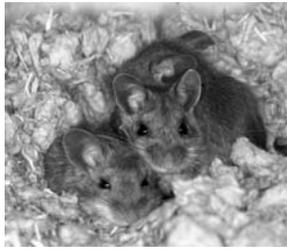
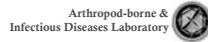


Hantaviruses

What can rodent reservoirs teach us about human disease?



Tony Schountz, PhD
Arthropod-borne and Infectious Diseases Laboratory
Department of Microbiology, Immunology and Pathology
College of Veterinary Medicine and Biomedical Sciences
Colorado State University



Reservoirs of Zoonotic Agents

- To perpetuate, viruses must:
 - Persistently infect their reservoirs without
 - substantial pathology
 - eliciting a sterilizing immune response
 - Infect another susceptible host before
 - the immune response controls the virus
 - the host dies

Hantaviruses

- Negative stranded RNA viruses
- Global distribution
- Enveloped
- Trisegmented
 - S - nucleocapsid (NSs?)
 - M - Gn and Gc glycoproteins
 - L - RNA-dependent RNA polymerase
- Zoonotic reservoirs (no pathology)
 - Rodents
 - Shrews Pathogenic hantaviruses have only been found in rodents
 - Moles
 - Bats
- Reservoirs remain infected, perhaps for life, despite an immune response
- In rodent reservoirs a regulatory T cell response occurs (Easterbrook et al., PNAS, 2007; Schountz et al., PNAS, 2007)



Factors Governing Emerging and Re-emerging Infectious Diseases

- Human demographics and behavior
 - Population growth, density and distribution
 - Immunosuppression
 - Sexual activity and substance abuse
- Technology and industry
 - Modern medicine
 - Food processing and handling
 - Water treatment
- Economic development and land use
 - Dam building
 - Reforestation
 - Climate change
- International travel and commerce
- Microbial adaptation and change
 - Natural variation and mutation
 - Selective pressure and development of resistance
- Breakdown of public health measures
 - Inadequate sanitation
 - Complacency
 - Vaccination rates

Adapted from Nathanson, Viral Pathogenesis and Immunity, 2007

Hantavirus Disease

- Two diseases with many similarities
 - Hemorrhagic Fever with Renal Syndrome (Eurasia)
 - Hantavirus Cardiopulmonary Syndrome (Americas)
- Both are thought to have immunopathologic components
 - No virus damage to the vasculature
 - Pronounced inflammatory immune response 10 to 35 days post exposure
 - The immune response is thought to be a significant contributor to hantavirus disease
- About 200,000 cases and about 10,000 deaths per year

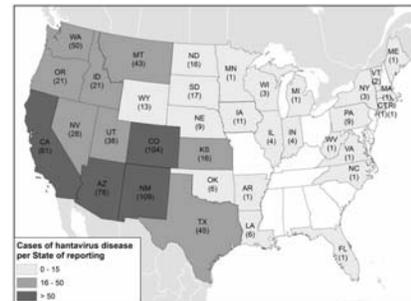


Schmaljohn and Hjelle, EID, 1997



Hantaviruses in the United States

- Sin Nombre virus (deer mouse, *Peromyscus maniculatus*)
- Black Creek Canal virus (cotton rat, *Sigmodon hispidus*)
- Bayou virus (rice rat, *Oryzomys palustris*)
- New York virus (white-footed mouse, *Peromyscus leucopus*)



Cumulative Case Count through January 2017 per State



Model Systems for the Study of Zoonotic Viruses

Model	zoonosis research
Animal model, cell (e.g., laboratory)	Limited
Animal model, rodent (natural reservoir)	limited to few species
Cell culture (E6, tum, monoc)	limited, but less questions fraction
Cell culture	available

Studying immunity to zoonotic diseases in the natural host — keeping it real

Andrew G. D. Bean¹, Michelle L. Baker¹, Cameron R. Stewart¹, Christopher Cowled¹, Celine Deffrasnes¹, Lin-Fa Wang^{2,3} and John W. Lowenthal^{1,4}

Cell

Reservoir Host Immune Responses to Emerging Zoonotic Viruses

Judith N. Mandi^{1,2}, Rafi Ahmed², Luis B. Barreiro³, Peter Daszak⁴, Jonathan H. Epstein⁴, Herbert W. Virgin⁵ and Mark B. Feinberg⁶

¹Lymphocyte Biology Section, Laboratory of Systems Biology, NIAID, National Institutes of Health, Bethesda, MD 20892, USA
²Emory Vaccine Center, Emory University School of Medicine, Atlanta, GA 30322, USA
³Saint-Justine Hospital Research Centre, Department of Pediatrics, University of Montreal, Montreal, QC H3T 1J4, Canada
⁴EcoHealth Alliance, New York, NY 10001, USA
⁵Department of Pathology & Immunology, Washington University School of Medicine, St. Louis, MO 63110, USA
⁶Merck Vaccines, Merck & Co. Inc., West Point, PA 19486, USA

Hantaviruses and Their Rodent Reservoirs

Why don't reservoir rodents have pathology when infected with their hantaviruses?

Why are they unable to clear the virus?

