# Capitalizing on mid-infrared to improve nutritional and environmental quality of milk

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# Introduction



- Changes of consumer's perception
  - Improvement of the nutritional quality of food
  - Limitation of the environmental impact of food production and consumption
- Milk quality can be improved:
  - Nutritional quality:

E.g., unsaturated fatty acids, calcium, lactoferrin

– Environmental quality:

E.g., milk production linked to urea, to methane

# Introduction



- Acquisition of phenotypes needed!
- Development of practical tools:
  - Cheap: to be used on a large scale
  - Robust: to adapt to different breeds, sampling methods and dates...
  - Fast: more and more cows/farm
  - Reliable
- Use of mid-infrared spectrometry (MIR) on milk
  ⇒ promising technology



# Part I: Usefulness of MIR spectrometry



- Ranged between 1,000 and 5,000 cm<sup>-1</sup>
- Used routinely by milk laboratories to quantify major milk components:
  - Fat, protein, lactose, urea...
- But: currently under-utilized technology







Milk samples

(milk payment, milk recording)

Every 2 or 3 days Bulk milk samples Managed by dairy companies





Milk samples

(milk payment, milk recording)

Regularly (mostly 4 or 6 weeks) Individual cows Managed by milk recording organizations









*Milk samples* (milk payment, milk recording)









**MIR** analysis

Milk samples (milk payment, milk recording)



Raw data = MIR spectra

#### **MIR spectrometry** 1000 (Foss, 2008) **MIR** analysis Milk samples (milk payment, milk recording) **Quantification:** 0.20 (a) FT-MIR spectrum of milk **Calibration equations** 1072 1045 0.15 145/ fat 2854 1649 0.10 1744 Absorbance 0.05 protein 0.00 lactose -0.05

...

Wave number, cm<sup>-1</sup>

2000

2500

1500

1000

3500

3000

Raw data = MIR spectra









*Milk samples* (milk payment, milk recording)



Traditional data flow (no MIR spectra stored)



# MIR calibration equations: Nutritional quality



- Milk fatty acid (FA) equations:
  - First equations developed in 2005
  - Improved through international collaborations:
    - Belgium, France, Germany, Ireland, UK, Luxembourg
  - Multiple breeds, countries and production systems

# Accuracy of fatty acids calibration equations



Calibration equations were developed from at least 1,600 milk samples

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# Accuracy of milk minerals calibration equations



Calibration equations were developed from at least 465 milk samples

# Accuracy of milk minerals calibration equations



 $R^2 \ge 0.80$  for Na and Ca  $\Rightarrow$  potential practical uses

- Ca: milk fever, osteoporosis

- Na: indicator of mastitis

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- Lactoferrin equations:
  - Cooperative effort of Belgium, Ireland and UK

### Lactoferrin



- Glycoprotein present naturally in milk
- Involved in the immune system
- Interests:
  - Potential indicator of mastitis
  - Help to maintain a good immune system in Humans
- R<sup>2</sup> of internal validation = 0.71

#### ⇒ MIR indicator of lactoferrin

## Lactoferrin



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R<sup>2</sup> < 0.80 ⇔ MIR indicator of lactoferrin Improves slightly detection of mastitis compared to just using somatic cell score

# MIR calibration equations: Environmental quality



#### Methane reference trait

- Measured by the SF6 method

- Indirect link with milk FA (predicted by MIR)
  ⇒ Direct prediction of methane by MIR ?
- If possible can be used for:
  - Inventory of methane emissions
  - Environmental labeling of food
  - Reducing methane produced by individual cows

#### Methane





#### Methane





#### Conclusions



• MIR spectrometry under-utilized in practice

 Potential to predict new traits with real economic and societal interests

• However, this is not always easy ...

### Not so easy...



- Developed MIR equations
  - Must be validated on used dairy population (even if the equation was built internationally)
  - Because of differences in breeds and production systems affecting prediction
- Adding specific samples needed!
  - Variability of calibration set **7**
  - Adaptation of equations to new population **7**
  - Therefore: general robustness of equations **7**

#### Not so easy...



- If spectral data was recorded, should be easy to implement new equations in milk laboratories?
- However
  - Specific spectrometers were used to develop given calibrations
  - To avoid any additional bias all the spectral data need to be standardized with those used in calibration

### Not so easy...



- Accuracy of the MIR prediction must be tested regularly based on reference samples
- Creation of reference samples needs
  - Reliable reference values (traits to predict from MIR), potential difficult to obtain (e.g., methane)
  - Conservation and distribution of fresh milk samples (needed to be analyzed by MIR)
- Many logistical challenges



# Part II: Capitalizing on MIR traits for dairy cattle breeding and management

# **MIR spectral databases**



- Creation of spectral databases related to milk recording needed
  - Already in Walloon Region of Belgium and in Luxembourg
- In August 2012, available spectral records:
  - 2,305,838 test-day records from Walloon Region of Belgium
  - 1,262,190 test-day records from Luxembourg
- This allows
  - Large scale studies of genetic and phenotypic variability

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⇒ Development of selection and management tools

# **Capitalizing for breeding**



Daily  $h^2$  for saturated FA = 0.59 Daily  $h^2$  for monounsaturated FA = 0.26

Soyeurt et al. (2012), EAAP

Bastin et al. (2012), EAAP

Daily h<sup>2</sup> for lactoferrin = 0.35 Previous estimates: 0.20 – 0.44 Daily  $h^2$  for calcium = 0.50 Daily  $h^2$  for sodium = 0.34 Daily  $h^2$  for magnesium = 0.52 Daily  $h^2$  for potassium = 0.48 Daily  $h^2$  for phosphorus = 0.55

Soyeurt et al. (2012), EAAP

Lactation	Records	Holstein cows	Daily h <sup>2</sup> CH <sub>4</sub> (g/day)	Daily h <sup>2</sup> CH <sub>4</sub> (g/L milk)
1	270,902	54,355	0.37	0.45
2	209,663	42,306	0.36	0.42
3	145,540	29,749	0.36	0.39

First results obtained by Purna Badhra Kandel (ITN Marie Curie, GreenHouseMilk project)

# **Capitalizing for breeding**







# **Capitalizing for breeding**



- Potentially useful also for other countries which have not access to these phenotypes...
- Different opportunities:
  - Collaboration in genomic prediction
    - Sharing of phenotypes and genotypes ⇒ up to joint evaluations
    - Creating and capitalizing on local prediction equations
  - Collaboration in genome wide association studies
    - Combining station and MIR predicted field data (e.g., bull EBVs)
    - Example for fatty acids done in **RobustMilk** project, more details given by Catherine Bastin (EAAP, 2012)

# **Capitalizing for management**



- Thanks to available large milk recording databases:
  - Study the phenotypic variability of MIR new traits
- Define **best practices**, potentially useful:
  - To mitigate the CH<sub>4</sub> emissions
  - To decrease the release of urea in milk
  - To improve the FA content of milk
- New step: direct use of MIR variability
  - OptiMIR project (www.optimir.eu)

#### Conclusion



• MIR interesting for breeding purposes

However ....

#### However ...



- Position of novel MIR traits in future breeding (and production) goals still uncertain
  - Need to discuss with all stakeholders to know what will be the future of dairy products and production
  - Better knowledge of relationships of these traits with other traits having economical and societal interests (e.g., production, health and fertility, longevity) needed
- Therefore:
  - Definition of new breeding programs and management
    objectives taking into account all these aspects needed

# Collaborations



 If you are interested in joining the consortium to improve the MIR equations:

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 If you are interested in sharing phenotypes and genotypes:

nicolas.gengler@ulg.ac.be

Thank you for your attention