

One Health In Action: From the **Field** to the Classroom



UC DAVIS

VETERINARY MEDICINE

Karen C. Drayer Wildlife Health Center



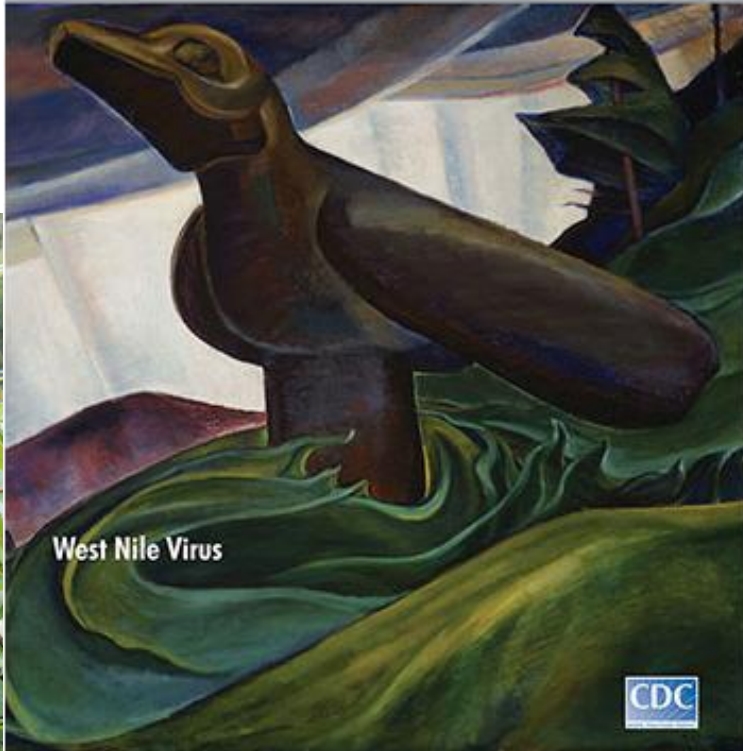
Majority of human infectious diseases are of animal origin

