

monoguthealth

Optimal gut function in monogastric livestock

Differences between low and high body weight broilers and strategies for catch up growth of low-performing chickens

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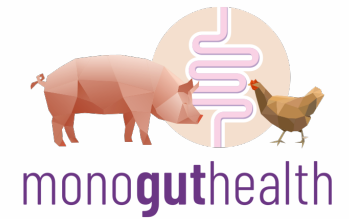
EAAP Annual Meeting, 1/5 September 2024



This project has received funding from the European Union's Horizon 2020 research and innovation programme under the Marie Skłodowska-Curie grant agreement No 955374.

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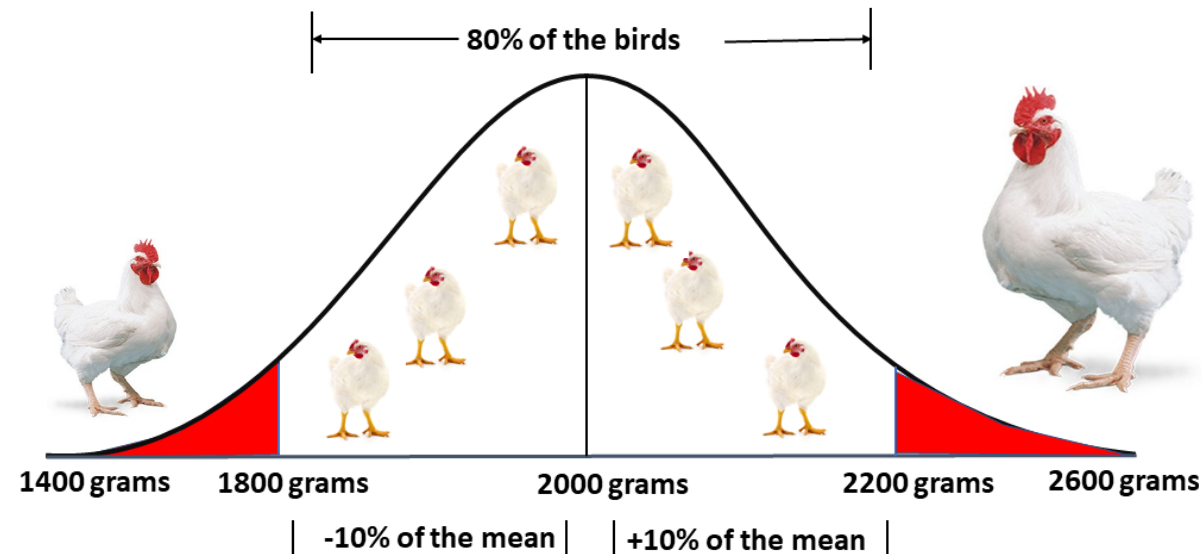
04

Study 3: Effects of different feed structures on growth performance and gastrointestinal development of low-weight broilers



Introduction

- Intensive selection has reduced the genetic variation in modern broilers and improved their growth performance.
- **Flock weight uniformity:** Percent individuals within a specific range of BW, measures in coefficient of variation (CV). CV is inversely proportional to flock weight uniformity.
- Broiler flocks with $\pm 8\text{-}10\%$ **CV** are considered uniform.



Akram et al., 2024; Animal Microbiome

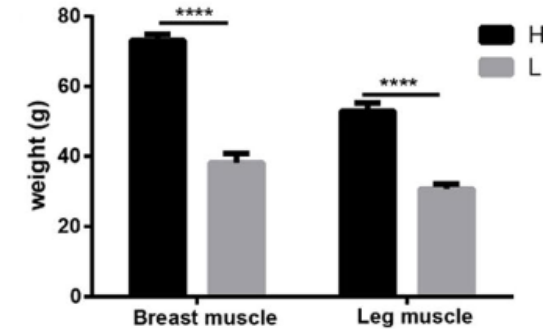
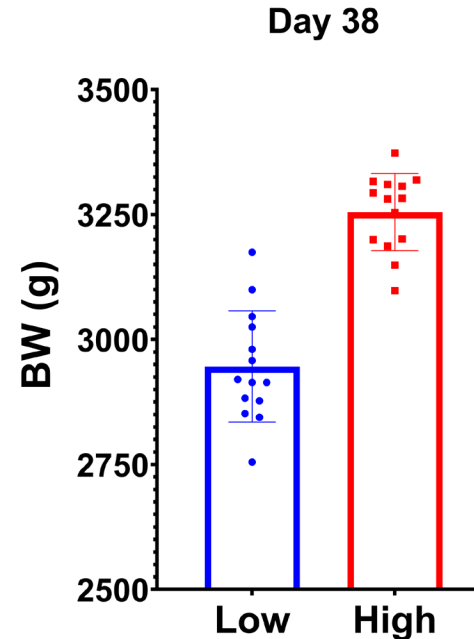
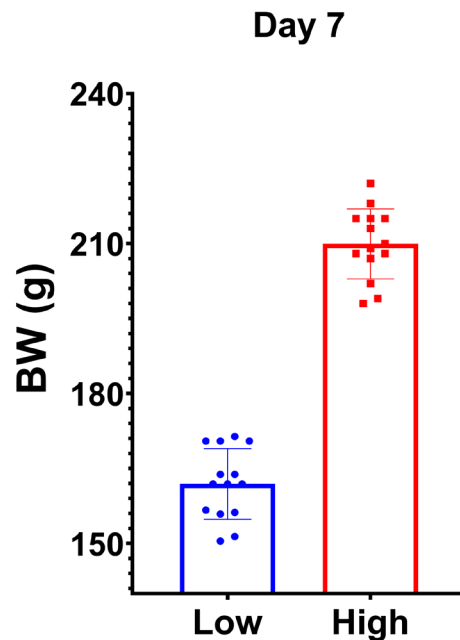
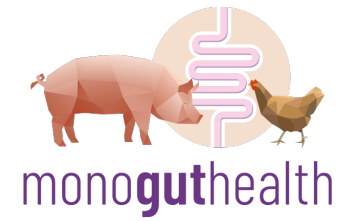


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Introduction

- In commercial settings, broiler chickens have **different growth rates within a flock** under uniform rearing conditions.
- What are the biological factors causing intra flock body weight (BW) variability?



Zhang et al. Microbiome (2022)



Akram et al., 2024; Animal Microbiome

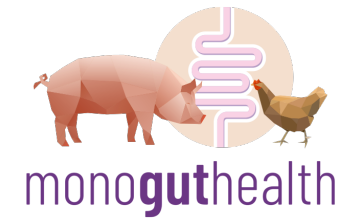


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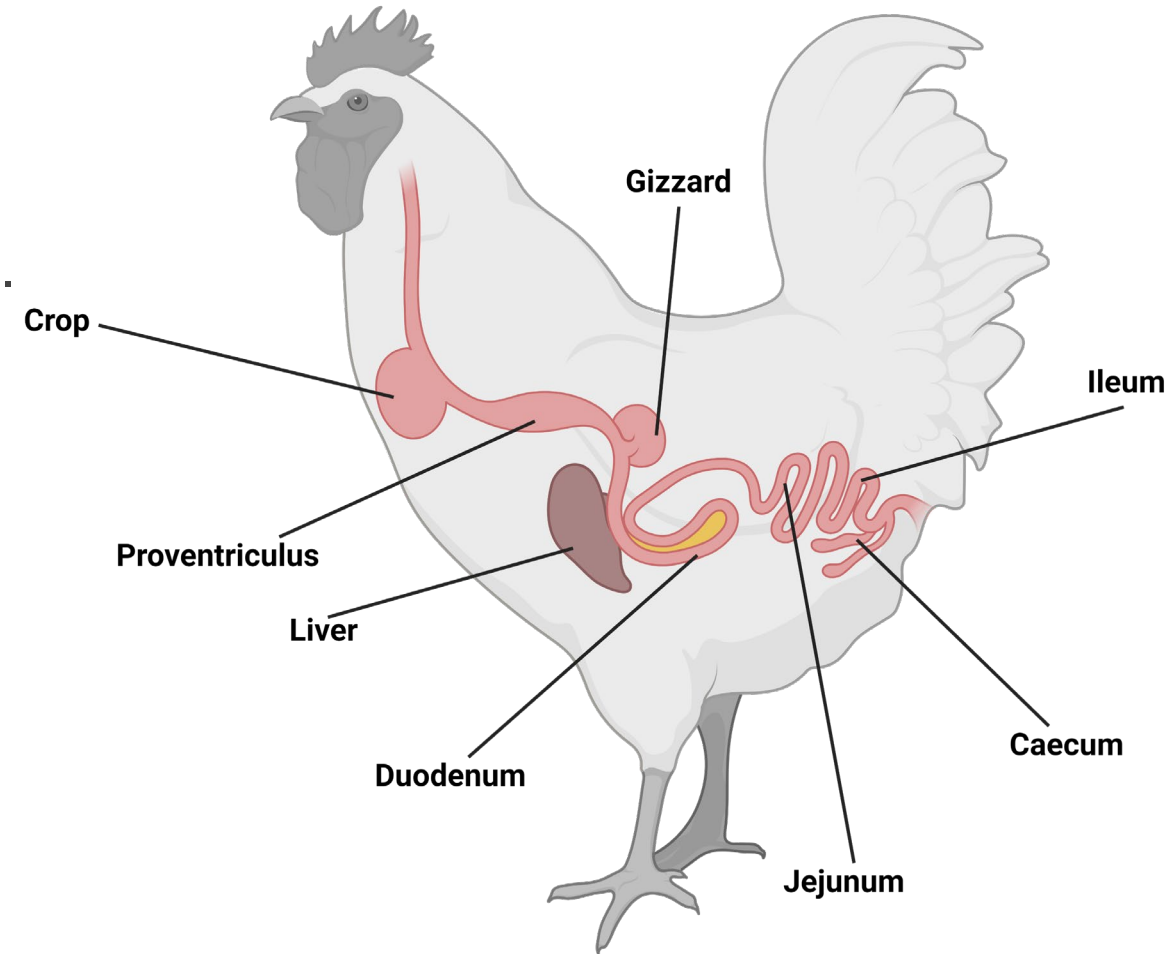
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Introduction



- **Gastrointestinal tract** is more proliferating organ in early life.
- **Early development** of gastrointestinal tract is important for optimal growth and development.
- Helps in meeting **nutritional and immunological demands** in early life.
- Boosts **immune function** and disease resistance.
- Reduces early **mortality and health issues**.

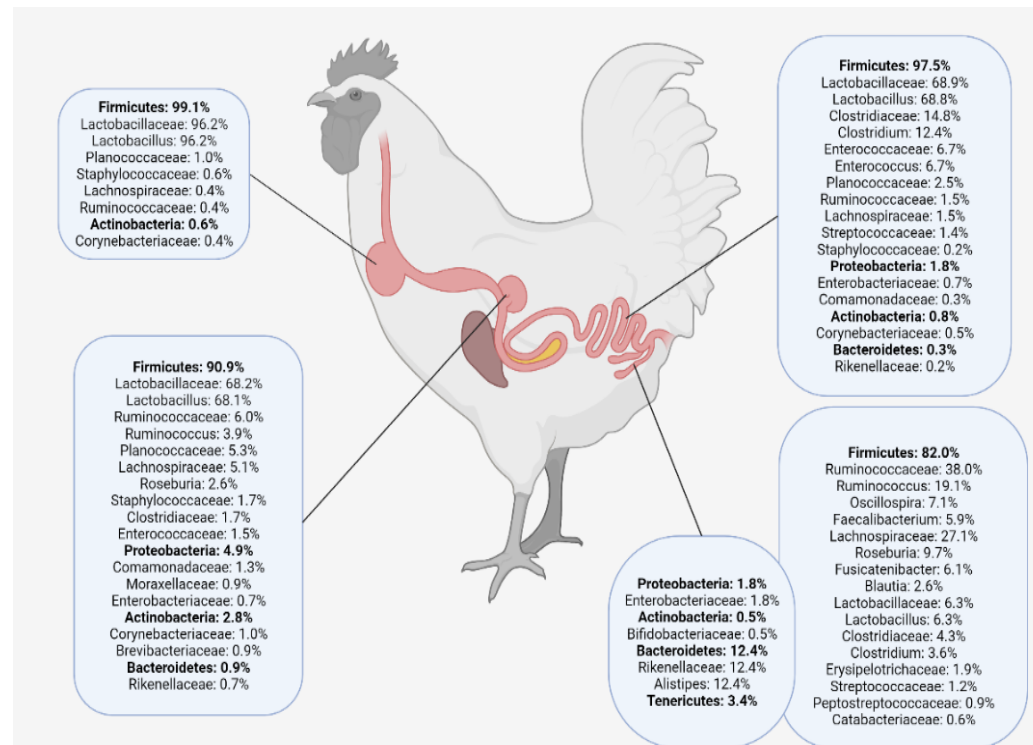
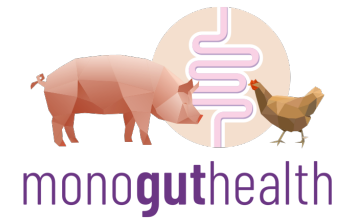


Based on <https://doi.org/10.3390/ani11102795>

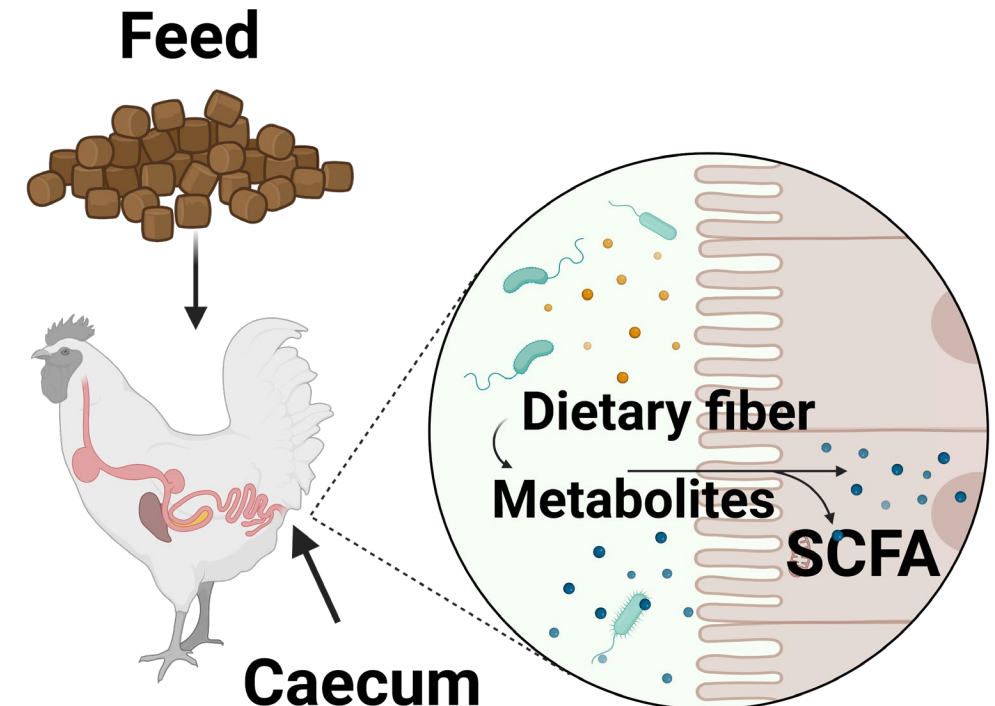


Introduction

- **Gut microbiota** residing in the digestive tract, contribute to digestion, nutrient absorption, metabolism, and immune regulation
- Intestinal microbiota produces **high-energy metabolites (short-chain fatty acids)** that benefit the host



Based on <https://doi.org/10.3390/vaccines11030595>



Created with BioRender.com



This project has received funding from the European Union's Horizon 2020 research and innovation programme under the Marie Skłodowska-Curie grant agreement No 955374.

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Introduction

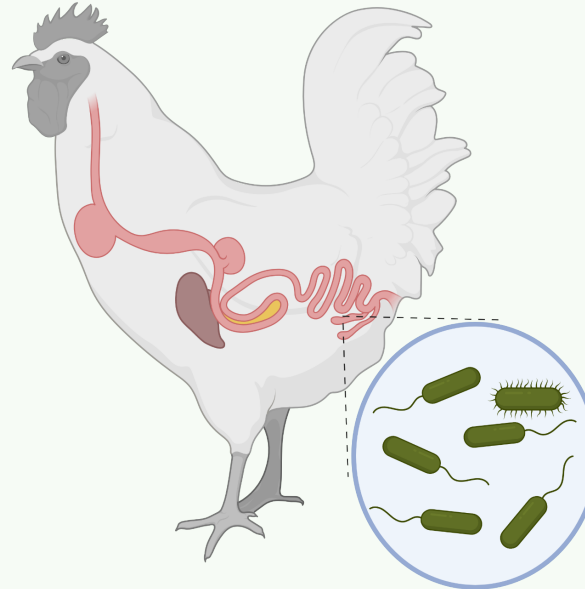
'EUBIOSIS'

Growth and health
conferring balanced
microbiota
community

'DYSBIOSIS'

microbial community
involving disbalances
or pathologies occur

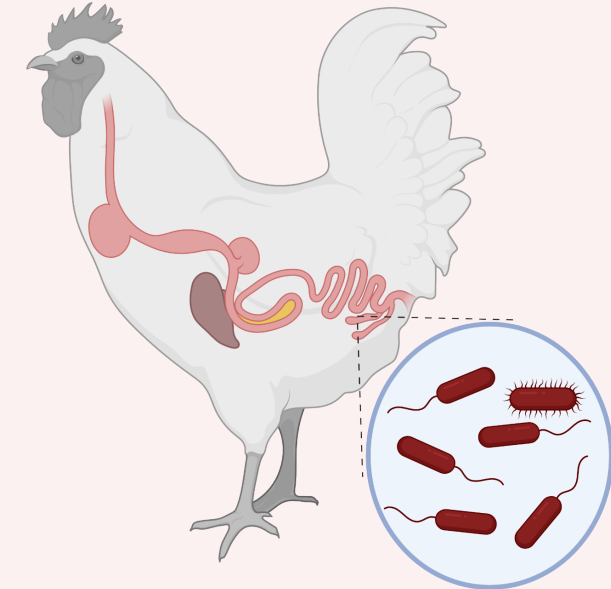
Eubiosis



↑ Firmicutes to Bacteroidetes ratio
Lactobacillus, *Bifidobacterium*, and
Bacteroides....

- Reduced Pathogen Load
- Balanced Immune Function

Dysbiosis



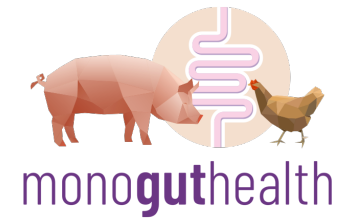
↓ Firmicutes to Bacteroidetes ratio
↑ *Campylobacter*, *Salmonella*, and
Escherichia.....

- Increased Disease Susceptibility
- Retarded growth performance

Based on <https://doi.org/10.1016/j.aninu.2023.03.005>

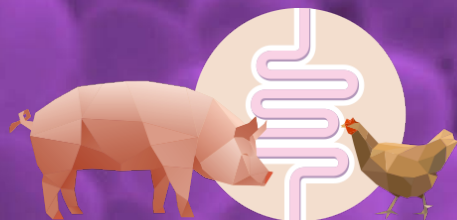
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Objectives of the project



- Characterization of **gastrointestinal related factors** contributing to body weight variation in broiler chicken flocks.
- Development of **supplementary interventions** to improve enteric microbial colonization in lo-performing broiler chicks to improve intestinal health and function.
- Development of **dietary strategies** to improve early life performance and health in low-performing broiler chicks and to reduce flock body weight variations.





