

Standardisation of MIR milk spectra, a step to build an international spectral database



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PARTNERS

Investing in Opportunities



This project has received European Regional Development Funding through INTERREG IV B.



INTERREG IV B

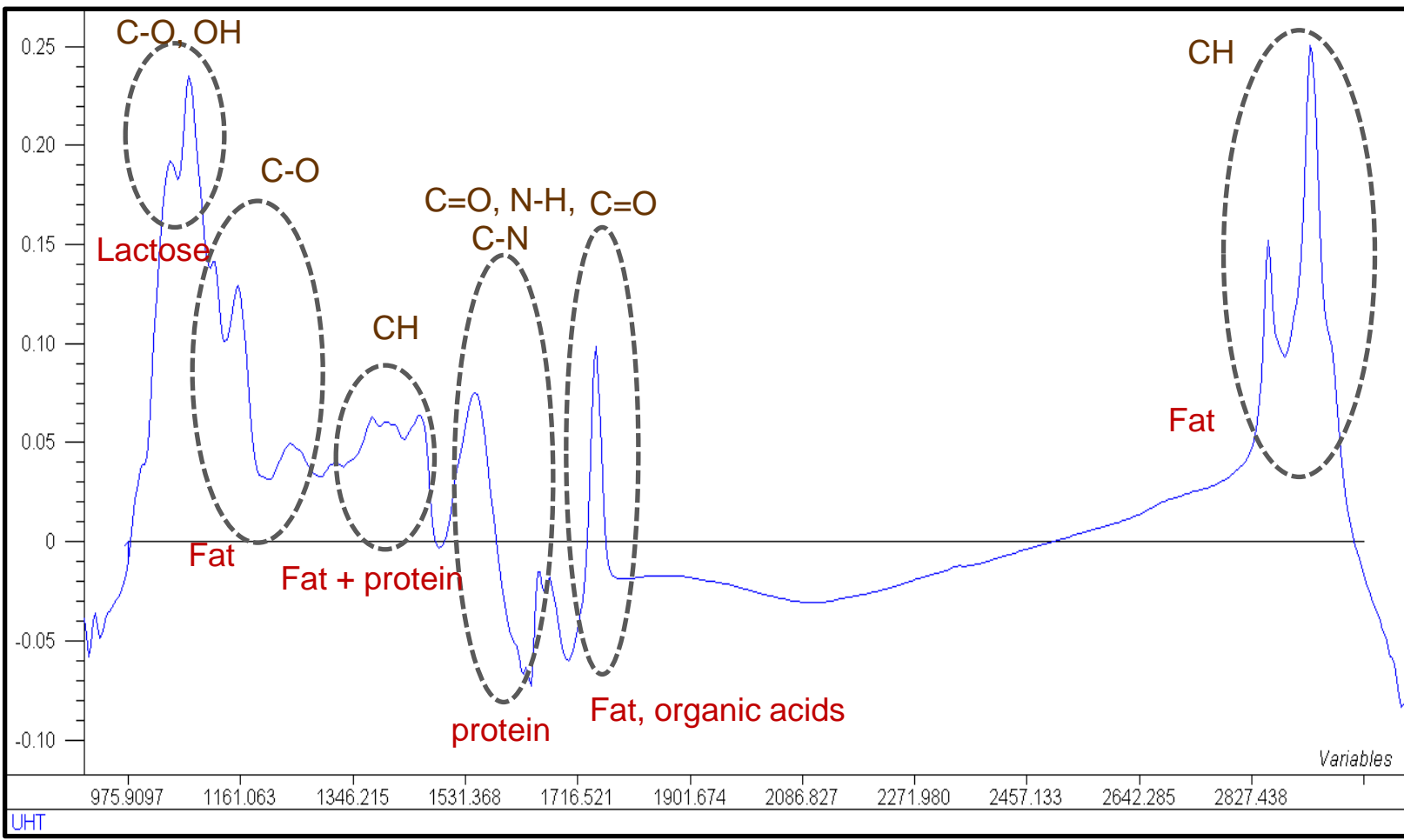
5 Research units + 1 Laboratoire

| Name | Country |
|--|---------|
| <i>Institut de l'Elevage</i> | FR |
| <i>Gembloux Agro-Bio Tech (Université de Liège)</i> | BE |
| <i>Centre wallon de Recherches agronomiques (Département Qualité Productions Animales)</i> | BE |
| <i>Comité du Lait</i> | BE |
| <i>Teagasc</i> | IR |
| <i>Scottish Agricultural College</i> | UK |

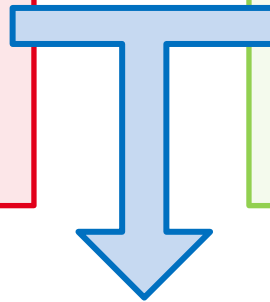
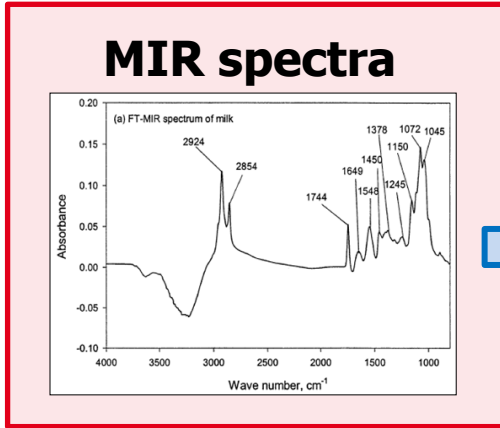


