

Research on Zoonotic Disease in NL

Research Exchange Group on Human-Animal Interaction & Wellness

November 15, 2019

Dr. Hugh Whitney,
Adjunct Professor MUN





Chief Veterinary Officer for NL – 1985-2015
Adjunct Professor (MUN) – 1995 to present

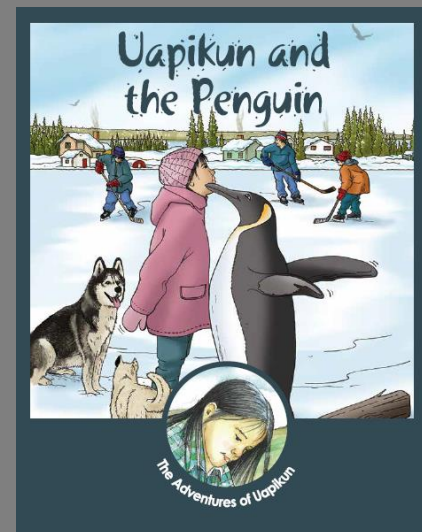
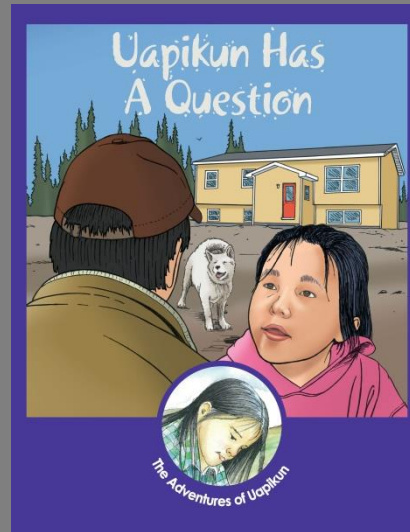
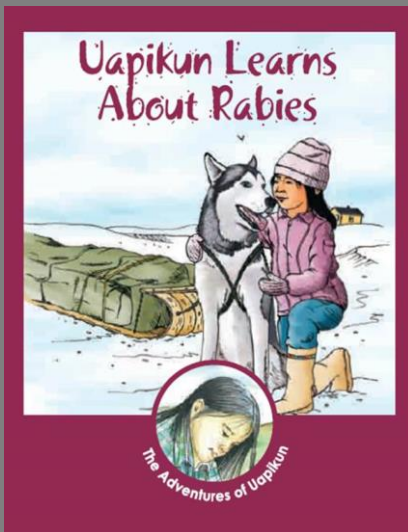
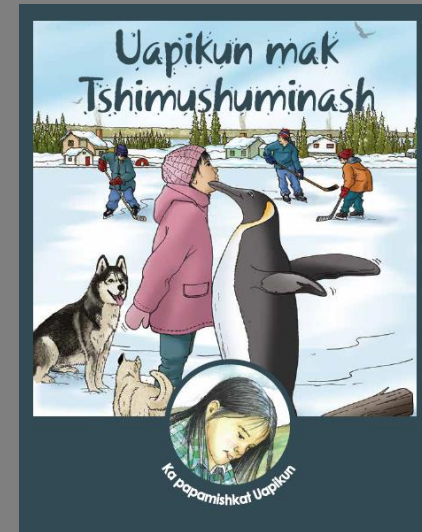
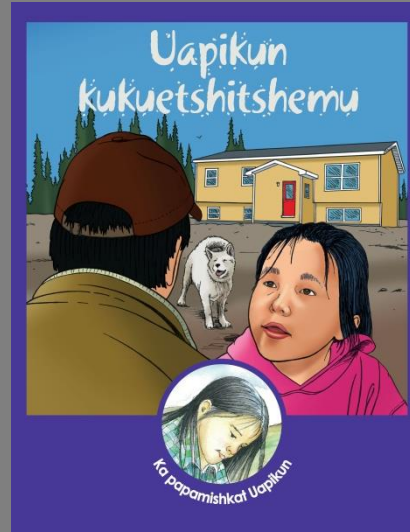
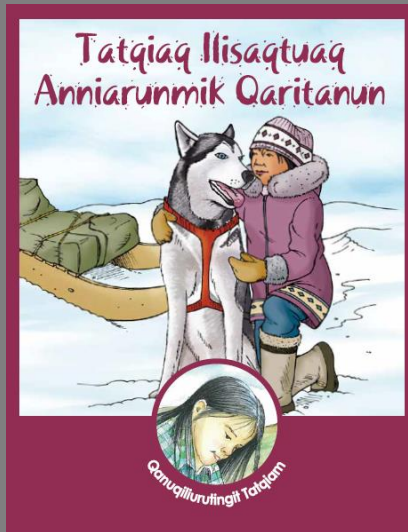
A few relevant highlights:

- Eradication of rabies from island of Newfoundland in 1988 and 2002-04
- Surveillance for West Nile virus, Lyme disease, rabies and other zoonoses
- Research into numerous animal diseases with MUN

DMV – Université de Montréal

MSc (Veterinary Microbiology) – University of Saskatchewan

MA (History) – Memorial University



Use of animals for teaching children

Uapikun Learns How To Be Safe Around Dogs

Dogs can be good friends.

Just like us, dogs don't like being:



Dogs can also be dangerous.



From the book
Uapikun Has A Question



The Importance of Zoonotic Disease Research in NL

1. Of economic interest to the commercial agricultural industry
2. Of interest to public health protection in the province
3. Of interest to wildlife managers and domestic animal owners
4. For public policy purposes
5. To encourage animal disease research that otherwise may not be done
6. To interest young people in research opportunities in this province
7. To assist in public education

