
 166 City of Shanxi Province, and Xinyang City of Henan Province. SFTSV antibodies were detected in  
 167 9/35 (25.7%), 2/6 (33.3%) and 2/8 (25%) of hedgehogs from Weifang City, Linfen City and  
 168 Xinyang City, respectively; 11.1%, 16.7% and 0 of hedgehogs were tested positive for SFTSV  
 169 RNA, respectively; and 11.1%, 50% and 12.5% of hedgehog were infected by ticks positive for  
 170 SFTSV RNA (Fig. 1E and 1F). We believe these results strongly support our hypothesis that  
 171 hedgehogs play an important role in the natural circulation of SFTSV.  
 172



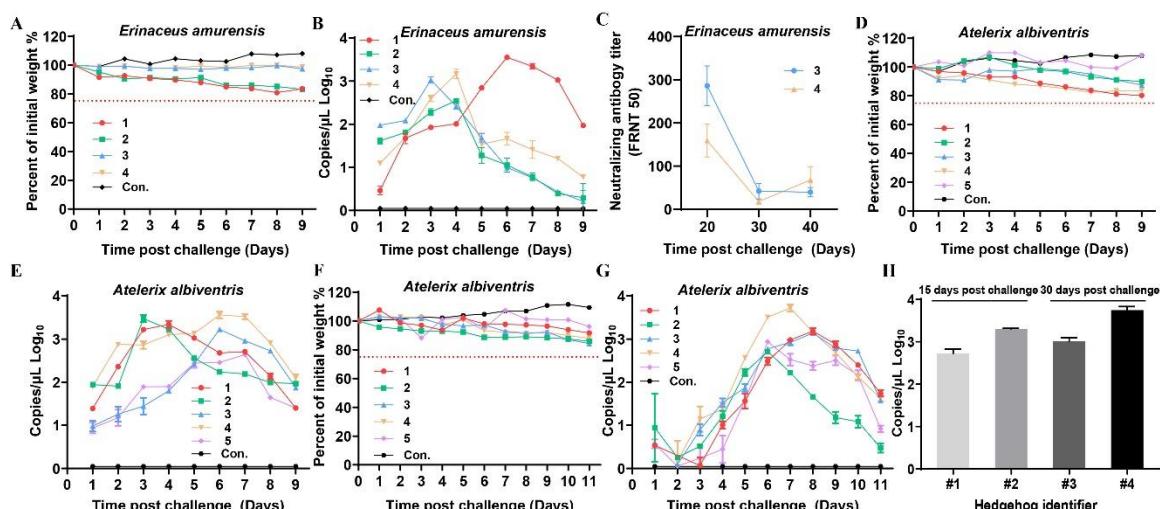
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174  
 175 **Figure 1.** The association between hedgehogs and SFTSV endemic. (A) Locations of the major  
 176 islands of Daishan County, Zhejiang Province. (B) Land area and SFTS case numbers for major  
 177 islands in Daishan County. DS, Daishan Island; QS, Qushan Island; XS, Xiushan Island; CT,  
 178 Changtu Island. (C) Species and relative rate of wild animals collected on Xiushan Island (left)  
 179 and Daishan Island (right). (D) Seroprevalence of SFTSV in wild animals captured in Xiushan  
 180 Island and Daishan Island. N, number of sampled animals; n, number of sampled animals  
 181 positive for SFTSV antibody; %, percentage of sampled animals positive for SFTSV antibody. (E)  
 182 Locations of Weifang City of Shandong Province, Linfen City of Shanxi Province, Xinyang City of  
 183 Henan Province, where hedgehogs were collected. (F) Epidemiological analysis of trapped  
 184 animals. Seroprevalence of SFTSV in hedgehogs was measured by ELISA against SFTSV  
 185 nucleocapsid protein. Viral RNA was tested by PCR. Pos. is the abbreviation of positive. "SFTSV  
 186 RNA Pos. Tick Rate" indicates the ratio of hedgehogs infested with SFTSV RNA positive ticks.  
 187

### The susceptibility of hedgehogs to experimental infection with SFTSV

188 Four male and female *E. amurensis* hedgehogs (6-12 months old) were inoculated with  $4 \times 10^6$   
 189 FFU of SFTSV by intraperitoneal (i.p.) route. A viremia of around 9 d was observed in all animals,  
 190 with peak titers of  $3.1 \log_{10}$  RNA copies/ $\mu$ l at d 3-6, suggesting viral multiplication. Two *E.*  
 191 *amurensis* hedgehogs showed a mild weight loss of less than 25% by 9 d (Fig. 2A-2B).  
 192 Neutralizing antibody titer against SFTSV in the sera of two *E. amurensis* hedgehogs were  
 193 measured at 20, 30 and 40 d post-infection (dpi). In contrast to the stable humoral immune  
 194 response of experimental dogs(Niu et al., 2013), the neutralizing antibody titer decreased quickly  
 195 and was almost eliminated by d 40 (Fig. 2C).  
 196

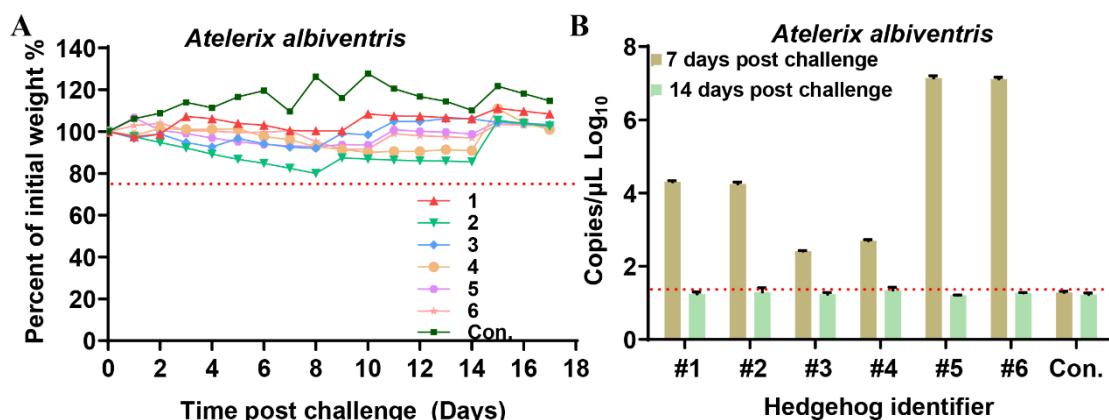
197 Groups of five male and female *A. albiventris* hedgehogs (6-12 months) were inoculated with  $4 \times$   
 198  $10^6$  FFU of SFTSV by intraperitoneal (i.p.) and subcutaneous (i.s.) route, respectively. A viremia  
 199 of around 9 to 11 d was observed in all ten animals, with peak titers of  $3.2 \log_{10}$  RNA copies/ $\mu$ l at  
 200 d 3-7 for i.p. route and titers of  $3.1 \log_{10}$  RNA copies/ $\mu$ l at d 6-8 for i.s. route, respectively (Fig.  
 201 2D-2G). Most animals showed mild weight loss of less than 20%. These results suggest that *E.*  
 202 *amurensis* and *A. albiventris* hedgehogs could develop similar viremias independent of  
 203 inoculation routes.



