



Effects of nonsynonymous SNPs at *GH2-N* and *GHR* genes on coagulation properties of Assaf ewes' milk

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Introduction



Milk production potential is a function of the number of mammary epithelial cells

Lactation performance depends on:

- Mammary cell proliferation (or decrease apoptosis)
- Structural and biochemical differentiation of mammary epithelium
- Synthesis and secretion of milk components

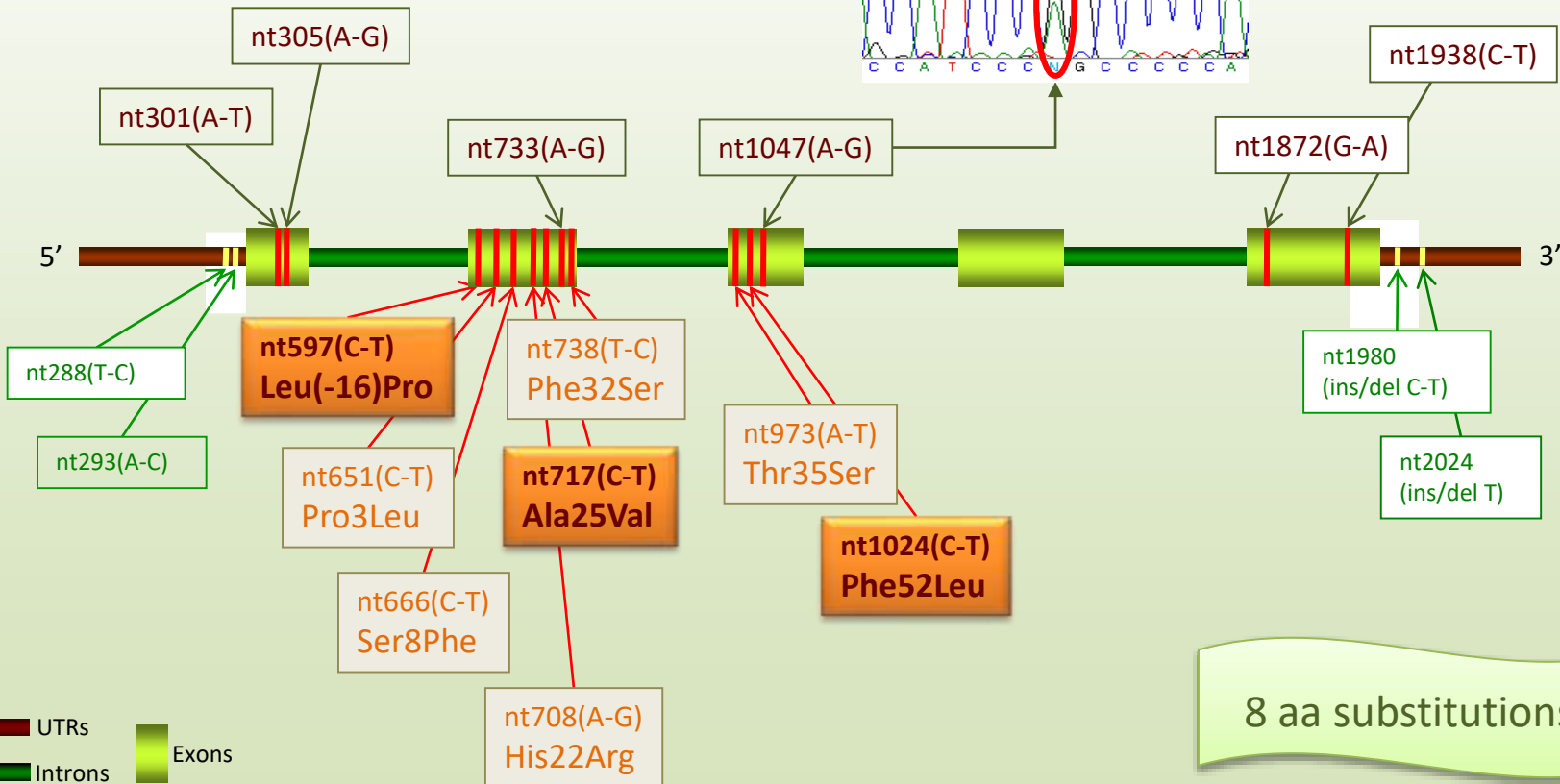
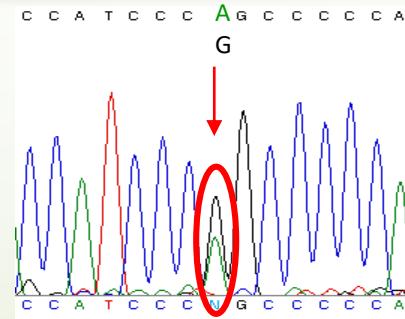
Various hormones are involved in these processes

Introduction

Growth hormone gene (*GH2-M*)



- Polymorphisms at the ***GH2-N*** copy



8 aa substitutions!

