

Genetic improvement of cooking bananas for high levels of pro-vitamin A

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INTRODUCTION

- Vitamin A Deficiency (VAD) is a global public health threat
- Vitamin A deficiency puts 140 million children at risk of illness and death – UNICEF
- VAD impairs proper immune function and increases risk of mortality

VAD intervention strategies

-Supplementation

-Food fortification

-Biofortification

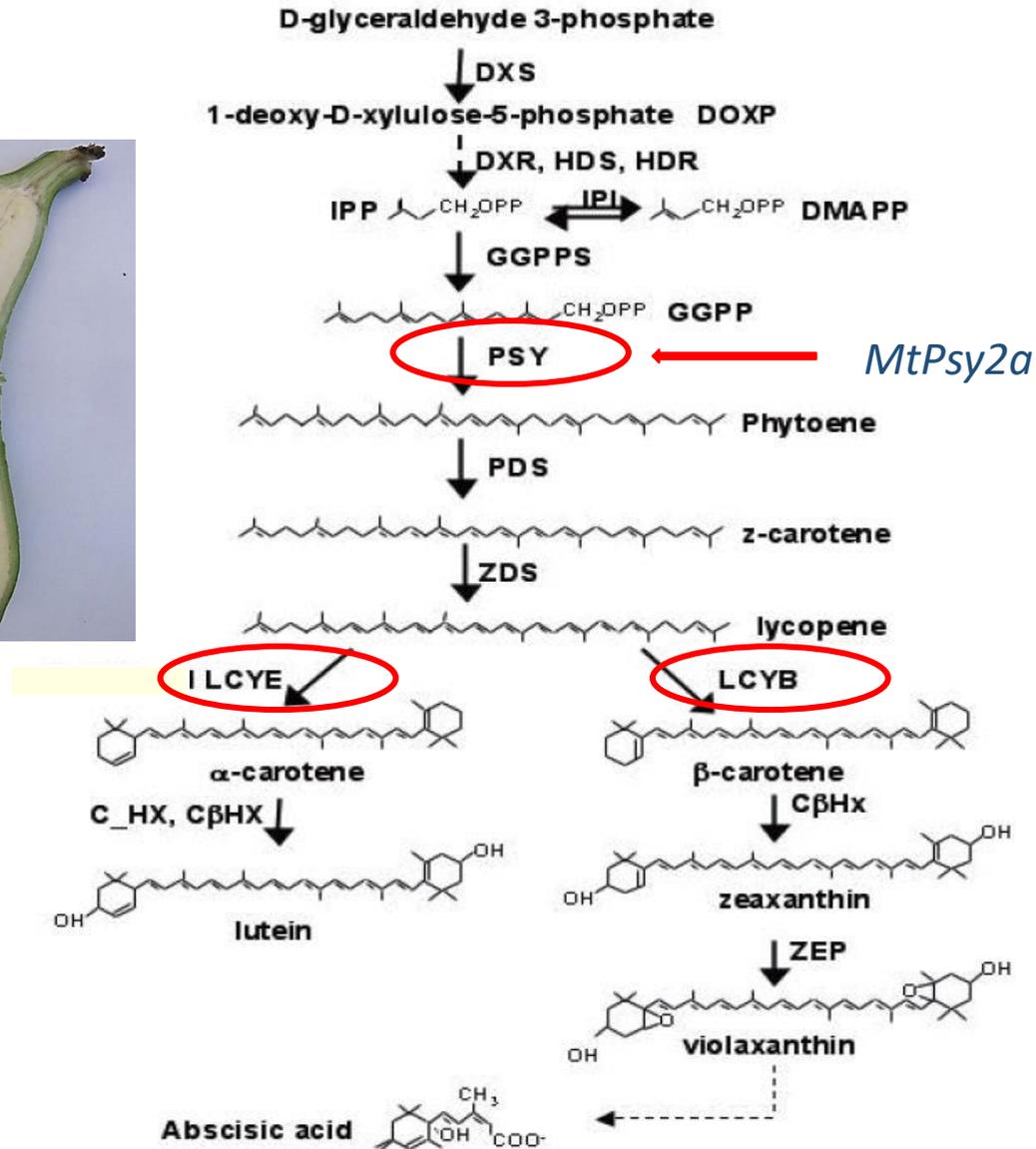
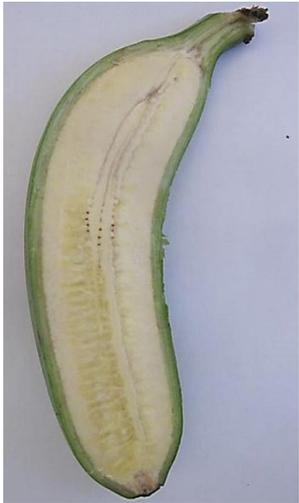


