

# Maternal Influences on the Calf Rumen Microbiome and Subsequent Impacts on Performance and Efficiency

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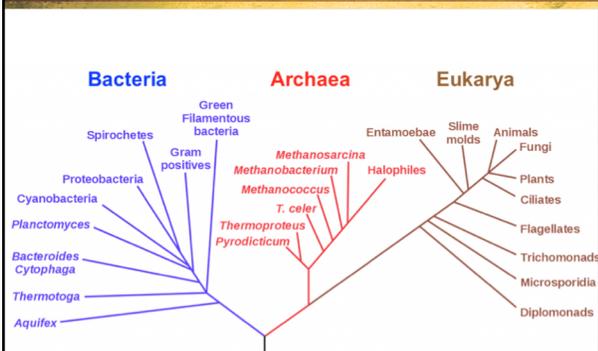
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## The Rumen Microbiome

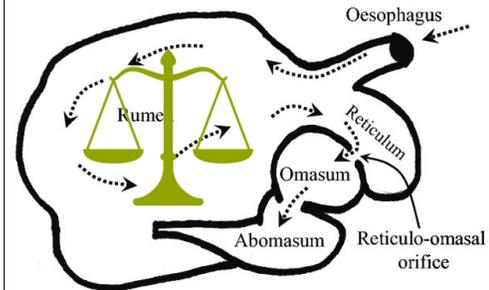
- What is it?
  - The microbial community (and its genetic material) that inhabits the rumen
- 1950's – Robert Hungate



## Types of Rumen Microbiota



## The "Healthy" Rumen Microbiome



## Why do we care?



## What is our goal?

- How is the rumen colonized?
- What factors affect colonization?
- What makes for a "healthier", more efficient, and productive host?
- Can we program the microbiome?



## Inoculation of the Rumen

- Believed to be “sterile” at birth
- Source of inoculation
  - Birth canal
  - “Maternal” factors
    - Licking, suckling, etc.
    - Colostrum
  - Consumption of bedding, hay, etc.
- What about pre-parturition factors?

## Pre-Parturition Factors

Aagaard et al., 2016

## Evidence in Ruminants

### Fecal/Meconium

### Rumen

Alipour et al., 2018 Malmuthuge et al., 2019

## Importance of the Early Microbiome

- Setting up fermentation patterns
- Production of VFA that are critical for rumen development (especially the papillae; Church, 1988)
- Immune system
  - Ruminal microbe/immune cell homeostasis (Garcia et al., 2017)
- Optimum stage for intervention?
  - Programming potential (Abecia et al., 2014a, 2014b, 2013; Yáñez-Ruiz et al., 2010)

Malmuthuge et al., 2019

## Establishment of the Rumen Microbiome

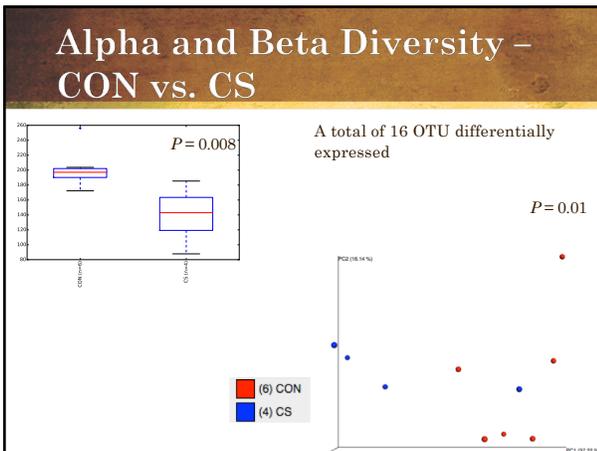
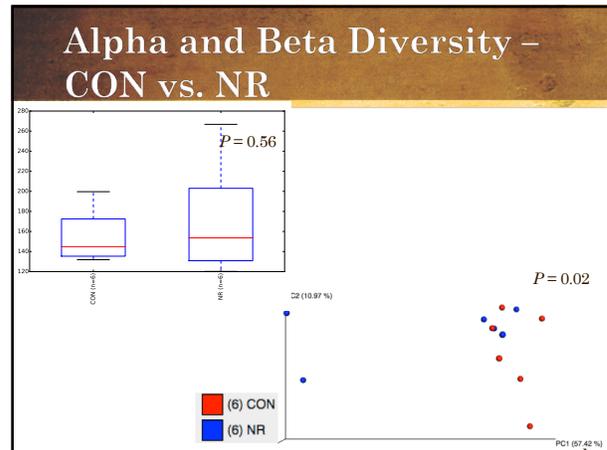
## What we have done thus far...