Objectives



To uncover SNPs in ovine somatotrophic axis associated with high yielding dairy ewes.

- Genotype SNPs in growth hormone copy *GH2-N* and *GHR* genes in Assaf ewes
- to identify candidate causative mutations for milk production and composition, to be used in Marker-Assisted Selection programs by dairy ewes breeders

Aiming at an increase in the sheep's breeding value for milk traits, and the flocks' profitability.

Methodology



➤ Genotyping and phenotyping



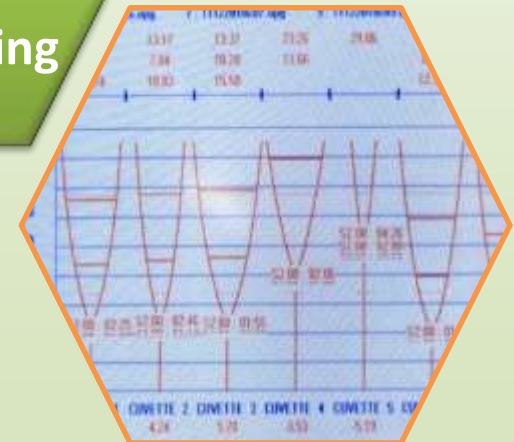
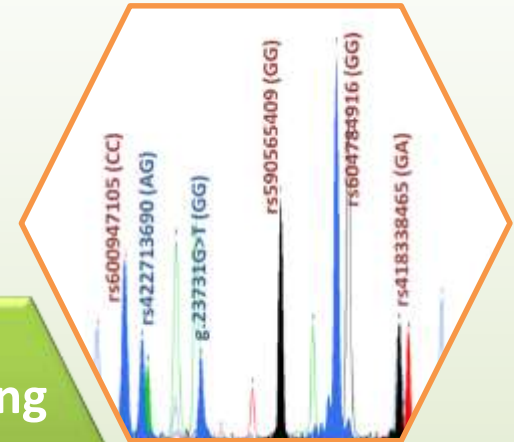
430 ewes

Blood
collection
and DNA
extraction

Milk
samples
collection

184 ewes

Genotyping
and
phenotyping



Methodology



- 1. Six SNPs genotyped by SnapShot analysis**
- 2. Milk production (monthly)**
- 3. Milk composition analysis by Milk-o-Scan:**
 - ❖ **Fat, protein, lactose, total solids and fat free total solids content (monthly)**
- 4. Coagulation properties evaluated at 1st and 3th month of lactation by Optigraph:**
 - ❖ **Clotting time (R), gel firmness after 20 min. (A20) and after a 2R (AR) period, and rate of firming (OK20)**



4. Statistical analysis

MIXED model procedure from SAS®

Y_{ijklmn}

$$= \mu + NLact_i + SNP_j + Contr_k + \beta_1(x_{ijkl} - \bar{x}) + \beta_2(x_{ijkl} - \bar{x})^2 + Ewe_{ijklm} + \varepsilon_{ijklmn}$$

$NLact_i$ – effect of the lactation number i (1 to 6)

SNP_j – effect of the genotype for each SNP j

$Contr_k$ – effect of lactation month k (1 to 5)

$\beta_1(x_{ijkl} - \bar{x})$ – linear and quadratic effect of the age of the ewe at lambing

