

Effect of tannins and medium-chain fatty acids on rumen bacterial community of beef calves during the transition period to a high-concentrate diet

Yuste, S.¹, Amanzougarene, Z.¹, de la Fuente, G.²,
Fondevila, M.¹, de Vega, A.¹.

¹Dept. de Producción Animal, Universidad de Zaragoza, Instituto Agroalimentario de Aragón (IA2) (Spain).

²Dept. de Ciència Animal, Universidad de Lleida (Spain).

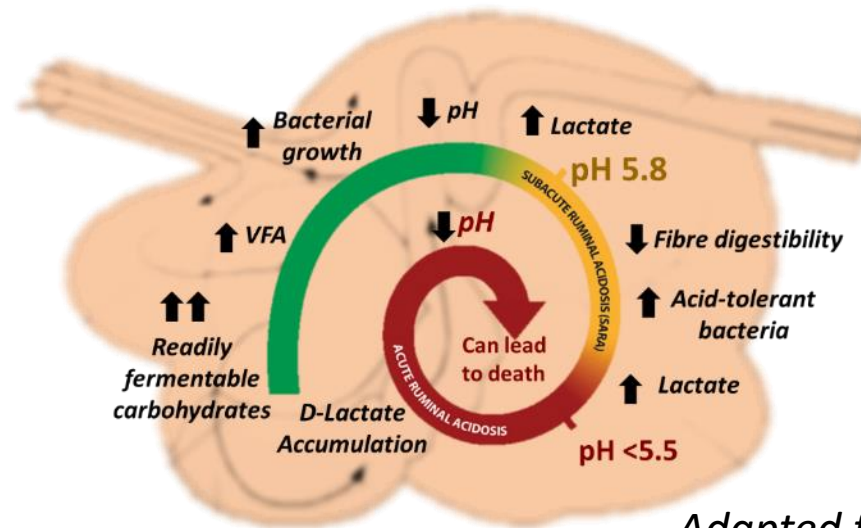
Introduction

Beef calves are weaned at around 6 months
Milk and pasture → High-concentrate ration



Transition period: 14-21 days after the feedlot arrival
Inadequate transition = **NUTRITIONAL DISORDERS**

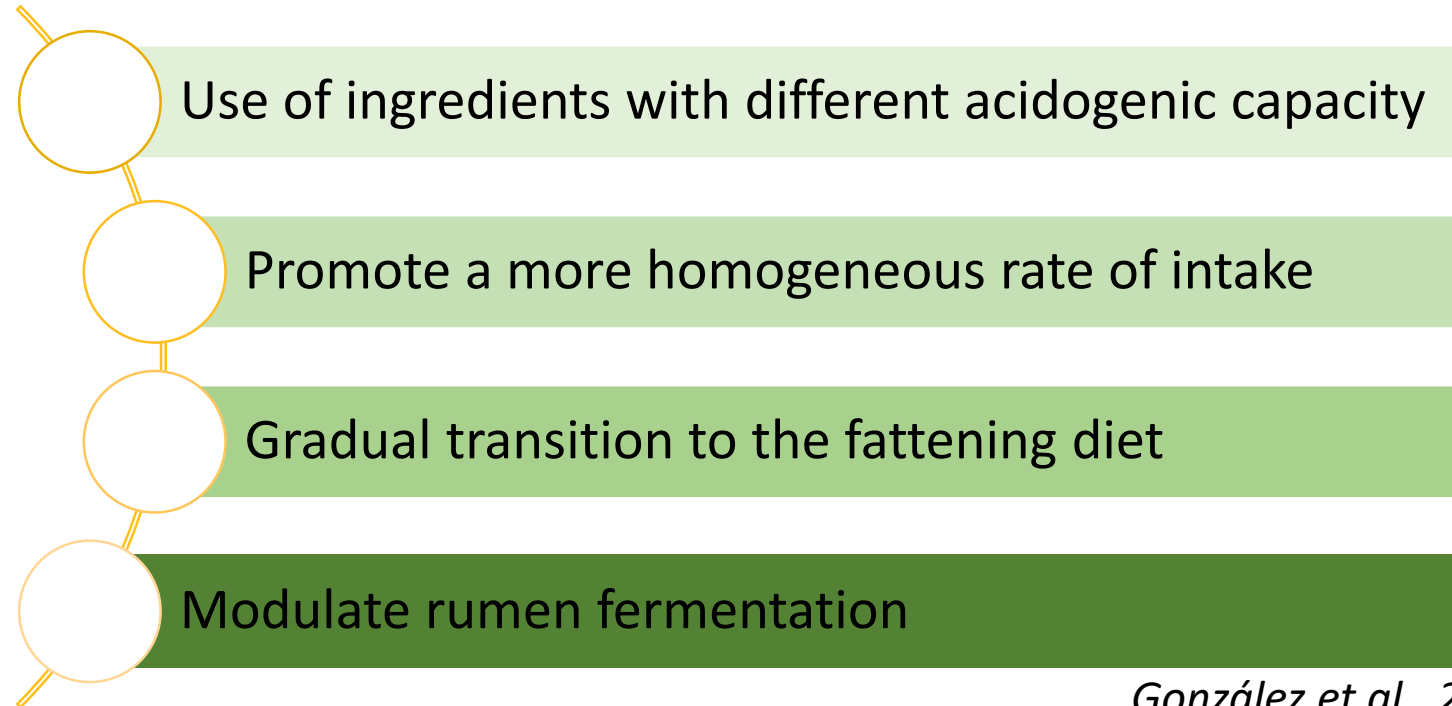
Brown et al., 2006



Adapted from Nocek, 1997

Introduction

Strategies to
improve rumen
health
and animal
performance



González et al., 2012

Feed additives such as tannins and medium-chain fatty acids (MCFA) modulate rumen fermentation



