



## Identification of microbiome tailored strategies to improve safety and sustainability of poultry meat-based products

### Objectives

- To characterize the microbiome dynamics associated with poultry houses of different production scores.
- To identify and test microbiome-based interventions that can support the microbiome characterized

### Materials and Methods

- Observational phase: DNA from chickens cecal content were extracted and bioinformatic analysis of the sequenced data were performed.
- Literature review: conducted to provide evidence to support identified microbiome from the observational phase.

### Results

- *Psychrobacter* and *Ruminiclostridium 9* were the dominant genera identified in high performing poultry houses studied in the observational phase.
- **Synbiotic PoultryStar**<sup>®</sup> and the enzyme **Muramidase** were found to support the proliferation of identified microbial dynamics.

### Conclusion:

- **PoultryStar**<sup>®</sup> and **Muramidase** can be used as the microbiome-based strategies for the interventions phase.

### Future Proposal

- To test the identified microbiome-based nutritional strategies tailored to the positive microbiomes identified.

### Period Abroad/at Company

- Planned to go to University of Hohenheim, Germany

