

Maize lethal necrosis in East Africa : tracking an emerging disease using NGS

Neil Boonham, Ian Adams, Adrian Fox and Julian Smith

Detection and Surveillance Technologies team

Fera Science Ltd

Maize diseases in East Africa

- New disease noted in 2011
- Samples tested in various labs in Africa/US
- No conclusive agent detected

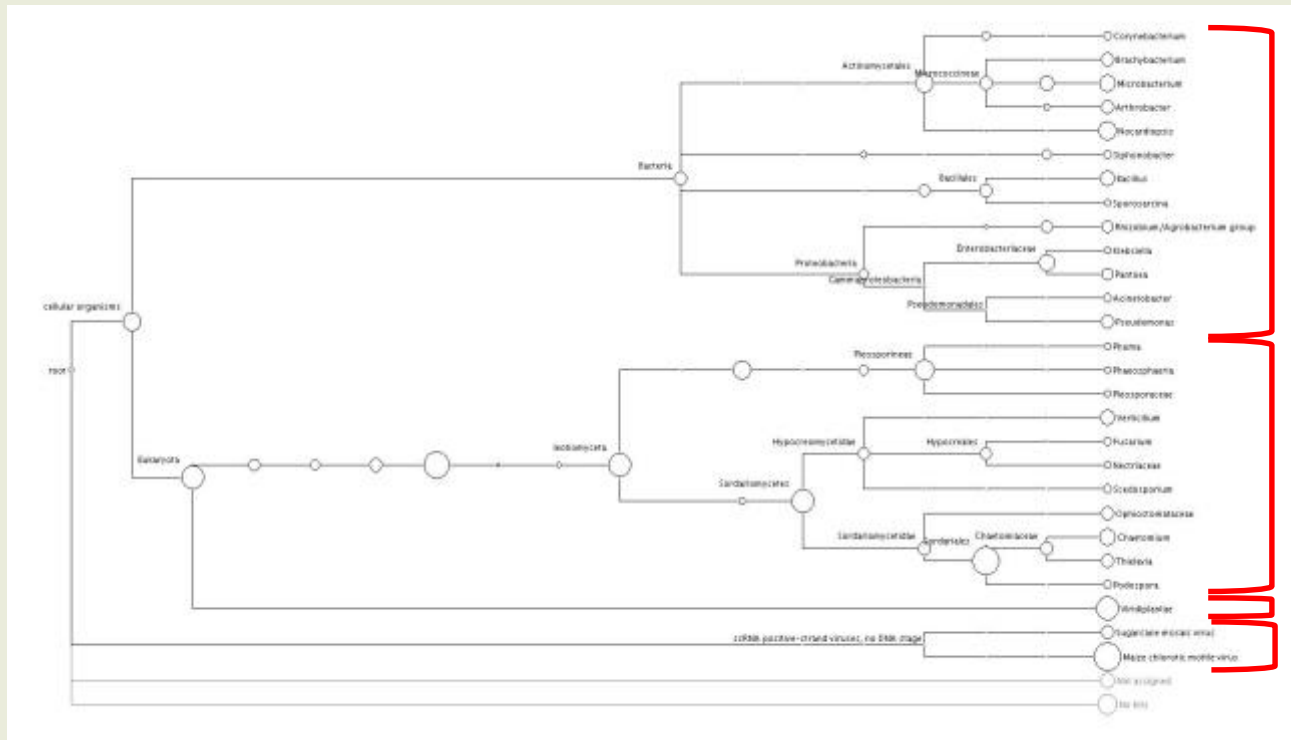


Samples tested using NGS



Adams IP1, Glover RH, Monger WA, Mumford R, Jackeviciene E, Navalinskiene M, Samuitiene M, Boonham N. (2009) Next-generation sequencing and metagenomic analysis: a universal diagnostic tool in plant virology. *Mol Plant Pathol.* 2009 Jul;10(4):537-45. doi: 10.1111/j.1364-3703.2009.00545.x.

