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Lumpy skin disease (LSD) An update on LSD situation and its control

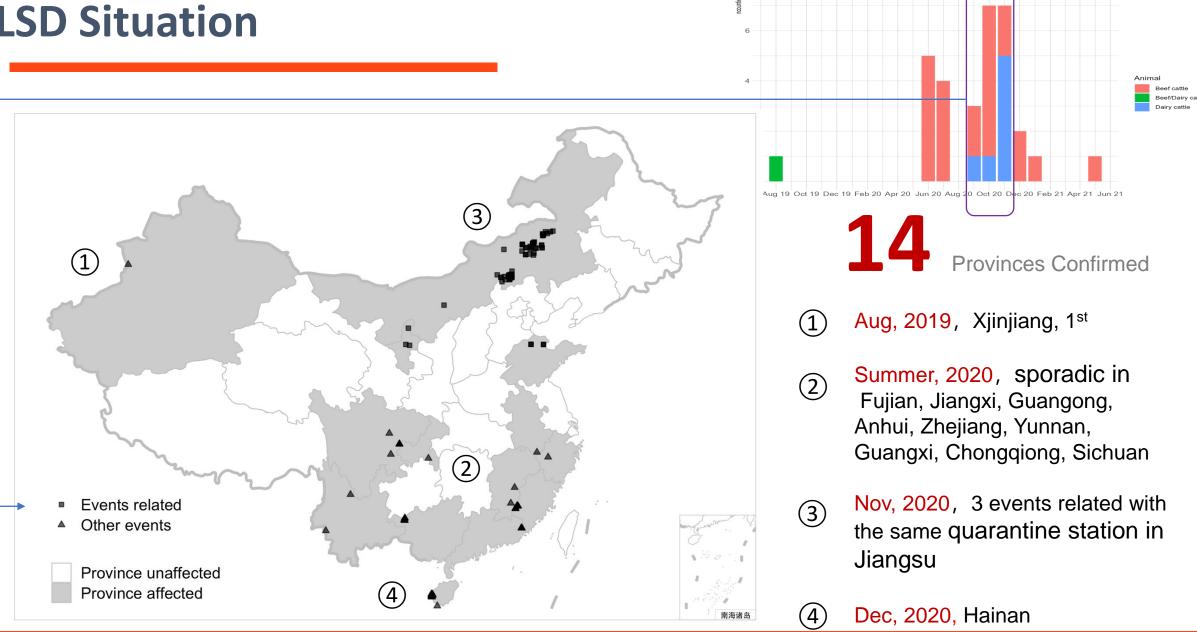
LI Jinming



Deputy Director

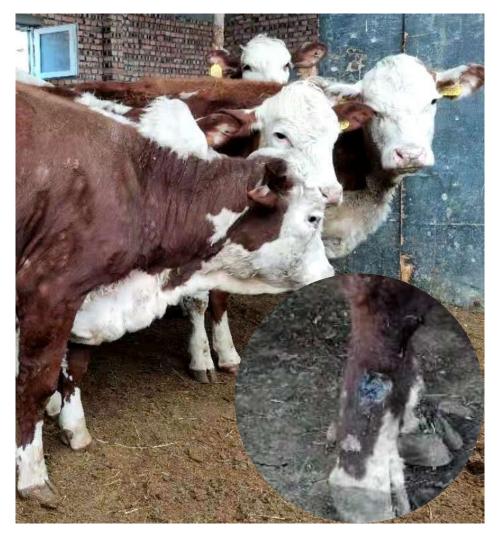
China Animal Health and Epidemiology Center