204  
 205 **Figure 2.** SFTSV viremia in experimentally infected *E. amurensis* and *A. albiventris* hedgehogs.  
 206 Hedgehogs were challenged (i.p. or i.s.) with  $4 \times 10^6$  FFU of SFTSV Wuhan strain and then  
 207 monitored for weight change (A and D and F) and viremia (B and E and G), tested by Real-time  
 208 PCR as RNA copies per  $\mu$ l of serum (B and E) (error bars represent SD). Control (Con.) was  
 209 mock infected with PBS. (C) SFTSV neutralizing antibody titer in two *E. amurensis* hedgehogs  
 210 were monitored at 20, 30, 40 dpi. (H) Hibernation extended the course of SFTSV viremia in *A.*  
 211 *albiventris*. Four hedgehogs were challenged (i.p.) with  $4 \times 10^6$  FFU of SFTSV and then kept at 4

212 °C to trigger hibernation. Viremia in #1 and #2 was monitored 15 dpi, while #3 and #4 at 30 dpi  
213 (error bars represent SD).

214  
215 There was a possibility that the observed weight loss could have been iatrogenic, since the blood  
216 samples were taken directly from the heart. Heart sampling was necessary because rapid blood  
217 coagulation makes sampling from superficial veins very difficult in hedgehogs (Lewis, 1976). To  
218 test whether heart sampling was the cause of the weight loss, a further six *A. albiventris* were i.p.  
219 inoculated with SFTSV as above and bled only twice on d 7 and d 14, instead of daily. We found  
220 that only one hedgehog showed a weight loss of 20% and recovered by d 16 (Fig. S1A). In these  
221 six hedgehogs, SFTSV viremia was detected at d 7 and disappeared at d 14 (Fig. S1B). Overall,  
222 these results demonstrate that both *E. amurensis* and *A. albiventris* can develop a similar SFTSV  
223 viremia after experimental infection, without significantly compromising their overall health.  
224 However, *E. amurensis* hedgehogs are shy and easy to die during transport because of the  
225 stress response. Thus, we performed most of the following experiments with *A. albiventris* due to  
226 the stable supply in the local pet store.



227  
228 **Figure S1.** SFTSV viremia and weight change in experimentally infected *A. albiventris*  
229 hedgehogs bled twice. Hedgehogs were challenged (i.p.) with  $4 \times 10^6$  FFU of SFTSV Wuhan  
230 strain and then monitored for weight loss everyday (A) and viremia on d 7 and 14, measured by  
231 Real-time PCR as RNA copies per  $\mu\text{L}$  of serum (B) (error bars represent SD). Each line and bar  
232 i n d i c a t e o n e h e d g e h o g .

### 233 234 **SFTSV viremia during the hibernation of hedgehogs**

235 Four *A. albiventris* were inoculated with  $4 \times 10^6$  FFU of SFTSV and kept at 4 °C to trigger  
236 hibernation. Two of the hedgehogs came out of hibernation at d 15 with viremias of 2.7 and 3.3  
237 log<sub>10</sub> RNA copies/ $\mu\text{L}$  respectively, whilst the other two hedgehogs continued in hibernation until d  
238 30 with viremias of 3.0 and 3.7 log<sub>10</sub> RNA copies/ $\mu\text{L}$ . All the viremias measured in these  
239 hibernating hedgehogs were comparable to the peak virus titers previously measured in the non-  
240 hibernating hedgehogs (Fig. 2H). However, the duration of viremia in these 4 hibernating  
241 hedgehogs was much longer than that recorded in the non-hibernating hedgehogs, suggesting  
242 that hibernation could potentially extend the course of SFTSV viremia in hedgehogs and  
243 contribute to the overwintering of SFTSV in the field.

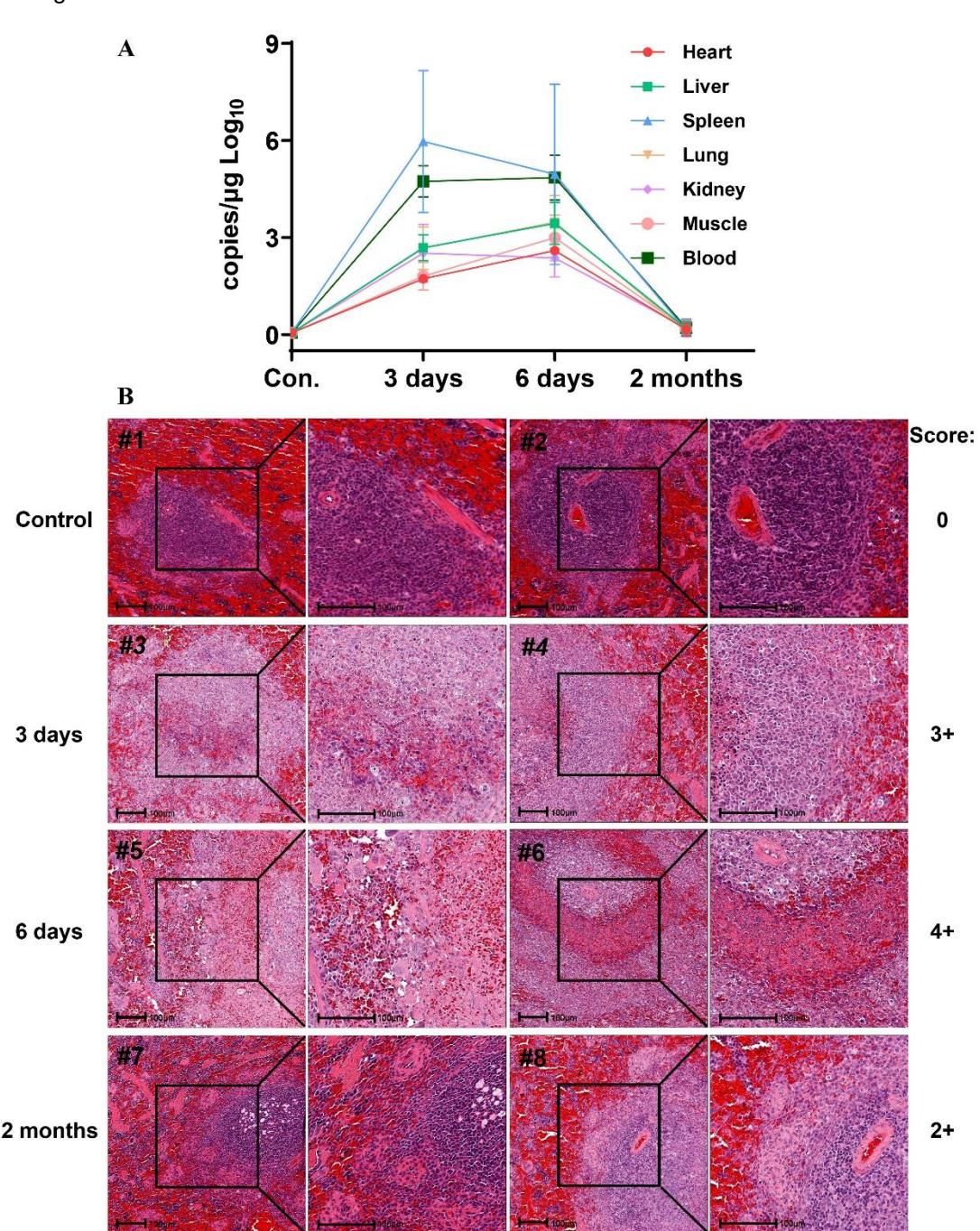
### 244 **SFTSV induced pathology in hedgehogs**

245 To assess the pathological changes in hedgehogs resulting from SFTSV infection, six *A.  
246 albiventris* were i.p. inoculated with  $4 \times 10^6$  FFU of SFTSV. Two animals were sacrificed at 3  
247 days, 6 days and 2 months post-infection and their organs were removed for viral RNA evaluation  
248 and hematoxylin and eosin (H&E) staining. A robust viremia was detected on d 3 and d 6 but was  
249 negative at 2 months after infection. The highest level of viral RNA was observed in the spleen,  
250 followed by the blood, while the lowest was in the heart (Fig. 3A). H&E-stained slides from the

251 spleen showed hemorrhagic necrosis and lymphopenia at d 3 and d 6. The severity of the lesions  
252 was assessed as +++ and ++++ on day 3 and 6 respectively but the lesions had largely  
253 recovered by 2 months with a severity score of ++ (Fig. 3B). These results further confirmed that  
254 hedgehogs show a high tolerance to SFTSV without obvious long-term or permanent pathological  
255 changes.

