

Influence of feed efficiency and physiological state on rumen VFA and microbial profiles in cattle

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Y Montanholi²



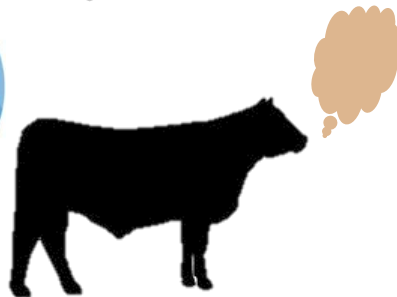
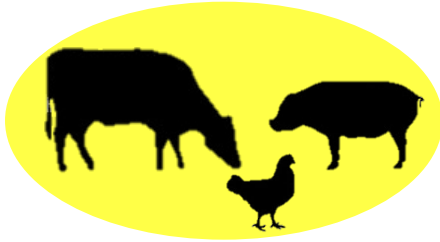
Outline

- Introduction
- Hypothesis
- Objectives
- Materials and methods
 - Grain-fed study
 - Grass-fed study
- Results and discussion
- Conclusion
- Acknowledgements



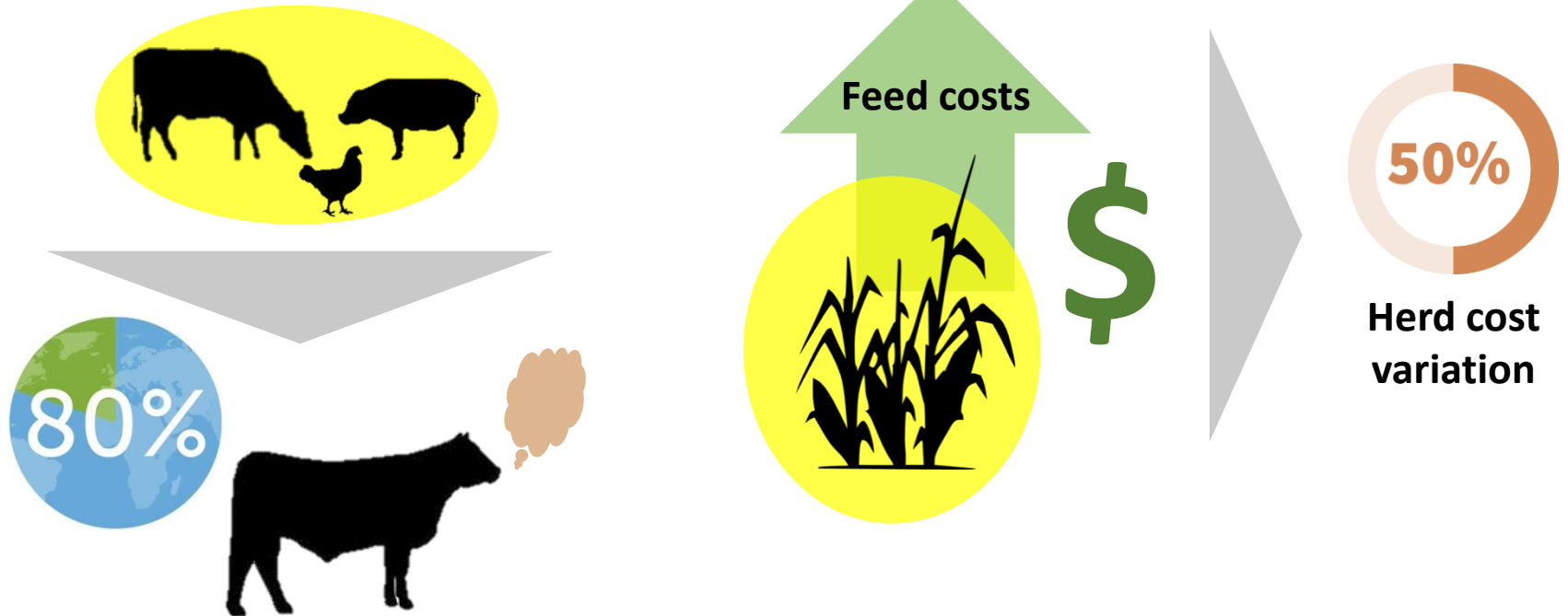
Introduction: Industry challenges

30-50% agriculture GHG



Introduction: Industry challenges

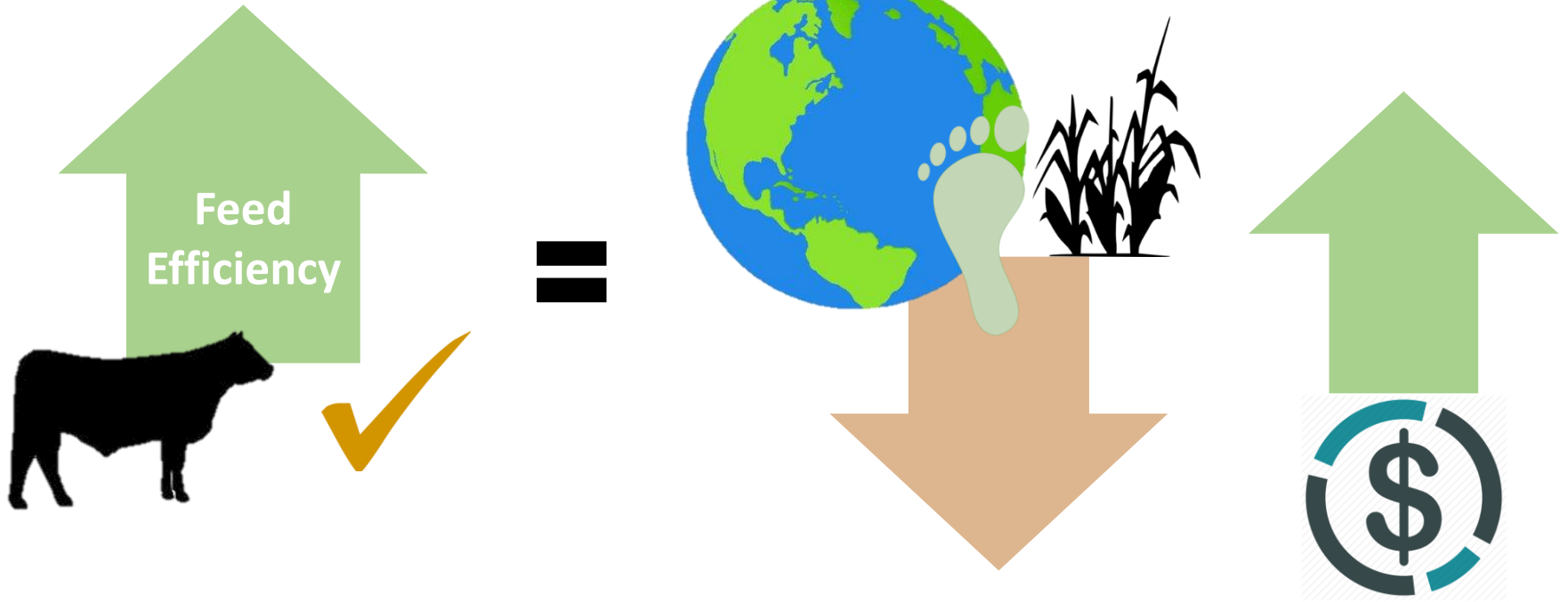
30-50% agriculture GHG



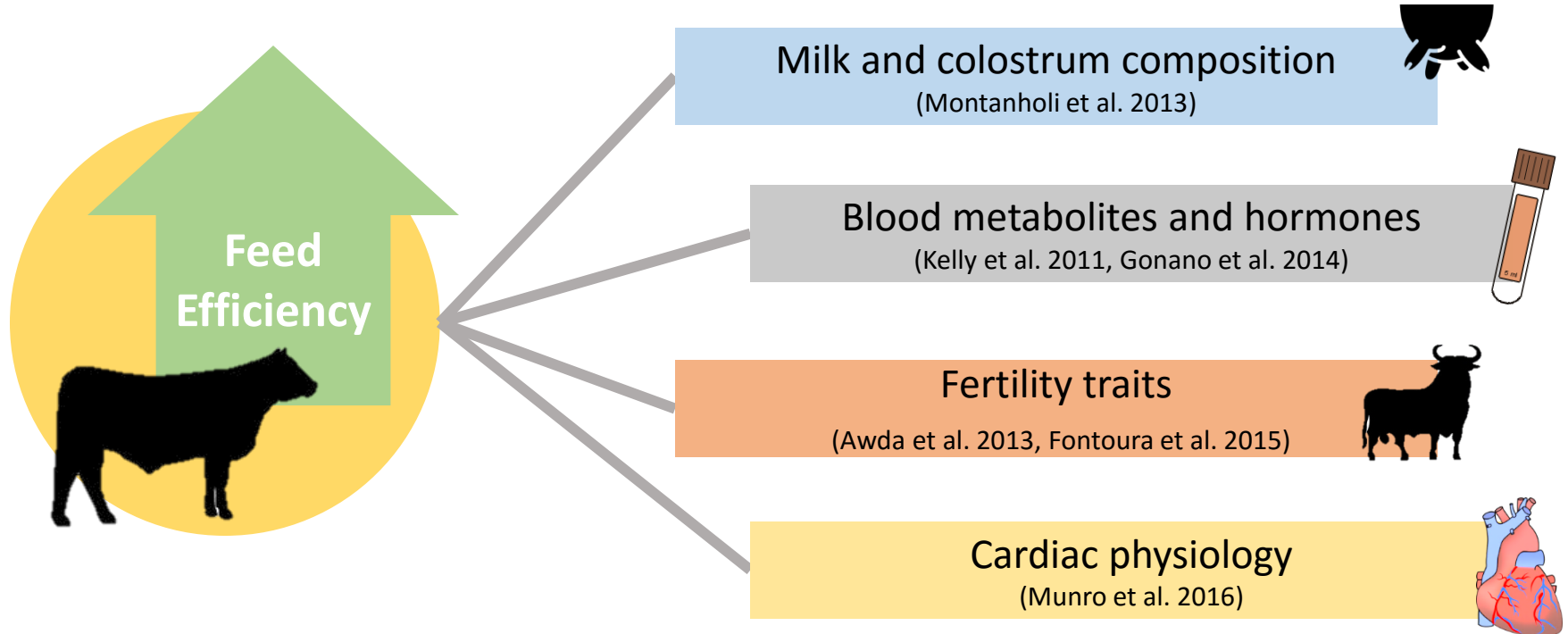
Introduction: Feed efficiency



Introduction: Feed efficiency



Introduction: Feed efficiency



Introduction: Rumen metabolism

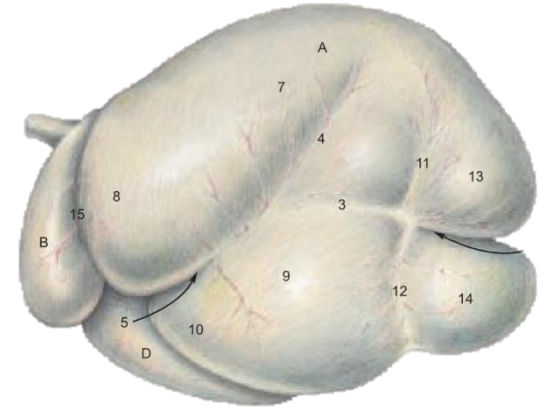
% 10% of the biological variation of RFI due to digestibility
(Richardson and Herd 2004)

