

CHARACTERIZATION OF ANASTOMOSIS GROUPS AND EVALUATION OF PATHOGENICITY OF *Rhizoctonia solani* ISOLATES INVOLVED IN FLAX SEEDLING BLIGHT.

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ABSTRACT

Forty eight isolates of *Rhizoctonia solani* were obtained from flax seedlings showed seedling blight. Anastomosis tests revealed that 12 isolates (25%) belonged to AG-2, while 36 isolates (75%) belonged to AG-4. Pathogenicity test on flax cultivar Shkha 1, under greenhouse conditions, showed that the pathogenic isolates in the pre-emergence stage represented 91.67 and 77.72% within AG-2 and AG-4, respectively. However, the pathogenic isolates of AG-4 representing 58.23% of the total isolates as well as the highest percentage of the pathogenic isolates (71.79%). This observations held true in the case of plant height and dry weight. These results indicate that both AGs 2 and 4 are important in the etiology of flax seedling blight. The importance of AG-2 is due to its high virulence, while the importance of AG-4 is due to its high prevalence. Cluster analysis suggested that, within each AG, isolates could be separated into subgroups with specific virulence patterns.

Keywords: *Rhizoctonia solani*, Anastomosis groups, Flax, Pathogenicity.

INTRODUCTION

Rhizoctonia solani Kuhn (teleomorph: *Thanatephorus cucumeris* (Fravk)Donk) is one of the more primitive basidiomycetes, *R. solani* exists in its vegetative form in nearly all agricultural soils. In this non-spore producing phase, the fungus lives saprophytically on dead plant remains but it can become vigorously parasitic when roots or other parts of a susceptible host penetrate the infested zone (Sneh *et al.*, 1991). *R. solani* is a major cause of flax seedling blight throughout most of the flax producing countries (Anderson, 1977, Krylova, 1981, Kumud *et al.*, 1997, and Andruszewska and Wusakowska, 1999). It attacks flax at early stages of development, destroying the roots and causing thinning or in severe infection, death of seedlings (Krylova, 1981). Under Egyptian conditions, *R. solani* is frequently isolated from blighted flax seedling (Personal communication); however, as far as we know, no attempts have been made to classify *R. solani* from flax into AGs. *Rhizoctonia solani* comprises a collection of noniterbreeding populations that are recognized through the anastomosis group concept (Ogoshi, 1987). Isolates of *R. solani* are currently grouped into 18 AG and subgroups on the basis of their anastomosis behaviour (Sneh *et al.*, 1991). Since virulence and host range of these groups differ, Knowledge of anastomosis group affiliation of an isolate involved in a particular disease has become very useful (Anderson, 1982; Sneh, *et al.*, 1991). Therefore, in the present study, we identified AGs of *R. solani* involved in flax seedling blight and evaluated pathogenicity of 48 isolates under greenhouse conditions.

MATERIALS AND METHODS

Isolation of *R. solani* from flax seedlings

Flax seedling showing seedling blight symptoms were collected from 48 fields in 6 governorates (Table 1) during the seasons of 2004 and 2005. Ten-fifteen seedlings were obtained from each field. Segments with *Rhizoctonia* lesions were washed in running tap water, disinfested in 1% sodium hypochlorite for 5 min, and rinsed in sterilized water. Small pieces, 0.5cm of diseased tissues were plated on water agar and kept for 48h at $25 \pm 1^\circ\text{C}$. Hyphal tips were transferred to acidified PDA slants. Pure isolates were stored at 5°C and served as stock cultures. Identification the anastomosis groups (AGs) of isolated *Rhizoctonia solani*.

Microscopic characterization

Individual isolates were examined microscopically to verify branching near the distal septum of cells as well as the constriction of the branch near the point of origin (Sherwood, 1969; Anderson, 1982; Ogoshi, 1985).

