

Herd'23:

March 16th, 2023, All Seasons Resort Hotel, Bendigo Australia

 @BaesC1



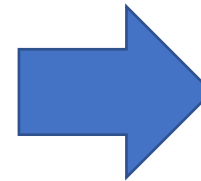
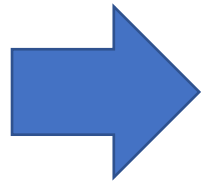
The Future of Genetic Evaluation in Canada: A Research Perspective

C. F. Baes, F. Schenkel, G. Kistemaker , K. Parker-Gaddis, R. Baldwin, A. Butty , J. Burchard, O. González-Recio, J. Lassen, M. VandeHaar, D. Segelke, R. Tempelman, F. Peñagaricano, K. Weigel, J. Koltes, F. Miglior, RDGP Consortium Partners, FARR Consortium Partners

What is a resilient cow?

an animal able to adapt rapidly to changing conditions without compromising its productivity, health or fertility while becoming more resource-efficient and reducing its environmental burden.

Past, Current and Potential Future Projects



FUNDED:

Alberta RDAR project ~\$2M

Sunalta Feed Bins & Sniffer prototypes
at Elora

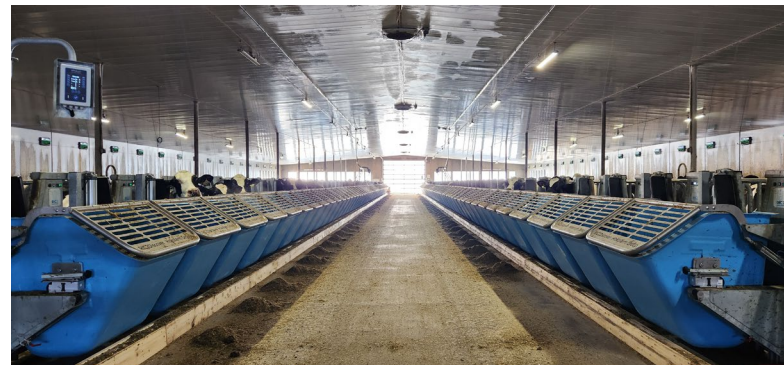
AFC Alberta Milk project \$892K

Alberta CH₄ Sniffers, KTT, weight scale
at Sunalta

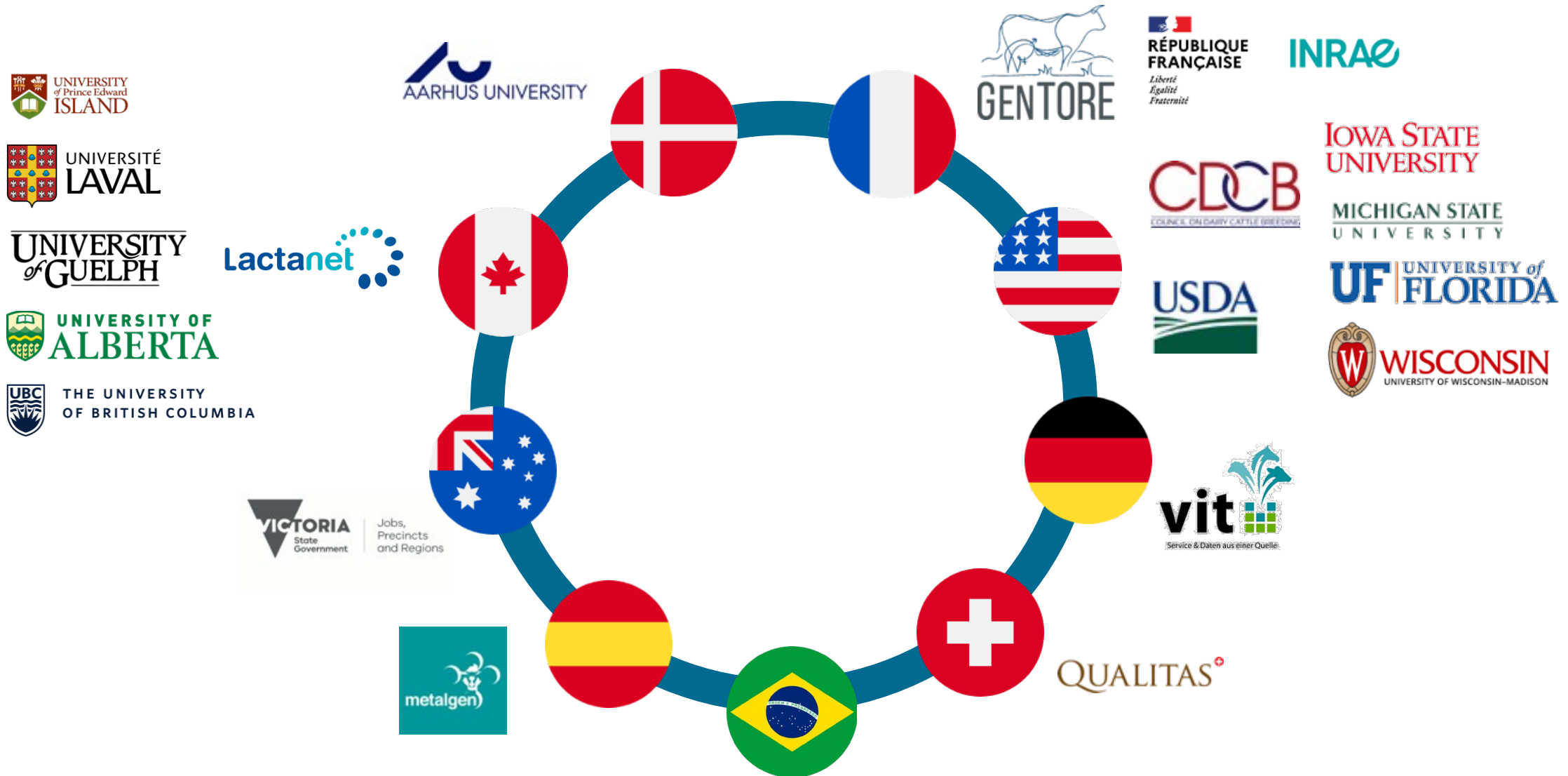
- International database for Feed Efficiency and Methane Emissions
- Genomic Evaluations for Feed Efficiency launched by Lactanet 2021
- Genomic Evaluations for Methane April 2023
- Resiliency Index (novel fertility, health, and efficiency traits) expected 2024

