

Reviewer Report

Title: "Genome sequencing of the sweetpotato whitefly *Bemisia tabaci* MED/Q"

Version: Revision 1 **Date:** 12/19/2016

Reviewer name: Denis Tagu

Reviewer Comments to Author:

Dear authors and editors,

With the recent publication of a *Bemisia tabaci* genome paper (BMC Biology, 2016 14:110) by a concurrent group, we have now two *Bemisia* genomes, released by two different consortia. More complicated, some authors are signing both papers! This situation is not well appropriate for the community. To which reference genome will the community refer to? It is really a pity that the two consortia did not work together to share means and ideas instead of duplicating sequencing and analyses.

I am thus forced to consider the submitted paper as a Data Note, since it does not afford strong novelty compare to the other genome. That's the stupid and unfortunate result of competition, instead of collaboration.

And at least, the two consortia should now communicate in order to define a reference genome, and to help the community to find their way. I do consider i5k as a central point for proposing access to insect genomes and I clearly invite the authors to contact them once/if the paper is accepted.

Concerning the re-submission, I thank the authors for having evaluating my suggestions. The addition of RNAi on some genes potentially involved in insecticide resistance is a clear added-value.

CEGMA versus BUSCO: I still think BUSCO is better than SEGMA but it is not a critical issue. The authors can stand with CEGMA, even if I suggest giving both analyses (CEGMA and BUSCO) to help the readers to estimate the chance to get a full sequence of genes.

So my rejection concerns a publication in GigaScience as a Research Report, but is an acceptance for a Data Note.

Sorry for being so severe, but I feel really bad in front of these situations.

All the best

Denis

Methods

Are the methods appropriate to the aims of the study, are they well described, and are necessary controls included? Yes

Conclusions

Are the conclusions adequately supported by the data shown? Yes

Reporting Standards

Does the manuscript adhere to the journal's guidelines on [minimum standards of reporting?](#) Yes

Statistics

Are you able to assess all statistics in the manuscript, including the appropriateness of statistical tests used? Yes, and I have assessed the statistics in my report.

Quality of Written English

Please indicate the quality of language in the manuscript: Acceptable

Declaration of Competing Interests

Please complete a declaration of competing interests, considering the following questions:

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