



An international initiative to decrease the environmental footprint of dairy cattle using genomics

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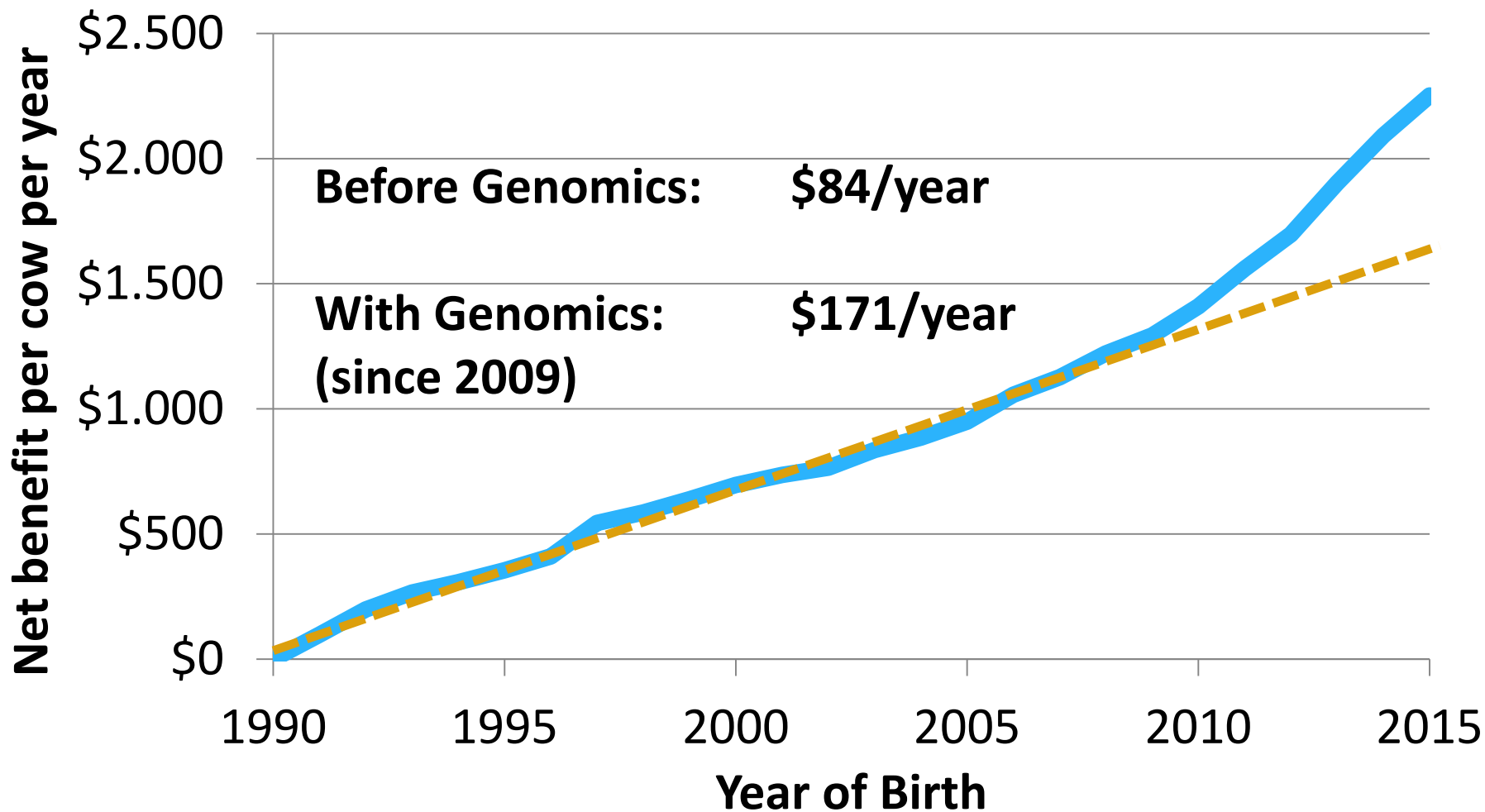
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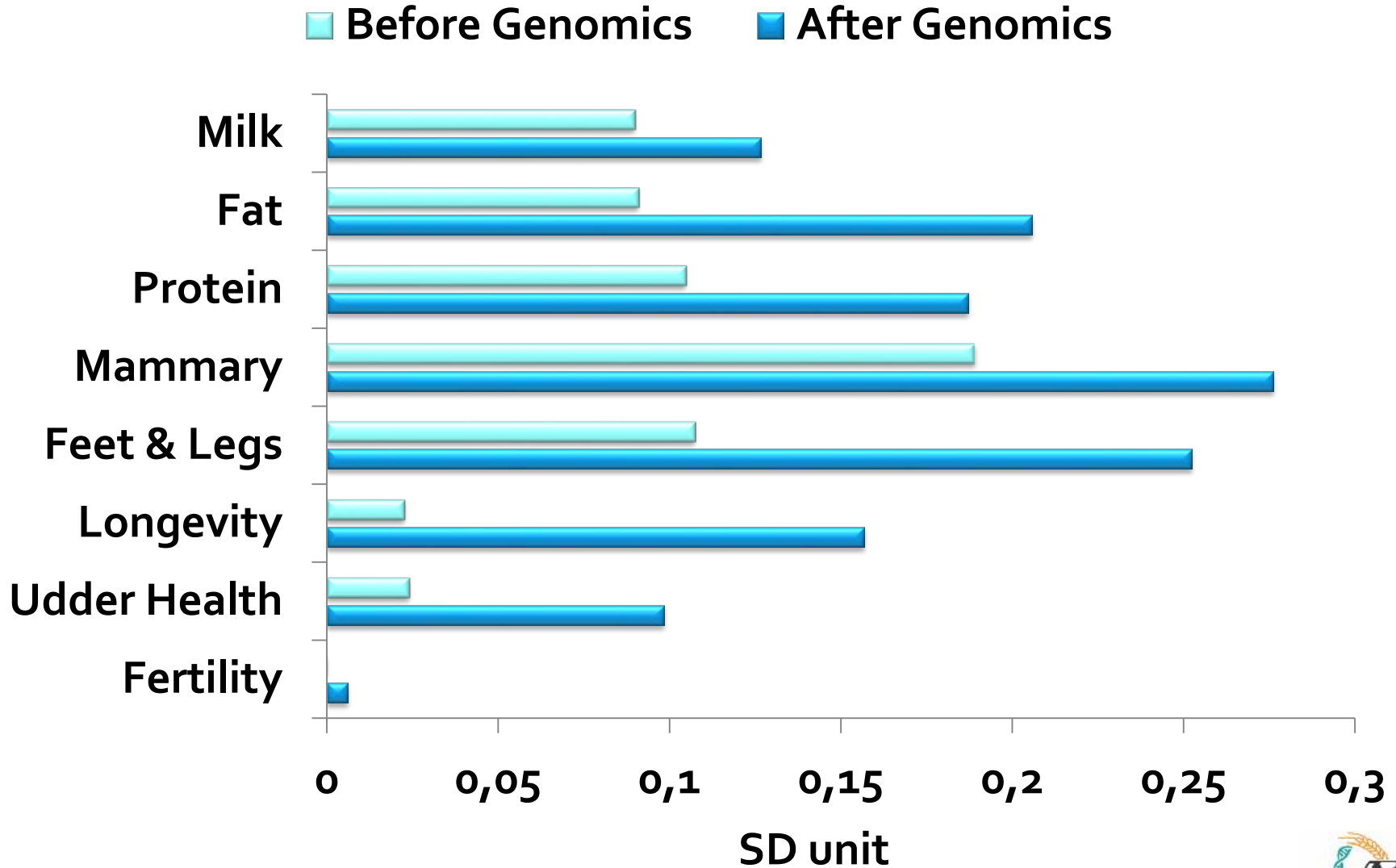
Impact of Genomics on Sire Evaluations

