

Emerging Pathogens and the Role of Natural History Archives: The Hantavirus Example

Emerging Pathogens

- newly appearing or spread of infectious diseases
 - unrecognized/underreported due to
 - ignorance of clinical expression
 - lack of reliable test
 - often associated with outbreak scenarios
- 1400 species known to be pathogens

Zoonotic Diseases: transmitted from animals to humans

Why are there more emerging diseases?

- there is an increase in human-animal contact
 - increase and spread of human population (2x the population since 1999)
- recognition of more pathogens

Pop-Growth Consequences

- need for space
 - sprawl
 - encroachment on natural areas
- need for food
 - habitat conversion

Habitat Conversion

- 50% of habitable land converted to farming
 - includes land with high biodiversity
- large-scale clearing for intensive monocultures
- human into natural areas
- decrease in biodiversity

Outbreak Scenarios

Ebola Example:

- reservoirs of long lived fruit bats
- maintain pathogen for long periods in natural areas
- human contacting virus in natural world

Hantavirus:

- reservoirs of short-lived rodents
- responds rapidly to environment change
- 2 types-rotates in natural and converted lands
- human to animal transmission

Initial 1999 Outbreak

- unknown disease emerged in SW killing 10 people in 8 weeks
- victims experienced flu-like symptoms
- caused the death of healthy individuals caused an initial level of panic

What was the disease agent?

- conspiracy theories
- massive collaboration
- virus isolation in human case
- reservoir of positive rodents that were carrying the virus

Why did this occur at this place and time?

- understanding reservoir ecology
- longitudinal studies
- 8 sites in US monitoring
 - small mammal population
 - hantavirus prevalence

Reservoir Studies lead to predictive models

- precipitation catalyst in SW → trophic scale
- delayed density-dependent rodent response

Today

- 639 cases
- mortality at 30-40%
- medical community recognition, treatment
- reservoir understanding

Many new Hosts for New Hantavirus

- new discoveries possibly with deep integrated specimen archives (moles, bats, etc)
- bats especially have a lot of contact with humans

Holistic Voucher

- build a large collection
- frozen tissue by UNM is the largest in the world, 20x the size of the Smithsonian's collection
- valuable samples for molecular biologists
- traditional specimen has skin, tag, and skeleton associated
- collection not only valuable to taxonomists
- tying data sets to survey ecology/evolution for these species
- many collections are digitized and made available online
- specimen is time-stamped to see window during the environmental conditions during the time the organism existed (geo-referencing)

Integrated Archives

- temporally deep (sampled every year)
- geographically broad, site intensive
- geo-referenced

Phylogenetic Tree

- virus tends to cluster with nonmammalian hosts
- virus co-evolving with the mammals for a long time

What went wrong?

- durable infrastructure lacking- deep/temporal and wide/spatial
- focus on hypothesis driven sciences
 - funding availability
 - narrower scope of new collections
- reluctance of museums to engage other communities
 - shift from field work to public programs in many museums' cases
- slow response of museums to build integrated resources
- communication as weak
- weak data models

Next Steps for Pathogen Discovery and Mitigation?

- information and technology needs
- increased growth of collections
- increased human capacity
 - broadening participation