



Microbiota clustering and energy balance in early lactation in Lacaune dairy sheep

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Introduction

- Lacaune sheep : 1st month = **Suckling milking** system
- **Energy deficiency** during this period → decrease in **milk production**
- Role of **microbiota** in ewes response

Is the ruminal microbiota of post-partum ewes linked to this energy deficit, and therefore to lower production?

Materials & methods

Animals

- ▶• 65 Lacaune dairy ewes
 - ▶• Multiparous
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- ▶ Diet:
 - 2/3 forages:
 - Alfalfa & cocksfoot hay
 - Grass silage
 - 1/3 concentrates

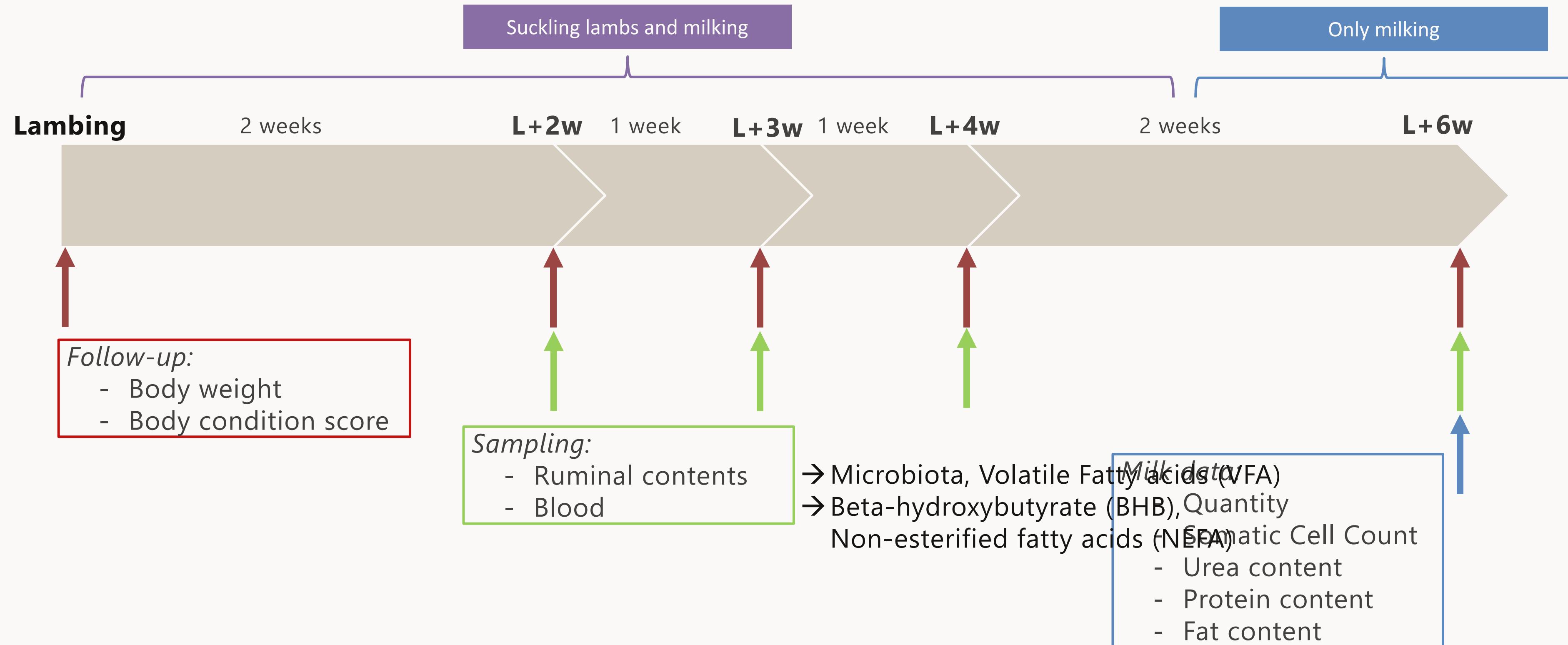


 Roquefort region

Materials & methods

04

Sampling and follow-up



Materials & methods

Microbiota analysis



Sequencing:

- V3-V4 regions of 16S rRNA



Genotoul Bioinformatics:

- Galaxy platform, FROGS
- SILVA base



• 258 samples (ie 4*65)

(2 ech removed - low number of sequences <6000)

