



**Preliminary results  
of the heterosis trial  
in Mozambique with  
clones derived from  
Ugandan inter- and  
intra- gene pool  
crosses**

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# Objective



- To demonstrate family means for inter genepool crosses 8A x 8B (64 families) and intra-genepool crosses 8A x 8A ( $(n * n-1) / 2 = 24$  families) as well as intra-genepool crosses 8B x 8B ( $(n * n-1) / 2 = 24$  families) under drought stress conditions – with other word heterosis increments under drought stress conditions
- **Note:** this is material in early breeding stage in which **no** selection has been carried out before

# Materials



A diversity study with 18 microsatellites (SSR markers) identified two gene pools in Ugandan parental material (polycross of Robert).

- Population A
- Population B

8 parents were chosen from each population to constitute the crosses.

- INTER\_GENE POOL CROSSES: 8 X 8 factorial design
- INTRA\_GENE A CROSSES: 8 X 8 DIALLEL without self- and considering reciprocals
- INTRA\_GENE B CROSSES: 8 X 8 DIALLEL without self- and considering reciprocals

# Parents selected



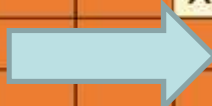
Population A			Population B		
Name	Code	SPVD resistance	Name	Code	SPVD resistance
Ejumula	A1	Susceptible	Resisto	B1	Susceptible
NASPOT1	A2	Moderate	Magabali	B2	Susceptible
Dimbuka-Buku	A3	Susceptible	NASPOT5	B3	Moderate
NASPOT5/58	A4	Susceptible	Wagabolige	B4	Moderate
NASPOT7	A5	Moderate	Mugande	B5	Moderate
SPK004	A6	Moderate	NASPOT11	B6	Moderate
NASPOT100	A7	Moderate	New Kawogo	B7	Moderate
NK259L	A8	Moderate	Huarmeyano	B8	HR SPFMV



# Number of clones per family obtained in Mozambique from seed from Uganda



	A1	A2	A3	A4	A5	A6	A7	A8	B1	B2	B3	B4	B5	B6	B7	B8
A1	X	45	47	42	49	49	41	42								
A2		X	0	38	44	44R	45R	16R								
A3			X	47	12R	0	16R	47								
A4				X	48	47	0	38								
A5					X	14R	35	44								
A6						X	42	33								
A7							X	42								
A8								X								
B1	50	44	45	24	46	48R	47	49	X	50	46	45	40	50	47	45
B2	49	42	46	47	46	49	50	48		X	47	48	44	42	44	47R
B3	49	43	48	0	48	49	49	49			X	49R	46	0	41R	49
B4	47	46	45	47	47	38	21	47R				X	48	49	43	49
B5	48	47	41	49R	48	48	46	49					X	50	49	46
B6	42	46	47	30	50	47	48	49						X	29	42
B7	35	48	49R	0	47	47	0	50							X	49
B8	48	47	47	43R	45	44	48	46								X



**INTRA\_A CLONES**

**INTRA\_B CLONES**



**INTER\_AxB CLONES**

# Experimental material



<b>Population</b>	<b>A</b>	<b>B</b>	<b>AxB</b>	<b>Checks</b>
Number of clones planted	898	1124	2287	Delvia Gaba-Gaba Irene Namanga Resisto

- A check was planted after every 50 clones
- 1 m row plots, 2 plot replications, 2 treatments (irri and no irri)
- RCBD with checks

# Irrigation regimes



## Irrigated treatment

Dates of planting: 8 – 22 September

- 40 mm water applied per each irrigation cycle
- Trials were irrigated 7 times

## Not irrigated treatment

Dates of planting: 21 August – 5 September 2014

- Trials were irrigated 5 times from planting to a month after planting.
- The trial was not irrigated from a month after planting.

