# WHO-convened Global Study of Origins of SARS-CoV-2: China Part

## Joint WHO-China Study

14 January-10 February 2021

Joint Report

### Stage 1 SARS-CoV-2 origins: Detailed reconstruction of current knowledge

#### Epidemiology

Studies on earliest known cases Studies investigating possibility of earlier circulation in Wuhan or elsewhere

#### Molecular epidemiology

Linking genomes with earliest known cases Analyse in combination with epi information



## Fever, ILI, ARI, and unspecified pneumonia in Wuhan, 1 Oct to 10 Dec 2019, and cases notified in December 2019



76,253 records, 233 health facilities, Wuhan. 92 COVID-19 compatible cases: 62 tested after one year – all seronegative

Retrospective testing of ILI cases: 9 positives in January, none in December







Trends of the pneumonia mortality rate in 2019-2020 compared to 2016-2018, Hubei Province (excl Wuhan), all ages.





#### Aquatic products





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#### Wild animal meat





#### Poultry

Meat

- 460 (frozen) Products from 9 provinces and 58 countries
- farmed wild animals and products from regions that are bat SARSr-CoV hotspots, including provinces where closest relatives to SARS-CoV-2 are found







Sequences of cases with onset of illness in December: Market may have served as amplification event, but already some diversity early december



ELISA testing of animals in & around Wuhan during Epidemic period:

Nov 2019-Mar 2020

Species	Number	Result
	tested	
Pig	187	Negative
Cow	107	Negative
Sheep	133	Negative
Horse	18	Negative
Chicken	153	Negative
Duck	153	Negative
Goose	25	Negative
Mice	81	Negative
Rat	67	Negative
Guinea pig	30	Negative
Rabbit	34	Negative
Monkey	39	Negative
Dog	487	Negative
Cat	87	Negative
Camel	31	Negative
Fox	89	Negative
Mink	91	Negative
Alpaca	10	Negative
Ferret	2	Negative
Bamboo rat	8	Negative
Peacock	4	Negative
Eagle	1	Negative
Tiger	8	Negative
Rhinoceros	4	Negative
Pangolin	17	Negative
Leopard cat	3	Negative
Jackal	1	Negative
Giant panda	14	Negative
Masked	10	Negative
civet		
Porcupine	2	Negative
Bear	9	Negative
Yellow-	4	Negative
throated		
marten		
Weasel	1	Negative
Red pandas	3	Negative
Wild boar	1	Negative

Also

1115 fecal samples fromdifferent bat species from Hubei(Nuclear Acid Testing (NAT))

Main livestock species: 11708 samples for serology 12092 samples for NAT

~27000 samples of different animals stored in 2019-2020 from 24 provinces for NAT

## Evidence syntheses pathways of emergence



## Recommendations

- Serosurveys using retained blood samples from Sept Dec 2019 from Wuhan/Hubei and other regions with evidence of early circulation
- Review of earliest cases from central genetic cluster Sichuan and elsewhere
- Review of notified cases, and potential cases from retrospective search, using less stringent case definitions
- Review of mortality data from other provinces with early cases as guided by phylogenetic analyses
- Retesting of suspected cases by serology, or following vaccination (primary or booster reponse)
- Further tracing-back of wild animal farms and testing (incl frozen products) based on supply chain analysis
- Fur animal farm screening (Asia)
- Further bat surveys within China, in Southeast Asia, and in other countries where *Rhinolophus* bats are found