

Incidence of Leaf Spot Disease Caused by *Cercosporidium personatum* in Resistant, Susceptible and Hybridized Population of Groundnut Cultivars

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Abstract

The disease incidence of *Cercosporidium personatum* in field was analysed in the hybridized population derived from the resistant mutant lines of the cultivar ICGV-87304 and TAG-24 & TG-26. Reciprocal crosses were performed using the same parents. The comparison of disease incidence in hybridized population was made with resistant cultivar Girnar-1. Susceptible parents TAG-24 and TG-26 showed disease incidence of 14.08 and 16.40%, respectively while Girnar-1 and the resistant mutant parents showed percentage infection ranging from 0.06-0.96%. The plants raised from the hybridized population showed the percent infection ranging from 0.01 to 0.02 %.

Keywords: groundnut, *Cercosporidium personatum*, susceptible, pathogen

1. Introduction

Groundnut, *Arachis hypogaea* L. is a major oilseed crop of India, accounting for about 27% of global area and has a contribution of 19% to world groundnut production (Dwivedi & Crouch, 2003). It is used for its oil as well as for food. It is an important cash crop of our country with 92 per cent fat, 46 per cent proteins, 24.2 per cent soluble carbohydrates, 8.4 per cent fiber and 5.8 per cent ash (Sheshadri, 1962). Early leaf spots are caused by *Cercospora arachidicola* Hori and late leaf spots are caused by *Cercosporidium personatum* (Berk.&Curt.). These are among the most important diseases of groundnut (Subrahmanyam et al., 1982; Subrahmanyam & Smith, 1989). The spots are first seen on the upper surface of lower leaves as small necrotic spots. These eventually enlarge and become light to dark brown or black circular spots, ranging from 1 to 10 mm in diameter. Gradually these spots coalesce and cause defoliation. Early leaf spots are characteristically brown to reddish brown in color with a yellow halo. The early leaf spot spores are formed on the upper leaf surface with a raised surface, and the lower leaf surface is usually smooth. Late leaf spots are seen as dark brown to black spots, usually without a yellow halo. The spores are formed on the lower surface with a rough appearance. The upper leaf surface is smooth (Tshilenge et al., 2010). These foliar diseases cause huge losses up to 25-43% as the photosynthetic process is disrupted. The pods thus produced are lesser and inferior in quality (Waliyar et al., 2000).

It is thus, very important to make attempts to transfer the resistant trait to the susceptible cultivars which are otherwise popular. The present paper deals with assessment of the selected cultivars in field and transfer of the resistant character from Mutant 10, 16, 22, 25, 26 and 29 (derived from ICGV-87304) to TAG-24 and TG-26 which are widely grown.

2. Material and Methods

Seeds of two widely grown cultivars of *A. hypogaea*, TAG 24 and TG 26, developed by BARC were procured from Panjabrao Krishi Vidyapeeth, Akola, as a pure genetic stock. These two cultivars were susceptible to late leaf spot disease. The cultivar Girnar-1 was also procured and used as a resistant check. In addition, five mutant lines derived from ICGV-7304, developed at the Department of Botany, Nagpur University, Nagpur (Kale, 1998) were used for the hybridization experiments. All the five lines were resistant to late leaf spot disease (Table 1).

Table 1. Resistance status of different groundnut cultivars

Cultivars	Status of resistance	
	Early leaf spot (<i>Cercospora arachidicola</i>)	Late leaf spot (<i>Cercosporidium personatum</i>)
Girnar-1	Resistant	Resistant
Mutant 10	Resistant	Resistant
Mutant 16	Resistant	Resistant
Mutant 22	Resistant	Resistant
Mutant 25	Resistant	Resistant
Mutant 26	Resistant	Resistant
Mutant 29	Resistant	Resistant
TAG 24	Susceptible	Susceptible
TG 26	Susceptible	Susceptible

The seeds of all plants selected above were sown in the experimental field of Department of Botany, Nagpur University, Nagpur, in plots of 10 × 10 size in the kharif season. Healthy plants (selected on the basis of percentage infection (calculated by measuring the diameter of the leaf spots in the infected area versus the total leaf area) and yield as compared to susceptible cultivars) from each cultivar in the population were selected for performing the crosses. The plants showed flowering after 34 days of germination. The anthers were emasculated on the previous day using sterile forceps from the female parent and these flowers were bagged. This was done to prevent any self pollination. The time of anthesis was found to be 6.50 AM. Post anthesis, the pollen grains from the male parent were carefully removed and dusted on the stigma of the female parent. The crossed flowers were bagged and labelled. The bagging was removed after formation of pegs.

In this investigation, Mutants 10, 16, 22, 25, 26 and 29 were crossed with TAG-24 and TG-26 (Table 2). Reciprocal crosses were also performed using the same parents. For identification of the hybridized flowers, they were tied with a thin nichrome wire. The hybridization process was performed up to 60 days of germination as the flowers failed to form pegs after this period. The total number of pegs, pods and seeds formed from the flowers which were used in hybridization experiment were calculated as a measure of successful hybridization. The population was termed the F1 population and the seeds of the F2 and F3 population raised from the F1 were also screened for disease resistance.