Research of provincial scope versus national and international scope



Geographic isolation - If it's not done here, it may not be done at all



Dr. Andrew Lang
Microbiology, MUN



Dr. Atanu Sarkar
Medicine, MUN



Dr. Tom Chapman
Entomology, MUN



Dr. Joel Finnis
Climatology, MUN



Dr. Kapil Tahlan
Microbiology, MUN



Dr. James Valcour,
Epidemiology, MUN



Dr. Dawn Marshall
Genetics, MUN



Dr. Patrick Leighton,
UdeM



Dr. Robbin Lindsay,
PHAC



Dr. Nick Ogden,
PHAC



Dr. Susan Nadin-Davis,
CFIA

Research Collaborators



MUN



Atlantic Vet College



Environment Canada



CFIA Rabies
Centre of Expertise



U Calgary



Faculté de médecine
vétérinaire, UdeM



Ontario Vet. College



National Microbiology
Laboratory

Collaborating Institutions



Avian influenza



Rabies



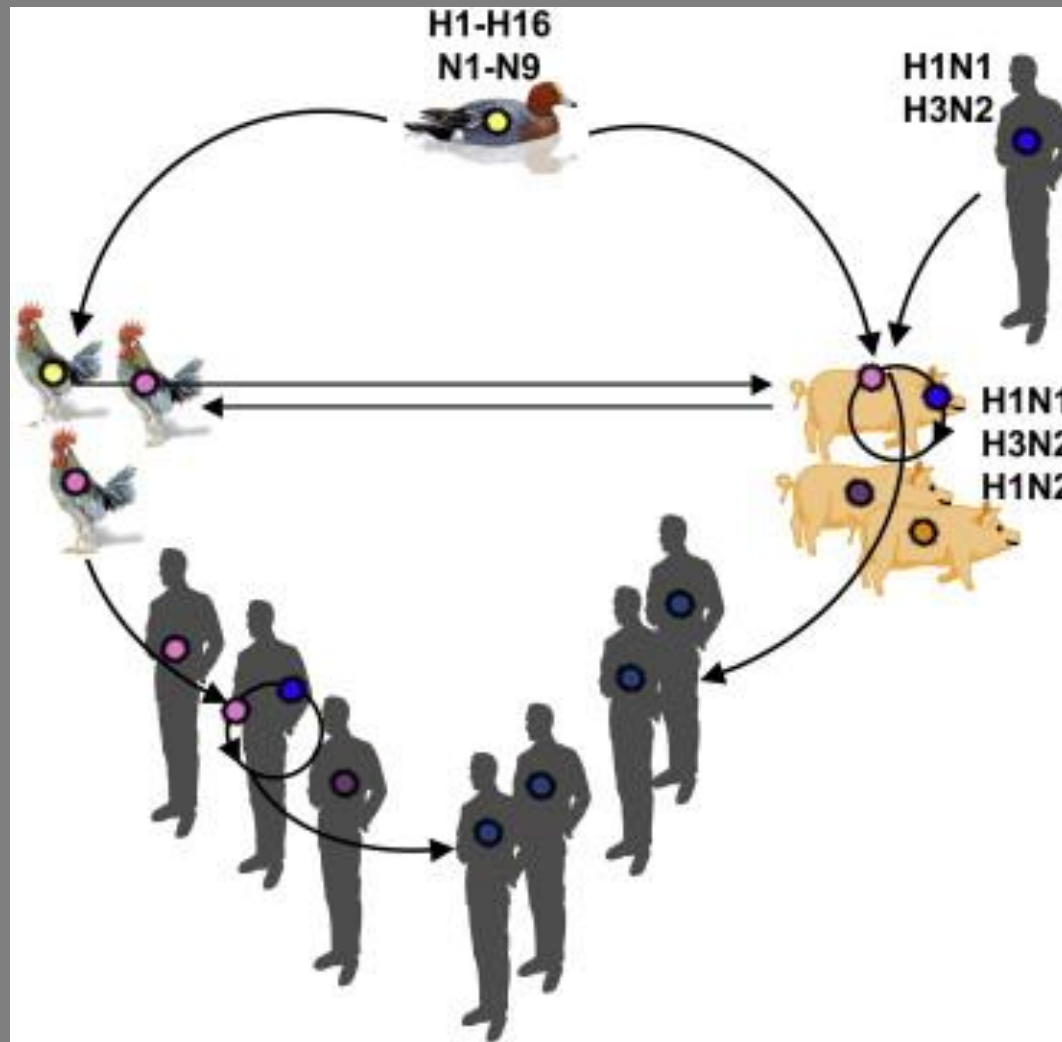
MRSA/MRSP



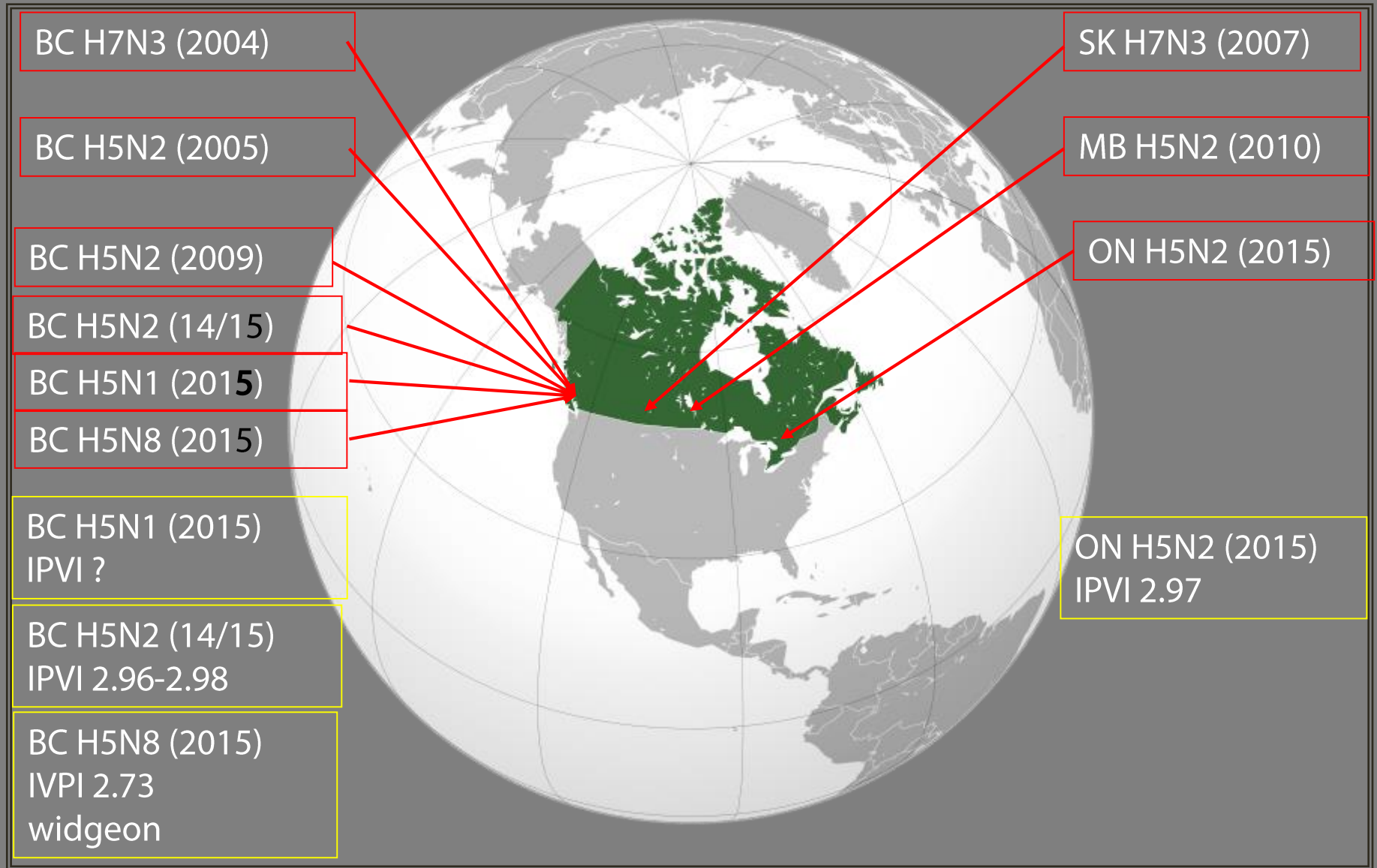
Arboviruses



Lyme disease

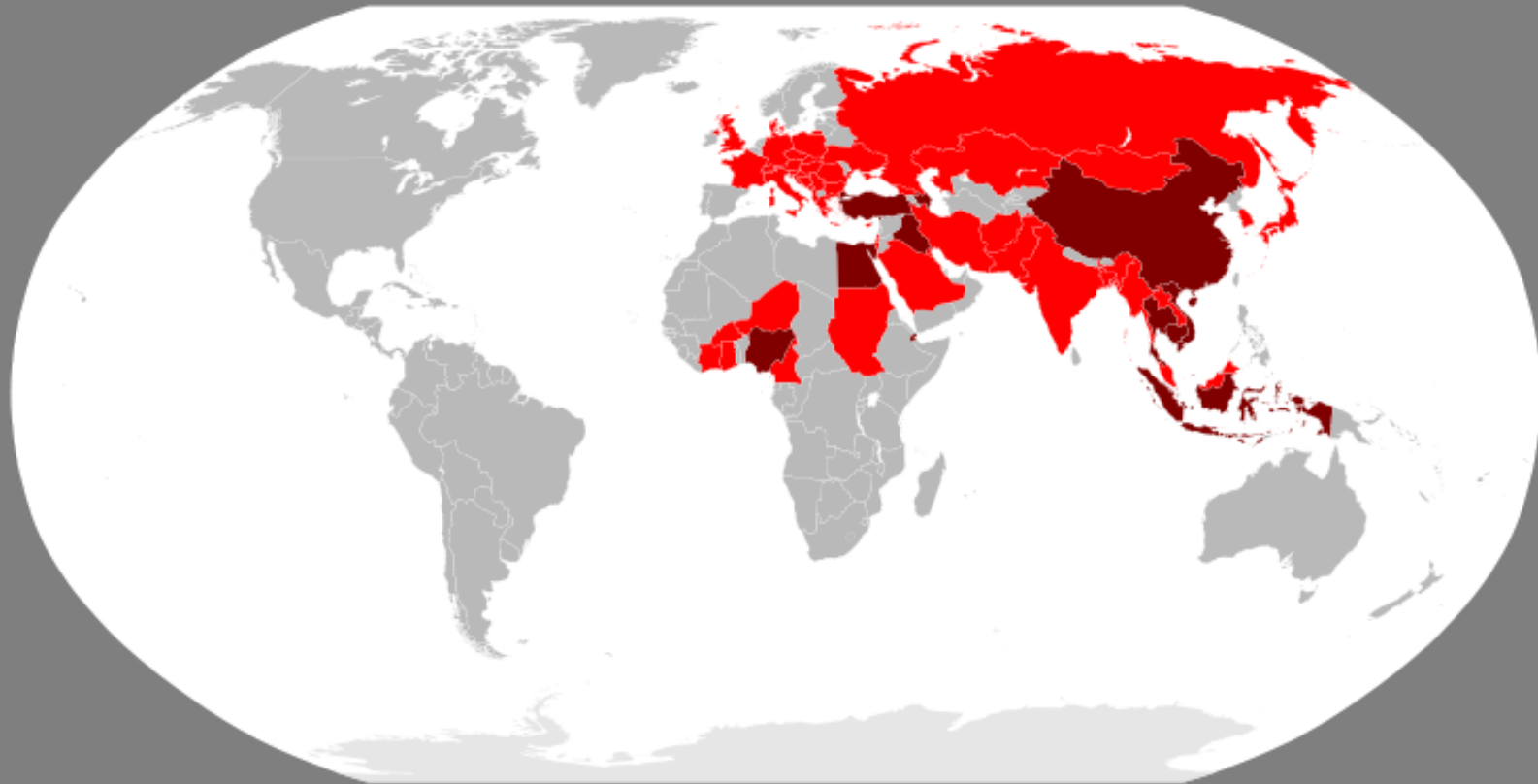


Influenza A – wild birds as reservoir



Influenza A – impact on the Canadian poultry industry

When the research started at MUN

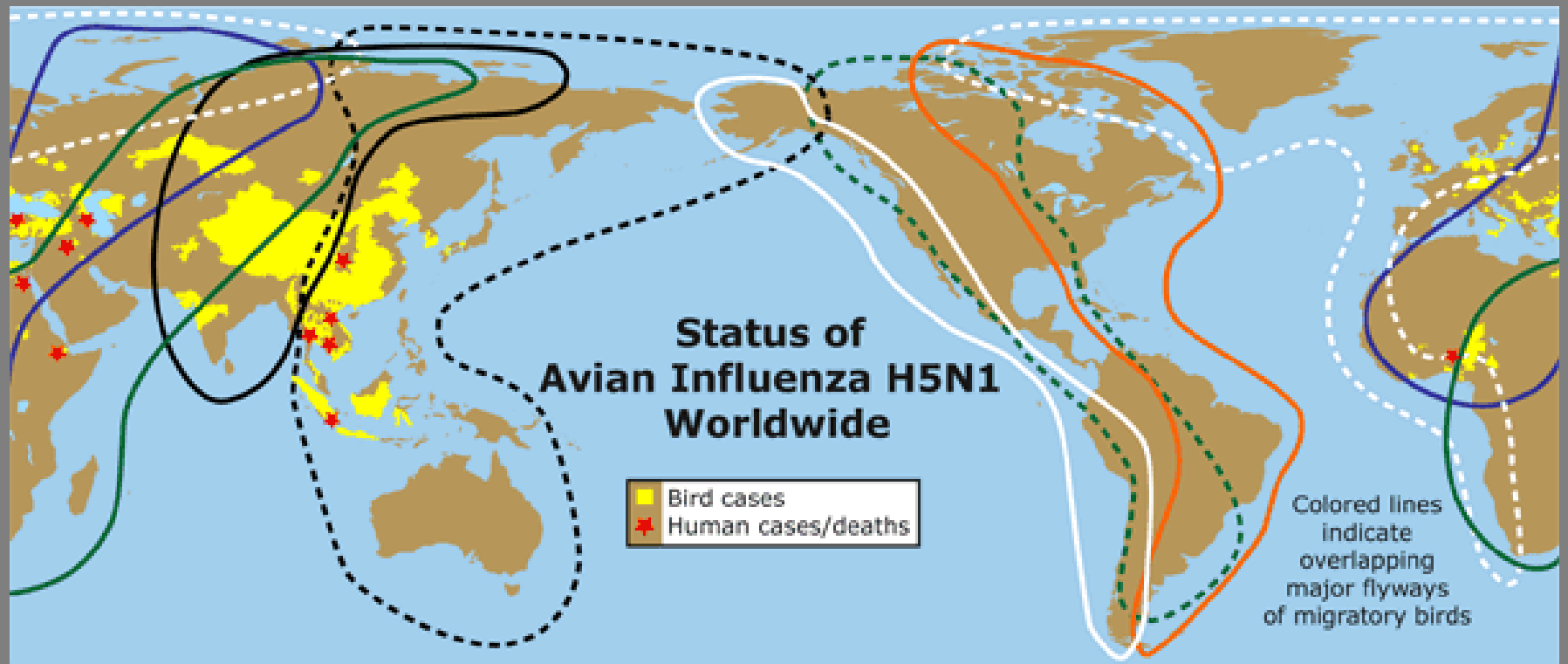


■ Countries where poultry died of H5N1

■ Countries where people died of H5N1

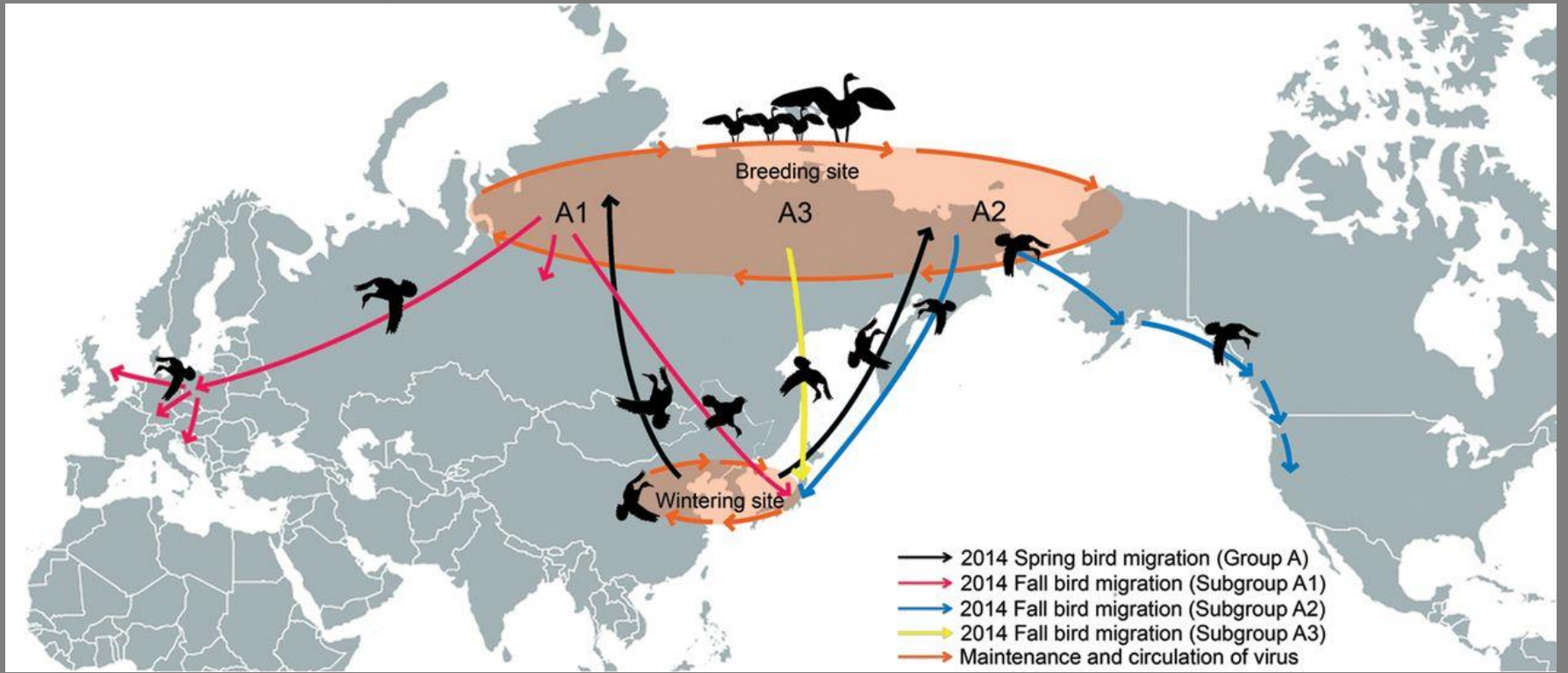
2003+

Avian Influenza – highly pathogenic H5N1 spread



North Pacific

Avian Influenza – risks to North America?

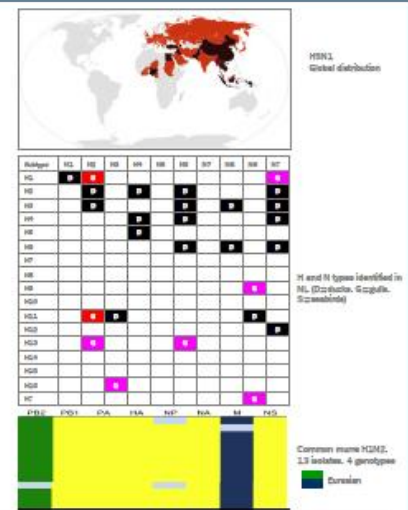


H5N8 presumably arrived in North America with migratory birds then reassorted into both H5N1 and H5N2, with H5N2 causing the most damage to the poultry industry.