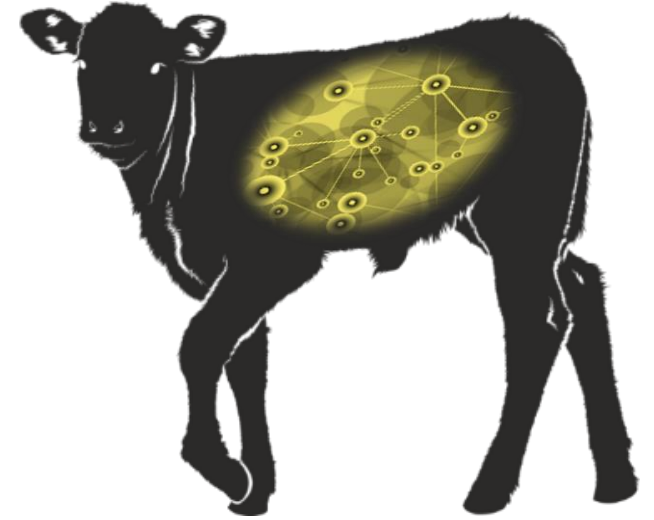
Machmüller, 2006

Patra and Saxena, 2011

Introduction

The rumen is a complex and dynamic ecosystem that changes in response to many factors.

(i.e. animal, age, **diet**, etc.) *Weimer, 2015*



Objective

To study the effect of the inclusion of tannins and MCFA on rumen bacterial community of beef calves during the transition from milk and pasture to a high-concentrate diet.

Materials and methods



18 cannulated calves / 212kg LW / ~ 6 months
Milk and grass → high-concentrate diet

Diet C

60% barley
15% maize
17% soybean
meal

Diet T

C + 20 g/kg of
65:35 chestnut
and quebracho
tannins



Diet M

C + 6 g/kg of a
MCFA mixture
(C₆-C₁₂)



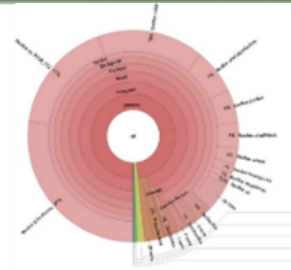
Doses were chosen based in previous in vitro studies (Amanzougarene et al., 2017,2018).

Concentrate + wheat straw were given *ad libitum* for **28 d**

Rumen fluid samples at 0h on days 0, 7, 14, 28

Storage at -80°C

Materials and methods



**Taxonomic
Classification**

**Bioinformatic
Analysis**

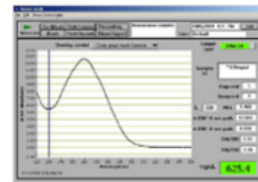
Ion Reporter Analysis

PCR and 16S DNA gene sequencing

DNA Extraction



ION TORRENT (PGM)



Materials and methods

- Log₁₀-transformed (n^o reads +1)
- Bioinformatic analysis with R software ('vegan', 'mixOmics'):
 - Multivariate analysis: PERMANOVA (ADONIS), analysis of similarity (ANOSIM)
 - Spatial ordination: Non-metric multidimensional scaling (NMDS) with Bray-Curtis distance
 - Shannon Index and genera richness
- Bacterial taxa (Log₁₀): PROC MIXED SAS. Diet, day of sampling and their interaction as fixed effects. Animal as random
 - Repeated measures. Mean comparison with Bonferroni's test
 - Differences were considered significant if $P < 0.05$. Trend was considered when $0.05 \leq P \leq 0.10$



Results



- Similar (DIET: $P > 0.10$)

