



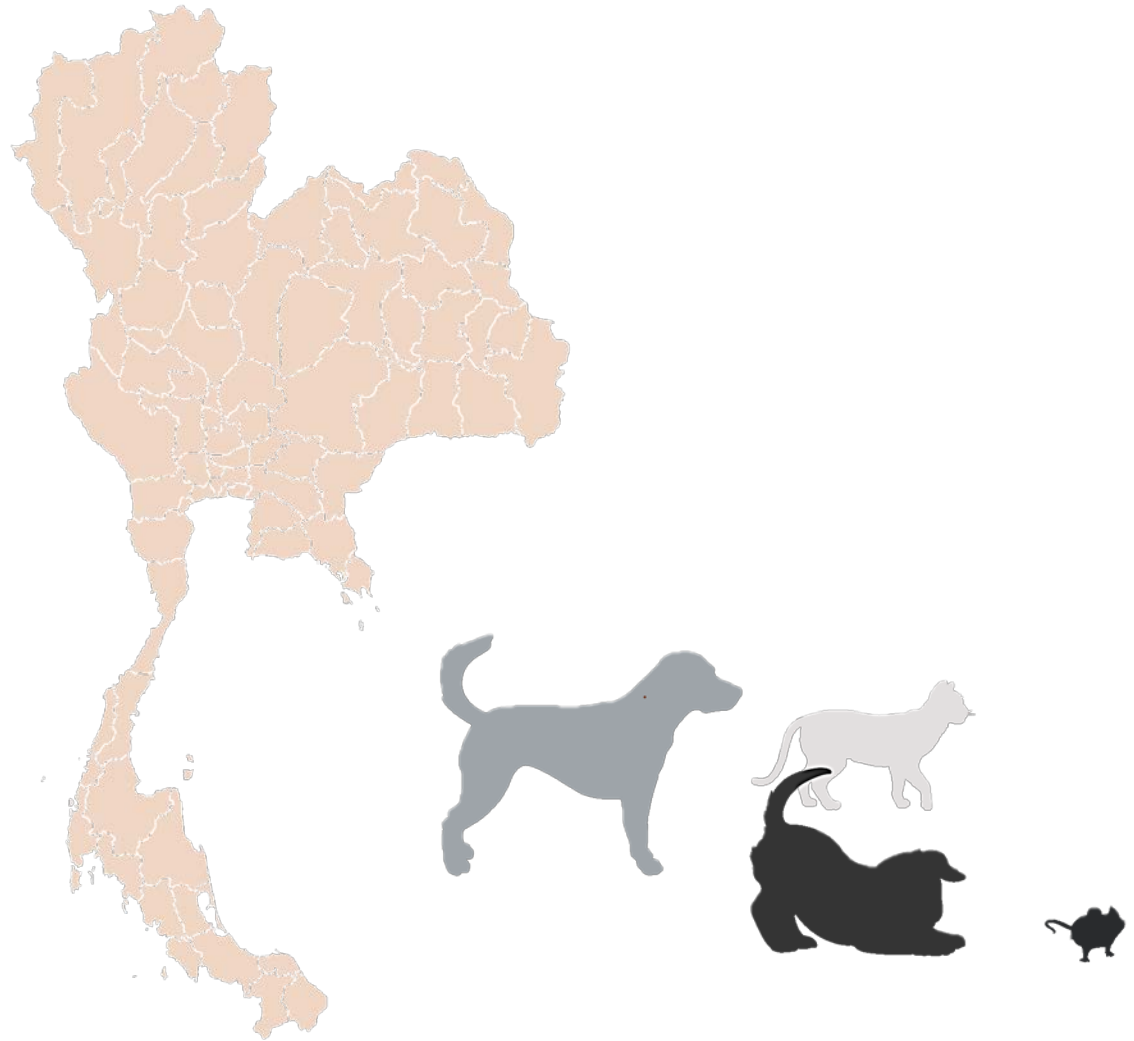
# “Severe Fever with Thrombocytopenia Syndrome (SFTS)”

**Prof. YONG POOVORAWAN, MD.**

ศ.นพ.ยง ภู่วรรณ

Center of Excellence in Clinical Virology, Faculty of Medicine, Chulalongkorn University

# Emerging SFTSV Threat in Thailand: Human Infections, Animal Reservoirs, and Vector Ecology



# Arthropod-borne viruses (Arboviruses)

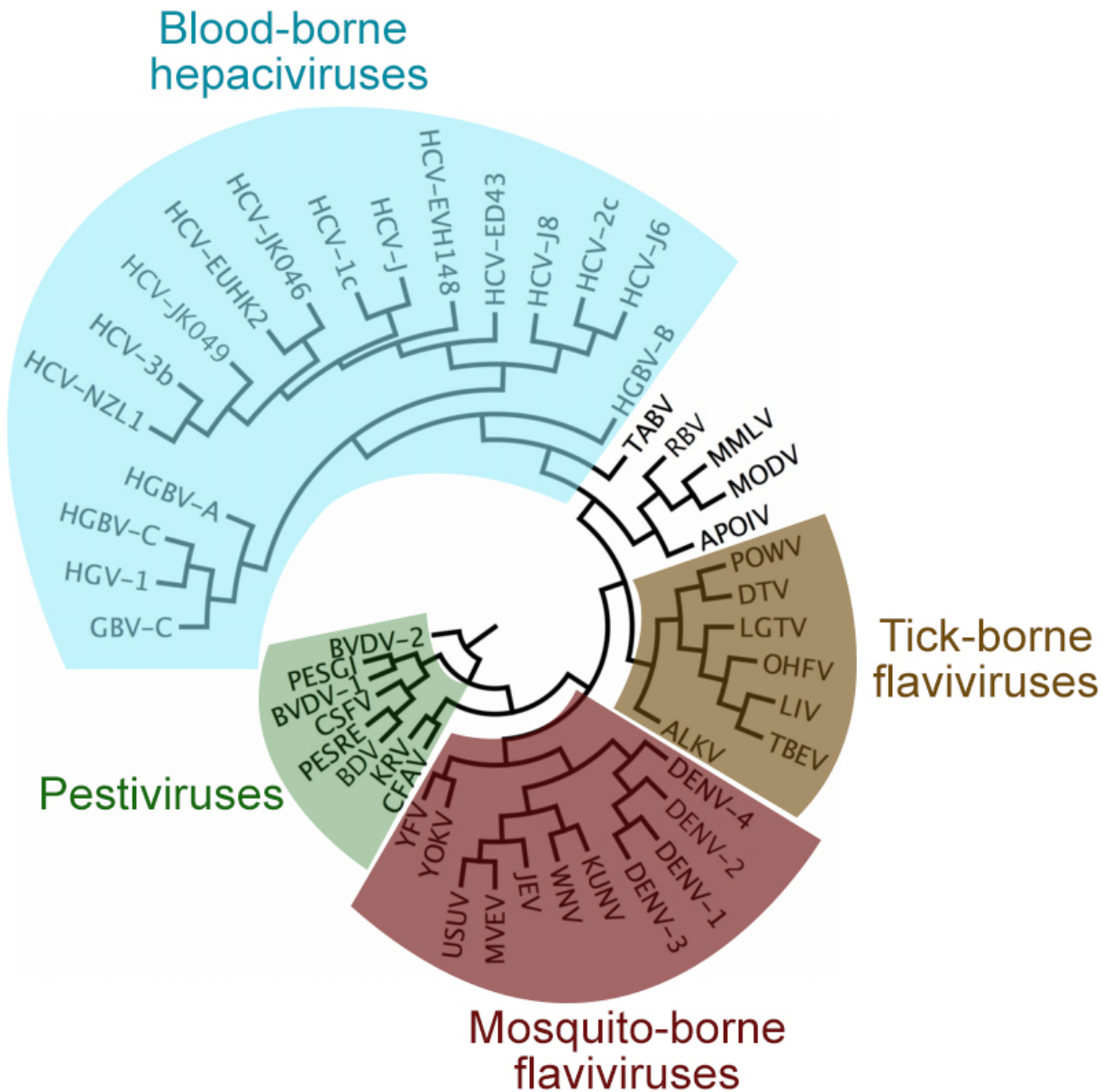
- Existed throughout human history
- Causes of human and animal diseases
- Emergence of arbovirus infections
  - international trade & transportation, migration, climate change, difficulties with treatment, vaccine development, vector control etc.

# Arthropod-borne viruses (Arboviruses)

Human pathogens:

- Dengue Virus
- West Nile virus
- Yellow fever
- Japanese encephalitis virus
- Chikungunya Virus
- Zika virus
- Oropouche virus
- Mayaro virus
- SFTSV (Dabie bandavirus)





# Phylogenetic analysis of Flaviviridae family

<http://www.utmb.edu/discoveringdenguedrugs-together/Diseases/Flaviviridae%20Family.asp>

วัยรุ่นชาย อายุ 16 ปี อยู่อำเภอเมือง (บ้านสวน)

ชลบุรี นักเรียนร่างกายแข็งแรงดี

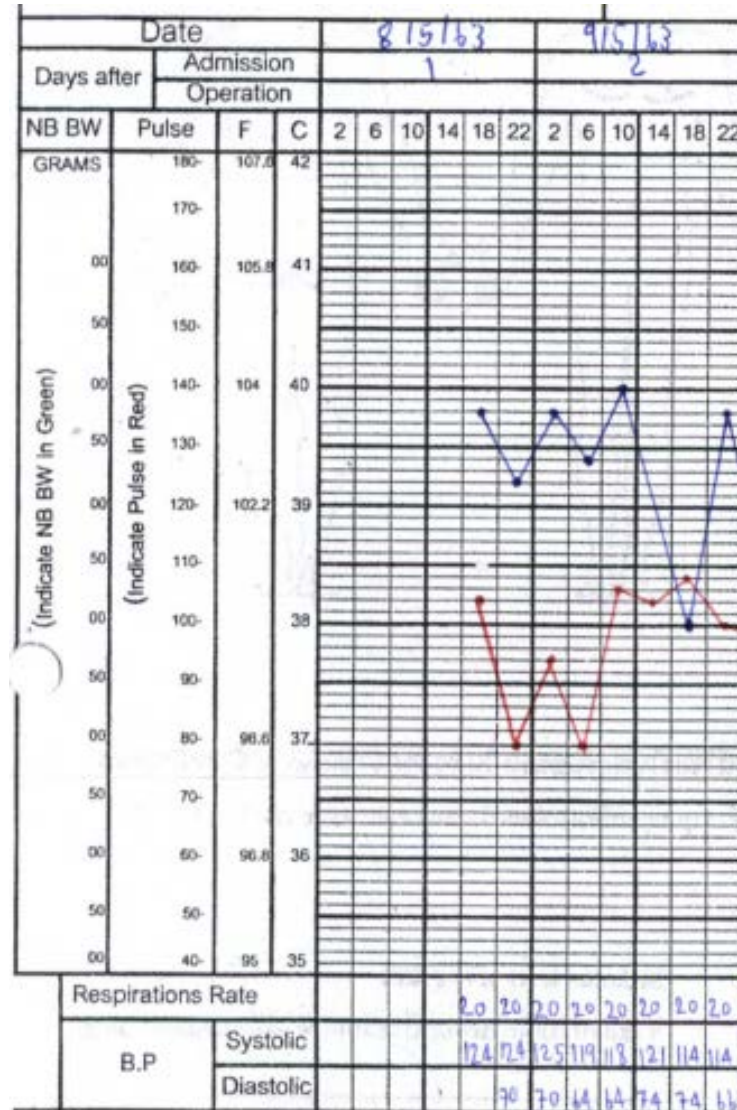
- ไข้สูงมา 3 วัน ตรวจร่างกาย คลำตับได้ อย่างอื่นไม่พบ  
อะไรผิดปกติ
- ตรวจ CBC Hb 14.9 gm%, Hct 44% WBC 2500  
/cumm Platelets 92,000 N71% L18% M8%
- Repeated the day after Platelets 172,000/cumm
- BUN, Cr normal, LFT mild elevation AST, ALT

## Day 4

Date				8/5/63					
Days after		Admission		1					
		Operation							
NB BW	Pulse	F	C	2	6	10	14	18	22
GRAMS  									

# Case 3

Day 4      Day 5



# Repeated CBC



- Hb 14.7 gm Hct 44%
- WBC 2300 cumm
- Platelets 130,000 cumm



# Case 3

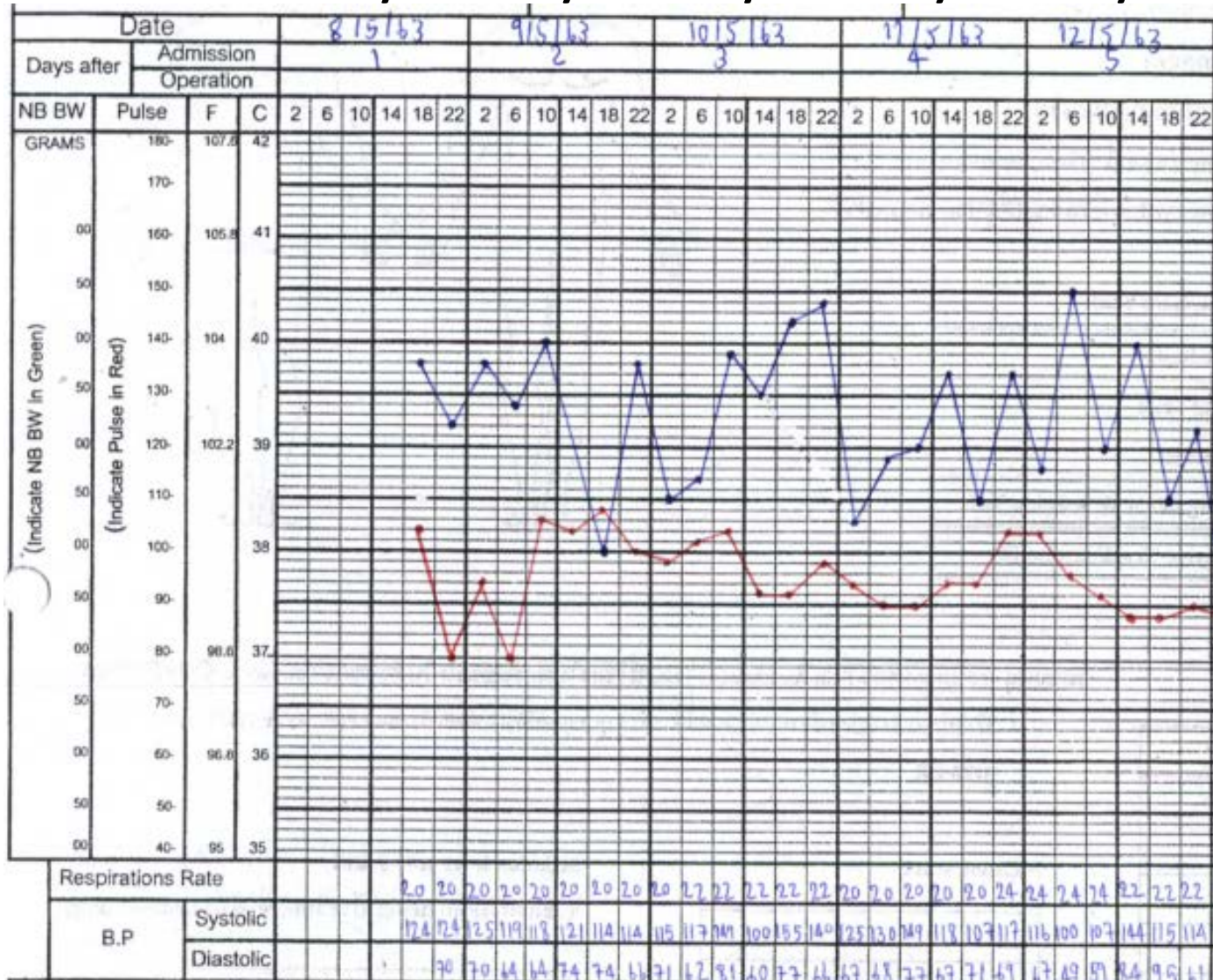
Day 4

Day 5

Day 6

Day 7

Day 8



# Lab investigation



**Flu A/B negative, Covid-19 neg**

**Dengue NS1 neg**

**EBV IgM neg, VZV neg**

# Consult ID



- \* Chikungunya IgG, IgM. neg, RT-PCR neg.  
UA negative
- \* Hemo culture x3D no growth x 2 ၁၁၁
- \* Scrup typhus IgM negative
- \* Stool c/s for enteropathogenic bacteria neg
- \* Ultrasound hepatomegaly 15.7 cm



