









# Lumpy Skin Disease in China

Jinming LI CAHEC

### **Outline**



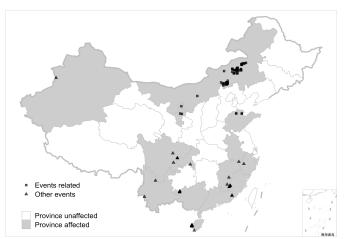






- Cattle industry in China
- LSD Reported outbreaks
- Control measures





### 农业农村部文件

农牧发[2020]30号

#### 农业农村部关于印发《牛结节性皮肤病防治 技术规范》的通知

各省、自治区、直辖市及计划单列市农业农村(农牧、畜牧兽医)厅(局、委),新疆生产建设兵团农业农村局:

为做好牛结节性皮肤病防控工作,保障养牛业持续健康发展, 我部组织制定了《牛结节性皮肤病防治技术规范》。现印发给你们,请遵照执行。





# 1. Cattle industry in China



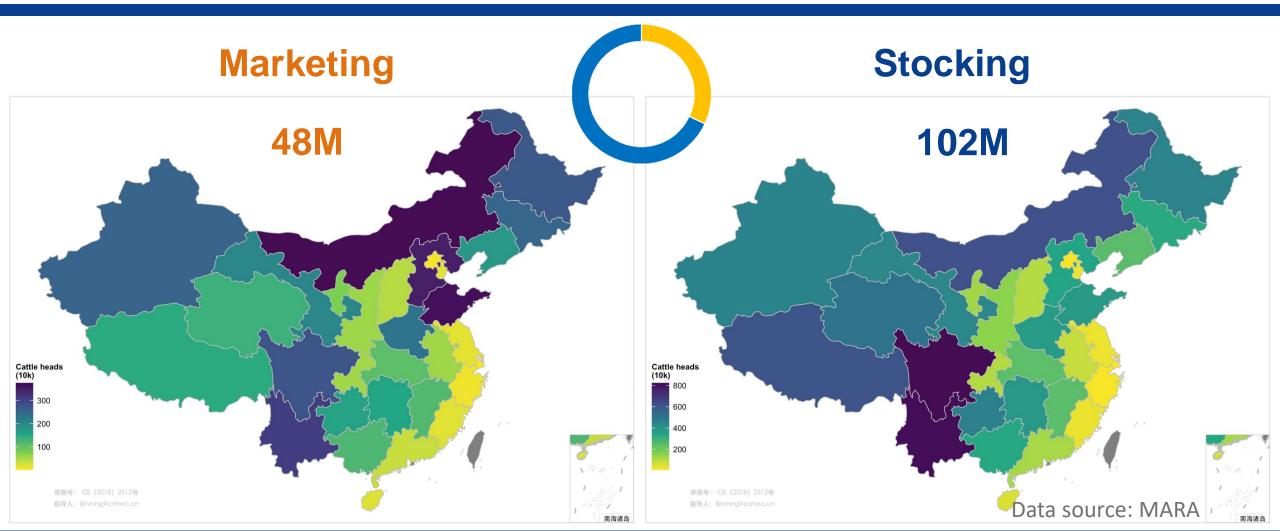
### **Cattle Industry in China**













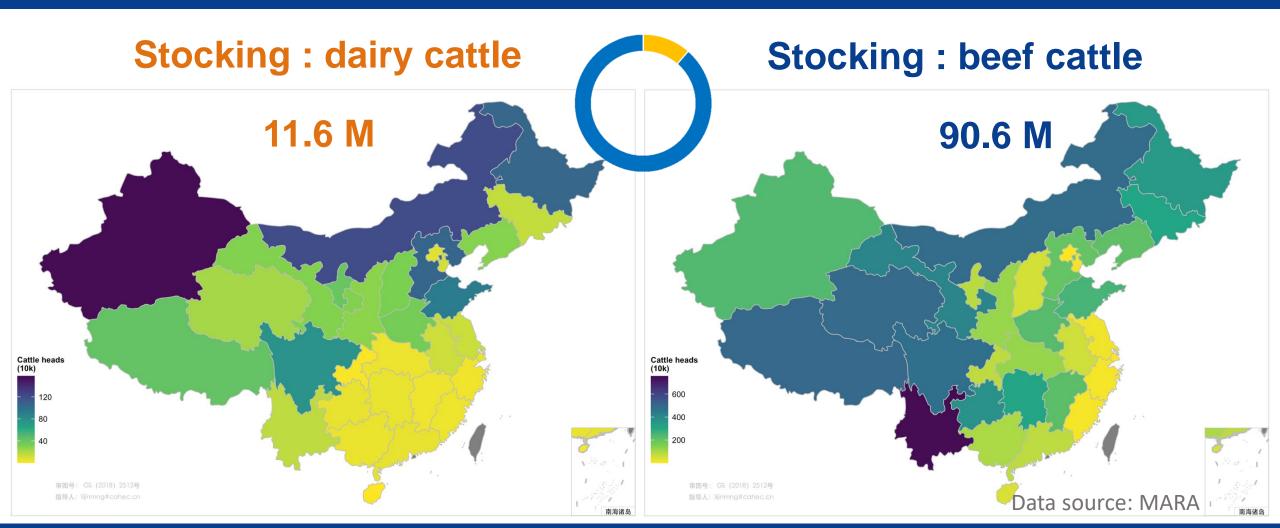
### **Cattle Industry in China**













### **Cattle Industry in China**









1~9	50~99	100~499
42.10%	13.40%	12.40%
10~49		
23.10%	1000~	500~999
	4.90%	4.10%

**7.8 M** farms

**90%** 1-9



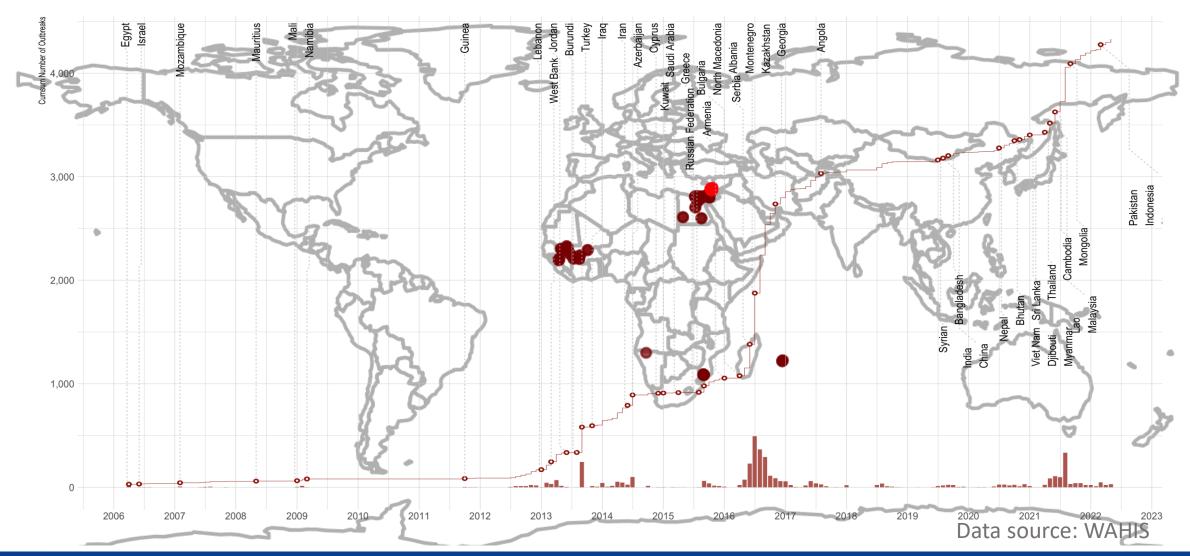
Data source: MARA





### **World Map of Lumpy skin disease**

Jan 2013







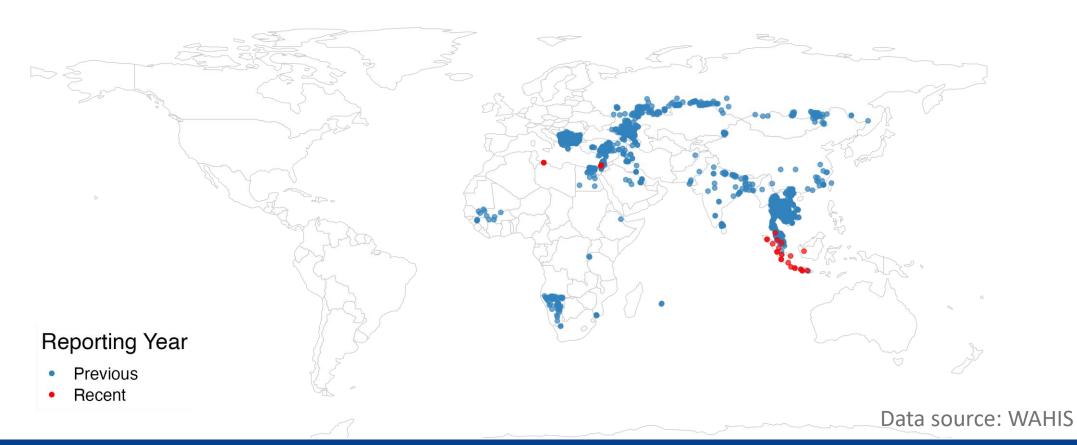






### World Map of Lumpy skin disease

as of 24 Jul, 2023