Project goal

- The aim of this program is to alleviate vitamin A deficiency (VAD) in Uganda through biofortification of a major staple food crop, banana



Project strategy



Asupina: a naturally high pVA banana from Papua New Guinea



Promoters:

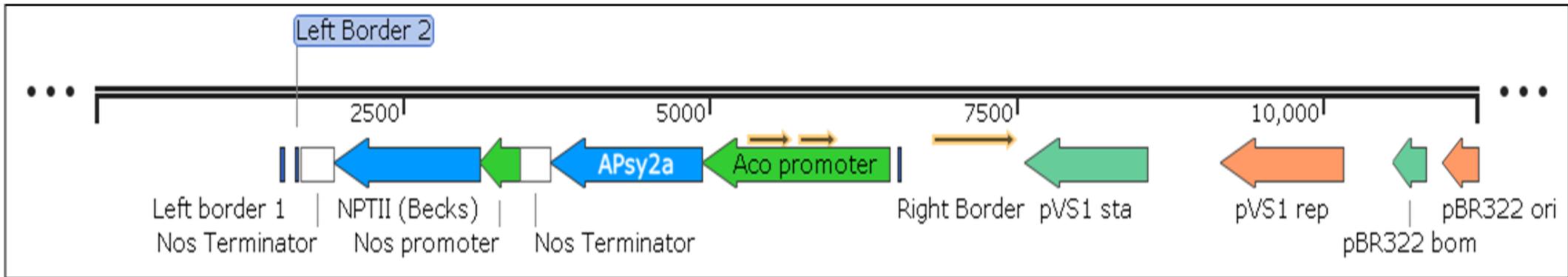
Aminocyclopropane-1-carboxylate oxidase (ACO): Banana Fruit preferred

Maize Ubiquitin (Ubi) : Constitutive

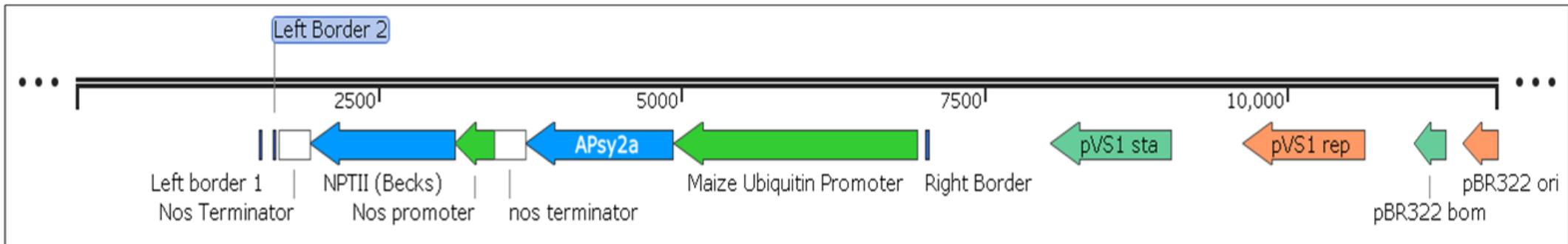
1. ACO-*MtPsy2a*
2. Ubi-*MtPsy2a*

Constructs used in product development

ACO-MtPsy2a



Ubi-MtPsy2a



Key milestones achieved

- Transformable and regenerable embryogenic cell suspensions established for both hybrid M9 and Nakitembe
- Efficient transformation protocols developed
- Transformants generated from both cultivars with both constructs
- Early event selection trial completed at NARL, Kawanda



