Gut microbiome and incidence of foodborne pathogens are affected by diet in pasture-raised chickens

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The pasture-raised system

Conventional chicken operations



Pasture-raised chicken operations



The pasture-raised system





The pasture-raised system used in this study:



In this system, we tested 2 different supplements:

1) Soy-containing supplement (**SB**) (18.1% soybean)

2) Soy-free supplement (SF)

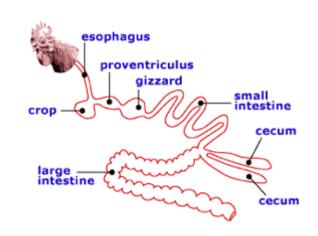
- 3 flocks of birds were fed SB
- 2 flocks of birds were fed SF



Multiple samples were collected using a farm-to-fork approach:









GIT from 1-day-old chicks

Feces from the pasture at 4, 7, and 12 weeks of age

Cecal contents at the slaughterhouse (12 weeks-old)

WCR at 12 weeks-old (day of processing)

WCR after frozen for 1 month (final product)

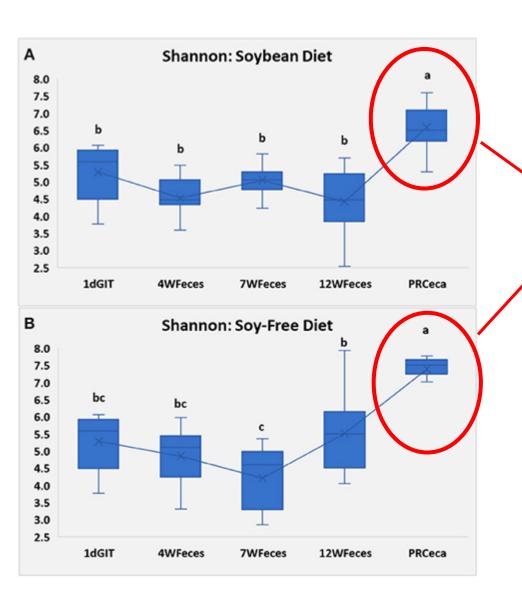
Sample processing

- DNA was extracted from all of those samples (GIT from 1-day-old chicks, feces, ceca, WCRs)
- Samples were then submitted for sequencing of the 16S rRNA gene (hypervariable V4 domain)

 Sequencing results were processed and analyzed using the QIIME pipeline (v1.9.1)

