

Ovine mastitis: Does early live nutrition influence immunity response in later life?





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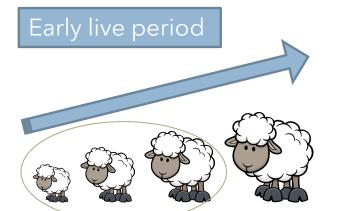
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MEGA

EARLY LIFE CONDITIONS INFLUENCING IMMUNITY STATUS



Sensitive life stage Hardship conditions



Animal fitness Immunity status

Food composition and availability



Immune system



(Fragkou et al., 2021)



(Mugabo et al., 2010)



(Schmidt et al., 2015)

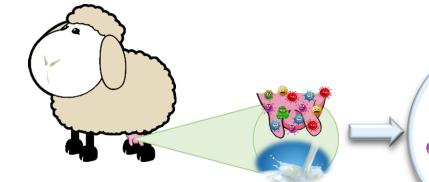




LACTATION, MASTITIS AND RNA-SEQ

Differences in immune response or immune system vulnerability can be more accused under stress situations, such as lactation in livestock animals

MASTITIS



- Inflammatory disease of the mammary gland
- Most commonly caused by environmental or contagious pathogenic microorganisms (Escherichia coli, Streptococcus uberis and Staphylococcus aureus)
- Economic losses

RNA-Seq

 Aims to characterize the transcriptome of a specific tissue at a specific time or condition





- RNA-Seq studies on Mastitis in dairy livestock:
 - More studied in dairy cows (Fang et al., 2017; Asselstine et al., 2019; Wang et al., 2020)
 - Mycoplasma agalactiae Infection in sheep (Chopra-Dewasthaly et al., 2017)

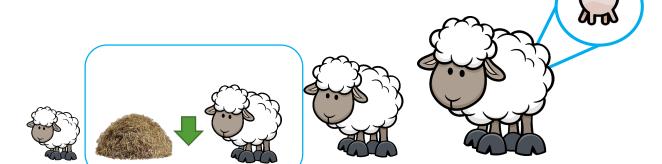


OBJECTIVE

The aim of this study was:

To evaluate mammary gland transcriptomic changes after an intramammary lipopolysaccharide (LPS) infusion in lactating sheep subjected to a nutritional challenge at the prepuberal stage (3-5 months of age)







EXPERIMENTAL DESIGN= NUTRITIONAL CHALLENGE



Nov-18	Dec-18	Jan-19	Feb-19	Mar-19	Apr-19	may-19	jun-19	Jul-19	Aug-19	Sep-19	Oct-19	Nov-19	Dec-19	Jan-20	Feb-20	Mar-20	Apr-20	May-20	Jun-20	Jul-20
MI	M2	M3	M4	M5	M6	M7	M9	MI0	MII	MI2	MI3	MI4	MI5	MI6	MI7	MI8	MI9	M20	M21	M22

Ewes' birth

N = 40

Allometric mammary gland







Ad libitum barley straw + 1.3 kg of granulated feed/animal and day

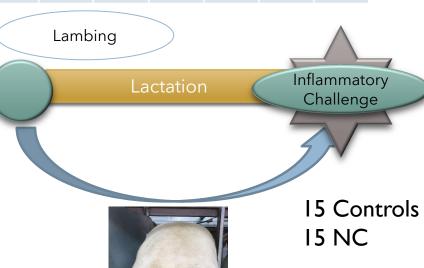
CSIC

Nutritional challenge= Differences in granulated feed

• Control (**n=20**)= 16% raw protein (soy cake)

• Challenge (n=20)= without soy cake, containing ~9% raw protein

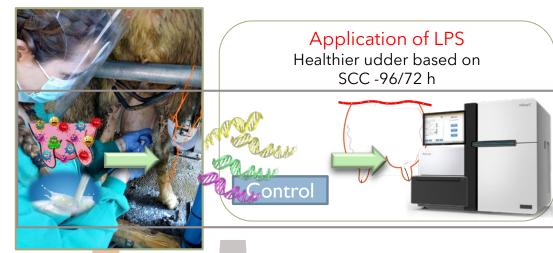






EXPERIMENTAL DESIGN= INFLAMMATORY CHALLENGE

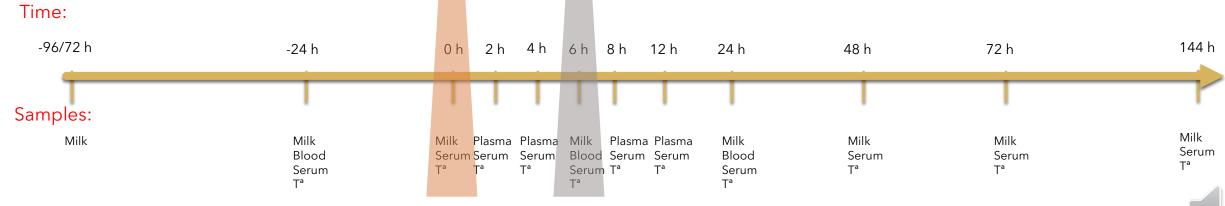




Animals:

12 Nutritional Challenge (NC

B Control (C





RNA-SEQ BIOINFORMATIC PIPELINE

RNA-Seq data

Quality Control

• FastQC

Mapping to
Ovine genome

- Rambouillet_v1.0
- STAR

Quantification

• RSEM

Differential Expression Analysis

• DESeq2

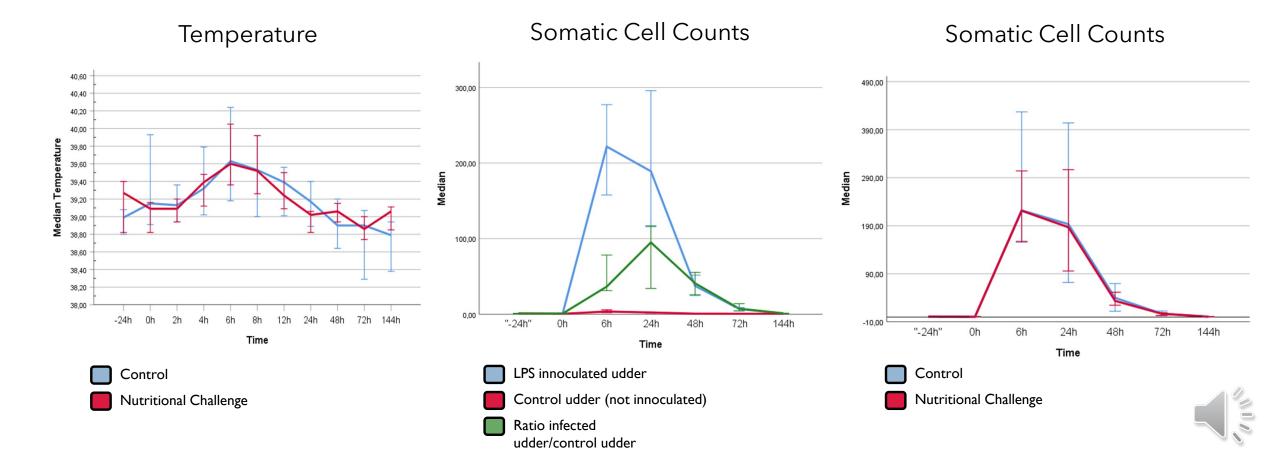
Enrichment Analyses

- ToppGene
- GOplot

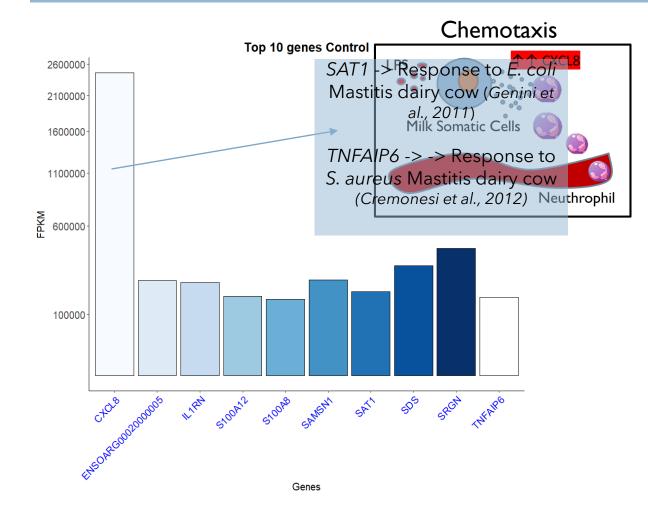


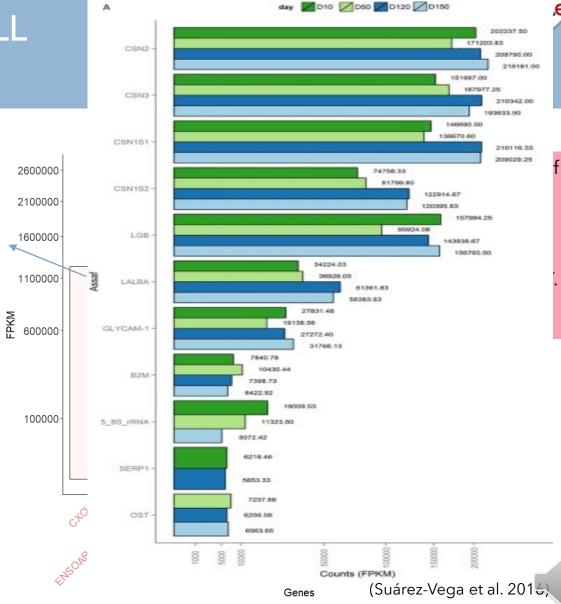


EFFECTS OF THE INFLAMMATORY CHALLENGE









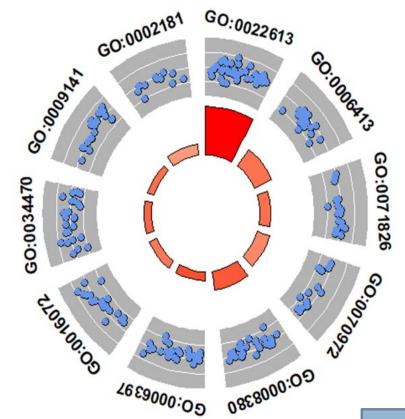


DIFFERENTIAL EXPRESSION ANALYSES: GENES UPREGULATED IN CONTROL

585 Differentially expressed genes (padj≤0.05)



495 upregulated in Control animals



	logFC	z-score	;			
•	upregulated					
	decre	asing	increasing			

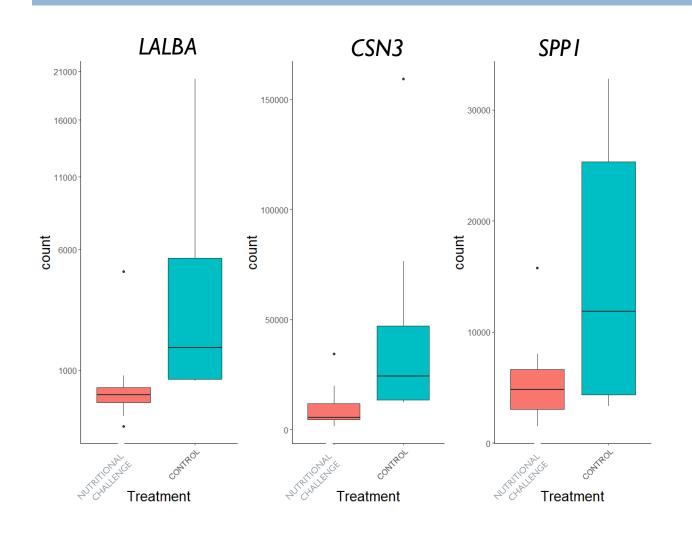
ID	Description
GO:0022613	ribonucleoprotein complex biogenesis
GO:0006413	translational initiation
GO:0071826	ribonucleoprotein complex subunit organization
GO:0070972	protein localization to endoplasmic reticulum
GO:0008380	RNA splicing
GO:0006397	mRNA processing
GO:0016072	rRNA metabolic process
GO:0034470	ncRNA processing
GO:0009141	nucleoside triphosphate metabolic process
GO:0002181	cytoplasmic translation

Transcriptional and translational regulation, protein synthesis





DIFFERENTIAL EXPRESSION ANALYSES: GENES UPREGULATED IN CONTROL



Transcriptional and translational regulation, protein synthesis



Synthesis of lactation proteins

Higher number of Milk Epithelial Cells in Controls??





stress-activated protein kinase

