

# Exploring trait-based approach for understanding and approximating pathogen spillover systems

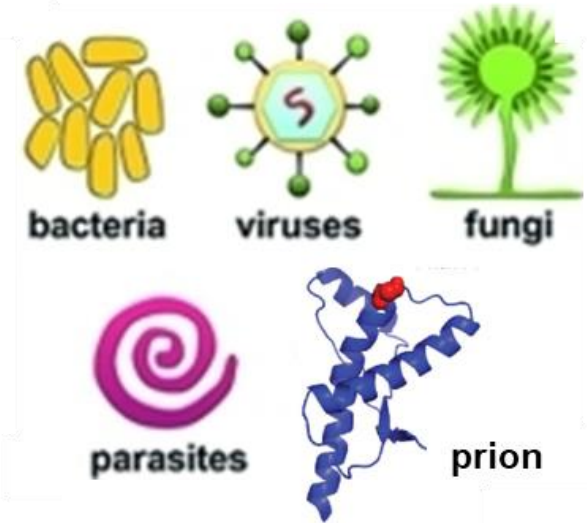
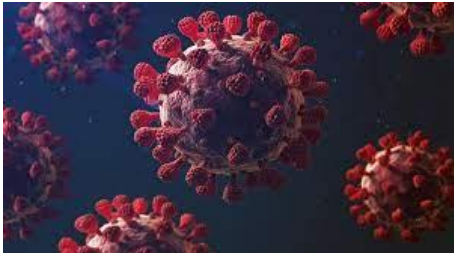
*NORDITA. Unifying the epidemiological and evolutionary dynamics of pathogens. May 29 – June 22. Stockholm. Sweden*

Marina Treskova

05 June 2023

# Zoonoses

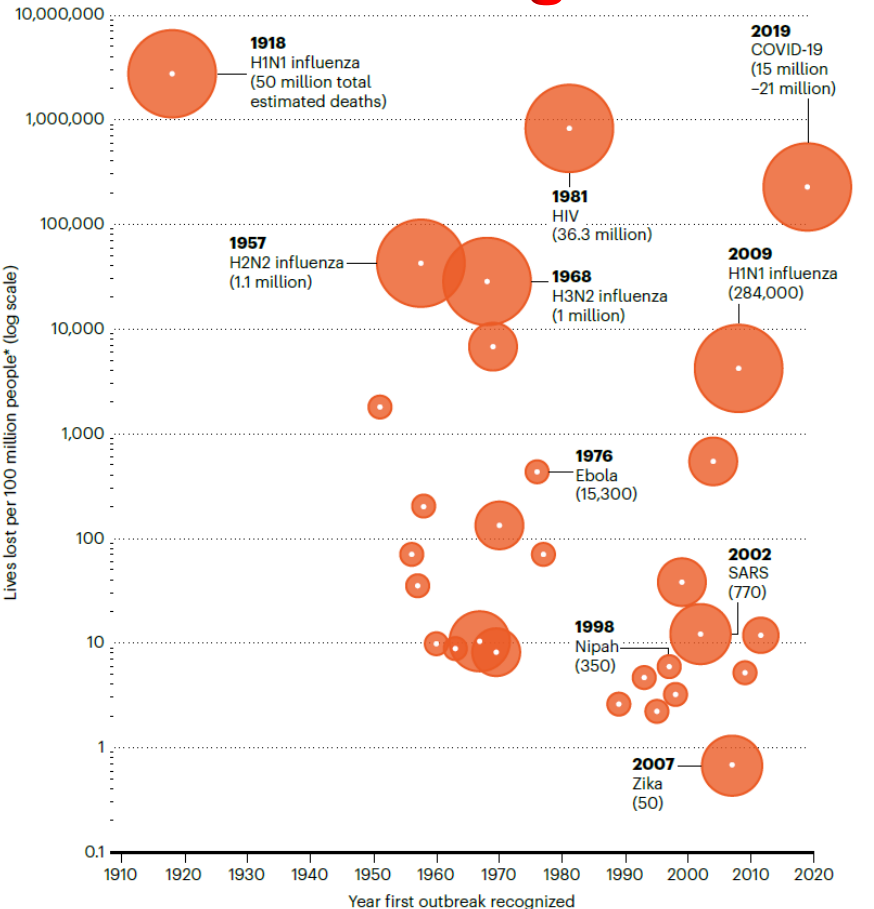
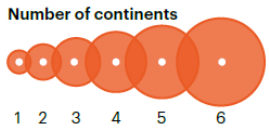
- Infectious disease caused by pathogens transmissible from a non-human animal to humans



# Disease emergence is strongly associated with ecological disturbance

## SPILLOVERS: A GROWING THREAT

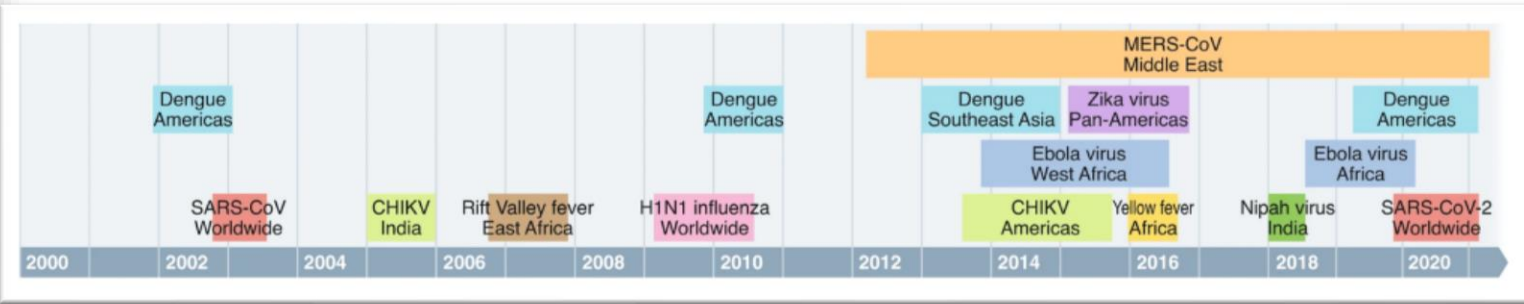
Deforestation and other changes have increased the likelihood of animal viruses jumping into people, with globalization and a higher density of human populations having increased the chance that such spillover events will be catastrophic. The annual economic loss from viral zoonoses since 1918 is US\$212 billion.



\*Data are from viral outbreaks of zoonotic origin that resulted in ten or more deaths; figures in parentheses are total estimated deaths spread over multiple years (and over multiple outbreaks, in some instances) rounded to the nearest ten.