11 MILK CONTROL ORGANISATIONS

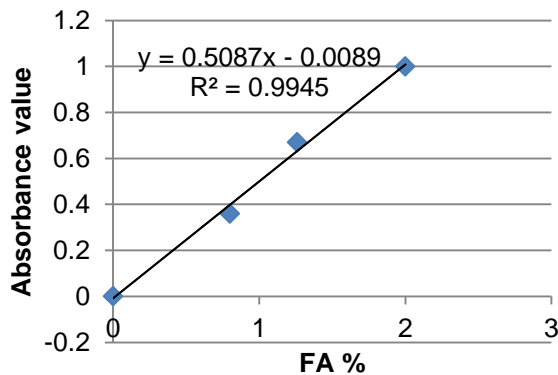
| Name | Country |
|---|---------|
| <i>Association Wallonne de l'Elevage</i> | BE |
| <i>Chambre régionale Agriculture Alsace</i> | FR |
| <i>ADECL62 (Pas-de-Calais)</i> | FR |
| <i>CLASEL (Sarthe & Mayenne)</i> | FR |
| <i>SCL du Doubs et du territoire de Belfort</i> | FR |
| <i>France Conseil Elevage</i> | FR |
| <i>LKV Baden-Württemberg</i> | DE |
| <i>LKV Nordrhein-Westfalen</i> | DE |
| <i>National Milk Recording</i> | UK |
| <i>Irish Cattle Breeding Federation</i> | IR |
| CONVIS | LU |



- **Position of the peaks** → **Qualitative analysis**
- **Intensity of the peaks** → **Quantitative analysis**



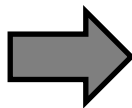
Saturated FA (g/100ml) = ax + b



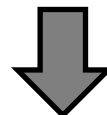
$$\begin{aligned}
 \text{Saturated FA (g/100ml)} = & 0.0985 + 4.6191 * X_1 + 1.1659 * X_2 + 1.4827 * X_3 \\
 & + 4.1684 * X_4 + 7.3294 * X_5 + 9.8991 * X_6 + 11.183 * X_7 + 8.0711 * X_8 \\
 & + 2.1599 * X_9 - 0.4619 * X_{10} - 1.7876 * X_{11} - 2.5708 * X_{12} - 2.8941 * X_{13} - \\
 & 2.9217 * X_{14} - 2.7392 * X_{15} - 2.2543 * X_{16} - 1.2677 * X_{17} + 0.0676 * X_{18} \\
 & + 1.0762 * X_{19} + 1.3228 * X_{20} + 1.0241 * X_{21} + 0.536 * X_{22} + 0.0177 * X_{23} - \\
 & 0.5265 * X_{24} - 1.1445 * X_{25} - 1.8178 * X_{26} + 2.212 * X_{27} - 2.0766 * X_{28} + \\
 & 8.3083 * X_{29} - 3.703 * X_{30} + 1.1999 * X_{31} + 0.5698 * X_{32} - 0.1674 * X_{33} \\
 & + 0.246 * X_{34} + 0.666 * X_{35} + 1.2938 * X_{36} + 2.0946 * X_{37} - 0.0689 * X_{38} - \\
 & 1.4774 * X_{39} - 1.7984 * X_{40} - 2.0553 * X_{41} - 2.9338 * X_{42} - 4.644 * X_{43} - 6.764 * X_{44} \\
 & - 8.1475 * X_{45} - 5.6904 * X_{46} + 2.6657 * X_{47} + 10.9883 * X_{48} + 14.4346 * X_{49} \\
 & + 13.8878 * X_{50} + 10.2135 * X_{51} + 4.8464 * X_{52} - 1.2081 * X_{53} - 7.4854 * X_{54} - \\
 & 11.6799 * X_{55} - 12.6849 * X_{56} - 10.7724 * X_{57} - 4.8936 * X_{58} + 0.4425 * X_{59} \\
 & + 3.583 * X_{60} + \dots + 0.9636 * X_{310} + 6.4566 * X_{311}
 \end{aligned}$$

Classical use of MIR spectra :

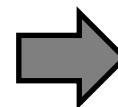
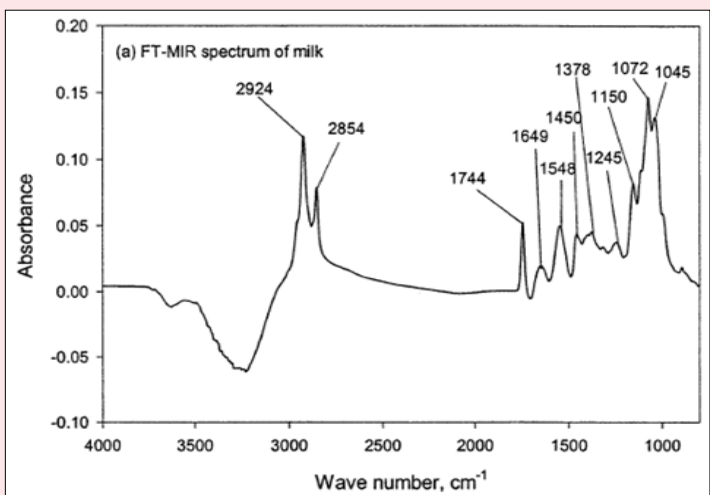
Milk control



