



# Parasite infection mediates intergenerational DNA methylation in the three-spined stickleback (*Gasterosteus aculeatus*)



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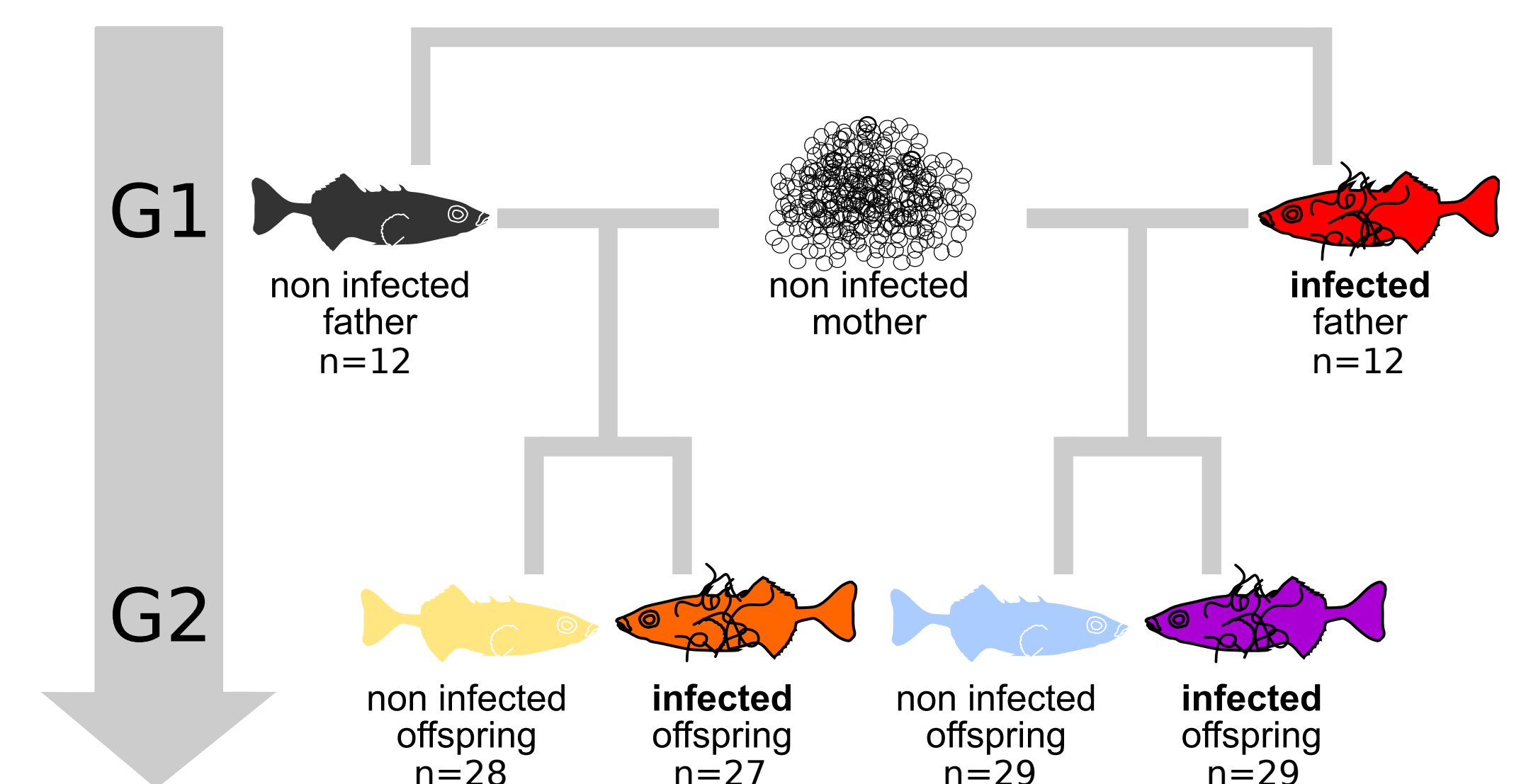
1. Paternal infection by the nematode *Camallanus lacustris* is associated with increased selection in offspring but also increased tolerance upon infection<sup>1</sup>
2. Genome-wide DNA methylation patterns differ between infected and control fish, demonstrating the link between infection and DNA methylation<sup>2</sup>