# Results



Drought Intensity Index (Fischer and Maurer 1978)

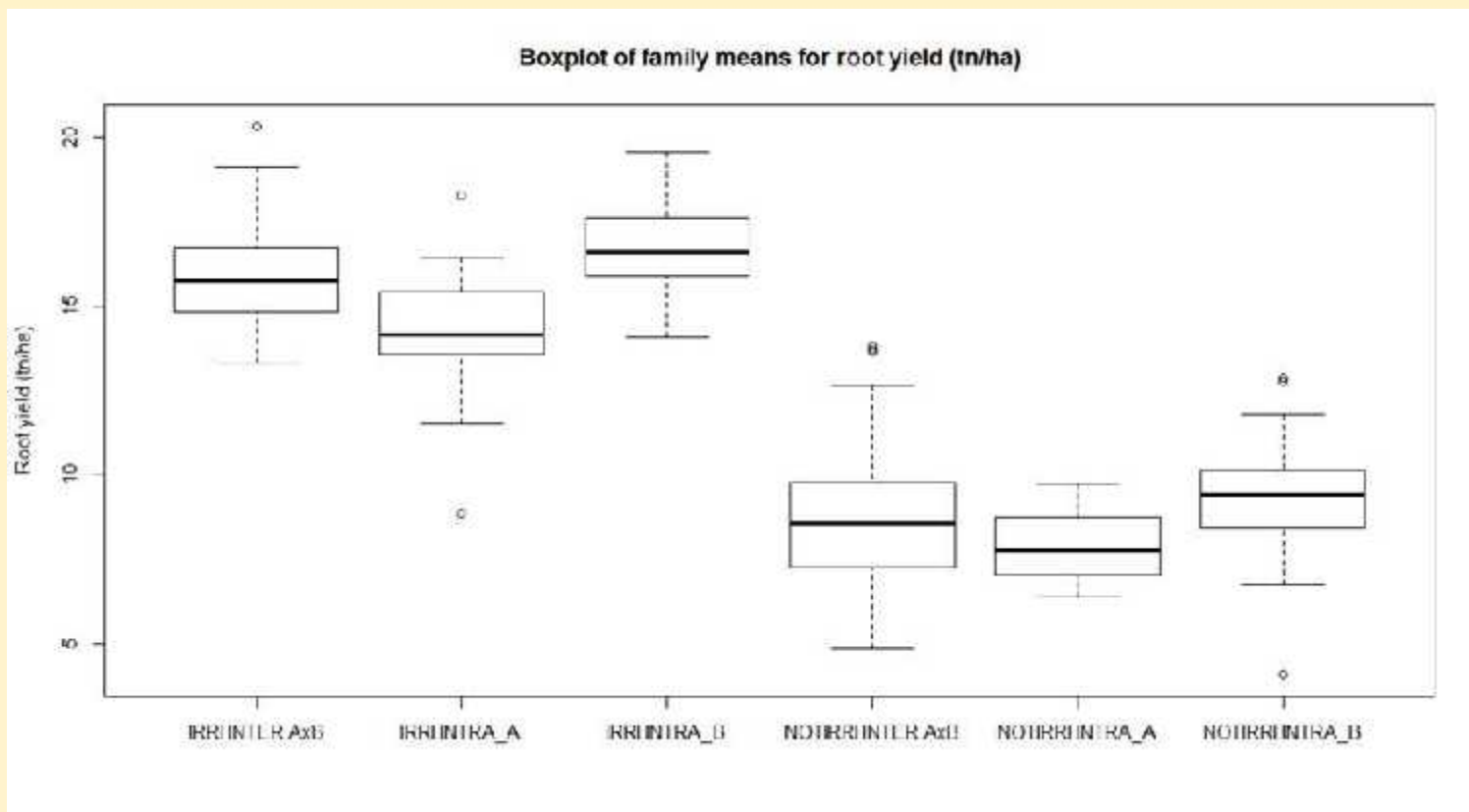
$$DII = 1 - (\text{Mean [NOIRRI]}/\text{Mean [IRRI]})$$

