Genomic ASSETS for Livestock: Changing the face of antimicrobial use decisions for livestock

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Laboratory and bioinformatic workflows for field nanopore metagenomic sequencing and recombinase polymerase amplification of BRD pathogens.

Canada's Beef Industry

Beef is Canada's largest livestock industry, contributing more than \$17 billion to the economy from more than 60,000 farms³. Most Canadian beef is finished in large feedlots in western Canada (capacities 5,000 to > 25,000 head, in pens of 200-300 animals).

Bovine Respiratory Disease (BRD) is the largest cause of morbidity/mortality in feedlot cattle⁴. BRD is the most common reason for parenteral antimicrobial use in the feedlot⁵.

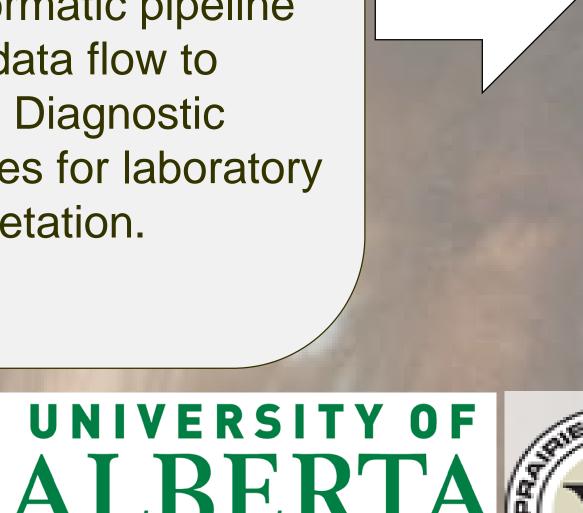


Laboratory Testing

Antimicrobial use is necessary for animal health and welfare. Diagnostic data informs precision pen medicine: ✓ Right Drug Antimicrobial ✓ Right Animals **Stewardship** ✓ Right Time

Phase 2

Optimize the feedlot pen-level sampling strategies and bioinformatic pipeline to for data flow to Prairie Diagnostic Services for laboratory interpretation.





Agriculture and Agri-Food Canada



Benefits to Canada

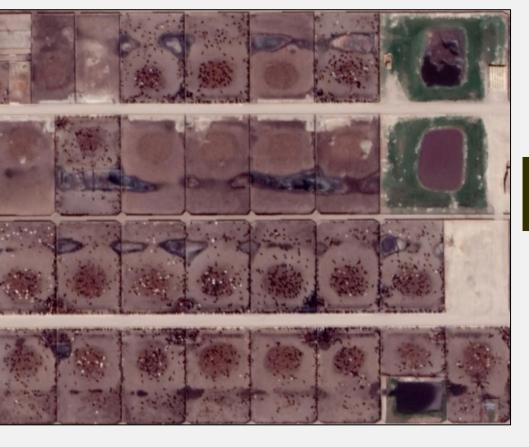
Enhanced Sustainability: improved antimicrobial use and welfare of beef cattle **Improved Disease Management:** informed prevention and treatment strategies Applications to all livestock: sampling strategies, genomics, bioinformatics and risk assessment (Prairie Diagnostic Services network laboratory hub) Market advantage: addresses World Health Organization guidelines¹ and prevents trade restrictions²

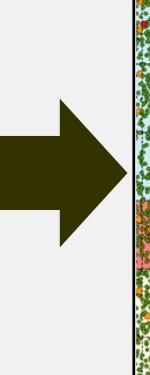
Phase 5



Economic analysis to inform the value of applying genomics to inform prudent AMU in beef feedlot







Dynamic risk assessment model incorporates genomic data to predict transmission of antimicrobial resistance and inform antimicrobial use for pens of feedlot calves.









GenomeCanada

GenomePrairie

Phase 4



Two-stage field roll-out to commercial feedlots in Alberta and Saskatchewan. Test field sequencing, bioinformatics and reporting.

References:

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[5] Brault SA, Hannon SJ, Gow SP, Warr BN, Withell J, Song J, et al. Antimicrobial Use on 36 Beef Feedlots in Western Canada: 2008–2012. Frontiers in Veterinary Science 2019;6:329

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[7] Ghosh KK, Ngeleka M, Wennekamp T, Campbell J, Waldner C, Trokhymchuk A. Evaluation of Mannheimia haemolytica antimicrobial resistance profile using broth microdilution method and nanopore whole genome sequencing. 17th Canadian Animal Health Laboratorians Network annual meeting 2018 June 11,:1. 2017 November 28;5:6-8.

