

# Monkeypox virus

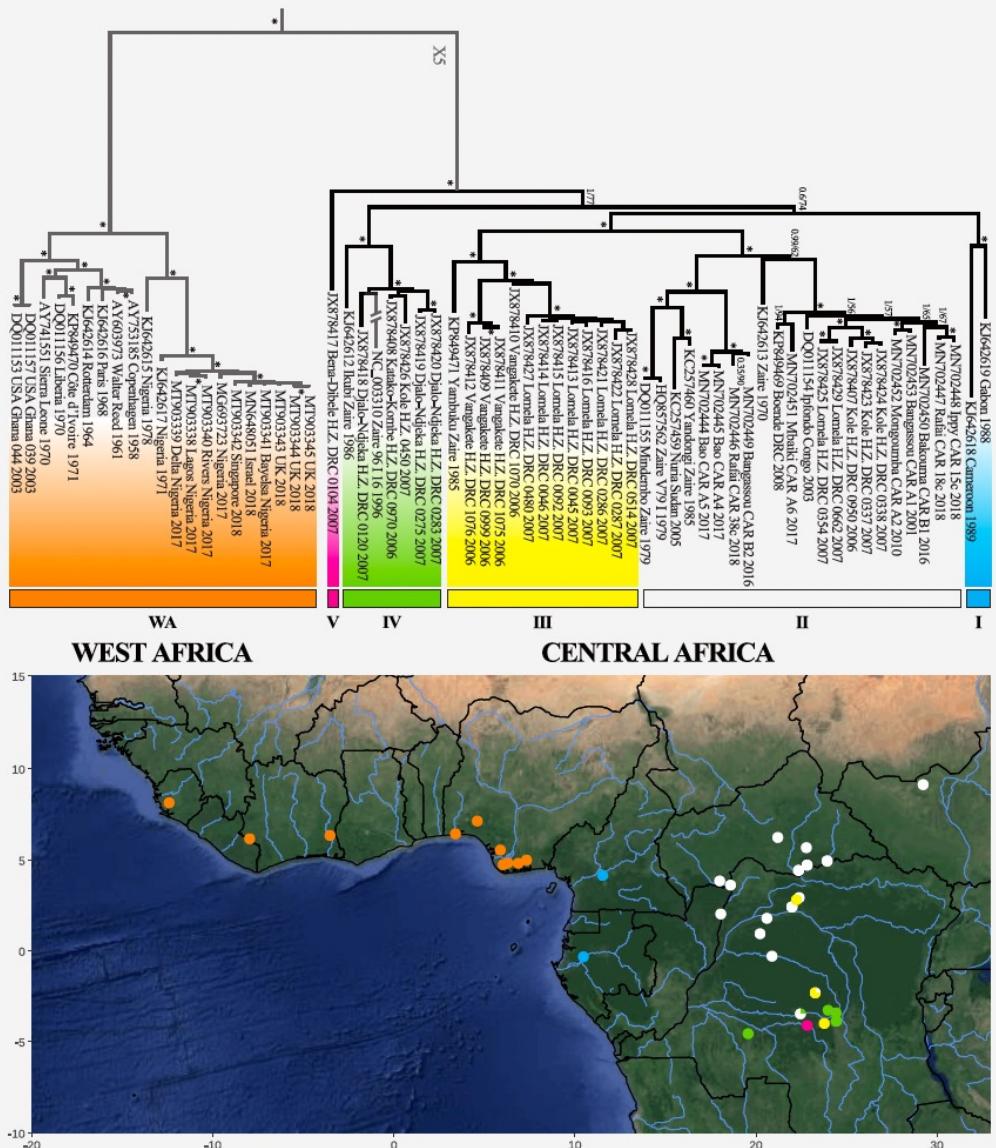
Monkeypox is caused by a virus: Monkeypox virus (MPXV)



- Poxviridae, like smallpox (Fenner and Nakano, 1988)
- Specifically *Orthopoxvirus* (Fenner and Nakano, 1988)
- Double-stranded DNA genome
- 197kb genome and 186 ORFs



# Phylogeny of Monkeypox virus



# Whole genome phylogeny of Monkeypox cases

- Human cases
  - Animal cases

2 main phylogenetic groups of MPXV

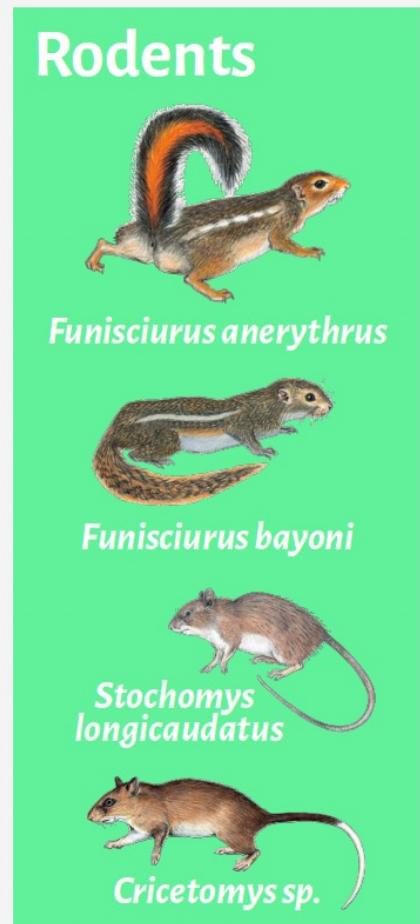
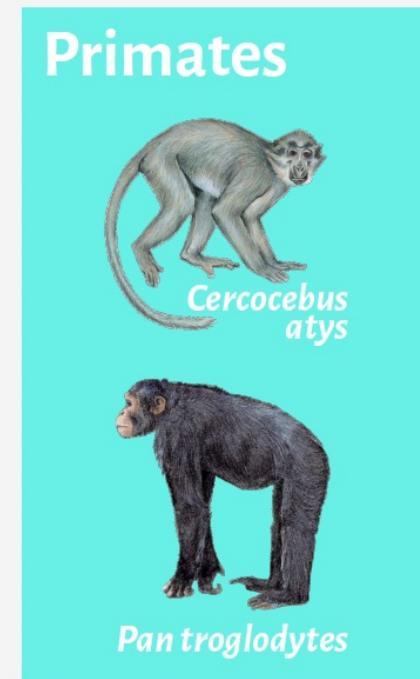
- West Africa
  - Central Africa/Congo Basin
  - Smaller groups within the two clades

