



Identification and characterization of *Pythium* spp. at CIAT-Kawanda, Uganda

Introduction



Pythium spp. cause severe bean root rots under cool and humid conditions. Poor soils and intensive cultivation compound the epidemics.

There are more than 120 species reported globally some of which are very pathogenic meanwhile others are saprophytic. Yield losses due to *Pythium* outbreaks can be as high as 100% on susceptible varieties in the absence of control measures.

Bean root rot caused by *Pythium* is the most devastating in Eastern & Central Africa where the intensity of bean production is high.

Pathogen isolation



Uproot Wash, blot dry



CMA amended with antibiotics, incubate at 24°C for 12 hrs



PDA plate



Preservation:
PDA agar slants at 4°C



V8 juice



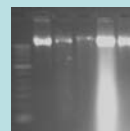
TWA

DNA extraction and PCR

DNA is isolated from a mat of mycelia growing on V8 juice.

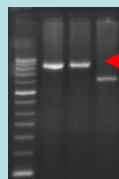


Maceration and DNA extraction



DNA quality and quantity

Amplification using *Pythium* specific primers



Pythium (700bp)
Others



Electrophoresis

Species identification

- PCR products of *Pythium* isolates are shipped for automated sequencing abroad.

- Chromatograms are verified and edited for correct peak scoring.

- Bioinformatics applications are used to identify the species through BLAST searches.

Pathology and screening tests

Newly identified *Pythium* species are:

1. Tested for pathogenicity using a resistant (RWR 719) and a susceptible (CAL96) varieties.
2. Used to screen Marker Assisted Selection lines to ascertain root rot resistance traits.

Procedure:



PDA, 4 days 24°C



Sterilized finger millet/sorghum, 14 days RT



Evaluation:
Uproot wash and score 21 days after planting



1-9 Scale



1 3 5 7 9



Mix inoculum : 8 sterilized soil and plant



Watering thrice a day in the 3rd week



This procedure is also applicable to *Fusarium* spp.