Staining nuclei

Aliquots of acidified (HCL) 0.5% trypan blue in lactophanol (Burpe *et al.*, 1978) was placed directly on young hyphae growing on agar media. Stained hyphae were covered with a slip and observed (400x) in a Petri dish, or a piece of agar infested with stained hyphae can be placed on microscope slide. Nuclei are stained dark blue to purple.

Hyphal fusion (Anastomosis)

Standard AG groups were kindly obtained from Cairo MERCIN, Faculty of Agriculture, Ain Shamis University. Isolates of *Rhizoctonia* are assigned to anastomosis group by pairing the isolates with 'tester' strains and observing the hyphae for fusion. Isolates paired on 2% water agar-coated slides or on agar in Petri dishes (Windels and Nabben, 1989). Mycelial transfers from the growing margins of young colony on PDA were plated 2-3cm apart in a slide in 9-cm Petri dish and incubated at 24 or room temperature until advancing hyphae made contact and slightly overlapped.

Pathogenicity test

Substrate for growth of each *Rhizoctonia solani* isolate was prepared in 500-ml glass bottle, each bottle contained 50 g of malt grains and 40 ml of tap water. Contents of bottle were autoclaved for 30 minutes. Inoculum, taken from one-week-old culture on PDA, was aseptically introduced into the bottle and allowed to colonize malt for 2 weeks. The present test was carried out by using autoclaved clay loam soil. Batches of soil were infested separately with inoculum of each isolate at a rate of 1g/kg of soil. Infested soil was dispensed in 10-cm-diameter clay pots and these were planted with 20 seeds per pot (cultivar Sakha1). In the control treatment, no fungal inoculum was added to the autoclaved soil. Pots were randomly distributed on a greenhouse bench under a temperature regime $24 \pm 6^\circ\text{C}$. Pre-emergence damping-off was recorded 15 day after planting, while, survival plant height (cm), and dry weight (mg/plant) were recorded 45 days after planting.

Table (1) : Isolates of *Rhizoctonia solani* from flax seedling used in this study.

| Isolats no. | Geographic origin | AG group | Previous crop |
|--------------------|--------------------------|-----------------|----------------------|
| 1 | Kafer-EL-sheikh | AG-4 | Rice |
| 2 | Kafer-EL-sheikh | AG-4 | Cotton |
| 3 | Kafer-EL-sheikh | AG-2 | Corn |
| 4 | Kafer-EL-sheikh | AG-4 | Rice |
| 5 | Kafer-EL-sheikh | AG-4 | Cotton |
| 6 | Kafer-EL-sheikh | AG-4 | Rice |
| 7 | Kafer-EL-sheikh | AG-4 | Rice |
| 8 | Kafer-EL-sheikh | AG-2 | Cotton |
| 9 | Kafer-EL-sheikh | AG-4 | Corn |
| 10 | Damiatta | AG-4 | Corn |
| 11 | Damiatta | AG-2 | Corn |
| 12 | Damiatta | AG-4 | Cotton |
| 13 | Damiatta | AG-4 | Corn |
| 14 | Damiatta | AG-4 | Rice |
| 15 | Damiatta | AG-4 | Cotton |
| 16 | Damiatta | AG-4 | Corn |
| 17 | Damiatta | AG-2 | Corn |
| 18 | Damiatta | AG-2 | Corn |
| 19 | Gharbiya | AG-4 | Rice |
| 20 | Gharbiya | AG-4 | Corn |
| 21 | Gharbiya | AG-4 | Rice |
| 22 | Gharbiya | AG-2 | Rice |
| 23 | Gharbiya | AG-4 | Coton |
| 24 | Gharbiya | AG-2 | Corn |
| 25 | Gharbiya | AG-4 | Coton |
| 26 | Gharbiya | AG-4 | Coton |
| 27 | Gharbiya | AG-4 | Coton |
| 28 | Beheira | AG-4 | Rice |
| 29 | Beheira | AG-2 | Corn |
| 30 | Beheira | AG-2 | Coton |
| 31 | Beheira | AG-4 | Corn |
| 32 | Beheira | AG-4 | Rice |
| 33 | Beheira | AG-4 | Coton |
| 34 | Beheira | AG-4 | Corn |
| 35 | Sharqiya | AG-2 | Rice |
| 36 | Sharqiya | AG-4 | Rice |
| 37 | Sharqiya | AG-4 | Corn |
| 38 | Sharqiya | AG-4 | Coton |
| 39 | Sharqiya | AG-4 | Coton |
| 40 | Sharqiya | AG-4 | Rice |
| 41 | Sharqiya | AG-4 | Rice |
| 42 | Sharqiya | AG-2 | Rice |
| 43 | Sharqiya | AG-4 | Corn |
| 44 | Daqahlyia | AG-4 | Corn |
| 45 | Daqahlyia | AG-4 | Coton |
| 46 | Daqahlyia | AG-2 | Corn |
| 47 | Daqahlyia | AG-4 | Corn |
| 48 | Daqahlyia | AG-4 | Rice |