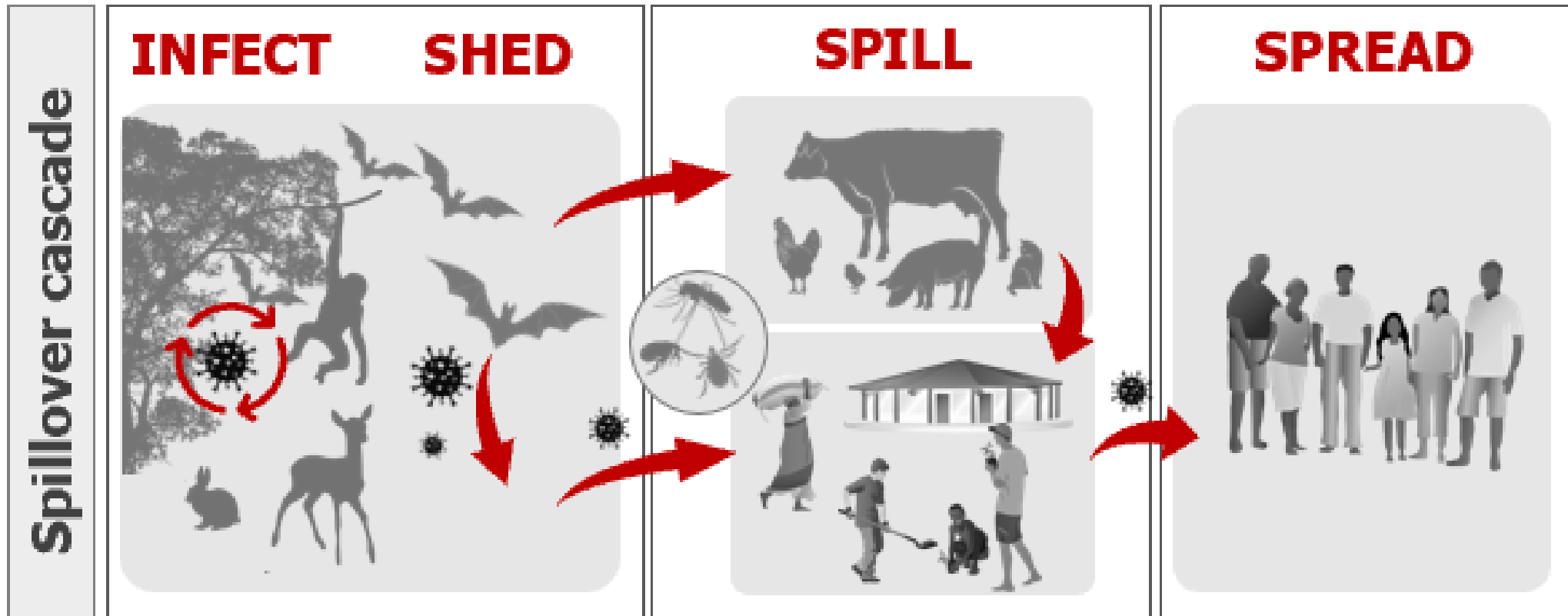
Source: Vora et al (2022)

The current issue is **increasing frequency** of disease emergence rather than the fact of it in humans



Timeline of twenty-first-century viral outbreaks, from 2000 to the present day. Source: Meganck et al (2022)

# Spillover is a key step in zoonotic disease emergence



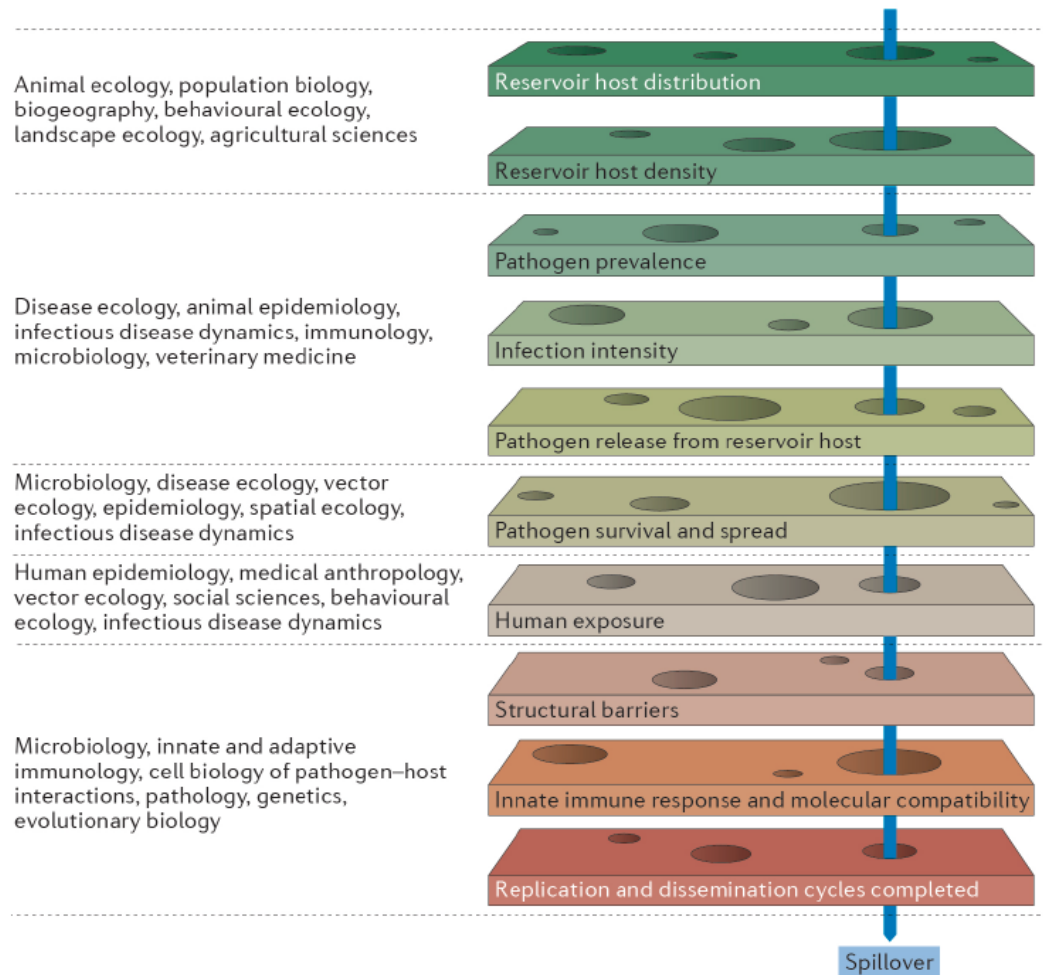
Spillover is defined as direct or indirect cross-species pathogen transmission from

- 1) source host to recipient host;
- 2) source host to vertebrate intermediate host to recipient host;
- 3) source host to invertebrate intermediate host to recipient host; and
- 4) source host to the environment to the recipient host.

