

Kenya Agricultural & Livestock Research Organization (KALRO)

Genetic improvement of Sugar Type Common Beans for Bean Scab (*Elsinoë phaseoli*) resistance through Marker Assisted Selection

Shamir Misango ABC- ACP Annual Meeting 2023 Livingstone, Zambia



Team

- Dr. Reuben Otsyula
- Shamir Misango
- Yona Masheti
- Shadrack Odikara
- Diana Gumba
- Faida Kelele





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BACKGROUND OF KALRO PROJECTS

Markers Assisted Selection in developing common bean varieties with multiple resistance to major diseases in Kenya	GENOTYPE	Yield T/Ha
	KK-BC4F3-27	2.118
	KK-BC4F3-13	1.975
	KK-BC4F3-10	1.957
BCMV/NV- MCM2001	KK-BC4F3-28	1.858
CBB- VAX3	KK-BC4F3-20	1.857
Anthracnose- G2333	KK-BC4F3-46	1.828
Pythium root rot Fixed in CAL 194	KK-BC4F3-32	1.827
47 lines with various combinations under AYT	KK-BC4F3-18	1.805
	CAL194	1.799
	RED16	1.799
	KK-BC4F3-31	1.725
	KK-BC4F3-7	1.653
ALL AVERTICAL STREET	KK-BC4F3-44	1.645
	KK-BC4F3-37	1.607
	KK-BC4F3-21	1.605
	KK-BC4F3-23	1.59
	KK-BC4F3-33	1.58
	KK-BC4F3-3	1.533
	KK-BC4F3-6	1.47
	KK-BC4F3-2	1.455
	KK-BC4F3-30	1.417
	KK-BC4F3-45	1.37



Performance Of Bean Genotypes Under Disease Pressure In Different Environments And Planting Dates In Western Kenya

- Conducted by Yona for his Masters
- Screened for common bean disease across environments
- Studied the effect of planting dates on performance
- Disease pressure across different planting times

Prevalence, Variability and Management af Bean Scab (*Elsinoë phaseoli*) of Common Beans (*Phaseolus vulgaris*) in Kenya

- > Prevalence of *Elsinoë phaseoli* in Kenya
- Variability of Elsinoë phaseoli in Kenya
- Effective management options for the control of bean scab



Identification of Novel Candidate Genes Associated with Scab Disease Resistance in Common Bean

- To determine the phenotypic variation among common bean accessions for scab resistance.
- To identify genetic variants and genes associated with scab resistance using a diverse set of common bean accessions using GWAS analysis approach.
- To develop high-throughput PCR-based markers for MAS targeting scab resistance improvement in common bean.



Characterization and Selection for Anthracnose and *Pythium Root* Rot Resistance in Common Bean Landraces Grown in Kenya

- Genetic diversity of landraces in Kenya (89 landraces)
 - Origin and relationships between landraces have been documented
- Screening for Pythium root rot and Anthracnose resistance
 - Significant differences observed in disease reaction
- > Introgression of diseases resistance in landraces- PRR (KK8) and Anthracnose (G2333)
 - Lines developed are under PYT

Publication: Misango S., Otsyula R. & Arunga E. E. (2022): Resistance to *Pythium* root rot and anthracnose among Kenyan common bean genotypes and marker-assisted introgression of resistance genes, Journal of Crop Improvement, DOI: 10.1080/15427528.2022.2158978





PROPOSED RESEARCH Introduction

- Scab disease is a fungal disease of common beans caused by the pathogen *Elsinoë phaseoli* (Philips, 1994).
- The pathogen was first identified by Jenkins in 1933 which was observed on lima beans
- Bean yield losses of as high as 100% under favorable conditions (Philips, 1994; KALRO Annual Report, 2018).
- Bean scab is the most important disease in western Kenya





Problem statement

- Bean Scab is major constraint to common bean production in western Kenya
- 100% yield losses of common bean (Masheti, 2019, KALRO Annual Report 2018)
- One study conducted on ADPs
- No reported resistant genotypes







Justification

- Breeding resistant varieties to scab disease is the most practical, ecofriendly and economical method to manage the disease
- Need to screen more accessions to identify different reactions
- Developed PCR-based markers need to be validated for use in MAS





Objectives

Main Objective

To develop target market class common bean lines resistant to bean scab

Specific Objectives

- 1. To determine bean scab reaction among common bean accessions
- 2. To validate developed PCR-based markers targeting scab resistance
- 3. To introgress bean scab resistance in sugar type beans



MATERIALS AND METHODS

To determine bean scab reaction among common bean accessions
Plant materials

- 1. 47 lines developed for multiple disease resistance through Markers Assisted Selection; BCMV/NV, CBB, Anthracnose, PRR. (Phase 1)
- 2. ADPs
- 3. Landraces

Exerimental Sites

- ➤3 agroecological zones
 - Lower midland
 - Upper midland
 - Lower highlands

Screen house experiment

• Isolates identified from previous study will be used



Cont'

- Yona's focus:
 - Assessment of phenotypic characteristics
 - Genotypic variations
 - Pathogenicity of *Elsinoë phaseoli* isolates
 - Will aid in screening through inoculation

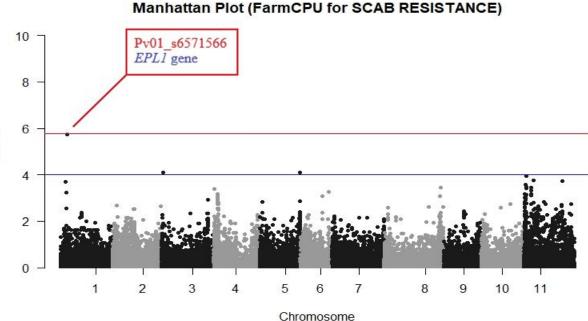






To validate developed PCR-based markers targeting scab

- Shadrack's identification of novel candidate genes associated with scab disease resistance and development of PCR based markers will aid in molecular screening
- Resistant gene locus associated with scab disease resistance to be in chromosome on 1, LOC0003 (MCM 2001)
- primer pairs have been designed and 6 synthesized for each of the targeted genes EPL1, ABC transporter and the PHD-finger genes.



EPL1 gene and ABC transporter associated with scab disease resistance was identified on chromosome one of *Phaseolus vulgaris*.

 $-\log_{10}(p)$

•A sample of the population screened in objective 2 will be used in validation of the primers targeting the *EPL1* gene (enhancer of polycomp-like 1)



Cont'

Plant materials

Selection of 3 sets from the screened panel

- Resistant
- Moderate resistant
- Susceptible
- Primer sets
- 6 primer pairs developed from the previous study



To introgress bean scab resistance in sugar type beans



- Direct crosses between identified potential donors and susceptible sugars with resistance to PRR and anthracnose
- A series of backcrosses to the recurrent parent
- Application of MAS at every stage





- Identification of potential resistant sources among various genotypes
- Development of farmer preferred scab resistant lines
- Potential release of resistant lines
- Enhance the used of designed primer sets in molecular breeding





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