



# 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**



# 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



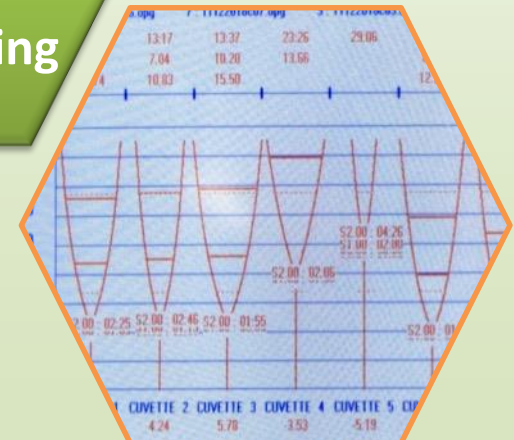
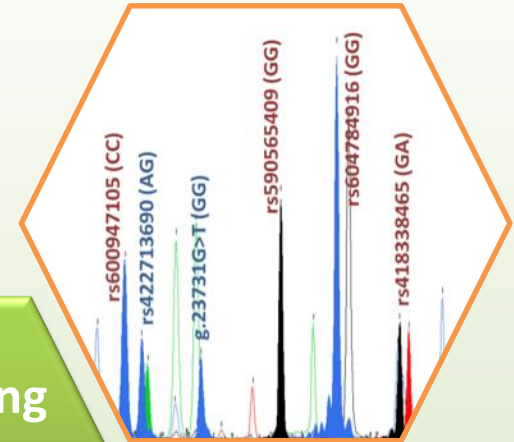
**430 ewes**

**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 1<sup>st</sup> and 3<sup>th</sup> 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  $ijklm$

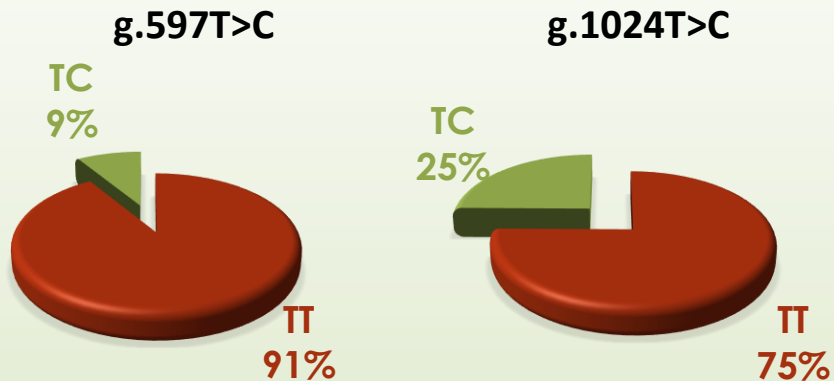
$\varepsilon_{ijklmn}$  - random error

# Results

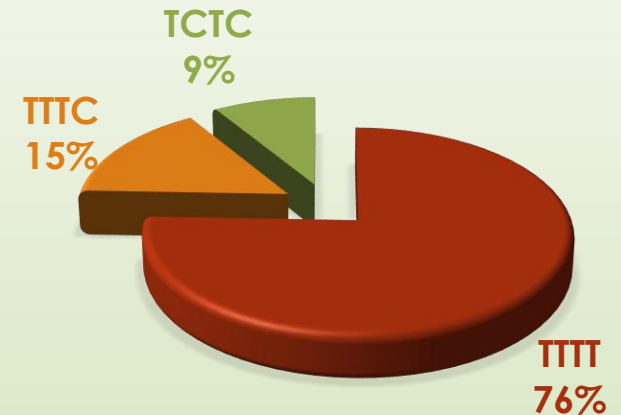
## *GH2-N* frequencies



### Genotypes frequencies



### Haplotypes frequencies



	SNP*	aa**	Alleles <sup>b</sup>		P <sub>HWE</sub>
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