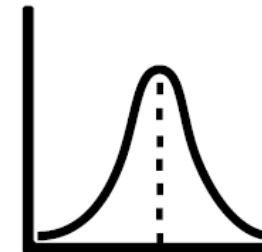
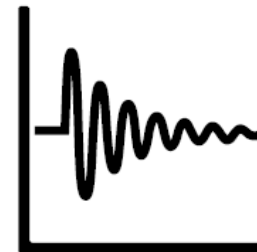


# Zoonotic spillover: barriers in space and time

Pathogen pressure -> dose and route of pathogen exposure -> probability of infection



- Wildlife/Reservoir-host ecology.
- Animal and human behaviour.
- Within host processes and viral evolution.
- Zoonotic and human transmission.
- Strategies for spillover/pandemic prevention.

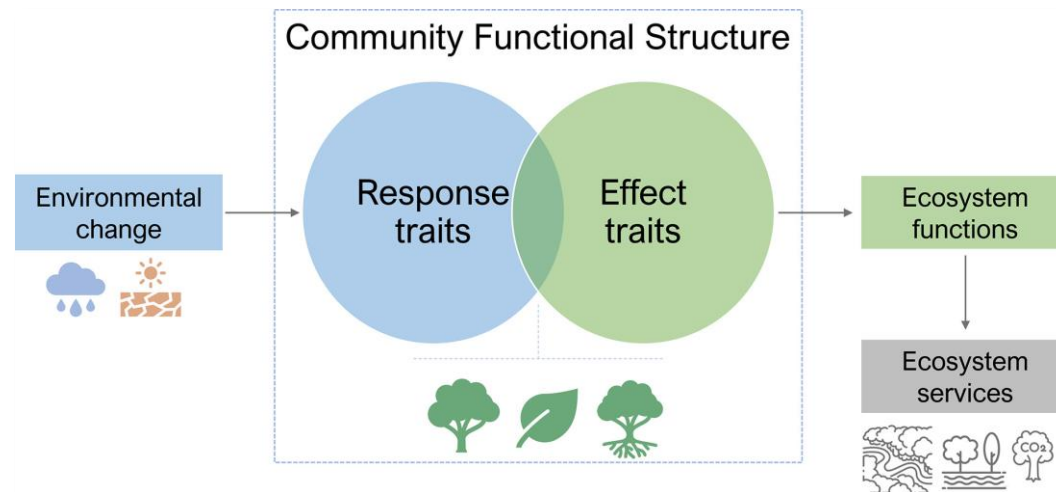


# Trait-based approach

- *Objective:*
  - understand and predict changes in patterns of transmission potential driven by environmental change and intervention
  - support evidence-based management and planning decisions
- *About:*
  - Comes from: functional diversity approach of community ecology and conservation biology developed over the last decades
    - Identifying key functional attributes of organisms (size, shape, nutrient requirements, etc.) that are important for community structure and function
  - Group hosts based on having a set of attributes not as belonging to certain species
  - -> Models with hosts communities represented as trait distributions without regard to species identity

# Functional traits

- Functional traits: any morphological, physiological, behavioral, or phenological characteristic that,
  - *impacts the fitness* of an organism
  - reflects *interactions with the environment* either:
    - mirroring the impact on some ecosystem process **effect traits**
    - and/or by responding to environmental changes **response traits**
    - Response and effect traits can overlap to different degrees, from being closely correlated to being random.



# Example of trait topology for primates

Ecological traits of the world's primates

Home range size



Locomotion type



Diel activity



Trophic guild



Body mass



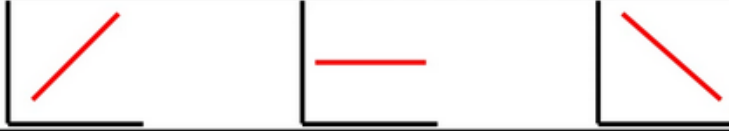
Habitat type



Conservation status



Population trend



Geographic realm



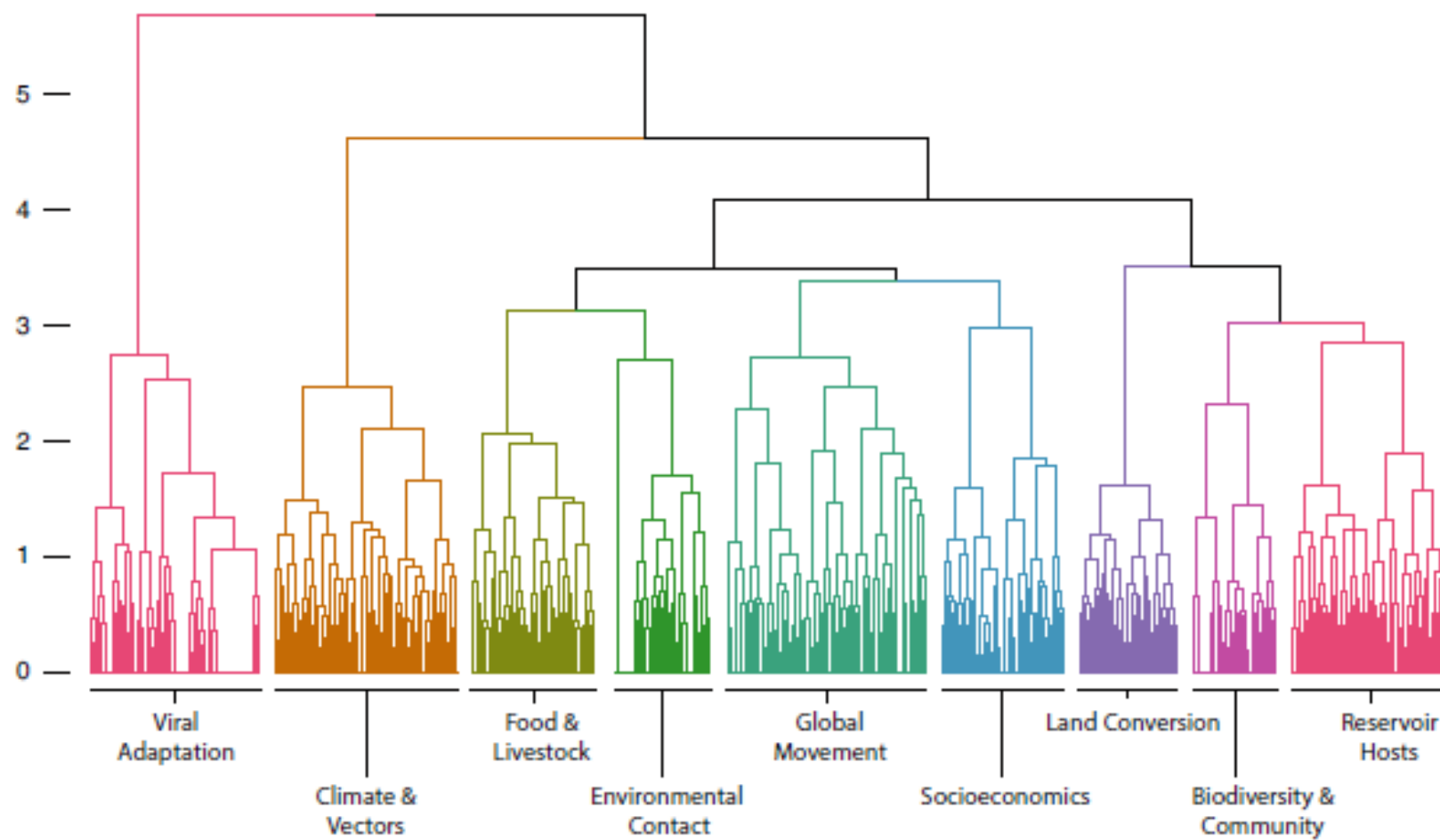
Source: Galán-Acedo, C., Arroyo-Rodríguez, V., Andresen, E. *et al.* Ecological traits of the world's primates. *Sci Data* **6**, 55 (2019). <https://doi.org/10.1038/s41597-019-0059-9>