- 21600 seq/sample
- 1932 ASVs / sample
- 104 genera



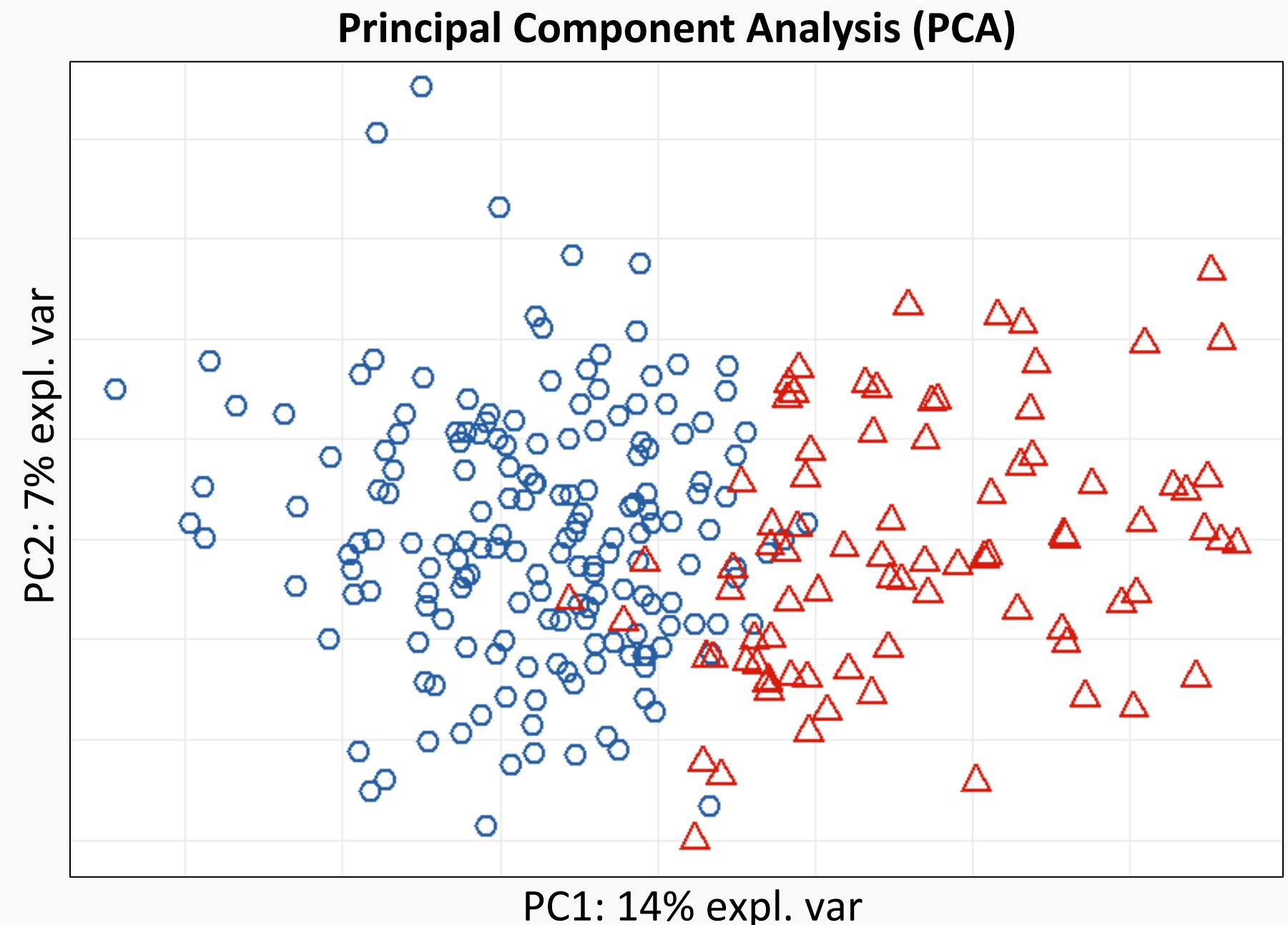
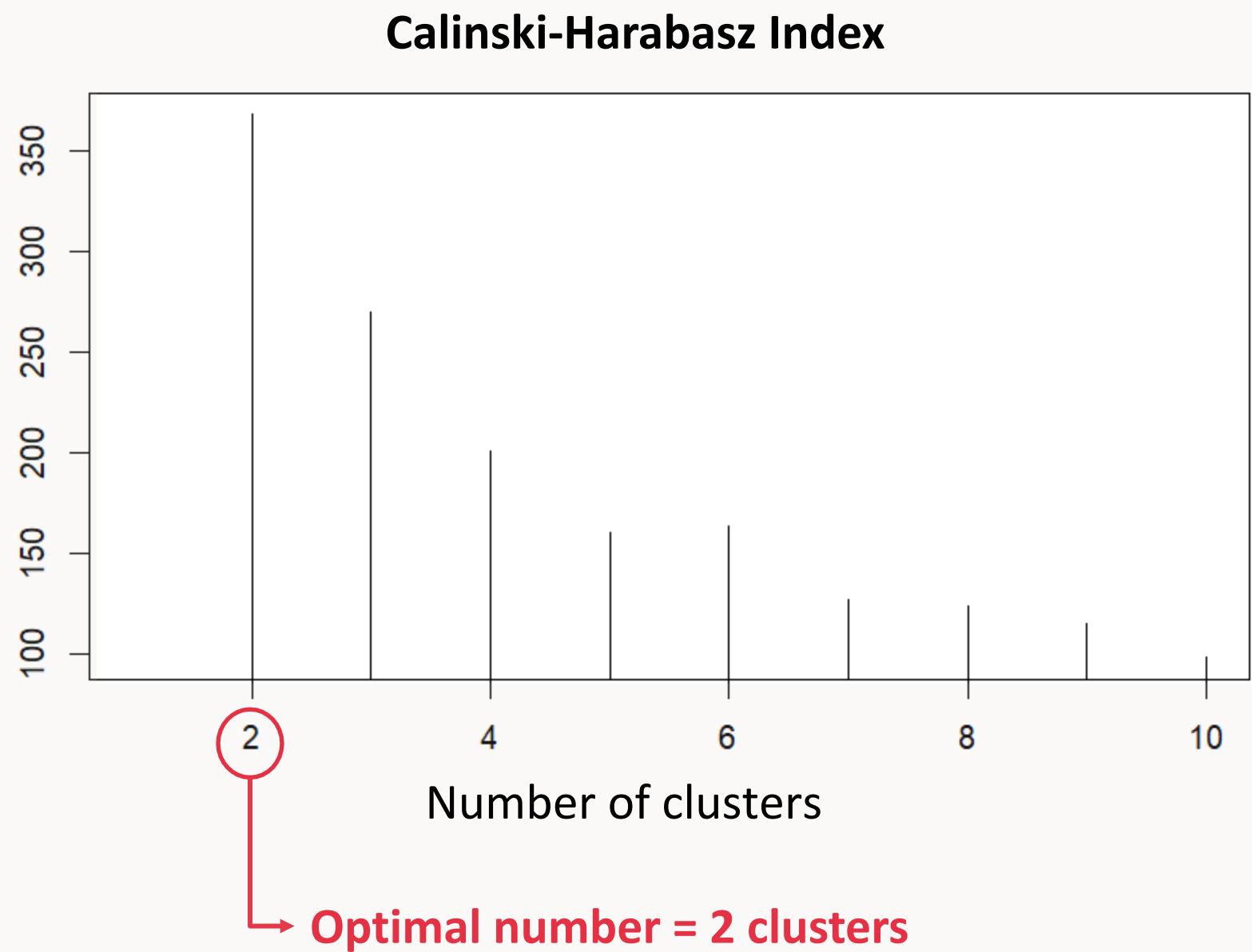
To reduce information

