

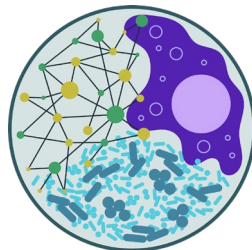


SIMON FRASER
UNIVERSITY

NanoVengers:

Oxford Nanopore Technologies sequencing takes on antimicrobial resistant pathogens and vaccine development

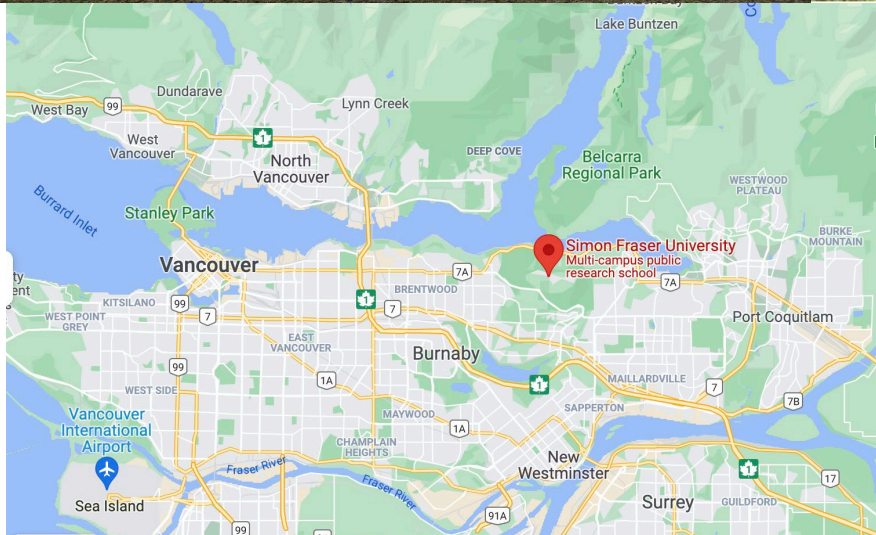
Amy H.Y. Lee, Ph.D. and Michael Trimble, Ph.D.
Department of Molecular Biology and Biochemistry
Simon Fraser University



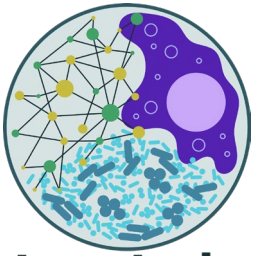
Lee Lab

SFU

SIMON FRASER UNIVERSITY



*Simon Fraser University acknowledges the unceded Traditional Coast Salish Lands including the Tsleil-Waututh (*səlilw̓ataʔt*), Kwikwetlem (*kʷikw̓əłəm*), Squamish (*Sḵw̓xw̓ú7mesh Úxwumixw*) and Musqueam (*xʷməθkʷəy̓əm*) Nations upon which our campus resides.*



Lee Lab

Systems biology approaches provide novel diagnostic, **preventative** and therapeutic strategies

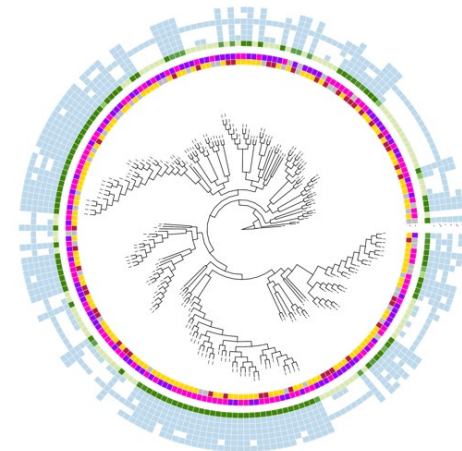
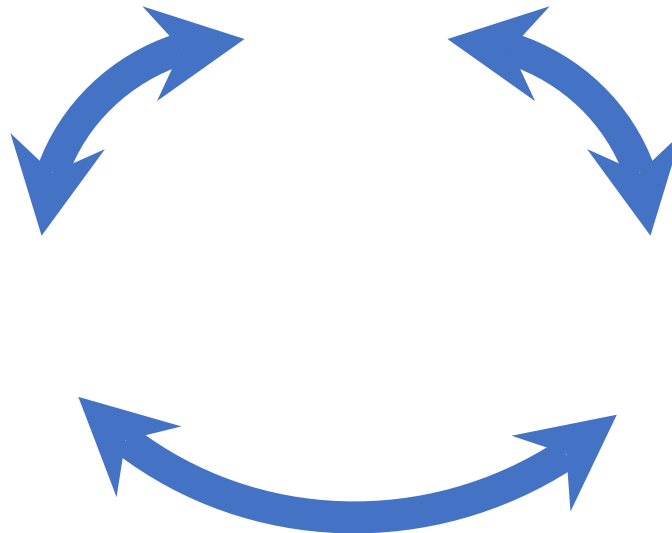
Host Immune Responses