## All cases in African rainforests

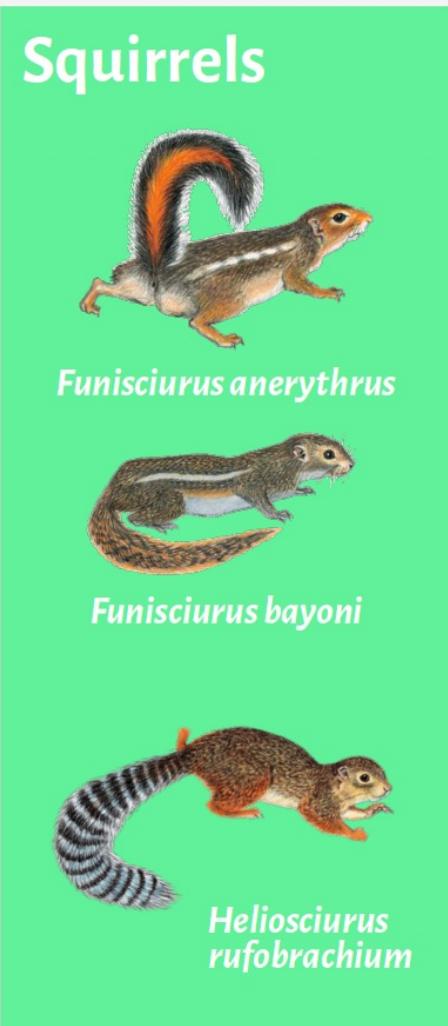
## → Which regions are suitable for the virus?

# What is Monkeypox virus animal reservoir?

1. MPXV belongs to *Orthopoxvirus*, a genus exclusive to mammals  
→ MPXV reservoir is a **mammal**
2. All human cases in African rainforests  
→ MPXV reservoir is an **African rainforests-dwelling mammals**
3. Isolated from seven wild species of African rainforests
  - All of them reservoirs?
  - Some of them intermediate hosts?
  - Some of them victims of the virus?
- **Exhaustive list of reservoirs**
  - **Niches of African rainforests-dwelling mammals**



# African squirrels as a potential reservoir of Monkeypox virus



1. MMPXV isolated from two African squirrels
    - *Funisciurus anerythrus* (Khodakevich *et al.*, 1986)
    - *Funisciurus bayonii* (Mariën *et al.*, in review)
  2. MPXV DNA in African squirrel museum specimens (Tiee *et al.*, 2018)
    - Five species of *Funisciurus* including two new species
  3. Anti-OPXV antibodies in African squirrels (Khodakevich *et al.*, 1988)
    - *Funisciurus*
    - *Heliosciurus*
- **African squirrels are good candidates for the reservoir of Monkeypox virus**
- **Focus on African squirrels**

# Monkeypox

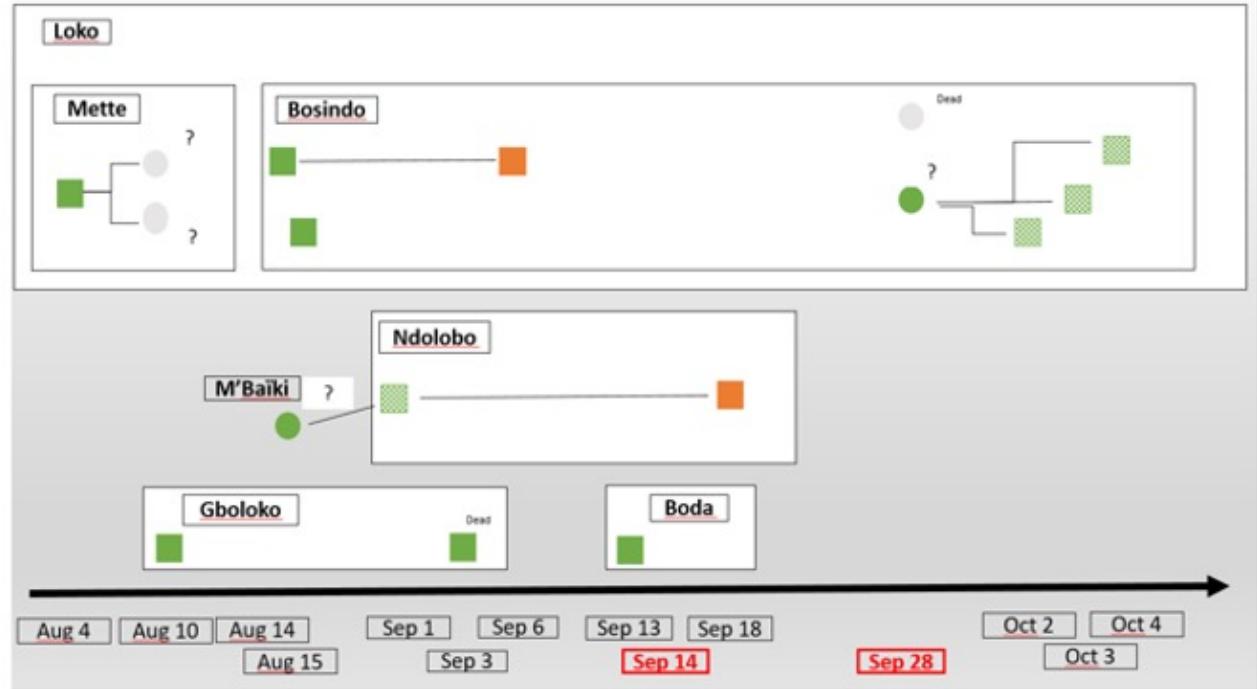
- Monkeypox is a viral zoonotic disease
- Part of the *Orthopoxvirus* genus which includes variola virus (smallpox) and cowpox virus
- Endemic in 9+ African countries
- There are two main strains, one which typically has caused more severe illness (Congo clade) than the other (West African clade). Only the West African clade has been identified in the multi-country outbreak.
- The reservoir host is still unknown, although rodents incidental hosts and play a part in transmission, typically through hunting, preparation or consumption of meat (game)