Statistical analysis of data

Pathogenicity test was carried out in a randomized complete block design of three replicates. Percentage data were transformed into $\sqrt{x + 0.5}$ or arc sine angles to produce approximately constant variance before carrying out the analysis of variance (ANOVA). Duncan's multiple range test was used to compare between isolate means. ANOVA and correlation analyses were carried out by MSTAT statistical package. Cluster analysis of *R.solani* isolates was performed with the software package spss 6.0.

RESULTS AND DISCUSSION

Number of nuclei

Determination of the number of nuclei in vegetative hyphal cell is an important process in the identification of *Rhizoctonia solani* (Parmeter and Whitney 1970). Colonies of *Rhizoctonia solani* were stained with trypan blue and observed to determine numbers of nuclei in individual cells. All the isolates were multinucleate (Fig. 1), contain from 3 to 8 nuclei per cell.

Hyphal anastomosis

Microscopic examination showed that the isolates were divided into two anastomosis groups. 36 isolates (75%) belonged to AG-4, while 12 isolates (25%) belonged to AG-2 (Table 1). Perfect fusion between tester isolate and tested isolate is shown in Figs. 2 and 3.

Pathogenicity test

Pathogenicity of 48 isolates of *Rhizoctonia solani* was evaluated on flax cultivar Sakha1 under greenhouse condition (Table 2). Twelve isolates (25%) belonged to AG-2, while 36 isolates (75%) belonged to AG-4 (Table 3). The pathogenic isolates in the pre-emergence stage represented 91.67 and 77.78% within AG-2 and AG-4, respectively. However, the pathogenic isolates of AG-4 represented 58.23% of the total isolates as well as the highest percentage of the pathogenic isolates (71.79%). These observations held true in the case of plant height and dry weight. These results are in agreement with the results of Anderson (1977) who found that *Rhizoctonia solani* isolates belonging to AGs 2 and 4 caused seed decay of flax. These findings may indicate that the prevailing environmental conditions during flax growing season are more favorable for the prevalence of AG-4. The results also indicate that both AGs 2 and 4 are important in the etiology of flax seedling blight. The importance of AG-2 is due to its high virulence, while the importance of AG-4 is due to its high prevalence. There were no clear differences between AGs 2 and 4 regarding their isolation frequencies from the different regions in the Nile Delta (Table 4). Thus, the isolation frequencies of AGs 2 and 4 were 50 and 47%, respectively from East Delta region (Damiatta, Sharqiya, and Daqahlyia). As to West Delta region (Gharbiya and Behira), the isolation frequency was 33.33% for each group. AGs 2 and 4 were isolated from Kafr EL-Sheikh (North Delta region) at isolation frequencies of 16.67 and 16.44%, respectively. Within each governorate, the isolation frequency of AG4 was much greater than that of AG-2. Thus, while the isolation frequency of AG-4 ranged from 66.67 to 80% it ranged from 33 to 33.20% for AG-2.

