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Uncovering environmental effects and genetic parameters on the mineral composition of sheep milk

**Johanna Ramírez-Díaz¹, Giorgia Stocco², Claudio Cipolat-Gotet², M. Ablondi², A. Summer², Alessio Negro³,
Alessandro Lotto⁴, Simone Blotta⁴, Francesco Tiezzi⁵, Stefano Biffani¹**

*¹National Research Council, Institute of Agricultural Biology and Biotechnology, Milano, Italy, ²University of Parma, Dep. of Veterinary Science, Parma, Italy,
³Associazione Nazionale della Pastorizia (Asso.Na.Pa.), Ufficio Studi, Roma, Italy, ⁴Nutristar spa, Reggio Emilia, Italy, ⁵Department of Agriculture, Food,
Environment and Forestry, Università degli Studi di Firenze, Firenze, Italy*

INTRODUCTION

Cheese Industry

- The global dairy sheep cheese production reached 1.5 million metric tons in 2021-2022



- The sheep cheeses production in Italy in 2022 was approximately 37 tonnes

Rural context and development

The sheep production systems:

- Importance in the rural context:
Challenges to produce other species
- Nutritional composition and taste of dairy products
- Higher quality composition



INTRODUCTION



Average values of sheep milk composition:

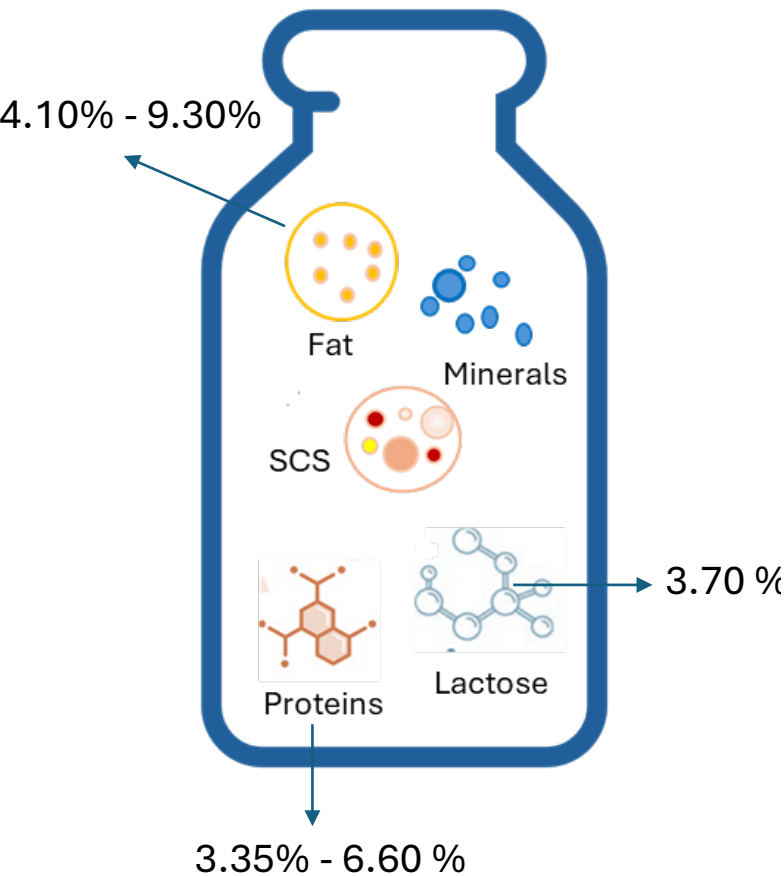
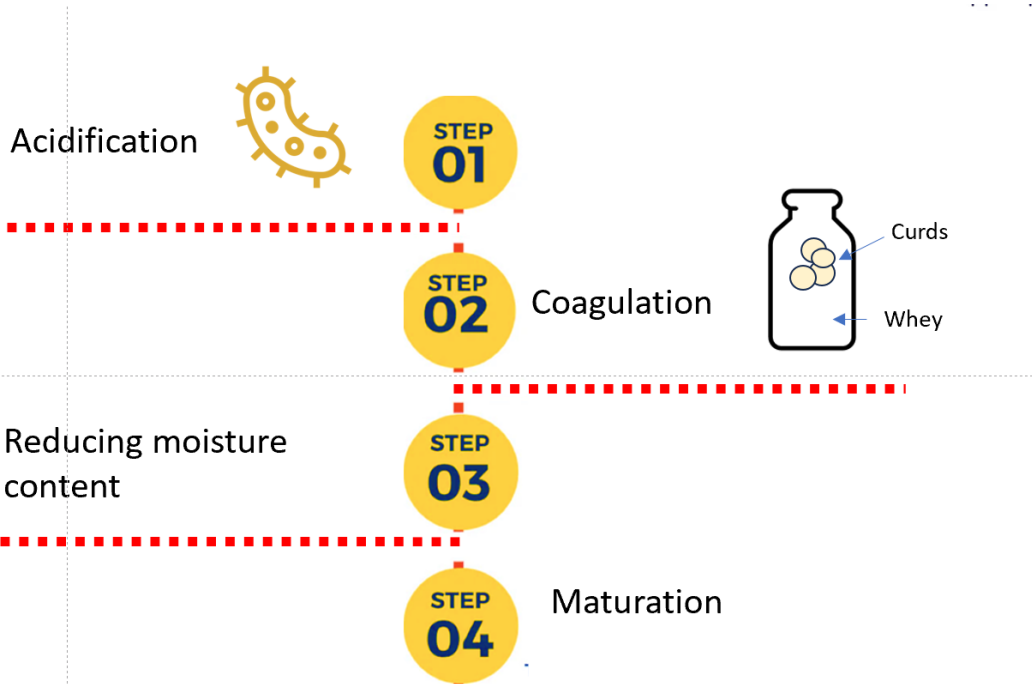