Assumed movement of H5N8 to North America

[illegible]

With the discovery of the highly pathogenic H5N1 AI subtype and acceleration of its global spread in 2005, and with the increased occurrence of non-H5N1 AI outbreaks on Canadian poultry farms, research was undertaken to examine AI spread within Canada. With the confirmation of H5N1 in western Europe in 2007, research into the cross-Atlantic spread of AI viruses and their North American entry through Newfoundland and Labrador was expanded. Results indicate a very significant entry of Eurasian AIV genes into this province through a variety of wild bird species.

[illegible]

Researchers

Graduate students

Funding agencies



Avian influenza - findings to date



Analysis of influenza A viruses from gulls: an evaluation of inter-regional movements and interactions with other avian and mammalian influenza A viruses

Jessica Benkaroun, Dany Shoham, Ashley N.K. Kroyer, Hugh Whitney, and Andrew S. Lang.
Accepted for publication in Cogent Biology, 2016.

In North America, only two wholly Eurasian avian IAVs have been found to date. One was an H16N3 virus from an American herring gull (*Larus smithsonianus*) in Eastern Canada, (Huang et al., 2014).



The other was the highly pathogenic avian influenza (HPAI) H5N8 virus that recently emerged in China, spread to South Korea, Japan and Siberia, and is proposed to have been transmitted through the Beringia region by migratory waterfowl into Pacific North America (Lee et al., 2015). This H5N8 arrival ended the long speculation about whether or not virulent Asian avian strains, such as H5N1, could be transmitted into the Americas by migrating birds.



H5N8

H16N3

Demonstrated entry points for Eurasian IA virus into NA

Wild Bird Influenza Survey, Canada, 2005. E. Jane Parmley, Nathalie Bastien, Timothy F. Booth, Victoria Bowes, Peter A. Buck, Andre Breault, Dale Caswell, Pierre-Yves Daoust, J. Chris Davies, Seyyed Mehdy Elahi, Madeleine Fortin, Fred Kibenge, Robin King, Yan Li, Norman North, Davor Ojkic, John Pasick, Sydney Paul Pryor, John Robinson, Jean Rodrigue, Hugh Whitney, Patrick Zimmer, and Frederick A. Leighton. *Emerging Infectious Diseases* (2008) 14(1), 84-87

The genome sequence of an H11N2 avian influenza virus from a Thick-billed Murre (*Uria lomvia*) shows marine-specific and regional patterns of relationships to other viruses. Alissa Granter, Michelle Wille, Hugh Whitney, Gregory J. Robertson, Davor Ojkic, and Andrew S. Lang. *Virus Genes* (2010) 41:224–230.

Reassortment of American and Eurasian genes in an influenza A virus isolated from a great black-backed gull (*Larus marinus*), a species demonstrated to move between these regions. Michelle Wille, Gregory J. Robertson, Hugh Whitney, Davor Ojkic, and Andrew S. Lang. *Archives of Virology* (2011) 156:107–115

Extensive Geographic Mosaicism in Avian Influenza Viruses from Gulls in the Northern Hemisphere. Michelle Wille, Gregory J. Robertson, Hugh Whitney, Mary Anne Bishop, Jonathan A. Runstadler, Andrew S. Lang. *PLoS ONE* (2011) 6(6):1-10

Diverse Inter-continental and Host Lineage Reassortant Avian Influenza A Viruses in Pelagic Seabirds. Yanyan Huang, Gregory J. Robertson, Davor Ojkic, Hugh Whitney, Andrew S. Lang. *Infection, Genetics and Evolution* (2014) 22:103–111.

A 4-year Study of Avian Influenza Virus Prevalence and Subtype Diversity in Ducks of Newfoundland, Canada. Yanyan Huang, Michelle Wille, Ashley Dobbin, Gregory J. Robertson, Pierre Ryan, Davor Ojkic, Hugh Whitney, and Andrew S. Lang. *Canadian Journal of Microbiology* (2013) 59:701–708.

Evaluation of Seabirds in Newfoundland and Labrador, Canada, as Hosts of Influenza A Viruses. Michelle Wille, Yanyan Huang, Gregory J. Robertson, Pierre Ryan, Sabina I. Wilhelm, David Fifield, Alexander L. Bond, Alissa Granter, Hannah Munro, Rachel Buxton, Ian L. Jones, Michelle G. Fitzsimmons, Chantelle Burke, Laura McFarlane Tranquilla, Megan Rector, Linda Takahashi, Amy-Lee Kouwenberg, Anne Storey, Carolyn Walsh, April Hedd, William A. Montevecchi, Jonathan A. Runstadler, Davor Ojkic, Hugh Whitney, and Andrew S. Lang. *Journal of Wildlife Diseases* (2014) 50(1):98–103.

Genetic Structure of Avian Influenza Viruses from Ducks of the Atlantic Flyway of North America. Yanyan Huang, Michelle Wille, Ashley Dobbin, Natasha M. Walzthoni, Gregory J. Robertson, Davor Ojkic, Hugh Whitney, Andrew S. Lang. *PLoS ONE* (2014) 9(1):1-10.

Perpetuation and reassortment of gull influenza A viruses in Atlantic North America. Yanyan Huang; Michelle Wille; Jessica Benkaroun; Hannah Munro; Alexander L Bond; David Fifield; Gregory J Robertson; Davor Ojkic; Hugh Whitney; Andrew Lang. *Virology* (2014) 456-457:353-363

Assessing the role of seabirds in the ecology of influenza A viruses. 2016. Lang, A.S., C. Lebarbenchon, A.M. Ramey, G.J. Robertson, J. Waldenström, and M. Wille. *Avian Diseases* 60: 378-386.

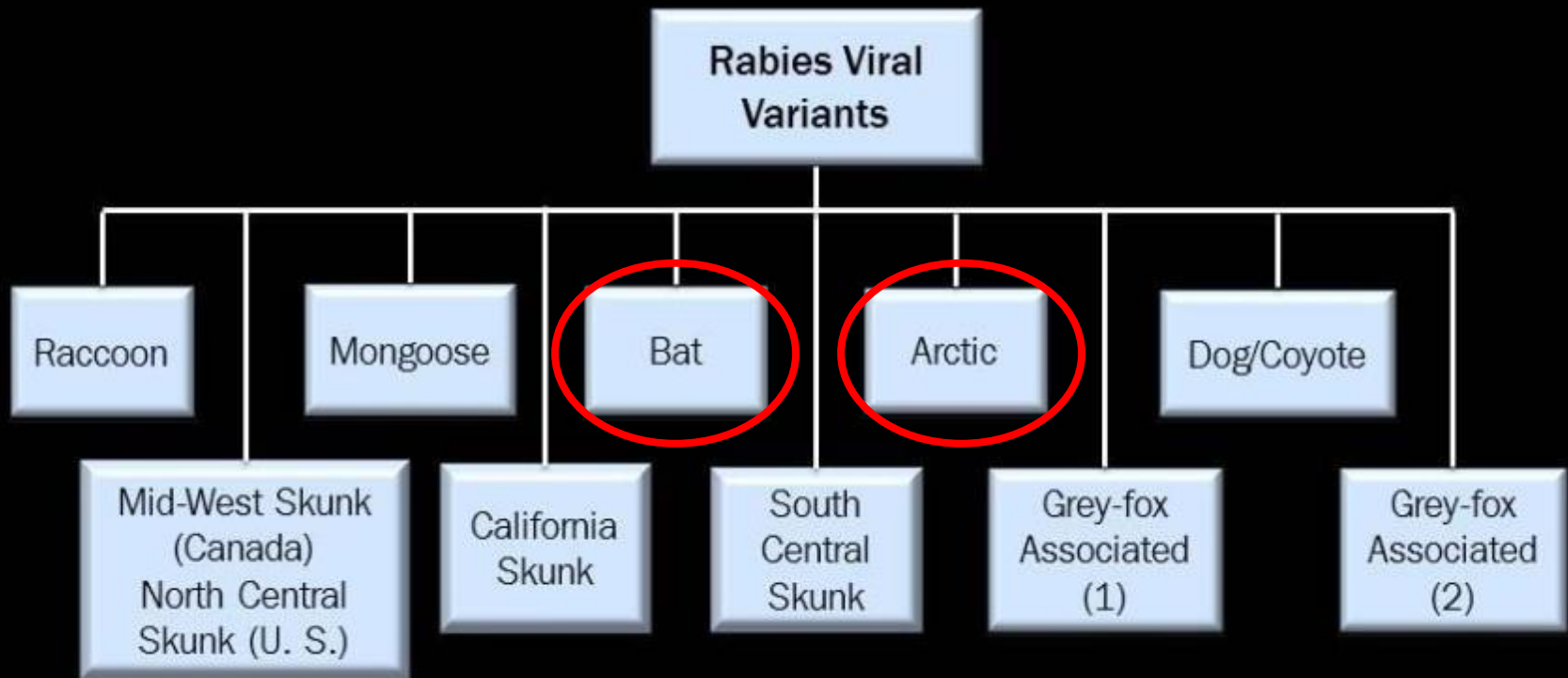
Analysis of influenza A viruses from gulls: an evaluation of inter-regional movements and interactions with other avian and mammalian influenza A viruses. Jessica Benkaroun, Dany Shoham, Ashley N.K. Kroyer, Hugh Whitney, and Andrew S. Lang. 2016 *Cogent Biology*. DOI: 10.1080/23312025.2016.1234957

Analysis of the variability in the non-coding regions of influenza A viruses. 2018. Benkaroun, J., G.J. Robertson, H. Whitney, and A.S. Lang. *Veterinary Sciences* 5: 76

RABIES



Rabies - a fatal disease spread by an infected bite



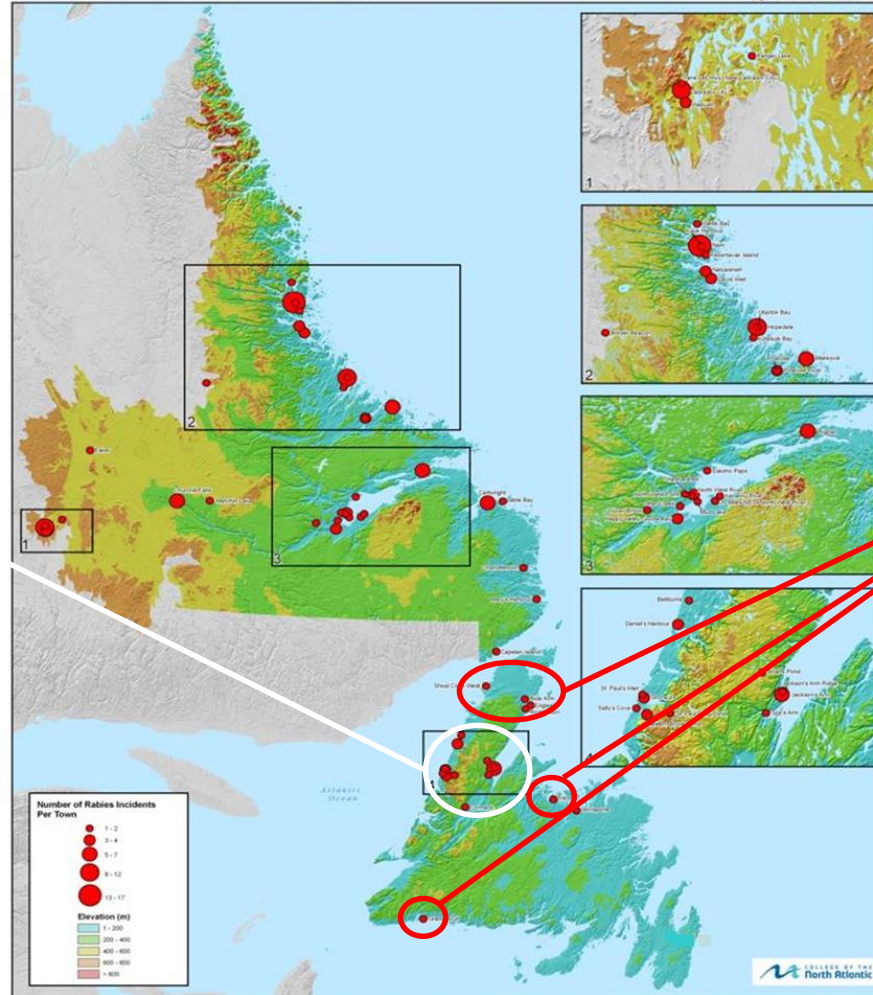
Different viral variants (lineages) are associated with different primary host species, originally determined by monoclonal antibody analysis.