UNDER REVIEW:

- **MAPAQ & Dairy Farmers of Quebec PLQ project ~\$3.17M**
 - Quebec CH₄ Sniffers, Nutrition R&D & trainees
- **BC Dairy project \$136K**
- **Dairy Cluster project \$980K**
- **DFO Cash support \$160K + \$10K IK**
- **Genome Canada ICT ~\$16M**



National and International Research Partnerships



1. 'Closer-to-biology' fertility



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Dr. Ronaldo Cerri

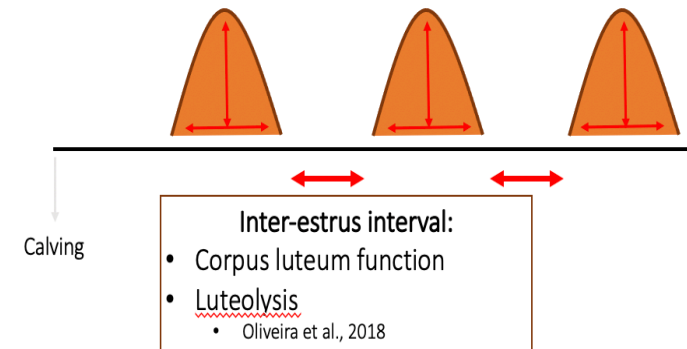
1. "Closer-to-Biology" Fertility



- Standardized phenotypes based on automated sensors
- Physiological factors affecting estrous expression and embryo survival
- Genomic markers of estrus expression and fertility
- Size and Position Score (SPS)
- Transmission Ratio Distortion

1st estrus event (timing):

- Already known for high heritability
- Transition – Health related
 - Madeira et al., 2018,



Intensity of estrus:

- Ovulation failure and timing
 - Burnett et al., 2018
 - Madeira et al., 2019
- Progesterone concentrations
 - Denis-Robichaud et al., 2018
 - Madeira et al., 2018, 2019
- Uterine receptivity
 - Davoodi et al., 2016
 - Cooke et al., 2019

