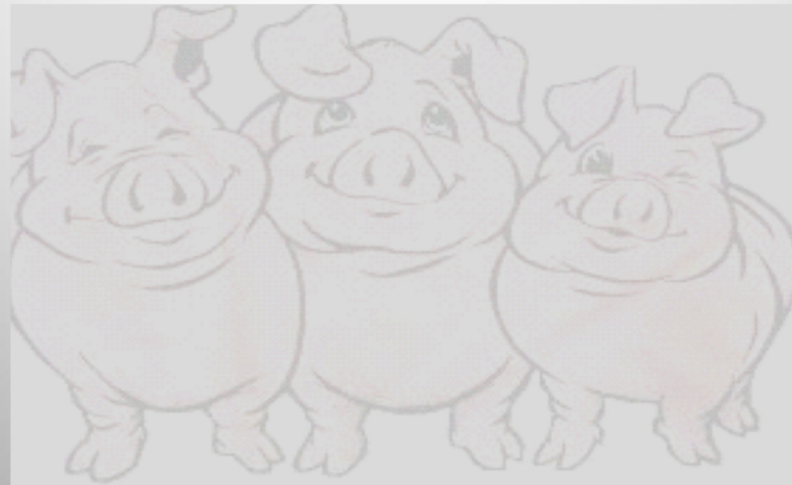
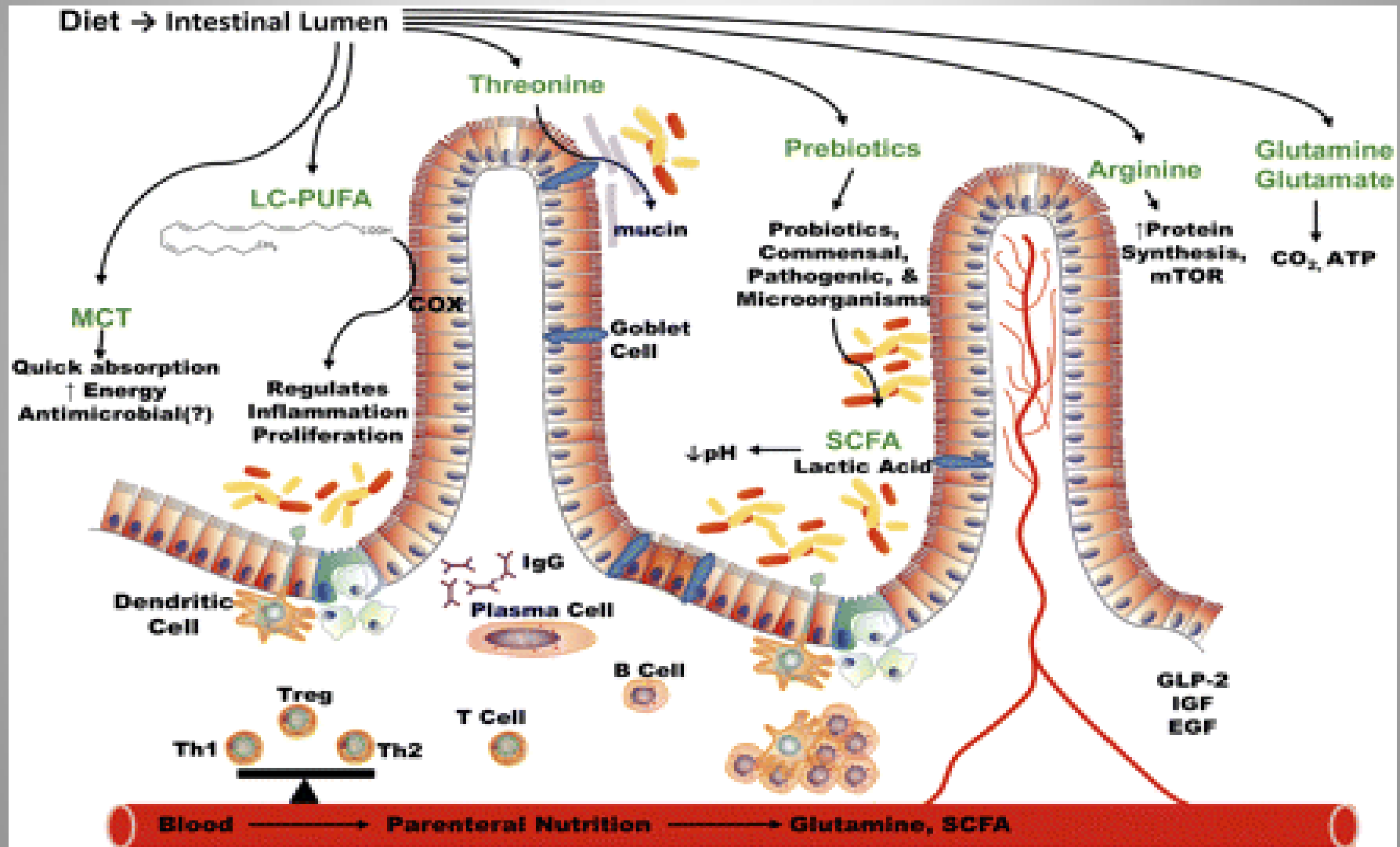


Influence of feeding and management on gut microbiota and barrier function

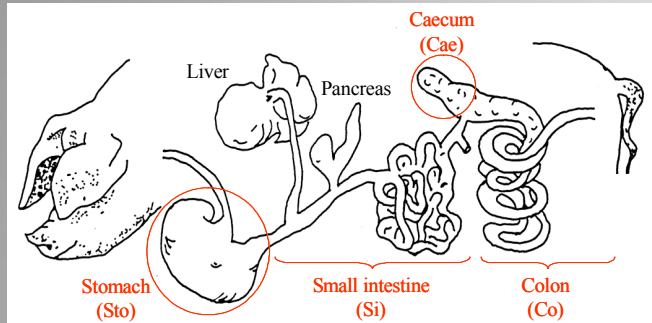
Bent Borg Jensen
Dept. Animal Science
Aarhus University



Mechanisms by which diet and feed additives may modulate the intestinal ecosystem and barrier function



Microbiological toolbox



Pig gastrointestinal tract (GI-tract)
Pig Intestine

Sampling



Microbiological Toolbox

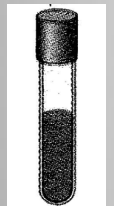
DNA Extraction & Purification



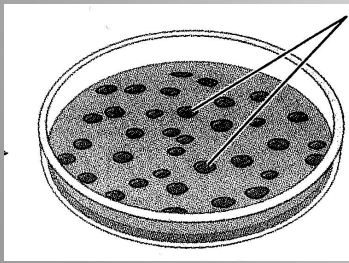
Direct analysis

DNA analysis

Community fingerprints

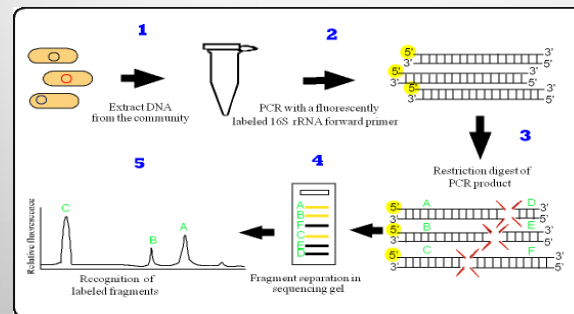


Metabolite analysis

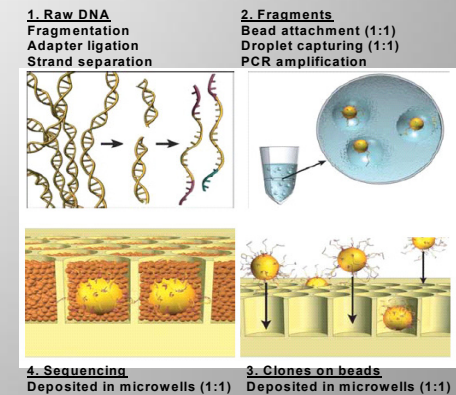


Selective culturing

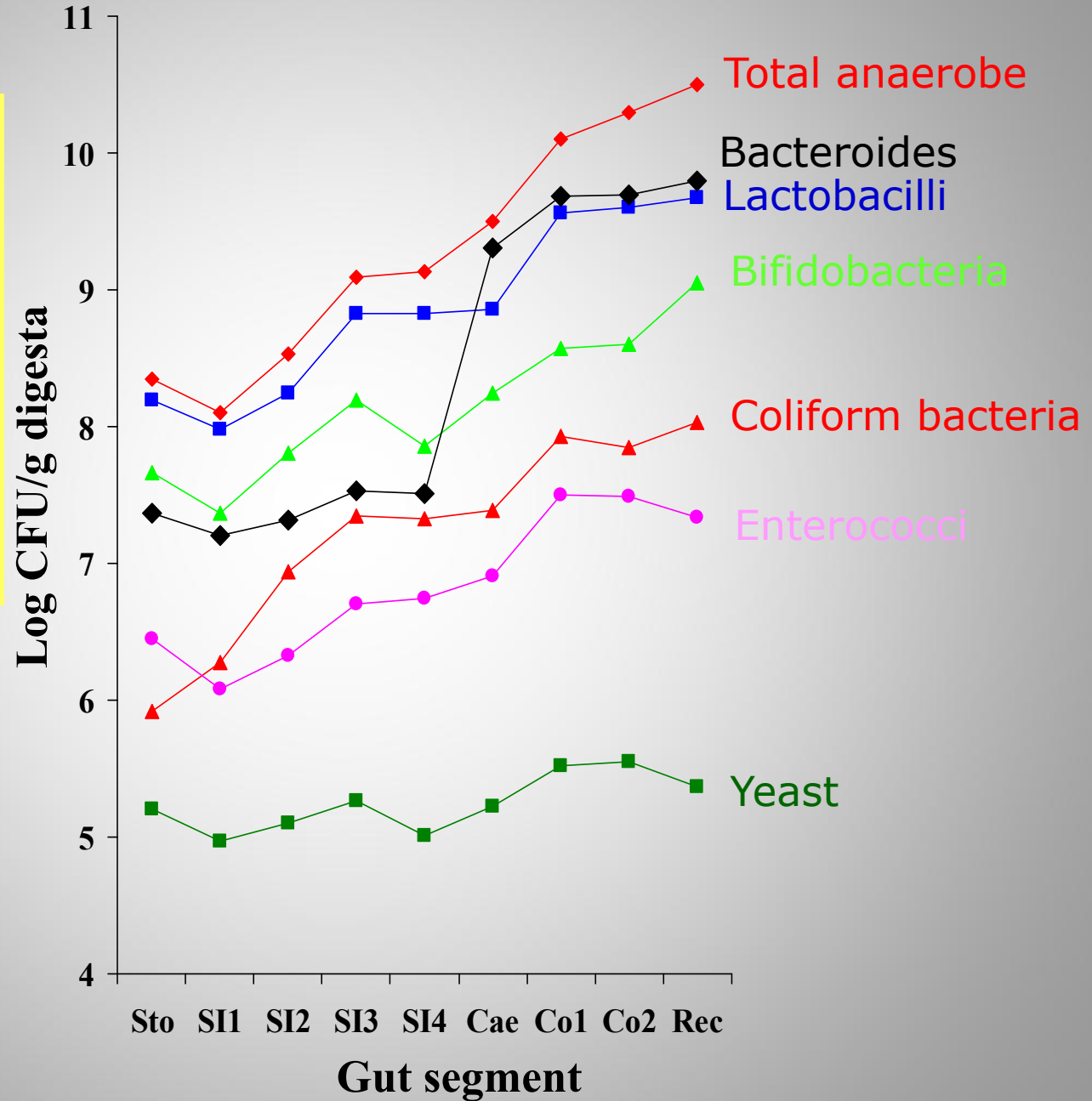
T-RFLP



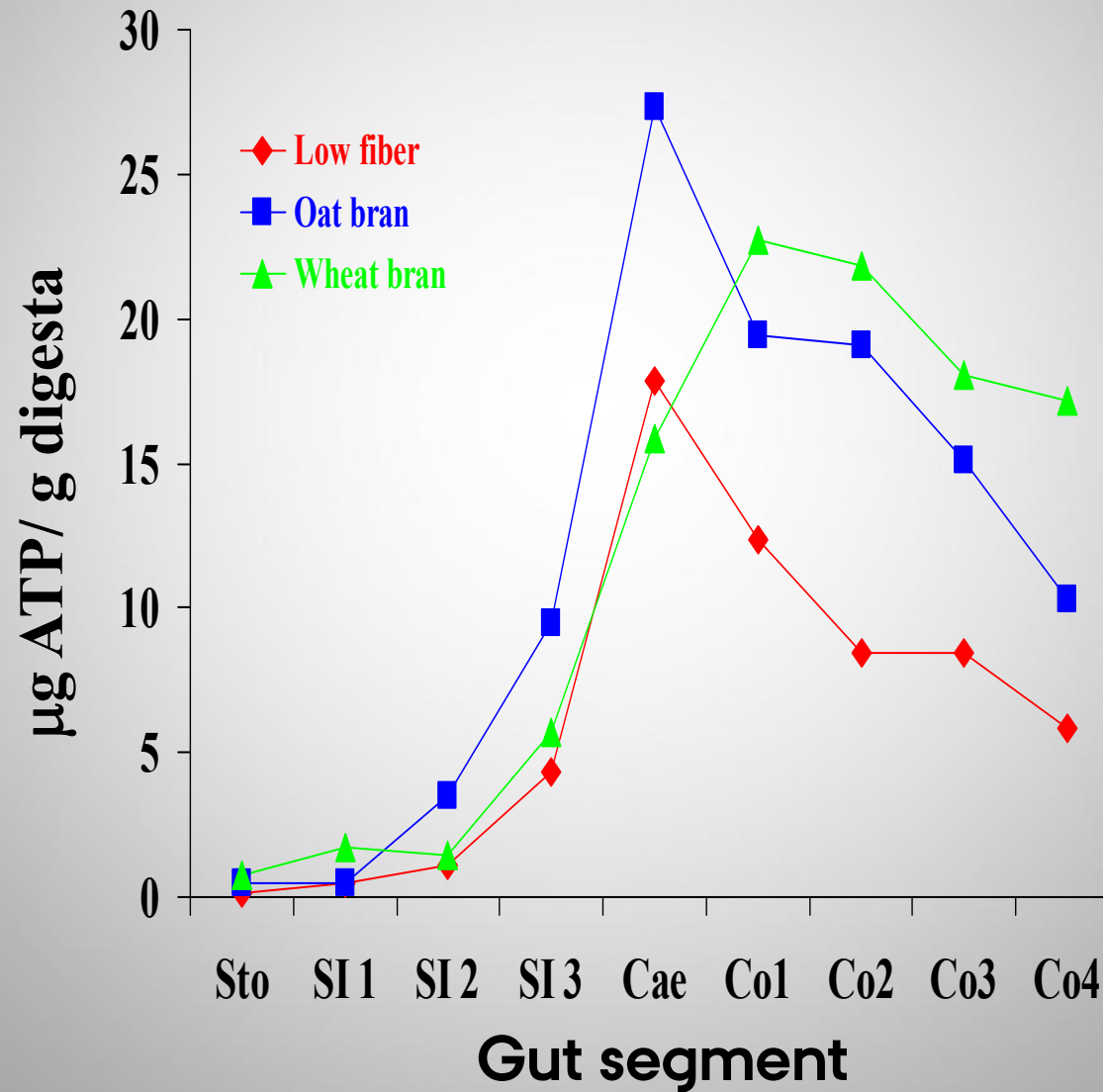
Pyrosequencing



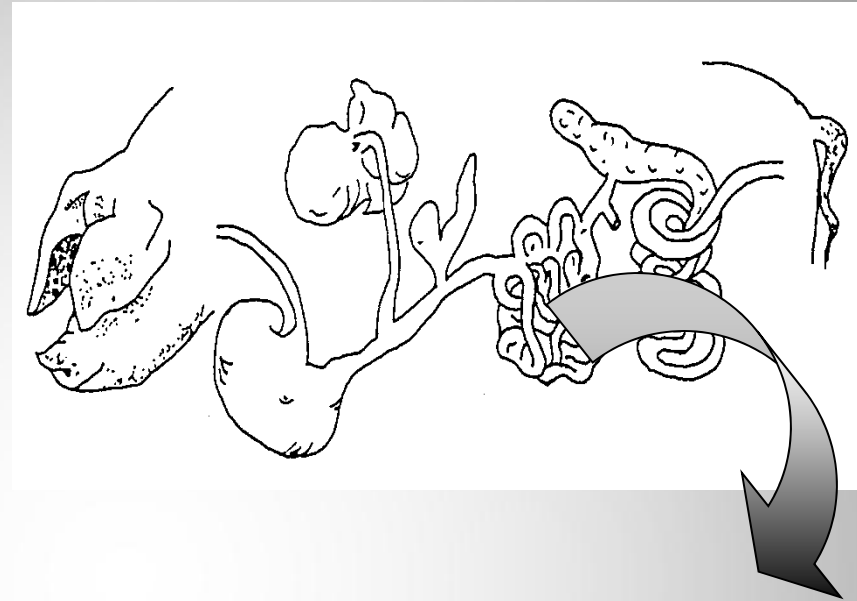
Density of selected groups of cultivable bacteria in various section of the gastrointestinal tract of slaughter pigs



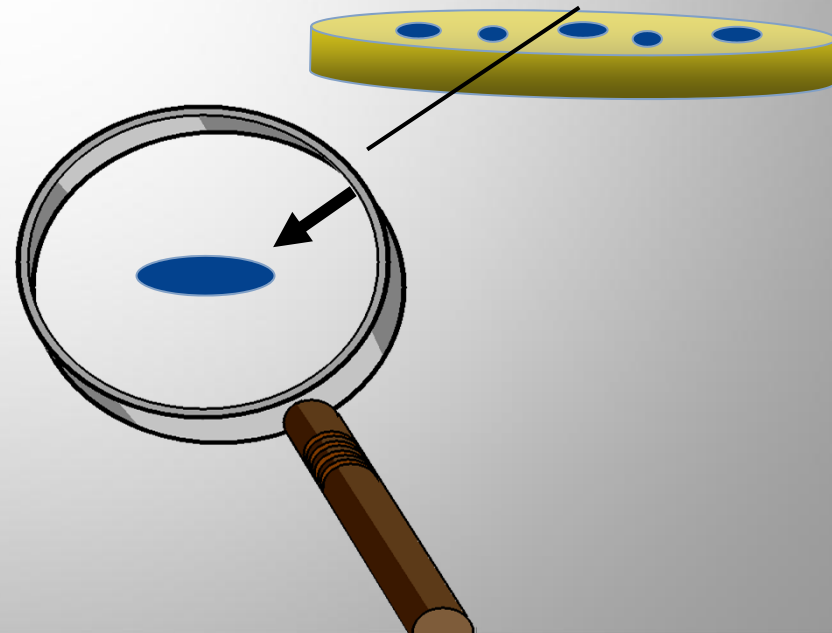
Microbial activity in various regions of the gastrointestinal tract of pigs



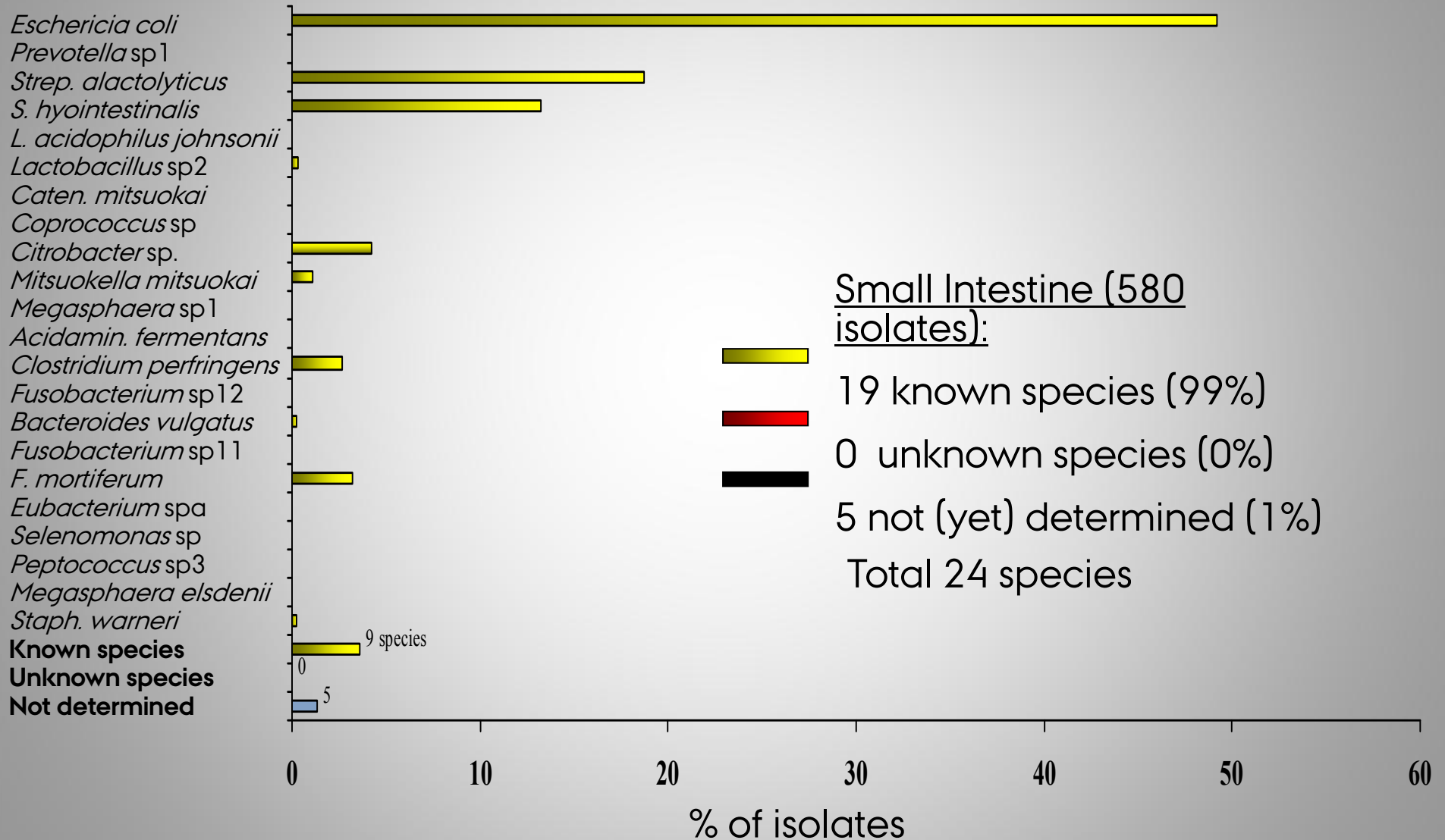
Characterise the
culturable bacteria
in the gastrointestinal
tract of pigs



