

Reviewer Report

Title: Comparison of the two up-to-date sequencing technologies for genome assembly: HiFi reads of Pacbio Sequel II system and ultralong reads of Oxford Nanopore

Version: Revision 1 **Date: 7/14/2020**

Reviewer name: Jason Chin

Reviewer Comments to Author:

Thanks for address most of the points I raised. The revised manuscript is a good improvement. Thanks. One minor thing, I am not sure the term "one suite of a diploid genome" is the right way to describe one single haplotype of the homologous chromosomes, please consider to the revise that for the manuscript.

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1. No, although I was an employee of one the company which developed the related technologies 2. No
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