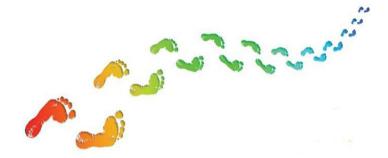






Lethbridge

From Virus Hunting To Vaccine Design; Steps To Prepare For Future Epidemics Or Pandemics



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University of











- BTV
- EHDV
- SBV
- SADS-CoV



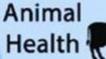




Sustainability of

wildlife biodiversity

Reduced wildlife-livestock contact





Improved livestock health











Preservation of wildlife, reduced anthropogenic effects

Disease surveillance and control, food security

- WNV
- DFV
- YFW
- ZIKV
- Nipah virus
- Hanta virus
- Ebola virus
- Marburg virus
- Lassa virus
- SARS-CoV-2



Health



Improved human health, global security

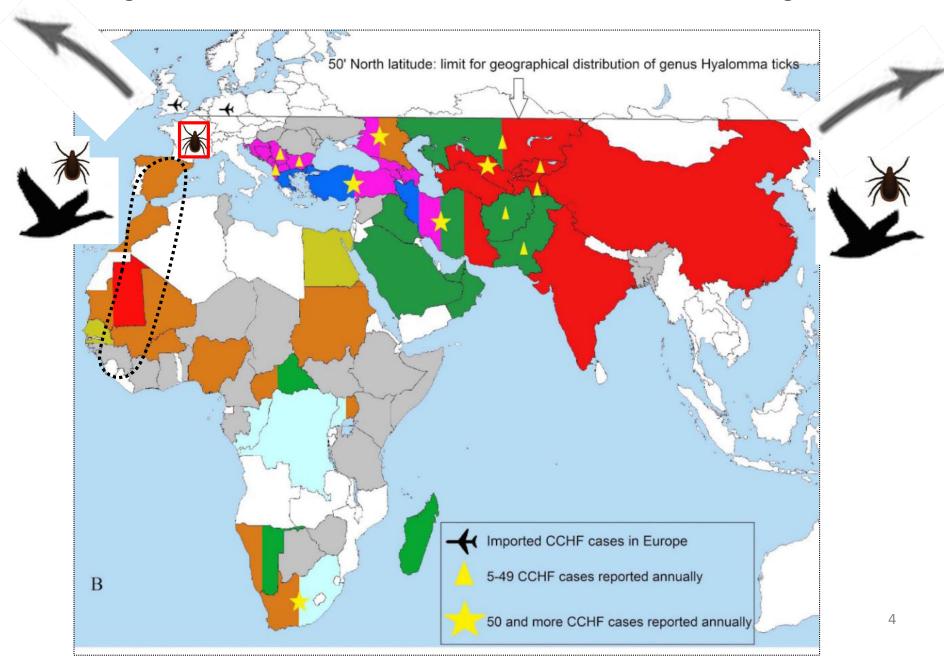


- RVFV
- JEV

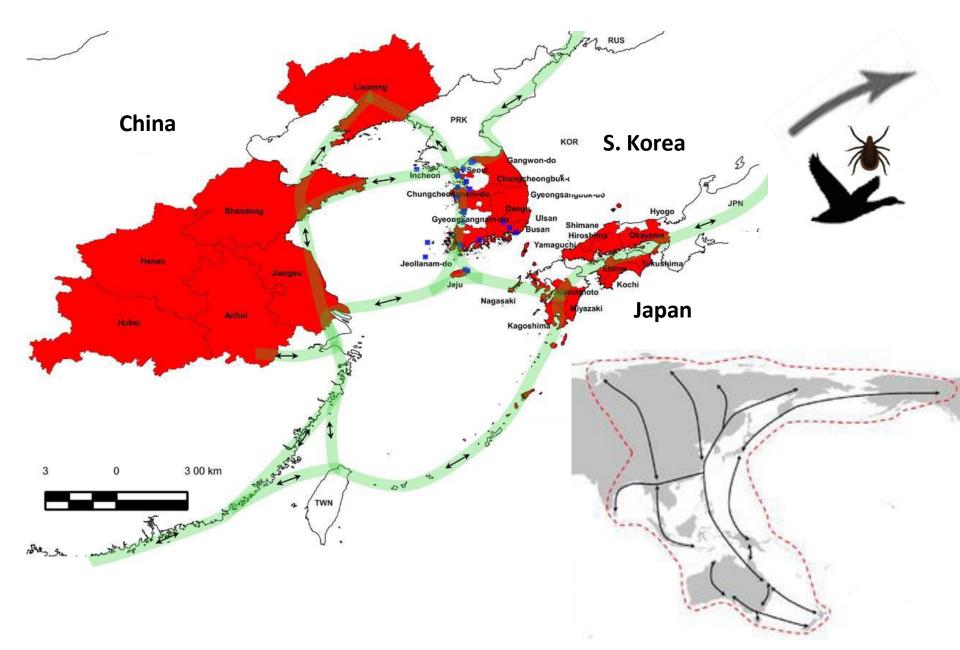




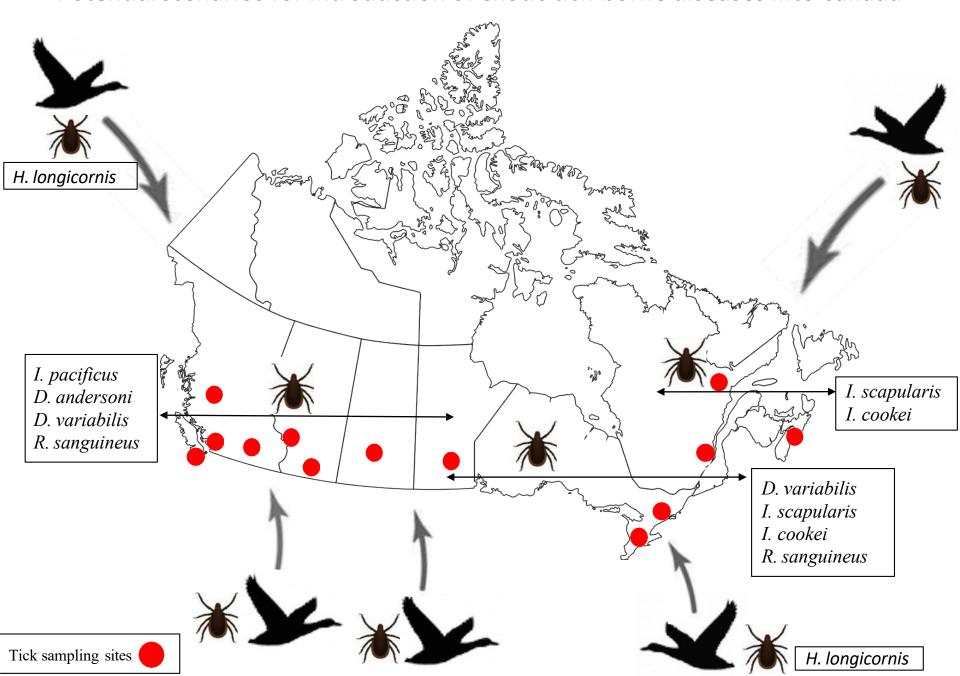
Emergence of tick-borne diseases on the rise due to climate change



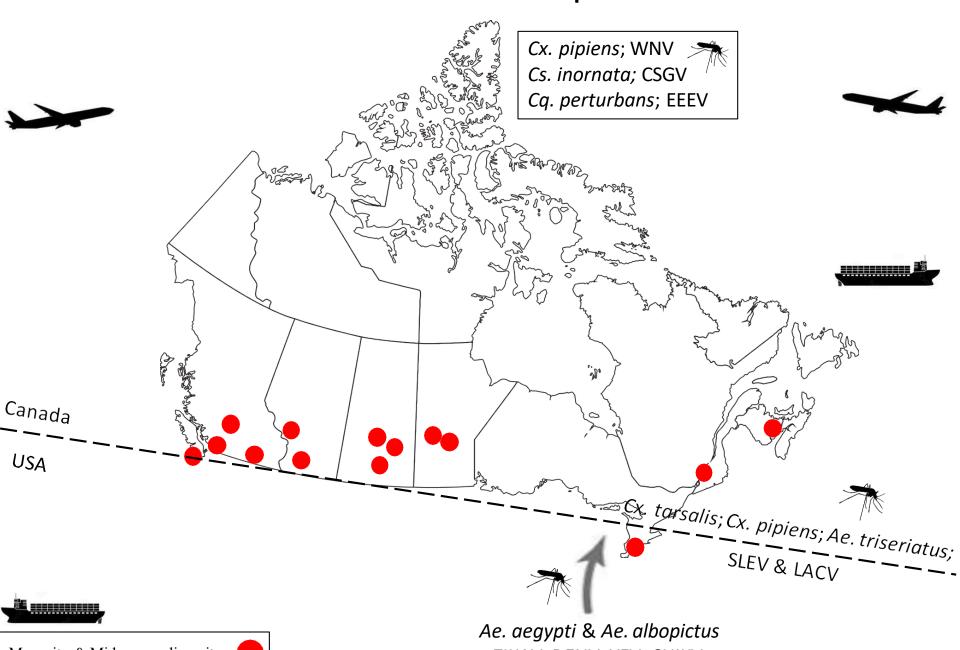
Emergence of tick-borne diseases on the rise due to climate change



Potential scenarios for introduction of exotic tick-borne diseases into Canada



Potential scenarios for introduction of exotic mosquito-borne diseases into Canada



