

Enhancing Phytosanitary Systems for Healthy Plants, Safe & Sustainable Trade"



Sub-theme:

Pest Surveillance in Phytosanitary Systems

Title:

Incidence and Distribution of Cassava Mosaic and Cassava Brown Streak Diseases in Western Kenya

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Introduction

Cassava (*Manihot esculenta* Crantz) belongs to the *Euphorbiaceae* family and the genus *Manihot*.

≻Number one root crop followed by yam and sweetpotato (FAOSTAT 2017).

➢Used as food ,feed and industrial purposes(Chikoti *et al.*, 2015).

➢ Propagated through stem cuttings.







➢ Pests and diseases are the major cassava biotic stress.

CMD and CBSD are the two most economically important viruses in cassava.

▶ Result in losses worth more than US\$1 billion every year (IITA, 2014).

>Vectored by whitefly (*Bemisia tabaci*).

Cassava production is challenged by lack of suitable varieties with farmer preferred traits









Problem Statement









Cassava has a great potential to sustainably change African economic

development and improve income-earning for rural farmers.

➤To tap this potential ,cassava productivity must be improved by addressing the main viral diseases.







➢To determine Incidence and Distribution of Cassava Mosaic and Cassava Brown

Streak Diseases in Western Kenya.







Methodology

- Survey in 5 counties
 Kakamega,
 Bungoma, Busia,
 Migori and
 Homabay.
- 201 household were interviewed
- 255 stakes of landraces collected











- Collected stakes planted in pots at KALRO Kakamega and KEPHIS PQBS.
- After sprouting leaves collected for RNA extraction and DNA extraction (Lodhi *et al.*,1994).
- PCR-CMD; ACMV & EACMV Primers ; (Alabi et al.,2008 and Fondong et al.,2000)
- ➢ qPCR- CBSD;CBSV & UCBSV Primers; (Abarshi et al.,2012).
- > 59.8% of landraces CMD positive
- 22.3% of landraces CBSD Positive























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- CMD and CBSD are wide spread in the five counties
- > Need for Farmers to access clean/virus free planting material.
- Need to have cultivars with dual resistance with CMD and CBSD that have farmers preffered traits







Acknowledgements





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