Ewe_{ijklm} - random effect of the ewe $ijklm$

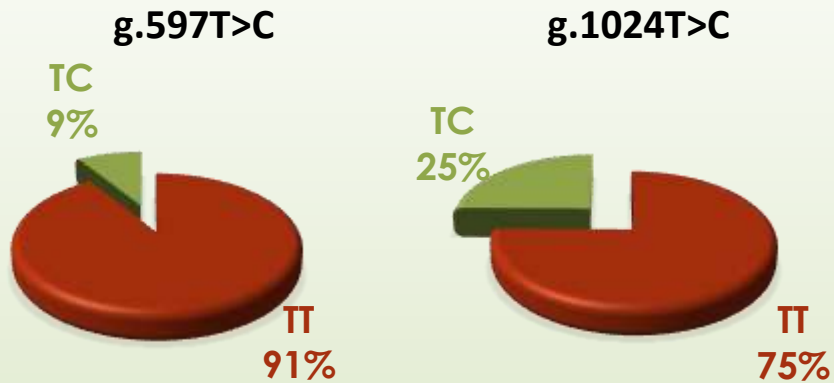
ε_{ijklmn} - random error

Results

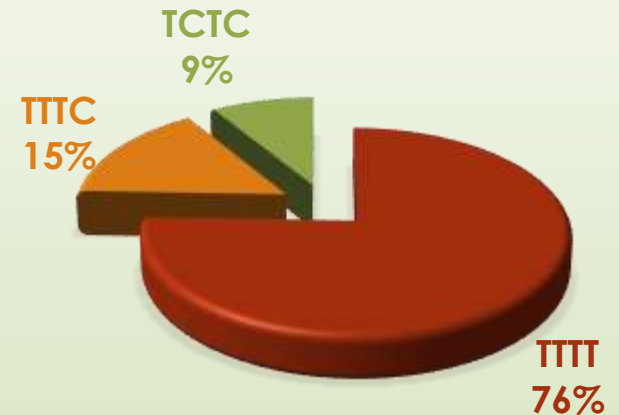
GH2-N frequencies



Genotypes frequencies



Haplotypes frequencies



	SNP*	aa**	Alleles ^b		P _{HWE}
GH2-N	g.597T>C	Leu11Pro	T (0.956)	C (0.044)	***
	g.717C>T	Ala51Val	C (1.000)		-
	g.1024T>C	Phe78Leu	T (0.876)	C (0.124)	***

* GenBank accession number X12546;

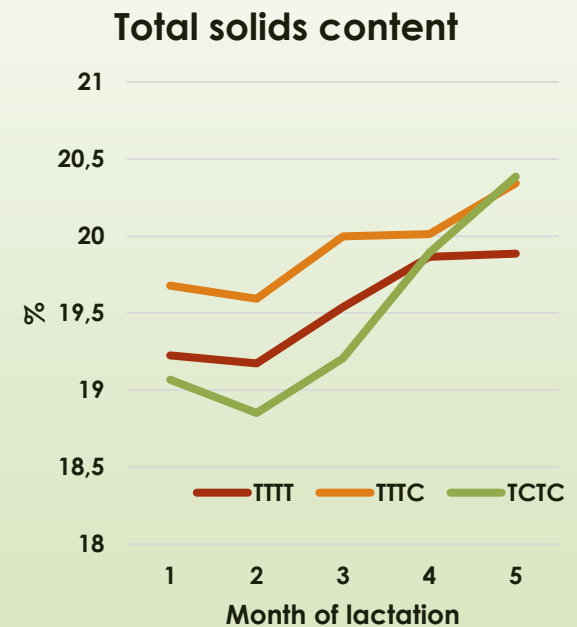
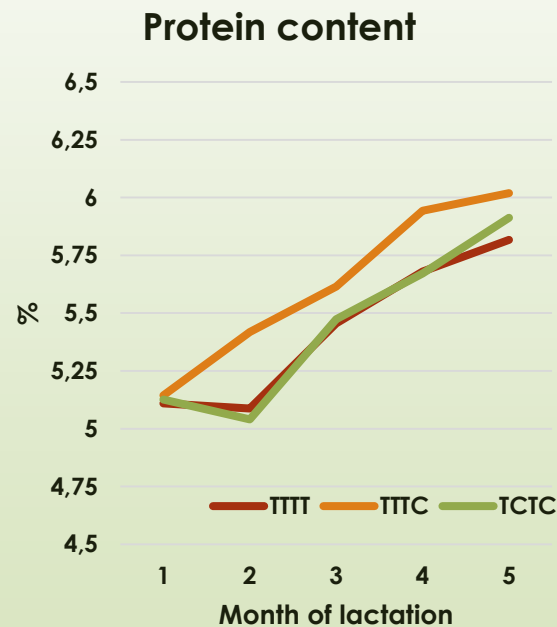
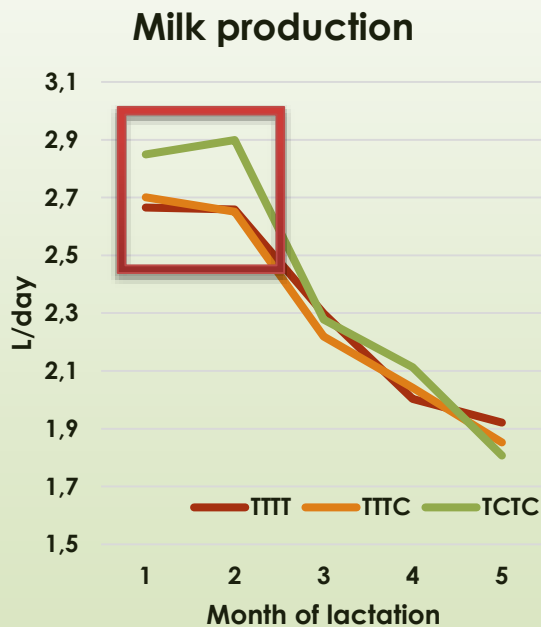
** GenPept accession number P67930