Rumen and reticulum = 75% total digestive tract
(Baldwin 1980)



Large energetic sink with high energy demand
(Hungate 1960)

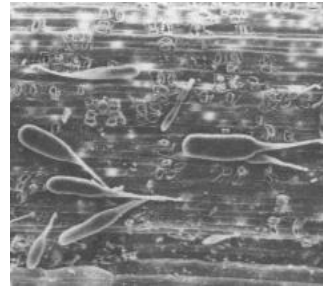
Area of energy absorption - 75% total VFA
(Bergman 1990)



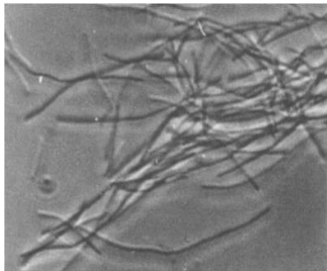
Introduction: Microbiology



Bacteria (10^{11} cells/ml)
75% feed particle digestion



Fungi (10^3 - 10^4 cells/ml)
Fibrolytic particle digestion



Methanogens (10^4 - 10^6 cells/ml)
Methanogen ecology associated with methane emissions



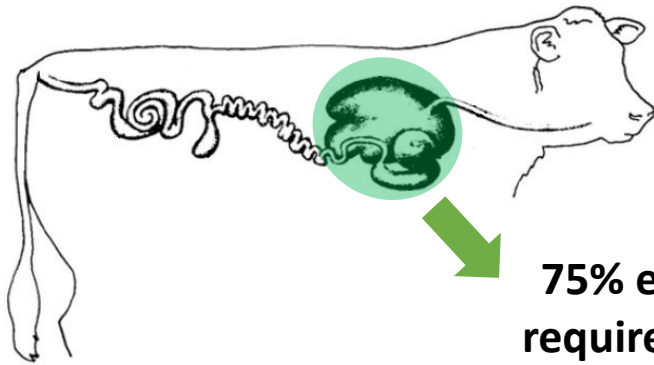
Protozoa (10^4 - 10^6 cells/ml)
Ciliate species digesting suspended and colonized feed particles



Feed efficiency

(Guan et al. 2008, Zhou et al. 2009, Hernandez et al. 2012, Carberry et al. 2014)

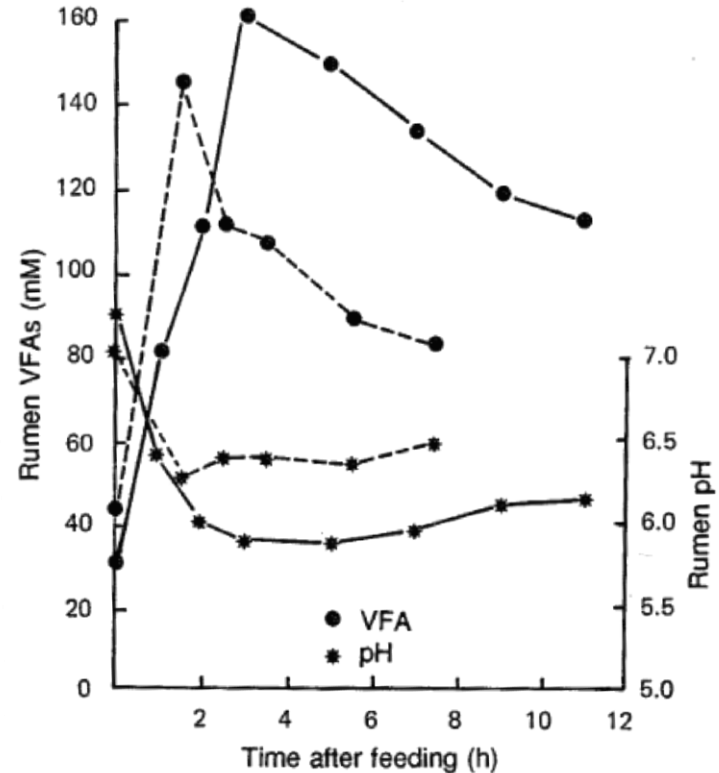
Introduction: Volatile fatty acids (VFA)



75% energy requirements



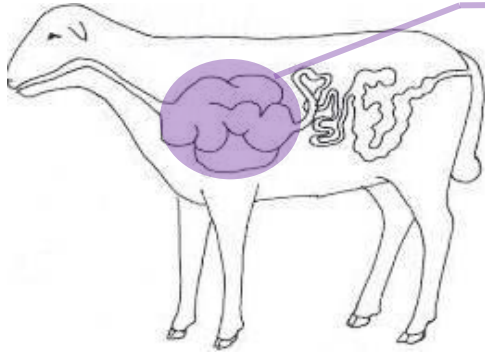
Feed efficiency



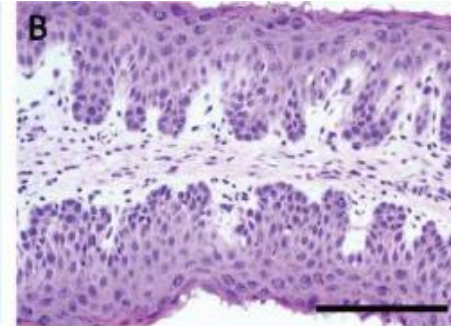
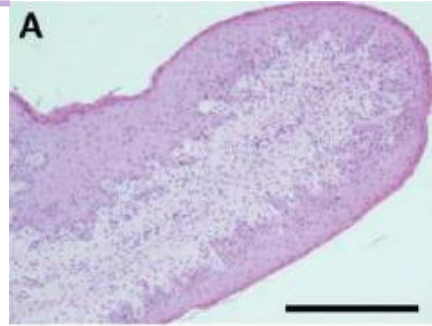
(Briggs et al. 1957, Bergman et al. 1990)

Introduction: Rumen microstructure

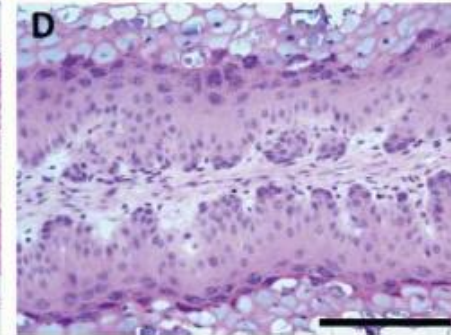
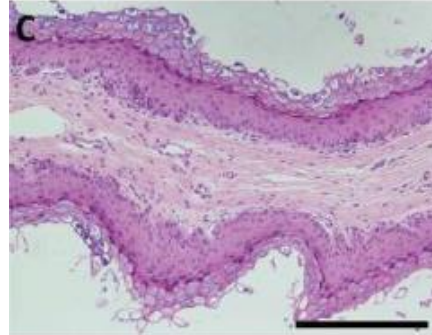
Sheep papillae microstructure



Low energy diet:



High energy diet:



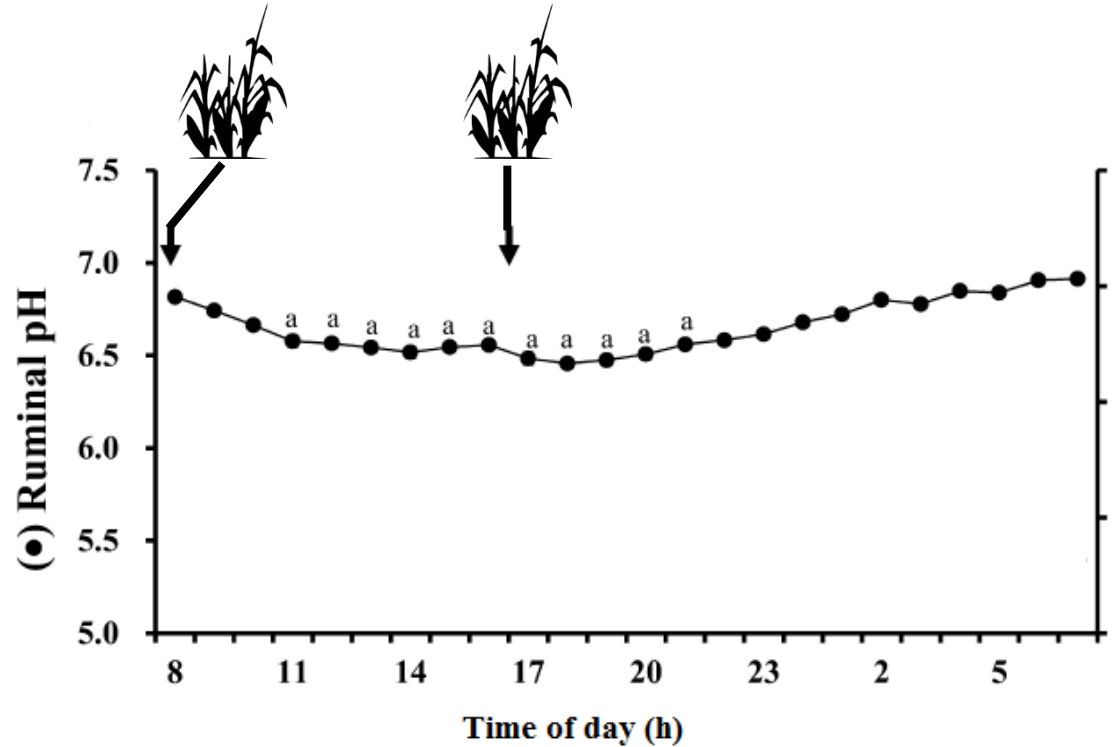
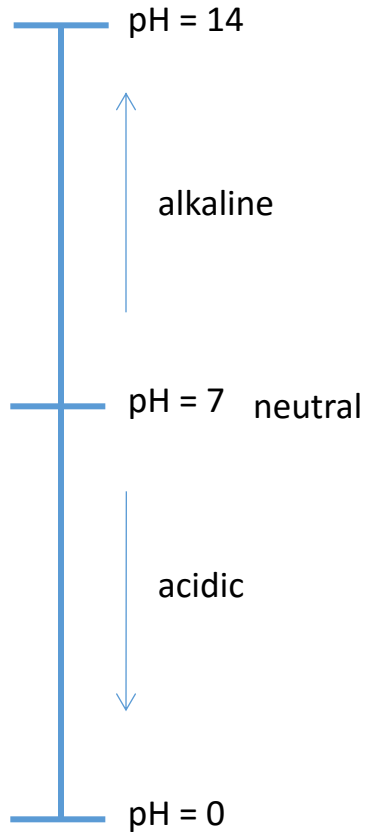
Stratum corneum thickness



Energy in diet

(Steele et al. 2012)

Introduction: Rumen pH



Hypothesis

Feed efficiency is associated with energetic processes and the rumen is a highly metabolically active organ. Therefore, the variability in rumen metabolism across feed efficiency phenotypes and dietary treatments may be featured through rumen functional and structural assessments.