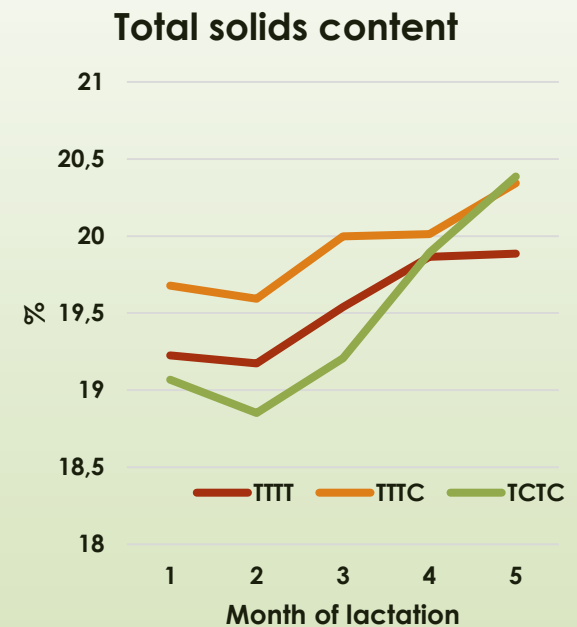
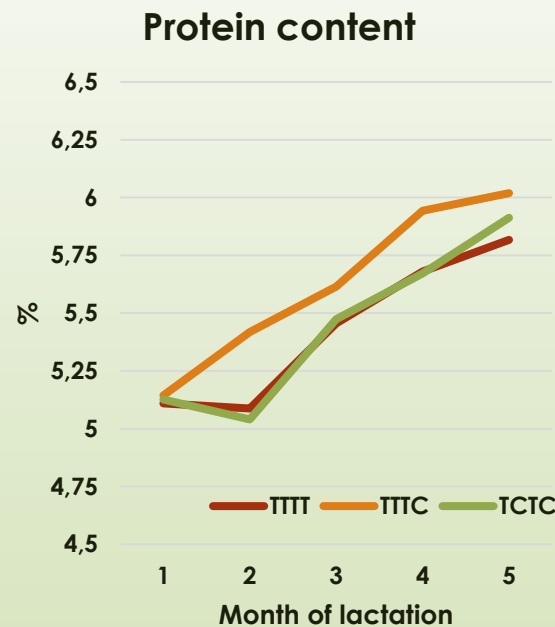
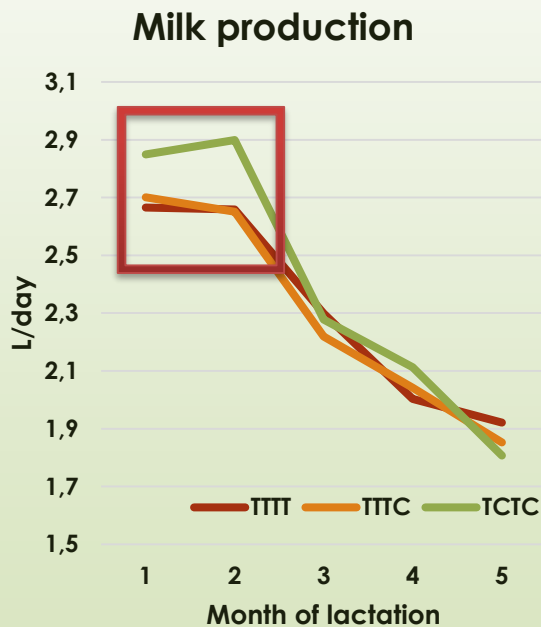