Best able to persist in the primary host species.

Rabies – numerous geographic & species specific variants

Rabies Incidents Per Town in Newfoundland and Labrador 1954 - 2011*

Newfoundland
Labrador
Government of Newfoundland and Labrador
Department of Natural Resources



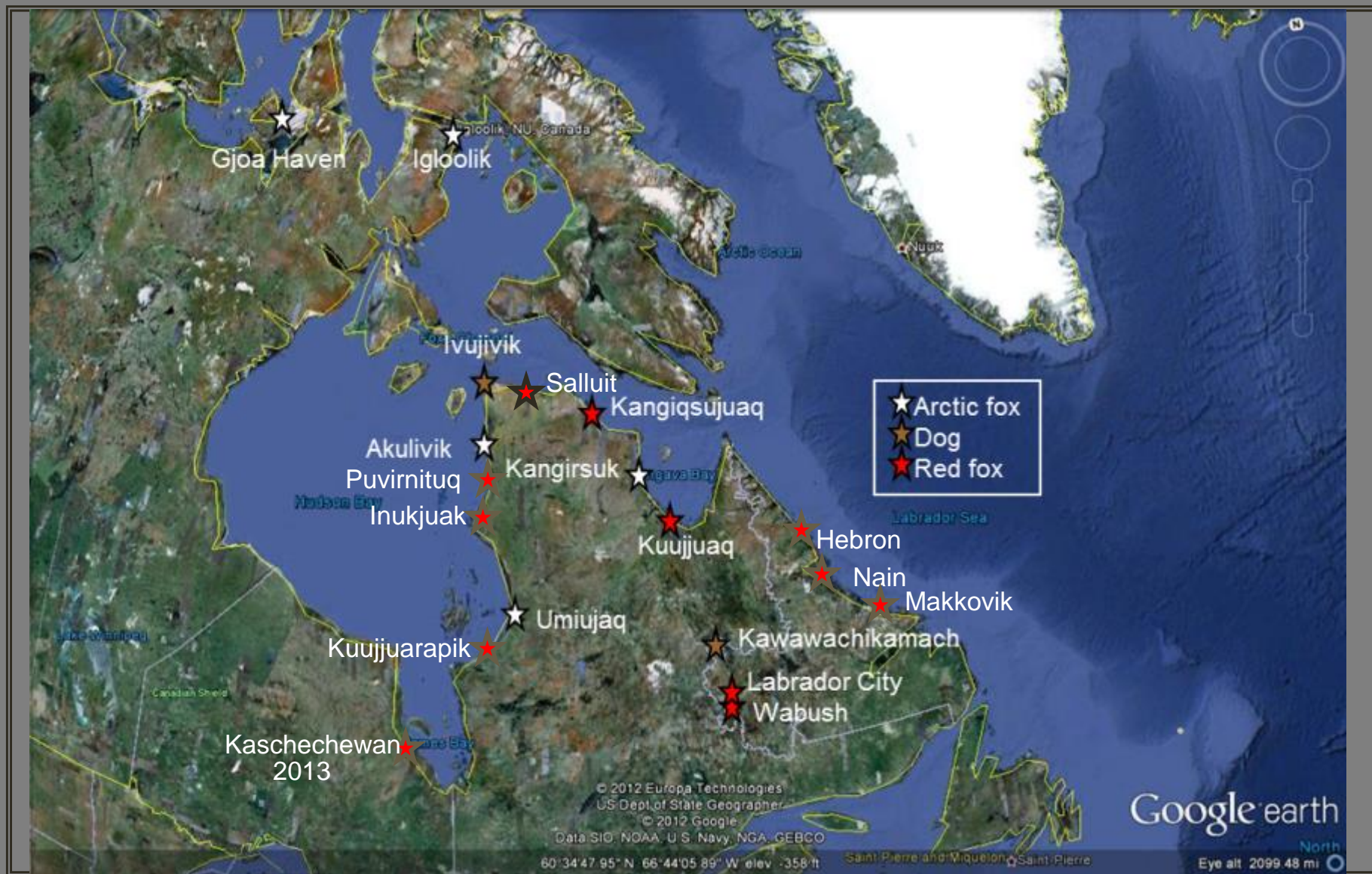
2002-03 outbreak

1988 outbreak

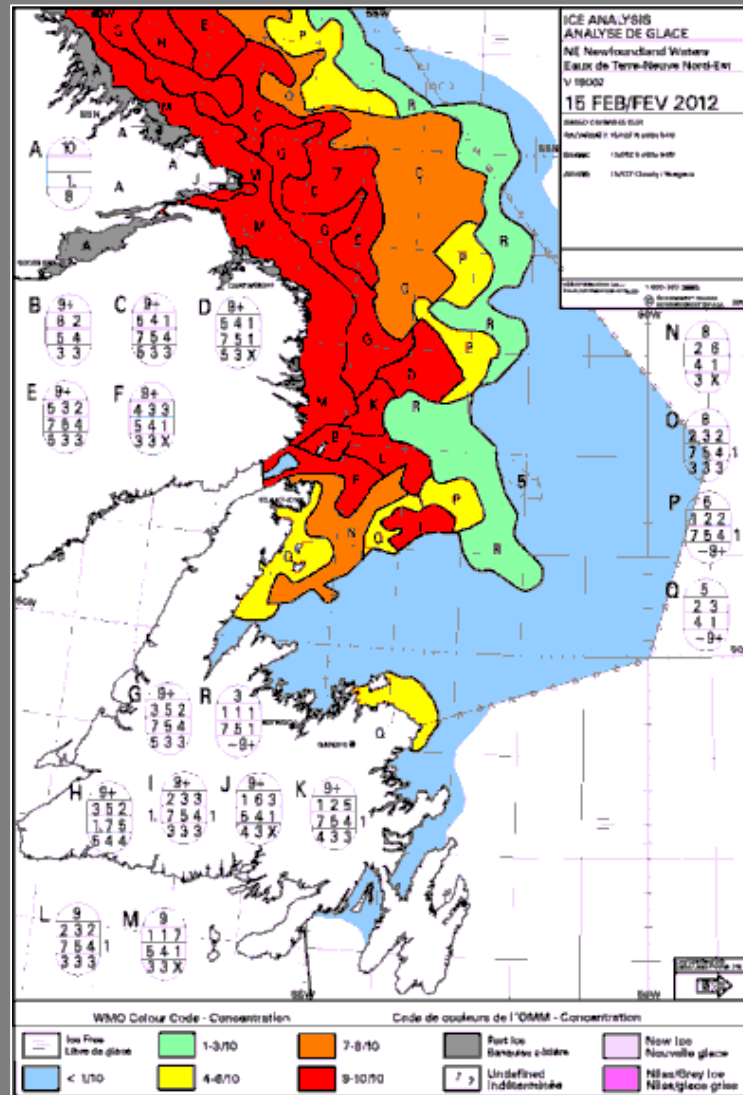
Rabies - cases in NL



Rabies – Arctic fox lineage - a pan-Arctic reality

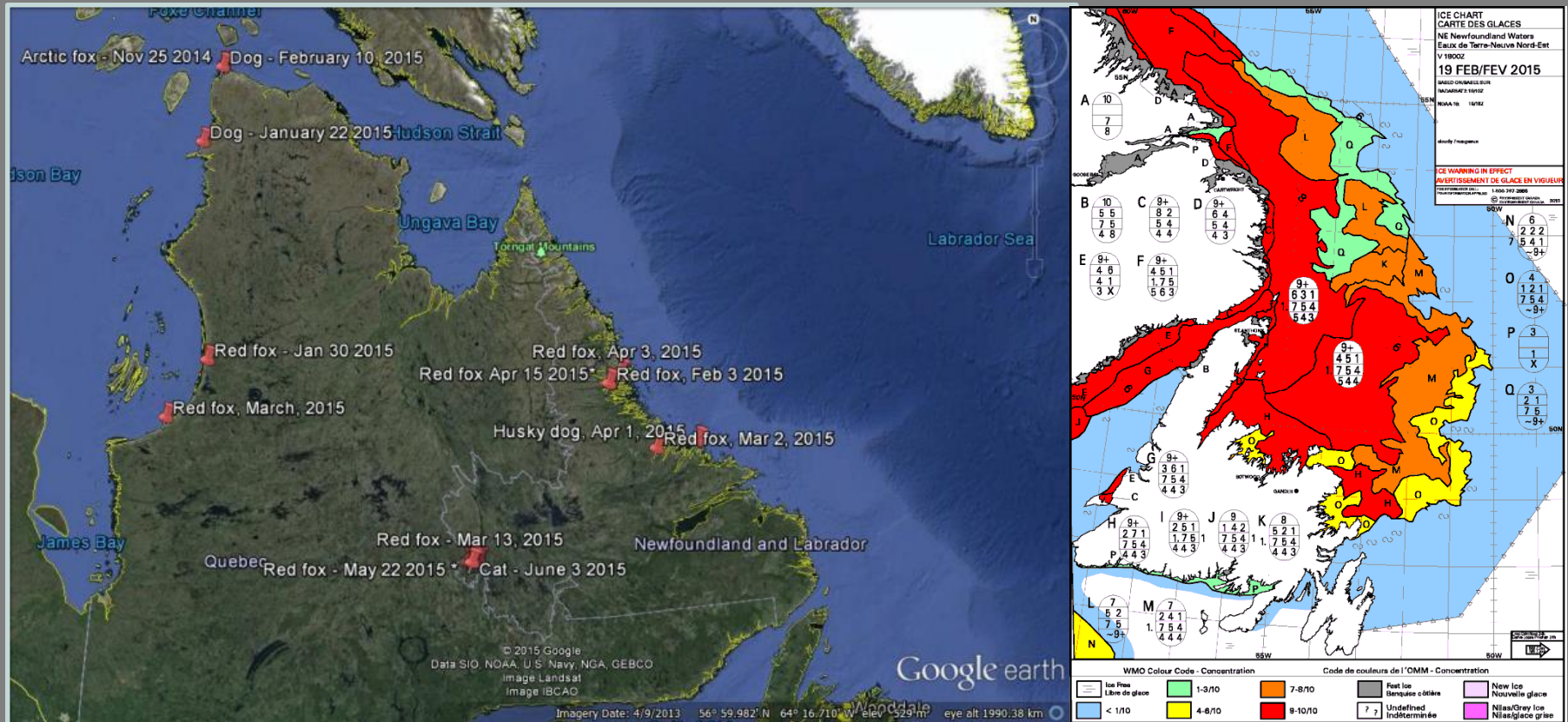


Rabies – Northeast Canada – 2011/2012

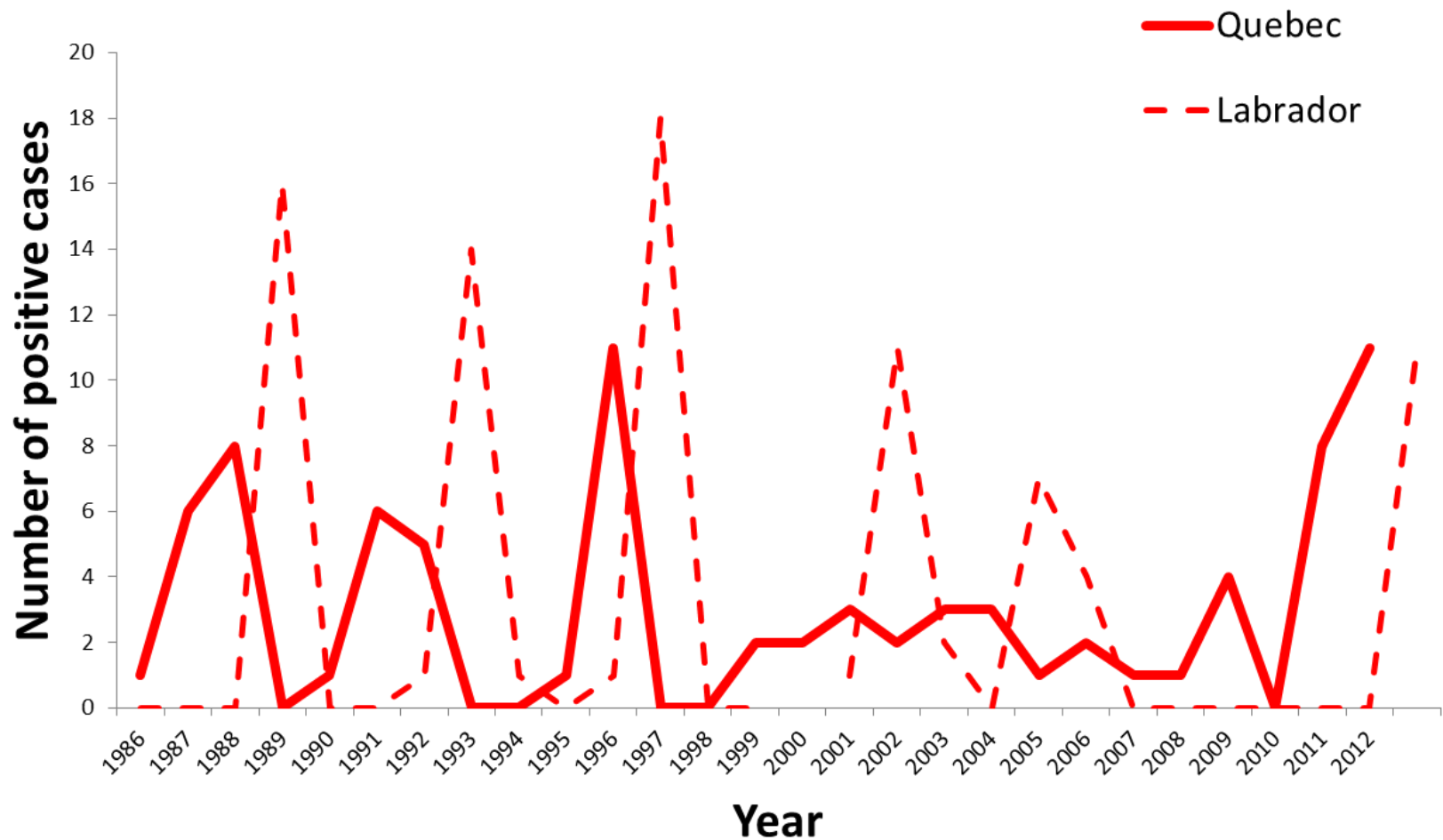


Rabies – the perpetual risk from ice bridges

2015



Rabies – the perpetual risk from ice bridges



Rabies - Synchronization of Que/Labrador outbreaks

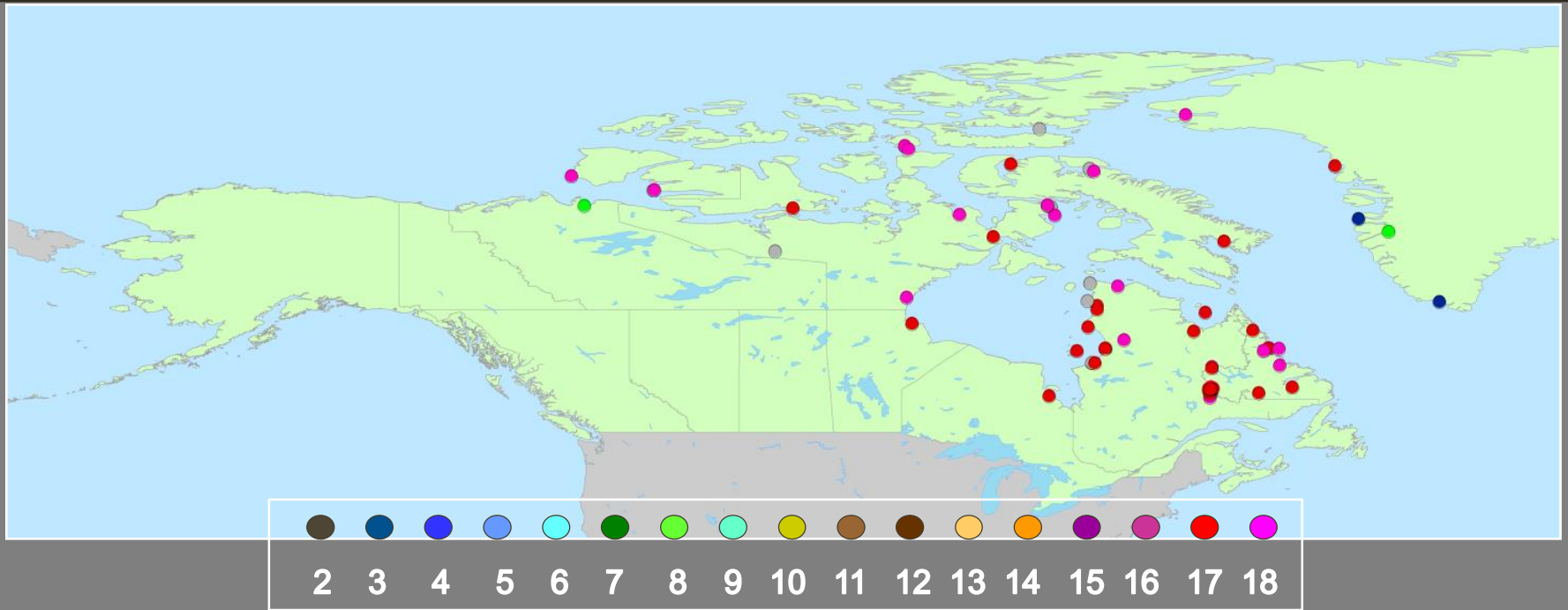


Rabies – but how does it move through the North?



Rabies - Exploring the movement of Arctic fox rabies in N Canada

Rabies - Exploring the movement of Arctic fox rabies in N Canada



The arctic fox variant has four sub-lineages across the Arctic, A1-A4, with A3 being the most common.

A-3 has been divided into 18 groups based on whole genome sequencing.

Since 2010, all cases have been from viral groups 2, 8, 17 and 18; with 17 and 18 being the most dominant.

Rabies - Exploring the movement of Arctic fox rabies in N Canada

Red foxes



Arctic foxes



Mitochondrial control region (MCR) and microsatellite (MS) analysis of red and arctic fox samples.

Arctic foxes
33 MCR subtypes

Red foxes
18 MCR subtypes



Rabies – looking for variability in fox genetics

Relationships between fox populations and rabies virus spread in northern Canada

S.Nadin-Davis, E.Fallardeau, A.Flynn & H.Marshall, manuscript, 2019



Dr. Kapil Tahlan



Dr. James Valcour



MRSA/MRSP in NL & relationship with pet ownership



Methicillin resistant *Staphylococcus aureus* – human infection



Methicillin resistant *Staphylococcus pseudintermedius* – dog infection

MRSA/MRSP in NL & relationship with pet ownership

Epidemiology

- prevalence in humans and dogs
- human risk associated with employment, source of dog, behaviour
- dog risk associated with medical history, source, age, activities