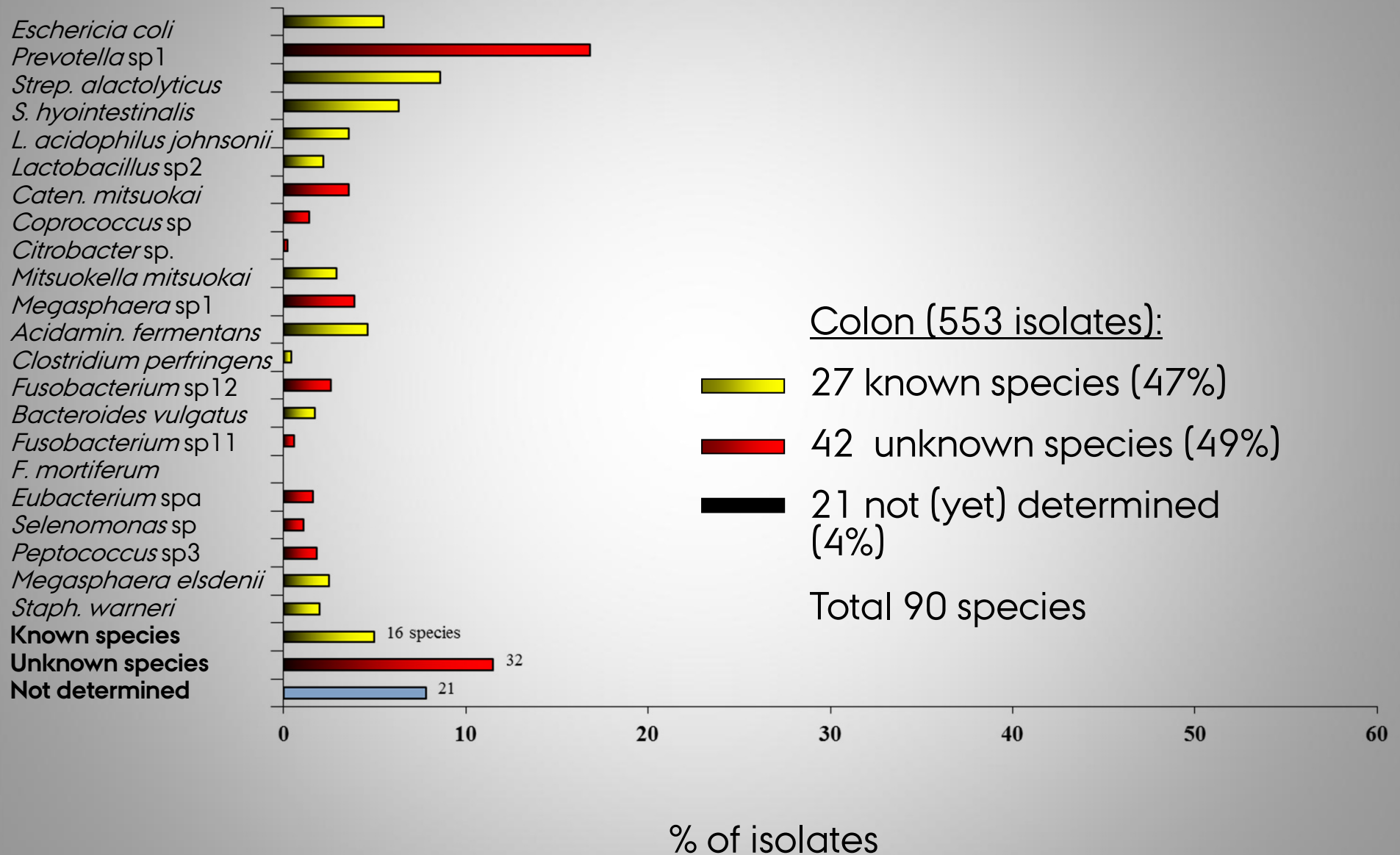
Are we able
to cultivate gut bacteria ?



Relative distribution of isolates on species



Relative distribution of isolates on species



16S rDNA match of predominant colon isolates

| Isolate/species | Proportion of isolates (%) | GenBank 16S rDNA match | Similarity (%) |
|--------------------------------------|----------------------------|--------------------------------------|----------------|
| <i>Prevotella</i> sp1 | 16.8 | OTU 16 | 98.8 |
| <i>Streptococcus alactolyticus</i> | 8.6 | <i>Streptococcus alactolyticus</i> | 100 |
| <i>Streptococcus hyointestinalis</i> | 6.3 | <i>Streptococcus hyointestinalis</i> | 99.8 |
| <i>Escherichia coli</i> | 5.5 | <i>Escherichia coli</i> | 99.8 |
| <i>Acidaminococcus fermentans</i> | 4.6 | <i>Acidaminococcus fermentans</i> | 99.4 |
| <i>Megasphaera</i> sp1 | 3.9 | <i>Megasphaera elsdenii</i> | 96.0 |
| <i>Lactobacillus acid. johnsonii</i> | 3.6 | <i>Lactobacillus johnsonii</i> | 98.7 |
| <i>Selenomonas</i> sp1 | 3.6 | URB 4C28d-14 | 98.1 |
| <i>Mitsuokella multiacidus</i> | 2.9 | <i>Mitsuokella multiacidus</i> | 98.7 |
| <i>Prevotella</i> sp22 | 2.6 | OTU 18 | 97.2 |
| | 58.4 | | |

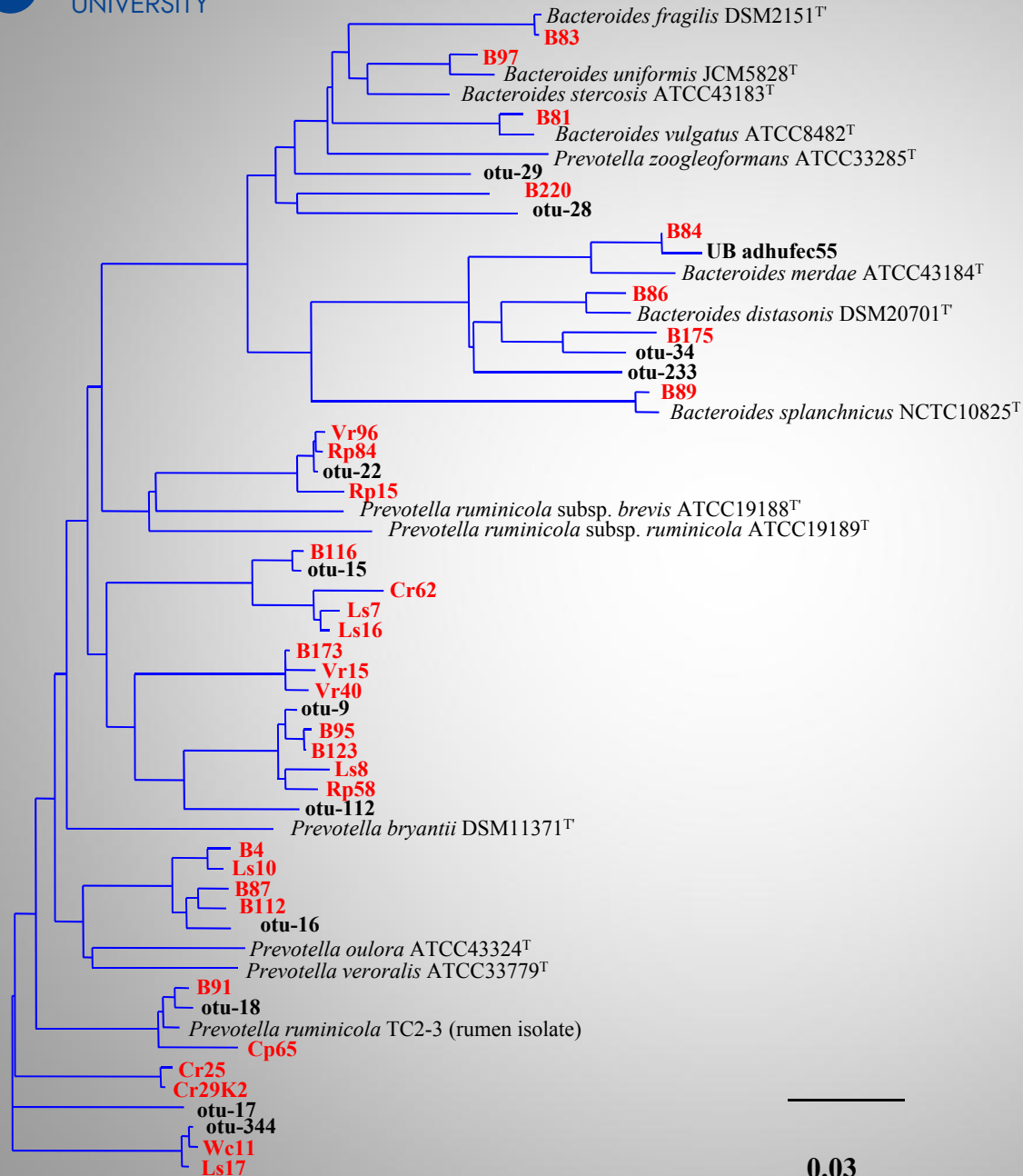
OTU: Uncultured Bacterium (Pig) URB: Uncultured Rumen Bacterium

Comparison between isolates and most dominant clones*

| Clones (9270) | | | Isolates** (791) |
|---------------|---|-------------|------------------|
| Rank | Closest related species (similarity) | % of clones | % of isolates |
| 1 | <i>Streptococcus alactolyticus</i> (99.7) | 7.9 | 22.8 |
| 2 | <i>Escherichia coli</i> (100.0) | 4.8 | 12.2 |
| 3 | <i>Lactobacillus sorbius</i> (99.2) | 4.5 | 12.0 |
| 4 | <i>Faecalibacteria prausnitzii</i> (92.6) | 4.0 | <0.1 |
| 5 | <i>Lactobacillus johnsonii</i> (99.8) | 3.2 | 5.3 |
| 6 | <i>Faecalibacteria prausnitzii</i> (97.6) | 2.9 | <0.1 |
| 7 | <i>Sporobacter termitidis</i> (90.7) | 2.4 | |
| 8 | <i>Gemella haemolysans</i> (87.1) | 2.1 | |
| 9 | <i>Lactobacillus reuterii</i> (99.5) | 1.9 | 4.7 |
| 10 | <i>Eubacterium rectale</i> (92.6) | 1.6 | |

* Data from Leser et al. 2002, ** Similarity to clones >97%

Bacteroides-Prevotella cluster



Isolates

Type strains (Strains)

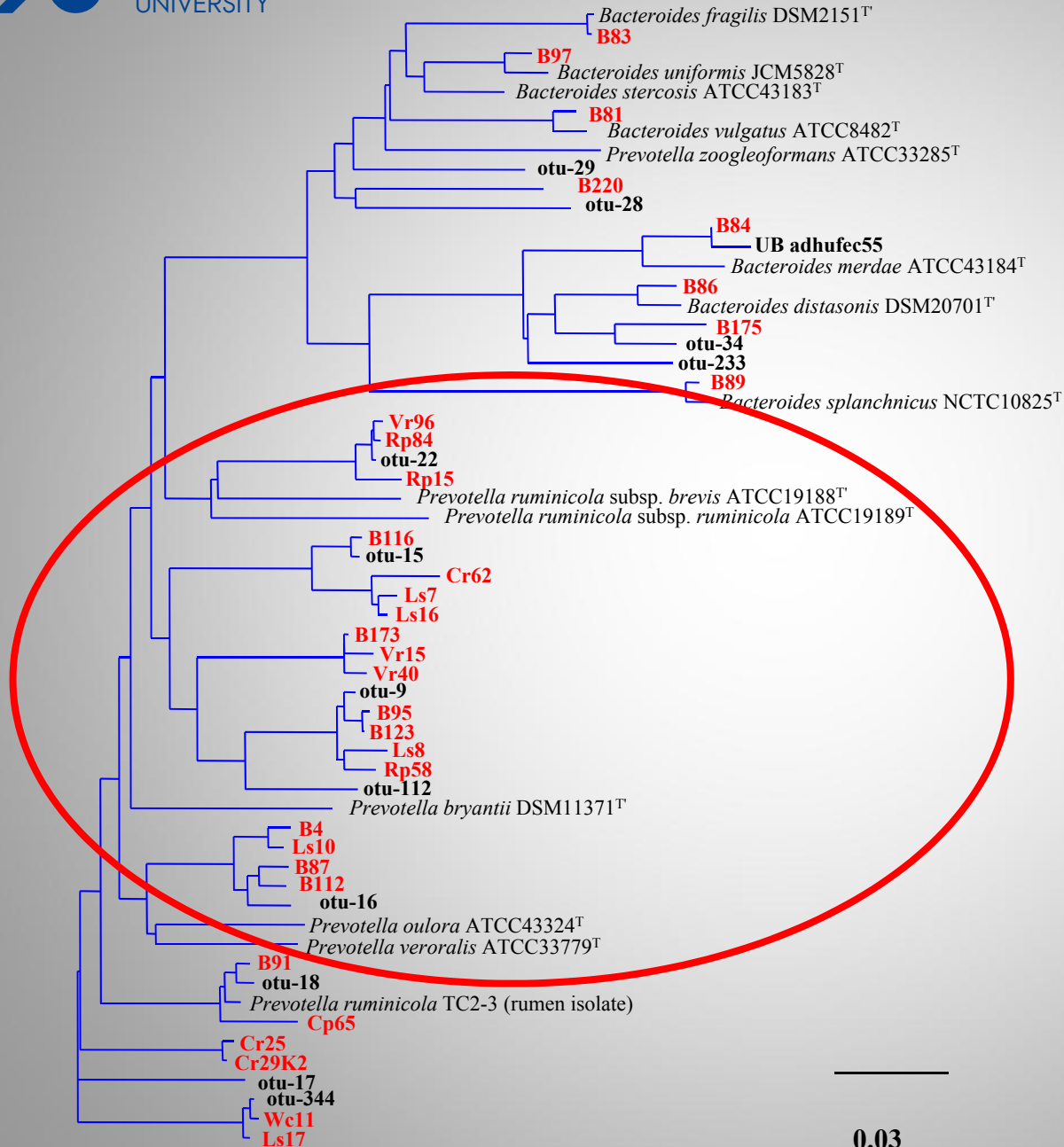
Uncultured bacteria clones

Leser et al. 2002 (otu)
Suau et al. 1999 (adhufec)

0.03

(Algorithm: Neighbor-Joining)

Bacteroides-Prevotella cluster



Isolates

Type strains (Strains)

Uncultured bacteria clones

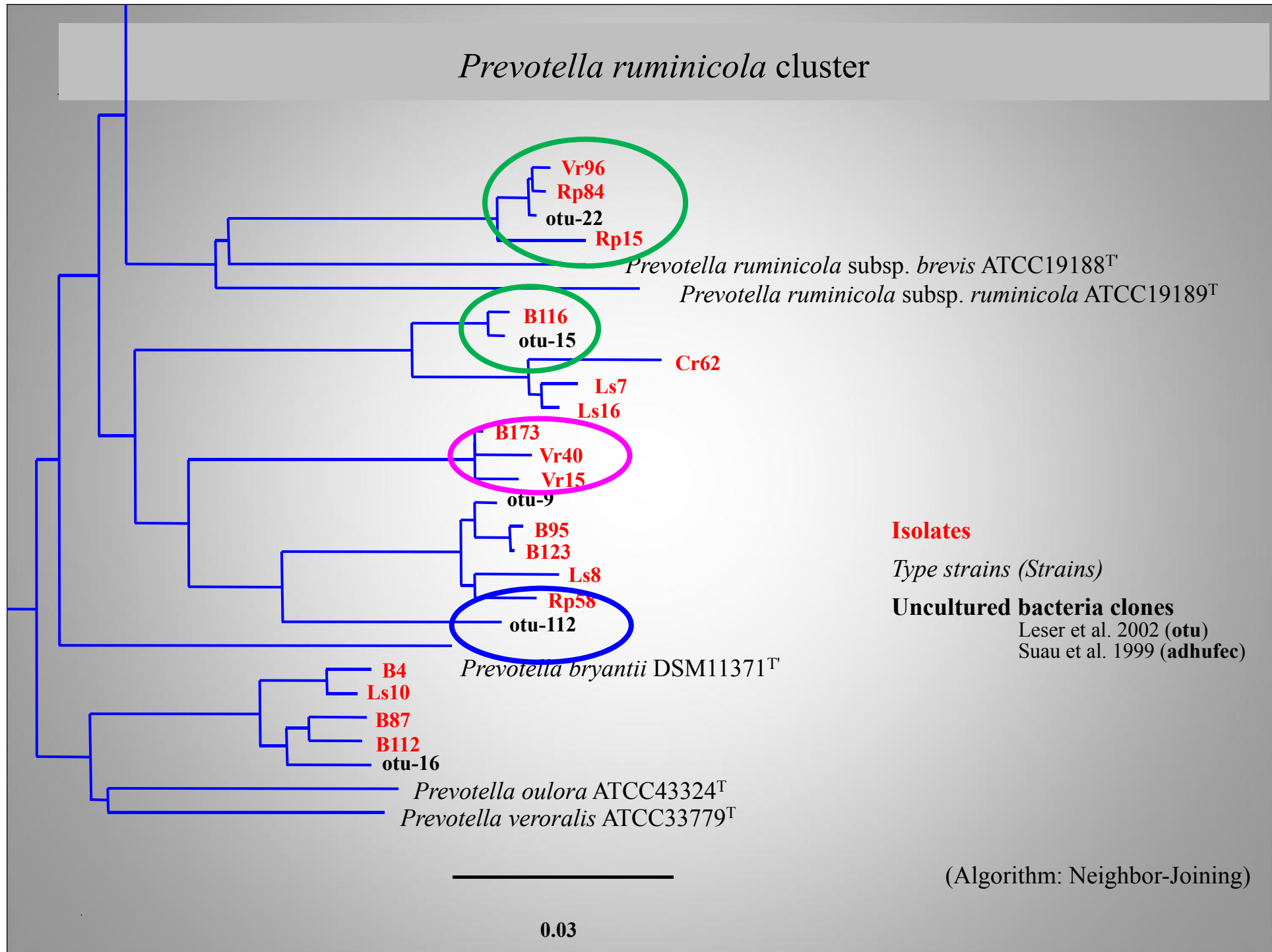
Leser et al. 2002 (otu)

Suau et al. 1999 (adhufec)

(Algorithm: Neighbor-Joining)

0.03

Prevotella ruminicola cluster



Vr96
Rp84
otu-22
Rp15

Prevotella ruminicola subsp. *brevis* ATCC19188^T
Prevotella ruminicola subsp. *ruminicola* ATCC19189^T

B116
otu-15

Cr62
Ls7
Ls16

B173
Vr40
Vr15
otu-9

B95
B123
Ls8

Rp58
otu-112

Prevotella bryantii DSM11371^T

B4
Ls10
B87
B112
otu-16

Prevotella oulora ATCC43324^T
Prevotella veroralis ATCC33779^T

Conclusion

- ➔ Defined >120 species from 2200 isolates
> 60% no match to any known species
- ➔ Despite high diversity, 75% of the isolates were covered by eighteen OTUs (<1% of isolates each)
- ➔ Good agreement between our isolates and OTUs obtained from pig clone library

however some dominant pig clone library OTUs were not between our isolates

and some of our isolates could not be related to any of the OTUs in the pig clone library

Most pig gastrointestinal bacteria are cultivable

Local Strain Collection of Gastrointestinal Bacteria

127 'Species' Operational Taxonomic Units (OTU)

