

DETECTION OF *BANANA MILD MOSAIC VIRUS* AND BANANA VIRUS X BY POLYVALENT DEGENERATE OLIGONUCLEOTIDE REVERSE TRANSCRIPTION PCR (PDO-RT-PCR)



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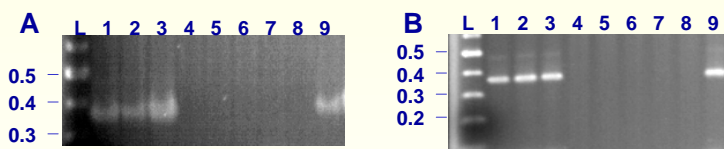
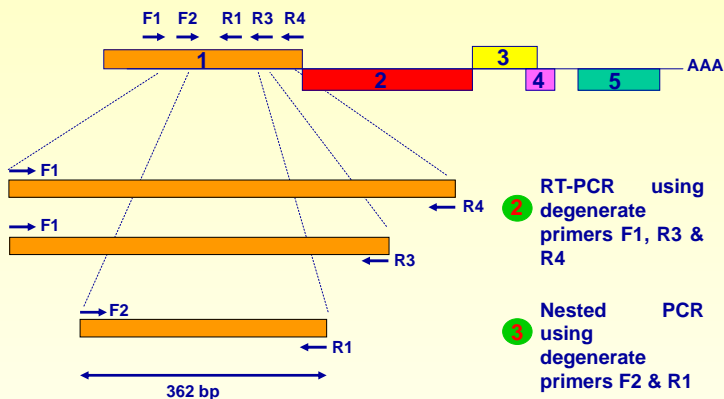
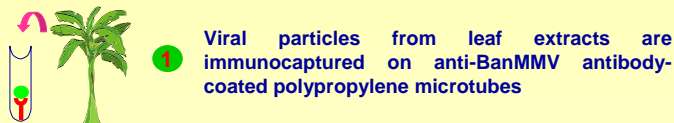


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An existing polyvalent degenerate oligonucleotide RT-PCR (PDO-RT-PCR) assay [1] was adapted for the detection of *Banana mild mosaic virus* (BanMMV) and *Banana virus X*, two *Flexiviridae* infecting *Musa* spp. PDO inosine-containing primers were found to be well suited for the detection of BanMMV, despite its high sequence variability [2], but not for that of the highly conserved BVX [3], for which species-specific primers were therefore designed. The sampling and sample processing steps were optimized in order to avoid nucleic acid purification prior to the reverse transcription step. A **polyclonal anti-BanMMV antiserum** was raised and successfully used for the immunocapture (IC) of BanMMV viral particles from leaf extracts, leading to the development of a **PDO-IC-RT-nested PCR assay** [4]. Although the anti-BanMMV antiserum could to some extent recognize BVX particles, **direct binding (DB)** was shown to be a more efficient method for processing BVX-infected samples and a **PDO-DB-RT-nested PCR assay** was developed for the detection of BVX from leaf extracts [4]. This work completes existing PCR-based detection of other viral species infecting *Musa* spp. [5, 6] and will benefit movement and propagation of *Musa* germplasm, for which viruses are important constraints.

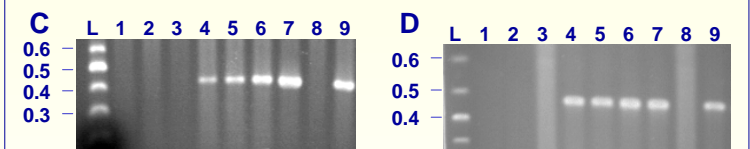
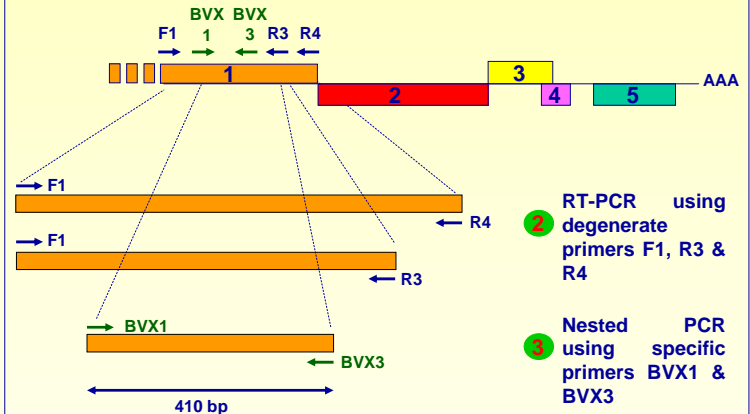
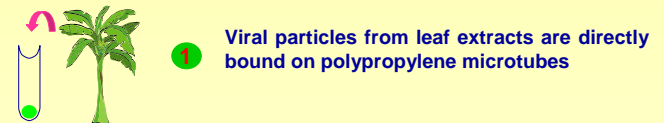
Detection of BanMMV by PDO-IC-RT-nested PCR



Detection of BanMMV by PDO-RT-nested PCR (A) and PDO-IC-RT-nested PCR (B)

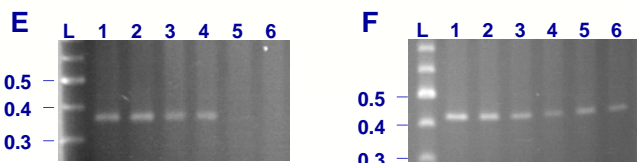
RT-PCR reactions were carried out on 50 ng of total RNAs purified from leaf tissues (A) or on immunocaptured viral particles (B) from leaf samples of plants from *Musa* accessions Moto Ebanga (lane 1), Kwa (lane 2), Kelong mekintu (lane 3), Som (lanes 4, 5 and 6) and Buitenzorg (lane 7). Lane 8 : uninfected control. Lane 9 : 10 ng of plasmid DNA containing part of the BanMMV genome used as a positive control for nested PCR. L: 100 bp DNA ladder (Promega) with indicated marker sizes in kbp.

Detection of BVX by PDO-DB-RT-nested PCR



Detection of BVX by PDO-RT-nested PCR (C) and PDO-DB-RT-nested PCR (D)

RT-PCR reactions were carried out on 50 ng of total RNAs purified from leaf tissues (A) or on directly trapped viral particles (B) from leaf samples of plants from *Musa* accessions Moto Ebanga (lane 1), Kwa (lane 2), Kelong mekintu (lane 3), Som (lanes 4, 5 and 6) and Buitenzorg (lane 7). Lane 8 : uninfected control. Lane 9 : 10 ng of plasmid DNA containing part of the BanMMV genome used as a positive control for nested PCR. L: 100 bp DNA ladder (Promega) with indicated marker sizes in kbp.



Sensitivity thresholds of IC-PDO-RT-nested PCR for the detection of BanMMV (E) and DB-PDO-RT-nested PCR for the detection of BVX (F)

IC-PDO-RT-nested PCR (E) and DB-PDO-RT-nested PCR (F) were performed using leaf extracts prepared from accession Kelong mekintu (E) or Som (F). Lane 1 : undiluted leaf extracts; lanes 2 to 6 : 1:10 to 1:100,00 dilutions, respectively of leaf extract. L: 100 bp DNA ladder (Promega) with indicated marker sizes in kbp.