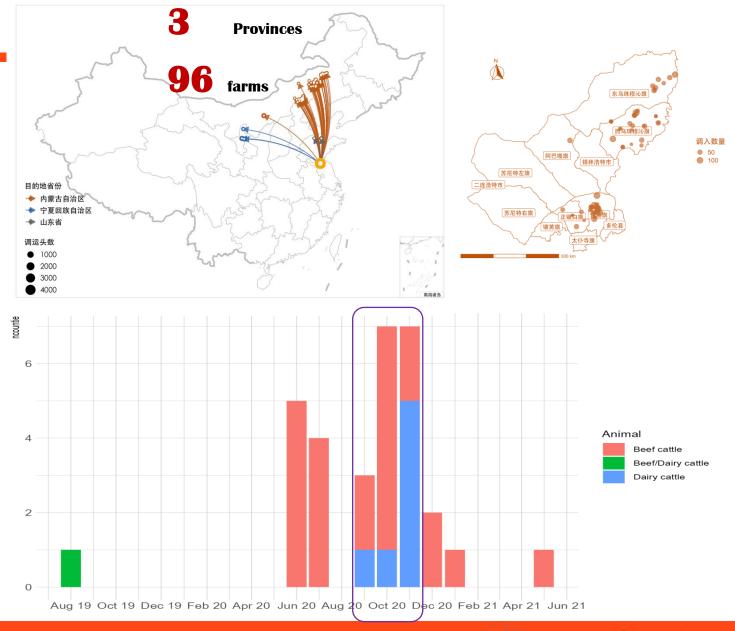
Fourth LSD Coordination Meeting for South-East Asia 28 – 29 November 2023, Bangkok, Thailand





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28 Counties **1445** Cases **34** Deaths

Animal	No. Provinces	No. Cases	No. Deaths
Beef Cattle	12	1259	30
Dairy Cattle	4	121	4
Beef/Dairy Cattle	1	65	0
Total	14	1445	34





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isolated from China

Genetic analysis of genome sequence

characteristics of two lumpy skin disease viruses

RESEARCH

Circulating virus types (confirmed by CAHEC)

Neethling vaccine LW 1959 Lin Li^{1†}, Zhenzhong Wang^{1,2†}, Chuanxiang Qi^{1,2}, Shan Liu¹, Mingxia Gong¹, Jinming Li¹, Xiaodong Wu^{1*} and ON152411.1|LSDV72/PrachuapKhiriKhan/Thailand/2021 OM793603.1 LSDV Russia Khabarovsk 2020 Zhiliang Wang^{1*} $\downarrow \uparrow \downarrow \uparrow$ MW355944.1|China/GD01/2020 OM793602.1ILSDV Russia Tomsk 2020 **KSGP0240** LSDV XJ201901 Abstract LSDV FJ2019 Genome LSDV Recombinant Background: Lumpy skin disease (LSD) is an acute or subacute infectious disease caused by lumpy skin disease MW732649.1|LSDV/HongKong/2020 virus (LSDV) of genus Capripoxvirus. The outbreaks of LSD were confirmed in the Yili area of the Xinjiang autonomous OL752713.2|LSDV/KM/Taiwan/2020 region in August 2019 and the Fujian province in June 2020. We detected LSDV in our daily monitoring work, then MZ577073.1|20L42 Quyet-Thang/VNM/20 isolated, identified and sequenced the virus, and analyzed the whole genome characteristics of the isolated strain. 99 MZ577076.1|20L81 Bang-Thanh/VNM/20 Results: Whole genome sequencing revealed that the strains isolated were all LSDV and were named as LSDV MZ577074.1|20L43 Ly-Quoc/VNM/20 XJ201901 and LSDV FJ2019. The results showed that the identity based on whole genome sequences between LSDV MZ577075.1|20L70 Dinh-To/VNM/20 LSDV Recombinant MH646674.1|LSDV/Russia/Saratov/2017 MK441838.1|Herbivac LS batch 008 76 KX764645.1|Neethling-LSD vaccine-OBP 99 AF409138.1|Neethling vaccine LW 1959 LSDV Cluster 1.1 100 KX764644.1 Neethling-Herbivac vaccine 99 MG972412.1|Cro2016 KX764643.1|SIS-Lumpyvax vaccine LSDV Recombinant MT134042.1|LSDV/Russia/Udmurtiya/2019 99 MT992618.1|KZ-Kostanay-2018 LSDV Recombinant MN072619.1|Kenya AF325528.1|Neethling 2490 KX683219.1|KSGP 0240 100 AF409137.1 Neethling Warmbaths LW KY829023.3|Evros/GR/15 LSDV Cluster 1.2 KY702007.1|SERBIA/Bujanovac/2016 MN642592.1|Kubash/KAZ/16 KX894508.1|155920/2012 MH893760.2|LSDV/Russia/Dagestan/2015 MH381810.1|GTPV AV41 GTPV KC951854.1|GTPV FZ 100 KT438551.1|SPPV-GL SPPV KT438550.1|SPPV-GH 100 0.0050

