

Network, Guidelines, Certification,

Why IDF-ICAR ExtraMIR Joint Project Plays a Crucial Role in Advancing the Global Adoption of FT-MIR-Based Methane

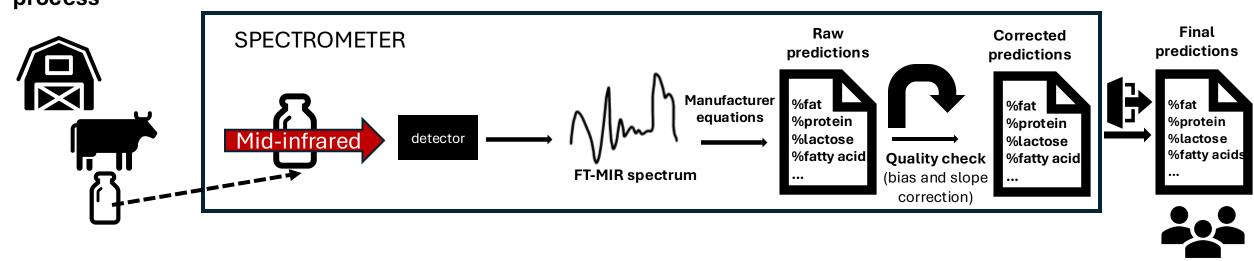
Proxy?

Prof. Hélène Soyeurt

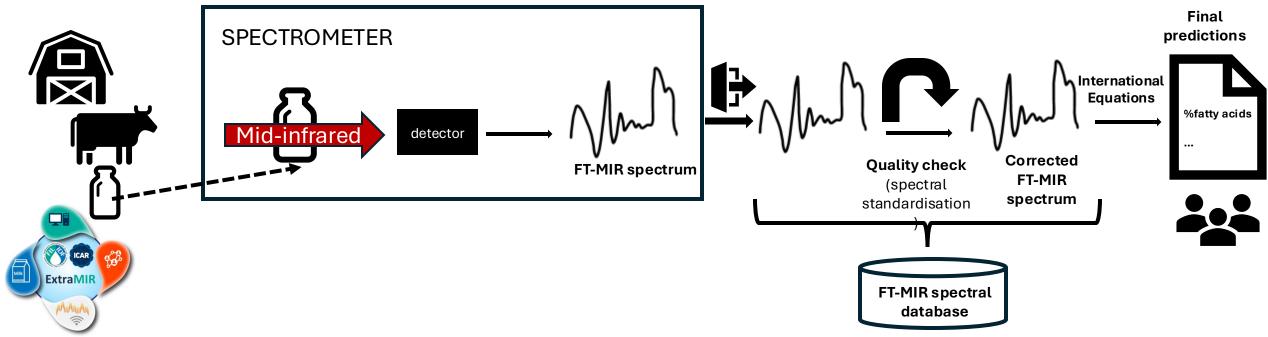
Gembloux Agro-Bio Tech, University of Liège, Belgium