PMID: 39472236
PMID: 38769383
PMID: 30862783
PMID: 32479514

Microbiome

Pathogen

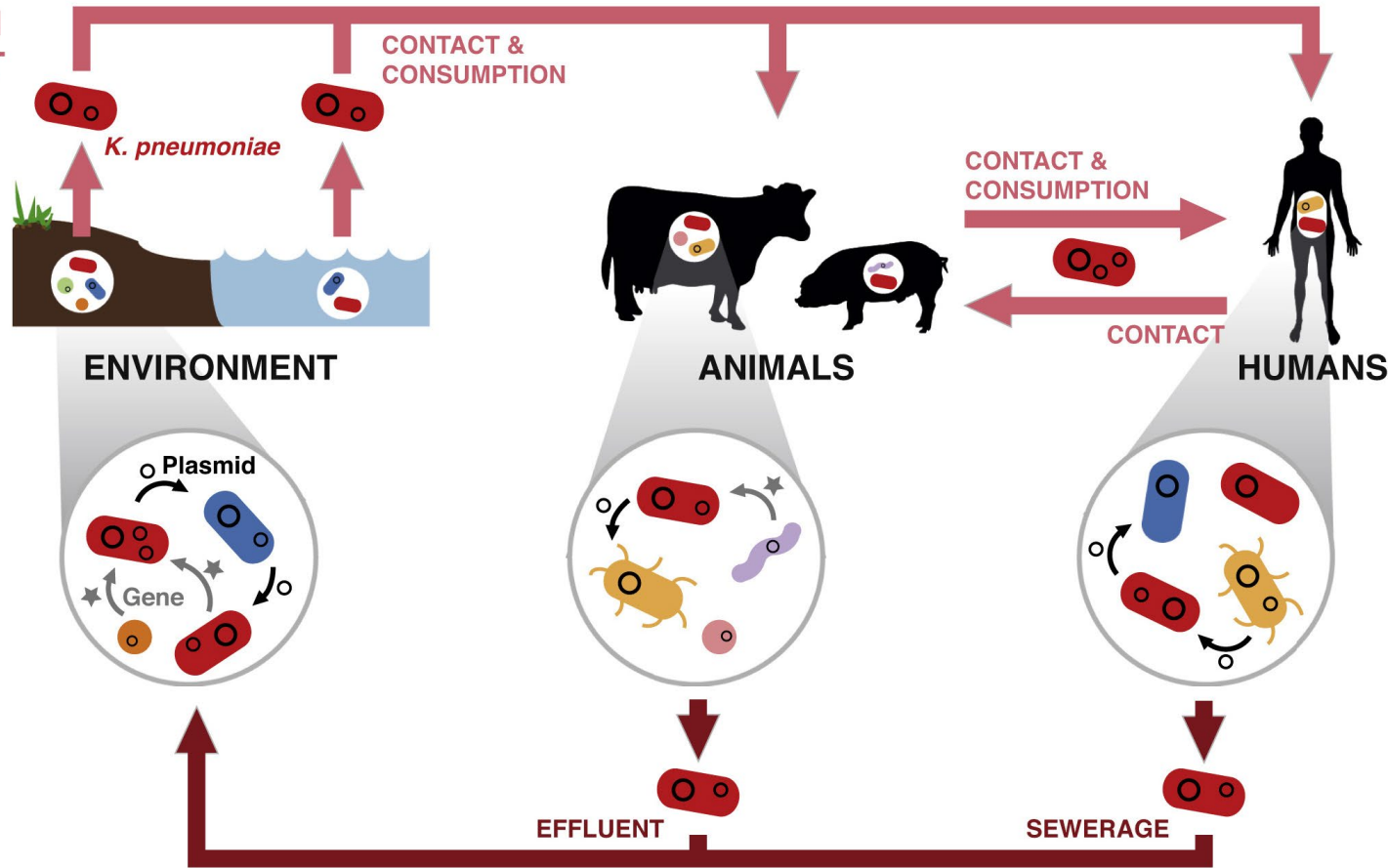


PMID: 34424159

PMID: 33329547

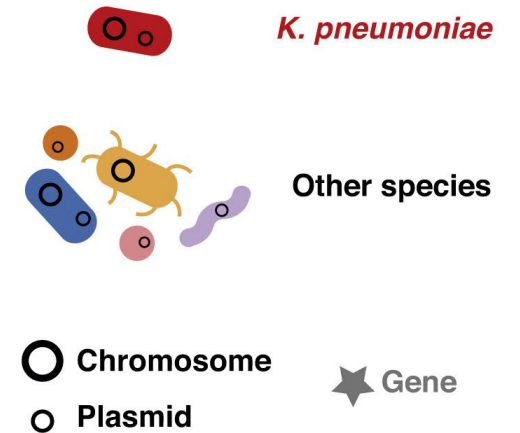
K. pneumoniae is an AMR One Health pathogen

STRAIN ACQUISITION FROM ENVIRONMENT



K. pneumoniae strains acquire AMR genes and plasmids in all niches, which they can carry between niches and donate to other human pathogens.

