

- behavior among US adults. *Cogent Soc Sci.* 2017;3:1302785. <https://doi.org/10.1080/23311886.2017.1302785>
3. Piltch-Loeb R, Merdjanoff AA, Abramson DM. How the US population engaged with and prioritized sources of information about the emerging Zika virus in 2016. *Health Secur.* 2018;16:165–77. <https://doi.org/10.1089/hs.2017.0107>
 4. Sell TK, Watson C, Meyer D, Kronk M, Ravi S, Pechta LE, et al. Frequency of risk-related news media messages in 2016 coverage of Zika virus. *Risk Anal.* 2018;38:2514–24. <https://doi.org/10.1111/risa.12961>
 5. Wolfe JH. Pattern clustering by multivariate mixture analysis. *Multivariate Behav Res.* 1970;5:329–50. https://doi.org/10.1207/s15327906mbr0503_6
 6. Goodman LA. The analysis of systems of qualitative variables when some of the variables are unobservable. Part IA modified latent structure approach. *The American Journal of Sociology.* 1974;79:1179–259. <https://doi.org/10.1086/225676>
 7. Nylund KL, Asparouhov T, Muthén BO. Deciding on the number of classes in latent class analysis and growth mixture modeling: a Monte Carlo simulation study. *Struct Equ Modeling.* 2007;14:535–69. <https://doi.org/10.1080/10705510701575396>
 8. Agarwal NK. Information source and its relationship with the context of information seeking behavior. In: *iConference 2011: proceedings of the 2011 iConference*; 2011 Feb 8–11; Seattle, Washington, USA. New York: Association for Computing Machinery; 2011. p. 48–55.

Address for correspondence: Rachael Piltch-Loeb, New York University, Global Public Health, 715 Broadway Rm 1229, New York, NY 10003, USA; email: rpl5@nyu.edu

Severe Fever with Thrombocytopenia Syndrome Virus in Ticks and SFTS Incidence in Humans, South Korea

Jeong Rae Yoo,¹ Sang Taek Heo,¹ Sung Wook Song, Seung Geon Bae, Seul Lee, Sungho Choi, Chaehyun Lee, Sugyeong Jeong, Myeongseop Kim, Woojin Sa, Yeongrim Lee, Haseon Choi, Sun-Ho Kee, Keun Hwa Lee

DOI: <https://doi.org/10.3201/eid2609.200065>

¹These authors contributed equally to this article.

Author affiliations: Jeju National University, Jeju, South Korea (J.R. Yoo, S.T. Heo, S.W. Song, S.G. Bae, S. Lee, S. Choi, C. Lee, S. Jeong, M. Kim, W. Sa, Y. Lee, H. Choi); Korea University, Seoul, South Korea (S.-H. Kee); Hanyang University College of Medicine, Seoul (K.H. Lee).

During 2016–2018, we collected 3,193 ticks from rural areas in South Korea to investigate the prevalence of severe fever with thrombocytopenia syndrome virus (SFTSV). We detected SFTSV in ticks at an infection rate (IR) of 11.1%. We noted increases in the human IR associated with the monthly SFTSV IR in ticks.

Severe fever with thrombocytopenia syndrome (SFTS) is a tickborne zoonosis caused by the SFTS virus (SFTSV) (1); >1,000 SFTS cases have been reported in South Korea (2). The SFTS prevalence rate was 2.26/100,000 inhabitants on the mainland and 13.66/100,000 inhabitants on Jeju Island, South Korea (2). SFTSV has been detected in several species of ticks, including *Haemaphysalis longicornis*, *Amblyomma testudinarium*, and *Ixodes nipponensis* (3). A previous study reported that the minimum infection rate of SFTSV in infected ticks was lower (0.37%) on Jeju Island than in other collection areas (1.97%) (4). However, 7%–14% SFTSV seropositivity was identified in domestic and wild animals (4–5), and 2%–5% SFTSV seropositivity was identified in a healthy population in South Korea (6). Therefore, our aim was to investigate the SFTSV infection rate (IR) in ticks in the region with the highest endemicity, Jeju Island, and to analyze the relationship between the geographic distribution of ticks and SFTSV and human cases of SFTS.