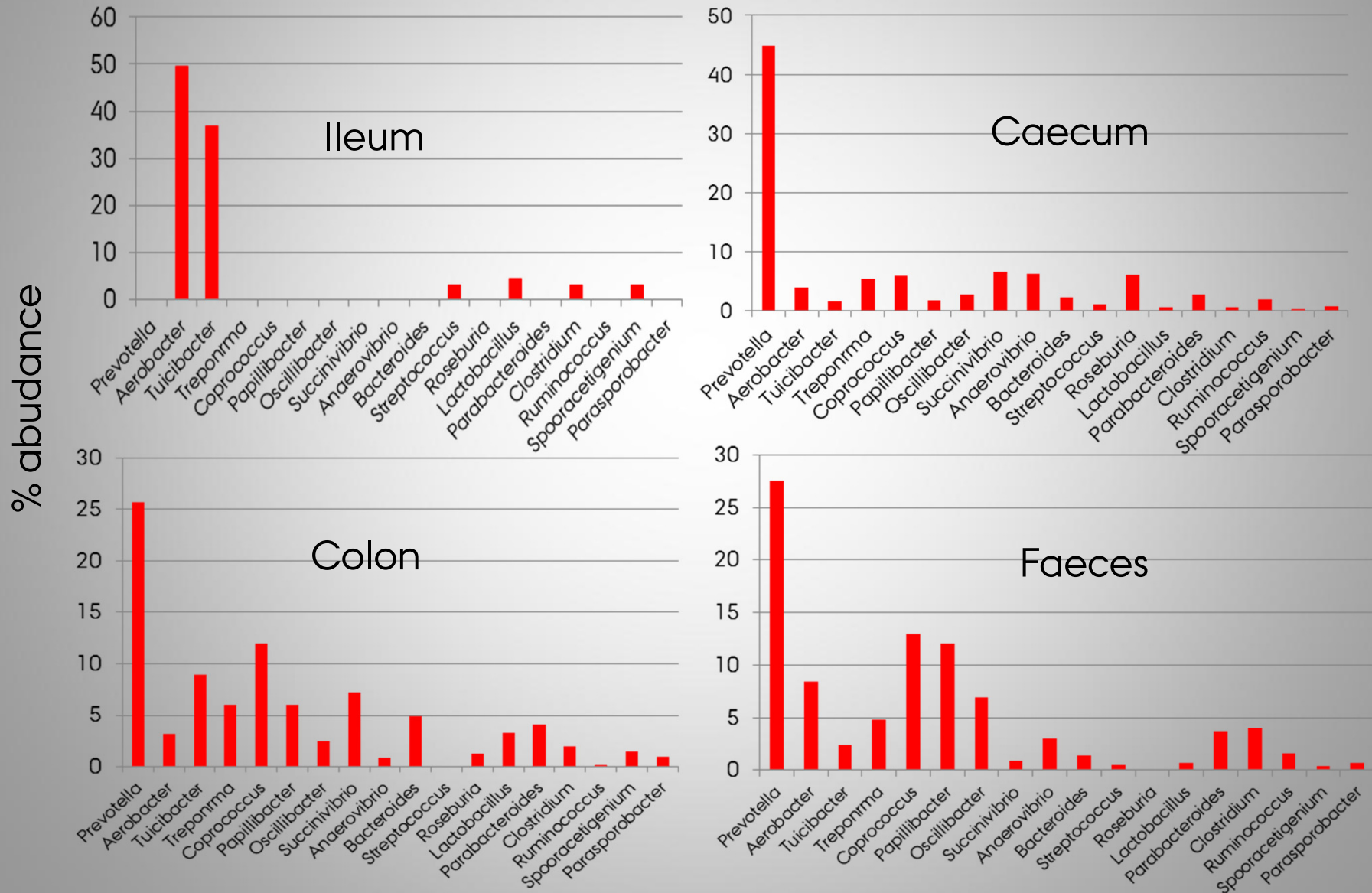
100% already known species

Species

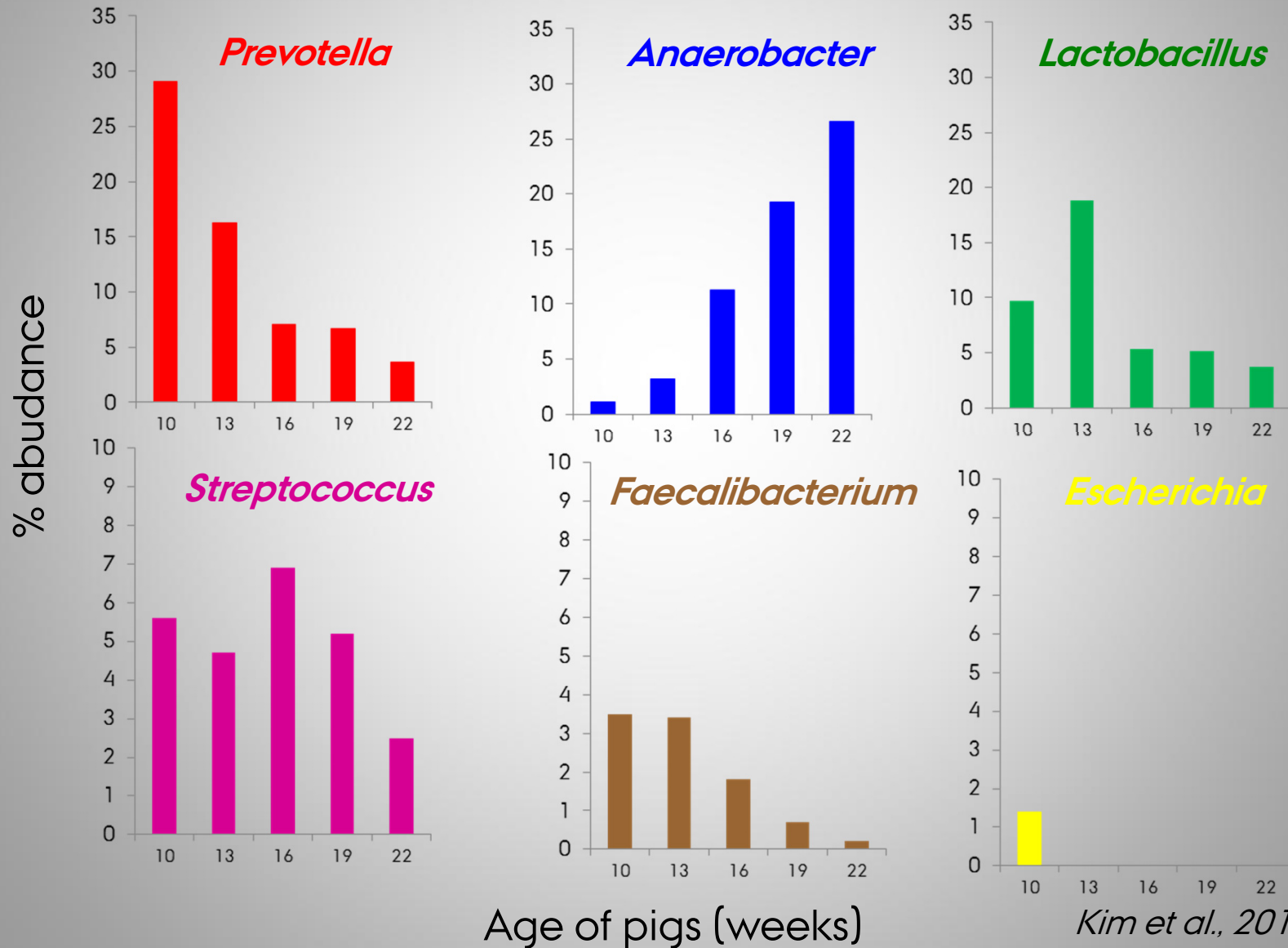
22 yet

Potential for studies on e.g.:
Probitotics
Host-microbe interactions

Abundance in most dominating bacteria genera in various sections of the gastrointestinal tract of pigs using pyrosequencing (Looft et al., 2014)



Age related change in bacterial populations in faeces of pigs



Important differences between the microbial ecosystem in the stomach/small intestine and the caecum/colon

Stomach and small intestine

- › Few bacterial species (20-30)
- › Low density/high activity
- › Compete with the host for digestible components (use around 6% of the energy)
- › Produce toxic compounds
- › Prevent growth of pathogenic bacteria

Caecum and colon

- › Many bacterial species (>400)
- › High density/low activity
- › Support the host with energy (around 16%)
- › Produce toxic compounds
- › May support growth of pathogenic bacteria

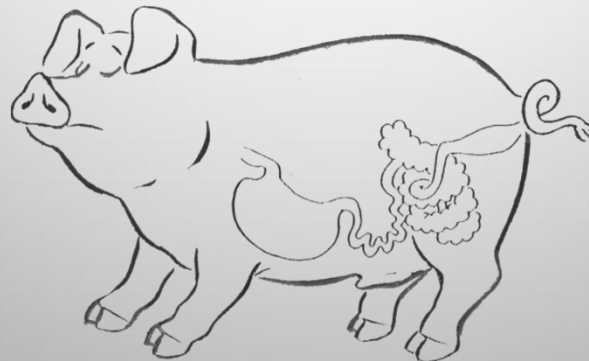
Factors that modulate the microbiota in the gastrointestinal tract

Feed

- Diet composition
- Feed processing
- Fermented liquid feed
- Prebiotics
- Low protein diets

Additives

- In-feed antibiotics
- zinc oxide
- organic acids
- probiotics
- Plant extracts/species



Feed additives Results from Danish weaner trails

| Product type | Number of trails | Trails with significance | ADG (% increase) | FCR (% increase) |
|---|------------------|--------------------------|------------------|------------------|
| Antibiotics | 13 | 8 | + 8,9 | - 3,5 |
| Acids/salts | 53 | 16 | + 5,1 | - 1,2 |
| <i>Acid/salts with significant effect</i> | <i>16</i> | <i>16</i> | <i>+ 9,4</i> | <i>- 3,8</i> |
| Probiotics | 17 | 1 | + 2,2 | - 1,1 |
| Aroma compounds | 27 | 2 | + 2,2 | -1,1 |
| Enzymes | 9 | 0 | + 2,1 | 0,0 |
| Oligosaccharides | 7 | 1 | + 2,3 | - 1,2 |

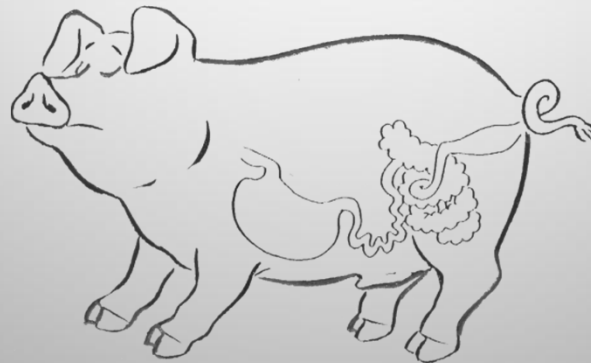
Factors that modulate the microbiota in the gastrointestinal tract

Feed

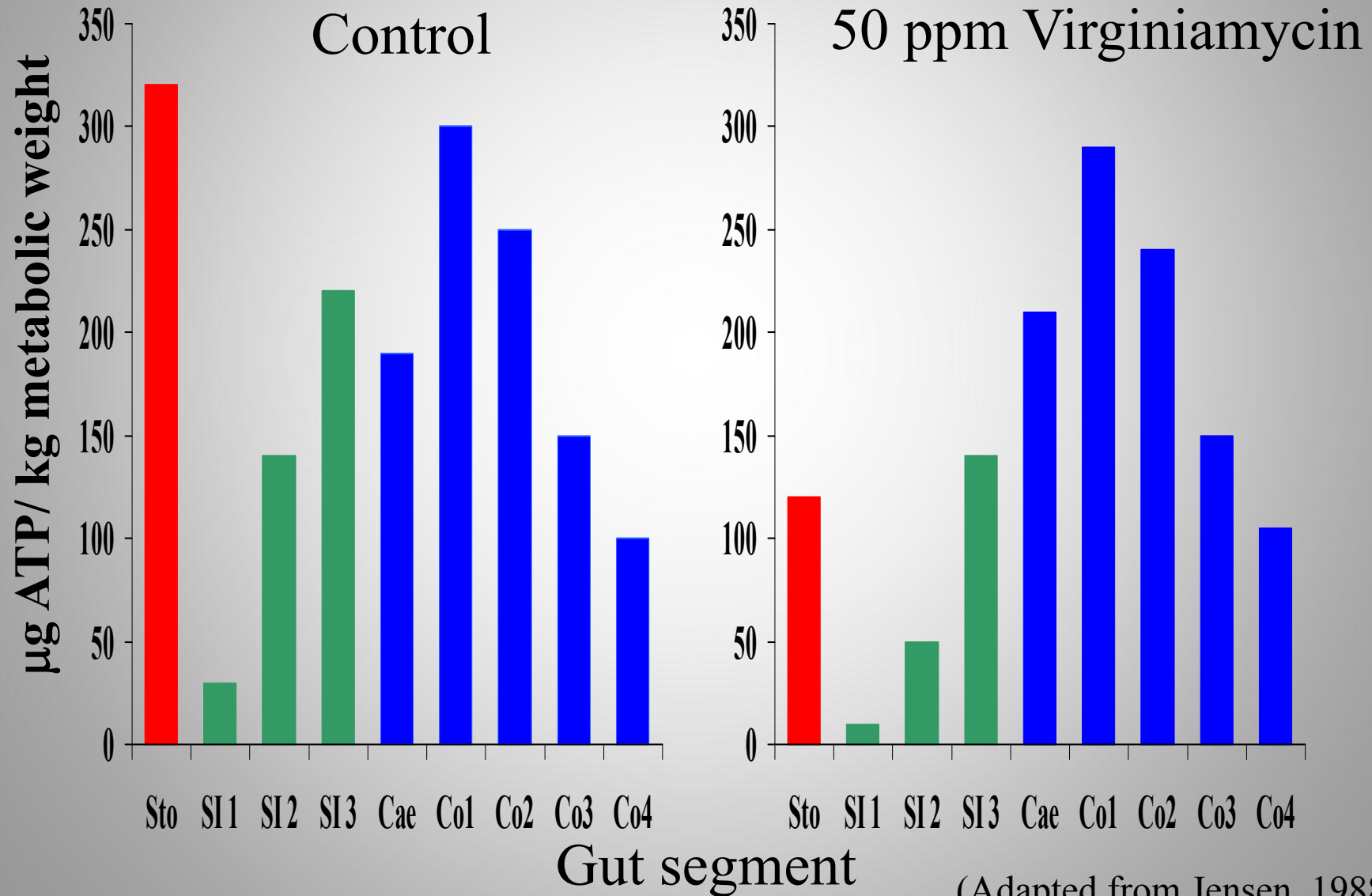
- Diet composition
- Feed processing
- Fermented liquid feed
- Prebiotics
- Low protein diets

Additives

- **In-feed antibiotics**
- zinc oxide
- organic acids
- probiotics
- Plant extracts/species

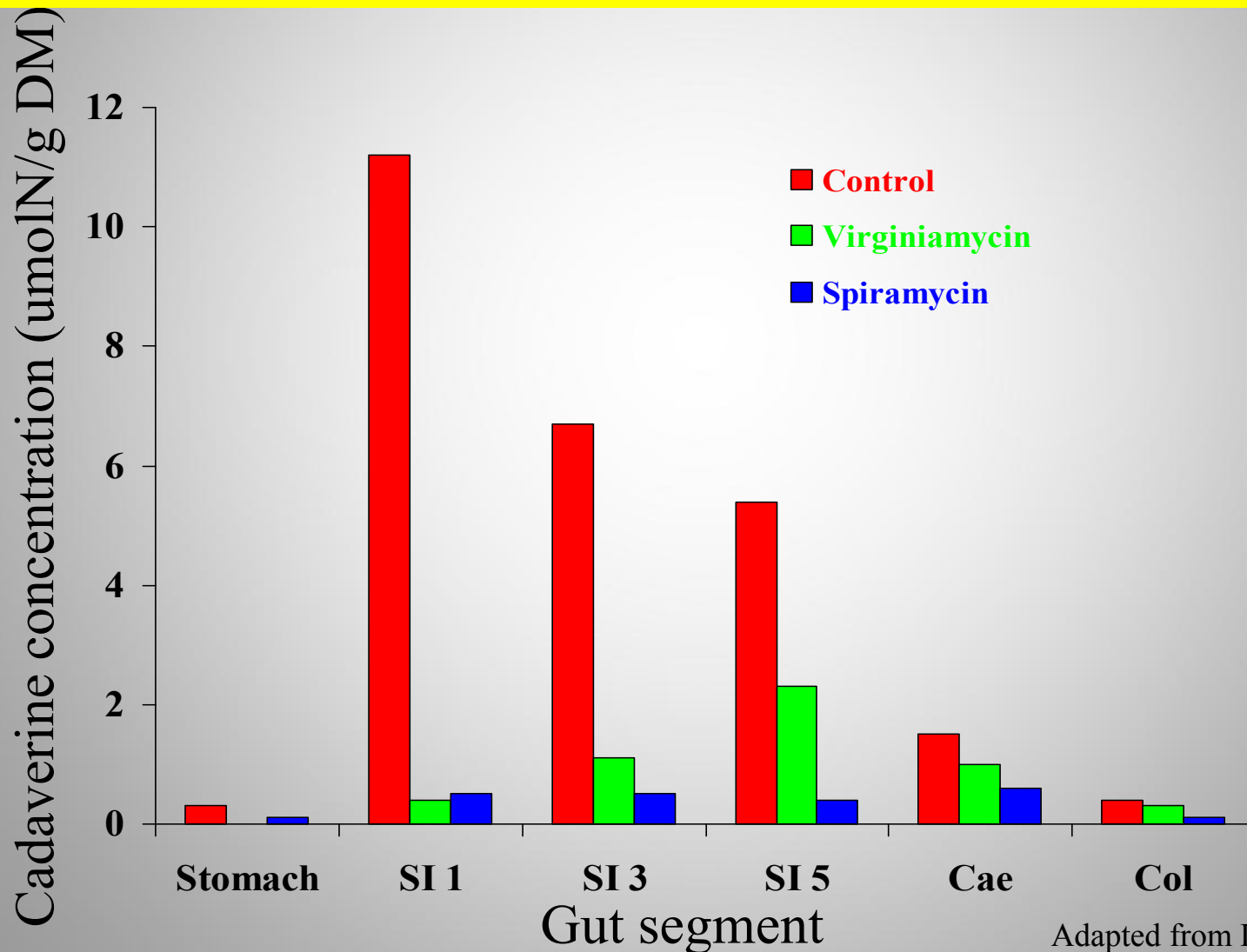


Effect of Virginiamycin on the microbial activity in various regions of the gastrointestinal tract of pigs



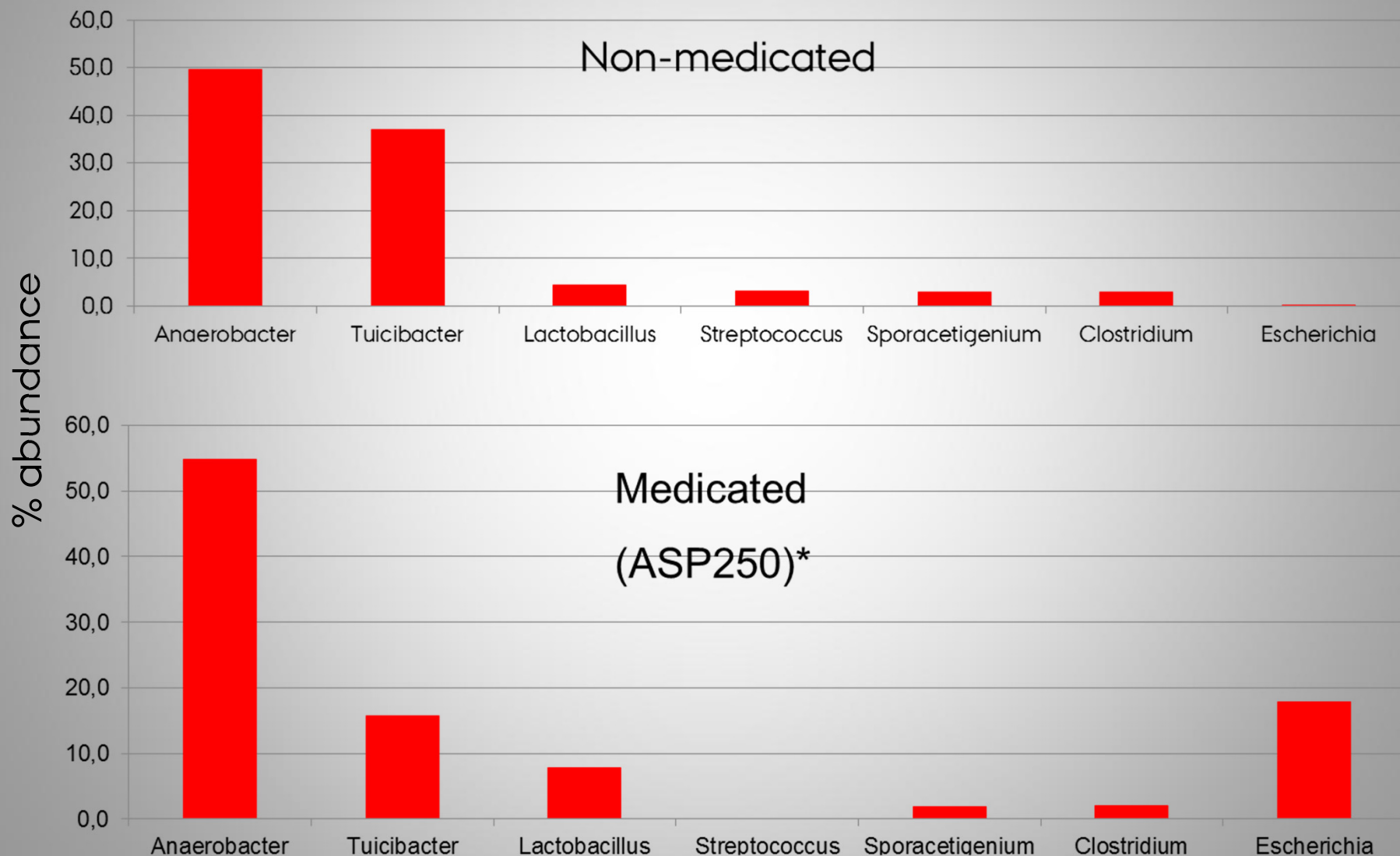
(Adapted from Jensen, 1988)