256

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changes.



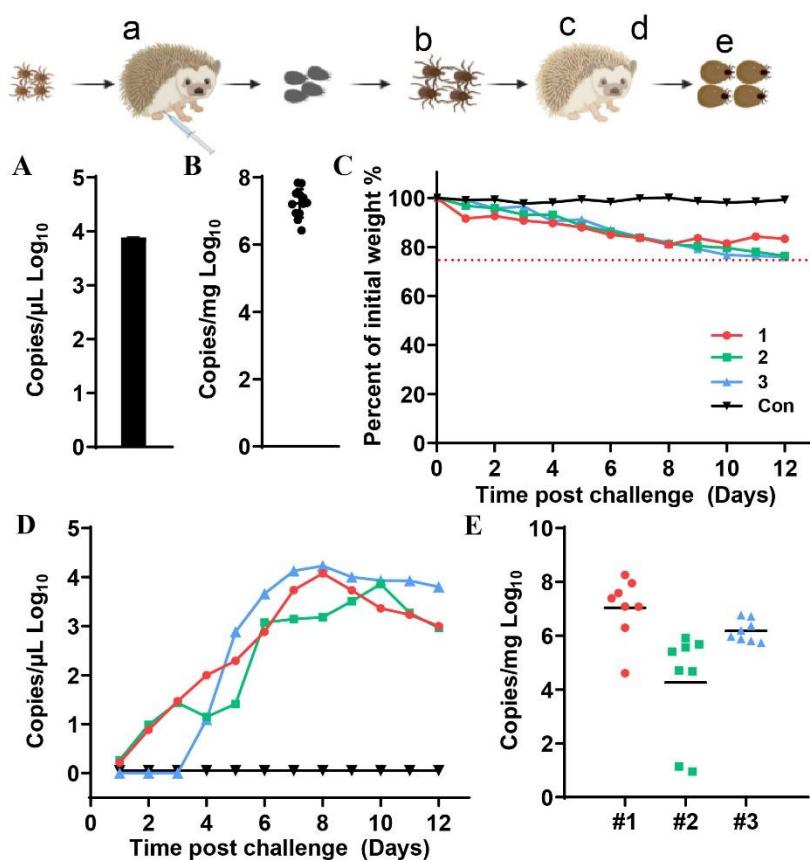
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258 **Figure 3.** Pathology of SFTSV infected *A. albiventris* hedgehogs. Six hedgehogs were  
259 challenged (i.p.) with  $4 \times 10^6$  FFU of SFTSV Wuhan strain and two were mock infected by PBS as  
260 control (Con.). Two hedgehogs were killed at indicated time points to test the viral load in the  
261 organs (A) and pathology of the spleen (B). (A) SFTSV viral load in organs was measured by  
262 Real-time PCR. (B) Spleen samples were H&E stained for the pathological interpretation, with the  
263 severity of pathological changes shown beside the image. Size bars indicate 100  $\mu$ m.  
264

265 **Transmission of SFTSV by *H. longicornis* between ticks and hedgehogs**

266 Lab-adapted *H. longicornis* ticks and *A. albiventris* hedgehogs were used to model the natural  
267 transmission of SFTSV hypothesized to occur in the wild. Naïve *H. longicornis* nymphs were fed  
268 on hedgehogs infected by i.p. inoculation with  $4 \times 10^6$  FFU of SFTSV at d 0. Viremia of  $3.8 \log_{10}$   
269 RNA copies/ $\mu$ L was detected in hedgehogs at d 5 (Fig. 4A) and fully engorged nymphs dropped  
270 off between d 4 to 8. After molting, the adult ticks tested 100% positive for SFTSV, with a level of  
271  $7.2 \log_{10}$  RNA copies/mg tick (Fig. 4B).  
272

273 Three naïve hedgehogs were each fed on by eight SFTSV-carrying adult ticks at d 0. Weight and  
274 viremia were monitored for 12 d. A slow weight loss of less than 25% was observed by d 12 and  
275 the viremia peaked between d 8-10 at  $4.1 \log_{10}$  copies/ $\mu$ L. After peaking, the viraemia decreased  
276 slowly until the 3 hedgehogs were euthanized on d 12 (Fig. 4C and 4D). The fully engorged ticks  
277 were collected between d 7 to 10 and then tested. All 24 ticks were still positive for SFTSV RNA  
278 (Fig. 4E). We believe that these data strongly suggest that SFTSV can be efficiently transmitted  
279 between hedgehogs and *H. longicornis* ticks, and that transstadial transmission occurs within *H. longicornis* ticks.



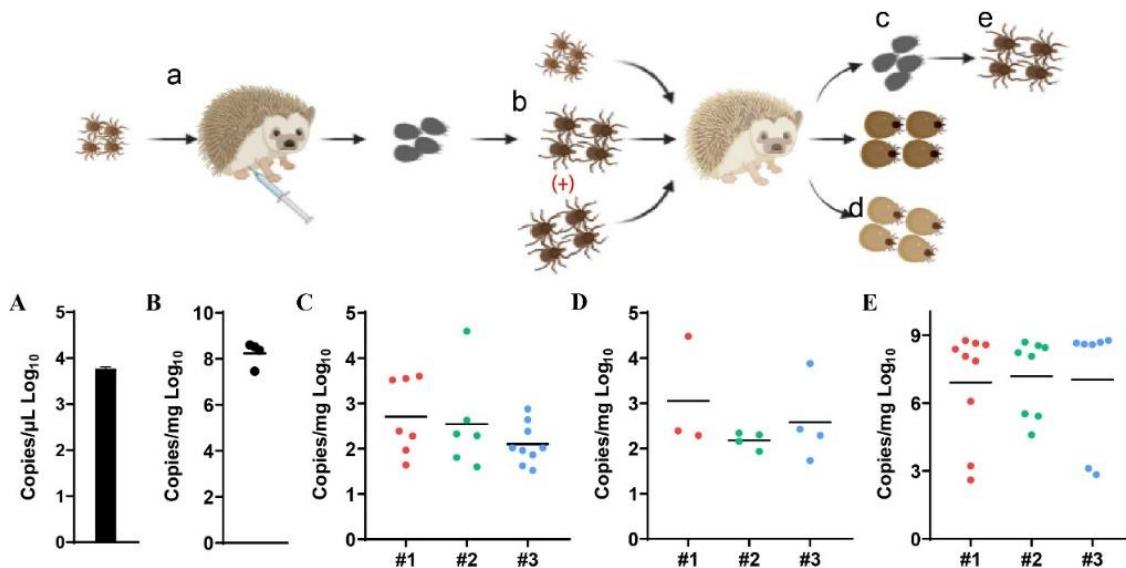
280 **Figure 4.** Transmission of SFTSV between *H. longicornis* ticks and *A. albiventris* hedgehogs.  
281 Hedgehogs were i.p. inoculated with  $4 \times 10^6$  FFU of SFTSV Wuhan strain and naïve nymphs were