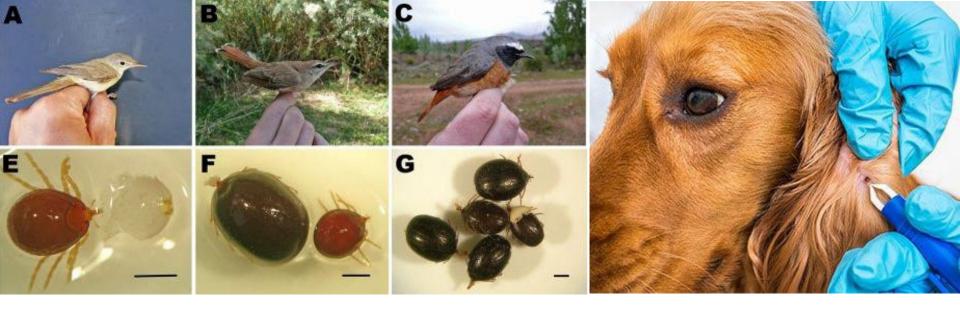
Mosquito & Midge sampling sites

ZIKAV, DENV, YFV, CHIKV

Travel without restriction



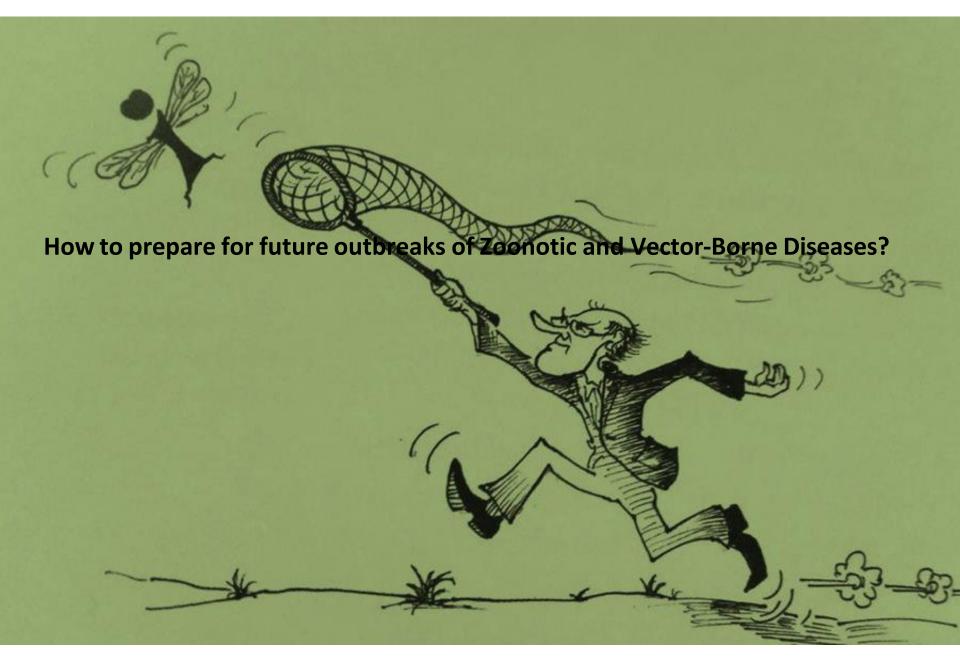








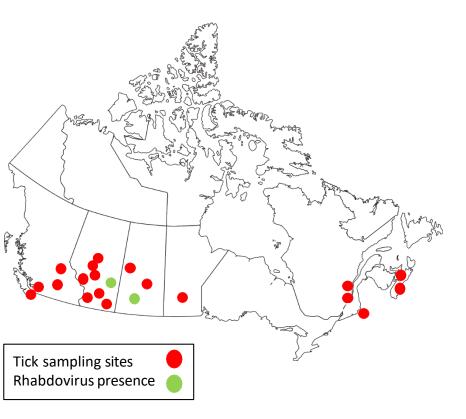
Crimean-Congo Hemorrhagic Fever Virus in Ticks from Migratory Birds, Morocco, D. 2013 https://www.mondou.com/en-CA/blogs/advice/dog/protect-your-pet-against-ticks-ad26.html



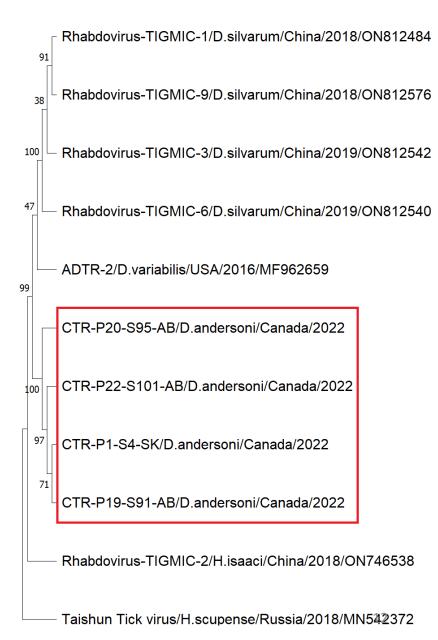
Metagenomics for novel/emerging pathogen discovery and characterization