-feed intake and rumen fermentation pattern

Additives intake: T: 100 g tannins/d M: 31 g MCFA/d

- PERMANOVA

Diet: $P = 0.92$, $R^2 = 0.007$

Day: $P < 0.001$, $R^2 = 0.32$

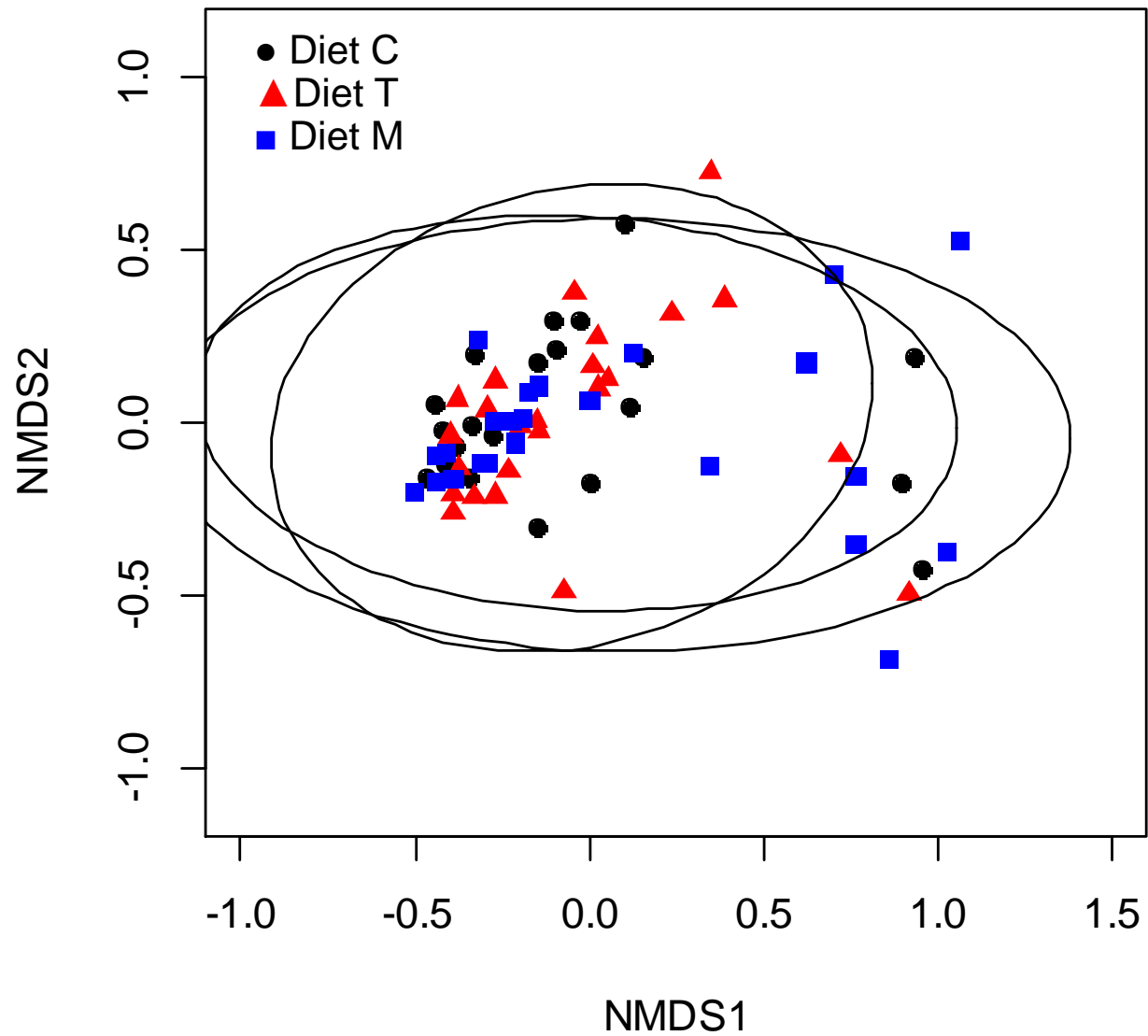
- ANOSIM

Diet: $P = 0.97$, $R = -0.03$

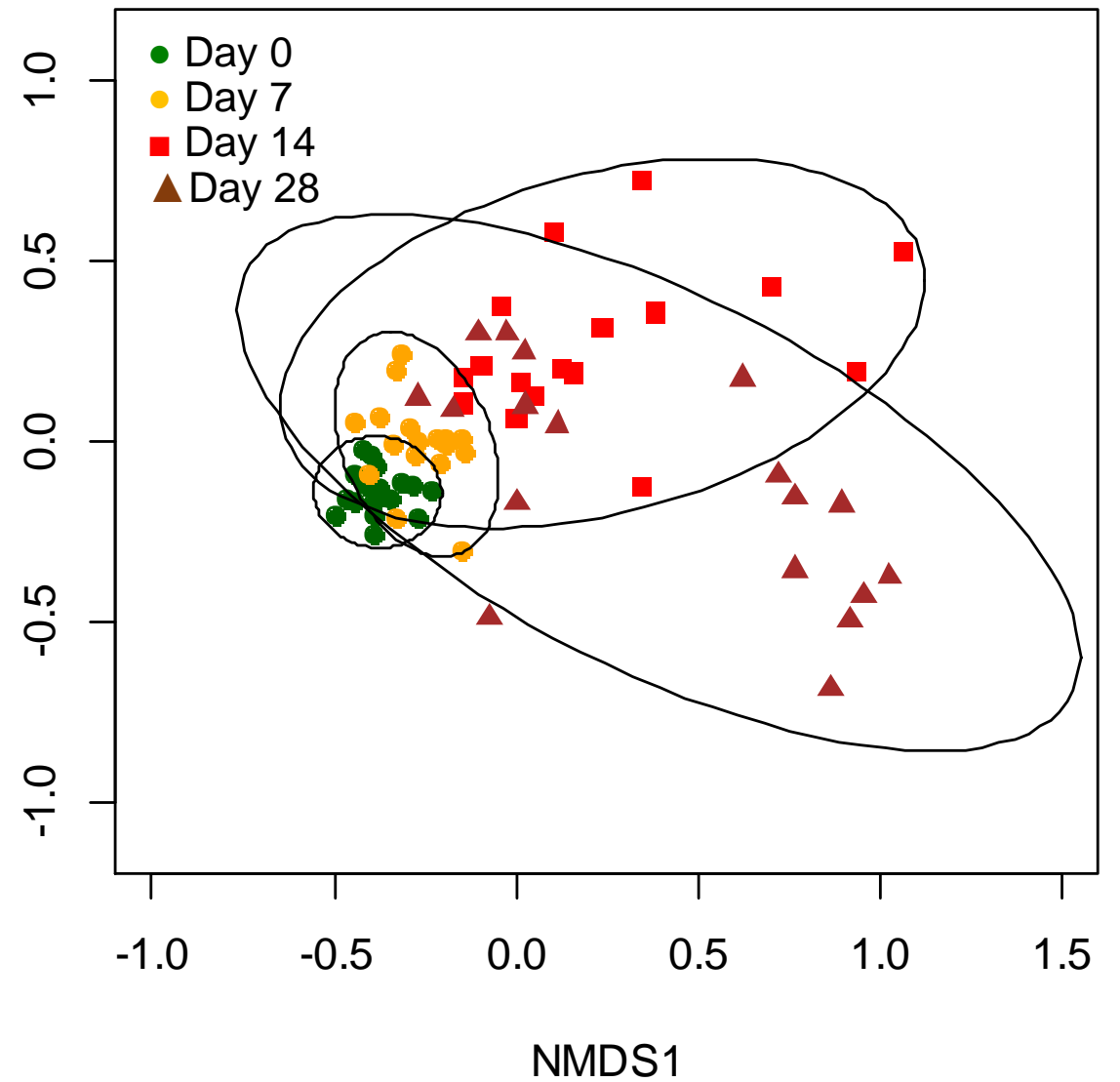
Day: $P < 0.001$, $R = 0.40$

The general structure of bacterial community changed over time as a result of the concentrate inclusion

