



Genetic correlation between methane emissions and nitrogen use efficiency proxies in Walloon dairy cows

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Greenhouse gases emissions (GHG) in dairy cattle

- ✓ Dairy cattle account for 20% of the global livestock sector's GHG



- ✓ GHG emissions from dairy cattle are in the form of **CH₄** (44%), **N₂O** (29%), and **CO₂** (27%)

Reducing CH₄ emissions from dairy cows

✓ CH₄ from cows represents **70%** of total CH₄ emissions of agricultural sector

✓ **Reducing CH₄ emissions from dairy cows is important at two aspects:**

1. Environmental concerns
Climate change



2. Economical perspective
6% to 10% of gross energy is lost as CH₄

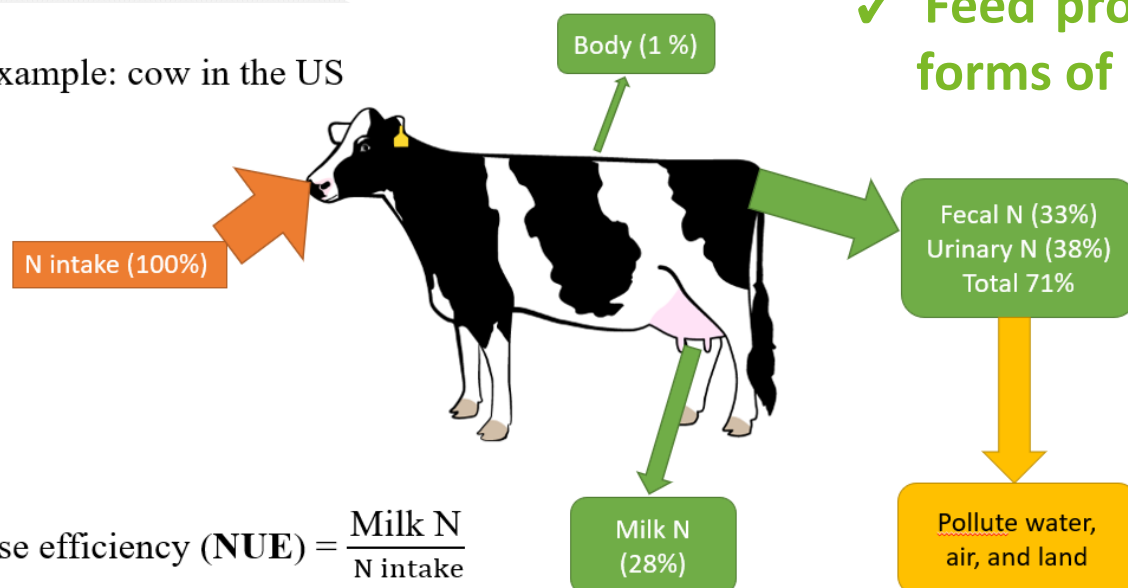


Nitrogen use efficiency in dairy cattle

✓ Protein is the most expensive part of a diet

✓ Feed protein is mainly metabolized into different forms of nitrogen in the cow's body

A example: cow in the US



$$\text{N use efficiency (NUE)} = \frac{\text{Milk N}}{\text{N intake}}$$

✓ Nitrogen use efficiency of dairy cows ranges from **20%** to **44%**

(Adapted from Spek et al., 2013)

Improving nitrogen use efficiency of dairy cows

✓ N_2O is **310** times more powerful greenhouse gas than CO_2

✓ Improving nitrogen use efficiency of dairy cows is important at two aspects:

1. Environmental concerns

Climate change



2. Economical perspective

> 50% intake nitrogen is not used



Regulation of CH₄ emissions and nitrogen use efficiency of dairy cows

✓ Improvement of feeding strategies

✓ Genetic selection

✓ cost-effective

✓ permanent strategy

✓ For genetic selection, **traits** need

✓ to be well defined

✓ to be recordable (milk MIR predicted)

✓ to be genetically variable and heritable

Research Objectives

The aims of this study were to estimate

- ✓ **Genetic parameters** of two MIR-based “proxies” for CH₄ features
- ✓ **Genetic correlations** of the CH₄ features with nitrogen use efficiency proxies

Material and Methods

Methane emissions traits:

- ✓ Milk MIR spectra were used to predict CH₄ emissions (**PME**, g/d)
Vanlierde et al. (2021)
- ✓ Log-transformed CH₄ intensity (**LMI**) was defined as log of PMI
CH₄ intensity (PMI) defined as PME per kg of milk yield (kg/d)

Material and Methods

- ✓ Data and model for **CH₄** traits:
 - ✓ 1,529,282 test-day records in first parity
 - ✓ Random regression test-day model



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Research

Estimation of genetic parameters for predicted nitrogen use efficiency and losses in early lactation of Holstein cows