EMERGING INFECTIOUS DISEASES

A Peer-Reviewed Journal Tracking and Analyzing Disease Trends

EID
Online
www.cdc.gov/eid

Vol. 10, No. 8, August 2004



75% of emerging infectious diseases in humans have wildlife origins

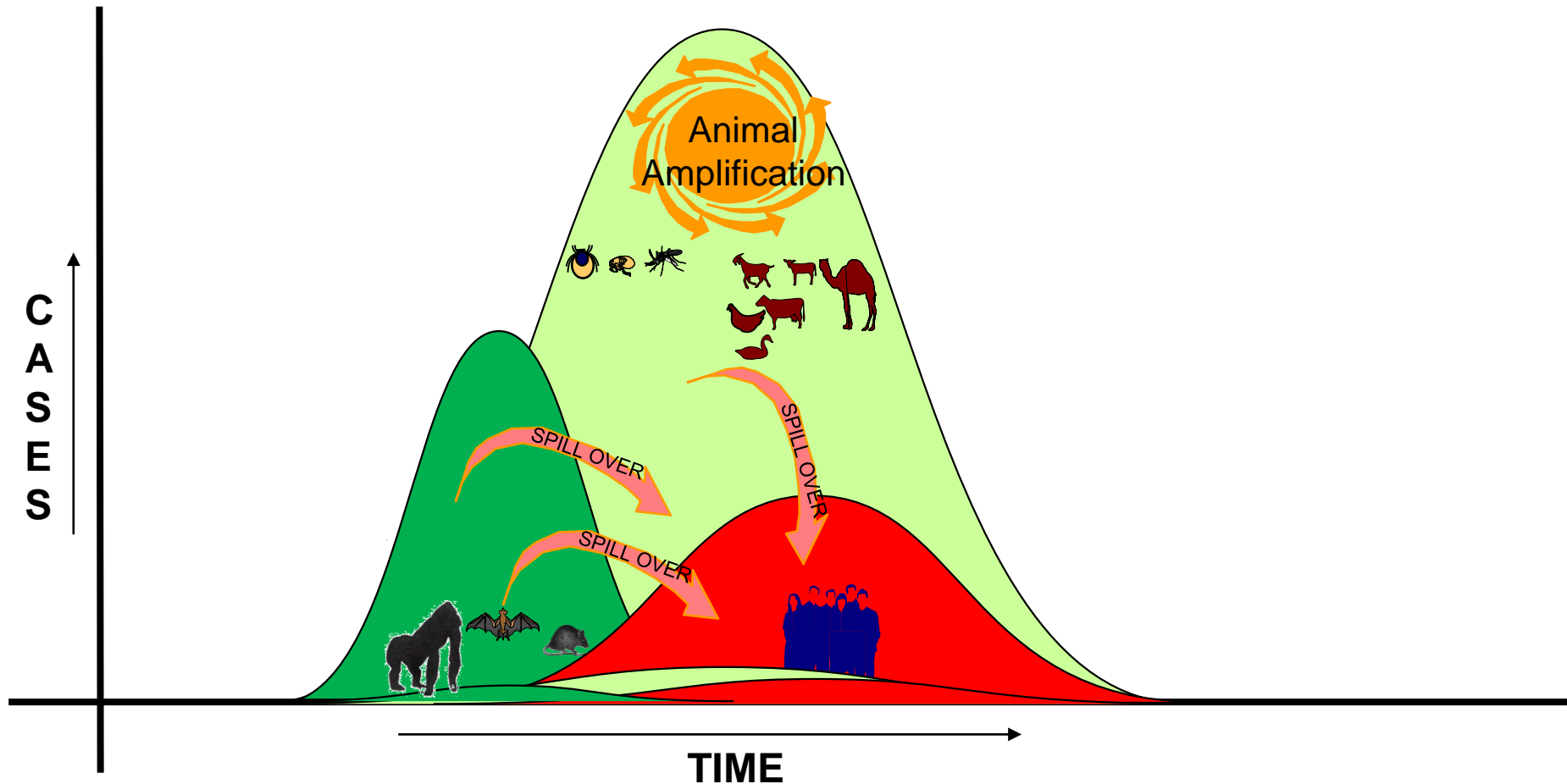
Human contact with wildlife has been linked to recent emergent diseases

Human population growth is highest near diverse wildlife areas

Zoo and wildlife veterinarians uniquely positioned to discover EIDs and mitigate the potential for pandemics



USAID Emerging Pandemic Threats PREDICT Project







P. Telfer, 2005



PREDICT Diagnostic Strategy

Targeted screening for different wildlife taxa

Family level primers

primates

Retro, Filo,
Flavi,
Influenza,
Paramyxo,
Henipa, Pox,
Bunya,
Herpes,
Seadorna,
Astro, Corona,
Arena

bats

Flavi, Corona,
Henipa,
Paramyxo,
Rhabdo,
Arena, Filo,
Reo,
Influenza,
Bunya,
Seadorna,
Astro

rodents

Arena,
Hanta, Pox,
Alpha, Flavi,
Reo,
Paramyxo,
Influenza,
Bunya,
Seadorna,
Astro

human

Retro, Filo,
Flavi,
Influenza,
Paramyxo,
Henipa, Pox,
Bunya,
Herpes,
Seadorna,
Astro,
Corona,
Arena, Hanta,
Entero, Lyssa

birds

Orthomyxo,
Paramyxo,
Flavi

Refine with specific primers or sequencing
Advanced pathogen discovery

20 PCR protocols
developed and going out
to countries

Follow-up on-the-ground field investigations
develop diagnostics
screen local human cases
screen local wildlife