- Need for large reference populations to achieve accurate reliabilities of young stock parent averages
 - 0.70 Reliability of GPA vs. 0.35-0.40 reliability of PA
- Competing AI organizations that share genotypes
 - **InterContinental Genomics Exchange Consortium**
Canada, US, Italy, UK, Switzerland and Japan
 - **EuroGenomics**
France, Germany, Netherlands, Denmark, Sweden, Finland, Poland and Spain
- Over 32,000 Holstein reference bulls per consortium

Strong impact due to immediate industry uptake



Annual Genetic Progress – Major Traits



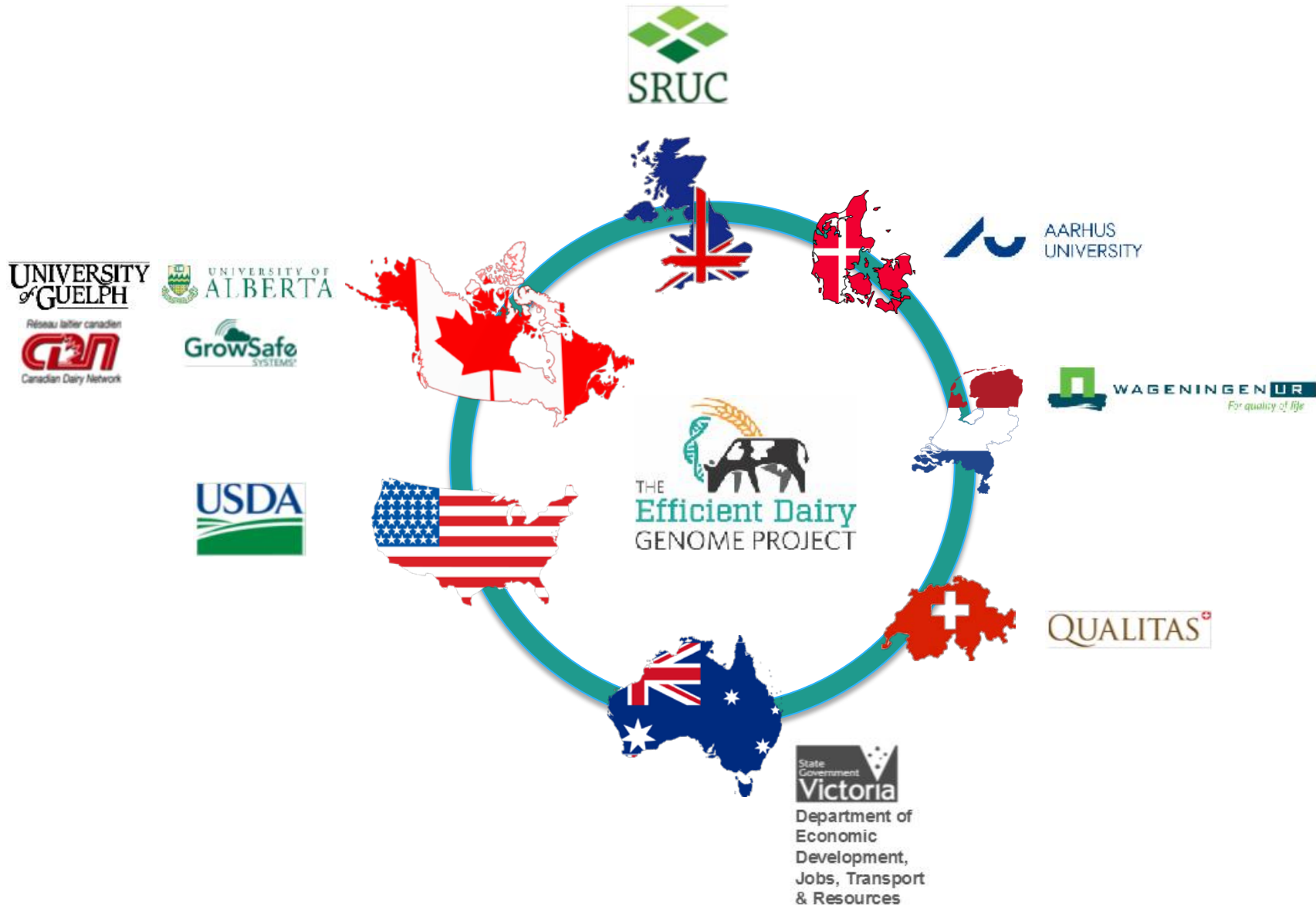
Integrated Solution for Greener Dairy

- Genomics provides an opportunity to select for higher feed efficiency and decreased methane emissions
- Will lessen the environmental impact of dairying
 - Reduction of emissions and feed waste
 - Less feeds (less land) used for dairying
- Compelling opportunity to make dairy greener worldwide, through strong collaboration
- Whereas at same time benefiting dairy producers
 - Reduction of feed costs
 - Potential credits for cutback in methane emissions

Efficient Dairy Genome Project

- Genome Canada Large Scale Applied Research Project
- Cash and in-kind funding from international and national partners to total CAD \$10.3 million
 - Genome Canada, Genome Alberta, Ontario Genomics
 - ALMA, Ontario MRI, CDN
- Start date Oct 2015 – End date Sep 2019
 - Led by Filippo Miglior (Guelph) & Paul Stothard (Alberta)
- **Overall objective to improve feed efficiency (FE) and reduce methane emissions (ME) in dairy cattle using genomics**

A Fully Integrated Partnership






Experimental Design



- Cow reference population for SNP calculation for FE and ME
- Continuous sharing of SNP solutions among partners
- Using MIR to predict FE & ME
- Genomic evaluations in each country using cow FE & ME
 - Adding relevant SNP
 - Multiple-trait evaluation adding MIR predicted FE and ME

Secure data base routinely updated
and accessible by all partners

Activities	Year 1	Year 2	Year 3	Year 4