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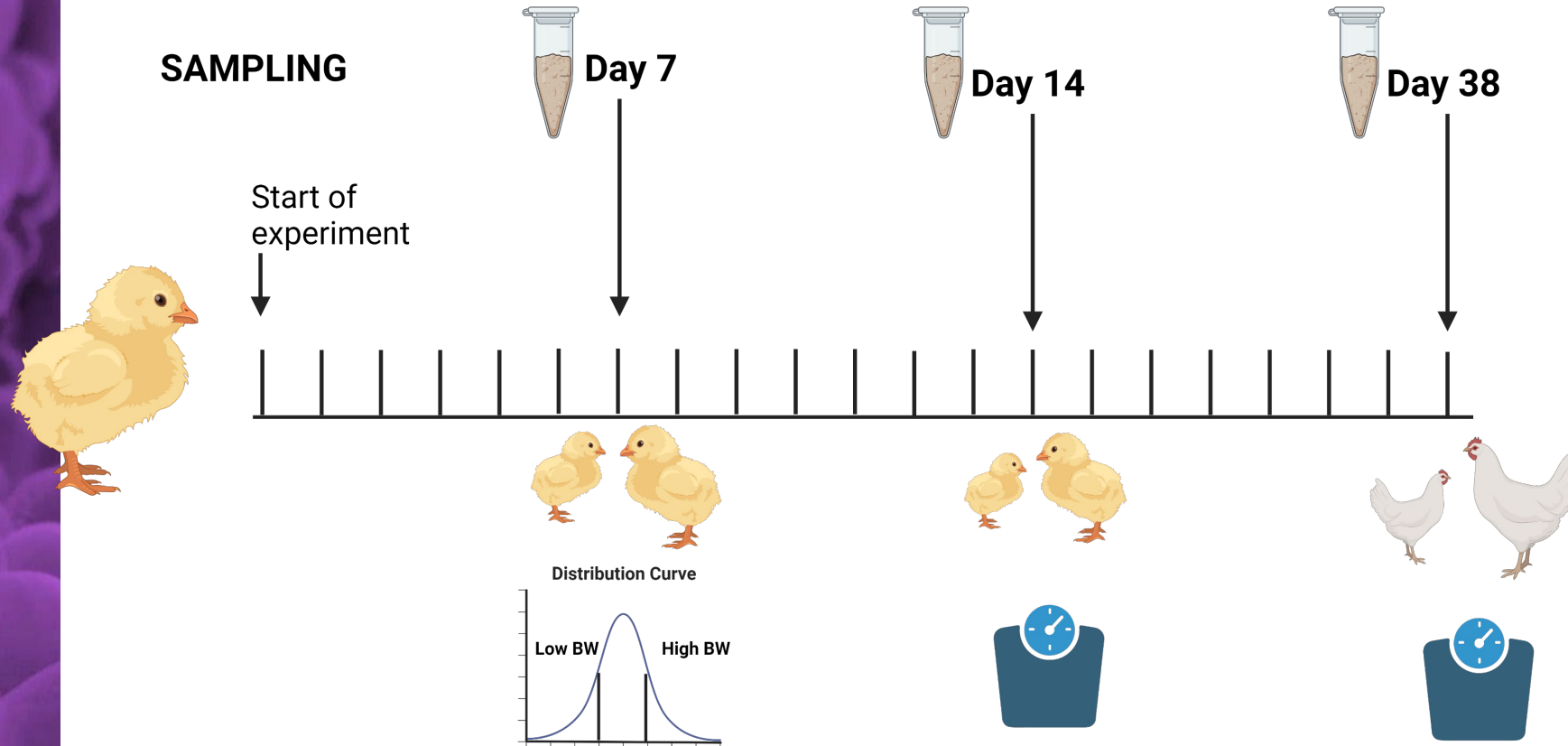
Optimal gut function in monogastric livestock

01

Study 1: Identification of gastrointestinal related factors affecting body weight variability in broiler flocks



Trial design



- Ross 308 male broilers
- Floor pens, 21 birds/pen
- 14 replicate pen/BW group
- Bodyweight categorization at day 7

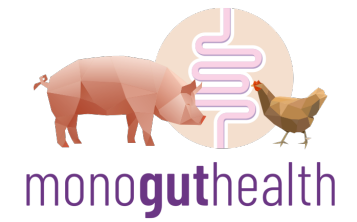
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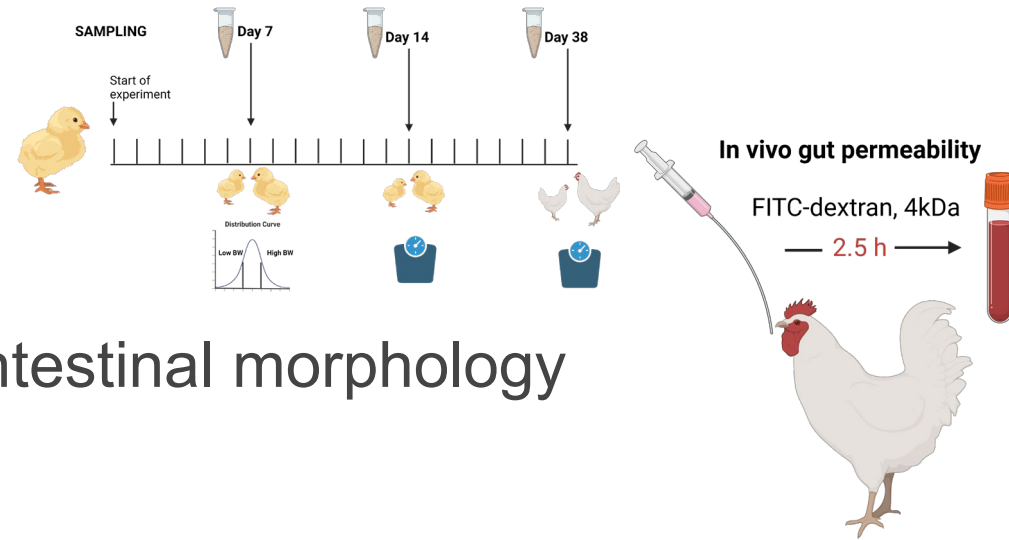
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Overview of analysis

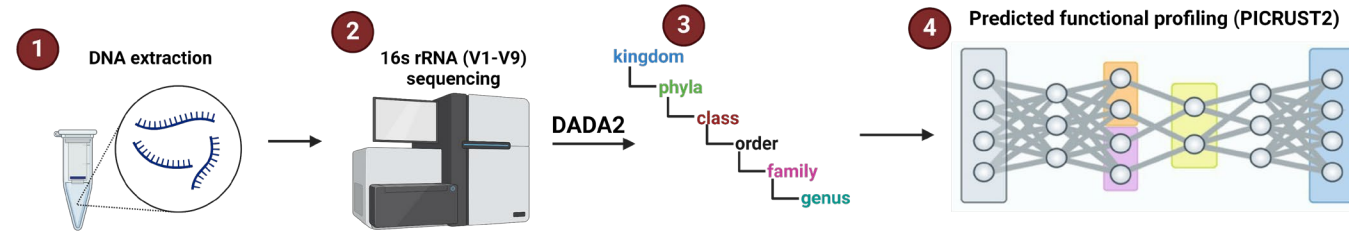


1. Growth performance

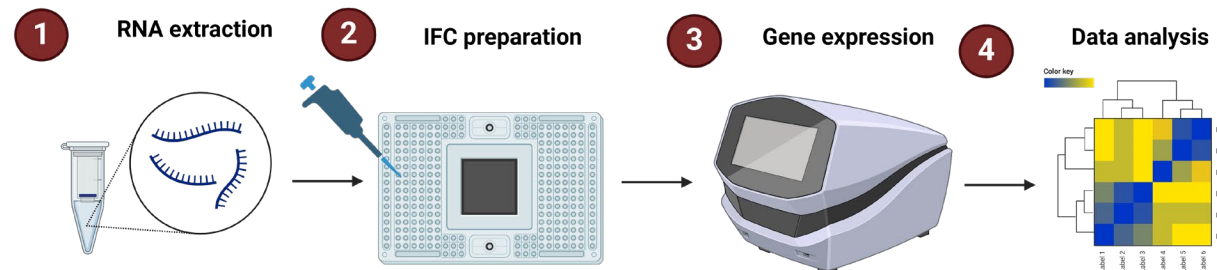


2. *In vivo* permeability, intestinal morphology

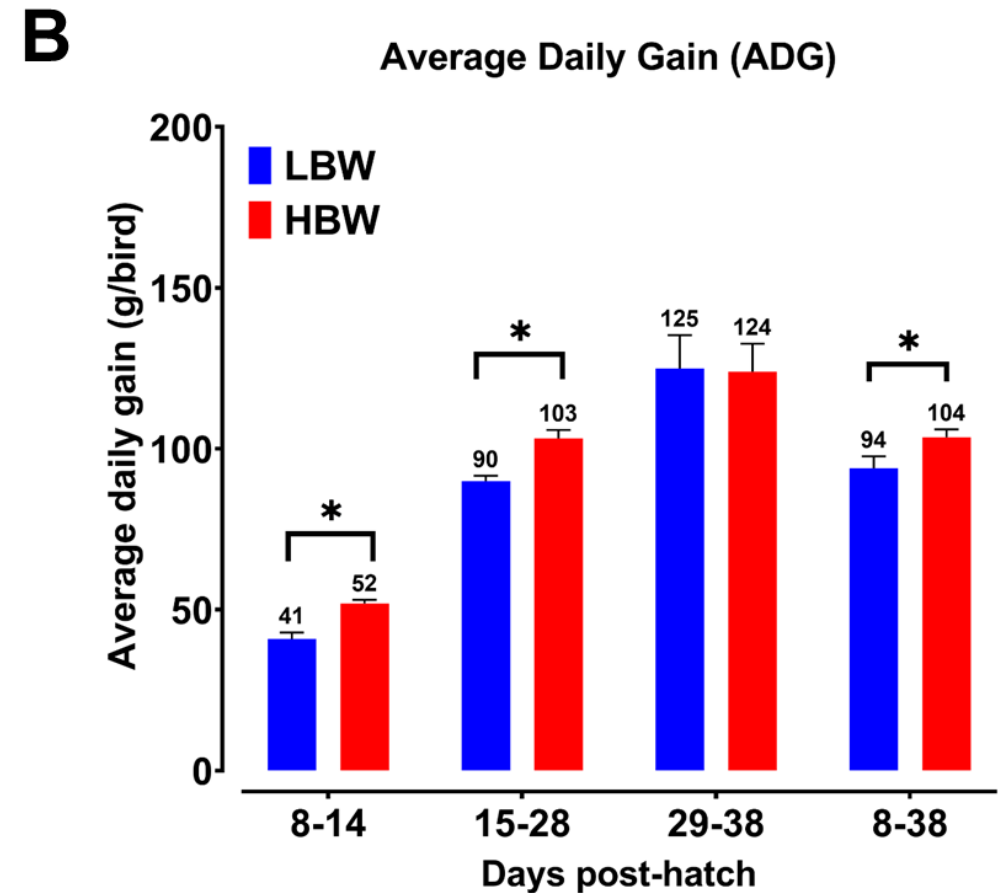
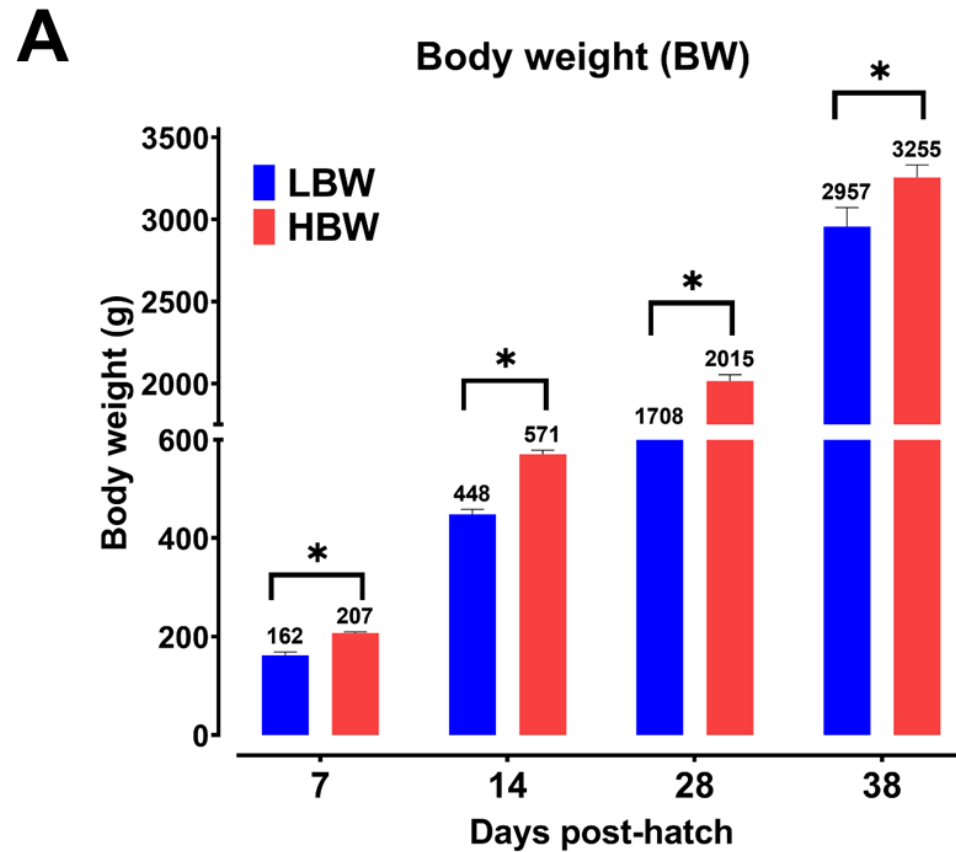
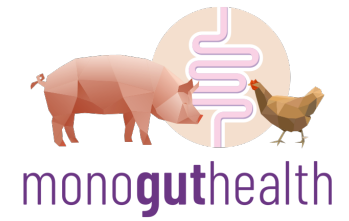
3. Caecal microbiota composition



4. Ileal gene expression



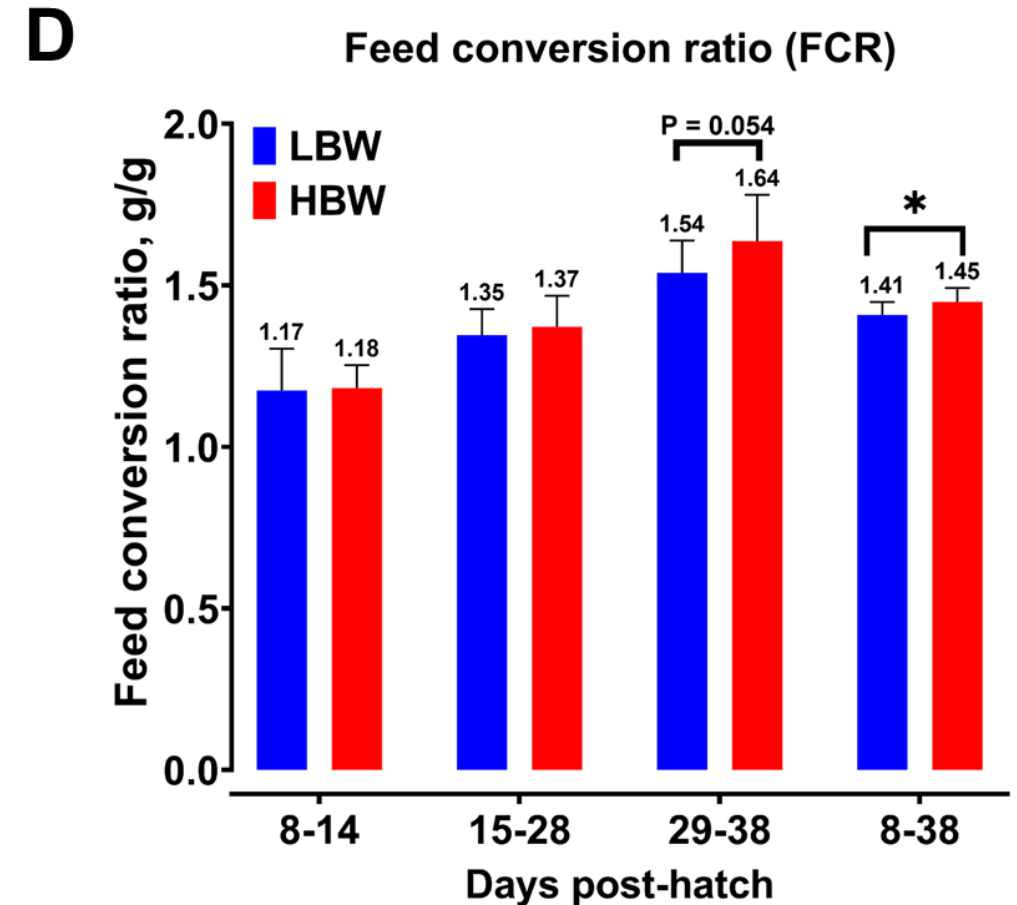
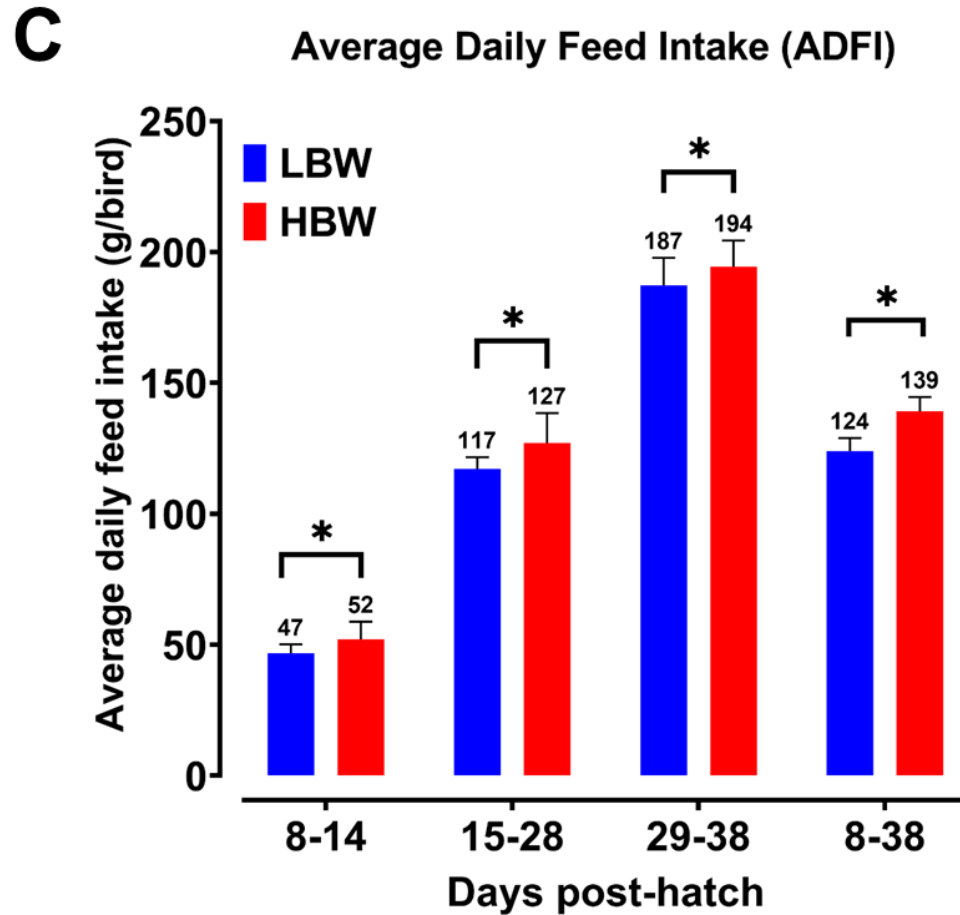
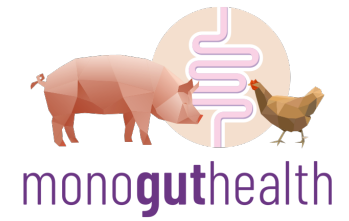
1. Growth performance



