

Evaluation of Blackgram Germplasm for Resistance against YMV

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Abstract

The experiment carried out at the R&D Farm, NRI Agritech and ANU, Nagarjuna Nagar, Guntur. To identify genetic sources of resistance to yellow mosaic virus (YMV) in blackgram vigna mungo (L), 49 germplasm lines are collected from different geographical regions were evaluated under field conditions during period rabi season November 2014 - February 2015, 2 entries exhibited resistance (R) reaction with rating of 1.0 to 2.0. Six genotypes fall in the category of moderately resistant (MR) with rating scale of 2.1 to 4.0, three were moderately susceptible (MS) with rating of 4.1 to 5.0; two genotypes were susceptible (S) with rating of 5.1 to 7; and 35 genotypes were highly susceptible (HS) with rating 7.1 to 9. This study reveals new source of resistance for use breeding programme aimed at developing YMV resistant varieties.

Keywords: Blackgram (urdbean); Yellow mosaic virus; MYMV Resistance breeding

Introduction

Blackgram (minumulu) is a very important grain legume crop grown in Andhra Pradesh in all the seasons but predominantly during rabi under rice fallows. The Guntur District ranks first in Andhra Pradesh for the production of blackgram. Blackgram is very nutritious as it contains high levels of proteins, potassium, calcium, iron, niacin (B3), thimine (B₁), Riboflavin (B2) [1]. Blackgram has been shown to be useful in mitigating elevated cholesterol levels. Blackgram has received prominence in Indian dities especially for culinary preparation of dal, Idli, Vada, Dosa, papad (Table 1).

Although blackgram has been traditionally cultivated after rice in krishna delta, it was considered only as subsistent crop with yields usually below 0.5 t/ha. Following comprehensive constraint analysis in 1980's cultivar improvement programme was initiated Satyanarayana 1994 and the resistant disease resistant varieties LBG-17 (powdery mildew disease), LBG -402 (wilt), LBG - 645 banda polish (wilt) etc... With yields exceeding 2.5 t / ha under with minimal management conditions. The above cultivars catalyzed commercialization of crop on large scale and economy of the farmers as well as sustainability of the production system. Blackgram production in rice fallows contributed to area and production increase in the state from 410 kg / ha on 219 00 ha in 1981-82 to 737 kg/ha on 560,000 ha in 1991-92.

In period of 2012-14, YMV was found in black gram. Due to this reason production of black gram was gradually declined. The grain achieved through the location specific technology developed have diminished for past decade due to YMV disease problem on Black gram in this area resulting in lower returns, shift to exploitative crops and un sustainability of system. Due to YMV, effect black gram crop area is diverted to maize, sorghum, due to non-availability of resistant varieties. Yellow mosaic disease is the most destructive disease on black gram both in Kharif and Rabi seasons. YMV is most destructive

Disease not only in India but also in Pakistan, Bangladesh, Srilanka and adjacent area of south East Asia [2,3].

YMV was first reported in 1960. During 1970's YMV has emerged as serious problem on Blackgram in India in northern plains [4-6]. The infected plants show alternating green and yellow patches on leaves. Leaf size is generally not affected but some time the green area are slightly raised and the leaves shown slightly puckering and reduction in size. The leaves become papery white and thin. The disease is transmitted through white fly by *Bemisia tabaci* [7].

Use of disease resistant crop varieties is regarded as an economical and durable method of controlling viral diseases. Identifying source of resistance to YMV on blackgram is a priority area towards research efforts screening blackgram germplasm against YMV for the Identification of resistance source under natural environmental conditions and a number of resistant genotypes have been reported by workers [8-10]. 49 genotypes of blackgram germplasm collected from diverse geographic region were collected and evaluated for resistance under natural environmental conditions in Rabi 2014-2015.

Materials and Methods

The study was carried out at disease screening field established at R&D farm, NRI Agritech; and Acharya Nagarjuna University, Nagarjunanagar, Guntur district of A.P., India during Rabi; 2015. Resistance screening was conducted in the natural epidemic conditions using alternative rows of highly susceptible varieties. About 49 varieties were evaluated at R&D Farm; NRI Agritech Guntur. The disease was scored on a 1-9 arbitrary scale [8].

Rating scale for yellow mosaic virus disease (1-9 scale)

- No visible symptoms on leaves or minute yellow specks on leaves.
- Small yellow with restricted spread covering 0.1 to 5% leaf area.

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Year	Area lakh/ha	Production lakh/t	Productivity (kg/ha)
1981-82	2.19	0.9	410
1991-92	5.60	3.57	739
2012-13	3.76	2.34	631
2013-14	3.20	2.00	558
1988-89	4.27	3.97	* 928

* indicates peak productivity year

Table 1: Area, production and productivity of Blackgram in A.P.

Rating	Reaction
1.0 to 2.0	Resistant (R)
2.1 to 4	Moderately Resistant (MR)
4.1 to 5	Moderately susceptible (MS)
5.1 to 7	Susceptible (S)
7.1 to 9	Highly Susceptible (HS)

Table 2: Categories are used in assessing the resistant reaction for yellow mosaic virus disease.