Effect of Virginiamycin (20 ppm) and Spiramycin (20 ppm) on cadaverine concentration in various regions of the gastrointestinal tract of piglets



Adapted from Dierick et al., 1985

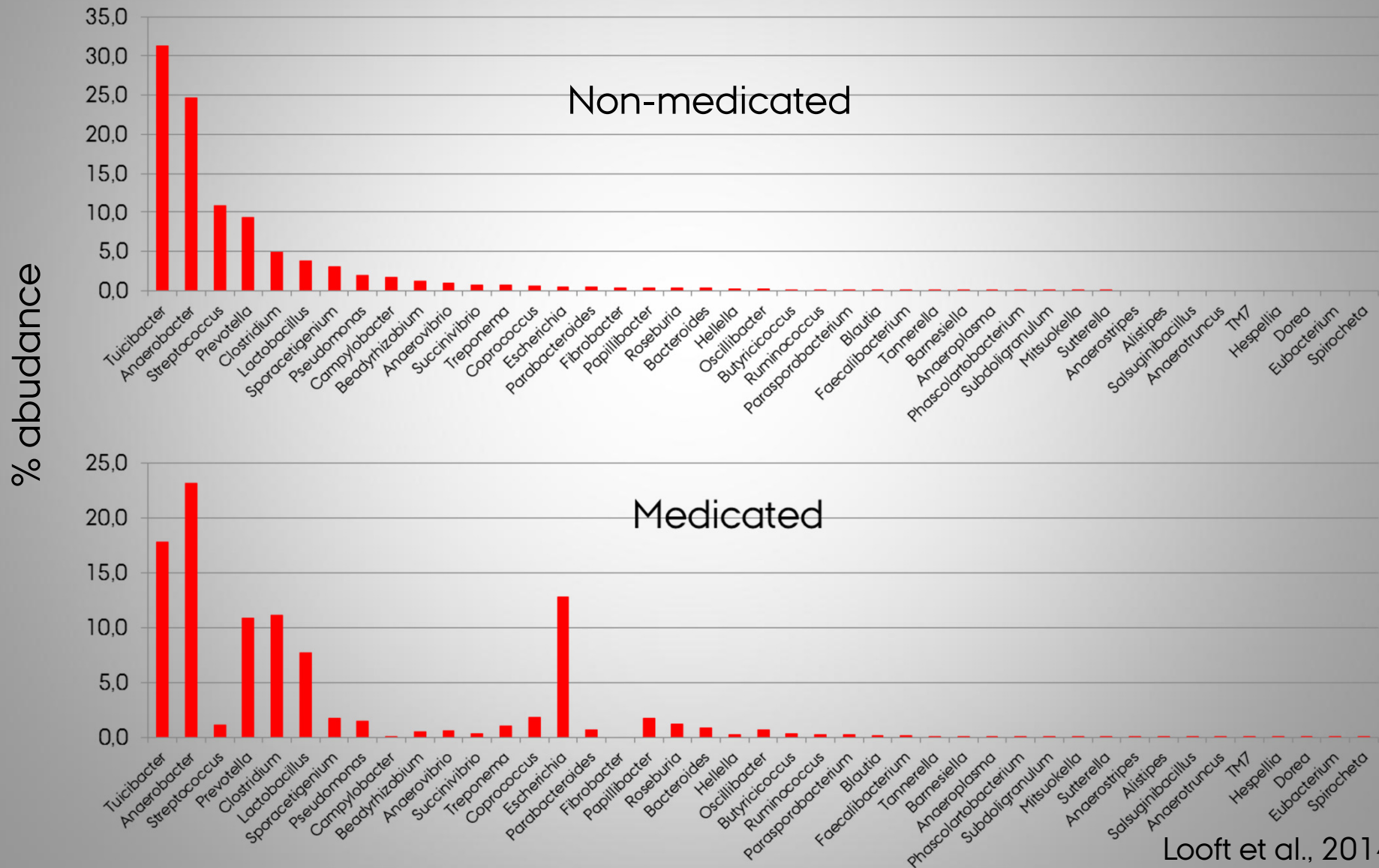
Effect of in-feed antibiotics on abundance of most dominating bacteria in ileal lumen samples



*100 ppm chlortetracyclin, 100 ppm sulfamethazine, 50 ppm penicillin)

Looft et al., 2014

Effect of in-feed antibiotics on most dominating bacteria in ileal mucosa samples



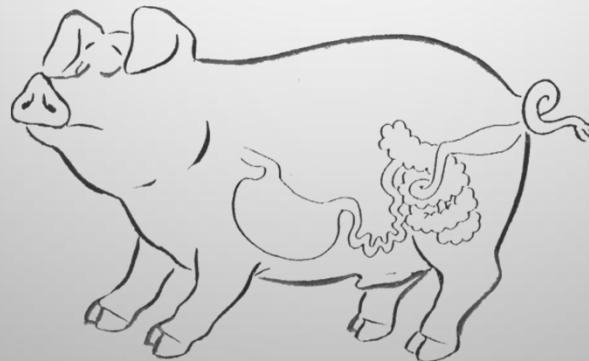
Factors that modulate the microbiota in the gastrointestinal tract

Feed

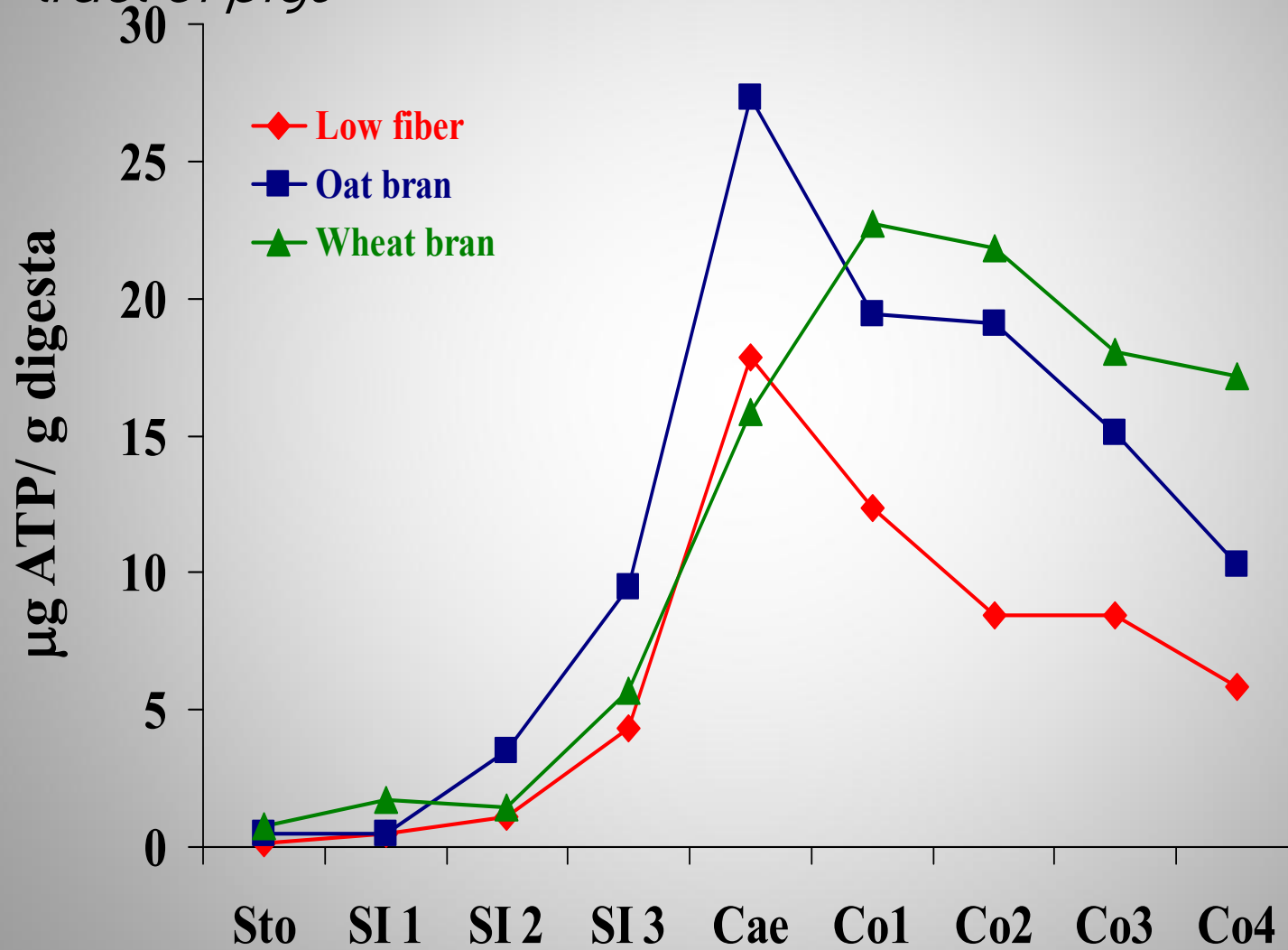
- Diet composition
- Feed processing
- Fermented liquid feed
- Prebiotics
- Low protein diets

Additives

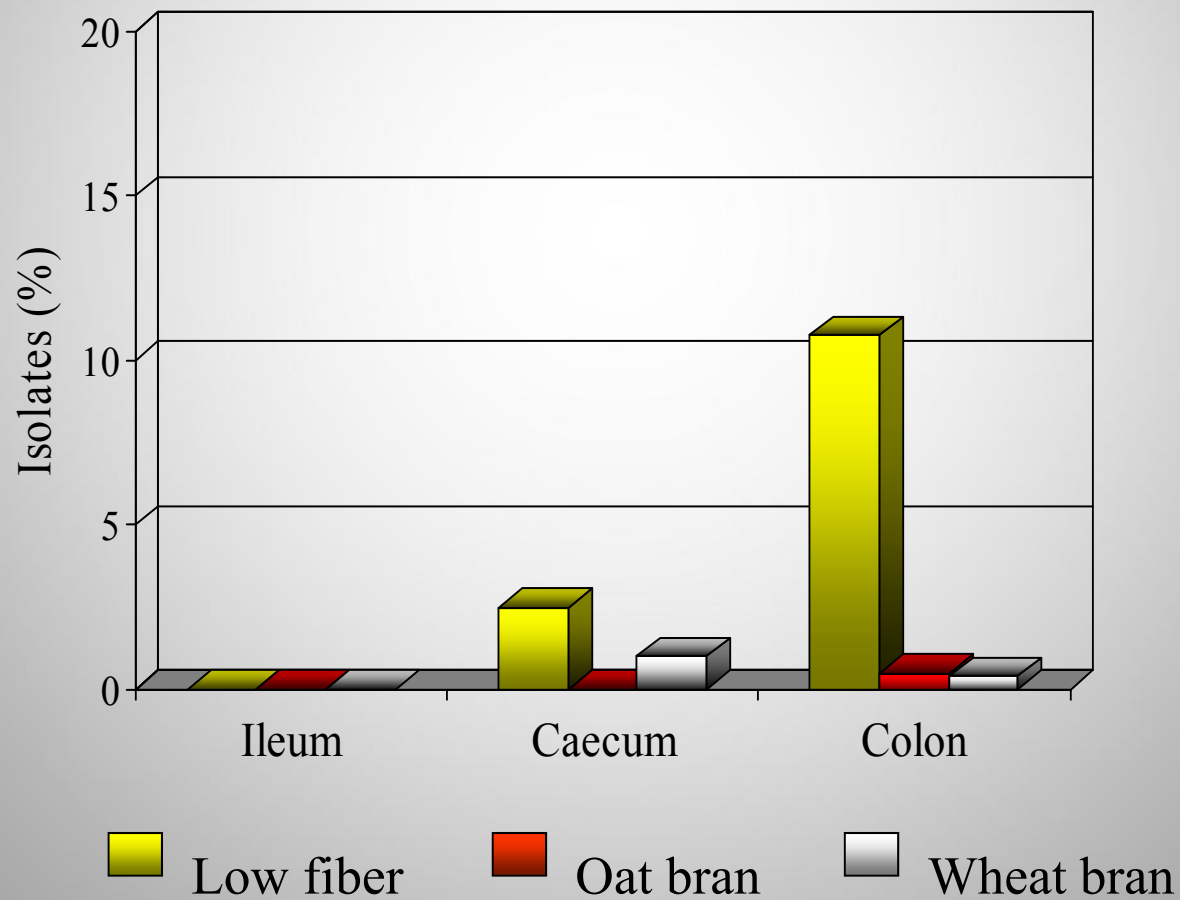
- In-feed antibiotics
- zinc oxide
- organic acids
- probiotics
- Plant extracts/species



Microbial activity in various regions of the gastrointestinal tract of pigs



Relative distribution (diets and segments) *Acidaminococcus fermentans*



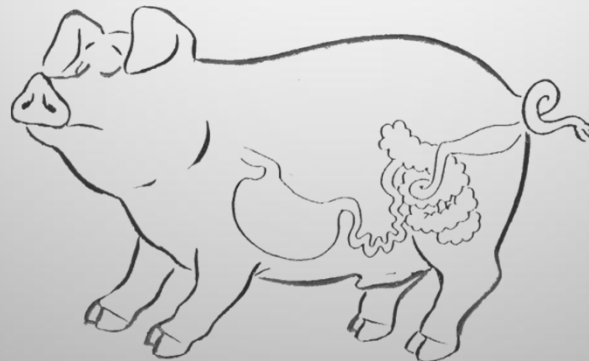
Factors that modulate the microbiota in the gastrointestinal tract

Feed

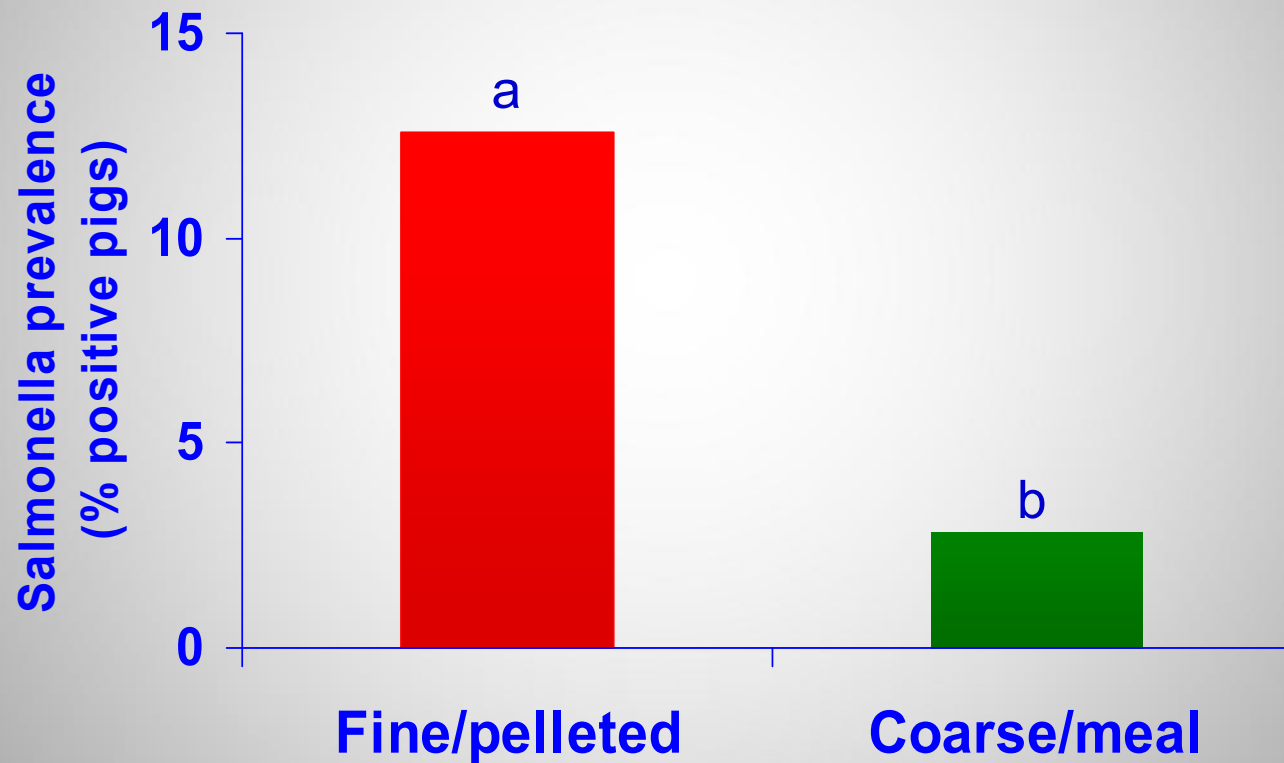
- Diet composition
- Feed processing
- Fermented liquid feed
- Prebiotics
- Low protein diets

Additives

- In-feed antibiotics
- zinc oxide
- organic acids
- probiotics
- Plant extracts/species

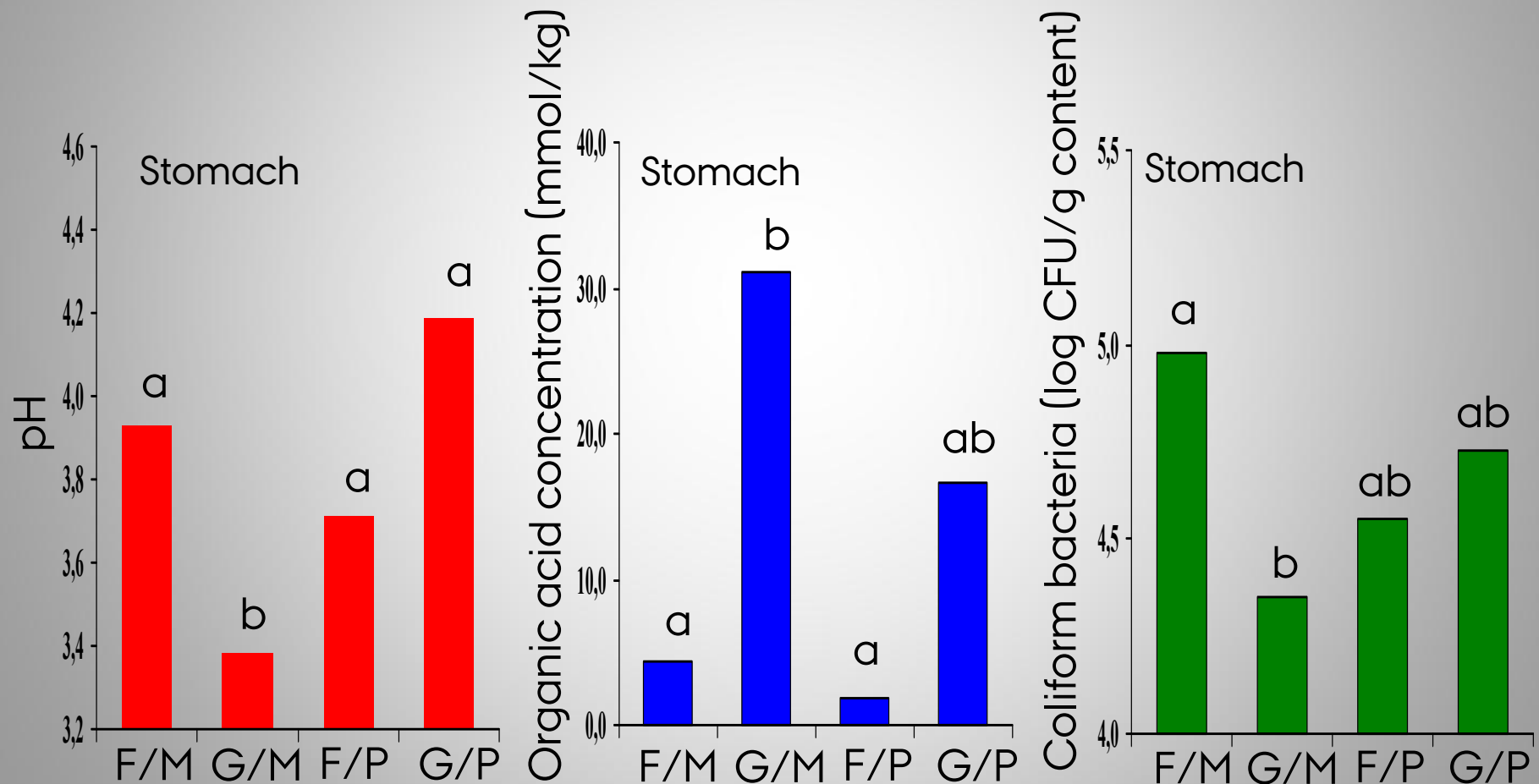


Effect of feed structure on Salmonella prevalence



Jørgensen et al., (1999)