HORIZONTAL GENE TRANSFER



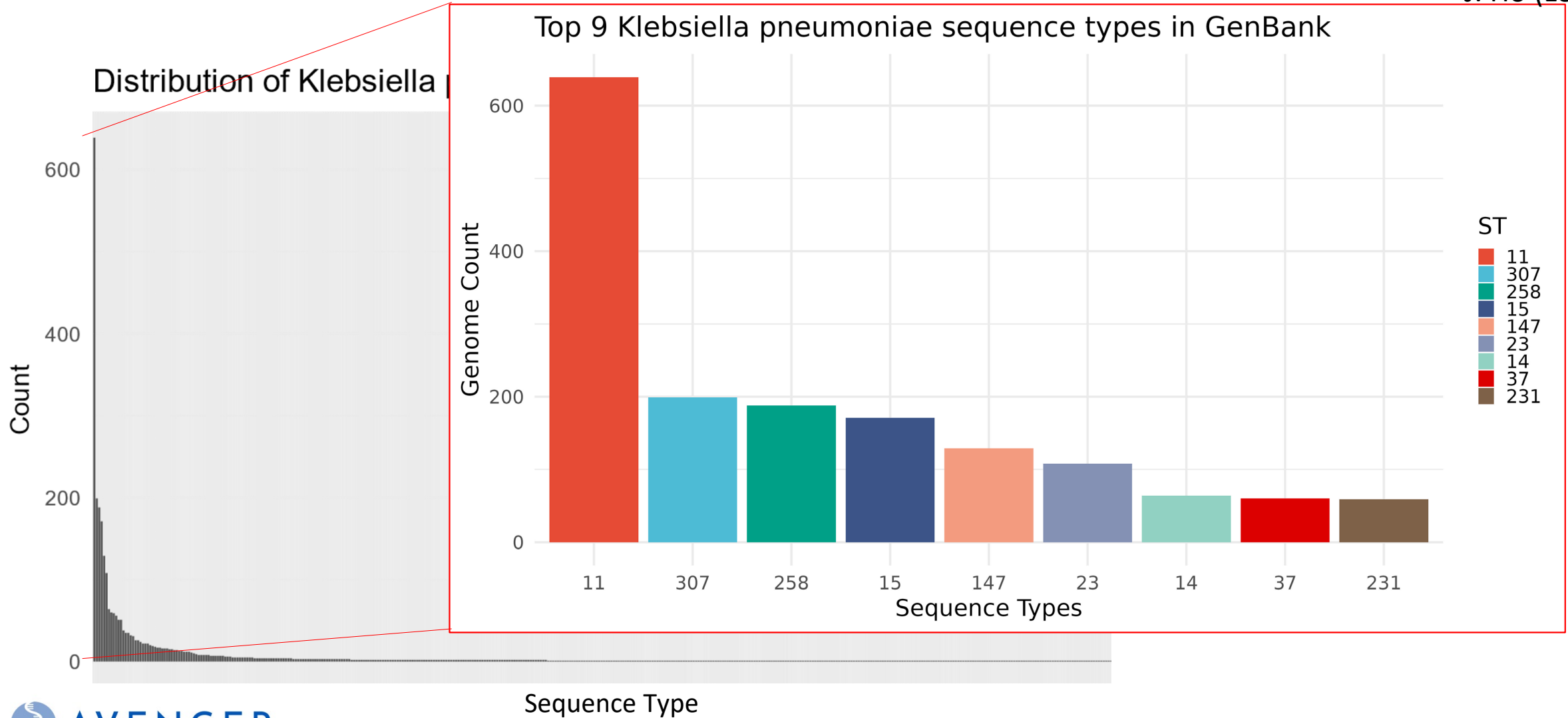
STRAIN DEPOSITION IN ENVIRONMENT

Current Opinion in Microbiology

NCBI reference genomes do not capture sufficient genetic diversity



J. Ho (Lee Lab)

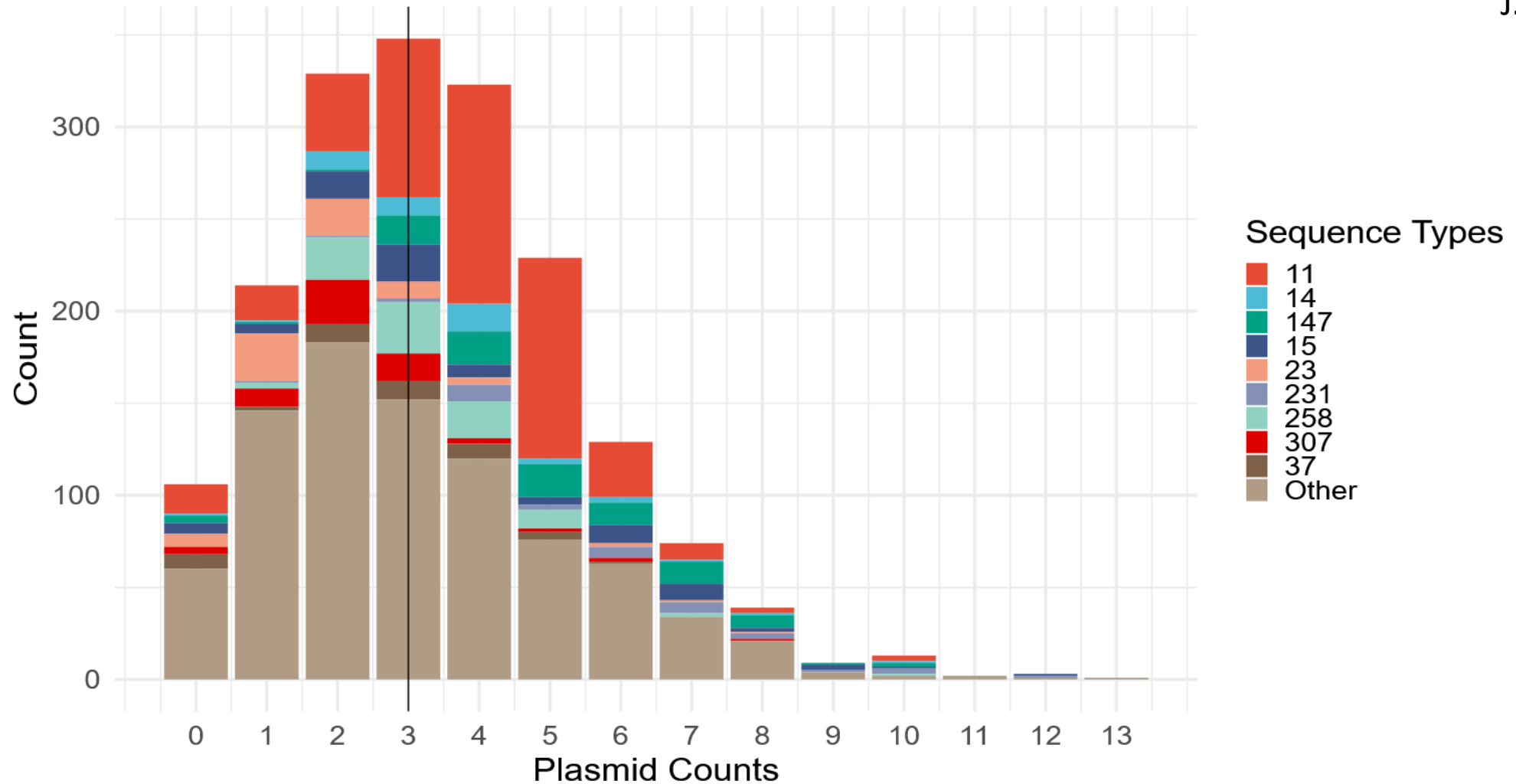


K. pneumoniae: AMR trafficker with lots of plasmids



J. Ho (Lee Lab)

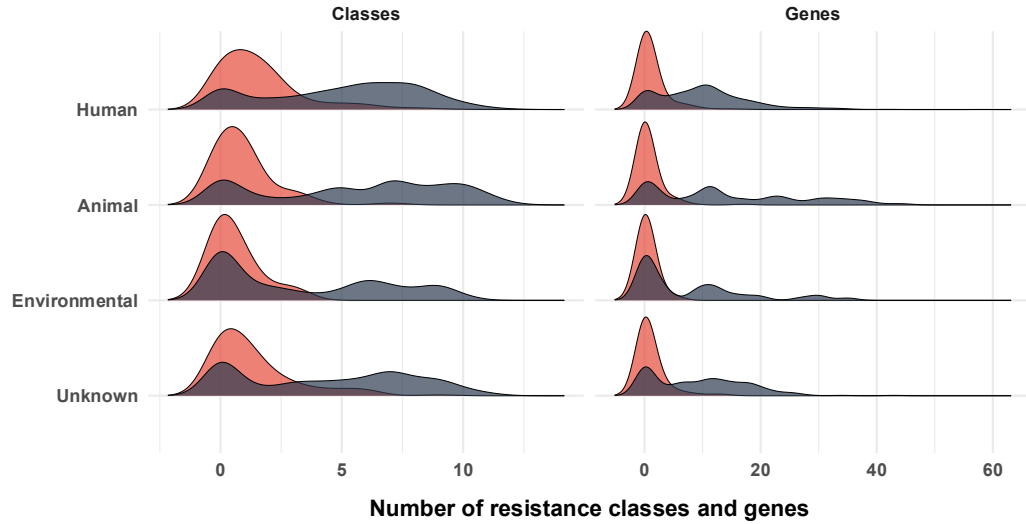
Distribution of *Klebsiella pneumoniae* plasmid counts in RefSeq