GE³LS Ethical, environmental, economic, legal and social aspects of genomics	Social benefits and costs of selection for FE and reduced ME			
	Societal acceptance and value of use of this new technology			
Data Consolidation	Feed Efficiency			8,000-10,000 cows 
	Methane Emissions			3,500 cows 
	Milk MIR data			1,000,000 cows 
Genomics	Imputation & GWAS			
	Sequence	Variant Discovery	RNA-Seq & Functional Studies	
Implementation	Genetic Evaluation & Software Development			
	Profit Index		GEBV Delivery & Translation	

Tangible Deliverables to End Users

- **Weekly release of GEBV for FE and ME to producers and the AI industry integrated by**
 - *Newly-identified mutations or markers*
 - *New developed FE and ME predictions from MIR spectra*
 - *World's first database to routinely validate genomic predictions and continuous exchange among partners*
 - *Measurement of the farm level benefits from adopting the use of the GEBV for novel traits*
 - *Societal cost benefit analysis results from incorporating the new traits into national breeding programs*

Governance

Genome Canada, Genome Alberta, Ontario Genomics

Research Oversight Committee

*Alison van Eenennaam, USA
Chris Reynolds, GBR
Christian Maltecca, USA
Dirk-Jan de Koning, SWE
Hermann Swalve, DEU (Chair)
Rémy Lambert, CAN*

**Project Leader
and Co-Leader**

Industry Advisory
*DairyGen Council of CDN
Joint Industry--Academia*

Research Management Committee

*International Partners & Activity Leaders
Student Representative, Project Manager ex-officio,
Genome Alberta and Ontario Genomics ex officio*

Financial Committee

Project Manager and Leaders

Highly Qualified Personnel Committee

*Project Manager and Leaders
Student Rep & Post-Doc Rep*

Team



Mary De Pauw
Project Manager



Dan Hailemariam
Post-Doc, Canada



Allison Fleming
Post-Doc, Canada



Luiz Brito
Post-Doc, Brazil



Pauline Martin
Post-Doc, France



Adrien Butty
PhD, Switzerland



Stephanie Lam
PhD, Canada



Dave Seymour
PhD, Canada



Caeli Richardson
MSc, Canada



Angela Wilson
MSc, Canada

Plans for Guelph Research Herd

Calves (n = 180/year)



