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Context

Titratable acidity (TA)

- TA influences all phases of milk coagulation
- Developed acidity results from bacterial activity
 - 🖌 Lactic acid
 - ✓ Collection, transportation, and transformation of milk
- Fresh milk
 - ✓ Some components: carbondioxide, citrates, casein, albumin/globulin, and phosphates
 - ✓ Buffer action

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Objective

 To investigate the potential use of MIR spectrometry in order to predict TA

- TA recorded as Dornic degree
- Walloon Region of Belgium
- Multibreed

Material and methods

Sampling

- Walloon Region of Belgium
- Large variability: several criteria
 - ✓ Milk sampling: individual or bulk milk
 - ✓ Breed: Dual Purpose Belgian Blue, Holstein, Red-Holstein, Montbeliarde, and Jersey
 - ✓ Time of sampling: morning milking, evening milking or mix of 50 % morning & 50 % evening milk samples

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- ✓ Analysed traits: fat, protein, free fatty acid (FFA), urea, lactose, dry matter (DM), somatic cell count (SCC), and pH
- ✓ SCC → Somatic Cell Score (SCS)

Material and methods

Analysis

- Titratable acidity
 - ✓ Recorded as Dornic degree (D°)
 - ✓ 0.1 N NaOH solution
 - ✓ Consumption of NaOH to shift the pH value from 6.6 to 8.4 (phenolphthalein)

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Calibration procedure

- First derivative pretreatment
- Partial least square regressions
- 22 outliers
- Statistical parameters
 - ✓ Mean and standard deviation (SD)
 - ✓ Standard error of calibration (SEC)
 - ✓ Calibration coefficient of determination (R²_c)

Material and methods

Calibration procedure

- Cross-validation
 - ✓ To determine the number of factors
 - ✓ To assess the accuracy of equation
 - ✓ Partitioning randomly the calibration set: 102 groups
- Statistical parameters to assess the accuracy
 - ✓ Standard error of cross-validation (SECV)
 - ✓ Calibration coefficient of determination (R²_{CV})
 - ✓ RPD = SD / SECV

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Results		
Characterization of the	samples	
Trait	Mean	SD
Fat (%)	3.88	1.03
Protein (%)	3.49	0.52
FFA (mmol/100 g of Fat)	5.63	8.62
Urea (g/100 mL)	0.023	0.011
Lactose (g/100 mL)	4.85	0.35
DM (%)	12.66	1.25
SCS	3.31	1.90
		0.00
рН	6.69	0.09

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SCS	2 21	1 00
рН	efficient of Var	iation = 14 %
TA (D°)	16.27	2.27
 * FFA = Free Fatty Acid; DM = Dr D° = Dornic degrees. 	y matter; SCS = so	omatic cell score;

R	es	u	ts

Observed	correlations	among m	ilk components
		- 0	

	Fat	FFA	Protein	Urea	Lactose	DM	SCS	рН
TA (D°)	0.04 ^{NS}	0.13 *	0.39***	0.18**	0.21**	0.26***	-0.16*	-0.32***
Fat		0.41***	0.42***	0.13*	-0.19**	0.89***	0.18**	-0.18**
FFA			0.68***	0.41***	-0.17**	0.50***	0.04 ^{NS}	-0.38***
Protein				0.30***	-0.07 ^{NS}	0.69***	0.10 ^{NS}	-0.26***
Urea					0.18**	0.25***	-0.18**	-0.01 ^{NS}
Lactose						0.11 ^{NS}	-0.40***	0.66***
DM							0.07 ^{NS}	-0.06 ^{NS}
SCS								-0.19**
FFA = Fre	e Fatty A	cid: DM =	Drv matter:	SCS = son	natic cell sc	ore: D° =	Dornic deg	rees.

* = P-value < 0.05; ** = P-value < 0.01; *** = P-value < 0.001; NS = non significant.

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• Obs	serve	d cor	relatior	ns am	ong mi	ilk coi	mpone	ents
	Eat	EEA	Protein	Uroa	Lactose	DM	272	nH

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Perspectives

- Validation with new set of samples
- Use of this equation
 - Walloon Database: 900,000 spectra
 - Study of TA variability in the Walloon dairy cattle
 ✓ Detection of potential effects of breed, season, DIM...
 - Development of a genetic evaluation
 - TA breeding values + others traits = a new economic index for cheese making abilities ?

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