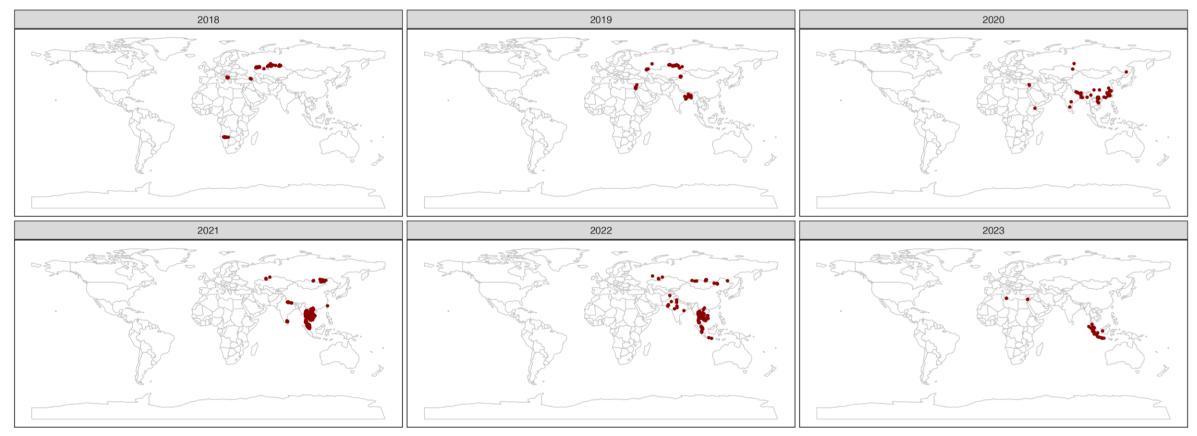




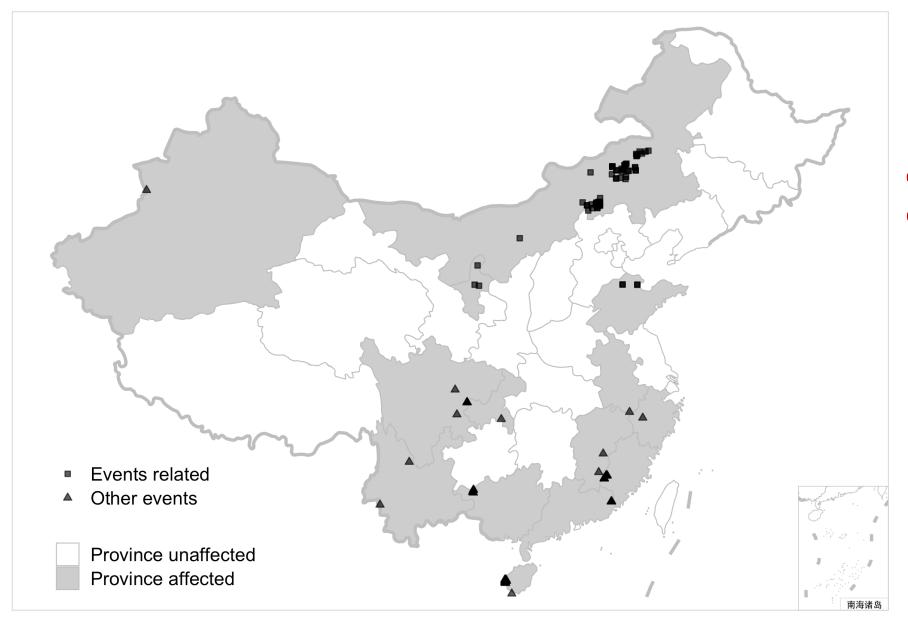


### **New Outbreaks of Lumpy Skin Disease (WAHIS)**

2018 ~ 2023















14 Provinces Confirmed

Aug, 2019, Xjinjiang, 1st

Summer, 2020, sporadic in Fujian, Jiangxi, Guangong, Anhui, Zhejiang, Yunnan, Guangxi, Chongqiong, Sichuan

Nov, 2020, 3 events related with the same quarantine station in Jiangsu

Dec, 2020, Hainan

Data source: MARA





































畜牧兽医局

请输入关键字

搜索

机构职能

工作动态

政策解读

政务服务

疫情发布

监测预警

国际合作

当前位置: 首页 > 机构 > 畜牧兽医局 > 疫情发布

#### 农业农村部畜牧兽医局关于浙江发生牛结节性皮肤病疫情有关情况的通报

日期: 2020-07-15

作者:

来源:农业农村部畜牧兽医局

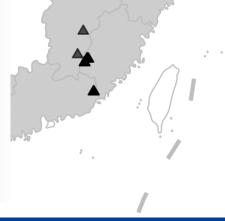
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🗇 打印本页

7月13日,农业农村部接到中国动物疫病预防控制中心报告,经中国动物卫生与流行病学中心确认,浙江省金华市婺城区发生一起牛结节性皮肤病疫情。该批牛共26头,发病3头,死亡1头。疫情发生后,当地按照有关预案和防治技术规范要求,切实做好疫情处置工作,已扑杀2头发病牛且无害化处理,并进行全面彻底消毒。截至目前,今年以来,全国共有福建长汀、江西赣州、广东潮州、安徽黄山、浙江金华等地确诊发生牛结节性皮肤病疫情(汇总表附后)。请各地按照我部前期通知要求,切实做好牛结节性皮肤病防控工作。