M9 CONTROL

M9
TRANSGENIC



5 μg

22 μg

NAKITEMBE
CONTROL

NAKITEMBE
TRANSGENIC



12 μg

35 μg

AVERAGE
 β -CAROTENE
EQUIVALENTS



Stage 1: Early event selection

- **Selection based on:**
 - Fruit beta-carotene equivalent (β -CE) concentration
 - 20 μ g/g (dry weight) or above
 - Minimum bunch weight of 20% of the respective control
 - Hybrid M9 - 16 kg
 - Nakitembe - 10 kg

Selection	Hybrid M9	Nakitembe
Total number of lines in the field	356	162
Number of lines harvested (at least one crop)	314	121
Number of lines analysed by HPLC (at least one crop)	310	99
Number of lines meeting PVA target (at least one crop)	234	85
Number of lines meeting yield target (in at least one crop)	153	57
Total No. of lines meeting both β-CE and yield targets (at least one crop)	61	52



Stage 2: Advanced molecular event selection

- **Detailed molecular and chemical analysis of the lines selected from Stage 1**
 - Number of T-DNA insertions (copy number)
 - Plasmid backbone integration into host genome
 - Identification of insertion sites: Illumina whole genome sequencing
 - Fruit compositional analysis



Elite M9 events selected for ML-CFTs

No.	Plant ID	Vector	Copy #	Plant crop		Ratoon crop 1		Ratoon crop 2	
				Bunch (kg)	β -CE ($\mu\text{g/g dw}$)	Bunch (kg)	β -CE ($\mu\text{g/g dw}$)	Bunch (kg)	β -CE ($\mu\text{g/g dw}$)
1	M9-12083	P	2	23	31.3	14	25.4	7.5	26.2
2	M9-12104	P	1	17	21.8	2	34.9	22	12.8
3	M9-12113	M	1	21	25.6	10	20.7	8	6
4	M9-12152	P	1	15.5	23.5	7.5	27.8	14.5	24.1
5	M9-12141	P	2	15	31.4	10	24.5	8	17.2



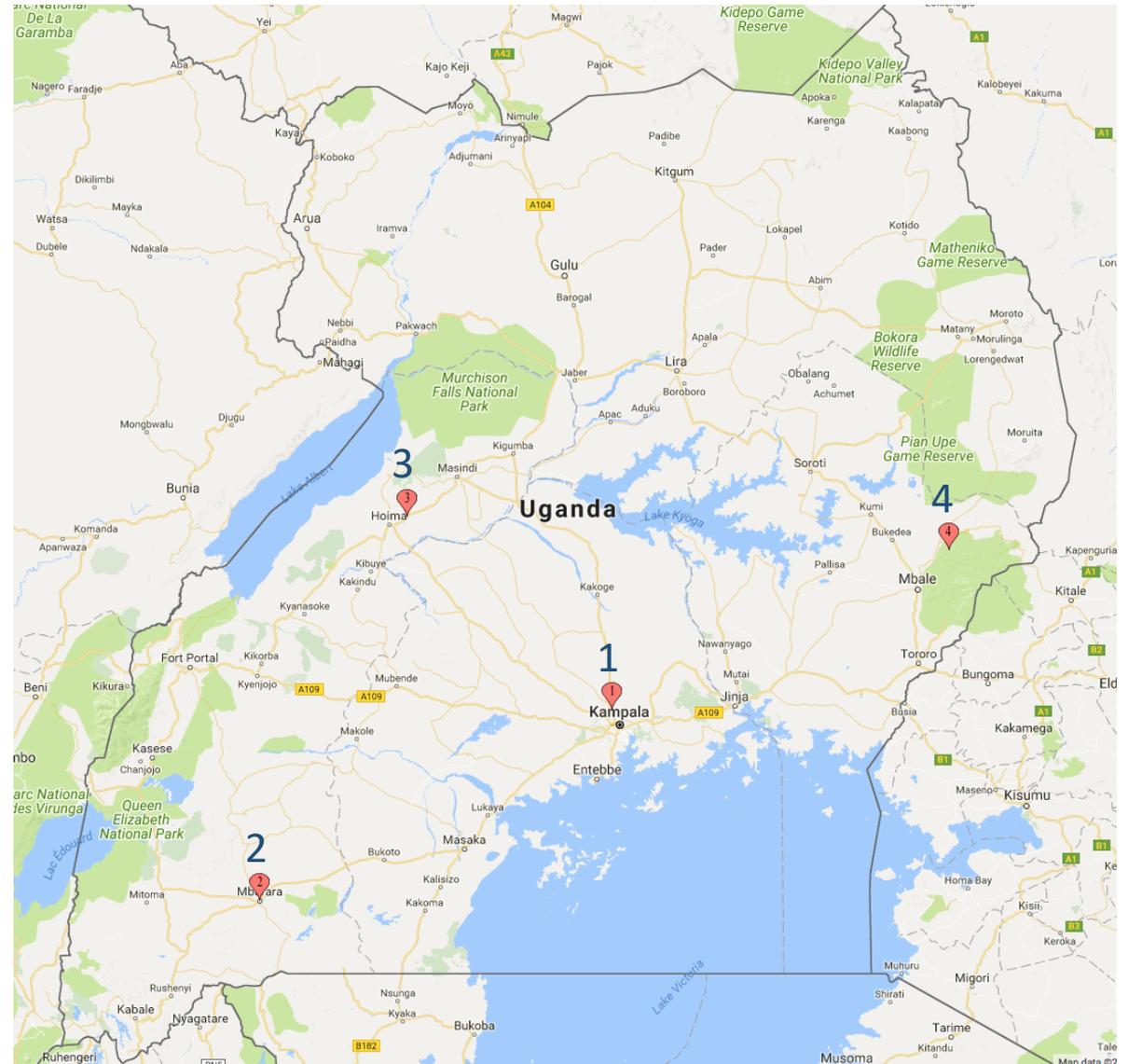
Elite Nakitembe events selected for ML-CFTs