The F1 seeds were collected from the pods which were obtained from the pegs tied with nichrome wire. The F1 plants were screened for incidence of disease and percentage of infected area per leaflet in the field (Table 3). The seeds from F1 plants showing minimum amount of infection as compared to the susceptible cultivars were selected and F2 population was raised. These F2 plants were screened for resistance and other desirable characters.

3. Results and Discussions

The seeds of the selected cultivars were grown in the experimental field and monitored carefully for germination. The hybridization experiments were carried out between the resistant and susceptible cultivars and the plants were screened for the appearance of pegs from the hybridized flowers and for pod and seed setting (Table 2). The F1 plants and the subsequent F2 and F3 generations were assessed for incidence to *Cercosporidium personatum* on the basis of percentage of infected area per leaflet in the field grown plants.

Table 2. Data on the number of pegs and pods formed in the crosses performed between cultivars, TAG-24 and TG-26, susceptible to late leaf spot and Mutant 16, 26, 29, 10, 22 and 25, resistant to late leaf spot disease

Crosses	Number of crosses performed	Number of pegs formed	Number of pods formed	Number of crossed seeds obtained
TAG 24 × Mutant 26	196	44	20	20
TAG 24 × Mutant 16	190	60	21	19
TAG 24 × Mutant 29	392	198	83	78
Mutant 16 × TAG 24	158	57	20	25
Mutant 26 × TAG 24	258	86	28	30
Mutant 29 × TAG 24	211	55	19	27
Mutant 10 × TG 26	103	29	25	34
Mutant 22 × TG 26	165	63	15	13
Mutant 25 × TG 26	54	19	12	18
TG 26 × Mutant 22	220	54	25	23
TG 26 × Mutant 25	35	07	05	06
TG 26 × Mutant 10	172	66	20	21

In F₂ population of the crosses involving TAG-24, thirteen lines were selected, while in the cross involving TG-26, only two selections were made, while the rest of the crosses did not yield resistant plants. The selected plants, however did not show absolute resistance but disease incidence was lesser than the susceptible parents in these selections. The cross between TAG-24 and mutants, M16, 22, 26 and 29 yielded % infection in the range of 0.01-0.02 while the cross between TG-26 and M22 showed % infection of 0.01 (Table 3). In the resistant check Girnar-1, the disease incidence in terms of % infection was 0.17%. Thus, in selections made in this investigation showed resistance better than the resistant check. These results were also correlated with the accumulation of different biochemical markers like the phytoalexin-reveratrol and hydrolases, which accumulated in higher content in the resistant cultivars (data not presented). Thus, these results in terms of appearance of leaf spot disease indicated that all the fifteen selections made in the hybridized population were resistant to *Cercosporidium personatum*.

Table 3. Data of percentage infection and yield of different lines derived from crosses between resistant cultivars and resistant mutants selected in F2

Cross	Lines resistant to late leaf spot disease	Plant type	Percentage of infected area/leaflet	Number of pods/plant	Seed with plant
TAG 24 × M26	CA/2/10	Bunch	0.01	05	2.32
	CA/2/12	Bunch	0.02	08	3.54
	CA/3/2	Bunch	0.02	11	4.20
	CA/3/10	Bunch	0.02	05	2.10
	CA/6/7	Bunch	0.01	05	3.86
TAG-24 × M16	CB/4/5	Bunch	0.01	04	3.17
	CB/6/6	Bunch	0.02	05	2.63
	CB/7/1	Bunch	0.02	05	2.48
	CB/8/2	Bunch	0.02	12	4.36
	CB/11/1	Bunch	0.02	12	4.27
	CB11//5	Bunch	0.02	08	3.42
	CB/12/1	Bunch	0.01	15	4.44
TAG 24 × M29	CC/14/3	Bunch	0.01	06	3.39
TG 26 × M22	CD/2/4	Bunch	0.01	08	3.34
	CD/17/2	Bunch	0.01	09	3.53
<i>Parents</i>					
TAG 24		Bunch	14.08	10	4.12
M26		Bunch	0.96	08	3.28
M16		Bunch	0.14	11	4.46
M29		Semi-erect	0.25	10	4.18
TG26		Semi-erect	16.40	17	5.43
M 22		Bunch	0.06	2	1.78
Girnar-1 (Used as resistant Check)		Bunch	0.17	13	5.62

Note. Rest of the crosses did not yield the resistant plants.

Bunch variety: The plant grows erect, possesses light-green foliage, produces pods in cluster at the base of the plant and has round, plump non-dormant seeds with light-rose testa. Semi-erect variety: Green to dark green leaves. It has 3-seeded (occasionally, 4-, 2-, or 1-seeded) pods, with slight to moderate ridges, slight reticulation, and with slight to moderate beaks and constrictions.

The transfer of resistant trait from related species to cultivated groundnut was successfully attempted by Smart (1978), and Gibbons (1980). They claimed that hybrids were resistant to early and late leaf spot. Similar investigations were made by many researchers in different crops (Alberschlem & Valent, 1978; Vidyashekharan, 1997; Kale, 1998; Koche et al., 2015). However, this is the successful attempt to transfer the resistant trait from mutant lines to popular cultivar TAG-24 and TG-26.

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