Identified two viruses



Bacteria

Fungi

Plant Virus

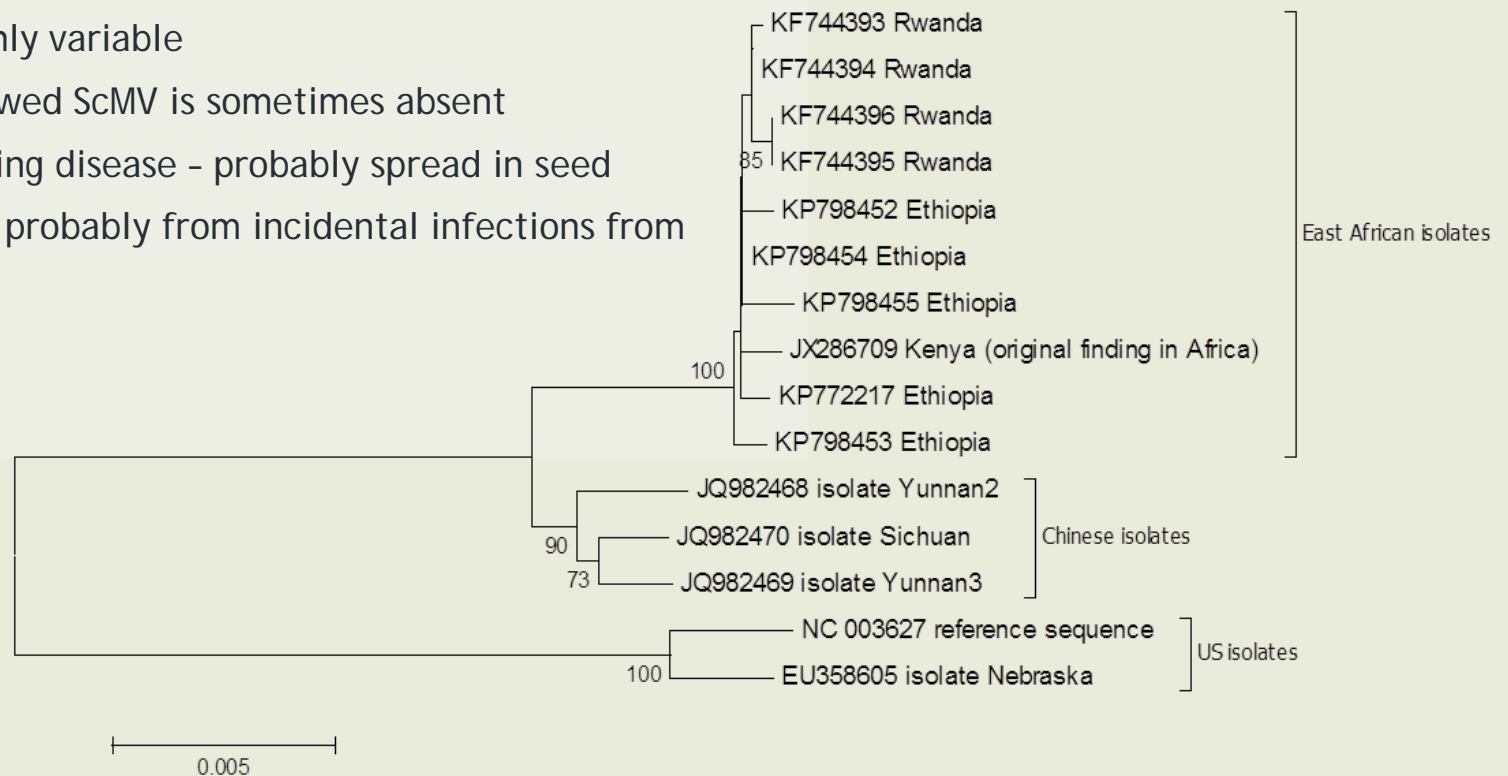
Confirmed the cause

- Back testing confirmed the presence of two viruses
- Satisfied Kochs postulates by mechanical inoculation
- Maize lethal necrosis
 - Maize chlorotic mottle virus (MCMoV)
 - Sugarcane mosaic virus (ScMV)
- Helped identification of likely vectors
- Developed a seed test and trained Kephis/Kari staff