No.	Plant ID	Vector	Copy #	Plant crop		Ratoon crop 1		Ratoon crop 2	
				Bunch (kg)	β -CE ($\mu\text{g/g dw}$)	Bunch (kg)	β -CE ($\mu\text{g/g dw}$)	Bunch (kg)	β -CE ($\mu\text{g/g dw}$)
1	NKT-12418	P	1	21	18.6	22	41.5	40	34.9
2	NKT-12431	P	1	15	19.7	16	57.9	28.5	44.7
3	NKT-12468	P	1	14	39.9	28	45.9		
4	NKT-12477	M	1	16	41	25	34.4		
5	NKT-12478	M	1	19	62.6	23.5	49.7		

Multi-location confined field trails

Approved ML-CFTs sites:

- 4 locations:
 - Kawanda (1)
 - Mbarara (2)
 - Bulindi (3)
 - Buginyanya (4)





Regulatory field trial design

- 4 locations: Kawanda (1), Mbarara (2), Bulindi (3) and Buginyanya (4)
- 4 x 5 randomized block design

Cultivar	Number of Lines	Number of copies	Total per site	Total for 4 sites
M9 Transgenics	5	20	100	400
M9 TC controls	1	20	20	80
M9 Cell Line Control	1	20	20	80
Sub-total (M9)			140	560
NKT Transgenics	7	20	140	560
NKT TC controls	1	20	20	80
NKT Cell Line Control	1	20	20	80
Sub-total (NKT)			180	720
Grand total (M9 and NKT)			320	1280

- Planted M9 in October 2019, and Nakitembe in February 2020



Regulatory field trials – Data collection

Data being collected for the deregulation dossier

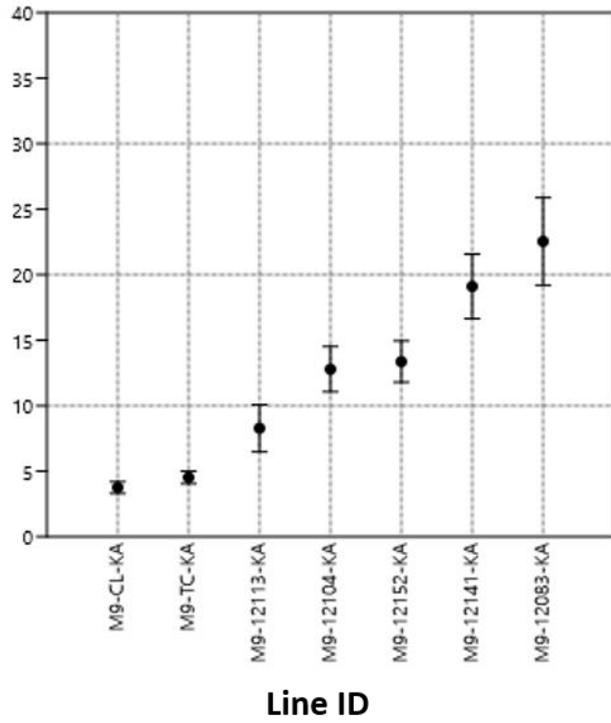
The following data for plant crop and ratoon 1 are being collected