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PREDICT



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METABIOTA



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Institution



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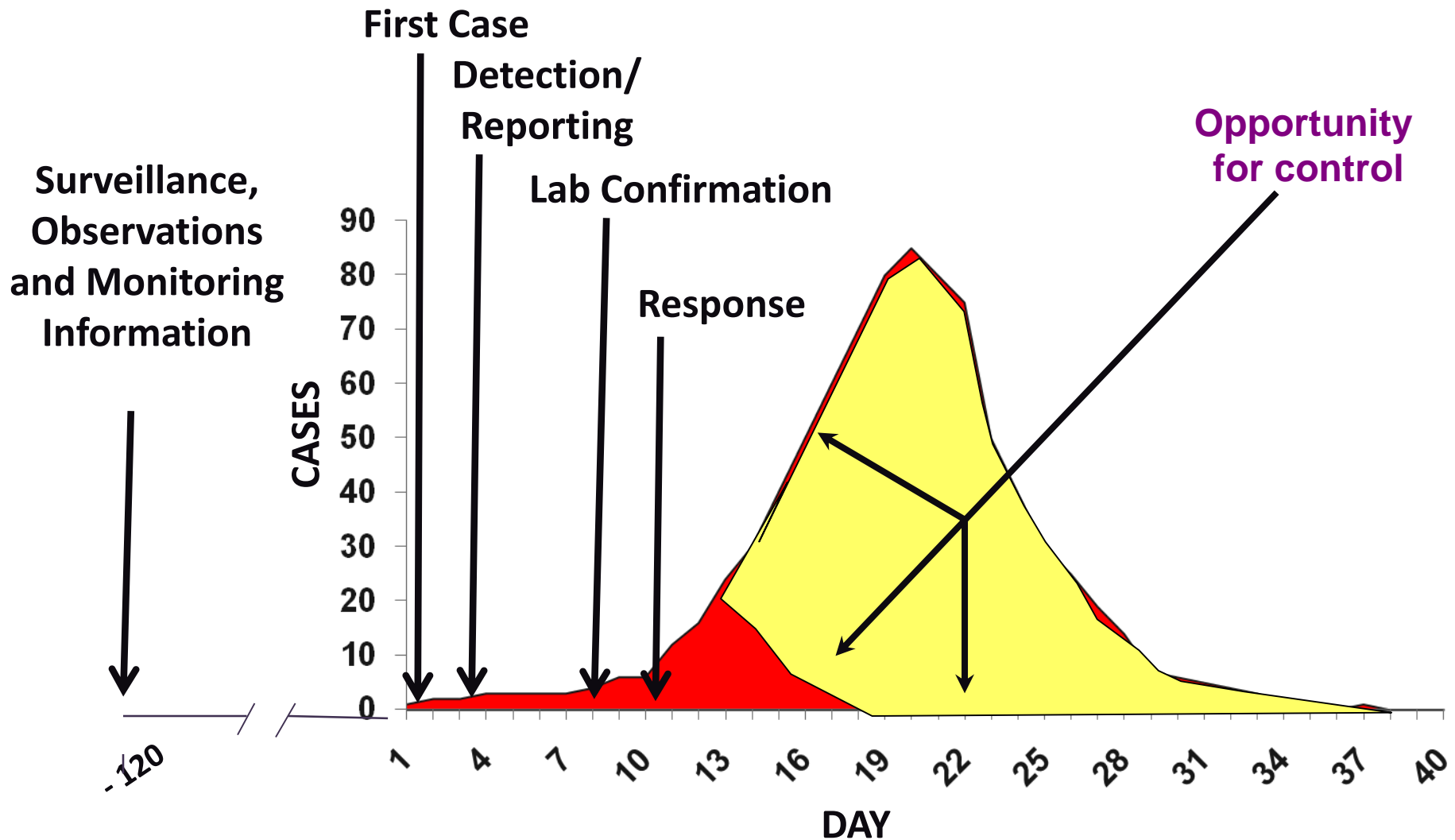