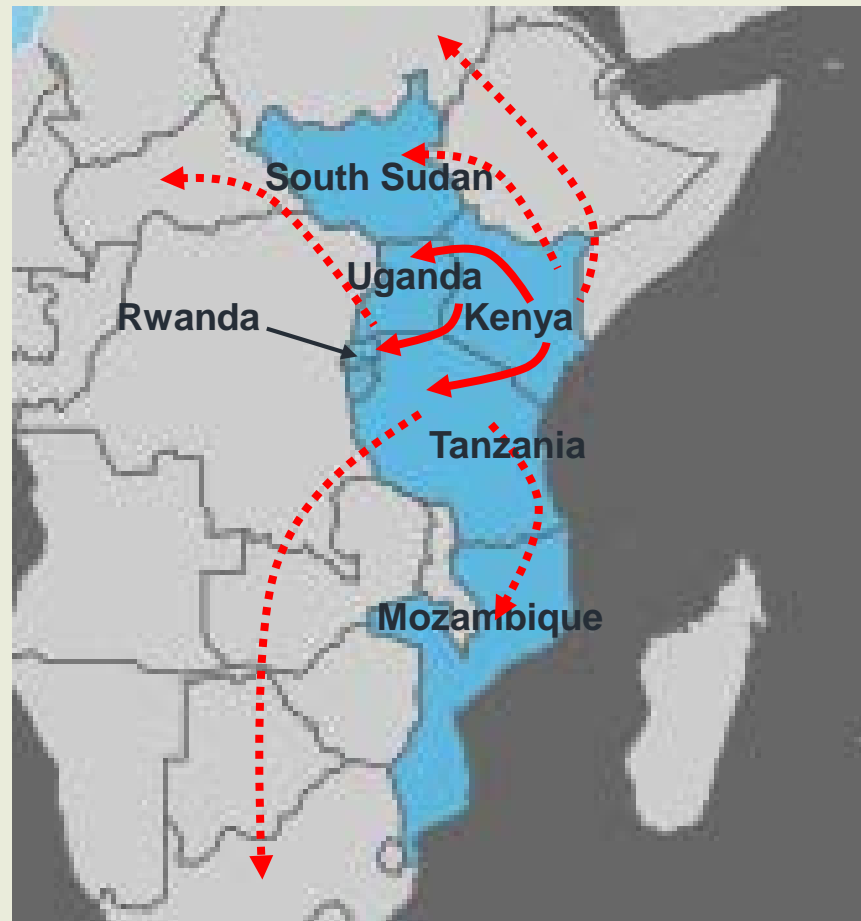
Tracking the disease

- MCMoV is highly conserved
- ScMV is highly variable
- Testing showed ScMV is sometimes absent
- MCMov driving disease - probably spread in seed
- Potyviruses probably from incidental infections from weeds



Adams, I. P., Harju, V. A., Hodges, T., Hany, U., Skelton, A., Rai, S., Deka, M., Smith, J., Fox, A., Uzayisenga, B., Ngaboyisonga, C., Uwumukiza, B., Rutikanga, A., Rutherford, M., Ricthis, B., Phiri, N. & Boonham, N. 2014. First report of maize lethal necrosis disease in Rwanda. *New Disease Reports*, 29.

Disease spread



Confusion with other diseases

- *Sugarcane mosaic virus* not always present in MLN plants
- *Maize yellow mosaic virus* commonly found

Independent site	Maize chlorotic mottle virus	Sugarcane mosaic virus	Other Potyvirus	Maize yellow stripe virus	Maize streak virus	Maize yellow mosaic	Field Diagnosis	Lab Diagnosis
Kenya	Present	Present	Present			Present	MLN	MLN
Kenya	Present						MLN	
Kenya	Present					Present	MLN	
Kenya	Present					Present	MLN	
Ethiopia	Present	Present		Present		Present	MLN	MLN
Ethiopia	Present	Present				Present	MLN	MLN
Ethiopia	Present	Present				Present	MLN	MLN
Ethiopia	Present	Present					MLN	MLN
South Sudan					Present	Present	MLN	
Rwanda	Present	Present					MLN	MLN

Deconstruct the maize virome

	Crop	samples	known plant viruses	Novel plant viruses	total samples
Farm 1	maize	4	Maize chlortic mottle virus, Sugarcane mosaic virus, Maize yellow dwarf mosaic virus	Tombusvirus, Carmovirus, Foveavirus, Closterovirus, betaflexivirus, positive strand ssRNA virus	30
	others	26	bean comon mosaic virus, Beet pseudoyellows virus, Maize yellow dwarf mosaic virus, SCMV, Potato virus S	Caulumoviridae virus, Chrysovirus, Crinivirus, Potyvirus(es), Tombusvirus, unclassified ssRNA positive strand virus, Varicosavirus, Filoviridae virus	
Farm 2	maize	9	Maize chlortic mottle virus	chrystovirus luteovirus ,Carmovirus, tombusvirus, virus, positive strand ssRNA virus, unclassified virus	29
	others	20	Shallot latent virus, Cauliflower mosaic virus,	Chrysovirus, Crinivirus, Cytorhabdovirus, Waikavirus, Varicosavirus polerovirus, polerovirus associated RNA, Tymoviradae virus, positive strand ssRNA virus	
Farm 3	maize	6	Maize chlortic mottle virus, Maize yellow dwarf mosaic virus	Badnavirus, polerovirus associated RNA, Tymoviradae virus	26
	others	20	Turnip mosaic virus,	Badnavirus, Chrysovirus, Cytorhabdovirus, unclassified ssRNA positive strand virus	
Farm 4	maize	4	Maize chlortic mottle virus, Maize yellow dwarf mosaic virus	none	30
	others	26	Banana streak virus, Apple stem grooving virus, Citrus tristeza virus	Badnavirus, potyvirus(es), Tombusvirus, Rhabdoviridae virus, positive strand ssRNA virus, unclassified virus, Varicosavirus	
				total	115