Table 5–Concentration of minerals in milk from various sources

	Cattle ^a	Sheep ^a
(mg/100 g)		
Calcium	122	195 to 200
Phosphorus	119	124 to 158
Potassium	152	136 to 140
Magnesium	12	18 to 21
Sodium	58	44 to 58
(µg/100 g)		
Zinc	530	520 to 747
Iron	80	72 to 122
Copper	60	40 to 68
Manganese	20	5.3 to 9
Iodine	2.1	10.4
Selenium	0.96	3.1

^aSchryver and others (1986); ^bMehaia and others (1995); ^cSalimei and others (2010).

Barłowska et al., 2011



Ca, Na, K and S -> Dairy Cattle

INTRODUCTION



Limited and almost non-existent in local breeds -> Cheese production -> environmental adaptation, traditions and cultural heritage

NEW ZEALAND JOURNAL OF AGRICULTURAL RESEARCH
<https://doi.org/10.1080/00288233.2024.2368505>



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RESEARCH ARTICLE

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Estimation of genetic parameters for production, composition and processability of milk from dairy sheep in a New Zealand flock

Ana Carolina Marshall ^{a,b}, Nicolas Lopez Mike Weeks ^d and Warren McNabb ^b

^aDepartment of Animal Science, School of Agriculture New Zealand; ^bThe Riddet Institute, Massey University Scientific and Industrial Research Organisation (CSIRO) Foods & Bioproducts Group, AgResearch Ltd, Massey University, Palmerston North, New Zealand



Journal of Dairy Science
Available online
In Press, Journal Pre-proof

Detailed mineral profile of cheese from cows, buffaloes, goats, and dromedary camels, and efficiency of recovery of minerals in their cheese

Nicolò Amalfitano ¹ , Nageshvar Patel ¹, Mohamed-Laid Haddi ², Hamida Benabid ³, Michele Pazzola ⁴, Giuseppe Massimo Vacca ⁴, Franco Tagliapietra ¹, Stefano Schiavon ¹, Giovanni Bittante ¹

710. Heritability and genomic analysis of coagulation event in sheep milk

G. Gaspa , A. Cesarani , F. Correddu , M. Congiu , C. Dimauro , A. Pauciuolo , N.P.P. Macciotta

Pages: 2928 - 2931

https://doi.org/10.3920/978-90-8686-940-4_710

Published Online: February 09, 2023

Abstract

References

Full-text

Research

Estimation of genetic parameters for cheese-making traits in Spanish Churra sheep

R. Pelayo ¹, B. Gutiérrez-Gil ¹, A. Garzón ², C. Esteban-Blanco ¹, H. Marina ¹, J.J. Arranz ¹

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PMCID: PMC8453569
PMID: [34014003](#)

Underlying milk production traits in Valle del Belice
ability mapping

Anna Maria Sutera, ¹ Marco Tolone, ² Salvatore Mastrangelo, ² Rosalia Di Gerlando, ² Maria Teresa Sardina, ² Baldassare Portolano, ² Ricardo Pong-Wong, ³ and Valentina Riggio ^{3, 4}