Student's t-test, $P < 0.05$



1. Growth performance

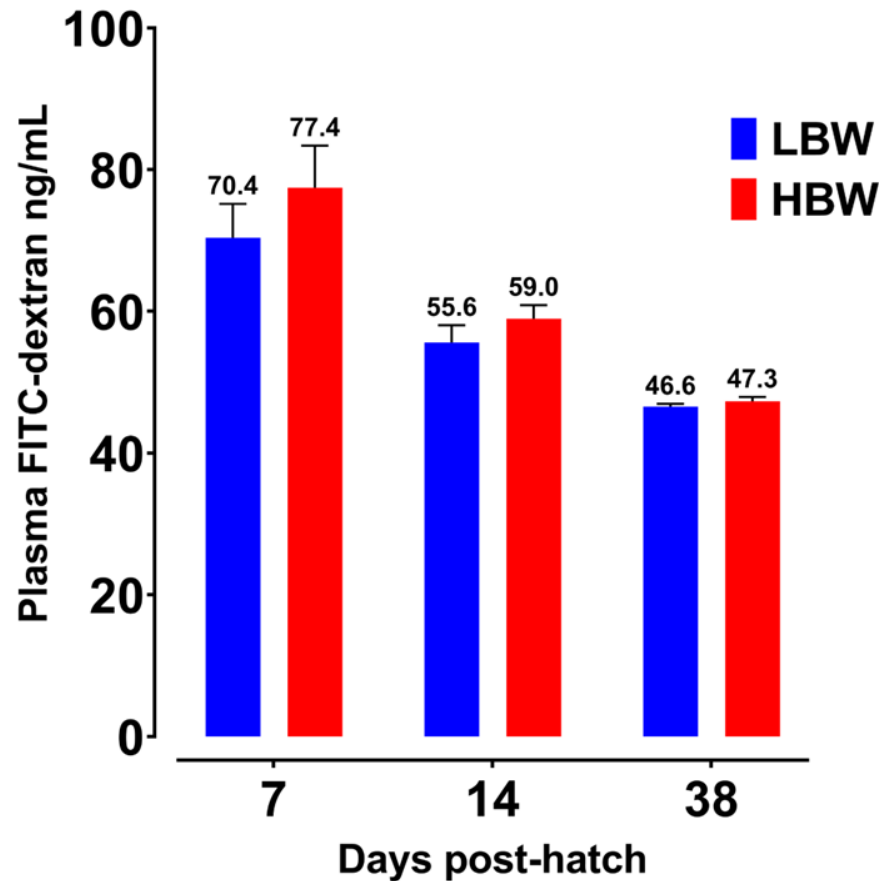


Student's t-test, $P < 0.05$

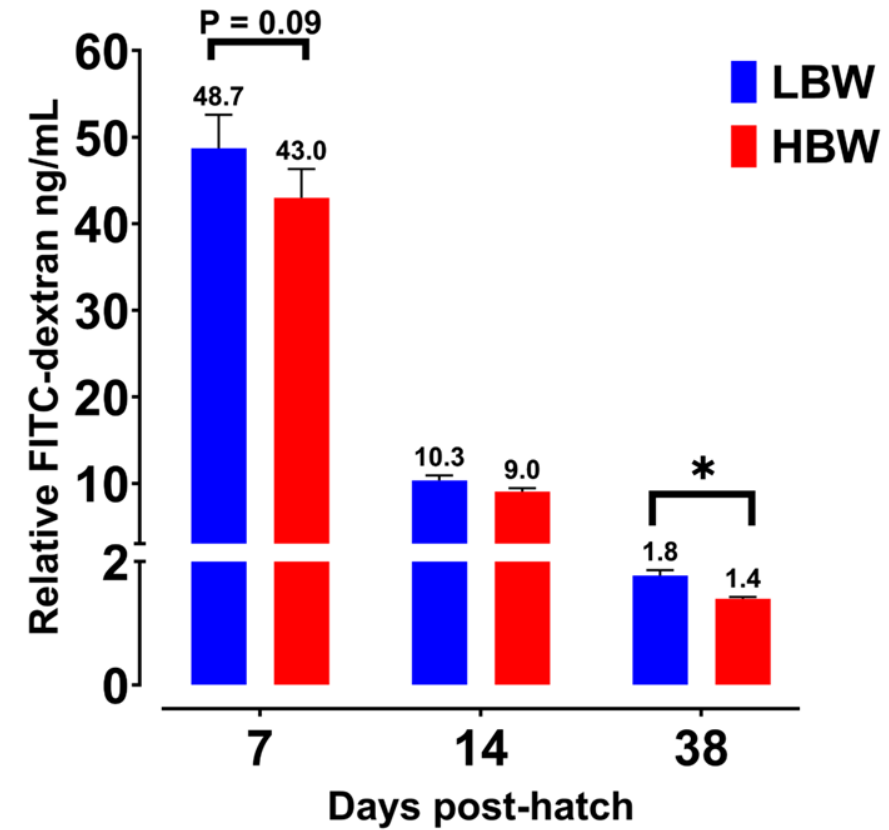


2. Intestinal permeability

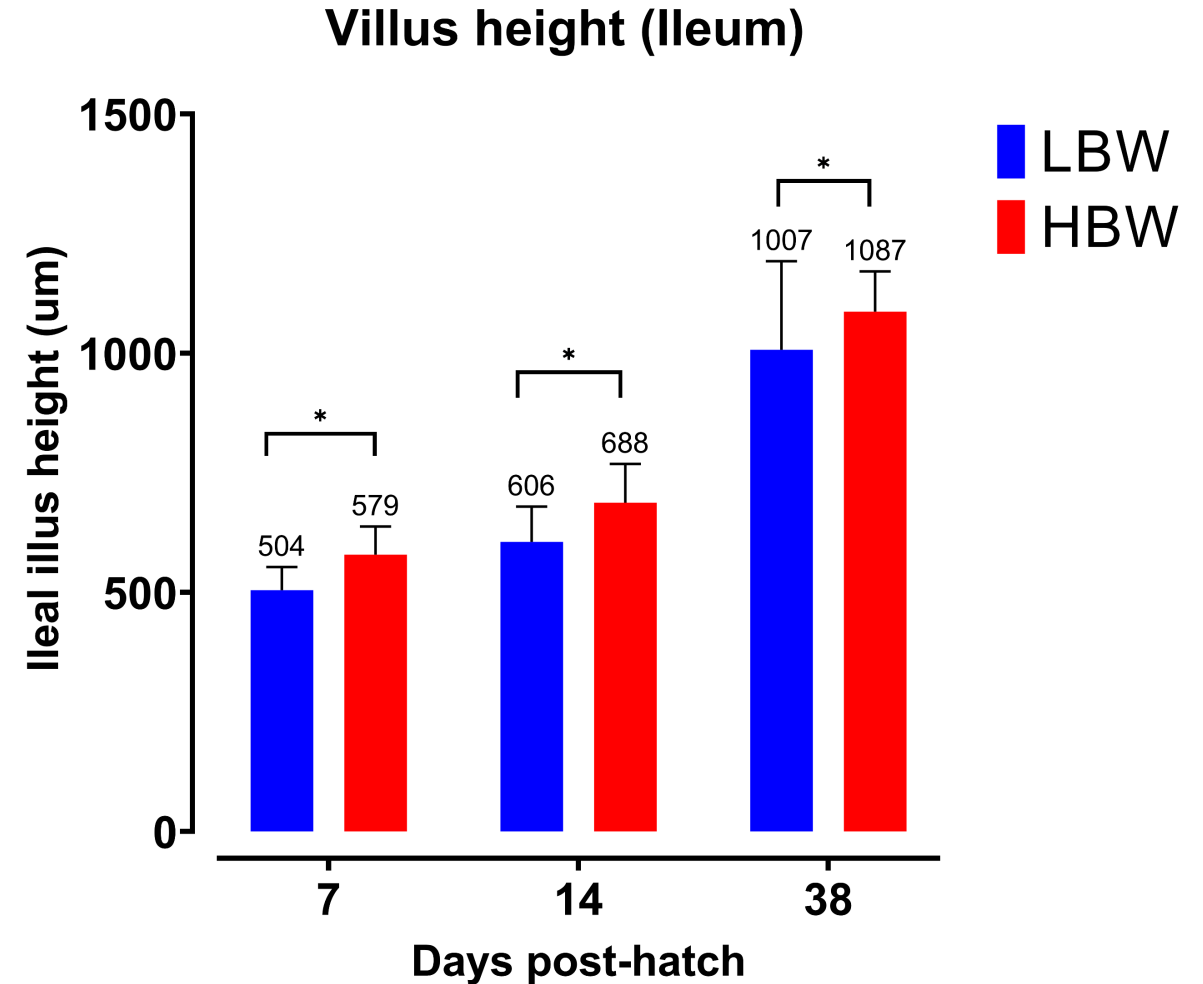
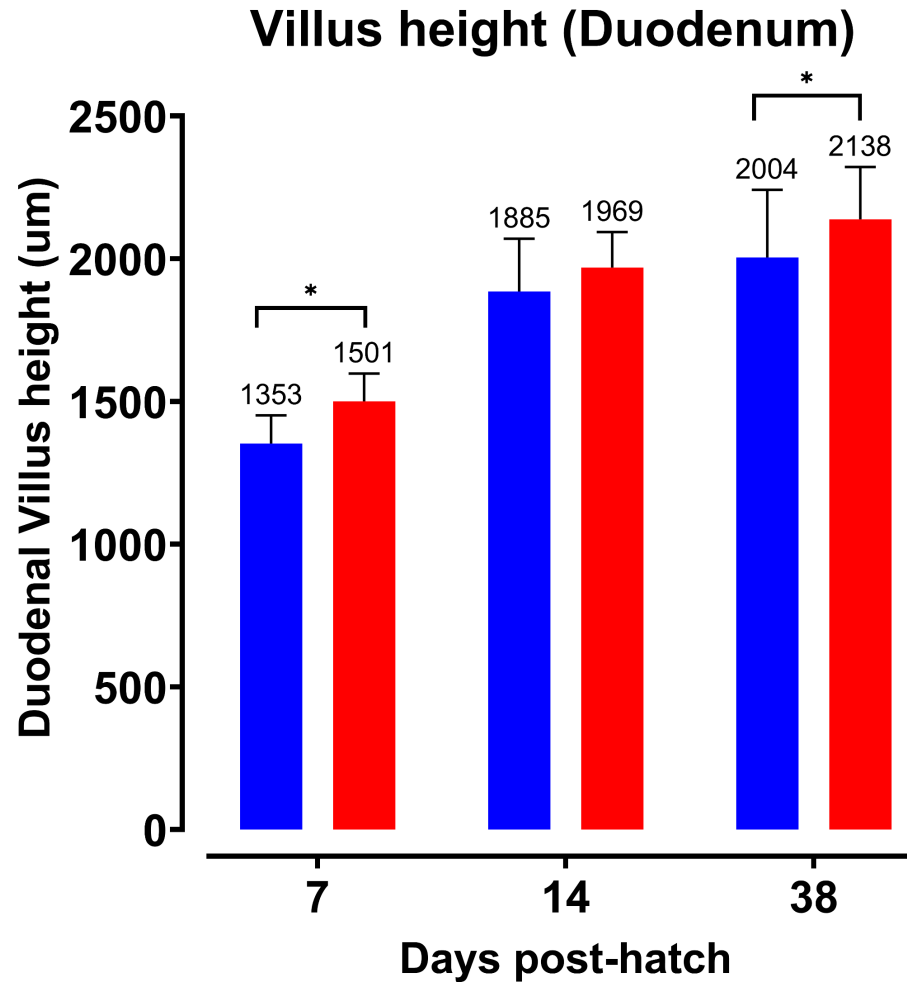
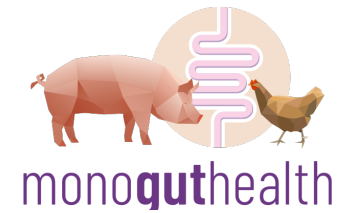
A Absolute plasma FITC-dextran



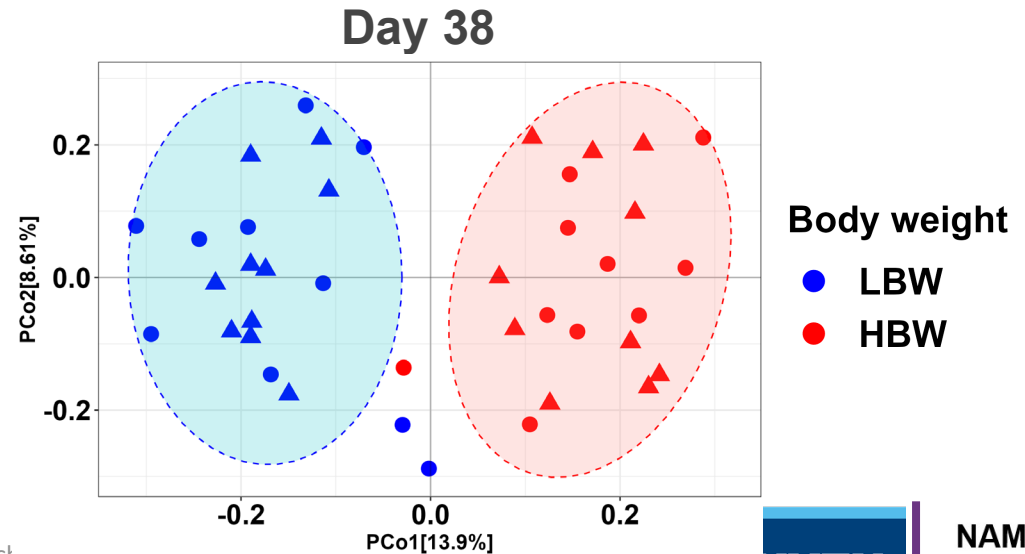
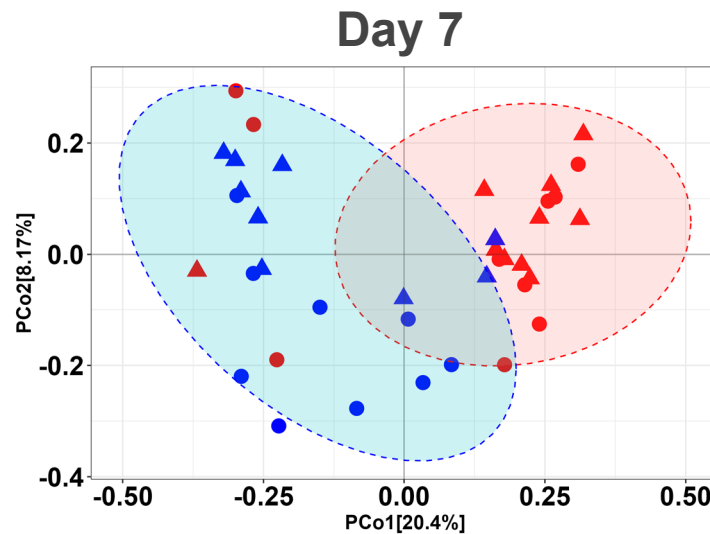
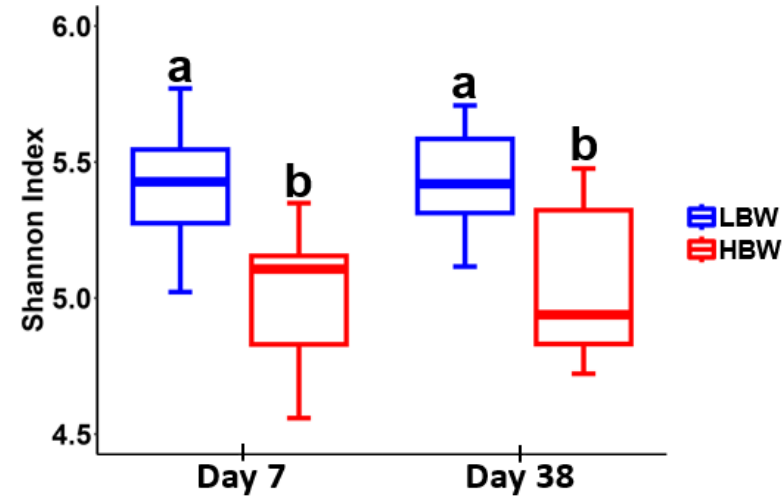
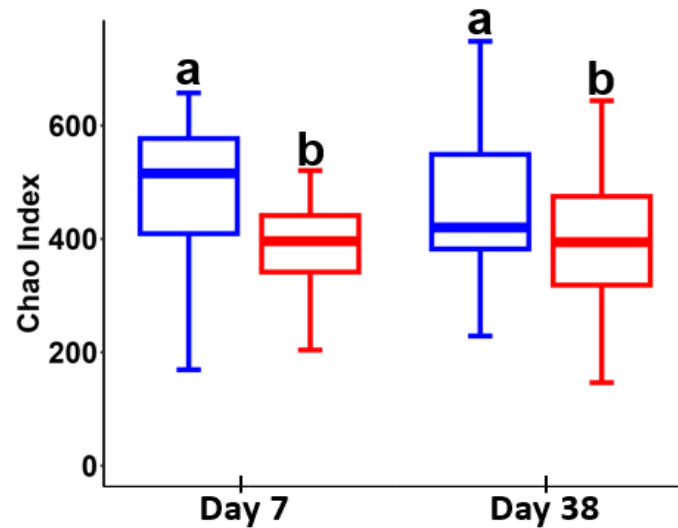
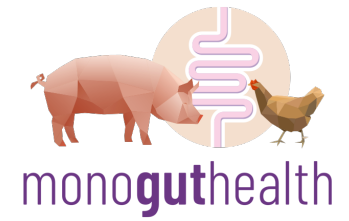
B Relative plasma FITC-dextran



