

# Identification of established genetic variants associated with milk traits

Michael Orford<sup>1</sup>, Ouranios Tzamaloukas<sup>1</sup>, Christakis Papachristoforou<sup>1</sup>, Georgia Hadjipavlou<sup>2</sup>, Alkis Koumas<sup>2</sup> & Despoina Miltiadou<sup>1</sup>

<sup>1</sup>Department of Agricultural Sciences, Biotechnology and Food Science, Cyprus University of Technology, P. O. Box 50329, 3603, Lemesos, Cyprus; <sup>2</sup>Agricultural Research Institute, P.O. Box 2016, 1516 Lefkosia, Cyprus

### THE MESSAGE

- \* The presence of  $\beta$ -lactoglobulin ( $\beta$ -LG) genotypes and single nucleotide polymorphisms (SNPs) known to affect bovine milk traits was examined in small ruminant breeds of Cyprus
- Major intra-species genotypic differences were found between the Cyprus fat-tailed and Chios sheep breeds at the β-LG locus
- The β-LG genotype seems fixed in Cyprus caprine breeds

### INTRODUCTION

- ➤ Dissecting the genetic basis of milk production traits would be of great benefit to the small ruminant dairy industry
- Genetic variants at several loci that associate with milk production traits have been mainly identified in dairy cattle populations

#### AIM:

Investigate the presence of  $\beta$ -lactoglobulin ( $\beta$ -LG) variants, growth hormone receptor (GHR) and DGAT1 single nucleotide polymorphisms (SNPs) in the four major purebred Cyprus sheep and goat breeds

## METHODS

**Sample:** Genomic DNA from a total of 366 animals

- -246 Chios and 40 Cyprus fat-tailed sheep
- -40 Damascus and 40 Machaeras Goats

The animals were bred at the Agricultural Research Institute experimental station and at private farms

Genotyping (PCR, gel electrophoresis and DNA sequencing):

- A)  $\beta$ -lactoglobulin most common variants (A & B)
- B) Growth Hormone Receptor (GHR) variant F279Y
- C) DGAT1 K232A SNP

#### AKNOWLEDGEMENTS

- ➤ This work was supported by the Cyprus Research Promotion Foundation and the Cyprus University of Technology
- ➤ We thank the technical staff of the Animal Production section at the Agricultural Research Institute and the private breeders for management of the sheep and goat resource flocks

## RESULTS

Significant differences in  $\beta$ -LG allele frequencies detected for the Chios and Cyprus fat-tailed sheep

Table 1. β-LG allele and genotype frequencies in Chios and Cyprus fat-tailed sheep breeds

Breed	No. of animals	<i>β-lactoglobulin</i> Genotype (%)			Allelic Frequency			
		AA	AB	BB	A-allele	B-allele		
Chios sheep	246	56.9	38.6	4.5	0.76	0.23		
Cyprus Fat- tailed sheep	40	12.8	48.7	38.5	0.37	0.63		
Chi-square (χ²) test (2 d.f.)*		56.4 (P<0.001)						

\*Chi square test of independence between genotype frequency and sheep breed

The β-LG A allele appears to be fixed in the two Cyprus goat breeds

Table 2. β-LG allele and genotype frequencies in Damascus and Machaeras goat breeds

		<i>β-lactoglobulin</i> genotype			Allelic frequency	
Breed	No. of animals	AA	AB	BB	A-allele	B-allele
Damascus Goat	40	40	0	0	1	0
<b>Machaeras Goat</b>	40	40	0	0	1	0

The GHR F279Y and DGAT1 K232A polymorphisms were not detected in samples from the four Cyprus sheep and goat breeds studied

#### KEY POINTS

- Two genetic variants (A and B) at the  $\beta$ -LG locus were identified in animals from the two purebred Cyprus sheep breeds
- > Only the β-LG variant A was detected in the Cyprus Damascus and Machaeras goat breeds
- The Cyprus fat-tailed sheep was predominantly of the  $\beta$ -LG B type, with an unusually high frequency of the  $\beta$ -LG BB genotype compared to the Chios or other Mediterranean sheep breeds
- ➤ Polymorphisms GHR F279Y and DGAT1 K232A appear not to be present in the four major purebred sheep and goat breeds of Cyprus