Madureira et al., 2022, 2021
Martin et al., 2021, 2022

2. Enhanced disease resistance

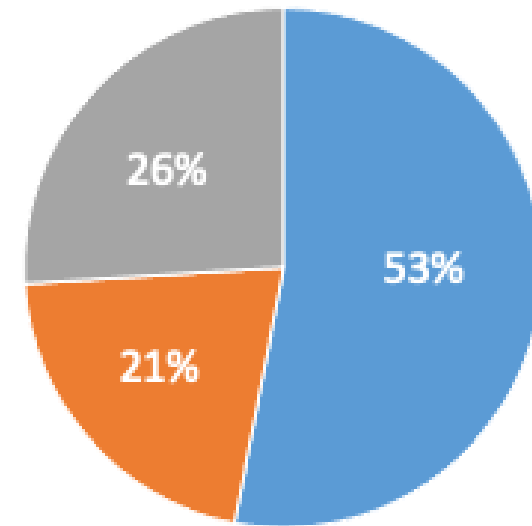


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Causes of pre-weaning mortality



■ Diarrhea ■ Respiratory problems ■ Other

Lynch et al., 2021; 2022; 2023

Bongers et al., 2023

Fong et al., 2023

- Fertility disorders in routine genomic analyses (Lactanet, 2020)
- Develop methods for routine phenotyping of
 - Calf health (Colin Lynch)
 - Leukosis (Renee Bongers)
 - Feed efficiency of calves (Kyle Hoeksema)
 - Johne's disease (Aisha Fong / Chrissy Rochus)
 - Effects of homozygosity (Makanjuola / Obari)

3. Feed efficiency and methane emissions

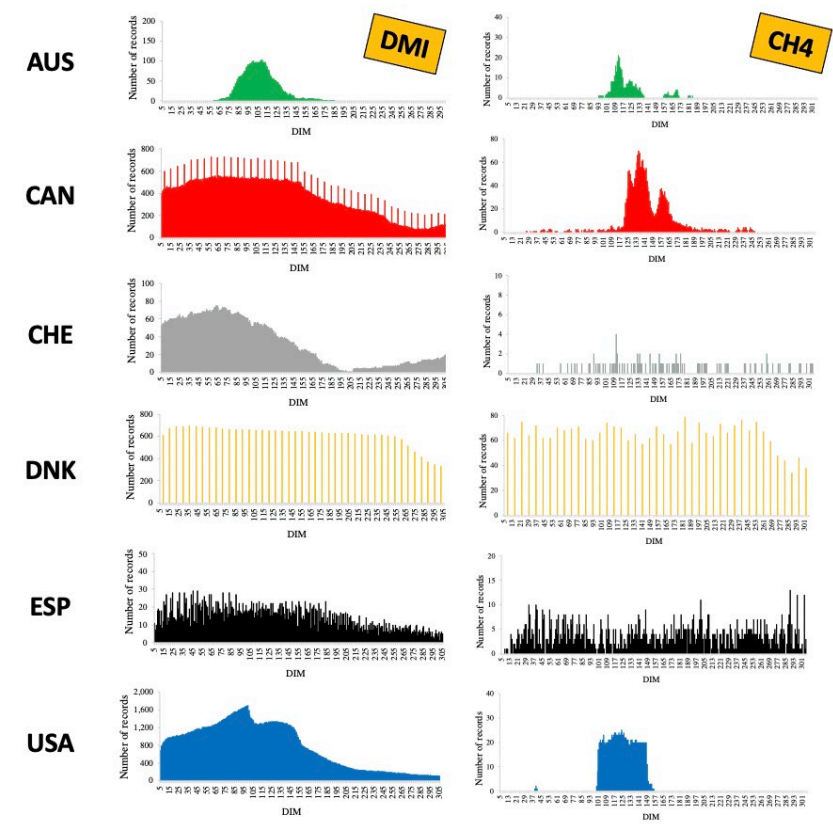
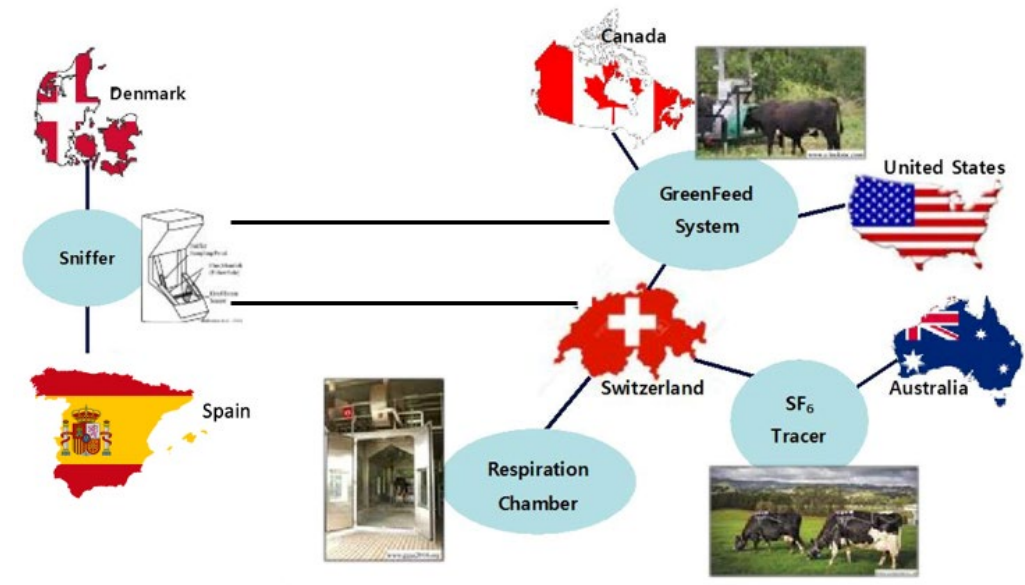
Lactanet
Dr. Gerrit Kistemaker