MIR



MIR spectra of each cow



Milk composition

Fat

Proteins

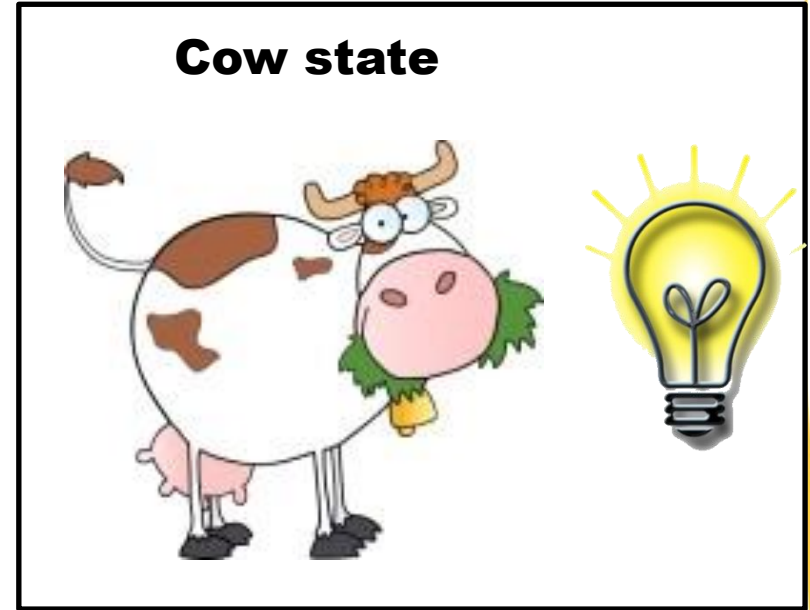
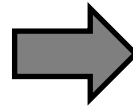
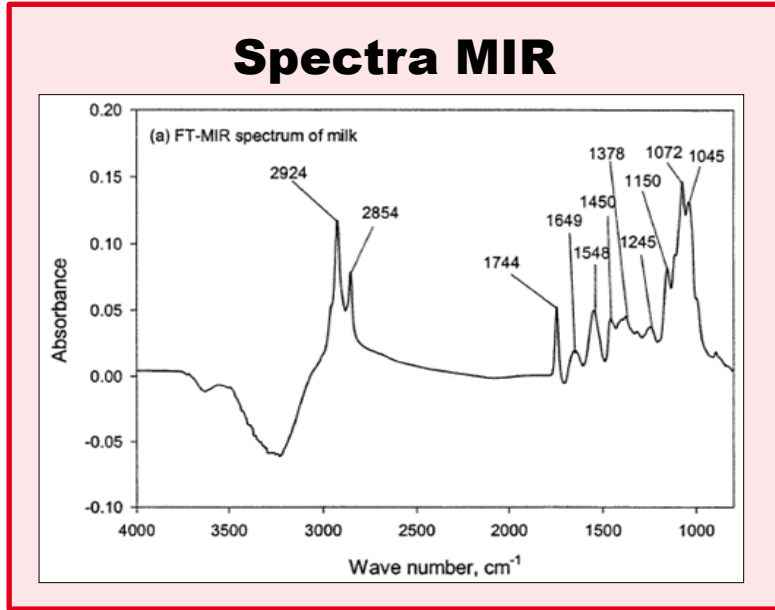
Urea

FA

...

**Fast
Cheap**

Innovative view of OptiMIR:



Prediction tools fast, cheap, via milk control organisations

Informations on :

- fertility**
- feeding**
- health**
- environmental impact**

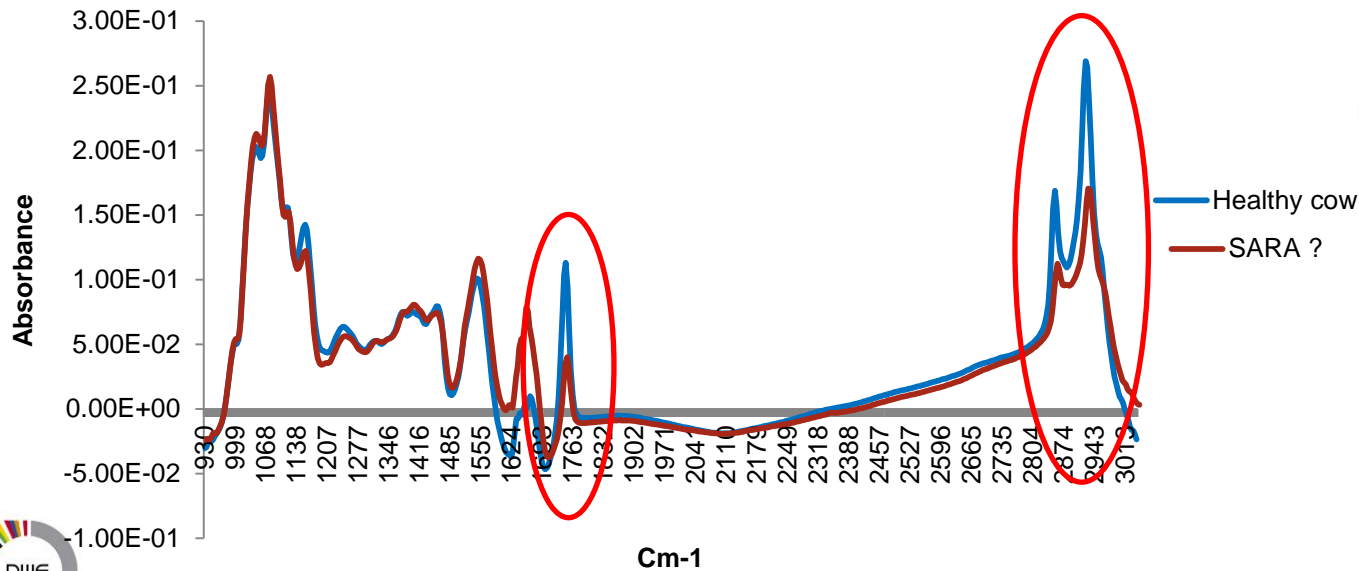
Concrete example: SARA



Impact of SARA on fat rate, fatty acid profile... (Sauvant et al., 1999; Colman et al., 2010)



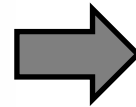
Hypothetic spectra of healthy cow and cow suffering of SARA



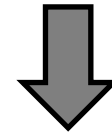
STEP N°1

Combine data from each MRO :

Spectra MIR + Fertility, health, feeding and environmental data

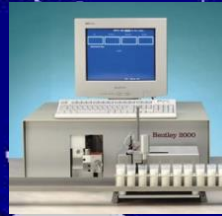
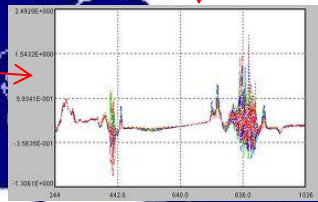