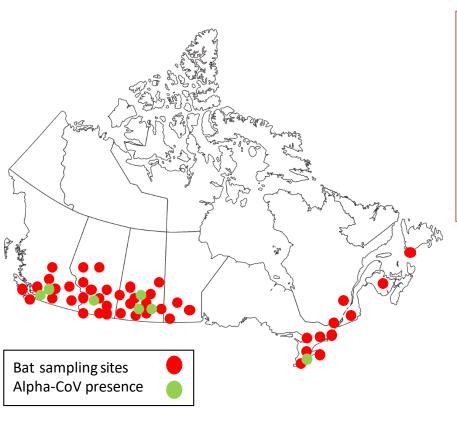
Characterization of a novel Rhabdovirus from ticks in Canada



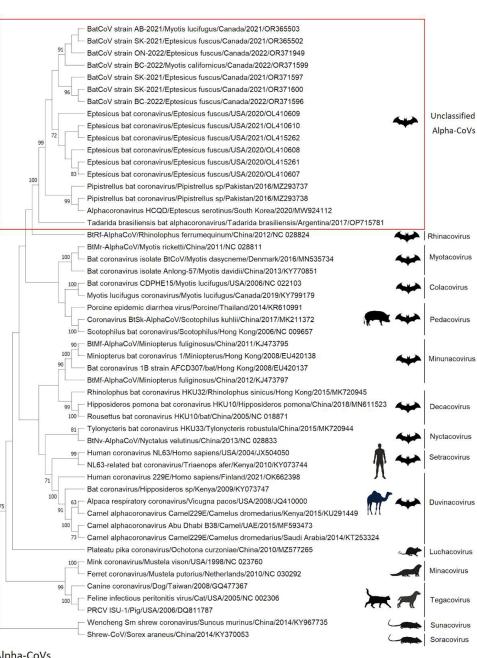




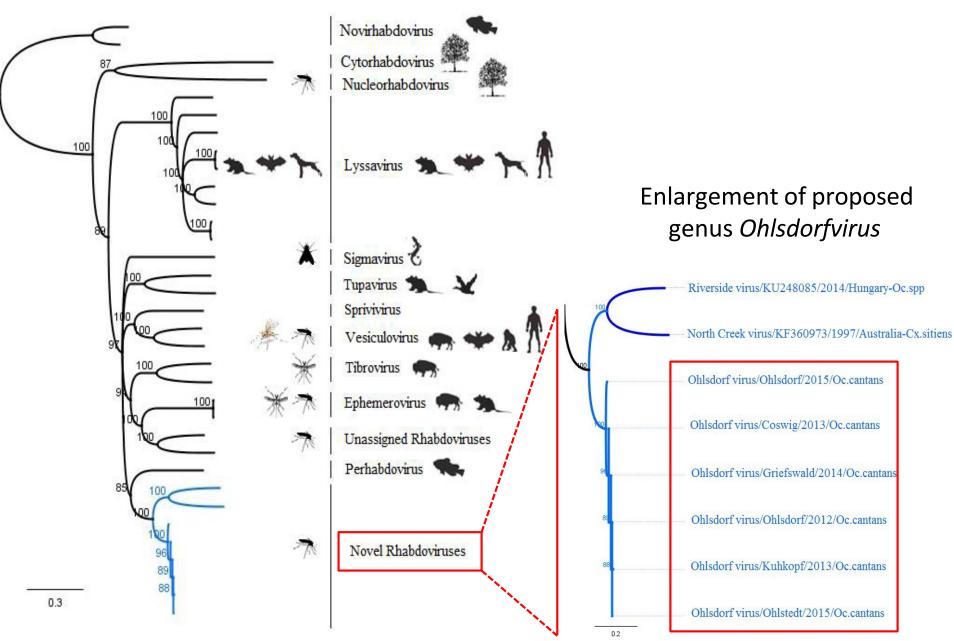
Characterization of an unclassified Coronavirus in bat samples in Canada