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INTRODUCTION



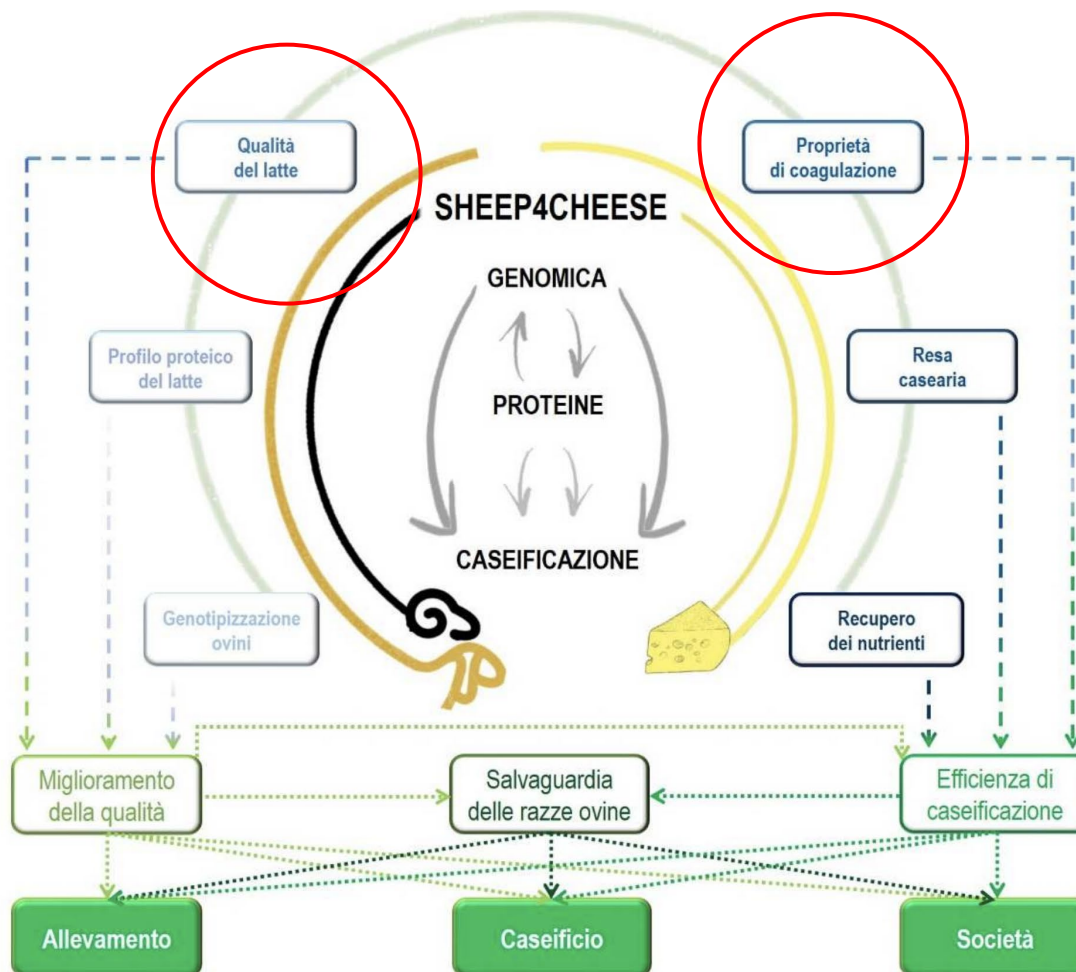
- In general, sheep breeding programs are related to Milk yield, Meat and growth traits and disease resistance → Programs to improve the cheese making process don't exist → mineral composition
- The role of minerals in improving milk composition and quality → for cheese production
- Genetic programs → Have permanent and cumulative changes in economic important traits:
 - Step 1: To guide effective breeding strategies is necessary to estimate variance components and genetic parameters



Aim:

Improving the milk quality and efficiency of cheese-making process

To improve the quality of the sheep milk through genetic selection of the animals



Objectives



General plan of activities to assess the composition and mineral content:

Genotyping



Parental information from
Massese and Comisana Sheep

Phenotyping



Evaluations of Mineral contents and
milk composition



Evaluation of coagulation and
cheese making process

1. Data analysis

2. Statistical modelling

3. Genetic parameters
Single and multitrait models

4. Genome wide association analyses
Single, multitrait models and pleiotropic effects

5. GEBVs and EBVs estimation
Selection to economical important traits and
genetic gain prediction

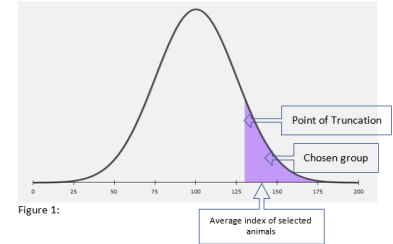


Figure 1:

MATERIAL



Italian Apenines



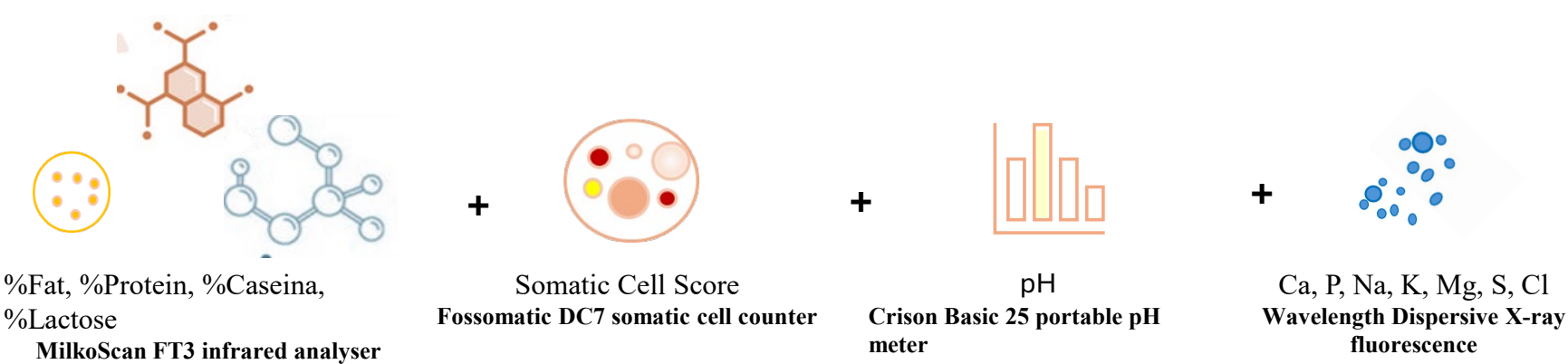
Massese sheep
n = 311



Comisana sheep
n = 429

Central and
Southern Italy

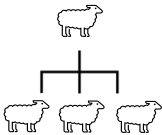
Milk samples
Morning milking
11 sampling season
Laboratory analysis



To explain this:

**Inclusion of environmental factors:
animal status, sampling and other traits**

Inclusion of genetic information



Pedigree:
Comisana: 742,851 animals
(sires: 15033 and 172,481 dams)
Massese: 132,685
(sires: 3014 and 45626 dams).

METHODS

$$y_t = X\beta + Wc + Zu + e$$

Sampling

Additive genetic Effects



Quality Control

Missing information
Biological Thresholds
....
Distribution on PO
Distribution on DIM

DIM:

Class 1: 8 -30d ,
Class 2: 31 -50d ;
Class 3: 51-70d ;
Class 4: 71-90d ;
Class 5: 91-110d;
Class 6: 111 -130d ;
Class 7: 131-148d

Milk Yield

class 1: 0.3 – 0.55 kg;
class 2: 0.6 – 0.79 kg;
class 3: 0.8 – 0.97 kg;
class 4: 1 – 2.33 kg

Parity Order:

1;2;3;4; > 5

Fat %

Casein %

Lactose %

pH

SCS

RESULTS



	Co	Ma	Co	Ma	Co	Ma	Co	Ma	Co	Ma	Co	Ma	Co	Ma
	Ca		P		Mg		S		Na		K		Cl	
Systematic effects														
Parity Order		*		*	*	*		*	*		*	*	*	*
DIM	*		*		*		*		*			*	*	
MY		*	*	*		*		*	*	*	*	*		*
Fat	*	*			*	*	*		*	*				*
Casein	*		*	*		*	*	*	*	*				
Lactose		*			*		*	*	*		*	*	*	*
pH		*		*		*	*	*		*		*		*
SCS ¹			*	*	*				*	*		*	*	*
Random Effects														
Direct additive genetic effects	*	*	*	*	*	*	*	*	*	*	*	*	*	*
Sampling	*	*	*	*	*	*		*	*	*		*	*	*
Residual	*	*	*	*	*	*	*	*	*	*	*	*	*	*

