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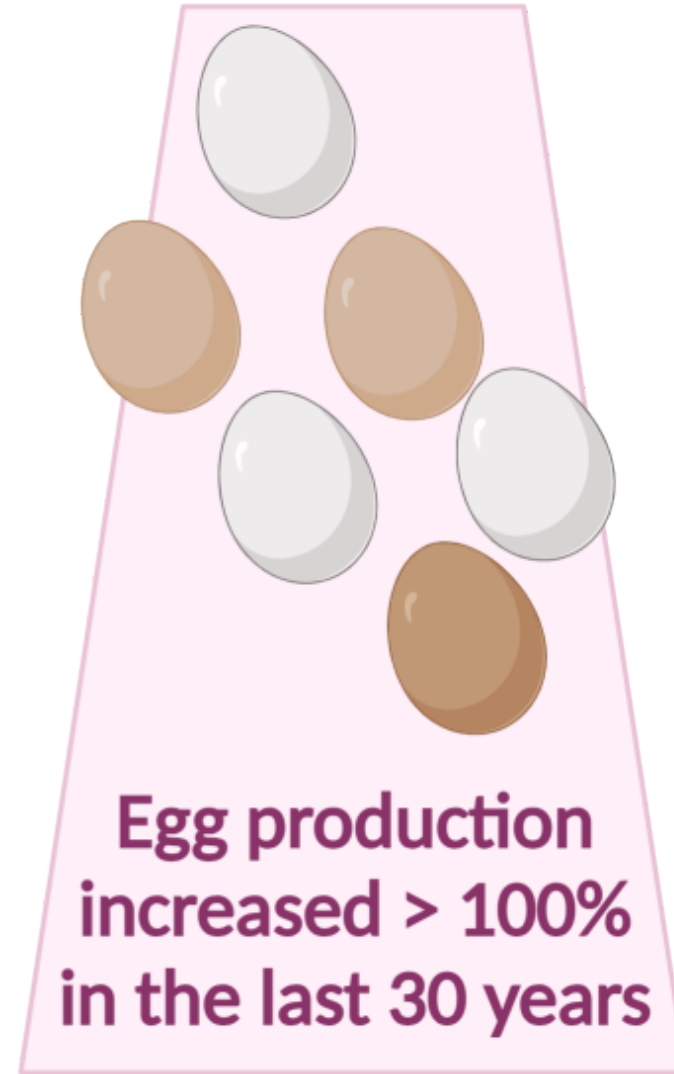
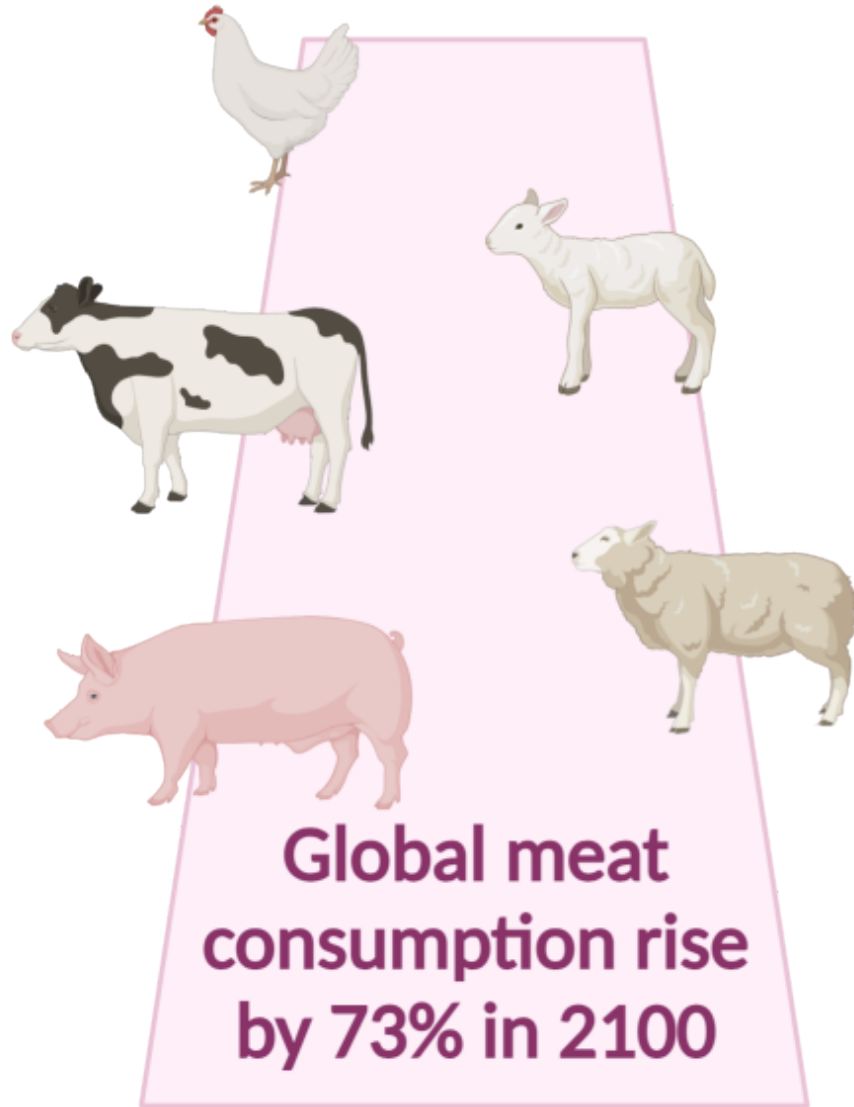