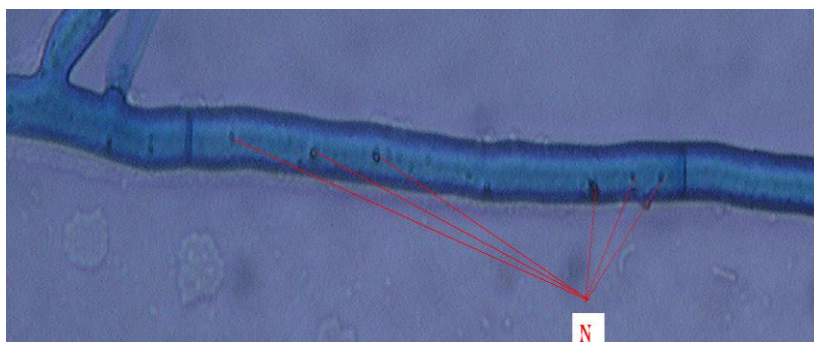


Fig.(1): Staining Nuclei (N) in vegetative hyphae of *Rhizoctonia Solani*

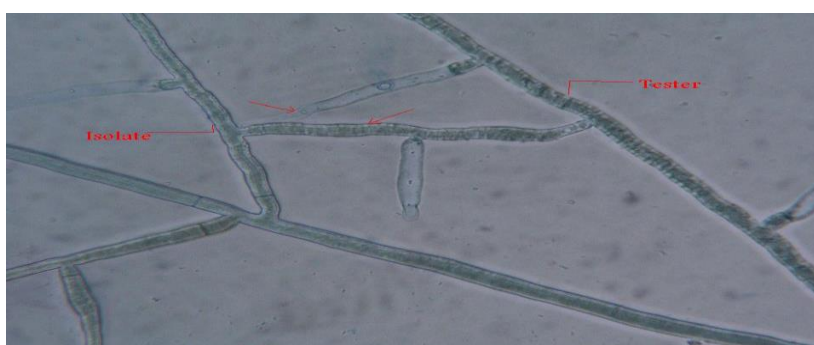


Fig.(2): Perfect fusion between hyphae of *Rhizoctonia solani*

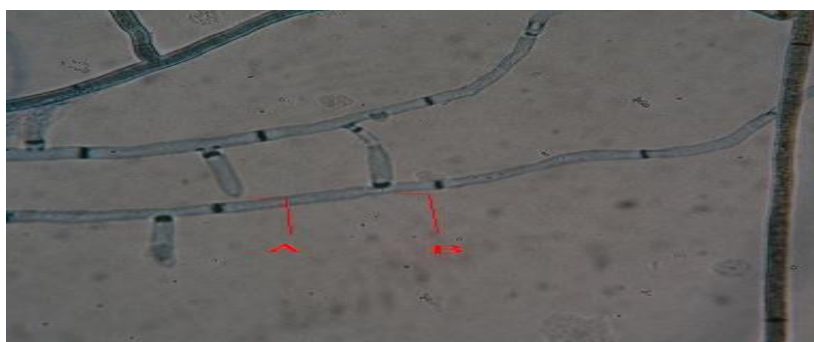


Fig.(3): Perfect fusion between hyphae of *Rhizoctonia solani*

- contact of hyphae.
- fusion of cells.
- Note lack of plasmolysis of fused cells.