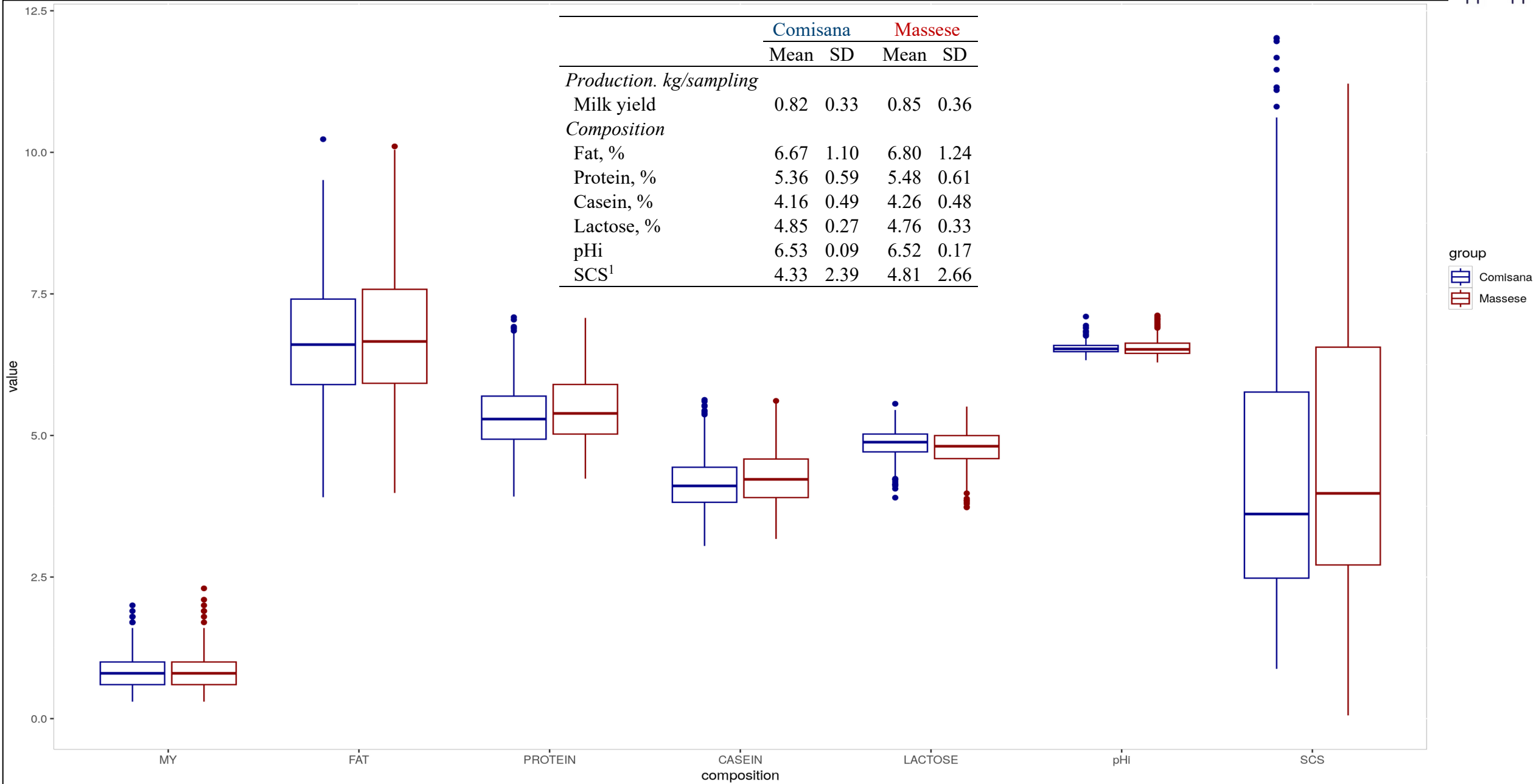
Co: Comisana

Ma: Massese

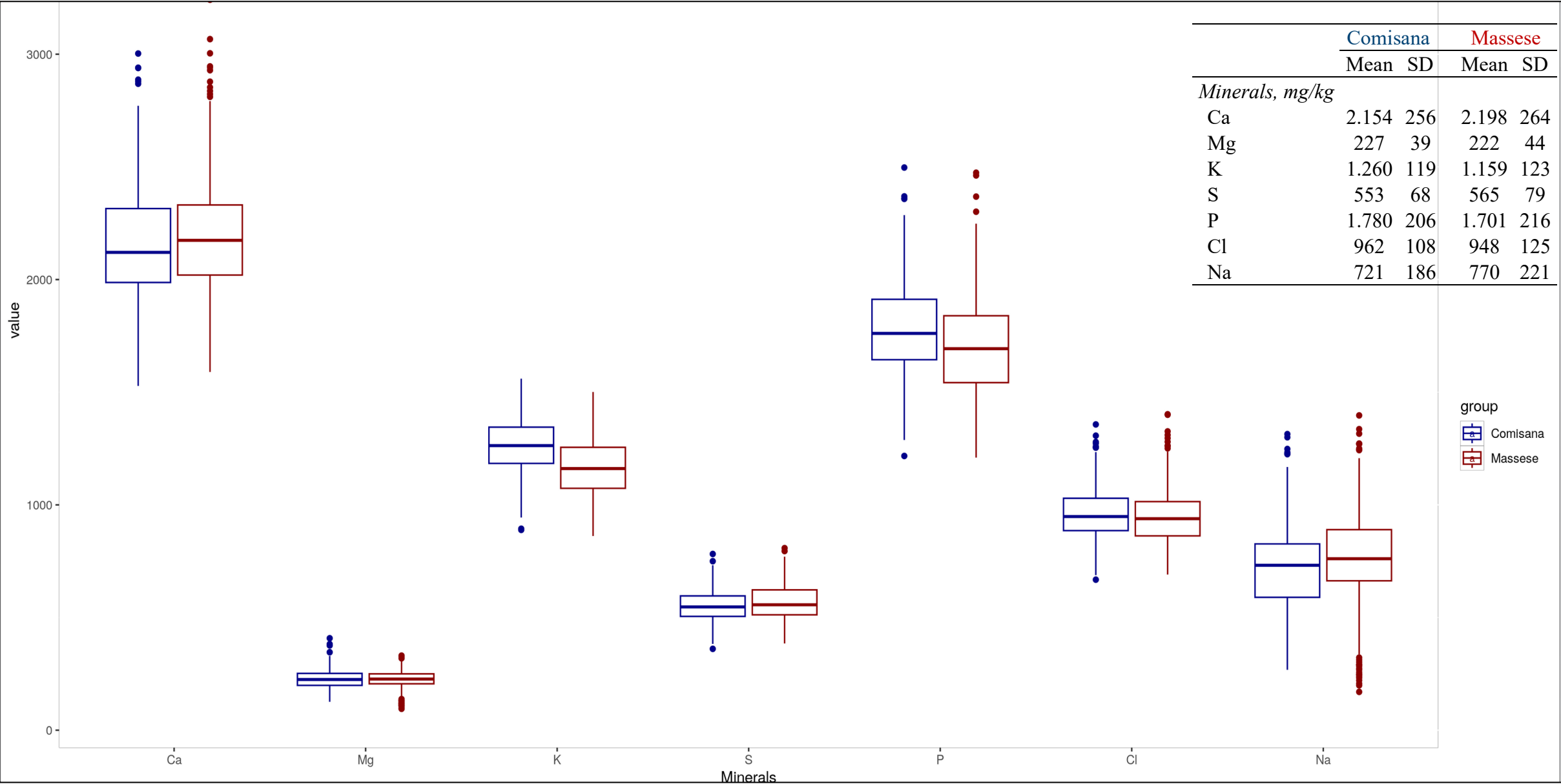
*p > 0.05

The specific significant effects on individual minerals were not exactly the same!!!

RESULTS



RESULTS



RESULTS



	Comisana	Massese
	$h^2 + SE$	$h^2 + SE$
<i>Production, kg/sampling</i>		
Milk yield	0.14 ± 0.10	0.29 ± 0.14
<i>Composition</i>		
Fat, %	0.22 ± 0.11	0.15 ± 0.10
Protein, %	0.10 ± 0.06	0.34 ± 0.14
Casein, %	0.30 ± 0.11	0.28 ± 0.14
Lactose, %	0.31 ± 0.10	0.29 ± 0.13
SCS ¹	0.081 ± 0.06	0.28 ± 0.12

	Comisana	Massese
	$h^2 + SE$	$h^2 + SE$
<i>Minerals, mg/kg</i>		
Ca	0.45 ± 0.11	0.41 ± 0.17
P	0.49 ± 0.12	0.17 ± 0.11
Mg	0.25 ± 0.12	0.16 ± 0.11
S	0.34 ± 0.10	0.14 ± 0.11
K	0.40 ± 0.11	0.28 ± 0.12
Na	0.29 ± 0.12	0.14 ± 0.14
Cl	0.42 ± 0.10	0.17 ± 0.12

- ✓ Differences highlight the need for breed-specific investigations to characterize the complex relationships between milk composition and mineral content