**Transnational
database**

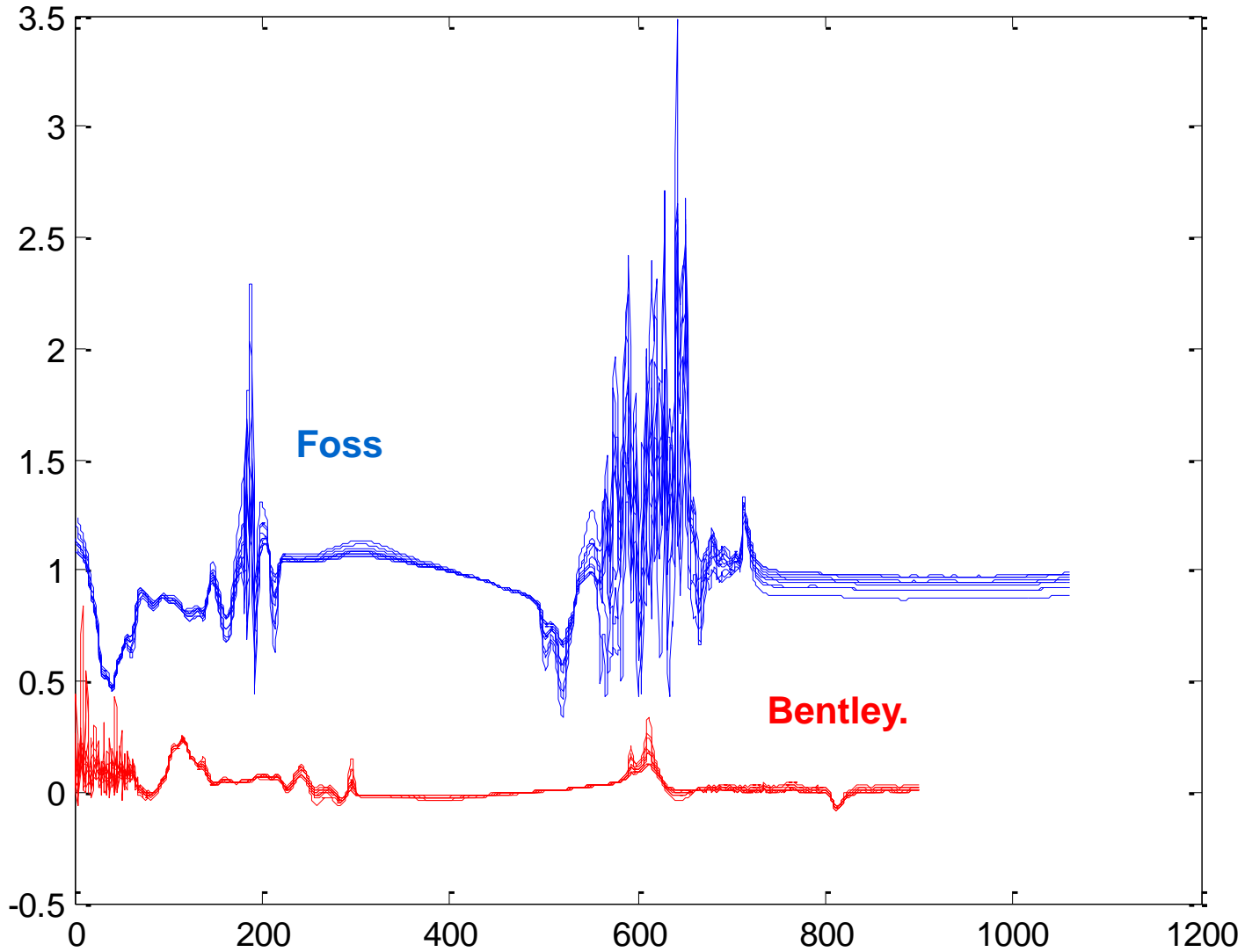


**Prediction tools of
cows states**

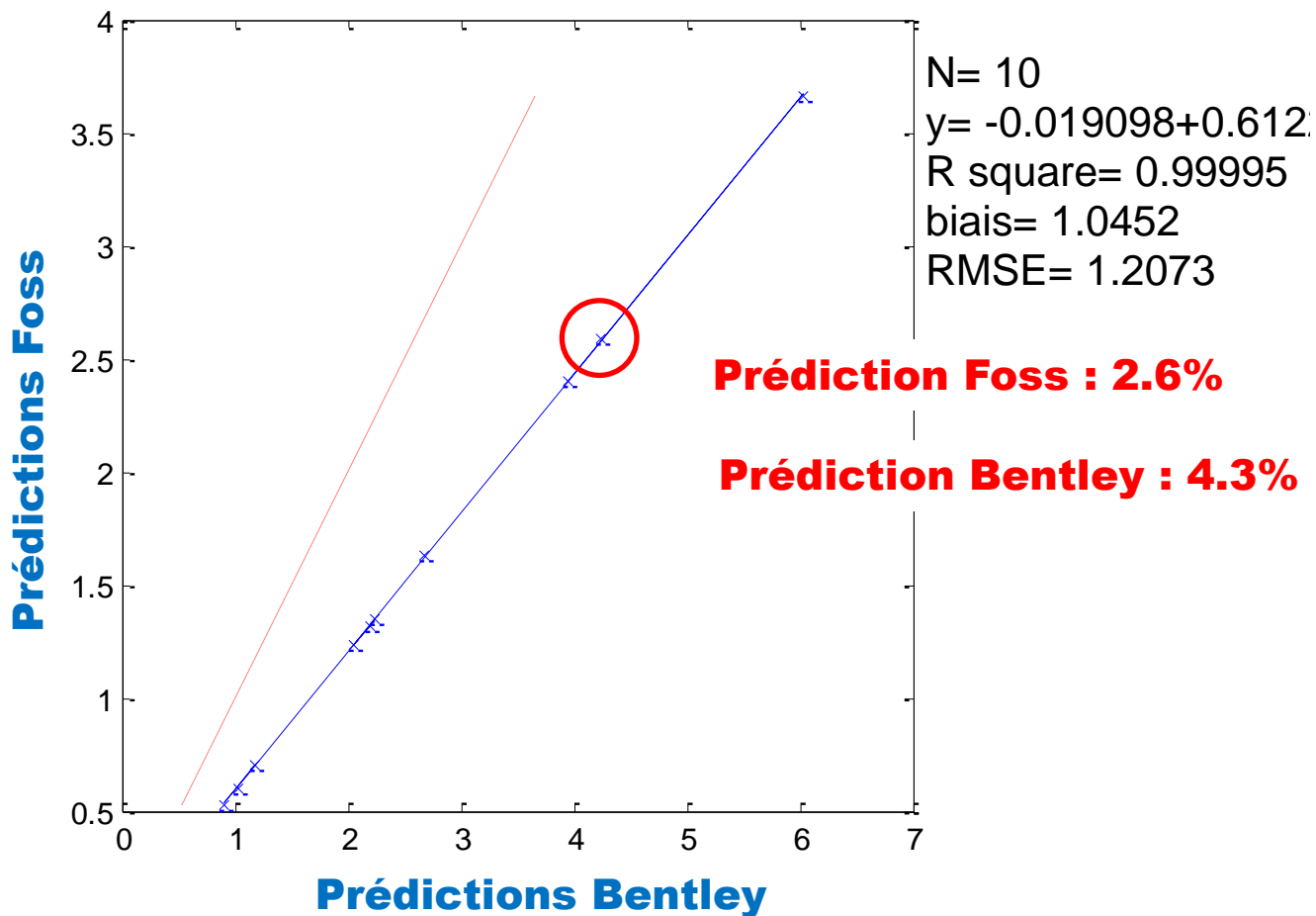
No standard format !!