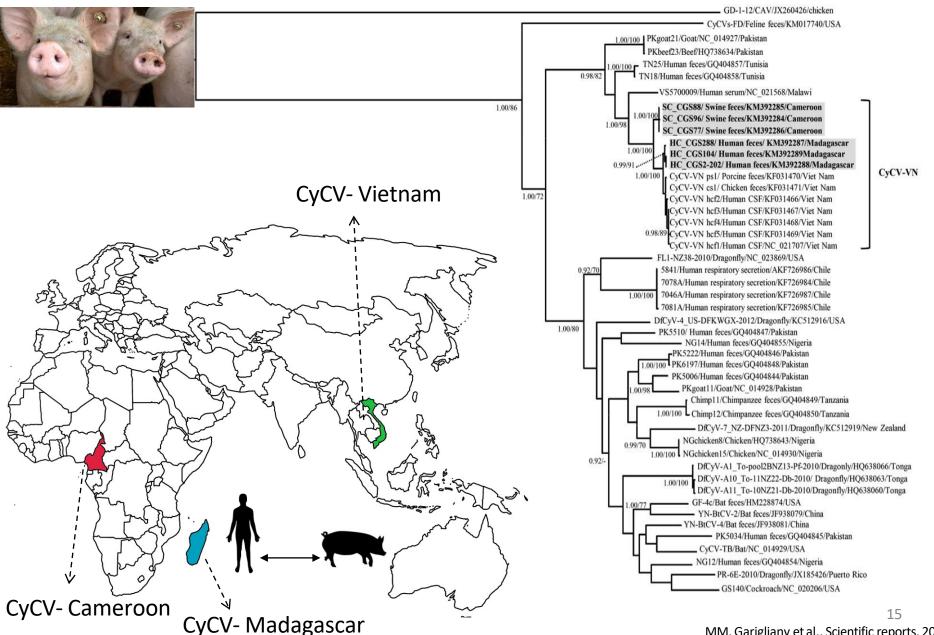


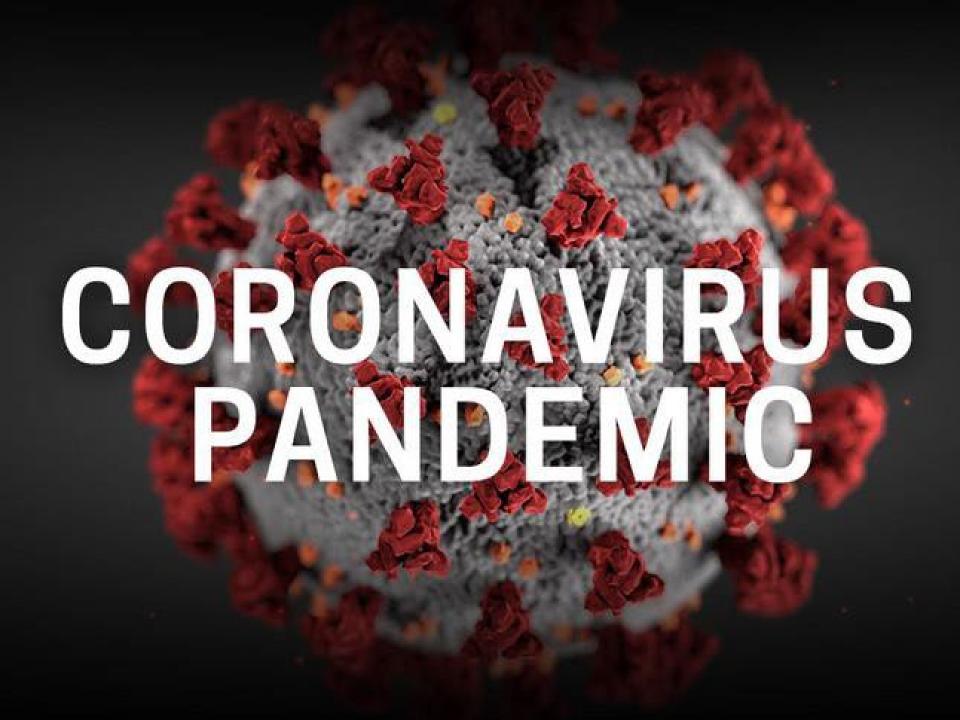


Characterization of a novel Rhabdovirus from mosquitoes in Germany

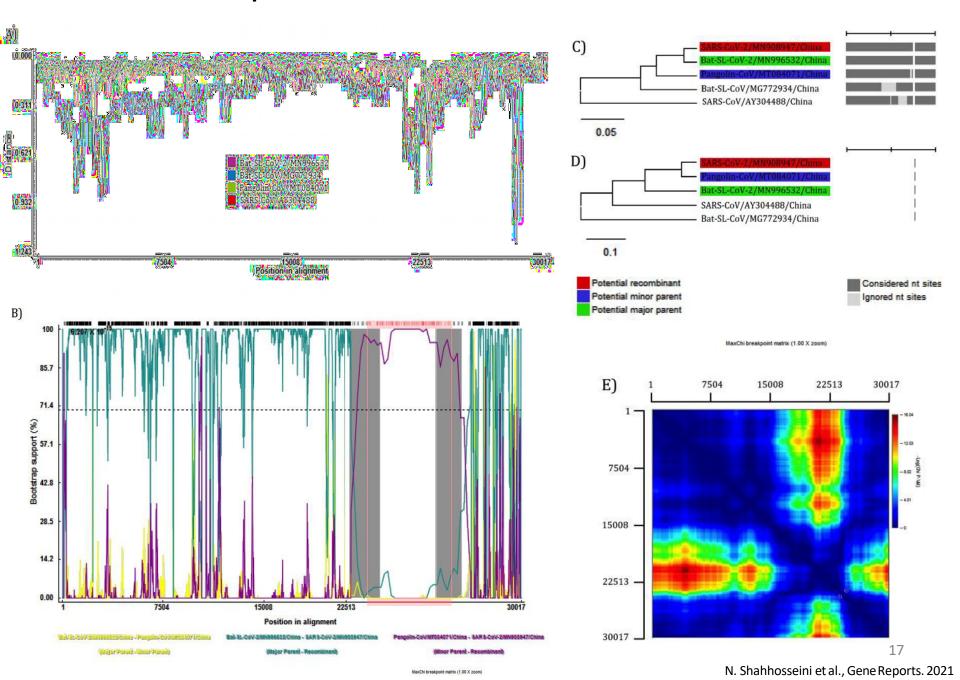


Characterization of an exotic pathogenic Cyclovirus in human and pigs in Africa



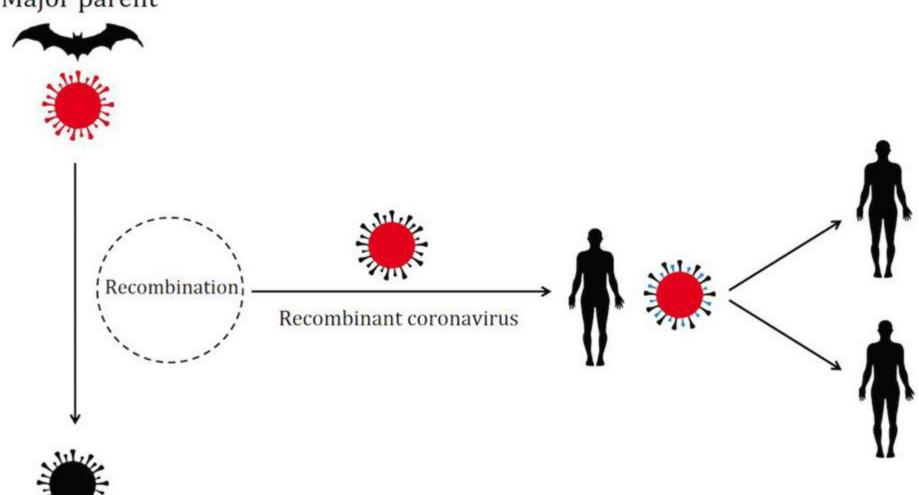


SARS-CoV-2 spillover transmission due to recombination event