附件

2020年牛结节性皮肤病疫情统计表















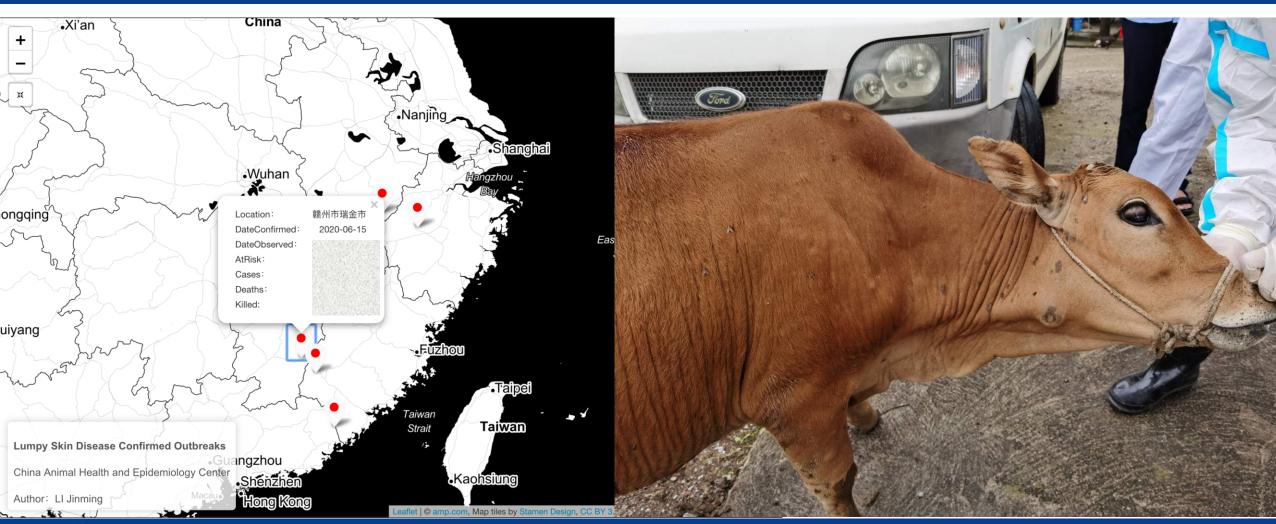










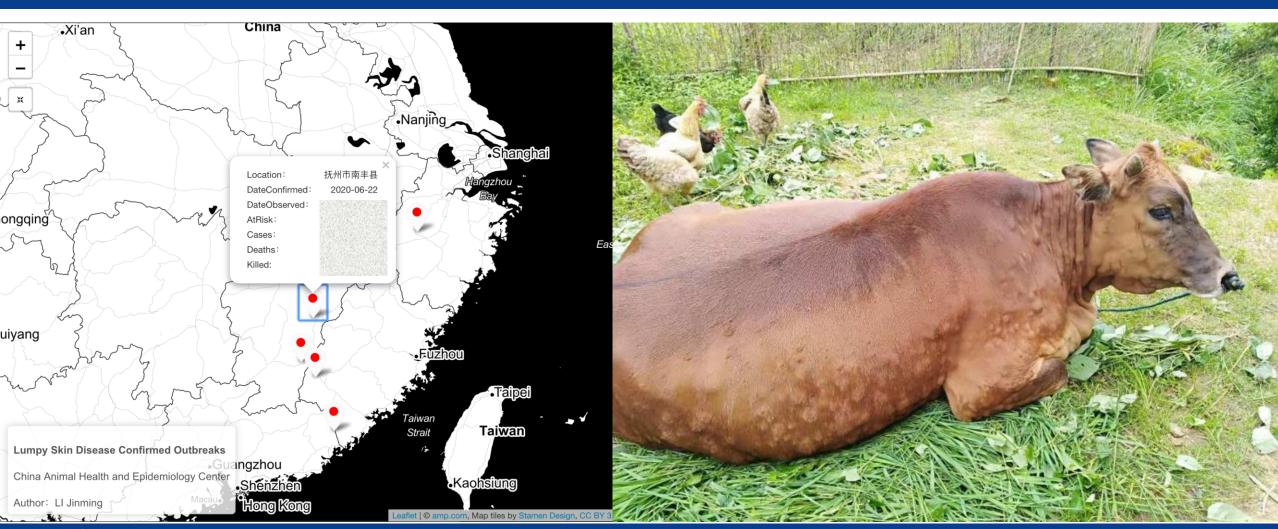






















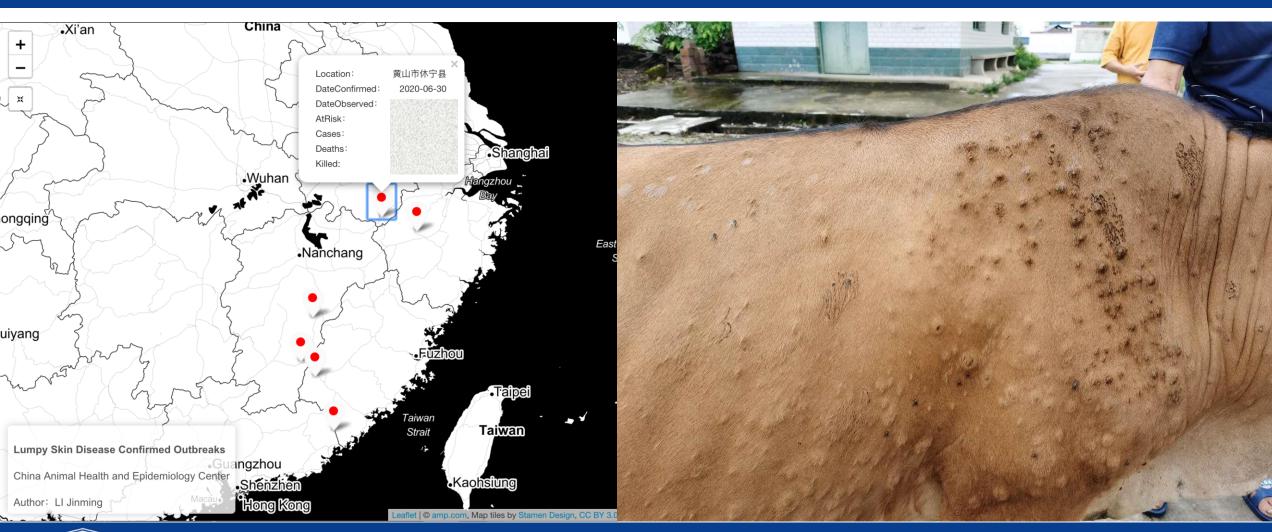












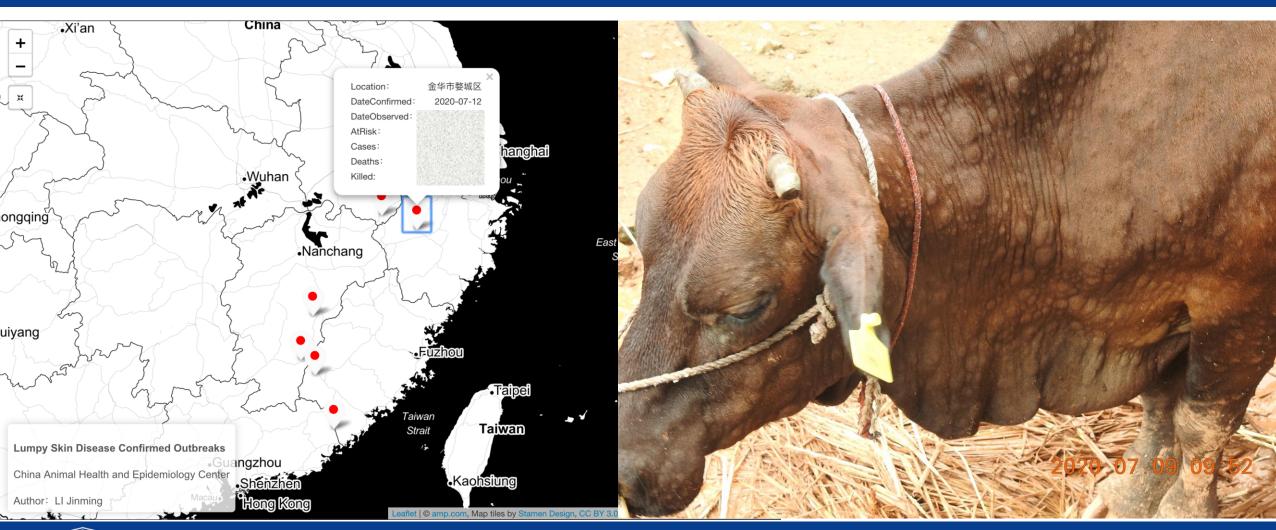