Results- Spatial ordination

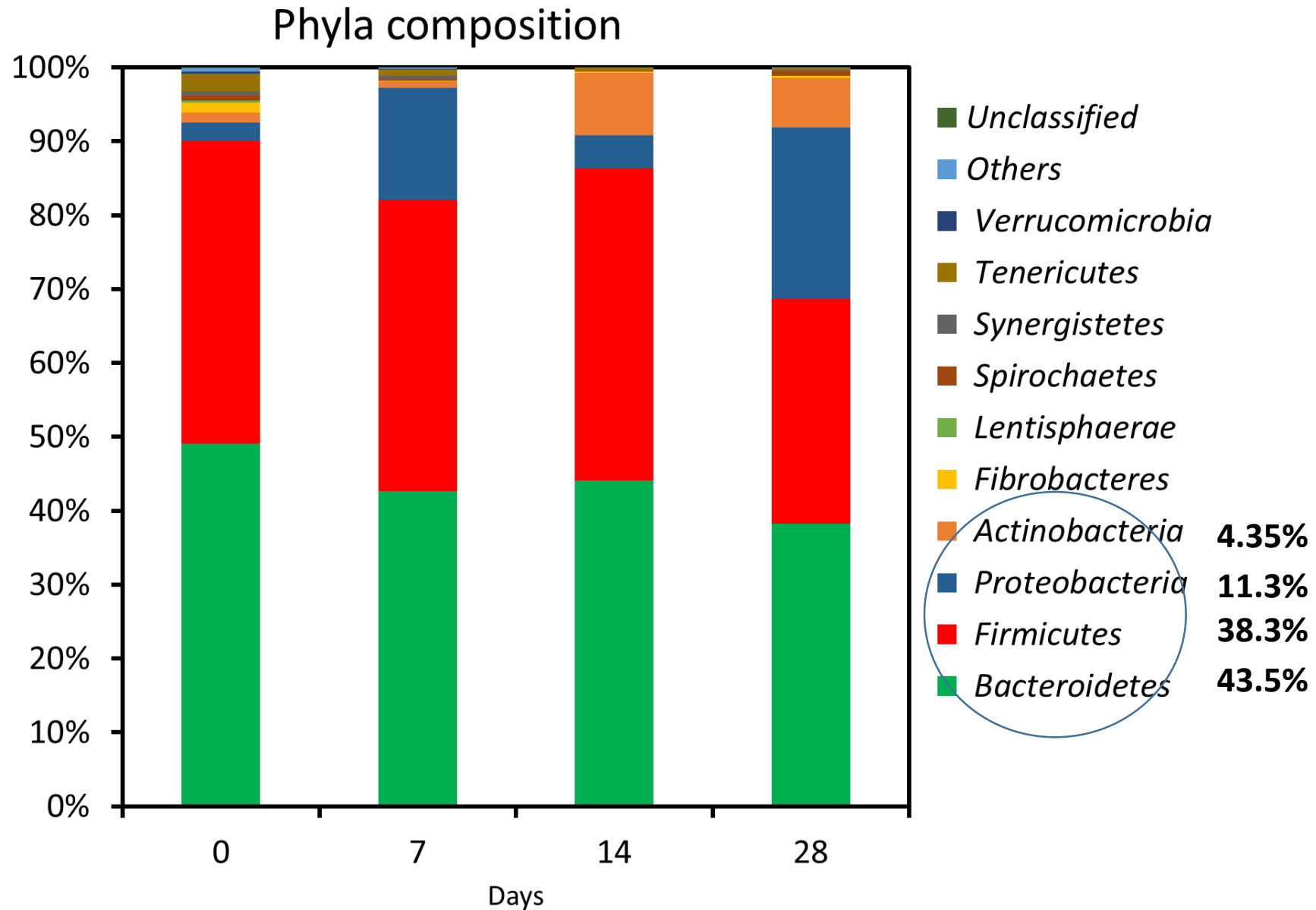


Stress 0.13



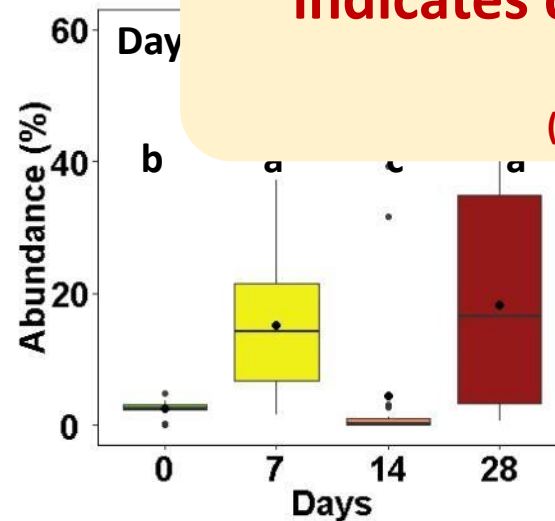
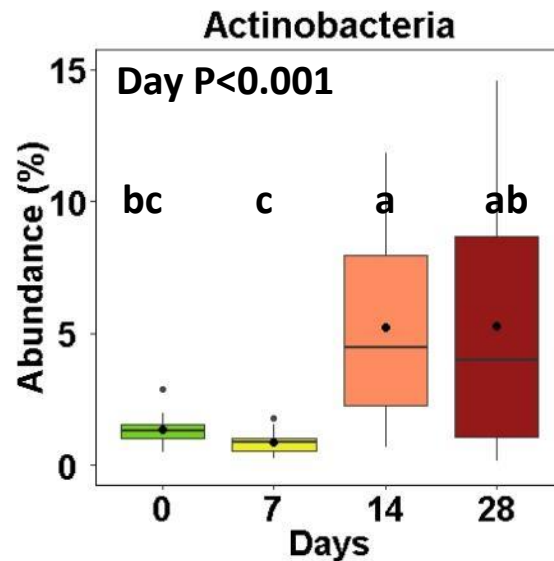