RESULTS

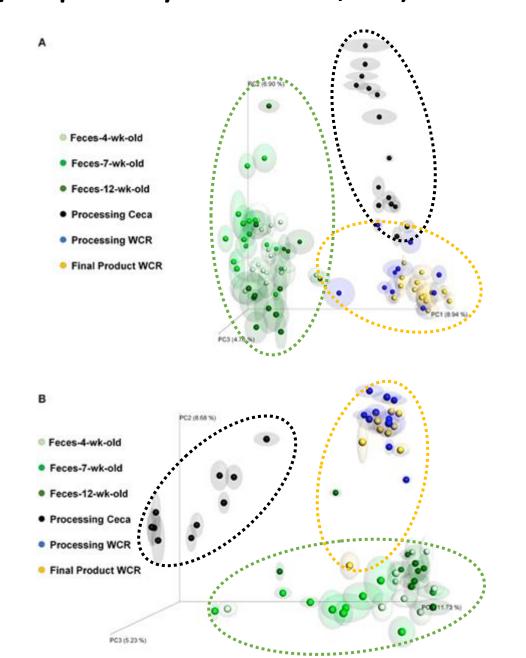
Microbial richness and diversity



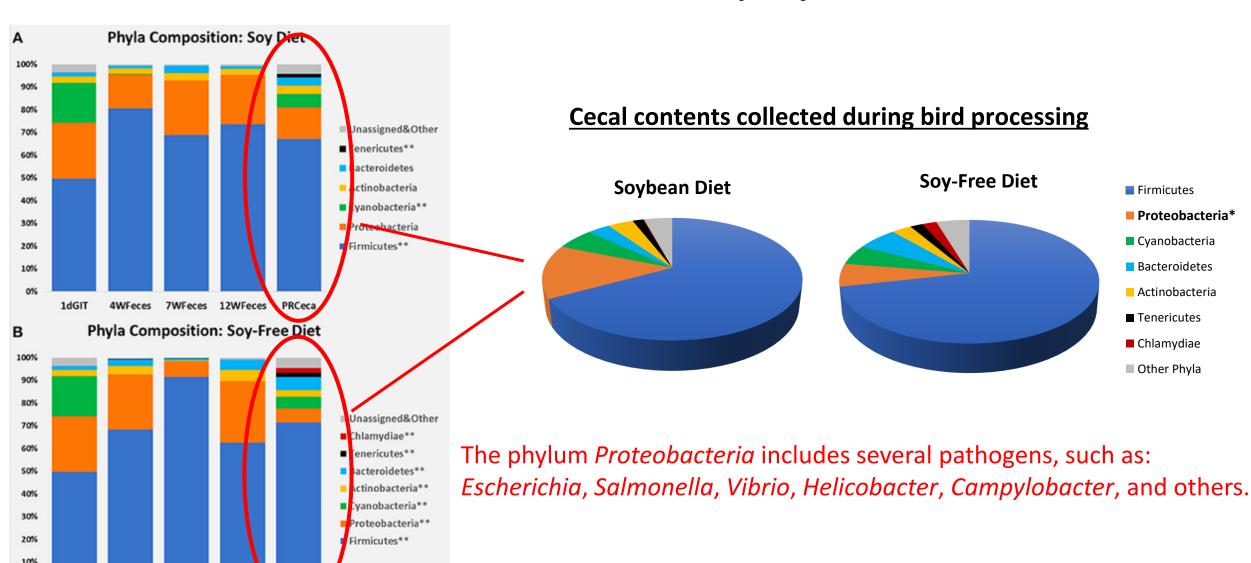
Effect of diet on α diversity indices				
	Suppl			
Item	Soybean	Soy-free	<i>P</i> -value	
Shannon Index (diversity)				
Cecal Contents from bird processing	6.6	7.4	0.001	

Effect of diet on α diversity indices				
	Supplement			
Item	Soybean	Soy-free	<i>P</i> -value	
Chao1 (richness)				
Cecal Contents from bird processing	647	1,065	0.001	

Beta-diversity: A) Top = Soybean diet; B) Bottom = Soy-free diet



Results: Main phyla



4WFeces 7WFeces 12WFeces

1dGIT

Foodborne pathogen groups found in the final product (frozen chicken carcass)

Effect of diet on abundance of foodborne pathogens (parts-per million) in the whole carcass rinses of the final product

	Suppl	Supplement	
Pathogen Group	Soybean	Soy-free	<i>P</i> -value
Genus Salmonella	1,170	321	0.38
Genus Acinetobacter	208,000	118,300	0.05
Genus Campylobacter	3,940	28	0.04

Foodborne Pathogens: <u>Campylobacter</u>

Effect of diet on relative abundance of *Campylobacter* (parts-per million)

	Supplement		
Type of Sample Evaluated	Soybean	Soy-free	<i>P</i> -value
Feces from 4-week-old birds	1,160	278	0.25
Feces from 7-week-old birds	52	46	0.83
Feces from 12-week-old birds	63	7	0.003
Processing Stage - Cecal Contents	1,220	499	0.28
Processing Stage - Whole Carcass Rinse	10,000	3,720	0.24
Final Product - Whole Carcass Rinse	3,940	28	0.04

Conclusions

- Regardless of which supplement was used, microbial diversity was greater in the cecal contents, compared to the feces
- Microbial richness and diversity were greater in the cecal contents of birds fed the soy-free supplement
- Supplementing broilers with the soy-free feed resulted in decreased populations of *Campylobacter* and *Acinetobacter* in the final product





For additional information on this study, please refer to these 2 publications:

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ORIGINAL RESEARCH published: 15 May 2019 doi: 10.3389/fsufs.2019.00035



The Successional Changes in the Gut Microbiome of Pasture-Raised Chickens Fed Soy-Containing and Soy-Free Diets

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The succession of bacterial species as birds mature can impact their growth efficiency, health, and food safety due to the ability of a mature gastrointestinal microbial population doi: 10.3389/fsufs.2019.00036



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The Effects of Feeding a Soybean-Based or a Soy-Free Diet on the Gut Microbiome of Pasture-Raised Chickens Throughout Their Lifecycle

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Edited by:

Soybean is one of the primary ingredients in poultry diets, but it causes problems in some consumers with allergies. Thus, production of poultry without soybean in their diets has increased in recent years. In addition, consumers are increasingly supporting

Thank you!

Any questions?



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