Fig. 4 Potential gene recombination events. All global LSDV strains were divided into 3 groups to analyse the gene recombination events involved in LSDV isolates using RDP4 and Simplot. Possible reorganization events and locations are marked in the figure

Li *et al*, 2022

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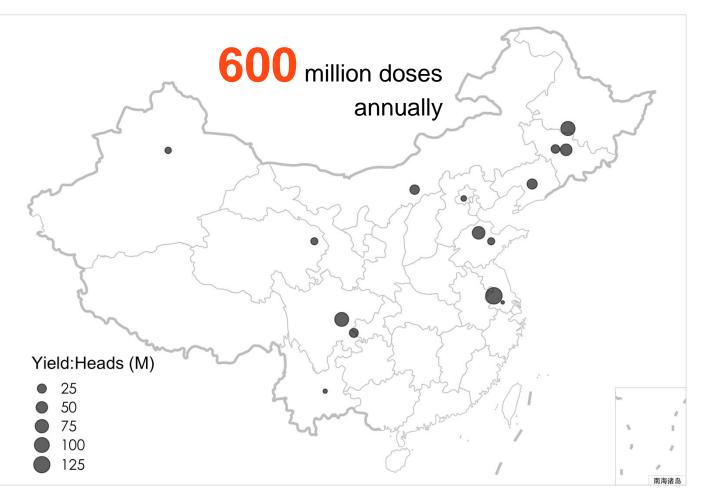
• Live-attenuated vaccines against goat pox (AV41 strain)





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• Live-attenuated vaccines against goat pox (AV41 strain)



Goat Pox Vaccine Manufactures

- 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



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• 2 inactivated vaccines were approved,

but not yet commercially available



Manufacturer	Virus strain	Titre	Dose	booster Administration	Immunity duration
<u>Jinyu Baoling Bio-</u> <u>Pharmaceutica</u> <u>Co.,Ltd.</u>	LSDV NMG strain	10 ^{6.5} TCID ₅₀	2.0mL	21d	6 months
<u>China Animal</u> <u>Husbandry</u> Industry Co.,Ltd.	GPTV AV41 strain	10 ^{6.2} TCID ₅₀	2.0mL	14	6 months



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







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- Emerging response for affected county:
 - Live cattle movement is restricted for 30 days.
 - Separate and quarantine the introduced cattle.
 - Vaccination campaign for unaffected herds.
 - Trace forward/backward the contact cattle within the last 30 days.
 - Vector control.





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



National Surveillance Plan 2023 Provinces: 28 Samples: 14,108 swabs: 6,048 serum: 7,160

Post Vaccination Evaluation 2023

- 2914 samples, 264 farms (vaccinated)
- ID Screen Capripox double antigen

Multi-species

- **ELISA** kit (ID Vet, Montpellier, France)
- Positive rate 28.4% (827/2914)

Data source: CAHEC



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Main challenges and recommended solution

Main challenges

- Timely outbreak reporting
- Animal movement control
- Vector control
- Post vaccination evaluation
- Recombinant Strains
- Wild animal infection

Recommended solution

- Encourage farmers to report disease
- Biosafety control measures
- Vaccination coverage
- Surveillance and research on new virus strains
- Cooperation and coordination with other sectors





THANK YOU



Reference Laboratories fo



- African Swine Fever
- Bovine Spongiform Encephalopathy
- Newcastle Disease



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



Peste des petits ruminants

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