Dr. James Valcour

Microbiology

- genotypes isolated
- geographic distribution of genotypes
- possibility of resistant gene transfer between MRSA & MRSP



Dr. Kapil Tahlan



West Nile virus
Snowshoe hare virus
Jamestown Canyon virus

Roles of Host Species, Geographic Separation, and Isolation in the Seroprevalence of Jamestown Canyon and Snowshoe Hare Viruses in Newfoundland

Gregory Goff,^a Hugh Whitney,^b and Michael A. Drebot^c

Environmental Sciences Department, Grenfell Campus of Memorial University, Corner Brook, Newfoundland, Canada^a; Animal Health Division, Newfoundland and Labrador Department of Natural Resources, St. John's, Newfoundland, Canada^b; and Viral Zoonoses, National Microbiological Laboratory, Public Health Agency of Canada, Winnipeg, Manitoba, Canada^c

Significant pathogens do exist in NL

And the list of important vector species gets longer

THE
CANADIAN
ENTOMOLOGIST



1

Aedes japonicus japonicus (Diptera: Culicidae) arrives at the most easterly point in North America

Miles A. Fielden, Andrew C. Chaulk,¹ Kate Bassett, Yolanda F. Wiersma, Mardon Erbland, Hugh Whitney, Thomas W. Chapman

Journal of Medical Entomology, 2016, 1–6
doi: 10.1093/jme/tjw105
Research article

OXFORD

Sampling, Distribution, Dispersal

The Arrival of the Northern House Mosquito *Culex pipiens* (Diptera: Culicidae) on Newfoundland's Avalon Peninsula

Andrew C. Chaulk,^{1,2} Kate P. Carson,¹ Hugh G. Whitney,³ Dina M. Fonseca,⁴ and Thomas W. Chapman¹

and more broadly distributed

Arboviruses – the need for sustained research



Dr. Atanu Sarkar



Dr. Andrew Lang



Dr. Tom Chapman



Dr. Joel Finnis



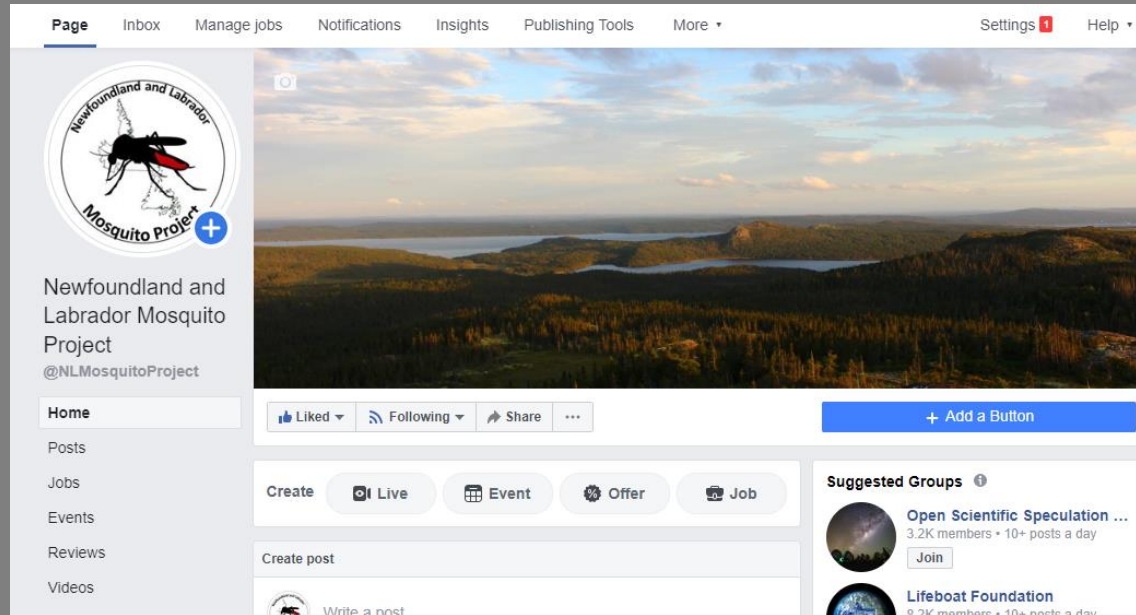
Dr. Hugh Whitney



Tegan Padgett



Dr. Marta Canuti



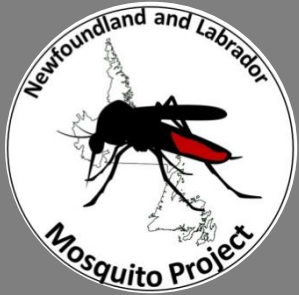
Facebook Page: @NLMosquitoProject



Kate Carson



Courtney White

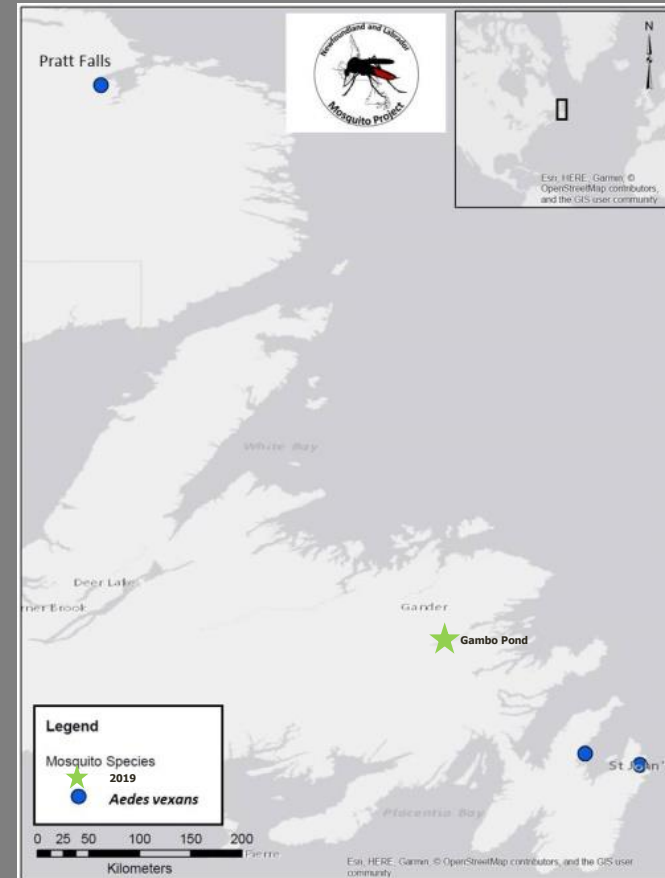


Aedes vexans inland floodwater mosquito



image credit: Sean McCann via University of Wisconsin

Its name means unpleasant, or troublesome (*Aedes*), and annoying (*vexans*). Voted as the greatest mosquito pest by the American Mosquito Control Association.