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- ✓ EBV and its reliability for **Predicted nitrogen use efficiency (PNUE) and losses (PNL)** 

Material and Methods

- ✓ Approximate genetic correlation by EBV of selected bulls

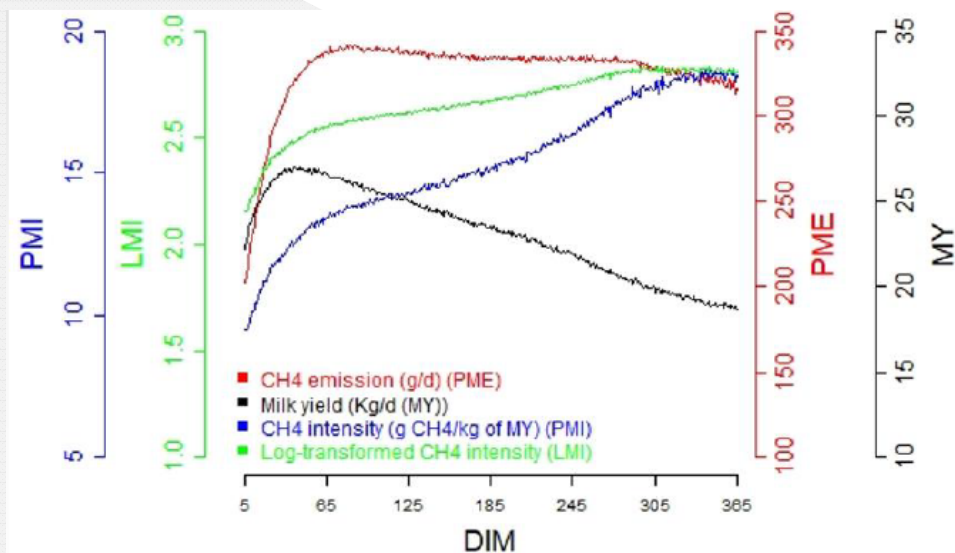
$$\hat{\rho}_{aa'} = \frac{\sqrt{(\sum b_i)(\sum b'_i)}}{\sum b_i b'_i} r(\text{EBV}, \text{EBV}')$$

b_i is the reliability of EBV for trait a (Blanchard et al., 1983)

- ✓ Conditions for selecting bulls

Reliability of EBV for used traits > 0.30

Results 1 — Data description



Traits	Mean	SD	Peak value	Peak time
MY	23.57	6.13	27.02	42
PME	326.82	67.65	341.62	80
PMI	14.97	5.70	18.65	344
LMI	2.64	0.35	2.83	344

Results 2 — Heritability

Range of daily heritability for

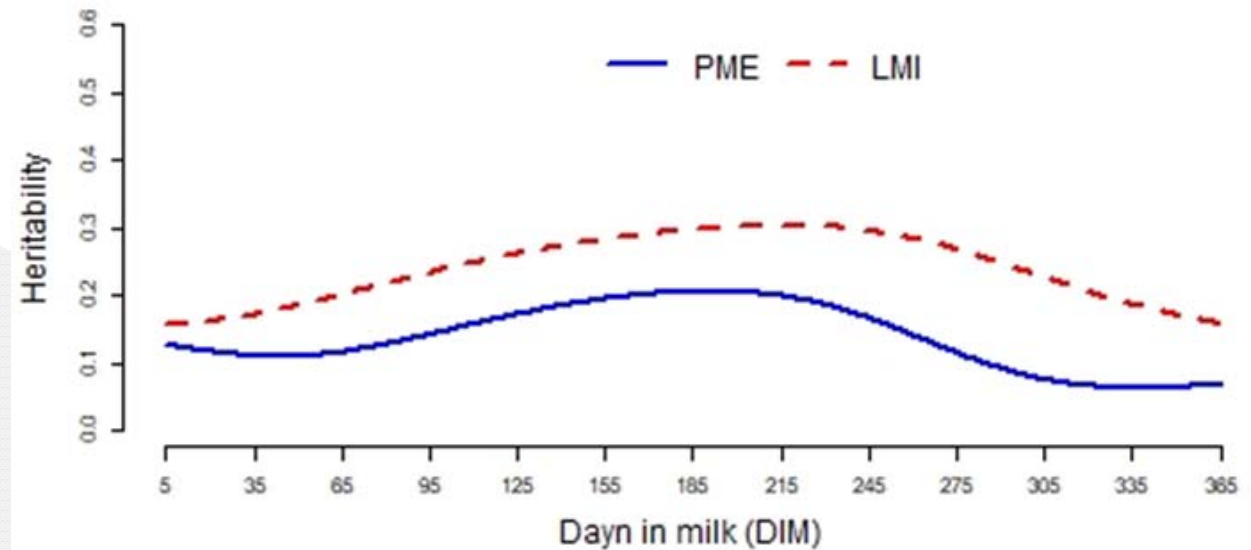
✓PME → 0.07 — 0.21

✓LMI → 0.16 — 0.30

Mean (SD) daily heritability for

✓PME → 0.14 (0.05)

✓LMI → 0.24 (0.05)



Results 3 – Selected Bulls

A total of **420 bulls** were used to estimate approximated genetic correlation (AGC)



137 (USA)



111 (NLD)



52 (CAN)

...