283 fed on the hedgehogs at the same time. (A) SFTSV viremia in the hedgehogs at 5 dpi as  
284 measured by Real-time PCR. (B) SFTSV RNA copies in the adult ticks after molting as shown by  
285 RNA copies per mg of tick. (C) Weight change and (D) SFTSV viremia in naïve hedgehogs bitten  
286 by SFTSV-carrying adult ticks were monitored for 12 d. (E) SFTSV RNA level in the engorged  
287 adult ticks from three hedgehogs as shown by RNA copies per mg of tick. Each dot indicates one  
288 tick (error bars represent SD). # indicates hedgehog identifier. The experimental process is  
289 graphically displayed above the plots and read from left to right, with the lower-case letters (a-e)  
290 corresponding to the upper-case letters of the main panels (A-E).  
291

## 292 Hedgehogs are the amplifying host for SFTSV

293 SFTSV can be transmitted both transovarially and transstadially in *H. longicornis*, however, a  
294 decreased efficiency has been observed during passaging (Zhuang et al., 2018). Thus, an  
295 amplifying host will be necessary to improve the transmission efficiency. SFTSV-positive adult *H.*  
296 *longicornis* ticks were prepared as described above with 100% efficiency (Fig. 5A and 5B).  
297 Next, 5 of the SFTSV-carrying adult *H. longicornis* ticks were fed together with 14 to 16 naïve  
298 nymphs and 3 to 4 naïve adult ticks on each of three naïve *A. albiventris* hedgehogs. The fully  
299 engorged ticks were collected between 7 to 10 d post biting and tested for the viral RNA levels.  
300 The viral load in the engorged nymphs and previously naïve adults were 2.5 and 2.7 log<sub>10</sub> RNA  
301 copies/mg tick respectively (Fig. 5C and 5D). After the nymphs molted, the adult ticks tested  
302 100% positive for SFTSV, with a level of 6.9 log<sub>10</sub> RNA copies/mg tick (Fig. 5E). Thus, these  
303 results suggest that hedgehogs could be acting as an amplifying host for SFTSV.  
304

305



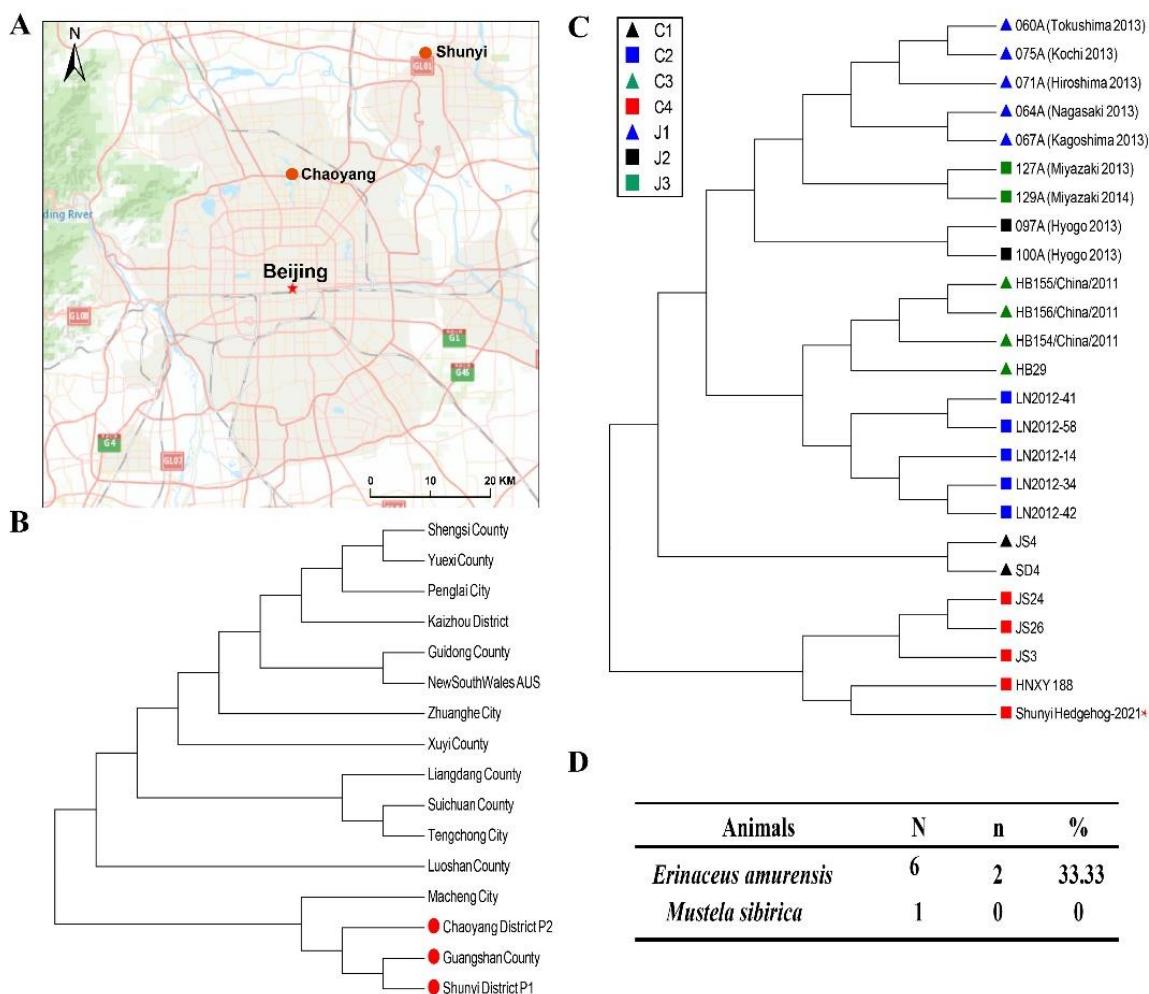
306

307 **Figure 5.** Naïve *H. longicornis* ticks infected by SFTSV through co-feeding with SFTSV positive  
308 ticks on naïve *A. albiventris* hedgehogs. Hedgehogs were i.p. inoculated with  $4 \times 10^6$  FFU of  
309 SFTSV Wuhan strain and nymph ticks were fed on the hedgehogs at the same time. (A) SFTSV  
310 viremia in hedgehogs at 5 dpi as measured by Real-time PCR. (B) SFTSV RNA copies in adult  
311 ticks after molting as shown by RNA copies per mg of tick. After molting, the SFTSV-carrying  
312 adult ticks and naïve nymph/adult *H. longicornis* ticks were fed on three naïve *A. albiventris*  
313 hedgehogs. The fully engorged ticks were collected 7 to 10 d post biting. (C-E) SFTSV RNA level  
314 was monitored in ticks as shown by RNA copies per mg of tick. Each dot indicates one tick. (C)  
315 Engorged nymph ticks. (D) Engorged adult ticks. (E) Adults molted from (C). The experimental

316 process is graphically displayed above the plots and read from left to right, with the lower case  
317 letters (a-e) corresponding to the upper case letters of the main panels (A-E). (+) indicates  
318 SFTSV infected ticks.  
319