Results

GH2-N vs milk traits



➤ *GH2-N* SNPs had no effects on milk production and composition



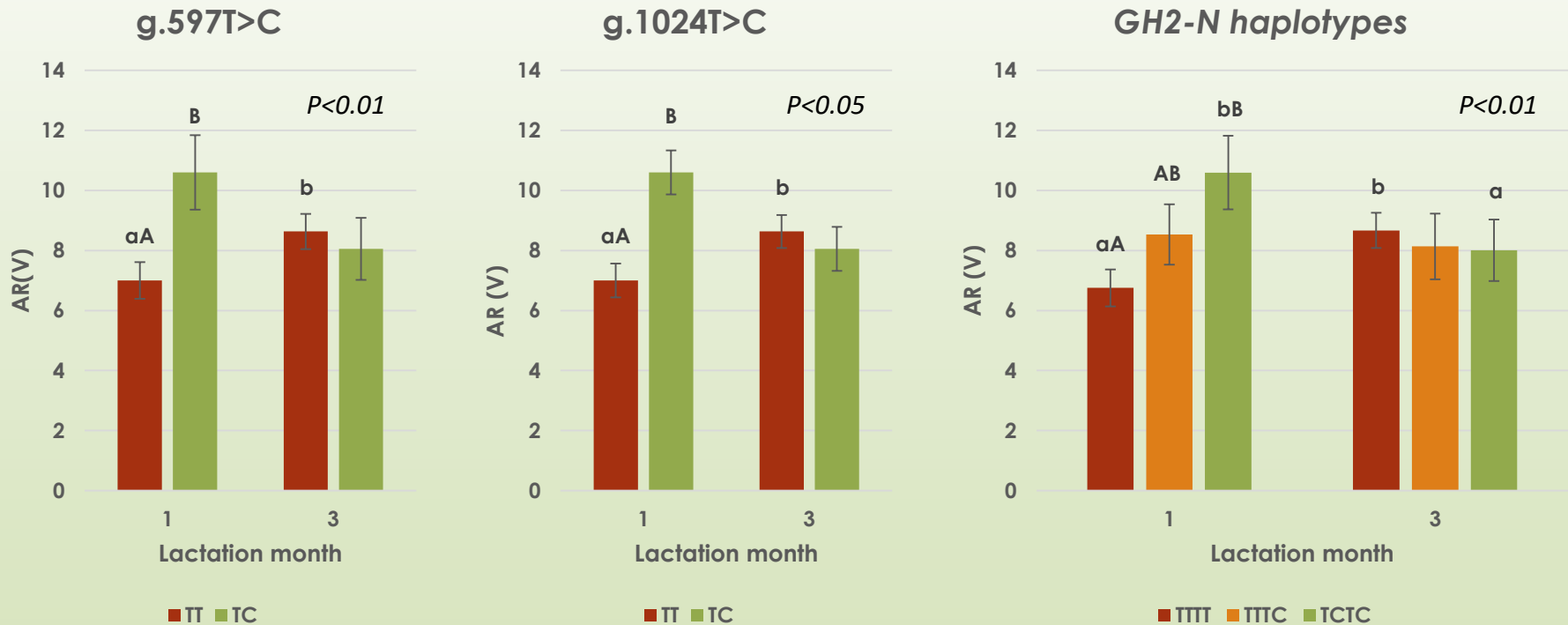
Note: haplotypes results

Results

GH2-N vs milk traits



➤ SNPs affected gel firmness after a 2x clotting time (AR) period



a, b – different lowercase letters correspond to significant differences between lactation month within genotypes ($P < 0.05$)
A, B – different capital letters correspond to significant differences between genotypes within lactation month ($P < 0.05$)