# Repeated CBC



**Hb 14.5 gm Hct 44.3%**

**WBC 1000 cumm**

**Platelets 78,000 cumm**

# Case 3

Day4

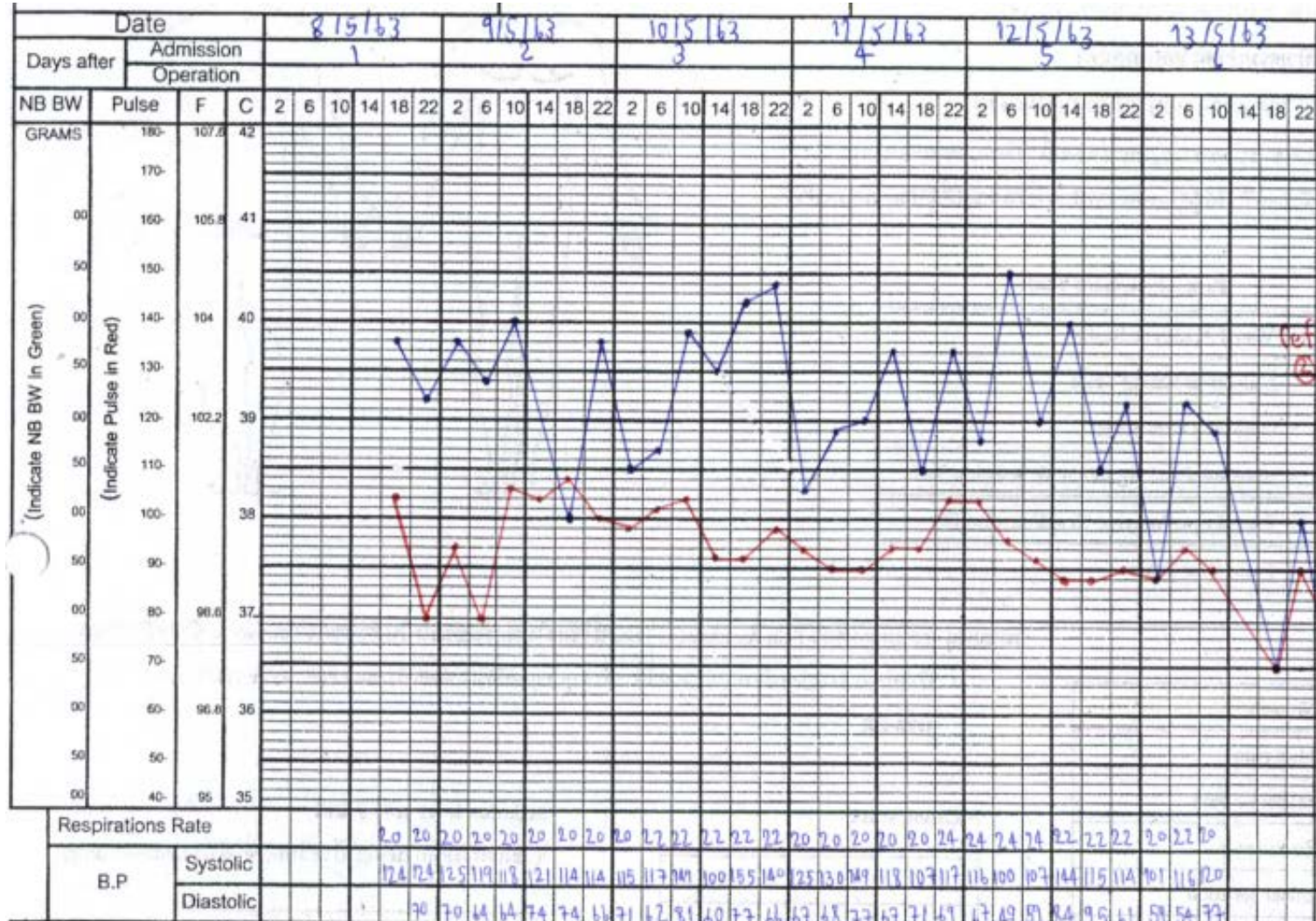
Day5

Day6

Day7

Day8

Day9



# Repeated CBC

**Hb 15.9 gm Hct 47.9%**

**WBC 5400 cumm**

**Platelets 68,000 cumm**



# Suspect hemophagocytic syndrome

**Consult Hemato**

**Repeated CBC**

**Hb 15.0 gm Hct 47.9%**

**WBC 5400 cumm**

**Platelets 68,000 cumm**





# Case 3

Day1

Day2

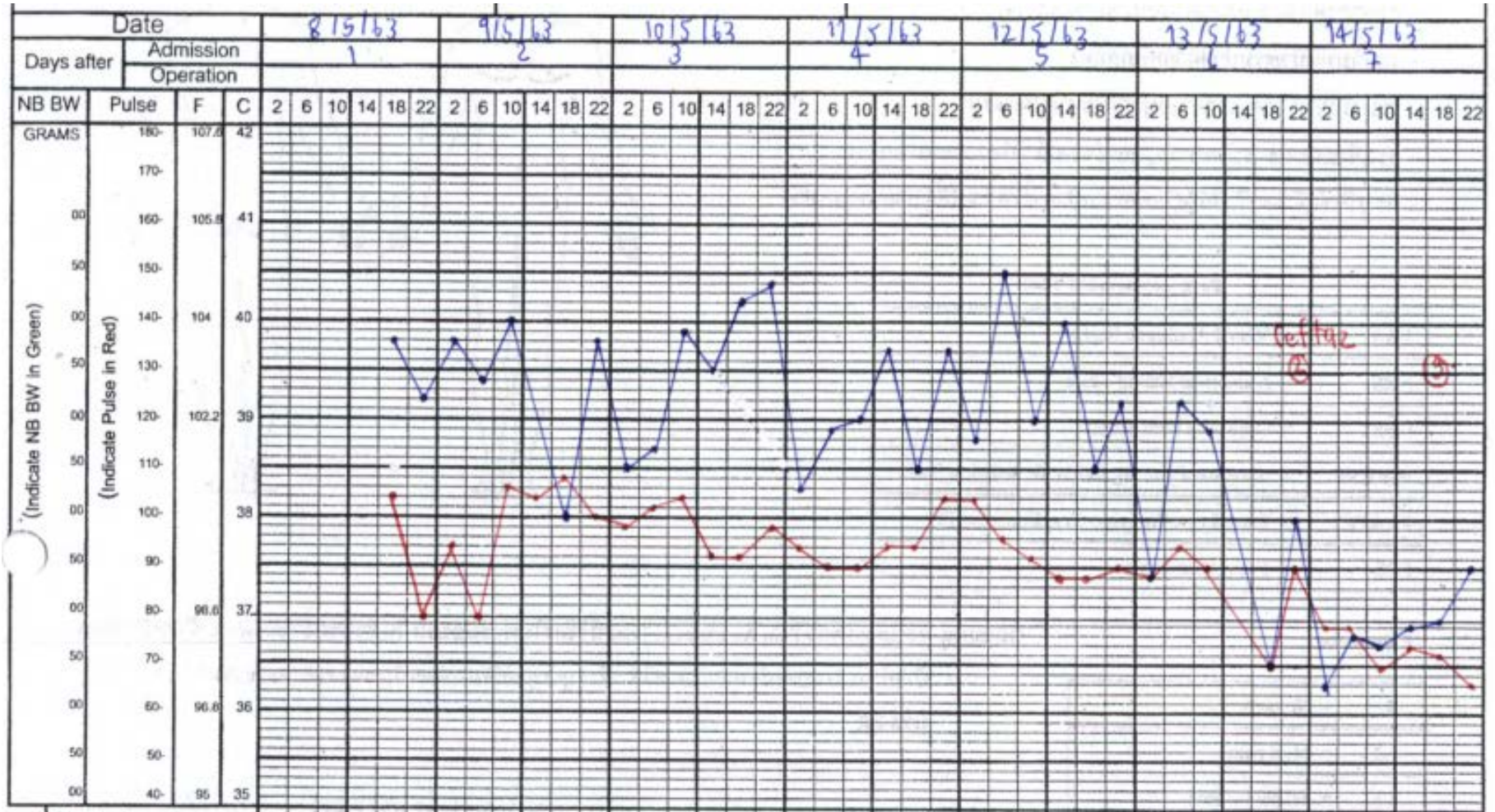
Day3

Day4

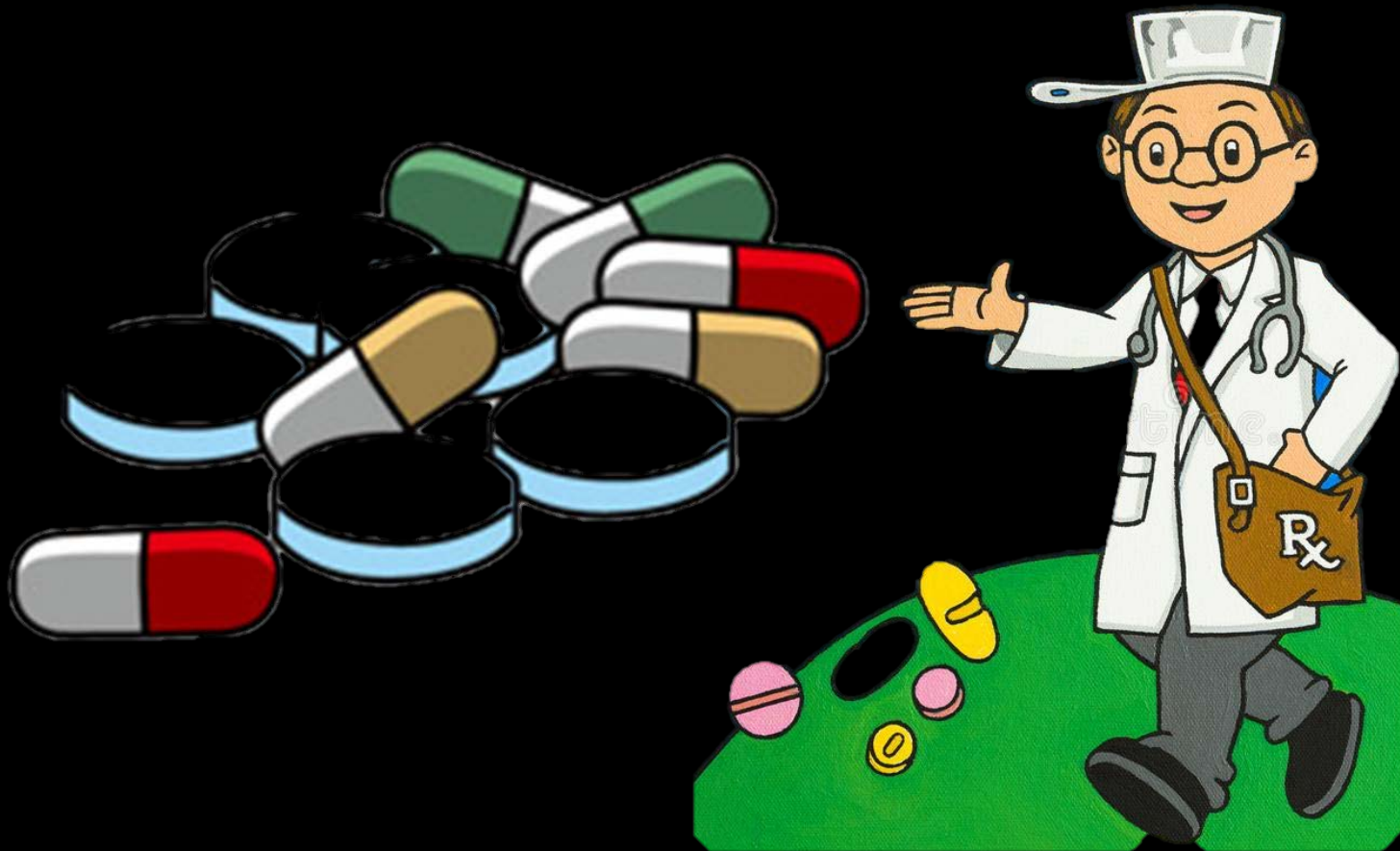
Day5

Day6

Day7



ไข่มลง หลังให้ยาปฏิชีวนะหลายตัว



น่าจะ consult อ. ยง มีตับโต  
จาก ultrasound





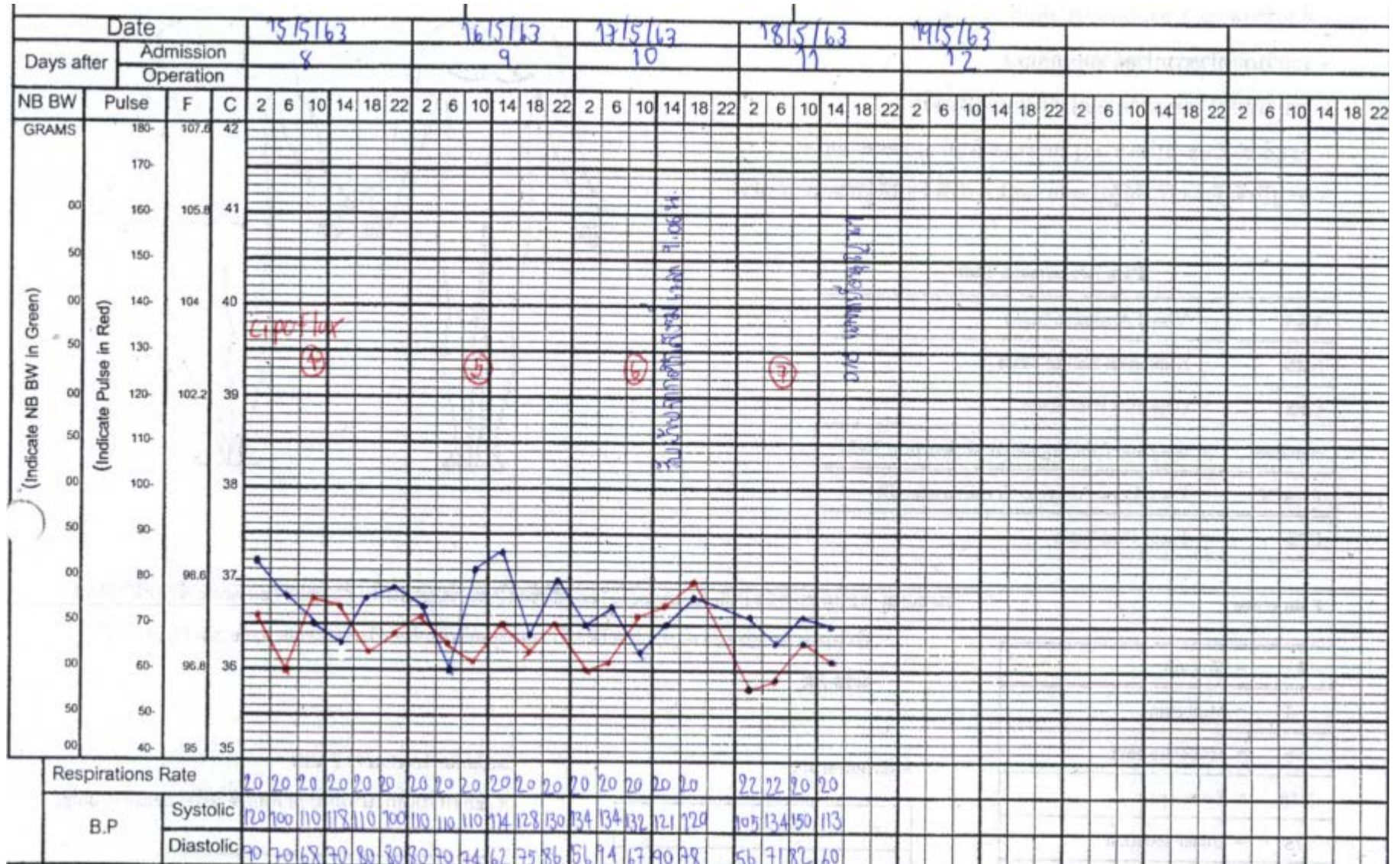
## Case 3

## Day8

## Day9

## Day10

# Day11







# **Severe Fever with Thrombocytopenia Syndrome Virus, Thailand.**

# Severe Fever with Thrombocytopenia Syndrome Virus: The First Case Report in Thailand

Supitcha Ongkittikul, MD<sup>1</sup>; Ruedeerat Watanawong, MD<sup>2</sup>; Photchana Rompho, RN<sup>3</sup>

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- Patient: 70-year-old Thai woman
- Begins with fever and nonspecific prodromal symptoms
- Progresses to encephalitis and multi-organ failure
- Laboratory Findings:
  - WBC: 2,100/mm<sup>3</sup> (45% neutrophils, 50% lymphocytes)
  - Platelet: 88,000/mm<sup>3</sup>
  - D-dimer: 2.96 µg/mL → Coagulopathy/DIC
  - Peripheral smear: ↓ platelets, few schistocytes, polychromasia
  - Liver enzymes: AST 814, ALT 142 IU/L
  - CPK: 1,336 U/L
  - LDH: 2,407 U/L
  - Dengue PCR and IgM: Negative
  - Specimens collected for rabies testing (saliva, serum, hairline, urine, CSF) → All results negative
  - Serum PCR detected SFTSV RNA → Confirmed diagnosis of Severe Fever with Thrombocytopenia Syndrome
- Upon further history taking, it was revealed that:
  - Her seven pet cats had fallen ill and died within one week, one month before her symptoms.

## EMERGING INFECTIOUS DISEASES®

EID Journal > Volume 28 > Number 12—December 2022 > Main Article

Volume 28, Number 12—December 2022

Research Letter

### Severe Fever with Thrombocytopenia Syndrome Virus Infection, Thailand, 2019–2020

Patthaya Rattanakomol, Sarawut Khongwicht, Piyada Linsuwanon, Keun Hwa Lee, Sompong Vongpunsawad, and Yong Poovorawan✉

Author affiliations: Chulalongkorn University, Bangkok, Thailand (P. Rattanakomol, S. Khongwicht, S. Vongpunsawad, Y. Poovorawan); US Army Medical Directorate–Armed Forces Research Institute of Medical Sciences, Bangkok, Thailand (P. Linsuwanon); Hanyang University, Seoul, South Korea (K.H. Lee)

[Cite This Article](#)

#### Abstract

Infection with severe fever with thrombocytopenia syndrome (SFTS) virus, which can cause hemorrhagic febrile illness, is often transmitted by ticks. We identified 3 patients with SFTS in or near Bangkok, Thailand. Our results underscore a need for heightened awareness by clinicians of possible SFTS virus, even in urban centers.

Severe fever with thrombocytopenia syndrome (SFTS) is a tickborne viral disease associated with acute fever, possibly accompanied by vomiting, diarrhea, fatigue, myalgia, and leukocytopenia (1). Most reports of infection have come from studies in South Korea, Japan, and China, although Taiwan, Vietnam, and Myanmar have had confirmed cases in recent years (2). Severe infections can cause hemorrhagic fever and multiple organ failure leading to death. SFTS results from infection by the SFTS virus (SFTSV, newly renamed *Dabie bandavirus*), an RNA virus in the family Phenuiviridae, genus *Bandavirus* (3). More frequent arbovirus infections in Thailand, primarily dengue and chikungunya, often confound diagnosis of febrile illness caused by other viruses such as SFTSV because most clinicians lack awareness.

Archives of Virology (2023) 168:271  
<https://doi.org/10.1007/s00705-023-05897-1>

#### ANNOTATED SEQUENCE RECORD



### Severe fever with thrombocytopenia syndrome virus genotype B in Thailand

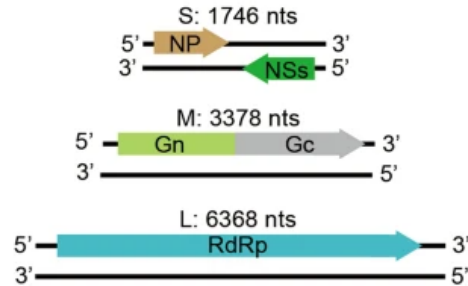
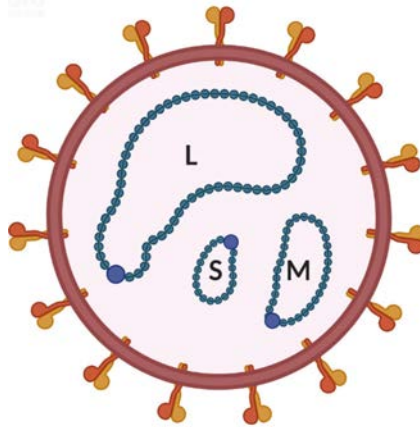
Patthaya Rattanakomol<sup>1</sup> · Sarawut Khongwicht<sup>1</sup> · Watchaporn Chuchaona<sup>1</sup> · Sompong Vongpunsawad<sup>1</sup> · Yong Poovorawan<sup>1</sup>

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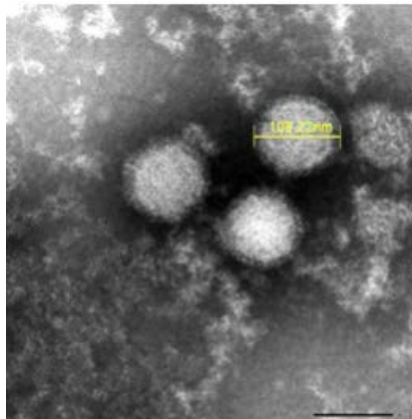
#### Abstract

Severe fever with thrombocytopenia syndrome virus (SFTSV) has been reported in many countries in Southeast Asia, which expands the original geographic range of China, Korea, and Japan. Here, we report the complete genome sequences of two Thai SFTSV strains previously identified in patients with undifferentiated febrile illness in 2020. Phylogenetically, both clustered with SFTSV genotype B strains and were most closely related to those previously reported in central China ( $\geq 99.0\%$  nucleotide sequence identity) in the L, M, and S gene segments. Nine amino acid residues encoded by one or more Thai SFTSV genomes differed from those found in global strains. Interestingly, the observed differences in numerous residues between the Thai strains suggest possible separate introductions of different variants into the region.