World Health Organization

HEALTH  
EMERGENCIES  
programme

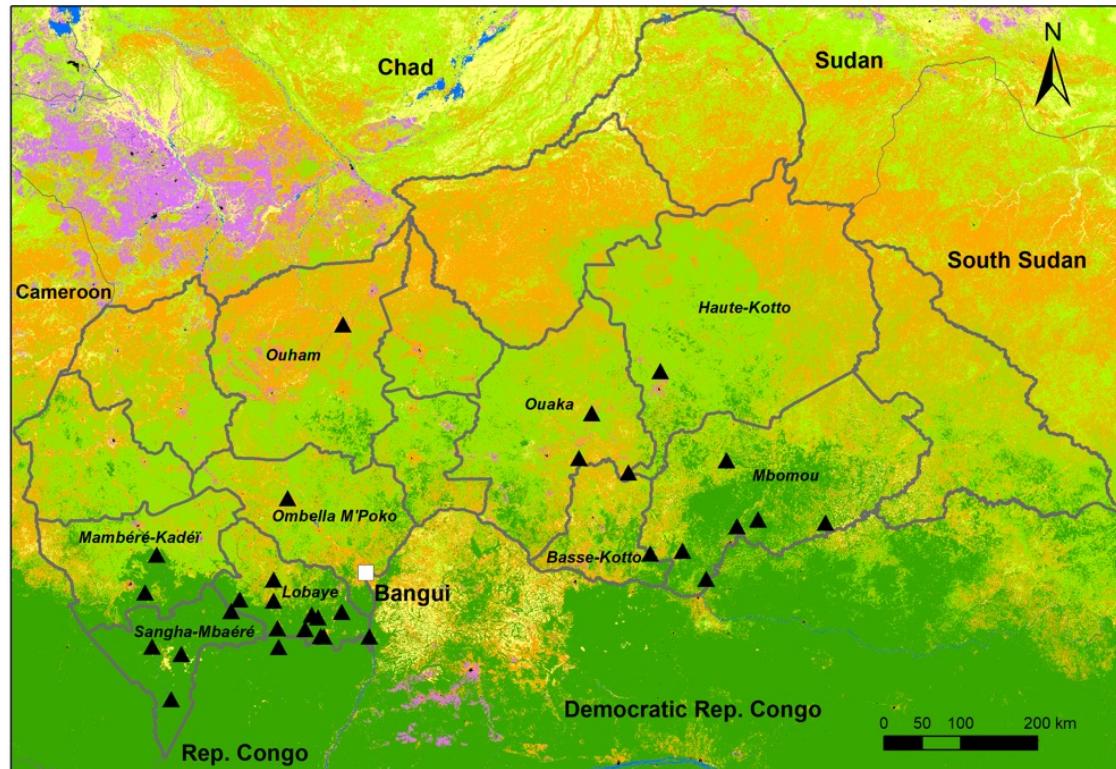
# Monkeypox : Outbreak investigation



Pattern of monkeypox virus transmission hypothesized to have occurred during these outbreaks



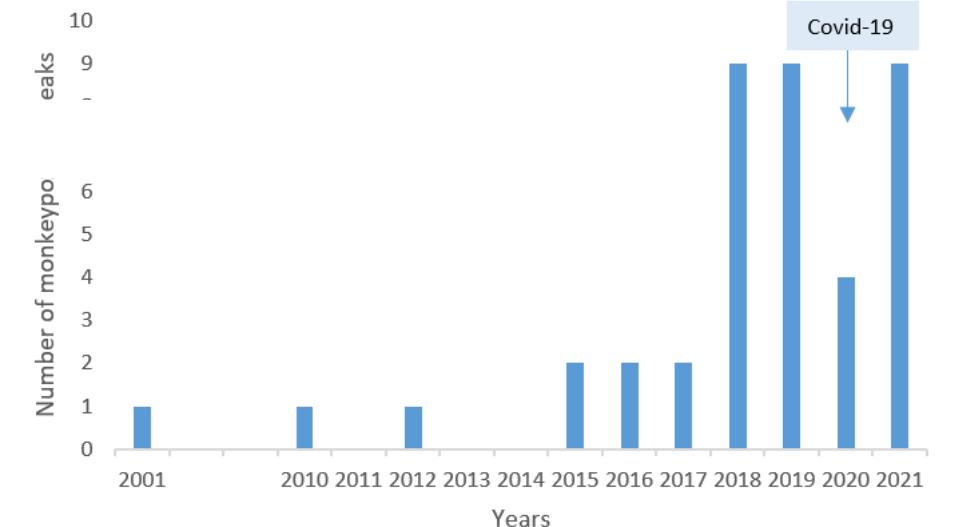
# EPIDEMIOLOGY



Landuse/Landcover data source: Copernicus 2019 Global 100m Landcover

Buchhorn, M. ; Lesiv, M. ; Tsendsazar, N. - E. ; Herold, M. ; Bertels, L. ; Smets, B. Copernicus Global Land Cover Layers — Collection 2. Remote Sensing 2020, 12, Volume 108, 1044. DOI 10.3390/rs12061044

Annual distribution of monkeypox outbreaks in the CAR, 2001-2021 (n=40)