During June 2016–January 2019, well-equipped trained researchers collected ticks from the natural environment of Jeju Island. The tick sampling sites included 5 rural areas: Aewol-eup (AW); Seon Hul-ri (SH); Jeo Ji-ri (JJ); and Ha Do-ri (HD) and Bo Mok-ri (BM) (Figure). These 5 areas were chosen to compare SFTSV IR in ticks in areas with the highest rates of human SFTS cases, SH, HD, and AW, and SFTSV IR in ticks in areas with lower human SFTS rates, JJ and BM. Ticks were manually collected 2 times per month, during the first and third weeks, by dragging a white cloth in woodlands for 2 hours in each area. We morphologically identified tick species and developmental stages by using an Olympus SD-ILK-200–2 stereomicroscope (Olympus Corporation, <https://www.olympus-lifescience.com>) (7) and extracted viral RNA by using a QIAamp Viral RNA Mini kit (QIAGEN Inc., <https://www.qiagen.com>) according to the manufacturer's instructions (Appendix, <https://wwwnc.cdc.gov/EID/article/26/9/20-0065-App1.pdf>).

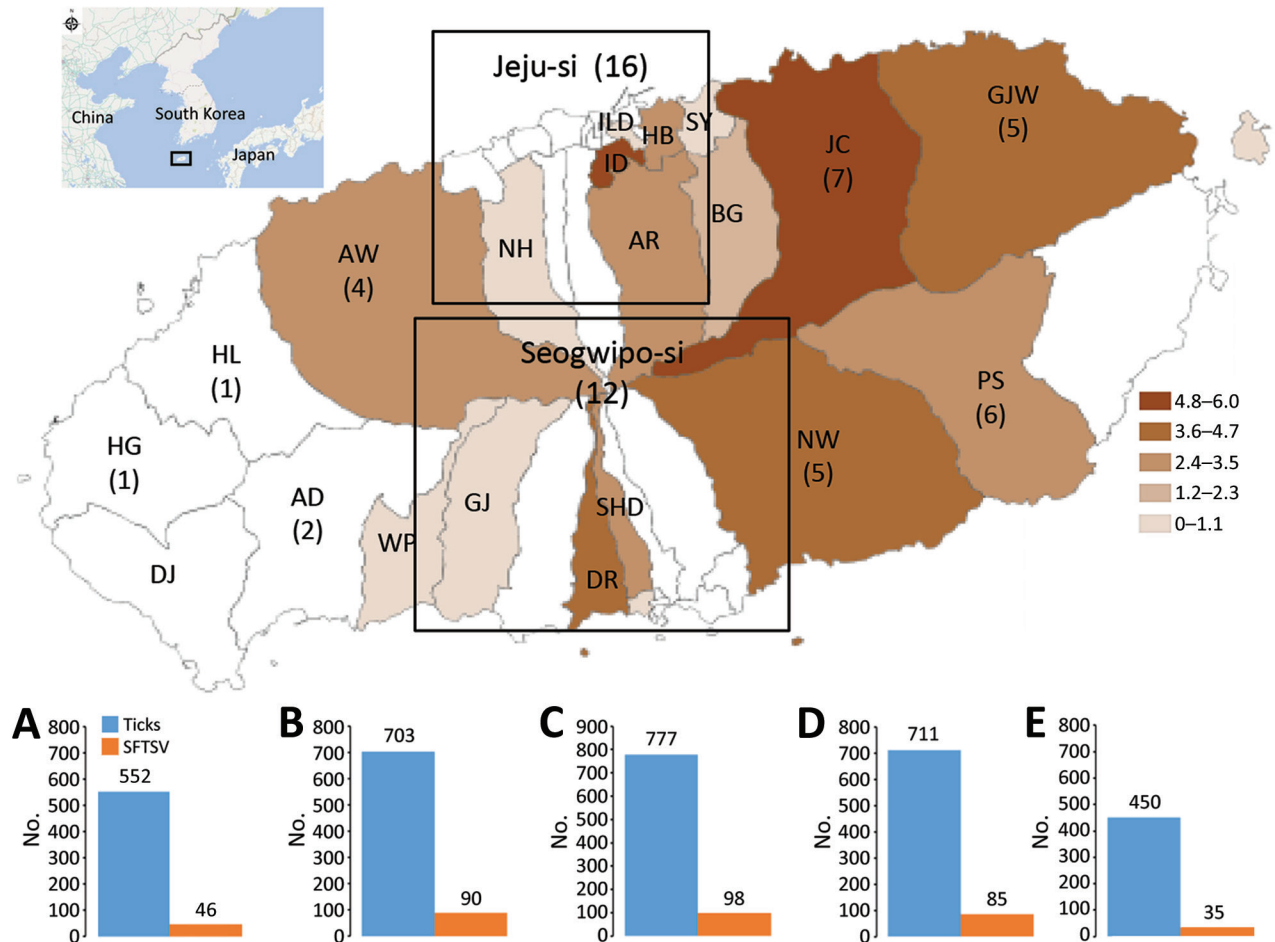


Figure. Geographic distribution of patients with severe fever with thrombocytopenia syndrome (SFTS) during January 2013–January 2019 and yearly incidence rates of SFTS virus (SFTSV) from June 2016–January 2019 on Jeju Island, South Korea. Inset shows location of Jeju Island near the coast of South Korea. Orange indicate regions of patients with SFTS in 2013–2019. Graphs show SFTSV detected in ticks and cases