# Severe Fever with Thrombocytopenia Syndrome Virus (SFTSV)



<https://doi.org/10.1186/s12985-023-02033-y>



IASR  
Infectious Agents Surveillance Report

Enveloped, negative-sense, single-stranded, three-genomic segmented, RNA virus

**Order:** Bunyavirales (formerly the Bunyaviridae family)

**Family:** Phenuiviridae

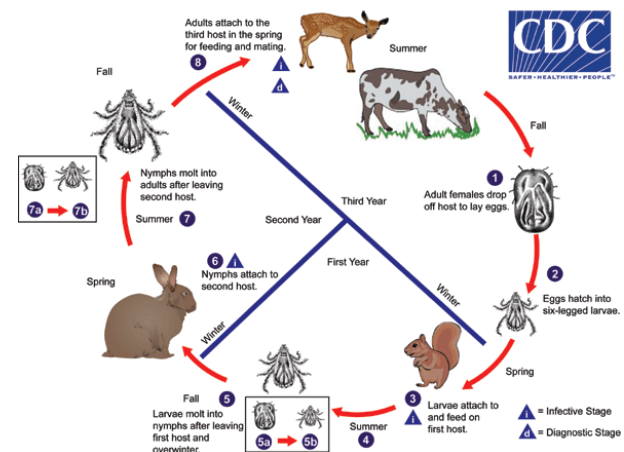
**Genus:** Bandavirus

Renamed as Dabie bandavirus

**Potential vectors:** ticks (*Haemaphysalis longicornis*, *Amblyomma testudinarium*, *Ixodes nipponensis*, and *Rhipicephalus microplus*)



## Three-Host Ixodid Tick Life Cycle



# Severe Fever with Thrombocytopenia Syndrome (SFTS)

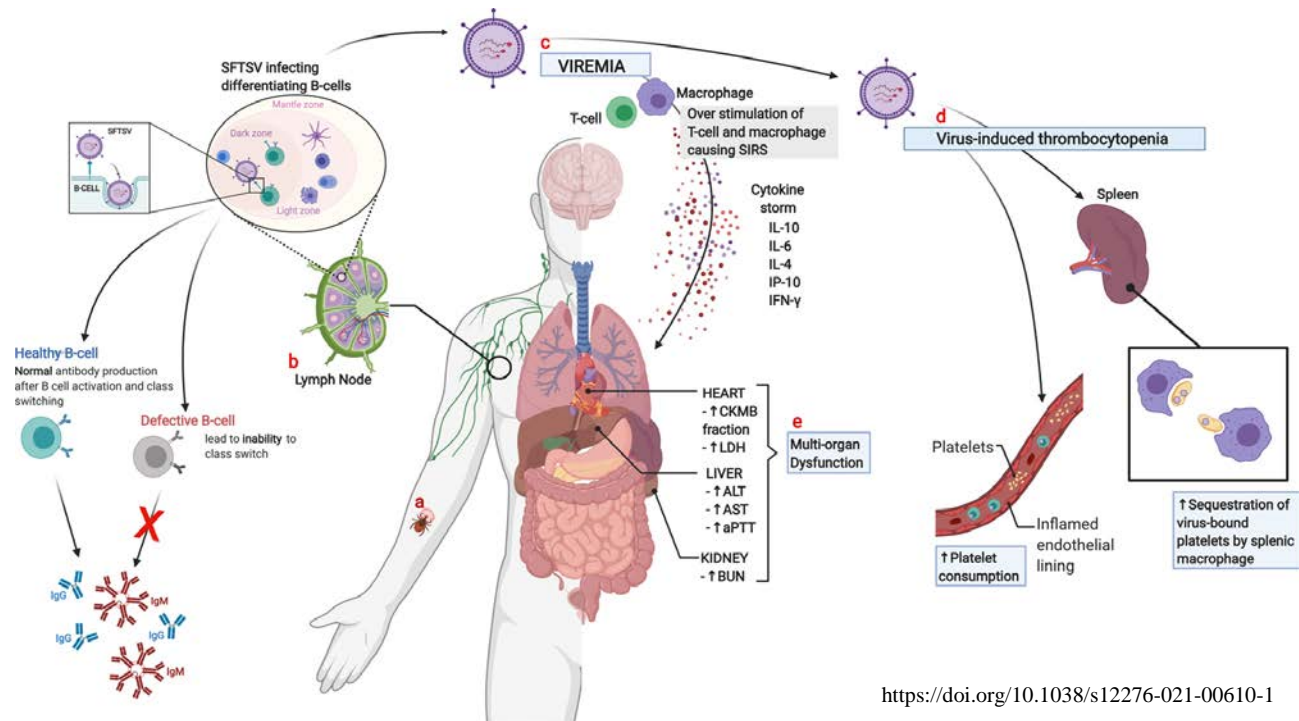
a tick-borne viral disease, caused by SFTSV, associated with acute fever, which may accompany vomiting, diarrhea, fatigue, myalgia, rash, and leukocytopenia



## Clinical disease course

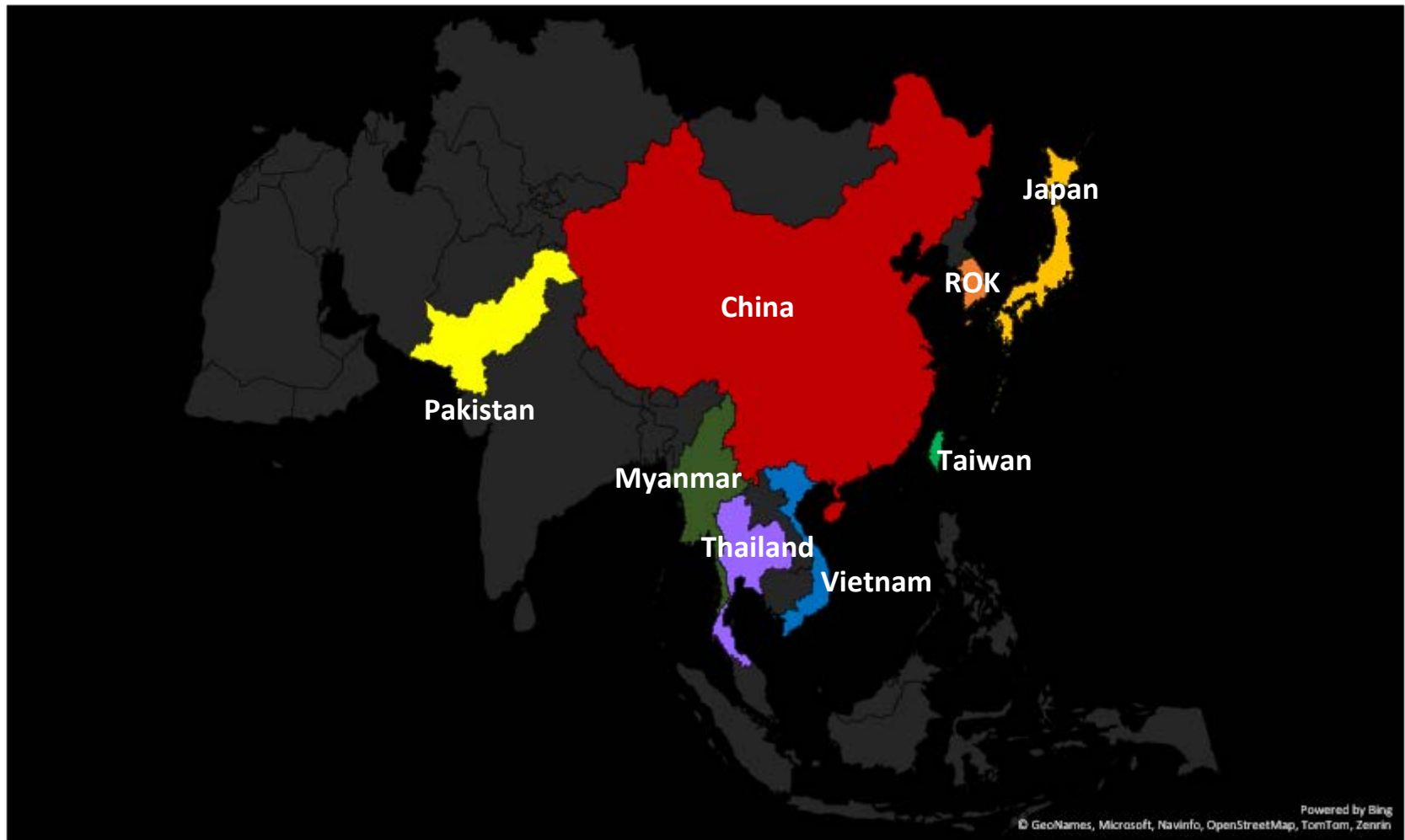
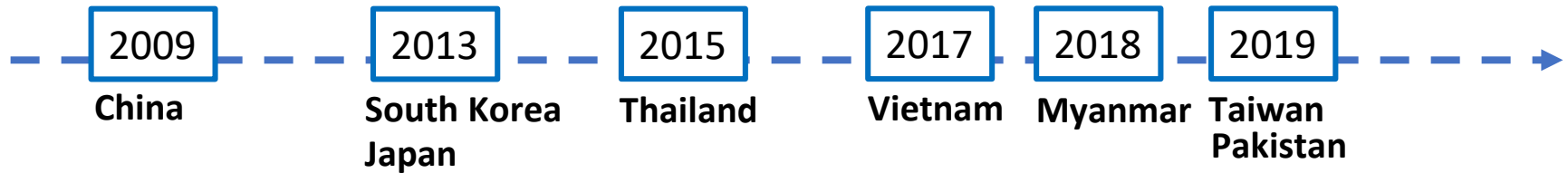
1. Incubation: 5 to 14 days since tick bites - onset
2. Fever: 1 to 7 days -- fever, headache, and GI symptoms  
When virus load ↑

- **Progressive PLT and WBC**  
↓
- Serum biochemistry parameters; **ALT, AST, LDH, BUN and CPK** ↑
- 3. Multiple organ failure
- 4. Convalescence



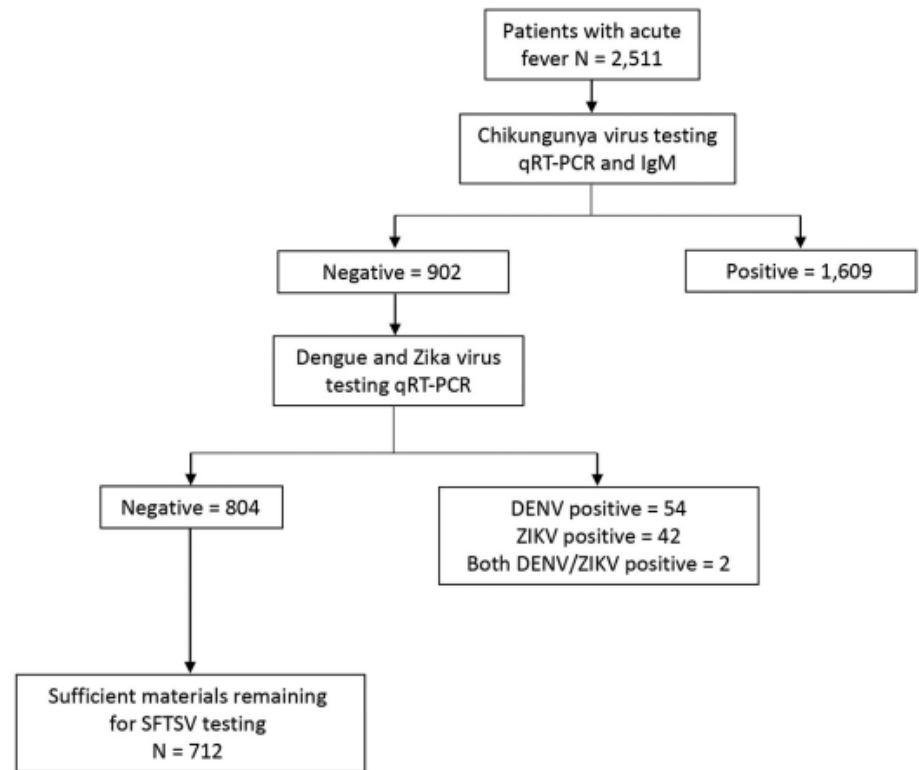
<https://doi.org/10.1038/s12276-021-00610-1>

# SFTSV infection has increased rapidly in both incidence and geographical range since 2009



# Most frequent arbovirus infections in Thailand, primarily dengue and chikungunya, often confound febrile illness caused by other virus infections

- At the end of 2018, an upsurge of chikungunya virus (CHIKV) infection in Thailand found that >70% of acute febrile illnesses were laboratory-confirmed CHIKV (Khongwichit S *et al.*, 2021).
- The eventual decreasing proportion of CHIKV-positive samples led to screening for other viral etiologies of acute fever including dengue and Zika.
- At the time, SFTSV was reported in Vietnam.
- Therefore, samples tested negative for all three viruses were examined for possible SFTSV infection.



**Appendix Figure.** Febrile illness samples chosen for SFTSV testing

P. Rattanakomol *et al.*, 2022



# A possible SFTSV in urban centers

**Table.** Detection of SFTSV in Thai patients, 2019-2020.

	Patient 1	Patient 2	Patient 3
<b>Information</b>			
Age and gender	60, male	16, male	52, female
Location	Bangkok	Chachoengsao	Bangkok
Collection date	14 Nov 2019	10 May 2020	19 Oct 2020
<b>Clinical manifestations</b>			
Temperature and symptoms	37.2 °C Myalgia, arthralgia, cough, nausea, vomiting, abdominal pain, diarrhea	40.6 °C Myalgia	38.1 °C Myalgia Arthralgia
<b>Laboratory findings (most extreme)</b>			
White blood cells (4,100-10,900 cells/ $\mu$ L)	1,790	900	2,770
Neutrophil (40-72 %)	45	31	121,000
Lymphocyte (18-49 %)	42	59	62
Platelet (140,000-400,000 cells/ $\mu$ L)	107,000	45,000	34
Aspartate aminotransferase (<40 U/L)	No done	102	1,758
Alanine aminotransferase (<41 U/L)	24	63	973
<b>Pathogens not detected</b>			
Rickettsia/Orientia spp. and	Influenza A/B	Influenza A/B, Epstein-Barr, Hepatitis B/C, SARS-CoV-2, Malaria, Leptospira, Burkholderia pseudomalle	Hepatitis A/B

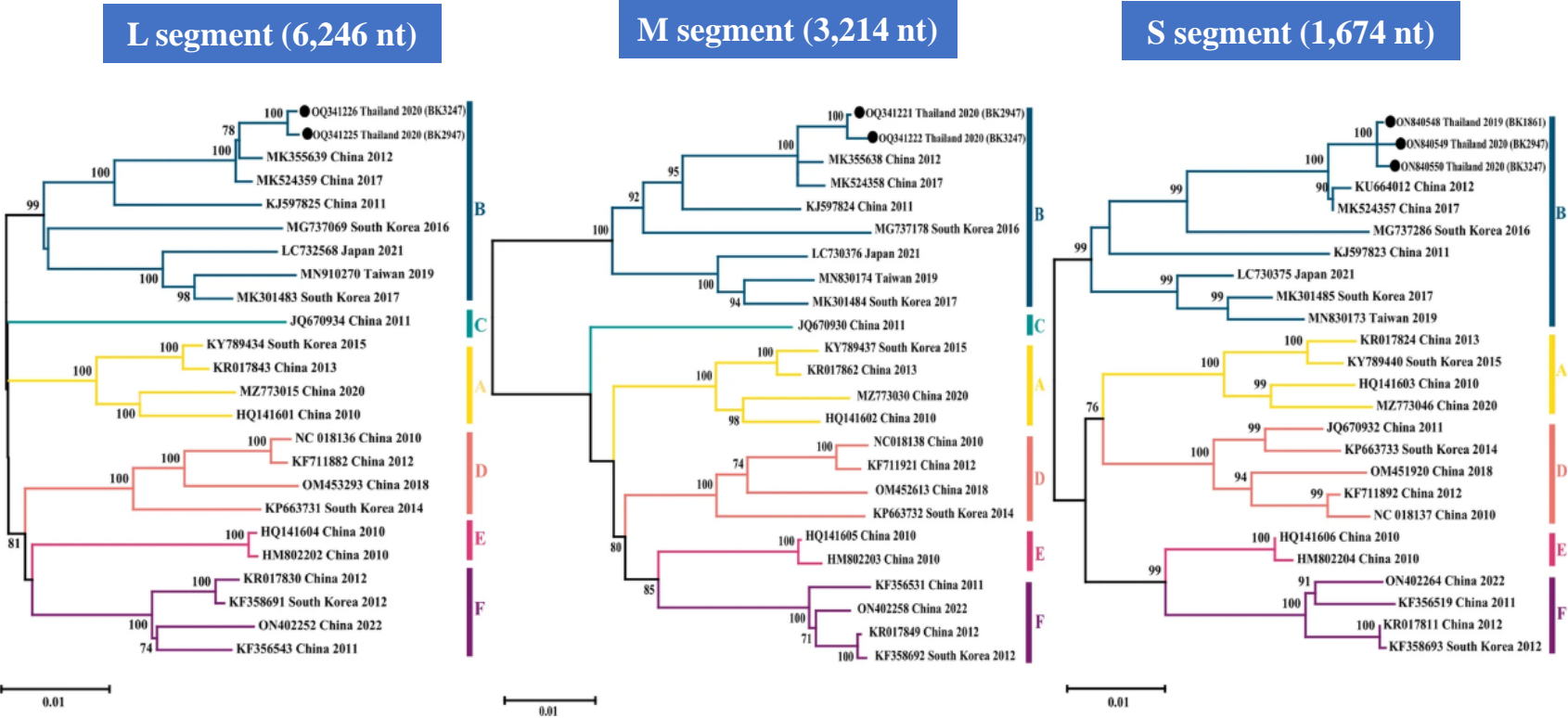




Severe fever with thrombocytopenia syndrome virus genotype B  
in Thailand

Patthaya Rattanakomol<sup>1</sup> · Sarawut Khongwicht<sup>1</sup> · Watchaporn Chuchaona<sup>1</sup> · Sompong Vongpunsawad<sup>1</sup> ·  
Yong Poovorawan<sup>1</sup>

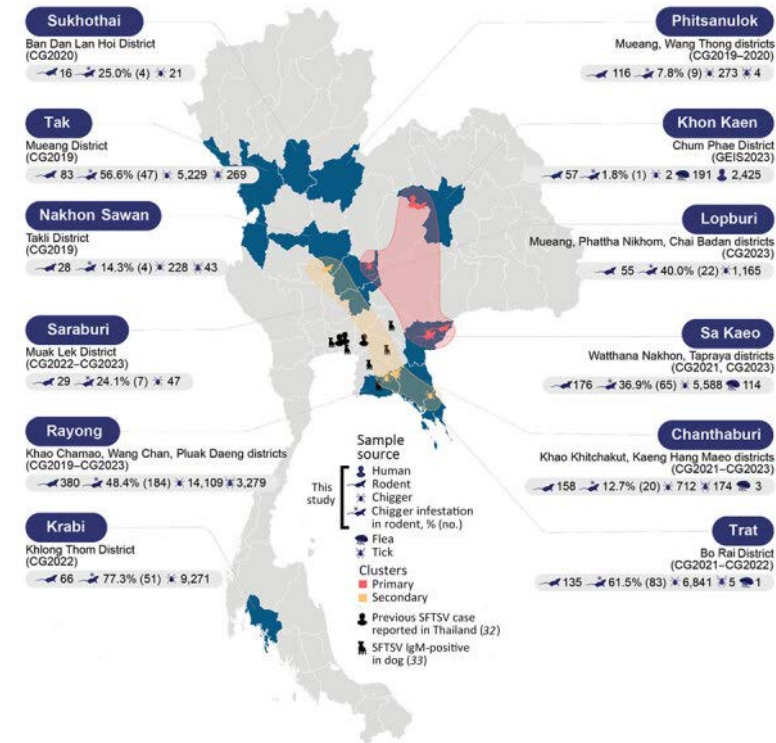
Phylogenetic analysis of the SFTSV strains in Thailand



Phylogenetic analysis of the S, M, and L segments confirmed that Thai strains clustered together and were most closely related to previously reported Chinese strains.