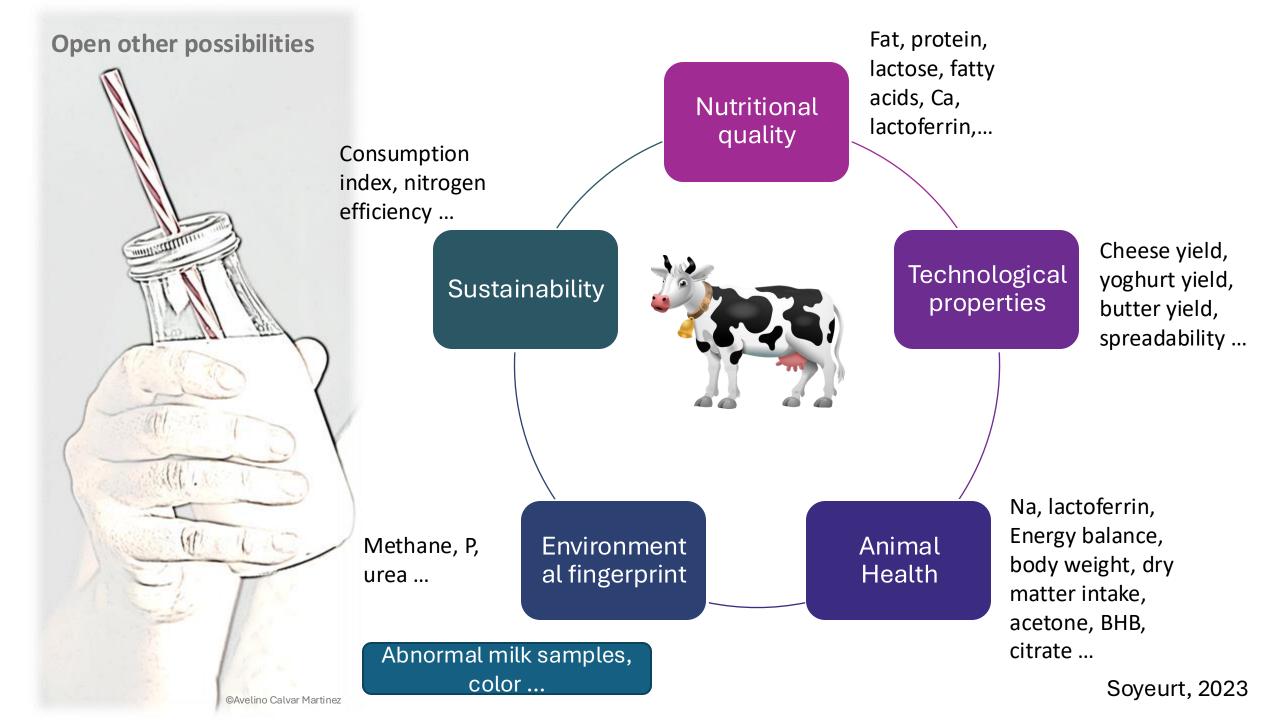


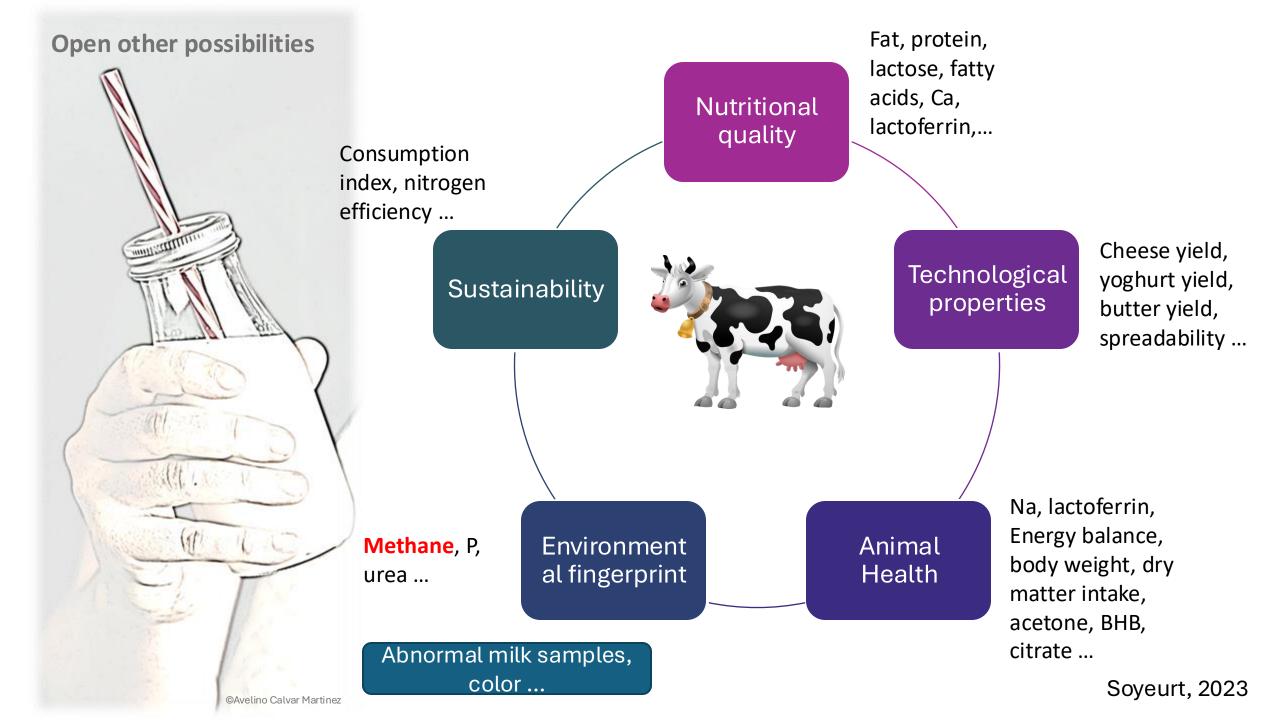
### **Existing Milk analysis** process

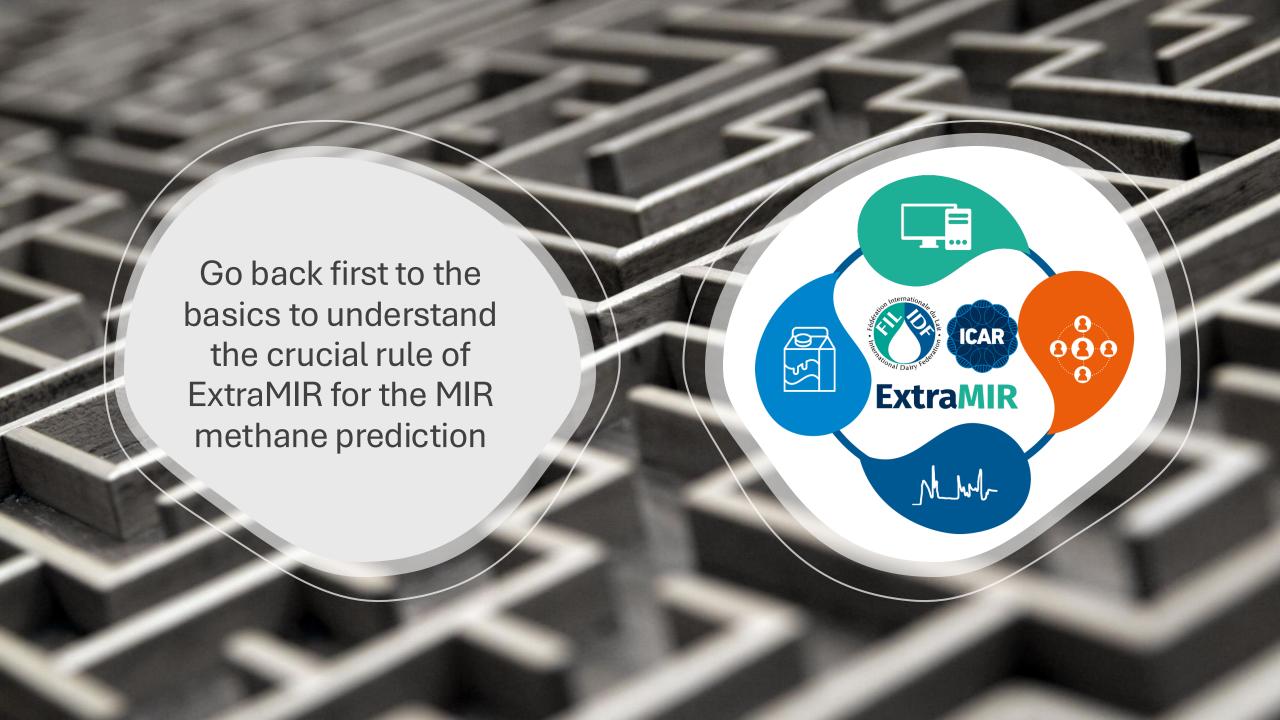