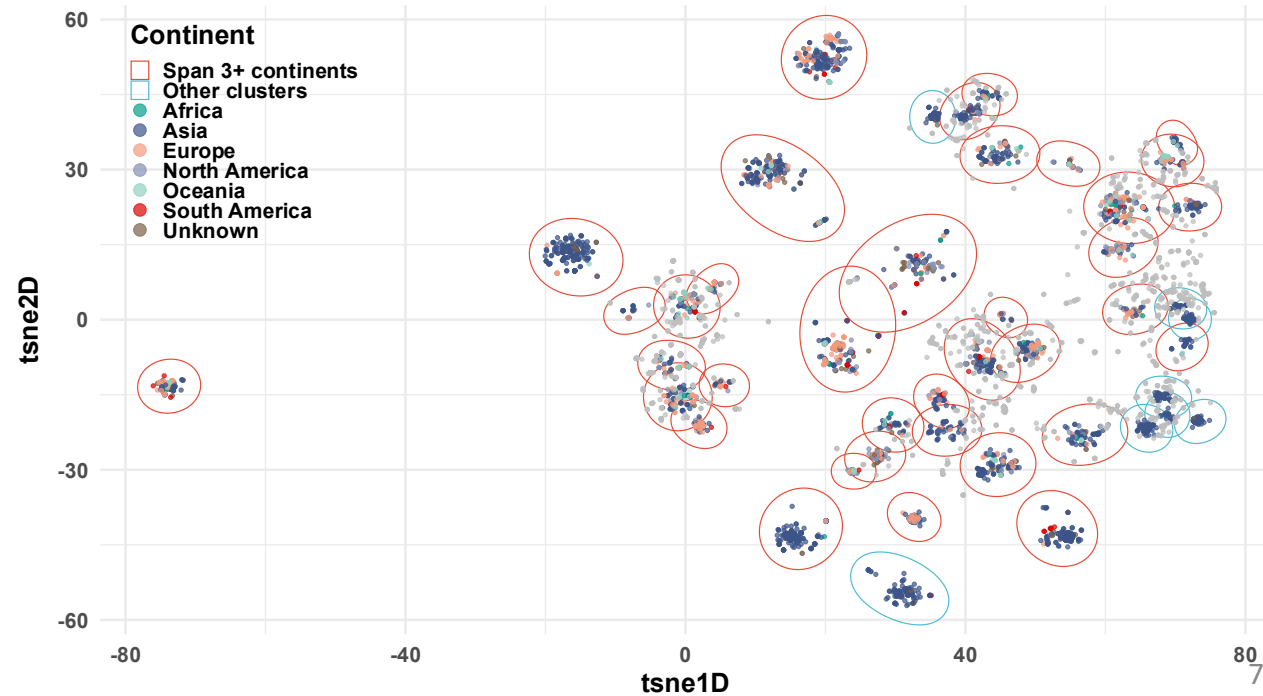
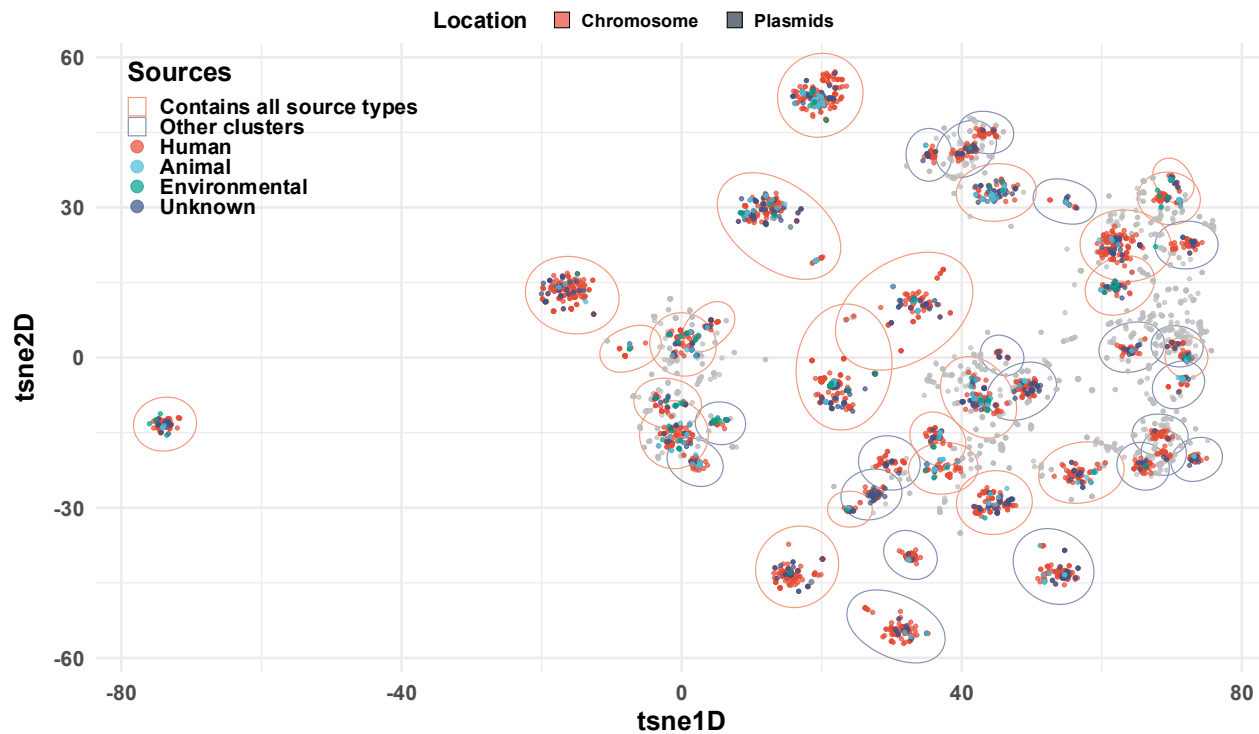
AMR and plasmids in *Klebsiella pneumoniae* are widespread



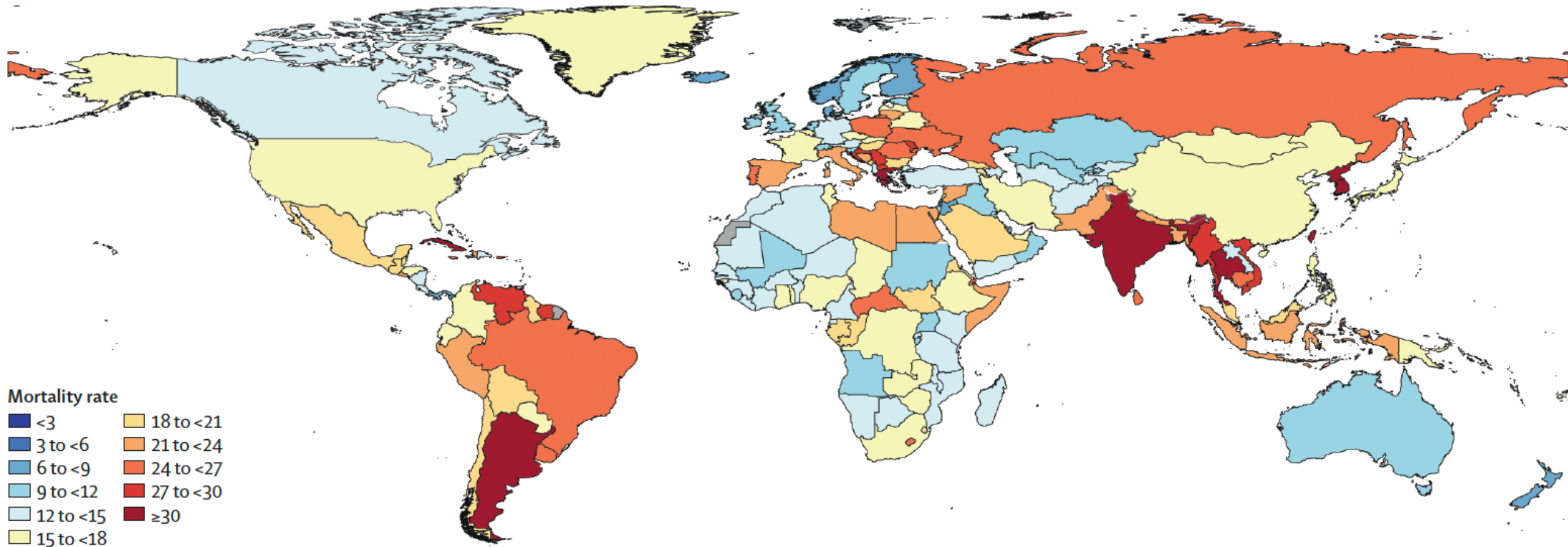
J. Ho (Lee Lab)



- Resistance is primarily carried on plasmids regardless of isolation source
- Similar plasmids are found across all isolation sources and around the globe