Comprehensive Surveillance of Severe Fever with Thrombocytopenia Syndrome Virus in Patients with Acute Febrile Illness, Wild Rodents, and Trombiculid Larval Mites, Thailand

Flayda Limswan <sup>1,2\*</sup>, Jong Poonvongwan <sup>1</sup>, Keun-Hwa Lee <sup>3</sup>, Nutthanun Aoyasawadi <sup>1</sup>, Sirima Wornpaiboon <sup>1</sup>, Chawin Limswan <sup>1</sup>, Vilasornak Vuttitanachot <sup>1</sup>, Sarachai Lerolabrat <sup>1</sup>, Somnong Yongsamarn <sup>1</sup>, Porajalin Nilvanimit <sup>1</sup>, Yossamong Paladsing <sup>1</sup>, Erica Lindoth <sup>1</sup>



# Molecular and Serologic Evidence of SFTSV Infection in Acute Febrile Illness Cases in Khon Kaen, Thailand

SFTSV RNA was detected in **1.6% (38/2,425)** of AFI patients at Chum Phae Hospital, Thailand (2015–2021).

**Sex ratio (M:F): 1.4:1.0**

**Median age: 47.2 years (range: 15.9–86.7; IQR: 29–64)**

**No co-infections with other pathogens**

**Clinical features:**

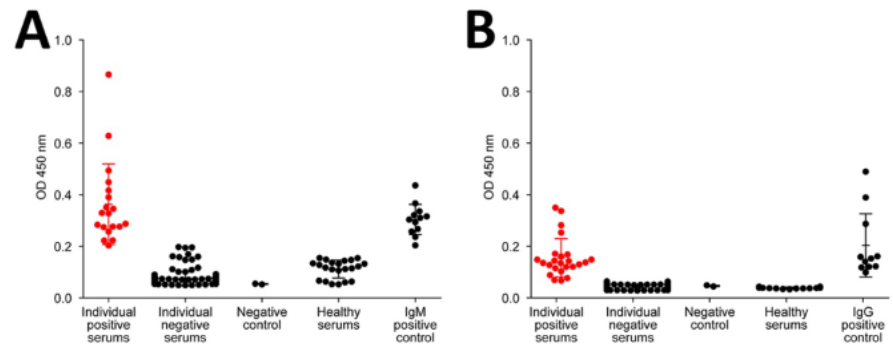
- Fever (mean 38.0°C, range 36.1–40.3°C)
- Headache (20.5%), dizziness (12.8%)
- Thrombocytopenia in 3 cases

**All patients recovered without severe complications**

**SFTSV seropositivity:** 16 patients (0.7%) IgM, 54 patients (2.2%) IgG, 3 patients (0.1%) both → **total 73 of 2,425 (3%)**

Average OD450 of SFTSV IgM positive control: 0.233

Average OD450 of SFTSV IgG positive control: 0.172



Appendix Figure 2. ELISA results showing optical density at 450 nm of the tested samples.

# Comprehensive Surveillance of Severe Fever with Thrombocytopenia Syndrome Virus in Patients with Acute Febrile Illness, Wild Rodents, and Trombiculid Larval Mites, Thailand

Piyada Linsuwanon<sup>1,89</sup>, Yong Poovorawan<sup>1</sup>, Keun Hwa Lee<sup>1</sup>, Nutthanun Auyasawadi<sup>1</sup>, Sirima Wongwalrot<sup>1</sup>, Chawin Limsuwan<sup>1</sup>, Yiboonsak Vuthitanachot<sup>1</sup>, Surachai Leepitakrat<sup>1</sup>, Sompong Vongpunsawasdi<sup>1</sup>, Pornjarim Nilyanimit<sup>1</sup>, Yossapong Paladsing<sup>1</sup>, Erica Lindroth<sup>1</sup>

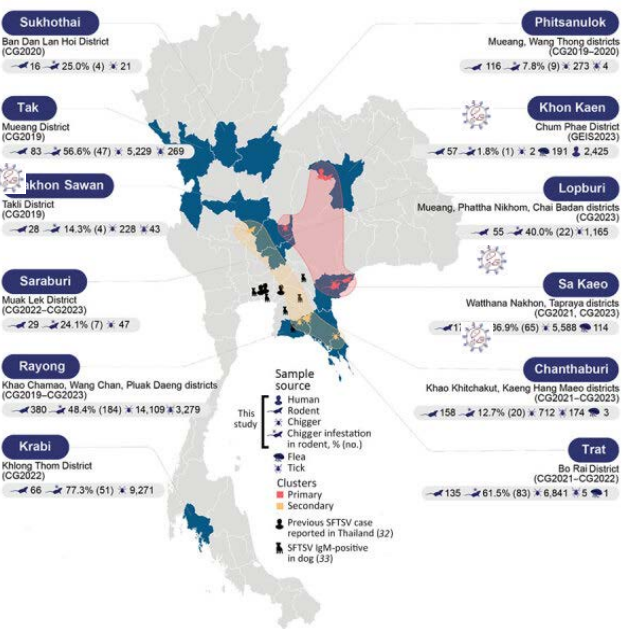
# Assessing Rodents in SFTSV Transmission: qRT-PCR of 1,019 Rodents

Table 2. Positivity rates of SFTSV RNA detected in wild rodents, Thailand, 2015–2021\*

Family	Species	No. (%) positive	SFTSV RNA–positive, by tissue			Average RNA level, copies/mL
			Lung	Liver	Spleen	
Muridae	<i>Rattus tanezumi</i> rat	1/559 (0.2)	–	1	–	$4.07 \times 10^4$
	<i>R. exulans</i> rat	1/98 (1)	–	–	1	$3.05 \times 10^3$
	<i>R. novogicus</i> rat	1/16 (6.3)	1	–	–	$5.73 \times 10^3$
	<i>Mus cervicolor</i> mouse	0/6	–	–	–	–
	<i>M. caroli</i> mouse	1/7 (14.3)	1	–	–	$4.03 \times 10^3$
	<i>Bandicota indica</i> rat	2/113 (1.8)	1	1	–	$4.89 \times 10^3$
	<i>B. savilei</i> rat	2/121 (1.7)	–	2	–	$1.01 \times 10^4$
	<i>Maxomys surifer</i> rat	0/23	–	–	–	–
	<i>Niviventer fulvescens</i> rat	0/5	–	–	–	–
	<i>Berylmys berdmorei</i> rat	1/23 (4.3)	–	–	1	$1.71 \times 10^4$
	<i>B. bowersi</i> rat	0/1	–	–	–	–
Tupaiaidae	<i>Chiromyscus chiropus</i> rat	0/1	–	–	–	–
	<i>Tupaia belangeri</i> shrew	0/19	–	–	–	–
Sciuridae	<i>T. glis</i> shrew	0/9	–	–	–	–
	<i>Menetes berdmorei</i> squirrel	2/19 (10.5)	1	–	1	$4.06 \times 10^3$
3 families	15 species	11/11,019 (1.1)	4	4	3	

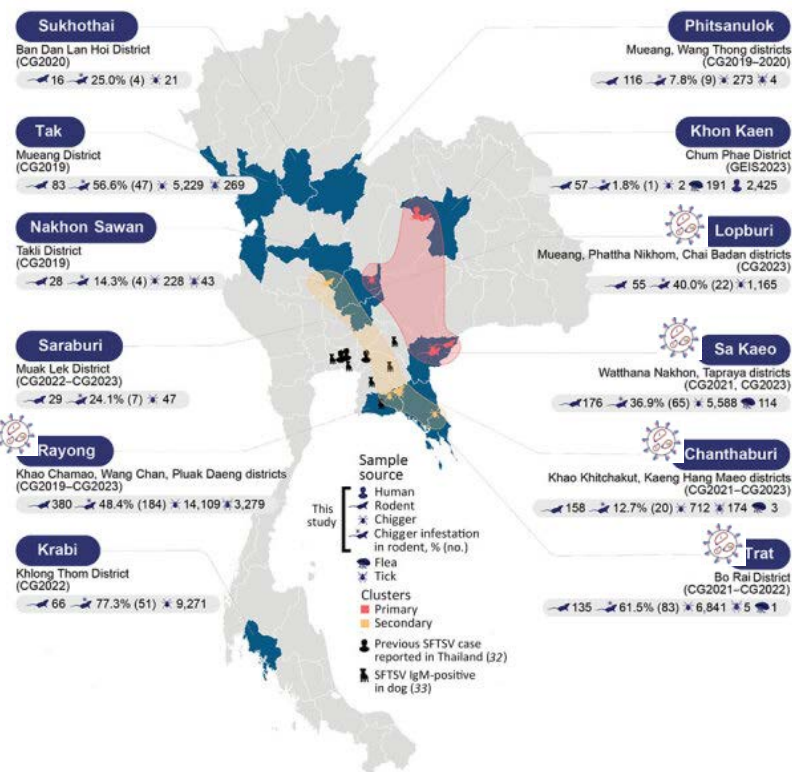
\*SFTSV, severe fever with thrombocytopenia syndrome virus; –, negative result.

Eleven rodents from 8 species were positive for SFTSV RNA, indicating an overall positivity rate of 1.1%.



## Comprehensive Surveillance of Severe Fever with Thrombocytopenia Syndrome Virus in Patients with Acute Febrile Illness, Wild Rodents, and Trombiculid Larval Mites, Thailand

Piyada Linsuwanon<sup>1,6\*</sup>, Yong Poovorawan<sup>1</sup>, Keun Hwa Lee<sup>1</sup>, Nutthanun Auyasawadi<sup>1</sup>, Sirima Wongwairat<sup>1</sup>, Chawin Limsuwan<sup>1</sup>, Viboonsak Yuthitanachot<sup>1</sup>, Surachai Leepitakrat<sup>1</sup>, Sompong Vongpunsawasdi<sup>1</sup>, Pornjarim Nilyanimit<sup>1</sup>, Yossapong Paladsing<sup>1</sup>, Erica Lindroth<sup>1</sup>



# SFTSV RNA Detected in Chiggers from Wild Rodents

Appendix Table 2. SFTSV RNA positivity rates in chiggers

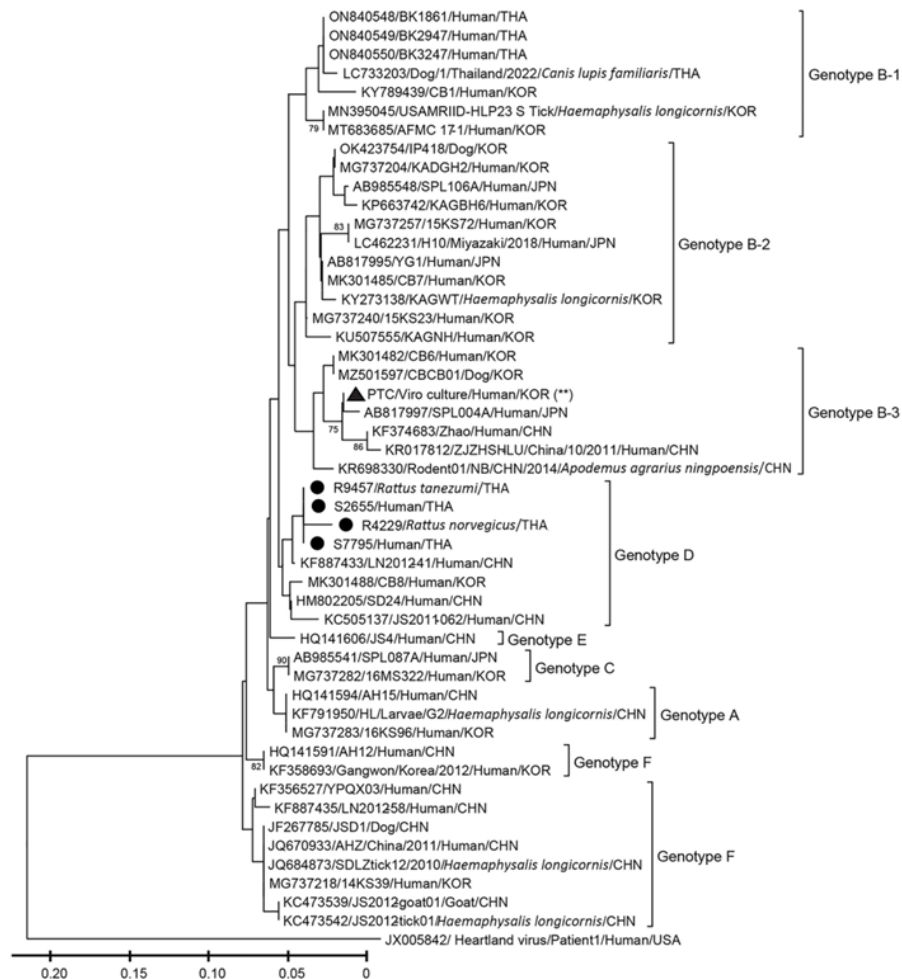
				N positive by host species															
Family	Genus	N chigger	N positive (%)	<i>Rattus tanezum</i>	<i>Rattus exulans</i>	<i>Rattus norvegicus</i>	<i>Mus cervicolor</i>	<i>Mus caroli</i>	<i>Bandicota indica</i>	<i>Bandicota saviei</i>	<i>Maxomys surifer</i>	<i>Niviventer fulvescens</i>	<i>Berylmys berdmorei</i>	<i>Berylmys bowersi</i>	<i>Chinomyscus chiropus</i>	<i>Tupaia belandieri</i>	<i>Tupaia alis</i>	<i>Menetes berdmorei</i>	Average RNA level (copies/ml)
<i>Trombiculidae</i>	<i>Ascoschoengastia</i> sp.	178	1 (0.6%)	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	$1.85 \times 10^4$
	<i>Leptotrombidium</i> sp.	73	1 (1.4%)	-	-	-	-	-	-	-	-	-	-	-	-	1	-	-	$1.49 \times 10^4$
	<i>Walchia</i> sp.	163	5 (3.1%)	1	-	-	-	-	1	3	-	-	-	-	-	-	-	-	$2.89 \times 10^4$
	<i>Blankaertia</i> sp.	1	1 (100%)	-	-	-	-	-	1	-	-	-	-	-	-	-	-	-	$1.38 \times 10^4$
	<i>Gahriepia</i> sp.	136	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
	<i>Walchiella</i> sp.	6	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
	<i>Schoengastiella</i> sp.	3	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
	<i>Eutrombicula</i> sp.	2	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
	<i>Helenicula</i> sp.	1	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
	Unidentified	Other mites	10	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-
	10 genera	573	8 (1.4%)	2	-	-	-	-	2	3	-	-	-	-	-	1	-	-	$2.39 \times 10^4$

8 of 573 chiggers (1.4%) from 155 wild rodents were SFTSV RNA–positive (avg.  $2.4 \times 10^4$  copies/chigger); positive chiggers were collected from 6 rodents and belonged to 4 chigger genera.



**Comprehensive Surveillance of Severe Fever with Thrombocytopenia Syndrome Virus in Patients with Acute Febrile Illness, Wild Rodents, and Trombiculid Larval Mites, Thailand**

Piyada Limsawann<sup>1,2\*</sup>, Yong Poovorawan<sup>3</sup>, Kuan Hwa Lee<sup>4</sup>, Nutthanun Aursawasdi<sup>5</sup>, Sirima Wongkajorn<sup>6</sup>,  
Chawin Limswan<sup>7</sup>, Yongsak Yuthitachot<sup>8</sup>, Suwacha Leelakul<sup>9</sup>, Sompong Yongsawasdi<sup>9</sup>, Pornjerm  
Niyonim<sup>9</sup>, Yossasong Paladang<sup>9</sup>, Erika Lindoth<sup>9</sup>



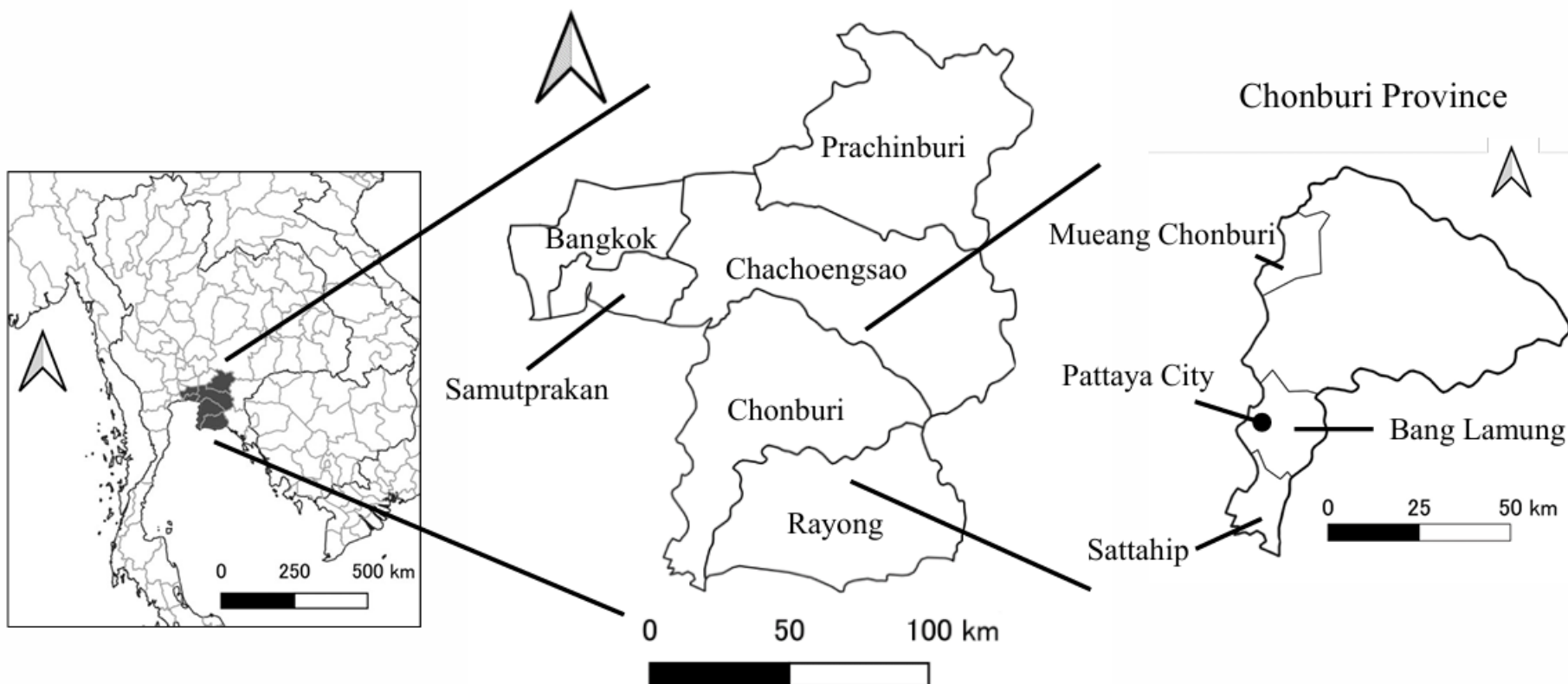
Phylogenetic analysis of the partial NS gene sequence (124 bp) from two SFTSV RNA–positive patients and two rodents revealed a close genetic relationship to genotype D.