**H o L M i R**  
Hohenheim Center for  
Livestock Microbiome Research

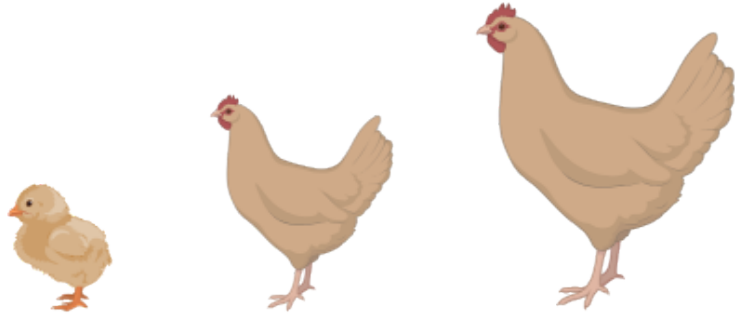
# GUT MICROBIOME VARIATIONS DURING THE PRODUCTIVE LIFESPAN OF TWO HIGH-YIELDING LAYING HEN STRAINS

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# INTRODUCTION

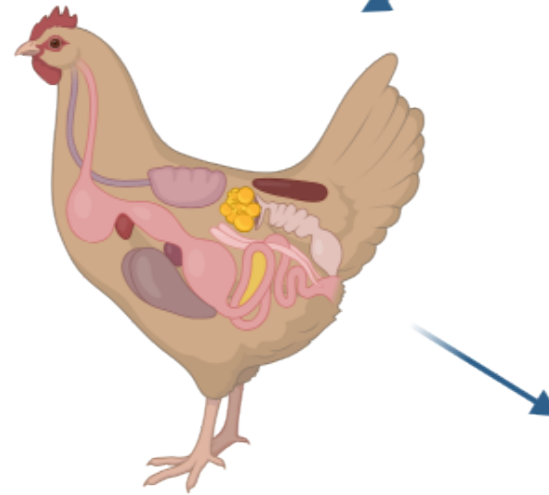


# INTRODUCTION



## LIFESPAN

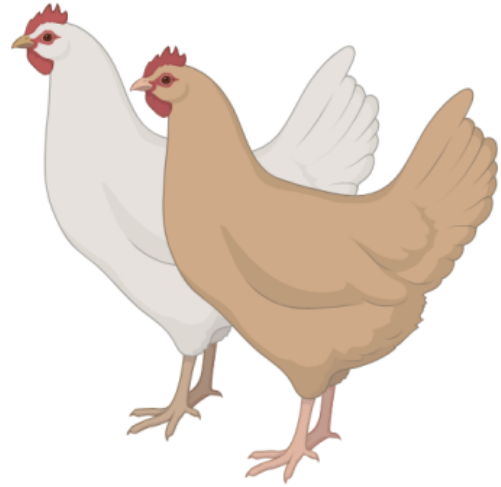
- Physiological changes
- Diet changes (energy and crude protein content, P, Ca)
- Onset of egg production (week 16 and 24)



