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Research Article

Screening of BC₄F₁, BC₅F₁ and F₁ Generations for *Fusarium* Wilt and SMD Diseases Reaction in Pigeonpea [*Cajanus cajan* (L) Millsp.]

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ABSTRACT

The efforts were made to evaluate BC_5F_1 , BC_4F_1 and $12 F_1$ generations of pigeonpea against two deadly diseases, Fusarium wilt (FW) and sterility mosaic virus (SMD) causing considerable yield losses to the crop. A minimum range of variability in the crosses was observed with respect to the reaction against FW and SMD diseases. The crosses (ICPA-2101-4 × ICPB-2101-4) × ICPB-2101-4, (ICPA-2161-5 × ICPB-2161-5) × ICPB-2161-5 and (ICPA-2043 × Maruti) x Maruti of above generations recorded moderately resistant (MR) and resistant (R) reaction against FW and SMD respectively. This paper reports information for significant use to the breeders and plant pathologists by physical elimination of diseased ones at the early stage, particularly for development of resistant hybrid or variety in pigeonpea.

Key words: Pigeonpea, Fusarium wilt, Sterility mosaic disease, Backcross, Disease reaction.

INTRODUCTION

Pigeonpea (*Cajanus cajan* (L) Millspaugh) is a short lived perennial shrub traditionally cultivated as an annual crop in developing countries. Based on the vast natural genetic variability in local germplasm and the presence of numerous wild Relatives¹, concluded that India is probably primary centre of origin. Globally, pigeonpea is cultivated on 4.92 million hectares (M ha) with an annual production of 3.65 million tons and productivity of 898 kg/ha². India is a major pigeonpea producer in the world with a It plays an important role in food security, balanced diet subsistence agriculture because of its diverse usage in food, fodder, soil

contribution of 75–80 per cent. In India, pigeonpea is cultivated in an area of 3.75 m ha with production of 2.46 million tons and productivity of 656 kg/ha³. In Karnataka, pigeonpea occupies second place in area (0.65 m ha) and ranks second in production (0.26 m tons) with a productivity of 406 kg/ ha³. Gulbarga is very important potential district in the country for extensive cultivation of pigeonpea and hence known as 'pulses bowl' of Karnataka. It is also grown in Bidar, Bijapur, Dharwad, Raichur, Bellary and Belgaum districts of northern Karnataka. conservation, integrated farming systems and symbiotic nitrogen fixation⁴.

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Lower productivity of pigeonpea in India is attributed to factors such as biotic (e.g., *Fusarium* wilt, sterility mosaic and pod borers) and abiotic (e.g., drought, salinity and waterlogging) stresses. Among the biotic stresses, Fusarium wilt and sterility mosaic diseases are considered to be the most important diseases of pigeonpea in India. Fusarium wilt (FW), caused by Fusarium udum Butler, is the major constraint for limiting pigeonpea production in all pigeonpea growing regions^{5,6}. The disease symptoms usually appear when plants are at the pre- flowering and podding stage (100% loss), at maturity (67%), and at pre-harvest stage (30% loss) but sometimes symptoms also appear in 1-2 month-old plants. The FW incidence increases in the ratoon and perennial crops⁷ and causes serious yield losses in susceptible cultivars. The another disease sterility mosaic disease (SMD), often referred to as "Green Plague", as the affected plants are green with excessive vegetative growth but with no flowers or pods, under congenial conditions spreads rapidly leading to severe epidemics⁸. SMD infection at an early stage (<45-day-old plants) results in a 95-100% loss in yield⁹. The disease is confined to Asia and apart from India it has been reported from Nepal, Bangladesh and Myanmar, Thailand and Sri Lanka¹⁰. SMD is caused by Pigeonpea Sterility mosaic virus and is transmitted by an eriophyid mite (Aceria cajani)¹¹. The loss in yield is greater by SMD disease than the Fusarium wilt and even other biotic stresses.Control of Sterility Mosaic Disease (SMD) and Fusarium wilt (FW) by chemical methods though effective, is not feasible

friendly¹². ecoeconomically and non Breeding resistant varieties is considered to be one of the most effective and economic methods of reducing crop losses and has received top priority. There are only few sources of resistance reported to FW and SMD^{13,14}. This paper reveals about the priority research for the development of Fusarium wilt and SMD resistant cultivar by using back cross and F_1 generations in pigeonpea as a source of resistance to both FW and SMD.