**Can parental DNA methylation induced by the parasite infection be transmitted to the next generation, and is it an underlying mechanism of the observed phenotypic differences?**

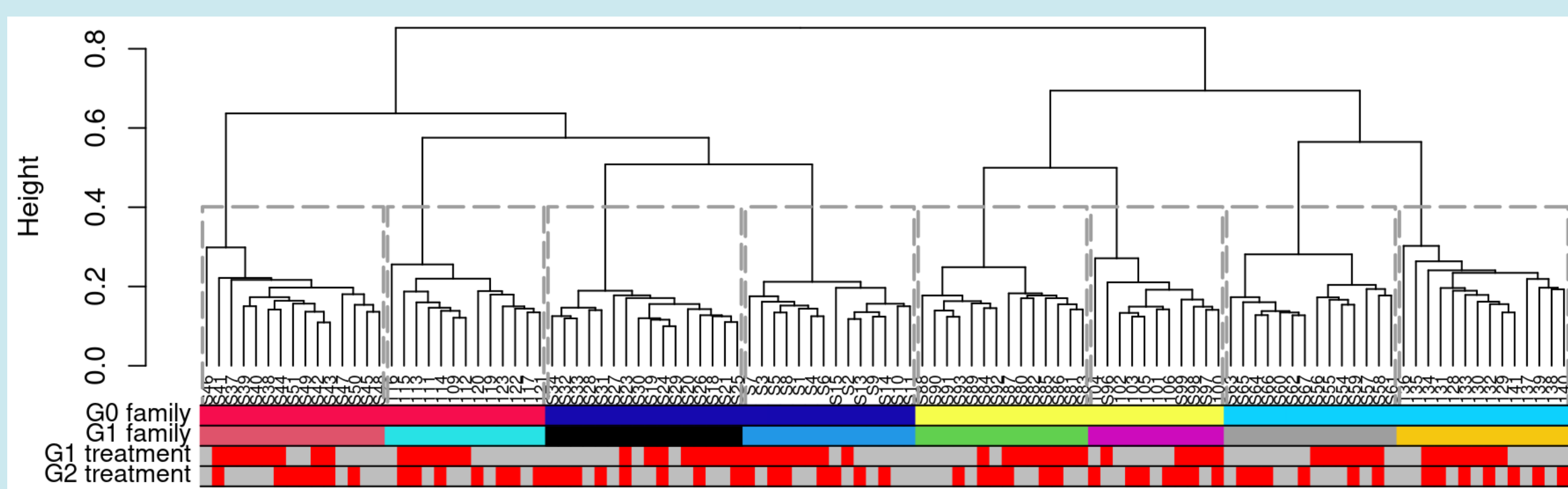
## Material & methods

- Methylome sequencing: **Reduced Representation Bisulfite Sequencing** single-end reads of 100bp long, Illumina HiSeq 2500. Alignment on a European gynogen genome<sup>3</sup> and methylation call with BSBolt. Downstream analyses with Methykit.
- Positional methylation:
  - Q1. Is genetics or treatment a stronger predictor of methylation?**
  - Q2. Is the methylation pattern affected by paternal/offspring treatment?**
- Differential methylation in half the brother pairs (4/8): **Q3. What are the specific differences between paternal/offspring treatment groups?**
- PCA of methylation at differentially methylated sites, linear model of Body Condition Index explained by the 2 first axes, the number of worms, paternal treatment, and interactions: **Q4. Can we find a link between methylation and phenotype?**