# Opportunities for trait-based approaches

- Advantages for ecosystem model of infectious disease:
  - Ecological questions associated with community assembly and habitat filtering and biodiversity patterns along environmental gradients, and ecological processes
  - Considers within-species variation
  - Approach to scale down parameter space for multi-host pathogen systems
  - Allows merging of trait-based models with network modeling to understand multi-host transmission across space and time and effects of environmental drivers within a process-based model
  - Results have the potential for across-community generality

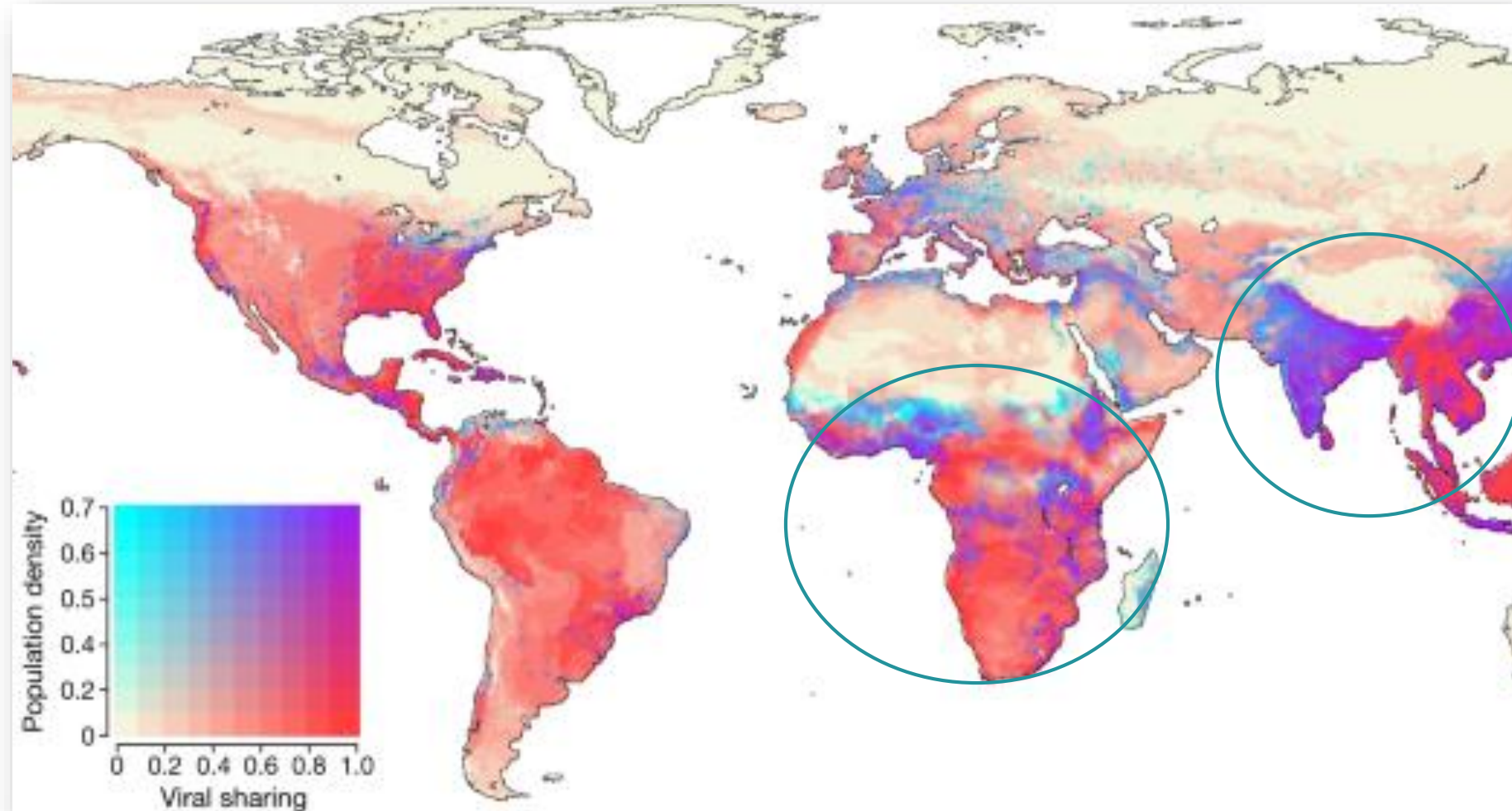
# Anthropogenic global environmental change relevant for spillover



- Climate change
- Land use
- Habitat fragmentation
- Urbanization
- Deforestation
- Pollution
- Biodiversity loss

# Mammals adaptation can change the virome and drive new zoonoses

- In 2070 species will aggregate in new combinations at high elevations, in biodiversity hotspots, and in areas of high human population density in Asia and Africa
- Cross-species transmission of their associated viruses to increase an estimated **4,000 times**