10 spectra from identical milks :



Predictions fat with raw spectra :



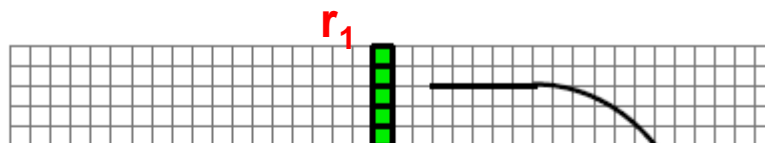
Need of spectral standardisation

PIECE-WISE DIRECT STANDARDIZATION (PDS)

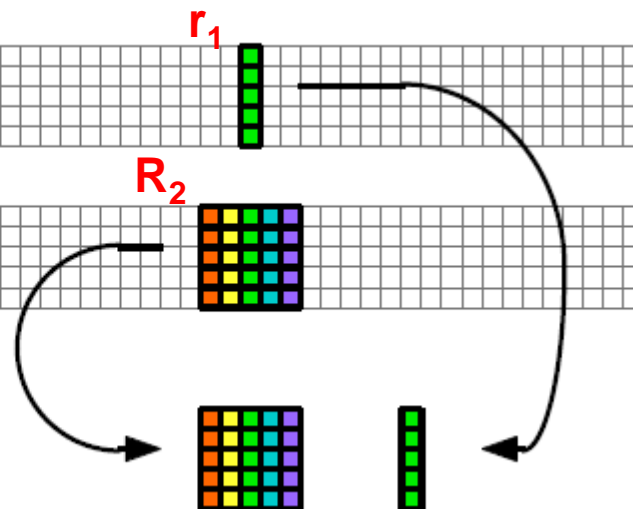
Absorbance in the area of r_1 (master)

→ correlated to R_2 (slaves)

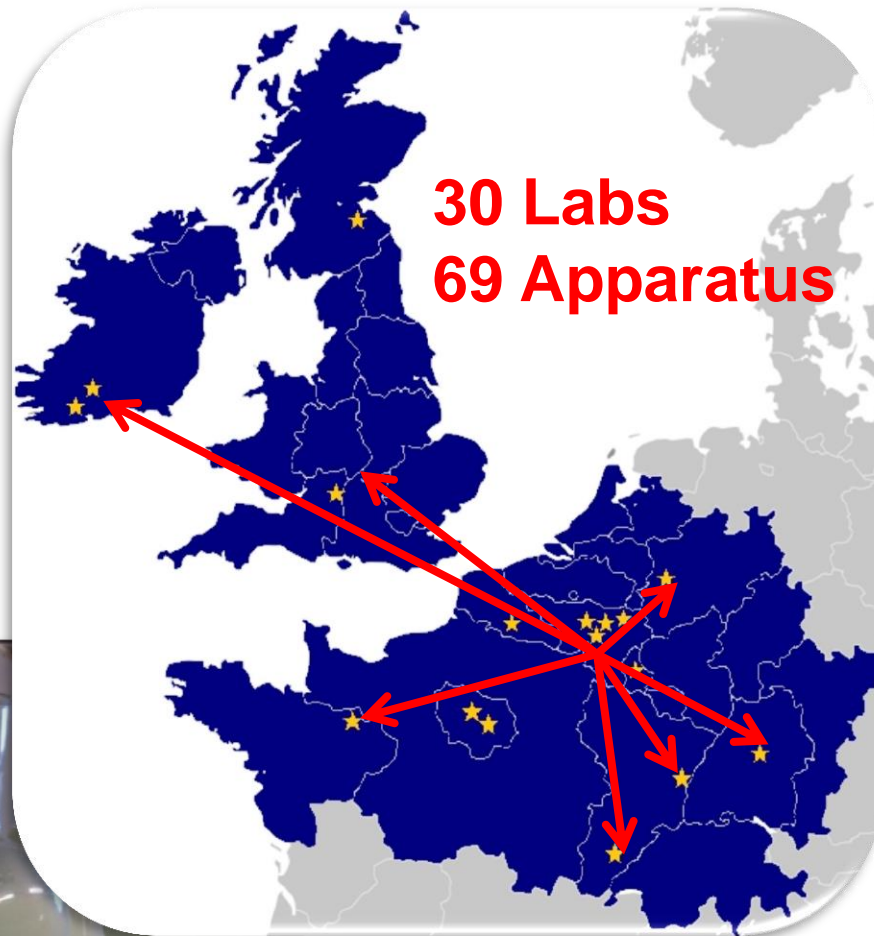
« Master »