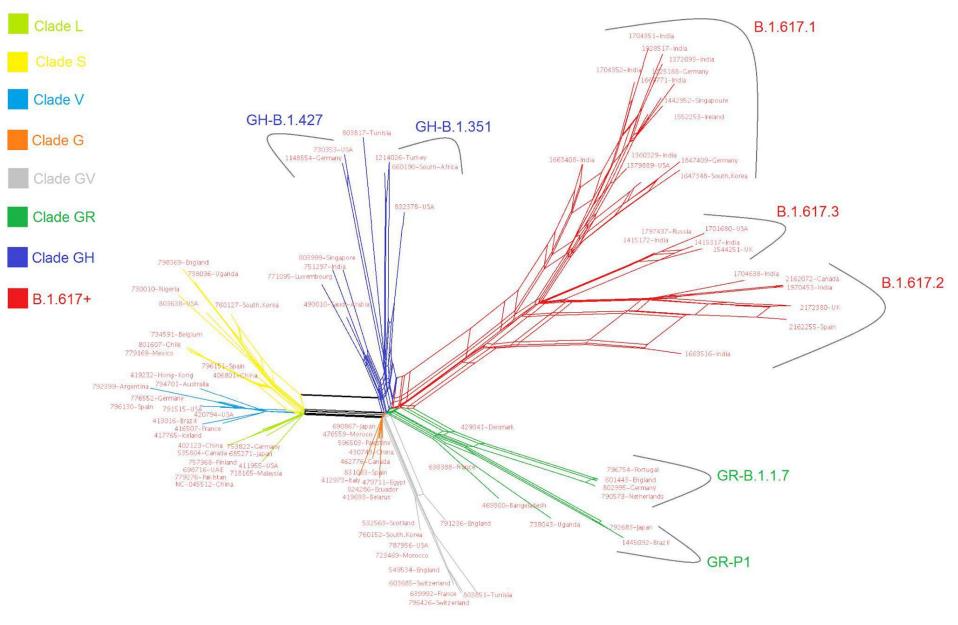
Major parent

Minor parent

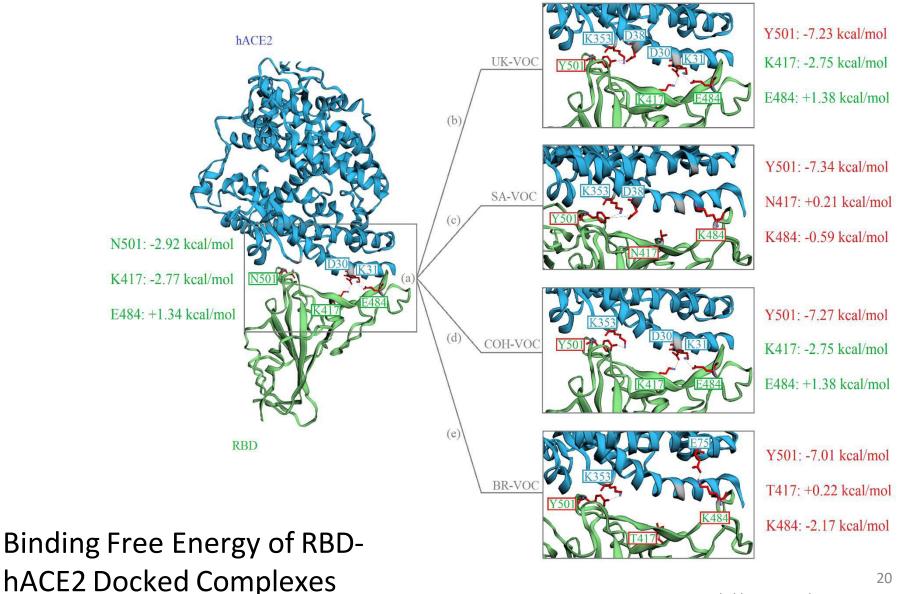




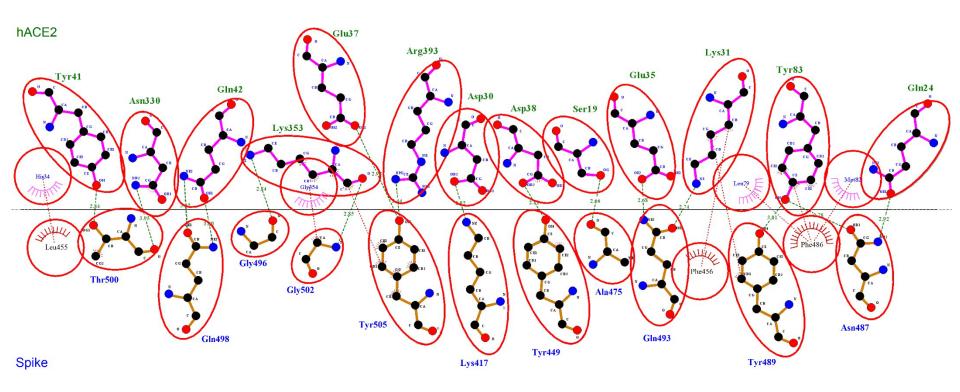
Split Tree of SARS-CoV-2 Variants of Concern



In Silico Docking of Novel SARS-CoV-2 Variants of Concern



Molecular Interactions Between hACE2 with WT and Delta Variant Spike



(A) The interacting residues Interacting residues Residue accessibility and residue accessibility Amino acid BFE (kcal/mol) (WT; A, Delta variant; D) hACE2 (C) (D) Interacting residues Residue accessibility Amino acid BFE (kcal/mol) hACE2 Structures of hACE2 docked with WT (B) and (F) Delta variant (E) 22 N. Shahhosseini et al., Gene Reports. 2022