S.no	Entry/variety	Disease scoring scale (1-9scale)
1	PU -19	4
2	NARENDRA -1	8
3	TVM -1	8
4	BARChart -1	6
5	PU -13	3
6	PU -10	4
7	ADT -5	9
8	KU -301	8
9	MARCH -404	6
10	LBG -623	9
11	CO -5	8
12	T -91	4
13	T -9	4
14	TAU -1	7
15	NRI, LBG -20	3
16	TU -9814	8
17	TU -651	8
18	TU -3113	8
19	IU -98843	9
20	IU -652	9
21	TAU -1-1	8
22	IU -92-3	8
23	TPU -4	8
24	TPU -4-1	7
25	TU -9814	8
26	IU 86	8
27	IU 835	9
28	IU -67219	8
29	IU -834	9
30	IU -861	9
31	IU -943	9
32	IU -8810	8
33	PDU -1	8
34	TAU -2	8
35	PU -17	8
36	PU -30	3
37	PU -35	1
38	TU 942	8
39	JU -3	5
40	PU -31	1
41	LBG -752	9
42	LBG -685	9
43	LBG -402	9
44	LBG -645	9
45	LBG -17	9
46	LBG -22	9
47	LBG -787	8
48	JU-2	8
49	LBG-754	8

Table 3: Screening of blackgram genotypes against YMV during Rabi, 2015.

Rating	Reaction	Genotypes
1.0 to 2	Resistant (R)	PU-31, PU-35.
2.1 to 4	Moderately Resistant (MR)	PU-13, PU-10, T-9, T-91 NRILBG-20, PU-30
4.1 to 5	Moderately Susceptible (MR)	PU-19, Barch chart-1, JU-3
5.1 to 7	Susceptible (S)	March-404, TAU-1
7.1 to 9	Highly Susceptible (HS)	TMV-1, ADT-5, KU-301, LBG-623, CO-5, TU-9814, Narendra-1, TU-651, TU-3113, IU-98843, IU-652, TAU-1-1, IU-92-3, TPU-4, TPU-4-1, TU-9814, IU-86, IU-835, IU-67219, IU-834, IU-861, IU-943, IU-8810, PDU-1, TAU-2, PU-17, TU-942, LBG-752, LBG-685, LBG-645, LBG-402, LBG-17, LBG-725, LBG-787, JU-5

Table 4: Grouping of genotypes screened against YMV in blackgram during Rabi, 2015.

- Yellow mottling leaves covering 5.1 to 10% leaf area.
- Yellow mottling leaves covering 10.1 to 15% leaf area.
- Yellow mottling leaves covering 15.1 to 30% leaf area.
- Yellow discoloration of 30.1 to 50% leaf area.
- Pronounced yellow mottling and discoloration of leaves and pods reduction in leaf size and stunting of plant covering 50.1 to 75% of foliage.
- Severe yellow discoloration of entire leaves covering above 75.1 to 90% of foliage, stunting of plants and no pod formation.
- Severe yellow discoloration of entire leaves covering above 90.1% of foliage, stunting of plants and no pod formation.

Observations on the disease incidence were taken on randomly selected five plants of each entry and took mean of each entry to assign the category. The following categories are used in assessing the resistant reaction for yellow mosaic virus disease (Table 2).

Results and Discussion

A total of 49 genotypes screened against YMV to identify the sources of resistance revealed that 2 entries exhibited resistance (R) reaction with rating of 1.0 to 2.0. Six genotypes fell in the category of moderately resistant (MR) with rating scale of 2.1 to 4.0, three were moderately susceptible (MS) with rating of 4.1 to 5.0; two genotypes were susceptible (S) with rating of 5.1 to 7; and 35 genotypes were highly susceptible (HS) with rating 7.1 to 9. The data in Table 3 presents the actual disease resistance /susceptibility reaction of different blackgram genotypes/entries towards YMV disease. Different blackgram genotypes/entries that fall into each category were grouped in Table 4.

Conclusion

Evaluation of germplasm for disease resistance is crucial step in controlling plant diseases through host plant resistance. Genes conferring resistance can be to a certain extent identified through routine screening procedures such as germplasm evaluation. In the case of YMV in blackgram, of 49 genotypes screened, 2 entries have been identified exhibiting promising reaction to YMV resistance. Identification of resistant lines is essential in the ambit of integrated

disease management which is upcoming concept in the field of agriculture. Earlier studies identified controlling the viral disease. Similar type of genotype evaluations were previously documented by several workers [9-13]. Our genotypes LBG-17, LBG-402, LBG-645, LBG 685, are highly susceptible to YMV but agronomical and quality wise were superior. Hence improvement of these varieties for YMV resistance is undertaken by developing DNA markers for YMV resistance, resistance for screening molecular aided selection in blackgram improvement.

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