Results

GHR frequencies



➤ Genotypes and alleles frequencies

SNP*		aa	Genotypes ^a			Alleles ^b		P _{HWE}
GHR	rs1086611503	Ser380Pro	TT (0.909)	TC (0.091)		T (0.955)	C (0.045)	***
	rs595567866	Glu392Lys	GG (0.905)	GA (0.093)	AA (0.002)	G (0.951)	A (0.049)	***
	rs597181420	Ala529Thr	CC (0.910)	CT (0.083)	TT (0.007)	C (0.952)	T (0.048)	***

* dbSNPs;

** GenPept accession number Q28575



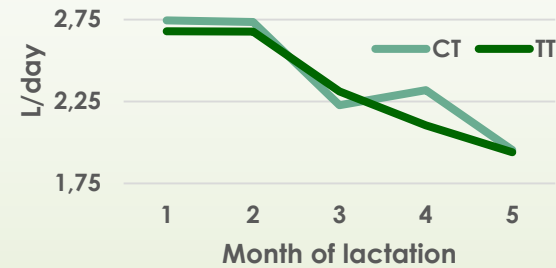
**Nine haplotypes
identified!**

Results

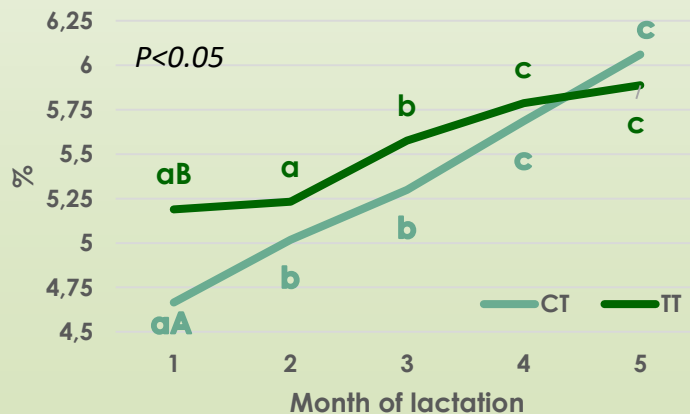
GHR - rs1086611503



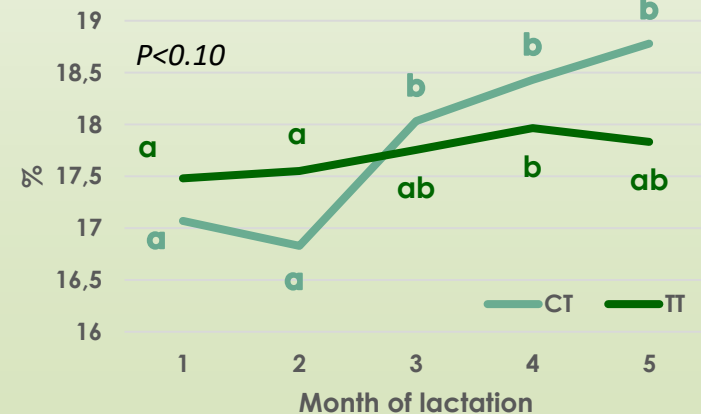
Milk production



Protein content



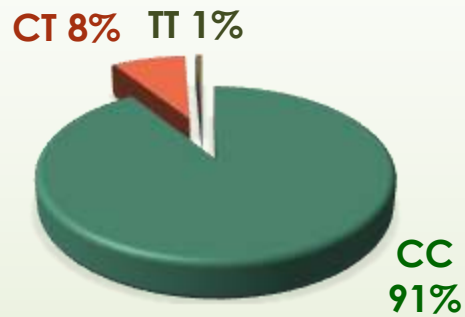
Total solids content



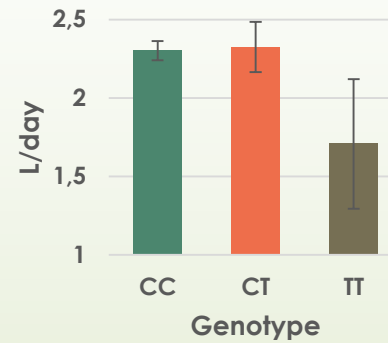
a, b – different lowercase letters correspond to significant differences between lactation month within genotypes ($P < 0.05$)
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Results