	PME	LMI	PNUE	PNL
Mean reliability (SD)	0.84 (0.16)	0.84 (0.16)	0.48 (0.15)	0.49 (0.15)

Results 4 – Approximated genetic correlation

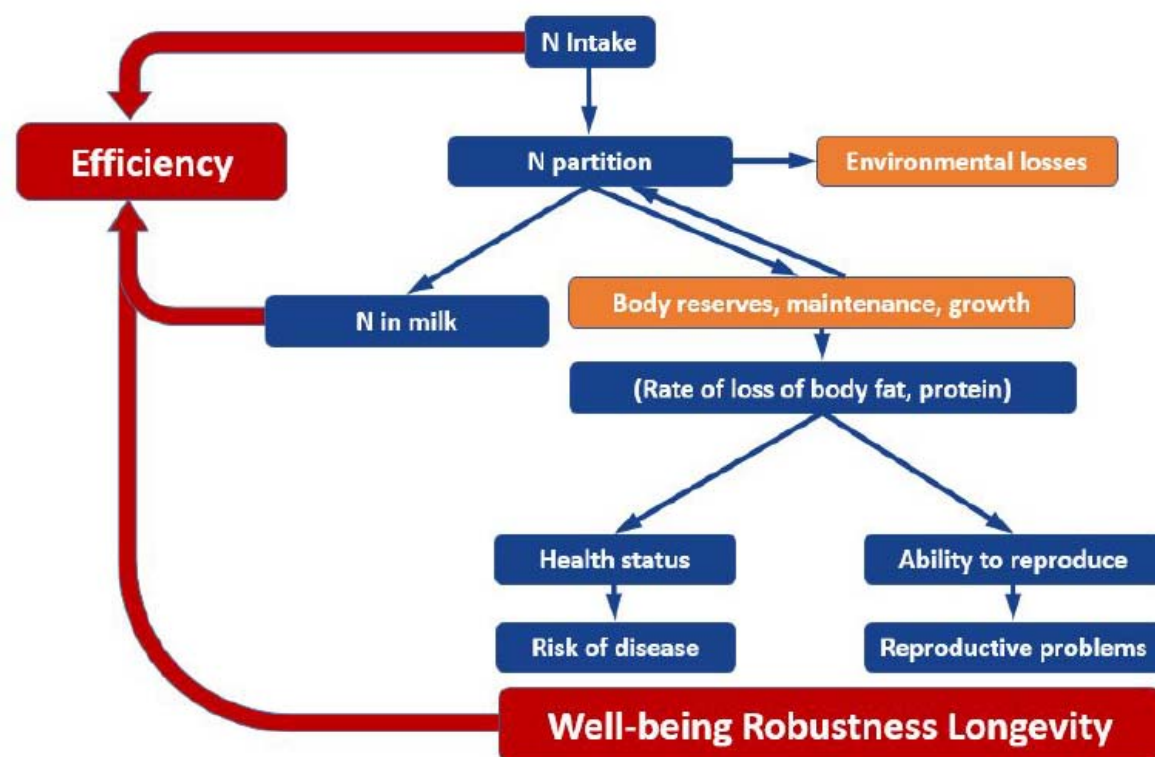
Approximated genetic correlation between four traits in the first parity (SE, n=420)

	Predicted nitrogen use efficiency	Predicted nitorgen losses
PME	-0.33 (0.07)	0.43 (0.08)
LMI	-0.60 (0.07)	0.32 (0.08)

Conclusions

- ✓ Two MIR-based “proxies” for CH_4 features were validated to be genetically variable and heritable → Genetic selection
- ✓ Genetic selection → CH_4 emissions ↓ + N loss ↓ + Nitrogen use efficiency ↑ in dairy cows → simultaneously selection
- ✓ Before introduction of these traits in a selection index, further studies needed:
 - ✓ Genetic correlation between these proxies (index traits) and direct trait (breeding objective trait)
 - ✓ Evaluate effect of inclusion in selection index on all traits already present in the current one

Future perspectives



Holistic breeding goals

- ✓ CH₄
- ✓ Nitrogen use efficiency
- ✓ Resilience
- ✓ Longevity
- ✓

(Chen et al., 2021)

Thank you for your attention!

Questions?

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