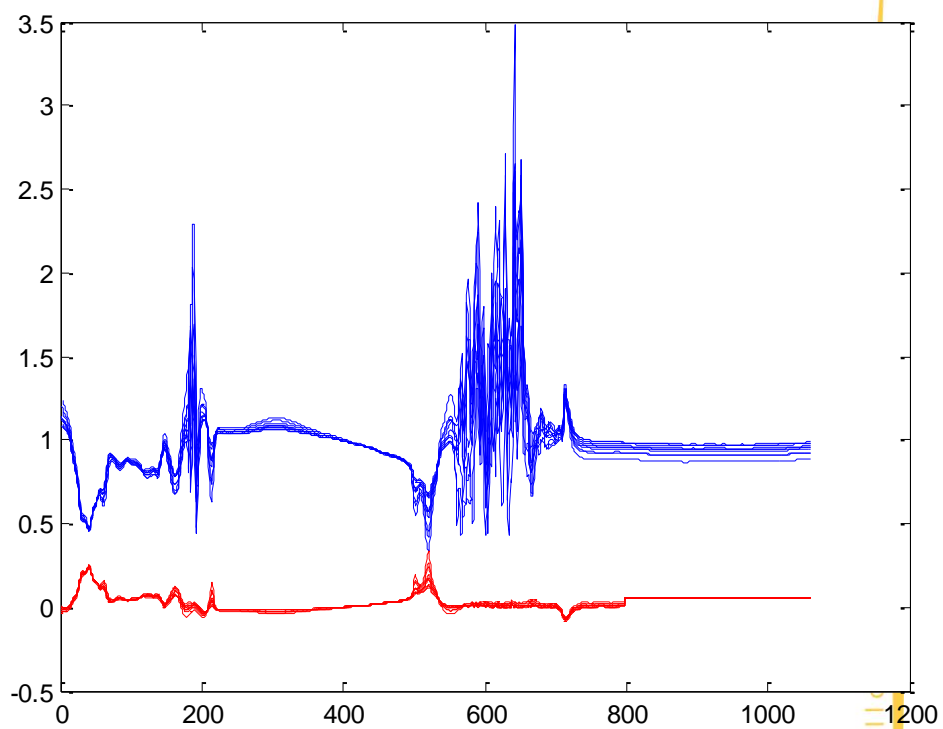
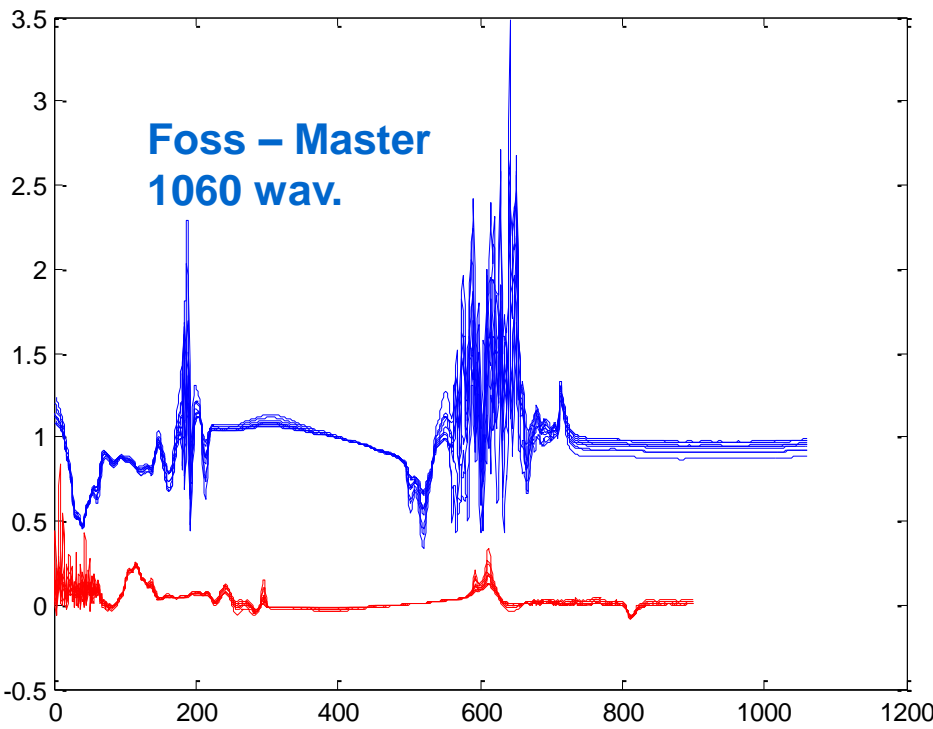


« Slave »