#### **ExtraMIR** proposal

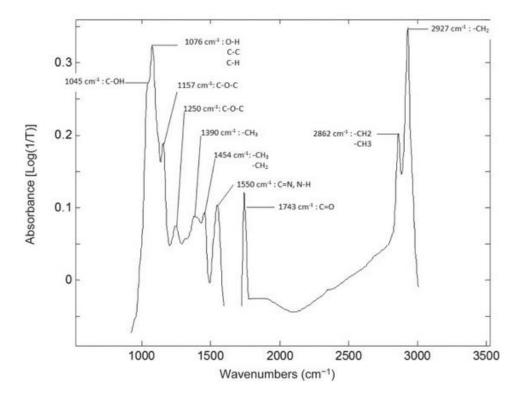




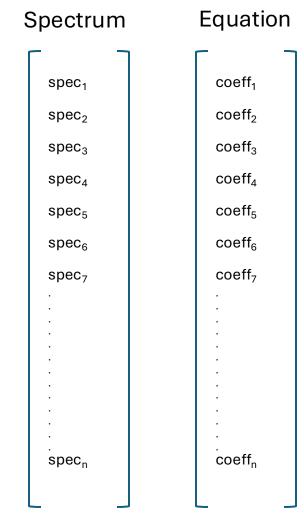




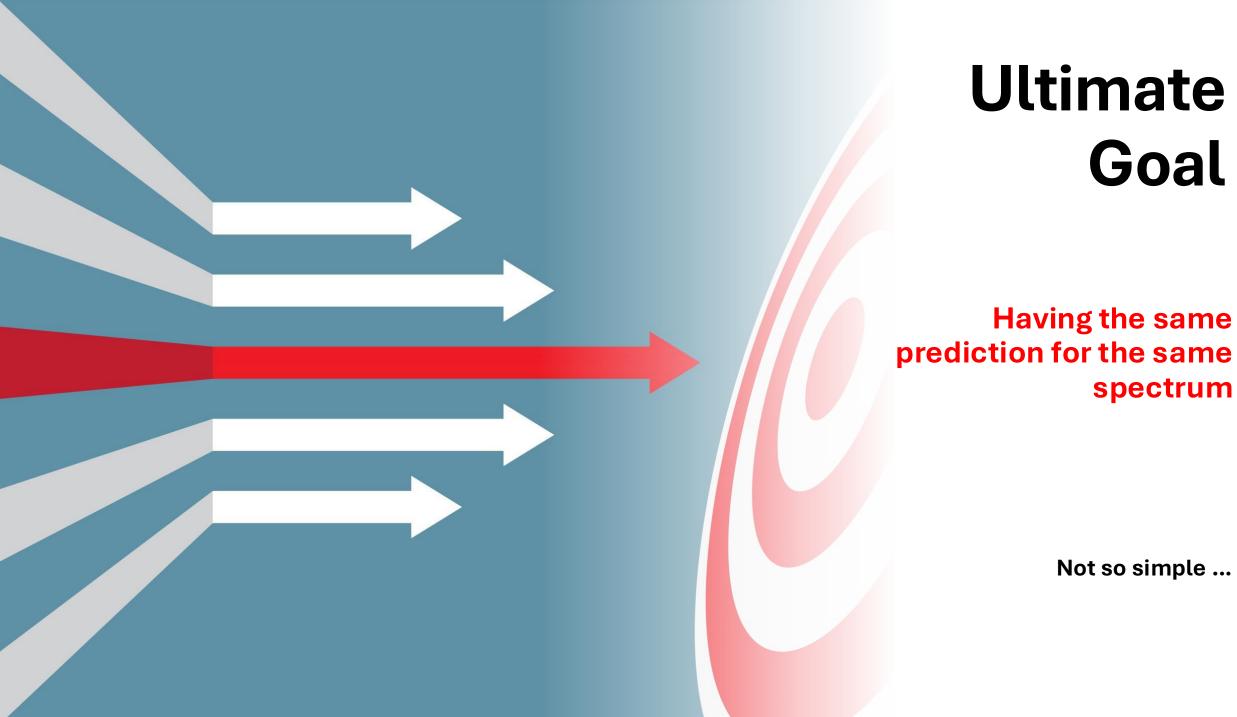
## MIR Prediction



#### 2 components:



Prediction = intercept + sum(spec\*coeff)



# Goal

prediction for the same spectrum

Not so simple ...

