

Latest developments of the use of midinfrared spectral data in the north-western region of Europe: from OptiMIR to GplusE

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Breeding for Efficiency – 2017 EDGP Symposium

My Presentation Today

- Overview of activities and projects of our group and affiliated groups since 2005
 - Earlier developments (e.g. RobustMilk, GreenhouseMilk)
 - Mostly focus on recent (e.g. OptiMIR and GplusE)
- Focus on impact for industry and producers
- Before I start
 - Many thanks to numerous collaborators, without them this work would not be possible, especially the Walloon Agricultural Research Center (CRA-W)



Mid-Infrared as Technology

Also often called FTIR

• Mid-infrared (MIR) spectrometry is not new

➔ Has been in use for a long time

- But far away from the sight of the general industry and producers
- Use for all major milk components (except SCC)
 - Fat %
 - Protein %
 - Urea
 - Lactose

Major Milk Components (except SCC)



First Steps to Extend Impact of MIR

 Around 2005 first ideas immerged in the European industry

Let's try to predict novel traits

- Possible because hidden in the "black box" → rather simple process of combining spectral data to get predictions!
- Based on prediction (often called calibration) equations

Major Milk Components (except SCC)





Impulse from Local and EU Projects

- Best known: **RobustMilk** (www.robustmilk.eu)
 - "Innovative and Practical Breeding Tools for Improved Dairy Products from More Robust Dairy Cattle"
- Focusing on
 - Milk quality (fatty acids) and some health related traits (as lactoferrin and minerals)



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 - "Innovative and Practical Breeding Tools for Improved Dairy Products from More Robust Dairy Cattle"
- Focusing on
 - Milk quality (fatty acids) and some health related traits (as lactoferrin and minerals)
- Large impact as it showed
 - Basic validity of approach
 - Possibilities to used MIR for novel traits

But Milk Composition Ideal Tool for More



Metabolic Status Subclinical Changes



Also: $CH_4 \leftarrow Milk$ Composition



An Important Remark!

- Spectrometers MIR data potentially unstable
- As long as clearly defined compound

possible to correct post-calibration

- Using reference milk with known composition
- As done in routine for fat%, protein%,...
- However if the target becomes a status or a non-milk phenotype (as CH₄)

impossible to correct post-calibration

Important Issue Different Machines

For each CH₄ measurement Milk MIR spectra Spectra after log **Standardization** Machine X Absorbance Machine Y 100 400 500 Wavenumber (cm-1) J. Dairy Sci. 100:1-12 https://doi.org/10.3168/jds.2017-12720 © American Dairy Science Association®, 2017 Standardization of milk mid-infrared spectrometers for the transfer and use of multiple models C. Grelet,* J. A. Fernández Pierna,* P. Dardenne,* H. Soyeurt,† A. Vanlierde,* F. Colinet,† C. Bastin,‡ N. Gengler.⁺ V. Baeten.^{*} and F. Dehareng^{*1} *Valorization of Agricultural Products Department, Walloon Agricultural Research Center, 5030 Gembloux, Belgium

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From January 2011 to September 2015

www.optimir.eu



WITH THE SUPPORT OF







OptiMIR

- Important project
 →large scale DHI industry collaboration in NW Europe
- Created a forum of exchange of knowledge and know-how for industry and science
- For industry:
 - Acquisition of MIR spectral data
 - ➔ knowing their spectrometers
 - Storing MIR data (and not loosing it)
 - Doing MIR "standardization"
 - Applying MIR equations
 - Reporting results

Creation of the European Milk Recording (EMR) group



OptiMIR Research

- Several innovative aspects (as direct use of MIR data)
- Also facilitator for exchange and data consolidation
 - For some traits consolidated into the OptiMIR research

database (e.g., for Ketosis related biomarkers, like BHB, citrate, ...)



 For other traits facilitator of contacts, harmonization of MIR and reference data and exchange (e.g. as for methane)

➔ For CH₄ also very important GreenHouseMilk and Methagene

CH₄ - SF₆ equation

- Research supported by local and international projects (e.g., GreenHouseMilk)
- Prediction of daily CH₄ from milk MIR spectra
 - Based on SF₆ references
 - From Belgium and Ireland

Ν	SD	R ² c	R ² cv	SEC	SECV
532	129	0.74	0.70	66	70









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Evolution of CH_4 equations \leftarrow Chambers

- "Open initiative" New members
 - Starting collaboration with Australia, potentially Canada
 - Through many contacts and projects as Methagene and GplusE
 - MIR and CH₄ data to be added in calibration
 - Get MIR equation and updates
- Evolution from SF₆ to respiration chambers
 - Independent datasets
 - Gold standard reference technique
- Confront the statistics with the SF₆ version



Results

Measured CH₄ emissions in function of the predicted CH₄





Genotype plus Environment Integration for a more sustainable dairy production system

- European project focusing on biomarkers for sustainable dairy cattle breeding and management (see <u>www.gpluse.eu</u>)
 - Lead partner UCD, Dublin, Ireland
 + BE, DE, DK, NL, IT, UK + USA + China associated to industry (CRV, QLIP, AWE, EMR)
 - 1. Objective: more classical work on energy balance and RFI and DMI



MIR Prediction of Feed Related Phenotypes

	Predictor traits	n	#LV	R ² cv	RMSECV
Energy balance (Mcal/d)	MIR	1069	15	0.43	5.1
Energy balance (Mcal/d)	MIR, DIM, FPCM	1098	12	0.51	4.8
RFI (kg/d)	MIR	1115	14	0.46	2.9
RFI (kg/d)	MIR, DIM, FPCM	1097	14	0.53	2.7
DMI (kg/d)	MIR	1052	14	0.47	3.2
DMI (kg/d)	MIR, DIM, FPCM	1098	8	0.59	2.9
				Grelet et al.	, ICAR 2017

Only 131 cows -> need for broader collaboration



Genotype plus Environment Integration for a more sustainable dairy production system

- European project focusing on biomarkers for sustainable dairy cattle breeding and management (see <u>www.gpluse.eu</u>)
 - Objective shown here: making blood based biomarkers useful
 - Creating "health" clusters
 - Predicting membership of clusters from milk MIR data



Blood Based Biomarkers as Reference





Creating Health Clusters

Phenotypes of interest (Grelet et al., ICAR 2017)

Blood metabolites and hormones are usefull when used togheter

 \rightarrow Clusters:

 Combination of the blood metabolites and hormones into 3 groups of cow status

K-means nearest neighbour clustering K=3



Creating Health Clusters



Creating Health Clusters



Predicting Health Clusters from MIR Data



Only 354 cows 🗲 need for broader collaboration

Conclusion

• In the era of novel phenotypes....



collaboration is imperial

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Thank you for your attention!

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