SmartCow

an integrated infrastructure for increased research capability and innovation in the European cattle sector

Proxies to predict

feed efficiency and its determinants in cattle

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INRAE - CRAW

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Global objective

- To reduce constraints on experimental animals (3Rs rules) in research infrastructures (RIs)
- To strengthen cattle phenotyping capacity for large scale studies
- To find alternative methods to estimate key phenotypes measured with gold standard method (GSM)



➤ To evaluate proxies* of feed efficiency and its determinants in dairy & beef cattle across diets and individuals



^{*}defined as indicators measurable in easily accessible matrices, and easier to implement than the GSM

Strategy

•To build a large and representative database with individual observations [phenotypes + proxies] for beef & dairy cattle

- To share data (samples) among SmartCow collaborators
 - > SRUC, UREAD, AU, CRA-W, INRAE, FBN Leibniz, IRTA, WUR
 - > TNA applicants (including private companies)
 - > Agroscope, Luke





Which proxies for which phenotypes?

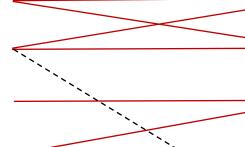




PHENOTYPES

PROXIES

- Feed efficiency
- □ N use efficiency & N excretion □
- □ Total tract OM digestibility
- □ CH₄ emission
- ☐ Rumen parameters (VFA, ammonia, pH)





- **Urea-N**
- **Near-infrared spectra (NIRS)**
- Mid-infrared spectra (MIRS)
- **Blood** metabolites
- Volatile metabolome













Proxies to predict Feed efficiency in beef cattle

Feed Conversion
Efficiency (FCE)

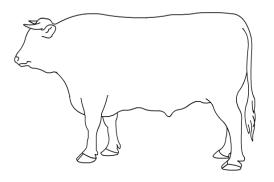
Residual Feed Intake (RFI)

INDIVIDITAL INTAKE



BODY WEIGHT GAIN





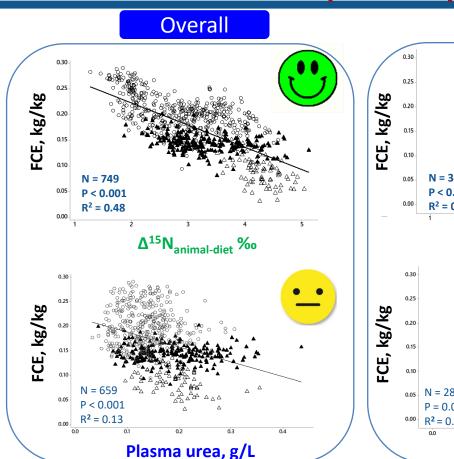
15N abundance Urea-N



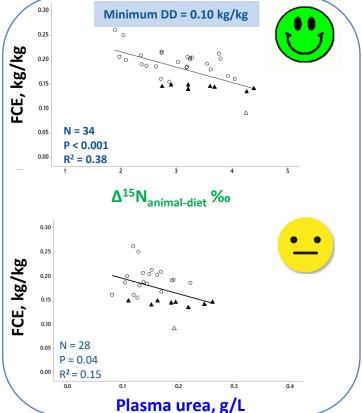


Proxies to predict Feed efficiency (FCE, RFI) in beef cattle Better models with Δ¹⁵N compared to plasma urea





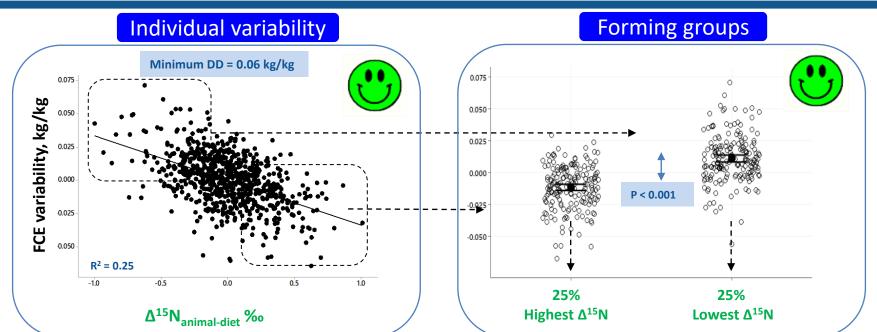
Dietary means





Proxies to predict Feed Efficiency (FCE, RFI) in beef cattle Δ¹5N allows to discriminate individuals





The minimum detectable difference in FCE between 2 individuals from ¹⁵N values was almost 0.06 kg/kg => too high for assisting genetic selection Mean difference in FCE, between 2 groups formed from ¹⁵N values, was significant and around 0.03 kg/kg => for precision feeding?