The most important vector of West Nile virus in eastern Canada. NL is the only province never to have recorded a single isolate in wild or domestic animals or humans. In the summer of 2018, all three Maritime provinces reported cases in birds.

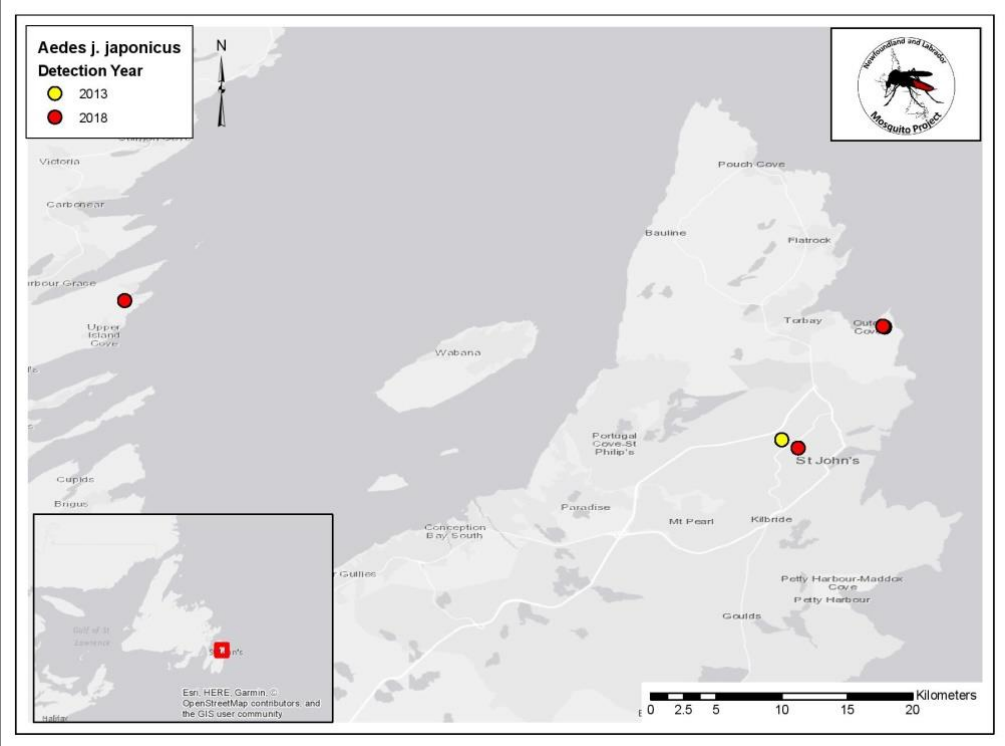




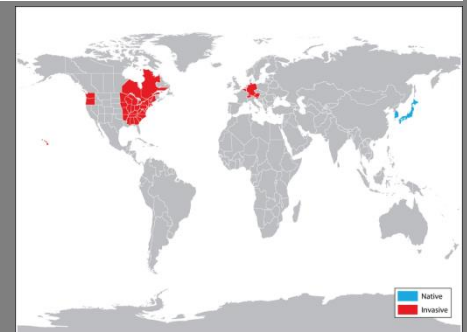
Aedes japonicus japonicus Asian bush mosquito



image source: US CDC

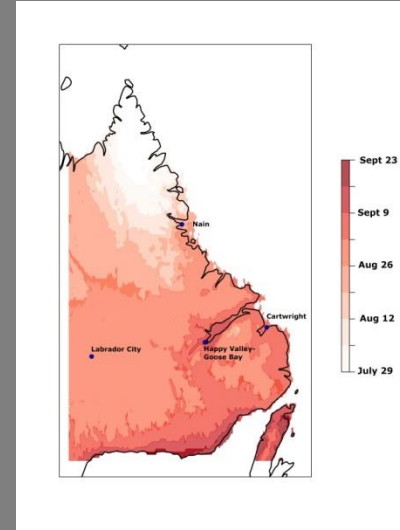
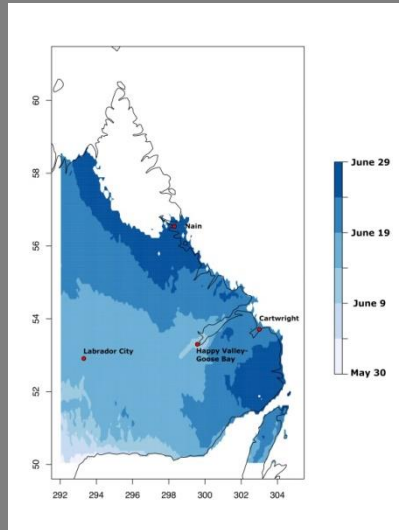


Aggressive biter, very important disease vector and globally invasive species.

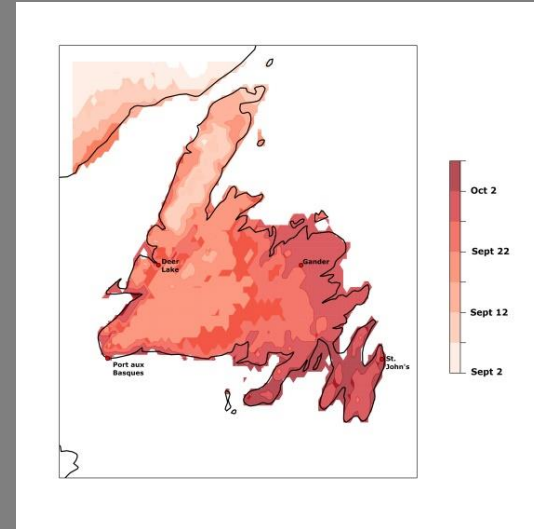
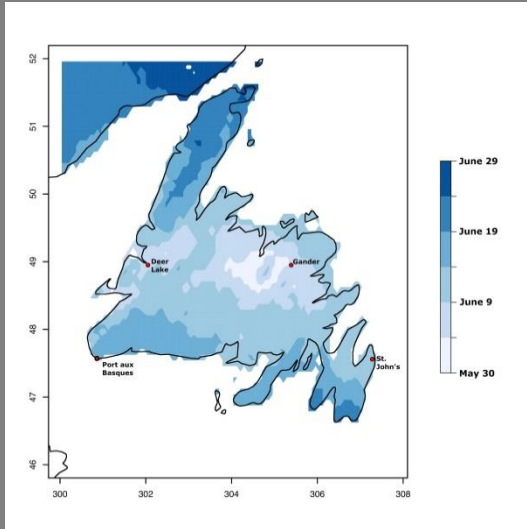


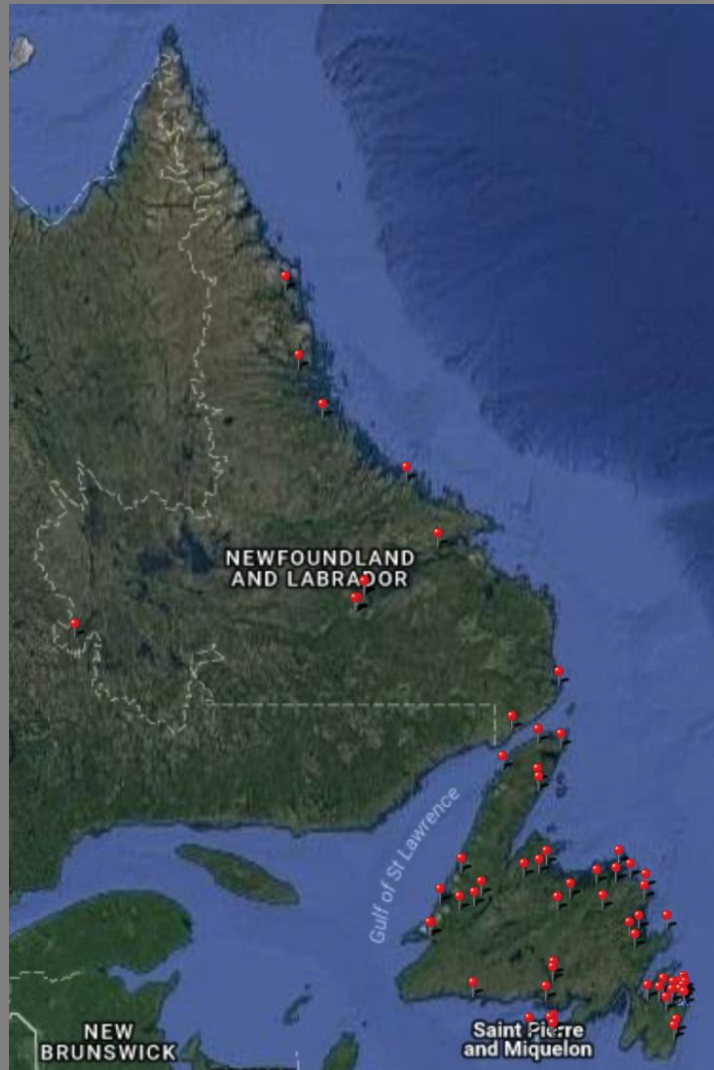
Kaufman MG, Fonseca DM. 2014.
Annu. Rev. Entomol. 59:31-49

2014



The **beginning** and **end** of mosquito season





NL Mosquito Project – participants in the project



Ticks and Lyme disease



Haemaphysalis leporis-palustris
The rabbit tick



Ixodes muris
The mouse tick



Ixodes uriae
The seabird tick



Ixodes angustus
The vole tick

Ticks – permanent populations



Ixodes marxi
The squirrel tick

Ticks – permanent populations?



Ixodes scapularis
The black-legged tick



Rhipicephalus sanguineus
The brown dog or kennel tick



Ixodes ricinus
The sheep tick



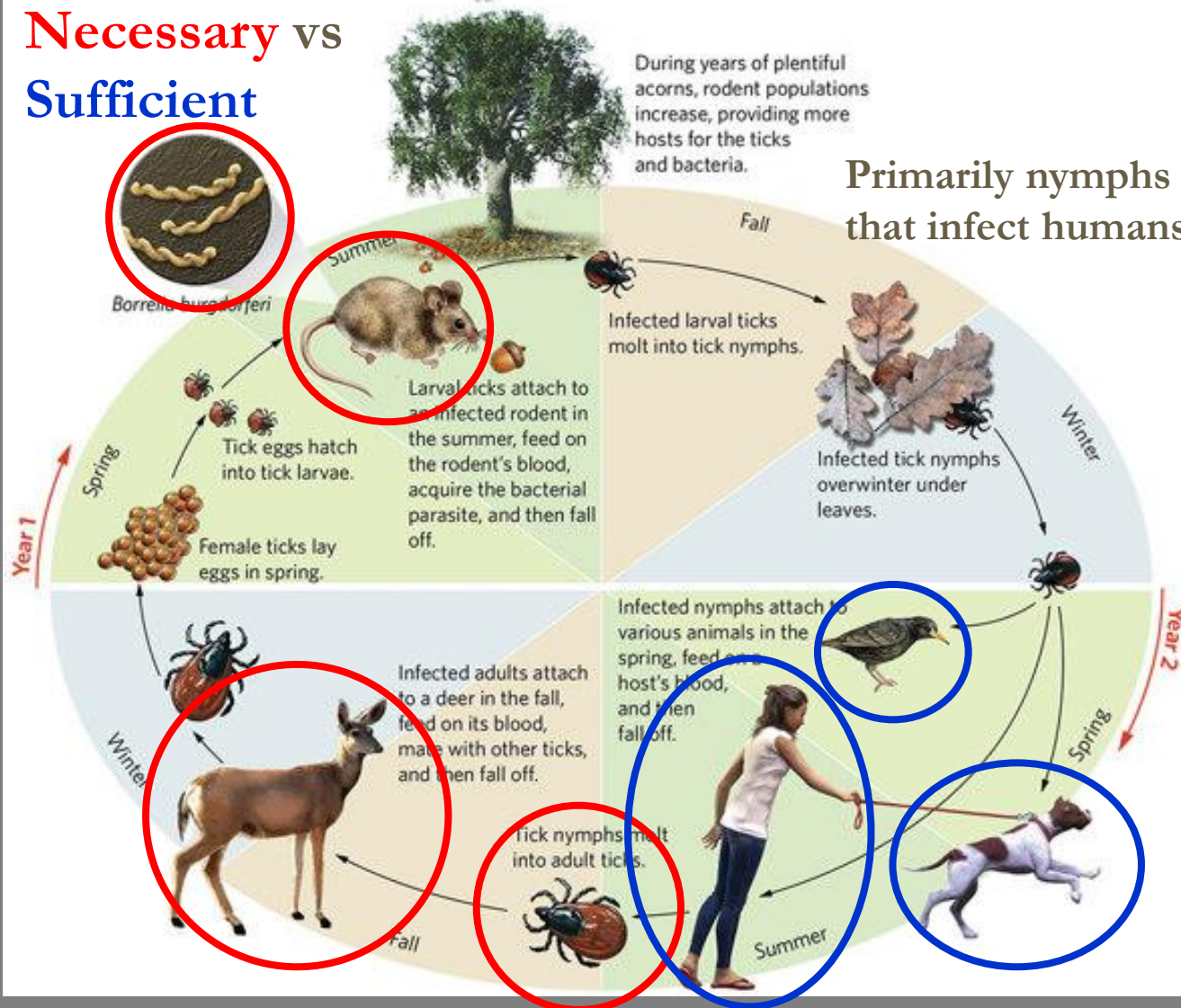
Dermacentor variabilis
American dog tick