40 outbreaks, size range: 1 to 13 (1 to 25)  
 99 confirmed cases, (160 with suspected cases)  
 Case-fatality rate: 12/160 (7.5%)

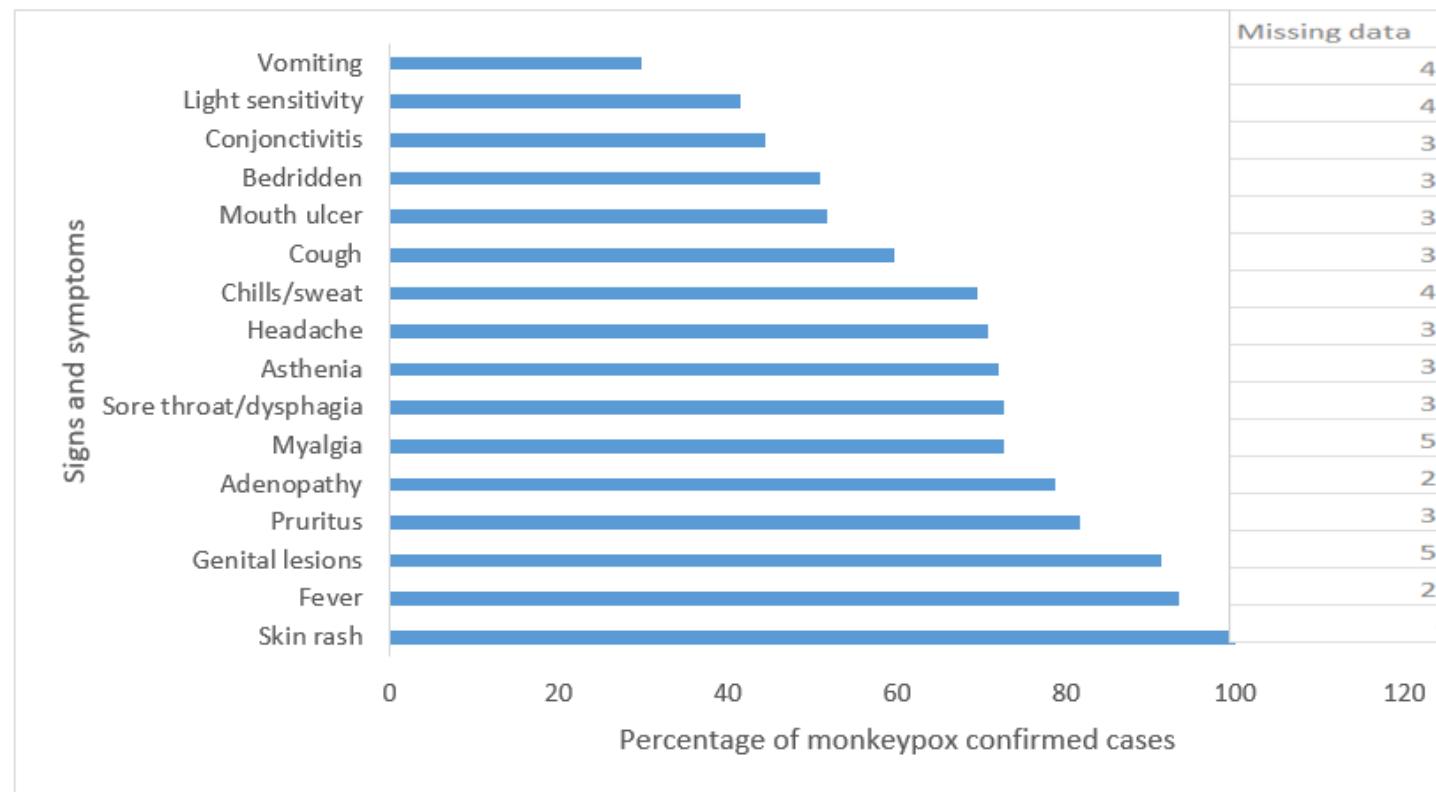
(Besombes et al, in preparation)

# Clinical characteristics

**Incubation= 7 days (IQR : 1-13, range : 0 - 17)**

**Genital lesions : 91,3% (n=42/46 with available information)**

Figure 4: Frequency of signs and symptoms in people with confirmed monkeypox virus infection in the CAR, 2001-2021 (n=99)