of human SFTS in A) Jeo Ji-ri; B) Aewol-eup; C) Seon Hul-ri; D) Ha Do-ri; and E) Bo Mok-ri. AR, Ara-dong; AW, Aewol-eup; BG, Bonggae-dong; BM, Bo Mok-ri; DR, Daeryun-dong; GJ, Ganjeong-dong; GJW, Gujwa-eup; HB, Hwabug-dong; HD, Ha Do-ri; ILD, Ildo-dong; ID, Ido-dong; JC, Jocheon-eup; JJ, Jeo Ji-ri; NH, Nohyeong-dong; NW, Nanwon-eup; PS, Pyoseon-myeon; SHO, Seohong-dong; SH, Seon Hul-ri; SY, Samyang-dong; WP, Wolpyeong-dong.

A total of 3,193 ticks were collected; most (99.9%) were *H. longicornis* and 81.3% of all ticks were nymphs. We detected SFTSV in 11.1% (354/3,193) of ticks (Appendix Table). Among the 5 areas, the average IR of SFTSV in ticks was 12.8% in AW, 12.6% in SH, 12.0% in HD, 8.3% in JJ, and 7.8% in BM. Adult ticks had a higher SFTSV IR (12.4%) than nymphs (9.5%). SFTSV was detected mainly in adult ticks (Appendix Table).

The monthly IR of SFTSV in ticks increased in May, peaked in July, and then slowly decreased (Appendix Figure 1). In addition, changes in the incidence of SFTS in patients were associated with increases in the monthly IR of SFTSV in ticks at a rate of 19.8% (95% CI 2.3%–40.2%) increase of SFTS in patients per 1% increase in monthly SFTSV IR in ticks ($p = 0.02$). SFTSV-infected ticks also were observed in the winter

season. SFTSV sequences from infected ticks in our study and SFTS patients on Jeju Island were consistent with each other but differed from viruses in other regions of South Korea (Appendix Figure 2).