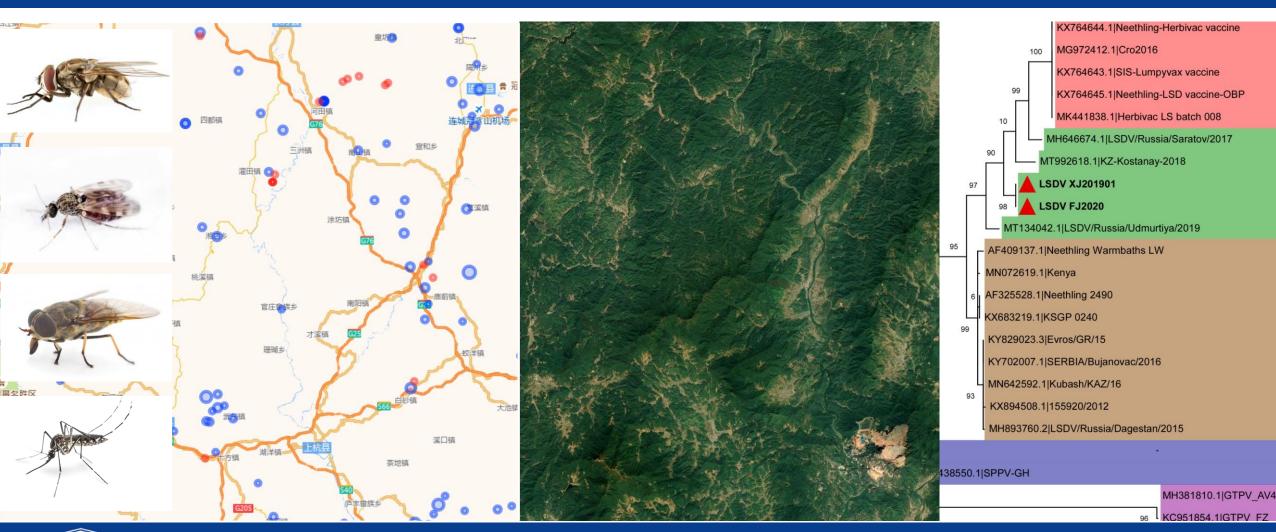












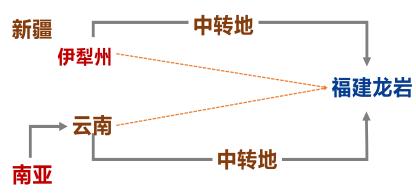




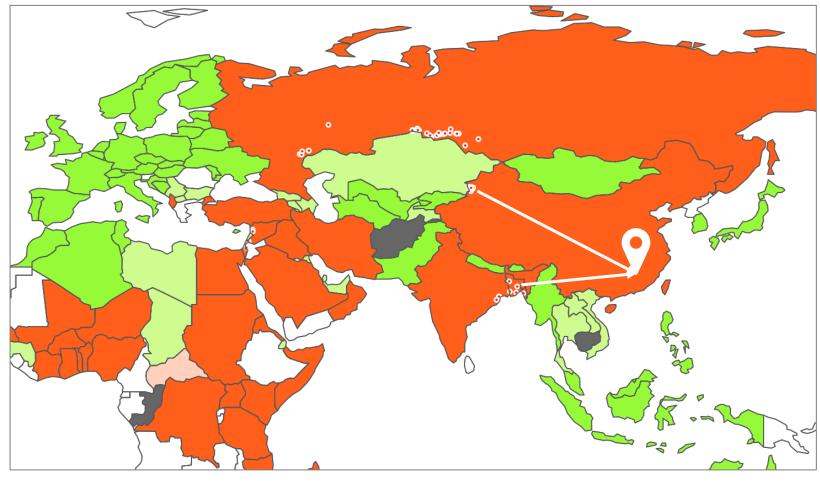














### **Reported Outbreaks**













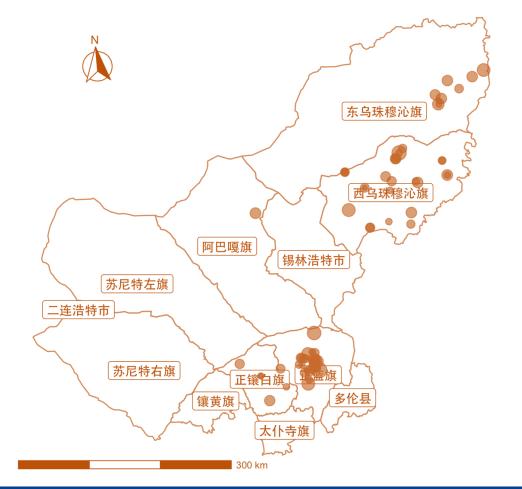