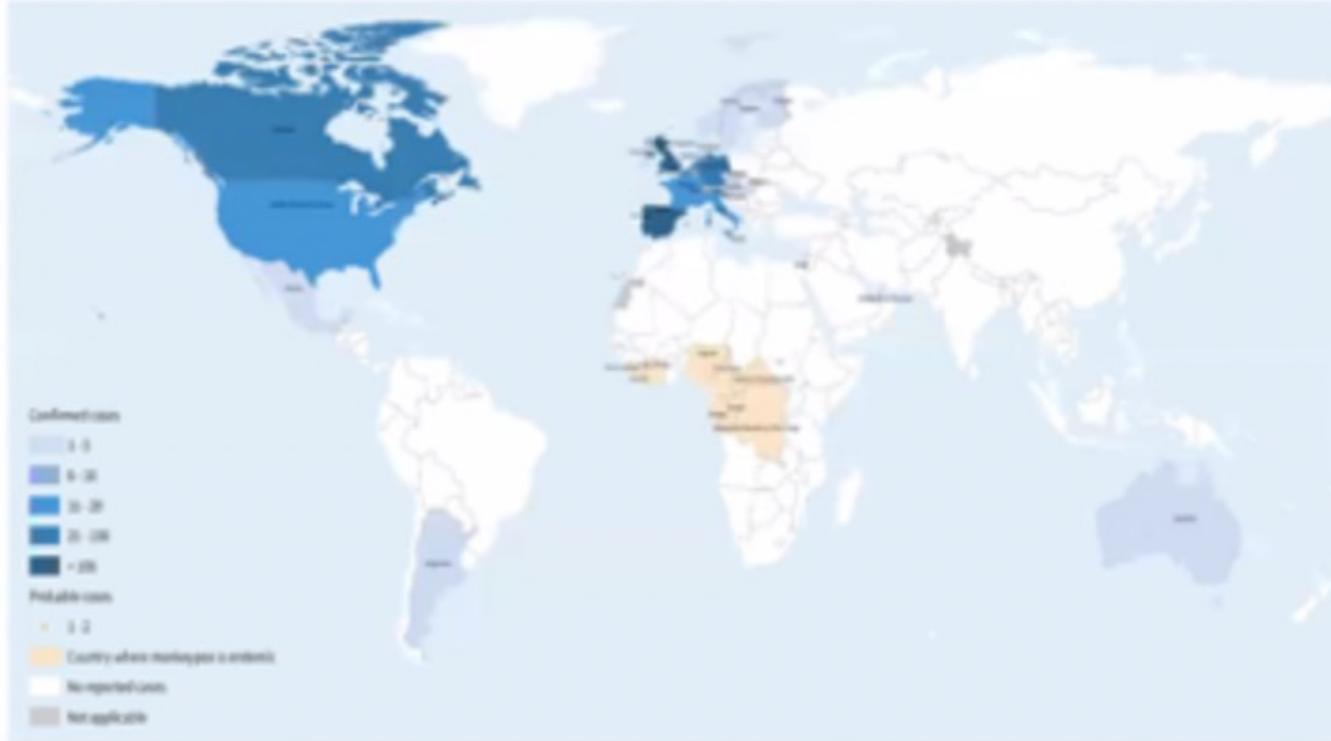


# Cases of monkeypox in non-endemic countries

13 May to 1 June 2022

Region	Country	Confirmed	Probable
AMRO	Argentina	2	
	Canada	26	
	Mexico	1	
	United States of America	18	
EMRO	United Arab Emirates	4	
	Austria	1	
	Belgium	10	2
	Czechia	5	
	Denmark	2	
	Finland	1	
	France	17	
	Germany	44	
	Hungary	1	
	Ireland	4	
EURO	Israel	2	
	Italy	14	
	Malta	1	
	Netherlands	26	
	Norway	1	
	Portugal	119	
	Slovenia	2	
	Spain	142	
	Sweden	4	
	Switzerland	4	
	The United Kingdom	190	
WPRO	Australia	2	
Total	26 countries	643	2

Confirmed and probable cases of monkeypox in non-endemic countries  
(since 13 May 2022, as of 1 June 2022 16:00 CEST)

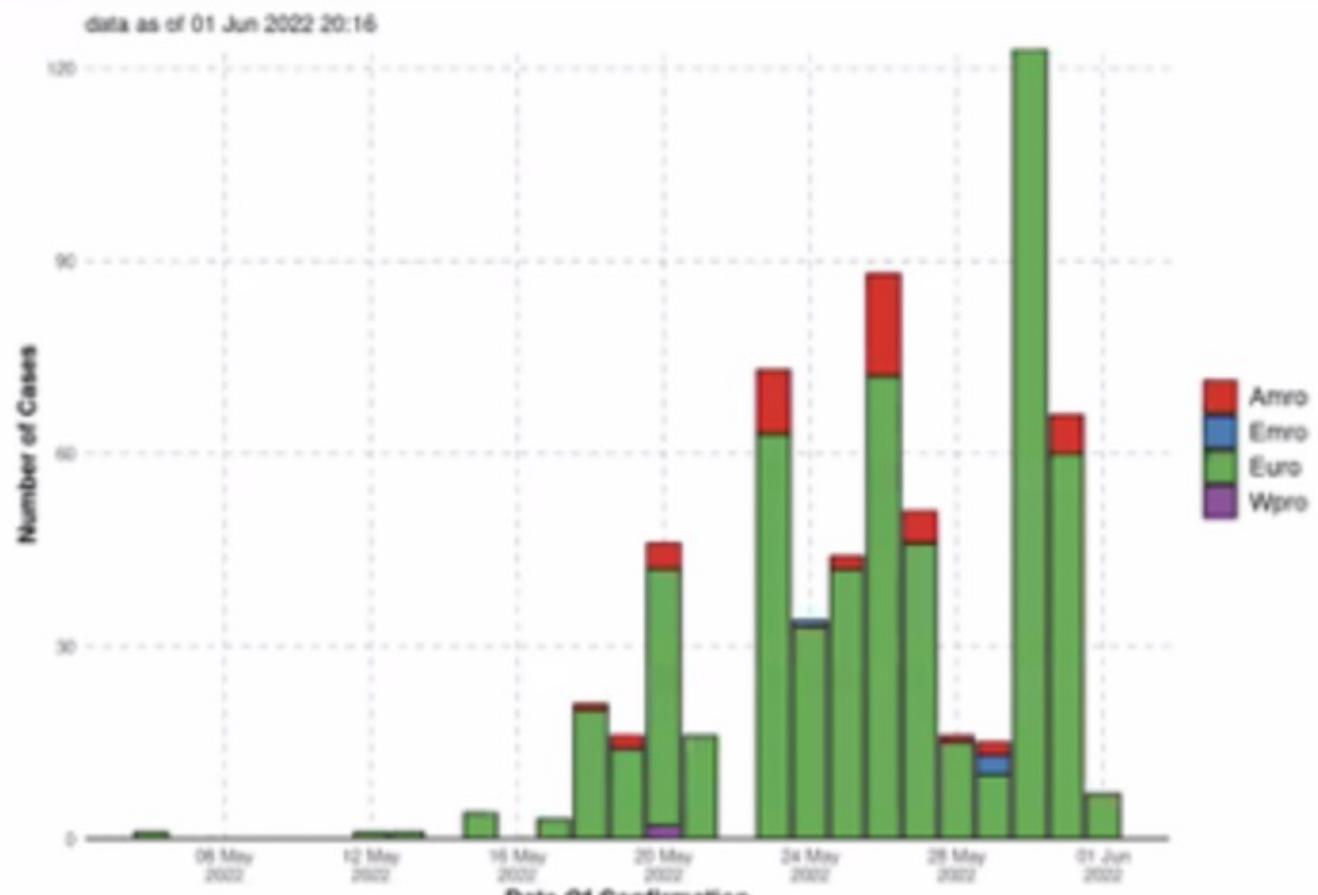


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Data Source: World Health Organization  
Map Production: WHO Health Emergencies Programme  
Map Date: 1 June 2022