Unsupervised clustering:

- On total genera table with all samples from all dates
- Distance matrix based on Jensen-Shannon divergence
- Partition around medoids clustering
- Validation: silhouette coefficient

Results

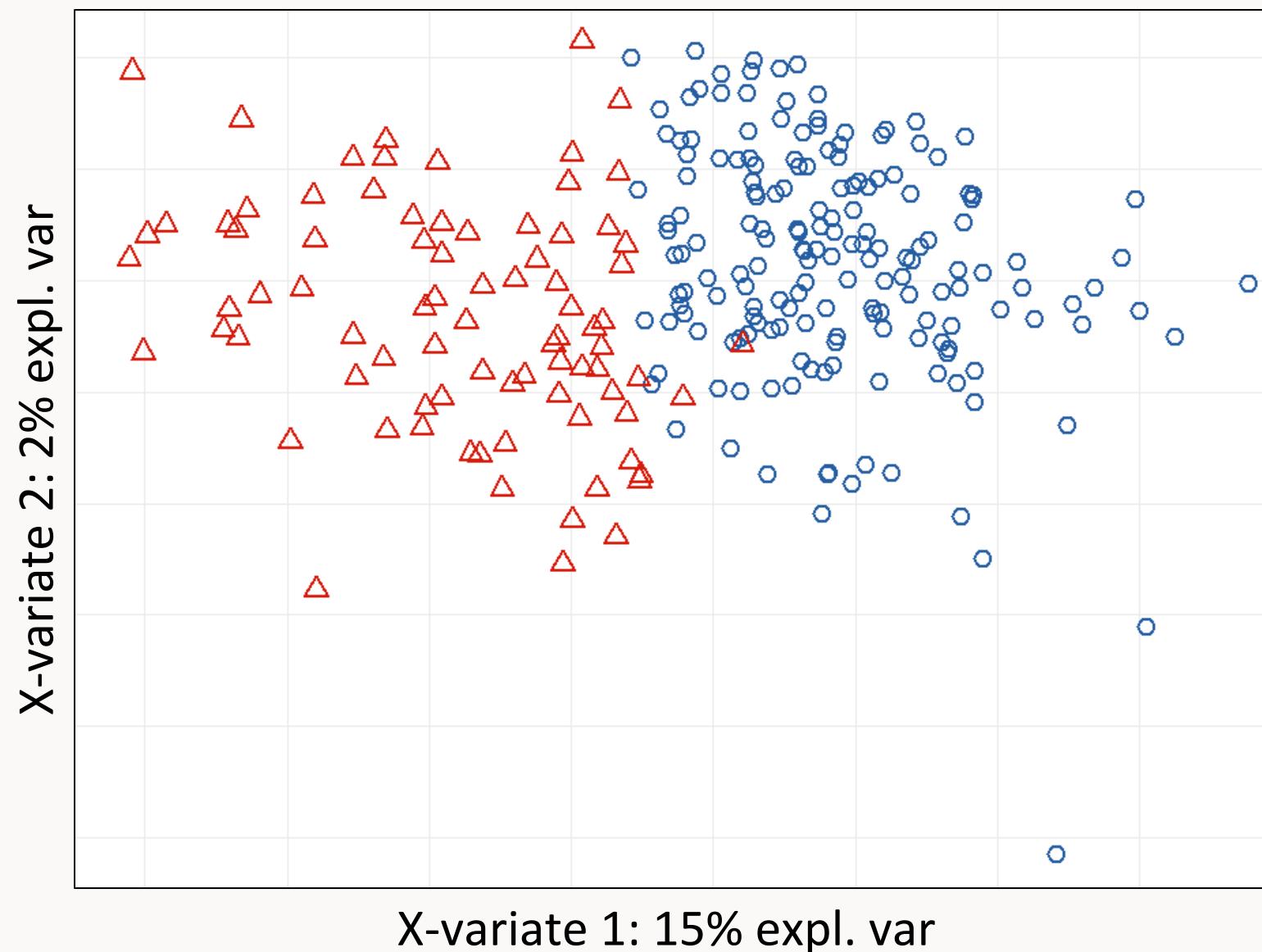
Microbiota clustering



Results

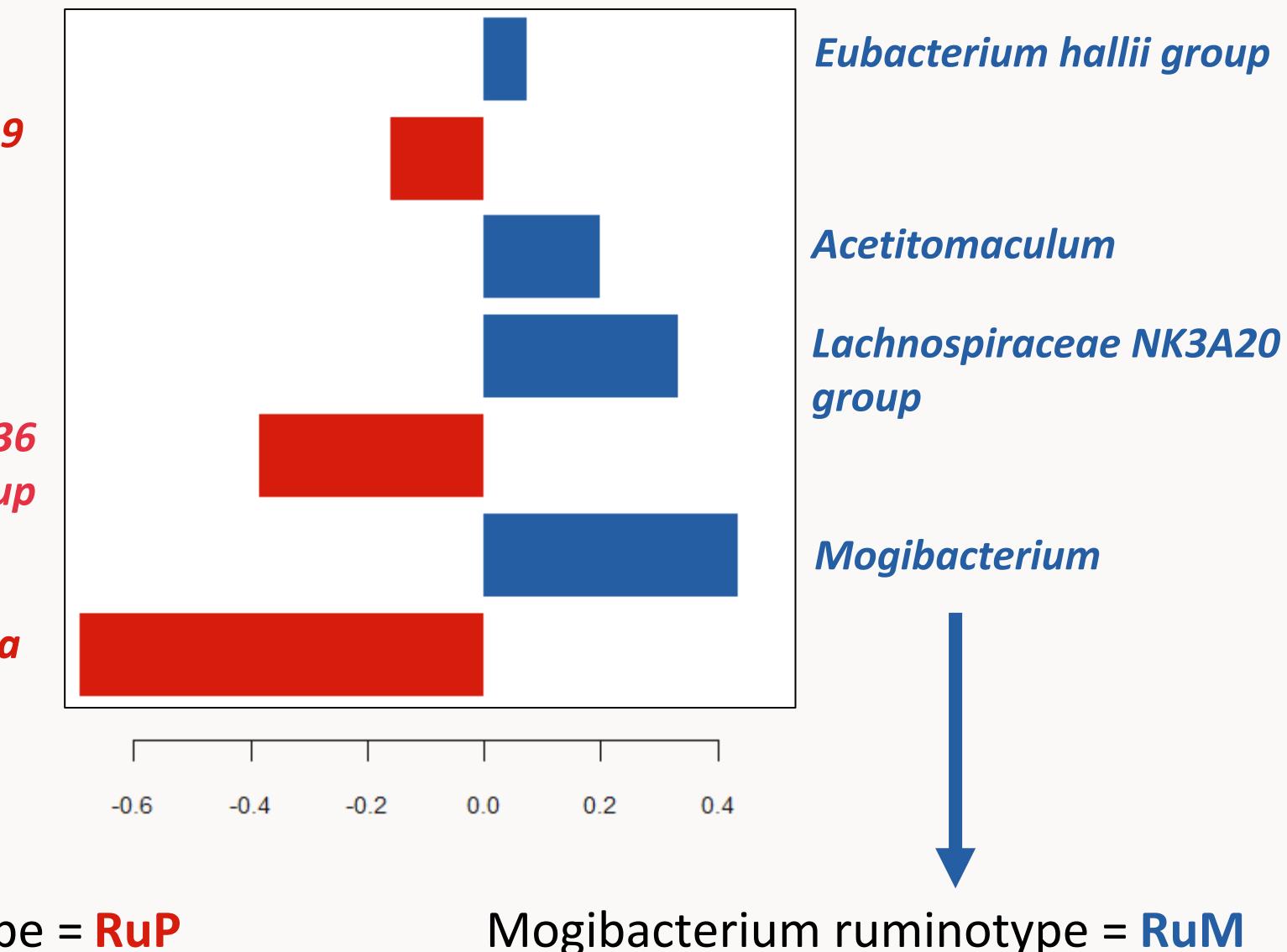
Microbiota clustering

Sparse Partial Least Squares Discriminant Analysis (SPLS-DA)