UNIVERSITY of GUELPH
Dr. Christine Baes



3. Feed Efficiency & Methane Reduction



- Enlarging the reference population for
- **Feed efficiency** (17,000 animals)
 - **Methane emissions** (7,800 animals)

4. Genomic and environmental relationships



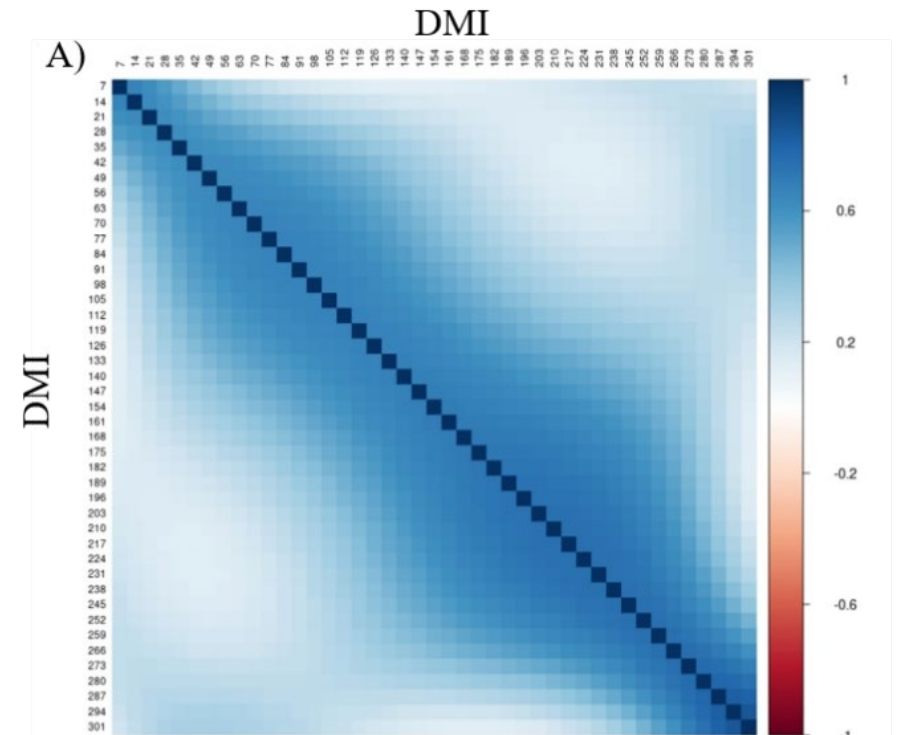
Dr. Flavio Schenkel

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Genetic parameters and prediction of EBVs of resilience traits

- Multi-trait analysis to **identify genomic regions** with pleiotropic effects on resilience traits
- Genomic predictions for resilience indicator traits using **copy number variants**
- Investigate the effects of **heat stress** on important traits



Houlahan et al., submitted

4. Genomic and environmental relationships

Fertility

- **Alcantara et al., 2022.** Machine learning classification of hormonal synchronization protocols for Canadian Holsteins cows. *JDS* (in review)
- **Martin et al. 2022.** Reproductive tract size and position score: Estimation of genetic parameters for a novel fertility trait in dairy cows. *JDS* (in review).
- **Oliveira et al. 2022.** Genome-Wide Association Analyses Reveals Copy Number Variant Regions Associated with Fertility and Disease Traits in Canadian Holstein Cattle, *PAG 2022*

Heat Stress

- **Campos et al. 2022.** Using publicly available weather station data to investigate the effects of heat stress on milk production traits in Canadian Holstein cattle, *CJAS*
<https://doi.org/10.1139/cjas-2021-0088>
- **Rockett et al. 2022.** Estimation of genetic parameters and prediction for heat tolerance in Holsteins using test-day production records and NASA POWER weather data. *JDS* (in review).