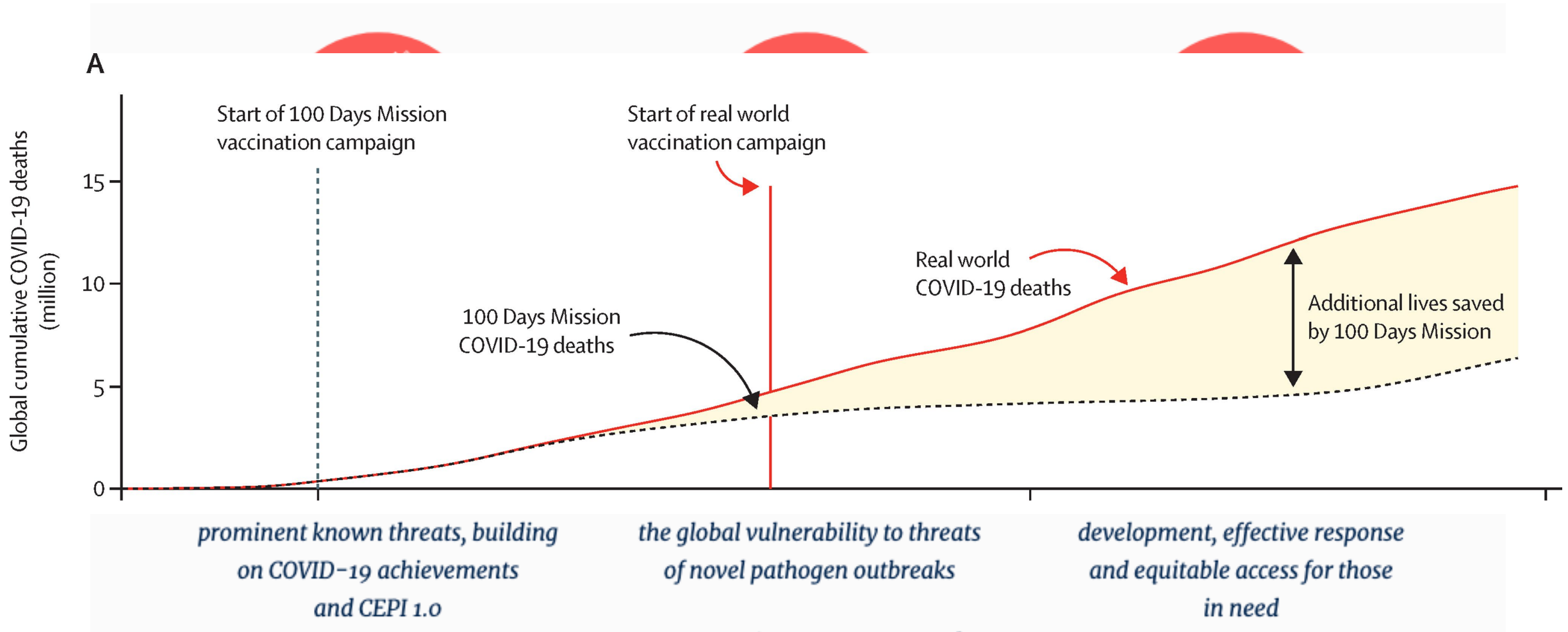


The Hidden Pandemic: Antimicrobial Resistance (AMR)



- ~ 1.2 million people died in 2019 from antibiotic-resistant bacterial infections
- More than 39 million deaths from AMR infections estimated between now and 2050

The 100 Days Mission – Vaccine Development



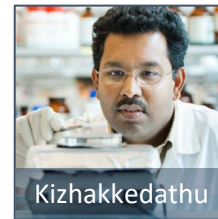
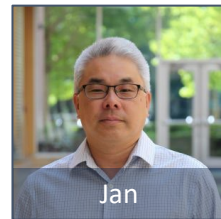
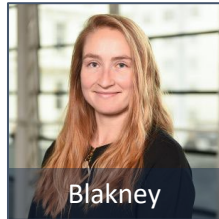


Advanced RNA LNP Vaccines Engineered with Next-Generation designs to Enhance pandemic Readiness

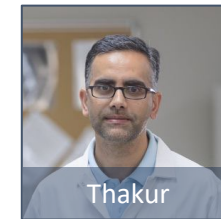
Program Directors: Dr. Pieter Cullis and Dr. Anna Blakney



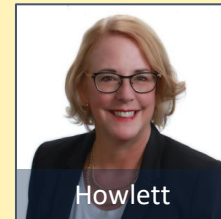
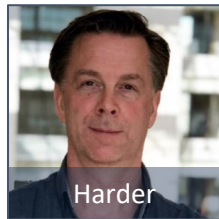
RNA LNP Technology



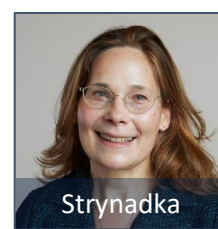
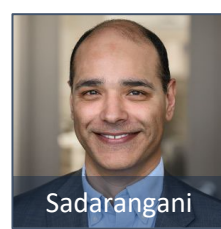
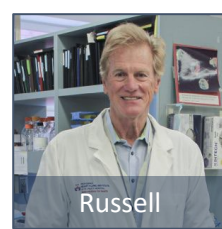
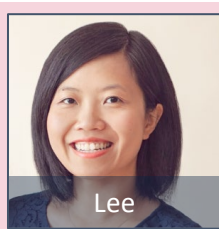
Translation



Immunology



Antigen Design



Biomufacturing

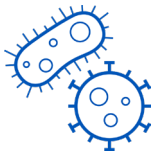


AVENGER will build domestic capacity for a 100-day response using next-generation RNA LNP technology



- **Aim 1: Develop high-performing, RNA LNP vaccines with improved efficacy**

- **Pillar – Research and innovation, Infection prevention and control**
 - LNP with improved potency, stability, and safety
 - saRNA and mRNA optimized for low-dose vaccines
 - Formulations that induce long-lived adaptive responses
 - Formulations for aged and frail populations



- **Aim 2: Develop vaccine candidates against antimicrobial-resistant (AMR) bacteria (*K. pneumoniae*, *S. pneumoniae*, *S. aureus*)**

- **Pillar – Surveillance; R&I, Infection prevention and control**
 - Build a strategy and extensive database for bacterial vaccine antigen selection and optimization
 - Obtain and sequence clinical isolates – long read sequencing (Nanopore)
 - At least one RNA LNP vaccine for an AMR bacterial pathogen

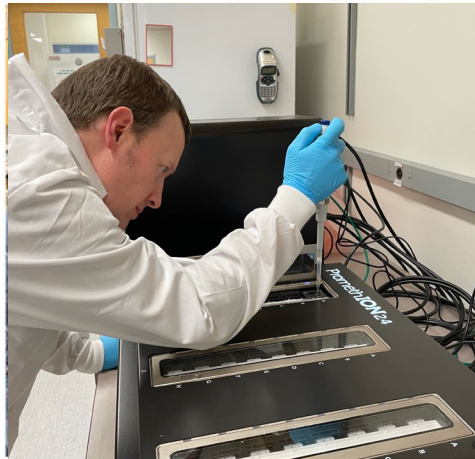


- **Aim 3: Bio-manufacture a candidate vaccine and test it in an NHP infection model**

- **Pillar – R&I, Infection prevention and control**
 - Logistics and implementation of a biomanufacturing pipeline
 - Adaptable platform technology

Sequencing of clinical isolates:

Dr. O. Pacios (Russell/Lee Labs)



Dr. M. Trimble (Lee Lab)



