

Reviewer Report

Title: Genome sequence of Japanese oak silk moth, *Antheraea yamamai*: the first draft genome in family Saturniidae

Version: Revision 2 **Date:** 9/18/2017

Reviewer name: Simon Baxter

Reviewer Comments to Author:

Dear authors, Thank you for your detailed response to my review. It's encouraging to see some additional gene curation alter the number of predicted genes from >21,000 to 15,400. I'm sure this effort will make the *A. yamamai* genome a more useful resource for the scientific community. I would recommend changing the abstract to include the curated number, rather than automated gene number. The website <http://lepbase.org> hosts numerous lepidopteran genomes, and may be a useful additional repository for this genome.

Level of Interest

Please indicate how interesting you found the manuscript: An article of importance in its field

Quality of Written English

Please indicate the quality of language in the manuscript: Needs some language corrections before being published

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