Diet-induced changes in the jejunal microbiota of broilers during development, greatly affecting *Enterococcus hirae* and *Enterococcus faecium* 

#### September 2024, Dirkjan Schokker et al.











#### **Acknowledgements**

#### **Knowledgebase project**

"Towards a circular climate positive today society" (KB-33-006-005)



Wageningen Bioveterinary Research

Wageningen Livestock Research

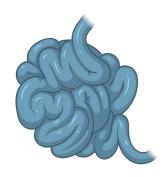




## **Early Broiler Development Critical**



## **Important Role for the Gut**



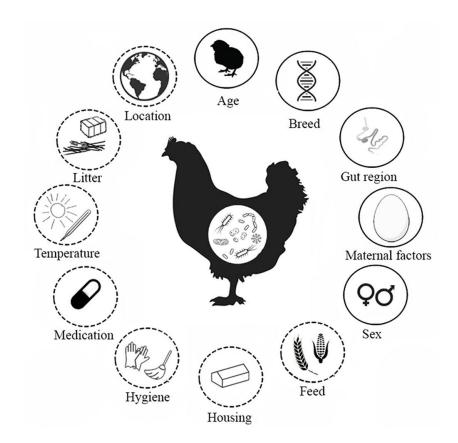
- Nutrient uptake
- Physical barrier
- > Gatekeeper of health

**Animal performance** 





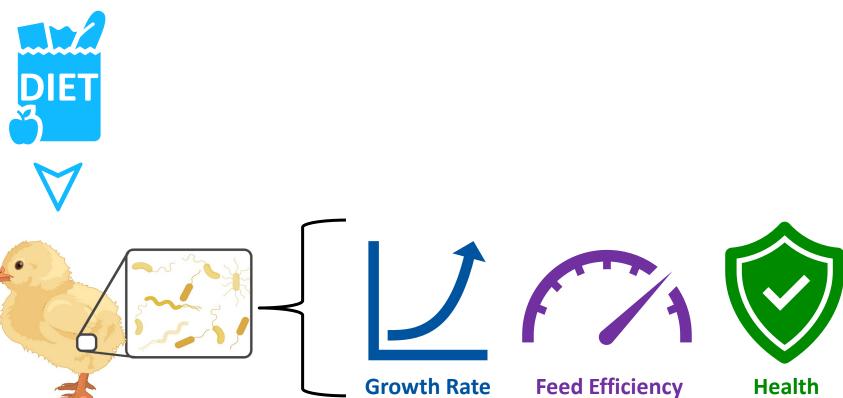
# **Characteristics Influencing Gut Microbiota**