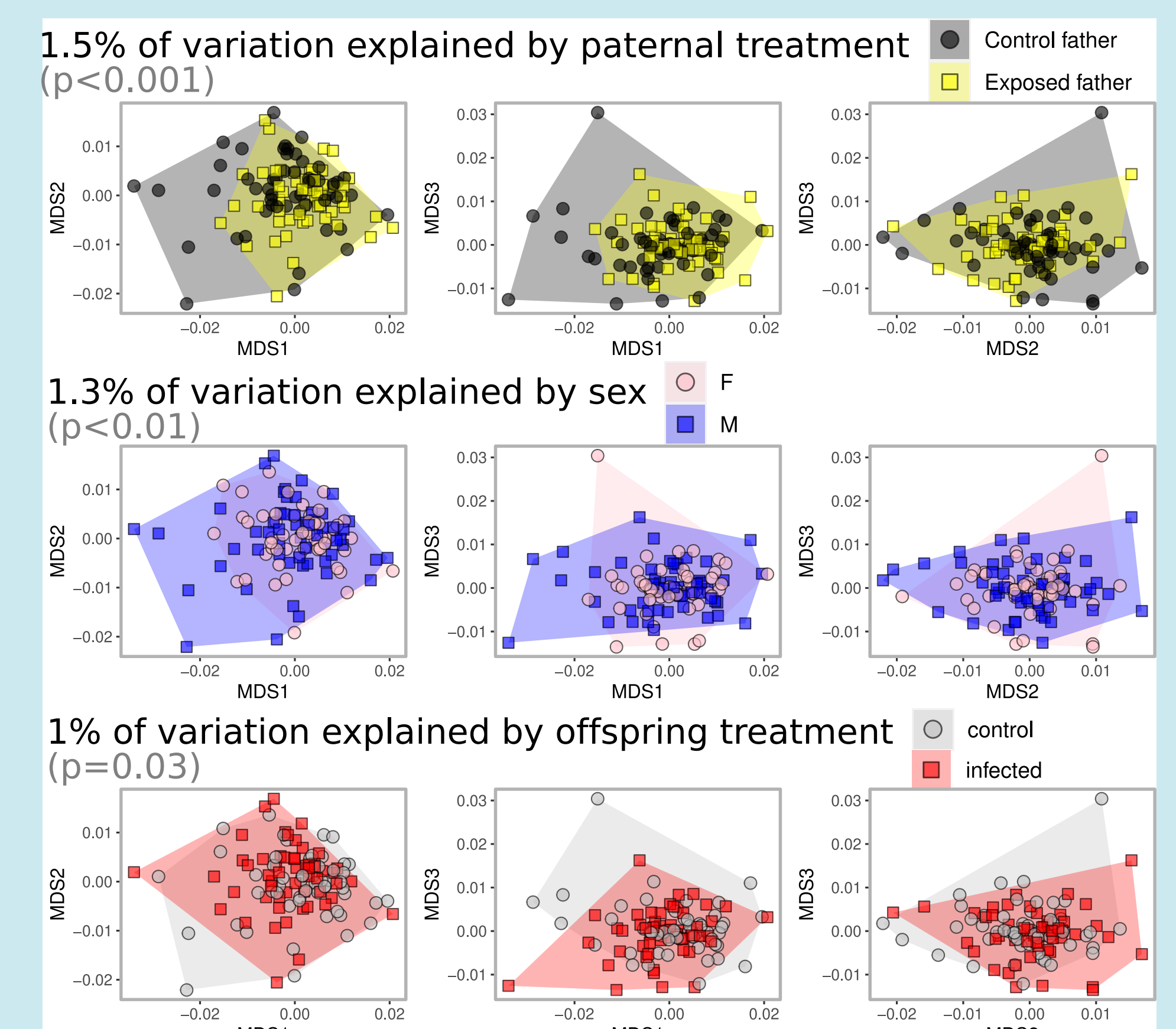
## Experimental design