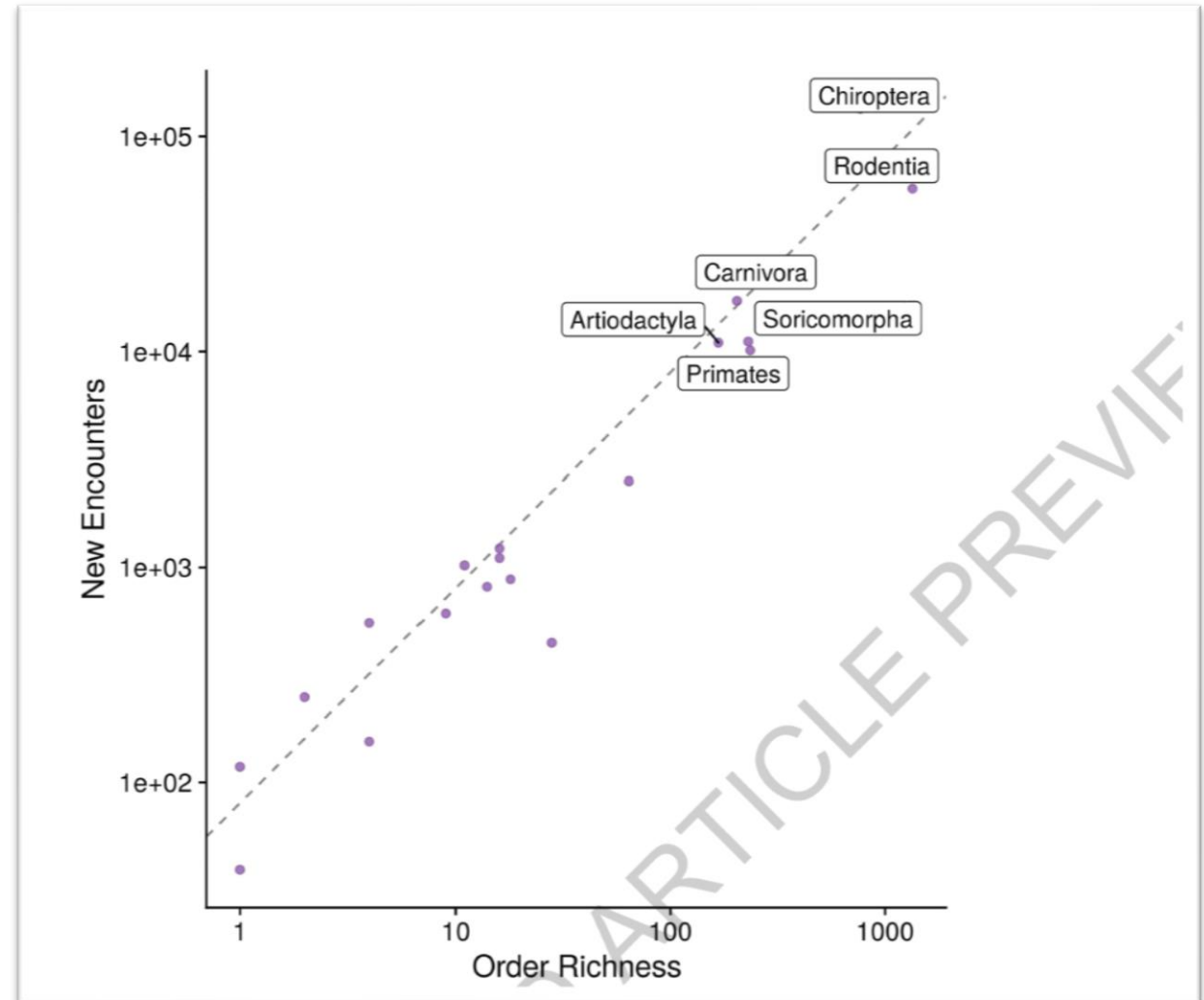
Source: Carlson et al (2022)

# Climate change increases cross-species viral transmission risk (Carlson et al.)

- Climate change linked to cross-species viral transmission
- Environmental change alter mammal communities in ways that expose hosts to novel viruses, altering the structure of the whole mammal-virus network
- Modelling: species distribution models and viral sharing model
- Cross-species transmission depends on *opportunity* and *compatibility*
- Opportunity:
  - Ecological or geographical overlap among the hosts
  - Encounter:
    - Potential interactions (direct contact) depend on behavioral characteristics
    - Transmission via vector (indirect contact) requires spatial proximity
- Compatibility:
  - Given a contact a virus requires certain host characteristics to be able to enter cells and reproduce
  - -> compatibility w.r.t. host cell receptors and immunity
- Because closely-related species share both ecological and immunological traits through identity by descent, *phylogeny is a strong predictor of pathogen sharing and of susceptibility to invasion by new viruses*