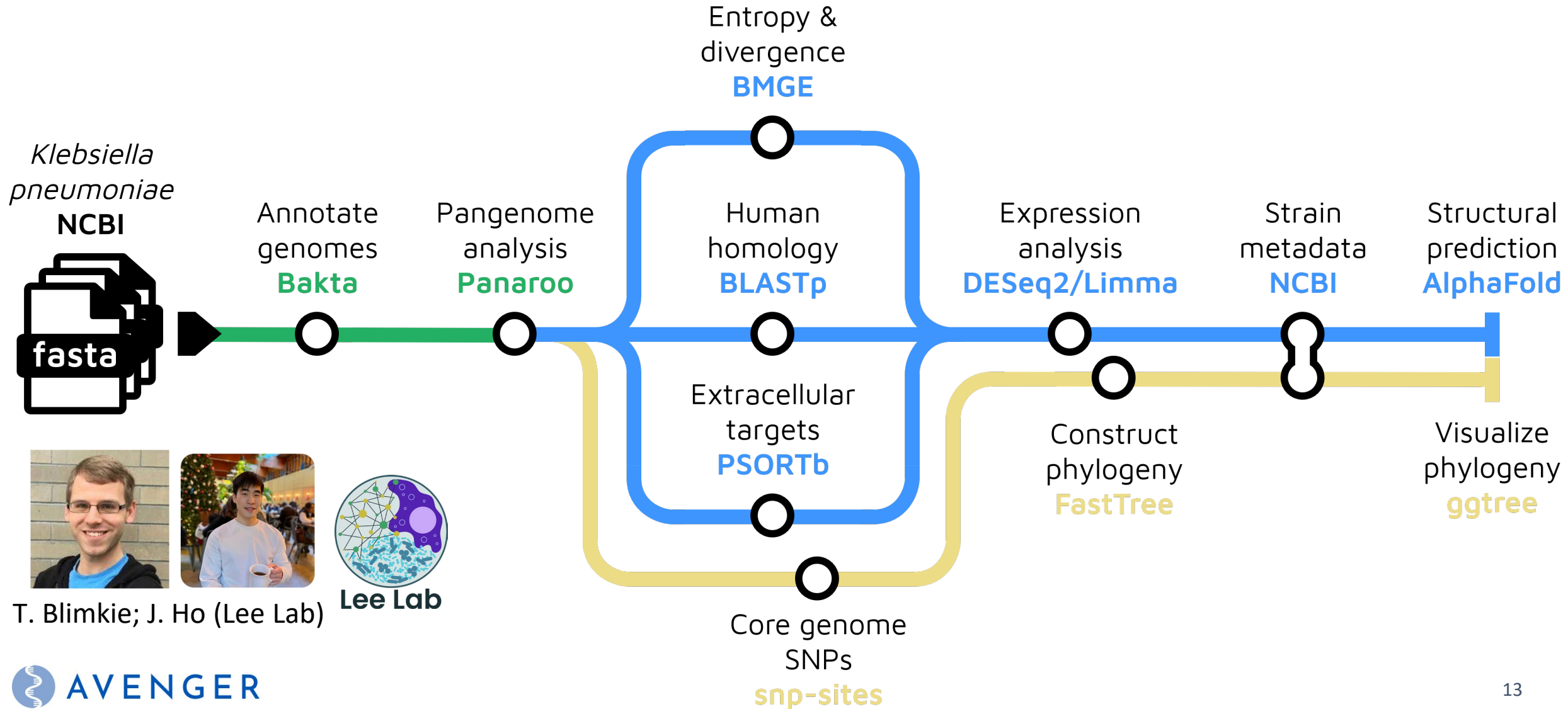
Dr. B. Raymond; A. Redey (Lee

➤ Thanks to Carmen Li and

Species	To be sequenced	Sequenced
<i>K. pneumoniae</i>	101	> 400
<i>S. aureus</i>	356	> 60
<i>S. pneumoniae</i>	114	-
MRSA	17	-
Total	588	



Reverse Vaccinology to the Rescue!

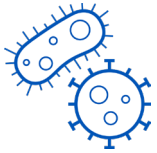


AVENGER Next Steps



RNA LNP technology development


- Identified PEG substitutes
- Designed new alphaviral vectors for saRNA backbones
- Establishing immunological assessments of benchmark vaccines
- Establishing immune alterations in variably aged and frail mice
- Standardizing immunoassays for evaluating new formulations





Bacterial antigen discovery and engineering

- Candidate antigens for *K. pneumoniae* and *S. aureus* have been engineered from existing data sets
- Acquiring clinical isolates of AMR bacteria (~900)
- Building a whole-genome assembly and reverse vaccinology pipeline
- Establishing murine infection models

ADVANCED FUNCTIONAL MATERIALS

Research Article | [Open Access](#) | 

An Ultrahydrating Polymer that Protects Protein Therapeutics and RNA-Lipid Nanoparticles Against Freezing, Heat and Lyophilization Stress



Anna Herrmann, Srinivas Abbina, Nuthan Vikas Bathula, Peyman Malek Mohammadi Nouri, Irina Chafeeva, Iren Constantinescu, Emaan Abbasi, Usama Abbasi, Matthew Drayton, Haiming Daniel Luo, Haisle Moon, Arshdeep Gill, Yu Xi, Allan K. Bertram, Caigan Du, Rainer Haag, Anna K. Blakney , Jayachandran N. Kizhakkedathu 

First published: 01 August 2024 | <https://doi.org/10.1002/adfm.202406878>

Molecular Therapy

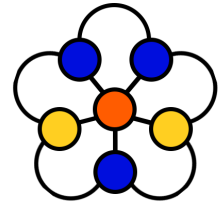
Original Article

Alphaviral backbone of self-amplifying RNA enhances protein expression and immunogenicity against SARS-CoV-2 antigen

Irafasha C. Casmil ^{1,2}, Nuthan V. Bathula ^{1,2}, Cynthia Huang ^{1,2}, Christopher J. Wayne ^{1,2}, Evan S. Cairns ^{1,2}, Josh J. Friesen ^{1,2}, Shekinah K. Soriano ^{1,2}, Suiyang Liao ^{1,2,3}, Chia H. Ho ^{1,2}, Kristen Y.S. Kong ¹, Anna K. Blakney ^{1,2}  