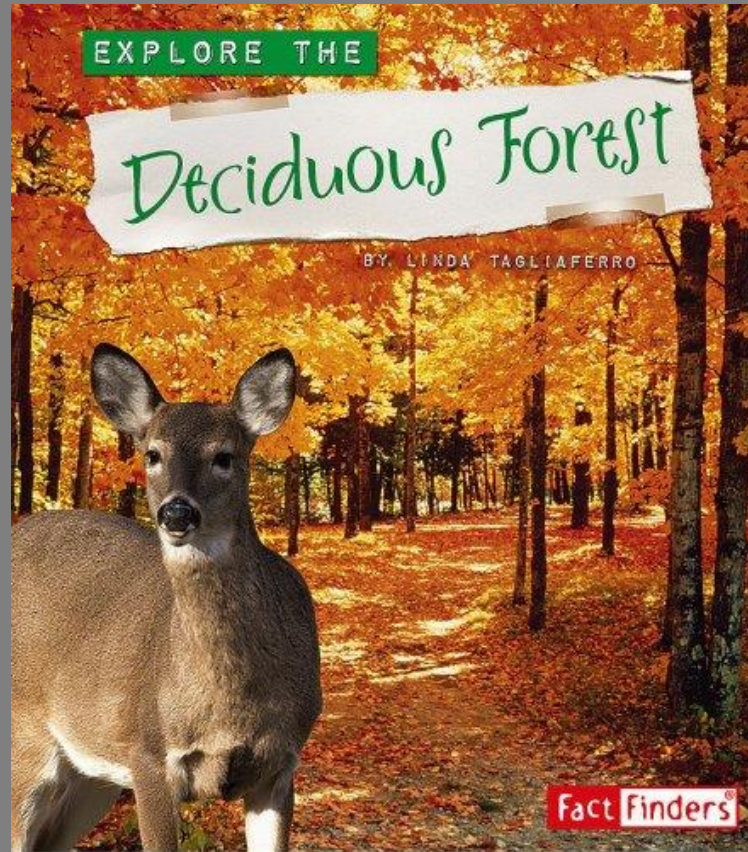
Ticks – transient populations

Necessary vs Sufficient

Primarily nymphs that infect humans



Life cycle of *Ixodes scapularis*



Deciduous forest (coniferous in Maritimes)

- Refuge from weather extremes
- Suitable hosts
- Suitable micro-climate for survival and host-seeking

Habitat of *Ixodes scapularis*



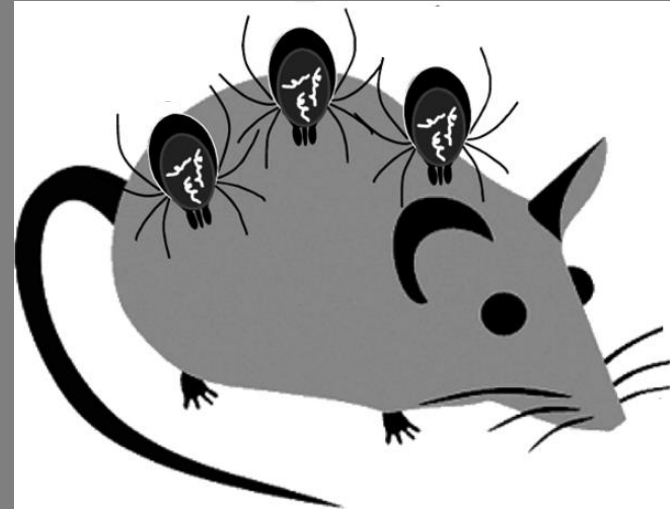
White-footed mouse



Mouse tick



Mouse tick



Maintenance of *B. burgdorferi* in nature

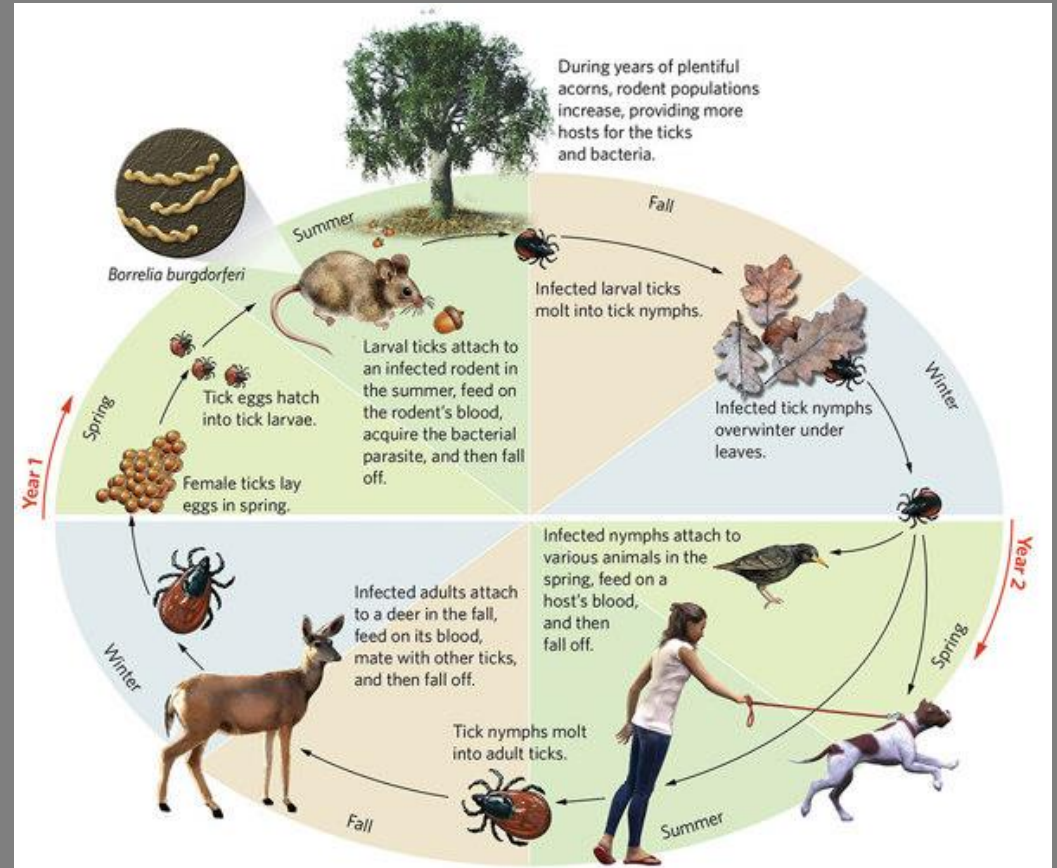
Rodent reservoirs for *Borrelia burgdorferi*