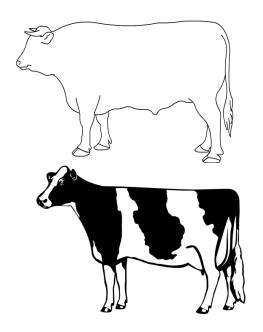


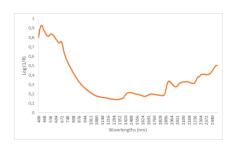
Proxies to predict Total tract digestibility in cattle

OM digestibility (OMD) = (ingested-excreted) ingested









Faeces NIRS



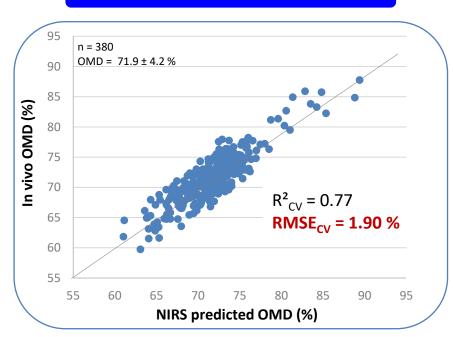


Proxies to predict Total tract digestibility in cattle

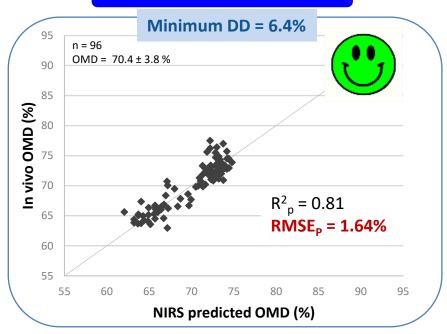




Smartcow calibration model



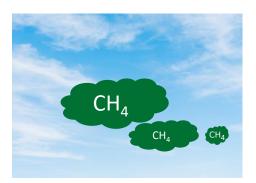
Validation dataset



- ➤ Minimum DD (6.4%) is close to that of GSM (5.2%)
- Good potential of fecal NIRS as proxy for OMD prediction in cattle
- Good alternative to GSM (stalls)

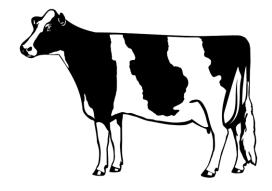


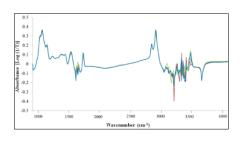
Proxies to predict daily CH₄ emissions in dairy cows











Milk MIRS



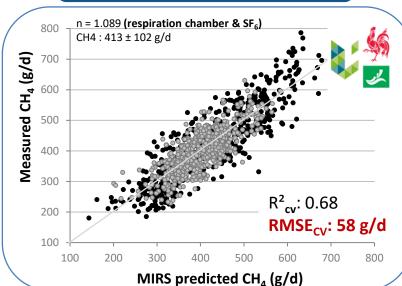


Proxies to predict daily CH₄ emissions in dairy cows Model based on milk MIRS : validation and next steps



Existing predictive model

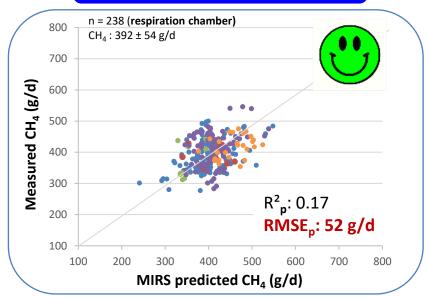
Vanlierde et al, 2021 (JSFA)



- Good potential of milk MIRS as proxy for CH₄
- ➤ High throughput approach allows CH₄ phenotype to be incorporated in dairy cow breeding programs