320 **Natural circulation of SFTSV in the urban area**

321 The density of hedgehogs in two rural villages from Daishan County in Zhejiang Province and in  
322 two urban communities from Chaoyang and Haidian District in Beijing City were found to be very  
323 similar (Table 1). To investigate the potential natural circulation of SFTSV in the urban setting, we  
324 carried out field surveys on small mammals and parasitic ticks at two locations in Beijing City in  
325 2021, one small park surrounded by up-market gated communities in Shunyi District and the  
326 Olympic Forest Park in Chaoyang District, where parthenogenetic *H. longicornis* ticks were  
327 discovered in a previous survey in 2019 (Fig. 6A)(X. Zhang et al., 2022). Six *E. amurensis*  
328 hedgehogs were caught in the Shunyi location and showed 2/6 (33%) SFTSV seroprevalence  
329 (Fig. 6D). *H. longicornis* ticks collected from vegetation at the same location tested positive for  
330 SFTSV RNA, clustering into lineage C4, similar to the strains of SFTSV collected in Xinyang City  
331 (Fig.6C). In contrast, no SFSTV RNA or antibody was detected in animal serum samples and  
332 parasitic ticks collected at the Chaoyang sample site. Through phylogenetic analysis of the whole  
333 mitochondrial sequences, we further found that the parthenogenetic *H. longicornis* ticks collected  
334 from the Shunyi and Chaoyang districts were closely related to those from Guangshan County,  
335 Xinyang City in Henan Province (Fig. 6B). These results imply that both the tick and SFTSV  
336 collected in Shunyi Distict might have originated from Xinyang City, one of the original SFTS hot  
337 spots. Although no local SFTS cases have been reported in Beijing yet, our results suggest that a  
338 population of hedgehogs and *H. longicornis* ticks could maintain the circulation of SFTSV in the  
339 urban ecosystem, which might result in urban SFTSV epidemics in the future.  
340



341  
342 **Figure 6.** Natural circulation of SFTSV in the urban Beijing. (A) Locations studied in Shunyi  
343 District and Chaoyang District. (B) Phylogenetic analysis of the parthenogenetic population of  
344 Asian long-horned ticks. Maximum likelihood tree was established with the mitochondrial  
345 genomes of *H. longicornis* collected in Chaoyang District and Shunyi District (Chaoyang  
346 DistrictP2, accession number, OL335942 and ShunyiDistriceP1 accession number, OL335941)  
347 (red) and from SFTS endemic areas(X. Zhang et al., 2022). (C) Maximum likelihood tree was  
348 established with the L Segments of SFTSV isolate in Shunyi (Shunyi-hedgehog-2021) and  
349 isolates from SFTS endemic areas(Shi et al., 2017; Yoshikawa et al., 2015). HNXY188 was  
350 isolated from Xinyang City, Henan Province. SFTSV lineages were illustrated by colors and  
351 shapes. (D) Seroprevalence of animals caught in Shunyi district. N, number of sampled animals;  
352 n, number of sampled animals positive for SFTSV antibody; %, percentage of sampled animals  
353 positive for SFTSV antibody.

354

## 355 Discussion

356

357 Viremia in the vertebrate host is important for the arbovirus to transmit from host to vector.  
358 Previous epidemiological surveys and experimental infections have revealed that many wild and  
359 domesticated animals are susceptible to SFTSV infection(Chen et al., 2019). However, these  
360 studies had similar findings that most vertebrate animals were only sub-clinically infected with  
361 SFTSV, with a limited viremia(Casel, Park, & Choi, 2021). For example, 80% of goats developed  
362 a viremia, after i.s. inoculation with  $10^7$  PFU of SFTSV, which lasted for only less than 24 h(Jiao

363 et al., 2015). Similarly, beagle dogs intramuscularly inoculated with  $2.51 \times 10^7$  TCID50 of SFTSV,  
364 only had a detectable viremia at d 3(S. C. Park et al., 2021). Furthermore, the efficient  
365 transmission of SFTSV between tick vectors and these potential wild animal hosts has not been  
366 proven. In this study, robust viremias of about  $10^3$  RNA copies/ $\mu$ l were consistently detected in  
367 both native *E. amurensis* and exotic *A. albiventris* hedgehogs after i.p. or i.s. inoculation with  $4 \times$   
368  $10^6$  FFU of SFTSV at 100% efficiency and lasted for nine to eleven days, which provides basis for  
369 the effective transmission of SFTSV from host to tick.  
370

371 Hedgehogs were highly tolerated to SFTSV infection, with slight weight loss and pathology which  
372 recovered after the clearance of virus. SFTSV seroconversion was observed. In contrast to the  
373 stable humoral immune response of experimental dogs and convalescent patients(L. Hu et al.,  
374 2021; Niu et al., 2013), the antibody titers in hedgehogs decreased quickly, suggesting that  
375 studies measuring the seroprevalence of wild hedgehogs may have underestimated the true  
376 prevalence of infection and that hedgehogs might even be vulnerable to reinfection by SFTSV.  
377

378 *H. longicornis* overwinters mostly as nymphs, but with an SFTSV positive rate of only 4% as  
379 measured by pool(Kim et al., 2020). Thus, we speculate that their role in overwintering of disease  
380 may be quite limited. Hedgehogs are involved in the overwintering of many pathogens during  
381 hibernation (Nosek & Grulich, 1967; Simkova, 1966) which could include SFTSV. Our results  
382 suggest that the SFTSV viremia can be extended from nine days, when non-hibernating, to at  
383 least one month during hibernation, and with a similar peak viremia to that seen in non-  
384 hibernating hedgehogs.  
385

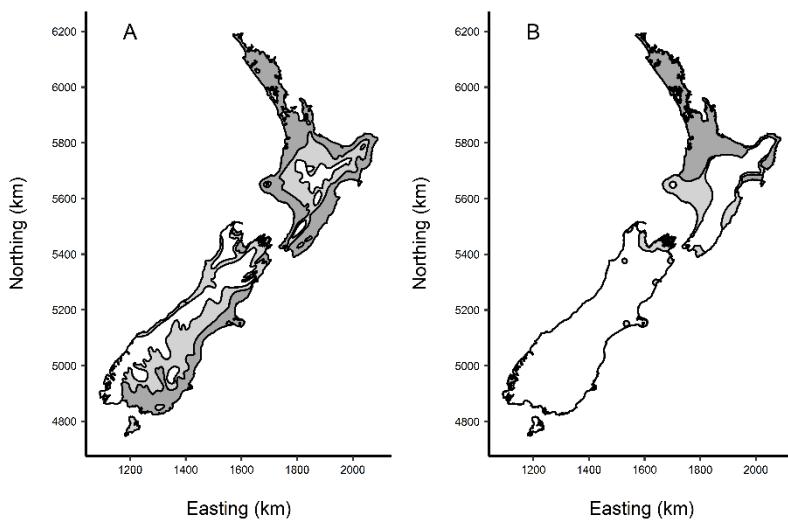
386 To meet the requirement for hedgehogs to be considered as important maintenance hosts for  
387 SFTSV, the transmission cycle between vector and host needs to be established. Using lab-  
388 adapted *H. longicornis* ticks and *A. albiventris* hedgehogs, this study conclusively showed  
389 efficient infection transmission from nymph or adult ticks to hedgehogs, efficient infection  
390 transmission from hedgehogs to nymph or adult ticks and transstadial infection transmission from  
391 nymph to adult tick. It is important to emphasize that these results were observed in 100% of  
392 tested subjects. Naïve nymph and adult *H. longicornis* ticks co-feeding with SFTSV-infected adult  
393 ticks on naïve hedgehogs were also 100% infected. We believe our results clearly show that  
394 hedgehogs fulfill the requirements to be considered the competent amplifying hosts for SFTSV.  
395 It's still plausible that other animals or birds could also maintain the natural circulation of SFTSV.  
396 For example, experimentally inoculated spotted doves (*Streptopelia chinensis*) can develop  
397 SFTSV viremia, however, the transmission between *H. longicornis* ticks and spotted doves is not  
398 proven(Li et al., 2019).  
399