Phosphorus - limited resource in environment

Animal health and performance

# METHODOLOGY



**LSL**  
(n=50)

**LB**  
(n=50)



Crop  
Gizzard  
Duodenum  
Ileum  
Caeca

**Weeks 10, 16, 24, 30, 60**



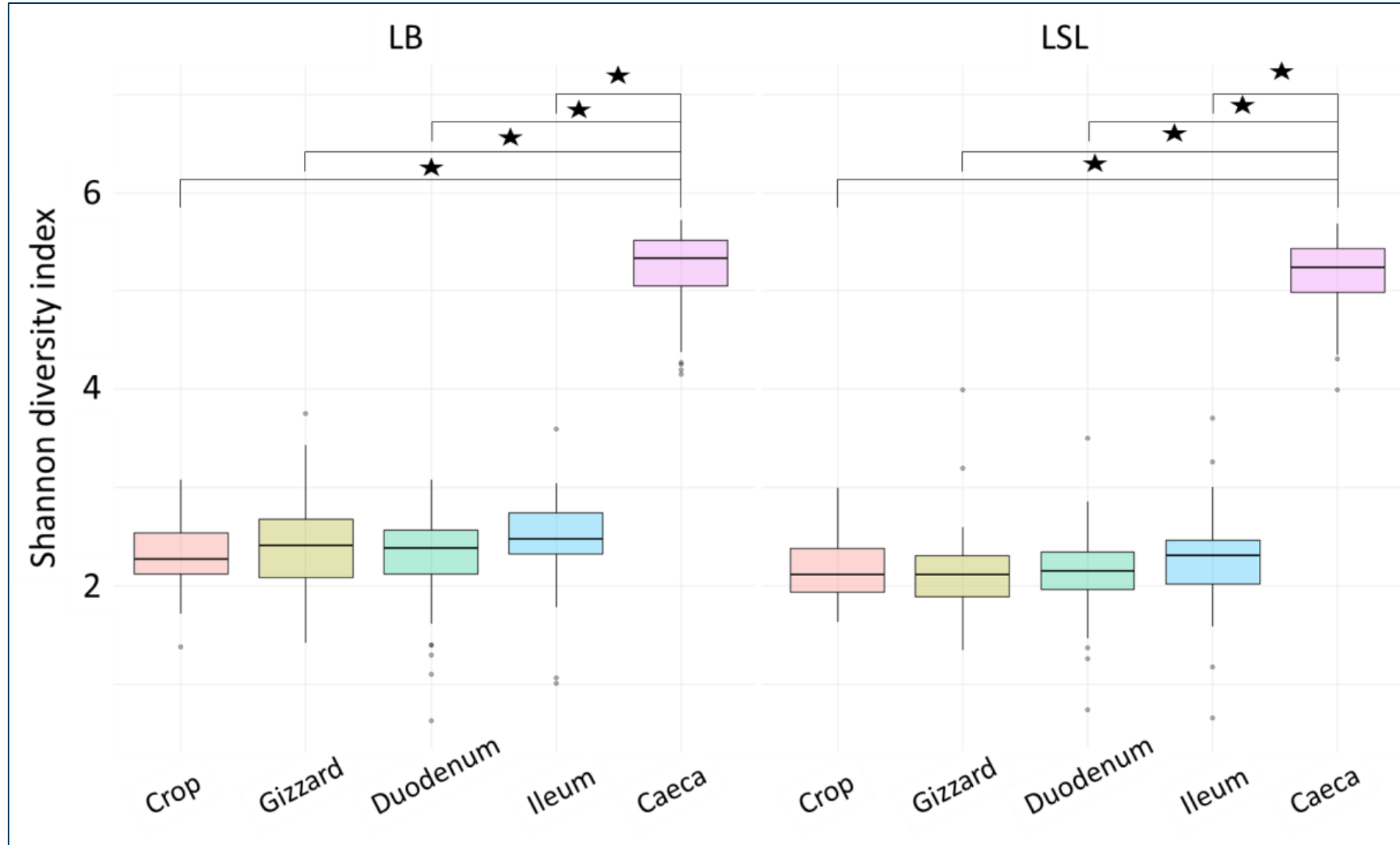
**Target  
amplicon  
sequencing**  
(n=500)