Genes=9; FDR=2.633E-3

signaling cascade

leukocyte activation involved in immune

Genes=13 FDR=3.132E-3

response

DIFFERENTIAL EXPRESSION ANALYSES: GENES UPREGULATED IN NUTRITIONAL CHALLENGE

585 Differentially expressed genes (padj≤0.05)



90 upregulated in Nutritional Challenge animals

immune effector process
Genes=19; FDR=2.234E-3

JNK cascade
Genes=8; FDR=2.234E-3

PYGL

PYGL

RAB31

TOP 5 GO BP

Genes=20; FDR=2.234E-3

Susceptibility/tolerance to mastitis in dairy cows and sheep (Banos et al., 2017; Oget et al., 2019; Yang et al., 2019)



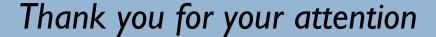
(Mugabo et al., 2010)



TAKE HOME MESSAGE

- The LPS inflammatory challenge has induced a change in the transcriptome profile of the milk somatic cells
 - Top 10 genes are related to immune response and not to milk protein synthesis
 - The majority of genes differentially expressed were upregulated in the control group.
 - Genes overexpressed in controls were implicated in transcriptional and translational processes, and main lactation proteins were upregulated. Consequently, we hypothesize controls had a higher proportion of mammary epithelial cells that nutritional challenge sheep at 6 hours post LPS inoculation
 - In the nutritional challenge animals, the differential expressed genes were mainly related to immunity processes suggesting that protein restriction during the growth period could lead to a higher inflammatory immune response







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