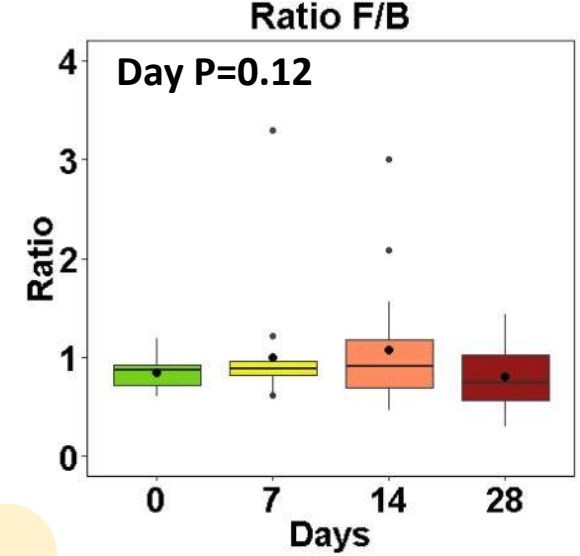
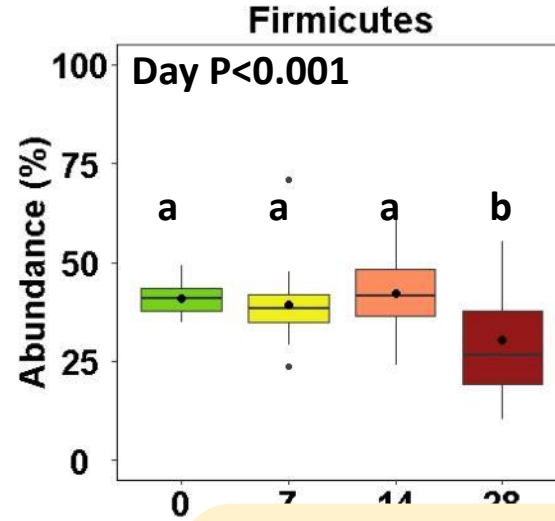
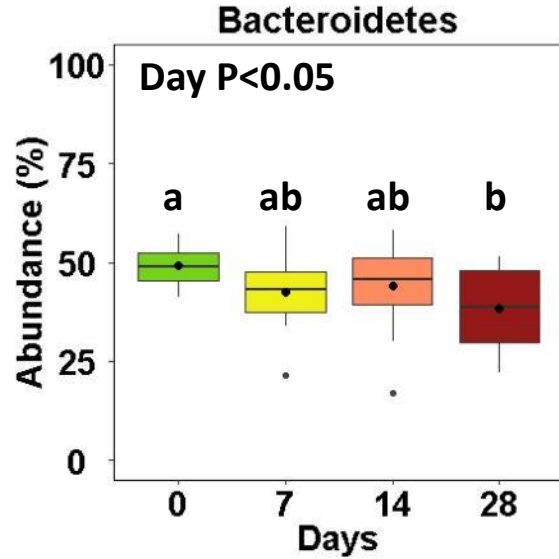
Ellipses show the confidence interval of each group at a 95% level

