

#### Clustering and genetic analysis of body condition score and body weight variations in meat sheep

Tiphaine Mace, Dominique Hazard, Sebastien Douls, Didier Foulquie, Eliel González García

#### ▶ To cite this version:

Tiphaine Mace, Dominique Hazard, Sebastien Douls, Didier Foulquie, Eliel González García. Clustering and genetic analysis of body condition score and body weight variations in meat sheep. 2017, 27 p. hal-01607238

HAL Id: hal-01607238

https://hal.science/hal-01607238

Submitted on 5 Jun 2020

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.









# CLUSTERING AND GENETIC ANALYSIS OF BODY CONDITION SCORE AND BODY WEIGHT VARIATIONS IN MEAT SHEEP

MACE Tiphaine <sup>1</sup>,

Hazard D.<sup>1</sup>,

Carriere F.<sup>2</sup>, Douls S.<sup>2</sup>, Foulquié D.<sup>2</sup>

González-García F. 3

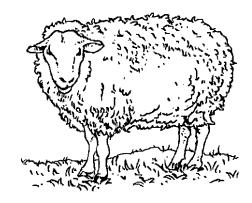
<sup>1</sup>INRA UMR1388 GENPHYSE, 31326 Castanet-Tolosan, France

<sup>2</sup>INRA UE321 La Fage, 12250 Roquefort-sur-soulzon, France;

<sup>3</sup> INRA UMR868 SELMET, 34060 Montpellier, France

### Plan

- Context Robustness
- Cluster analysis
  - Materials and methods
  - > Results
- Genetic analysis
  - > Creation of new traits
  - ➤ Materials and methods
  - > Results
- Discussion Conclusion
- Perspective



## Robustness

> Robust animals to assure the farm sustainability in harsh condition

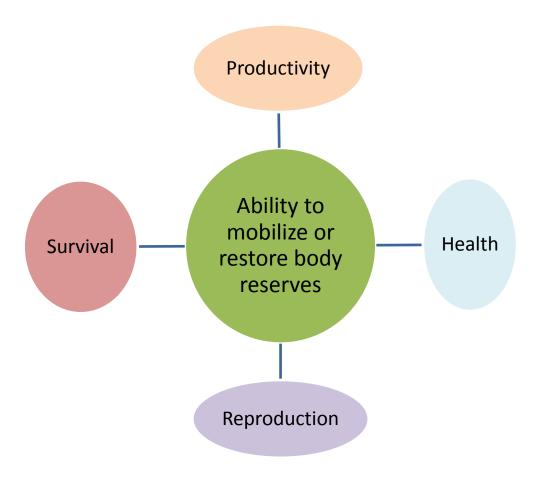
Ability to mobilize or restore body reserves

Russel *et al.,* 1971 Blanc *et al.,* 2006



## Robustness

> Robust animals to assure the farm sustainability in harsh condition



Russel *et al.,* 1971 Blanc *et al.,* 2006



# Robustness – Body Condition Score and Body Weight

- Body Condition Score (BCS) and BW to characterize robustness
  - BCS: grid from 1 to 5
  - Capacity of mobilization or accretion of body reserves (BR)
- > Heritability
  - BCS
    - From 0.10 to 0.30 for Targhee ewes
  - Body Weight (BW)
    - From 0.30 to 0.56 for adult sheep
- No study on genetic for BCS and BW variations across productive cycle

Borg *et al.,*Gizaw *et al.,*Loker *et al.,*Shackell *et al.,*



# Robustness – Body Condition Score and Body Weight

#### Objectives

- To characterize the different patterns of BCS and BW variation during productive cycle and career
- To analyze the genetic variability of BCS and BW changes throughout several productive cycles

# Materials and Methods

➤ La Fage: meat Romane flock (n=250 ewes) reared exclusively outdoors in harsh conditions





#### Materials and Methods

- ➤ La Fage: meat Romane flock (n=250 ewes) reared exclusively outdoors in harsh conditions
- > BCS and BW measured on 1448 ewes from 78 sires (2002-2015)