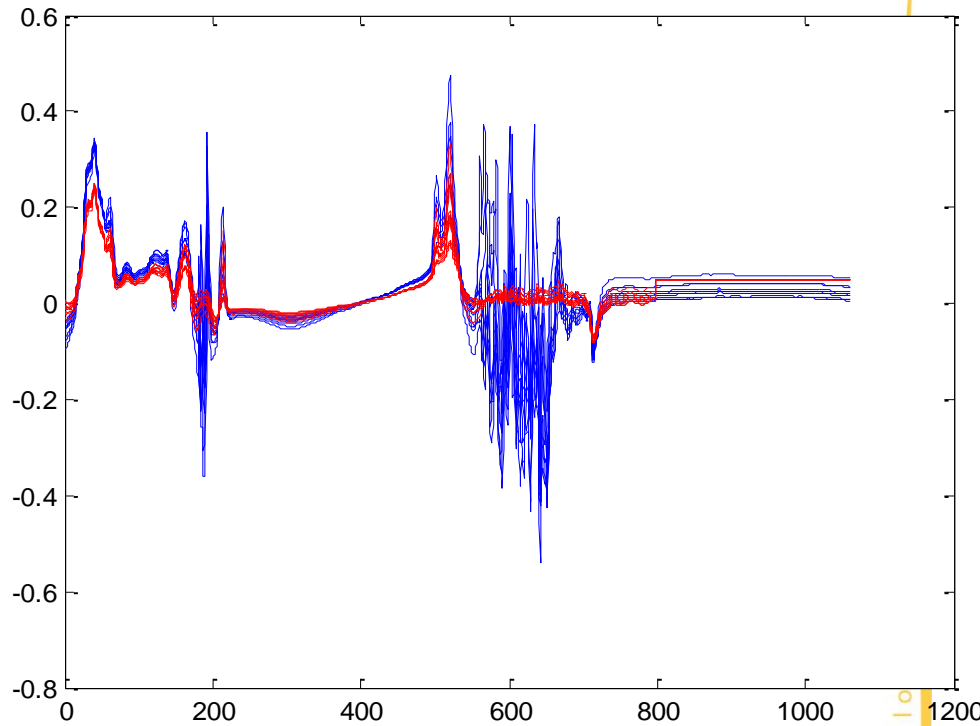
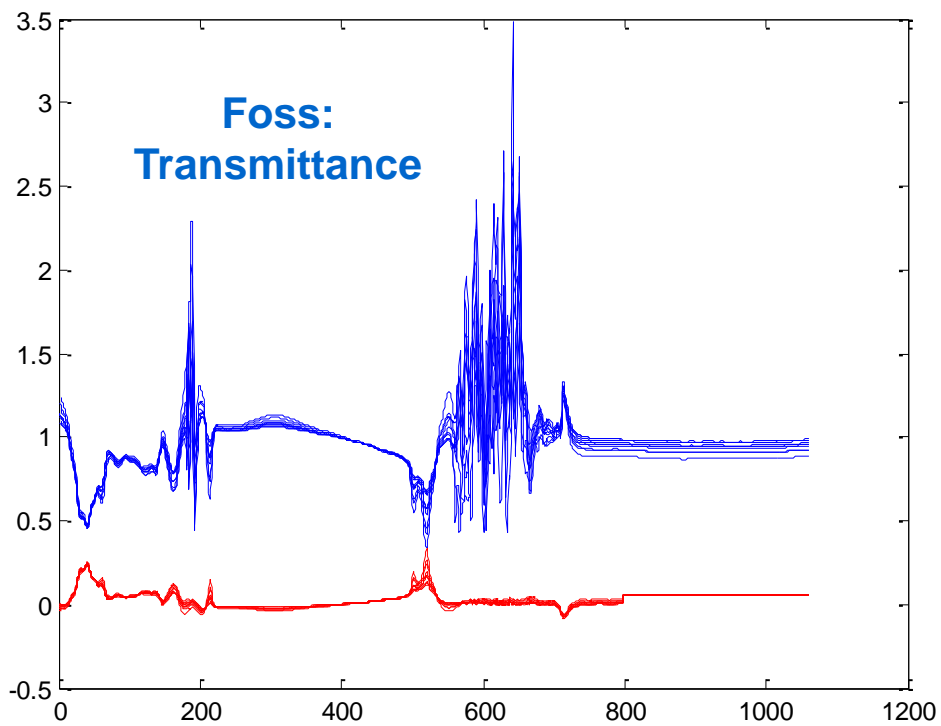