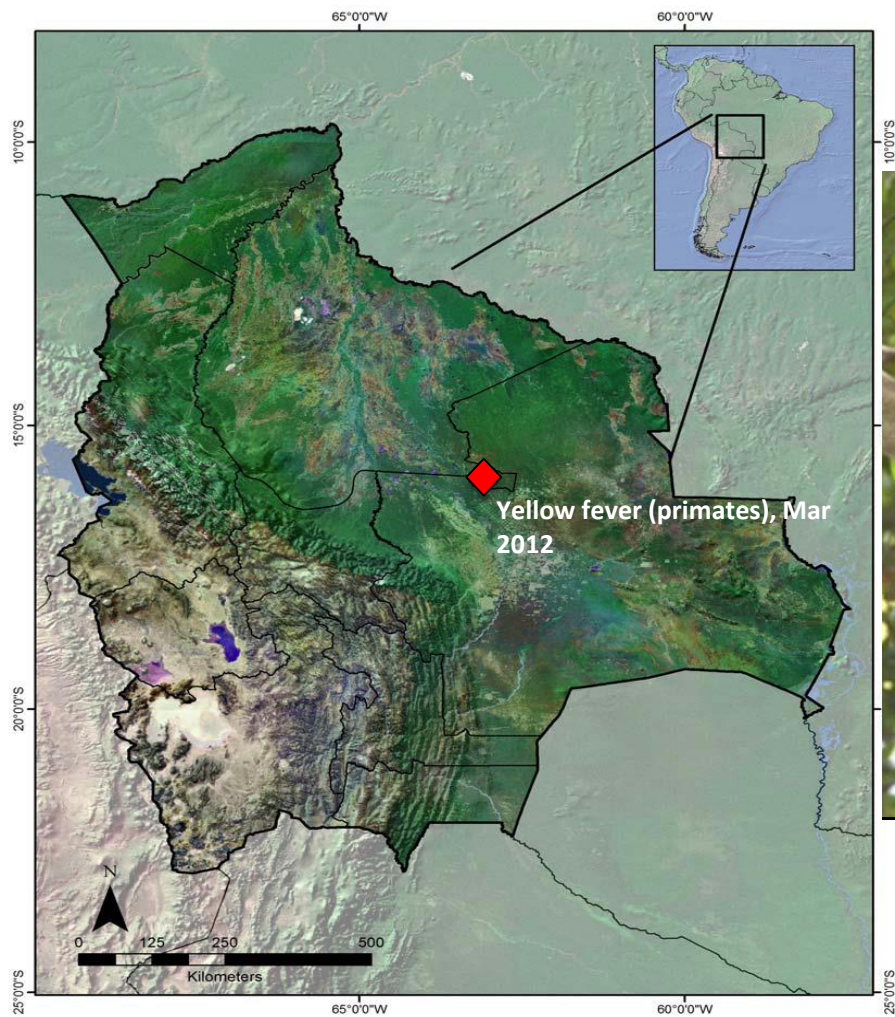
METABIOTA



Smithsonian
Institution

- Trained >2,000 field personnel, veterinarians, laboratory technicians, public health workers from 20 countries in 59 ministries
- Building capacity to test for viral families in 33 labs
- Collected samples from 50,000 animals
- Discovered >250 novel viruses in wildlife (e.g. corona, retro, adeno, rhabdo, herpes, boca)
- Documenting human pathogens in wildlife and animal-origin pathogens in humans