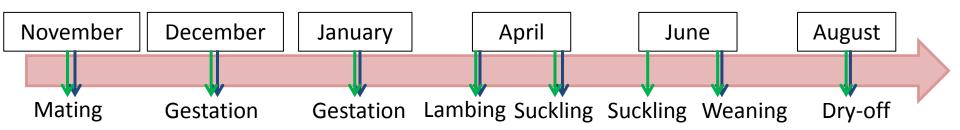
o Cycle 1: 1448

o Cycle 2: 1170

Cycle 3 and more: 665

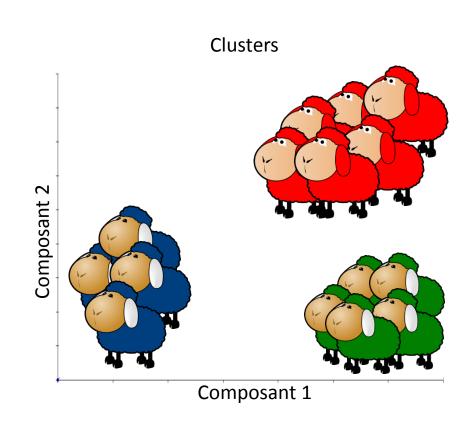
#### Materials and Methods

- ➤ La Fage: meat Romane flock (n=250 ewes) reared exclusively outdoors in harsh conditions
- BCS and BW measured on 1448 ewes from 78 sires (2002-2015)
  - o Cycle 1: 1448
  - o Cycle 2: 1170
  - Cycle 3 and more: 665
- 8 BCS and 7 BW measurements per production cycle



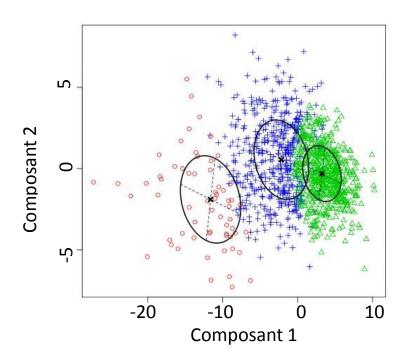
# Materials and Methods – Cluster analysis

- Clustering
  - To describe profiles
  - Smoothing
  - o PCA
  - Clusters construction
    - Expectation-Maximization Algorithm