Effect of grinding and pelleting on pH, organic acid concentration and the density of coliform bacteria in stomach content from slaughter pigs



Bacteria genera (T-RFs) in the stomach of pigs.
 Number of pigs in which the bacteria were found
 (out of 6)

| bp | Tentative identification | Fine | Coarse |
|-----|--|------|--------|
| 206 | <i>Unknown</i> | 3 | 5 |
| 223 | <i>L. delbreuckii subsp delbreueckii</i> | 0 | 6 |
| 253 | <i>L. delbreuckii subsp bulgaricus</i> | 0 | 4 |
| 267 | <i>L. mucosae</i> | 0 | 6 |
| 390 | <i>M. jalaludinii/multiacidia</i> | 0 | 6 |
| 406 | <i>L. reuteri</i> | 0 | 6 |
| 574 | <i>Unknown</i> | 0 | 6 |
| 581 | <i>S. alactolyticus/hyointestinalis</i> | 2 | 4 |
| 589 | <i>S. ruminantum</i> | 0 | 5 |
| 591 | <i>M. elsdenii</i> | 0 | 5 |
| 595 | <i>L. sorbius</i> | 2 | 5 |
| 611 | <i>Weisella sp.</i> | 0 | 6 |
| 605 | <i>L. curvatus</i> | 0 | 5 |

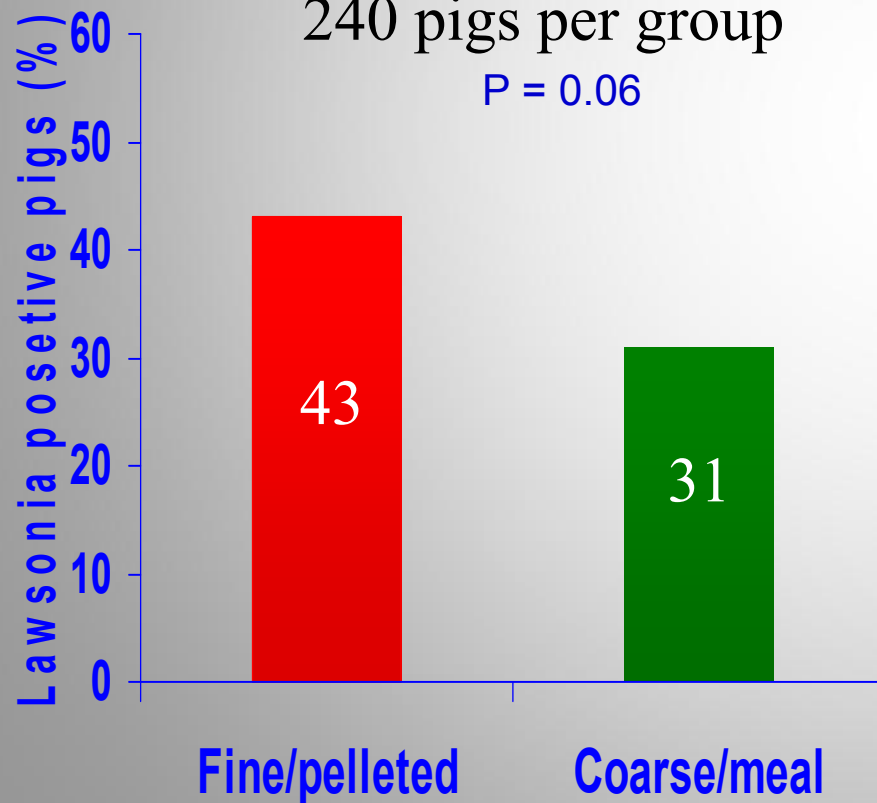
Effekt af feedstruktur og tylosin on infection with *Lawsonia intracellularis*

Feed structure

Johansen et al., (2003)

240 pigs per group

P = 0.06

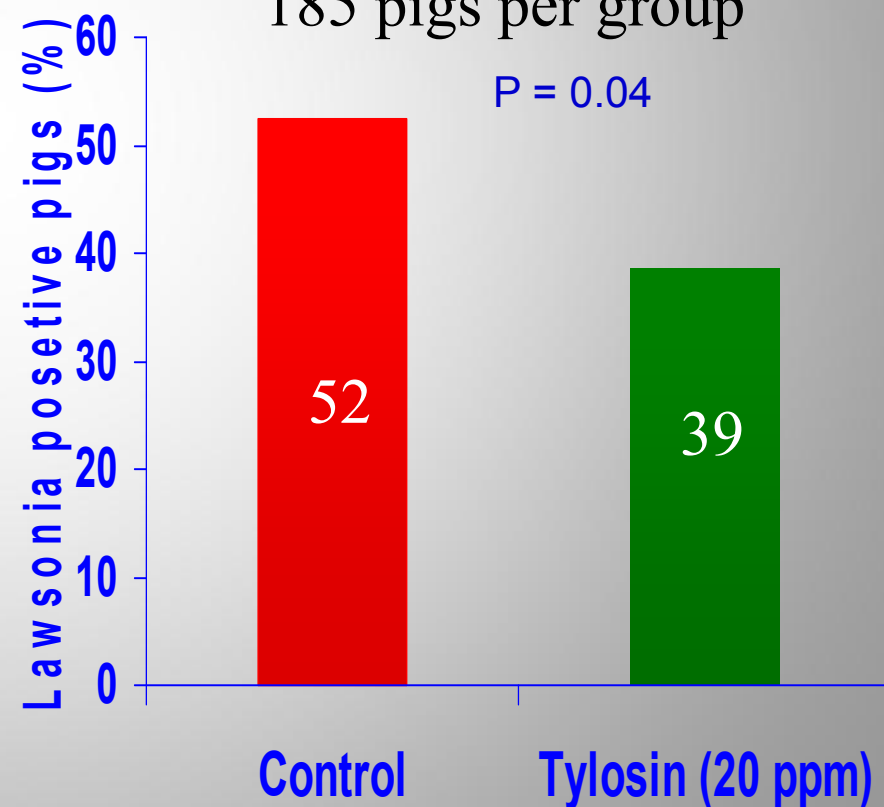


Tylosin

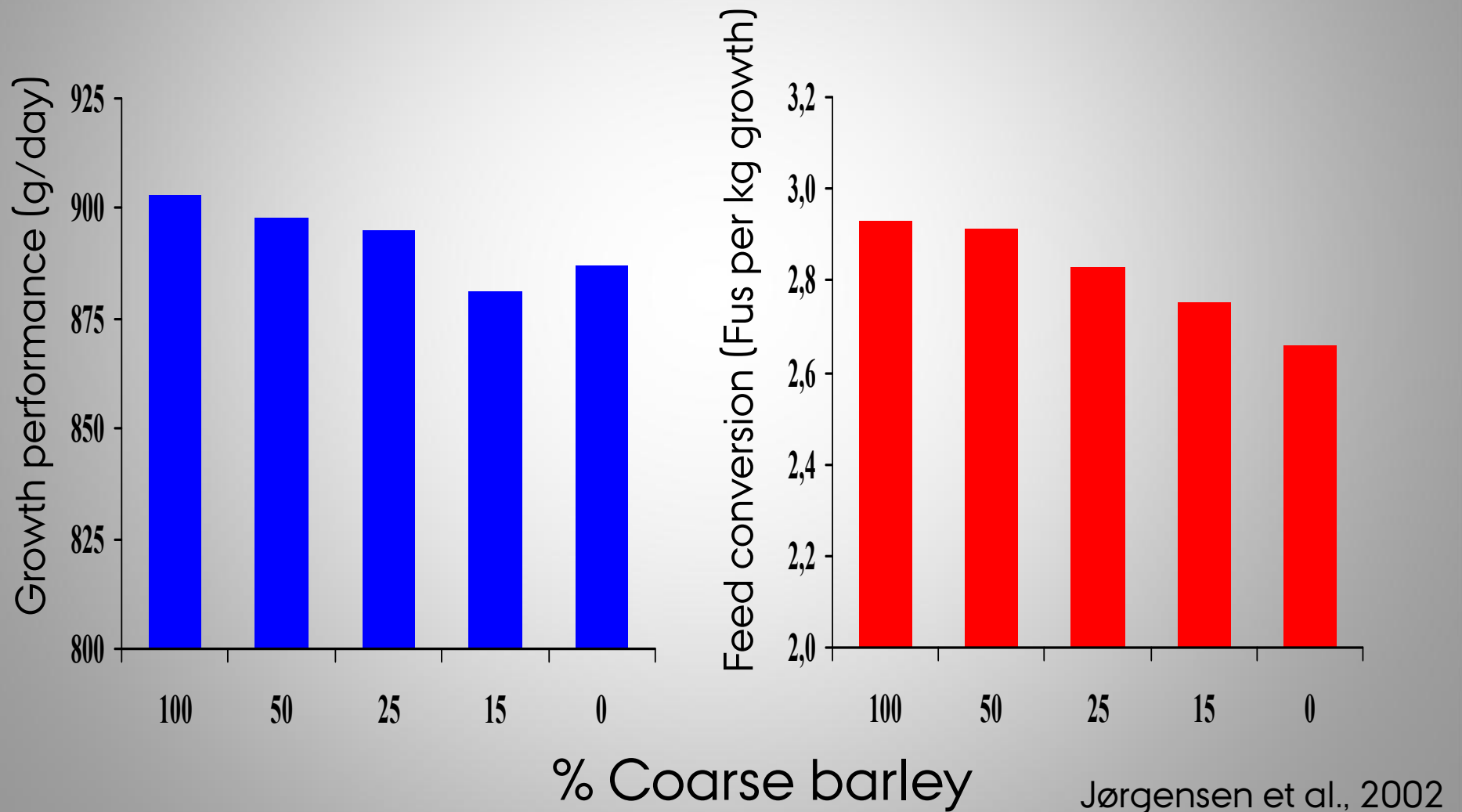
Pedersen et al., (2003)

185 pigs per group

P = 0.04



Effect of particle composition (fine vs. coarse barley) on growth performance and feed conversion.



The stomach as a barrier

Increased microbial activity in the stomach



Increased production of organic acids → decrease pH



Kill enterobacteria entering the stomach



Lower proliferation of enterobacteria in the small intestine



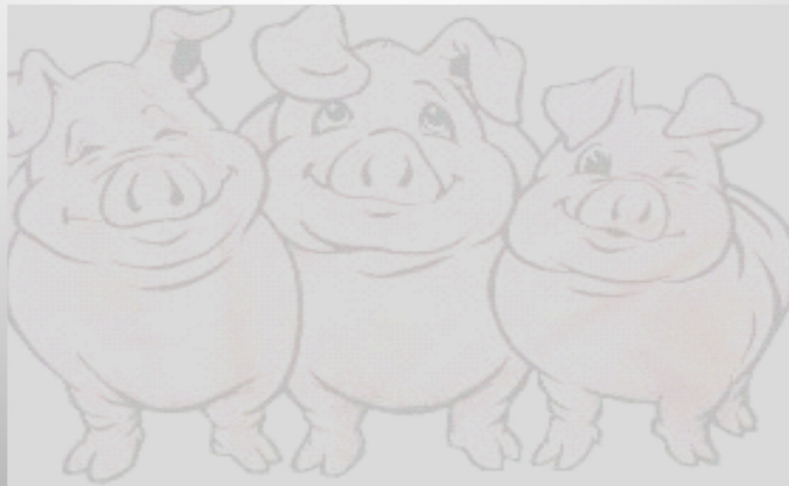
Fewer enterobacteria excreted with faeces



Breaking the vicious circle where pigs in problematic herds infect each other and are re-infected through consumption of faeces

Three ways to reduce pH and increase the concentrations in stomach content:

- › Addition of organic acids to the feed
- › Use coarse grounded feed
- › Use of fermented liquid feed



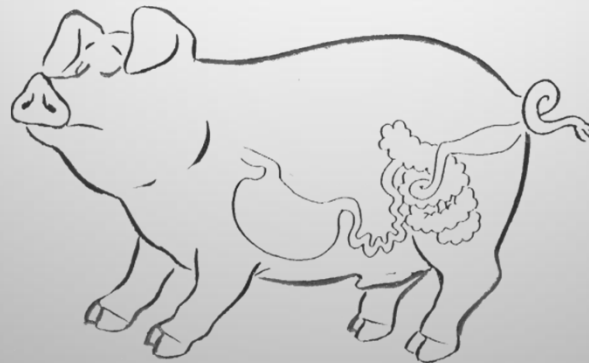
Factors that modulate the microbiota in the gastrointestinal tract

Feed

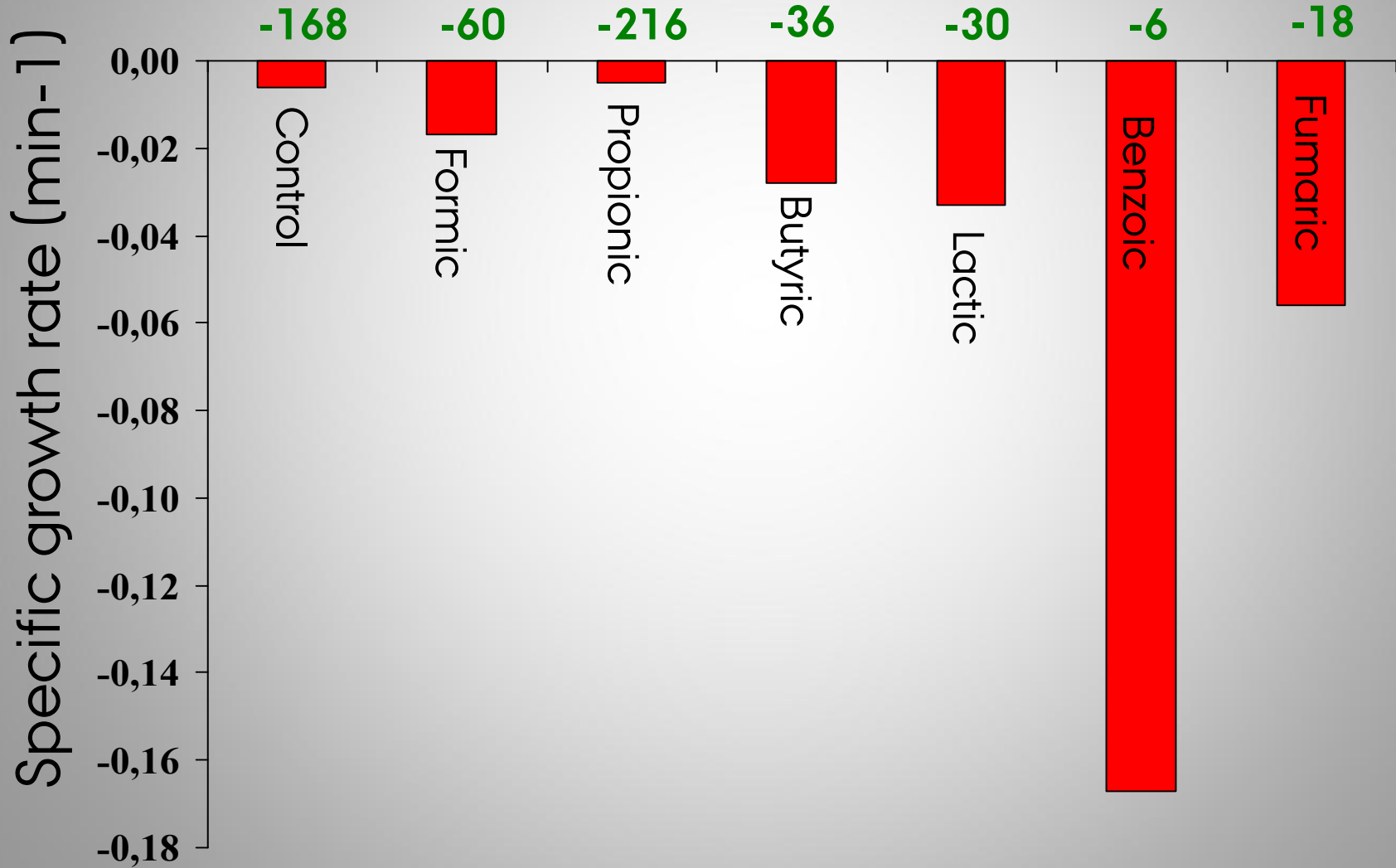
- Diet composition
- Feed processing
- Fermented liquid feed
- Prebiotics
- Low protein diets

Additives

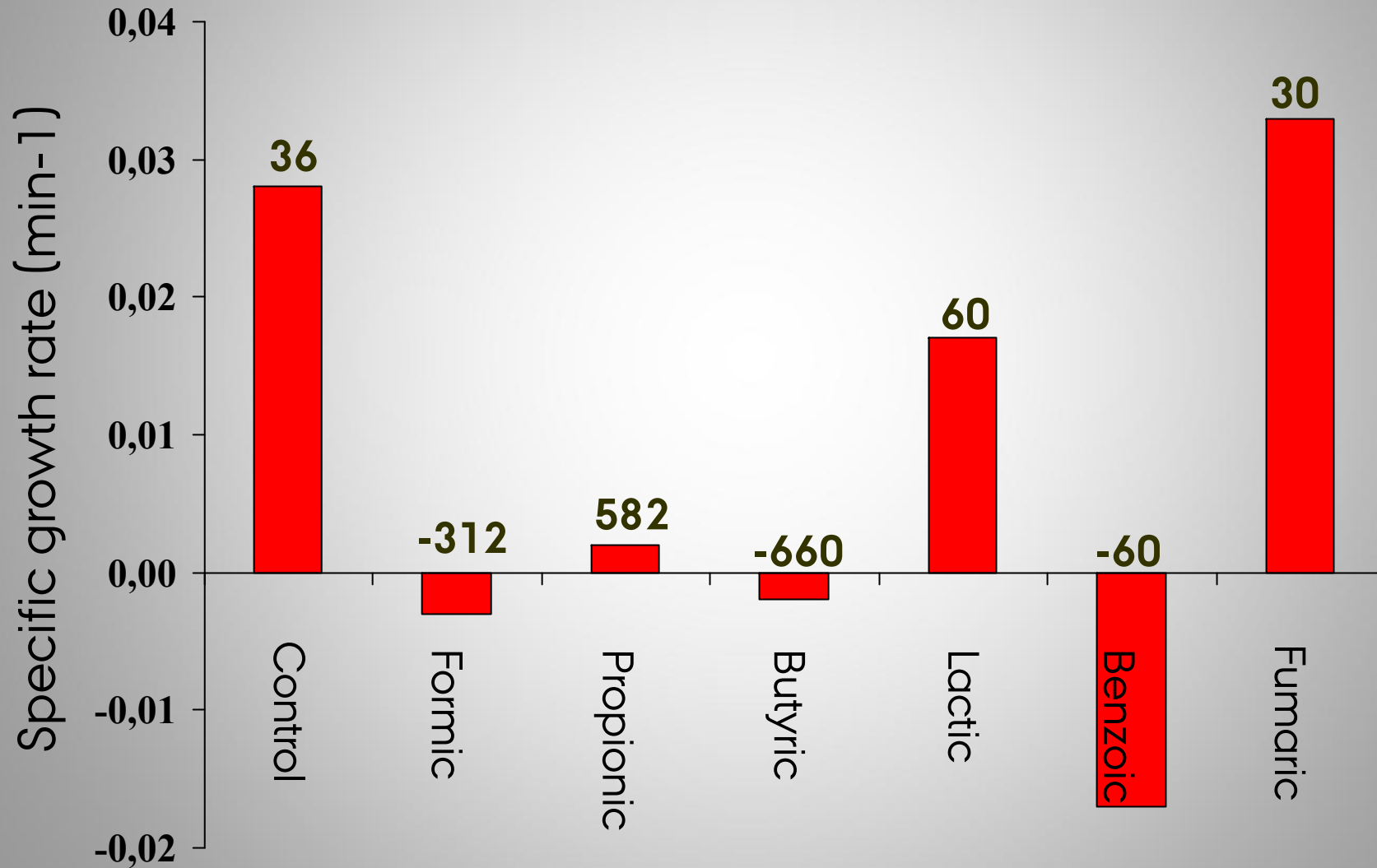
- In-feed antibiotics
- zinc oxide
- **organic acids**
- probiotics
- Plant extracts/species



Effect of various organic acids (100 mM) on specific growth rate of coliform bacteria in gastric content at pH 4.5