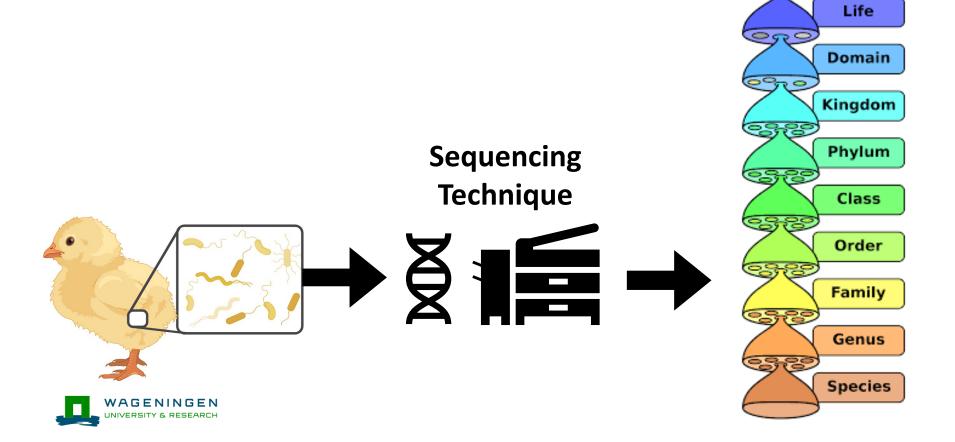
## **Impact of Diet on Gut Microbiota**







## **Defining the Microbiota**

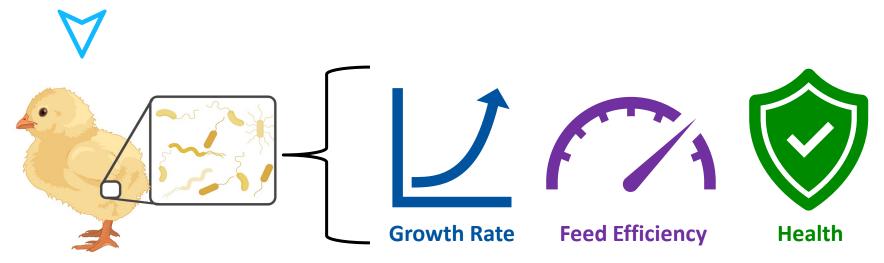




#### **Objective**



Assess dietary impact on jejunal microbiota composition at <u>species level</u> in broilers







#### **Experimental Design**



1,344 day-old Ross 308 male broiler chickens [average weight of 43.3 g]



Floor pens with wood shavings as substrate



ad libitum access to feed and water



Randomly allotted to 6 diets in a completely randomized block design

#### This study subset of 4 diets



**CON** control diet without any supplementation

control diet supplemented with micro-encapsulated sodium butyrate

MCFA control diet supplemented with a mixture of medium-chain fatty acids

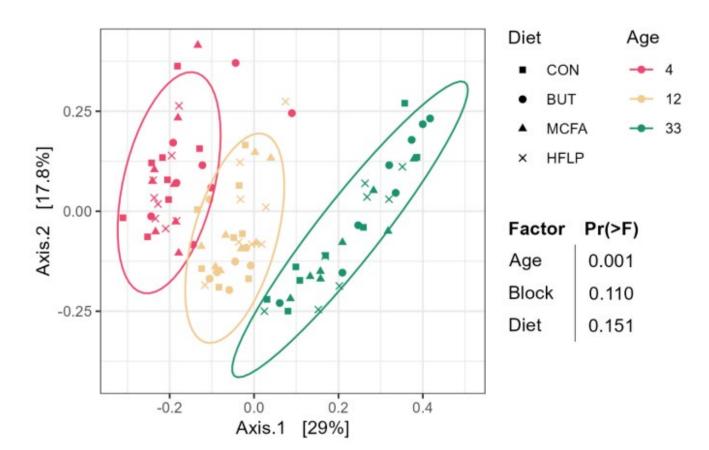
**HFLP** a diet with a higher fiber and lower protein content compared to the control diet