400 To conclude that hedgehogs are the major amplifying hosts of SFTSV in the real world,  
401 abundance, tick-association, geographic distribution in areas of transmission, and field exposure  
402 need to be investigated. Our initial survey in SFTSV endemic Daishan Island and non-endemic  
403 Xiushan Island reveals that the existence of hedgehogs was correlated to SFTSV transmission.  
404 The epidemiological surveys we conducted in four SFTSV endemic provinces consistently  
405 showed high SFTSV seroprevalence and that the population density of hedgehogs in SFTSV  
406 endemic areas can be much higher than 60 animals per  $\text{km}^2$ . Also, Hedgehogs are heavily  
407 infested by ticks including *H. longicornis* with a density of 145 ticks per animal as observed in  
408 Daishan Island. Hedgehogs are widely distributed across farms and rural communities, which  
409 contain the people most likely to be bitten by *H. longicornis* carrying SFTSV(Li et al., 2014; Y.  
410 Sun et al., 2017). Furthermore, hedgehogs share the same environment as domestic animals  
411 such as dogs, goats, and cows, which are also natural hosts for *H. longicornis* ticks and showing  
412 high seroprevalence to SFTVS. Thus, it is possible that humans and domestic animals get  
413 similarly infected by ticks which had previously fed on SFTSV-positive hedgehogs at an earlier  
414 stage in their life cycle. As previously stated, there are few large wild animals in SFTSV endemic  
415 areas in China and the most common animals are rodents and insectivores. Tests on rodents  
416 have shown that they are not capable of maintaining infection(Matsuno et al., 2017). Our results

417 conclusively show that of the mammals present in rural China, the hedgehogs meet all the  
418 requirements of major amplifying hosts for SFTSV.

419  
420 Beijing is the capital of China and is abundant in *H. longicornis* ticks and *E. amurensis*  
421 hedgehogs, between which local transmission of SFTSV were detected recently in an urban  
422 community in Shunyi District. Considering that there are no livestock, poultry, and stray dogs in  
423 this area, it is reasonable to infer that SFTSV circulation can be maintained by just hedgehogs  
424 and *H. longicornis* ticks in an urban area. So far, no human SFTS cases reported, but further  
425 surveillance is warranted.

426  
427 SFTSV may also spread to other countries with competent hosts and vectors. *E. europaeus*  
428 hedgehogs were introduced to New Zealand by human intervention(Brockie, 1975; ISAAC, 2005).  
429 The summer density of hedgehogs in three studies in New Zealand was estimated at between  
430 250 hedgehogs/km<sup>2</sup>(Brockie, 1957; Parkes, 1975) and 800 hedgehogs/km<sup>2</sup>(Campbell, 1973). In  
431 addition *H. longicornis* ticks are common in New Zealand and are all parthenogenetic (Figure  
432 S2)(Heath, 2016). New Zealand is also on the East Asian-Australian flyway. Therefore, New  
433 Zealand might be considered to have a high risk of SFTSV disease incursion, likely through  
434 SFTSV positive *H. longicornis* ticks infested in migratory birds.



435

436 **Figure S2.**(A) The distribution and relative abundance of hedgehogs (*Erinaceus europaeus* L.) in  
437 New Zealand modified from (Brockie, 1975). In the dark grey areas hedgehogs are numerous, in  
438 the light grey areas they are few and in the white areas they are rare or absent. (B) The  
439 distribution of *Haemaphysalis longicornis* in New Zealand modified from (Heath, 2016). The dark  
440 grey areas are high risk, the light grey areas are low risk, and the white areas are zero risk of *H.*  
441 *longicornis* Infestation. (Reproduced with permission of Elsevier and Copyright Clearance  
442 Center).  
443

444 In conclusion, our data strongly support our initial hypothesis that hedgehogs can maintain the  
445 natural circulation of SFTSV in rural areas. The high density and wide distribution, the high-level  
446 susceptibility and tolerance of hedgehogs to SFTSV, the heavy *H. longicornis* infestation rates  
447 and the ability to amplify the infection level of feeding ticks are all compelling evidence that  
448 hedgehogs are the major wildlife amplifying host of SFTSV. Furthermore, evidence that SFTV is  
449 already circulating in ticks and hedgehogs from one urban area of Beijing means that an urban  
450 epidemic of SFTS could happen quite soon. More research and surveillance are needed to  
451 reduce SFTSV risks in both rural and urban areas.

452

## 453 Materials and Methods

### 454 Ethics statement

455 All animal studies were carried out in strict accordance with the recommendations in the Guide for  
456 the Care and Use of Laboratory Animals of the Ministry of Science and Technology of the  
457 People's Republic of China. The protocols for animal studies were approved by the Committee on  
458 the Ethics of Animal Experiments of the Institute of Zoology, Chinese Academy of Sciences  
459 (Approval number: IOZ20180058).

### 460 Animal trapping and sample collection

461 Animal sampling took place in Daishan City (121°30'-123°25'E, 29°32'~31°04'N), Zhejiang  
462 Province, Weifang City (118°10'-120°01'E, 35°32'-37°26'N), Shandong Province, Xinyang City  
463 (113°45'-115°55'E, 30°23'-32°27' N), Henan Province, Linfen City (110°22'-112°34'E, 35°23'-  
464 36°57'N) ,Shanxi Province, and Beijing City (39°26'-41°03'E, 115°25'-117°30'N), China. The  
465 animals were captured using rodent capture cages (cage size: 14 x 14 x 26 cm) baited with fried  
466 bread sticks for three nights at each site (trappings varied between 30 and 50 traps/night  
467 depending on the availability of sites in the area). Cages were deposited into fields and collected  
468 the next morning (FERNANDO TORRES-PÉREZ, 2004). Animals were anesthetized by  
469 inhalation using Isoflurane with a dose of 1 mL per kilogram weight in a closed container. Blood  
470 samples were drawn from heart, and animals were released after blood collection. Blood samples  
471 were centrifuged at 3000 g for 10 minutes and the serum was transferred to small vials, which  
472 were kept at -80°C until analysis.

### 473 Virus and cells

474 SFTSV Wuhan strain (GenBank accession numbers: S, KU361341.1; M, KU361342.1; L,  
475 KU361343.1) and rabbit anti-SFTSV-NP polyclonal antibody were provided by Wuhan Institute of  
476 Virology, Chinese Academy of Sciences. Vero cells (African green monkey kidney epithelial cells)  
477 were obtained from American Type Culture Collection (ATCC) and maintained in Dulbecco's  
478 modified Eagle's medium (DMEM, Hyclone, US) supplemented with 8% FBS and penicillin (100 U  
479 mL<sup>-1</sup>), streptomycin (100 µg mL<sup>-1</sup>; Gibco) and L-glutamine in a 37°C incubator supplemented with  
480 5% CO<sub>2</sub>. SFTSV was propagated at 37°C in Vero cells at a multiplicity of infection of 0.1 and  
481 cultivated for 4 d. Cell culture supernatant was collected at 4 dpi and stored at -80°C as the  
482 working virus stock for animal studies.

### 483 Virus titration

484 Focus-forming assay was performed in Vero cells to titrate the viral titers. Cells were seeded in  
485 96-well plates at 10<sup>4</sup> cells/well in triplicates 24 h before infection. The virus samples were diluted  
486 10-fold in DMEM with 2% FBS. After the removal of culture media, a diluted viral solution was  
487 added to the cells. Three hours later, the cells were washed once and incubated with DMEM plus  
488 2% FBS and 20mM NH<sub>4</sub>Cl at 37°C. 2 d post-infection, the cells were fixed with cold methanol and  
489 stained using a rabbit anti-SFTSV-NP polyclonal antibody at 1:700 dilution and Alexa 488-labeled  
490 goat anti-rabbit IgG at 1:700 dilution. Viral titers were examined under a fluorescent microscope  
491 and calculated by Reed–Muensch method.

