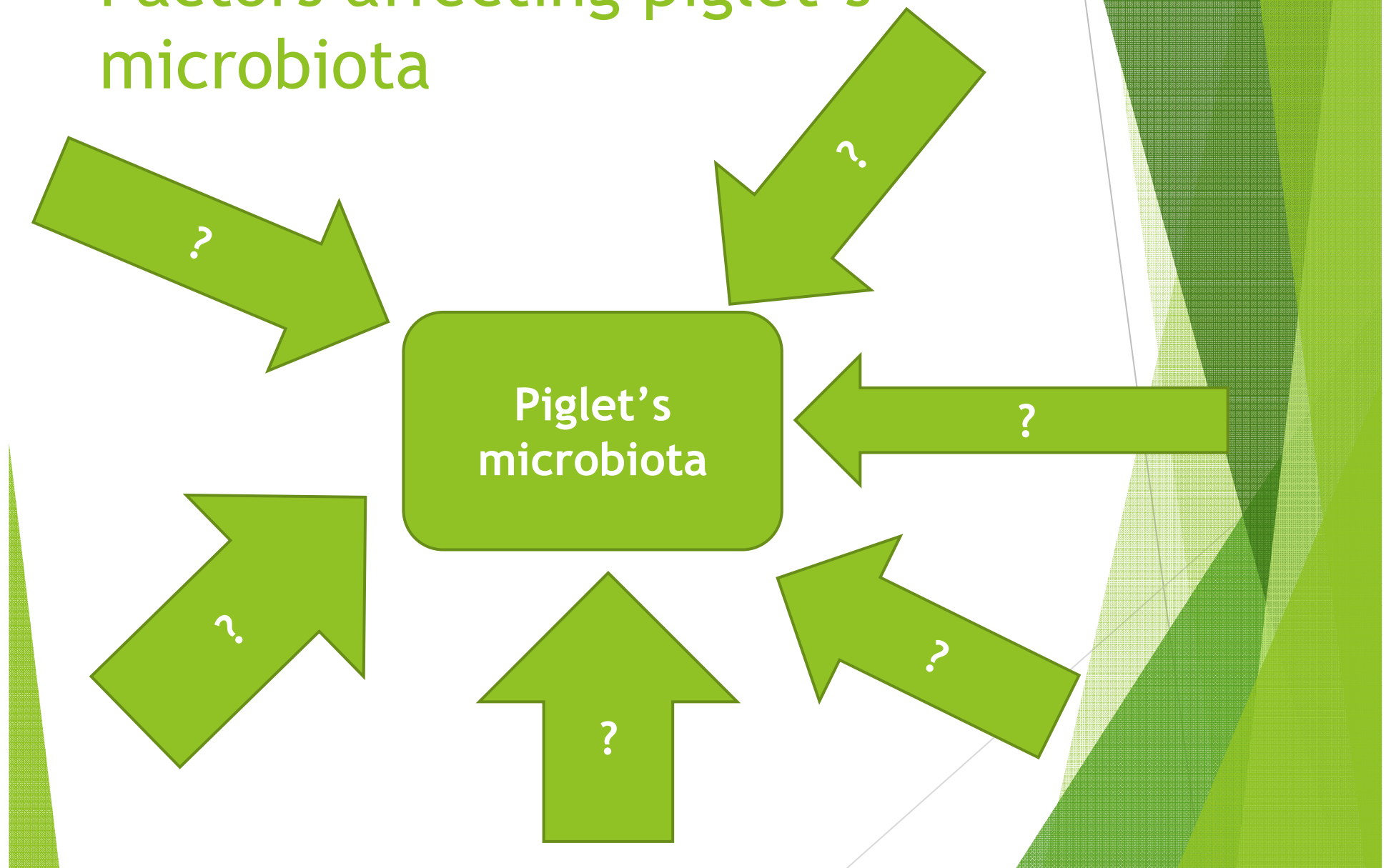


# Adding fermentable feed ingredients to pigs: changes or stability to the intestinal microbiota

Nadia Everaert, PhD, Associate Professor  
Liège Université- Gembloux Agro-Bio Tech  
Precision Livestock and Nutrition Unit  
Belgium

# Factors affecting piglet's microbiota



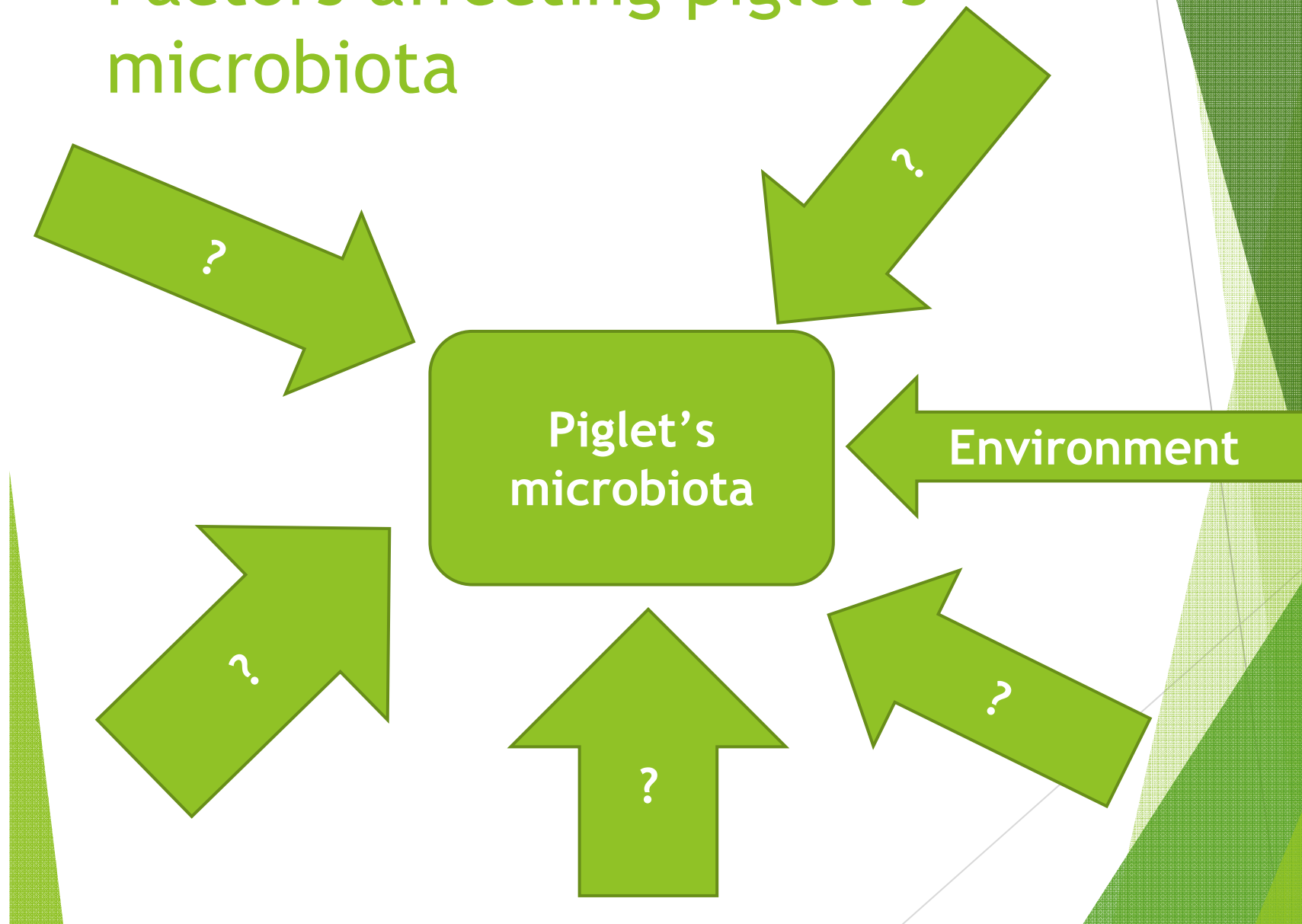
# Early life colonization

- ▶ Pigs separated from sow after birth and reared on milk replacer
- ▶ Non-siblings were co-housed in pairs
- ▶ The community of piglets **older than 31 days** was inferred to show **high stability relative to the first 28 days post birth**
- ▶ Significant **correlation** of microbial communities **between cohabiting piglets**, but not between siblings

**Gut colonization in piglets is greatly influenced by the immediate environment**

Thompson et al., 2008

# Factors affecting piglet's microbiota



# Early life colonization

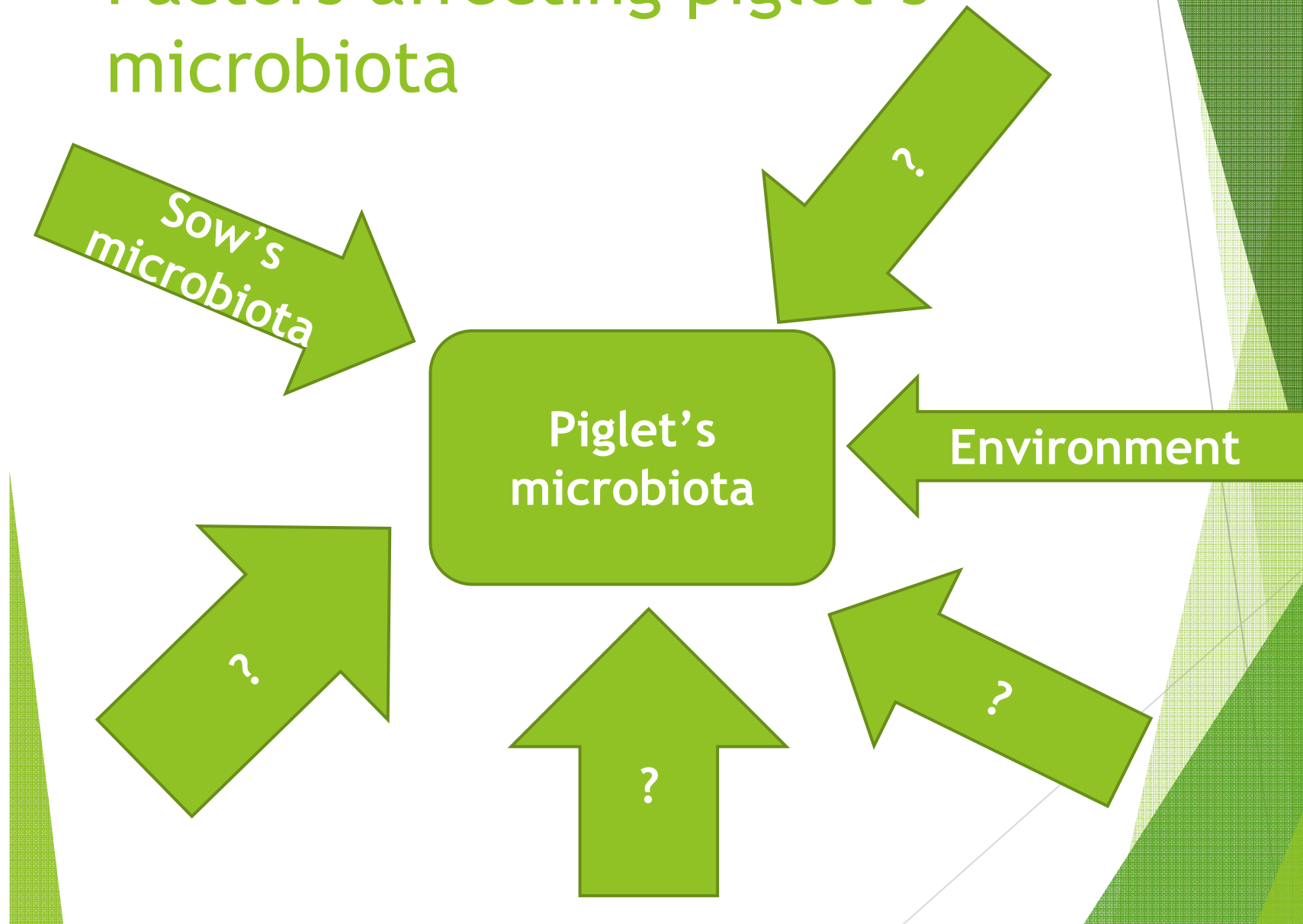
- ▶ Piglets born by caesarian section, kept in clean isolators during 28 d
- ▶ Two treatments:
  - ▶ Inoculum + saline inoculations (SA)
  - ▶ Inoculum + complex sow's faeces' inoculations (CA)
- ▶ *Inoculum: Lactobacillus amylovorus, Clostridium glycolicum, and Parabacteroides spp.*
- ▶ Faecal microbiota composition increased in diversity in time
- ▶ Clustering of CA faeces with sow's faeces
- ▶ SA treatments: a lower similarity in microbial composition (diarrhea)