**Shotgun  
metagenomics**  
(n=120, weeks  
16 & 24)

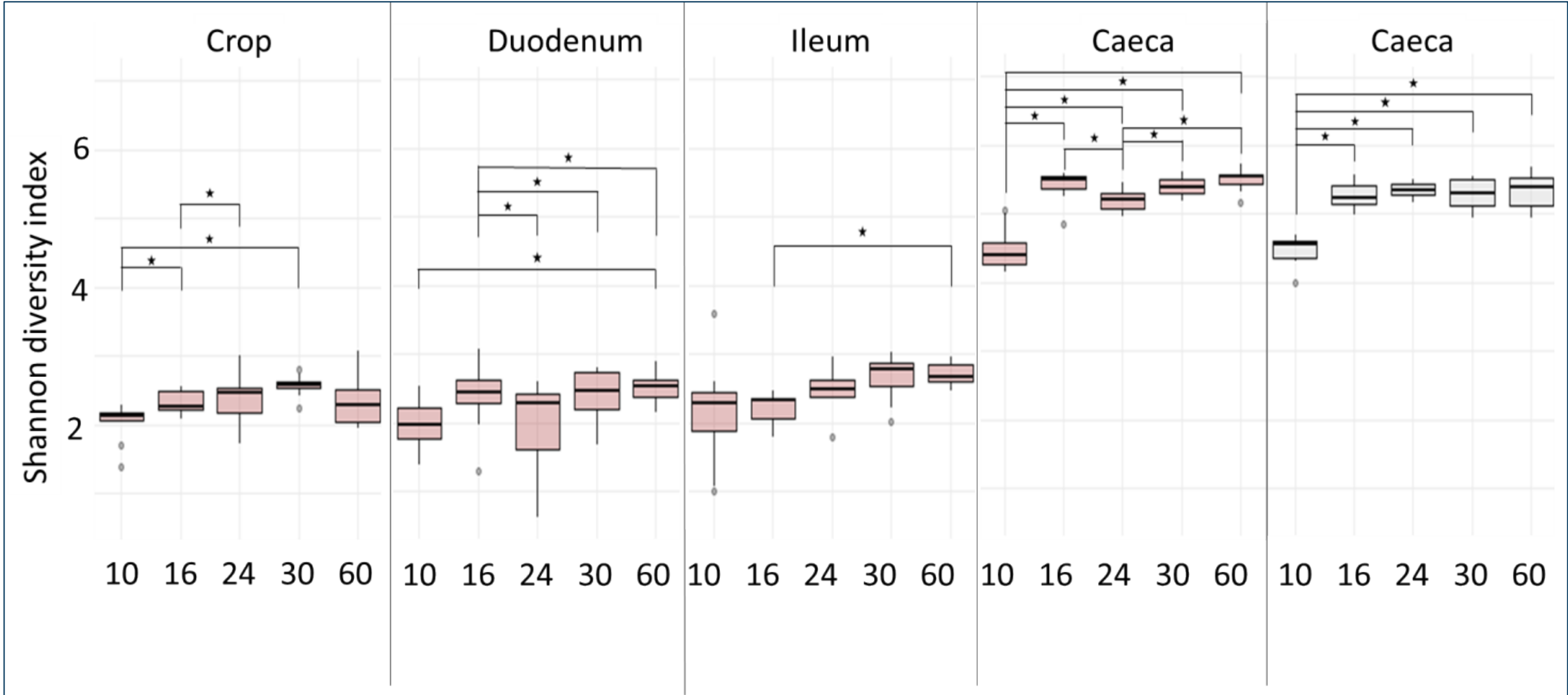
Lohmann Brown-Classic (LB)  
Lohmann LSL Classic (LSL)