Table (3): Classification of pathogenic *Rhizoctonia solani* from flax based on AG group .

| Seedling growth variable | AG Group | Total number of tested isolates | Total number of pathogenic isolates | Percentage of isolates within AG group | Percentage of total isolates ^a | Percentage of pathogenic isolates |
|--------------------------|----------|---------------------------------|-------------------------------------|--|---|-----------------------------------|
| Pre-emergence | AG2-2 | 12 | 11 | 91.67 | 22.92 | 28.21 |
| damping-off | AG-4 | 36 | 28 | 77.78 | 58.33 | 71.79 |
| Survival | AG2-2 | 12 | 12 | 100 | 25 | 25 |
| | AG-4 | 36 | 36 | 100 | 75 | 75 |
| Plant height | AG2-2 | 12 | 12 | 100 | 25 | 25 |
| | AG-4 | 36 | 36 | 100 | 75 | 75 |
| Dry weight | AG2-2 | 12 | 11 | 91.67 | 22.92 | 26.19 |
| | AG-4 | 36 | 31 | 86.11 | 64.58 | 73.81 |

^aTotal of 48 isolates of *Rhizoctonia solani* were tested for pathogenicity on seedlings of flax cultivar sakha1 under greenhouse conditions .

Table (4) : Distribution of *Rhizoctonia solani* AGs based on their geographic origin

| AG group | No of Isolates | Governorate | | | | | |
|----------|----------------|----------------|----------|----------|--------|----------|-----------|
| | | Kafr-EL-sheikh | Damiatta | Gharbiya | Behira | Sharqiya | Daqahlyia |
| AG2-2 | 12 | 2 | 3 | 2 | 2 | 2 | 1 |
| AG-4 | 36 | 7 | 6 | 7 | 5 | 7 | 4 |

AG-2 showed the highest isolation frequency after corn, while AG-4 showed its highest isolation frequency after cotton (Table 5). This result indicates that previous crop exerted a differential effect on the isolation frequency of the two groups. Within each previous crop, AG-4 was always the predominant group . Our results are in agreement with the other reports, which indicate that preceding crop plays an important role in enhancing or suppressing the different AGs of *R. solani*. For example, maize as a preceding crop in rotation may keep up the inoculum potential in the soil and therefore play an important role in the epidemiology of *Rhizoctonia* root rot of sugar beet (Ithurrant *et al.*, 2004). Schillinger and paulitz (2006) reported that *Rhizoctonia* bare patch in wheat was reduced with barely rotations.

Correlation among variables used for evaluating pathogenicity of *R. solani* isolates on flax seedlings are shown in Table 6 significant ($p < 0.01$) negative correlation was observed between pre-emergence damping-off and survival. Pre-emergence damping-off was correlated with plant height for isolates of AG-4 and for all isolates, while it was nonsignificant for isolates of AG-2. Correlation between pre-emergence damping-off and dry weight was nonsignificant. Post-emergence damping-off and plant height were negatively correlated for AG-4 isolates and for all isolates. Survival was negatively correlated with pre-emergence damping-off. The least number of significant

correlations was observed in the case of AG-2, this could be attributed to its small sample size (n=12).

Eight groups of similar isolates were identified by cluster analysis (Fig. 4). These groups were isolates 7,26,11,38,46,41; isolates of 6,12,44,33,24,1,8; isolates of 20,36,5,13,16,30,43; isolates 28,40,23; isolates 18,35,3,21,47,19; isolates of 9,25,2,14,37,42; isolates of 4,32,10,15,45,29, and isolates of 22,34,17,39,27,48,31. Neither AG nor geographic origin was associated with virulence pattern of the isolates because, most of the groups included isolates belonging to AG-4 and AG-2 and isolated from North, West, and East Delta regions. Similarly, previous crop was not associated with virulence pattern. Cluster analysis suggested that within each AG, isolates could be separated into subgroups with specific virulence patterns.