# Epidemic Curve by region and date of confirmation (public database)



Source: global health linelist - official reporting only.  
where dates are not available, showing date of database entry

# Epidemiological update as of 1 June 2022



- 373 cases in the EU/EEA since 18 May:  
Austria (1), Belgium (10), Czechia (5), Denmark (2), Finland (1), France (17), Germany (44), Hungary (1), Ireland (2)\*, Italy (14), Malta (1), the Netherlands (26), Portugal (100), Slovenia (2), Spain (142)\* and Sweden (4).
- Almost all cases have no travel history to endemic countries.
- The majority of cases are in young men, self-identifying as MSM.
- Most cases presented with lesions on genitalia or peri-genital area, indicating that transmission likely occurs during close physical contact during sexual activities.
- **No deaths** have been reported.
- First time that chains of transmission are reported in Europe **without known epidemiological links to West or Central Africa**.

# AFRIPOX

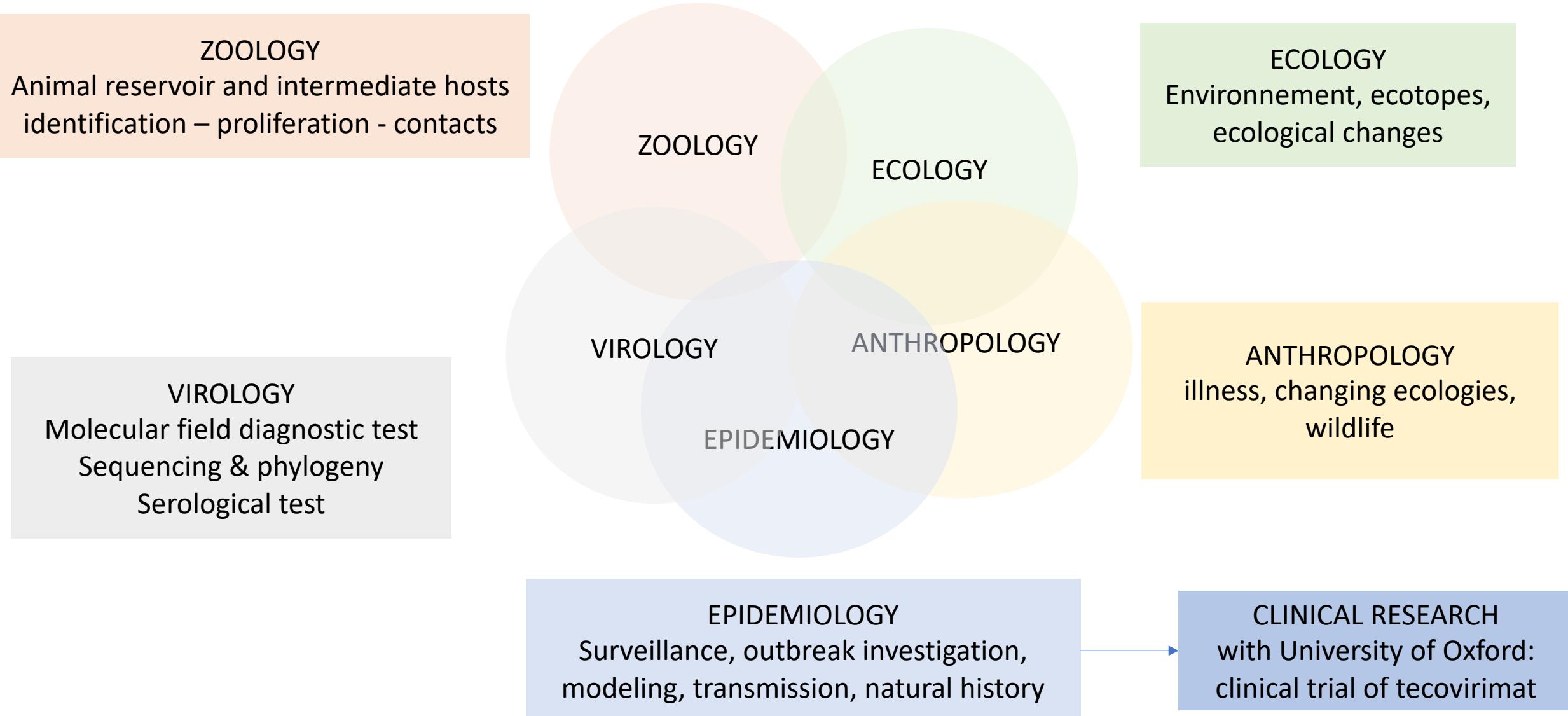
## A One Health approach of monkeypox in the Central African Republic

Emmanuel Nakouné & Arnaud Fontanet

2-3 June 2022 – WHO monkeypox research webinar



# One health – monkeypox - CAR



# VIROLOGY – diagnostic tools



## Molecular field diagnostics

- Monoplex (RT-LAMP) and multiplex (RT-LAMP QUASR) rapid tests
- Integrated test cartridges (Withings) (MPX lineage, VZV)
- Detection by isothermal amplification / RPA strip technology



## Serological diagnostic tests

- Multiplex test based on a library of more than 7891 viral peptides representative of the proteome of human pathogenic OPXV (MPXV, VACV, CPXV, VARV) (PhiP-Seq)
- Multiplex assay based on a combination of a selection of 10 MPXV proteins and peptides (MMIA)