How much the spectral variability is representative of the cow population?



References hard to obtain leading a limited dataset

Spectral variability

Spectral representativity

Equation performances

its own validation set

Validation
performances
are not
comparable

Sources of variation

Reference used to create the equation Different teams
develop
equations using
different
references
(respiration
chambers,
greenfeed, SF6,
laser gun ...)



How much are they comparable?

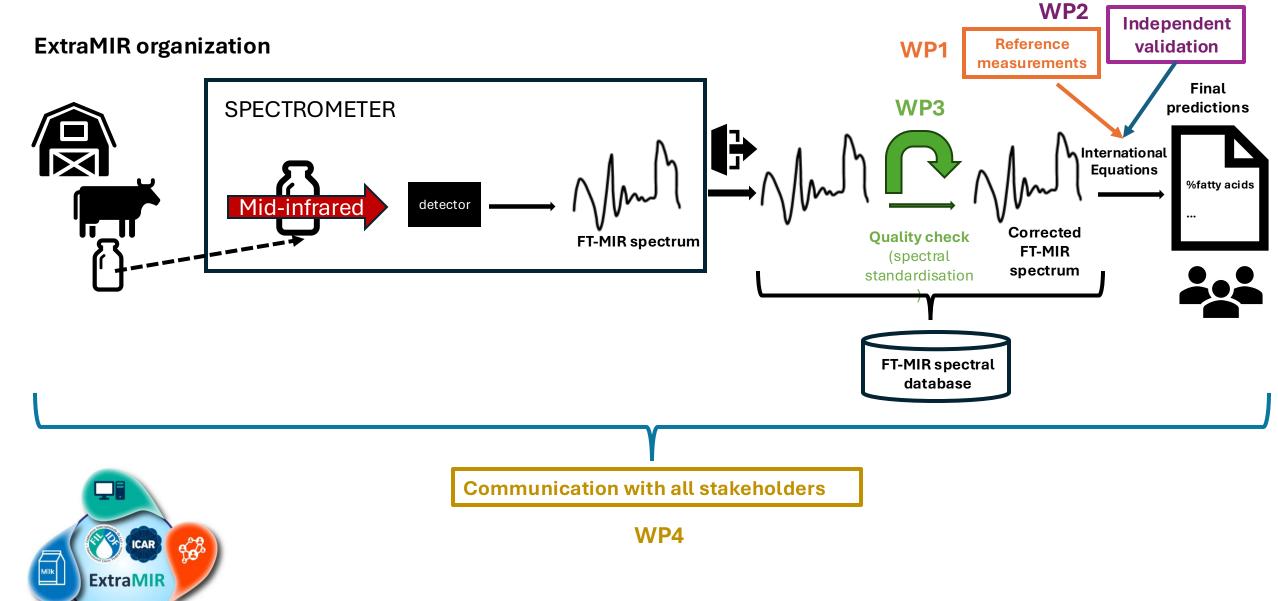
How to standardize the signal?