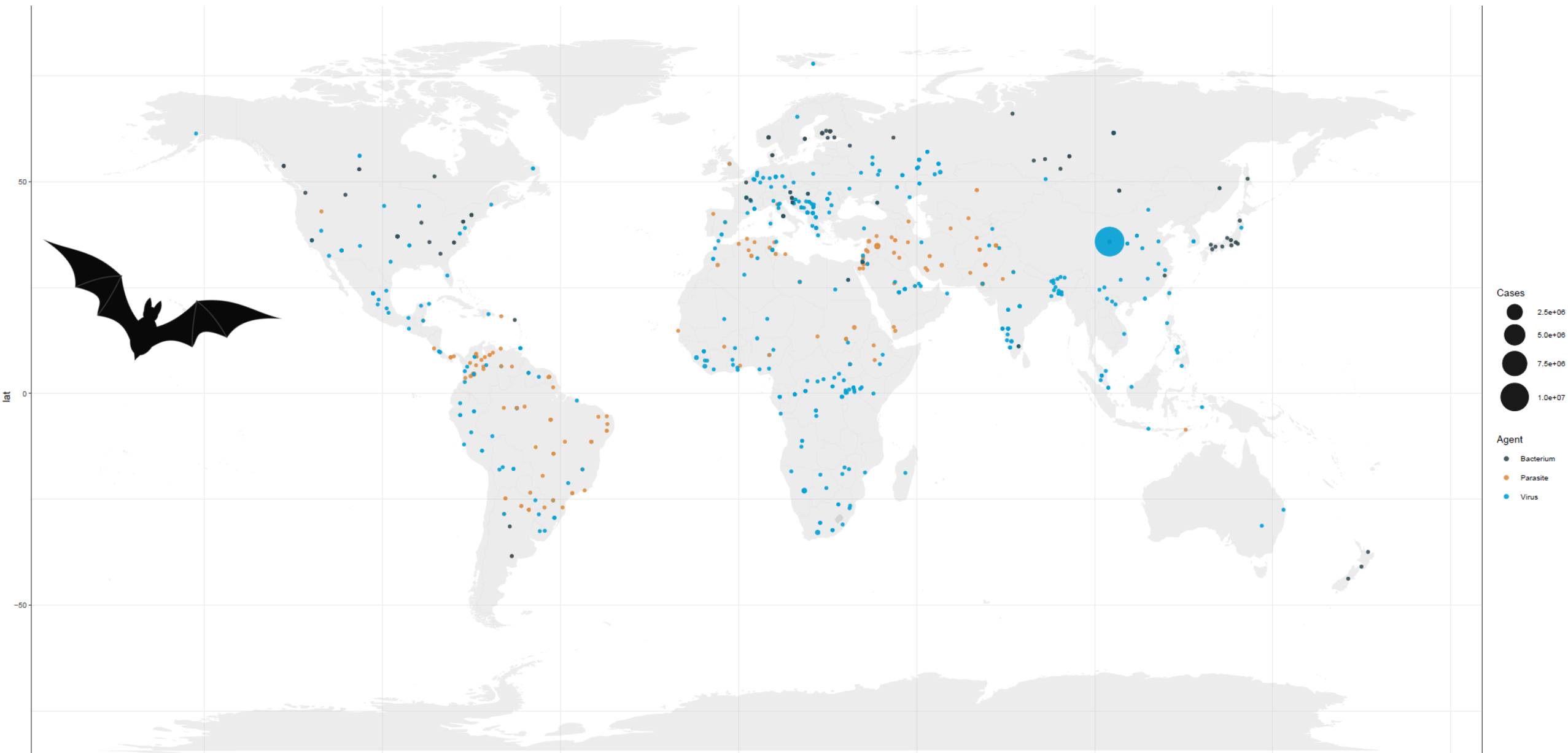
# Animal dispersal capacity as a driver

- **Animal dispersal capacity play a role** in first encounters and novel viral sharing.
- Trophic position and body size determine dispersal capacity
- **Viral sharing hotspot is driven disproportionately by bats.**
- Are some animal groups more common sources of zoonotic viruses than others ?
  - pose a greater zoonotic and pandemic risk?
- Why can some mammals be key hosts for zoonotic viruses?
  - Close proximity: live near human settlements, act as food sources
  - Closely related to humans that viruses face little challenge in host adaptation