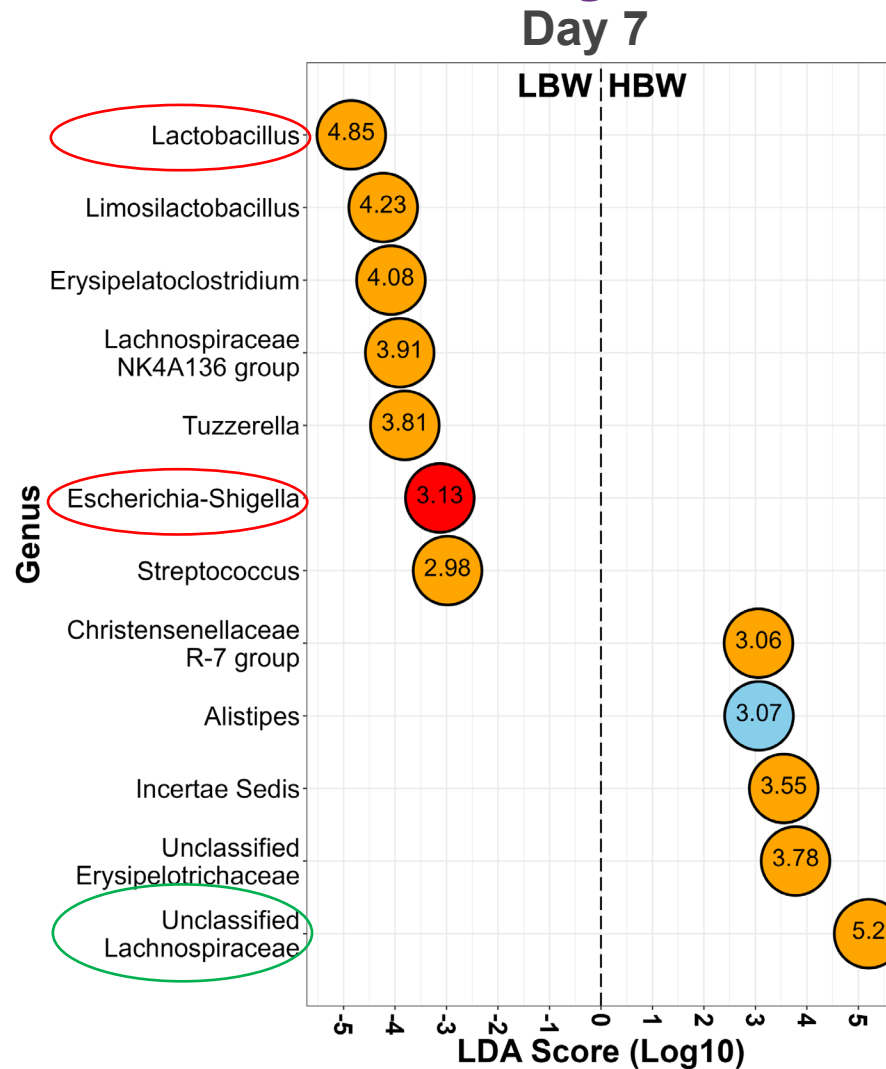
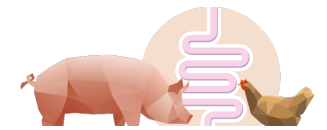
3. Intestinal histomorphology



3. Caecal Microbiota diversities

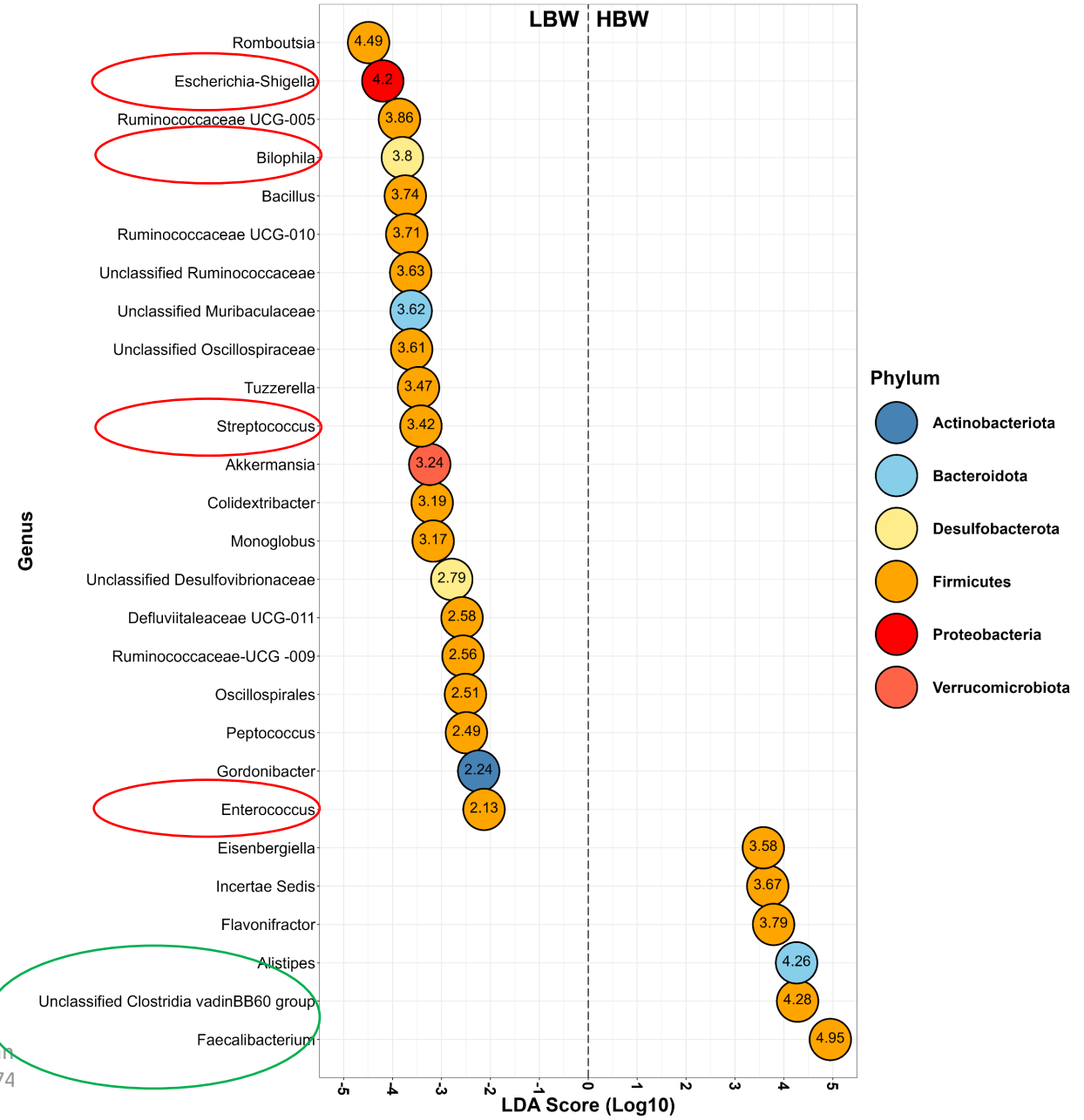


3. Differentially abundant bacterial genera

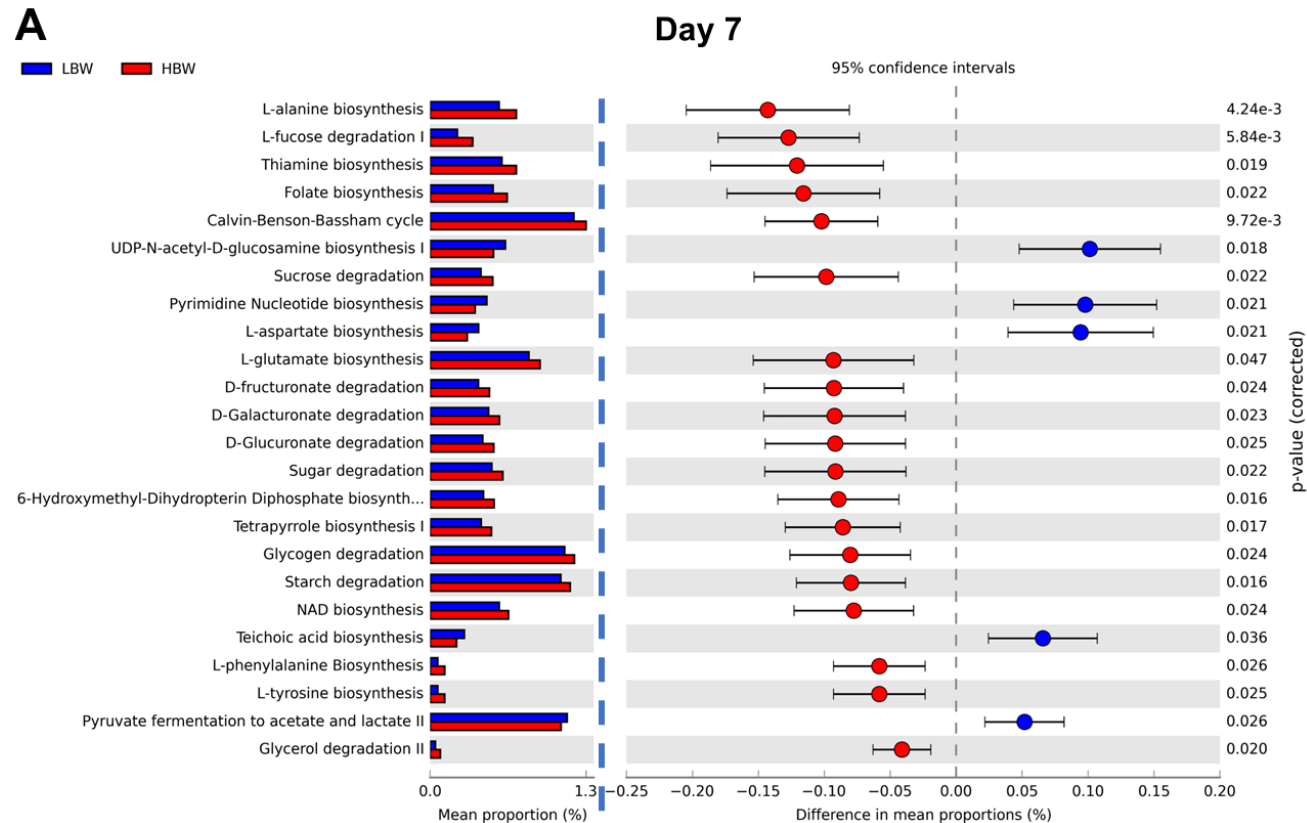
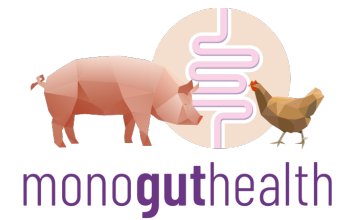


16S rRNA gene sequencing

LEfSe, LDA > 2.0, FDR < 0.05

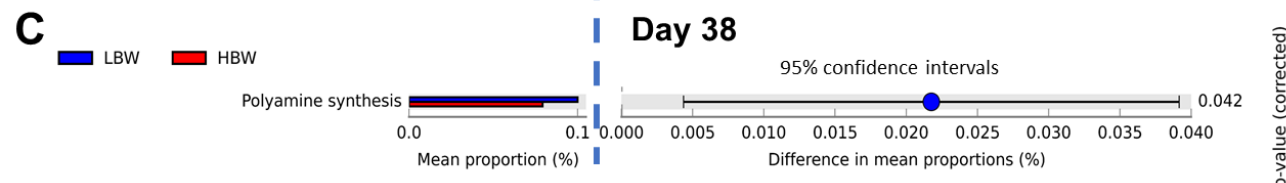
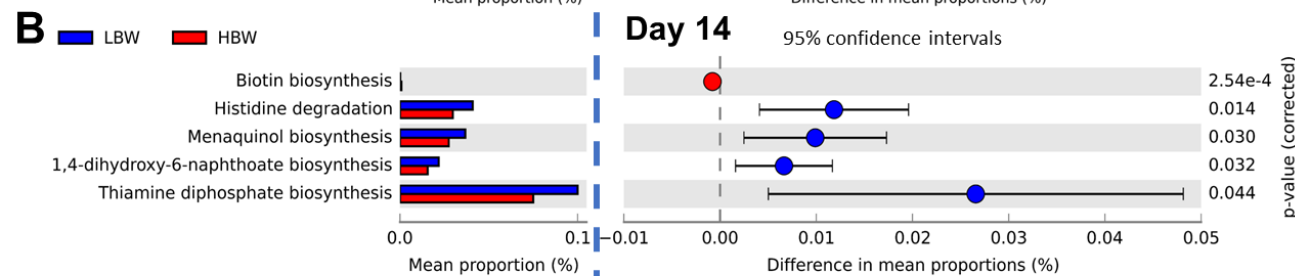


3. Predicted functionality – PICRUST2

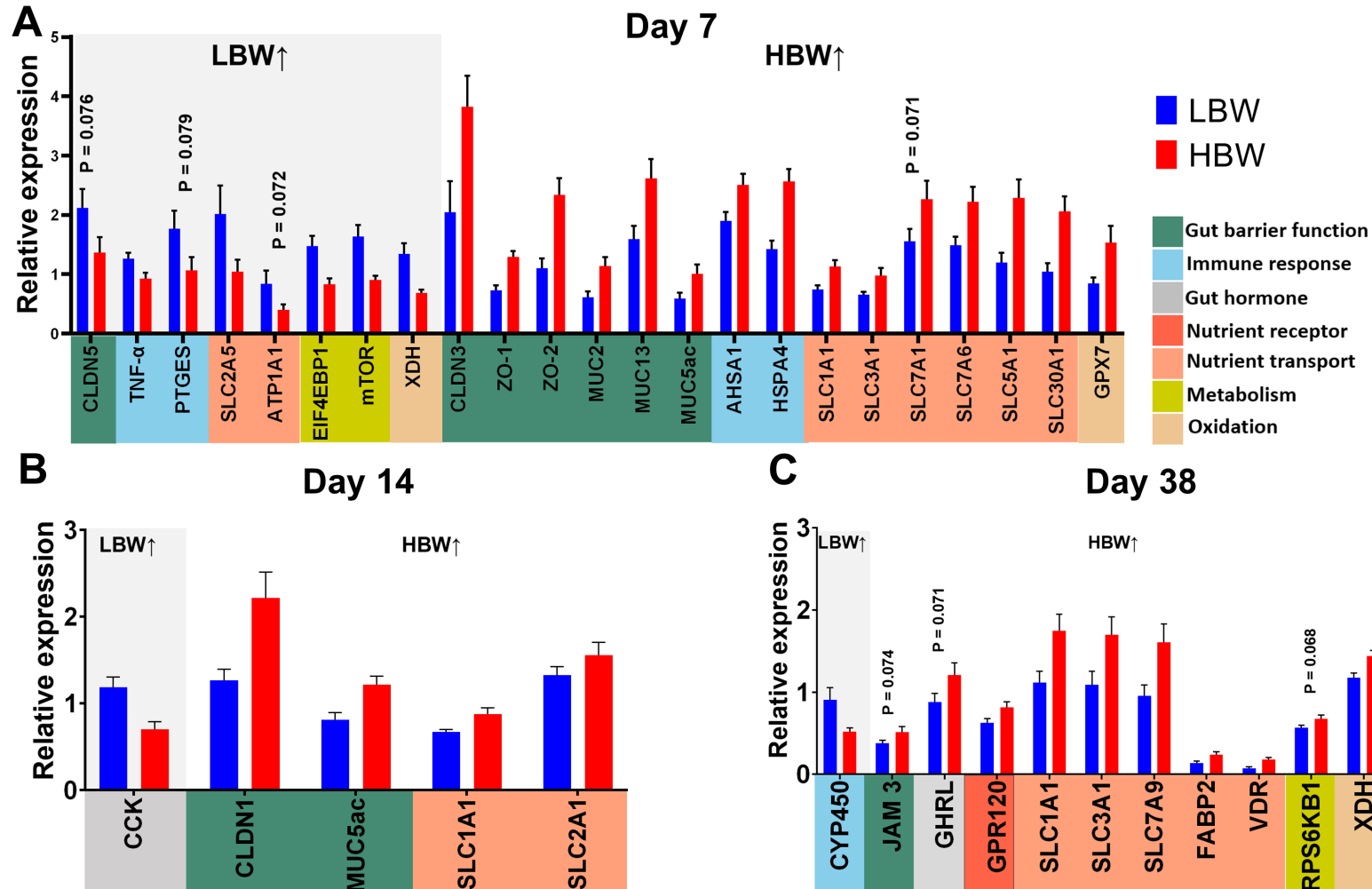


HBW: Microbial pathways enriched in the biosynthesis of amino acids, vitamins and degradation of complex carbohydrates.

LBW: Pathways enriched in cell wall antigen (UDP-N-acetyl-D-glucosamine)



4. Ileal gene expression



HBW → Barrier function, nutrient transport and antioxidant genes ↑

LBW → Immune response and neuropeptide hormone CCK ↑

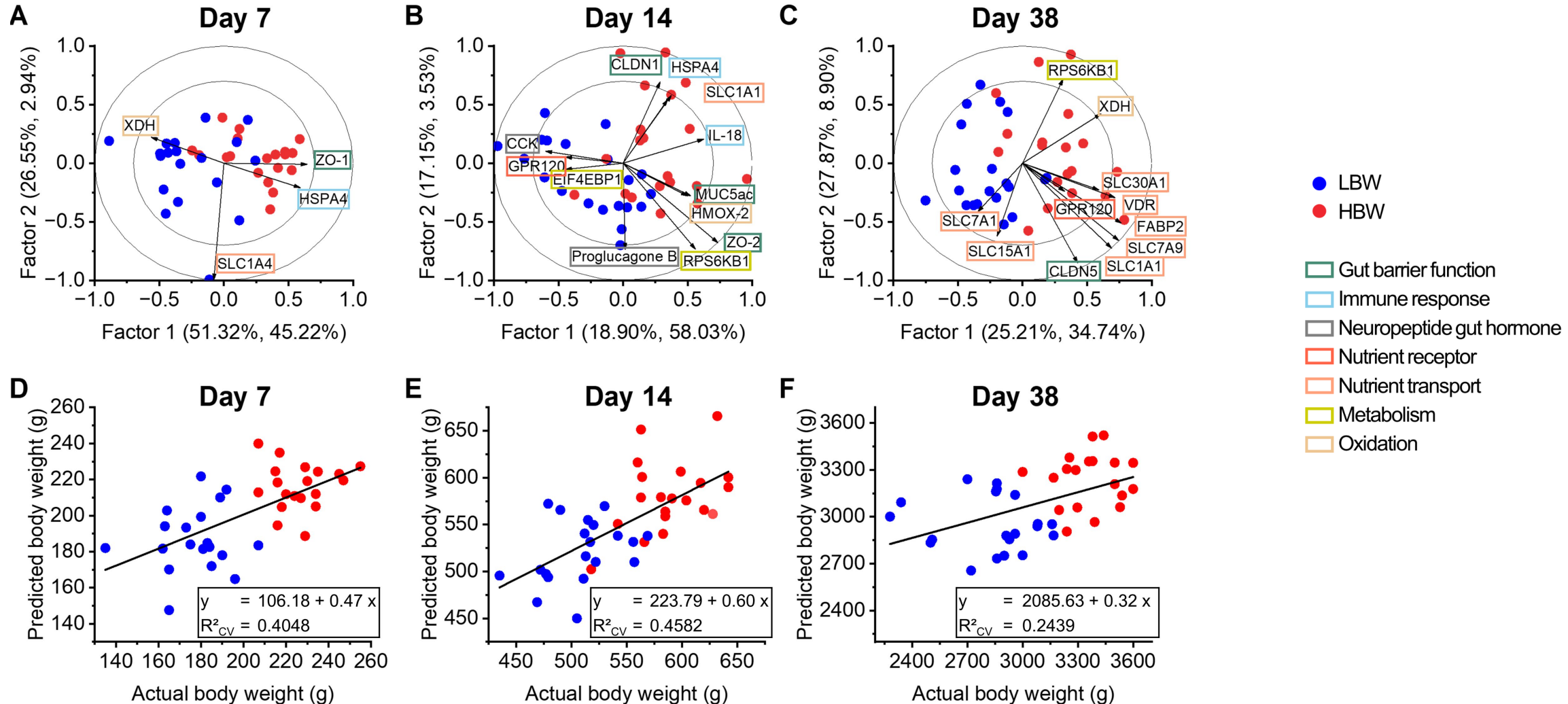
Student's t-test, $P < 0.05$



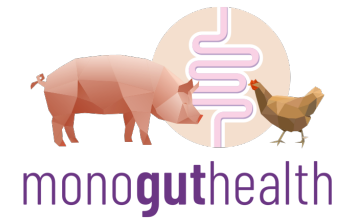
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4. Ileal gene expression – Partial least square regression



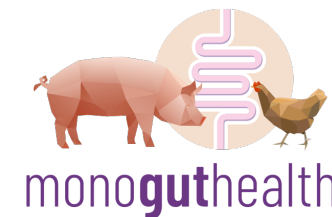
Conclusion



- Broiler chickens within a flock shows **divergence growth rates** despite uniform rearing conditions.
- **Differential gene expression** was observed, mainly related to nutrient transport, absorption and oxidation, and intestinal barrier function.
- **Feed intake** seemed to be controlled by **gut hormones**, which led to differences in weight gain.
- Variations in the **caecal microbiota composition and functionality** were observed starting from 7 days, coinciding with the divergence of performance.