Sin Nombre virus and the deer mouse (*Peromyscus maniculatus*)



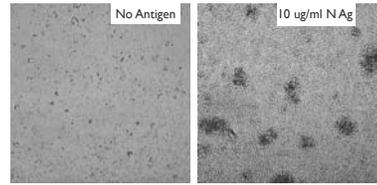
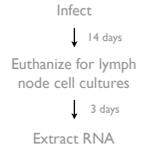
Rodent-Borne New World Hantaviruses



How do deer mouse immune responses differ to SNV and ANDV?

Experimental Design

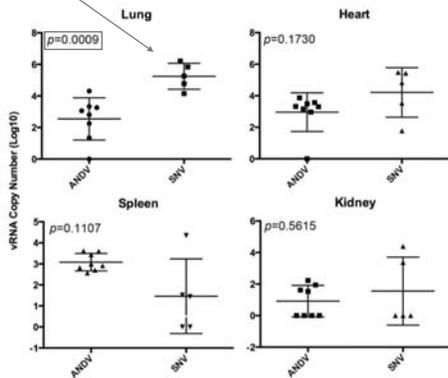
Infected deer mice with either SNV or ANDV
 Euthanize deer mice 14 days later
 Hematology (all normal, even platelets)
 Viral RNA detection
 Serology (ELISA for nucleocapsid)
 Gene expression profiling of virus-specific lymph node helper T cells (94 gene real-time PCR array)



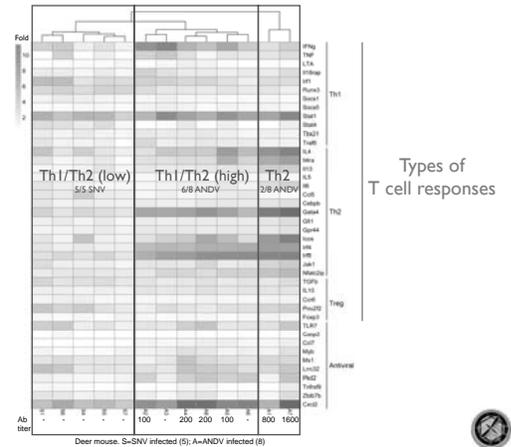
72 hour lymph node cultures from a deer mouse infected with ANDV 14 days before

Similar to levels in fatal human cases

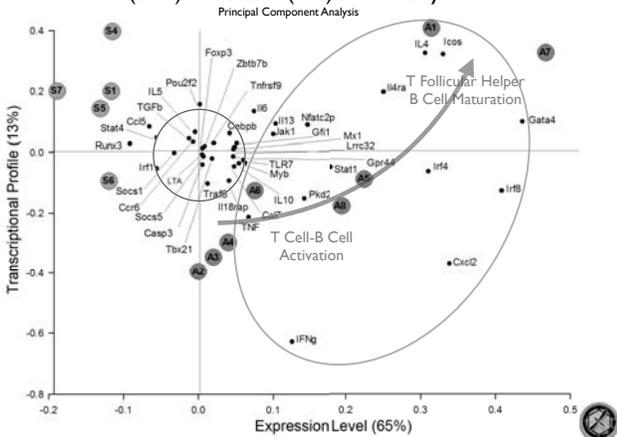
Viral Load (per mg tissue)



Gene Expression Cluster Analysis



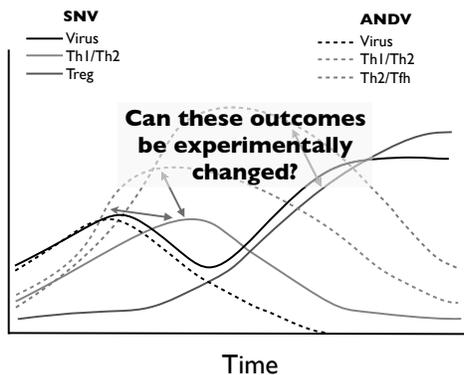
Gene Expression Profiles of Deer Mice Infected with SNV (blue) or ANDV (red) for 14 Days



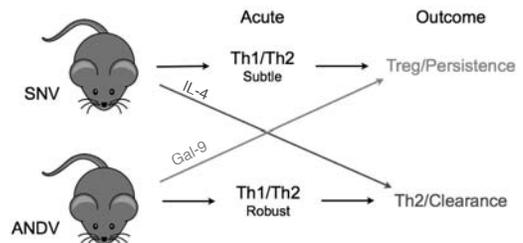
Clearance of Andes Virus Seems to be Mediated by Follicular Helper T Cells

- Follicular helper T cells are essential for robust antibody responses
 - Stimulate B cells to make antibodies
 - They facilitate class switching from IgM to IgG and IgA
 - The stimulate affinity maturation of antibodies
- The deer mouse immune response to Andes virus suggests follicular helper T cells are required for clearance of hantaviruses
 - Deer mice infected with Sin Nombre do not make a robust antibody response and remain persistently infected

An Adaptive Immune Response Model for Hantavirus Infection of Reservoirs



Altering the Outcome of Infection



Can Human Monoclonal Antibodies be used to Treat Hantavirus Disease?

- Chilean researchers examined B cells from 27 Andes virus survivors
- They identified B cells from two survivors that produced antibody that strongly neutralized Andes virus
- They genetically-engineered immortalized cells to express these antibodies in large amounts in cell culture
- Administration of these antibodies reduce mortality of hamsters infected with Andes virus to zero
- Clinical trials will begin soon
- If successful, it could be adapted to treat Sin Nombre virus infections



Garrido et al., Science Translational Medicine, 2018