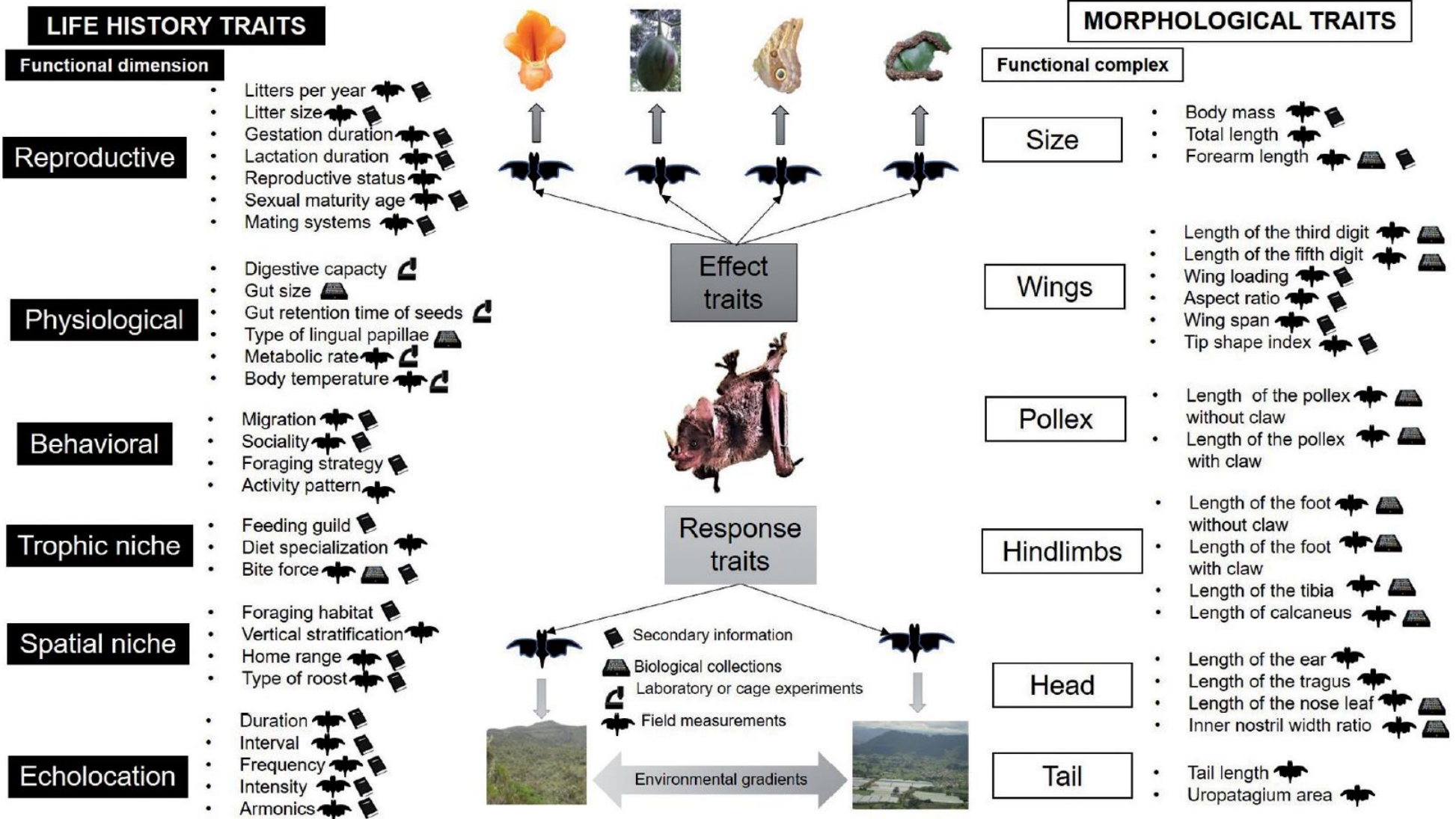




# GIDEON: Infectious diseases with bats stated in reservoir host designation

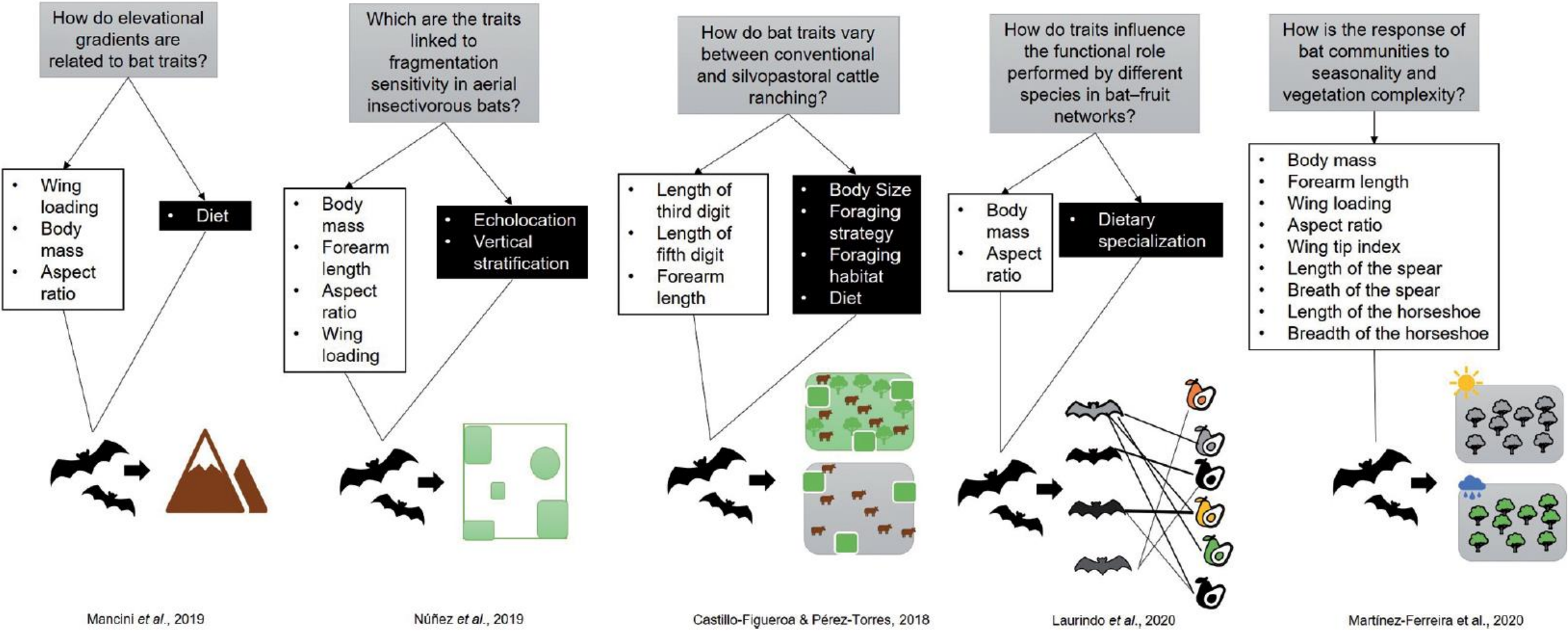


# Schematic model of biological traits of Neotropical bats



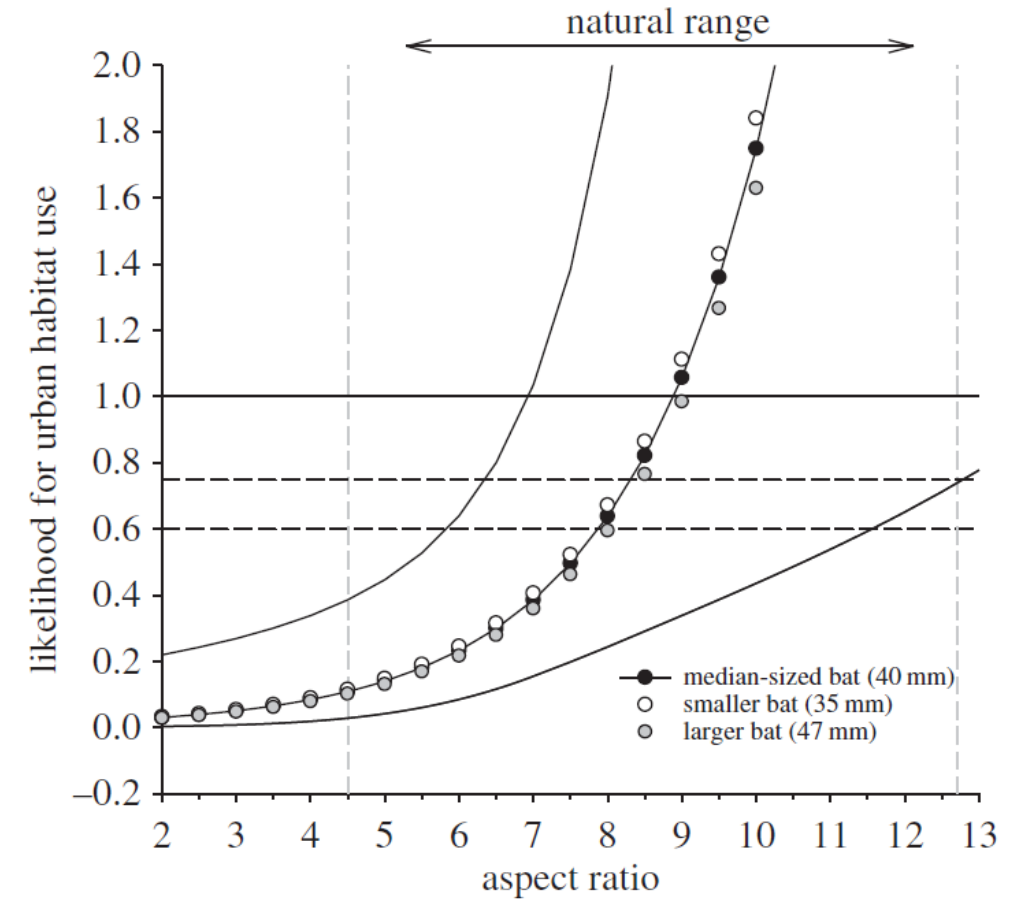
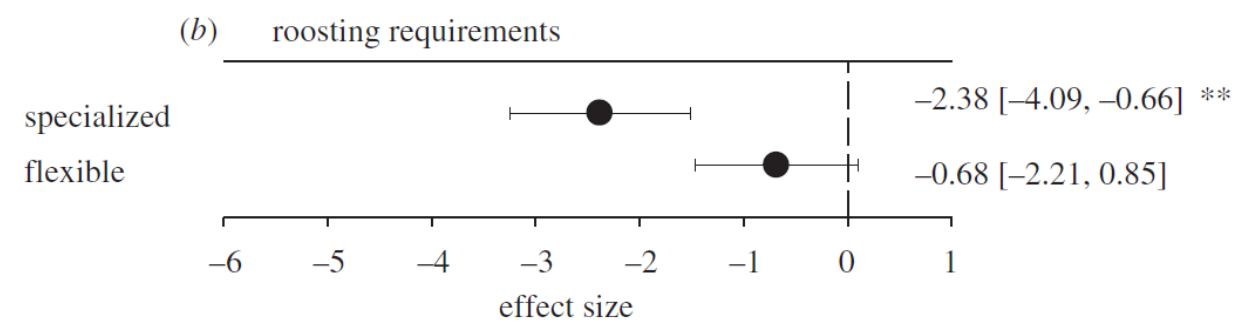
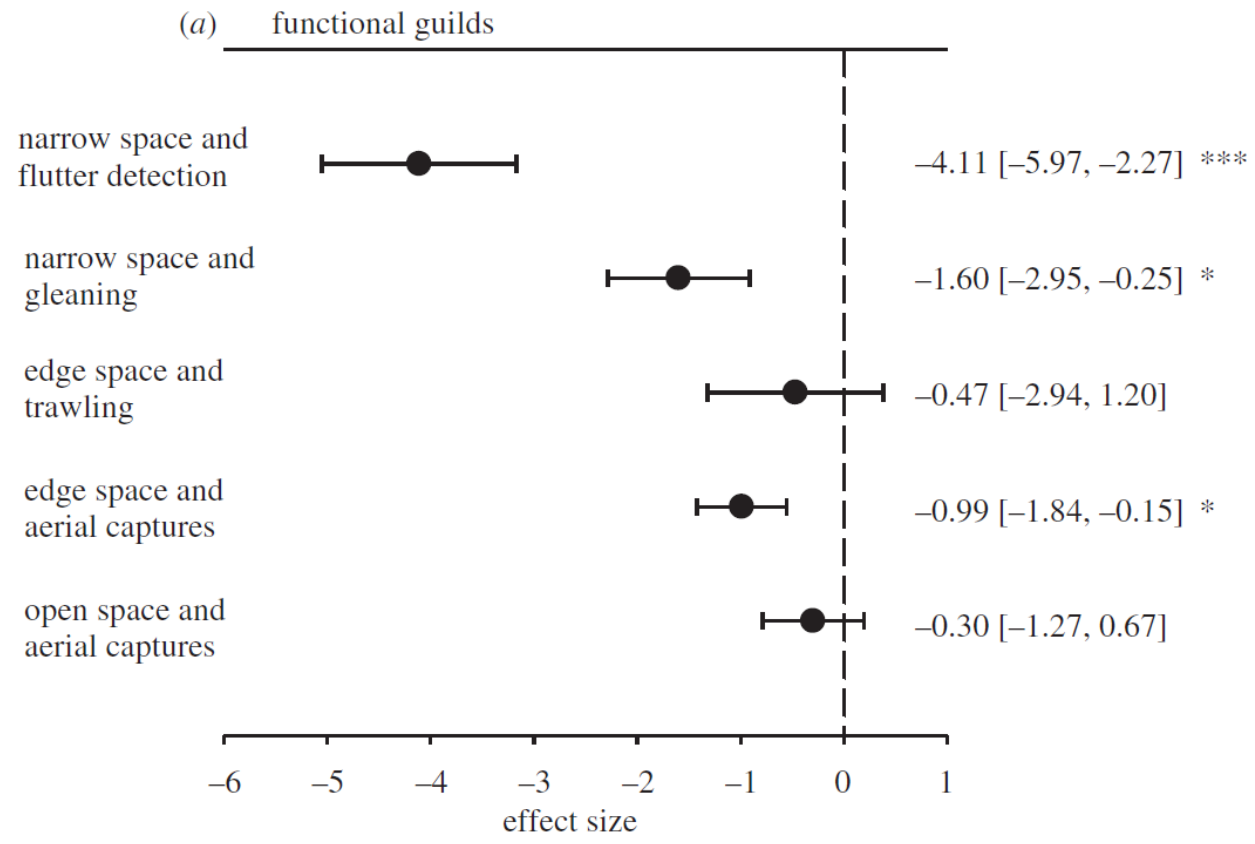
Source: Castillo-Figueroa & Pérez-Torres (2020). <http://doi.org/10.11606/1807-0205/2021.61.24>. Life history traits in black boxes whereas morphological traits are depicted in white boxes.