For further reading



Animal Microbiome

Research | [Open access](#) | Published: 24 July 2024

Assessing the impact of hatching system and body weight on the growth performance, caecal short-chain fatty acids, and microbiota composition and functionality in broilers

[Muhammad Zeeshan Akram](#), [Ester Arévalo Sureda](#), [Luke Comer](#), [Matthias Corion](#) & [Nadia Everaert](#)

Animal Microbiome **6**, Article number: 41 (2024) | [Cite this article](#)

546 Accesses | 1 Citations | [Metrics](#)



Poultry Science

Available online 2 August 2024, 104158

In Press, Journal Pre-proof [What's this?](#)



Linking gastrointestinal tract structure, function, and gene expression signatures to growth variability in broilers: a novel interpretation for flock uniformity

[Muhammad Zeeshan Akram](#) * , [Ester Arévalo Sureda](#) * , [Matthias Corion](#) * ,
[Luke Comer](#) * , [Nadia Everaert](#) *

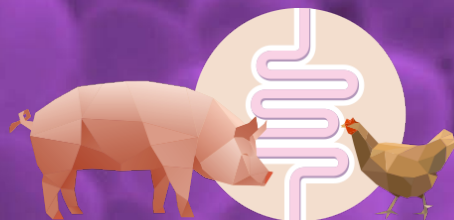
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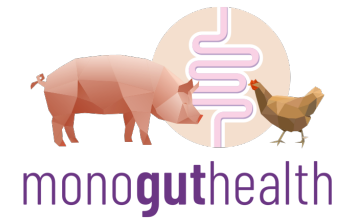
Optimal gut function in monogastric livestock

02

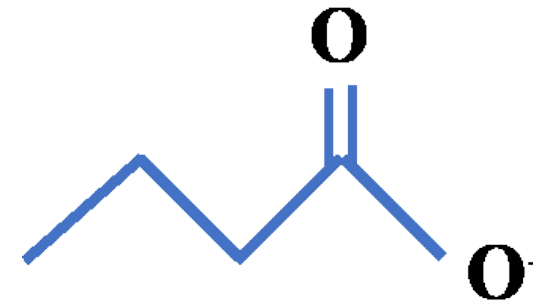
Study 2: Effects of in ovo sodium butyrate (SB) injection on growth performance, intestinal health and gut microbiota of chickens with varying hatch weight (HW)



Rationale for choosing sodium butyrate (SB)?



- Butyrate, a **major SCFA** produced by bacterial fermentation of undigested carbohydrate in the gut.
- It acts as energy source for enterocytes, improve **gut barrier function**.
- It enhance **mucin production**, which might modulate gut microbiota.
- It has been shown to increase **digestive and absorptive functions** of the intestine.
- It modulates immune function, improves intestinal health by reducing the intestinal **inflammatory response**.

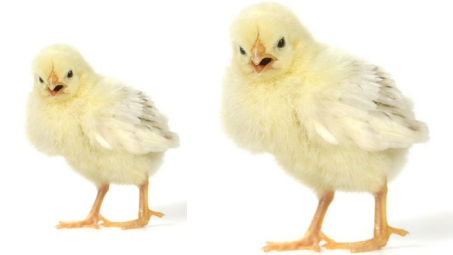
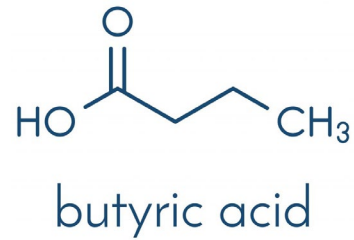


Butyrate

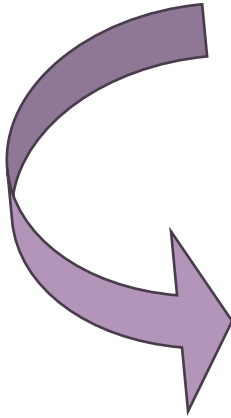
Based on <https://doi.org/10.1016/j.psj.2024.104108>



Objectives and Aim



**Phenotype of
chickens**



To examine the effect of in-ovo SB injection on growth performance, intestinal development and gut microbiota of chickens with different hatch weight (HW)

Trial design

4 x 2 factorial design

Control: normal saline (n=300)

LHW-control
HHW-control

SB1: 0.1% sodium butyrate (n=300)

LHW-SB1
HHW-SB1

SB3: 0.3% sodium butyrate (n=300)

LHW-SB3
HHW-SB3

SB5: 0.5% sodium butyrate (n=300)

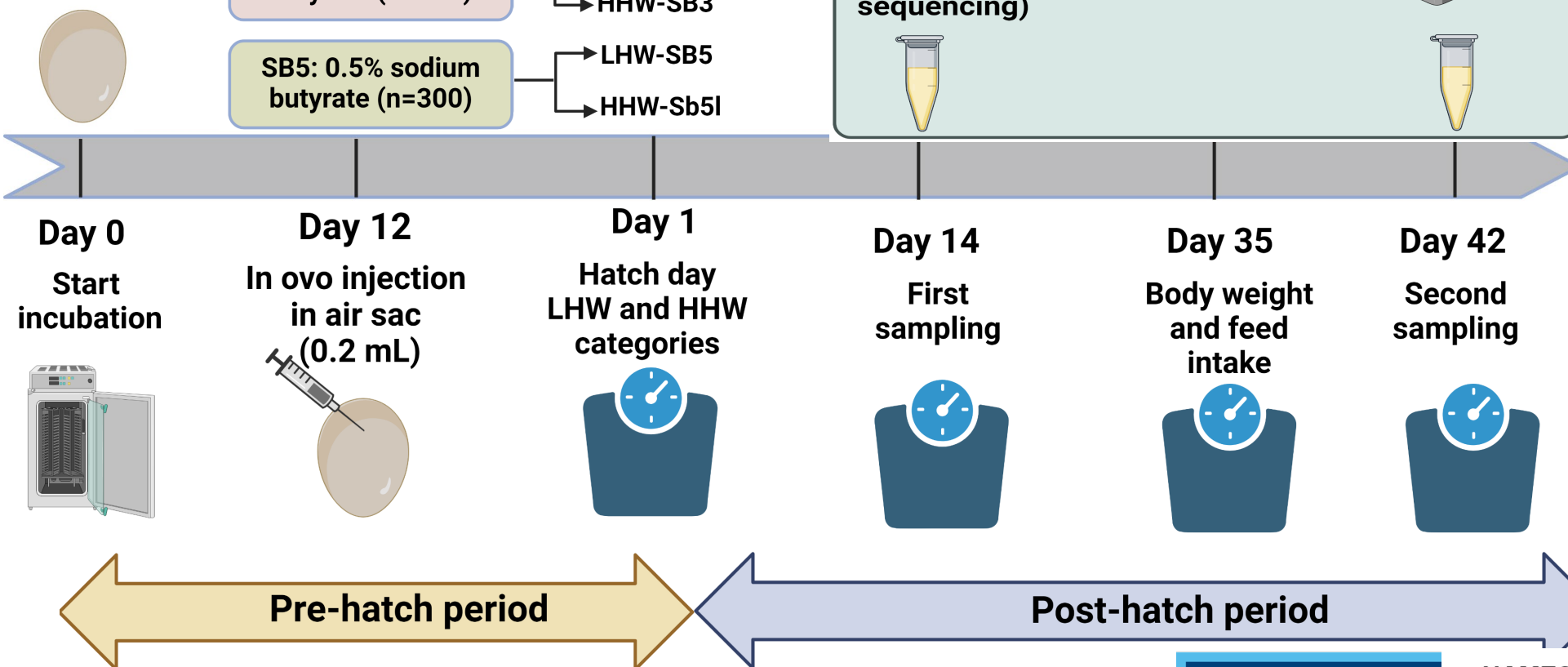
LHW-SB5
HHW-Sb5l

Analyses (n = 6 birds/group)

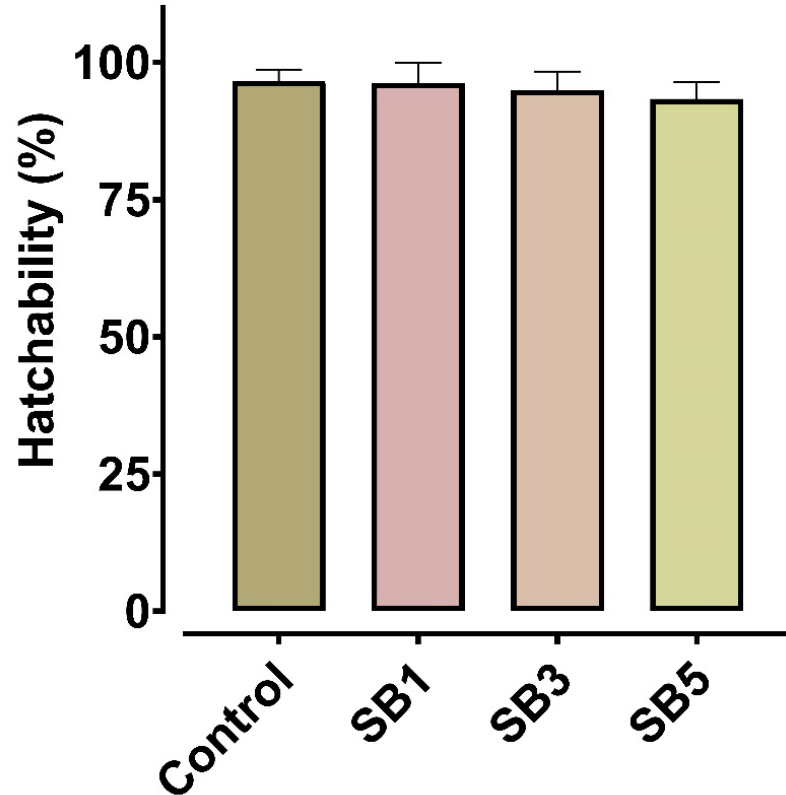
Relative intestinal weight and length

Expression of gene related to gut barrier function and immune response (qPCR)

Caecal microbiota feature including predicted metabolic pathways (16S sequencing)



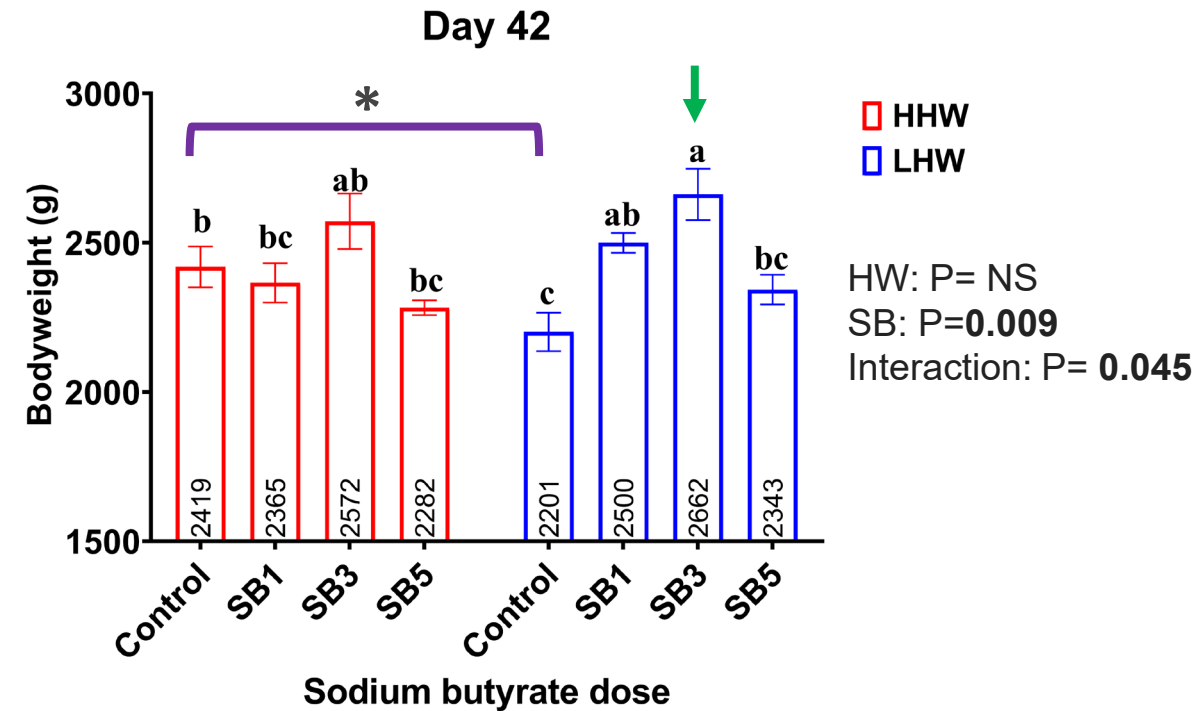
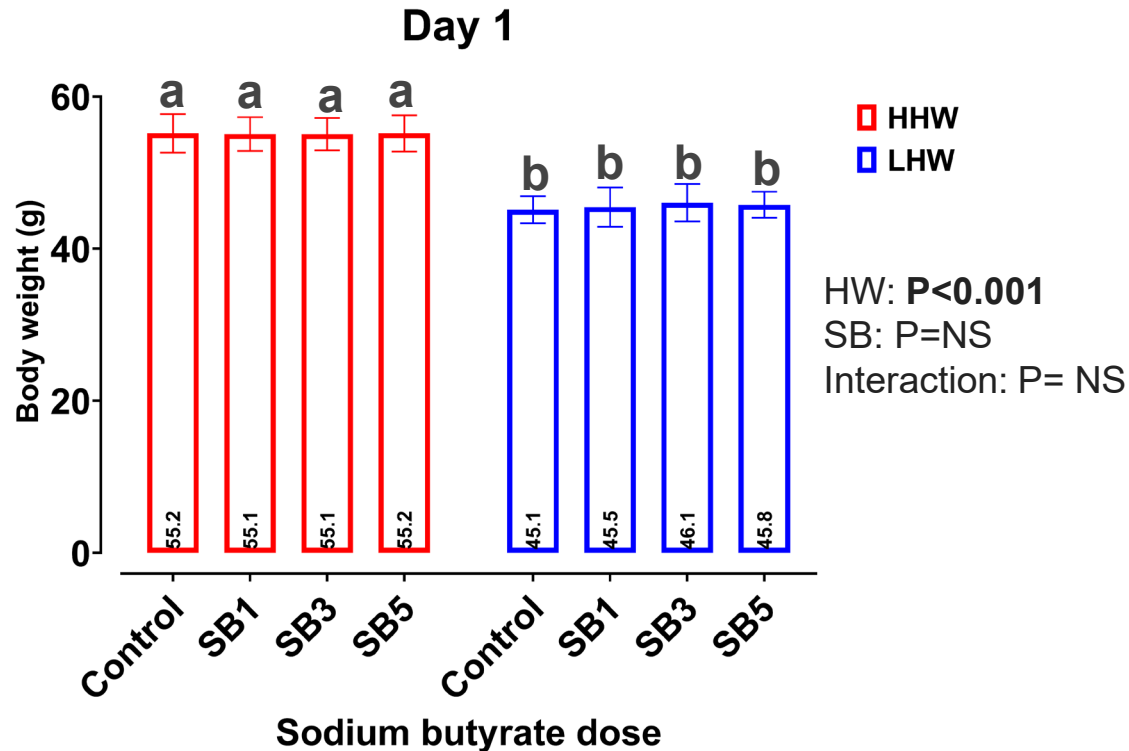
SB effects on embryonic development



- No negative effects on **embryonic development**
- No effect on **hatchability**