The strains shared high nucleotide similarity with strain LN2012–41 (GenBank accession no. KF887433), which was previously identified in a patient in China in 2012.

# High Seroprevalence of Severe Fever with Thrombocytopenia Syndrome Virus Infection among the Dog Population in Thailand

Keita Ishijima <sup>1,†</sup>, Thanaporn Phichitraslip <sup>2,†</sup>, Nattakarn Naimon <sup>2</sup>, Preeyaporn Ploypichai <sup>2</sup>, Benyapa Kriebkajon <sup>2</sup>, Torntun Chinarak <sup>2</sup>, Jirasin Sridaphan <sup>2</sup>, Anamika Kritiyakan <sup>2</sup>, Noppadol Prasertsinchaoen <sup>2</sup>, Sathaporn Jittapalpong <sup>2</sup>, Kanate Tangcham <sup>3</sup>, Worawut Rerkamnuaychoke <sup>4</sup>, Yudai Kuroda <sup>1</sup>, Masakatsu Taira <sup>1</sup>, Kango Tatemoto <sup>1</sup>, Eunsil Park <sup>1</sup>, Milagros Virhuez-Mendoza <sup>1</sup>, Yusuke Inoue <sup>1,2</sup>, Michiko Harada <sup>1,2</sup>, Tsukasa Yamamoto <sup>1,2</sup>, Ayano Nishino <sup>1,2</sup>, Aya Matsui <sup>1</sup> and Ken Maeda <sup>1,2,\*</sup>

**Dog serum samples (n=458) were collected from 16 districts across six provinces in Thailand between April 2021 and April 2022.**



# Detection of Anti-SFTSV Antibodies from Dogs

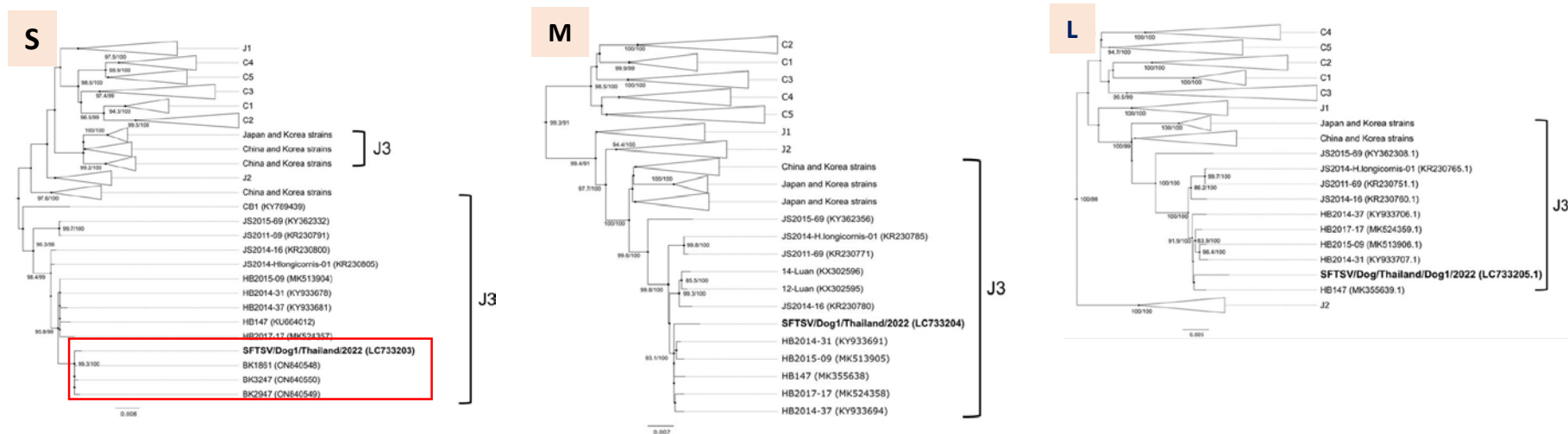
Results of the anti-SFTSV IgG ELISA and virus-neutralization test in dogs in Thailand.

Province	District	ELISA (OD > 0.129)		50% Focus Reduction Neutralization Test (≥1:10)		Minimum Positive Ratio (%)
		No. of Examined Dogs	No. of Positive Dogs	No. of Examined Dogs	No. of Positive Dogs	
Prachinburi		17	1	1	0	0.0
Bangkok		143	12	12	6	4.2
Chachoengsao		18	2	2	2	11.1
Samutprakan		81	19	16	10	12.3
Rayong		14	2	2	1	7.1
Chonburi	Mueang Chonburi	27	6	6	3	11.1
	Bang Lamung	56	5	5	3	5.4
	Sattahip	95	56	52	50	52.6
	Pattaya city	7	3	2	1	14.3
Total		458	106	98	76	16.6

- First screening showed 106 dogs (23.1%) were seropositive for anti-SFTSV IgG.
- 76 of 98 (**16.6%**) samples were confirmed positive by FRNT50
- Highest regional seropositive rate:
  - Sattahip, Chonburi with 52.6%
- Further analysis revealed:
  - Most positive samples came from a dog shelter in Sattahip
  - In February 2022, 50 of 64 dogs at this shelter were positive (78% seropositivity)

# SFTSV RNA Detection and Genome Sequencing

- RT-PCR performed on **57 dog serum samples** (Sattahip, Feb 2022)  
→ **7 samples** excluded due to insufficient volume
- **SFTSV RNA detected** in **1 sample** from a healthy dog
- Complete coding sequences of **S, M, and L segments** were obtained
- Virus designated:  
→ **SFTSV/Dog/Thailand/Dog1/2022**
- Nucleotide sequences closely related to:  
→ **SFTSV strains from Thailand and China**





# SFTSV Seroprevalence in Humans and PCR-Based Tick Surveillance in Thailand, 2019-2023

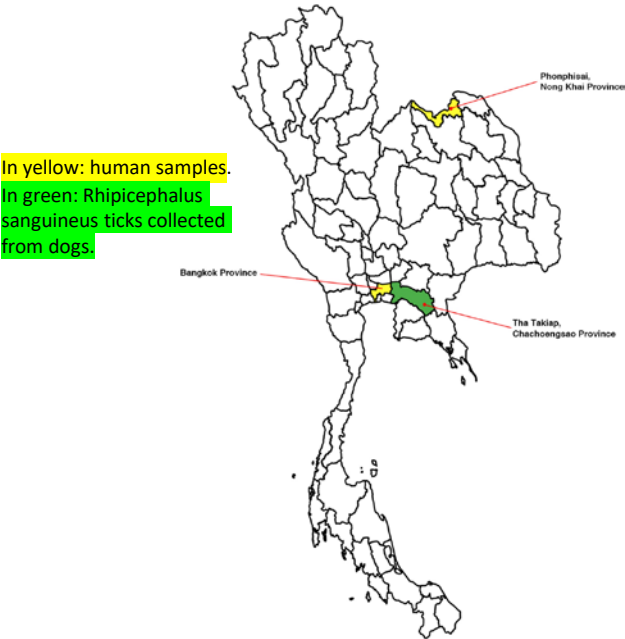
**Table 1 Human seroprevalence of severe fever with thrombocytopenia syndrome and molecular analysis of *Rhipicephalus sanguineus* ticks in Thailand.**

From: [Seroprevalence study in humans and molecular detection in \*Rhipicephalus sanguineus\* ticks of severe fever with thrombocytopenia syndrome virus in Thailand](#)

Sample	Collection	Location	Province	SFTSV					
				ELISA		VNT		qPCR <sup>a</sup>	
				No. total	No. pos	No. total	No. pos	No. total	No. pos
Human sera	Blood donors	Thai Red Cross	Bangkok	940	5	5	0	NA	NA
Human sera	Dengue suspected	Phonphisai Hospital	Nong Khai	222	7	7	0	NA	NA
<i>Rh. sanguineus</i> ticks	Dogs	Tha Takiap Subdistrict	Chachoengsao	NA	NA	NA	NA	433 (50 pools)	0
<i>Rh. sanguineus</i> eggs	<i>Rh. sanguineus</i> ticks	Tha Takiap Subdistrict	Chachoengsao	NA	NA	NA	NA	12 pools	0

ELISA Enzyme-linked immunosorbent assay, NA Not applicable, pos Positive, qPCR Real-time polymerase chain reaction, SFTSV Severe fever with thrombocytopenia syndrome virus, VNT Virus neutralization test.

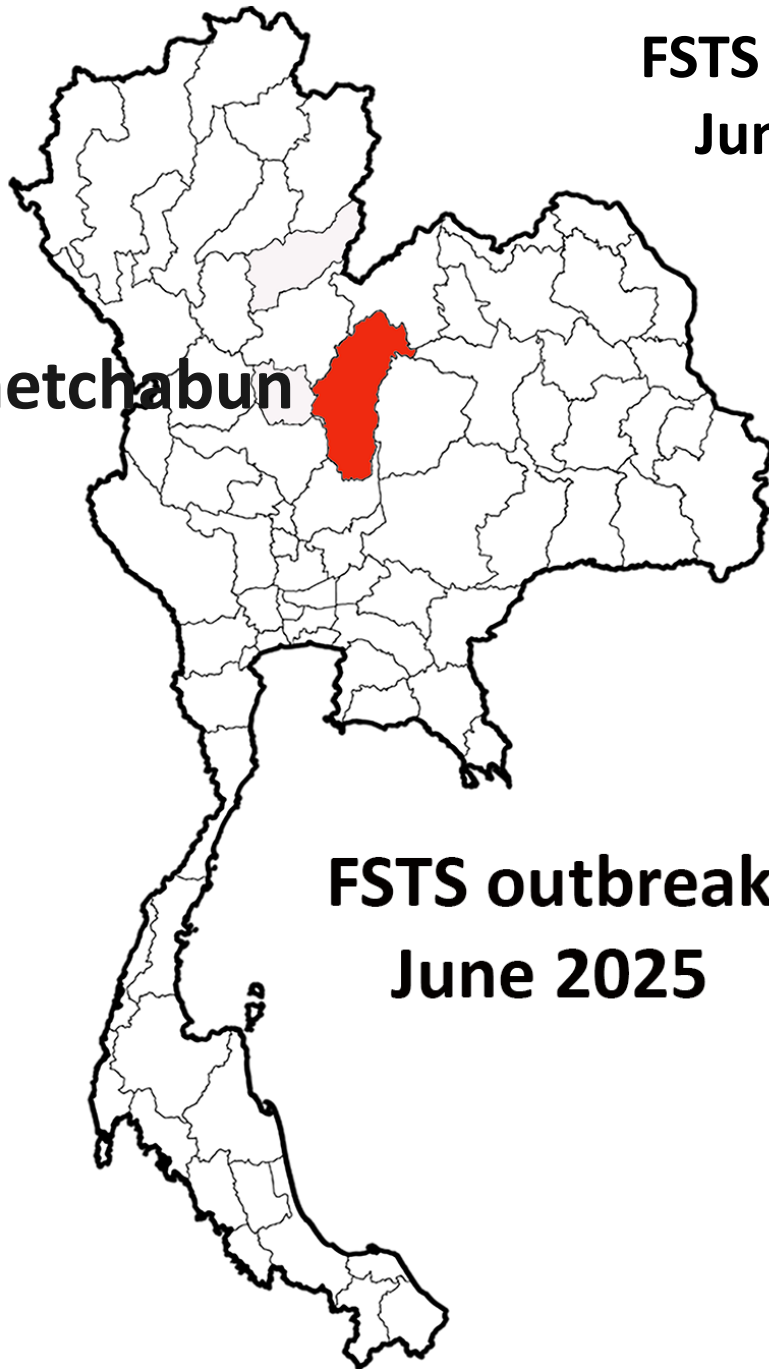
<sup>a</sup>Pathogens: SFTSV, Crimean-Congo hemorrhagic fever, *Coxiella* spp., *Bartonella* spp., and *Rickettsia* spp.



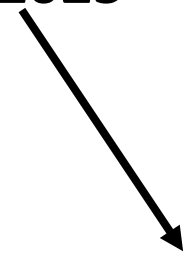
- 1,162 human serum samples from Bangkok (940 asymptomatic adult volunteer blood donors) and Nong Khai (individuals suspected of dengue virus) were tested for anti-SFTSV IgG
  - 12 samples (1.1%) were ELISA-positive
  - All 12 negative in virus neutralization test
- 433 *Rh. sanguineus* ticks from 49 dogs (Chachoengsao, 2023) tested for: SFTSV, CCHF, *Coxiella* spp., *Bartonella* spp., and *Rickettsia* spp.
  - No pathogens detected

# **Outbreak of SFTS with 2 Fatal Cases**

**Phetchabun**

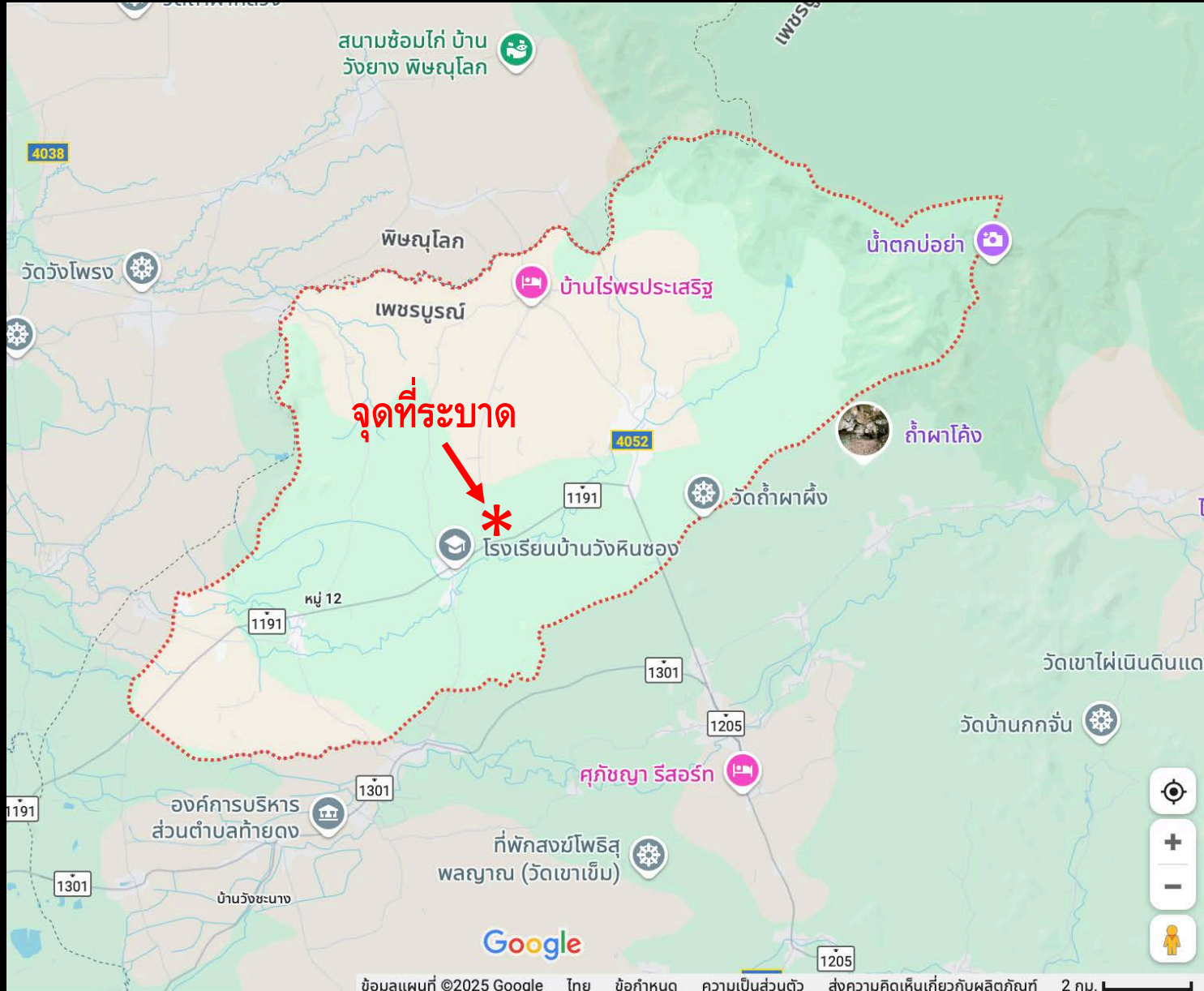


**FSTS outbreak  
June 2025**



**FSTS outbreak  
June 2025**

# จุดที่เกิดการระเบิด





บ้านหลังที่ 1

วัดเนินศิลาเพชร

วัดวังสมปดี  
สามัคคีธรรม

5.35 กม.

โรงเรียนบ้านวังหินทอง

วัดอรัญจิตสว่าง

โรงพยาบาล

5.35 กม.