In a previous study, the prevalence of SFTSV in ticks was very low (0.2%), implying that ticks alone might not be sufficient to maintain SFTSV in nature (8). However, the high IR of SFTSV in ticks could explain why Jeju Island had one of the highest rates of human SFTS infection in South Korea. Changes in the incidence of patients with SFTS showed a pattern similar to that of monthly SFTSV IR in ticks. The IR of SFTSV in SH was the highest, along with HD and AW, where rates of human SFTS cases also were high.

In East Asia, humans most frequently acquire SFTS during May–July, and shrub, forest, and rainfed

cropland areas are associated with high risk for infection(2,3,6). Northeast Jeju Island, which includes SH, AW, and HD, has many farms and wetlands, and the IR of SFTSV in ticks peaked there in July, August, and September. In addition, SFTSV was detected in ticks in winter on Jeju Island, but no SFTS cases were reported in South Korea during winter. The 62 confirmed SFTS cases were statistically significantly associated with higher ambient temperature ($22.5^{\circ}\text{C} \pm 4.2^{\circ}\text{C}$) compared with patients with negative RT-PCR results for SFTSV ($18.9^{\circ}\text{C} \pm 5.7^{\circ}\text{C}$; $p < 0.001$) (J.R. Yoo, unpub. data). The optimal temperature range for growth and reproduction of *H. longicornis* ticks is 20° – 24° . Jeju Island maintains a temperature $>20^{\circ}$ during May–October and is largely a rural and natural environment. We consider this area to have the highest prevalence of SFTS cases and ticks with SFTSV in South Korea.

The results of this study showed that Jeju Island has the highest IR of SFTSV in ticks compared with other regions of South Korea and endemic countries. In addition, we found that the partial small segment of SFTSV in ticks was highly homologous to SFTSV in patients on Jeju Island and that Northeast Jeju Island, which includes SH, is a high-risk area for human SFTS infections.

Acknowledgments

We thank the members of department of microbiology, Jeju National University, Graduate College of Medicine and the Safety for Agricultural Injuries of Famers cohort study, Jeju National University Hospital for assistance in collecting tick samples in various regions of Jeju Island, South Korea.

This work was supported by a grant from the Korea Health Technology Research and Development Project through the Korea Health Industry Development Institute and funded by the Ministry of Health & Welfare, South Korea (grant no. HG18C0037).

About the Author

Dr. Yoo and Dr. Heo are professors in the Department of Internal Medicine, Jeju National University, College of Medicine, Jeju, South Korea. Their research interests are infectious diseases.

References

1. Yun SM, Lee WG, Ryou J, Yang SC, Park SW, Roh JY, et al. Severe fever with thrombocytopenia syndrome virus in ticks collected from humans, South Korea, 2013. *Emerg Infect Dis*. 2014;20:1358–61. <https://doi.org/10.3201/eid2008.131857>
2. Korea Centers for Disease Control and Prevention. Infectious disease surveillance 2019, public health weekly report [in Korean] [cited 2020 May 1]. <https://is.cdc.go.kr>
3. Im JH, Baek J, Durey A, Kwon HY, Chung MH, Lee JS. Current status of tick-borne diseases in South Korea. *Vector Borne Zoonotic Dis*. 2019;19:225–33. <https://doi.org/10.1089/vbz.2018.2298>
4. Oh SS, Chae JB, Kang JG, Kim HC, Chong ST, Shin JH, et al. Detection of severe fever with thrombocytopenia syndrome virus from wild animals and Ixodidae ticks in the Republic of Korea. *Vector Borne Zoonotic Dis*. 2016;16:408–14. <https://doi.org/10.1089/vbz.2015.1848>
5. Yu KM, Yu MA, Park SJ, Kim YI, Robles NJ, Kwon HI, et al. Seroprevalence and genetic characterization of severe fever with thrombocytopenia syndrome virus in domestic goats in South Korea. *Ticks Tick Borne Dis*. 2018;9:1202–6. <https://doi.org/10.1016/j.ttbdis.2018.05.001>
6. Yoo JR, Heo ST, Kim M, Song SW, Boo JW, Lee KH. Seroprevalence of severe fever with thrombocytopenia syndrome in the agricultural population of Jeju Island, Korea, 2015–2017. *Infect Chemother*. 2019;51:337–44. <https://doi.org/10.3947/ic.2019.51.4.337>
7. Goff ML, Loomis RB, Welbourn WC, Wrenn WJ. A glossary of chigger terminology (Acari: Trombiculidae). *J Med Entomol*. 1982;19:221–38. <https://doi.org/10.1093/jmedent/19.3.221>
8. Luo LM, Zhao L, Wen HL, Zhang ZT, Liu JW, Fang LZ, et al. *Haemaphysalis longicornis* ticks as Reservoir and Vector of Severe Fever with Thrombocytopenia Syndrome Virus in China. *Emerg Infect Dis*. 2015;21:1770–6. <https://doi.org/10.3201/eid2110.150126>