# Results

## *GH2-N* vs milk traits



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



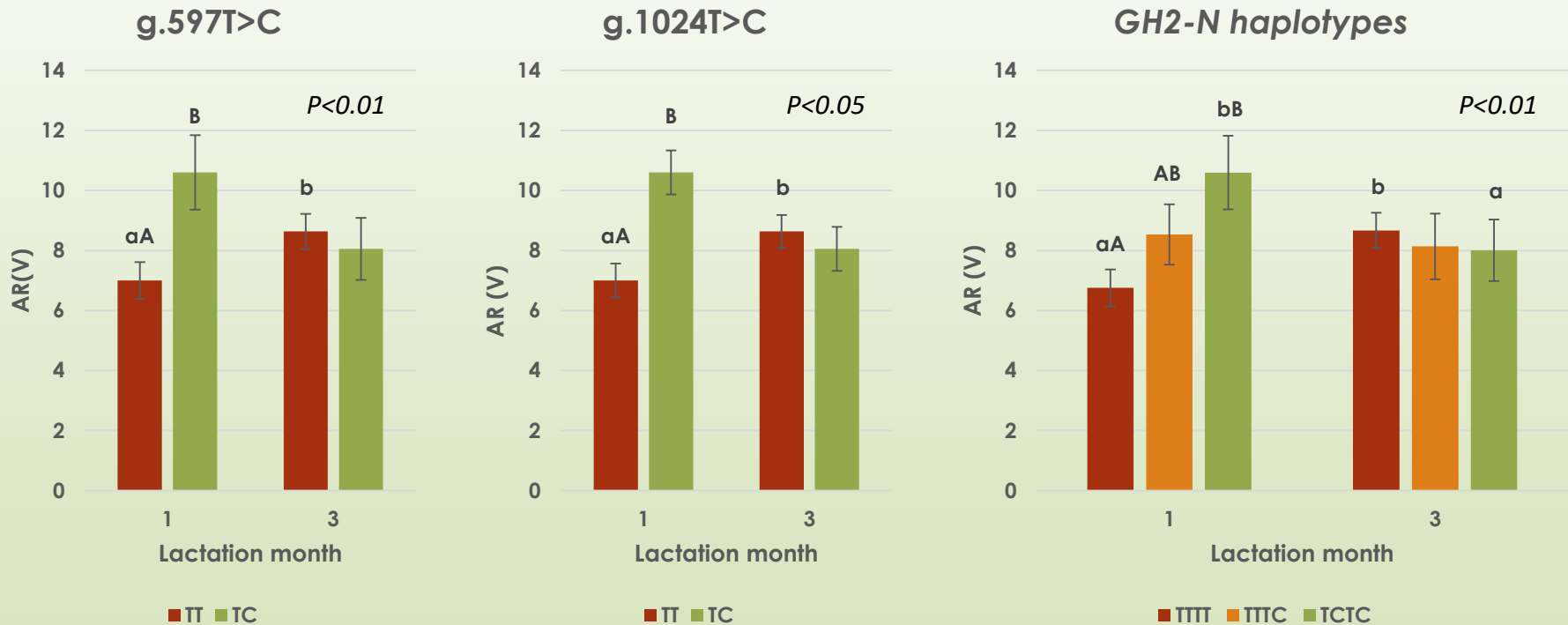
Note: haplotypes results

# 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 <sup>a</sup>			Alleles <sup>b</sup>		P <sub>HWE</sub>
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;