	INTER_AxB	INTRA_A	INTRA_B
DII	0,46	0,45	0,45

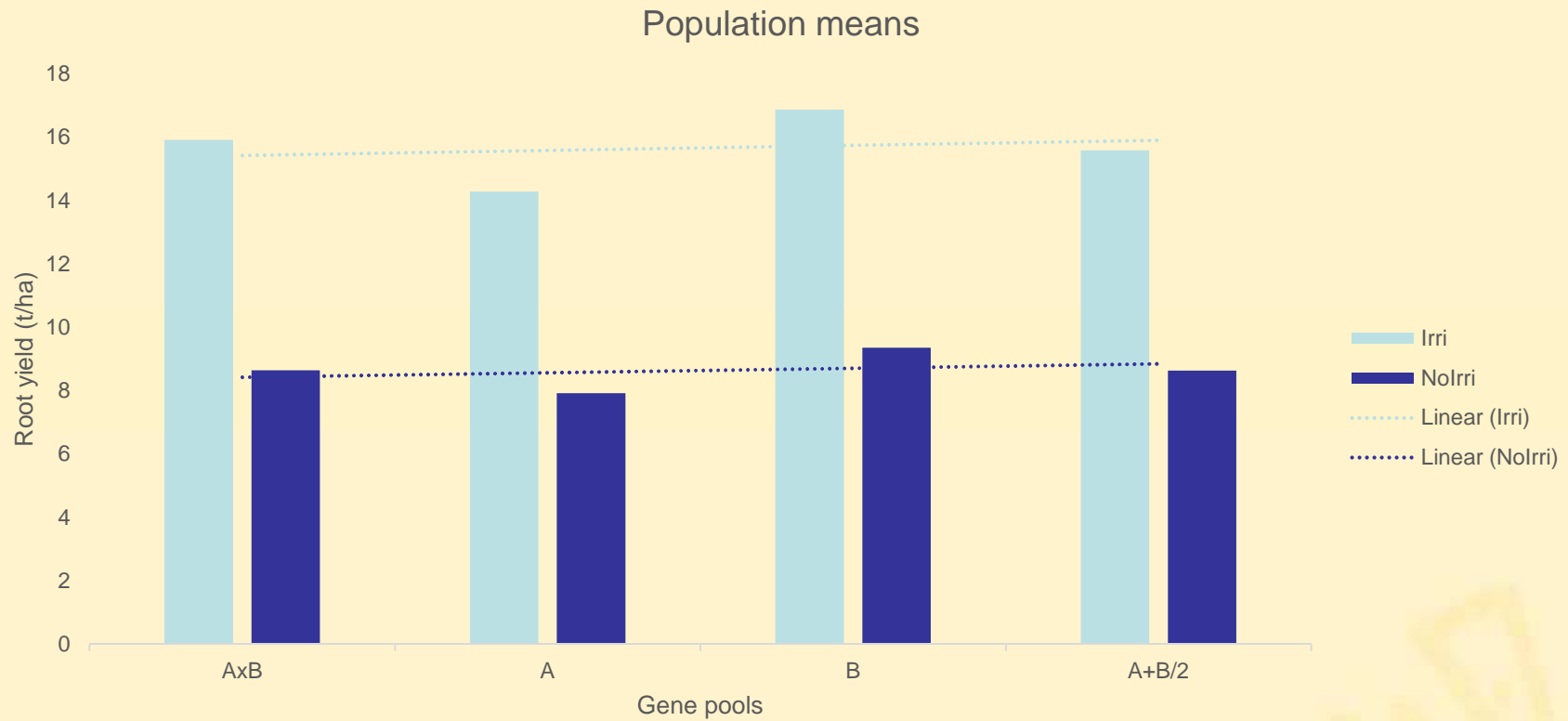
**NB:** All clones from the 3 populations were randomised and planted together



# Family means for root yield (tn/há)



# Comparison of population means among gene-pools and $(A+B)/2$



# Conclusions and perspectives



- In our case, we didn't plant parents for mid-parent mid-offspring heterosis
- The populations can be maintained as reference population for one further evaluation
- We are not sure on whether the parents were randomly selected for the study (question for Robert ?)
- We found heterosis increment on average for the AxB population and more "good" families in AxB than in A and B population
- The AxB population appeared to be more stable than the A and B population – especially under drought "good" families were observed

# What we do next?



- Further data checking
- For sure we will select a larger number of clones for preliminary yield trials from families which were performing well under drought stress
- We need more information on parents
- We still have all the clones in the greenhouses and can utilize the 2015 winter season for example for an efficiency study for the accelerated breeding scheme

# Acknowledgements



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- Maria Andrade for her support and patience
- Raul Eyzaguirre for checking data and the boxplots within few days
- The Bill and Melinda Gates Foundation – for such an opportunity to work on such an experiment – please note I have worked on breeding for drought tolerance before and this has a complete different dimension