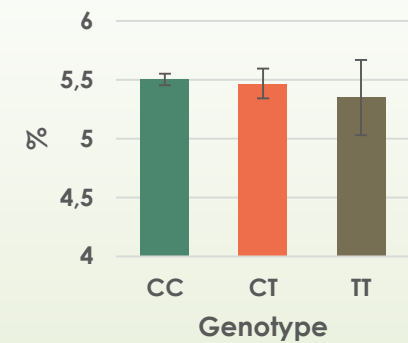
GHR - rs597181420



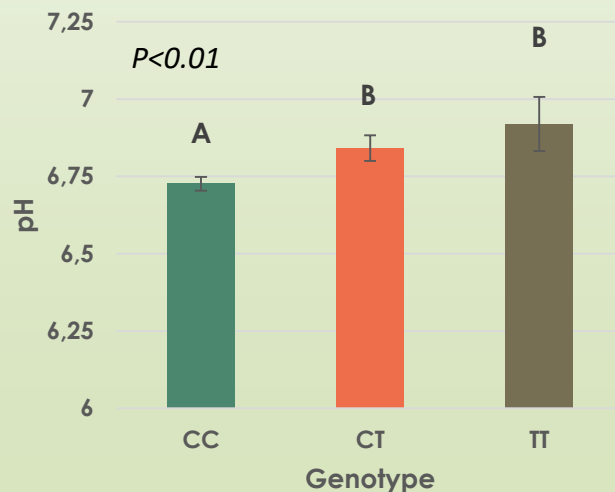
Milk production



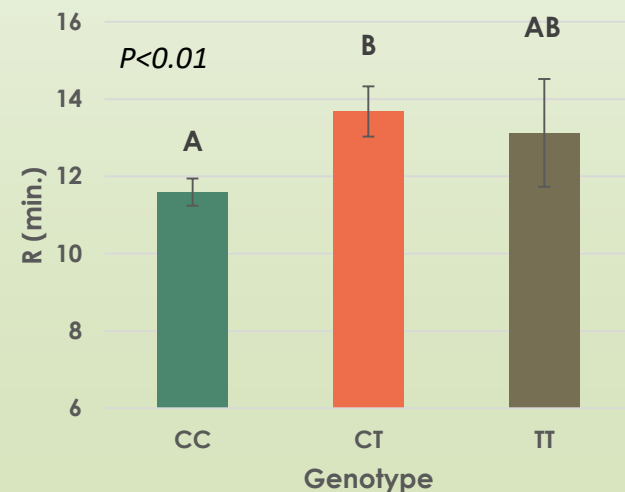
Protein content



pH



Clotting time (R)