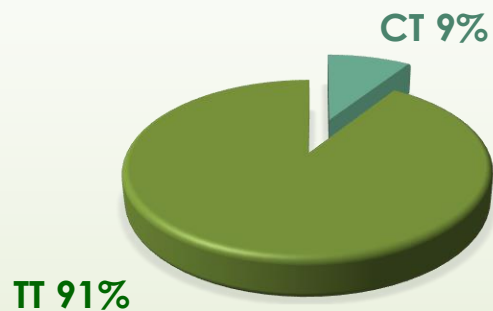
\*\* GenPept accession number Q28575



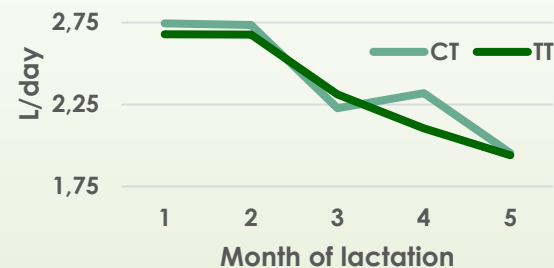
**Nine haplotypes  
identified!**

# Results

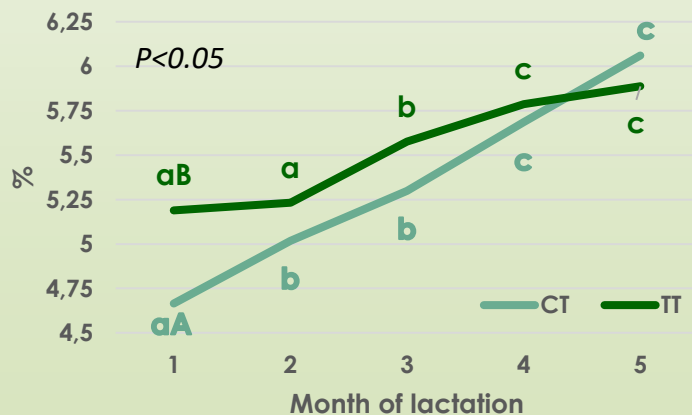
## GHR - rs1086611503



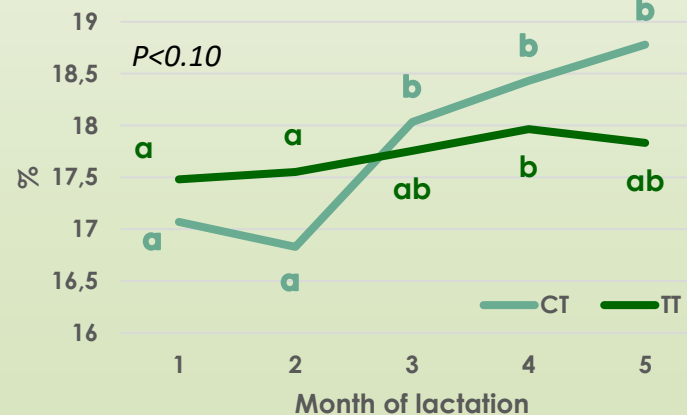
### Milk production



### Protein content



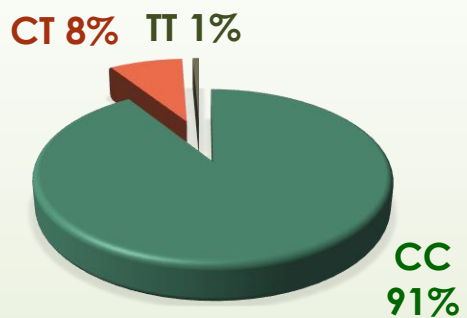
### Total solids content



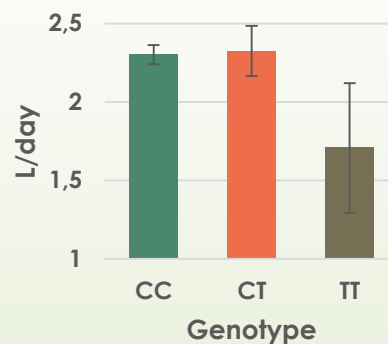
a, b – different lowercase letters correspond to significant differences between lactation month within genotypes ( $P < 0.05$ )  
A, B – different capital letters correspond to significant differences between genotypes within lactation month ( $P < 0.05$ )