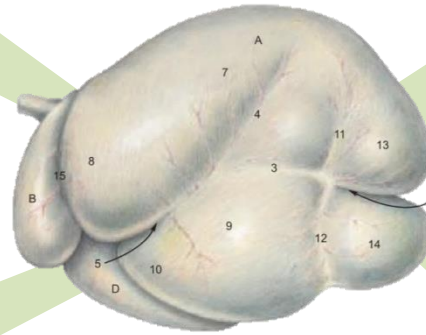
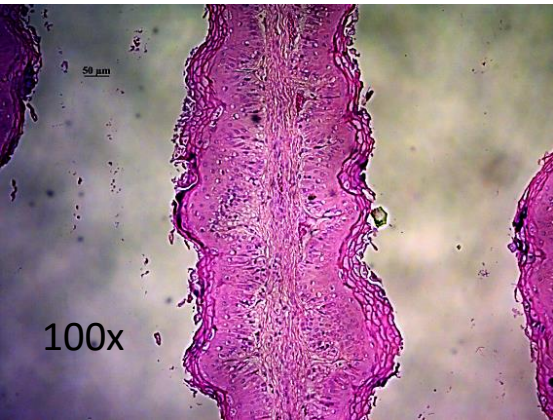
Objectives

Microbiology



Volatile fatty acids (VFA)

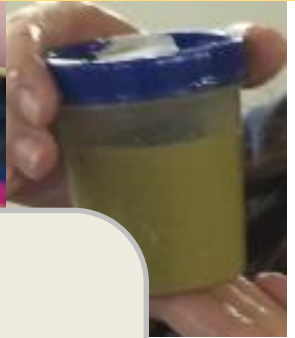
Papillae epithelium



Rumen pH

Objectives

Microbiology

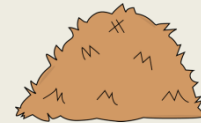


Efficient vs Inefficient

Grain-fed animals

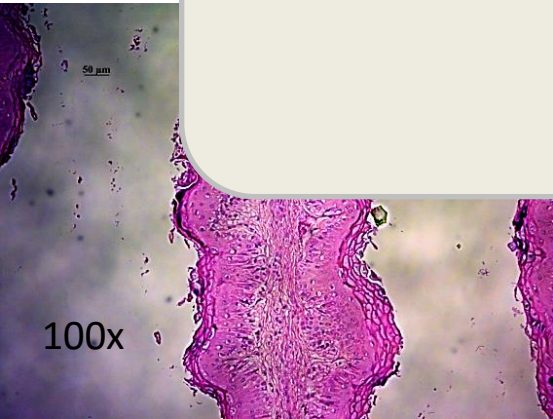


Grass-fed animals

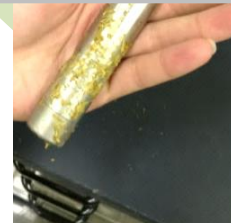


(VFA)

Pa



100x



Rumen pH

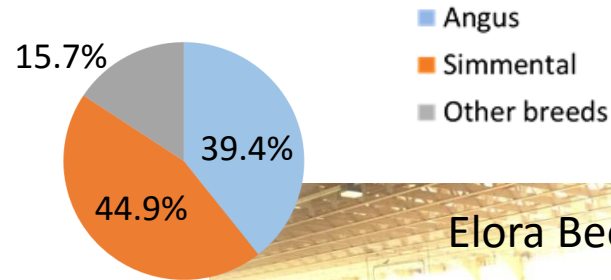
Materials: Experimental conditions



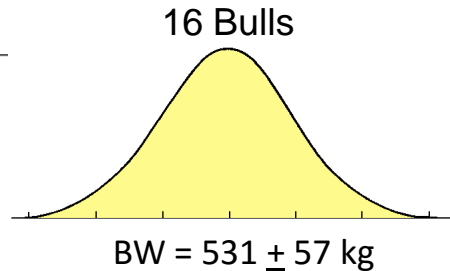
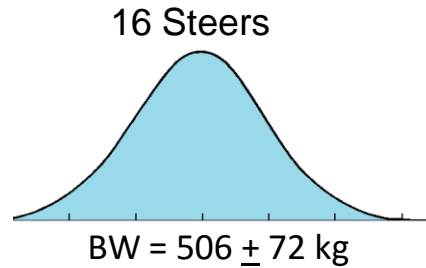
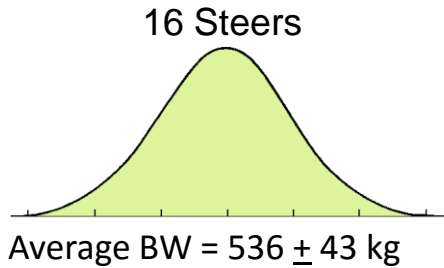
Feedlot study

- 48 crossbred cattle
- Trial length: 112 d

Overall breed composition



Elora Beef Research Centre



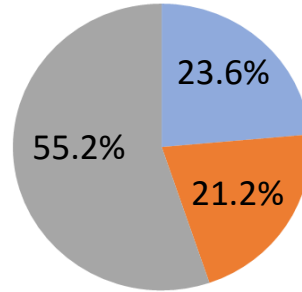
Materials: Experimental conditions



Grass-fed study

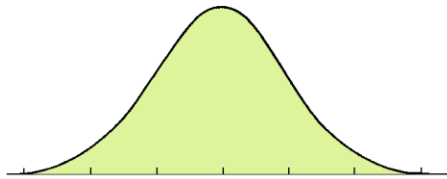
- 141 crossbred cattle
- Trial length: 124 d