Prevotella 9
Lachnospiraceae NK4A136 group
Prevotella

Prevotella ruminotype = **RuP**



Results

Ruminotypes

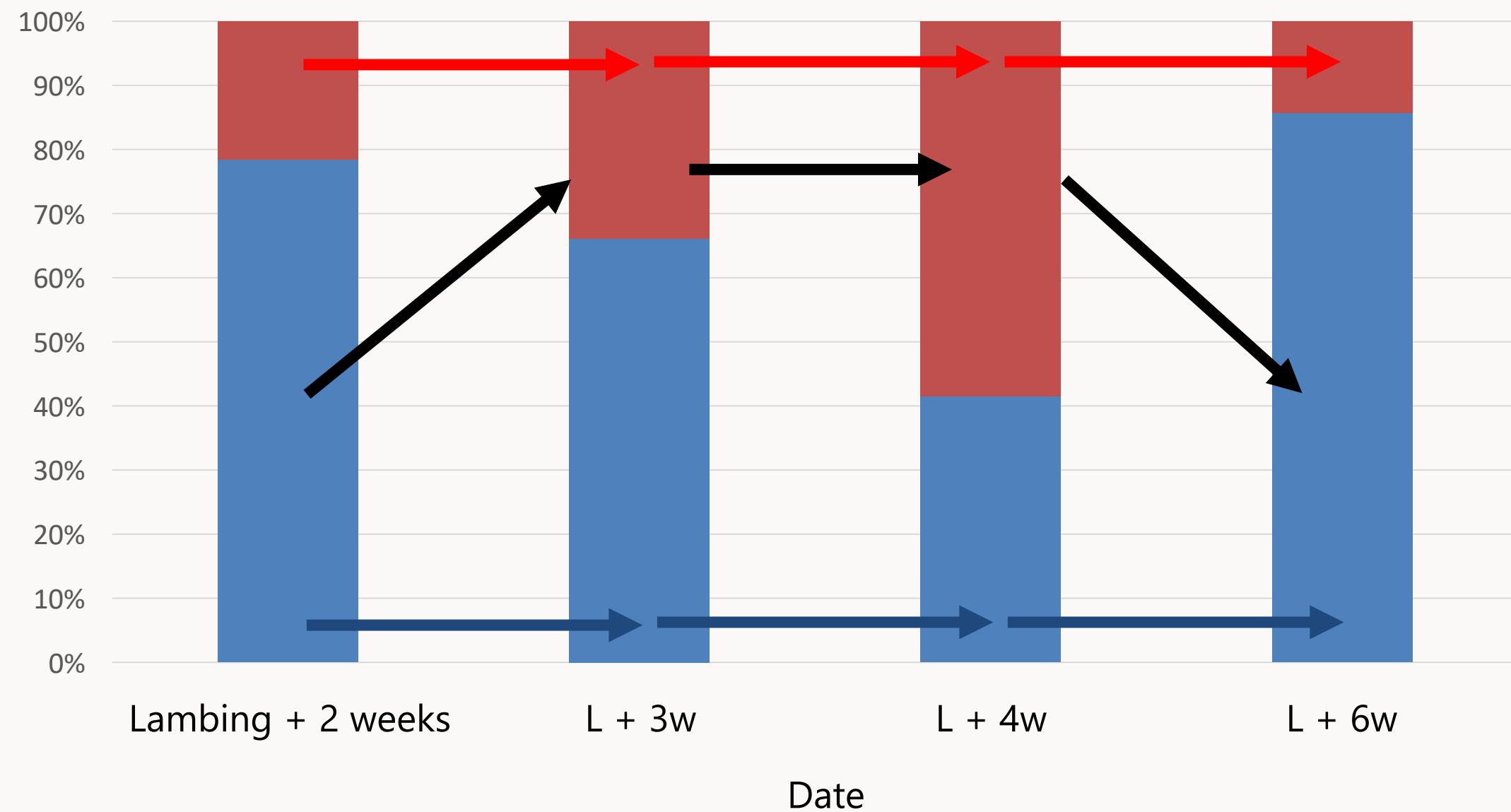
Or stay

Sheep may change

Or stay

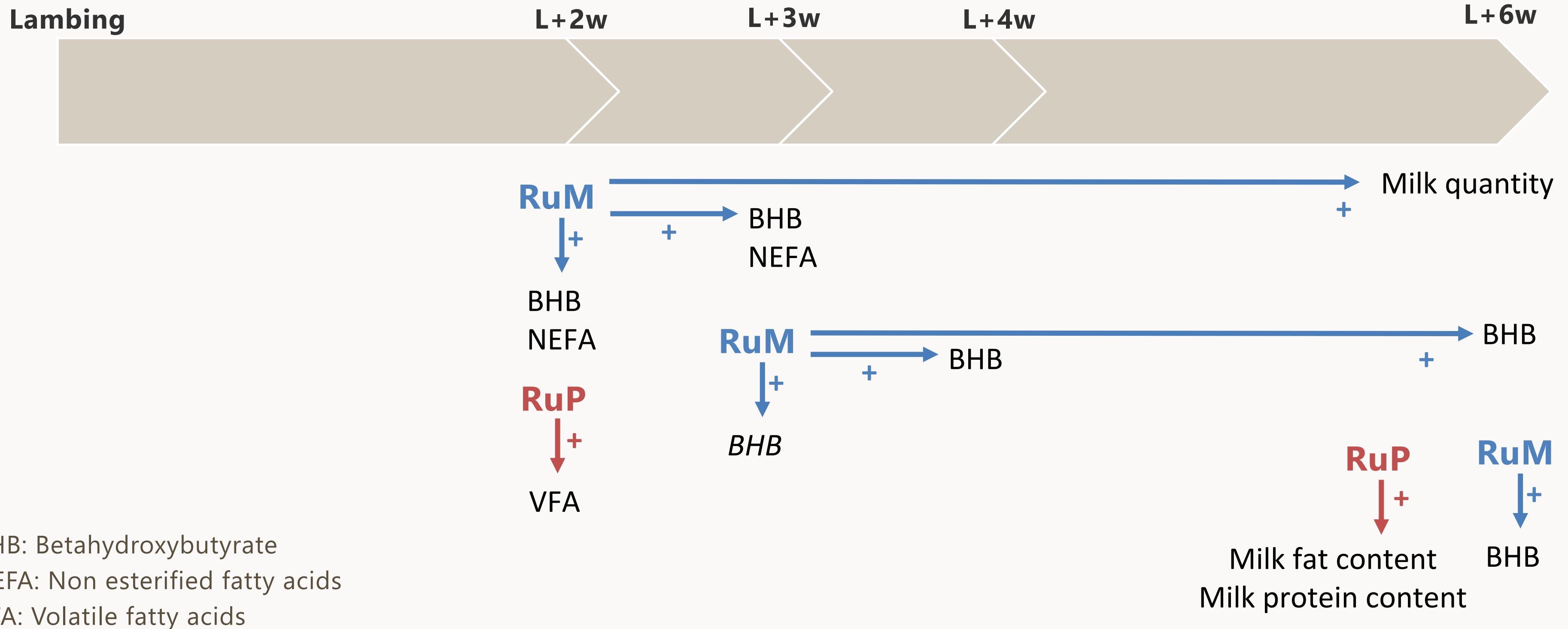