- Agronomic data
- PVA levels in FG green fruits
- Fruit compositional analysis
- Transgene mRNA expression
- ELISA to measure MtPsy2a and NPTII proteins in green, ripe and cooked fruits

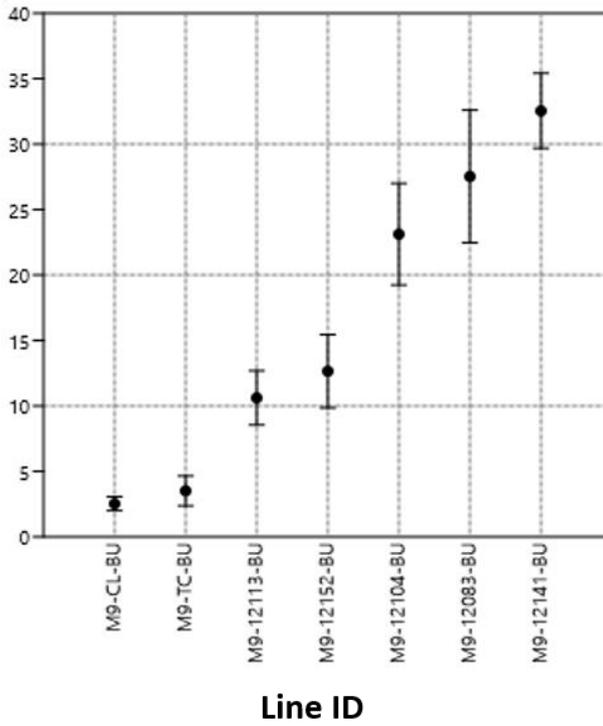


Mean BCE levels across sites – M9, PC

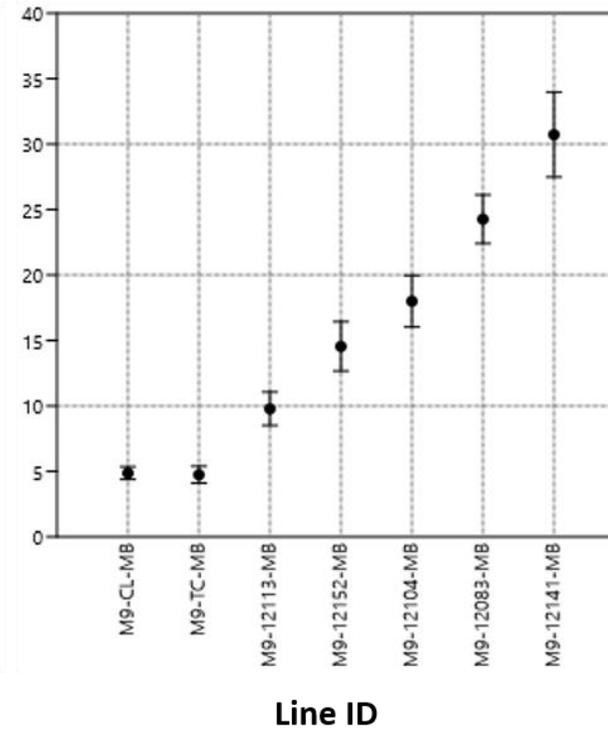
KAWANDA



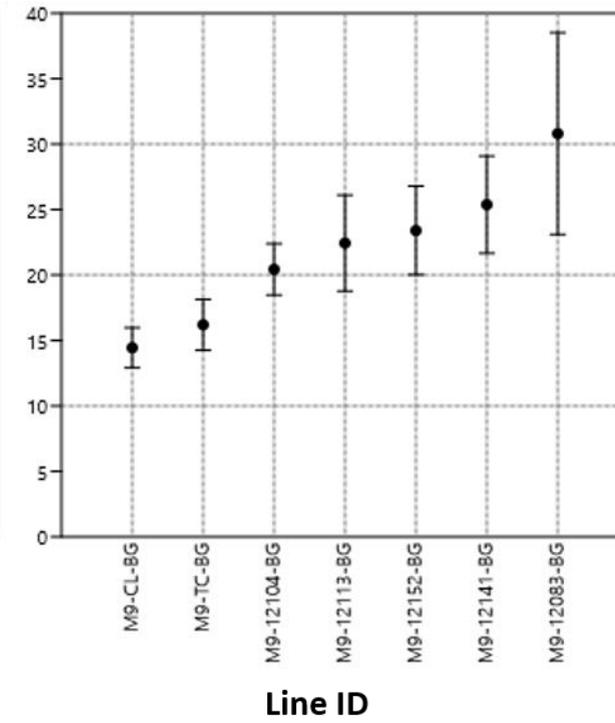
BULINDI



MBARARA



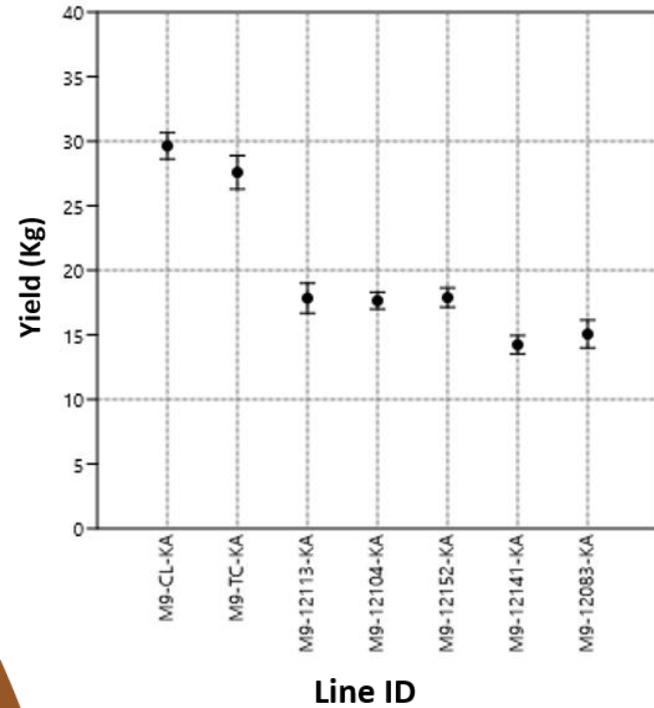
BUGINYANYA



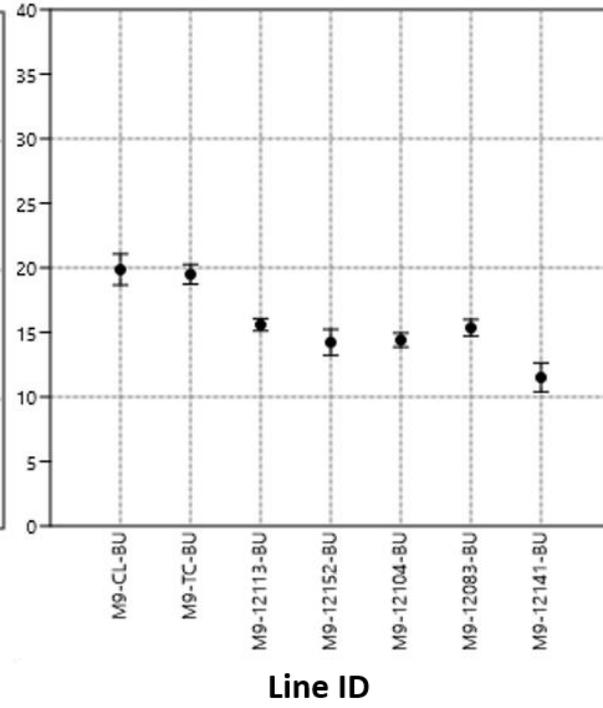


Mean yield levels across sites – M9, PC

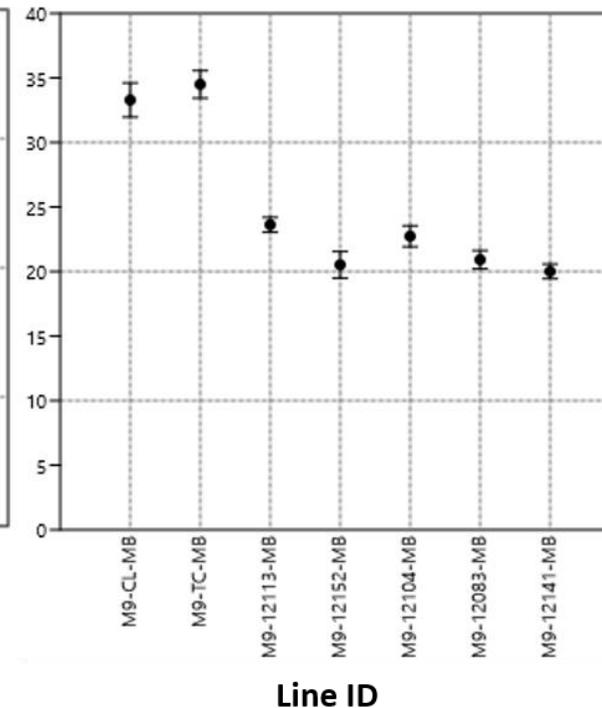