Different spectral resolutions

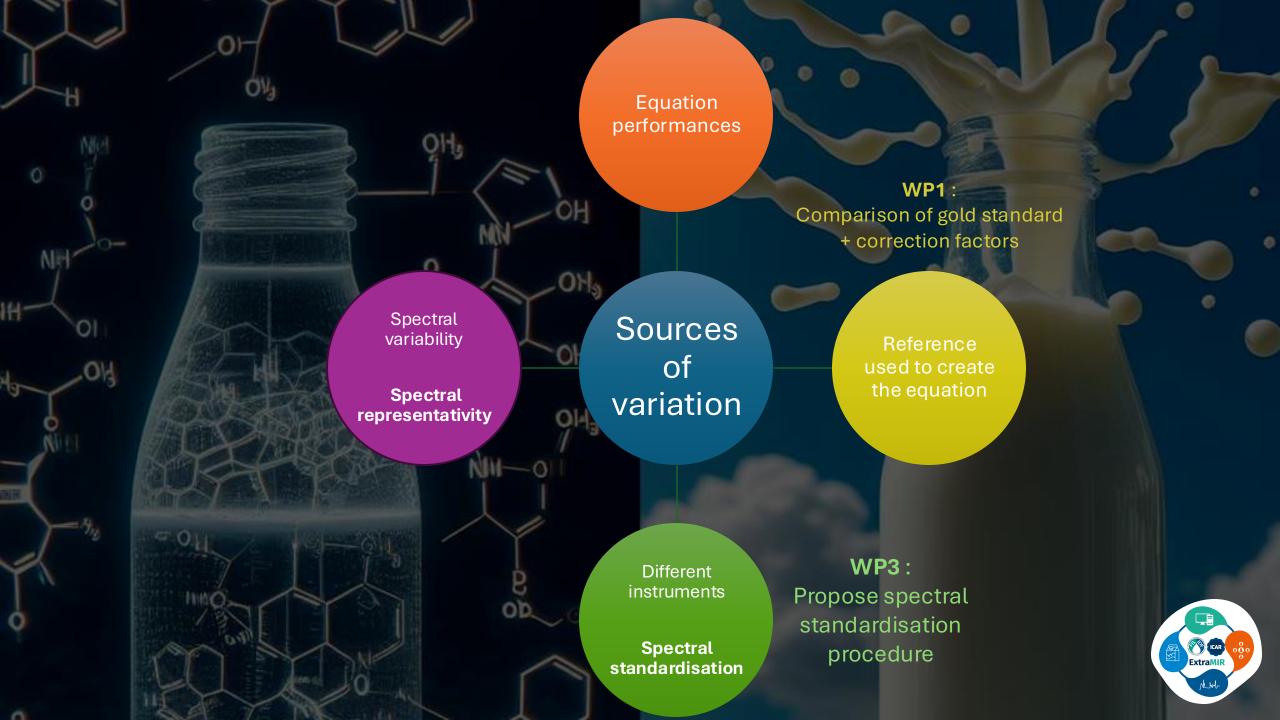
Calibration dataset used

Spectral standardisation



photony

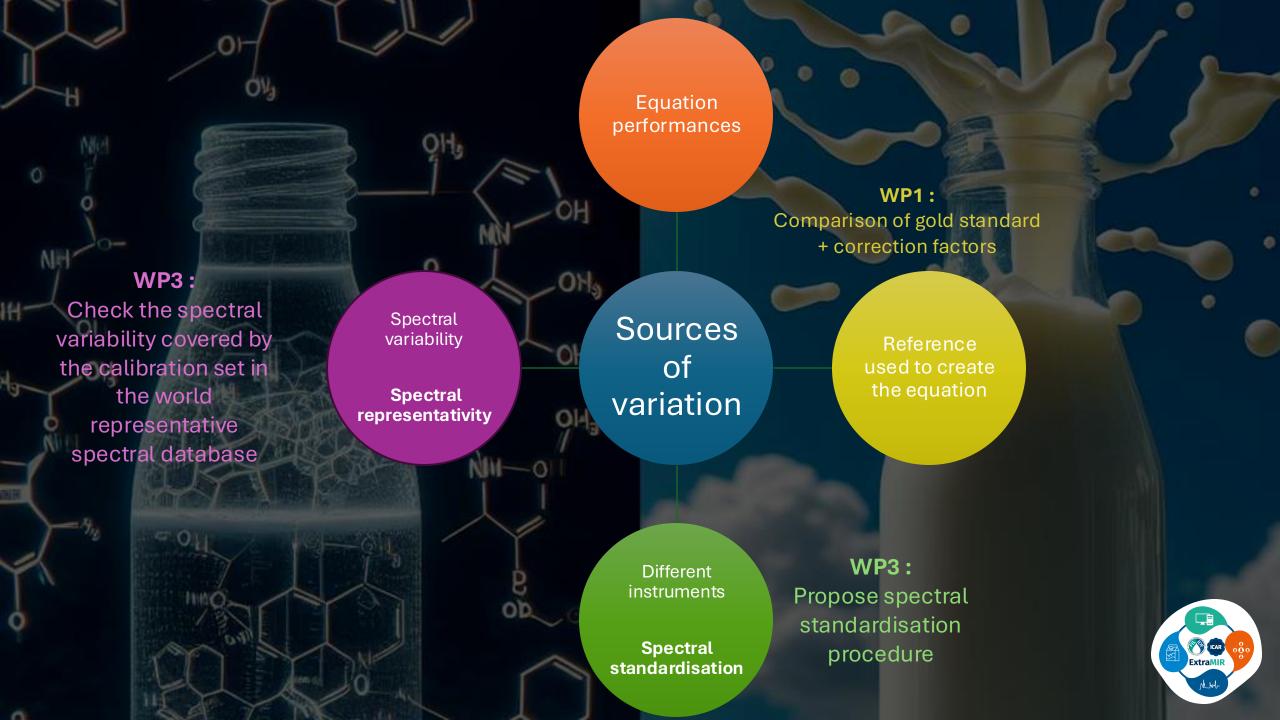
Focussing on Fatty Acids this year but many achievements are interesting for the MIR-based prediction of methane





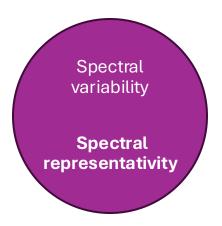
WP3:
Propose spectral standardisation procedure

