First report of Banana streak virus infecting banana in Burkina Faso

ABSTRACT

Banana plants (*Musa* spp), both food and economic crops, are potential hosts for a various range of badnavirus species considered a major constraint to banana improvement and a threat to *Musa* production worldwide. A survey and sample collection were carried out in two main banana-growing regions of Burkina Faso. The samples collected were subjected to biological, serological and molecular diagnosis of the reverse transcriptase/ribonuclease H (RT/RNase H) region using Badna FP/RP primers, followed by sequence comparison with the Genbank database. Analyses confirmed the presence of BSV in all symptomatic samples tested in both regions. Amplification bands of the expected size were obtained for the symptomatic samples tested positive in the serological test. These partial RT/RNase H gene sequences shared highest nucleotide identity ranging from 85.80% to 99.05% with DBV isolates in GenBank. These results are proof of the existence of BSV in Burkina Faso. It is therefore important to undertake studies that will provide basic information on the virus for the development of effective control strategies.

Comment [M1]: Full meaning of DBV?

Comment [M2]: State the meaning in either abstract or introduction and subsequently with the abbreviation as BSV

INTRODUCTION

Banana (*Musa spp*) is herbaceous plant belonging to the family *Musaceae* of the genus *Musa* and is native to the tropical region of South East Asia. Viruses are important constraints to the movement and propagation of plant germplasm, especially for vegetatively propagated crops such as banana and plantain. Banana production is threatened by the Banana streak disease (BSD), and its pathogen belongs to the genus Badnavirus, family *Caulimoviridae* (Alangar *et al.*, 2016). BSV is widely distributed in the main planting areas of banana industry in Southeast Asia and Africa, and it had seriously affected the yield and quality of bananas resulted in huge economic losses (Kumar *et al.*, 2015). BSD can manifest with wide-ranging symptoms, from complete lack of visible impacts to plant death, depending on BSV isolates, host cultivars, and environmental conditions (Dahal *et al.*, 2000). However, the major symptoms of BSD are chlorotic and necrotic streaks.

Comment [M3]: Report of 2016 as the latest report on BSV? Try to find the latest reports throughout the manuscript instead of using old references. Like 2020 to 2024.

Comment [M4]: Mention the meaning for the first time and then proceed with abbreviation as BSV!

In Burkina Faso, banana is produced in all agro-ecological zones and help combat food and nutritional insecurity and unemployment. From 2012 to 2014, production rose from 50.571

tonnes to 79.561 tonnes (TFB, 2015). In spite of this increase, production is struggling to cover national consumption needs. This is due to abiotic and biotic constraints including viral diseases causing drastic yield loses. Unfortunately, the viruses infecting bananas are not well documented in Burkina Faso's banana-growing areas. BSV has never been reported in banana growing areas in Burkina Faso. Alt~~T~~hough, viral diseases symptoms similar to badnavirus symptoms was observed on banana plant in many regions.

The objective of this study is to do what? Through which methodology? And what do you want to achieve all should be stated in this paragraph

MATERIALS AND METHODS

Twenty-five leaf samples were collected from symptomatic banana plants in two main production regions, Boucle du Mouhoun and Hauts-Bassins. A~~a~~mong those, 22 showed disease symptoms and 3 were symptomless. All samples were assayed to confirm BSV infection by Indirect Antigen Coated Plate Assay-ELISA with BSV polyclonal antisera (developed in-house at CIRAD, ~~kindly~~ provided by Serge GALZI). BSV was detected in all samples tested. Total DNA was extracted by the CTAB protocol (Permingeat *et al.*, 1998) from 25 samples. The fragment of RT/RNase H region was amplified by PCR using primers Badna FP (5'-ATGCCITTYGGIAARAAYGCICC-3') and Badna RP (5'-CCAYTTRCAIACISCICCCCAICC-3') (Yang *et al.*, 2003). Amplification products were sent for sequencing to Macrogen (Amsterdam, Netherlands). Sequences were compared with other viral sequences in the NCBI database using BLAST (BLAST, http://www.ncbi.nlm.nih.gov/blast).

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RESULTS AND DISCUSSION

During the survey, plants showing symptoms of discontinuous and continuous streaks, chlorotic streaks and necrotic streaks were observed in the plantations (Fig. 1). Based on the nature of the symptoms observed, infection by BSV was suspected. These symptoms are similar to the usual BSV symptoms reported by many authors (Gayral and Iskra-Caruana, 2009; Furuya *et al.*, 2012).

All symptomatic samples were positives for serological assay but any symptomless samples were positives. An amplification product of the expected size was obtained for all 22 symptomatic sample tested positive (Fig. 2). Five PCR products were directly sequenced. These five sequences showed identity ranging from 85.80% to 99.05% with each other BSV isolates. All these facts are evidence of the existence of BSV in the banana-growing areas

Comment [M5]: The reference sequences supposed to be shown and aligned with the obtained sequenced of the tested sample and al so indicate the similarity of the sequences through colour shadings!

surveyed in Burkina Faso. The presence of BSV in bordering countries such as Côte d'Ivoire (Kouadio *et al*., 2016), Benin (Pasberg-Gauhl *et al*., 1996), Togo (Lockhart, 1995) and Ghana, Nigeria (Agindotan *et al*., 2006) has already been reported. Exchanges of plant material between these countries could be cause of the spread of the virus in the sub-region.

Fig 1. Symptoms observed in the surveyed banana plantations. (A) Healthy leaves, (B) Discontinuous and discontinuous chlorotic streaks, (C) Chlorotic and necrotic streaks

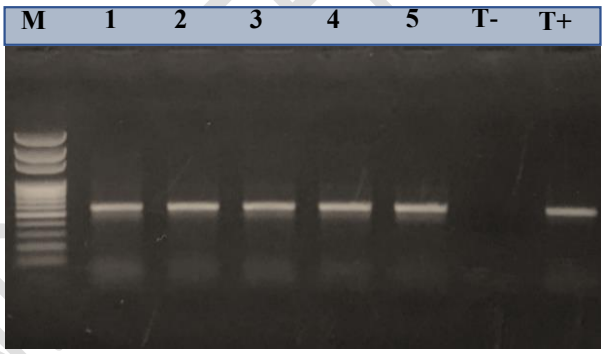

Fig 2. Agarose gel electrophoresis (1%) showing PCR amplified products: lane M = 100pb DNA ladder (Solis Biodyne); lanes 1, 2 = BSV infected samples, lane T- = negative control; lane T+ = Positive control

CONCLUSION

To the best of our knowledge, this is the first report of BSV infecting banana in Burkina Faso. Based on the GenBank repository, the sequenced virus revealed the similarity of greater than 70% which is enough to confirmed the virus as BSV. Therefore, the obtained identity of 85.80% to 99.05% validated the virus is 85.80% to 99.05%. This report is the primary step to

initiate research on the impact of the virus in banana production and germplasm exchange. Further research is needed to elucidate epidemiology and impact of the virus in banana production and germplasm exchange.

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