# Study cases using trait-based approach in Neotropical bats



Source: Castillo-Figueroa & Pérez-Torres (2020). <http://doi.org/10.11606/1807-0205/2021.61.24>. Life history traits in black boxes whereas morphological traits are depicted in white boxes.

# Trait-dependent tolerance of bats to urbanization



Source: Jung K, Threlfall CG. 2018  
<http://dx.doi.org/10.1098/rspb.2018.1222>.





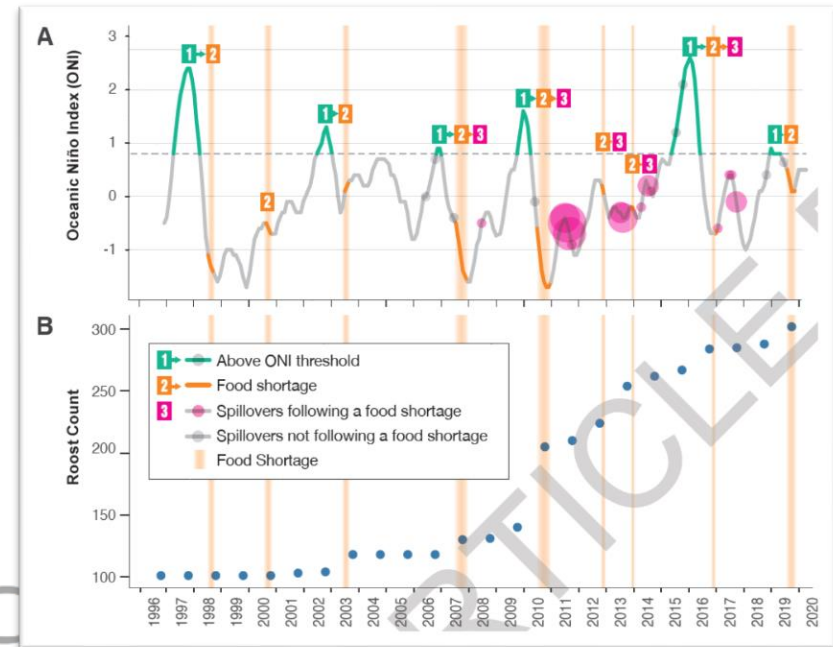


# Show-case: Hendra virus spillover in subtropical Australia (Eby et al 2022)

- Hendra: Australian flying foxes
  - transmitted to horses (the intermediate host) causing a highly lethal disease
  - capable of spilling over from horses to humans => a severe or fatal infection.
- interactions between land-use change and climate altered the behavior of wildlife reservoir hosts, increasing their proximity to domestic or human recipient hosts
  - spillover was coming from small populations of bats that had just formed in agricultural areas
- the highest risk of spillover was from bats in agricultural areas after a food shortage when no winter habitat produced food (nectar)
- potential solution to stop spillover is to restore the winter habitat of the bats and preserve what is left



Source: <https://www.newsweek.com>



- Can traits be a useful tool in rapid assessment and monitoring of changes in infectious disease due to anthropogenic ecological disturbance?
- Can they become indicators?
- Which bat traits are good predictors of sustainable ecological and conservation practices that provide reduction of spillover risk for people and livestock?

# Thank You!



—CSIDLlab—

Climate-Sensitive Infectious Diseases lab