MATERIALS AND METHODS

The experiment was conducted at Agricultural Research Station, Kalaburagi during Kharif 2009-2010. The parental material for the present study was obtained from the germplasm collection maintained at Agricultural Research Station, Kalaburagi and ICRISAT, Hyderabad are presented in Table 1 and Table 2. The disease scoring was made for Fusarium wilt and SMD diseases in BC_4F_1 , BC_5F_1 and 12 F_1 's. These populations were sown in single rows of 5 m length with spacing of 90 x 30 cm. The disease reaction was screened under natural epiphytotic condition (normal field), because the seeds generated are few and precious. A 9 point scale divided into five categories is used and adopted by AICRP on Pigeonpea for easy scoring (Table. 3) for both FW and SMD. Interpretation of scale is as follows: 1-Immune, 3-Resistant, 5-Moderately Resistant, 7-Moderately susceptible and 9-Susceptible. The scored values were converted in to per cent infection by using the formula for both FW and SMD.

Number of infected plants

Per cent Disease Index (PDI) = ------ x 100 Total Number of plants

RESULTS AND DISCUSSION

The data recorded on FW and SMD in $BC_4F_{1,}$ BC_5F_1 and the 12 F_1 's populations were screened under natural epiphytotic condition (Table 4 and Table 5). The back cross population comprised of two groups, the first group BC_5F_1 consists of four crosses obtained

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from ICPA-2101 × ICPB-2101. A minimum range of variability in the crosses was observed with respect to the reaction against wilt and SMD. Out of the four crosses observed, one cross (ICPA-2101-4 × ICPB-2101-4) × ICPB-2101-4 showed 5 rating while other two crosses showed 3 rating scale and

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recorded 16.6 per cent reactio	n considered as	mosaic disease and all are considered as
moderately resistant followed	by 6.6 per cent	resistant. A minimum range of variability in all
and 3.3 per cent are considere	d as resistant to	the crosses was observed with respect to the
wilt based on rating scale use	d and for SMD	reaction against wilt and SMD disease.The
no reaction was observed	hence all are	above identified Fusarium wilt and SMD
considered as resistant to steril	ity mosaic virus	resistant lines from natural epiphytotic
disease. The second group BC	₄ F ₁ consists of 6	condition (normal field) can further be tested
crosses obtained from ICPA-	-2161 × ICPB-	under wilt sick plot and SMD infested nursery
2161. Out of the six crosses, i	nterestingly one	(leaf staple method). To develop a resistant
cross (ICPA-2161-5 × ICF	PB-21611-5) ×	hybrid or variety along with crucial yield
ICPB-2161-5 showed 3 rat	ing scale and	parameter for the benefit of farming
recorded 3.3 per cent for the	wilt, while the	community. Hence, no such reports were
cross (ICPA2161-4 \times ICPB-2	161-4) × ICPB-	available. However, the recent reports for FW
2161-4 showed 3 rating and r	ecorded 5.7 per	and SMD by different researchers followed
cent for SMD reaction a	nd which are	wilt sick plot and leaf staple screening
considered as resistant. Amo	ng the 12 F_1 's	methods to identify resistant sources against
screened the only one cross	(ICPA-2043 ×	above diseases. Researchers ^{15,16,17} were
Maruti) \times Maruti recorded 3 i	rating scale and	identified wilt resistant accessions after
showed 3.7 per cent reaction a	against wilt and	screening a pigeonpea mini-core
for SMD 7 crosses recorded 3 i	rating scale with	collection/accessions/germplasm lines,
the variability ranged from 2	2.8-4.7 per cent	respectively. Whereas, for SMD disease,
was observed for reaction aga	inst the sterility	resistance sources were identified by ^{15,16,18} .

Table 1: List of parental	materials used in	BC_4F_1 and BC_5F_1
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Sl. No.	Parents	Source	Generation	Per cent male sterility		
Female parent						
1.	ICPA-2101	BP 13A	BC_4F_1	32		
2.	ICPA-2161	BP 13A	BC_3F_1	42		
Male parent						
1.	ICPB-2101	BP 13A				
2.	ICPB-2161	BP 13A				

Table 2: List of stable CMS lines and restorers used in Backcrossing programme

Sl. No.	Parents	Source	Pedigree			
Female parents						
1	ICPA-2043	ICDISAT Hydershad	Cajanus	cajanifolius		
1.		ICKISAT, Hyderabad	(A ₄ - Cytoplasm)			
2	ICDA 2078	ICDISAT Hydershad	Cajanus	cajanifolius		
2.	2. ICPA-20/8 ICRISAI, Hyderabad		(A ₄ - Cytoplasm), BP 13 ^a	(A ₄ - Cytoplasm), BP 13 ^a		
3	CT 288A	284 S. K. Nagar Guiarat	Cajanus	scarabaeoides		
5.	01-200A	S.K. Nagai, Oujarat	(A ₂ - Cytoplasm)			
		Male paren	ts			
1.	Maruti	ARS, Gulbarga	Selection from germplasm			
2.	Gullyal Red	ARS, Gulbarga	Local germplasm collection	from Gulbarga		
3.	WRP-1	ARS, Gulbarga	GS-1 x Maruti			
4.	TS-3	ARS, Gulbarga	ICP-87051 x PT-221			
5.	TS-3R	ARS, Gulbarga	TS-3 x Maruti			
6.	BRG-3	UAS, GKVK Bangalore	OGUK 3 x ICP 7036			