- A group is formed to compare different spectral standardization using samples sent with the ICAR proficiency test
- Results will be communicated soon

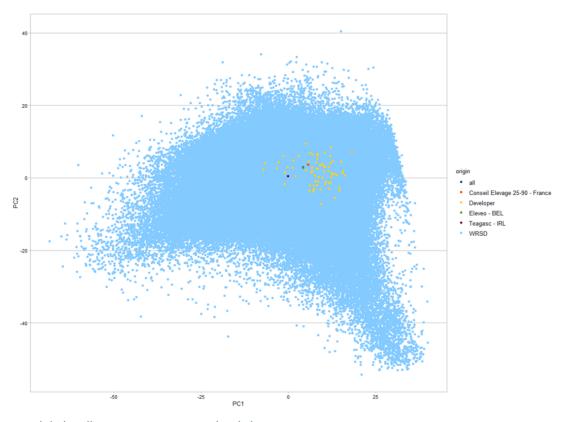


**WP3:** 

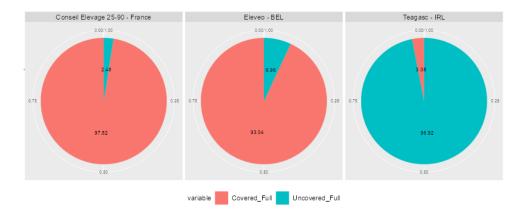
Check the spectral
variability covered by
the calibration set in
the World
Representative
Spectral Database
(WRSD)