<https://doi.org/10.1016/j.ymthe.2024.12.055>

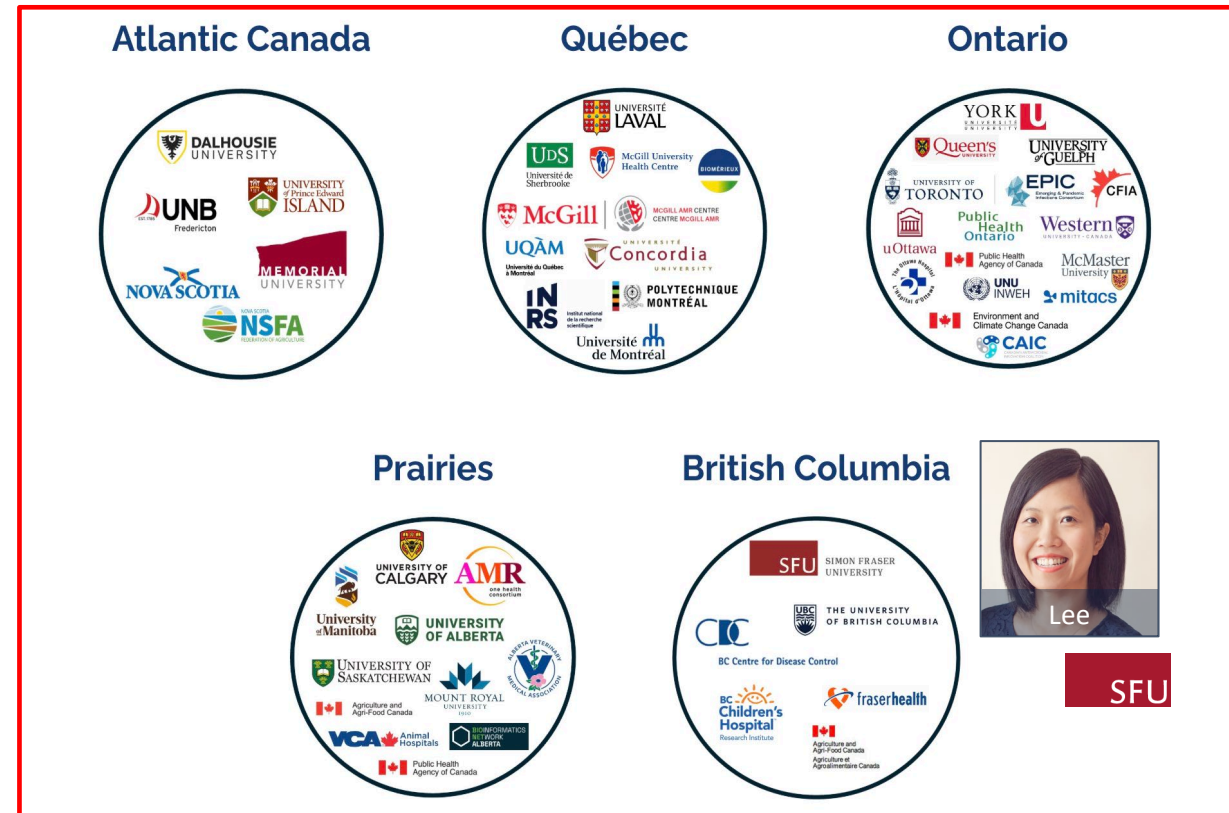
CIHR Funded One Health AMR Training Platform: CAN-AMR-NET



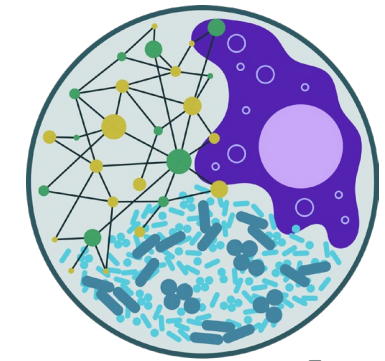
CAN-AMR-Net

<https://canamrnet.org/>

- Central Coordination through the University of Calgary
 - Encompassing **5 Regional** hubs under 4 themes
1. One Health/AMR Translational Competencies
 2. Transdisciplinary Training & Career Readiness
 3. Excellence in Knowledge Translation & Communication
 4. Integration of Indigenous Knowledge and Community-Led Research



Acknowledgements



Lee Lab



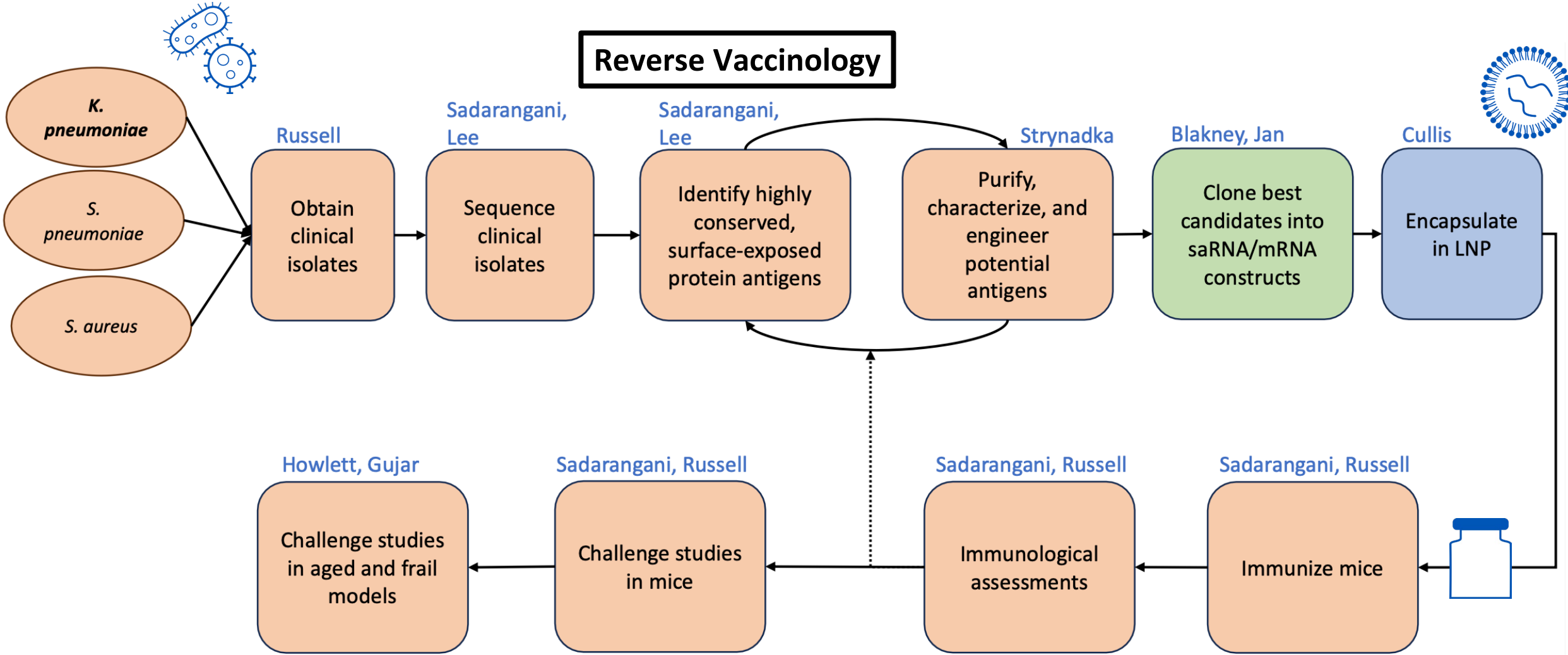
SFU Collaborators: BCCHRI (Malawi & 16S Diagnostics): Dalhousie (Neonatal Mouse Sepsis):



Funding Support:



AVENGER takes on 3 Key Priority AMR Pathogens



Klebsiella pneumoniae: a Global AMR Threat

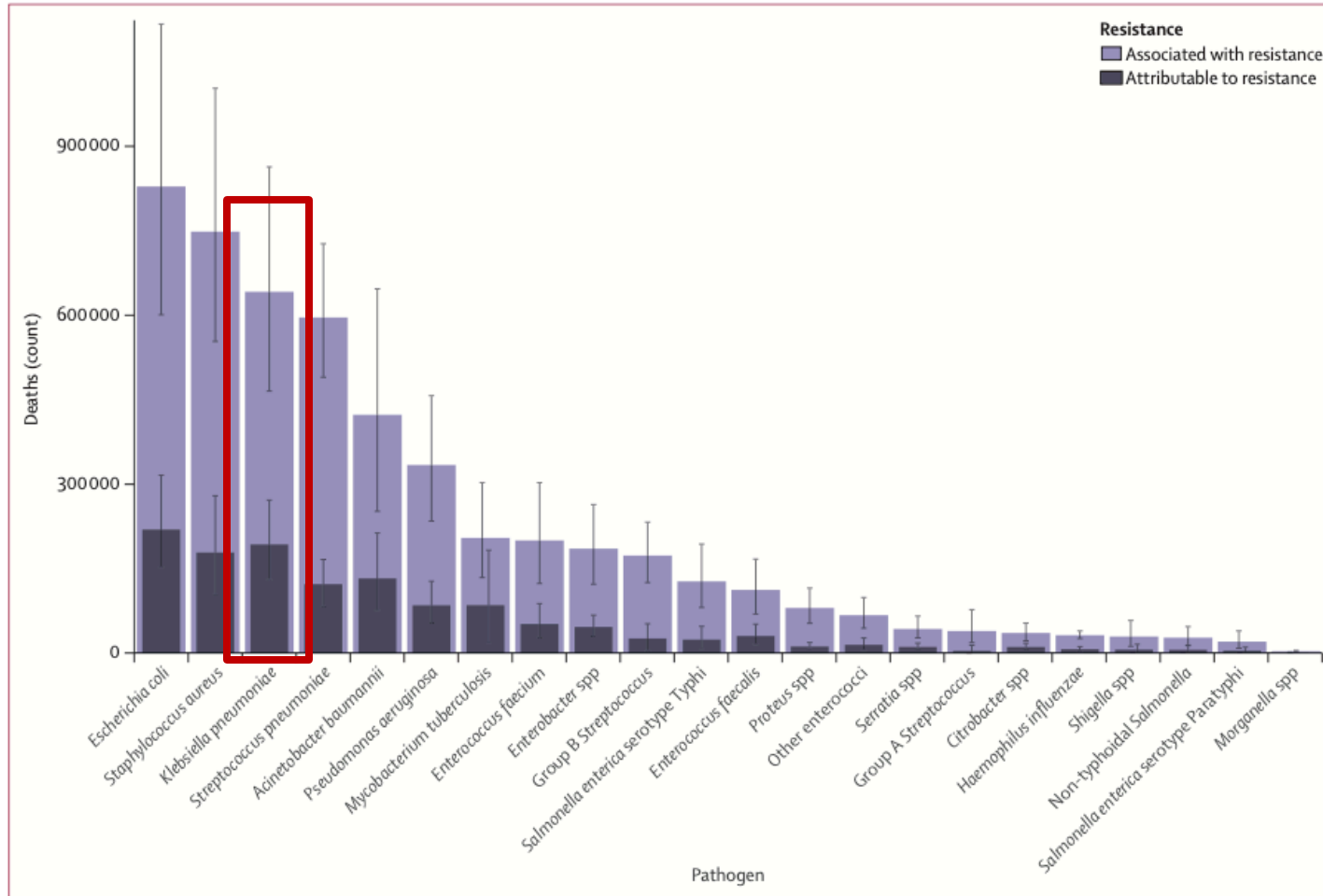
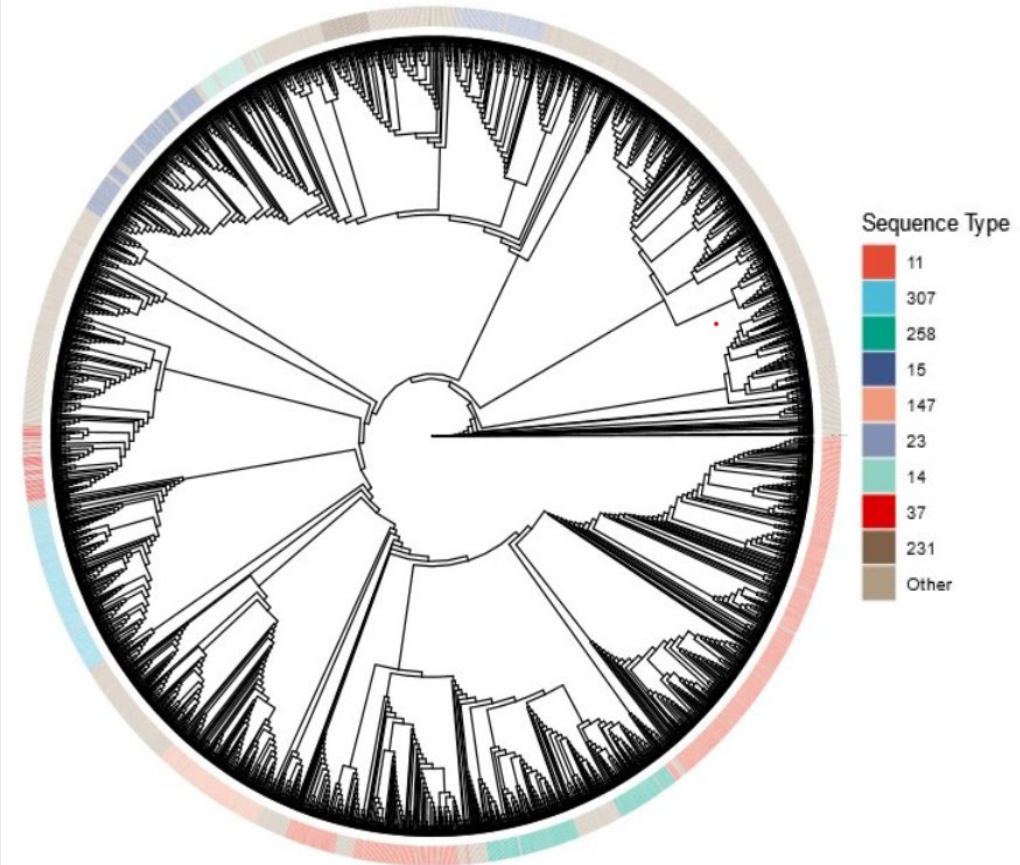


Figure 4: Global deaths (counts) attributable to and associated with bacterial antimicrobial resistance by pathogen, 2019
 Estimates were aggregated across drugs, accounting for the co-occurrence of resistance to multiple drugs. Error bars show 95% uncertainty intervals.



K. pneumoniae: Vaccine Evaluation Profile

KLEBSIELLA PNEUMONIAE	
<p>Health impact:</p> <p>Direct health impact</p> <p>1.0 Mortality</p> <p>0.0 Morbidity</p> <hr/> <p>Impact on AMR reduction</p> <p>1.0 Antibiotic use</p> <p>2.0 Urgency of AMR threat</p> <hr/> <p>Secondary health impact</p> <p>None identified</p> <hr/> <p>Sub-population benefits</p> <p>Individuals with recurrent urinary tract infections, HIV, chronic lung conditions or diabetes</p> <hr/> <p>Alternative interventions</p>	<p>Probability of R&D success:</p> <p>0.0 Pipeline robustness</p> <p>0.5 Pathogen biology</p> <p>0.5 Pre-clinical and clinical R&D</p> <hr/> <p>Combination potential</p> <p>None identified</p> <hr/> <p>Acceleration potential</p> <p>None identified</p> <hr/> <p>Major barriers to development</p> <p>Lack of understanding of disease epidemiology</p> <hr/> <p>Probability of uptake:</p> <p>1.0 Commercial attractiveness</p> <p>0.0 Expected policy stance</p> <p>1.0 Payer, government or Gavi support</p> <p>1.0 Barriers to uptake</p>