Table (5) : Distribution of AGs of *Rhizoctonia solani* based on previous crop.

| AGs | No.of Isolats | Previous crop | | |
|-------|---------------|---------------|------|------|
| | | Cotton | Corn | Rice |
| AG2-2 | 12 | 2 | 7 | 3 |
| AG-4 | 36 | 13 | 10 | 13 |

Table (6) : Correlation among variables used for evaluating pathogenicity of *R.solani* isolates on flax seedlings (cultivar Sakha 1) under greenhouse conditions.

| AG | Variable | Variable | | | | |
|-------------------------|------------------------------|----------|----------|----------|----------|--------|
| | | 1 | 2 | 3 | 4 | 5 |
| 2-2 ^a | 1-Pre-emergence damping-off | | 0.167 | -0.993** | -0.521 | 0.005 |
| | 2-Post emergence damping-off | 0.167 | | -0.280 | -0.011 | -0.064 |
| | 3-Survival | -0.993** | -0.280 | | 0.508 | 0.001 |
| | 4-Plant height | -0.521 | -0.011 | 0.508 | | 0.052 |
| | 5-Dry weight | 0.005 | -0.064 | 0.001 | 0.052 | |
| 4 ^b | 1-Pre-emergence damping-off | | 0.101 | -0.980** | -0.564** | -0.080 |
| | 2-Post emergence damping-off | 0.101 | | -0.288 | -0.476** | 0.231 |
| | 3-Survival | -0.980** | -0.288 | | 0.623** | 0.029 |
| | 4-Plant height | -0.564** | -0.476** | 0.623** | | 0.005 |
| | 5-Dry weight | -0.080 | 0.231 | 0.029 | 0.005 | |
| All ^c AGs | 1-Pre-emergence damping-off | | 0.101 | -0.983** | -0.544** | -0.057 |
| | 2-Post emergence damping-off | 0.101 | | -0.274 | -0.398** | 0.190 |
| | 3-Survival | -0.983** | -0.274 | | 0.588** | 0.018 |
| | 4-Plant height | -0.544** | -0.398** | 0.0588 | | 0.015 |
| | 5-Dry weight | -0.057 | 0.190 | 0.018 | 0.015 | |

