

## Transgenerational response to endocrine disruptor: phenotypic, genotypic and epigenotypic analyses in quail

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Stacy Rousse, Sophie Leroux, David Gourichon, Ludovic Calandreau, Chloé Cerutti, et al.. Transgenerational response to endocrine disruptor: phenotypic, genotypic and epigenotypic analyses in quail. Environmental and Agronomical Genomics Symposium, Feb 2024, Toulouse, France., pp.128, 2024, Book of abstracts. hal-04474289

### HAL Id: hal-04474289 https://hal.inrae.fr/hal-04474289v1

Submitted on 5 Sep 2024

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# Transgenerational response to endocrine disruptor ingestion: phenotypic, genetic and epigenetic analyses in quail

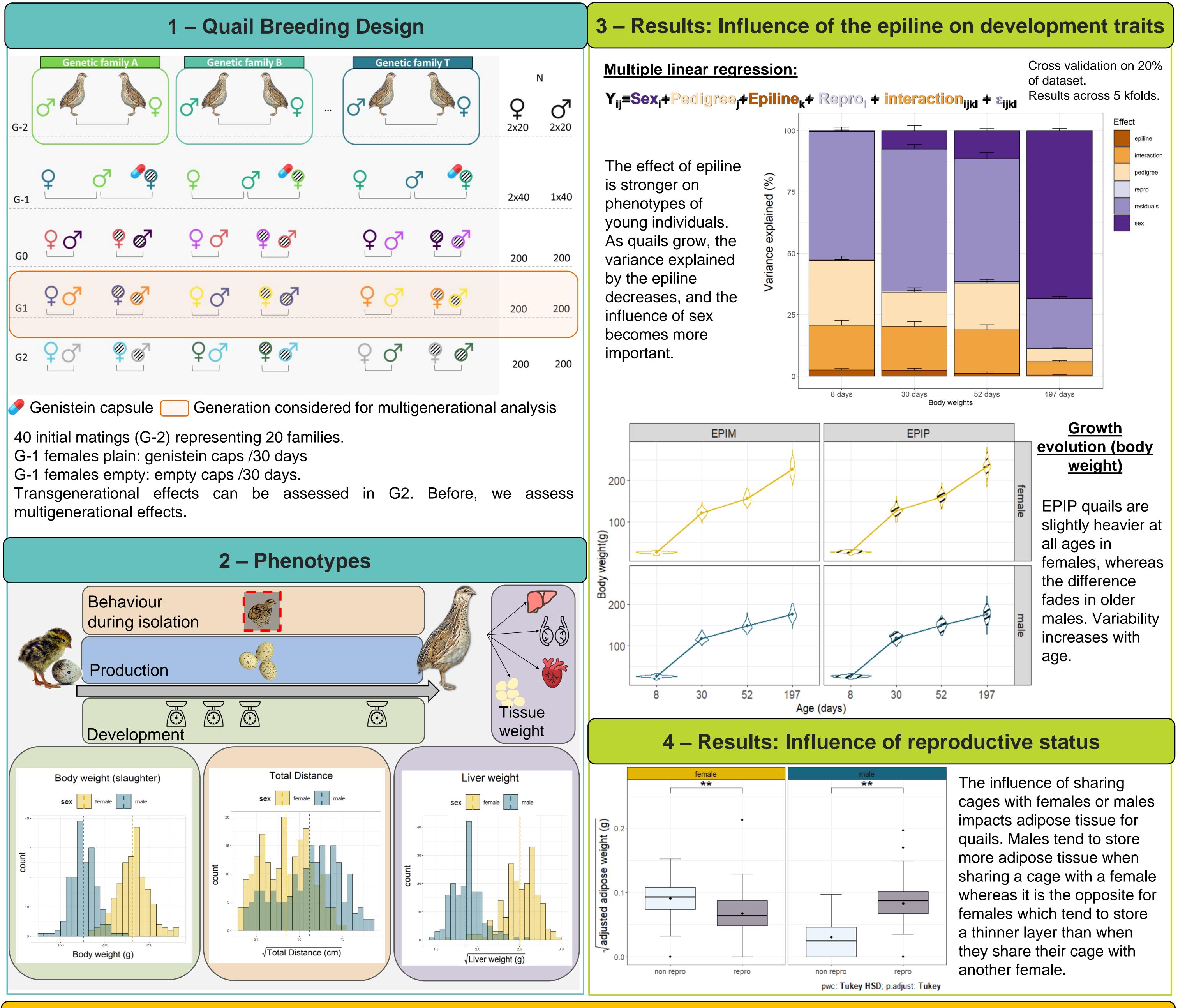
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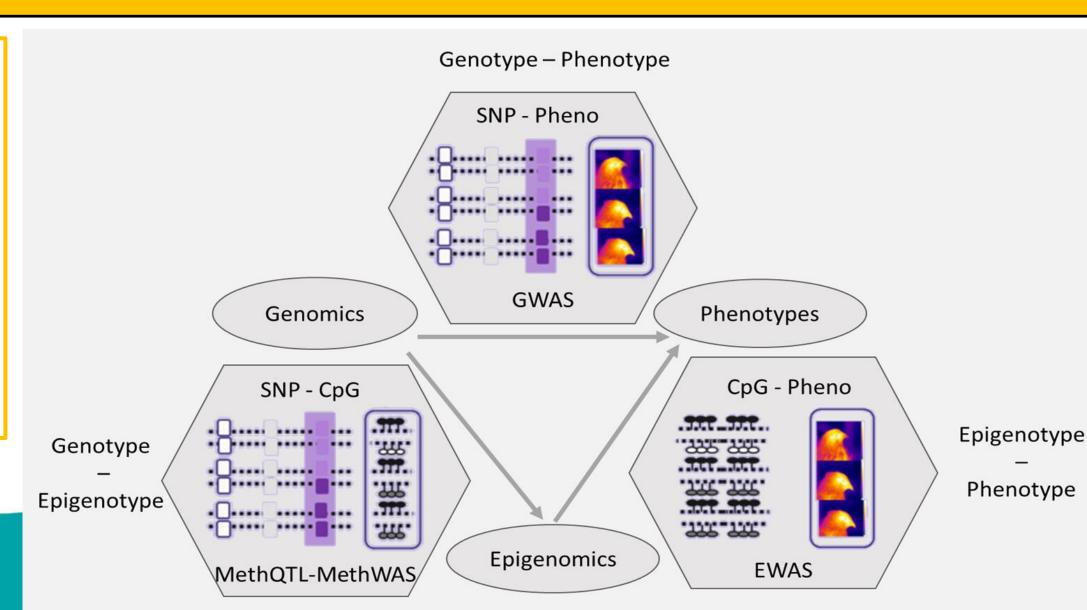
Typical evaluation of an offspring's phenotype usually focuses on the inheritance of parental alleles. However, variations across generations can also result from the transmission of non-genetic factors. Epigenetic marks like DNA methylation offer a dynamic molecular signature of an individual's history, as they can carry a memory of the individual's environmental past, and also of its parents. In this study, we aim to analyze the impact of an initial supplementation with an endocrine disruptor, genistein, on the phenotypes of subsequent generations. To better disentangle the complex interplay between genetic and epigenetic effects, a specific quail (Coturnix japonica) breeding plan was designed, where mirror crosses ensure a balanced genetic structure between epilines. By recording multiple phenotypes (development, breeding, production, behaviour), our goal is to investigate the transgenerational inheritance of an environmental effect through changes in DNA methylation profiles. Here, we present preliminary results on phenotypic traits two generations after the environmental disruption and thus, focus on multigenerational inheritance.



## 5 – Discussion and perspectives

Methylation patterns can be sensitive to environmental changes and can be transmitted to next generations. To evaluate the importance of non-genetic inheritance, one major difficulty lies in disentangling epigenetic effects from genetic effects.

We aim to answer this question by using methylome and genome data to better characterize multigenerational inheritance (RRBS data analysis to come) and analyze transgenerational effects (G2 data - May 2024).



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