Overall breed composition



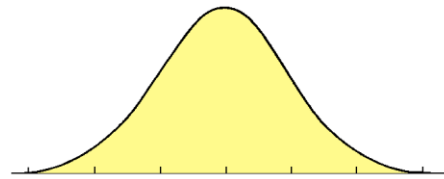
- Angus
- Simmental
- Other breeds

107 heifer calves



BW = 253 ± 38 kg
Age = 403 ± 27 d

36 pregnant heifers

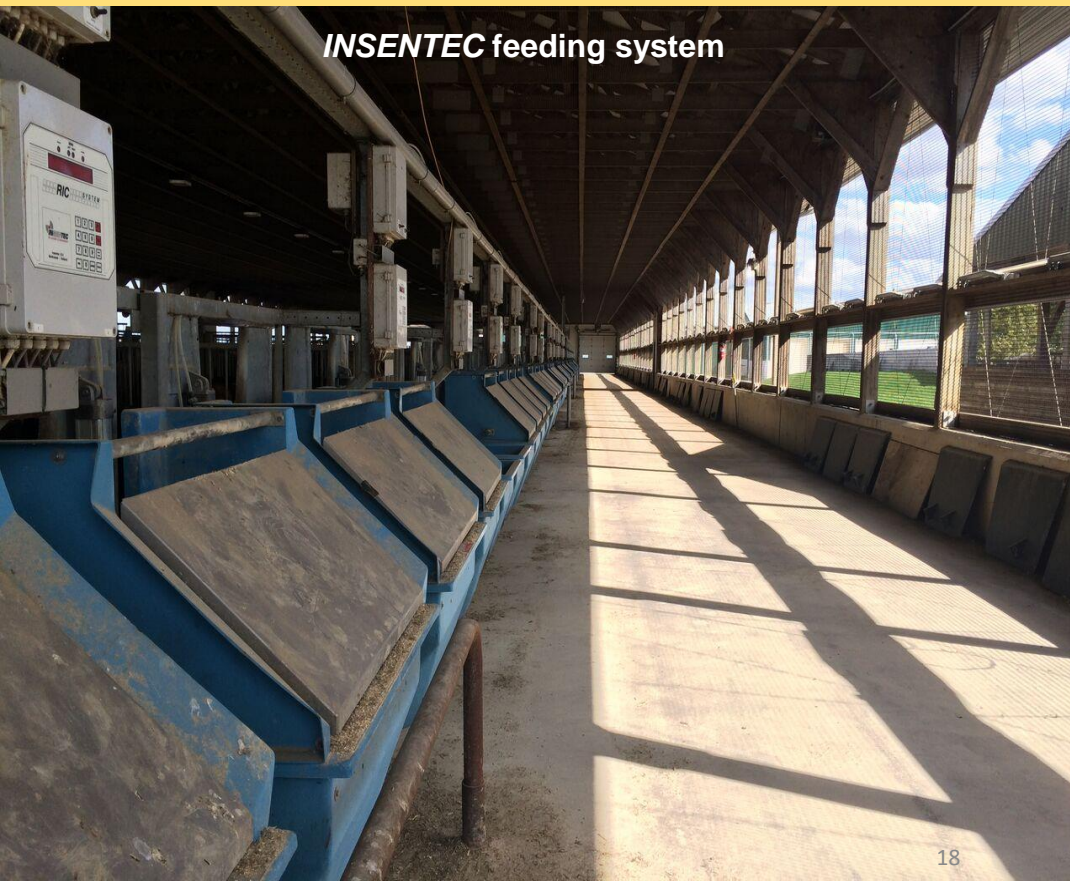


BW = 406 ± 42 kg
Age = 594 ± 95 d



Maritime Beef Test Station

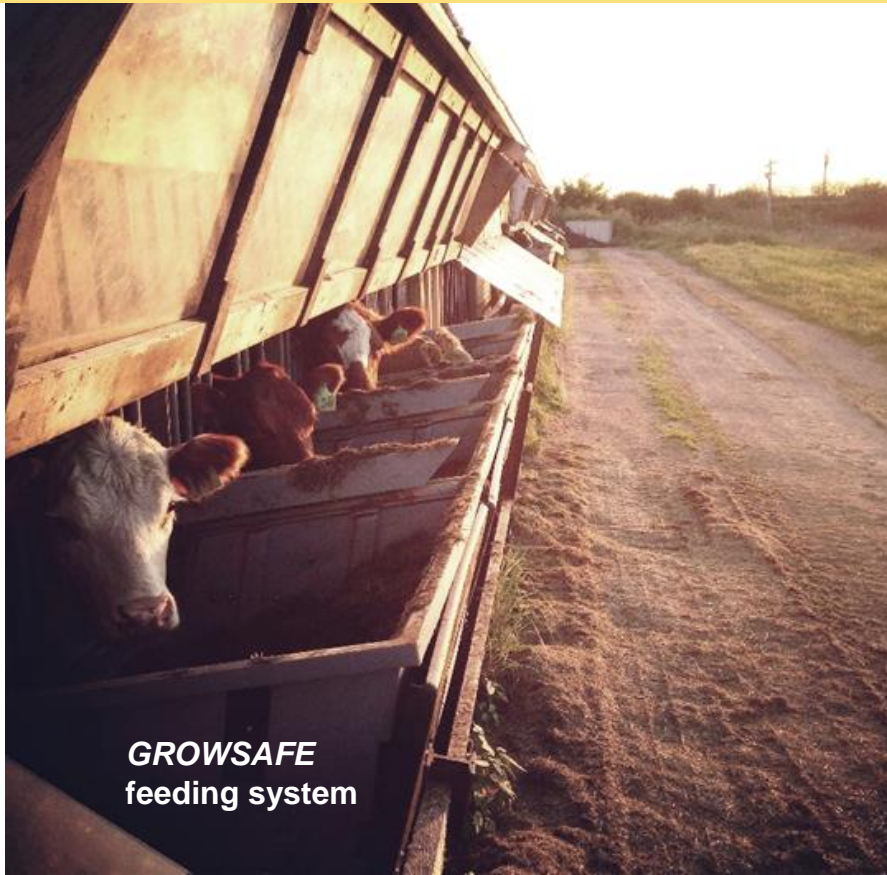
Methods: Diet



INSENTEC feeding system

<i>Chemical Composition</i>	<i>Dry Basis (%)</i>
Dry Matter	53.8
Crude Protein	13.9
Acid Detergent Fibre	10.9
Neutral Detergent Fibre	22.2
Starch	45.0
Total Digestible Nutrients	86.0
<i>Ingredient Composition</i>	<i>Dry Basis (%)</i>
High-moisture corn	52.5
Haylage	42.3
Soybean meal	3.5
Premix*	1.7

Methods: Diet

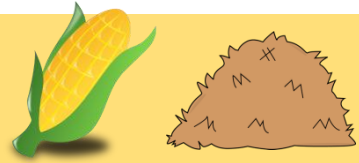


GROWSAFE
feeding system

<i>Chemical Composition</i>	<i>Dry Basis (%)</i>
Dry matter	36.1
Crude Protein	15.5
Acid Detergent Fibre	29.6
Neutral Detergent Fibre	53.7
Starch	6.5
Total Digestible Nutrients	70.3
<i>Ingredient Composition</i>	<i>Dry Basis (%)</i>
Haylage	99.5
Premix*	0.5

*Contains 37.4% of calcium phosphate, 62.7% trace mineralized salt

Methods: RFI models



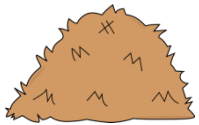
Grain-fed ($R^2 = 0.74$):



$$\text{Feed intake} = [\mu + (\beta_1 \times \text{body weight})] + (\beta_2 * \text{ADG}) - (\beta_3 * \text{ribeye area}) + (\beta_4 * \text{back fat}) - (\beta_5 * \text{marbling}) + \text{subpopulation} + \text{RFI}$$

Body Size *Rate of Gain* *Leanness*
Fatness

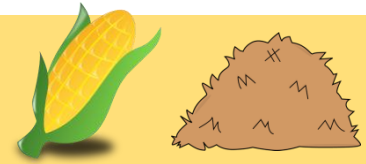
Grass-fed ($R^2 = 0.61$):



$$\text{Feed intake} = [\mu + (\beta_1 * \text{body weight})] + (\beta_2 * \text{ADG}) - (\beta_3 * \text{ribeye area}) + (\beta_4 * \text{rump fat}) - (\beta_5 * \text{age}) + \text{subpopulation} + \text{RFI}$$

Body Size *Rate of Gain* *Leanness*
Fatness

Methods: Sample collection



5.5±1 d prior slaughter



pH logger insertion



End of performance test

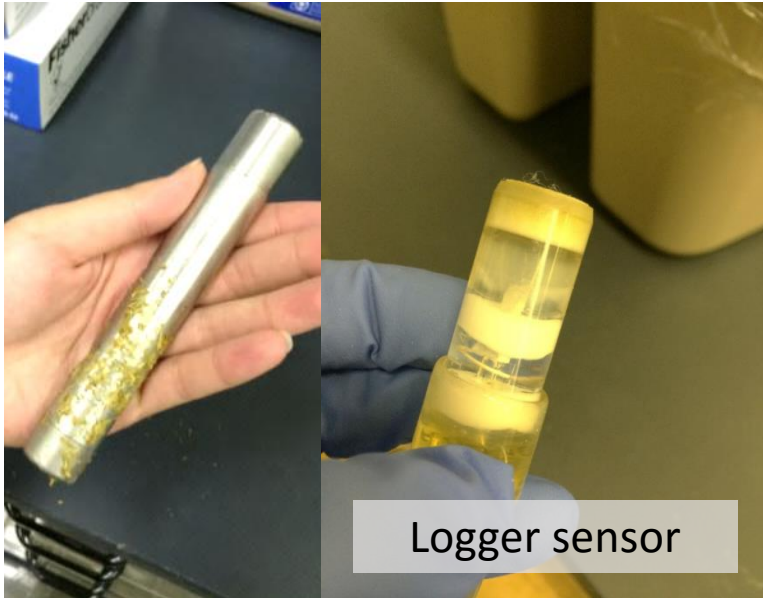


Rumen fluid



Rumen tissue

Methods: Logger insertion



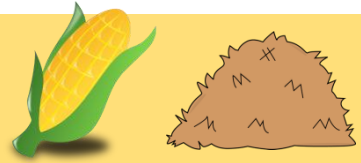
Rumen pH loggers

T9 LRCpH Data Logger Dascor

Method: esophageal tubing

Recording: 5 minute intervals
(~2,600 data points/animal)

Methods: Rumen fluid collection



Method:

Oro-ruminal probe with suction

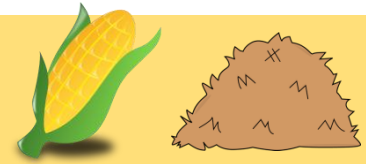
Evaluating:

Microbiology

Volatile fatty acid profiles



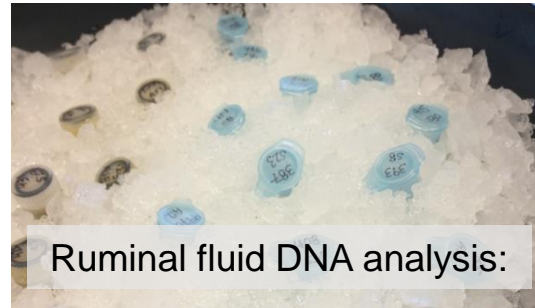
Methods: Microbiology



Method:

Rumen fluid DNA isolation

RT-qPCR



Evaluating:

Bacteria



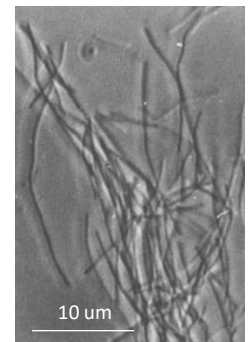
Protozoa



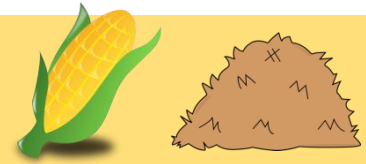
Fungi



Methanogen



Methods: VFA



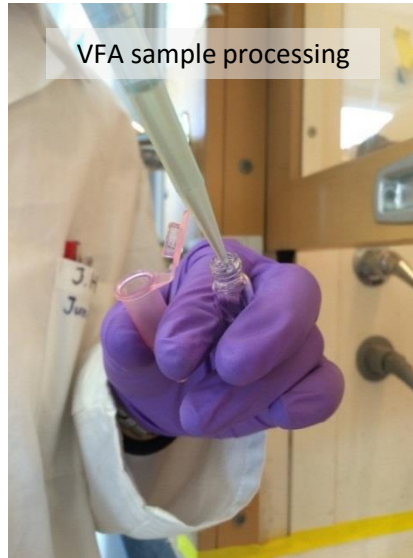
Evaluated:

VFA molar concentrations

- **Acetate**
- **Propionate**
- **Butyrate**
- Valerate
- Isovalerate
- Isobutyrate
- Caproate

Total VFA concentration

Method:



Methods: Rumen tissue collection

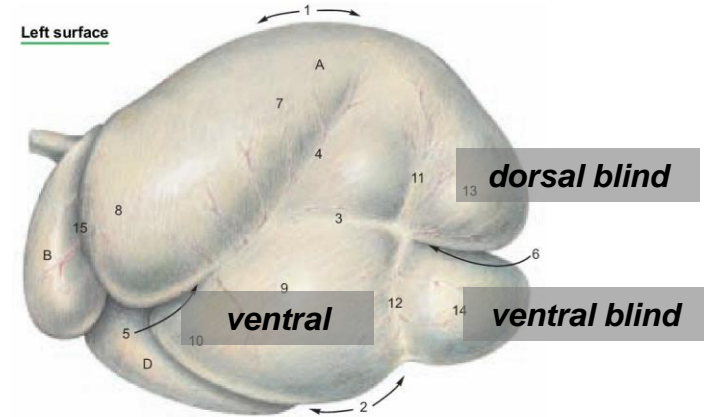


Method:

Tissue collection

Processed for histomorphology

Evaluating: Papillae epithelial thickness

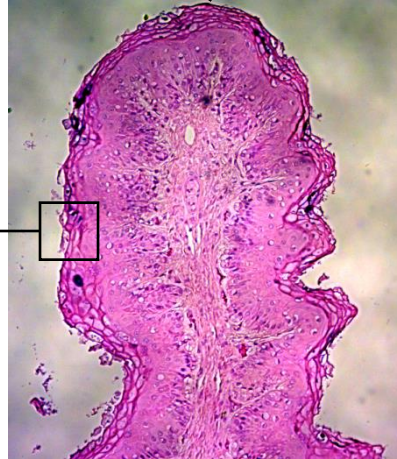
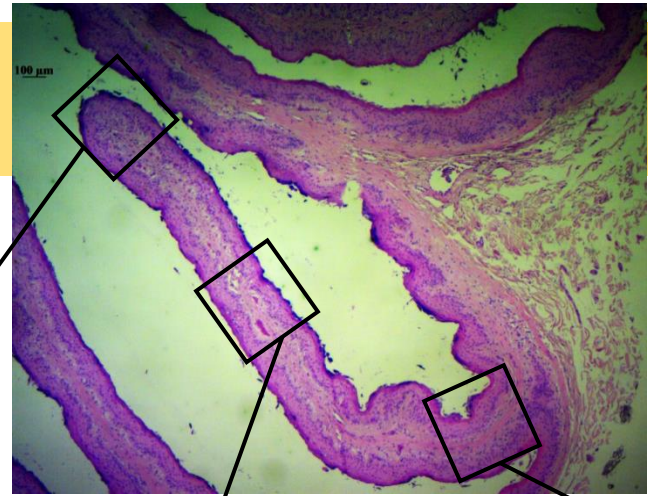


Methods: Histomorphometry

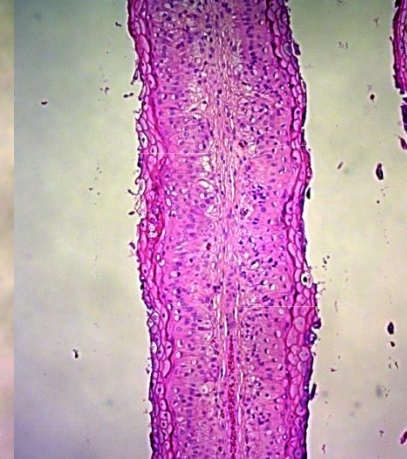


Histology traits:

- Stratum corneum ●
- Papillae width ●



Tip

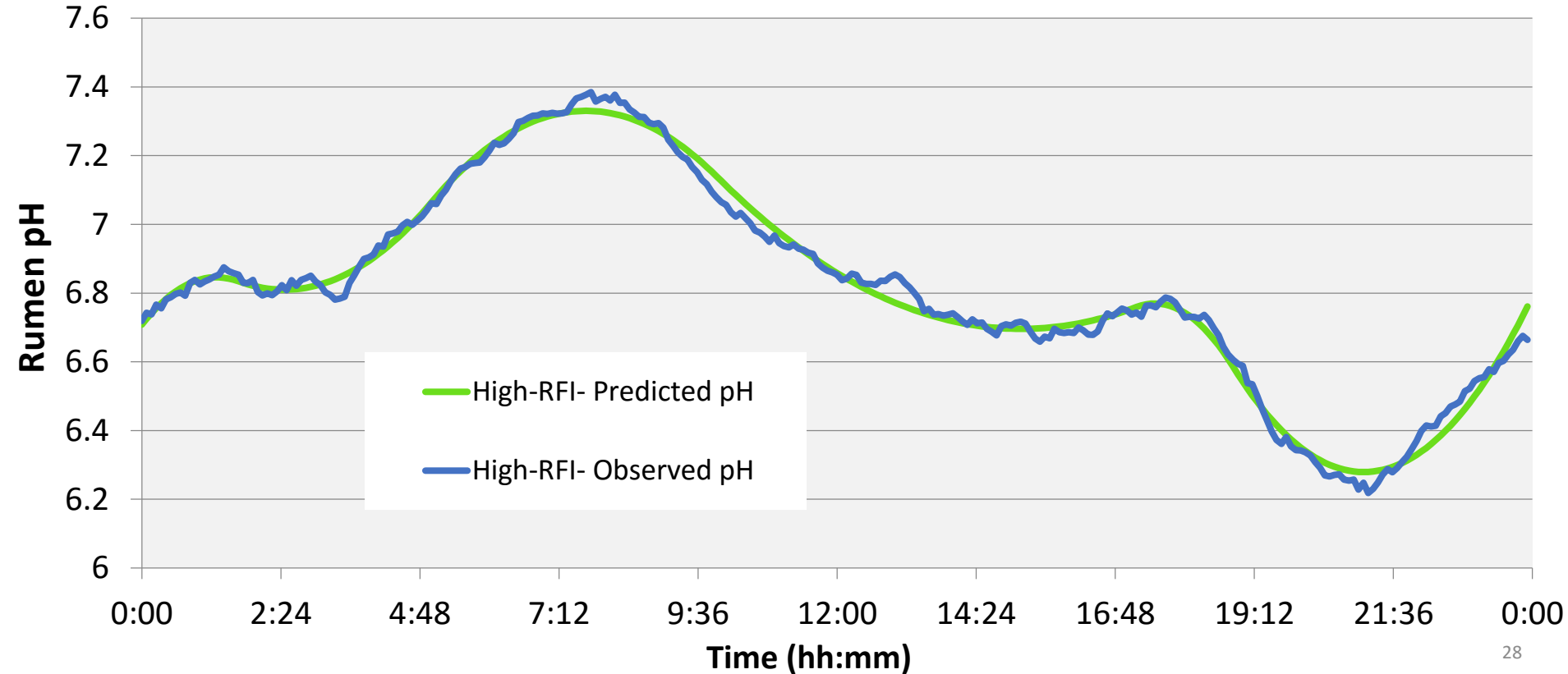


Middle piece

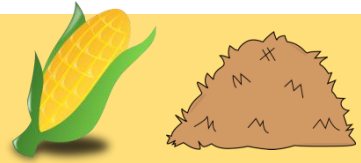


Base

Methods: pH measurements



Methods: Statistical analysis



Univariate Normality Procedure

- Skewness, Kurtosis, Anderson-Darling Test
- Transformations
 - logarithm
 - squared

GLM Select procedure

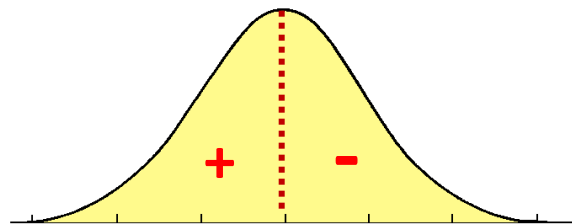
- Determine model effects

General Linear Model (GLM) Procedure

- Rumen traits

Partial Least Square procedure

- Determine % contribution to RFI



50% Efficient vs. 50% Inefficient

$$Y_{ijkl} = \mu + \text{efficiency group}_i + \text{subpopulation}_j + \text{breeds}_k + \epsilon_{ijkl}$$

Results: Physiological status



Trait (%/total VFA)	Heifer calves	Pregnant heifers	<i>P</i>-value
Acetate	71.68	74.62	<0.01
Propionate	18.30	16.53	<0.01
Isobutyrate	0.97	0.89	0.051
Butyrate	7.29	6.47	<0.01
Isovalerate	0.90	0.90	0.952
Valerate	0.58	0.41	<0.01
Caproate	0.28	0.18	<0.01
Total VFA (μmol/ml)	43.52	37.39	<0.01

Results: Physiological status



Pregnant heifers



Bacteria population

Metabolic activity
throughout gestation

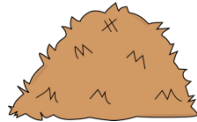
VFA metabolism and
energy demand

(Church, 1988; Drackley et al. 2001)

Trait (%/total VFA)	Heifer calves	Pregnant heifers	<i>P-value</i>
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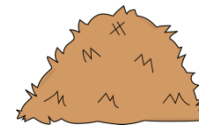
Results: RFI – Microbial profiles

Trait (16s rRNA copy/ml)	Inefficient	Efficient	<i>P</i> -value
Total bacteria	4.3x10 ¹¹	7.6x10 ¹¹	<0.05
Methanogen	4.9x10 ⁹	2.3x10 ⁹	<0.05
Protozoa	4.3x10 ⁷	1.5x10 ⁷	0.18
Fungi	6.3x10 ⁴	3.8x10 ⁴	0.37



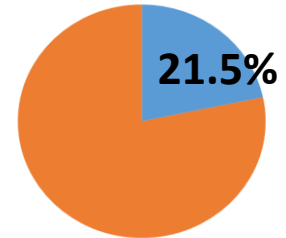
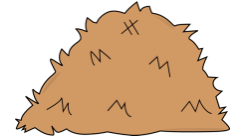
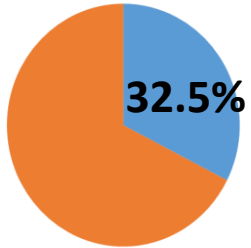
Trait (16s rRNA copy/ml)	Inefficient	Efficient	<i>P</i> -value
Total bacteria	6.0x10 ¹⁰	5.3x10 ¹⁰	0.16
Methanogen	2.6x10 ⁷	3.1x10 ⁷	<0.05
Protozoa	1.2x10 ⁵	1.6x10 ⁵	0.30
Fungi	1.3x10 ⁵	1.9x10 ⁵	<0.05