#### **Cattle Numbers**

调入数量

**5**0

100

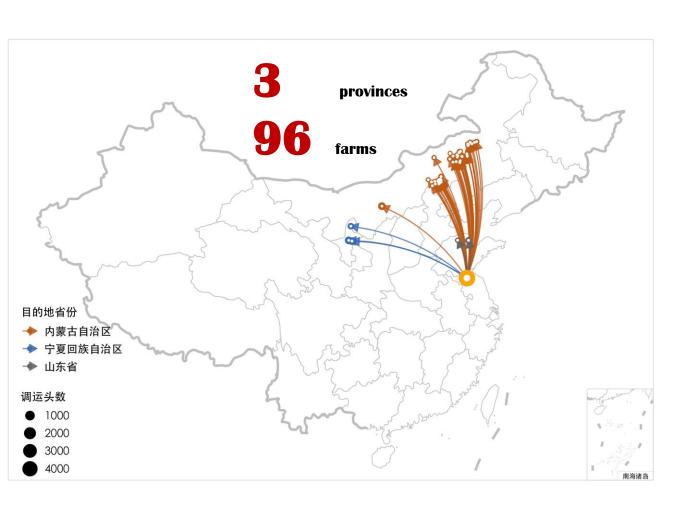














### 3. Control measures











### 农业农村部文件

农牧发[2019]26号

#### 农业农村部关于做好牛结节性皮肤病 防控工作的紧急通知

各省、自治区、直辖市及计划单列市农业农村(农牧、畜牧兽医)厅(局、委),新疆生产建设兵团农业农村局:

8月12日,经中国动物卫生与流行病学中心国家外来动物疫病研究中心确诊,新疆维吾尔自治区伊犁州发生牛结节性皮肤病疫情,这是我国首次确诊发生该病。经当地畜牧兽医部门排查,截至目前,共在伊犁州察布查尔县、霍城县、伊宁市发现病牛218头,死亡1头。为做好牛结节性皮肤病防控工作,现将有关事宜通知如下。

#### 一、高度重视牛结节性皮肤病防控工作

牛结节性皮肤病是由山羊痘病毒属结节性皮肤病病毒引起的

-1-

### 农业农村部文件

农牧发[2020]30号

### 农业农村部关于印发《牛结节性皮肤病防治技术规范》的通知

各省、自治区、直辖市及计划单列市农业农村(农牧、畜牧兽医)厅(局、委),新疆生产建设兵团农业农村局:

为做好牛结节性皮肤病防控工作,保障养牛业持续健康发展, 我部组织制定了《牛结节性皮肤病防治技术规范》。现印发给你们,请遵照执行。

.



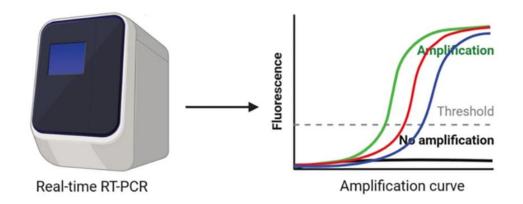








- LSDV specific Real-time PCR
- LSDV, GTPV, SPPV differential triplex Real-time PCR
- OIE recommended methods: PCR, Real-time PCR, ...









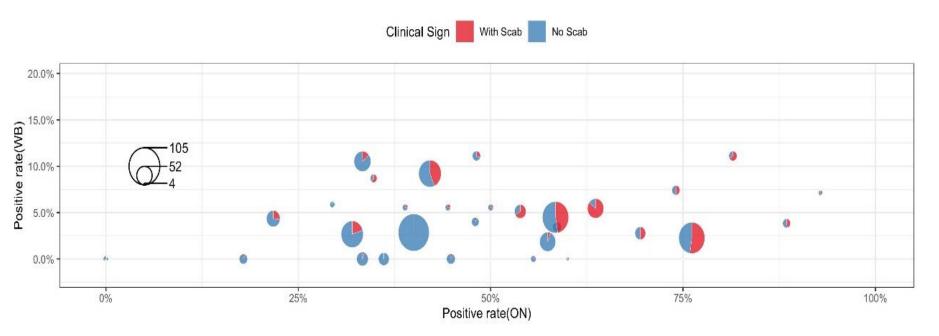






TYPE Brief Research Report
PUBLISHED 26 July 2022
DOI 10.3389/fvets.2022.936581

#### A total of 1206 cattle in 36 farms, 355 cattle with skin scabs



#### Check for updates

#### **OPEN ACCESS**

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real-time PCR detection and analysis
of a lumpy skin disease outbreak in
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Chinia. Front. Vet. Sci. 9: 936581.
doi: 10.1389/frvets.2022.936581

copprigner
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#### Quantitative real-time PCR detection and analysis of a lumpy skin disease outbreak in Inner Mongolia Autonomous Region, China

Lin Li<sup>11</sup>, Chuanxiang Qi<sup>1,21</sup>, Jinming Li<sup>1</sup>, Wenlong Nan<sup>1</sup>, Ying Wang<sup>1</sup>, Xing Chang<sup>1</sup>, Tianying Chi<sup>1</sup>, Mingxia Gong<sup>1</sup>, Da Ha<sup>3</sup>, Jide De<sup>3</sup>, Lifeng Ma<sup>4</sup> and Xiaodong Wu<sup>1\*</sup>

\*China Animal Health and Epidemiology Center, Oingdao, China, \*MOE Joint International Research Laboratory of Animal Health and Food Safety, MOA Key Laboratory of Animal Bacteriology, College of Veterinary Medicine, Narijing, Anicultural University, Narijing, China, \*Xingo League, Animal Disease Prevention and Control Center, Inner Mongolia, China, \*Inner Mongolia Animal Disease Prevention and Control Center Inner Mongolia, China, \*Inner Mongolia Animal Disease Prevention and Control Center Inner Mongolia, China.

Lumpy skin disease (LSD) is a severe disease of bovine characterized by nodules on the skin, mucous membranes, and profuse nasal discharge which causes severe economic losses. In October 2020, an LSD outbreak case was found in Inner Mongolia Autonomous Region, China. A total of 1,206 cattle were sold from the same imported animal quarantine field to 36 farms after the quarantine period finished, and over 30 farmers reported symptoms such as skin scabs found in newly arrived cattle shortly after that. A large-scale LSD outbreak investigation was launched after laboratory diagnosis confirmed LSD. The clinical samples of 1,206 cattle from 36 farms, including 1,206 whole blood, 1,206 oral and nose swabs, and 355 scabs, were collected for the qRT-PCR test. The result showed that 51 whole blood samples (4,23%), 580 swab samples (48,09%), and 350 skin scabs (98,59%) were lumpy skin disease virus (LSDV) positive, 33 of 36 farms were affected. This study aims to provide a basis for LSD epidemiological traceability, movement control, and measures for prevention and control.