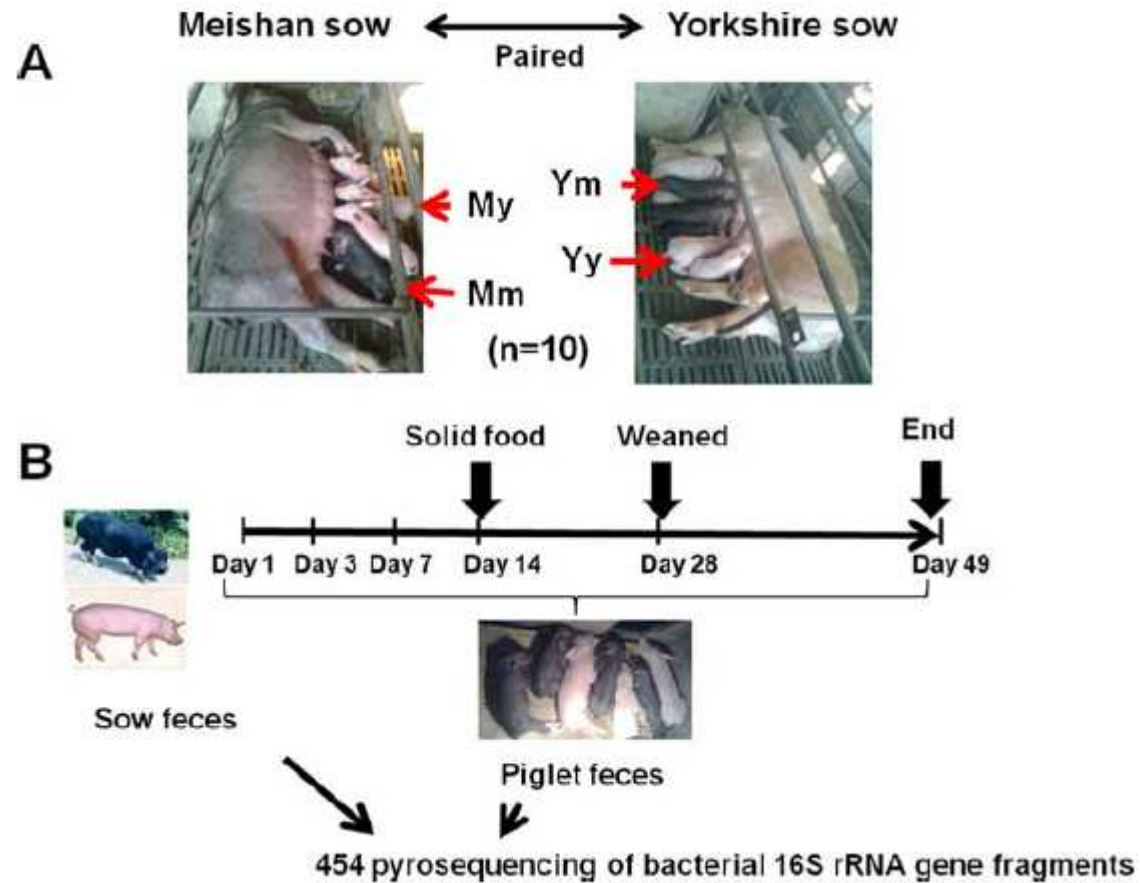
**Microbiota from piglets is influenced by the sow's microbiota and by the environment**

Jansman et al., 2012

# Factors affecting piglet's microbiota



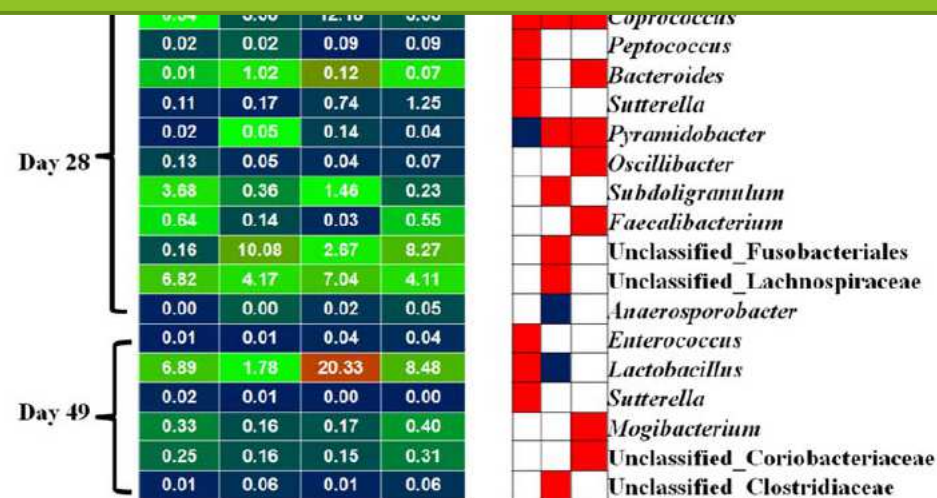
# Study on the effect of breed or nursing mother



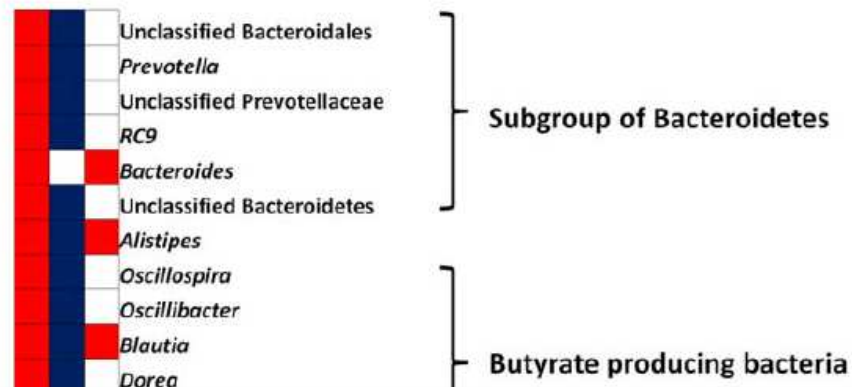
# Effects of breed and nursing mother



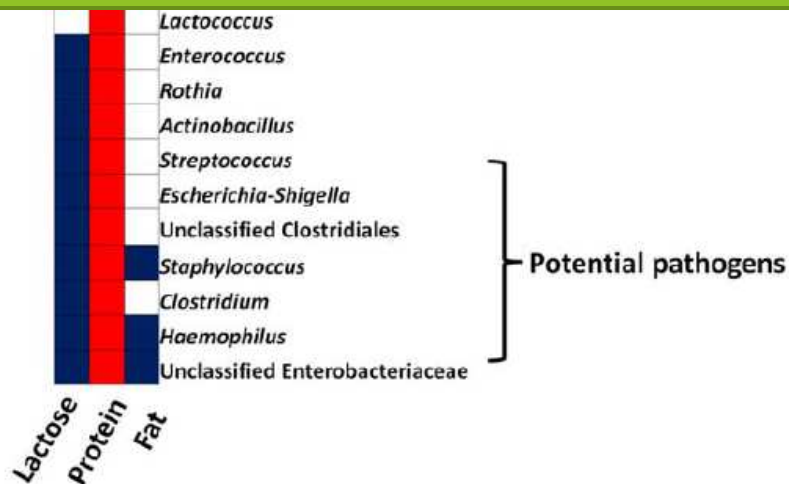
Effect of nursing mother and the breed were evident through the suckling period