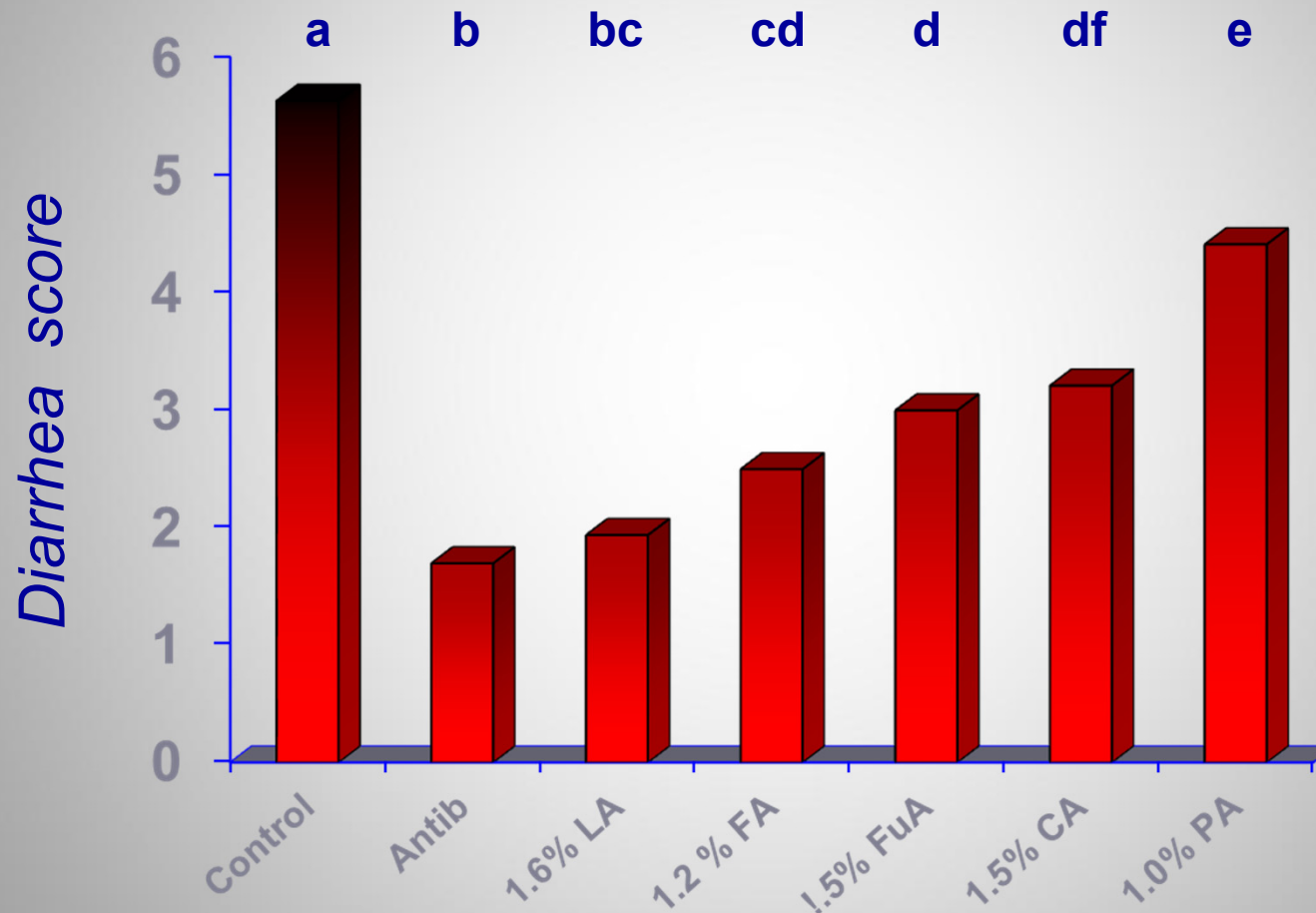


Effect of various organic acids (100 mM) on specific growth rate of coliform bacteria in content from the small intestine at pH 5.5



Effect of organic acid on diarrhea in a herd with ETEC problems

(Weaning at 25 days, 4 weeks in experiment)

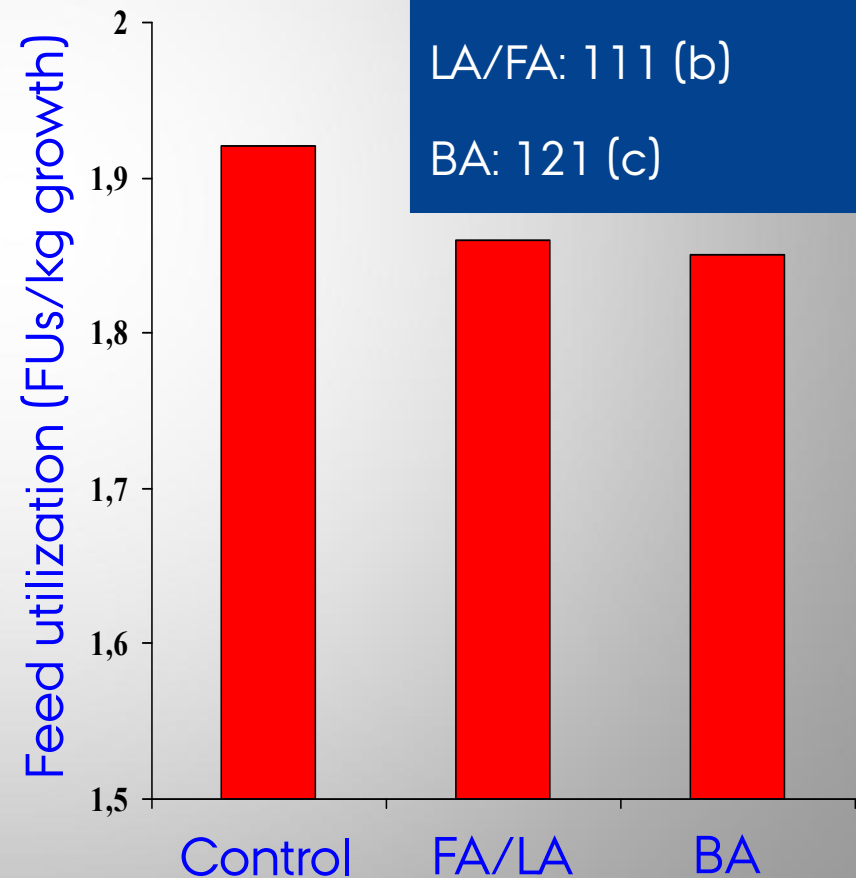
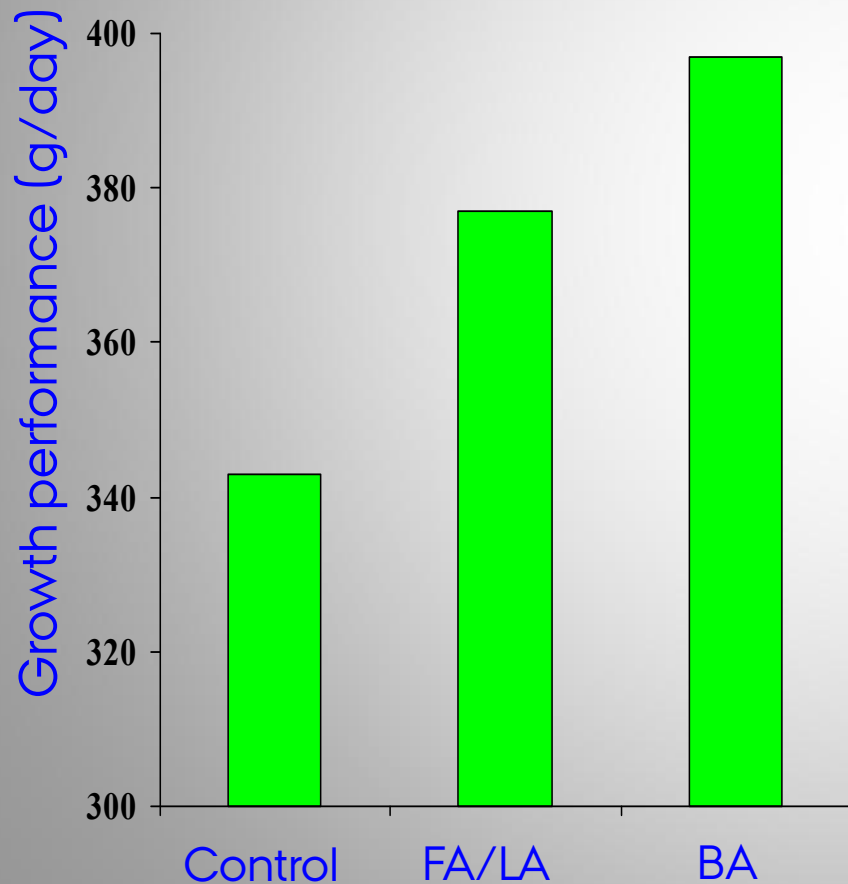


N= 4, one replicate is one pen with 12 pigs.

Antibiotics: 40ppm lincomycin og 44ppm spectinomycin

Tsiloyiannis et al., 2001

Effect of 0.7% formic acid /0.7% lactic acid and 2% benzoic acid on growth performance and feed utilisation by piglets after weaning (4-10 weeks)



Performance index:

Control: 100 (a)

LA/FA: 111 (b)

BA: 121 (c)

Effect of organic acids and zinc oxide on diarrhea and killing out range of weanears

| | Control | Acid* | Acid + ZnO** |
|-----------------------------------|---------|----------|--------------|
| Treatment against diarrhea (days) | 8,7 % a | 6,9 % b | 0,9 % c |
| Killing out range | 2,6 % a | 1,9 % ab | 1,2 % b |

*: 1% lactic acid 1% formic acid og 0,5% benzoic acid

** : 2500 zinc as zinc oxide

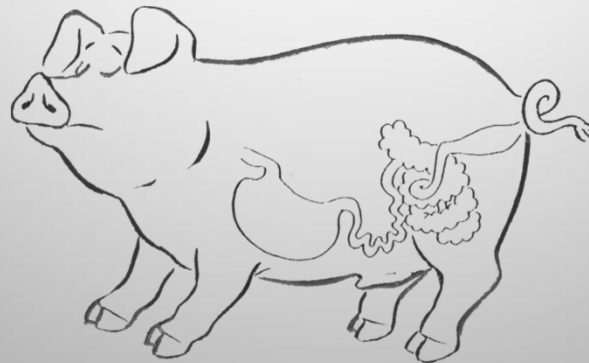
Factors that modulate the microbiota in the gastrointestinal tract

Feed

- Diet composition
- Feed processing
- **Fermented liquid feed**
- Prebiotics
- Low protein diets

Additives

- In-feed antibiotics
- zinc oxide
- organic acids
- probiotics
- Plant extracts/species



Definitions

Liquid feed (LF): water and feed are mixed shortly before feeding.

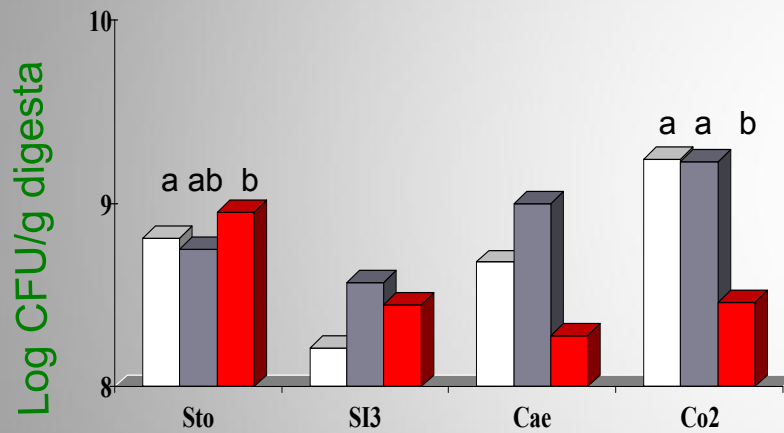
Fermented liquid feed (FLF): feed and water are mixed and soaked during a certain period of time, at a certain temperature before feeding.

Characteristics of dry feed, LF and FLF

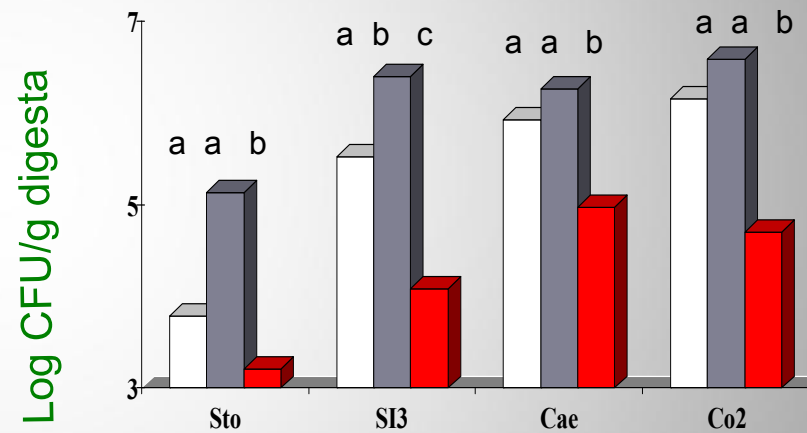
| | DF | LF | FLF |
|-------------------------------|------------|-------------|-------------|
| DM% | 89.3 | 27.3 | 24.6 |
| pH | - | 5.9 | 4.4 |
| Lactic acid bact. (log CFU/g) | <4.3 | 7.2 | 9.4 |
| Enterobacteria (log CFU/g) | <4.7 | 6.2 | <3.2 |
| Energy (MJ/kg DM) | 18.7 | 19.0 | 19.0 |
| Lysine (g/16 g N) | 6.0 | 5.8 | 4.8 |
| LMW-sugars (% DM) | 3.6 | 2.9 | 0.1 |

Microbial populations in the GIT

Lactic acid bacteria

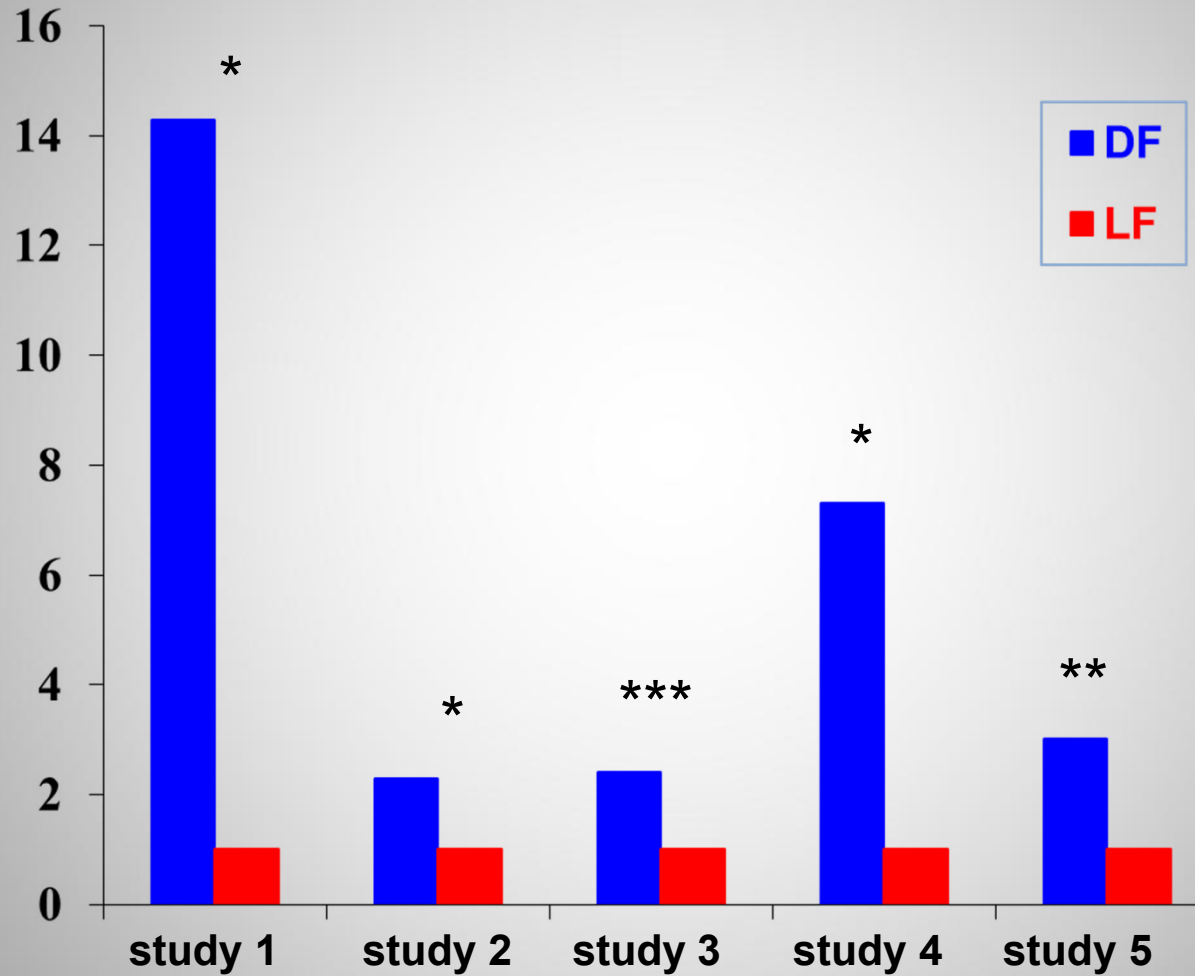


Coliform bacteria



DF NFLF FLF

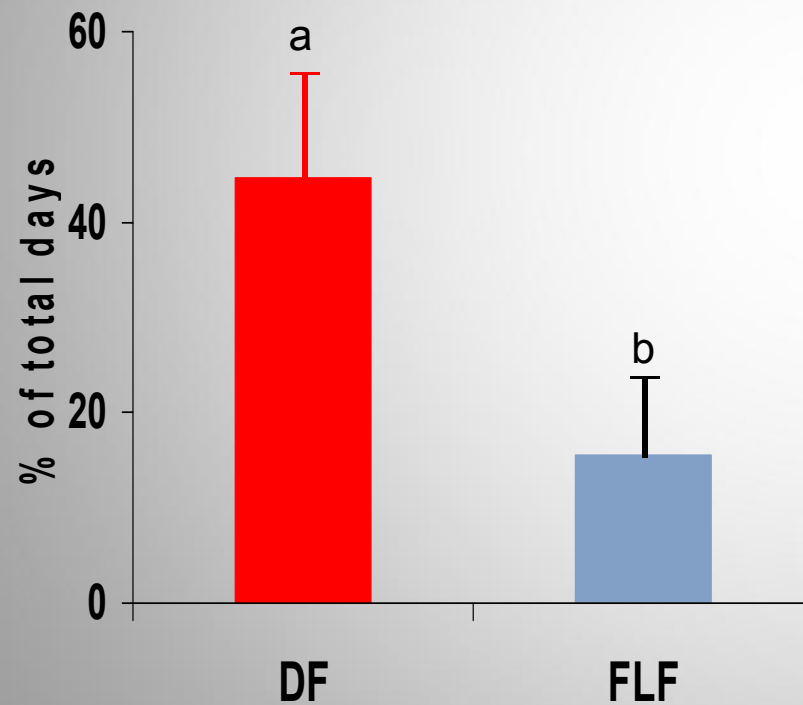
Dry feed as a risk factor for *Salmonella* incidence



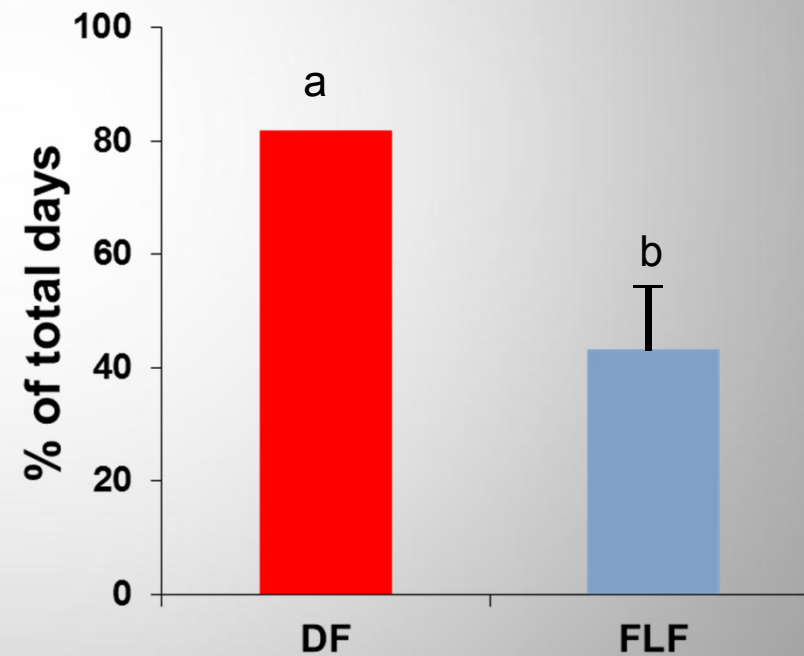
Bager, 1993; Bager & Emborg, 1994; Dahl, 1997, Stege et al., 1997; Dahl (unpubl.)