- Three studies
  - 1) Maternal Gestational Nutrition & Mode of Delivery
  - 2) Maternal Genotype & Pre- and Post-Parturition Factors
  - 3) Survey of the maternal microbiomes

### \*\*A brief methodology

- Collect rumen fluid
  - Oral lavage
- Isolate *microbial* DNA
- Sequence DNA
- Analyze sequences
  - Alpha-diversity
    - How "rich" or "even"
  - Beta-diversity
    - What does the composition look like?
  - Taxa abundances





### Maternal Breed – Two Divergent Breeds

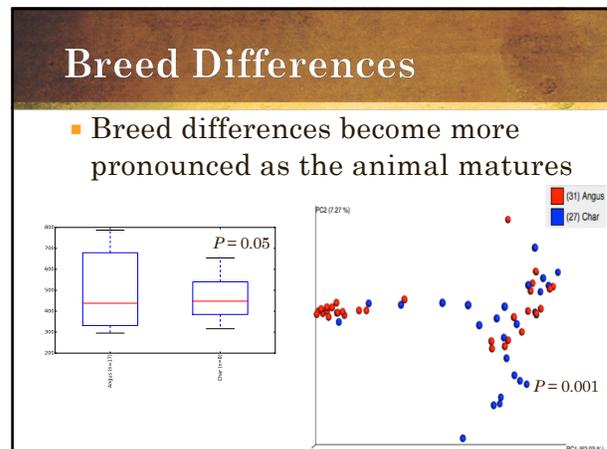
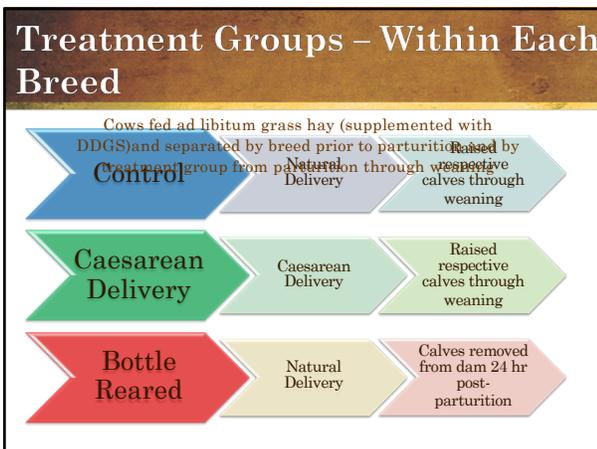