RuP

RuM



Results

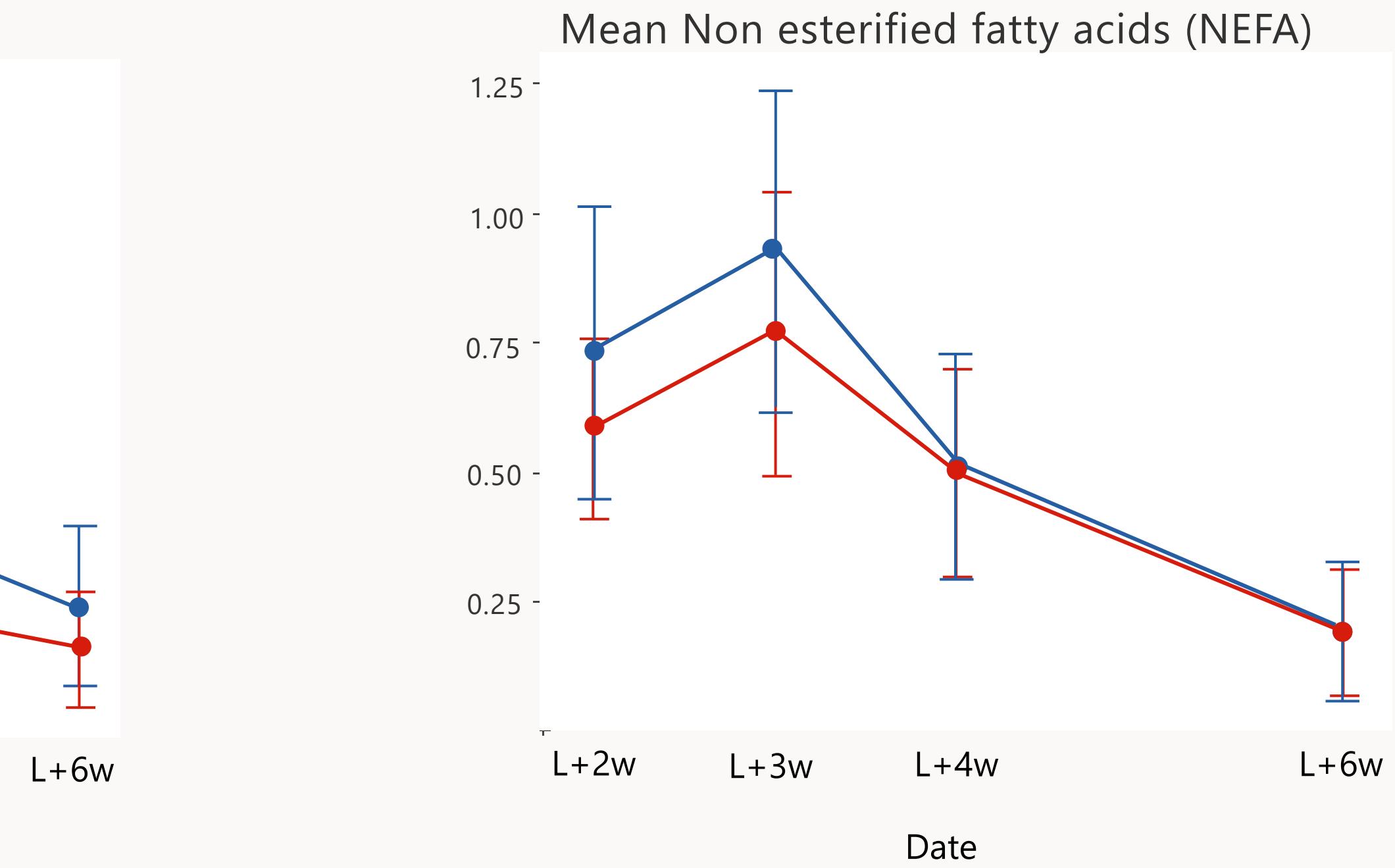
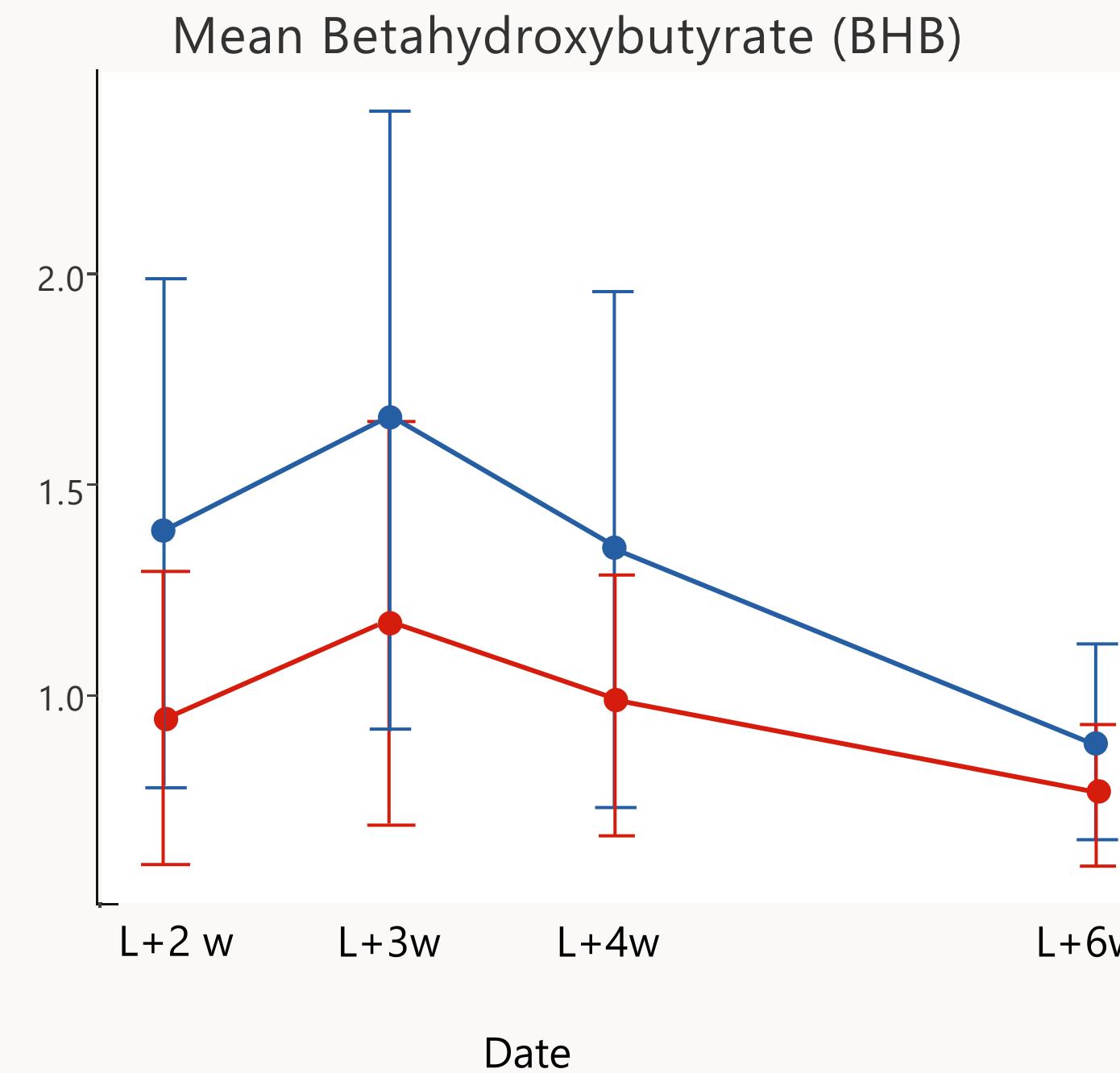
Ruminotypes



