Reviewer Report

Title: Chromosome-level genome assembly and annotation of the loquat (Eriobotrya japonica) genome

Version: Original Submission Date: 11/28/2019

Reviewer name: Soichiro Nagano

Reviewer Comments to Author:

This paper is worth publishing for the Data Note for GigaScience because the authors have constructed a highly accurate genome and gene sequence of loquat. The method is reasonable and the presentation is pretty good. Speaking of greed, since it is clear that the relationship with Malus is relatively close among the fruit trees of the Rosaceae family, so there should be presented some discussion about the traces on the genome that triggered the differentiation of morphological features between both species, I recommend that as it would be done for the future work. The following minor concerns should be corrected before acceptance.

Minor concerns

L58 How to extract the RNA from the collected samples should be described.

L63 double-end would be pair-end.

L71 the appearance of the equation is not clear. Please correct the format of the equation.

L86 "quality" have to be "quantity", because the Qubit 2.0 Fluorometer is the device to evaluate the quantity of the DNA/RNA with fluorescence.

L89 How many flow cells have been used with the PromethION platform to acquire about the 106.23 Gb. L205 "Sac1, 4, and 8" has been suddenly appeared at the text. Please describe about relationships between the former and later text to be clearly understood, although chromosome scale duplication is very interesting.

L349 "de novo" should be written in italic.

L364 Add period.

L366 Add period.

L367 E. japonica should be written in italic.

L375 "Hic" should be "Hi-C"

Figures Provide higher resolutional figures than current version. Because those are not clear.

Figure 4A Correct the overlap of the legends on a bar.

Figure 5A It is difficult to see the scale of the figure. Please provide higher resolutional figures.

Level of Interest

Please indicate how interesting you found the manuscript: Choose an item.

Quality of Written English

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