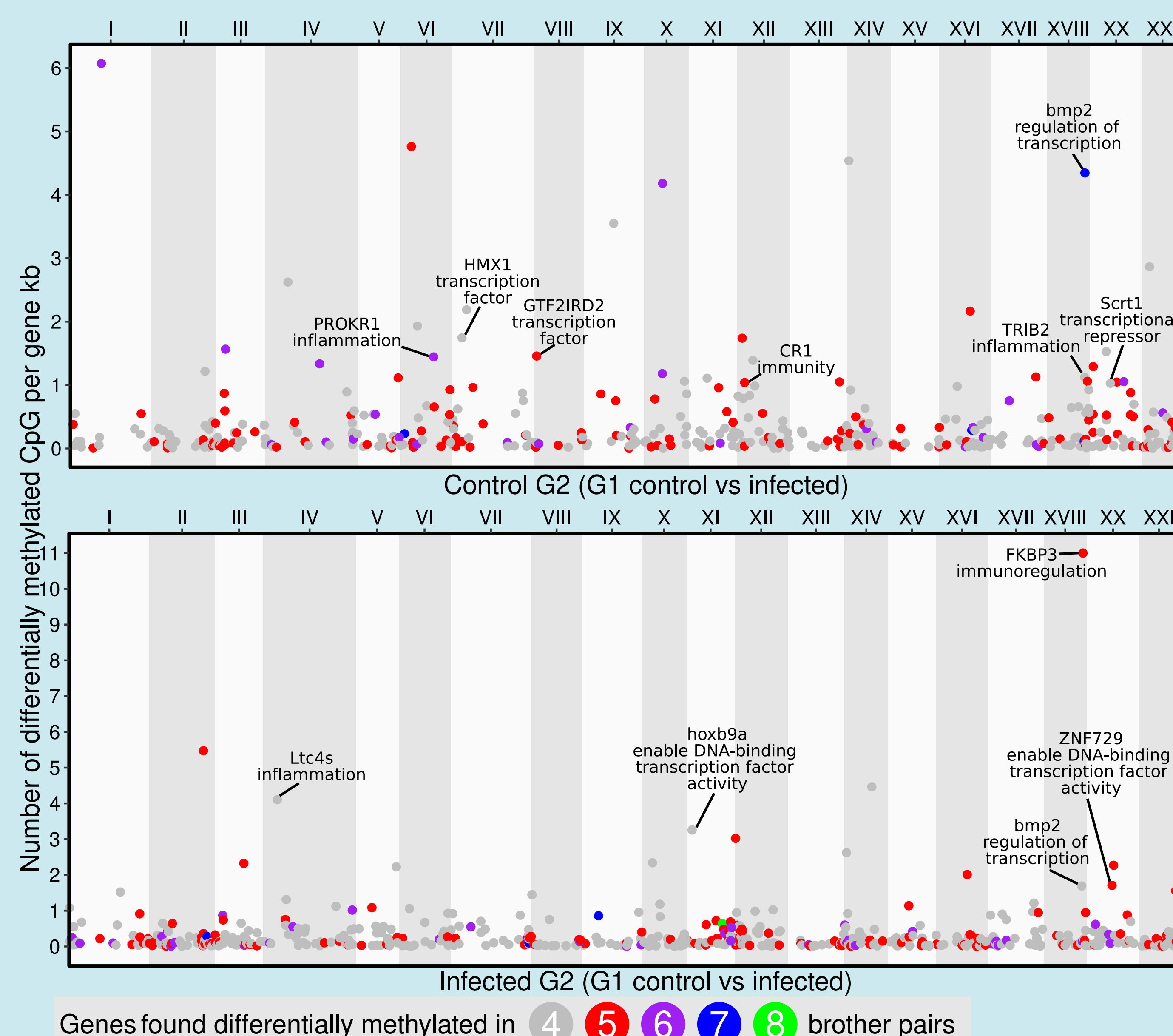
## Q1. DNA methylation profiles cluster by genetic background