# Results

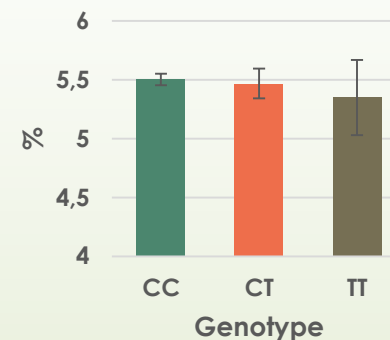
## GHR - rs597181420



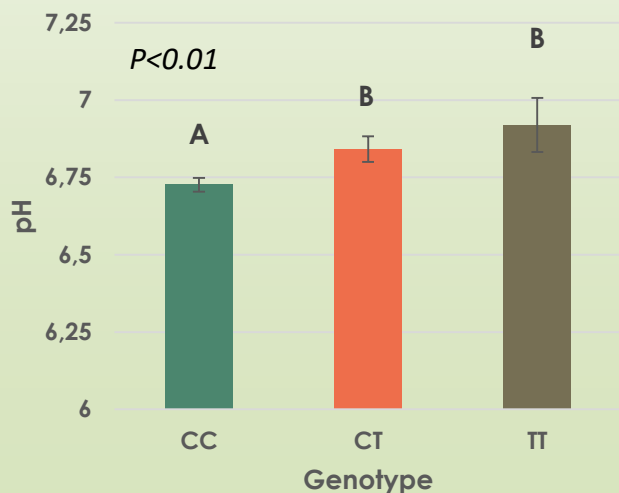
### Milk production



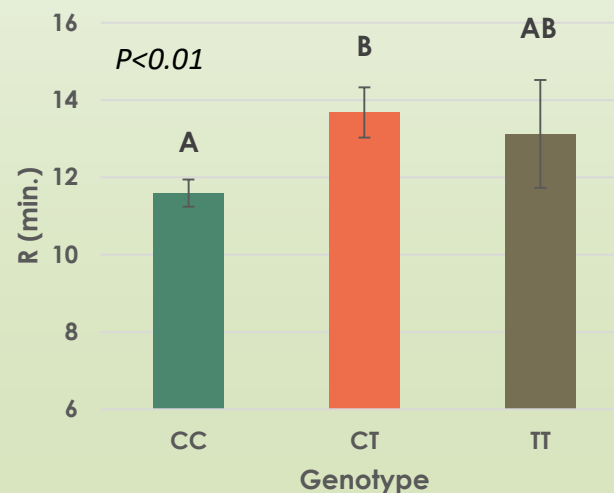
### Protein content



### pH



### Clotting time (R)



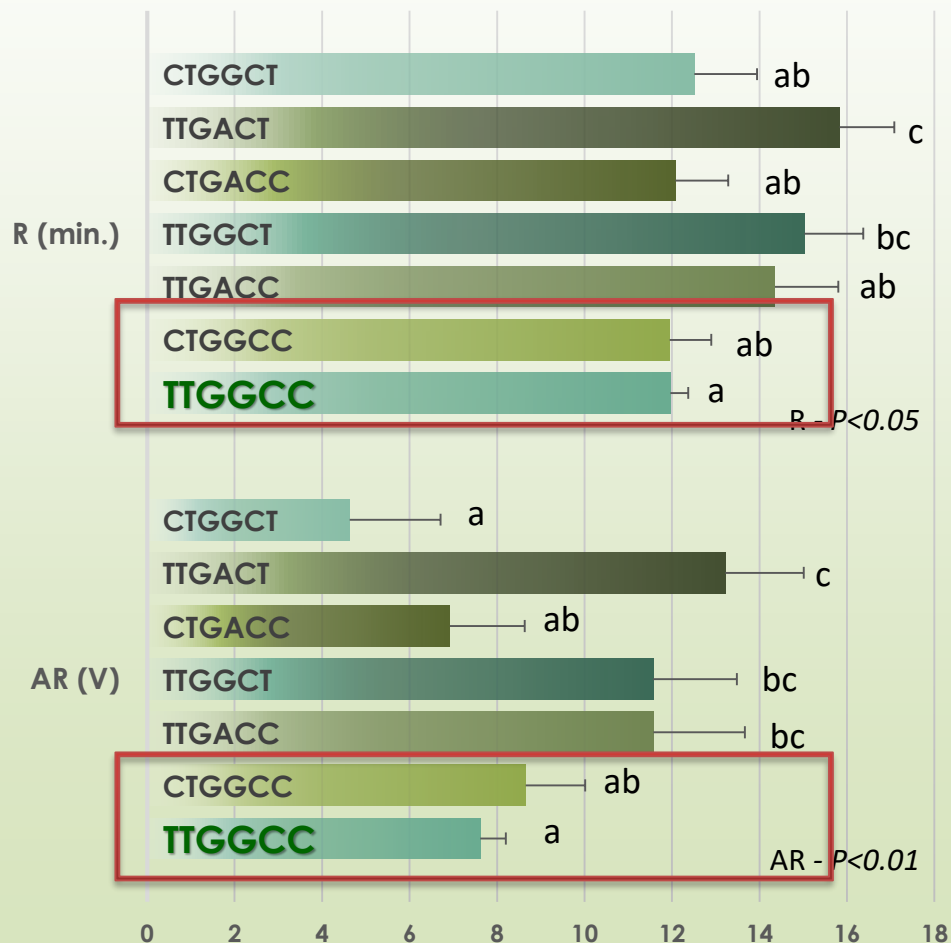
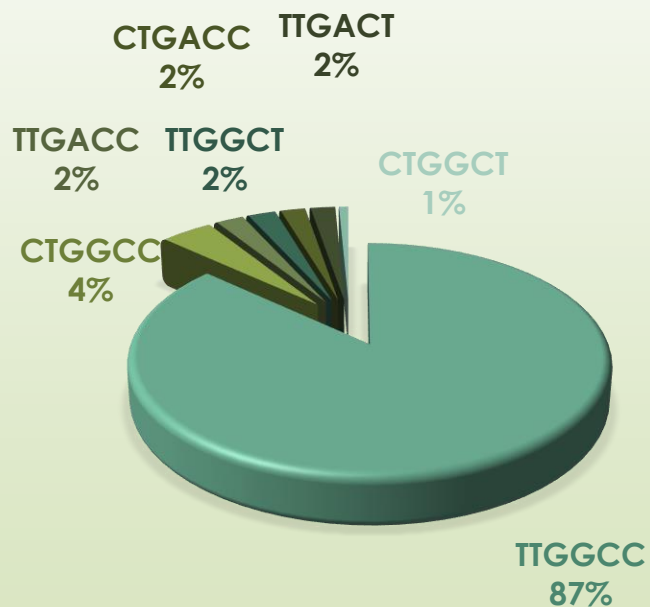
A, B – different capital letters correspond to significant differences between genotypes ( $P < 0.05$ )

# 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)

# Perspectives



## 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



Gene markers for  
Production and milk  
quality in Assaf  
OVine breed.

Project

ALT20-03-0145-FEDER-000019

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

Co-financed by



# Thanks for your attention!