**Correlation is significant at (P<0.01).

^aNo = 12

^bNo = 36

^cNo = 4

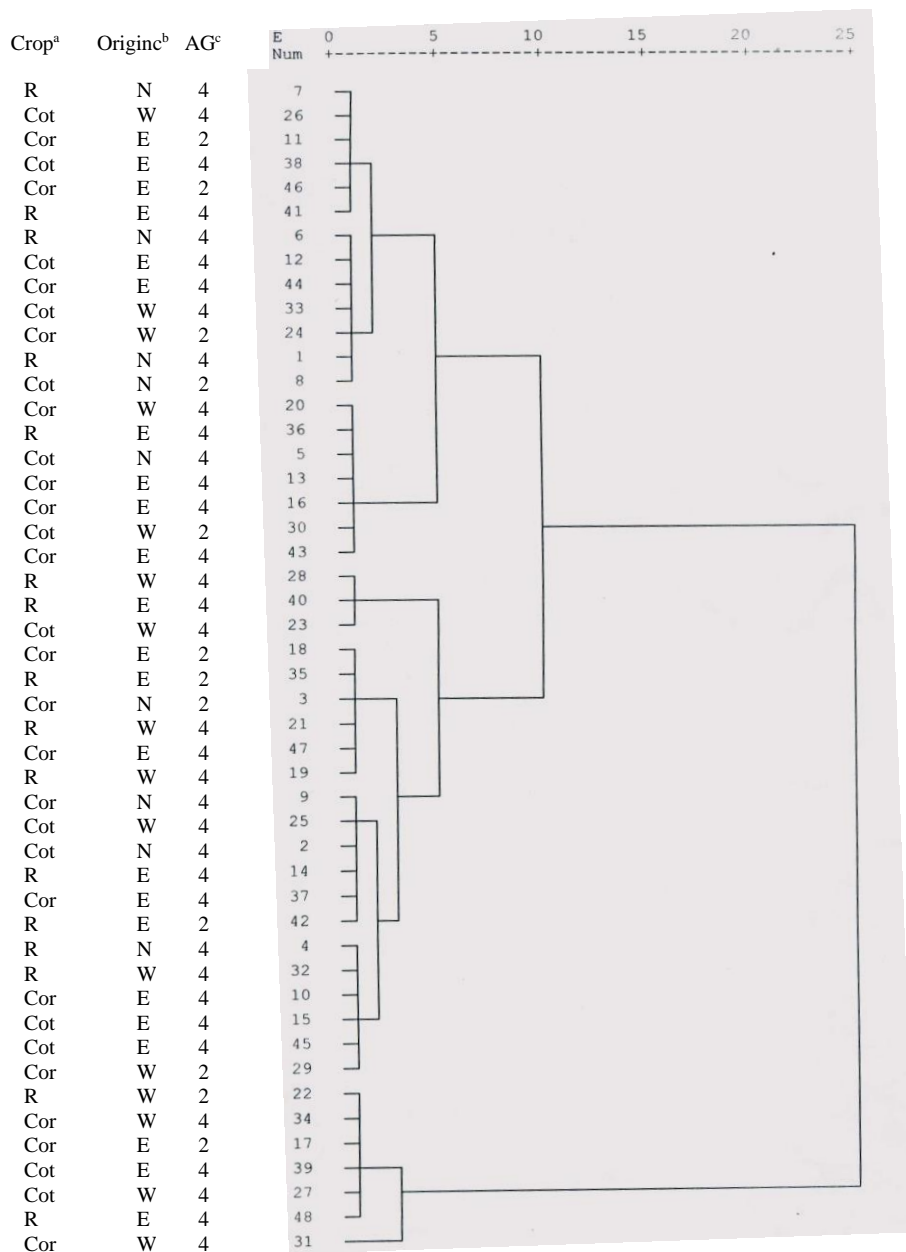


Fig.(4): Phenogram based on average linkage cluster analysis of virulence of 48 isolates of R.solani on flax cultivar Sakha 1

a Previous crop Rice (R), Cotton (Cot), and Corn (Cor).

b Geographic origins of isolates were East Delta (E), North Delta (E), North Delta (N), and West Delta (W).

c Anastomosis groups

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توصيف المجموعات الالتحامية وتقييم القدرة المرضية لعزلات فطر الريزكتونيا
سولاني المسببة لمرض لفحة بادرات الكتان .
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أمكن الحصول على ٤٨ عزلة لفطر الريزكتونيا سولاني من بادرات كتان مصابة
بمرض لفحة البادرات . أظهر الفحص الميكروسكوبي إن ١٢ عزلة (٢٥%) كانت تتبع المجموعة
الالتحامية AG2 في حين أن ٣٦ عزلة (٧٥%) كانت تتبع المجموعة الالتحامية AG4 . أظهر
اختبار القدرة المرضية لهذه العزلات على صنف سخا ١ تحت ظروف الصوبة أن ٦٧، ٩١ % من
عزلات المجموعة ٢ وكذلك ٧٢، ٧٢ % من عزلات المجموعة ٤ كانت ممرضة في مرحلة ما قبل
ظهور البادرات فوق سطح التربة . العزلات الممرضة التابعة للمجموعة ٤ مثلت ٢٣، ٥٨ % من
إجمالي العزلات المختبرة كما مثلت ٧٩، ٧١ % من إجمالي العزلات الممرضة . وكانت تلك النتائج
متوافقة مع تأثير العزلات على طول البادرات و وزنها الجاف . أكدت النتائج أن العزلات التابعة
لكل من المجموعتين ٢ و ٤ تلعب دورا هاما في إحداث لفحة بادرات الكتان حيث أتضح أن عزلات
المجموعة ٢ كانت ذات قدرة مرضية عالية في حين أن عزلات المجموعة ٤ كانت أكثر أنتشارا .
أظهر التحليل العنقودي إن كل من المجموعة الالتحامية ٢ و ٤ تنقسم إلى ٨ مجموعات فرعية من
العزلات التي تتميز كل منها بنمط محدد من حيث القدرة المرضية.

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