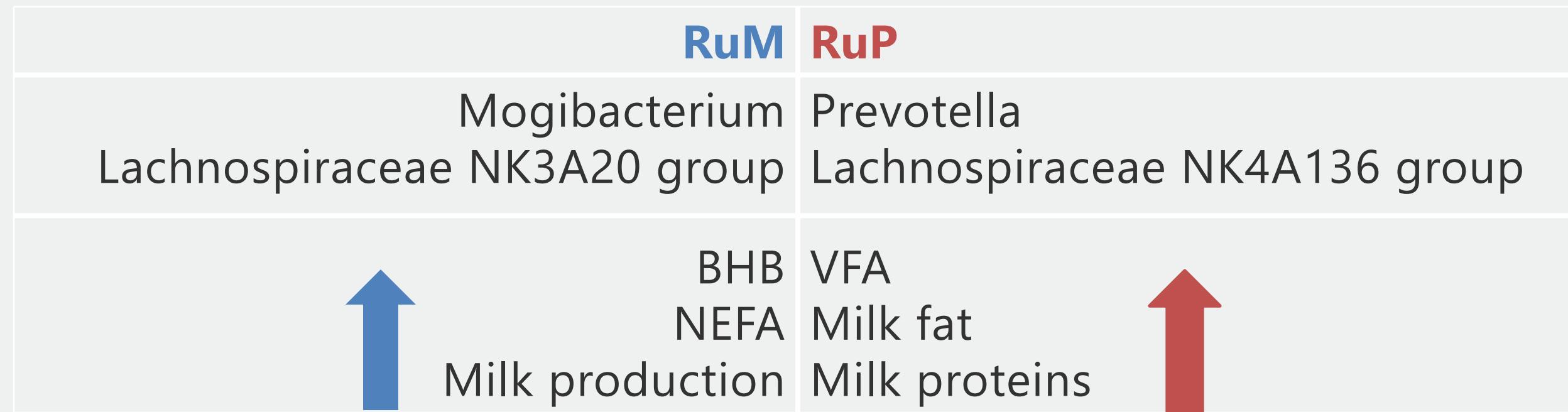
Results

Ruminotypes

● RuM at D1 & D2 (n= 31)
● Others (n= 32)



Conclusion



- Microbiota → Link with lipomobilization
- High lipomobilization → higher milk production?

BHB: Betahydroxybutyrate
 NEFA: Non esterified fatty acids
 VFA: Volatile fatty acids

Acknowledgements

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