

Sweetpotato Action for Security and Health in Africa

SASHA

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Introduction



- Third most important root crop after potato and cassava
- Seventh in global food crop production
- Fourth in developing world after rice, wheat, and maize
- Viral diseases are greatest production constraints

Major viruses infecting sweet potato SASHA

Sweetpotato Action for

Virus name	Abb	Family (genus)	Transmission
Sweet potato feathery mottle virus	SPEMV	Potyviridae (Potyvirus)	Aphid (non-persistent)
Sweet potato virus G	SPVG	Potyviridae (Potyvirus)	Aphid (non-persistent)
Sweet potato latent virus	SPLV	Potyviridae (Potyvirus)	Aphid (non-persistent)
Sweet potato leaf curl virus ^a	SPLCV	Geminiviridae (Begomovirus)	Whitefly (persistent)
Sweet potato mild speckling virus	SPMSV	Potyviridae (Potyvirus)	Aphid (non-persistent)
Sweet potato mild mottle virus	SPMMV	Potyviridae (Ipovirus)	Whitefly (persistent)
Sweet potato chlorotic stunt virus	SPCSV	Closteroviridae (Crinivirus)	Whitefly(non-persistent)
Sweet potato collusive virus	SPCV	Caulimoviridae(Cavemovirus)	*
Sweet potato virus 2	SPV2	Potyviridae (Potyvirus)	Aphid (non-persistent)
Sweet potato virus C	SPVC	Potyviridae (Potyvirus)	Aphid (non-persistent)
Sweet potato symptomless virus 1	SPSMV-1	Geminiviridae (Mastrevirus)	*
Sweet potato chlorotic fleck virus	SPCFV	Betaflexiviridae (Carlavirus)	*
Sweet potato vein clearing virus	SPVCV	Caulimoviridae (Solendovirus)	4:
Sweet potato pakakuy virus	SPPV	Caulimoviridae (Badnavirus)	40
Sweet potato C6 virus	SPC6V	Betaflexiviridae (Carlavirus)	40
Sweet potato leaf speckling virus	SPLSV	Luteoviridae (Polerovirus)	*
Cucumber mosaic virus	CMV	Bromovindae (Cucumovirus)	Aphid (non-persistent)

^aSweet potato leaf curl virus has been classified into seven species: Sweet potato leaf curl virus (SPLCV), Ipomoea yellow vein virus (IYVV), Sweet potato leaf curl Georgia virus (SPLCGoV), Sweet potato leaf curl China virus (SPLCV-CN), Sweet potato leaf curl Lanzarote virus (SPLCLaV), Sweet potato leaf curl Canary virus (SPLCCaV), and Sweet potato leaf curl Spain virus (SPLCESV) by ICTV. *Not reported.

Effect of specific virus unknown for the recently described viruses

Why begomoviruses



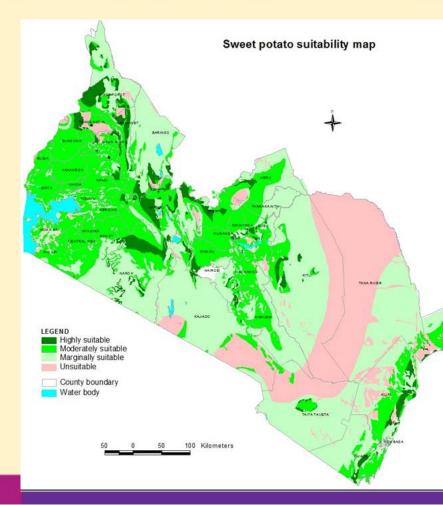
 Genus of economic importance to many crops

Widely distributed/diverse

Cause yield loses in sweetpotato up to 40%

 Occurrence, distribution and diversity in Kenya?

Conduct country-wide survey to assess the incidence and distribution of begomoviruses in Kenya



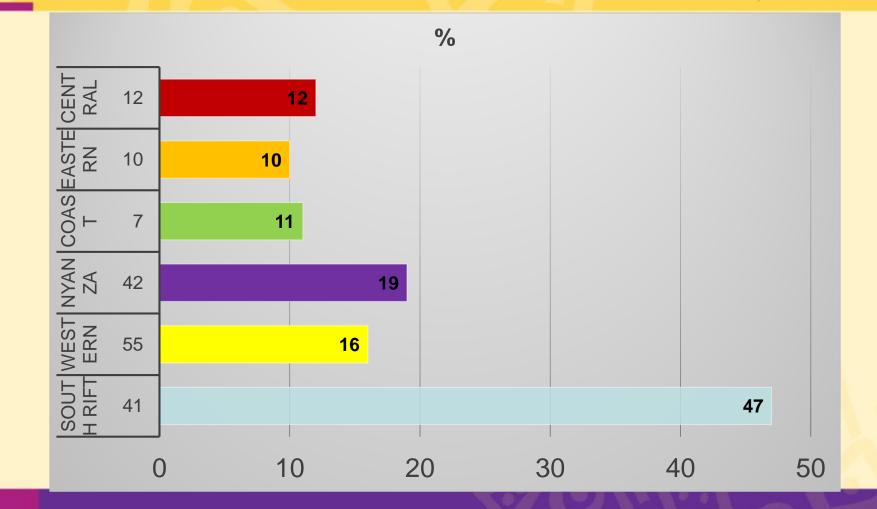
- Survey Western, RV, Eastern, Central and Coastal-25 Counties
- Jan-Dec,2016/2017
- GPS location, photographs
- Vines established in screenhouse
- Graft inoculation
- NCM-ELISA
- Molecular tools:

PCR,

 $\cap_D \cap R$

RT-PCR and

Percentage of begomovirus positive samples collected from different sweetpotato growing regions in Kenya



Security and Health in Africa

Yield impact on selected sweetpotato varieties

- Two season yield trial KALRO Kiboko
- Varieties: Kakamega and Ejumula
- Treatments combinations of: Begomo, SPFMV SPCSV, and controls
- Expt design: RCBD, 3 replicates, 4 rows/block, 10 plants
- Data collection: Stand count ,length of vines, weight of vines and weight of tubers, tuber quality
- Visual symptoms will be recorded biweekly
- Virus testing of will be done at end of trial

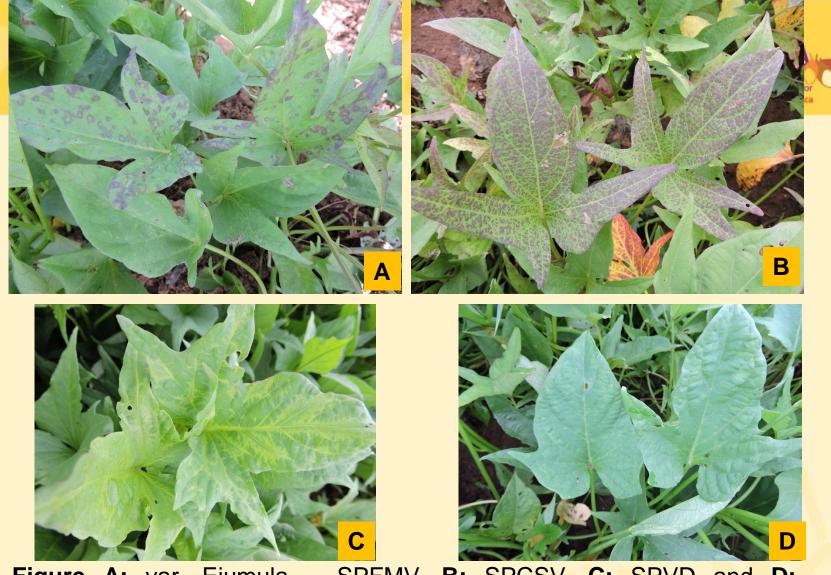


Figure A: var. Ejumula – SPFMV, B: SPCSV, C: SPVD and D: begomoviruses

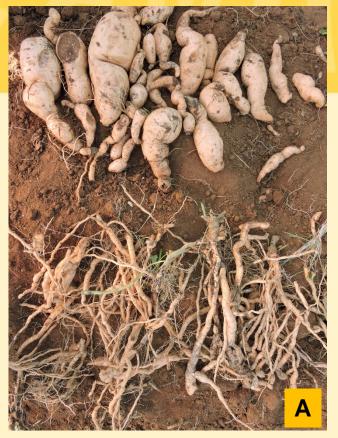






Figure A: Top marketable and bottom non-marketable roots of var. Ejumula infected with SPCSV + begomoviruses

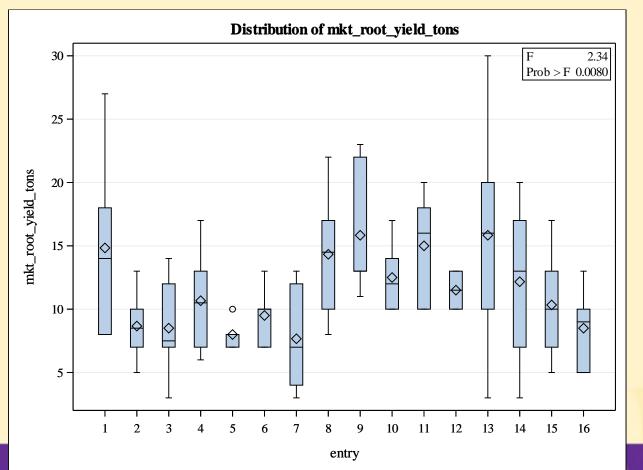
Figure B: Top marketable and bottom non-marketable roots of var. Kakamega infected with SPCSV + begomoviruses





Figure A: Roots of var. Ejumula infected with SPVD + begomo and B: healthy roots Ejumula

Figure . Biplot root yield for the two seasons. 1- Healthy Ejumula, 2- SPCSV infected Ejumula, 3- SPEMV infected Ejumula, 4- Begomo+SPCSV infected Ejumula, 5- Begomo+SPFMV infected Ejumula, 6- SPCSV+SPFMV infected Ejumula, 7- Begomo+SPFMV+SPCSV infected Ejumula, 8- Begomo infected Ejumula, 9- Healthy Kakamega, 10- Action for SPCSV infected Kakamega, 11- SPFMV infected, 12- Begomo+SPCSV infected Kakamega, 13- Begomo+SPFMV infected Kakamega, 14- SPCSV+SPFMV infected Kakamega, 15- Begomo+SPCSV+SPFMV infected Kakamega and 16- Begomo infected Kakamega



Findings



- Occurrence of begomo ranged from 10-50 %
- Seasonal variations in the trial rendered most of the differences statistically non-significant
- Yield reduction was observable in 'Kakamega' when begomoviruses were infecting it which was not evident for 'Ejumula'
- Var. 'Ejumula' was affected more by SPFMV and SPCSV infections
- Single infection with SPFMV/SPCSV/ begome had a high number of nonmarketable roots
- High above ground biomass did not translate to high root yield

Recommendations



Establish genetic diversity of begomoviruses

 Screen more varieties for begomovirus resistance

 Test for begomoviruses prior to import/export plant material



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