#### KEYWO

LSDV, quantitative real-time PCR (qPCR), China, outbreak, Inner Mongolia Autonomous Region, 2020

#### Introduction

Lumpy skin disease (LSD) is a severe disease of bovine characterized by multifocal cutaneous nodules which were caused by the lumpy skin disease virus (LSDV). LSDV is classified into the genus Capripoxvirus of the family Pavariridae. As a large double-stranded DNA virus, LSDV encodes 156 putative viral genes with a genome of 151 bb (1). Unlike other Capripoxirus which infect sheep and goats, LSDV is highly host-specific and mainly infects cattle and buffalo under natural conditions. The severity of clinical signs of LSD varies from subclinical to fatal depending on the virulence of the strains and the host susceptibility.

atters in Veterinary Science 01 frontiers in .org













scab

- swab
- blood







**RP030** 

MW355944.1|China/GD01/2020

▲ LSDV FJ2019

KY829023.3|Evros/GR/15 AF409137.1|Neethling Warmbaths LW KX894508.1|155920/2012

MH893760.2|LSDV/Russia/Dagestan/2015

**GPCR** 

OM793602.1|LSDV Russia Tomsk 2020

MZ577073.1|20L42 Quyet-Thang/VNM/20

MZ577074.1|20L43 Ly-Quoc/VNM/20 MZ577076.1|20L81 Bang-Thanh/VNM/20 MZ577075.1|20L70 Dinh-To/VNM/20

OM793603 1II SDV Russia Khaharovsk 2020

MW732649.1|LSDV/HongKong/2020

OL752713.2|LSDV/KM/Taiwan/2020

ON152411.1|LSDV72/PrachuapKhiriKhan/Thailand/2021

MK441838.1|Herbivac LS batch 008

94 KX764643.1|SIS-Lumpyvax vaccine

AF409138.1|Neethling vaccine LW 1959 MG972412.1|Cro2016

KX764644.1|Neethling-Herbivac vaccine

- MH646674.1|LSDV/Russia/Saratov/2017

KX764645.1|Neethling-LSD vaccine-OBP MT134042.1|LSDV/Russia/Udmurtiya/2019





MW355944.1|China/GD01/2020

OM793602.1|LSDV Russia Tomsk 2020

MZ577073.1|20L42 Quyet-Thang/VNM/20 MZ577074.1|20L43 Ly-Quoc/VNM/20

MZ577076.1|20L81 Bang-Thanh/VNM/20

OM793603.1|LSDV Russia Khabarovsk 2020

LSDV FJ2019 ON152411.1ILSDV72/PrachuapKhiriKhan/Thailand/2021

MZ577075.1|20L70 Dinh-To/VNM/20

MW732649.1|LSDV/HongKong/2020

OL752713.2|LSDV/KM/Taiwan/2020

KX764643.1 SIS-Lumpyvax vaccine

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▲ LSDV XJ201901

MH646674 1II SDV/Russia/Saratov/2017

MT134042.1|LSDV/Russia/Udmurtiva/2019

MT992618.1|KZ-Kostanay-2018

KX683219 1 KSGP 0240

KY829023.3IEvros/GR/15



Li et al. BMC Veterinary Research (2022) 18:426 https://doi.org/10.1186/s12917-022-03525-9

**BMC Veterinary Research** 

#### Genetic analysis of genome sequence characteristics of two lumpy skin disease viruses isolated from China

Lin Li<sup>1†</sup>, Zhenzhong Wang<sup>1,2†</sup>, Chuanxiang Qi<sup>1,2</sup>, Shan Liu<sup>1</sup>, Mingxia Gong<sup>1</sup>, Jinming Li<sup>1</sup>, Xiaodong Wu<sup>1\*</sup> and Zhiliang Wang<sup>1</sup>

Background: Lumpy skin disease (LSD) is an acute or subacute infectious disease caused by lumpy skin disease virus (LSDV) of genus Capripoxvirus. The outbreaks of LSD were confirmed in the Yili area of the Xinjiang autonomous region in August 2019 and the Fujian province in June 2020. We detected LSDV in our daily monitoring work, then isolated, identified and sequenced the virus, and analyzed the whole genome characteristics of the isolated strain

Results: Whole genome sequencing revealed that the strains isolated were all LSDV and were named as LSDV XJ201901 and LSDV FJ2019. The results showed that the identity based on whole genome sequences between LSDV X1201901 and LSDV F12019 was 100% and the identity based on whole genome sequences between the two isolated strains and the global LSDV strains was 97,28%-99,99%, with the strain LSDV72/PrachuapKhiriKhan/Thailand/2021 (99.99%) having the highest sequence identity. Analysis of potential recombination events revealed that a total of 18 potential recombination events were identified in strains LSDV XJ201901 and LSDV FJ2019. The two strains are a recombination of Neethling vaccine LW 1959 (GeneBank: AF409138.1) with KSGP 0240 (GeneBank: KX683219.1). It was observed that Neethling vaccine LW 1959 (11/18) and KSGP 0240 (10/18) are involved in most of the potential

Conclusions: The virus isolate in this study was LSDV and was identified as a vaccine recombinant strain. The most likely potential parent strains of the two strains in this study are Neethling vaccine LW 1959 and KSGP 0240. The strains in this study are very similar to those isolated in East and Southeast Asia since 2019.