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 Table 3: Reaction of Pigeonpea (Cajanus cajan) crosses to wilt and sterility mosaic disease (SMD)

Rating scale	Per cent	Disease reading	<i>Fusarium</i> wilt	Sterility Mosaic Disease		
(1-9)	infection					
1.	0	Immune	No symptoms on any plant	No symptoms on any plant		
2.	10	Resistant	10 or less mortality	Symptoms on 10% of fewer plants		
3.	11-20	Moderately resistant	11-20% mortality	Ring spot symptoms on most plants but disappearing with age; no sterility		
4.	20-50	Moderately susceptible	20-50% mortality	Mild mosaic symptoms on most plants causing plant sterility		
5.	> 50	Susceptible	51% or more mortality	Severe mosaic on most plants; almost completely sterility		

Table 4: Screening for wilt and Sterility mosaic disease in BC₄F₁and BC₅F₁ selected plant to plant crosses

Sl. No.	Crosses	Total No.	Wilted plants	Per cent	SMD	Per cent
		of plants		wilt	plants	SMD
BC ₅ F ₁ plant	to plant crosses					
1.	(ICPA-2101-3 × ICPB-2101-3) × ICPB-2101-3	30	2	6.6		
2.	$(\text{ICPA-2101-4} \times \text{ICPB-2101-4}) \times \text{ICPB-2101-4}$	30	5	16.6		
3.	$(\text{ICPA-2101-9} \times \text{ICPB-2101-9}) \times \text{ICPB-2101-9}$	32				
4.	(ICPA-2101-12 × ICPB-2101-12) × ICPB-2101-12	30	1	3.3	0	0
BC ₄ F ₁ plant	to plant crosses					
1.	$(\text{ICPA-2161-4} \times \text{ICPB-2161-4}) \times \text{ICPB-2161-4}$	35			2	5.7
2.	$(\text{ICPA-2161-5} \times \text{ICPB-2161-5}) \times \text{ICPB-2161-5}$	30	1	3.3		
3.	$(ICPA-2161-6 \times ICPB-2161-6) \times ICPB-2161-6$	34				
4.	$(ICPA-2161-6 \times ICPB-2161-6) \times ICPB-2161-7$	33				
5.	$(ICPA-2161-6 \times ICPB-2161-6) \times ICPB-2161-9$	33				
6.	$(\text{ICPA-2161-6} \times \text{ICPB-2161-6}) \times \text{ICPB-2161-10}$	32				
7.	GS-1 (Check-wilt susceptible)	30	4	13.3		
8.	Maruti (Check-SMD susceptible)	28			4	14.3

--: No disease symptoms, SMD: Sterility mosaic disease

Table 5: Screening for wilt and sterility mosaic disease in 12 $F_{1'}s$

Sl. No.	Crosses	Total No. of	Wilted	Per cent	SMD	Per cent
		plants	plants	wilt	plants	SMD
1.	$(GT-288A \times TS-3R) \times TS-3R$	21			1	4.76
2.	$(\text{GT-288A} \times \text{WRP-1}) \times \text{WRP-1}$	26			1	3.85
3.	$(\text{GT-288A} \times \text{MARUTI}) \times \text{MARUTI}$	22			1	4.54
4.	$(ICPA-2078 \times WRP-1) \times WRP-1$	33			2	6.06
5.	$(ICPA-2078 \times TS-3R) \times TS-3R$	35			1	2.86
6.	$(ICPA\text{-}2078 \times G.RED) \times G.RED$	30			1	3.33
7.	$(ICPA-2043 \times TS-3R) \times TS-3R$	34				
8.	$(ICPA\text{-}2043 \times G.RED) \times G.RED$	30				
9.	$(ICPA-2043 \times WRP-1) \times WRP-1$	28			1	3.57
10.	$(ICPA-2043 \times MARUTI) \times MARUTI$	27	1	3.7		
11.	$(ICPA-2043 \times TS-3) \times TS-3$	30				
12.	$(ICPA-2043 \times BRG-3) \times BRG-3$	28				
13.	GS-1 (Check-wilt susceptible)	30	4	13.3		
14.	Maruti (Check-SMD susceptible)	28			4	14.3

--: No disease symptoms, SMD: Sterility mosaic disease

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CONCLUSION

The identified resistance sources could become the pre breeding tools for the breeder to make use in disease resistance breeding programme for developing enhanced disease resistance in different elite but susceptible cultivar of pigeonpea. Further, these resistance sources can be utilized in gene pyramiding for multiple races of *Fusarium* wilt through advanced molecular technology using marker assisted backcrossing (MABC) in future.

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