- ✓ The synthesis of the minerals in milk depend several factors that include the diet and nutrition, stress and illness, hormonal levels, but also genetic factors that can be determinate the absorption, transport and metabolism

Heritability reported for composition milk sheep



Higher variability of heritability estimates in Sheep Breeds !!!

Table 2

Pooled estimates of heritability from meta-analysis in dairy sheep.

Trait ¹	Pooled h^2 ($\pm$ SE)	$\longrightarrow$	Min ² h^2	Max ³ h^2	N obs ⁴	N studies	I^2 between ⁵	I^2 within ⁶
MY	0.24 $\pm$ 0.02	$\longrightarrow$	0.06	0.46	29	19	83.98	15.03
FY	0.21 $\pm$ 0.06	$\longrightarrow$	0.14	0.28	7	5	0.26	96.7
PY	0.22 $\pm$ 0.04	$\longrightarrow$	0.12	0.30	7	5	0.64	95.77
FC	0.28 $\pm$ 0.11		0.04	0.68	16	12	54.79	45.13
PC	0.33 $\pm$ 0.07		0.10	0.77	25	17	62.58	37.2
SCS	0.13 $\pm$ 0.02		0.03	0.27	29	22	52.64	44.26
FEC	0.14 $\pm$ 0.04		0.09	0.35	6	3	0	57.97

SCS – somatic cell score, FEC – faecal egg count.

