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Context

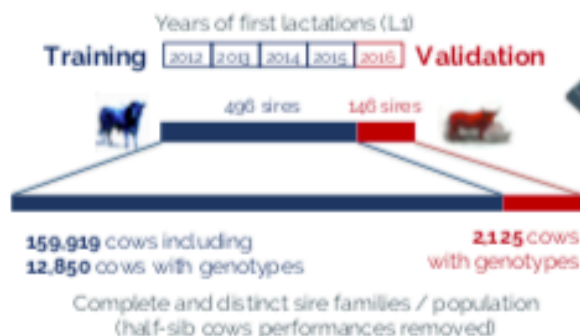
Milk cheese-making properties (CMP), strongly related to milk composition, are economically important BUT difficult and costly to measure

From MIR project 2015-2018

- Equations of prediction of CMP from mid-infrared (MIR) spectra¹
- Genetic analysis of CMP and milk composition traits (proteins, fatty acids and minerals) predicted from 6 million MIR spectra from 400,000 Montbéliarde cows^{2,3}

Objective 1 Reliability of ssEBV in a validation population

Training and validation sets



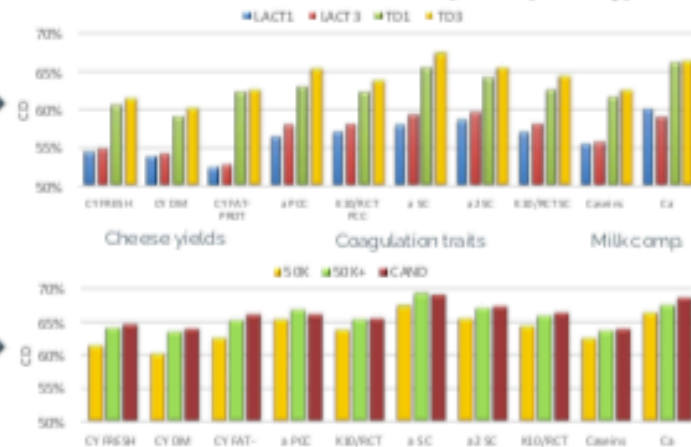
4 models tested

| | L1 | L1 - L3 |
|--------------------|----------------------|----------------------|
| Mean per lactation | LACT1 191,532 obs | LACT3 327,265 obs |
| Test-day records | TD1 1,422,782 obs | TD3 2,059,054 obs |

3 scenarios tested with TD3

| | | |
|---------------------|--------------------|---------------------------------|
| SNP number | 41,942 50K SNPs | 477,94 50K+ GWAS candidate SNPs |
| SNP effect variance | Equal for all SNPs | Higher for 5 to 14 SNPs |
| 3 scenarios | 50K | 50K+ CAND |

CD validation = correlation² (ssEBV, adjusted phenotype) / h²



Objectives

- Estimate the reliability of single step GBLUP breeding values (ssEBV)⁴ by testing 4 models and 3 scenarios (different SNP sets including or not candidate variants detected by GWAS) in a validation population
- Estimate genetic trends of CMP traits
- Simulate different breeding objectives including CMP traits

For 10 traits related to CMP:
3 cheese yields
5 coagulation traits
% caseins and Ca in milk

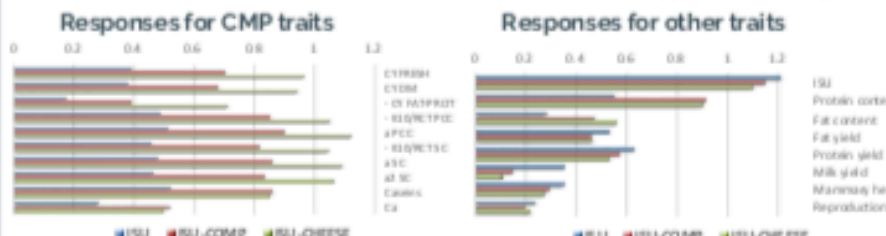
Objective 2 Genetic trends of CMP traits

Application of genomic prediction equations on genotyped Montbéliarde cows (311,761) and bulls (21,171) born between 2005 and 2018



Objective 3 Responses to 3 breeding scenarios

Current total merit index ISU (milk production and composition, mammary health, reproduction, longevity and conformation)
Alternative breeding objectives:
ISU-COMP = 0.7 ISU + 0.3 Caseins
ISU-CHEESE = 0.7 ISU + 0.1 (CY_{DM} + a_{PCC} + (-K10/RCT_{PCC}))



Conclusions

Reliable cheese-making genomic indexes, the most reliable and less biased obtained with:

- A test-day model applied to the first 3 lactations
- 50K SNP+ candidate variants detected by GWAS
- Higher weightings for candidate variants

Indirect favorable response for CMP traits with the current breeding objective but possibility to increase responses by directly including CMP traits in the breeding objective with a limited impact on other traits

In 2021: release of Single-Step genomic evaluation of CMP predicted from MIR spectra in Montbéliarde cows

References

- ¹El Jabri M, Sanchez M-P, Trossat P, Laithier C, Wolf V, Graspequin P. & al. 2019. J. Dairy Sci. 102:6943-6958
²Sanchez M-P, El Jabri M, Minéry S, Wolf V, Beuquier E, Laithier C & al. 2018. J. Dairy Sci. 101:10048-10061
³Sanchez Y, Ramayo-Caldas V, Wolf V, Laithier C, El Jabri M, Boussaha & al. 2019. Genet. Sel. Evol. 51:34
⁴Tribout, T., Ducrocq V., Boichard D. 2020. ICGG6, paper #47.