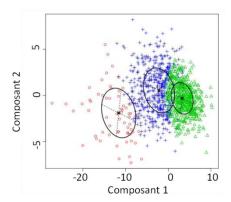




# **Results for BCS Clusters**

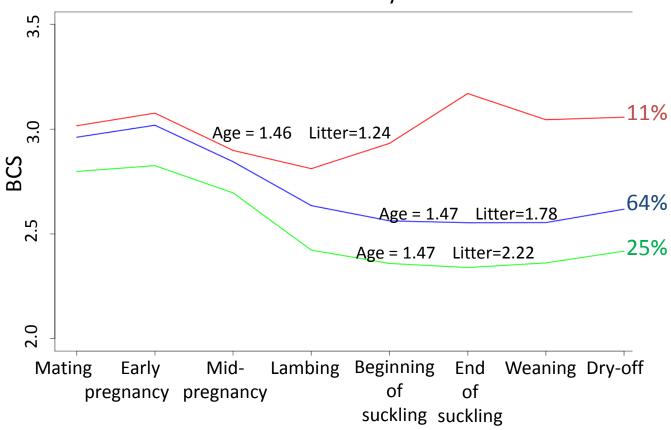






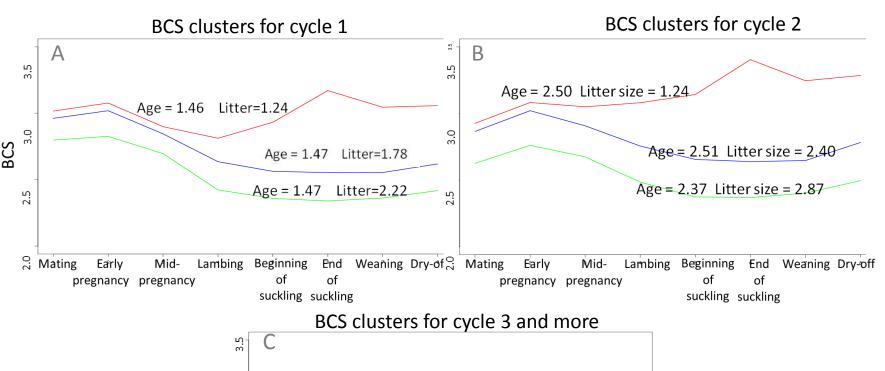
## **Results for BCS Clusters**

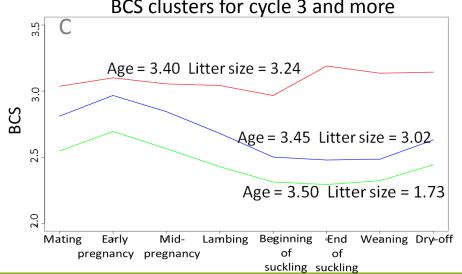




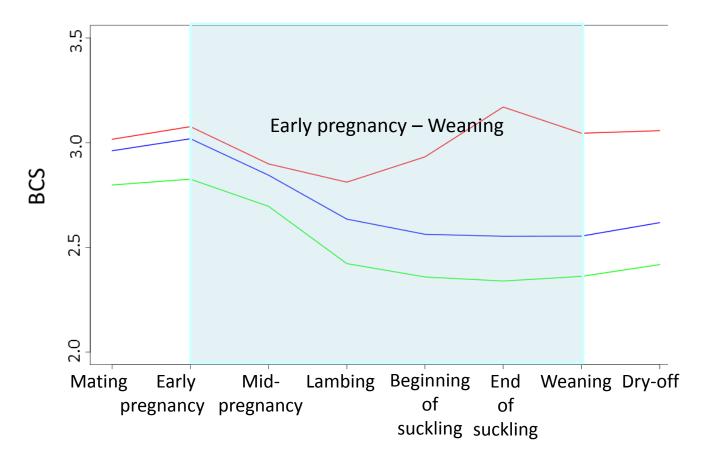


## Results for BCS Clusters



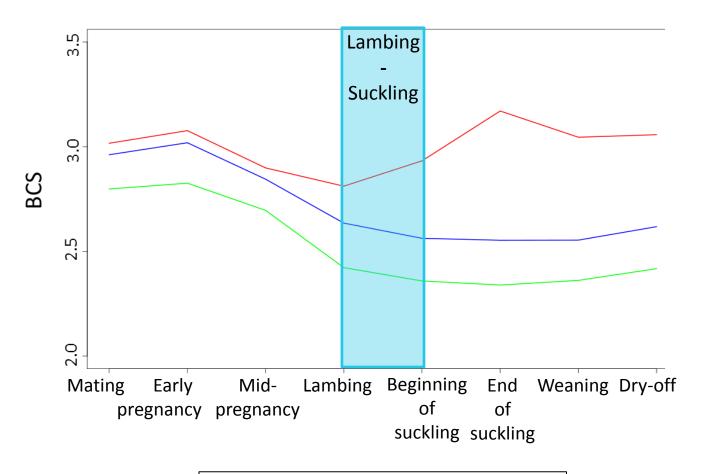






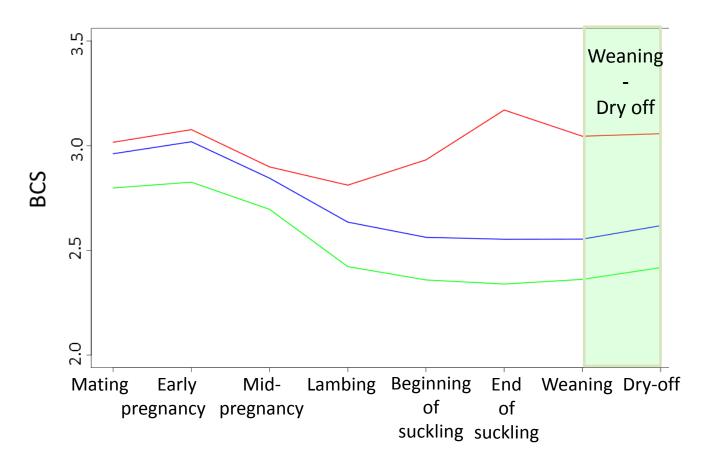
$$Y = BCS_{Weaning} - BCS_{Early Pregnancy}$$