¹ Trait: MY – milk yield, FY – fat yield, PY – protein yield, FC – fat content, PC – protein content.

² Minimum h^2 from individual studies included in meta-analysis.

³ Maximum h^2 from individual studies included in meta-analysis.

⁴ Number of observations used in meta-analysis.

⁵ Heterogeneity between clusters.

⁶ Heterogeneity within clusters.



Genetic (co)variances between milk mineral concentration and chemical composition in lactating Holstein-Friesian dairy cows

G. Visentin ¹, G. Niero ¹✉, D.P. Berry ², A. Costa ¹, M. Cassandro ¹, M. De Marchi ¹, M. Penasa ¹

Table 1. Mean, genetic standard deviation (s_g), heritability (h^2 ; standard error in parentheses), repeatability (t ; standard error in parentheses) and coefficient of genetic variation (CV_g) estimated using the repeatability animal model for calcium (Ca), potassium (K), magnesium (Mg), sodium (Na) and phosphorus (P) concentration, chemical composition, pH and somatic cell score (SCS) of cow milk

Traits	<i>n</i>	Mean	σ_g	h^2	<i>t</i>	CV_g (%)
Ca (mg/kg)	12 208	1295.29	85.58	0.54 (0.04)	0.66 (0.01)	6.61
K (mg/kg)	12 165	1684.48	56.09	0.19 (0.03)	0.25 (0.01)	3.33
Mg (mg/kg)	12 198	136.89	7.88	0.21 (0.03)	0.27 (0.01)	5.76
Na (mg/kg)	12 223	439.06	21.56	0.24 (0.04)	0.39 (0.02)	4.91
P (mg/kg)	12 220	995.04	62.25	0.42 (0.04)	0.53 (0.01)	6.23