Control: 0% SB, **SB1:** 0.1% SB, **SB3:** 0.3% SB, **SB5:** 0.5% SB

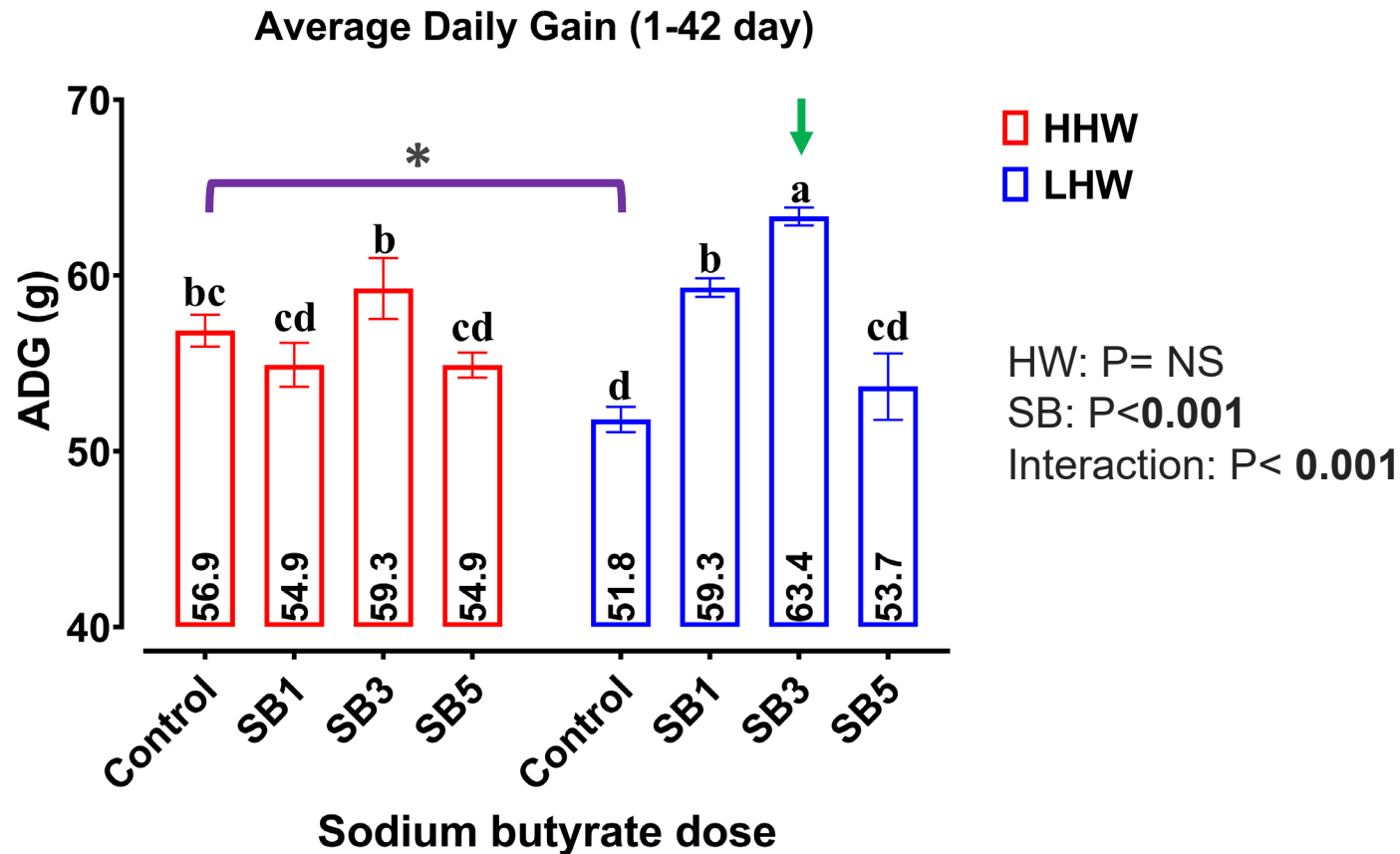
2. Body weight



abc: TukeyHSD, $P < 0.05$

Control: 0% SB, **SB1:** 0.1% SB, **SB3:** 0.3% SB, **SB5:** 0.5% SB

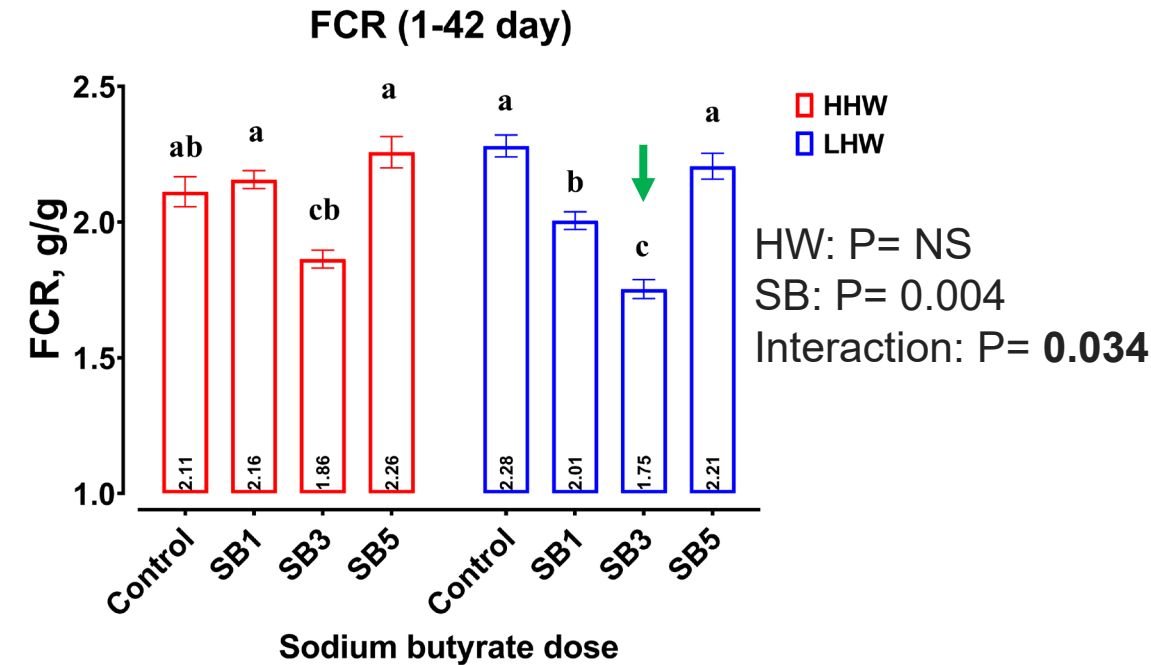
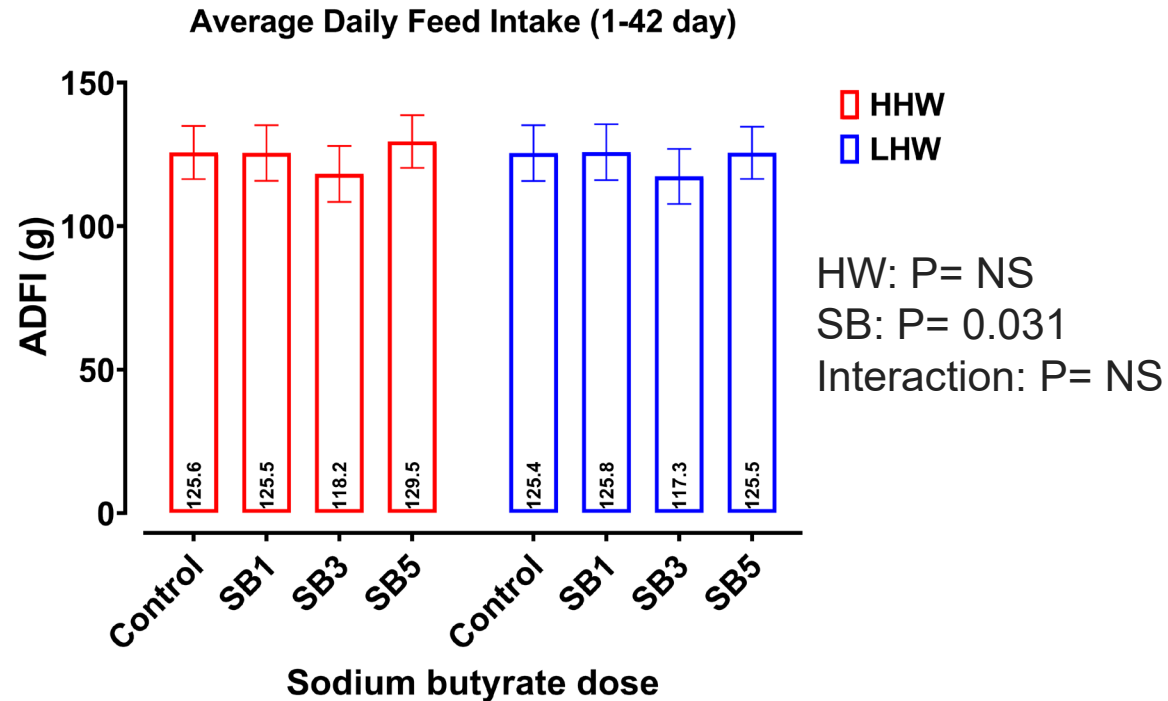
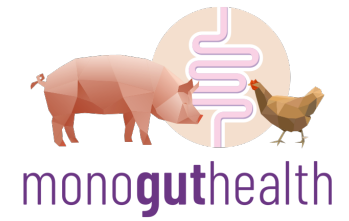
2. Average daily gain



abc: TukeyHSD, P<0.05

Control: 0% SB, **SB1:** 0.1% SB, **SB3:** 0.3% SB, **SB5:** 0.5% SB

2. Feed intake and Feed conversion ratio



abc: TukeyHSD, $P < 0.05$

Control: 0% SB, **SB1:** 0.1% SB, **SB3:** 0.3% SB, **SB5:** 0.5% SB

3. Ileal gene expression on day 14

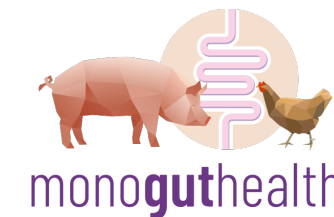
Genes	HHW				LHW				P-value
	Control	SB1	SB3	SB5	Control	SB1	SB3	SB5	HW x SB
CLDN1	1.00 ^c	2.40 ^{ab}	2.56 ^{ab}	2.61 ^{ab}	0.69 ^d	2.49 ^{ab}	3.31 ^a	1.24 ^c	0.016
TJP1	1.00 ^b	2.06 ^{ab}	1.69 ^{ab}	2.03 ^{ab}	1.90 ^{ab}	2.03 ^{ab}	2.56 ^a	1.56 ^{ab}	0.028
MUC6	1.00	3.05	3.09	3.61	2.15	3.29	3.58	4.76	0.633
IL-10	1.00 ^c	2.38 ^{ab}	2.44 ^{ab}	2.52 ^{ab}	2.51 ^{ab}	2.53 ^{ab}	3.01 ^a	2.10 ^b	0.048
IL-1B	1.00	1.55	1.95	0.99	1.71	2.07	1.78	1.19	0.658
IL-12P40	1.00	1.72	1.56	1.51	2.06	1.91	1.46	1.30	0.205

abcd: TukeyHSD, P<0.05

CLDN1: claudin1, **TJP1**, tight junction protein1, **MUC6**: mucin 6

IL-10: interleukin 10, **IL1B**: interleukin 1-beta, **IL-12P40**: interleukin 12 subunit 40

3. Ileal gene expression on day 42



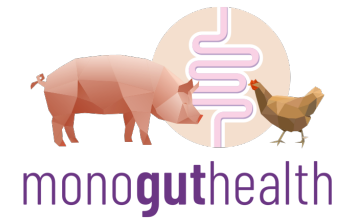
Genes	HHW				LHW				P-value
	Control	SB1	SB3	SB5	Control	SB1	SB3	SB5	HW x SB
CLDN1	1.00 ^{ab}	0.60 ^{bc}	0.30 ^c	0.52 ^{bc}	1.01 ^a	1.27 ^a	0.52 ^{bc}	0.23 ^c	0.002
TJP1	1.00	0.77	0.64	0.46	1.60	1.12	0.79	0.32	0.180
MUC6	1.00 ^{ab}	0.41 ^b	0.40 ^b	0.36 ^b	1.03 ^{ab}	0.51 ^b	1.42 ^a	0.14 ^c	0.024
IL-10	1.00	0.46	1.18	0.53	0.86	0.93	0.41	0.58	0.207
IL-1B	1.00	1.03	0.57	0.93	0.70	1.0	0.58	0.28	0.227
IL-12P40	1.00	0.55	0.96	1.01	0.92	1.27	0.80	0.48	0.109

abc: TukeyHSD, $P < 0.05$

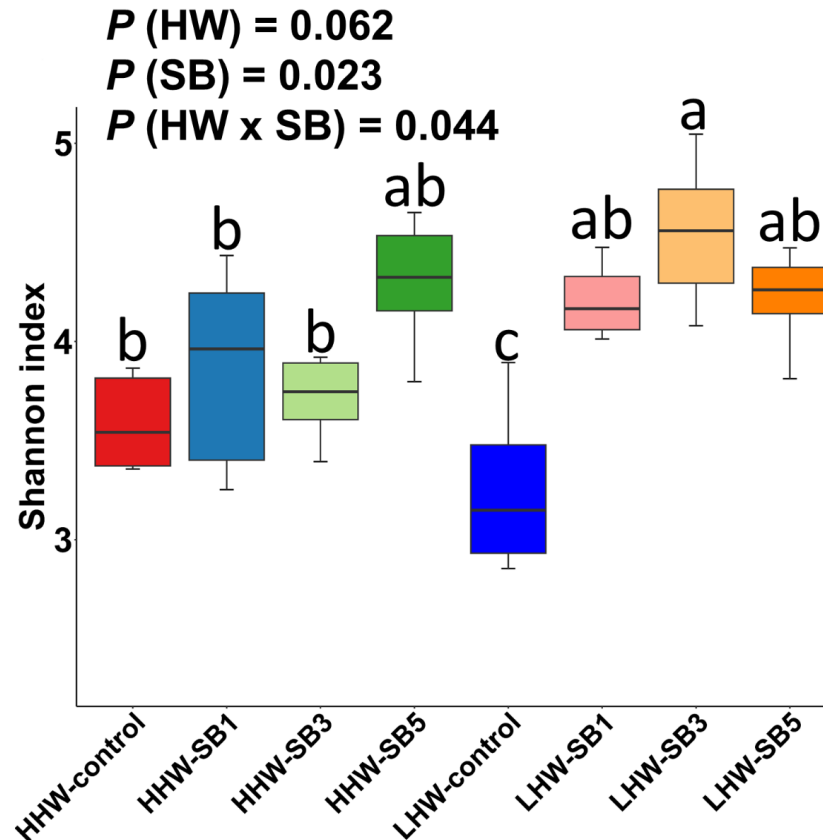
CLDN1: claudin1, **TJP1**, tight junction protein1, **MUC6**: mucin 6

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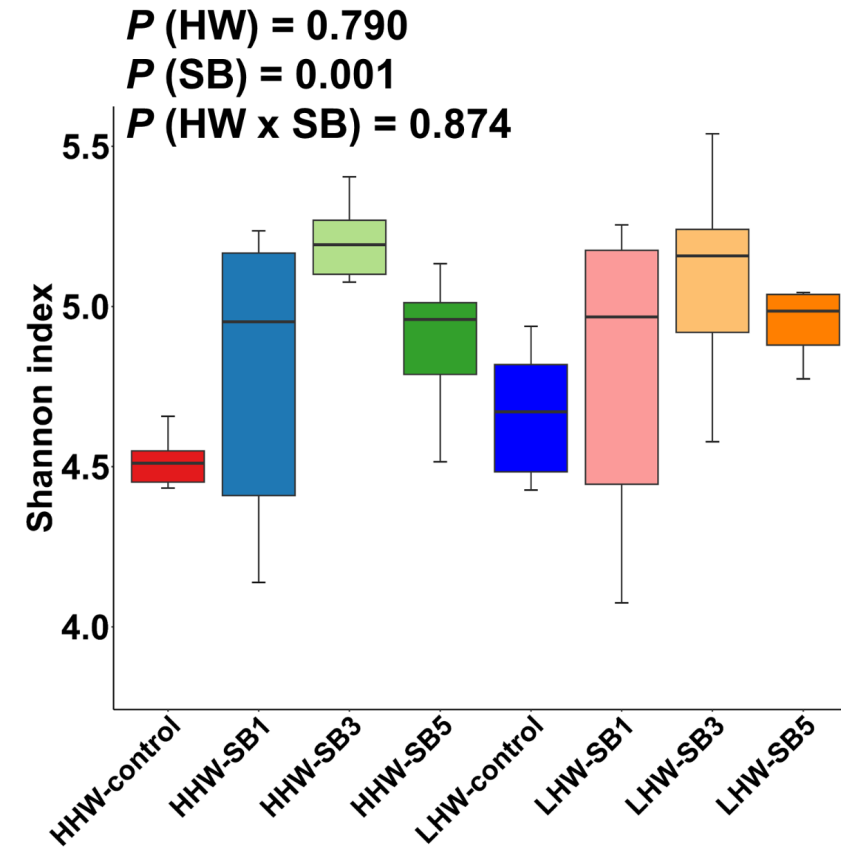
4. Caecal microbiota alpha diversity



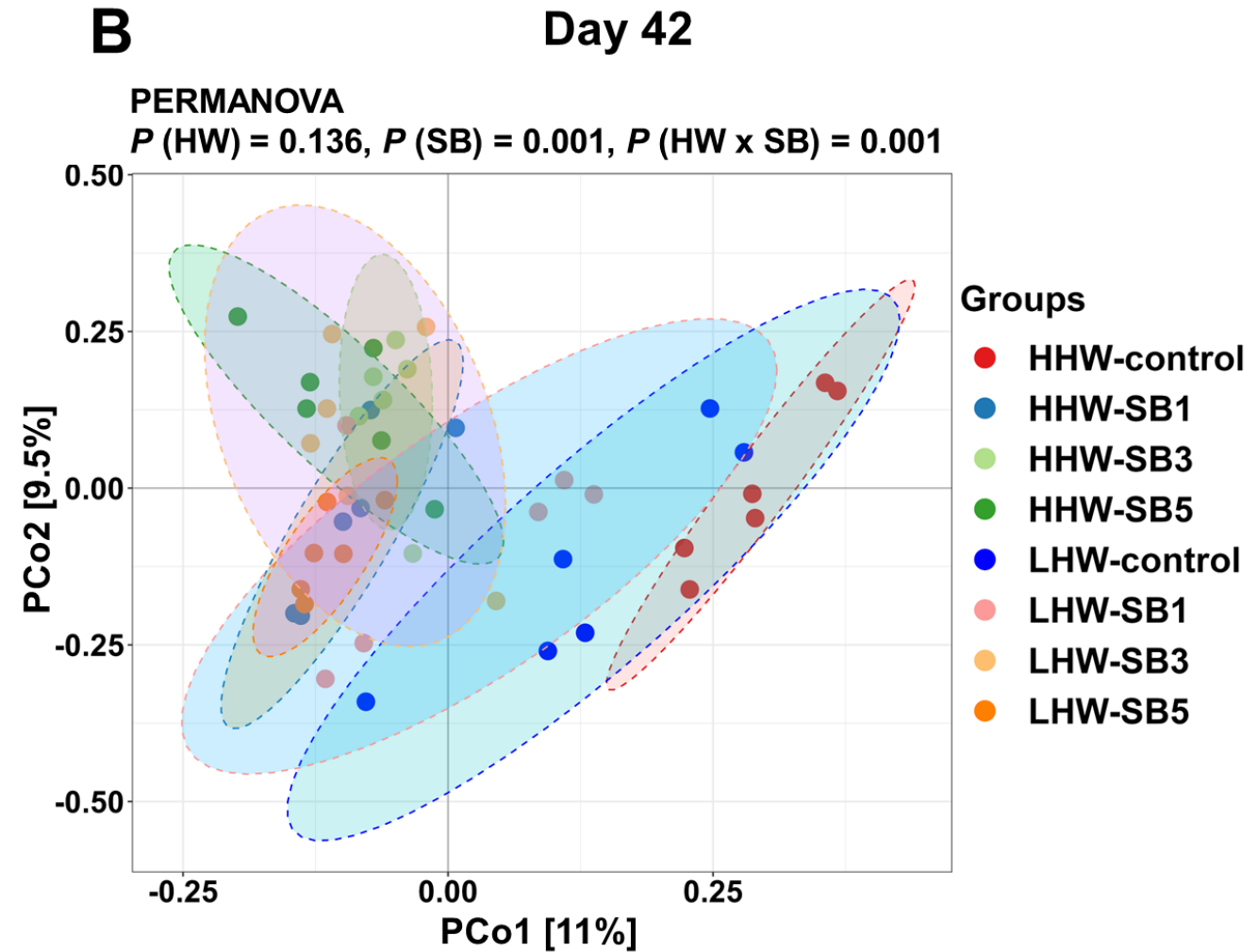
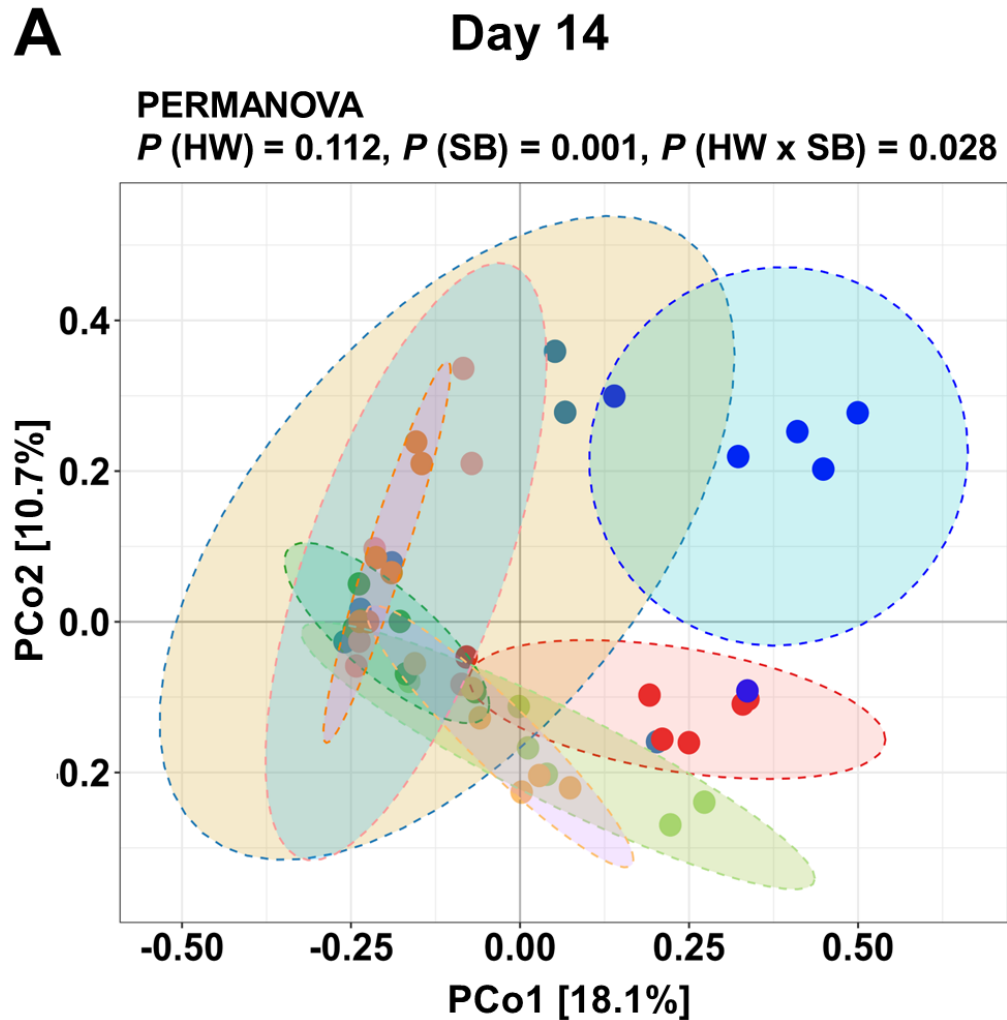
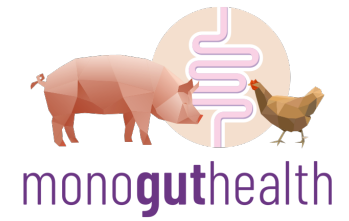
Day 14



