

The complete genome of Banana streak GF virus Yunnan isolate infecting Cavendish Musa AAA group in China

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1. Basic reporting

The manuscript has been updated for English language however, few sentences needs to be changed:

1. Line 35-36: " The result showed that *Banana streak GF virus* Yunnan isolate (BSGFV-YN) infecting Cavendish *Musa* AAA group was determined " should be " The result showed that *Banana streak GF virus* Yunnan isolate (BSGFV-YN) infecting Cavendish *Musa* AAA group was co-infecting this sample ".
2. Line 46: " In summary, a BSGFV-YN different from BSV-Acum was identified from the one sample " should be " In summary, a BSGFV-YN different from BSV-Acum was identified from the same plant ".
3. Line 118: " bio-sanger sequencing at Invitrogen (Guangzhou, China). " should be " Sanger sequencing at Invitrogen (Guangzhou, China). ".
4. Line 151: " each fragment was high similar to the BSGFV " should be " each fragment was highly similar to the BSGFV ".
5. Line 210: " DISCUSS" should be " DISCUSSION".
6. Line 215 to 217: "The badnaviruses have been concerned by people recently owing to they could cause significant agricultural economy loss." Should be rephrased for more clarity.

The literature references are fine and cited in appropriate context. The manuscript has an acceptable format with good sectioning structure and is self-contained as well.

2. Experimental design

The manuscript falls under the Aims and Scope of PeerJ. The concept and experimental design of the study is fine.

3. Validity of the findings

In my previous review report, I have commented as "Currently, the manuscript is limited to sequencing of an isolate, however, no work is presented for its prevalence in the area; whether and to which extant is it present in the region? Is it present only in association with BSACYNV or alone? Is it episomal or present as integrated in the field? The authors are encouraged to determine the presence of this isolates in the area using specific primers/PCR assay and sequence the genome of few more isolates". Though the authors have not added any data for these aspects, however, their reply is logical.