Validation dataset

FBN and WUR data



Next steps

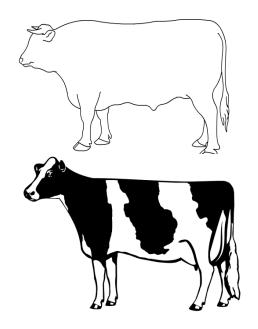
- To include Ref data not yet represented
- Not possible to merge GF data with other Ref datasets (RC, SF₆) → noise in the model
- A specific model based on GF values is on progress

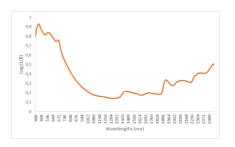
Proxies to predict daily CH₄ emissions in cattle



CH₄ EMISSIONS







Faeces NIRS

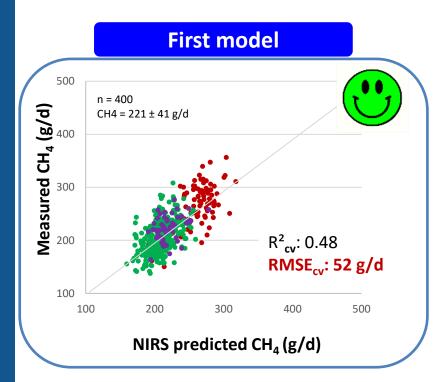




Proxies to predict daily CH₄ emissions in cattle

Faecal NIRS: an innovative proxy





- > A good example of complementary dataset
 - BE -> Belgian Blue; grass, grass silage
 - FR -> Charolaise; grass, grass silage
 - CH -> Simmental Angus Limousine; corn silage
- > Promising proxy especially for an indirect proxy
- ➤ **Useful for non-lactating cattle :** beef, calf heifers, dry dairy cows
- ➤ More Ref data not yet represented are needed
 - Other breeds, diets, physiological status, etc.
 - Standardized protocols for sampling and dataset management



Impact

- **For RI:** improvement of the cattle phenotyping capacity while implementing 3Rs principles & research recommendations
- -> instructions for using proxies according to common and standardized protocols / open access guidelines





For stakeholders:

- > Academic : production of knowledge (publications) and innovating concepts to test in pilot studies
- Industry : development of phenotyping tools for feed and breeding industry



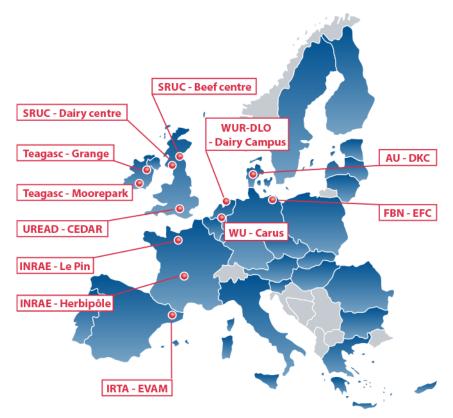
Future prospects

- To enlarge diversity of the reference databases to update models following research recommendations
- To continue investigating new proxies and their combinaison to improve predictions
- Adoption and implementation of these research proxies by the different actors of the livestock production

Key of success : collaboration network / common guidelines for measurements and data recording

Thank you for your attention





First-class Cattle Research Infrastructures (RIs) across Europe:

- 11 major RIs distributed in 7 EU countries
- 12 locations, which include 18 installations
- 2500 dairy and 1000 beef cows
- **Networking of RIs** to inventorize resources, harmonize procedures, and share data
- **Joint research activities** to improve experimental methods and phenotyping capability
- **Interaction with stakeholders** to stay in line with industry needs and improve dissemination

http://www.smartcow.eu/stakeholders/

TRAINING PROGRAM

For Scientists, Technicians, Stakeholders, PhD students

- Face-to-face training courses
- Free web-conferences
- One-day study tours in 4 different countries

http://www.smartcow.eu/resources/training/

TRANSNATIONAL ACCESS CALLS

Offers external users (academic and industry) free access to SmartCow RIs

- 30 projects during the 4 years of SmartCow
- Access to around 10,000 cow-weeks

http://www.smartcow.eu/calls/



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Questions



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