Effect of FLF on swine dysentery

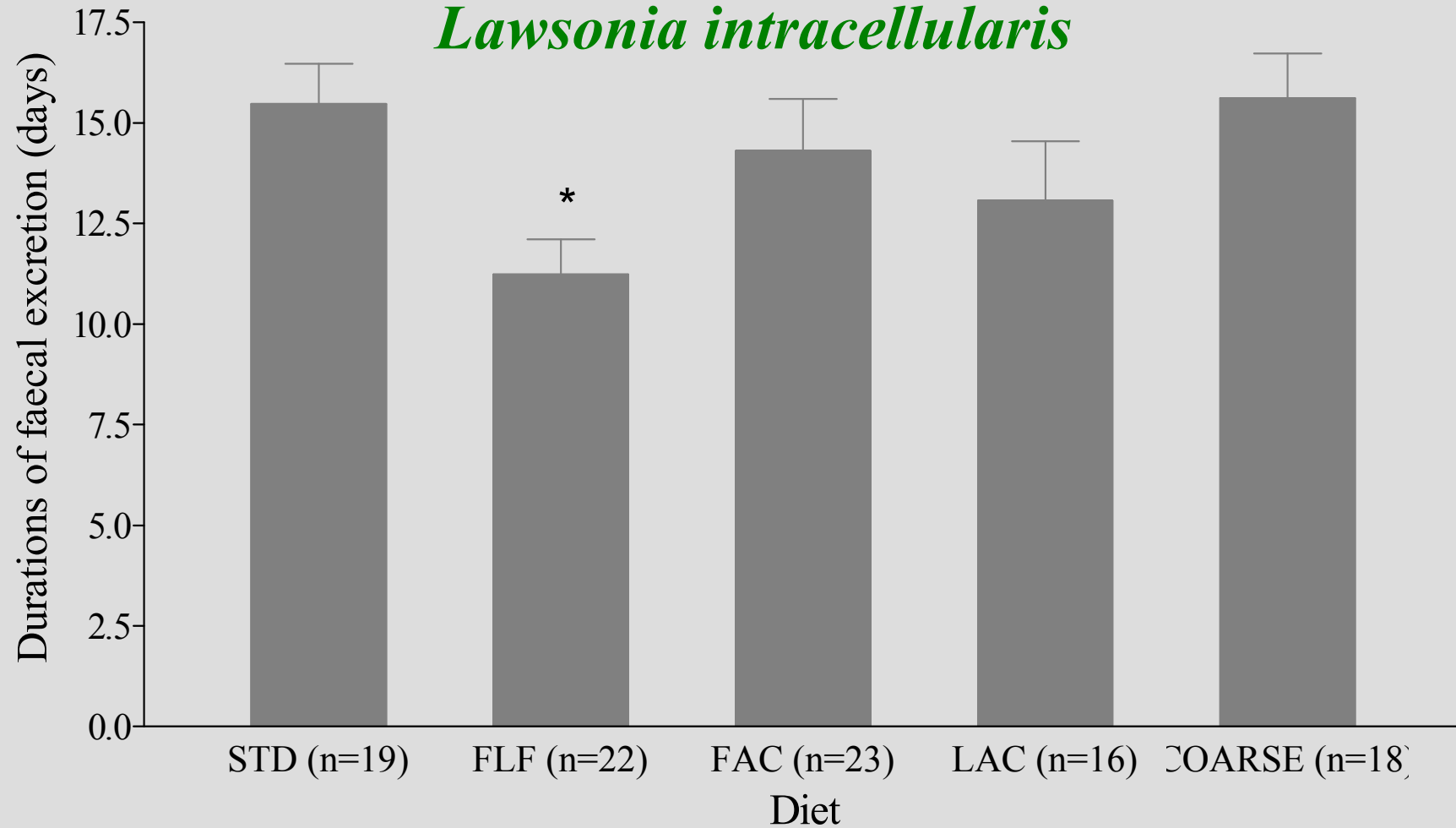
Positive culture from faeces
of *B. hyodysenteriae*



Clinical disease after infection
with *B. hyodysenteriae*



Effect of FLF on infection with *Lawsonia intracellularis*



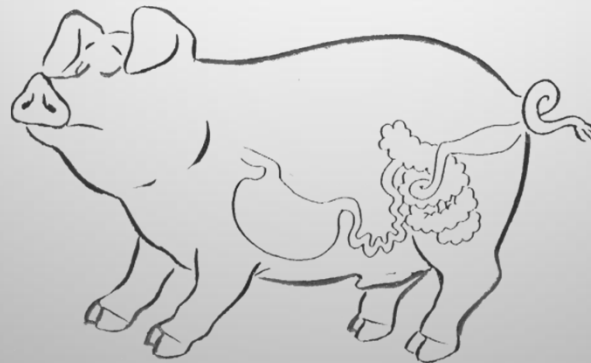
Factors that modulate the microbiota in the gastrointestinal tract

Feed

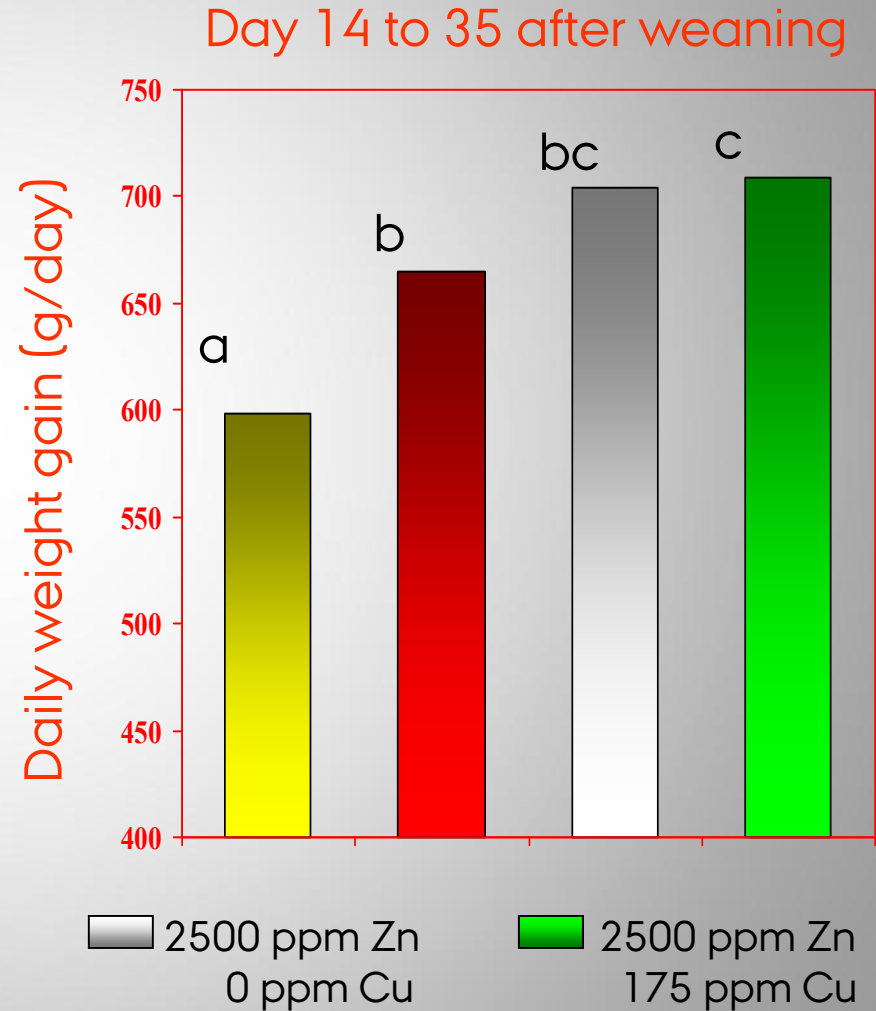
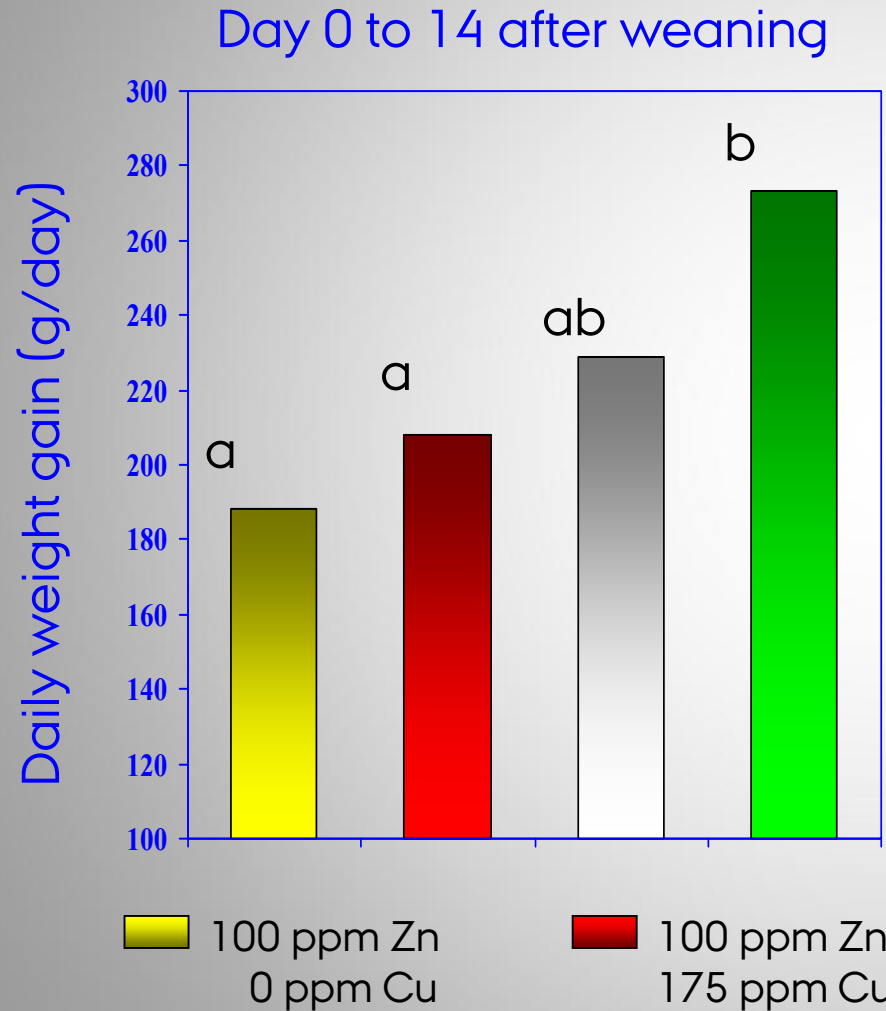
- Diet composition
- Feed processing
- Fermented liquid feed
- Prebiotics
- Low protein diets

Additives

- In-feed antibiotics
- **zinc oxide**
- organic acids
- probiotics
- Plant extracts/species

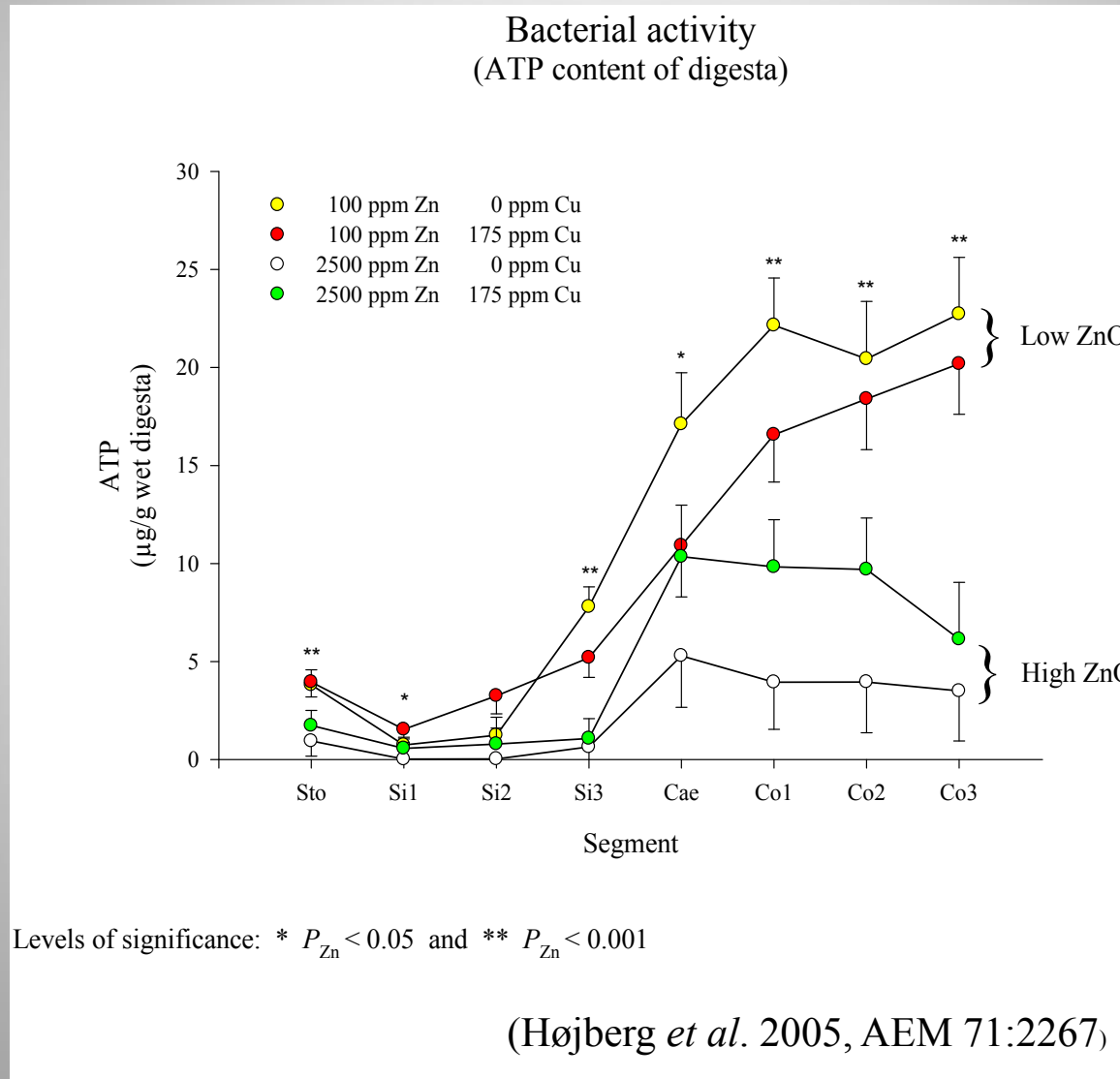


Effect of zinc and copper on daily weight gain in piglets after weaning (at 4 weeks)

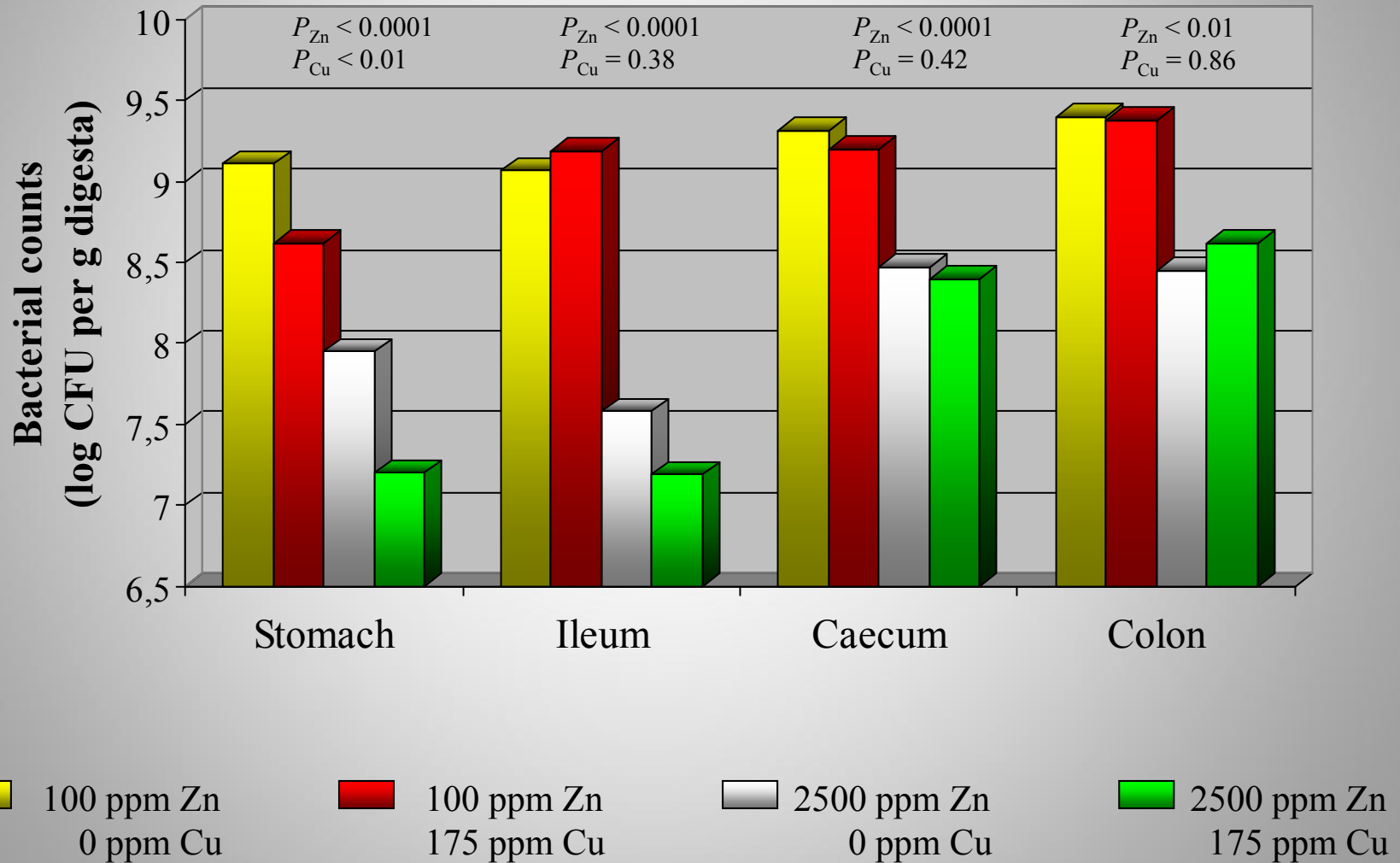


Poulsen et al. (2003)

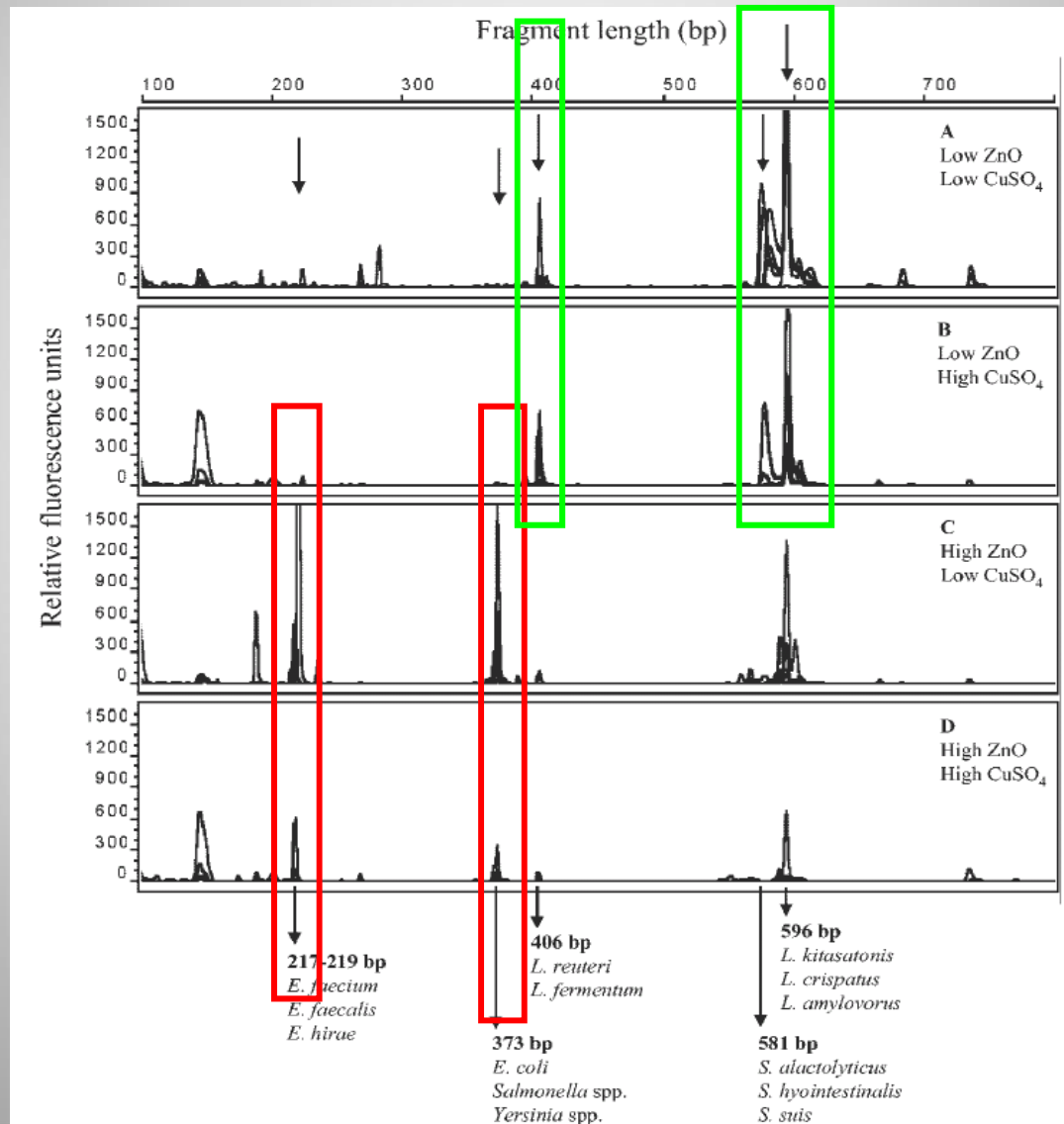
Bacterial activity



Lactic acid bacteria (MRS agar)