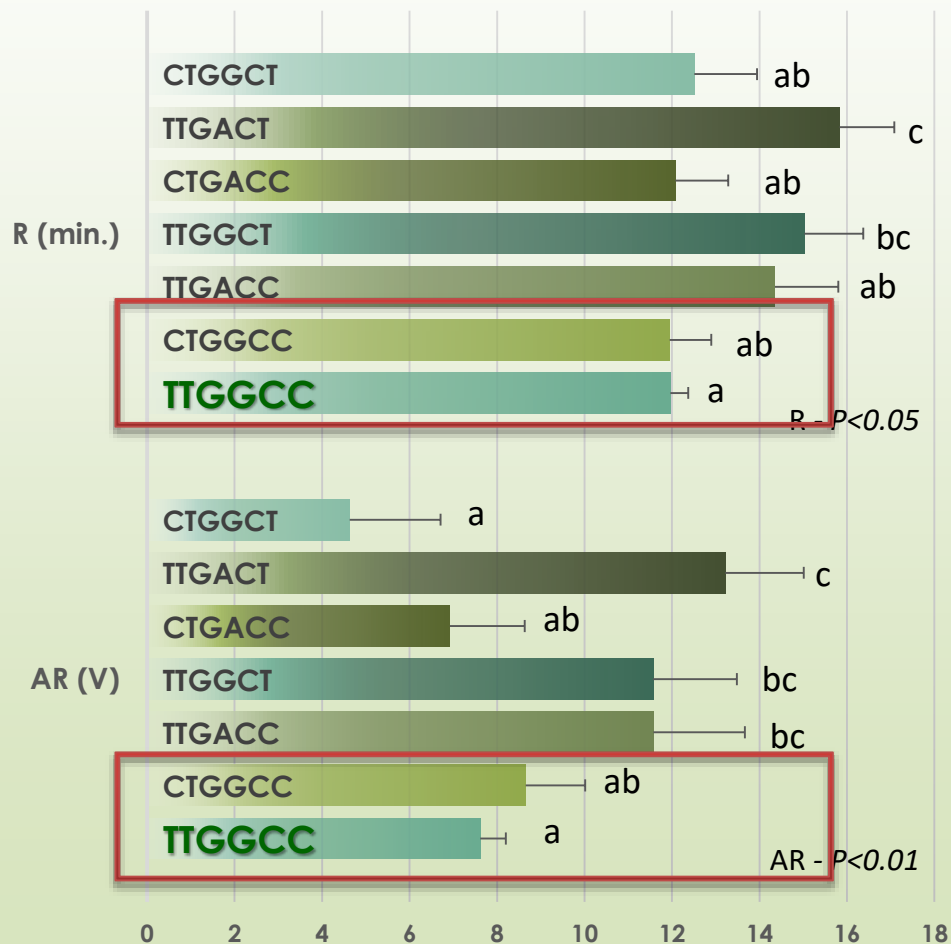
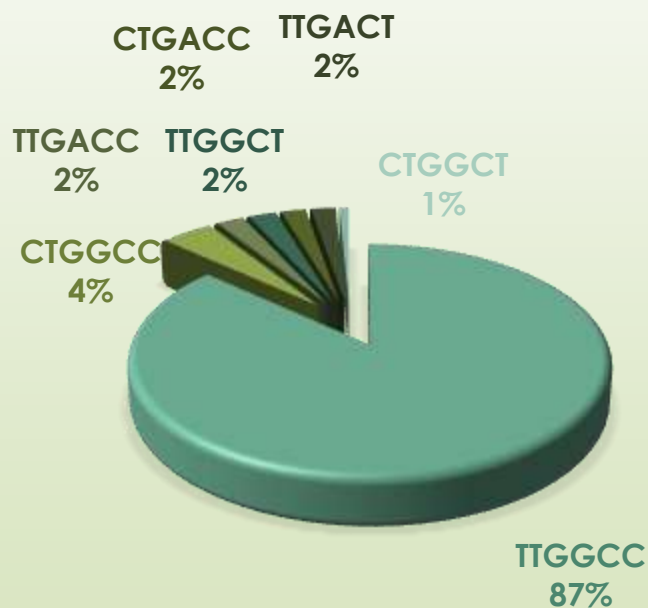
A, B – different capital letters correspond to significant differences between genotypes ($P < 0.05$)

Results

GHR haplotypes vs milk traits



Haplotypes frequencies



a, b – different capital letters correspond to significant differences between haplotypes ($P < 0.05$)

Conclusions



1. Studied SNPs at *GH2-N* and *GHR* genes were
 - ✓ Polymorphic – all but *GH2-N* g.717C>T
 - ✓ not associated with milk production traits
 - ✓ associated with milk quality and coagulation parameters

2. *GH2-N* SNPs and haplotypes ➡ gel firmness (AR)

3. *GHR* SNPs:

- ✓ rs1086611503 ➡ milk protein and total solids content throughout lactation
- ✓ rs597181420 ➡ pH and clotting time (R)
- ✓ Haplotypes ➡ clotting time (R) and gel firmness (AR)



Ongoing SNP genotyping:

- ❖ *GH2-N* and *GH2-Z* genes copies and GHR.

- ❖ Other

 -

Ongoing

- ❖ Colle

propo

Selection of the panel of SNPs that best allows to estimate breeding value of the ewes and rams to be bred for the production of quality milk

Associate these SNP with milk production traits:

- ❖ Twenty years of records for milk production traits.

Acknowledgment



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Production and milk
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OVine breed.

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<https://projects.iniav.pt/genprov/>



Thanks for your attention!