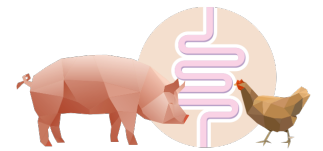
Day 42



4. Caecal microbiota beta diversity

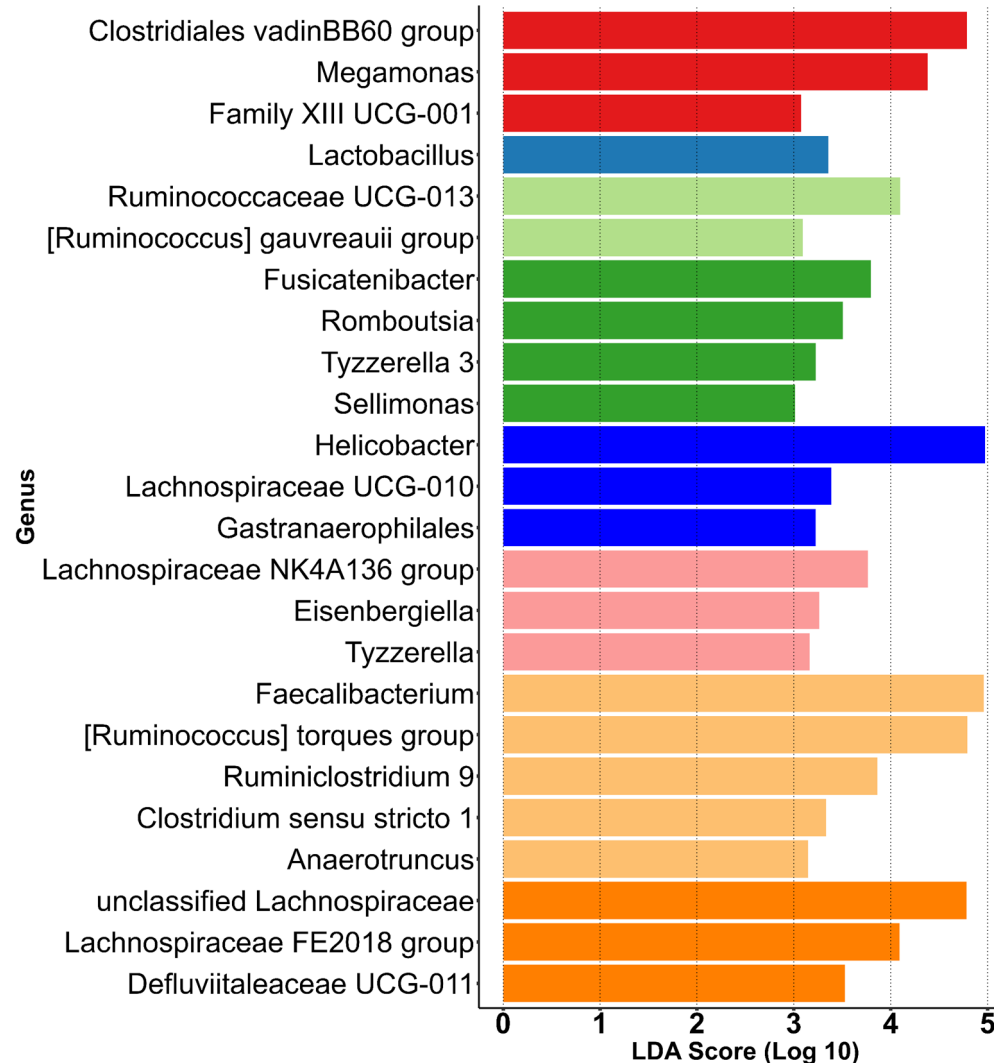


4. Differential abundance of bacterial genera



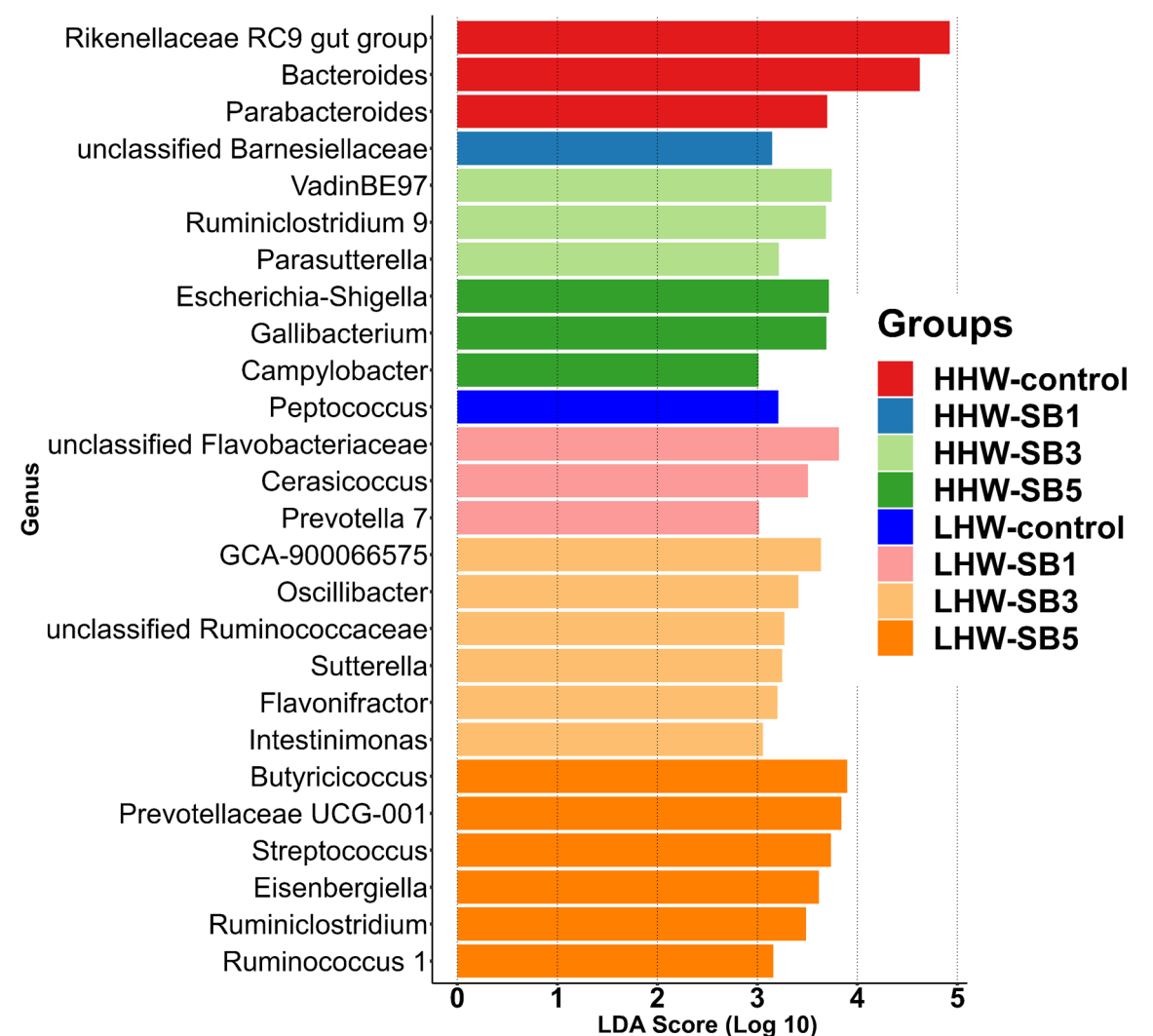
A

Day 14

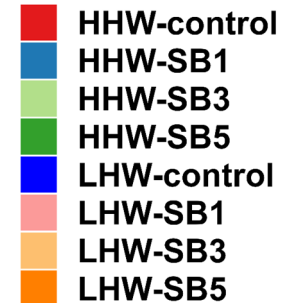


B

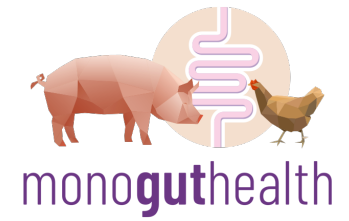
Day 42



Groups

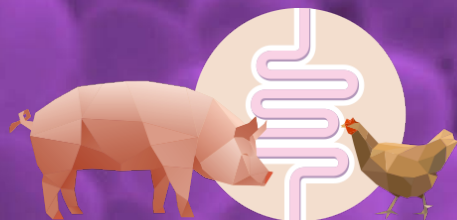


Conclusion



- **Hatch-weight** is an important factor influencing subsequent growth performance.
- Effects of SB were **dose-dependent**, with 0.3% significantly enhancing growth, gut barrier function and anti-inflammatory gene expression, and beneficial caecal microbiota.
- In ovo SB injection **enhances growth** in both HW categories, with **strong effects in LHW at 0.3% inclusion**.
- The findings offer **targeted intervention** for improving the growth performance of LHW chicks, leading to more uniform flocks and better production efficiency.





monoguthealth

Optimal gut function in monogastric livestock

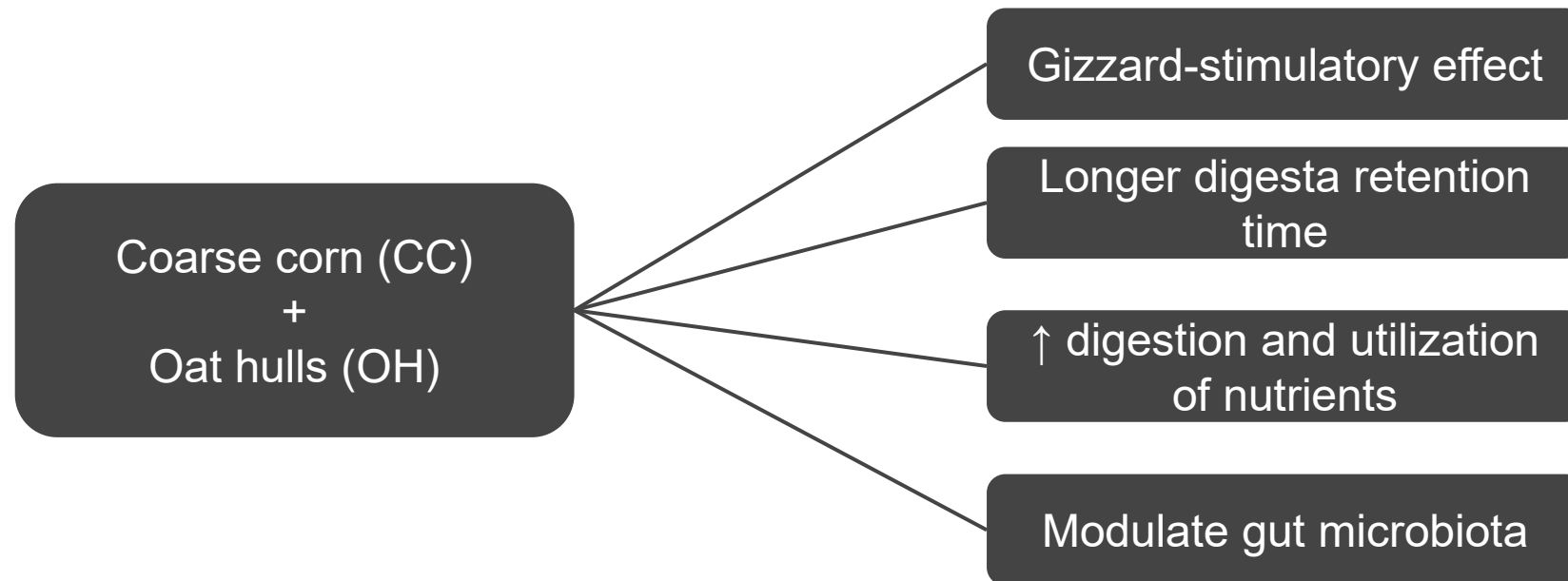
03

Study 3: Effects of coarse corn, oat hulls and their combination on growth performance, intestinal functions and microbiota of underperforming chickens



Selection of coarse corn and oat hulls?

- Chickens may have a requirement for **dietary physical structure** to enhance gut development and health.
- A **well-developed gizzard and GIT** have been shown to improve nutrient digestibility and growth performance of chickens



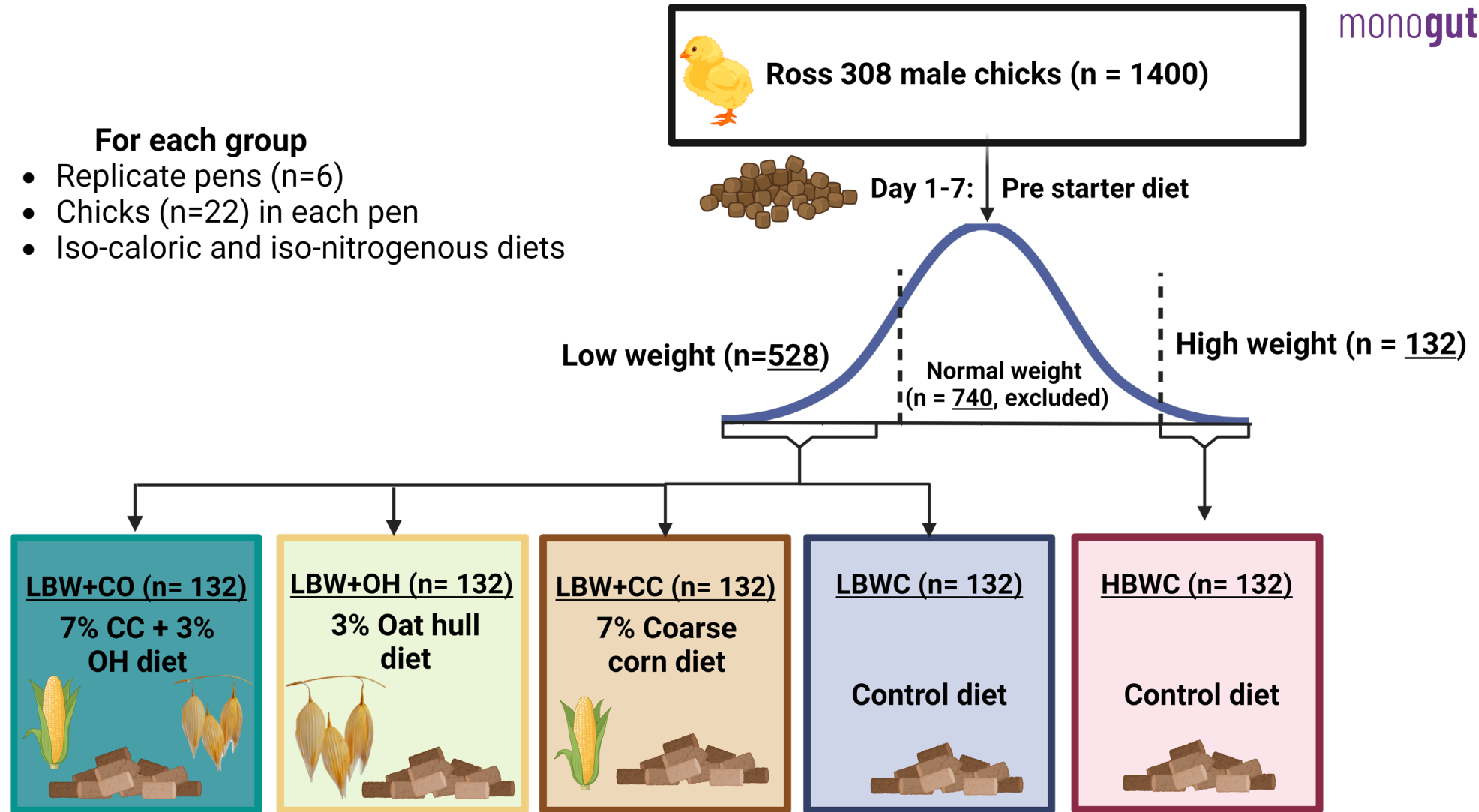
Kheravii et al., 2017; Li et al., 2016; Downs et al., 2023

Goal of the study

- To determine the effects of **coarse corn and oat hulls and their combination** on growth performance, GIT development and microbiota of underperforming chickens.
- To evaluate whether **low-weight broiler chickens**, under these dietary treatment, can catch up with their high bodyweight counterparts.