# VIROLOGY - sequencing

-Use of capture probes for long DNA fragments

- Microfluidic technology and droplet generation for targeted enrichment of droplets containing viral genome fragments

-Direct Illumina sequencing, or direct and real-time MinION sequencing on samples



[www.nature.com/scientificreports/](http://www.nature.com/scientificreports/)

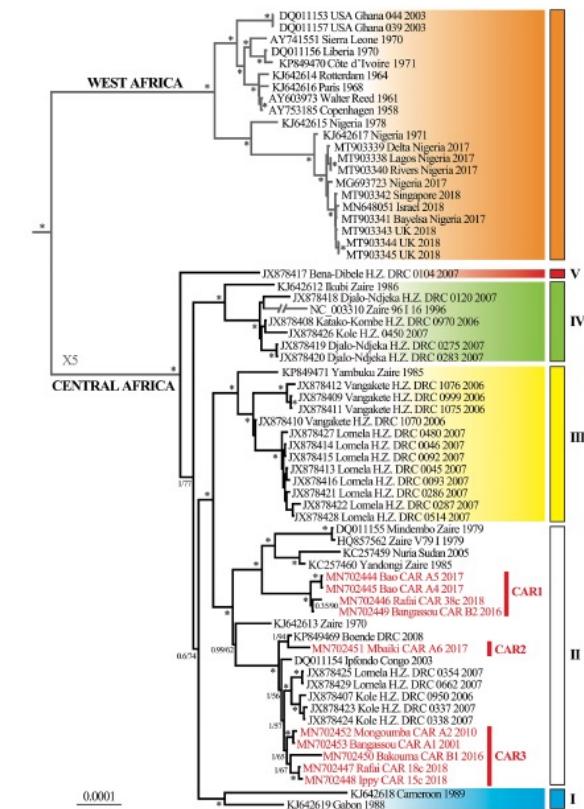
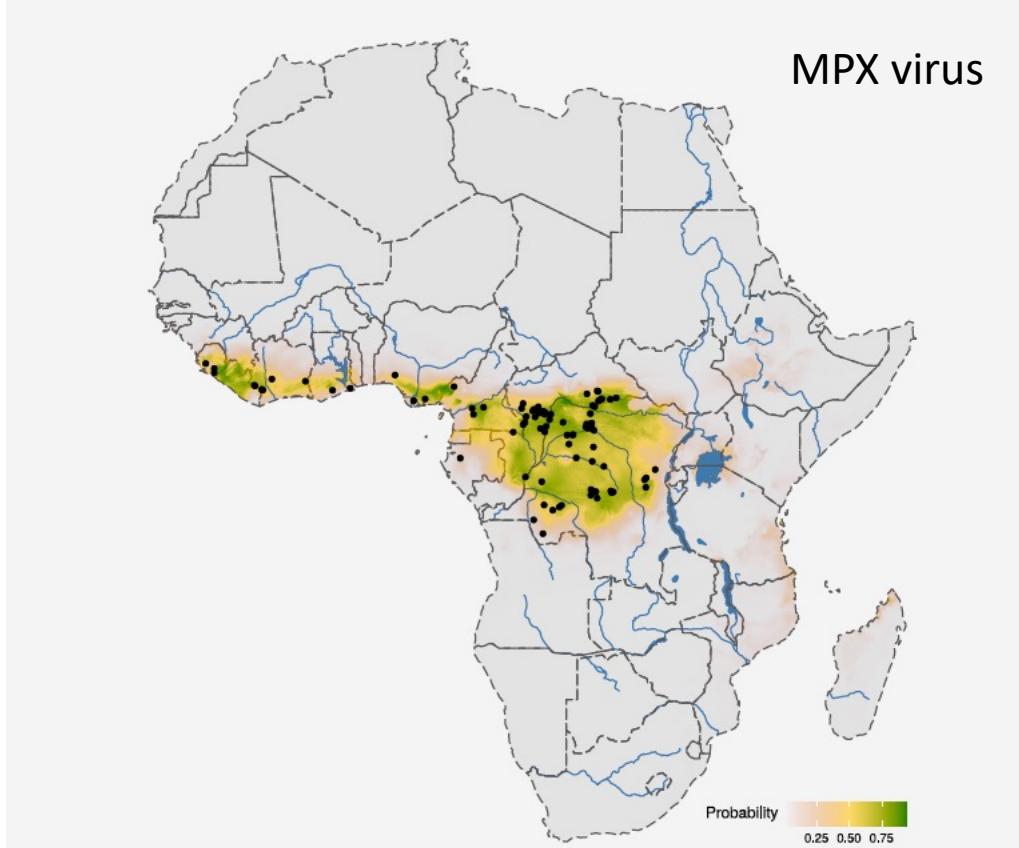


Figure 2. Phylogeny of monkeypox virus (MPXV) based on complete genomes. The Bayesian tree was

(Berthet et al, Scientific Reports, 2021)

# ZOOLOGY – Ecological niche



## African squirrels as a potential reservoir of Monkeypox virus



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- **African squirrels are good candidates for the reservoir of Monkeypox virus**
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Curaudeau *et al.*, in prep