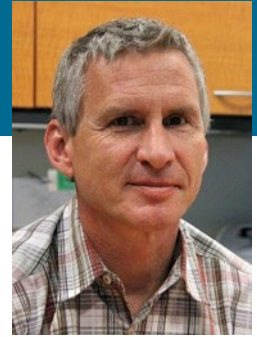
Calf Health

- **Bongers et al. 2022.** Incorporation of enhanced disease resistance into genetic evaluations. 2022 Interbull meeting.
- **Bongers et al. 2022.** Genetic analysis of leukosis milk ELISA test records in Holstein cows. 2022 ADSA meeting.
- **Lynch et al. 2022.** A Canadian genetic evaluation for calf health: preliminary analysis. 2022 WCGALP.

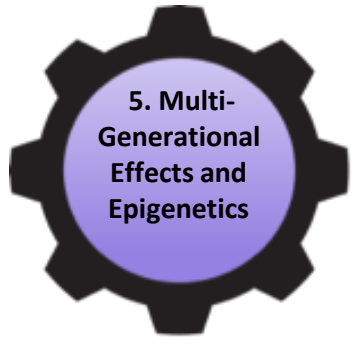
Feed Efficiency / Methane

- **Shadpour et al., 2022.** Predicting dry matter intake in Canadian Holstein dairy cattle using milk MIR and other predictors via ANN. *Accepted, JDS*
- **Shadpour et al., 2022.** Predicting methane emission in Canadian Holstein dairy cattle using milk MIR and other predictors via ANN. *Accepted, JDS*
- **Lopes et al. 2022.** Estimates of genetic parameters for environmental efficiency traits for first lactation Holsteins. *2022 ICAR & WCGALP meetings*
- **Houlihan et al. 2022.** The dynamic behavior of genomic predictions for feed efficiency over lactation. *2022 ICAR & WCGALP meetings*

5. Multi-generational effects and epigenetics



Dr. Marc-André Sirard



- Quantify effect of **early environment** (i.e., cow's production) on **resilience of daughters**
- Survey for **epigenetic signature** on precisely phenotyped animals
 - Whole Genome Bisulfite Sequence
 - 24 healthy (good)
 - 24 with mastitis, poor performance, infertility, lameness (bad)



Methylation level of C_16_22592086 (on the y-axis) and C_8_101832167 (on the x-axis)

6. Data management



Dr. Paul Stothard



Management of project database

- Whole-genome **sequence data** analysis for variants, genotypes, functional annotations
 - SnakeMake pipeline to call SNPs?
- **Genome browser integration** of GWAS findings, epigenetic signatures, & annotated sequence variation

SnakeMake pipeline developed for calling SNPs from methylation sequencing – Coverage now 50x

Sample	Average coverage	Bases with >10X coverage (Mbp)	Filtered SNPs
2258	25.73	2.49	3,813,056
2260	13.29	1.17	1,202,679
2261	29.19	2.52	4,109,577
2262	15	1.22	1,345,862
2267	5.88	0.37	274,853
2268	21.46	2.42	3,216,416
8761	14.5	1.72	1,634,613

Next steps: compare WGBS-called SNPs to SNPs from conventional WGS to gauge performance and utility of this approach.