Results: RFI – VFA profiles

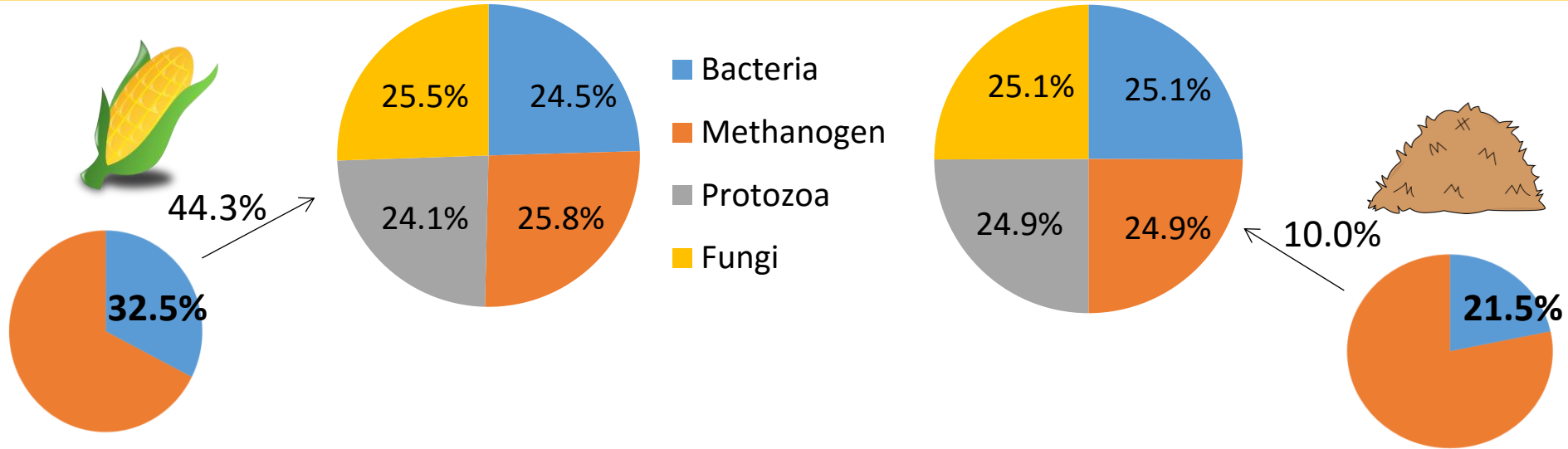


Trait (%/total VFA)	Inefficient	Efficient	<i>P</i> -value	Inefficient	Efficient	<i>P</i> -value
Acetate	54.4	54.5	0.97	73.48	72.95	0.55
Propionate	29.0	29.3	0.89	17.25	17.45	0.38
Isobutyrate	1.2	1.0	0.11	0.90	0.95	<0.10
Butyrate	8.9	8.6	0.75	6.81	6.91	0.44
Isovalerate	3.0	2.7	0.41	0.87	0.94	<0.05
Valerate	2.6	2.9	0.29	0.47	0.49	0.43
Caproate	0.4	0.5	0.40	0.22	0.22	0.87
Total VFA (μmol/ml)	78.0	80.3	0.66	38.94	41.83	<0.10

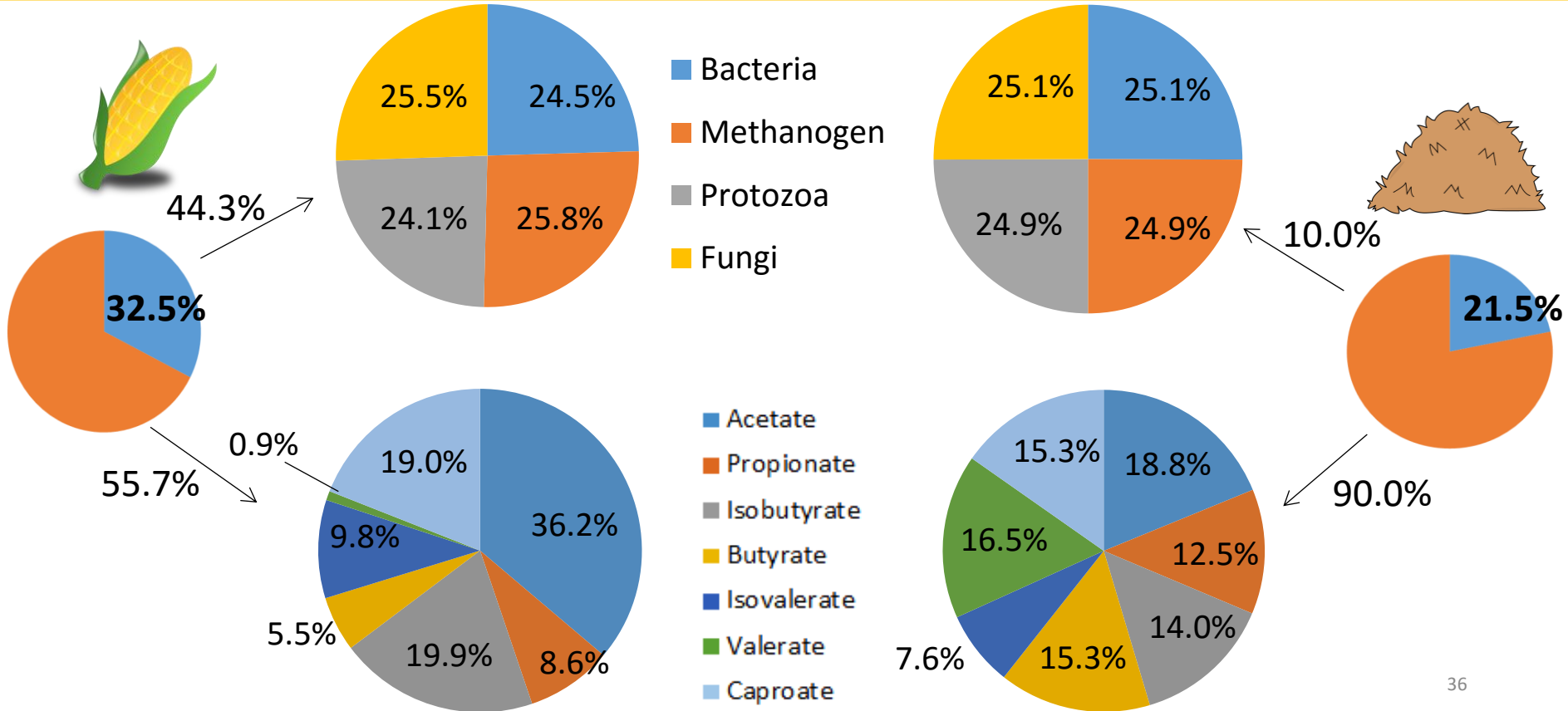
Results: Variance analysis



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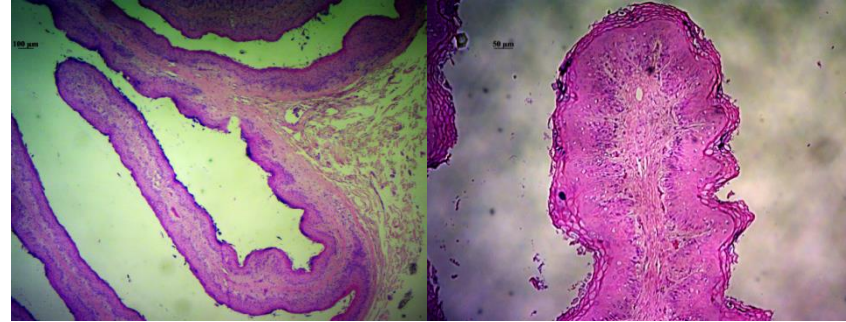
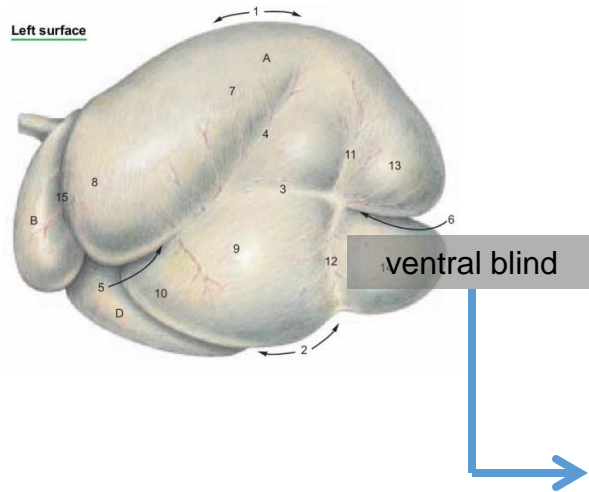