# Impact of milk composition

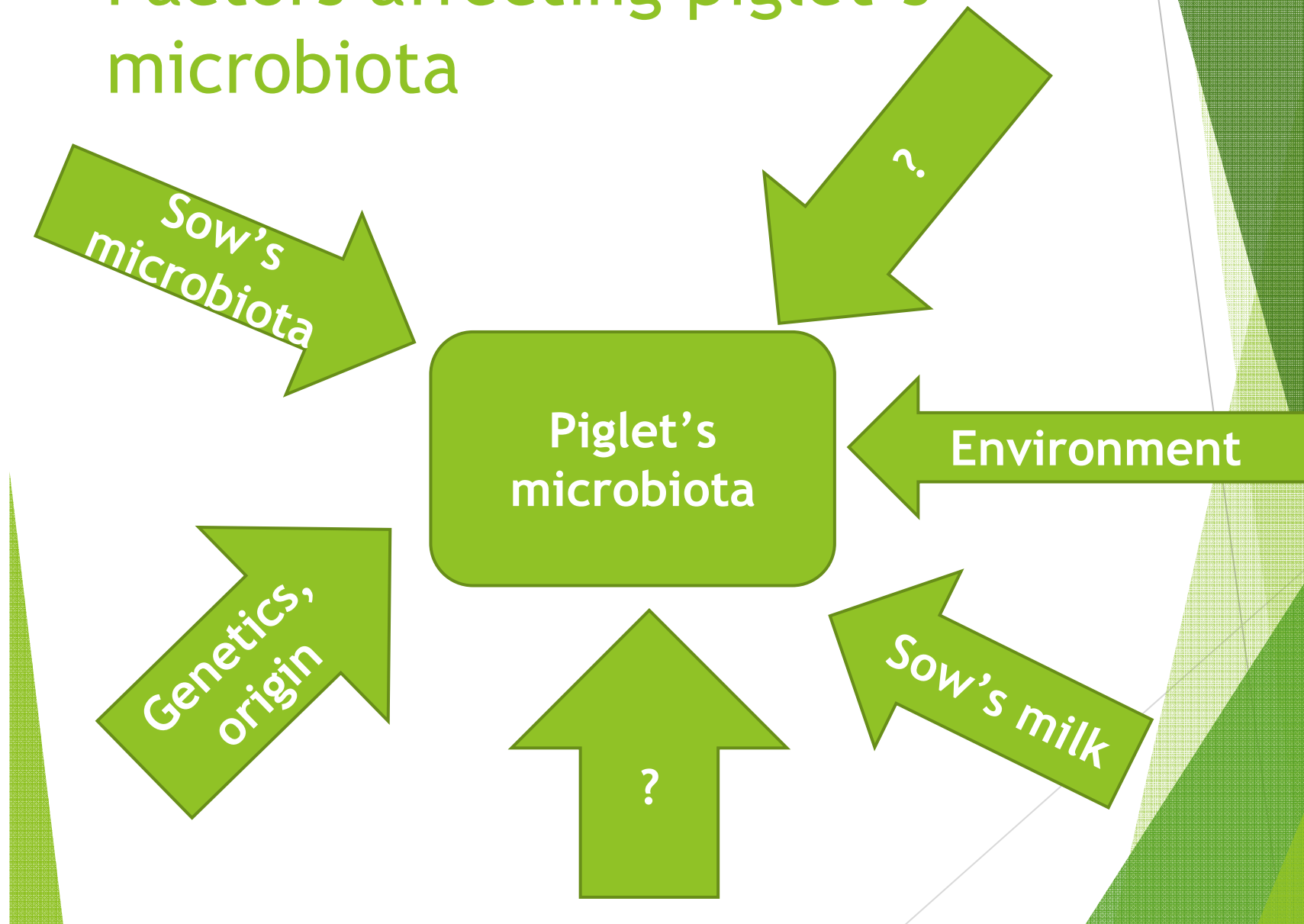


Milk lactose, protein and fat all significantly impacted the bacterial profile of piglets