Keywords: China, Genome sequence, Genetic analysis, LSDV

Lumpy skin disease (LSD) is an acute or subacute infectious disease caused by lumpy skin disease virus (LSDV) of genus Capripoxvirus in Poxviridae. Cattle are the natural host of LSDV, and all kinds of cattle are susceptible

<sup>†</sup>Lin Li and Zhenzhong Wang contributed equally to this work

\*Correspondence: wuxiaodong@cahec.cn; wangzhiliang@cahec.cn China Animal Health and Epidemiology Center, Qingdao 266032, Shandong,

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to LSDV [1]. LSDV infection can lead to weight loss, sig-

nificant reduction of milk production and even death in

cattle. The mortality rate of cattle infected with LSDV

was up to 10% and even 20%, causing serious economic

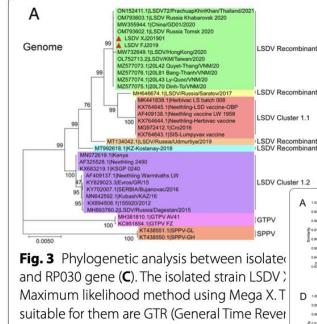
losses and affecting international trade [2]. Water buffalos [3], antelope [4] and giraffe can also be infected with LSDV. LSD is a notifiable disease stipulated by the

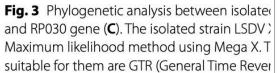
World Organization for Animal Health (OIE, 2017) and

1 of the 15 class I infectious diseases stipulated by List of

imported animal guarantine diseases of the China, LSD

was first observed in 1929 in Zambia in East Africa [5]

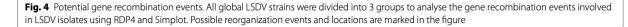




MN072619.1|Kenya KY702007.1|SERBIA/Bujanovac/2016

Neethling vaccine LW 1959

 $\downarrow \uparrow \uparrow \downarrow \uparrow \uparrow$ KSGP0240













- Emerging treatment for the infected farm:
  - Stamping out the affected cattle : clinical signs, LSDV positive.
  - Separate and quarantine other cattle.
  - Cleaning and disinfection.
  - Insecticide treatment.















- Emerging response for the affected county:
  - Live cattle movement is restricted for 30 days.
  - Separate and quarantine the introduced cattle.
  - Vaccination campaigns for unaffected herds.
  - Outbreak investigation, traceability forward/backward for last 30 days.
  - Vector control.



30days













- Emerging response for other counties:
  - Enhance surveillance, especially for related counties and neighbor counties.
  - Enhance quarantine inspection for animal movements.
  - Awareness campaigns.
  - Vaccination is required for animal movement at least 30 days before the date of departure.



30days















山东绿都生物科技有限公司

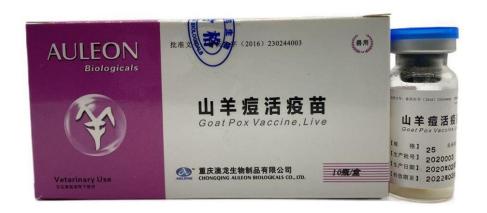






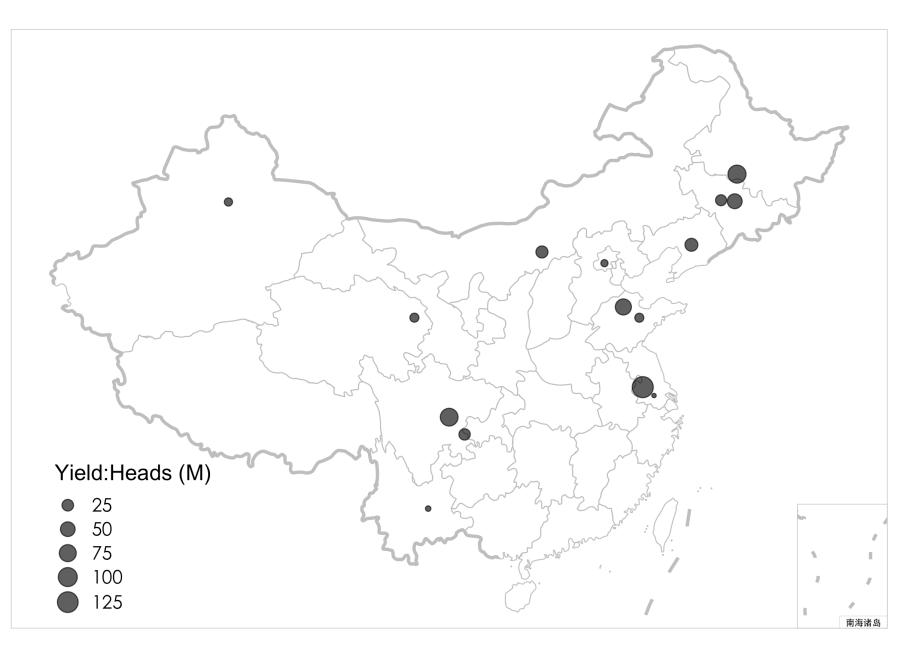


**AV41** 

















#### **Goat Pox Vaccine Manufacture**

- 哈药集团生物疫苗有限公司
- 山东绿都生物科技有限公司
- 吉林正业生物制品股份有限公司
- 辽宁益康生物股份有限公司
- 金宇保灵生物药品有限公司
- 重庆澳龙生物制品有限公司
- 吉林和元生物工程股份有限公司
- 青海生物药品厂有限公司
- 山东华宏生物工程有限公司
- 天康生物制药有限公司
- 北京华信农威生物科技有限公司
- 云南生物制药有限公司
- 江苏南农高科技股份有限公司













# National Surveillance Plan 2022

**Provinces: 27** 

Samples: 9498

swabs: 4423

serum: 4434

Farms: 427

Data source: CAHEC













# Post Vaccination Evaluation 2022

- 913 samples, 95 farms (vaccinated)
- ID Screen Capripox double antigen
   Multi-species ELISA kit (ID Vet,
   Montpellier, France)
- Positive rate 28.4% (259/913)

Data source: CAHEC



# Thank You









lijinming@cahec.cn



- African Swine Fever
- Bovine Spongiform Encephalopathy
- Newcastle Disease



- African Swine Fever
- Newcastle Disease
- Peste des petits ruminants



Peste des petits ruminants