**BUT** 



#### **Significant Age-Dependent Effect**

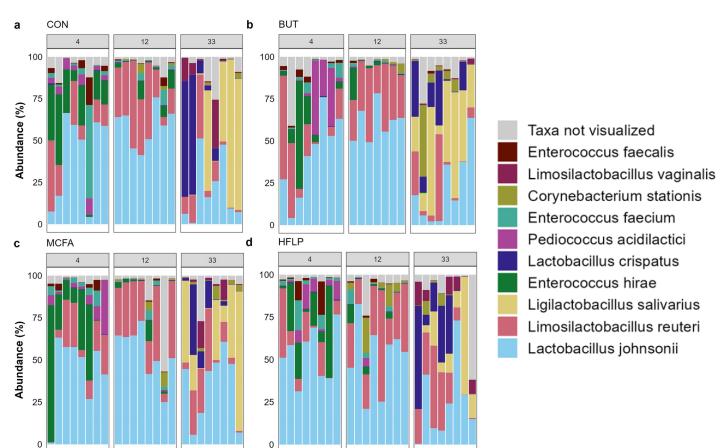






## For the purists;)

Individual broilers



Individual broilers





#### **Five Species with the Largest Effect Size**

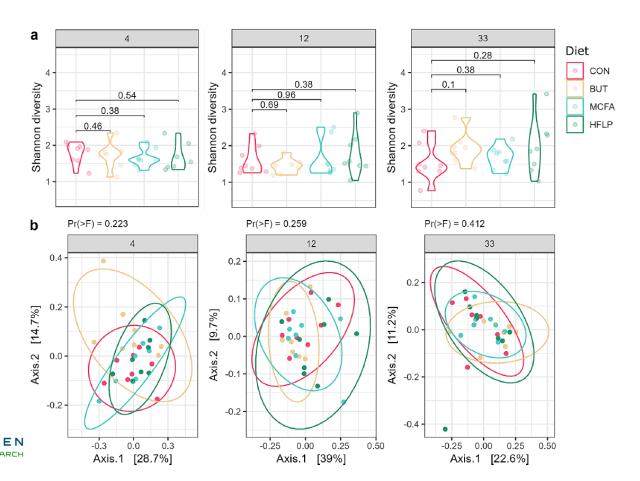
#### These genera show the largest change in relative abundance over time

- > Lactobacillus johnsonii
- Limosilactobacillus reuteri
- Corynebacterium stationis
- Lactobacillus crispatus
- Ligilactobacillus salivarius





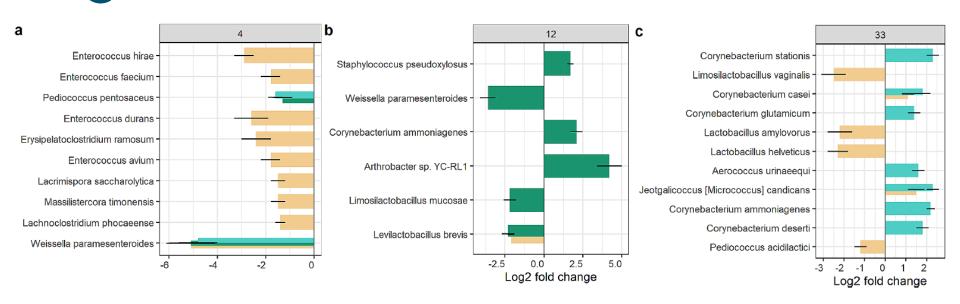
## Diet Not a Main Driver of Microbiota Composition





#### **Changes Specific to Age / Diet**

- 1 Across all diets → prevalence > 10% + abundance > 0.001%
- 2 **Per diet group** → abundance > 0.01% + an absolute fold change > 2





**BUT / MCFA / HFLP** 

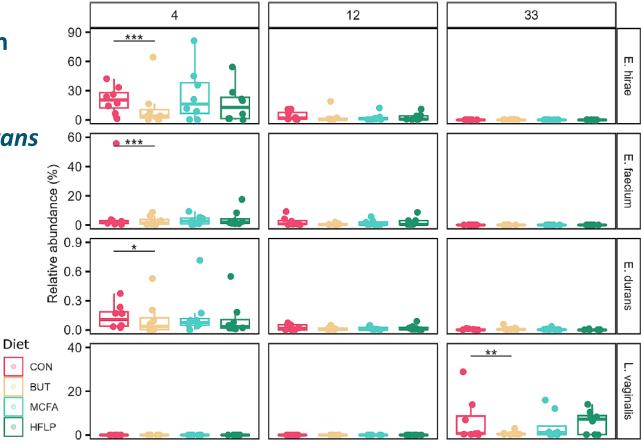


#### **Reduced Relative Abundance Pathogens**

High relative abundance in CON of opportunistic pathogens:

E. hirae, E. faecium, E. durans

Day 4 effects significant between BUT vs. CON







#### **Conclusions**

Metagenomic shotgun sequencing enabled us to study speciesspecific effects

BUT, MCFA and HFLP diets induced changes in the jejunal microbiota composition of specific bacterial species of broilers at d4, d12, and d33 post-hatch

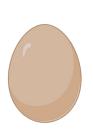
Most notable was the effect of BUT 4 days post-hatch, reducing the abundance of *E. faecium* and the opportunistic pathogen *E. hirae* 

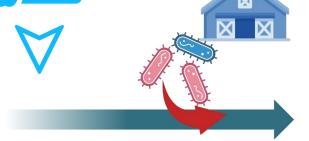




#### Take home message





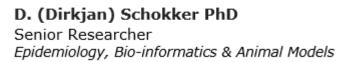






To explore the potential of nature to improve the quality of life





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