

RESPONSE OF Solanum lycopersicum L. (Tomato) TO Tuta absoluta AND Glomus clarum USING SIMPLE SEQUENCE REPEAT (SSR) MARKER

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INTRODUCTION

Tomato (Solanum lycopersicum L.) belongs to the family Solanaceae and the order Solanales (Singh et al., 2016).

It is sensitive to a number of diseases and attacked by devastating pests of which Tuta absoluta is one of them (Chhetri, 2018)

management of Tuta absoluta but there are clarum) and identification of molecular markers associated with resistance of tomato to Tuta absoluta.

This study aimed at investigating the response of tomato to Tuta absoluta and Glomus clarum using morphological and molecular techniques

MATERIALS AND METHODS

Eight tomato varieties were collected from TechnoServe, NACGRAB and Ojoor Market, Nigeria.

Complete Randomized Design was used with three replicates and three treatments (Glomus clarum + Tuta absoluta, Tuta absoluta alone and

screen house experiment was conducted involving 72 polypots where five (5) grams of G. clarum was inoculated to the varieties and Tuta absoluta was infested on the plants I week after.

Growth, agronomic, and vield characters were determined while percentage incidence was calculated according to the method described by Rasheed et al. (2018) and severity of infestation was assessed using the visual rating scale described by Anjorin et al. (2013).

Amplification of DNA and detection of polymorphisms from young apical leaves were rried out using SSR primers; LEGA003, LECA001, LEAAT005 and LETAA001in

RESULTS AND DISCUSSION

There was significant difference between Glo clarum and Tuta absoluta treated plants across growth, agronomic and yield characters.

Plants treated with G. clarum showed reduced percentage incidence (13%) and severity of infestation (27%) compared to T. absoluta alone (19% and 40%) and control (23% and 35%) as seen in figures 1 and 2 respectively. This could be due to G. clarum which induced activation of tomato defense mechanism, optimum nutrient uptake and damage compensation in accordance with the reports of Olawuyi et al. (2014) and Olowe et al. (2018).

Primer LEAAT005 had the highest allele number (4), gene diversity (0.73), and polymorphic information content (0.67) while the highest major allele frequency of 0.67 was observed in LEGA003.

CONCLUSION AND RECOMMENDATION

Plants treated with Glomus clarum reduced the infestation of Tuta absoluta.

Primers LEAAT005 and LECAA001 detected the highest genetic polymorphism and could be used in marker assisted breeding of tomato and other Solanum spp.

The tomato varieties; NGB00717, NGB00725 (NACGRAB) and Dan Eka Jibia (Technoserve), were tolerant to Tata absoluta and could be recommended for breeding of tomato against Tutu absoluta.

RESULTS

Table 1: Growth, agronomic and yield characters of Tomato as affected by Glomus clarum and Tuta absoluta

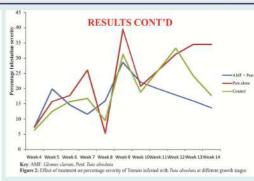
Treatments	Plant beight (cm)	Leaf length (cm)	Number of leaves	Stem girth (cm)	Leaf width (cm)	Presence of flowers	Number of flowers	Number of dead flowers	Presence of fruits	Number of fruits	Fruit fresh weight (g)
Glomus clarum + Tuta sksoluta	95.07*	5.55*	11.97*	0.58*	2.59*	3.64ª	3.78s	1.31*	1.76*	1.184	1.759
luta absoluta alone	85,68°	5,06°	11.60°	0.53°	2.46 ^b	2.97	3.27 ^b	1.27*	1.26*	0.45 ^b	1.13*
Control	89,25 ^b	5.23 ^b	11.64a	0.56 ^b	2,44 ^b	3,46*	3.25 ^b	1.13*	1.56°	0.28 ^b	0.976

RESULTS CONT'D

Table 2: Gene diversity, Allele number, and Polymorphic information content of SSR primers

Molecular Markers	Major Allele Frequency	Sample Size	Allele No.	Gene Diversity	Polymorphic Information Content
LEGA003	0.67	24	3	0.50	0.44
LECAA001	0.46	24	3	0.64	0.57
LEAAT005	0.33	24	4	0.73	0.67
LETAA001	0.63	24	3	0.50	0.41
Mean	0.52	24	3.25	0.59	0.52





RESULTS CONT'D







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