บ้านหลังที่ 2

โรงเรียนวัง  
โป่งพยอม

# Case 1. (Fatal case)

- ชายไทยอายุ 82 ปี อาศัยอยู่วังโป่ง จังหวัดเพชรบูรณ์ แข็งแรงดี
- 3 วัน ใช้ ไอเสมหะเหนียวๆ ไม่มีน้ำมูก ไม่เหนื่อย ไม่ปวดท้อง ยังรู้ตัวรู้เรื่อง ไม่มีแขนขาอ่อนแรง ไปตรวจคลินิกจิตยา อาการไม่ดีขึ้น
- 1 วัน สังเกตว่าพูดงๆ ซ้ำลงกว่าเดิม บางครั้งไม่ค่อยรู้เรื่อง แต่ขยับแขนขาพอๆ กัน 2 ข้าง พอลุกเดินเข้าห้องน้ำได้ ไม่เหนื่อย ไข้อยู่ ปวดท้องบริเวณท้องน้อย ไม่มีท้องเสียถ่ายเหลว ไม่มีคลื่นไส้อาเจียน



# ตรวจร่างกาย

- **Good consciousness with, agitation, not pale, no jaundice**
- **Heart lungs:** ปกติ
- **Abdomen:** soft, mild tender at mid lower abdomen
- **Ext.:** no edema
- **Neuro:** mild agitation, follow to one step command, disorientation to place

# CBC แรกครับ

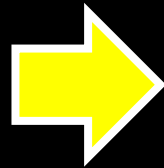
**HB 13.9 gm%**

**WBC 3210/mm<sup>3</sup>**

**Platelets 117,000/mm<sup>3</sup>**

# Platelets count แรกครับ

117000



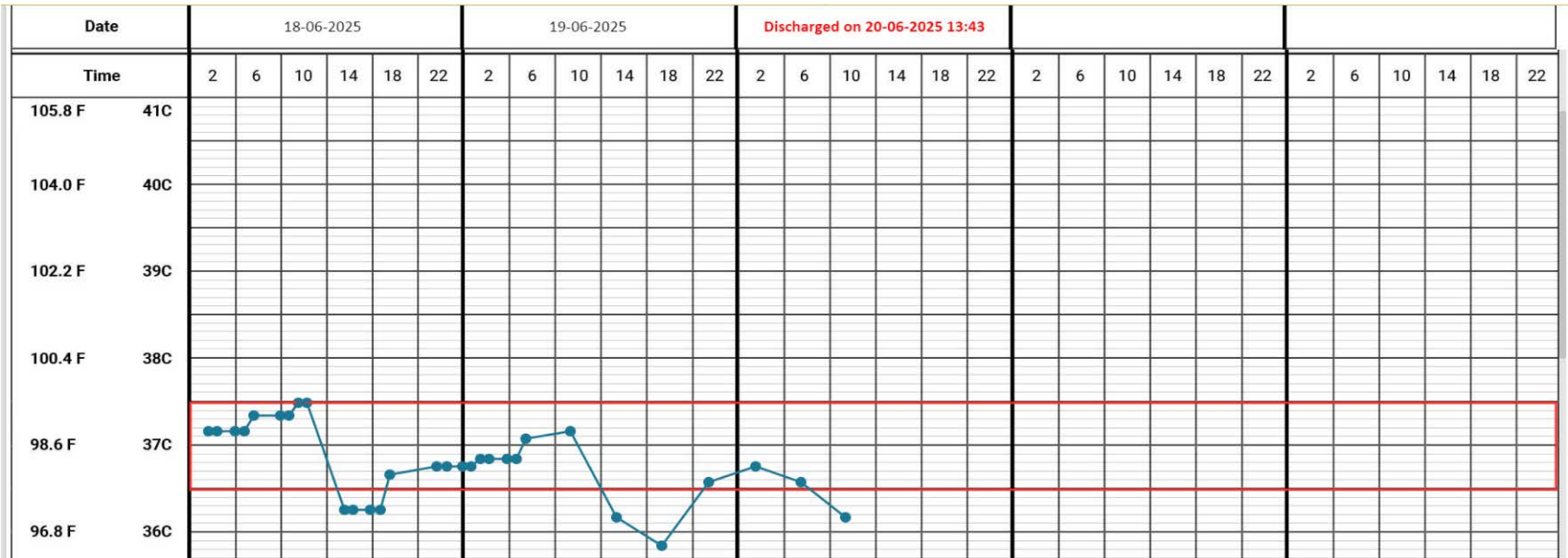
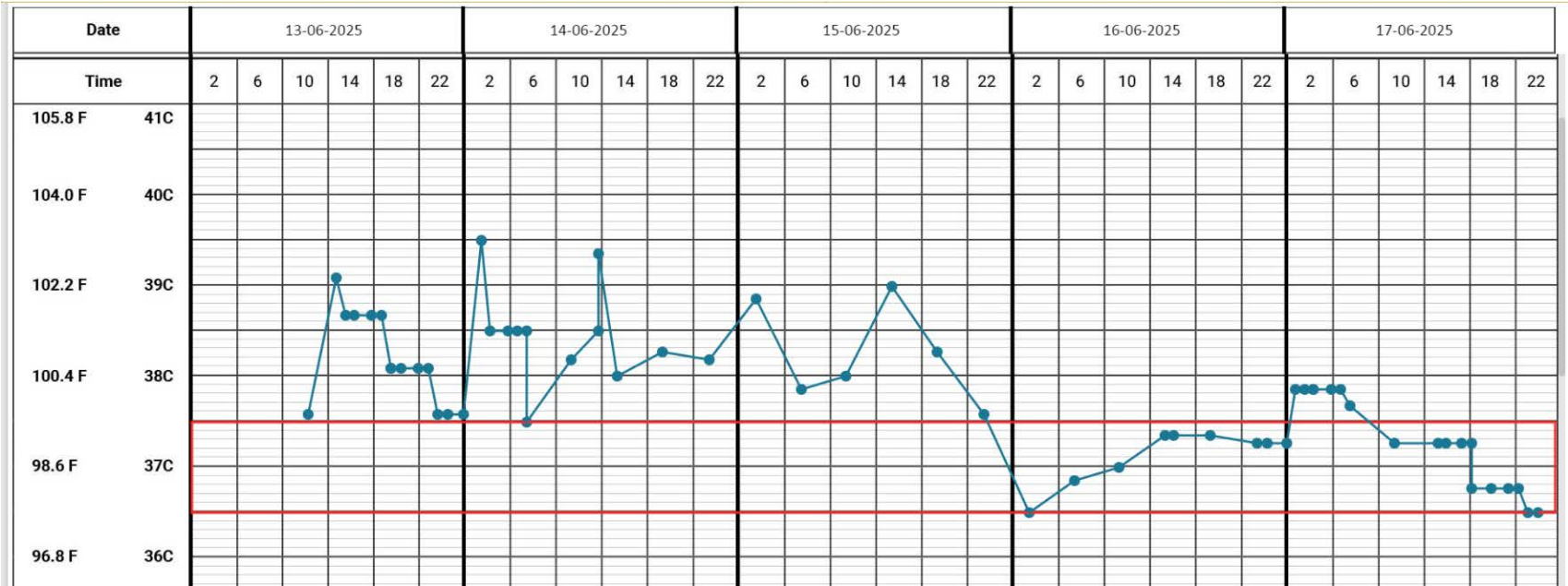
38000 cumm

(13-6-68)

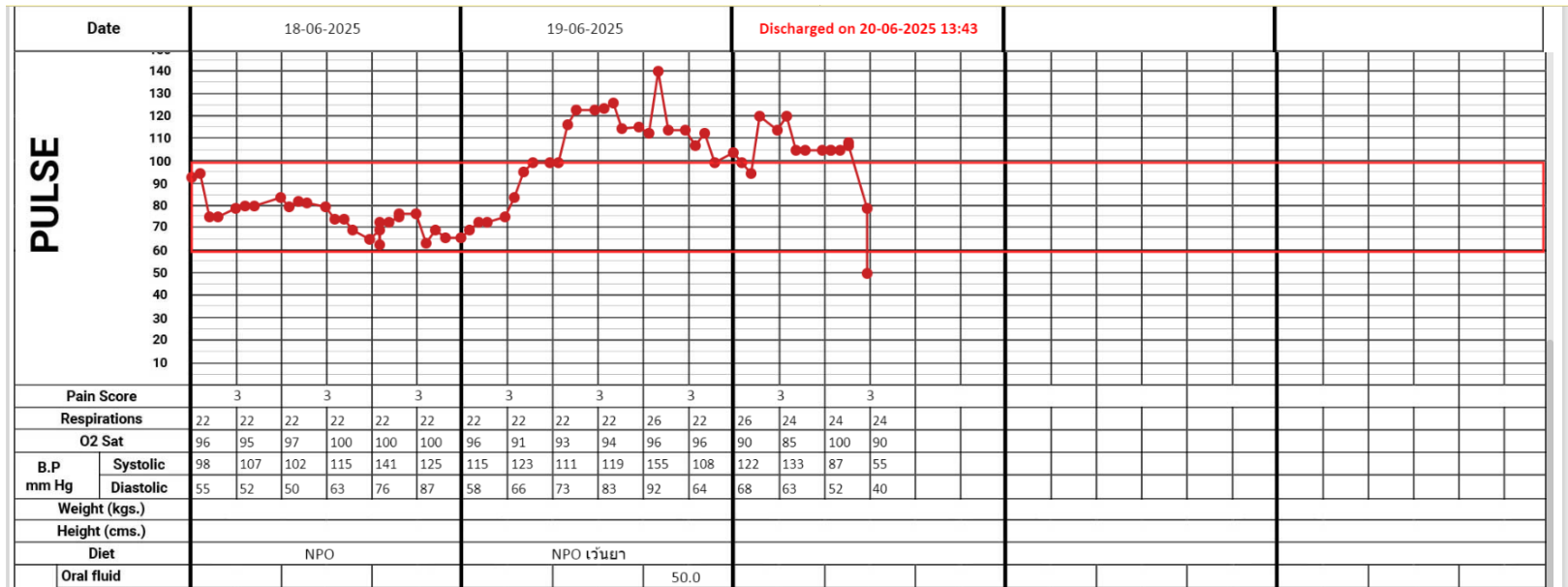
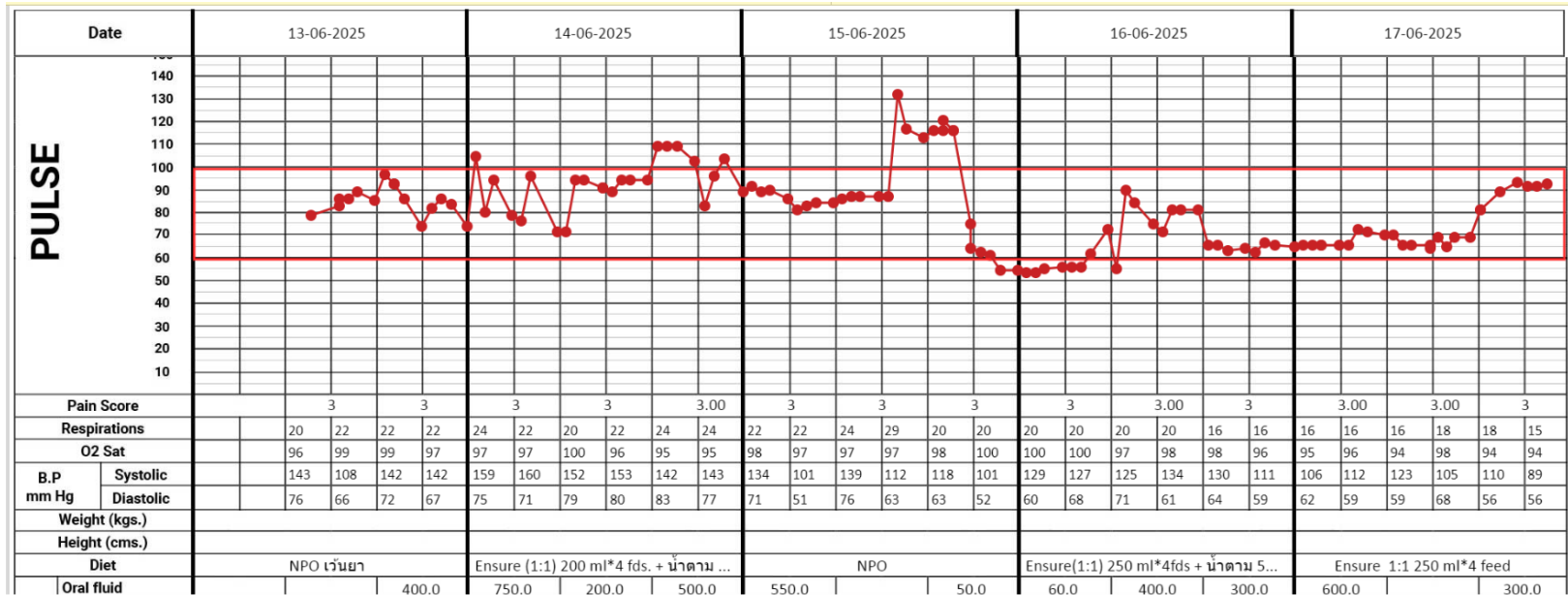


(18-6-68)

# Temperature



# Pulse rate



# Case 2 Severe Fever with Thrombocytopenic Syndrome

- ชายไทย 56 ปี Hypertension and BPH อาศัยอยู่วังโป่ง จังหวัดเพชรบูรณ์ He lived in the rural area in the farmland and exposure to stray and pet dogs.
- He was referred from Wang Pong Hospital. He had fever, dyspnea, cough, and malaise for 1 day and referred from district hospital. 1 day after admission, the patient had high fever and worsening respiratory distress despite mechanical ventilation.
- He had a dark lesion on the left foot.
- He had closed contact with the first case with confirmed SFTS case on 9/6/2568.



# ตรวจร่างกาย

- Vital Signs: BT: 38.1 °C PR: 148/min RR: 32/min BP: 74/55 mmHg SpO<sub>2</sub>: 100%
- BT 38.1 °C PR 148 /min. RR 32 /min. BP = 74/55 SpO<sub>2</sub> 100 %
- drowsiness on sedation, mild agitation, not follow to command.

# CBC แรกครับ

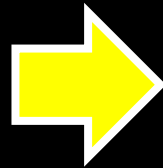
**HB 16.4 gm%**

**WBC 8210/mm<sup>3</sup>**

**Platelets 203,000/mm<sup>3</sup>**

# Platelets count แรกจับ

203,000



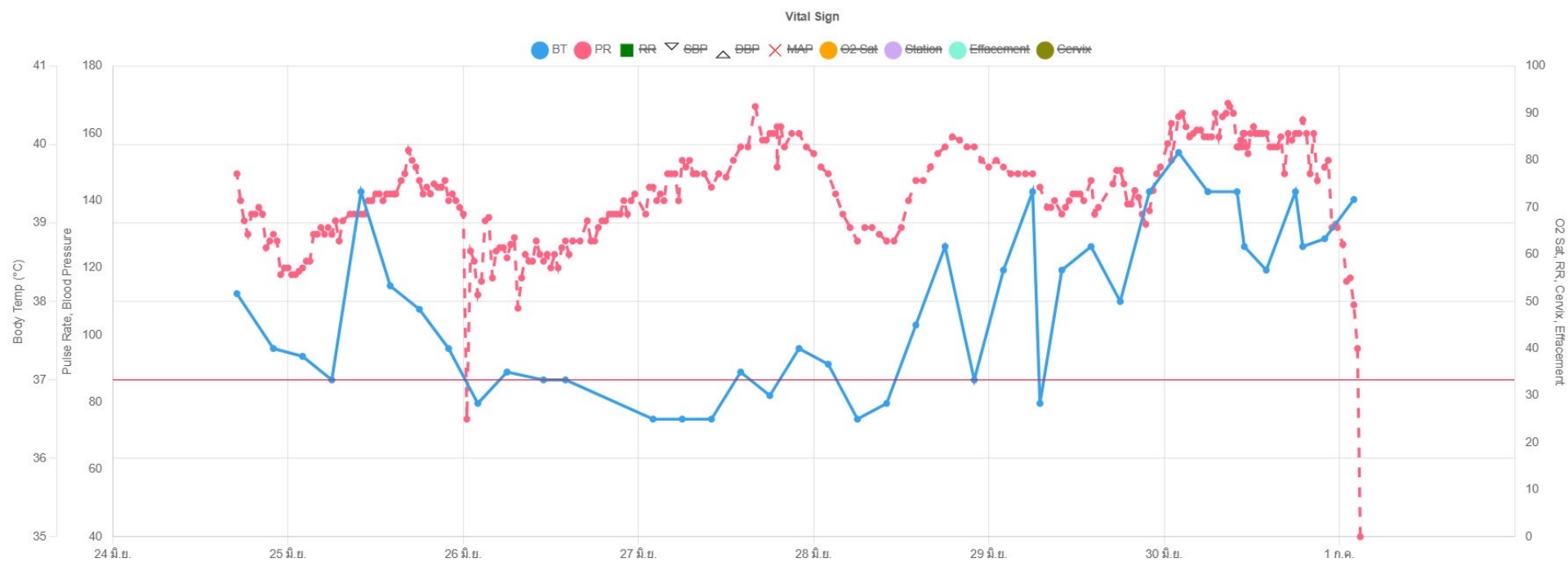
93,000 *cumm*

(13-6-68)



(18-6-68) *cumm*





# Characteristics of specimens positive for SFTSV, Thailand

Source	Age	Sex	Day of illness	Specimen types	qRT-PCR (Ct value)
Human, Case 1	82	Male	5	CSF	34.18
			7	Plasma	26.29
Human, Case 2	56	Male	3	Plasma	29.45
			7	Urine	32.25
			7	Stool	28.59
			7	Nasopharyngeal swab	30.00
			7	Throat swab	28.98