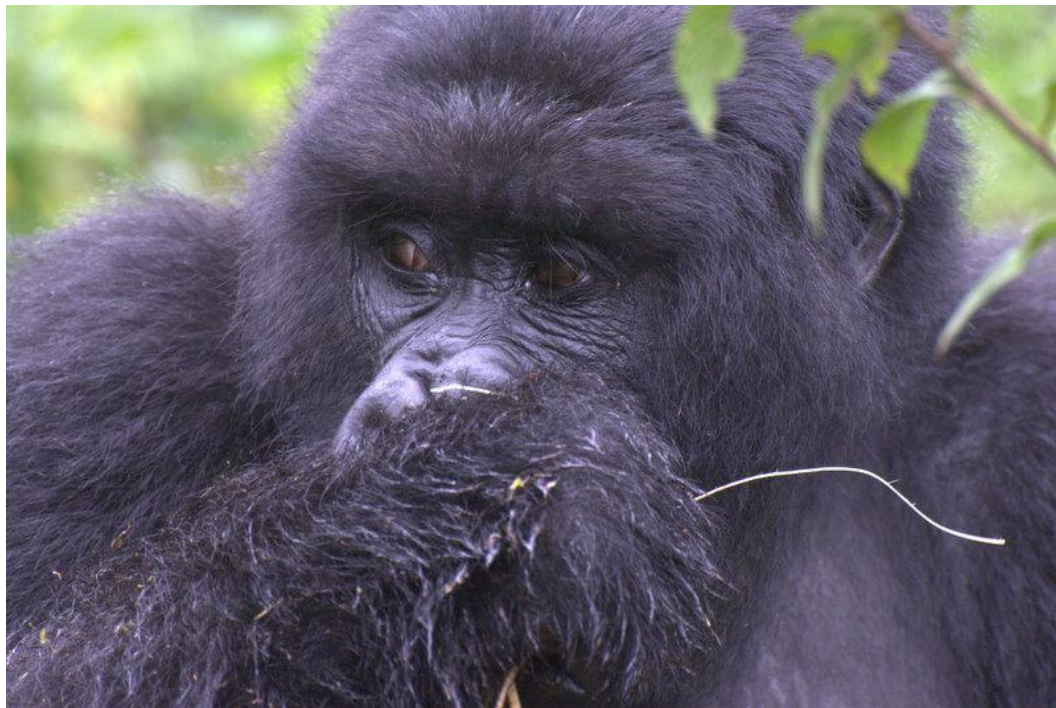






**11,370,425 people;
1,087 people/sq. mi**

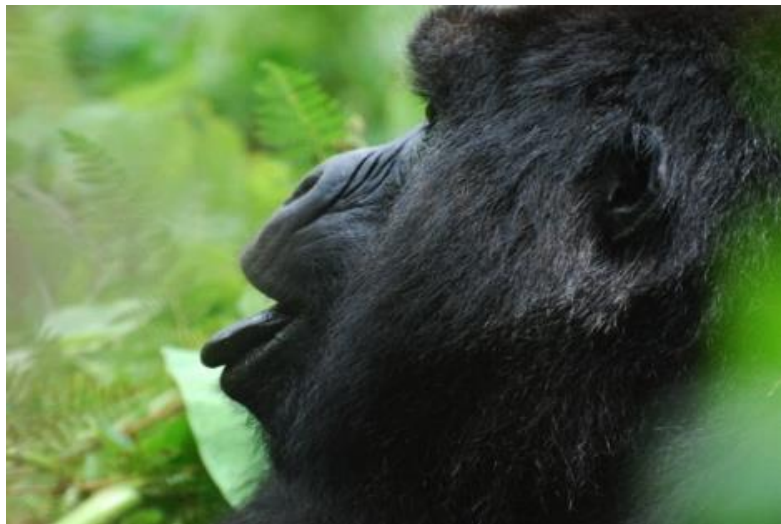




Human Metapneumovirus Infection in Wild Mountain Gorillas, Rwanda

**Gustavo Palacios, Linda J. Lowenstine,
Michael R. Cranfield, Kirsten V.K. Gilardi, Lucy
Spelman, Magda Lukasik-Braum,
Jean-Felix Kinani, Antoine Mudakikwa,
Elisabeth Nyirakaragire, Ana Valeria Bussetti,
Nazir Savji, Stephen Hutchison, Michael Egholm,
and W. Ian Lipkin**

The genetic relatedness of mountain gorillas and humans has led to concerns about interspecies transmission of infectious agents. Human-to-gorilla transmission may explain human metapneumovirus in 2 wild mountain gorillas that died during a respiratory disease outbreak in Rwanda in 2009. Surveillance is needed to ensure survival of these critically endangered animals.



Gorilla Conservation Employee Health Program

- Sensitization, consent forms
- Logistics (site, transportation)
- Pre-exam questionnaire
- Physician examination
 - CBC
 - TB intradermal test
 - HIV
 - Measles titer
- Laboratory testing



Gorilla Conservation Employee Health Program

- +/- vaccination: DPT, polio, MMR, tetanus
- De-worming and health/hygiene education
 - Quarterly, including families
 - Mebendazole
- Follow-up

2012/13

1,850 individuals





DOI: 10.7589/2012-09-229

Journal of Wildlife Diseases, 50(1), 2014, pp. 21–30
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GIARDIA IN MOUNTAIN GORILLAS (*GORILLA BERINGEI BERINGEI*), FOREST BUFFALO (*SYNCERUS CAFFER*), AND DOMESTIC CATTLE IN VOLCANOES NATIONAL PARK, RWANDA

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Extreme Conservation Leads to Recovery of the Virunga Mountain Gorillas

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