Address for correspondence: Keun Hwa Lee, Department of Microbiology, Hanyang University College of Medicine, 222 Wangsimni-ro Seongdong-gu, Seoul 04763, South Korea; email: yomust7@gmail.com

Typhus Group Rickettsiosis, Brazilian Amazon

Antonio H.H. Minervino, Marcelo B. Labruna, Salatiel R. Dias, Francisco B. Costa, Thiago F. Martins, Phablo N.S. da Silva, Álvaro A. Faccini-Martínez

Author affiliations: Federal University of Western Pará, Santarém, Brazil (A.H.H. Minervino, S.R. Dias, P.N.S. da Silva); University of São Paulo, São Paulo, Brazil (M.B. Labruna, T.F. Martins); Universidade Estadual do Maranhão, São Luís, Brazil (F.B. Costa); Asociación Colombiana de Infectología, Bogotá, Colombia (Á.A. Faccini-Martínez)

DOI: <https://doi.org/10.3201/eid2609.201305>

Severe Fever with Thrombocytopenia Syndrome Virus in Ticks and SFTS Incidence in Humans, South Korea

Appendix

Methods

Severe fever with thrombocytopenia syndrome viral RNA was extracted from ticks collected from Jeju Island, South Korea, by using a QIAamp Viral RNA Mini kit (QIAGEN Inc., <https://www.qiagen.com>) according to the manufacturer's instructions. Real-time reverse transcription PCR (RT-PCR) of the partial small (S) segment of SFTSV was performed for molecular diagnosis (1). The real-time RT-PCR products were sequenced by using a BigDye Terminator Cycle Sequencing kit (Perkin Elmer Applied Biosystems, www.perkin-elmer.com). Phylogenetic analyses of SFTSV partial S segment sequences from ticks and patients were conducted with MEGA6 (2), and phylogenetic trees were constructed using the maximum likelihood method.

Statistical analyses were performed by using SPSS 20.0 (IBM Corp., <https://www.ibm.com>) and $p < 0.05$ was considered statistically significant. We used the annual number of cases of SFTS from the Disease Web Statistics System, Korea Centers for Disease Control and Prevention (<http://is.cdc.go.kr>). Binomial negative regression was used to analyze the monthly infection rates of ticks with SFTSV and SFTS patients on Jeju Island.