Red: positive correlation  
Blue: negative correlation

# Factors affecting piglet's microbiota

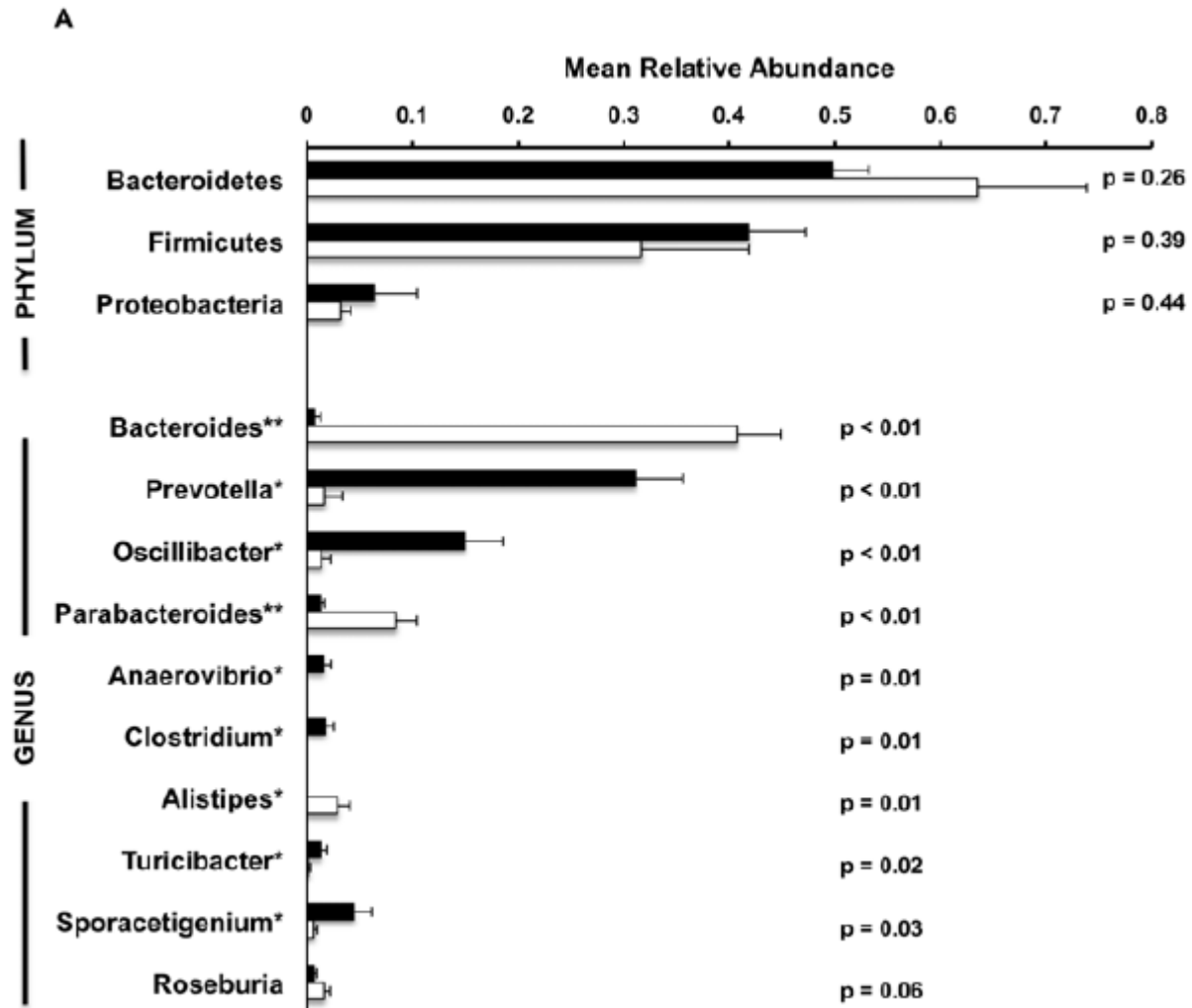


# Mother-fed vs formula milk

D21

Cecal content

Black: mother fed  
White: formula fed

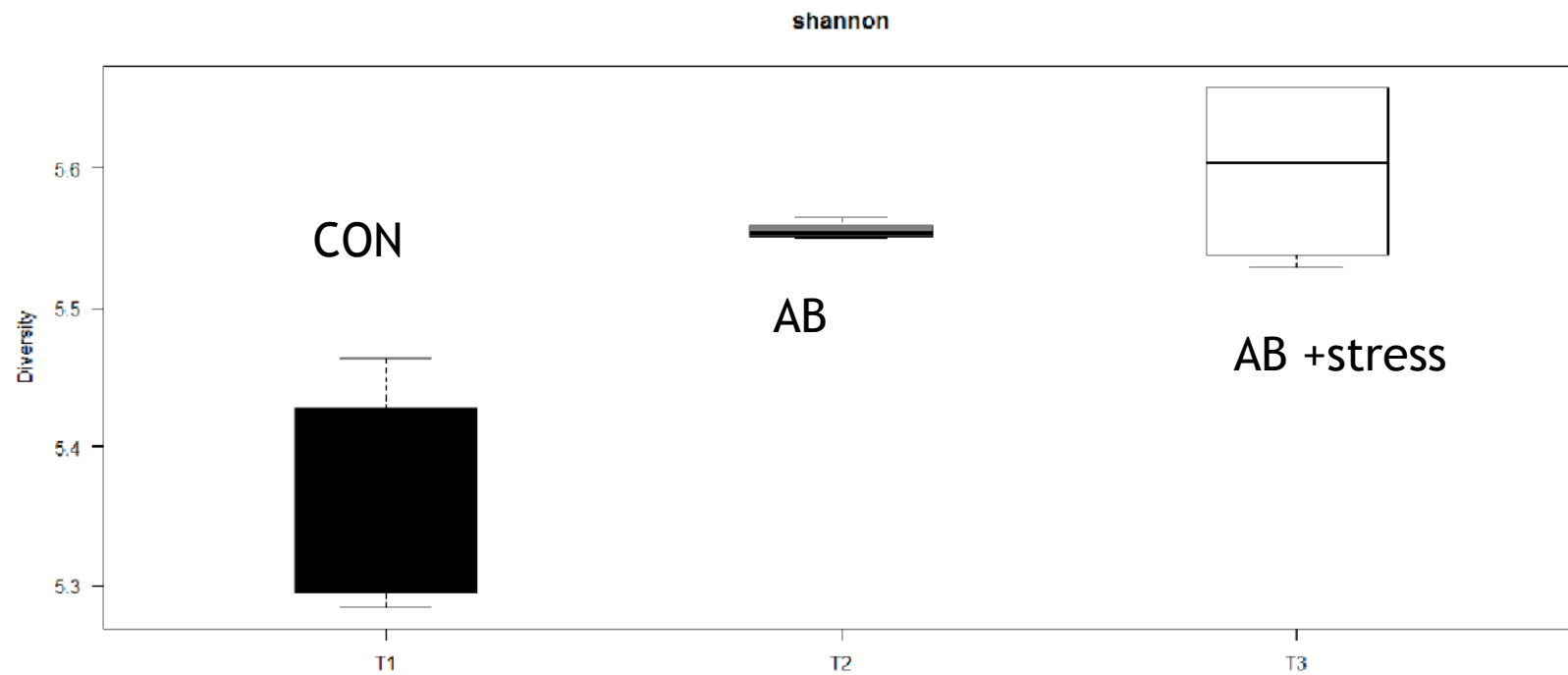


Poroyko et al., 2011

# Effects of antibiotic or stress-treatment on d4

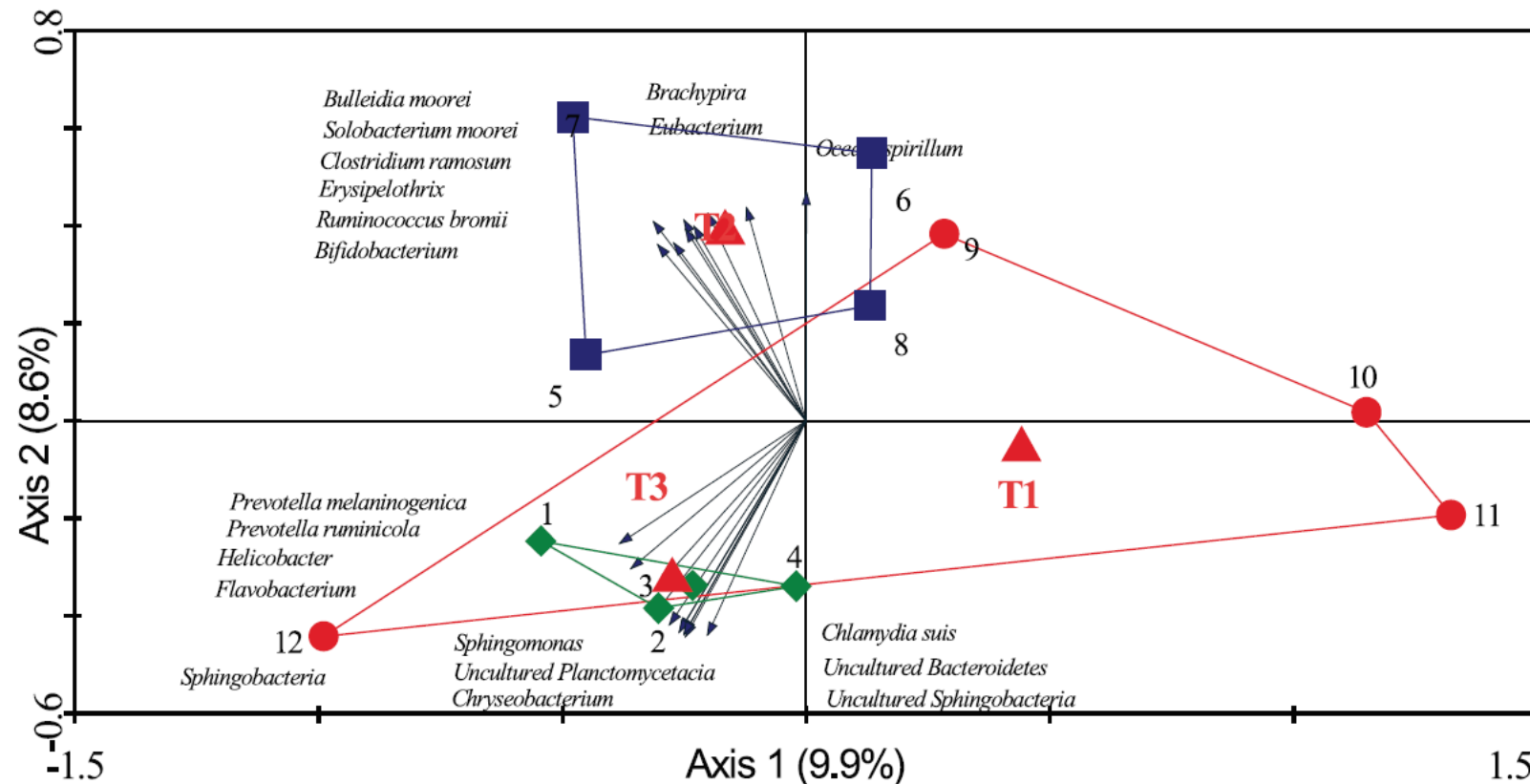
D8

Jejunal content



**Figure 2. Diversity in microbiota in the three treatment groups.** The Shannon index (y-axis) was calculated for all three treatments (T1, T2, and T3) (x-axis).  
doi:10.1371/journal.pone.0100040.g002

# Impact on microbiota composition



**Figure 1. Triplot for RDA analysis of jejunal microbiota composition.** Nominal environmental variables T1, T2 and T3 are represented by red triangles (▲). Samples are grouped by treatment: T1 (red; ○), T2 (blue; □) and T3 (green; ◇), each symbol represents a pool of four pigs, and numbers represent pool identity number. Microbial groups contributing at least 60% to the explanatory axes are represented as vectors. Both axes together explain 18.5% of the total variance in the dataset.  
doi:10.1371/journal.pone.0100040.g001

# Effects on transcriptome

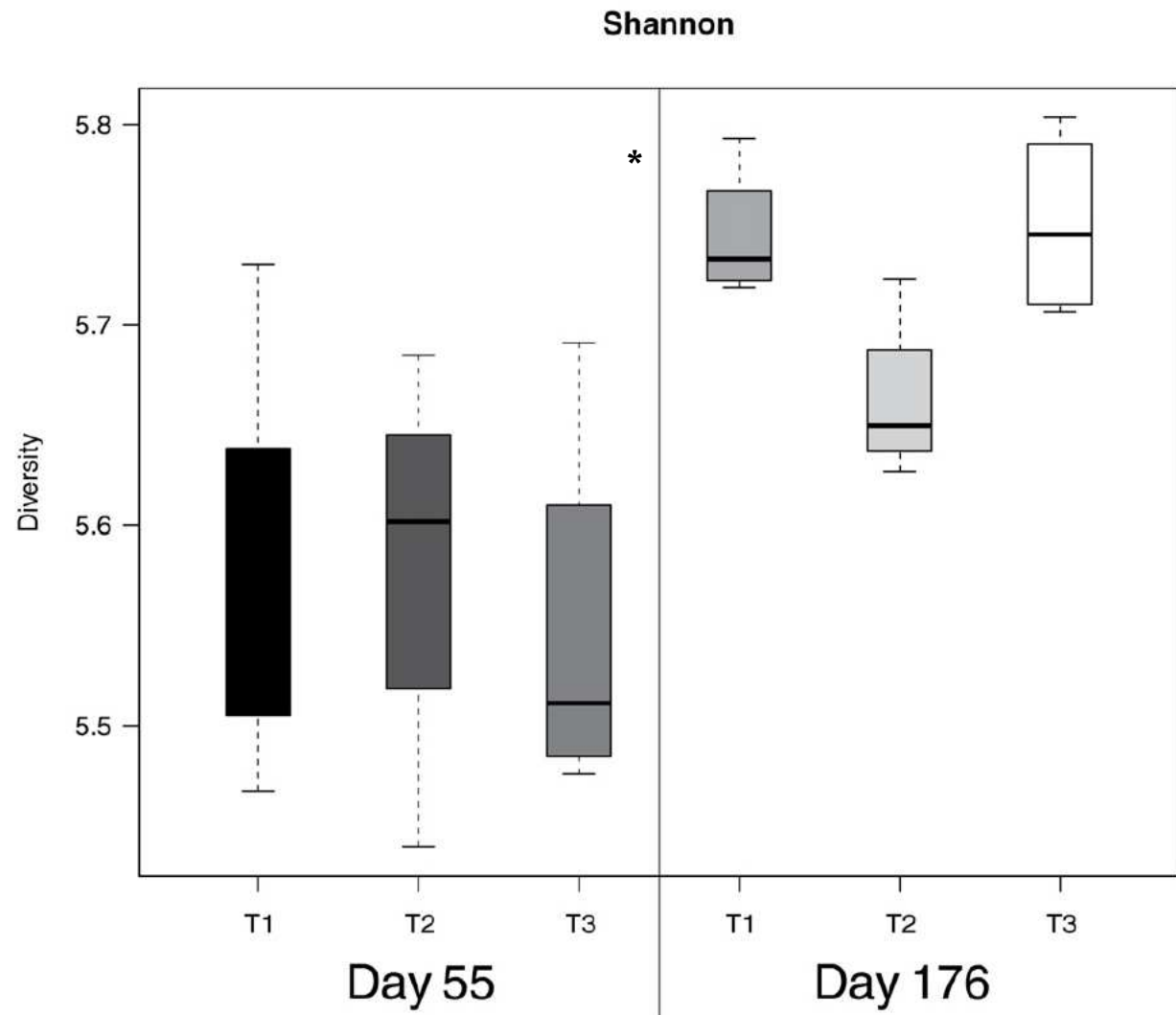
- Differences in immune programming in early-life
- Antibiotic treatment versus control

**Table 2.** Functional analysis of genes differentially expressed between treatment 2 versus 1.

DOWN		
JEJUNUM	#	Name
	1	chemotaxis
	2	cytokine activity
	3	chemokine activity
	4	regulation of secretion/immune effector process
	5	cell migration/motion (leukocyte)
ILEUM	#	Name
	1	cytokine activity
	2	chemotaxis
	3	second-messenger-mediated signaling (cAMP)
	4	chemokine activity
	5	response to bacterium/regulation of systemic process