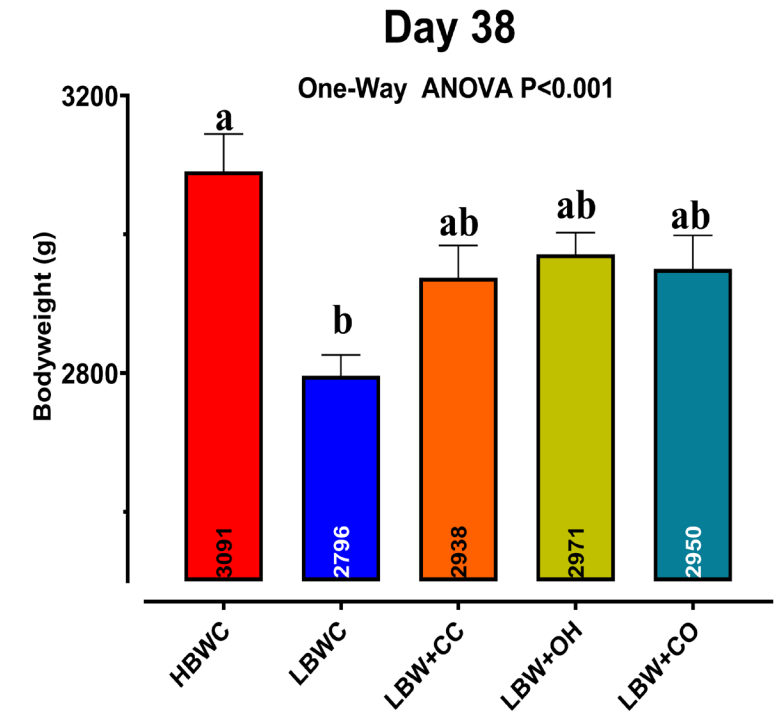
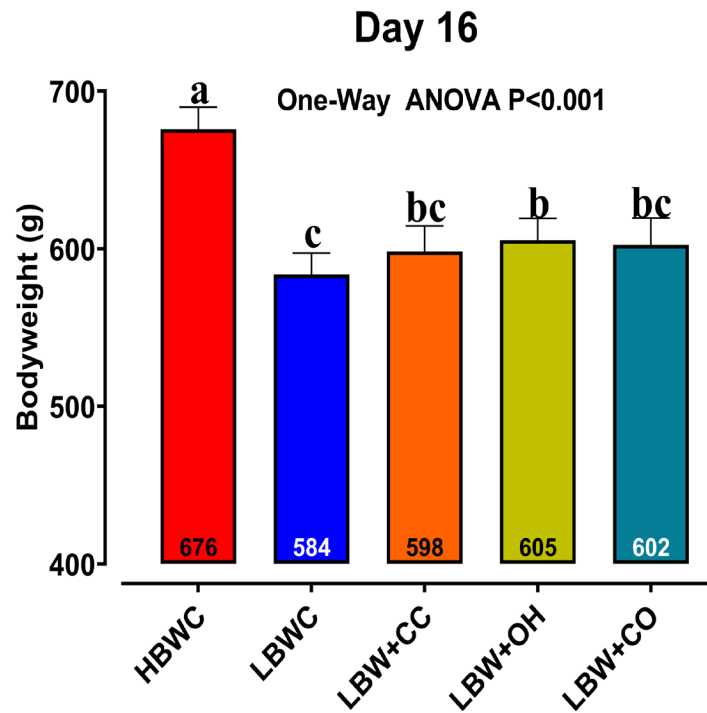
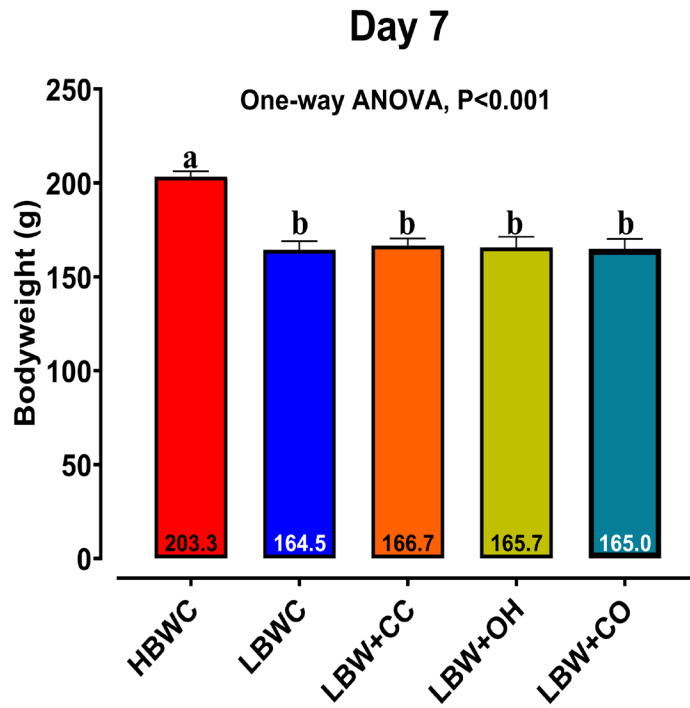
Trial design

- For each group**
- Replicate pens (n=6)
 - Chicks (n=22) in each pen
 - Iso-caloric and iso-nitrogenous diets

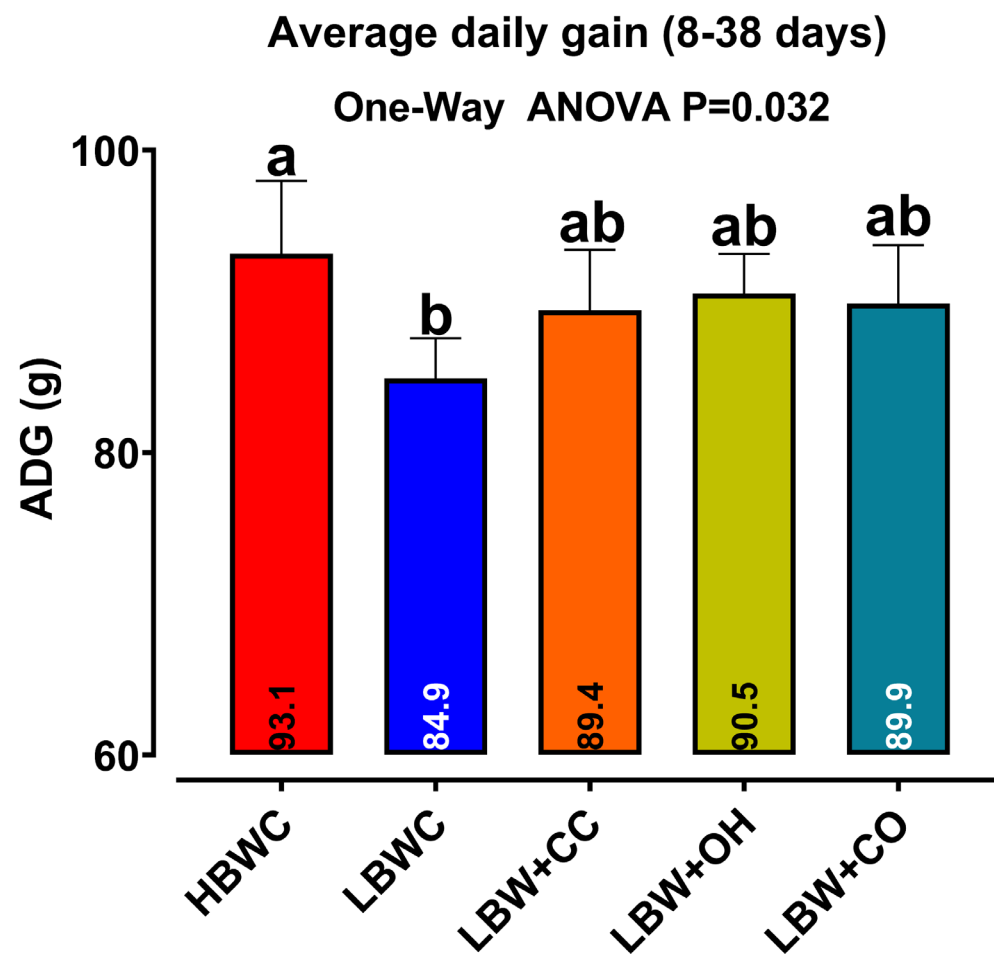




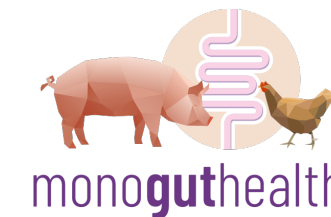
Body weight



Average daily gain

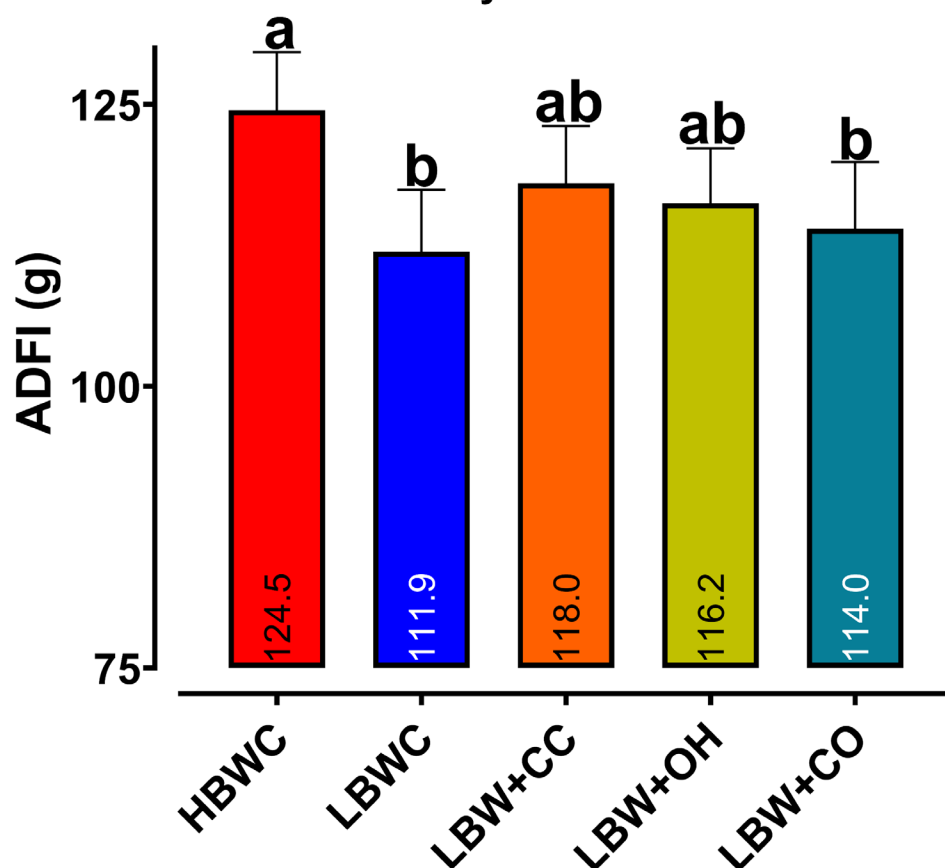


Feed intake and feed conversion ratio

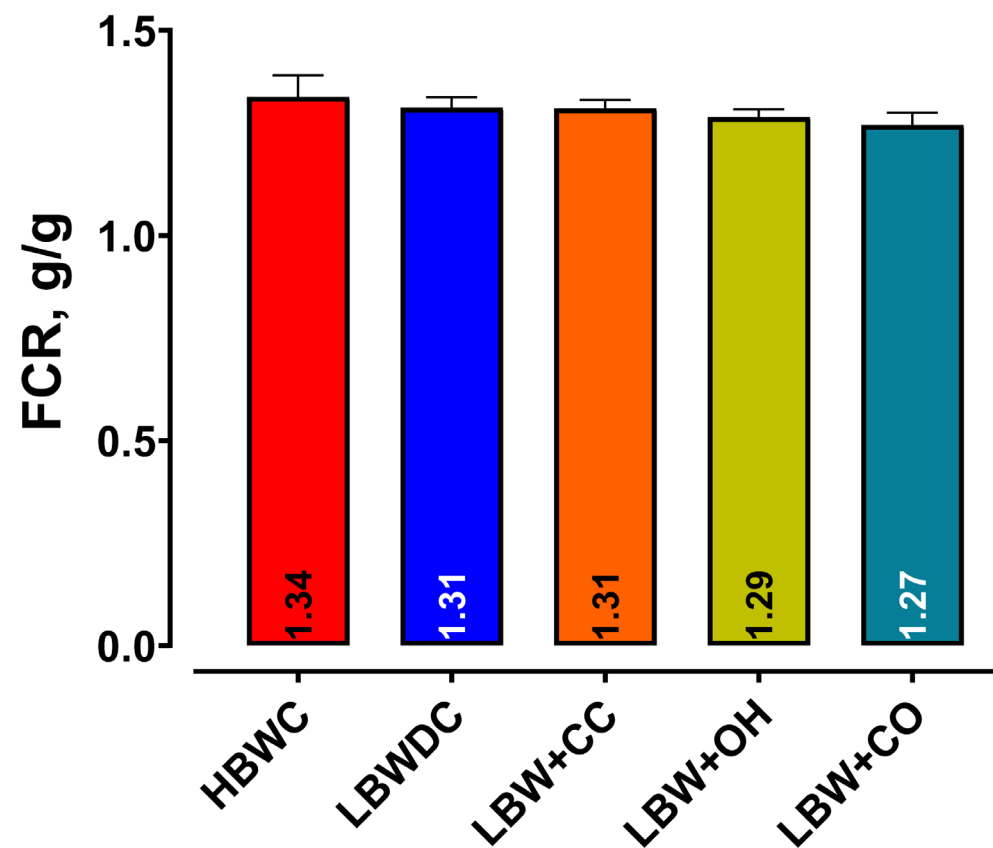


Average daily feed intake (8-38 days)

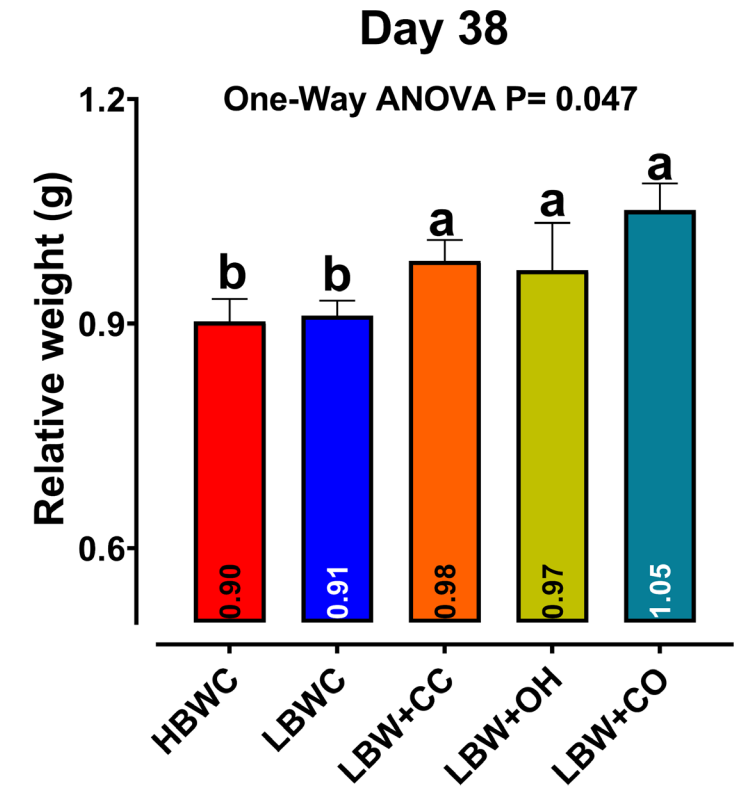
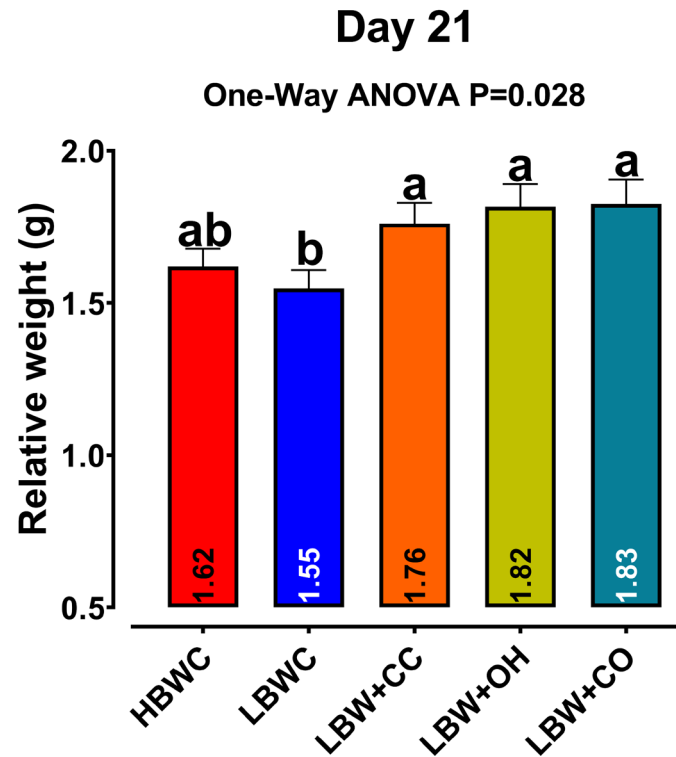
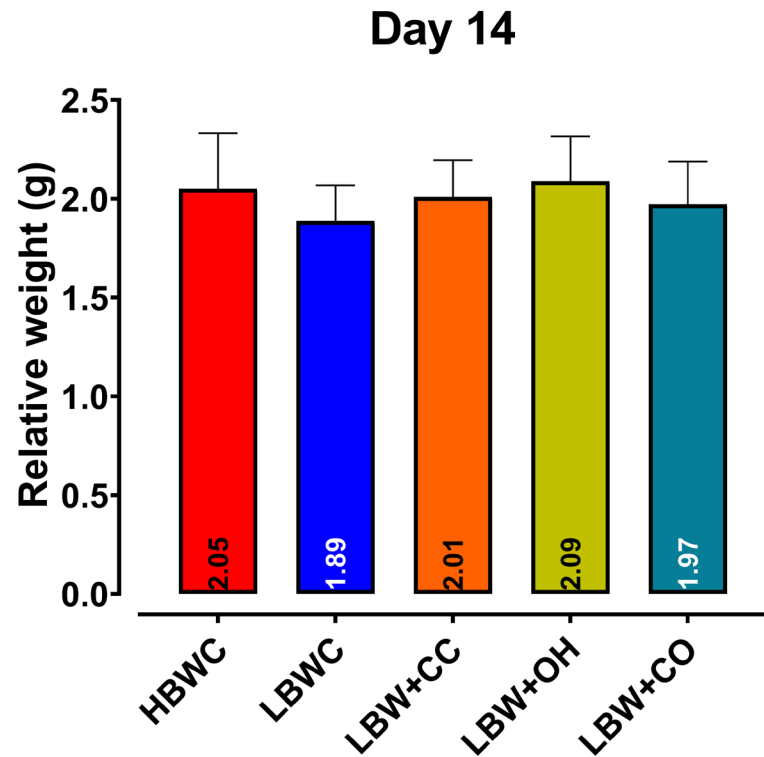
One-Way ANOVA P=0.005



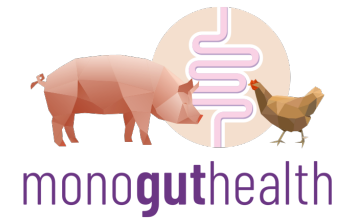
Feed conversion ratio (8-38 days)



Gizzard development



Conclusion thus far



- Coarse corn, oat hulls, and their combinations positively **impact GIT development and enhance the Gizzard size** in LBW chickens.
- The LBW groups on experimental diets showed numerically **better performance parameters** compared to LBWC group, paralleling the performance levels of HBWC group.

< Programme

Wednesday 4 September 2024 – Afternoon, 14:00

Session 98. Nutrition in health and welfare in monogastric animals

Room: Sarda – Palazzo Affari Below ground Floor

Chair: Trocino / Khiaosa-ard

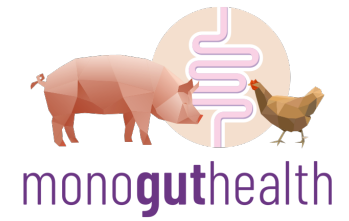
Session Type: Theme session

14:45 Boosting underperforming broilers: feed structure strategies for better gut development and health, and microbiota

M. Z. Akram, E. A. Sureda, N. Everaert



Take home message



- **Gut Microbiota and transcriptomic related factors** cause divergence in growth rates of broilers within a flock.
- In ovo sodium butyrate administration **differentially affects** broiler chickens with different hatch-weights.
- Optimal dose of 0.3% sodium butyrate significantly affects low hatch-weight chickens, suggesting a **targeted strategy to reduce bodyweight variation** among chickens in a flock.
- **Coarse corn and oat hulls** aided in improving gut development and growth performance of low-weight broilers, paralleling to performance of high-weight broilers – suggest a **potential dietary strategy for improving flock uniformity**.



THANK YOU

Do you have any questions?

Nadia Everaert

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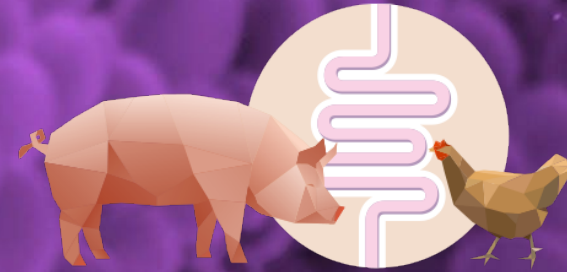
KU LEUVEN

NAMES Lab

Nutrition & Animal
Microbiota EcoSystems



This project has received funding from the European Union's Horizon 2020 research and innovation programme under the Marie Skłodowska-Curie grant agreement No 955374.



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Optimal gut function in monogastric livestock