Results- Taxonomic composition



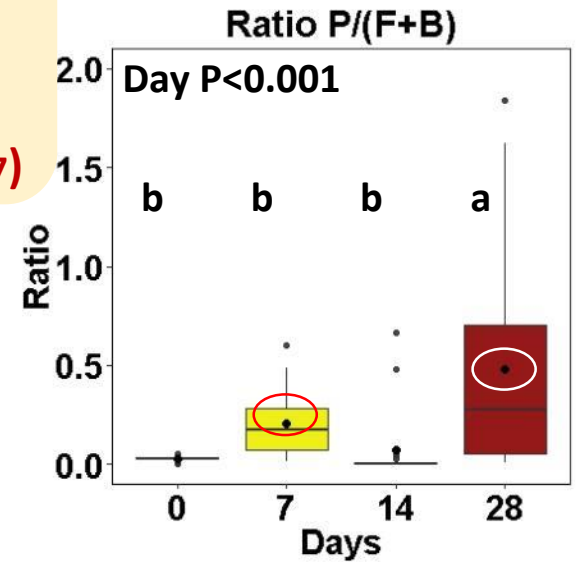
Results- Taxonomic composition

Diet P=NS
Day P<0.001



Ratio P/(F+B) > 0.19
indicates dysbiosis

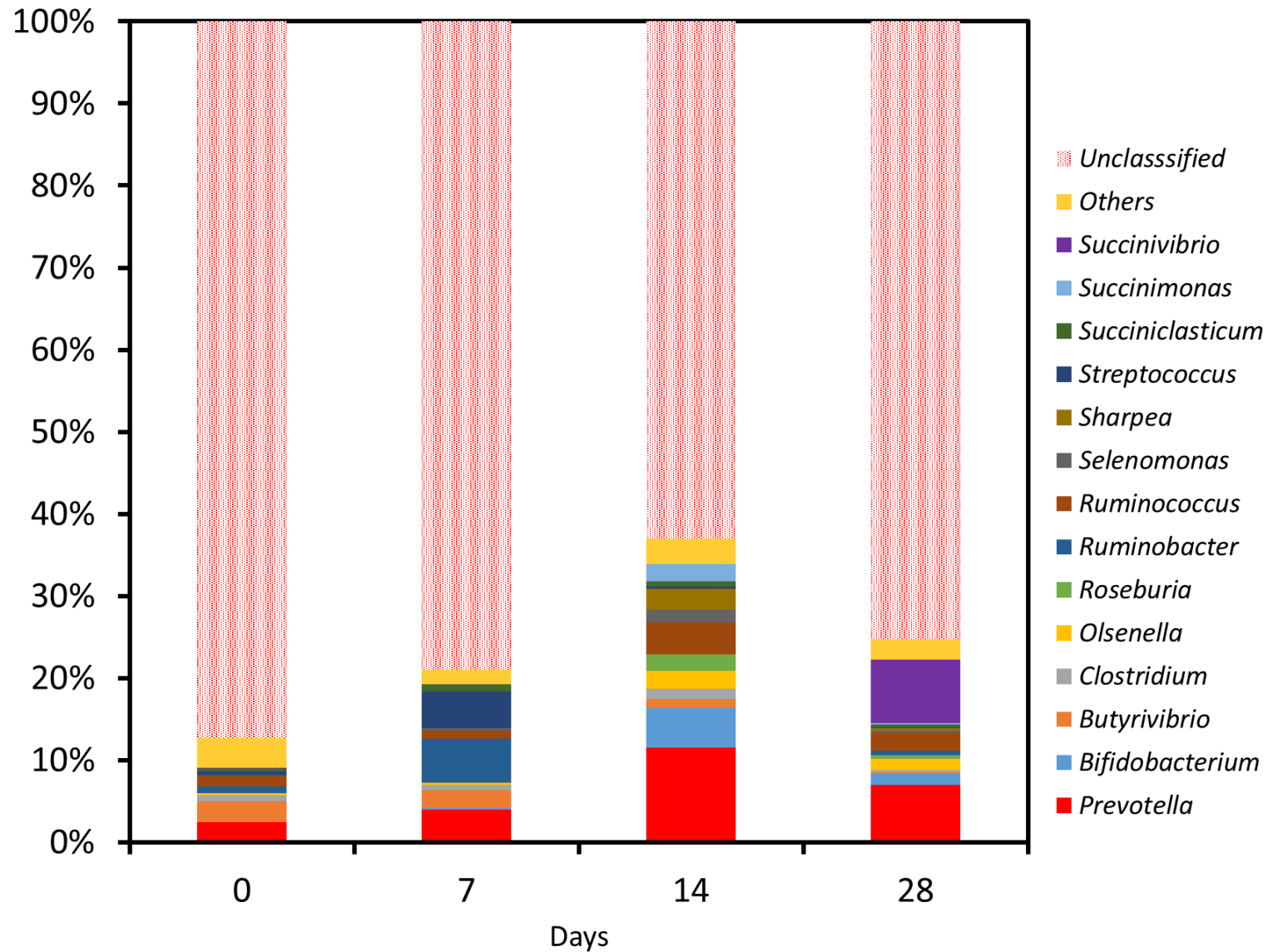
(Auffret et al., 2017)



ANOVA was performed with Log_{10} -transformed data.