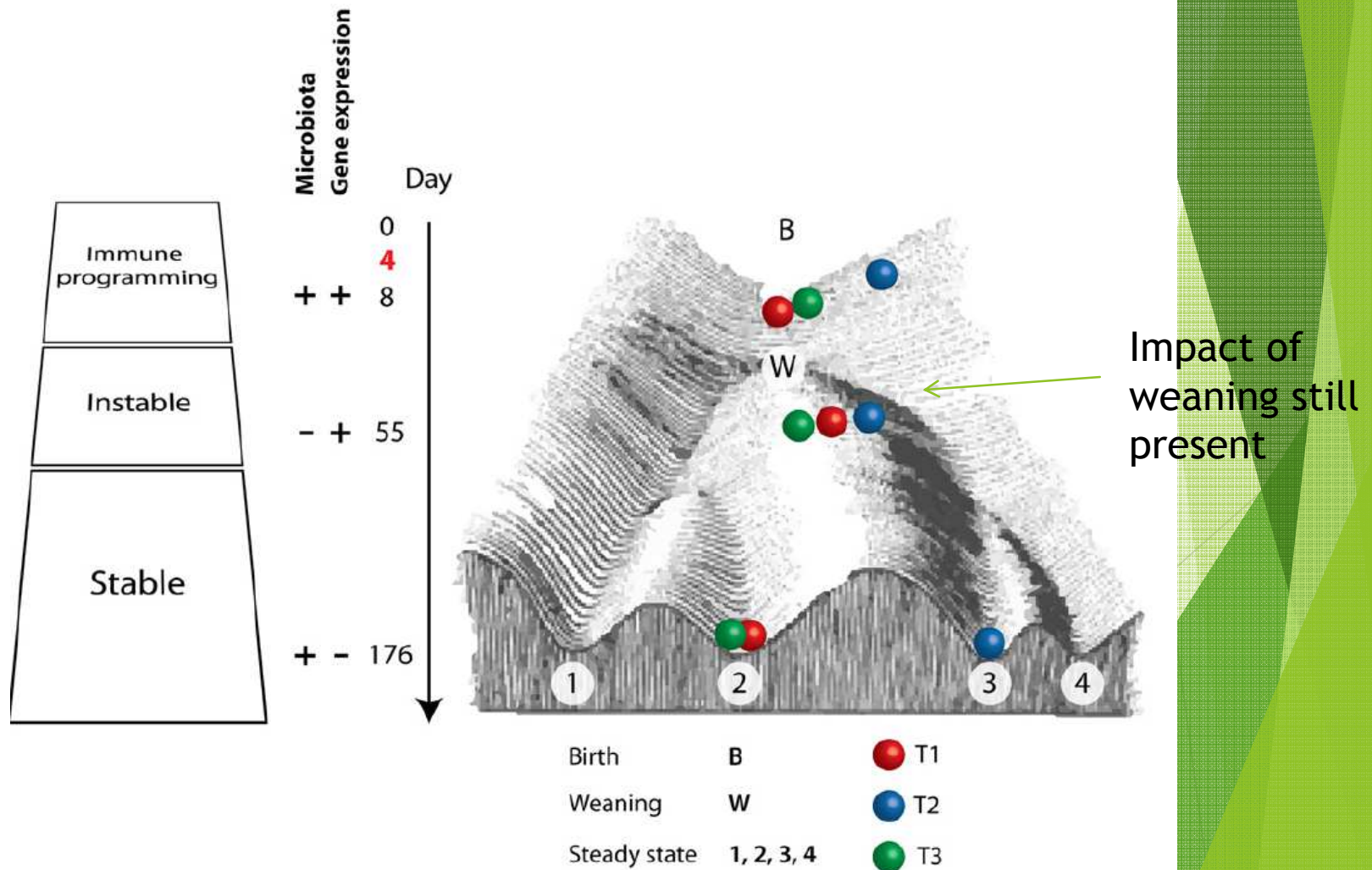
# Long-lasting effects on microbiota

Jejunal content

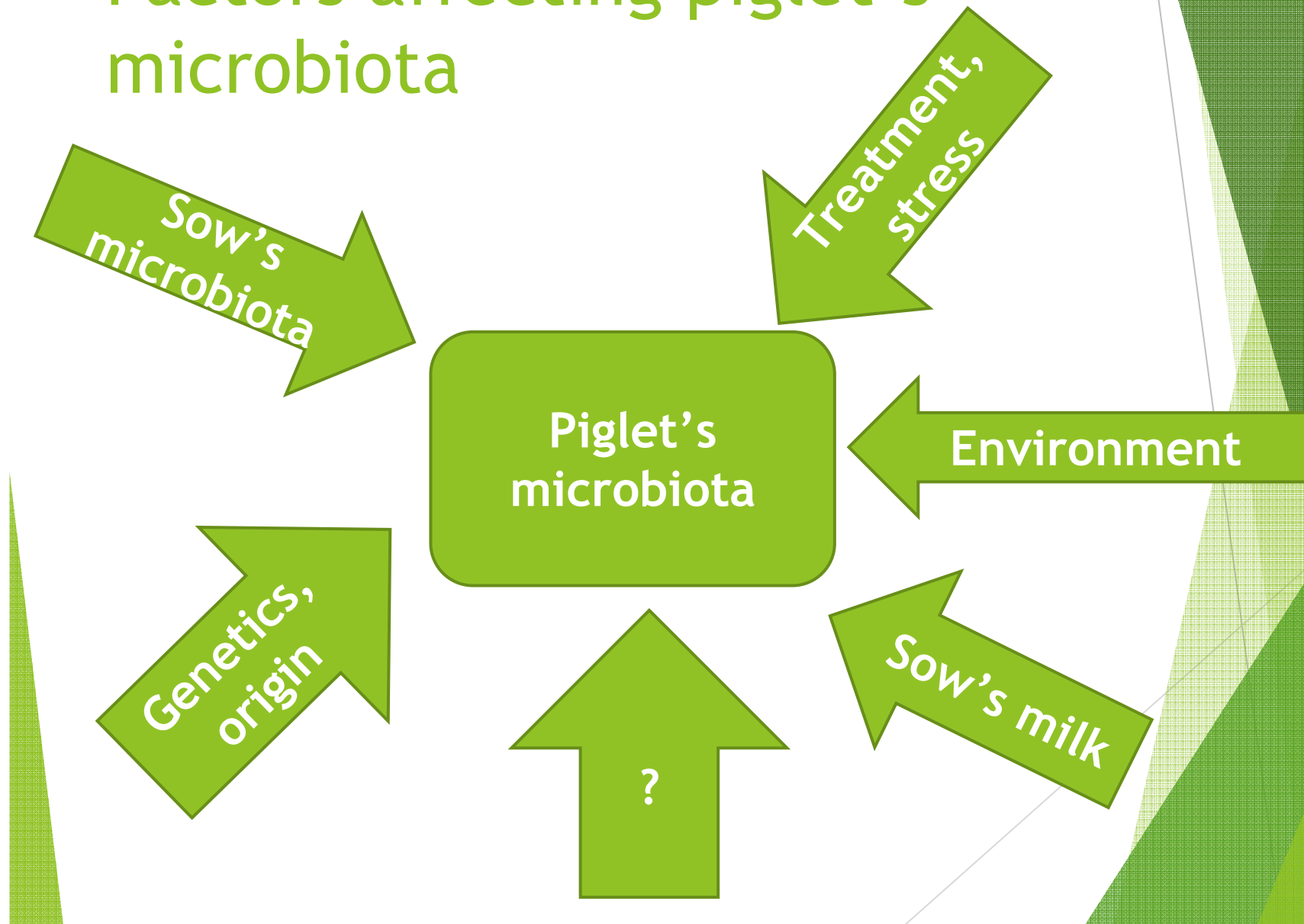


Schokker et al., 2015

# Towards a differential gut homeostasis

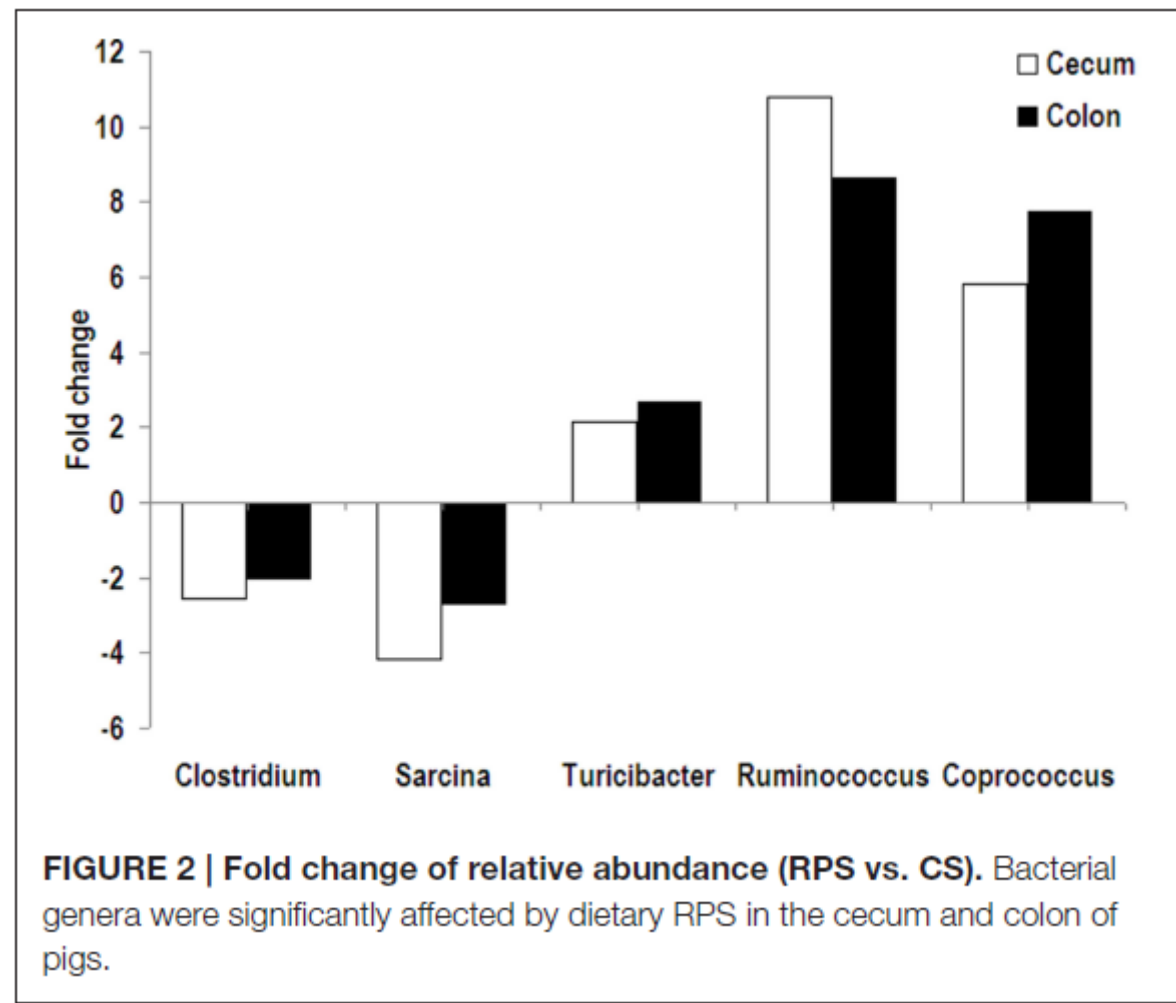


# Factors affecting piglet's microbiota



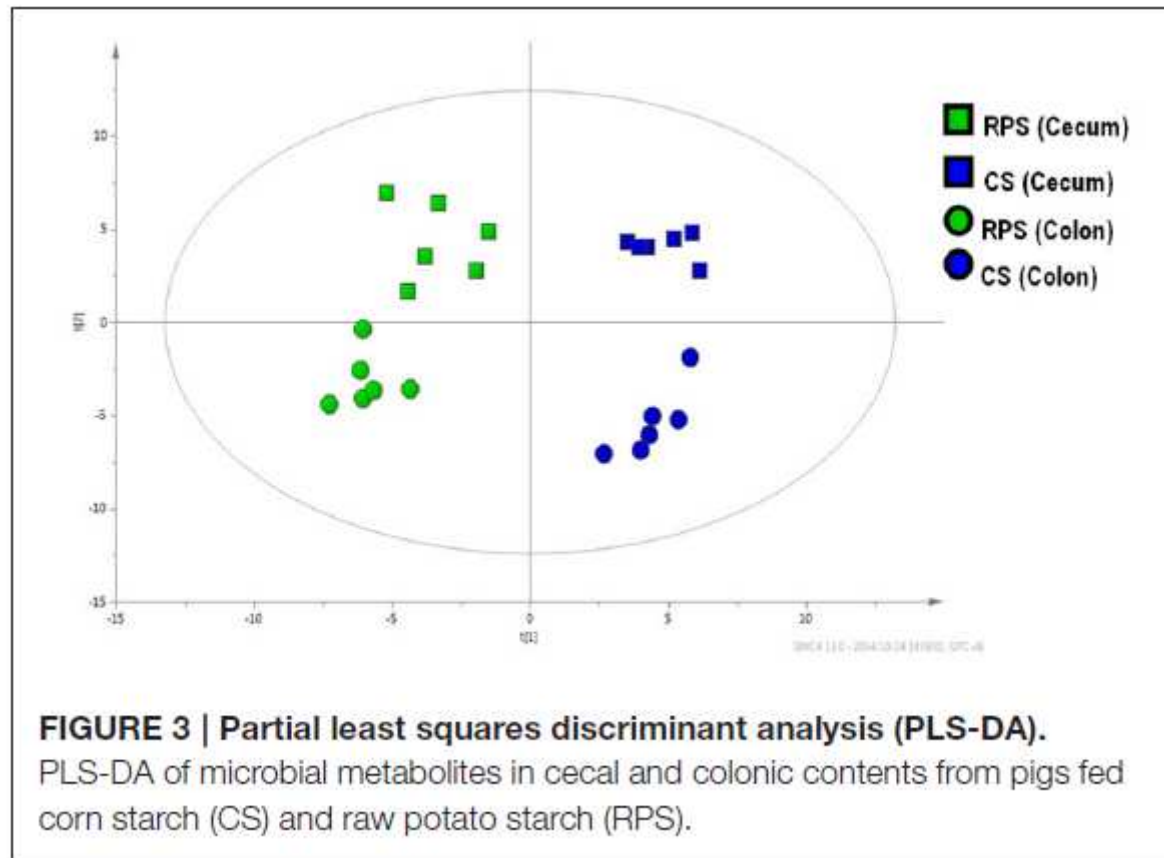
# Corn starch vs Raw potato starch

From d70-d170

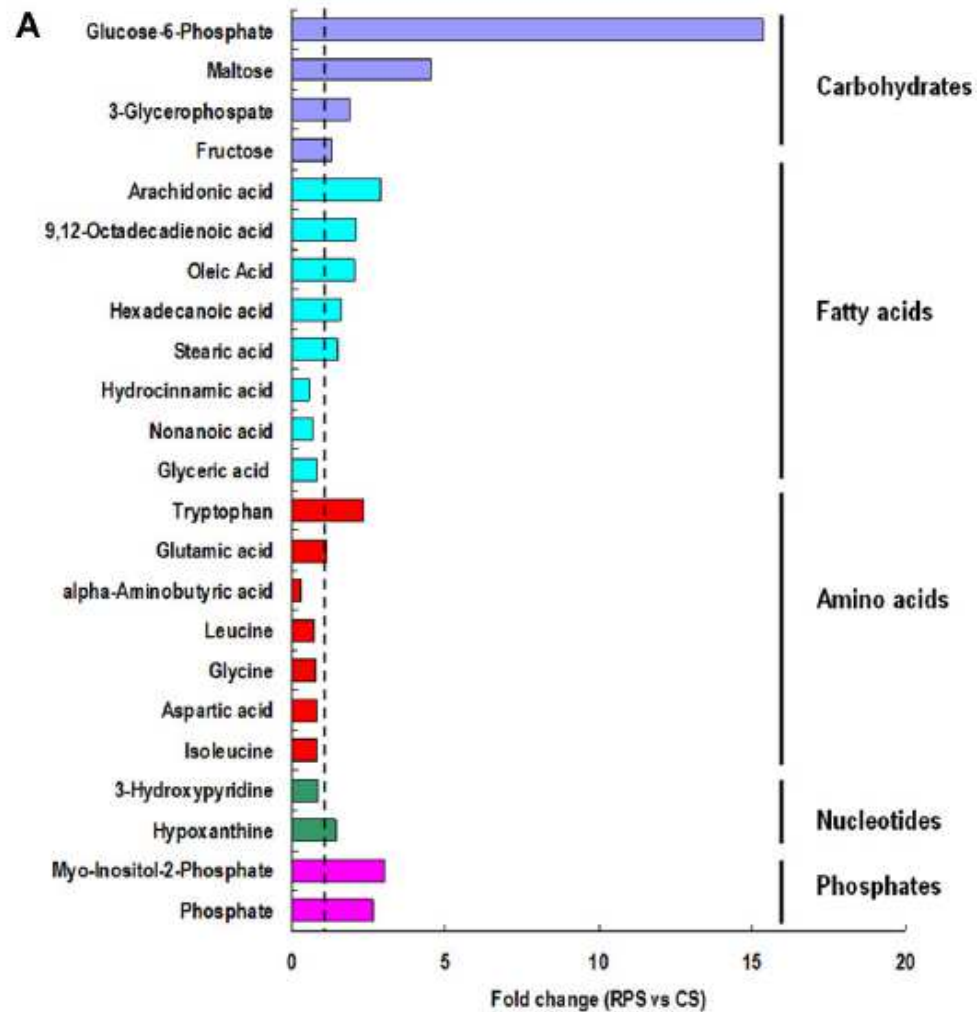


Sun et al., 2016

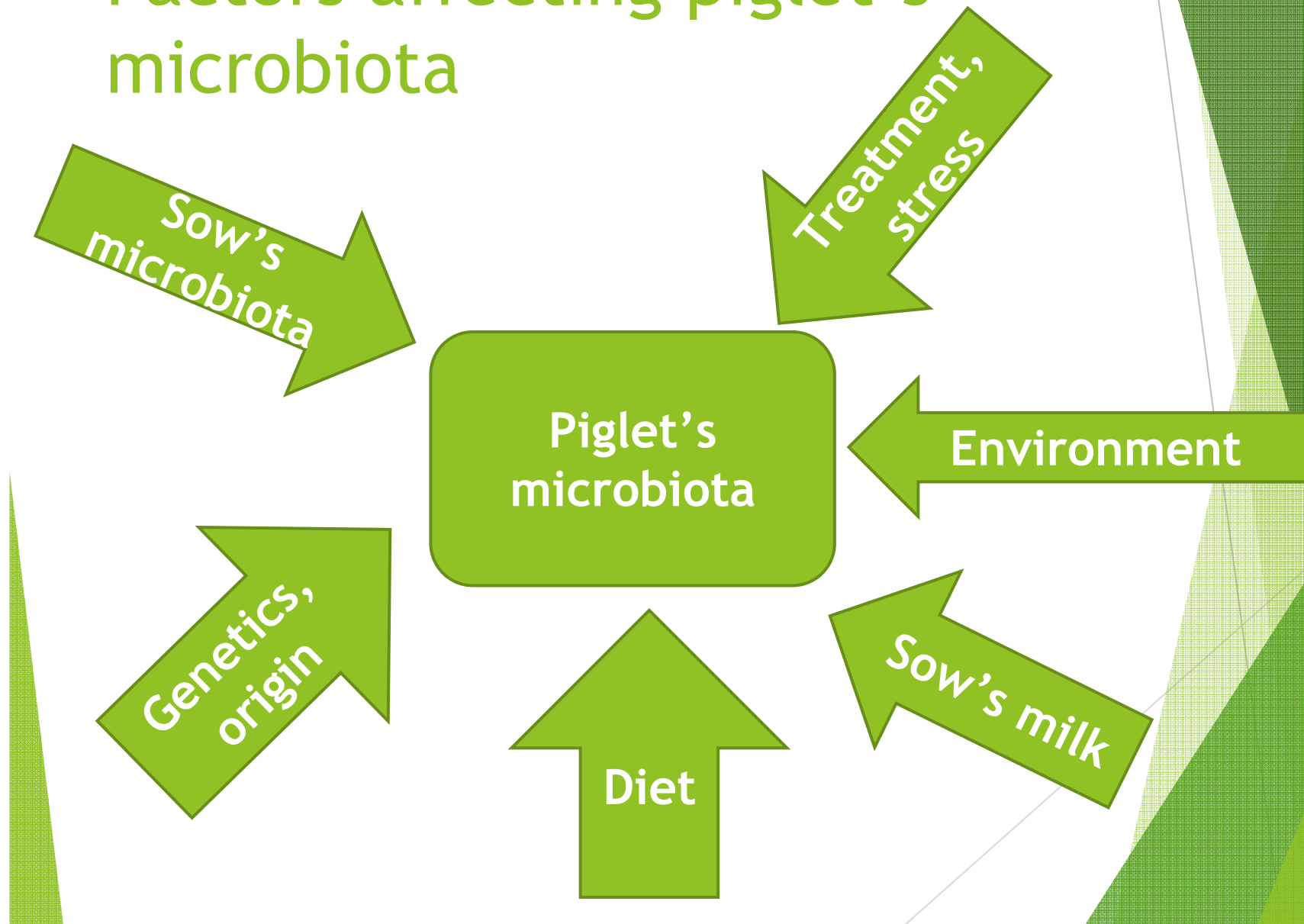
# Clustering according to intestinal segment and dietary treatment




# Resulting in different metabolites produced



# Factors affecting piglet's microbiota



The slide features two large, abstract green geometric shapes. On the left, a thin, elongated triangle points downwards. On the right, a more complex shape composed of several overlapping triangles and polygons is oriented vertically. The text is centered between these two shapes.

**Adding fermentable feed  
ingredients to pigs:  
changes or stability to the  