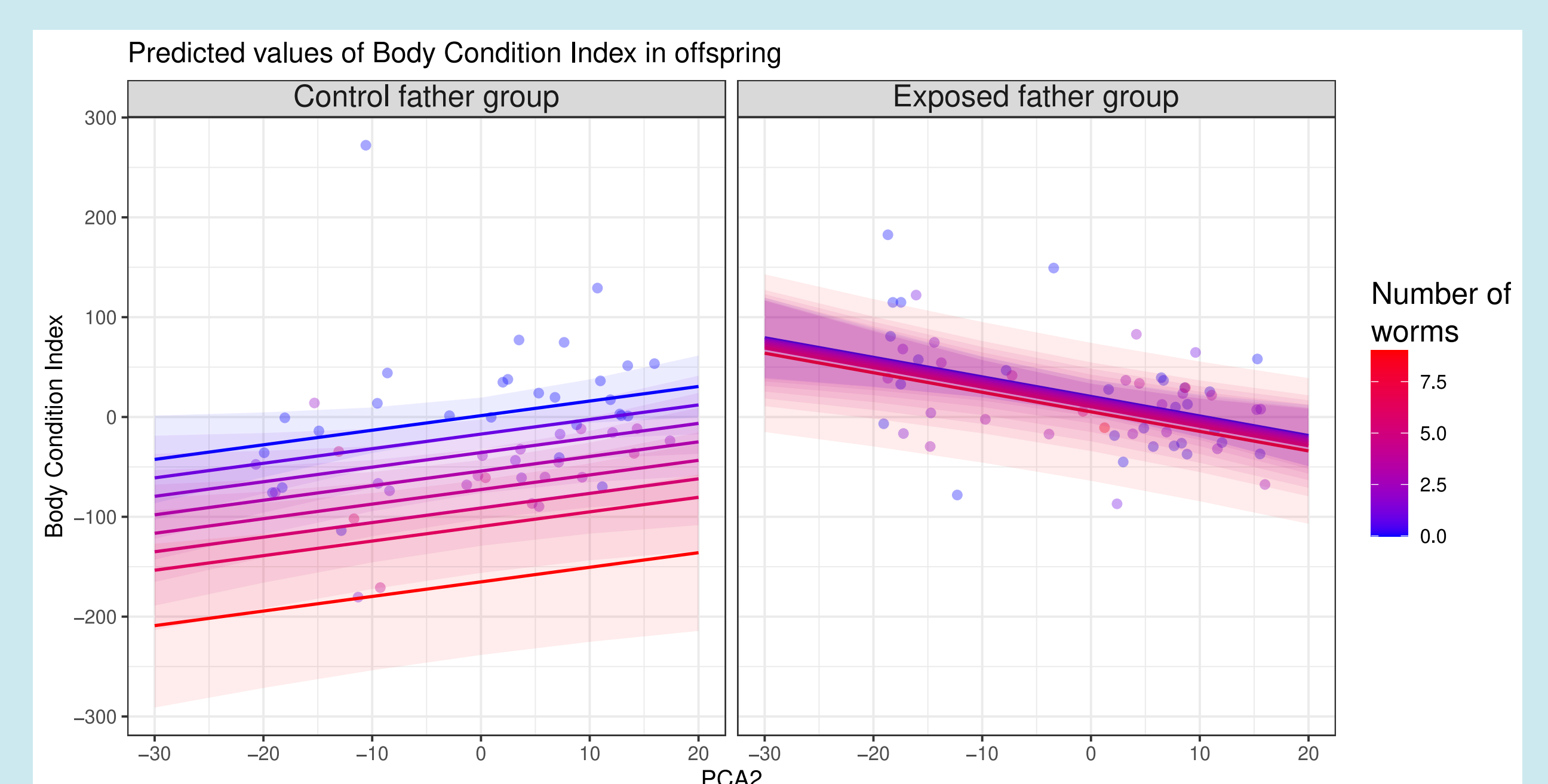
## Q2. Methylation is more affected by the paternal infection than by the offspring infection itself



## Q3. Specific methylated sites linked with paternal infection are associated with genes related to immunity and transcription



## Q4. Body condition correlates with methylation at differentially methylated sites, in different directions depending on the paternal treatment



<sup>1</sup>Kaufmann, J., Lenz, T. L., Milinski, M., & Eizaguirre, C. (2014). Experimental parasite infection reveals costs and benefits of paternal effects. *Ecology Letters*; <sup>2</sup>Sagonas, K., Meyer, B. S., Kaufmann, J., Lenz, T. L., Häslér, R., & Eizaguirre, C. (2020). Experimental parasite infection causes genome-wide changes in DNA methylation. *Molecular Biology and Evolution*; <sup>3</sup>Thornburn et al., in prep.