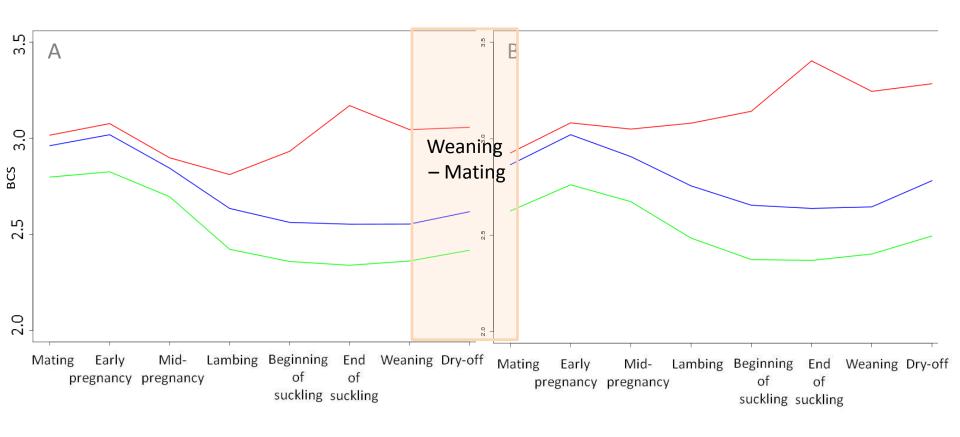
$$Y = BCS_{Suckling} - BCS_{Lambing}$$





$$Y = BCS_{Dry off} - BCS_{Weaning}$$

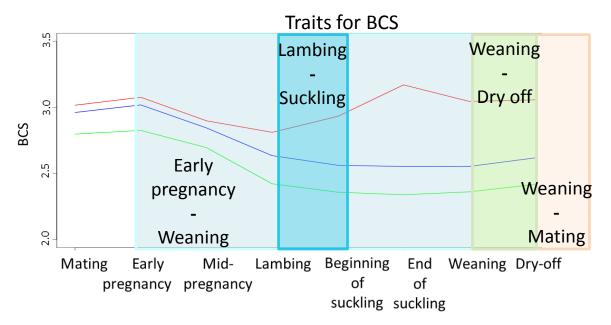




$$Y = BCS_{Mating} - BCS_{Weaning}$$



# Materials and Methods – Genetic Analysis



Univariate and multivariate analysis (animal model): Heritabilities and correlations estimation with (ASReml), use of all cycles

Y = mu + animal + perm + age + parity + litter + year + e

Y: trait

mu: mean of the trait

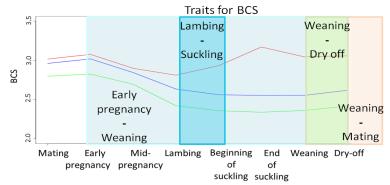
animal: additive genetic effect of the ewe

perm: environmental permanent effect of the ewe

age of the ewe, parity, litter, year of measurements as fixed effects

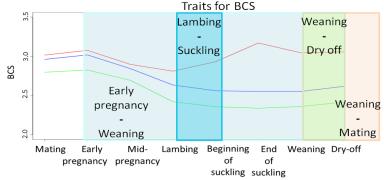
e: residual





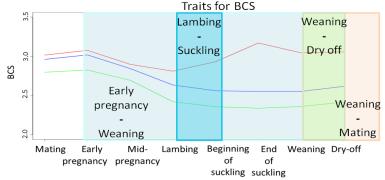
- $\blacktriangleright$  Low to medium heritabilities (0.06 ± 0.01 to 0.15 ± 0.01)
- Positive correlations between periods characterizing BR mobilization

Variables BCS	Lambing - Suckling
Early pregnancy - Weaning	0.49 (0.12)



- $\triangleright$  Low to medium heritabilities (0.06 ± 0.01 to 0.15 ± 0.01)
- Positive correlations between periods characterizing BR mobilization
- Positive correlations between periods characterizing BR accretion

Variables BCS	Weaning - Mating
Weaning – Dry off	0.75 (0.10)



- $\triangleright$  Low to medium heritabilities (0.06 ± 0.01 to 0.15 ± 0.01)
- Positive correlations between periods characterizing BR mobilization
- Positive correlations between periods characterizing BR accretion

Negative correlations between periods characterizing BR accretion and mobilization

Variables BCS	Weaning – Dry off
Early pregnancy - Weaning	-0.71 (0.08)
Lambing - Suckling	-0.46 (0.12)

#### **Discussion - Conclusion**

- > BCS: three similar profiles across productive cycles
- BCS and BW variations heritable: first demonstration
- Increase in weight during dry-off correlated to heavier lambs
- Accretion and mobilization of BR linked at the genetic level
  - Occurrence of the common genes?

#### **Discussion - Conclusion**

#### Perspectives

- Modelisation of the variations profiles (BCS and BW)
- Longitudinal genetic analysis with reaction norm model
- Joint analysis with climatic data (reaction norm model)
- Better phenotyping for BR variations (key metabolites /hormones /ultra sound...)