Calf Measurements

Duration: 0 – 60 d

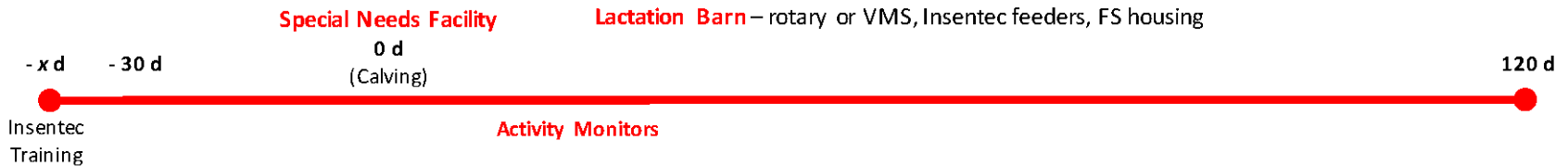
Genotyped at birth.

BW recorded at birth, 1 month and 2 months.

Milk replacer intake, starter intake (normally recorded)

Primiparous Heifers:

Feed Efficiency (n = 180/year)



Methane Emissions (n = 75 – 100/year)



Methane Emission Measurements

Duration: 20 – 30 d GreenFeed System

Daily (M–F)

CH4 emissions measured every 4 hours.

Measurements taken 1 hr later each day .

Other

Rumination collars

Feed Efficiency Measurements

Duration: -30 d to +150 d

Daily

Daily yield
Daily feed activity
Daily feed intake

BCS (every milking, using 3D cameras)

Weekly

DHI (alternating am/pm recording):
Milk yield
Milk fat, protein
Lactose

BHBA
MUN
SCC
Feed Samples

Two Weeks

Body weight
Visual body conformation

Ongoing Measurements

Individual Cows

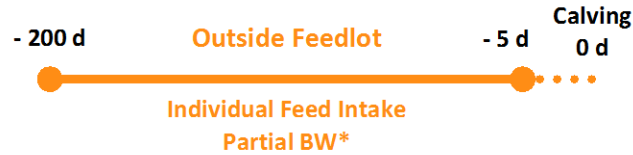
Health events
Reproduction events

Other

Environmental data (RH, temp, inside and outside CO2)

Plans for Partner Farm – 450 cows

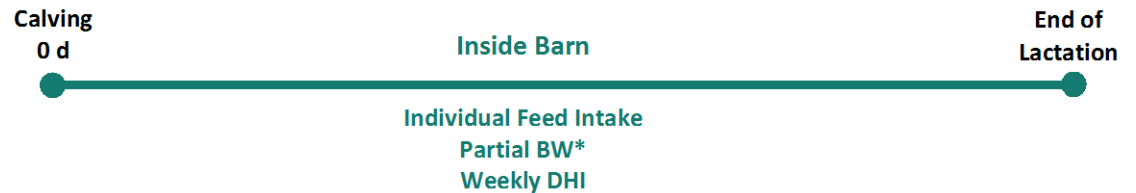
Pregnant Heifers (n = 150/year)



First Parity Cows (n = 150/year)



High-Producing, Later Parity Cows (n ≈ 150/year)