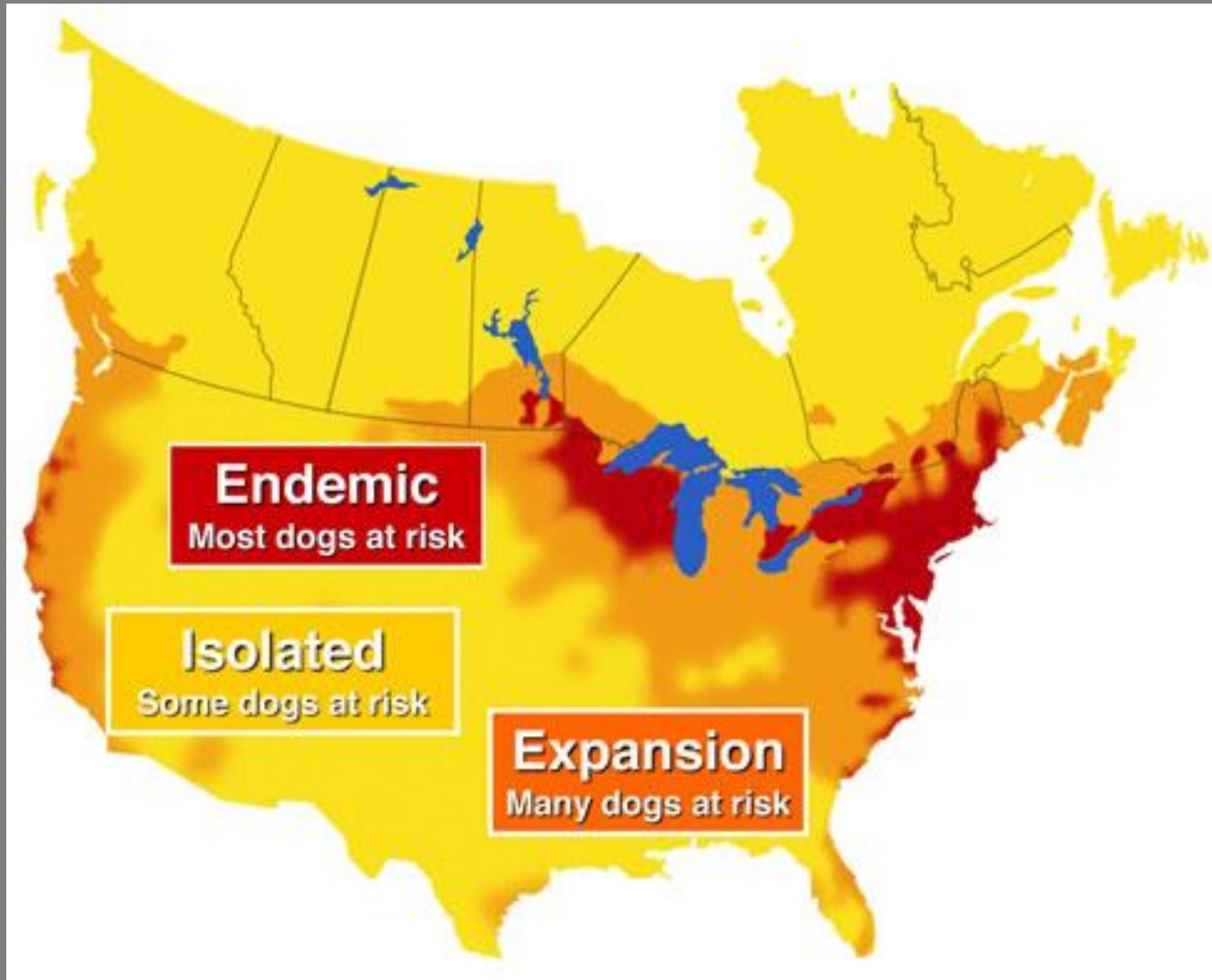
Ixodes scapularis
Borrelia burgdorferi



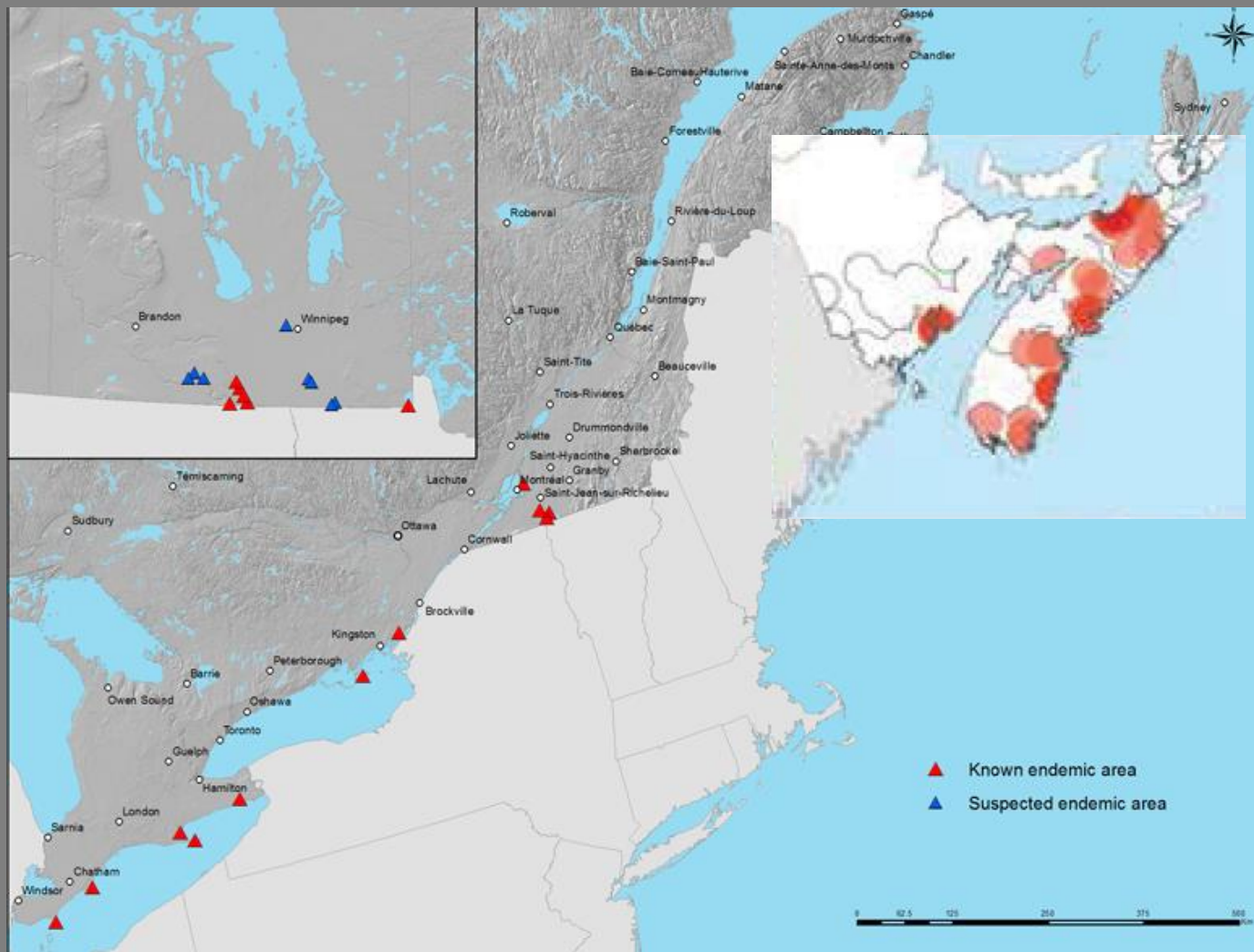
Mouse tick



Role of *Ixodes scapularis*



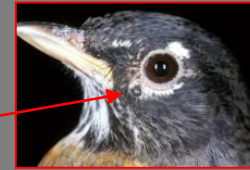
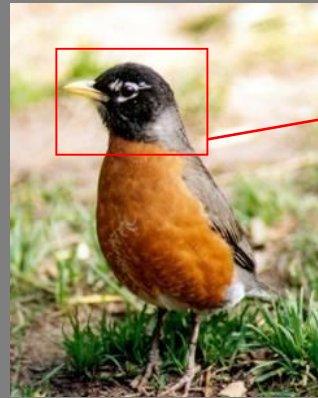
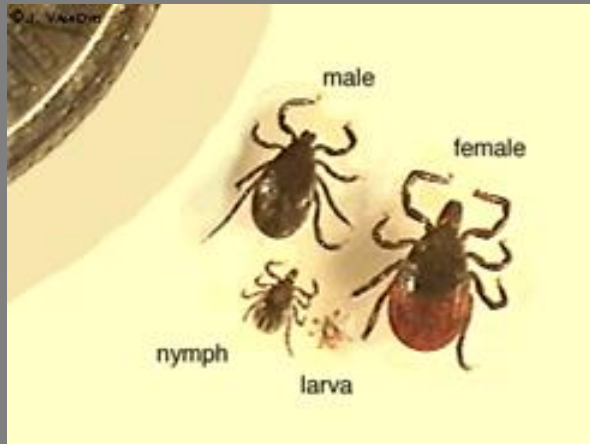
Range for *Ixodes scapularis*



Range expansion for *Ixodes scapularis*



Effect of climate change on range expansion



Ixodes scapularis

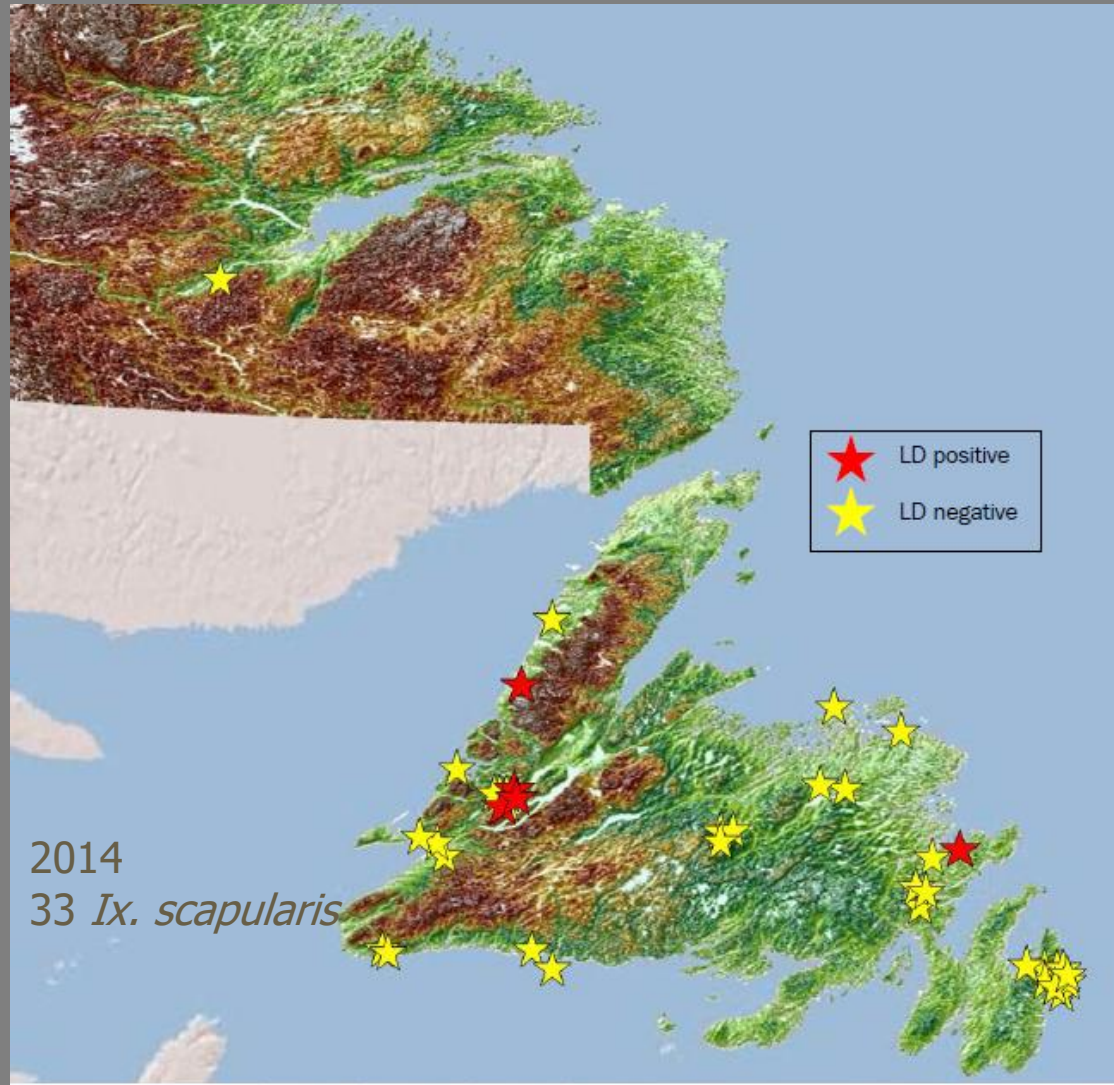


Adventitious ticks



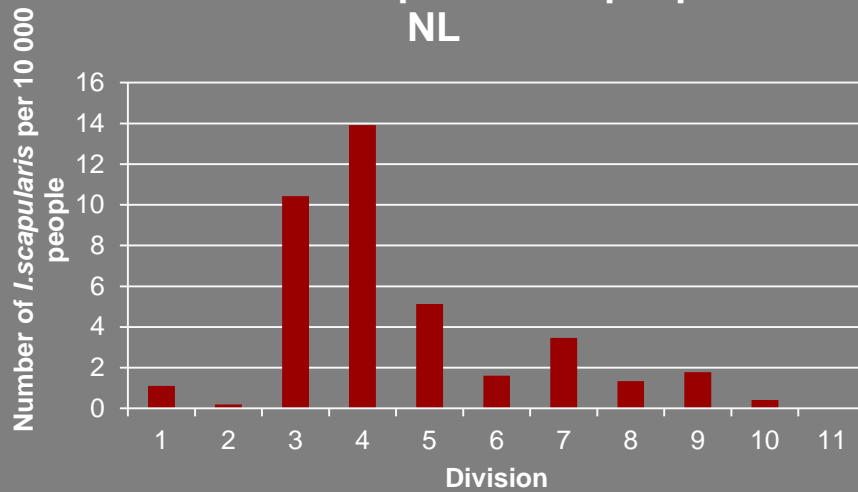
Adventitious ticks

"These ticks pose a low-level but geographically widespread potential risk of exposure to LD-infected ticks."

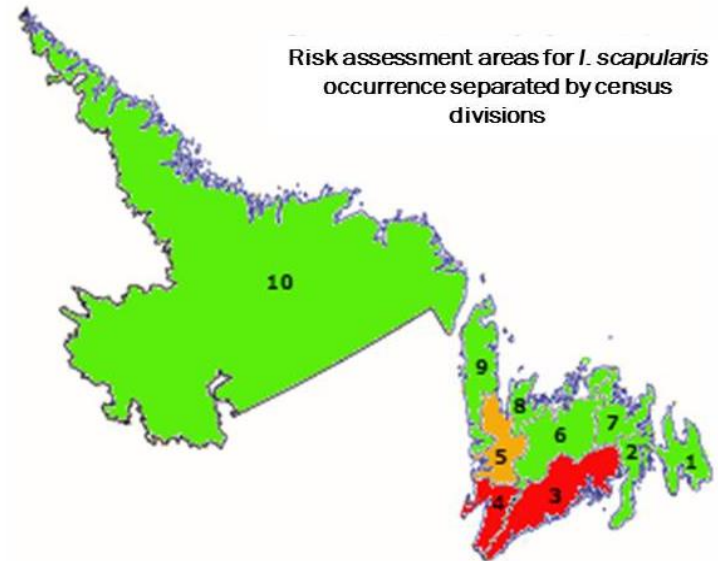


Adventitious ticks

Number of *I. scapularis* in each census division per 10 000 people in NL



Risk assessment areas for *I. scapularis* occurrence separated by census divisions







Hundreds of caribou grazing on N.L. farm
- GFW, July 2012

Caribou as an alternative host?



Dr. Hannah Munro



FIG. 1. Locations of seabird colonies where *I. uriae* ticks were collected. 1, Campbell Island, New Zealand; 2, Crozet Islands; 3, West Point, Falkland Islands; 4, Egg and St. Lazaria Islands, Alaska; 5, Gannet Island, Canada; 6, Flatey Island, Iceland; 7, Nólsoy, Faeroe Islands; 8, Cape Sizun, France; 9, Bonden Island, Sweden.



Tick infested
Atlantic puffin

Ixodes uriae



Borrelia garinii, an emerging health threat in Europe

Munro, H.J., N.H. Ogden, L.R. Lindsay, G.J. Robertson, H. Whitney, and A.S. Lang. 2017. Evidence for *Borrelia bavariensis* infections of *Ixodes uriae* within seabird colonies of the North Atlantic Ocean. Applied and Environmental Microbiology 83: e01087-17 **(Featured in Issue Spotlight)**

Hannah J.Munro, Nicholas H.Ogden, SamirMechai, L. RobbinLindsay, Gregory J.Robertson, HughWhitneyAndrew S.Lang. Genetic diversity of *Borrelia garinii* from *Ixodes uriae* collected in seabird colonies of the northwestern Atlantic Ocean. Ticks and Tick-borne Diseases, 10:6, 2019, 101255



Climate change

North Atlantic
connections

Emerging
zoonotic
diseases

Need to have
strong
multidisciplinary
research focus
at MUN



Thank you