# RESULTS

Significant differences for the factors **breed**, **age** and **GIT section**



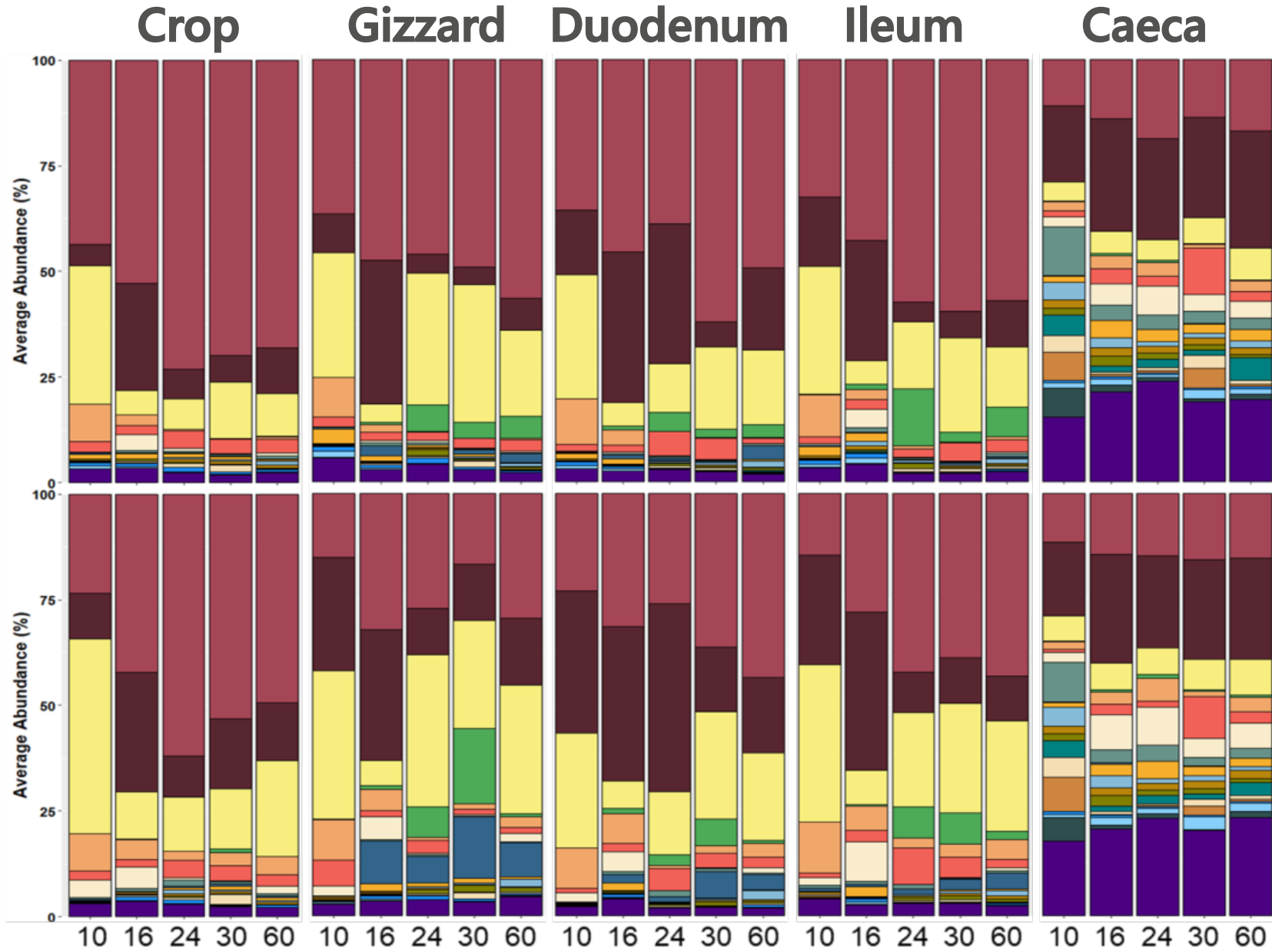
# RESULTS



# RESULTS

LSL

LB



## Genus

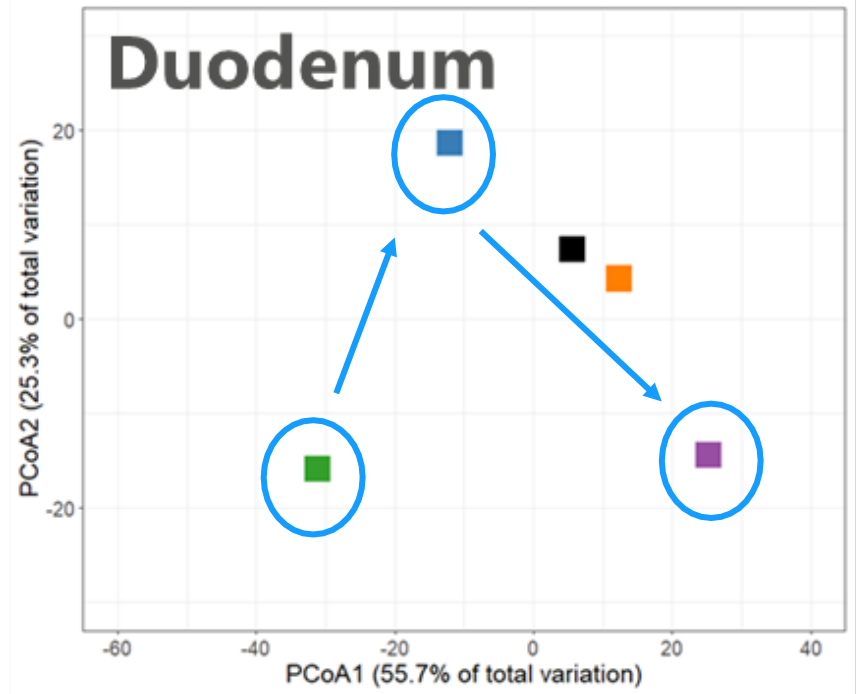
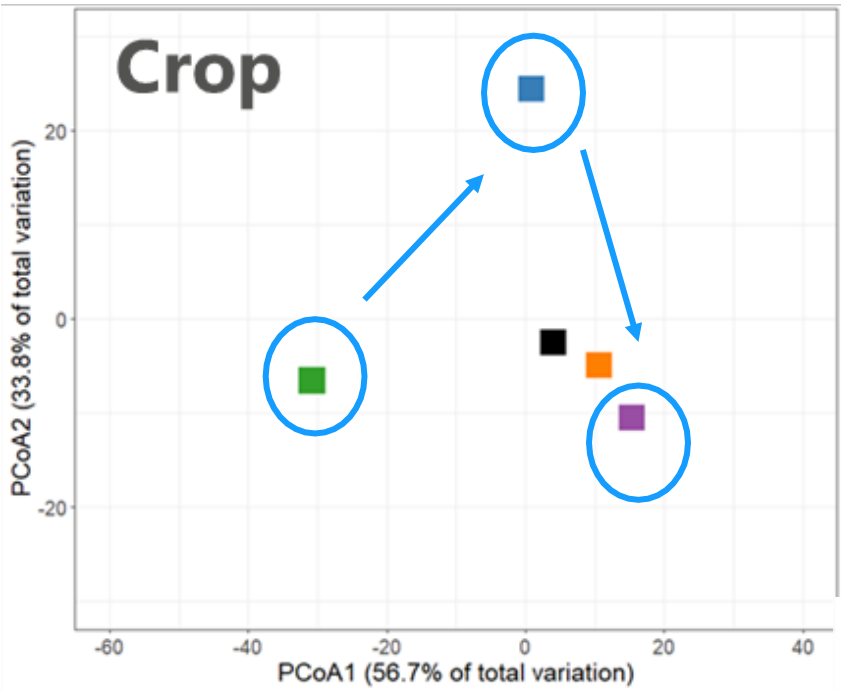
- *Ligilactobacillus*
- *Lactobacillus*
- *uncl. Lachnospiraceae*
- *Blautia*
- *Romboutsia*
- *uncl. Bacteroidales*
- *uncl. Ruminococcaceae*
- *uncl. Clostridiales*
- *Eisenbergiella*
- *uncl. Bacteria*
- *uncl. Bacteroidetes*
- *uncl. Lactobacillaceae*
- *Faecalibacterium*
- *Parabacteroides*
- *Mucispirillum*
- *Megamonas*
- *uncl. Bacteroidaceae*
- *Phocaeicola*
- *Enterococcus*
- *Others*

