# Covid and Wildlife

Slides (mainly) by (soon to be) Dr. Tricia Fry





### Covid and Wildlife Management

- Why?
- What we know?
- What don't we know?
- What is next?



**Pathogen:** Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)

### What we know...

Originated in bats, with evidence of bridge host being responsible for spillover into humans.

Rapid mutation.



Horseshoe bat (Rhinolophidae)



Bat

Sunda pangolin (*Manis javanica*)



Lytras et al. (2017) Science 373, Issue 6558 pp. 968-970 DOI: 10.1126/science.abh0117

### What we know...

Species that have been infected

- Domestic animals
- Felids
- Mustelids
- Rodents
- Bats
- Deer

Environmental contamination possible



American Mink (Neogale vison)



### Why do we care if SARS-CoV-2 is in wildlife?

- Understanding origins
- Does it cause pathologies?
- Spillback –transmitting infection back to a potential maintenance host
- Spillover transmission between new species
- 'New and improved' viruses





#### What we don't know....



A lot....

### So, we do

- Lab research including experimental infections
- Genomic analysis
- Think about population densities and risks
- Systematic Surveillance actively survey tissue and samples from a variety of species in many locations
- Risk Assessment and Structured Decision Making

#### Surveillance Strategies



#### World Organisation for Animal Health Founded as OIE









## What's next's...

- SARS-CoV-2 pandemic continues to be driven by human-to-human transmission with no evidence that domestic or wild animals are playing an important role, but ....
- reservoir of infection in a wild animal population poses a significant risk to public health if it had potential to spillback into communities the burden of infection had been reduced through con
- opportunity for evolutionary adaptation of the virus, which could potentially (positively or negatively) influence transmission dynamics and the effectiveness of diagnostics and vaccines
- Pay attention and learn for the future







### Ecology suggests that diseases will continue to become more prevalent

- Increased population density
  - more rapid transmission, selection against reduced virulence
  - more frequent invasion of new habitats  $\rightarrow$  more epizootics
- Widespread movement → more rapid transmission of new diseases
- Evolution of drug resistance across pathogen strains, antibiotic resistance leads to greater virulence
- Climate change → increased ranges of pathogens ad their vectors