RESULTS

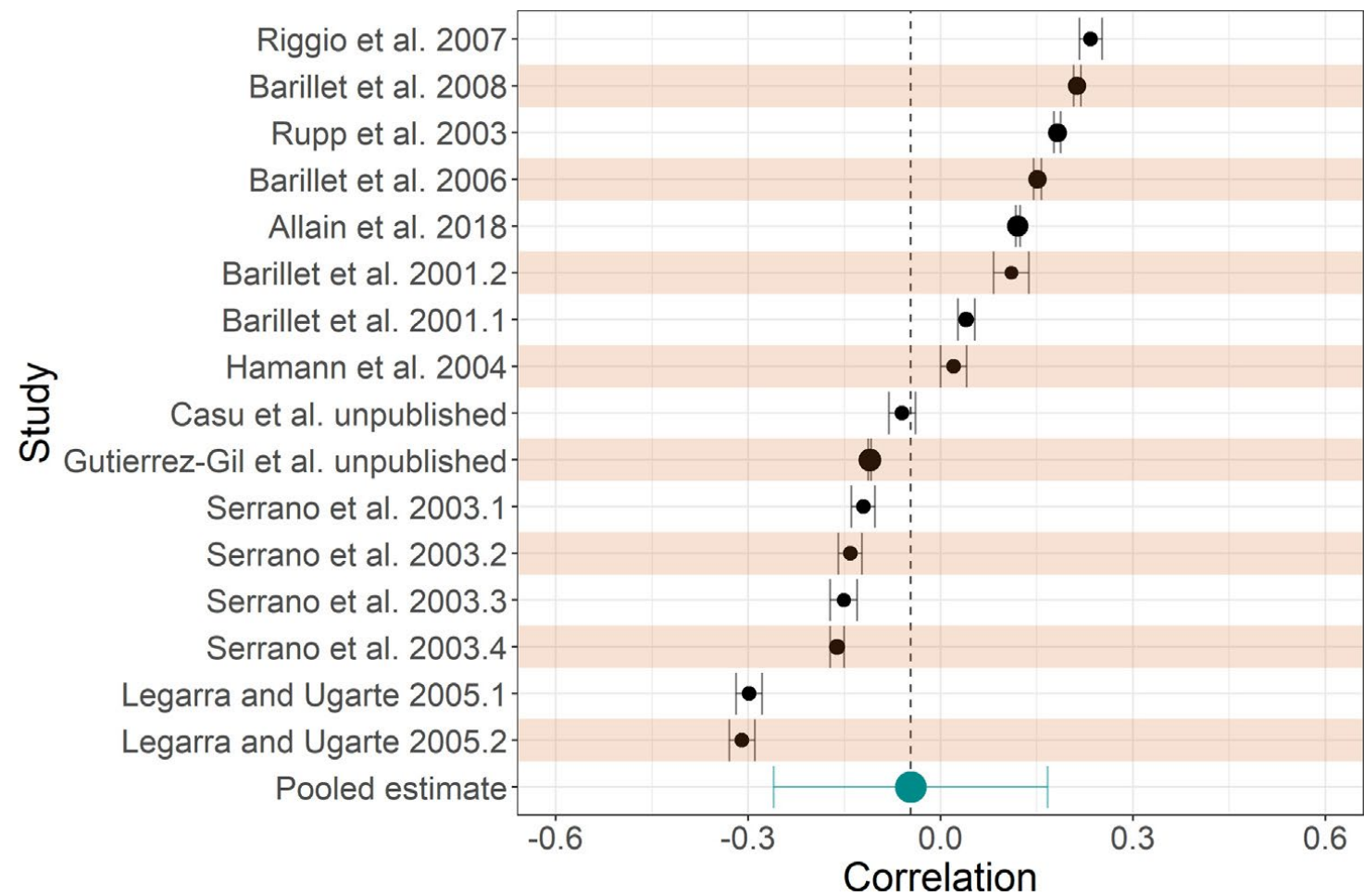


Genetic correlation matrix among milk yield (MY), composition and minerals in Comisana (above the diagonal) and Massese (below the diagonal) breeds.

		<i>Comissana</i>													
		<i>Composition %</i>						<i>Minerals, mg/kg</i>							
		MY	Fat	Protein	Casein	Lactose	<u>SCS¹</u>	Ca	P	Mg	S	K	Na	Cl	
<i>Massese</i>	<i>Composition</i>	MY	*	0.55	0.35	0.10	-0.07	0.51	-0.61	-0.19	-0.36	0.30	-0.09	0.04	-0.65
		Fat, %	0.38	*	0.51	NC	-0.89	NC	0.08	0.16	0.07	-0.10	-0.68	NC	0.13
		Protein, %	NC	0.21	*	NC	0.32	-0.66	0.33	0.33	0.33	NC	-0.51	0.22	-0.44
		Casein, %	NC	NC	NC	*	-0.13	-0.47	0.20	0.26	0.25	NC	-0.20	0.21	-0.41
		Lactose, %	NC	NC	NC	0.28	*	NC	0.08	-0.22	-0.40	-0.10	-0.89	NC	0.64
		pH													
		SCS ¹	NC	NC	- NC	0.19	NC	*	-0.22	-0.50	NC	-0.19	NC	NC	NC
	<i>Minerals, mg/kg</i>	Ca	-0.35	-0.51	0.071	0.33	0.49	-0.08	*	0.66	0.67	0.30	-0.30	0.62	0.27
		P	-0.40	-0.48	-0.038	0.016	-0.045	-0.32	0.25	*	NC	0.35	0.05	0.51	0.04
		Mg	-0.22	-0.73	-0.37	0.013	-0.26	0.23	-0.39	0.24	*	0.47	-0.023	0.74	0.10
		S	-0.34	-0.54	0.046	0.17	-0.006	-0.13	-0.33	0.66	0.57	*	-0.37	0.43	-0.24
		K	-0.11	-0.59	-0.42	-0.01	-0.78	NC	-0.10	0.010	0.34	-0.037	*	-0.31	0.52
		Na	-0.80	-0.53	-0.36	-0.38	-0.79	NC	-0.53	0.61	0.79	NC	0.35	*	0.11
		Cl	-0.30	0.03	-0.60	-0.60	-0.70	0.88	-0.74	-0.74	0.039	-0.18	0.01	0.26	*

¹SCS = $\log_2(\text{somatic cell count} \times 10^{-5}) + 3$

Genetic correlation reported for SCS and Milk yield



Mucha et al., 2022

Fig. 2. Forest plot showing genetic correlation estimates between somatic cell score and milk yield in dairy sheep.

Conclusions



- These differences highlight the need for breed-specific investigations to characterize the complex relationships between milk composition and mineral content. The synthesis of the minerals in milk depend several factors that include the diet and nutrition, stress and illness, hormonal levels, but also genetic factors that can be determinate the absorption, transport and metabolism

Heritability

- Composition: Low to medium heritability for both breeds
- Minerals: Medium to high heritability for both breeds

- Genetic Correlation

Generally negative genetic correlations between minerals and composition but highly variable

- Need to increase the number of records to more accurate estimates
- Need of the increase research in these area to obtain better insight about the relationship between this traits → Breeding programs



Thank you for your attention