$$r_{1j} = R_{2j} b_j + b_{0j}$$





Interpolation



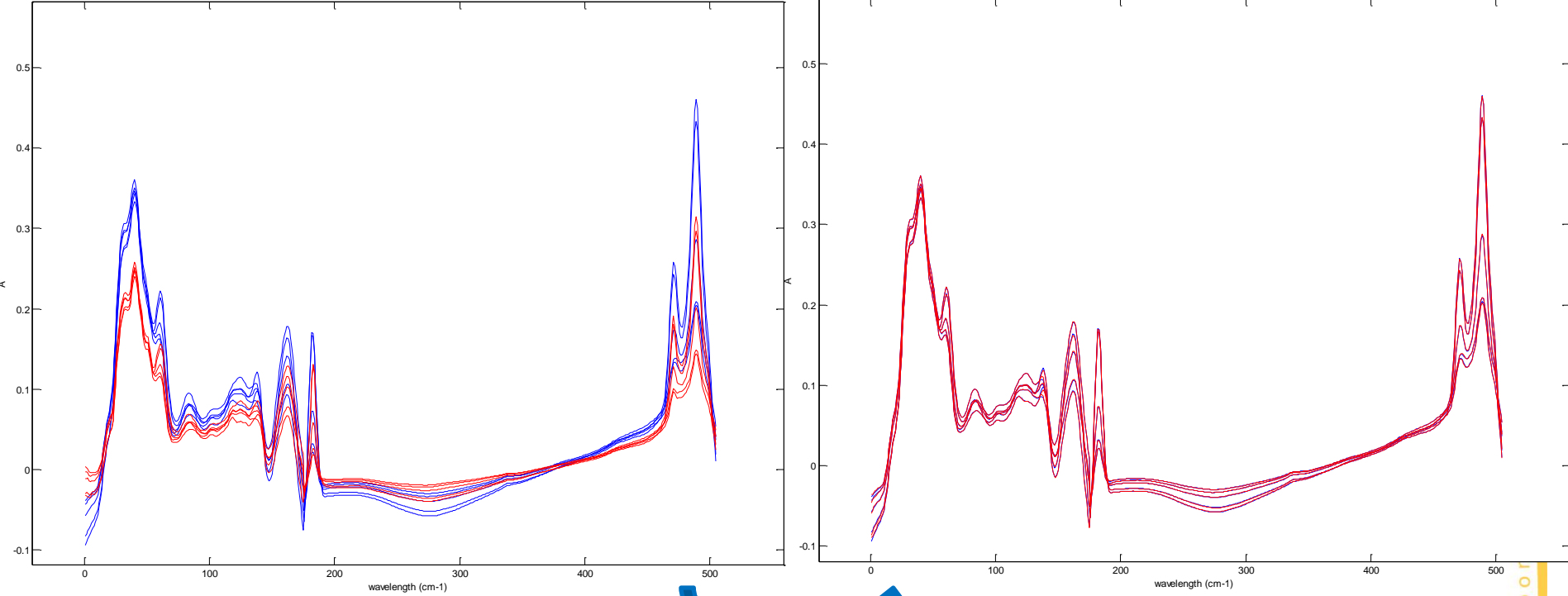
Bentley: Absorbance



**Logarithmic
transformation**

Spectra after log

Spectra after slave standardization

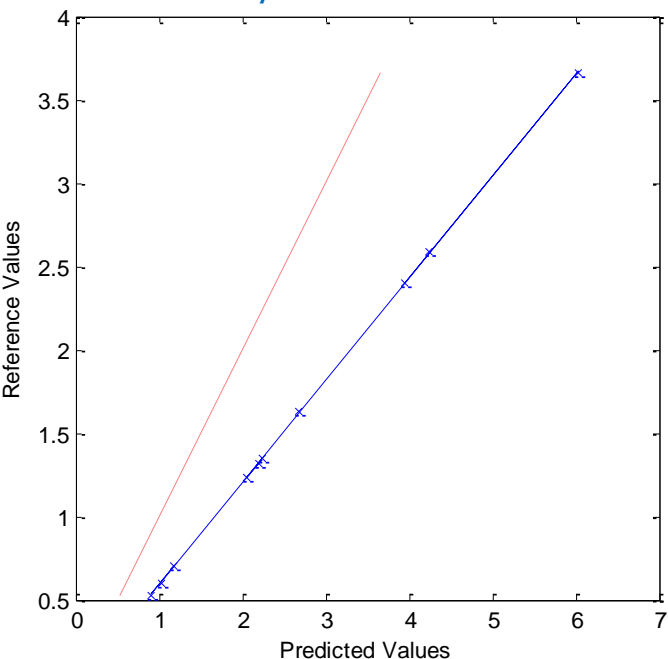


Possibility to merge spectra into a common database

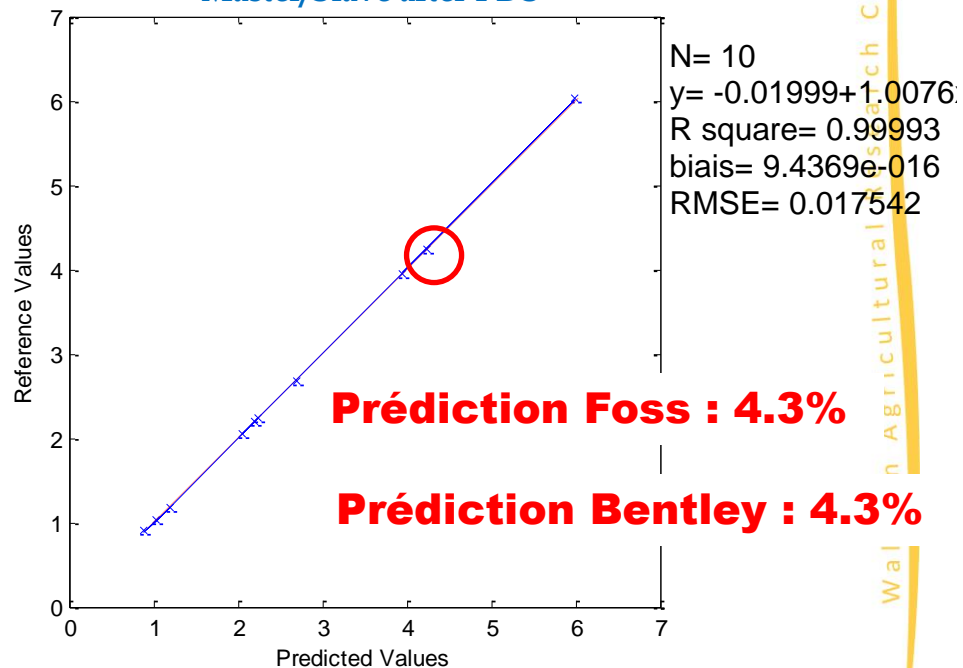
Test: application of fat prediction



Regression of fat prediction
Master/Slave before PDS



Regression of fat prediction
Master/Slave after PDS



Possibility to create and use models on all apparatus

Example of first tools: Methane prediction equation (A. Vanlierde, 2013)



368 Foss spectra

+

84 Delta spectra

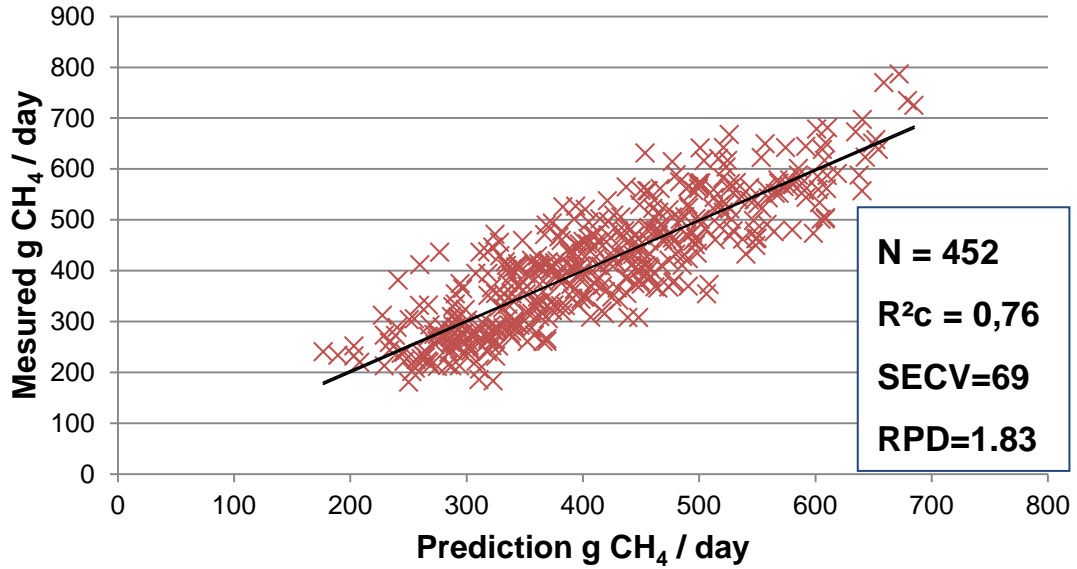


=

452 spectra in a common format

+

Reference values (SF6)



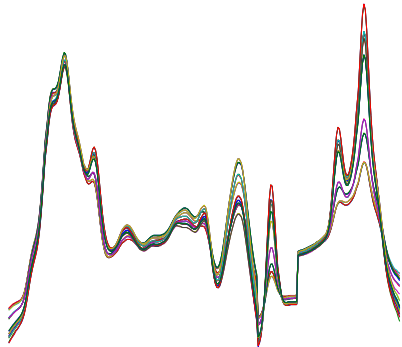
- ✓ PDS allows to group spectra from all apparatus in a common database



- ✓ Allows to create and use universal equation on all apparatus



2015



Thanks for your attention !



WITH SUPPORT OF