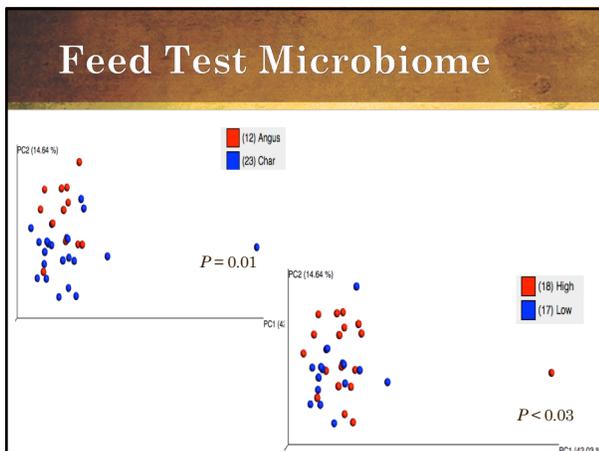
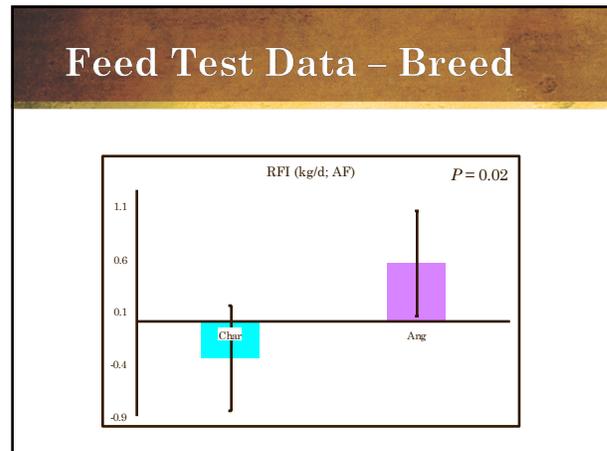
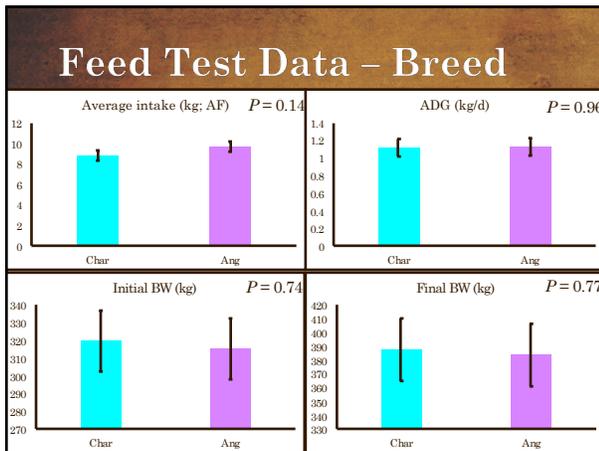
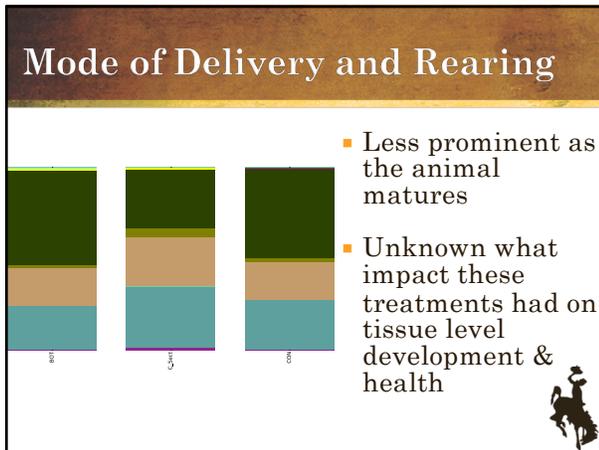
VS.

n = 42 (CON) vs. n = 40 (NR)

USDA NIFA

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### Conclusions & Implications

- Day is a driving factor in rumen physiology
  - Microbiome
    - Abundance, alpha- and beta-diversity
- Maternal factors affect the rumen microbiome, microbial fermentation, and performance
  - Mode of delivery and rearing type → some differences persist
  - Maternal breed → more pronounced as animal matures
  - Gestational nutrition → at least early on, calves microbiomes differ
- Potential for programming and intervention
  - But how does colonization happen?

## Ongoing Study

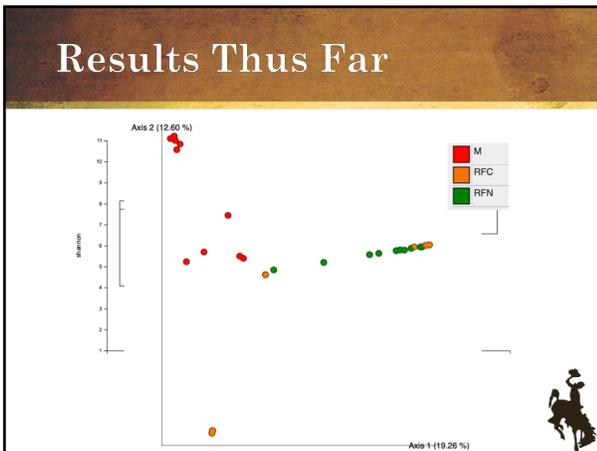


- Survey of maternal “sources” of inoculation to the rumen microbiome
  - Objectives:
    - 1) To evaluate several maternal microbiomes yet to be evaluated in beef cattle
    - 2) To compare those maternal microbiomes with the developing calf gut microbiome



## Methods – Cow Samples

- Prior to calving:
  - Rumen fluid
  - Vaginal Swab
  - Amniotic Fluid
- After calving:
  - Placental samples
  - Calf samples
    - Meconium, blood, rumen fluid
    - Feed efficiency



## Our objective:

- Phase 1: Investigate potential mechanisms of colonization/inoculation*
- Phase 2: Investigate intervention strategies that can alter/program the microbiome*



## Thank you!

- Kathy Austin
- Gwen Hummel
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- Ryan Knuth
- Beef Unit Crew
  - Travis Smith
- Dr. Kristi Cammack