Development of two DNA vaccines for SARS-CoV-2

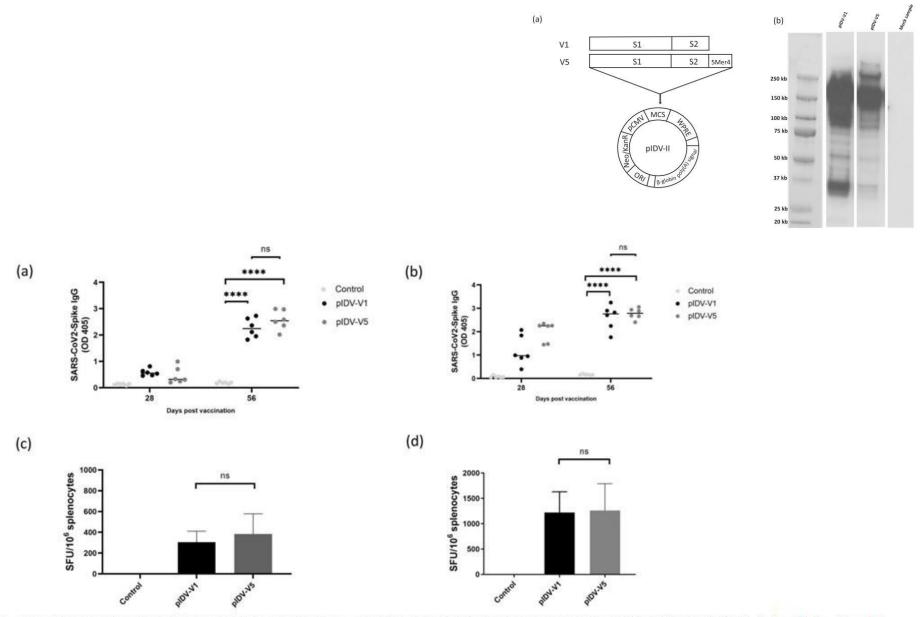


Fig. 2 Humoral and cell-mediated responses in vaccinated mice. a BALB/c and b C57BL/6 mice were immunized with pIDV-II-SARS-CoV2- Spike-Vectors 1 and 5



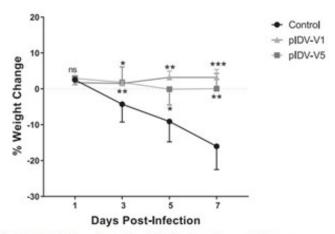


Fig. 3 Weight loss in animals following virus challenge.

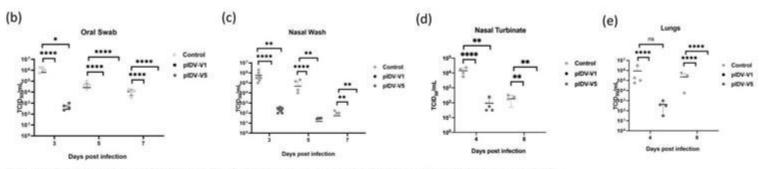


Fig. 4 Vaccination reduces viral shedding and viral RNA load in SARS-Cov2 challenged hamsters.

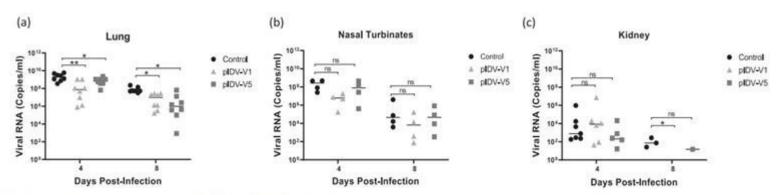
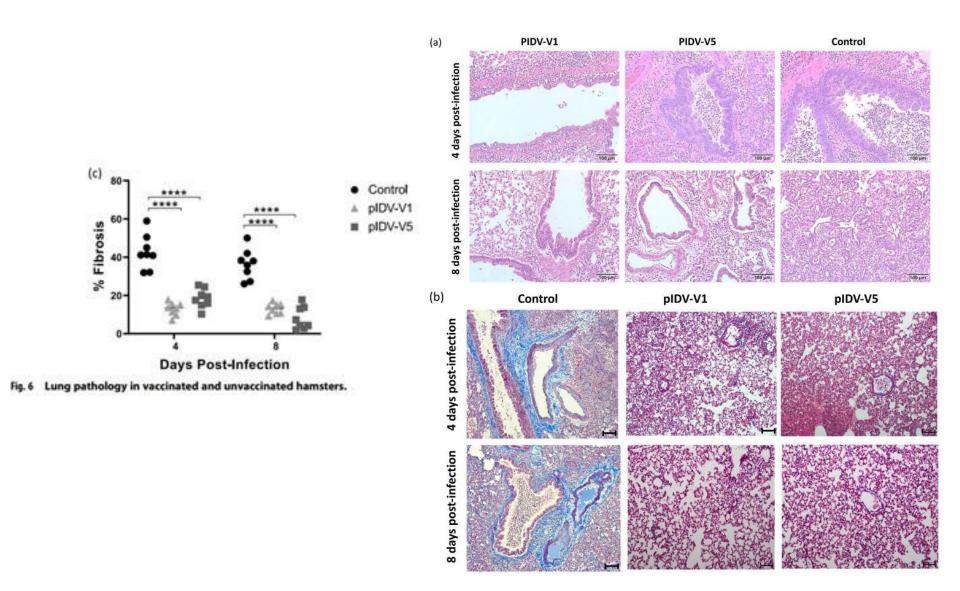


Fig. 5 Comparison of SARS-CoV-2 viral load in different tissues.





Acknowledgement









Thank you for your attention!

OPINION

What if the Next Pandemic Happens Tomorrow?



Questions?