7. GE3Ls: sustainability and social acceptance

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Dr. Getu Hailu

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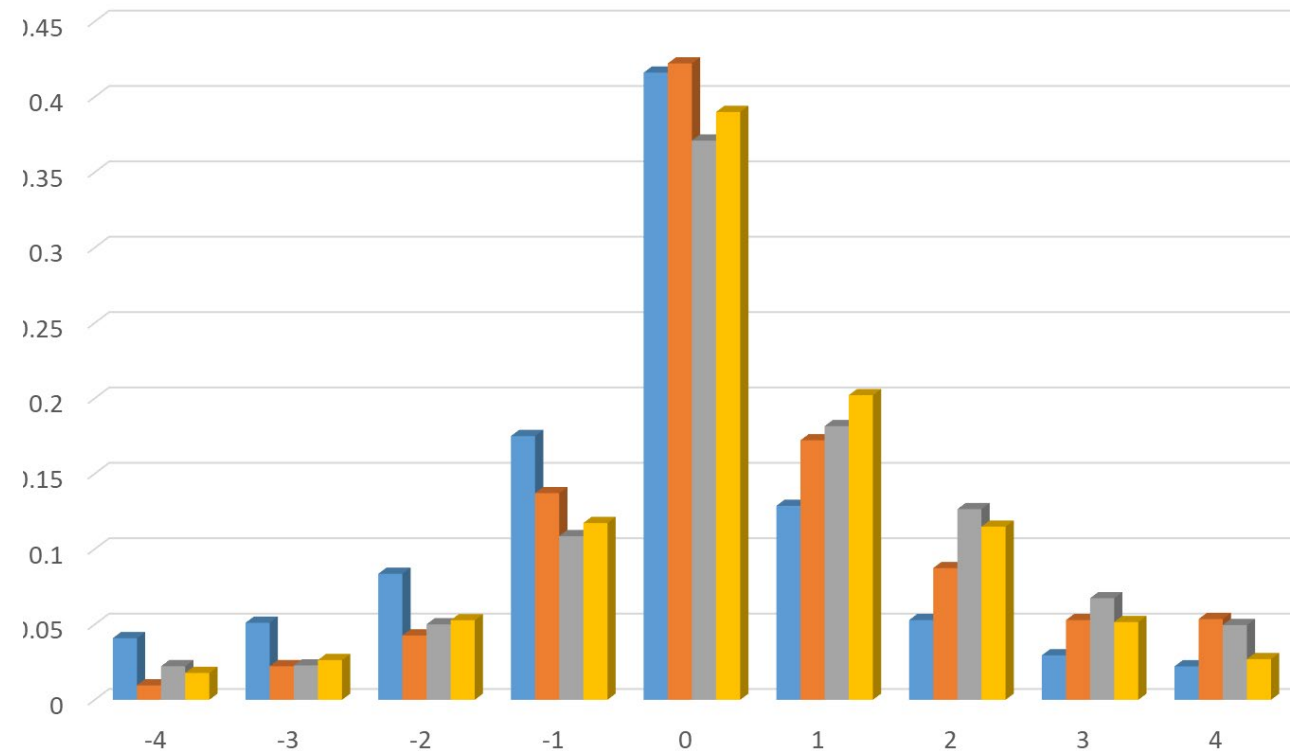
Dr. Ellen Goddard

7. GE³LS: Optimizing traits to maximize sustainability and societal acceptance

- Farm level decisions about tradeoffs between traits
- Farm/Market level outcomes from selection of resilience traits
- Public acceptance of dairy under different breeding strategies

