Transcriptominc divergence of a parasite-populations: two common garden experients in two hosts

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Abstract

Background:

Results:

Conclusions: Yeh!

Background Results

The populations differ

The gene expression does too

null device

1

Discussion Conclusions Methods General coding methods

The bulk of analysis (unless otherwise cited) presented in this paper was carried out in R [1] using custom scripts. We used a method provided in the R-packages Sweave [2] and Weaver [3] for "reproducible research" combining R and TeXcode in a single file. All intermediate data files needed to compile the present manuscript from data-sources are provided upon request. For visualistation we used the R-packages lattice [4] and ggplot2 [5].

Competing interests

The authors declare no competing interests.

Authors contributions Acknowledgments

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- Figures
 Figure 1 Figure 2 Figure 3 Figure 4 Figure 5 -

Tables

Table 1 -

Table 7 -

Additional Files