Results: Variance analysis



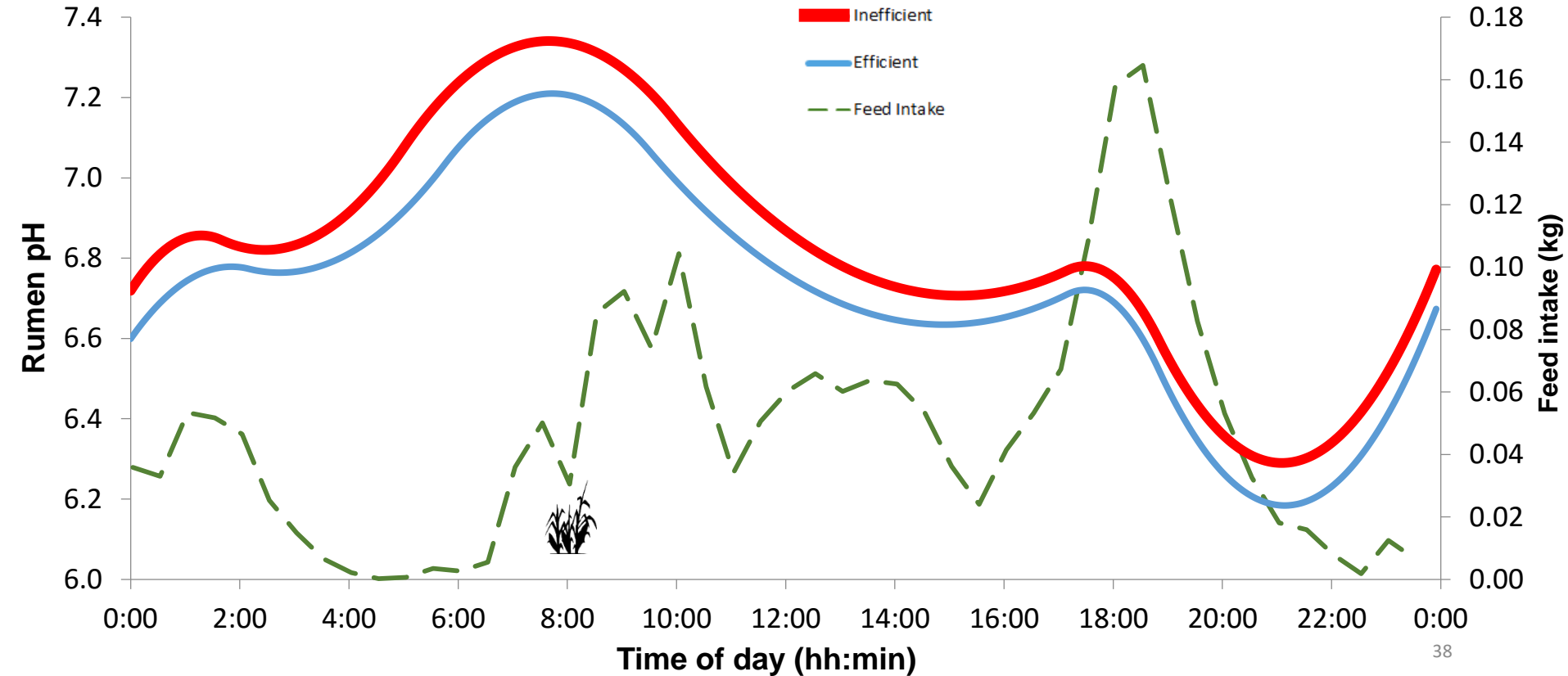


Results: Papillae histomorphometry

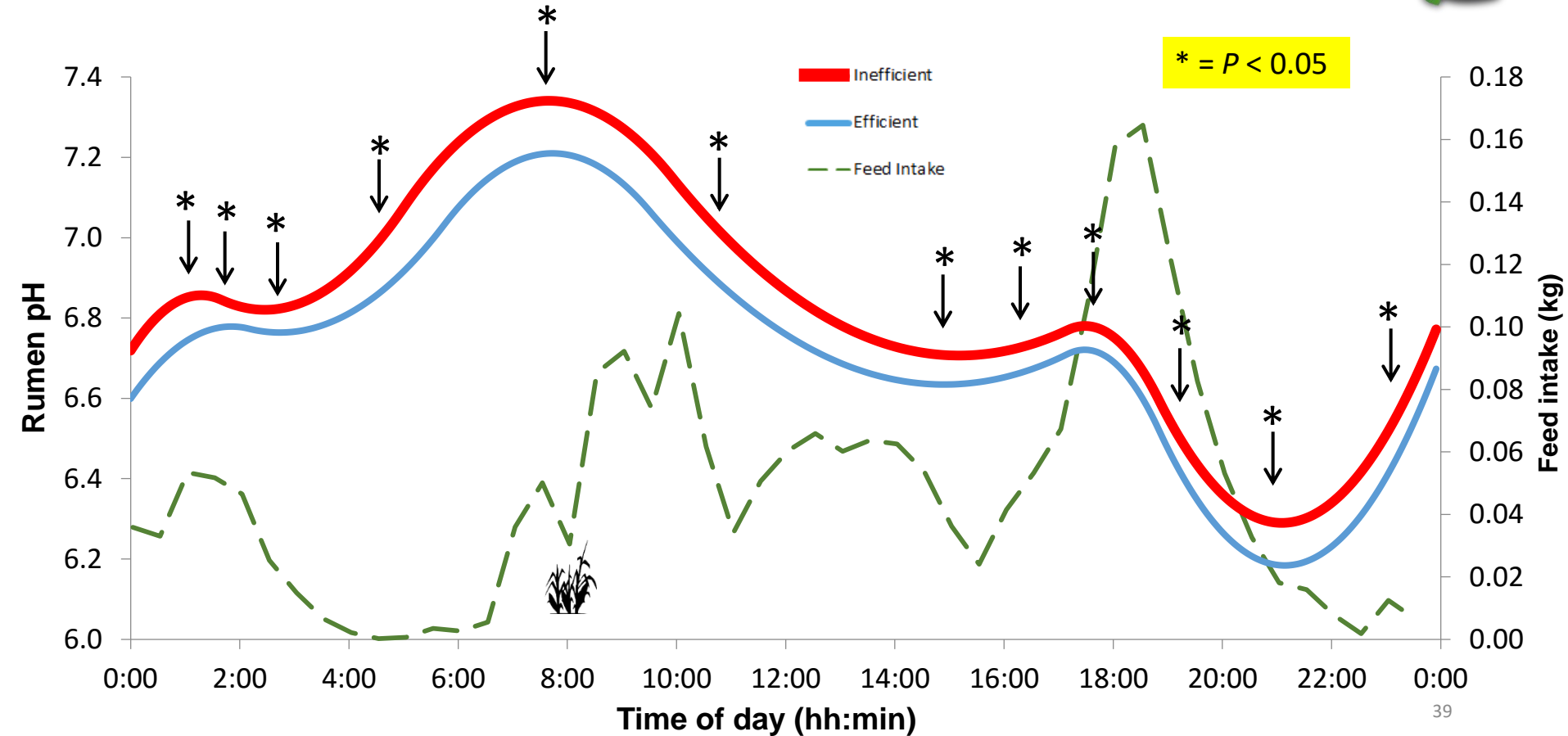


Papillae width	Papillae area	Inefficient	Efficient	<i>P</i>
(μm)	Base	110.1	113.4	0.39
	Middle	119.5	139.4	<0.05
	Tip	125.0	148.4	<0.01