# บ้านหลังที่ 1





# บ้านหลังที่ 2









# From house of patient case-2



เห็บสุนัข

# *Rhipicephalus sanguineus*

Nymph



# *Rhipicephalus sanguineus*

Adult male





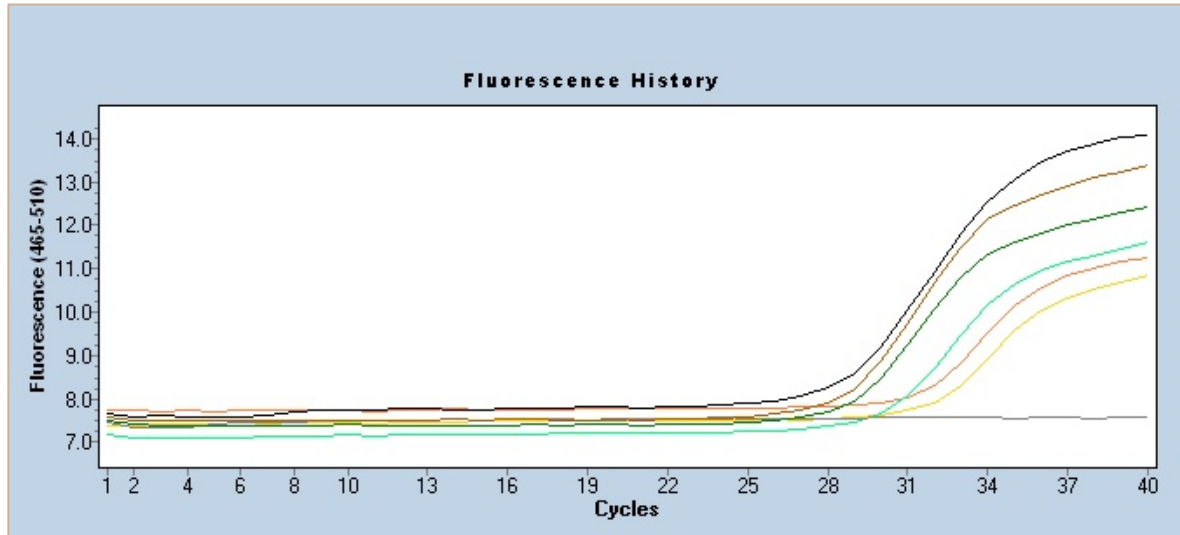
# *Rhipicephalus sanguineus*

Adult female





# Real-time RT-PCR Results for SFTSV Detection



- Negative control
- Positive control for SFTSV (Ct = 27.56)
- Sample ID: PHU12, urine collected at 30/06/2025 (Ct=32.25)
- Sample ID: PHU12.2, sediment (centrifuged) (Ct=31.91)
- Sample ID: PHN12, NP collected at 30/06/2025 (Ct=30.00)
- Sample ID: PHT12, throat swab collected at 30/06/2025 (Ct=28.98)
- Sample ID: PHF12, stool collected at 30/06/2025 (Ct=28.59)

## Case No.2

นาย สมชาย ชุนพล

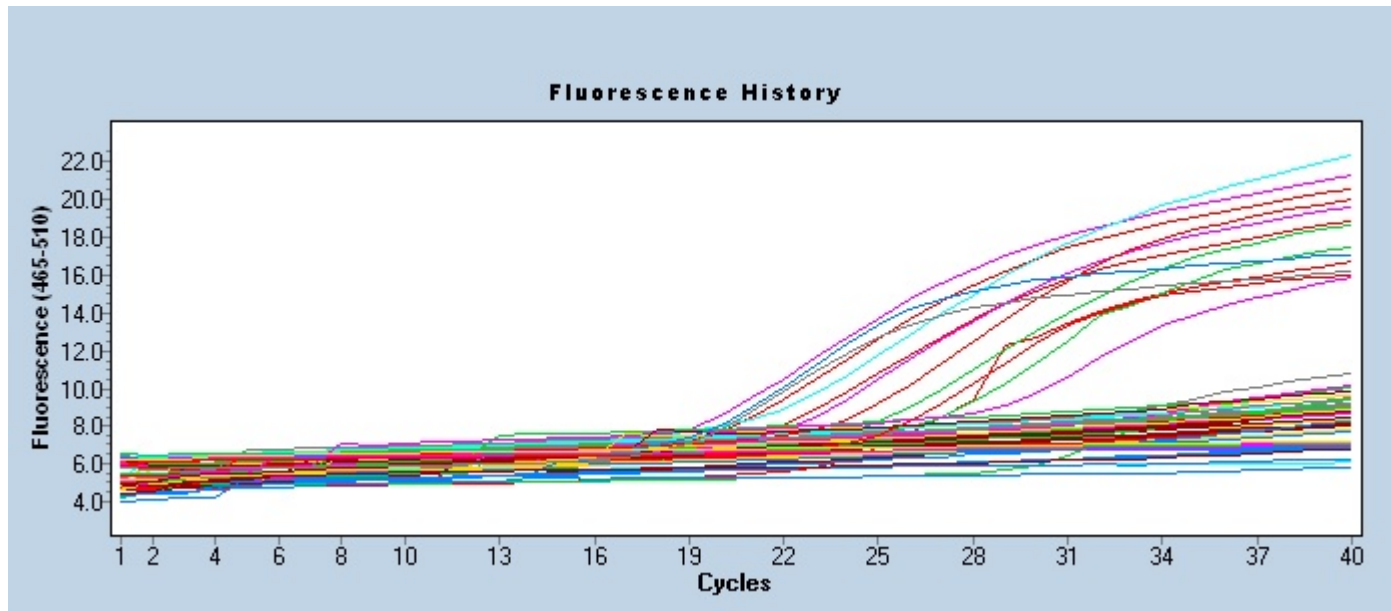
1) SFTSV RNA was detected in all sample types collected from the patient, including urine, nasopharyngeal swab, throat swab, and stool. 2) The lowest Ct values, indicating the highest viral load, were observed in the stool and throat swab specimens. 3) Centrifugation of urine slightly improved detection sensitivity compared to unprocessed urine. 4) These findings suggest that collecting multiple specimen types can enhance diagnostic accuracy in SFTSV infection.

# Partial S segment of the SFTSV



Partial S segment (477 bp) of SFTSV was successfully amplified from urine, nasopharyngeal swab, throat swab, and stool samples of the patient. This demonstrates that SFTSV RNA can be detected in multiple specimen types, supporting molecular confirmation across different clinical materials.

# Real-time RT-PCR Results for SFTSV Detection



## Partial S segment of the SFTSV

*Rhipicephalus sanguineus*

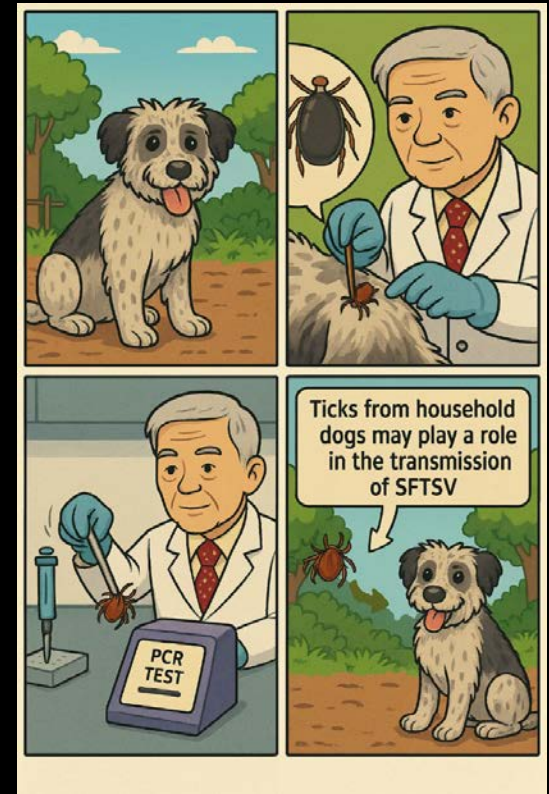
12525 12526 12529 12530 12532 12533 12534 12535 12536 12543 12546 12548 12557 NEG



← 477 bp

# Summary of SFTSV Detection in Ticks from Pet Dogs of the Second Fatal Case

- A total of 44 ticks were collected from 5 pet dogs belonging to the second patient who died from SFTSV infection in Thailand.
- 13 ticks tested positive for SFTSV RNA by qRT-PCR.
- These positive ticks were found on 4 dogs:
  - Satang สตางค์ (black and white) – 9 ticks
  - Makham มะขาม (white) – 2 ticks
  - Taew แท้ว (brown) – 1 tick
  - Ting ทิ้ง (brown) – 1 tick
- Life stages and sex of the SFTSV-positive ticks:
  - Adult females: 7 ticks
  - Adult males: 6 ticks
- The positivity rate was 29.5% (13 out of 44 ticks).





## Laboratory testing to determine the presence of neutralizing antibodies against SFTSV (HB29 strain) using the SFTSV RVP microneutralization assay

Host	No. of positive	Positive rate per species
Human (n=48)	8	16.67%
Dog/Canine (n=30)	11	36.67%
Cat/Feline (n=7)	0	0%
Cow/Bovine (n=2)	0	0%
Rodent (n=1)	1	100.00%

We tested 88 sera from close contacts and animals, all PCR-negative. Using the SFTSV microneutralization assay, 23% were positive overall, mainly dogs and one rodent, with some humans. This suggests the virus is already circulating locally, with dogs and rodents likely being part of its natural cycle.

# **SFTSV Detection in Animals and Ticks**

## **Phitsanulok and Phetchabun Provinces, Thailand**

### **Animal Samples (Dogs & Cats) – Phitsanulok**

Total tested: 139 samples (90 dogs, 49 cats)

PCR results:

3 dog samples positive for SFTSV RNA

- Bang Rakam District (Ct = 33.15, 35.39)

- Chat Trakan District (Ct = 37.23)

Serology:

13 sera tested for neutralizing antibodies (6 dogs, 7 cats)

2 dog sera positive for neutralizing antibodies against SFTSV

### **Tick Samples – Phitsanulok & Phetchabun**

Total ticks tested: 227

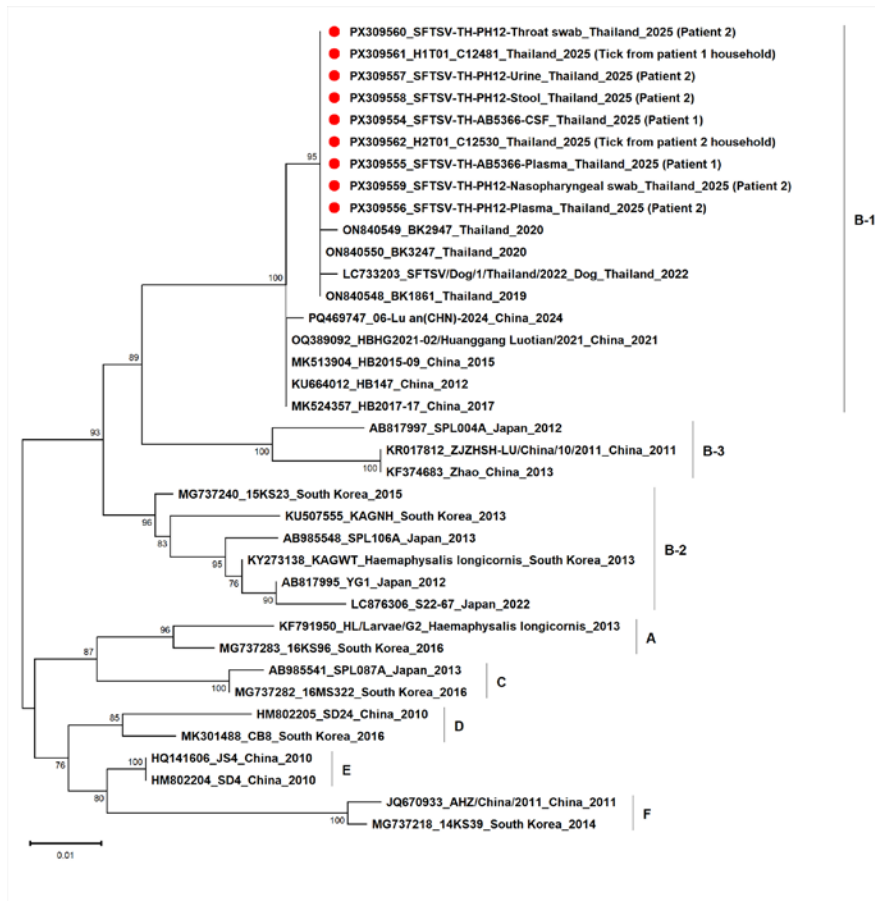
- Phitsanulok = 183

- Phetchabun = 44

PCR results:

No SFTSV RNA detected in any tick samples

# Maximum likelihood phylogenetic tree of SFTSV based on the partial S segment nucleotide sequences



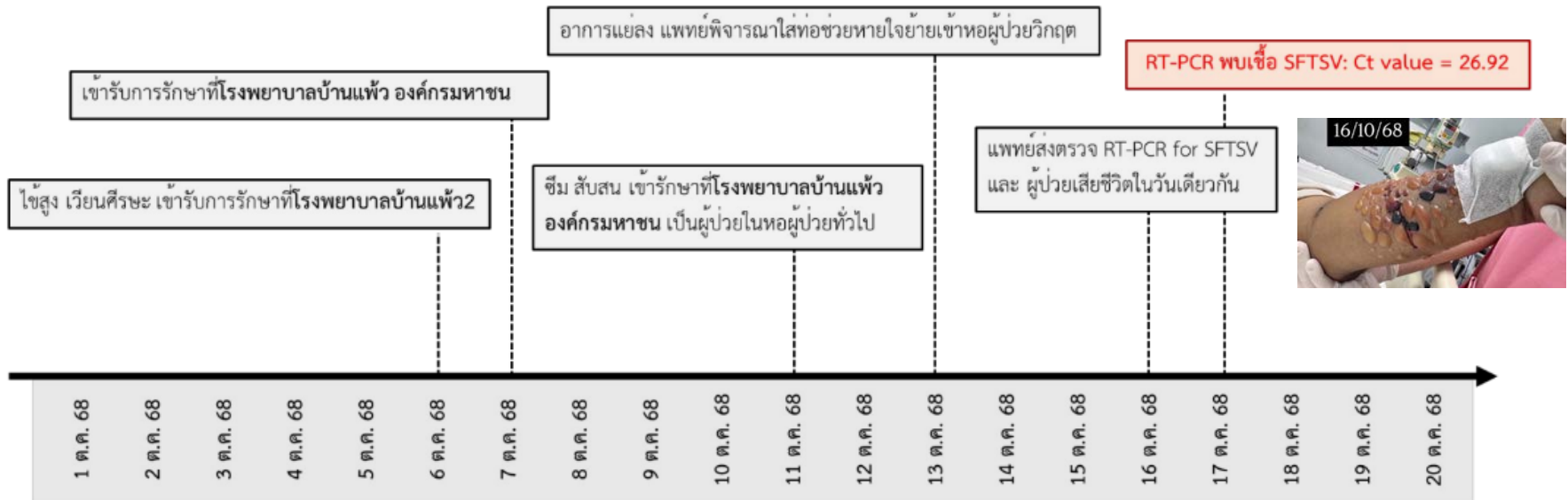
Phylogenetic analysis showed that all sequences obtained in this study (patients and representative tick samples) were identical and clustered within genotype B-1 together with previously reported Thai strains and closely related Chinese strains (2012–2024).

# **Outbreak of SFTS with Fatal Cases No. 3**

# Case 3: 54-year-old Male, Ban Phaeo, Samut Sakhon

## Timeline

ญาติให้ประวัติว่าผู้ป่วยมีอาการเบื่ออาหาร และถ่ายเหลวหลายครั้ง



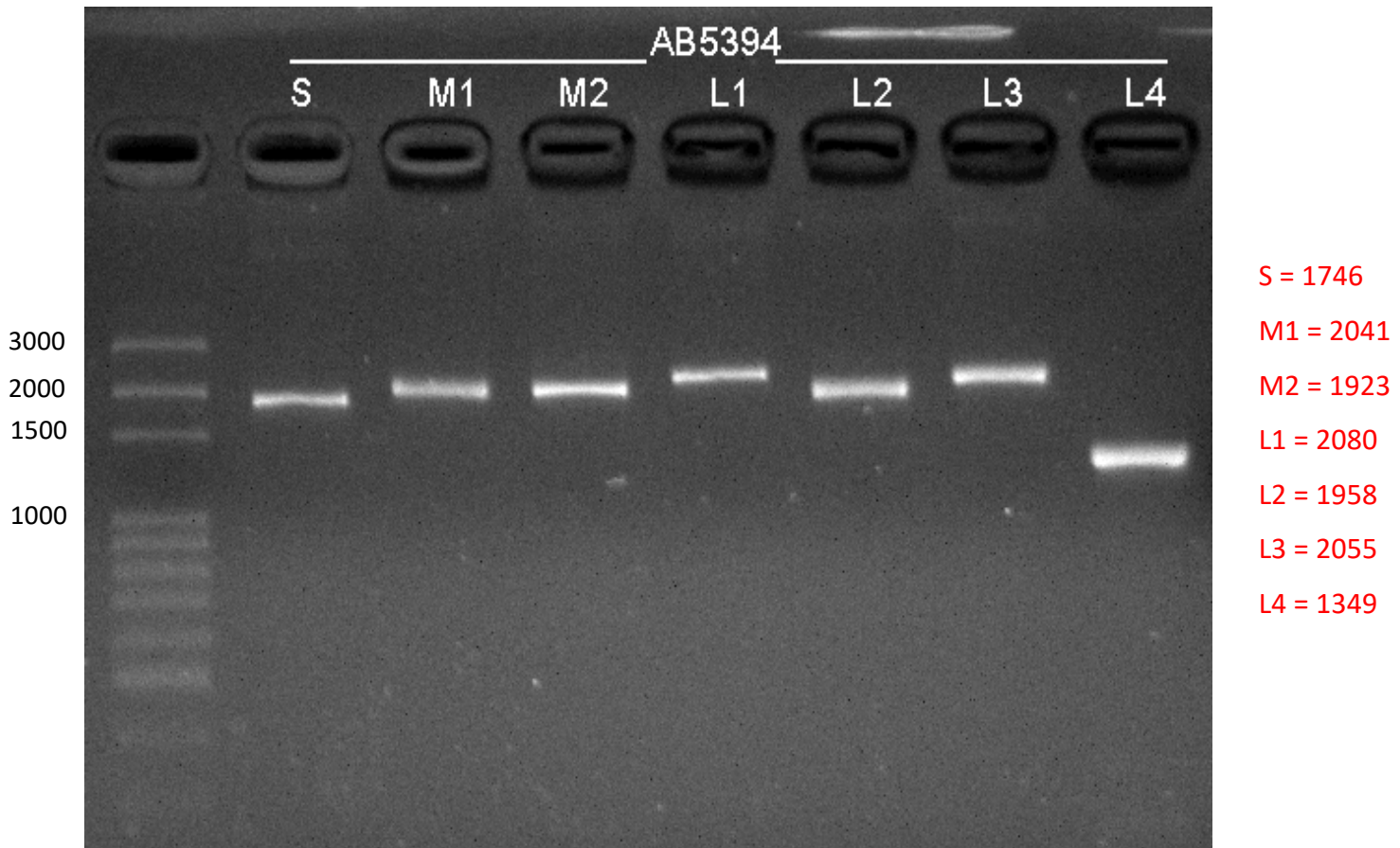
ที่มา: Slide คุณหมอแน sw.บ้านแพ้ว องค์การมหาชน