References

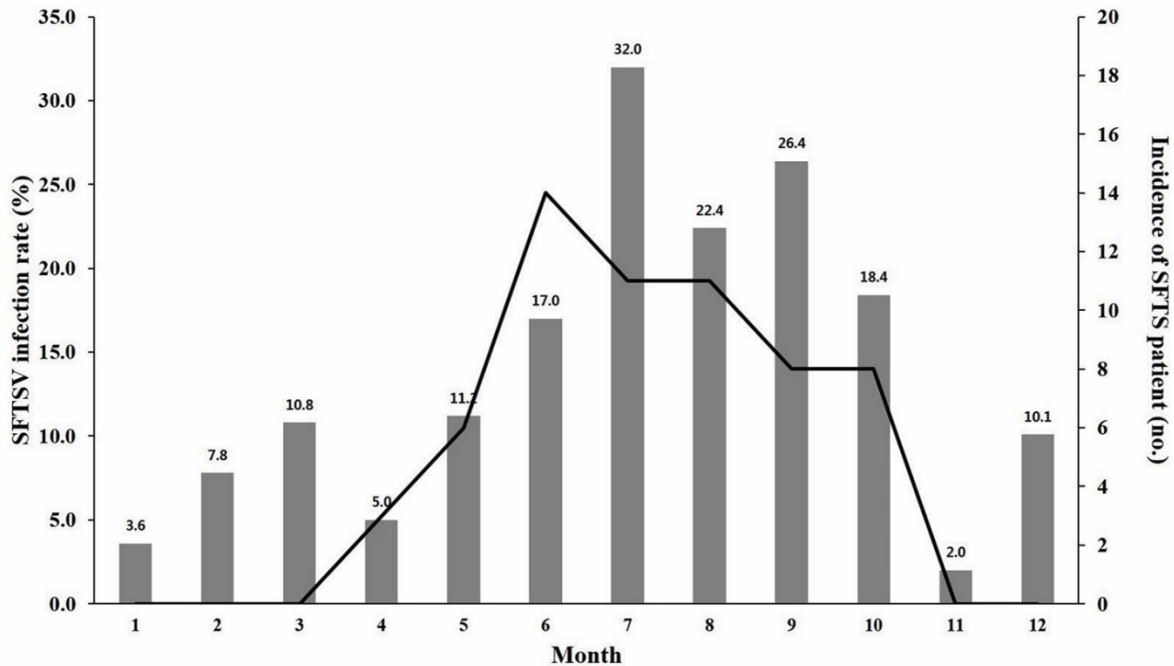
1. Yun Y, Heo ST, Kim G, Hewson R, Kim H, Park D, et al. Phylogenetic analysis of severe fever with thrombocytopenia syndrome virus in South Korea and migratory bird routes between China, South Korea, and Japan. *Am J Trop Med Hyg.* 2015;93:468–74. [PubMed](https://pubmed.ncbi.nlm.nih.gov/260065/)
<https://doi.org/10.4269/ajtmh.15-0047>

2. Tamura K, Stecher G, Peterson D, Filipski A, Kumar S. MEGA6: Molecular Evolutionary Genetics Analysis version 6.0. Mol Biol Evol. 2013;30:2725–9. [PubMed](https://doi.org/10.1093/molbev/mst197)
<https://doi.org/10.1093/molbev/mst197>

