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
```

```
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 visualistaion we used the R-packages lattice [4] and ggplot2 [5].

Competing interests

The authors declare no competing interests.

Authors contributions

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Figures

Tables

Table 1 -

Table 7 -

Additional Files