Results- Taxonomic composition

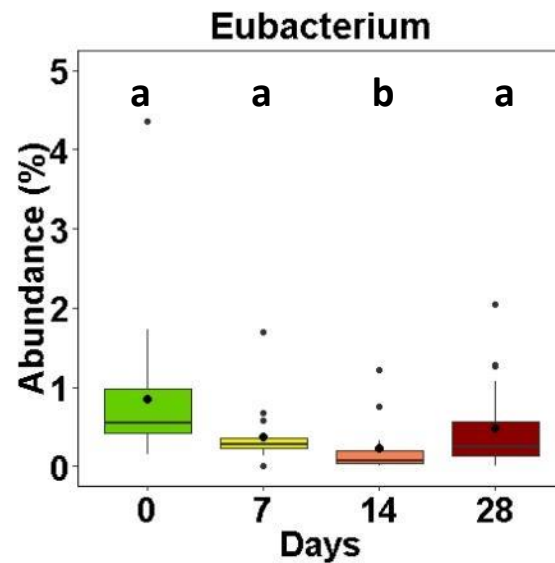
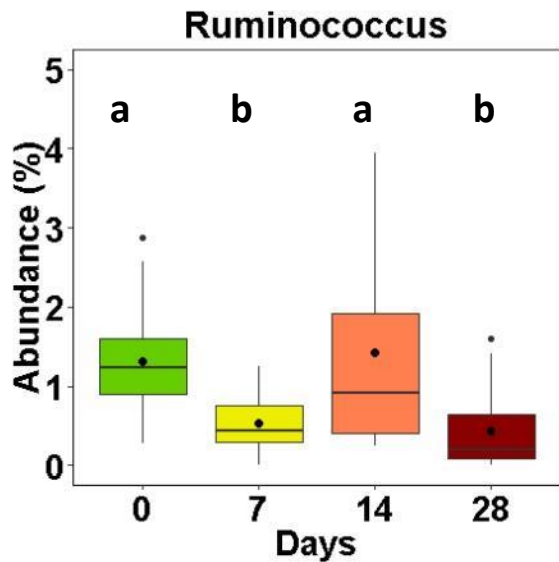
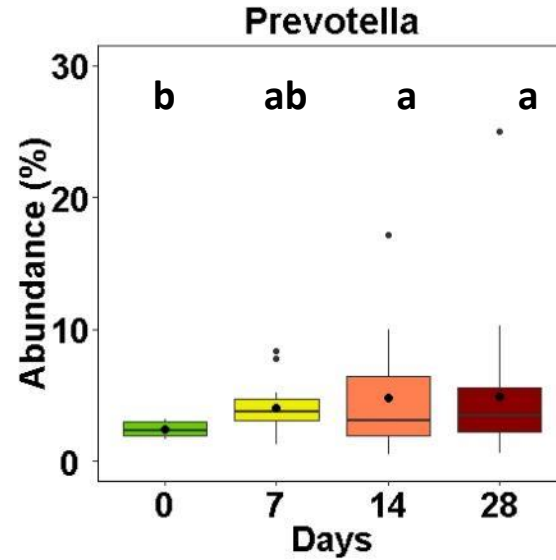
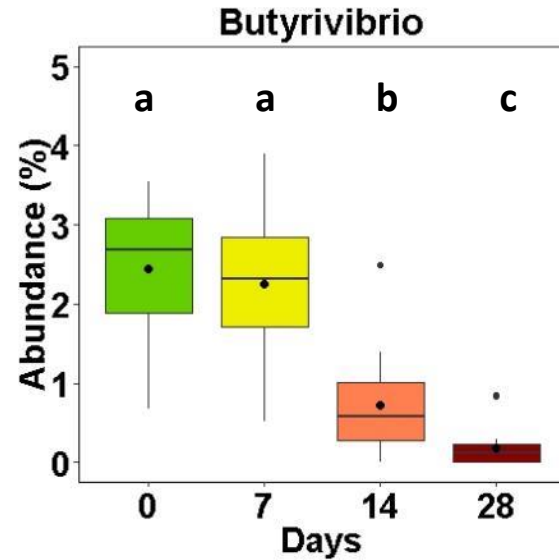
Genera distribution