intestinal microbiota**

# AIM

- ▶ Reduce the risk of infections at weaning

# STRATEGY

- ▶ Through action on microbiota
- ▶ Indirectly: Via sow
- ▶ Directly: Interventions on piglets



The slide features abstract green geometric shapes. On the left, a thin, elongated green triangle points downwards. On the right, a larger, more complex green shape with multiple overlapping layers and a fine grid texture is positioned. The main title is centered in a large, bold, green sans-serif font.

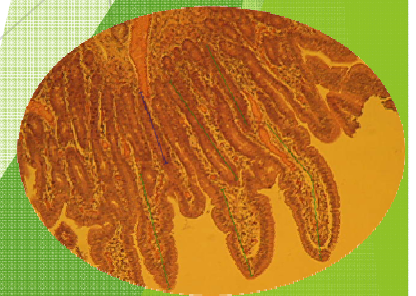
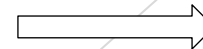
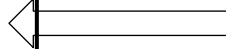
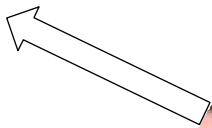
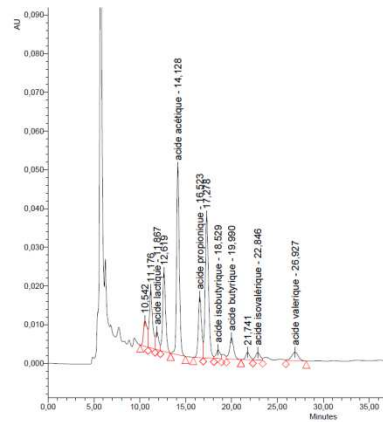
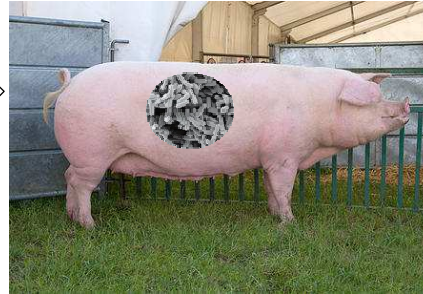
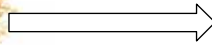
# Effects by the maternal diet

Inulin

Wheat bran

Resistant starch

# Hypotheses



# Inulin in the diet of gestating and lactating sows

- ▶ 3% inulin diet from 14d a.p.;
- ▶ change from gestation to lactation diet on 1d p.p.