Nanopore sequencing

- Inexpensive sequencer
- Long read lengths
- Cloud based analysis
- Potential in resource poor labs where a large sequencer is impractical/expensive



MinION against MiSeq

- Sequencing errors causing problems for *de-novo* detection of new viruses with BLASTx

Analysis	MiSeq	MinION
BLASTn/x	1 known, 4 novel	1 know, 2 novel

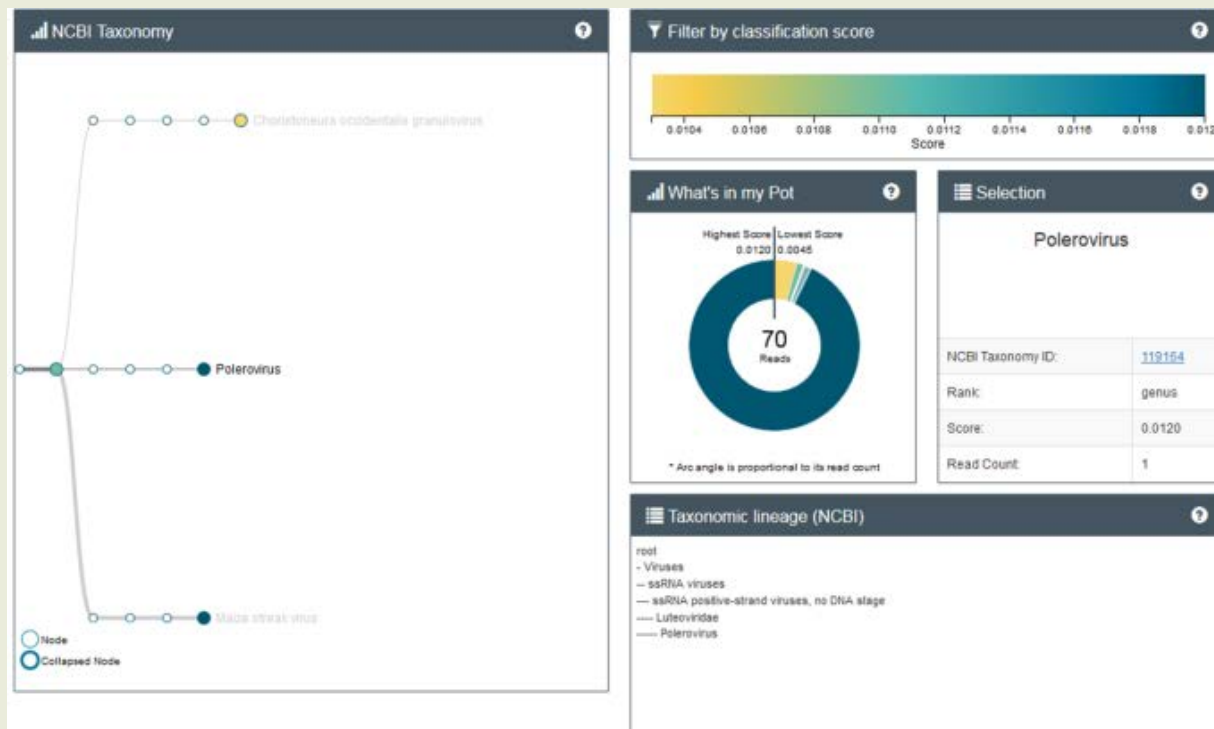
Reference mapping

- Virus sequences were all present in the Minlon data

Virus Sequence	RPKM MiSeq (BWA)	MinION (LAST)	
Maize Yellow Streak virus	7615	11289	Known maize virus
Polerovirus	347	381	Novel, related to Maize yellow dwarf virus
unclassified virus	1220	735	Novel virus
dsRNA virus RNA1	2847	656	Novel virus
dsRNA virus RNA2	24280	5943	
Totivirus RNA1	2619	2056	Novel virus
Totivirus RNA2	146	375	
Totivirus RNA3	211	291	

What's in my pot ?

- Simple analysis pipelines available (WIMP)
- Identified one known and one new virus in samples from Ethiopia



Future platforms

- Smaller - Smidgelon
- Larger - Promethlon



Automated solutions

- Sample prep - Zumbador
- Library set up - Voltrax



Collaborators

biosciences
eastern and central africa



Funders

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syngenta foundation
for sustainable
agriculture