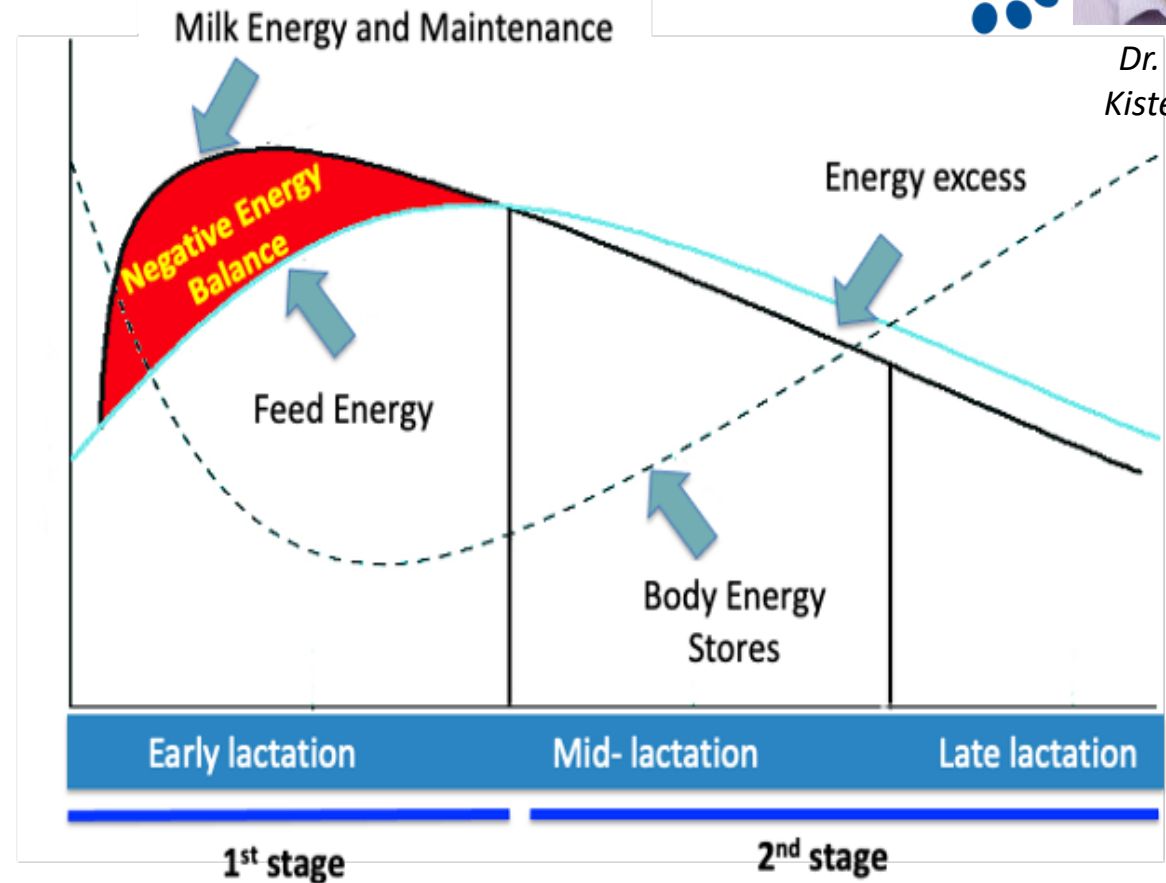
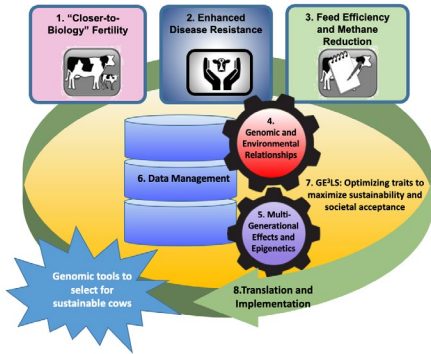
Goddard et al., 2022

Public Perceptions: Benefits of Genomic Selection minus Risks of Genomic Selection



■ Beef Feed Efficiency 2012 ■ Pork Disease Resilience 2012 ■ Dairy Feed Efficiency 2016 ■ Dairy Feed Efficiency Disease Resilience 2016

8. Translation and Implementation



Dr. Gerrit Kistemaker

- Implement fertility disorder evaluations (done 2020)
- Feed efficiency evaluations (2022)
- Methane evaluations (2023)
- Develop resiliency index

Overall aim is to select for cows that use less feed at the same level of production and body size **after peak of lactation**

Final remarks: Environmental Efficiency

- RDGP data base is sizeable and growing
 - Canada, US, Denmark, Switzerland, Germany, Spain, Australia
 - About 3,200 cows for methane emissions
- CH₄ sniffers installed soon in multiple commercial farms
- CH₄ emissions predicted **accurately** (~0.85) using milk MIR
 - Evaluations for **CH₄ emissions** ready to be launched in April 2022
- New Genome Canada proposal underway
 - GHG mitigation roadmap using genetic and nutrition strategies
 - Reduce GHG emissions by 54% (6.72 Mt CO₂-eq)

Climate-Smart Agriculture and Food Systems – Interdisciplinary Challenge Team: Leveraging Genomics to Achieve Dairy Net-Zero