**Table 1 Microbial cell counts (log<sub>10</sub>/g wet weight) in the faeces of sows fed a diet without (C) or with inulin (I)**

	Day 4 a.p.		Day 1 p.p.		Day 5 p.p.		P-value		
	C (n <sup>1</sup> = 10)	I (n = 10)	C (n = 10)	I (n = 9)	C (n = 10)	I (n = 10)	Diet	Time	Diet*time
Eubacteria	10.6 ± 0.51	10.8 ± 0.85	9.86 ± 0.10	9.75 ± 0.21	10.8 ± 0.15	10.8 ± 0.18	0.847	<b>0.005</b>	0.718
Enterobacteria	7.21 ± 1.01	6.66 ± 0.90	7.16 ± 0.98	7.77 ± 0.75	7.42 ± 0.68	7.54 ± 0.99	0.802	0.103	<b>0.017</b>
Enterococci	6.42 ± 0.45	6.83 ± 0.59	5.96 ± 0.31	7.00 ± 0.91	6.62 ± 0.48	6.98 ± 0.61	<b>0.014</b>	0.109	<b>0.028</b>
Bifidobacteria	7.92 ± 1.05	8.10 ± 1.91	7.24 ± 0.51	7.60 ± 1.08	8.05 ± 0.87	7.92 ± 1.51	0.976	0.052	0.357
Lactobacilli	9.39 ± 0.62	9.32 ± 0.77	8.89 ± 0.67	7.73 ± 1.42	8.98 ± 0.57	8.60 ± 0.89	0.109	<b>&lt;0.001</b>	0.051
<i>L. reuteri</i>	8.37 ± 0.86	8.25 ± 0.64	7.36 ± 1.12	6.83 ± 0.66	7.93 ± 0.73	7.57 ± 0.47	0.305	<b>&lt;0.001</b>	0.244
<i>L. amylovorus</i>	9.39 ± 0.40	9.32 ± 0.35	8.50 ± 0.96	8.22 ± 0.63	9.12 ± 0.61	9.03 ± 0.32	0.502	<b>&lt;0.001</b>	0.606
<i>L. johnsonii</i>	6.16 ± 1.10	5.81 ± 0.94	4.97 ± 0.79	5.20 ± 0.52	5.46 ± 0.42	5.25 ± 0.48	0.872	<b>0.002</b>	0.411
<i>L. mucosae</i>	8.22 ± 0.83	7.83 ± 0.41	7.08 ± 0.89	6.72 ± 0.47	7.56 ± 0.61	7.17 ± 0.96	0.088	<b>&lt;0.001</b>	0.898
<i>C. leptum</i>	10.5 ± 0.24	10.1 ± 0.55	9.50 ± 0.97	9.84 ± 0.62	9.89 ± 0.73	9.91 ± 1.11	0.987	<b>0.010</b>	0.266
<i>C. coccoides</i>	10.7 ± 0.53	10.8 ± 0.24	9.69 ± 1.05	9.79 ± 0.77	10.6 ± 0.41	10.5 ± 0.93	0.891	<b>0.002</b>	0.762
BPP	9.66 ± 0.53	9.69 ± 0.54	9.40 ± 0.25	9.31 ± 0.37	9.83 ± 0.52	9.55 ± 0.28	0.152	0.052	0.557

# Effects on the microbial metabolites of sows

## ► Change in microbial metabolites

**Table 2 Microbial metabolites and pH in the faeces of sows fed a diet without (C) or with inulin (I)**

	Day 4 a.p.		Day 1 p.p.		Day 5 p.p.		P-value		
	C (n = 11)	I (n = 10)	C (n = 11)	I (n = 9)	C (n = 11)	I (n = 10)	Diet	Time	Diet*time
pH	6.79 ± 0.23	6.63 ± 0.34	6.99 ± 0.34	6.59 ± 0.36	6.68 ± 0.12	6.63 ± 0.23	<b>0.007</b>	0.323	0.167
L-lactate (mmol/kg)	0.73 ± 0.55	0.70 ± 0.30	0.43 ± 0.28	0.28 ± 0.22	0.35 ± 0.18	0.24 ± 0.19	0.185	<b>0.002</b>	0.909
D-lactate (mmol/kg)	0.62 ± 0.64	0.54 ± 0.26	0.20 ± 0.20	0.14 ± 0.12	0.16 ± 0.12	0.11 ± 0.11	0.451	<b>0.002</b>	0.947
Ammonia (mmol/kg)	32.6 ± 19.5	47.0 ± 29.1	13.7 ± 9.64	20.2 ± 10.4	21.7 ± 10.2	27.3 ± 12.9	0.070	<b>0.001</b>	0.586
SCFA (mmol/l)	152 ± 44.4	155 ± 30.1	100 ± 31.4	123 ± 31.5	138 ± 38.3	158 ± 17.5	0.164	<b>&lt;0.001</b>	0.602
Acetic acid (mol. %)	54.4 ± 3.61	53.7 ± 3.78	60.6 ± 4.11	56.8 ± 4.19	60.9 ± 4.35	59.1 ± 3.16	0.081	<b>&lt;0.001</b>	0.388
Propionic acid (mol. %)	23.0 ± 1.08	22.6 ± 1.33	19.3 ± 2.03	20.9 ± 1.11	19.4 ± 1.40	19.9 ± 1.69	0.153	<b>&lt;0.001</b>	0.053
i-butyric acid (mol. %)	2.87 ± 0.56	2.79 ± 0.33	2.87 ± 0.40	2.61 ± 0.55	2.52 ± 0.27	2.51 ± 0.32	0.337	<b>0.037</b>	0.641
n-butyric acid (mol.%)	13.0 ± 2.85	13.9 ± 2.62	10.4 ± 4.24	12.8 ± 3.36	11.1 ± 3.54	12.4 ± 1.71	0.152	<b>0.025</b>	0.711
i-valeric acid (mol. %)	4.05 ± 0.91	3.92 ± 0.51	4.06 ± 0.62	3.88 ± 0.86	3.62 ± 0.43	3.56 ± 0.53	0.538	0.068	0.950
n-valeric acid (mol. %)	2.66 ± 0.50	3.09 ± 0.70	2.77 ± 0.51	3.02 ± 0.63	2.45 ± 0.51	2.61 ± 0.45	0.147	<b>0.012</b>	0.467

Abbreviations: a.p.: ante partum; p.p.: post partum; SCFA short chain fatty acids.

Boldface P-values indicate significant effects ( $P \leq 0.05$ ).

# Altering the microbiota and metabolites of the piglets

**Table 3 Microbial cell counts (log<sub>10</sub>/g wet weight) in the digesta of the stomach and caecum of suckling piglets, where the mother sows received either a diet without (C) or with inulin (I)**

	Stomach		Caecum	
	C (n <sup>1</sup> = 7)	I (n = 8)	C (n <sup>2</sup> = 8)	I (n = 8)
Eubacteria	10.3 ± 0.26 <sup>a</sup>	11.0 ± 0.25 <sup>b</sup>	11.7 ± 0.37	11.7 ± 0.33
Enterobacteria	4.56 ± 0.87	3.45 ± 0.50 <sup>b</sup>	9.02 ± 0.66	8.58 ± 1.20
Enterococci	7.24 ± 0.45	7.21 ± 0.74	6.76 ± 0.49 <sup>a</sup>	7.57 ± 0.46 <sup>b</sup>
Bifidobacteria	3.89 ± 0.70	5.04 ± 1.83	5.16 ± 0.91	5.57 ± 1.29
Lactobacilli	9.31 ± 0.88	9.06 ± 0.97	10.0 ± 1.23	9.94 ± 1.49
<i>L. reuteri</i>	8.23 ± 0.63	7.81 ± 0.84	8.40 ± 0.50	8.13 ± 0.62
<i>L. amylovorus</i>	9.28 ± 0.32 <sup>a</sup>	8.17 ± 0.66 <sup>b</sup>	9.26 ± 0.69	9.28 ± 0.79
<i>L. johnsonii</i>	7.05 ± 0.69	6.45 ± 0.93	6.76 ± 0.79	6.22 ± 1.09
<i>L. mucosae</i>	7.61 ± 0.82	6.78 ± 1.42	7.95 ± 0.89	7.97 ± 1.31
<i>C. leptum</i>	6.70 ± 0.31	6.40 ± 0.89	10.0 ± 0.78 <sup>a</sup>	10.8 ± 0.15 <sup>b</sup>
<i>C. coccoides</i>	7.02 ± 0.93	6.66 ± 1.02	10.3 ± 0.78	10.7 ± 0.43
BPP	6.14 ± 1.12	5.43 ± 1.72	9.87 ± 0.74	10.3 ± 0.32

On day 10

# Altering the microbiota and metabolites of the piglets

**Table 4 Microbial metabolites and pH in the digesta of suckling piglets, where the mother sows received a diet without (C) or with inulin (I)**

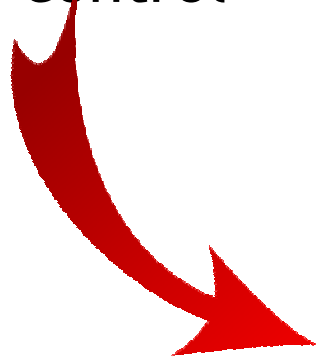
	Stomach		Small intestine		Caecum		Rectum	
	C (n <sup>1</sup> = 8)	I (n = 8)	C (n <sup>2</sup> = 8)	I (n = 8)	C (n <sup>3</sup> = 8)	I (n = 8)	C (n <sup>4</sup> = 8)	I (n = 8)
pH	3.37 ± 0.11	3.74 ± 0.75	6.78 ± 0.13	6.71 ± 0.64	6.97 ± 0.15	6.78 ± 0.34	6.15 ± 0.48	6.26 ± 0.62
Acetic acid (mol.%)	3.33 ± 1.75	3.52 ± 1.93	4.40 ± 2.08	3.28 ± 3.41	36.5 ± 9.78	33.9 ± 11.4	13.9 ± 6.75	9.68 ± 4.01
Propionic acid (mol.%)	0.12 ± 0.10	0.12 ± 0.20	0.31 ± 0.35	0.19 ± 0.30	10.8 ± 2.59	8.93 ± 3.26	3.83 ± 2.33	2.80 ± 2.36
i-butyric acid (mol.%)	0.01 ± 0.00 <sup>a</sup>	0.01 ± 0.02 <sup>b</sup>	0.02 ± 0.02	*	1.24 ± 0.49	0.90 ± 0.40	0.69 ± 0.48	0.47 ± 0.36
n-butyric acid (mol.%)	0.18 ± 0.04 <sup>a</sup>	0.06 ± 0.07 <sup>b</sup>	0.12 ± 0.11	0.11 ± 0.11	3.74 ± 1.68	3.10 ± 1.50	2.07 ± 1.61	1.21 ± 1.12
i-valeric acid (mol.%)	0.27 ± 0.52 <sup>a</sup>	0.03 ± 0.03 <sup>b</sup>	0.04 ± 0.03	0.03 ± 0.03	1.62 ± 0.56	1.13 ± 0.41	1.37 ± 0.87	0.86 ± 0.67
n-valeric acid (mol.%)	0.02 ± 0.01	0.00 ± 0.00	0.01 ± 0.01	0.01 ± 0.00	1.70 ± 0.72	1.16 ± 0.50	0.91 ± 0.57	0.47 ± 0.24

Inulin seems to have the potential to influence the gastrointestinal microbiota of suckling piglets through the diet of their mother

