





### Milk Metabolomics of Heat-Stressed Goats with and without Intramammary Lipopolysaccharide Challenge

S. Love<sup>1</sup>, A. Contreras-Jodar<sup>2</sup>, N. Mehaba<sup>2</sup>, X. Such<sup>2</sup>, G. Caja<sup>2</sup>, <u>A.A.K. Salama<sup>2</sup></u>

<sup>1</sup> University of Glasgow, Glasgow, UK
 <sup>2</sup> Universitat Autònoma de Barcelona, Barcelona, Spain

### Responses of goats to heat stress (HS)

#### Thermophysiological traits:

Rectal Temperature (+0.58°C) Respiratory Rate (+300%) Water consumption (+202%) Dry matter intake (-21%) Body weight (-125 g/d)

# Milk yield and milk quality: Milk yield (-4 to -10%) Milk quality (-8 to12% fat & protein) Cheese coagulation properties (> RCT)

Hamzaoui *et al.* (2012, 2013); Abdel-Gawad *et al.*, 2014; Castro-Costa *et al.*, 2015; Contreras-Jodar *et al.*, 2018)





### Transcriptomics of goat milk cells

RNA-seq revealed 699 differentially-expressed genes by HS.



#### **Objectives**

- 1. Whether heat stress would affect the immune response of the mammary gland to infection.
- The use of <sup>1</sup>H NMR-based milk metabolomics to identify inflammation markers under:
  - Thermal-neutral (TN) conditions
  - Heat stress (HS) conditions



### **Experimental design**

- 8 Murciano-Granadina dairy goats (42 ± 2 kg BW; 100 ± 5 DIM; 2.2 ± 0.1 L/d)
- Climatic chamber:
  - TN: 15 to 20°C, 45 to 55% HR; THI = 59 to 65
  - HS: 35°C-12h and 28°C-12h, 40% HR; THI = 83 (day) and 75 (night)
- Challenge with LPS :
  - LPS half:**10 µg** in 2 ml 0.09% sterile saline
  - CON half: 2 ml 0.09% saline

### **Experimental design**



**n = 8** 4 goats / trt

#### d 1 to d 15:

- Milk yield
- Milk composition
- Feed and water intakes
- RT and RR

#### d 12 to 15:

- Milk: 0, 2, 4, 6, 8, 12, 24, 48, 72 h
   for yield and composition
- Milk: 0, 4, 6,12, 24 h for NMR metabolomics analysis

#### Milk sample preparation and NMR analysis (n=80)

Sample Preparation (Beckonert et al., 2007)					
1.	Buffer	Na <sub>2</sub> HPO <sub>4</sub> , NaH <sub>2</sub> PO <sub>4</sub> , NaN <sub>3</sub> , D <sub>2</sub> C			
2.	Milk Filtration	4 mL in 10 kDa filtration tube			
3.	Tube centrifugation	12,000 × g, 20 min at 22°C			
4.	Sample	400 μΛ			
5.	Buffer	200 μΛ			
6.	Milk + Buffer mix	550 μΛ ιν 5 mm NMR tubes			



#### Milk sample preparation and NMR analysis (n=80)

Sample Preparation (Beckonert et al., 2007)				
1.	Buffer	Na <sub>2</sub> HPO <sub>4</sub> , NaH <sub>2</sub> PO <sub>4</sub> , NaN <sub>3</sub> , D <sub>2</sub> O		
2.	Milk Filtration	4 mL in <b>10 kDa</b> filtration tube		
3.	Tube centrifugation	12,000 × g, 20 min at 22°C		
4.	Sample	400 μΛ		
5.	Buffer	200 μΛ		
6.	Milk + Buffer mix	550 μΛ ιν 5 mm NMR tubes		
<sup>1</sup> H nuclear magnetic resonance (NMR) analysis				
1.	Bruker Avance-III	NMR 600 MHz <sup>1</sup> H; 298 <sup>o</sup> K		
	spectrometer (Bruker	NOESY pulse sequence		
	BioSpin, DE)			
2.	Bruker TopSpin	Data acquisition		
	software v2.1.	Data processing		



# **Bioinformatic and statistical analysis**

1.	Bioinformatics	<ul> <li>Baseline correction</li> </ul>		
		Region of interest:	$\delta$ 8.0 – 5.0 and 4.6 – 0.5 ppm	
		<ul> <li>Binning: interval width from 0.0003 to 0.0300 ppm</li> </ul>		
		<ul> <li>Normalization: mean centered, logarithm transf.</li> </ul>		
2.	Statistics	Multivariate analysis : PCA and PLS-DA		
		Volcano Plot		
3.	Metabolite Id	Previous bibliography (Sundekilde <i>et al.</i> , 2013) Human Metabolome Database (Wishart <i>et al.</i> , 2009)		



R software 3.2.3 version (R Core Team, 2015)

# 600 MHz <sup>1</sup>H NMR Metabolomics in Milk



- 1. 3-metylhistidine
- 2. Hippurate
- 3. Fumarate
- 4. Orotate
- 5. Maleate
- 6. Urea
- 7. Lactose
- 8. Carnitine
- 9. Creatinine
- 10. Choline
- 11. Citrate
- 12. N-acetylcarbohydrates
- 13. Acetate
- 14. Lactate
- 15. b-hydroxybutyrate
- 16. Butyrate

#### PLSDA Scores: TN-CON vs HS-CON (markers of HS)



#### **PLSDA plot: response to LPS throughout time**



# CON vs. LPS (inflammation markers)

Flux	Chemical shift (ppm)	Metabolite
+	3.19	Choline
+	1.32	L-lactate
+	2.05	N-acetyl CH
+	1.20	Beta-hydroxybutyrate
-	3.84	Lactose
-	4.16	Phosphocholine

## Response to LPS (markers of inflammation)





# **Response to LPS (markers of inflammation)**



### **Response to LPS (markers of inflammation)**



# Conclusions

- Milk metabolome changed throughout time after LPS administration, with values restored earlier in TN compared to HS goats.
- Milk citrate increased by heat stress as part of the heat shock response, enabling the synthesis of pro-inflammatory mediators.
- Inflammation markers included choline, phosphocholine,
   N-acetylcarbohydrates, L-lactate and ß-hydroxybutyrate.
- The importance of these markers varied between TN and HS indicating different mammary immune response.





#### Thank you for the attention!

Component 1 ( 96.8 %)

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