↓ *Lactobacillus* wk 16-24  
in crop gizzard and ileum

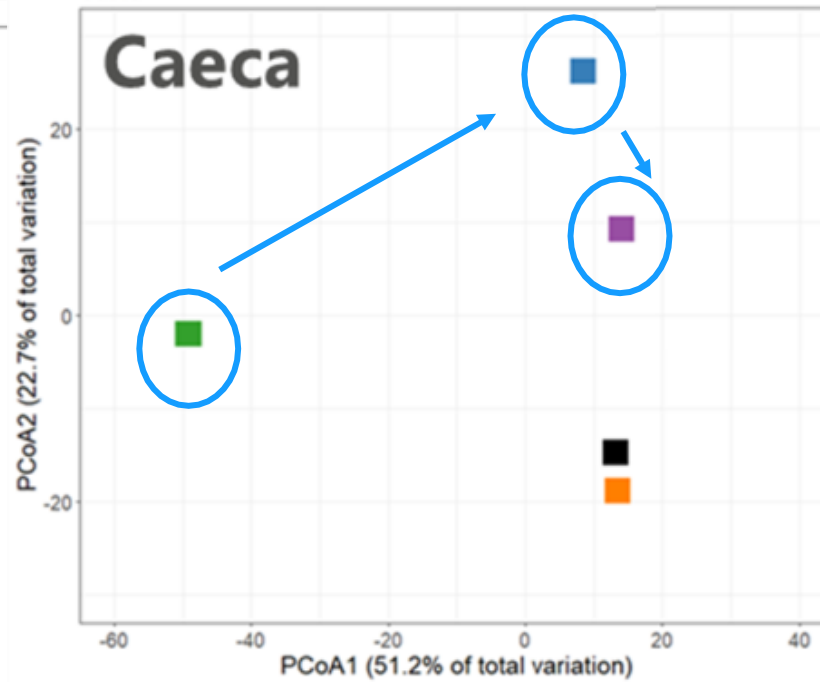
↑ *Blautia* wk 24 in SI

↓ *Romboutsia* after wk 10

# RESULTS



- Week 10 ■
- Week 16 ■
- Week 24 ■
- Week 30 ■
- Week 60 ■



Strongest microbiota shift between weeks 16 and 24

# RESULTS

## Which functions and metabolisms are affected by the maturing process?

- KEGG orthology: **11.489 Kos**
- High impact of the productive stage:
  - **decrease** in functions abundance
  - Some functions were **not detected** on week 24

### 16 vs 24 weeks


- protein metabolism
- carbohydrate metabolism
- co-factors and vitamins
- lipid metabolism
- amino acid metabolism
- digestive system


### Breed

- genetic information processing
- signal transduction
- membrane transport and metabolism