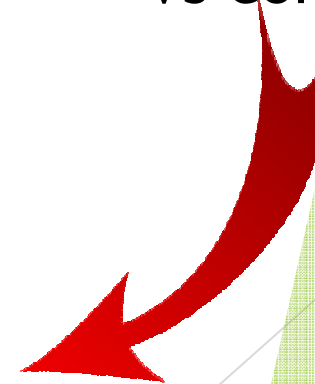
# Two animal experiments



Wheat bran (insoluble  
fibers)  
Vs. control



Pea starch (resistant starch)  
Vs control

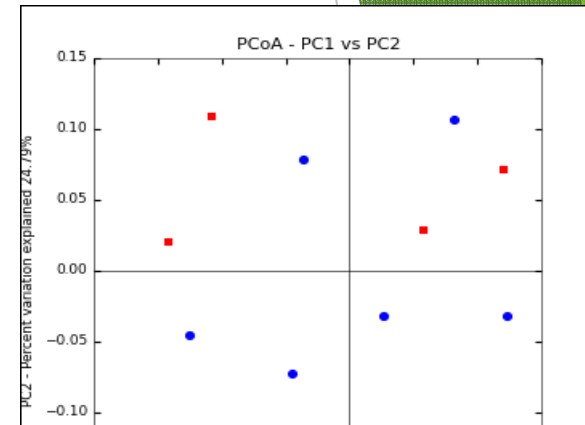
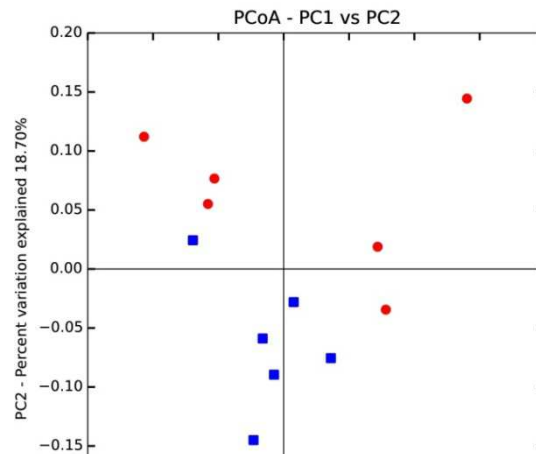


Gestation  
Lactation

## Gestation

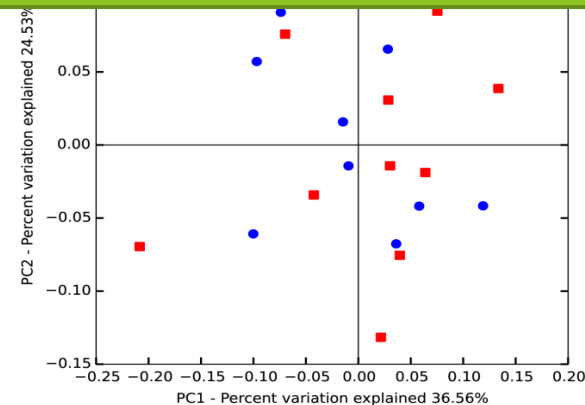
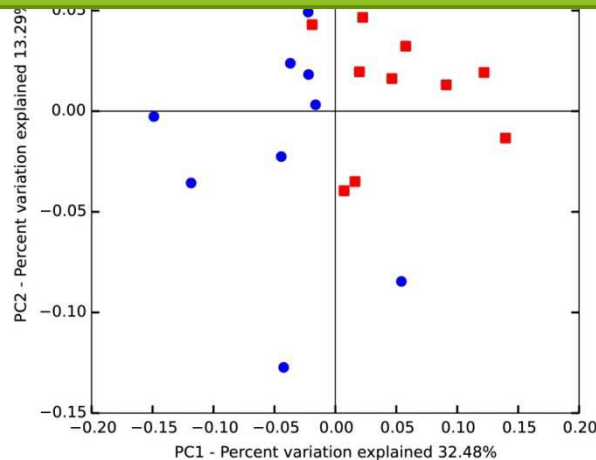
## Lactation

Wheat  
bran



Clustering per treatment during gestation, but not during lactation


Pea  
starch



Gestation Wheat bran				
Genus	CON	WB	P-values	FDR
<b>Bacteroidetes</b>				
Parabacteroides	0.36	0.14	<0.001	0.02
Unclassified_Bacteroidales	6.13	2.25	<0.001	0.02
Bacteroides	0.22	0.04	<0.005	NS
CF231	1.22	0.57	0.01	NS
Unclassified_RF16	2.38	0.79	0.03	NS
Prevotella	15.5	19.0	NS	NS

13 genera differed in relative abundance between the CON and WB groups

Unclassified_Erysipelotrichaceae OTU1	0.02	0.06	0.01	NS
Anaerovibrio	0.20	0.53	0.03	NS
Turicibacter	0.13	0.07	0.03	NS
Oscillospira	2.69	1.76	0.03	NS
Unclassified_Erysipelotrichaceae OTU2	0.08	0.03	0.06	NS
Unclassified_Mogibacteriaceae	0.75	0.44	0.07	NS
<b>Proteobacteria</b>				
Unclassified_Enterobacteriaceae	0.04	0.01	0.01	NS
Ruminobacter	0.02	0.03	NS	NS

Genus	Gestation Pea starch			
	DS	RS	P	FDR
<b>Actinobacteria</b>				
Bifidobacterium	0.92	1.36	0.02	NS
Bacteroidetes				
Unclassified_RF16	1.53	0.80	0.01	NS
<b>Firmicutes</b> 				
Unclassified_Ruminococcaceae	17.75	20.68	0.02	NS

Differences at the phylum level and the genus level during gestation

Parabacter	0.33	1.18	0.003	NS
Sharpea	0.21	0.79	0.03	NS

Dietary interventions on sows affect their fecal microbiota

ae	0.12	0.20	0.02	NS
Spirochaetes				
Treponema	4.20	3.10	0.01	NS
Sphaerochaeta	1.05	0.50	<0.005	NS

# WB exp, milk composition

Period	Treatment	Protein (%)	Fat (%)	Lactose (%)	IgA (mg/ml)	IgG (mg/ml)	IgM (mg/ml)
Colostrum	CON	19.1	6.33	2.60	13.9	63.5	4.84
	WB	19.0	6.45	2.62	13.4	68.5	4.22
	SEM	0.30	0.14	0.03	0.81	3.46	0.32
Milk W1 <sup>1</sup>	CON	6.17	9.70	4.66	2.10	0.40	1.08
	WB	5.96	9.48	4.74	2.53	0.41	1.15
	SEM	0.09	0.51	0.05	0.16	0.05	0.11
Milk W2	CON	6.03	9.84	4.81	2.43	0.30	1.10
	WB	5.62	9.62	4.89	2.66	0.25	1.08
	SEM	0.10	0.28	0.02	0.17	0.02	0.07
Milk W3	CON	6.22	9.72	4.82	3.41	0.20	0.97
	WB	5.86	8.85	4.93	3.57	0.16	1.01
	SEM	0.08	0.38	0.03	0.23	0.02	0.08
treatment		0.14	0.46	0.03	0.88	0.47	0.58
P-values	time	<.0001	<.0001	<.0001	<.0001	<.0001	<.0001
treatment*time		0.75	0.62	0.88	0.81	0.78	0.35

<sup>1</sup>W1= first week after farrowing, W2= second week, W3= third week

# RS exp, milk composition

- ▶ Resistant starch:
  - ▶ Decreased protein concentration (all time points)
  - ▶ Increased lactose concentration in colostrum
  - ▶ Decreased lactose concentration on W3

**Dietary interventions on sows affect milk  
macronutrient composition**

# WB exp, microbiota of piglets

	CON (N=7)	WB (N=7)	P-value	FDR
Actinobacteria	0.71	0.57	NS	NS
<b>Collinsella</b>	0.29	0.08	<b>0.04</b>	NS
Bacteroidetes	32.3	28.4	NS	NS
<i>Butyricimonas</i>	0.15	0.02	0.07	NS
<i>Odoribacter</i>	0.25	0.02	0.07	NS
<i>Bacteroides</i>	6.72	2.21	NS	NS
Unclassified_Bacteroidales	3.27	5.61	NS	NS
<i>Prevotella</i>	12.3	11.8	NS	NS
Euryarchaeota	0.01	0.02	<b>0.05</b>	NS
<b>Methanobrevibacter</b>	0.01	0.02	<b>0.05</b>	NS
Firmicutes	56.0	63.2	NS	NS
<b>Unclassified_Clostridiaceae</b>	1.57	2.82	<b>&lt;0.001</b>	<b>0.04</b>
<b>Unclassified_Lachnospiraceae OTU2</b>	1.91	4.14	<b>0.04</b>	NS
<i>Ruminococcus</i>	1.74	0.85	0.07	NS
<i>Phascolarctobacterium</i>	2.35	3.68	0.07	NS
<i>Roseburia</i>	0.11	0.57	0.09	NS
<i>Lactobacillus</i>	14.8	13.1	NS	NS
Unclassified_Clostridiales	6.57	6.97	NS	NS
Unclassified_Ruminococcaceae	11.7	14.3	NS	NS

# RS exp, microbiota of piglets

- ▶ The maternal diet did not affect colonic microbiota composition at weaning

**Dietary interventions on sows has rather limited effects on piglet's microbiota**

The background features abstract green geometric shapes. On the left, a thin, elongated green triangle points downwards. On the right, a larger, more complex green shape with multiple overlapping layers and a fine grid texture is visible. The main text is centered in the white space between these shapes.

# Treatment during the lactation period

Inulin

# Inulin supplementation during the lactation period



12 litters with  
6 piglets per  
litter

4 litters-Control  
(water)

4 litters-20% inulin  
solution

4 litters-30% inulin  
solution

After  
weaning  
d-28

Same diet  
(no inulin)  
for another  
3 weeks



\* Inulin was obtained by  
oral ingestion :  
1<sup>st</sup> week: 2.5ml per day  
2<sup>nd</sup> week: 5ml per day  
3<sup>rd</sup> week: 7.5ml per day  
4<sup>th</sup> week: 10ml per day

# Temporary effects on microbiota

Effects on microbiota did not remain in the early post-weaning period

									M	P
<i>Lactobacillus spp.</i>	0.69	0.69	0.55	0.16	0.007	7.65	5.45	4.76	0.78	0.410
<i>Clostridium spp.</i>	0.73	0.53	1.20	0.13	0.088	0.19	0.15	0.13	0.04	0.883
<i>Escherichia spp.</i>	1.22 <sup>b</sup>	0.40 <sup>a</sup>	1.24 <sup>b</sup>	0.17	0.047	0.01	0.02	0.02	0.00	0.883
<i>Enterobacteria spp.</i>	1.25 <sup>ab</sup>	0.40 <sup>a</sup>	1.48 <sup>b</sup>	0.17	0.014	0.04	0.05	0.02	0.02	0.730

# Take home message

- ▶ Piglet's microbiota is affected by:
  - ▶ Breed, origin
  - ▶ Environment
  - ▶ Sow's microbiota
  - ▶ Sow's milk composition
  - ▶ (Medical) treatments and stress
  - ▶ Diet
- ▶ Effects on gut homeostasis are observed
- ▶ Dietary interventions with fermentable feed ingredients affect microbiota, but sometimes to a limited extend