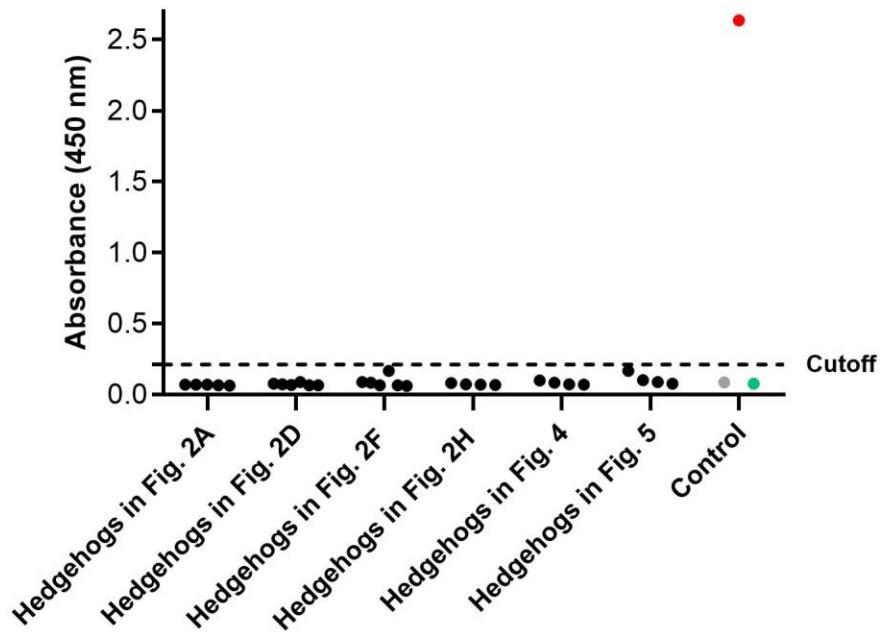
### 492 ELISA for SFTSV antibody detection.

493 Serum samples from animals were tested for SFTSV antibodies including IgG and IgM with a  
494 commercial double antigen sandwich ELISA kit from Nanjing Immune-detect Bio-tech Co., Ltd.  
495 (Jiangsu, China).

496

## 496 Experimental Infection

497 All experimental infection study was conducted in a Bio-safety Level-II Animal Laboratory in the  
498 Beijing Institute of Microbiology and Epidemiology, Academy of Military Medical Sciences. Six to  
499 twelve months old male and female (1:1) African pygmy hedgehogs were purchased from  
500 Longchong Pet in Beijing. Six to twelve months old male and female (1:1) Amur hedgehogs were  
501 purchased from Heze animal store in Shandong Province. All animals were tested for SFTSV  
502 seroprevalence by ELISA before experiments (Figure S3). Following acclimation, hedgehogs  
503 were challenged with  $4 \times 10^6$  FFU of SFTSV Wuhan strain via i.p. injection or i.s. injection, with  
504 the 200  $\mu$ l volume divided between two injection sites. Bodyweight and clinical symptoms were  
505 monitored. Hedgehogs were assigned a clinical score of increasing severity: 1, unfeeding; 2,  
506 hunched posture; 3. green faeces; 4, moribund. Hedgehogs with a score of 3 or a weight loss of  
507 more than 25% were humanely euthanized.



508

509 **Fig. S3** Seroprevalence of SFTSV in all hedgehogs prior to experimental infection. The serum  
510 samples were tested for SFTSV IgG and IgM antibodies prior to experimental infection with a  
511 commercial double antigen sandwich ELISA kit. Cutoff =  $0.748 \times \text{Negative OD} + 0.146$ . Red dot  
512 indicates positive sample, green dot indicates negative sample and gray dot indicates blank  
513 control.

514

515 For the virus transmission studies between host and vector, naïve ticks were fed on African  
516 pygmy hedgehogs which were i.p. inoculated with  $4 \times 10^6$  FFU of SFTSV Wuhan strain at d 0,  
517 and the ticks collected when they naturally detached. And for the transmission studies between  
518 vector and host naïve African pygmy hedgehogs were bitten by infected ticks. Bodyweight and  
519 clinical symptoms in bitten hedgehogs were monitored.

520 For the hibernation experiment, Hedgehogs were challenged with  $4 \times 10^6$  FFU of SFTSV Wuhan  
521 strain and then spent half month or one month in 4°C refrigerator. Two of the hedgehogs were  
522 waken up at d 15 and another two were waken up at d 30. The blood samples were taken from  
523 the heart for viral RNA detection.

#### 524 **SFTSV-RNA extraction and real-time RT-PCR**

525 Total RNA prepared from the homogenates of the ticks and the blood samples collected from  
526 hedgehogs' heart were extracted using TRIzol reagent (Thermo Fisher Scientific, USA) or the

527 RNeasy kit (Qiagen, Germany) according to the manufacturer's instructions. Samples were  
528 analyzed using a One-Step SYBR PrimerScript reverse transcription (RT)-PCR kit (TaKaRa,  
529 Japan) on Applied Biosystems QuantStudio. Each sample was measured by triplicate. The  
530 primers were designed as previously described(Dong et al., 2019). Conditions for the reaction  
531 were as follows: 42°C for 5 min, 95°C for 10 sec, 40 cycles at 95°C for 5 sec, and 60°C for 20  
532 sec.

### 533 **Pathological lesions in SFTSV-Infected hedgehogs**

534 Six African pygmy hedgehogs were inoculated with  $4 \times 10^6$  FFU of the Wuhan strain of SFTSV by  
535 i.p. route and two were killed at each time point of d 3, 6 and months 2 post infection for analysis  
536 of viremia and pathology. Two mock-infected control hedgehogs were killed at d 0. For  
537 histopathological evaluation, spleens were rapidly removed, fixed in 4% PFA at room temperature  
538 for 7 d, and routinely processed for paraffin embedding. Coronal, 4–5  $\mu$ m thick serial sections  
539 were performed and selected sections were stained with H&E for light microscopy examination.  
540 Images were obtained using a Nikon Eclipse 50i Light Microscope (Nikon, Tokyo, Japan) or  
541 Olympus BX60 microscope (Shinjuku, Tokyo, Japan)

### 542 **Identification of tick species and phylogenetic analysis**

543 Ticks were identified based on morphological characteristics, visualized through a light  
544 microscope, with further molecular confirmation in the laboratory by sequencing the mitochondrial  
545 16S ribosomal RNA (16S rRNA) gene. The primers were as follows: (16S-1)  
546 CTGCTCAATGATTTTTAAATTGCTGTGG (Forward primer) and (16S-2)  
547 CGCTGTTATCCCTAGAGTATT (Reverse primer). A single leg was removed from each tick for  
548 the molecular analysis to confirm identification. Phylogenetic analysis was performed using the  
549 whole mitochondrial genomes. Tick DNA was extracted using the MightyPrep reagent for DNA Kit  
550 (Takara, Japan) according to the manufacturer's instructions. The mitochondrial DNA were  
551 sequenced by next generation sequencing (Tsingke Biotech, Beijing, China) and deposited in  
552 GenBank (Parthenogenetic *H. longicornis* in Shunyi District: OL335941; Chaoyang District:  
553 OL335942). The phylogenetic tree was constructed using the maximum likelihood method,  
554 MEGA-X with the bootstrap value set at 1000.

