Reviewer Report

Title: Draft genome sequence of the Solanum aethiopicum provides insights into disease resistance, drought tolerance and the evolution of the genome

Version: Original Submission Date: 4/3/2019

Reviewer name: Kenta Shirasawa

Reviewer Comments to Author:

This paper reports the first genome assembly of Solanum aethiopicum. The description is easy to follow and the data would be useful for the breeding programs of eggplant. I recommend the authors to submit the data (genome, genes, protein, annotatoin, sequence variations etc) to Sol Genomics Network so that potential users can access them easily.

Minor comments:

The term "the reference genome" in the main text should be replaced by "the reference genome sequence".

Abstract: LTR-Rs should be spelled out.

Abstract: "closely" is ambiguous.

Introduction: "We also re-sequenced two ...". Is this 65 (not two) as mentioned in Abstract and other parts?

Data Description: While a total of 242.6 Gb raw reads were obtained, only 127.83 Gb were used for assembly. I assume that approximately 115 Gb reads were low quality. Correct?

Data Description: Only 80.4% complete BUSCOs were found in the assembly, whereas the total length of the assembly was 1.02 Gb covering 87% of the estimated genome size (1.17 Gb). Please clarify the reason for the low BUSCOs.

Increased resistance is facilitated by LTR-Rs amplification: What is the definition of "LTR-Rs captured"? It is unclear why the "LTR-Rs captured" genes enhance disease resistance.

NLR?

Polymorphisms in different S. aethiopicum groups: What's the difference between indels and SVs? Artificially selected genes in S. aethiopicum: What types of selections do the authors mention here? Potential implications: This part can be deleted because this is not based on the data.

Methods: What are the "standard BGI protocols"?

SNP calling: "samtools mpileup" and "VariantFiltration" are duplicated.

Methods

Are the methods appropriate to the aims of the study, are they well described, and are necessary controls included? Choose an item.

Conclusions

Are the conclusions adequately supported by the data shown? Choose an item.

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