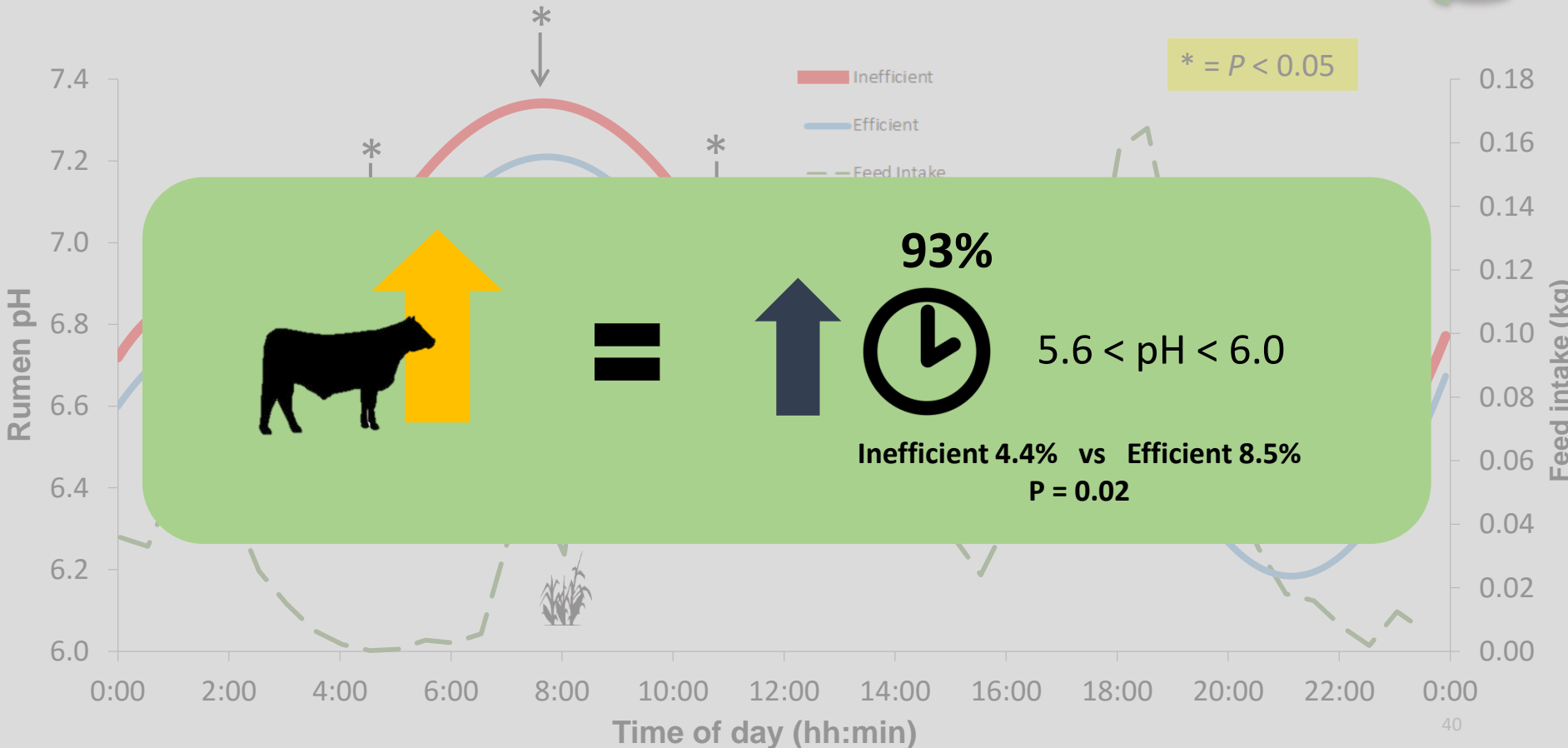
Results: Predicted rumen pH curves



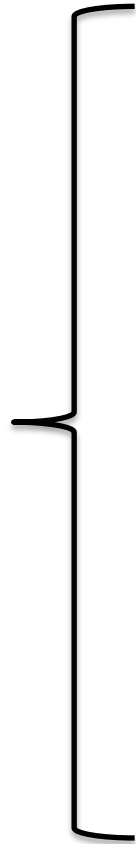
Results: Predicted rumen pH curves



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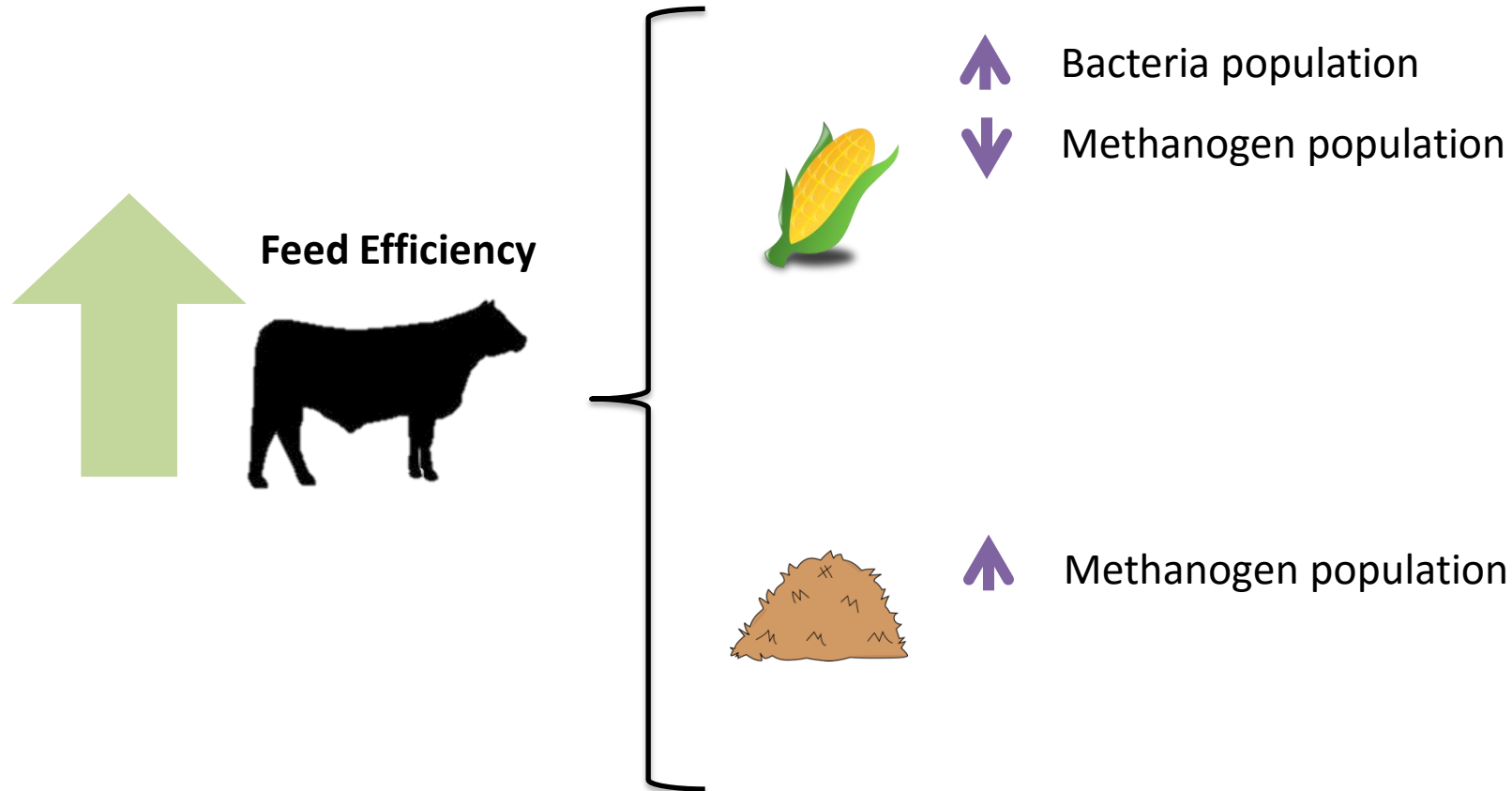


Summary

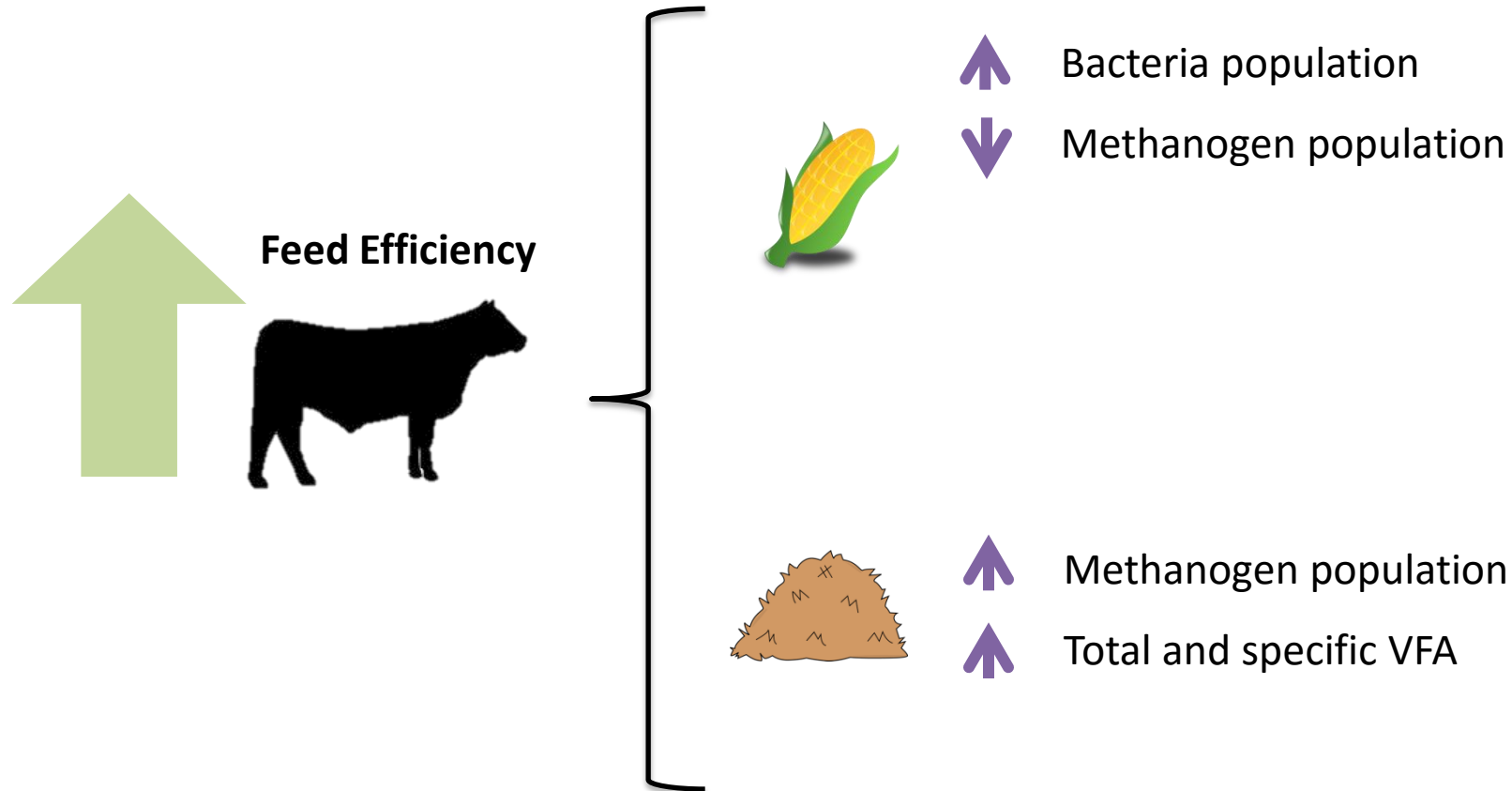


Bacteria population

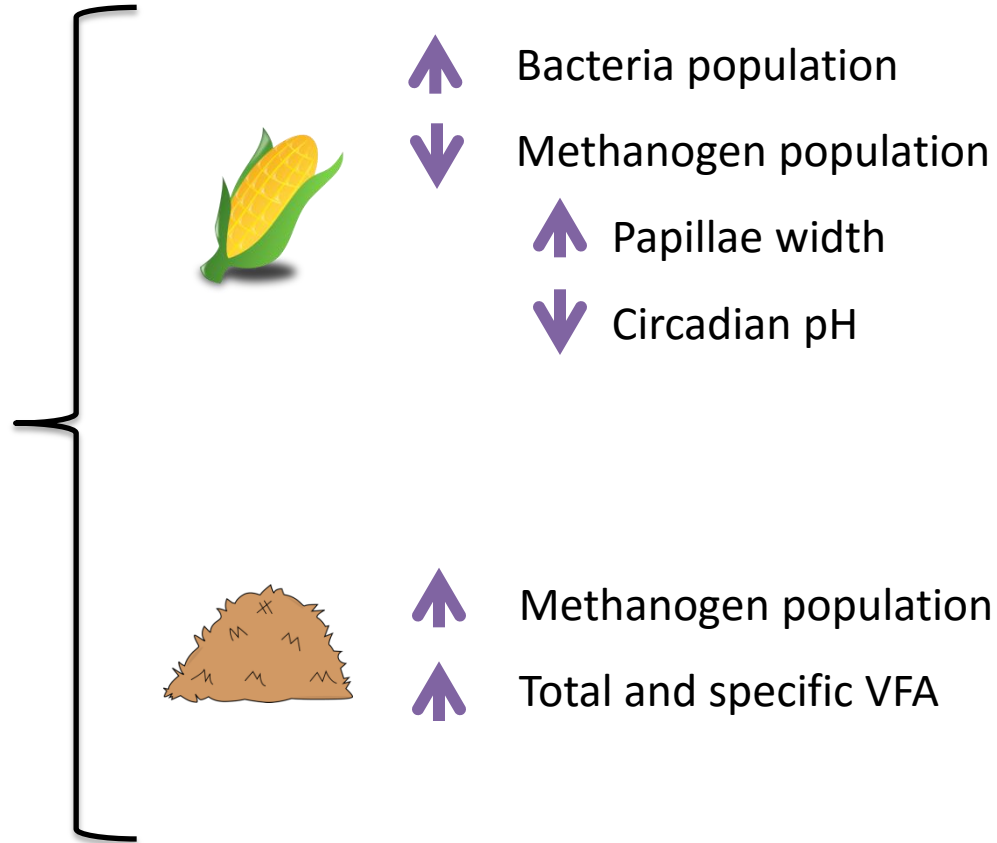
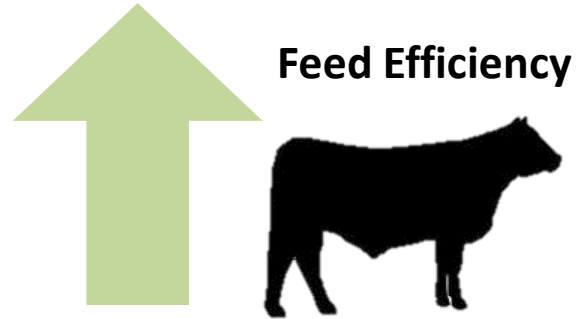
Summary



Summary



Summary



Conclusions

- Rumen microbiology, functional and structural parameters are important in assessing the underlying digestive biology of feed efficiency
- Dietary treatment has an impact on the relevance of rumen parameters used for indicating feed efficiency



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