### 555 **SFTSV sequencing and phylogenetics analysis**

556 The SFTSV sequences from Shunyi *H. longicornis* ticks were amplified using (L4010-  
557 F:GGAACCTCTAGCCACTCTGTT; L4963-R:GTAGAGAAGGCCTATGATC) and sequenced  
558 by Tsingke Biotech, Beijing, China, and then deposited in the GenBank (Accession number:  
559 OL518989). SFTSV sequences in the phylogenetics analysis were downloaded from  
560 GenBank(Shi et al., 2017; Yoshikawa et al., 2015). The phylogenetic trees were constructed  
561 using the maximum likelihood method, MEGA program. The confidence of the tree was tested  
562 using 1000 bootstrap replications.

563

### 564 **Acknowledgments**

565 We gratefully thank the following funders: the State Key Research Development Program of  
566 China (2019YFC12005004, 2019YFC12005001), the Strategic Priority Research Program of  
567 Chinese Academy of Sciences (Grant No. XDPB16), the key program of Chinese Academy of  
568 Sciences (CAS) (KJZD-SW-L11), Natural Science Foundation of Zhejiang Province (NO.  
569 LY21H100002), and the Open Research Fund Program of State Key Laboratory of Integrated  
570 Pest Management (IPM2109). We also thank Rachel Summers, Massey University for  
571 constructing the New Zealand hedgehog distribution map.

572

### 573 **Competing interests**

574 The authors declare no competing interests.

575

### 576 **References**

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## Supplementary Information for

### Hedgehogs are the major amplifying hosts of severe fever with thrombocytopenia syndrome virus

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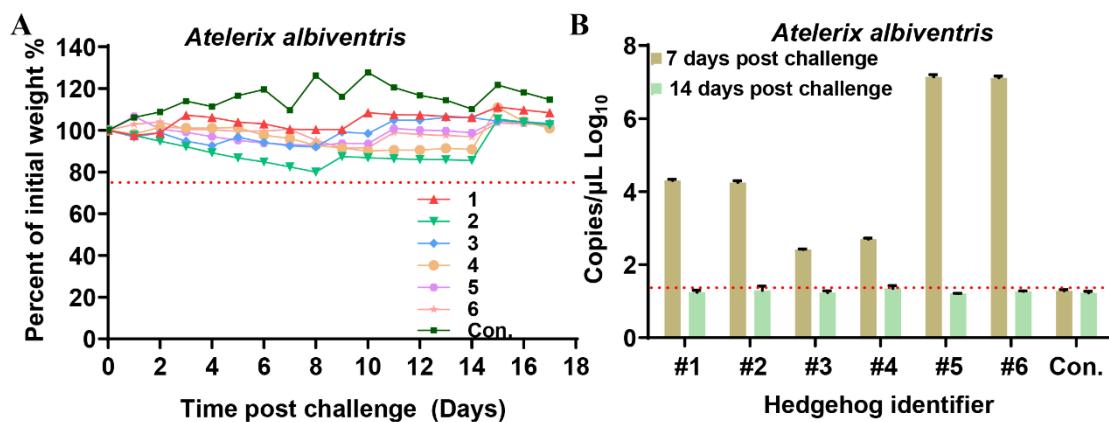
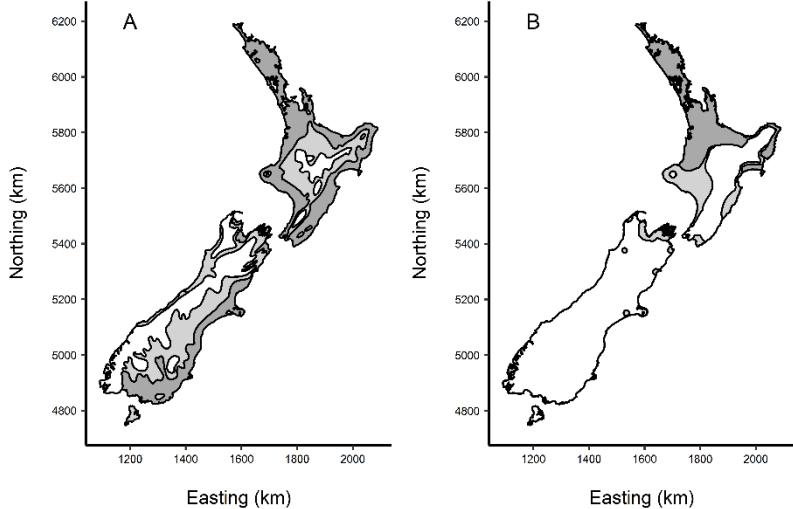


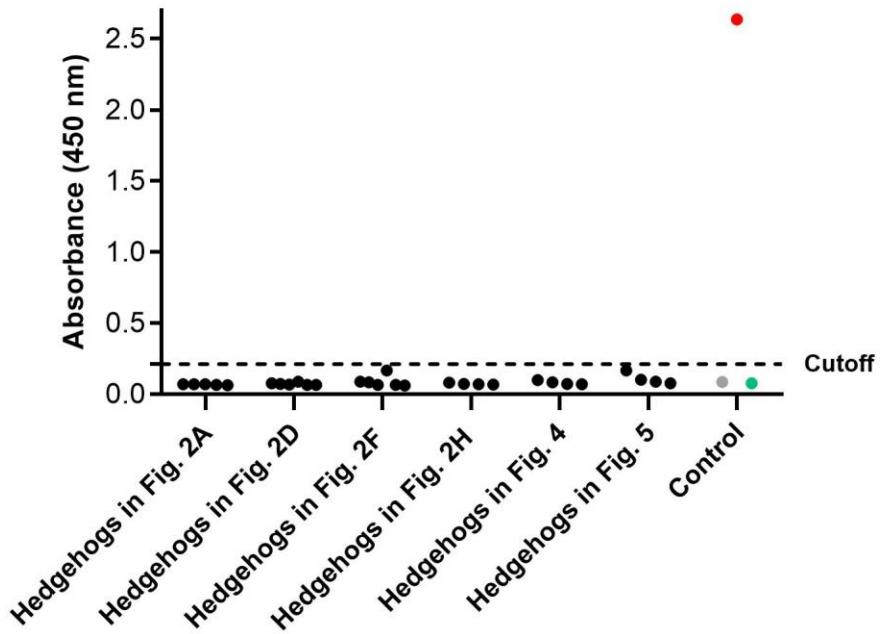
Figure S1.

SFTSV viremia and weight change in experimentally infected *A. albiventris* hedgehogs bled twice. Hedgehogs were challenged (i.p.) with  $4 \times 10^6$  FFU of SFTSV Wuhan strain and then monitored for weight loss everyday (A) and viremia on d 7 and 14, measured by Real-time PCR as RNA copies per  $\mu\text{L}$  of serum (B) (error bars represent SD). Each line and bar indicate one hedgehog.



**Figure S2.**

(A) The distribution and relative abundance of hedgehogs (*Erinaceus europaeus* L.) in New Zealand modified from (Brockie, 1975). In the dark grey areas hedgehogs are numerous, in the light grey areas they are few and in the white areas they are rare or absent. (B) The distribution of *Haemaphysalis longicornis* in New Zealand modified from (Heath, 2016). The dark grey areas are high risk, the light grey areas are low risk, and the white areas are zero risk of *H. longicornis* Infestation. (Reproduced with permission of Elsevier and Copyright Clearance Center).



**Fig. S3**

Seroprevalence of SFTSV in all hedgehogs prior to experimental infection. The serum samples were tested for SFTSV IgG and IgM antibodies prior to experimental infection with a commercial double antigen sandwich ELISA kit. Cutoff =  $0.748 \times \text{Negative OD} + 0.146$ . Red dot indicates positive sample, green dot indicates negative sample and gray dot indicates blank control.

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