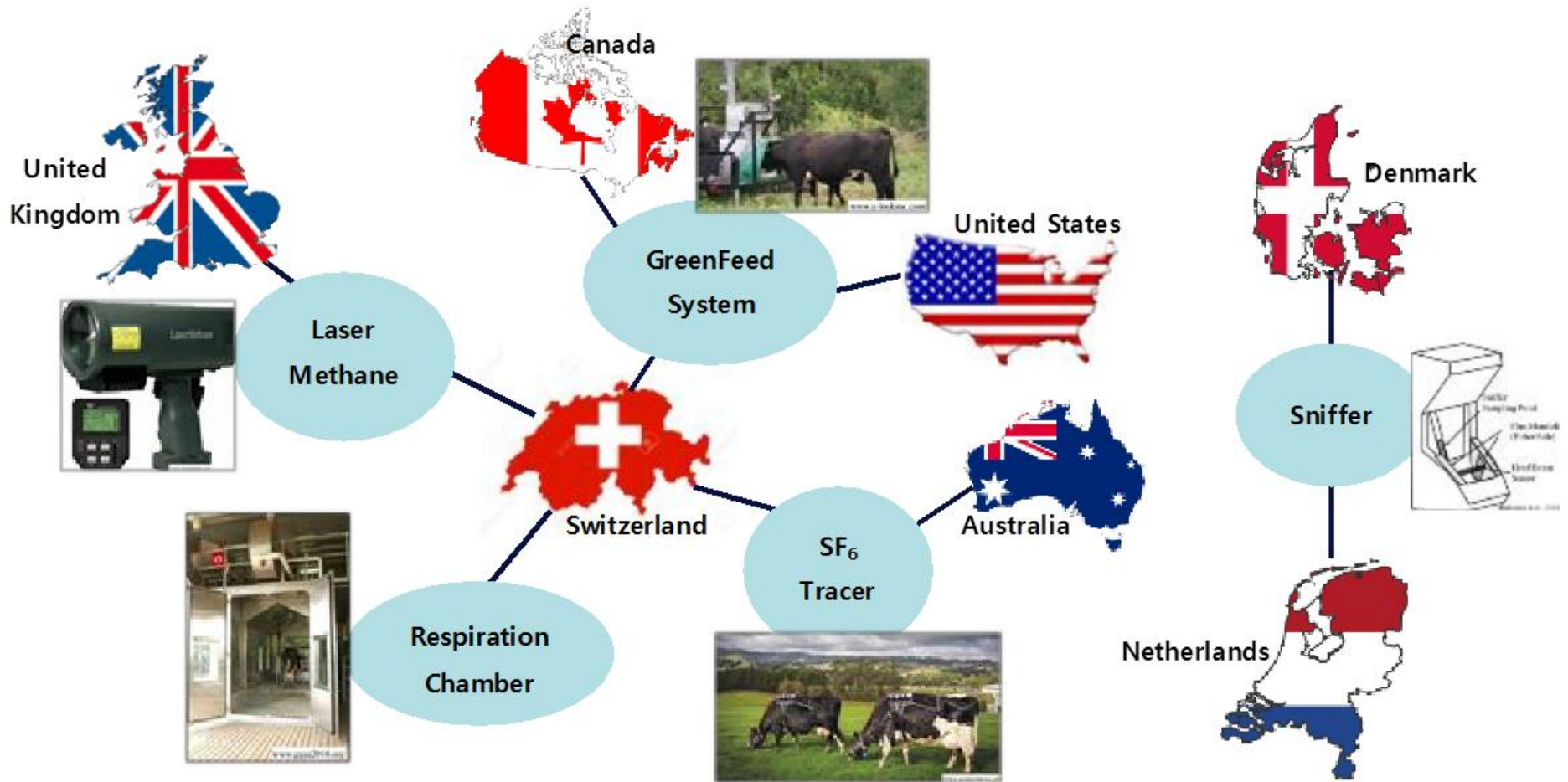
Possible Additions:

- Rumination Collars
- GreenFeed system

All animals genotyped with 50 K

*Body weight of cow measured from two front feet and is recorded multiple times per day

Methane Emissions



Progress

- Data routinely collected from each partner
- Survey to partners for data base development
- Survey to ICAR members
 - Trait definition and standardization
 - Selection goals definition
- Identification of bulls to sequence
- Setup of economic studies
- Strengthening of collaboration with gDMI 2
- Complementary projects recently approved
 - Microbiome and system biology (Dr. Angela Canovas)
 - Breeding Strategies (Dr. Christine Baes)

Participating Organizations



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