Baes, Miglior, Gervais and Stothard



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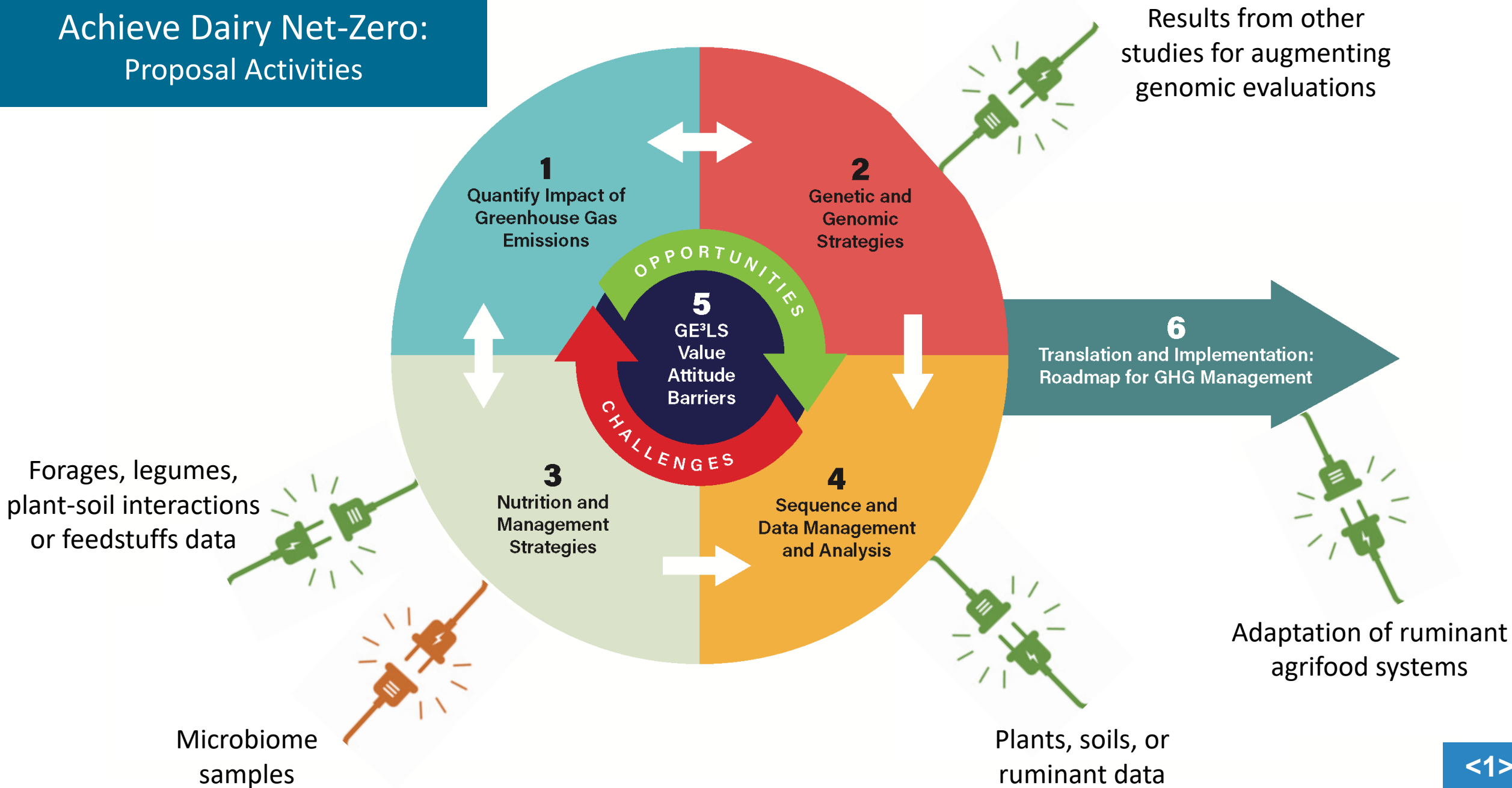


**Caeli
Richardson**

Plant and
Animal
Geneticist
AbacusBio



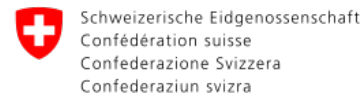
Leveraging Genomics to Achieve Dairy Net-Zero: Proposal Activities



Leveraging Genomics to Achieve Dairy Net-Zero: Deliverables

- A roadmap (overall goal) and embedded toolbox for GHG reduction
- Quantify impact and uncertainty surrounding GHG mitigation strategies
- Understanding biological architecture to deliver novel mitigation tools for methane emissions
- Reports on:
 - broad context of emissions reductions
 - public and wider stakeholder attitudes to such reductions
 - producer engagement to help ensure uptake of mitigation approaches
- Accurate and robust method for estimating individual animal and herd-level GHG emissions for use in national policy and GHG inventories

Acknowledgements



Thanks to a fantastic team!

www.resilientdairy.ca/