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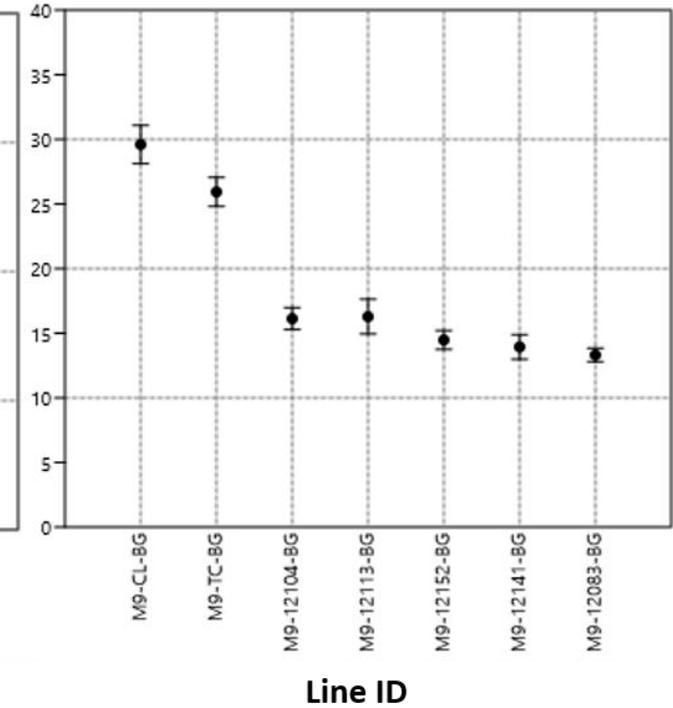
BULINDI



MBARARA



BUGINYANYA





Other ongoing/accomplished activities

- DUS data collection by MAAIF
- Communication - media and community engagement
- Environmental impact assessment

Community engagement





Quality control and assurance mechanisms

- Constant update and backup the BananaTracker and MusaBase, physically and iCloud-based
- Developed and applied line specific PCRs in the greenhouse and field early after planting
- Developed and applied a barcode-based electronic data collection system
- Developed/updated SOPs for all lab and field procedures and processes
- Continuous training of all staff in biotech stewardship, risk assessment and regulation

Acknowledgements



Wilberforce Tushemereirwe

**NARL-KAWANDA
BULINDI-ZARDI
BUGINYANYA ZARDI
MBARARA ZARDI**



**BILL & MELINDA
GATES foundation**



Australian Government
**Department of Agriculture,
Fisheries and Forestry**



James Dale



**National
Biosafety
Committee**



Thank you for listening