Ecological Niche Modelling with MaxEnt in R

# ZOOLOGY – Field work

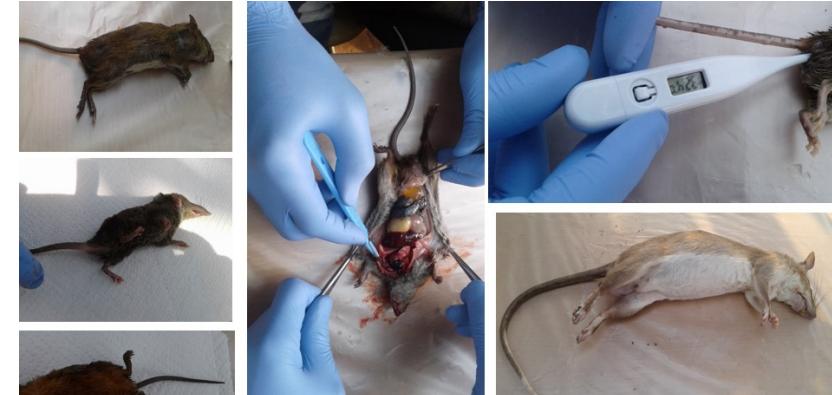
Animal samples: 250

2019 Toma outbreak

2021 Moloukou outbreak

2021 Grima outbreak

Rodents, duikers, squirrels,  
bats, pangolins



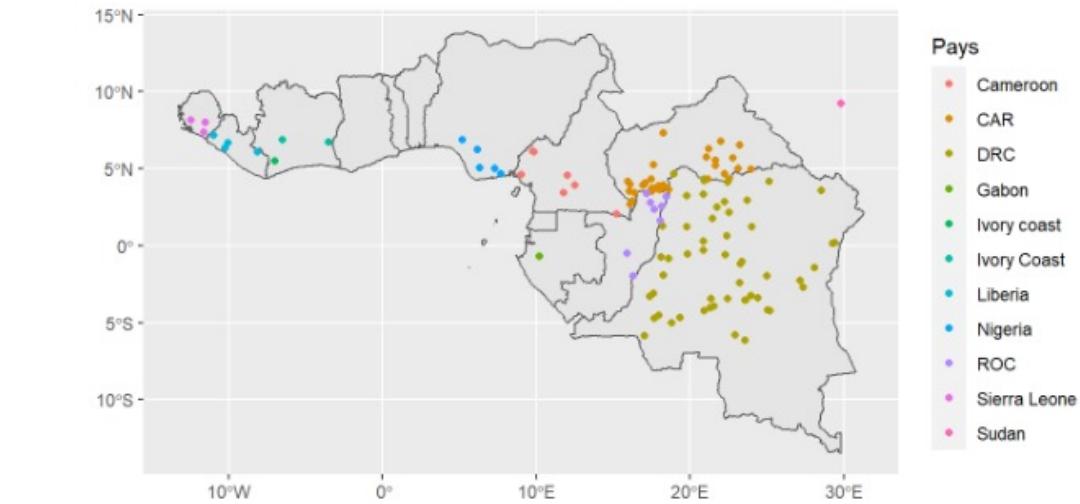
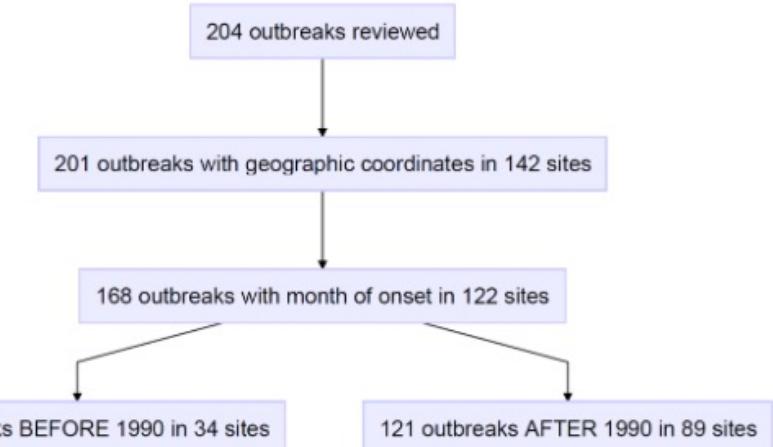
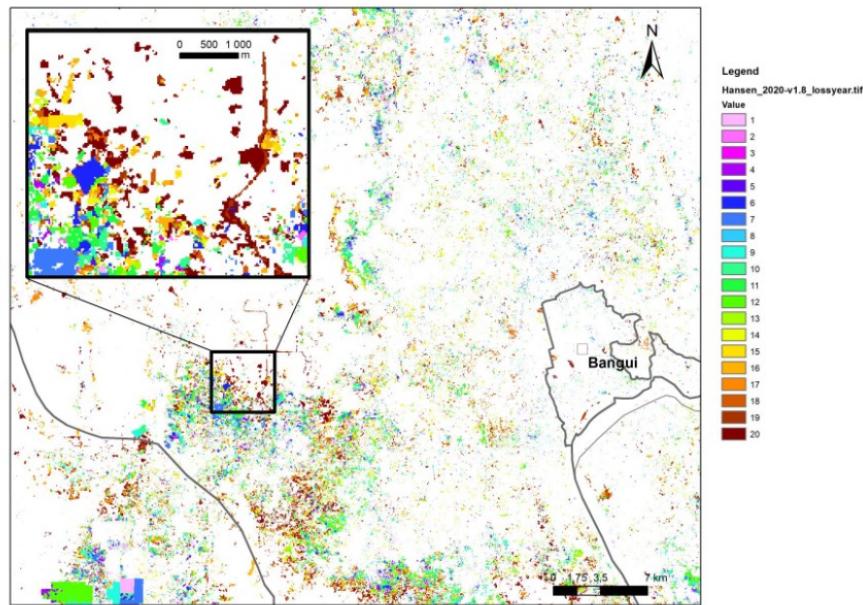
PCR  
Serological assay



# ECOLOGY

Environmental atlas  
Climate profile  
Human activities

DEFORESTATION



# ANTHROPOLOGY

- **Anthropological investigation** of monkeypox illness, diagnosis & treatment pathways, care: formal health care workers, traditional healers, former patients
- **Participatory investigation** of « local epidemiologies » (local understandings of origins, emergence, transmission)
- **Ethnohistorical** study of local ecological (forest, wildlife) & social changes implied in monkeypox emergence in the CAR since 1970
- **Ethnoecology:** local conceptions and observations of wildlife; current practices with wildlife



# Collaborators

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