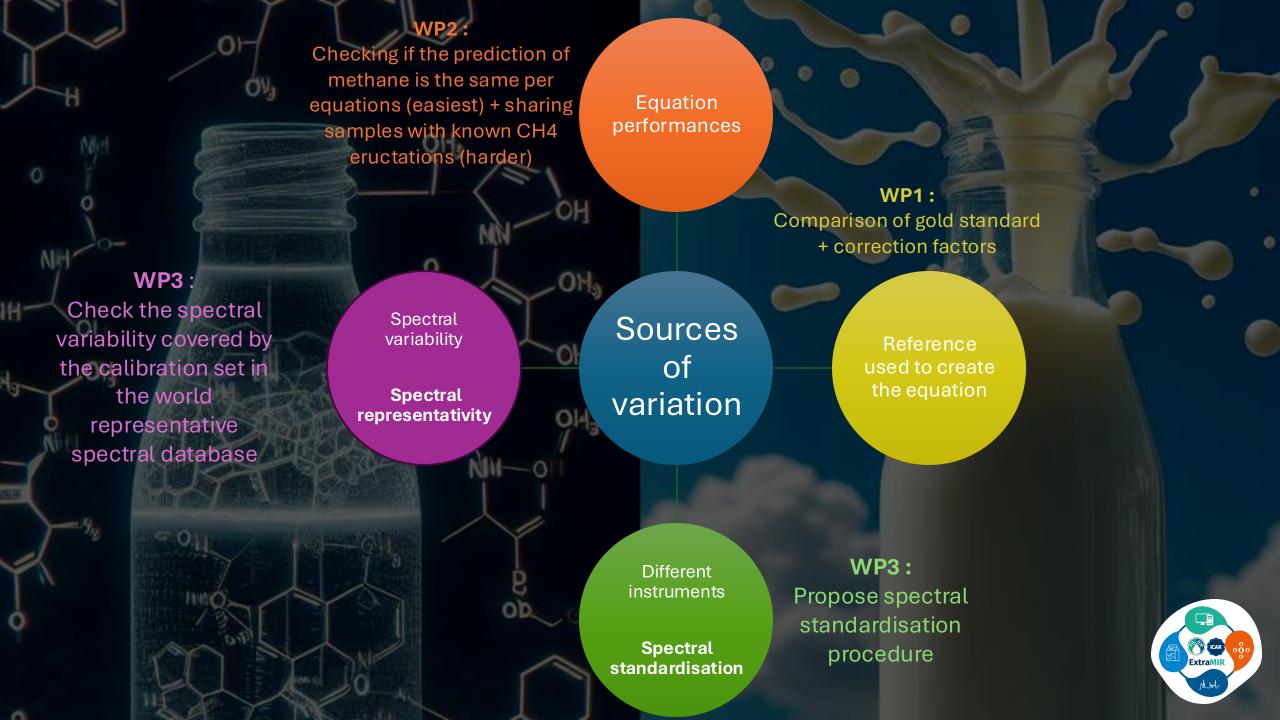


- A web application will be available soon
- The idea is to compare the spectral variability of a calibration set with the spectral variability existing in WRSD



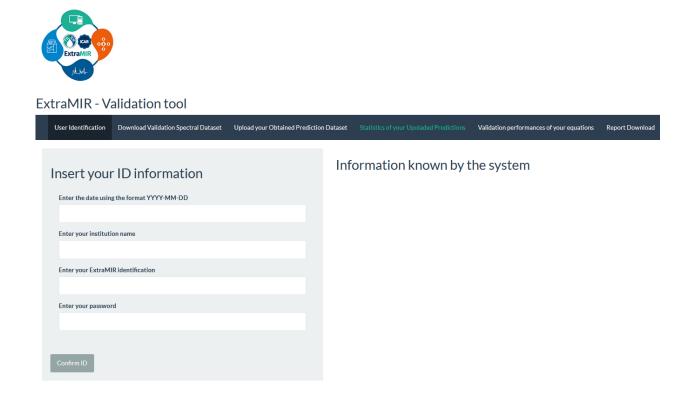
Global-H distance per partners - visual view





WP2:
Checking if the prediction of methane is the same per equations (easiest) + sharing samples with known CH4 eructations (harder)





- A web application developed in RShiny for Fatty Acids will be available soon.
- We can derive a tool for Methane which compare the predictions performances of equations



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Proxy?

Outputs from ExtraMIR can help to ensure high quality MIR-based prediction of methane

**Next chapter for ExtraMIR?** 