T-RFLP (caecum)



Low ZnO

High ZnO

(Højberg *et al.* 2005, AEM 71:2267)

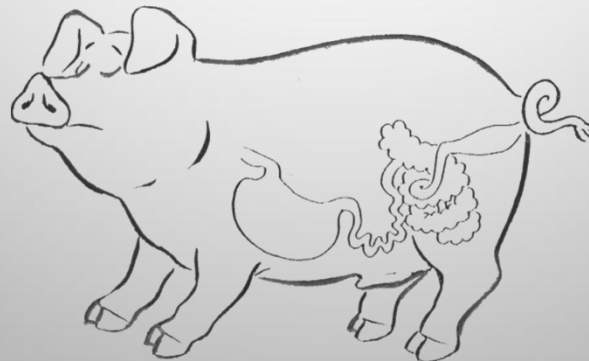
Factors that modulate the microbiota in the gastrointestinal tract

Feed

- Diet composition
- Feed processing
- Fermented liquid feed
- Prebiotics
- Low protein diets

Additives

- In-feed antibiotics
- zinc oxide
- organic acids
- **probiotics**
- Plant extracts/species



Probiotica

Viable defined microorganism, which alter the microbiota in a way that exert beneficial effects on the host

The three most commonly used organisms as probiotics:

bacillus, **yeast** and **lactic acid producing bacteria**
(lactobacillus, bifidobacterium and enterococcus)

Inconstancy in results properly due to:
dosage and type of strains, environment and diet type

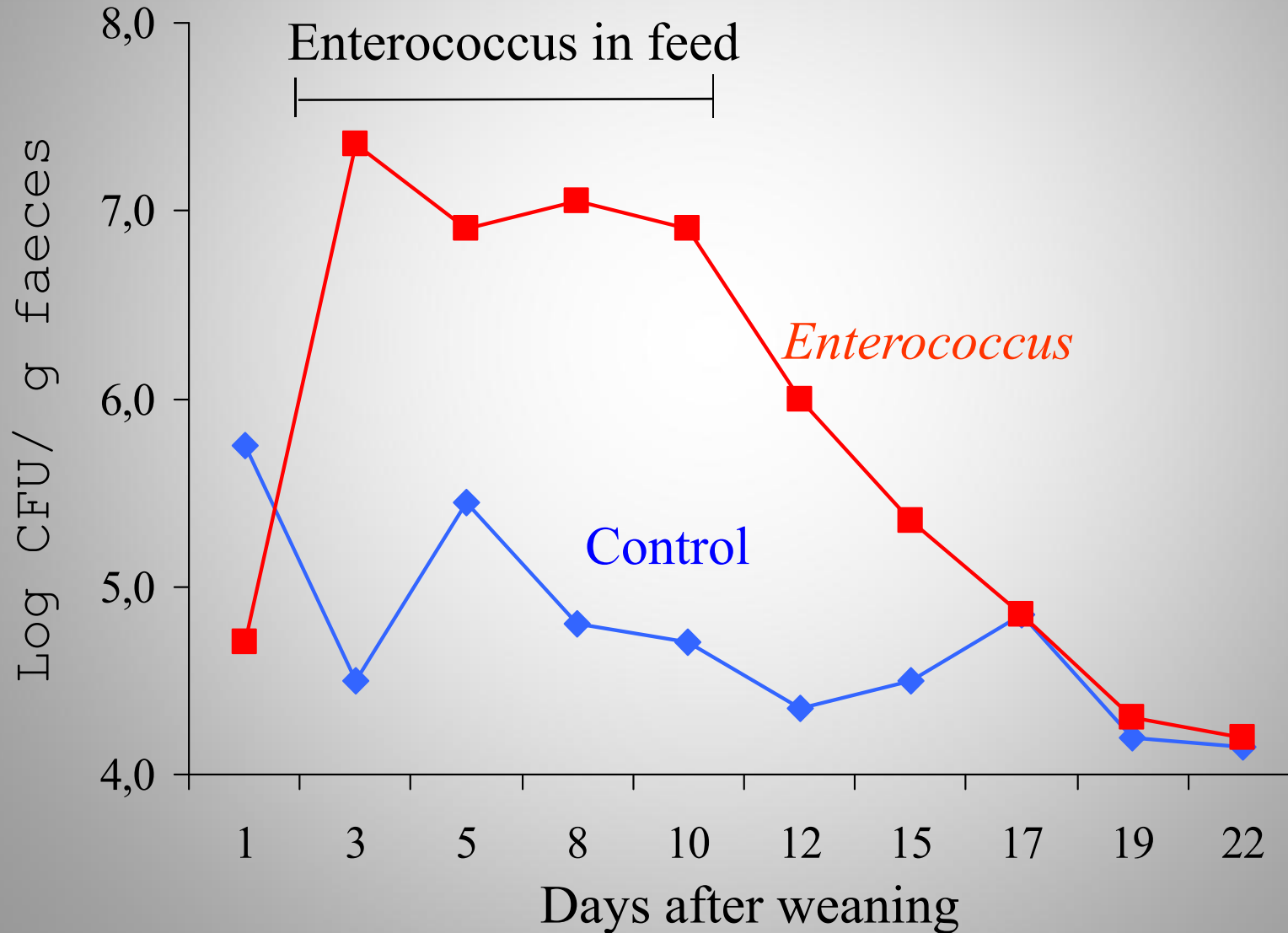
Needed:

strains as well of doses that gives consistently good results

Mechanism not completely understood, a few have been proposed:

- *inhibit pathogen adhesion by steric hindrance or competitive exclusion*
- *production of products with antibacterial activity (bacteriocins or OA)*
- *modulation of the host immune system*

*Development of the population of Enterococci in faeces from piglets fed a control diet or a diet supplemented with 10^9 *Enterococcus faecium**



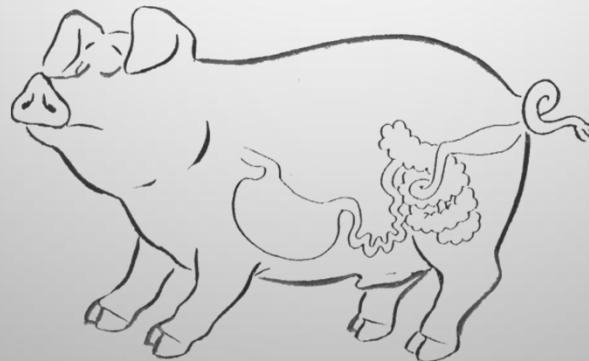
Factors that modulate the microbiota in the gastrointestinal tract

Feed

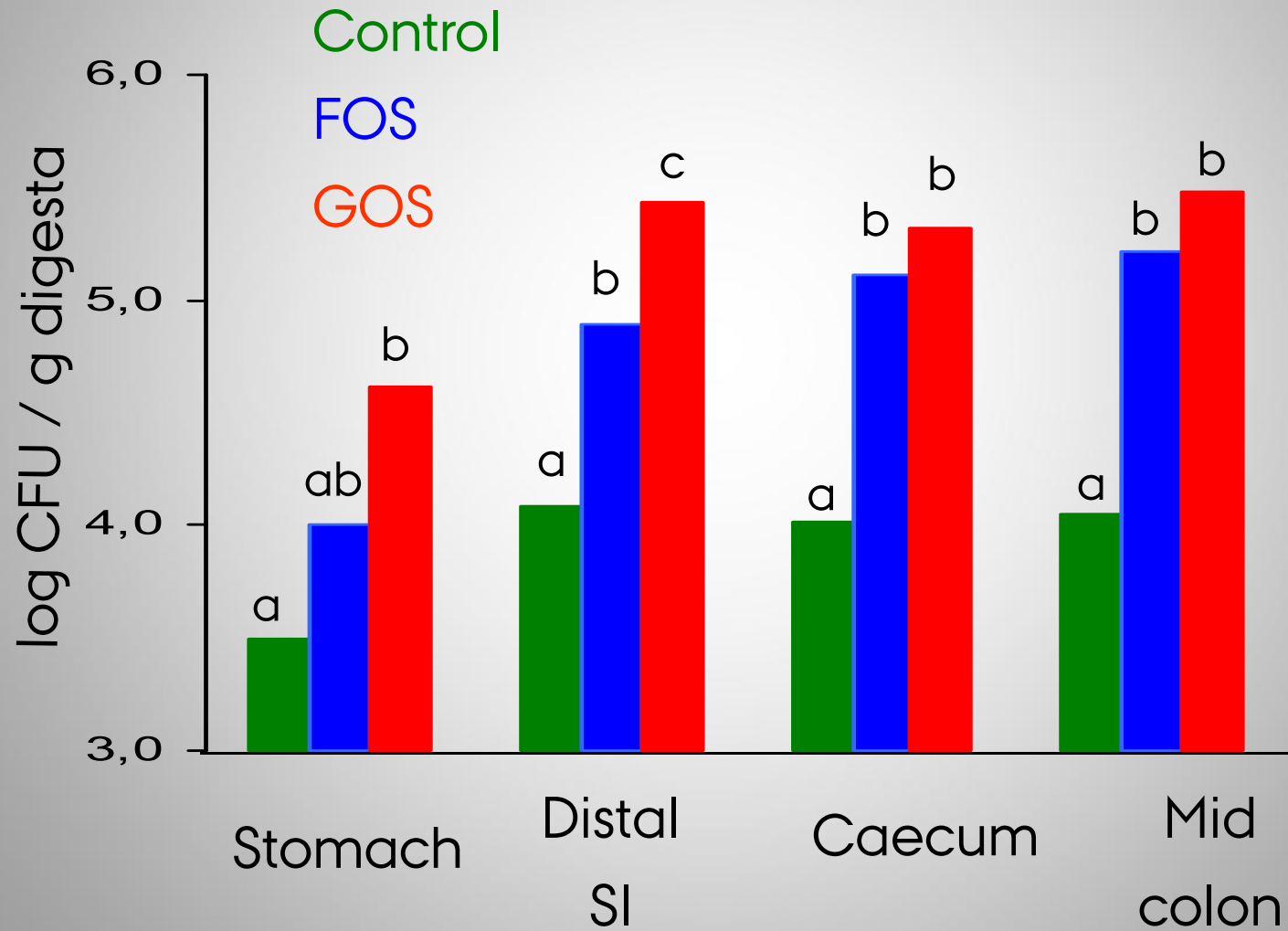
- Diet composition
- Feed processing
- Fermented liquid feed
- **Prebiotics**
- Low protein diets

Additives

- In-feed antibiotics
- zinc oxide
- organic acids
- probiotics
- Plant extracts/species



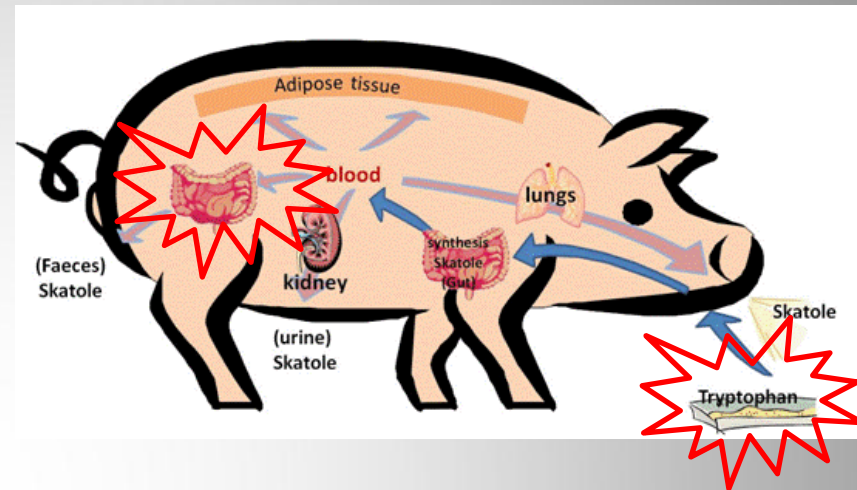
Effect of NDO on yeast



Boar taint

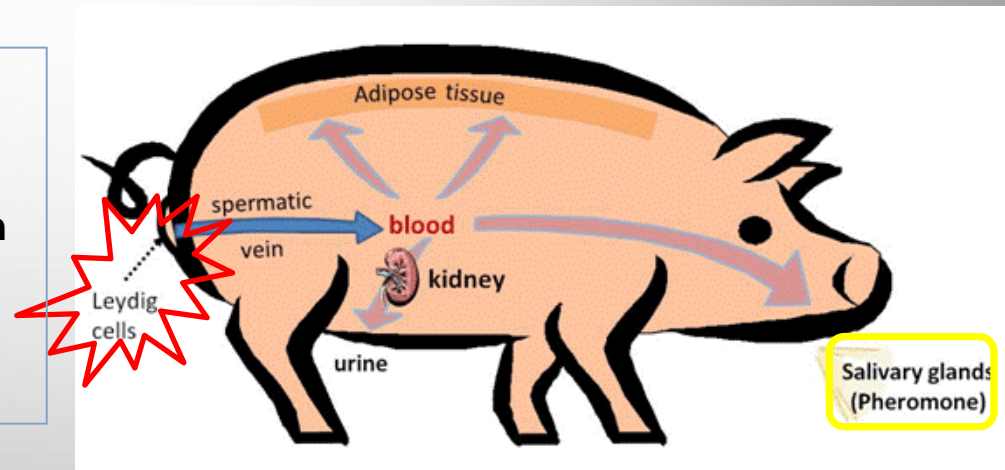
- **Skatole**

- Produced in the large intestine from microbial metabolism of tryptophan
- Can be reduced by dietary intervention (high fermentable fibre, protein type)



- **Androstenone**

- Produced in the testicles
- From there to blood, saliva, and from liver to (bile), urine, adipose tissue.



Task 2.6 Experiment with paraxites/chicory root



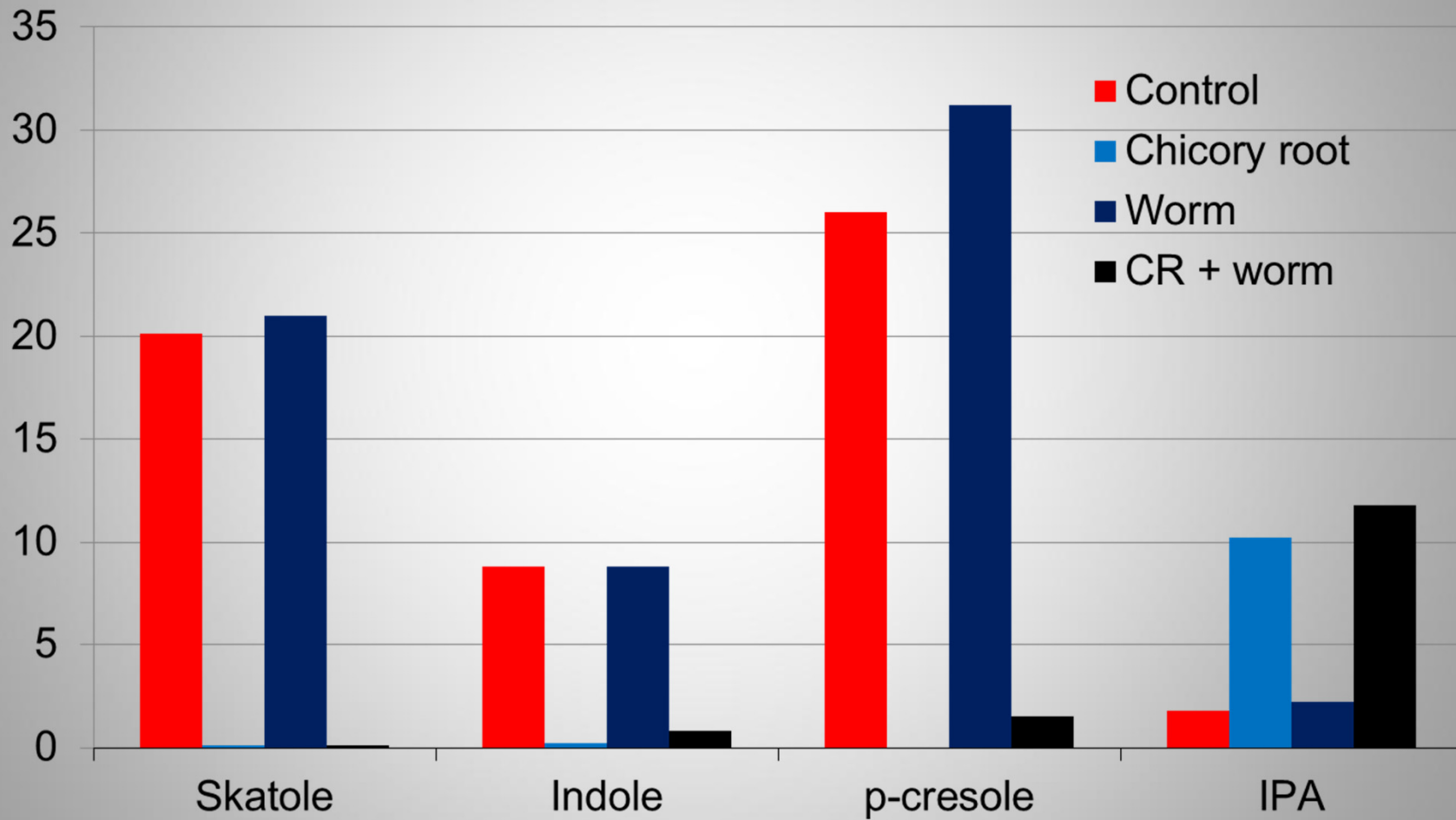
- › Two-factorial experiment (n=72)
- › Pigs slaughtered at three times (after 5, 9 and 12 weeks on the experimental diets)



Nudelworm

| | - Chicory root | + Chicory root (25%) |
|--------|-----------------|-------------------------|
| - worm | Control (n=3x6) | hickorye (n=3x6) |
| + worm | Worm (n=3x6) | Chickory + worm (n=3x6) |

Indols and p-cresol in colon content



Androstenone and skatole in back fat

| | Control | | Control Worm | | Chicory root | | Chicory root Worm | |
|------------------------|---------|--------|-----------------|--------|--------------|--------|----------------------|--------|
| | Male | Female | Male | Female | Male | Female | Male | Female |
| Androstenone (ng/g) | 1,48 | 0,01 | 1,21 | 0,00 | 1,24 | 0,01 | 1,53 | 0,01 |
| Skatole (µg/g) | 108,7 | 57,0 | 93,8 | 70,1 | 0,0 | 0,0 | 0,0 | 0,8 |
| Indole (µg/g) | 21,4 | 8,6 | 14,6 | 15,2 | 7,3 | 1,8 | 7,4 | 6,1 |

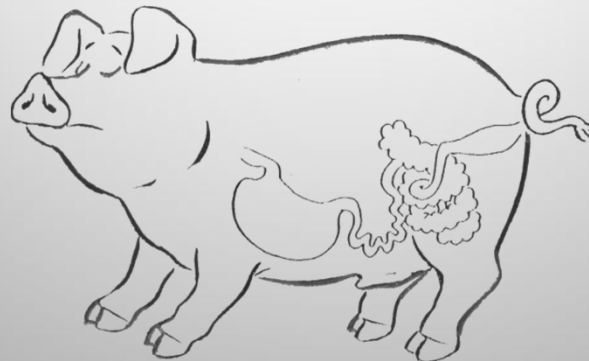
Factors that modulate the microbiota in the gastrointestinal tract

Feed

- Diet composition
- Feed processing
- Fermented liquid feed
- Prebiotics
- **Low protein diets**

Additives

- In-feed antibiotics
- zinc oxide
- organic acids
- probiotics
- Plant extracts/species



Low protein diets

Source and level known to affect enteric health of weaned pigs

Source:

- animal protein better than plant protein
- more knowledge on microbial composition and activity needed

Level:

- low protein diets supplemented with synthetic AA prevent post weaning diarrhoea without compromising growth performance

Conclusion

Feed

- *FLF*: Increase gut health but reduce growth performance may be a way to make probiotics cost effective.
- *Feed structure*: Increase gut health but reduce growth performance
- *Prebiotics*: Inconsistent results. More research needed. May not be cost effective.

Additives

- *ZnO*: increase growth performance and reduce PWD, but gives environmental problems
- *OA*: right doses and types have beneficial effects on growth performance and health
- *Probiotics*: Inconsistent results. More strains should be tested. May not be cost effective