Results- Taxonomic composition

Diet: P=NS

Day: P<0.05 in 95% of the genera



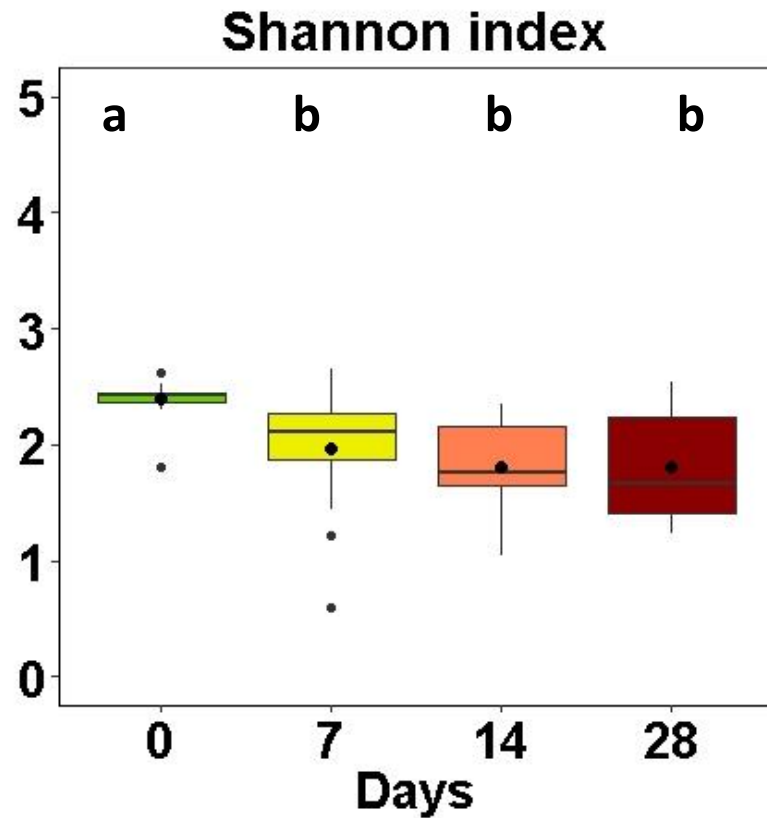
↑
Prevotella,
Ruminobacter,
Succinivibrio,
Selenomonas,
Roseburia

↓
Butyrivibrio,
Pseudobutyrvibrio,
Fibrobacter,
Ruminococcus

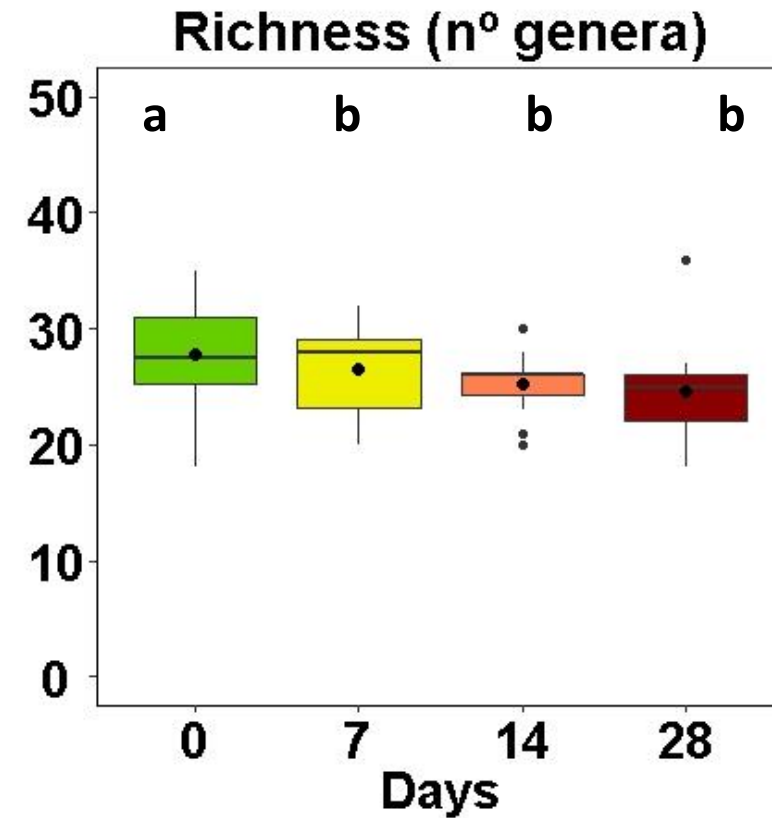
★
Bifidobacterium
Sharpea
Lactobacillus

ANOVA was performed with Log_{10} -transformed data

Results- Diversity



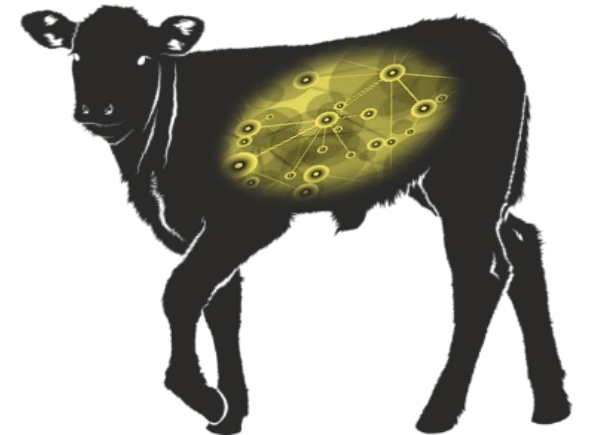
Diet: P=0.54
Day: P<0.01



Diet: P=0.48
Day: P<0.05

Conclusions

- Transition from a milk and grass diet to a high-concentrate ration strongly altered bacterial composition.
- Bacterial diversity decreased over time but some genera emerged after concentrate inclusion indicating a selection of more tolerant bacteria to the new environmental conditions.
- The inclusion of tannins and medium-chain fatty acids in the adaptation concentrate did not affect rumen bacterial community.



Thanks for your attention



Instituto Universitario de Investigación Mixto
Agroalimentario de Aragón
Universidad Zaragoza



Departamento de Innovación,
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