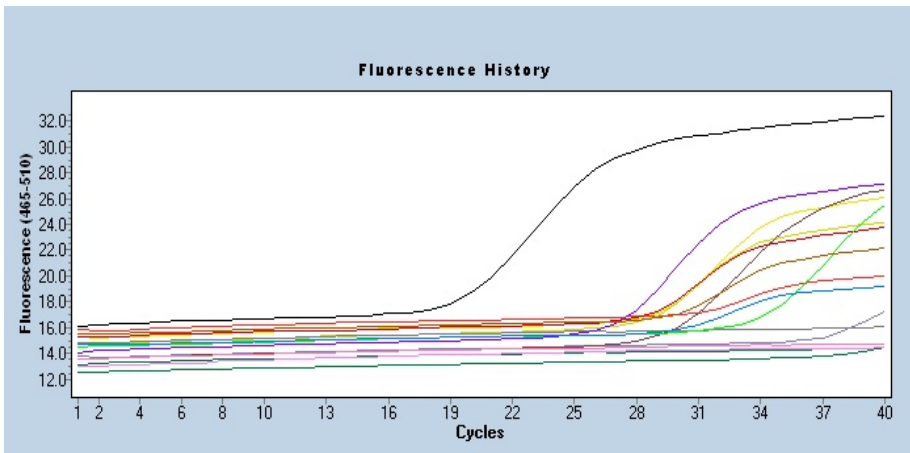




# RT-PCR Amplification of SFTSV Genome Segments (S, M, L) from Patient Sample



# Detection of SFTSV RNA in Multiple Clinical Specimens from Patient No. 3 (Real-Time RT-PCR and Conventional PCR)



**S1 = Urine (Ct = 29.31), collection date 13 Oct 2025**

**S2 = Urine sediment (Ct = 28.68)**

**S3 = CSF (Ct = 37.87), collection date 11 Oct 2025**

**S4 = Sputum (Ct = 29.47), collection date 14 Oct 2025**

**S5 = Hemoculture H665 (Not detected), collection date 15 Oct 2025**

**S6 = Hemoculture H666 (Not detected), collection date 15 Oct 2025**

**S7 = Pus for slide (Ct = 31.68), collection date 15 Oct 2025**

**S8 = Stool (Ct = 30.71), collection date 13 Oct 2025**

**S9 = Pus from rt ear 1 (Ct = 32.34), collection date 14 Oct 2025**

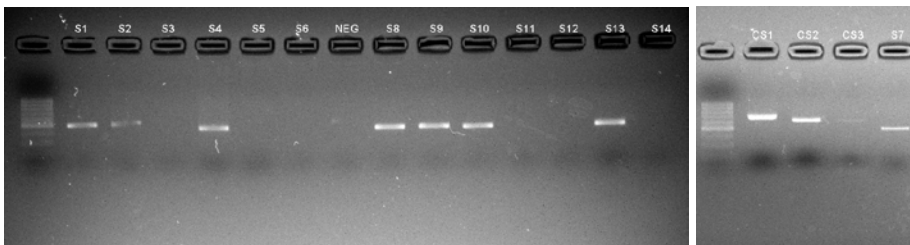
**S10 = Pus from rt ear 2 (Ct = 28.99), collection date 14 Oct 2025**

**S11 = Plasma (Heparin)(Not detected), collection date 14 Oct 2025**

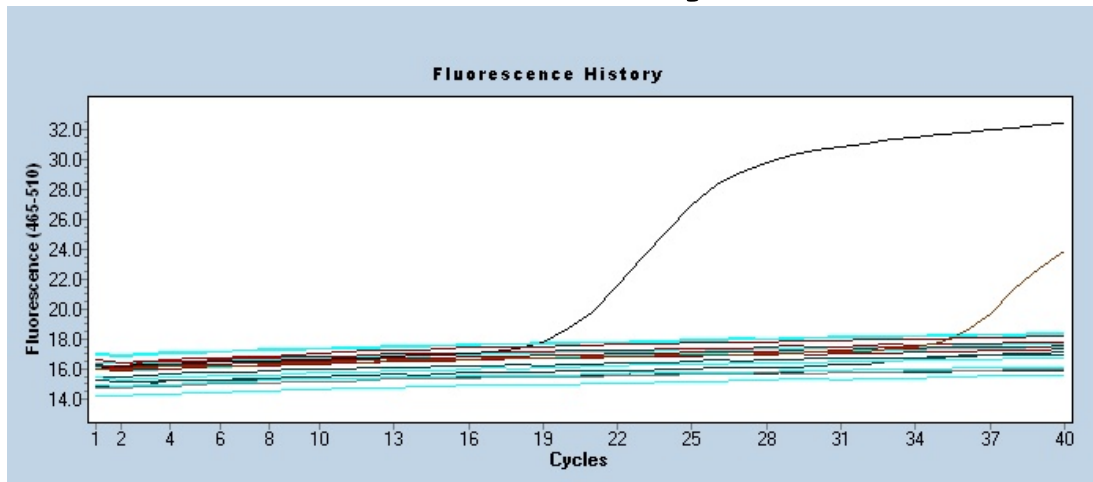
**S12 = Plasma (Heparin)(Not detected), collection date 15 Oct 2025**

**S13 = Plasma (EDTA)(Ct = 26.79), collection date 16 Oct 2025**

**S14 = Plasma (Heparin)(Ct = 33.64), collection date 16 Oct 2025**

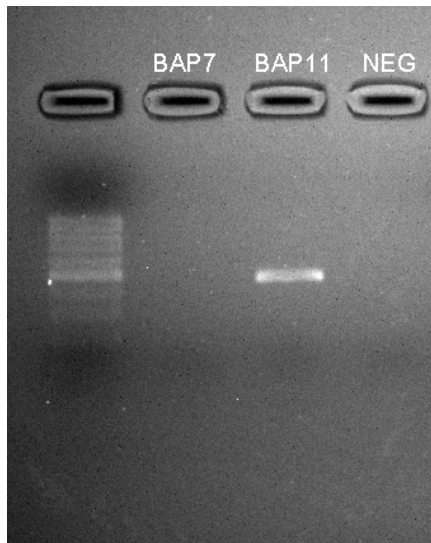


# Screening of Dog and Cat Samples for SFTSV RNA by RT-PCR



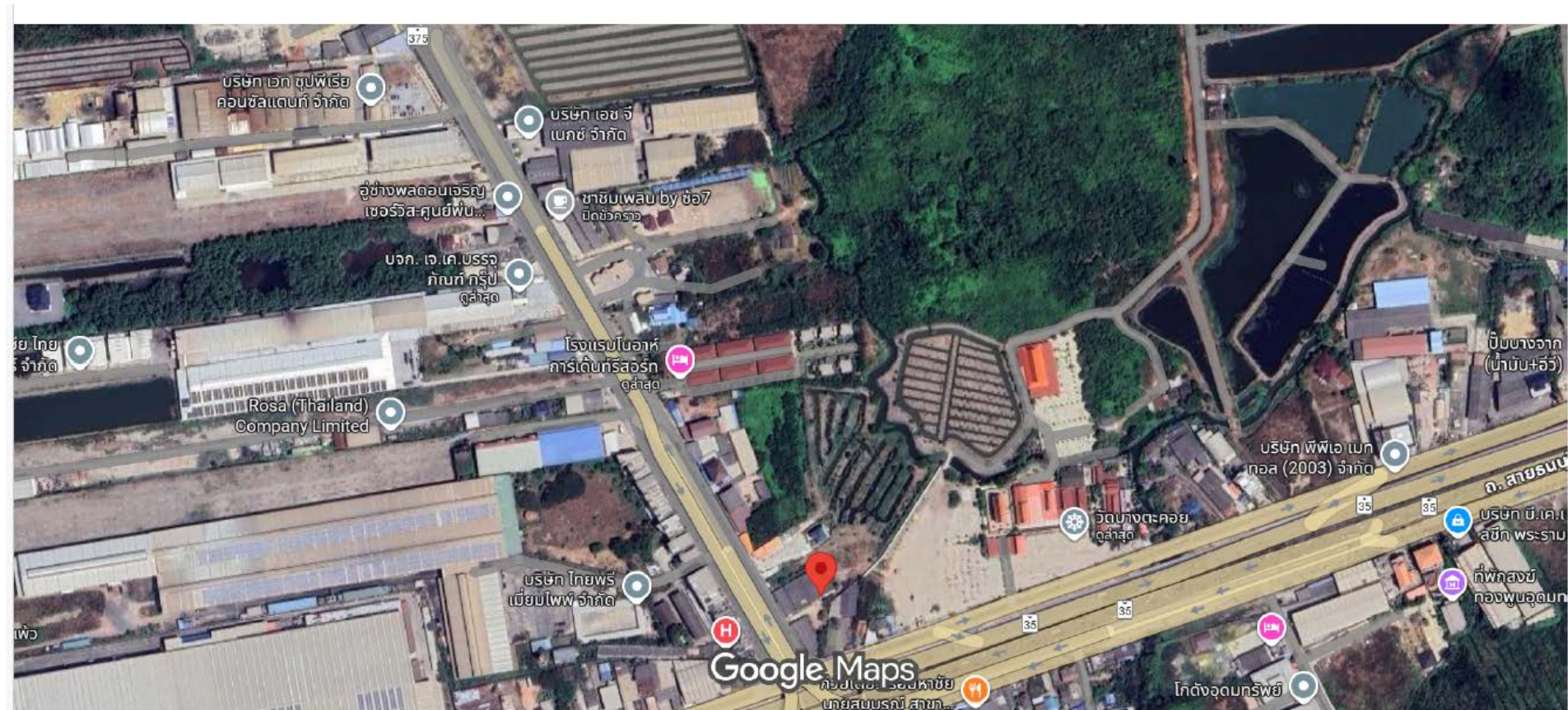
Dog No. 11 (Ct = 35.46)

น้องนาง



A total of 13 animal samples, including eight dogs and five cats, were tested for the presence of SFTSV RNA using real-time RT-PCR and confirmed by conventional PCR. One dog sample (Dog No. 11) tested positive





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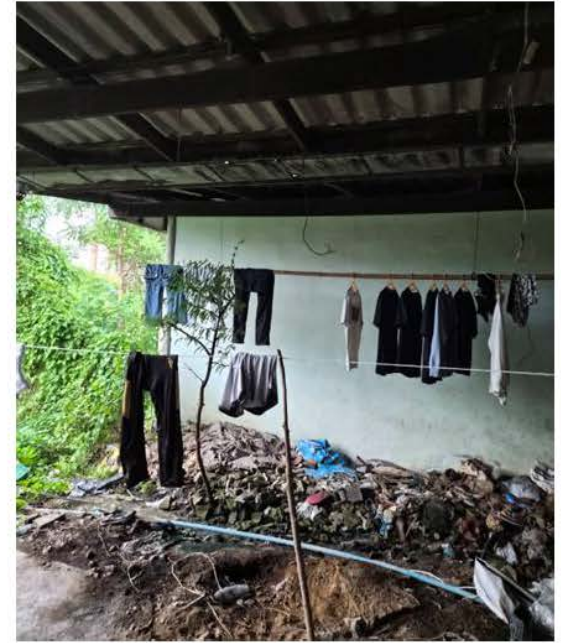
# การศึกษาสิ่งแวดล้อมที่พักอาศัยของผู้ป่วย



ช่องทางเดินระหว่างห้อง



พื้นที่รกร้างด้านหลัง



บริเวณตากผ้า

# การศึกษาสิ่งแวดล้อมที่พหุอาศัยของผู้ป่วย



บ้านดาวรุ่ง/  
เก่ง<sup>B</sup>



บ้านสมควร<sup>B</sup>

ที่ตาก ผ้า	16	15	14	13	12	11	พื้นที่ รกร้าง
3			6	9 <sup>A</sup>	10	บ้าน คุณ ยาย	
2			5	8	พื้นที่ โล่ง		
1 <sup>B</sup>			4	7 <sup>C</sup>			

A = ห้องผู้ป่วยชีวิต  
B = มีประวัติดื่มสุราด้วยกัน + มีโอกาสสัมผัสสุนัขสูง  
C = มีประวัติโดยเห็นกัก

# การศึกษาสิ่งแวดล้อมที่พักอาศัยของผู้ป่วย



ห้องผู้ป่วย



จุดที่มีเห็บปริมาณมาก



เห็บที่เก็บตัวอย่าง









# การศึกษาวัดที่ดำเนินการพิธีศพ

- ลูกสาวผู้เสียชีวิตนำสุนัขมาขังไว้ในกรงใกล้กับศาลาทำพิธีศพ
- พบสุนัขและแมวไม่มีเจ้าของจำนวนมาก โดยเฉพาะบริเวณกุฏิ



## Sample collection sheet SFTSV in Dog and Cat ณ วัด ตรงบริเวณศาลา



สมศรี (อยู่ใกล้ชิดกับมุกตลอดเวลา)

# Results Summary: SFTSV Detection in Humans and Animals (Ban Phaeo, Samut Sakhon)

## Human Samples:

- **Total tested:** 84 serum samples
- **Result:** All samples were **negative (Not detected)** by real-time RT-PCR
- **Interpretation:** No evidence of active SFTSV infection among human participants

## Animal Samples:

- **Total tested:** 13 samples
  - Dogs: 8
  - Cats: 5
- **Result:** **One dog sample** tested **positive** (Ct = 35.46)
- **All other samples:** Negative
- **Confirmation:** Positive dog confirmed by conventional PCR and gel electrophoresis

# Sample collection sheet SFTSV in Dog and Cat ณ บ้านพักผู้ป่วย



ถุงทอง



บี้ (สุนัขจร)



บุญรอด



ไทเกอร์



Tick จากในห้องผู้ป่วย



# Sample collection sheet SFTSV in Dog and Cat ณ วัด ตรังบริเวณศาลา (หน้าวัด) มีสุนัขอยู่บริเวณนี้ประมาณ 6 ตัว



# Sample collection sheet SFTSV in Dog and Cat

## ณ วัด ตระบริเวณกุฎี มีสุนัขและแมวอยู่รวมกัน



เด็กใหม่



น้องนาง



# Sample collection sheet SFTSV in Dog and Cat

ณ ร้านใกล้โรงงาน (บริเวณที่ผู้ป่วยมาทานอาหาร)



ส้ม



ก้อยจ๊ะ



ไฟ

# สิ่งที่ดำเนินการไปแล้ว

## คน

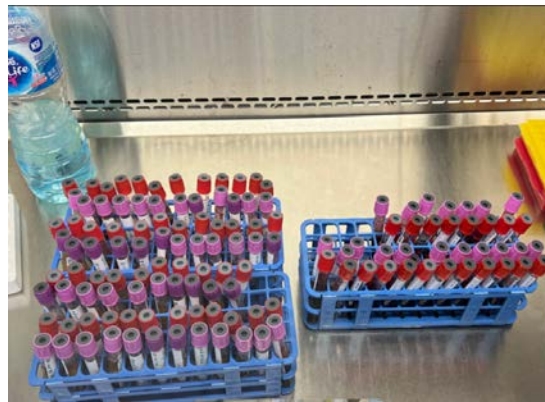
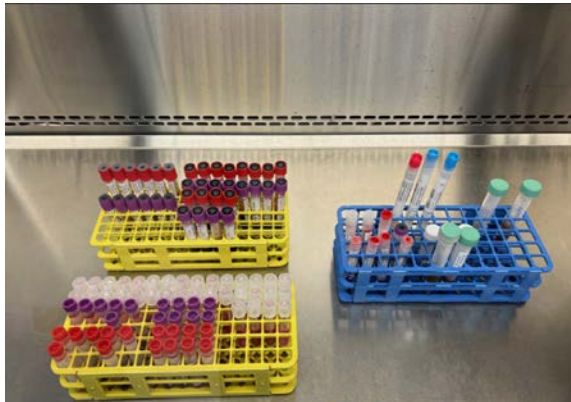
- ดำเนินการเก็บตัวอย่างเลือดผู้สัมผัสใกล้ชิดและผู้ป่วยตามนิยามส่งตรวจทางห้องปฏิบัติการ
- ติดตามผู้สัมผัสใกล้ชิดและผู้ป่วยเพิ่มเติมในชุมชน และสถานพยาบาล

## สัตว์

- ดำเนินการเก็บตัวอย่างเลือดสุนัขส่งตรวจทางห้องปฏิบัติการ
- ฉีดยากำจัดเห็บหมัด (Ivermectin) ให้กับสุนัขในบริเวณชุมชน

## สิ่งแวดล้อม

- เก็บตัวอย่างเห็บจากห้องเช่าผู้เสียชีวิตและในสุนัขบริเวณชุมชน
- ฉีดพ่นสารเคมีกำจัดแมลง (Cypermethrin) ตามสิ่งแวดล้อมภายในห้องของผู้เสียชีวิต และบริเวณรอบห้องเช่า











# Characteristics of specimens positive for SFTSV, Thailand

A

Source	Age	Sex	Collect date Day after the onset of fever	Specimen types	qRT-PCR (Ct value)	RT-PCR	Isolate	GenBank accession no.
Patient 2	54	Male	12	Plasma	26.79	+ve	SFTSV-TH	XXXXXXXXXXXX
			9	Urine	29.31	+ve	SFTSV-TH	XXXXXXXXXXXX
			7	CSF	37.87	Wkly.+ve	-	
			10	Sputum	29.47	+ve	SFTSV-TH	XXXXXXXXXXXX
			10	Pus (Rt. ear)	28.99	+ve	SFTSV-TH	XXXXXXXXXXXX
			9	Stool	30.71	+ve	SFTSV-TH	XXXXXXXXXXXX
			11	Vascular fluid	31.68	+ve	SFTSV-TH	XXXXXXXXXXXX

B

Sample Source	No. Tested	Positive n (%)	Remarks	Specimen types	qRT-PCR (Ct value)	RT-PCR	Isolate	GenBank accession no.
Dogs	18	1 (5.6)	+ve in temple dog	Serum	-	+ve	SFTSV-TH	XXXXXXXXXXXX
Cats	5	0	All -ve	Serum	-	-	-	
Ticks	268	6 (2.24)	Adult male (2) Adult female (4)	No.1 (neighboring dog) No. 2 (patient's room) Patient's room	19.7	-ve	SFTSV-TH	XXXXXXXXXXXX
					18.64	+ve	SFTSV-TH	XXXXXXXXXXXX
					18.23	+ve	SFTSV-TH	XXXXXXXXXXXX
					33.3	+ve	SFTSV-TH	XXXXXXXXXXXX
					19.39	+ve	SFTSV-TH	XXXXXXXXXXXX
					33.19	+ve	SFTSV-TH	XXXXXXXXXXXX



# Summary

## Human infections

Confirmed cases, including fatalities. Viral RNA detected in multiple specimens (plasma, CSF, urine, stool, respiratory), showing systemic spread and the need for testing beyond plasma.

## Animal reservoirs

Dogs show notable seroprevalence with RNA-positive samples; wild rodents also harbor SFTSV.

## Vector ecology

SFTSV detected in dog-associated *Rhipicephalus sanguineus* ticks and chigger mites; tick sequences identical to human strains, indicating a complex zoonotic cycle.

## Public health implications

Human cases, animal reservoirs, and competent vectors confirm SFTSV as an emerging threat in Thailand, requiring clinician awareness, multiplex diagnostics, and One Health surveillance.

เพราะงานวิจัย

เด็กเกิดวันนี้จะมีอายุ 120 ปี

# Acknowledgement

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