# RESULTS

## P assimilation: inositol-phosphate pathway

- 
- **Up-regulation of K13024** (inositol-hexakisphosphate/diphosphoinositol-pentakisphosphate 1-kinase) **LSL vs LB crop week 16**
  - **Up-regulation K22231 – K22233** (3-dehydro-scyllo-inosose hydrolase, scyllo-inosose 3-dehydrogenase, 5-keto-L-gluconate epimerase) and **K17237** (InsP transport system substrate-binding protein) **LB vs LSL caeca week 24**

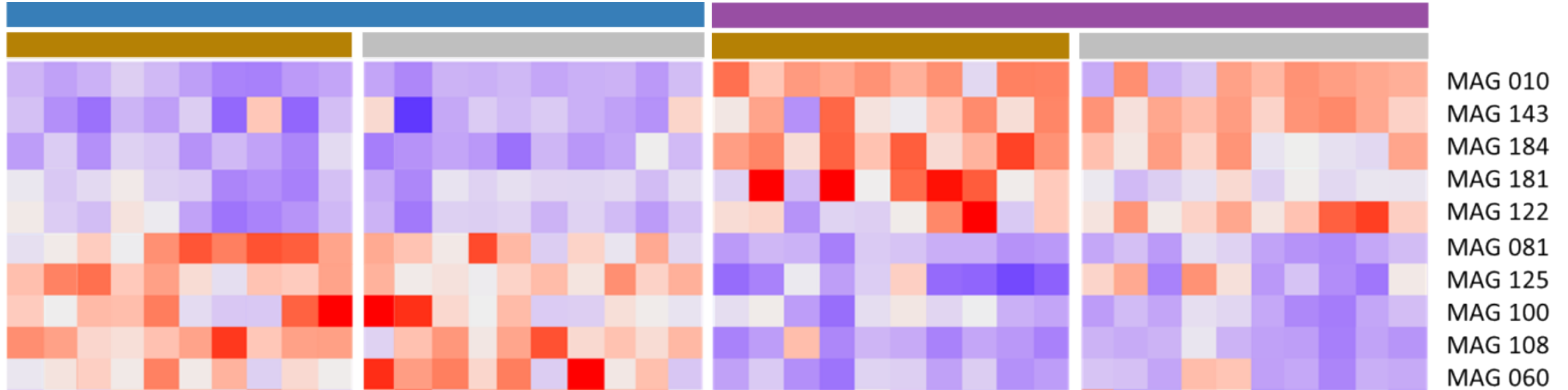


**Down-regulation of K22233** (5-keto-L-gluconate epimerase) (iolO) **LSL ileum from week 16 to 24**

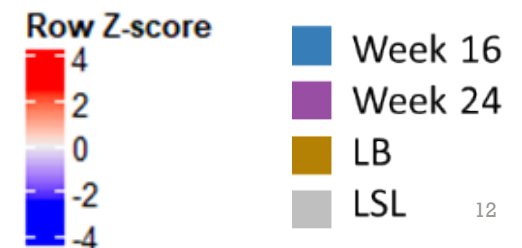
# RESULTS

- **204 MAGs** across 3 sections and 2 breeds
- **48 with high quality** (completeness >90%; contamination <5%)
- Average length of **2.5Mb** and **GC content of 46.6%**
- **MAGs differed between breeds, weeks 16 and 24 and GIT sections**
- Some **MAGS** could **not be assigned to a species**

# RESULTS



- Top 5 MAGs week 16: Amino acid utilization, Hydrogenesis and oxidative phosphorylation
- Top 5 MAGs week 24: Ethanol fermentation and complex carbon degradation



# CONCLUSION

- Same **diet** / housed under **similar conditions** – **breed and age effect on microbiota**
- **Major shift** in the microbiota between weeks **16** and **24**
  - bacterial fluctuations due to the starting of the laying period
- **Metagenomic** analysis confirmed the shift and highlighted **functional adaptations**
- It remains unclear if the changes in the feeding influenced the microbiota shifts or if the anatomical and physiological alterations affected the GIT microbiota



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THANK YOU FOR YOUR ATTENTION



**DFG**