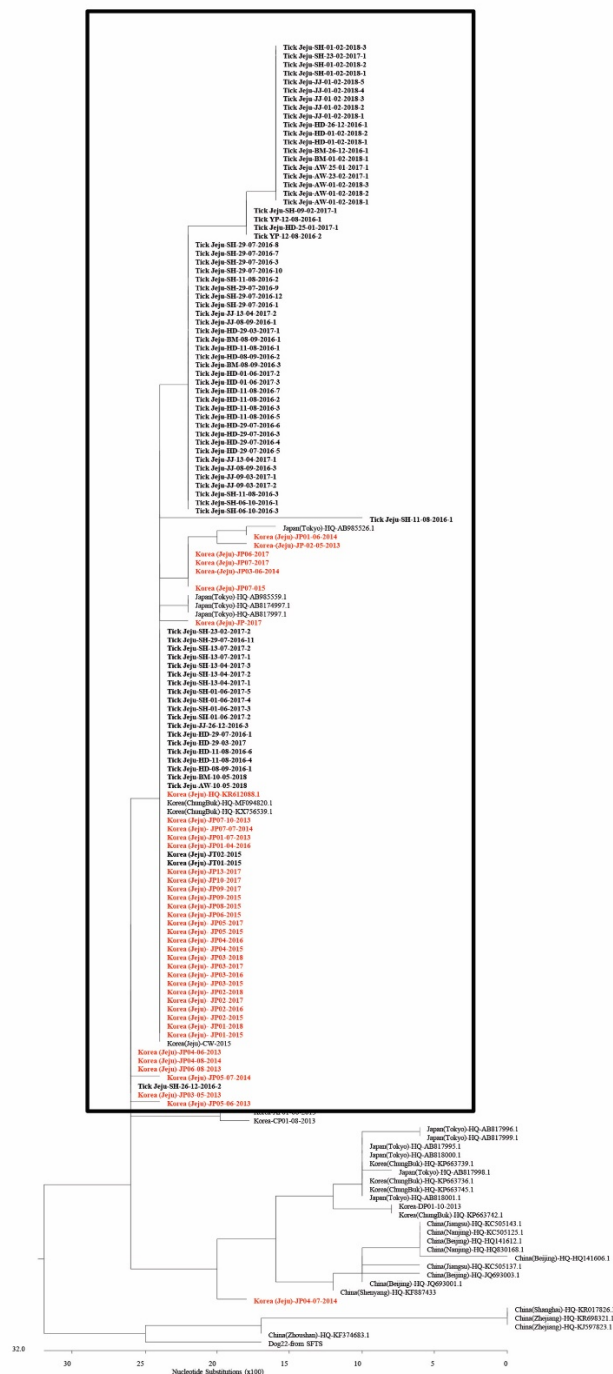
Appendix Table. Severe fever with thrombocytopenia syndrome virus detection in 3,193 ticks by developmental stage and region, Jeju Island, South Korea*

Stage	JJ		AW		HD		BD		SH		Total, %
	No.	SFTSV (IR, %)	No.	SFTSV (IR, %)	No.	SFTSV (IR, %)	No.	SFTSV (IR, %)	No.	SFTSV (IR, %)	
Adult, F	51	3 (5.9)	73	13(17.8)	76	15 (19.7)	55	4 (7.3)	98	15 (15.3)	12.7
Adult, M	25	3 (12.0)	52	7 (13.5)	52	9 (17.3)	32	0	80	13 (16.3)	12.0
Nymph	476	40 (8.4)	578	70 (12.1)	583	61 (10.5)	362	31 (8.6)	598	70 (11.7)	9.5
Larva	0	0	0	0	0	0	1	0	0	0	0
Total	552	46 (8.3)	703	90 (12.8)	711	85 (12.0)	450	35 (7.8)	777	98 (12.6)	11.1

*AW, Aewol-eup; BM, Bo Mok-ri; Ha Do-ri; HD, IR, infection rate of SFTSV in ticks; JJ, Jeo Ji-ri; SH, Seon Hul-ri; SFTSV, severe fever with thrombocytopenia syndrome virus.



Appendix Figure 1. Monthly severe fever with thrombocytopenia syndrome virus infection rate in ticks (2016–2018) and incidence of patients with SFTS (2013–2019) on Jeju Island, South Korea. Black line indicates the monthly incidence of SFTS in patients on Jeju Island (2013–2019). Gray bars indicate the monthly SFTSV infection rate in ticks on Jeju Island (2016–2018). SFTS, severe fever with thrombocytopenia syndrome; SFTSV, severe fever with thrombocytopenia syndrome virus.



Appendix Figure 2. Phylogenetic tree based on partial small (S) segment sequences severe fever with thrombocytopenia syndrome (SFTS) virus. The tree was constructed by using the maximum likelihood method in MEGA 6 (2). The partial S sequences were obtained from ticks during June 2016–January 2019. The partial S sequence data for the viruses identified in China, South Korea, and Japan were obtained from NCBI/BLAST (<https://blast.ncbi.nlm.nih.gov/Blast.cgi>).