Research Article

Morphological and Pathogenic Variability among *Macrophomina phaseolina* Isolates Associated with Mungbean (*Vigna radiata* L.) Wilczek from Pakistan

Umer Iqbal¹ and Tariq Mukhtar²

¹ Crop Diseases Research Institute (DPEP), National Agricultural Research Centre, Park Road, Islamabad 45500, Pakistan
² Department of Plant Pathology, Pir Mehr Ali Shah Arid Agriculture University, Rawalpindi 46300, Pakistan

Correspondence should be addressed to Tariq Mukhtar; drtmukhtar@uaar.edu.pk

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Macrophomina phaseolina is a serious pathogen of many crops. In the present studies, 65 isolates of *Macrophomina phaseolina* from different agroecological regions of Punjab and Khyber Pakhtunkhwa provinces of Pakistan were analyzed for morphological and pathogenic variability. Regardless of their geographic origins, significant differences were detected among 65 isolates in their radial growth, sclerotial size, and weight as well as in pathogenicity. Sixteen isolates were rated as fast growing, 11 as slow growing, and the rest of the isolates as medium growing. Nine isolates were classified as large sized, 26 as small sized, and the remaining 30 isolates as medium sized. Thirty-five isolates were ranked as heavy weight, 12 as low weight, and the rest of isolates were grouped as medium weight. Ten fungal isolates appeared to be least virulent, whereas eight isolates of diverse origin proved to be highly virulent against mungbean cultivars. The remaining isolates were regarded as moderately virulent. No relationship was found among the morphological characters and pathogenicity of the isolates. These morphological and pathogenic variations in various isolates of *M. phaseolina* may be considered important in disease management systems and will be useful in breeding programmes of mungbean cultivars resistant to charcoal rot.

1. Introduction

Mungbean (*Vigna radiata* L.) Wilczek is a well known summer pulse crop of Pakistan and is cultivated on an area of 245.9 thousand hectares with a total production of 1777 thousand tones [1]. The crop is grown in a wide range of agroecological zones. The average yield of mungbean in Pakistan is very low as compared to its yield in many other countries. The low yield of mungbean in Pakistan can be attributed to legions of biotic and abiotic constraints. Among biotic factors, diseases are the most dominant. Depending upon the crop variety, the losses due to diseases to pulse crops have been estimated to be as high as 44 percent [2]. Mungbean is vulnerable to about 26 diseases in the world [3]. Among these, charcoal rot caused by *Macrophomina phaseolina* (Tassi) Goid is of prime importance in reducing crop yield especially in arid regions of the world [4]. The pathogen is distributed in diverse climatic conditions from arid to tropical regions and has a broad host range [5, 6]. There are more than 500 hosts of the fungus including legume and cereal plants [7, 8]. *M. phaseolina* is a soil- and seed-borne pathogenic fungus and produces cushion shaped black sclerotia [9]. Its prevalence can be enhanced by different physiological and ecological factors such as low moisture contents, high temperature, and heat [10, 11]. Disease severity is correlated with viable sclerotia present in the soil.

Charcoal rot infects plants at almost all growth stages. Dark lesions appear on the epicotyls and hypocotyls followed by seedling death due to obstruction of xylem vessels. In plants, the pathogen causes red to brown lesions on roots and stems with production of dark mycelia and black microsclerotia. Ultimately the plant becomes defoliated and wilted [6] and perishes [2].

Among the main management strategies, use of cultivars resistant to *M. phaseolina* has gained wide popularity and acceptance amongst farmers as application of fungicides is
A thorough knowledge of pathogenic variability of *M. phaseolina* is essential to design disease management strategies for different agroecological zones of the country by breeding resistant cultivars. At this time no information on the variability among *M. phaseolina* isolates is available in the country. Hence, we investigated morphological and genetic variability among 65 isolates of *Macrophomina phaseolina* to adapt and survive in diverse environments. Furthermore, resistant cultivars outstrip fungicides in various respects and emphasis is being laid on the development of new resistant germplasm. However, it has been observed that control measures against pathogens become complicated and even ineffective due to the variability among populations of the same pathogen in different agroecological zones.

There are reports in other parts of the world that populations of *M. phaseolina* showed significant variations morphologically [12], physiologically [13], pathogenically [14–17], and genetically [14–16, 18–24]. These variations aid the pathogen to adapt and survive in diverse environments.

A thorough knowledge of pathogenic variability of *M. phaseolina* is essential to design disease management strategies for different agroecological zones of the country by breeding resistant cultivars. At this time no information on the variability among *M. phaseolina* isolates is available in the country. Hence, we investigated morphological and pathogenic variability among 65 isolates of *M. phaseolina* infecting mungbean, collected from six different agroecological zones (Table 1) of Pakistan. It has also been determined whether morphological variations among *M. phaseolina* isolates have any relationship with the pathogenic variability.

### Table 1: Isolates of *Macrophomina phaseolina* collected from mungbean plants from different agroecological zones.

| Agroecological Zone       | Soil type                                      | District             | Isolate          |
|---------------------------|-----------------------------------------------|----------------------|------------------|
| Sandy desert              | Sandy and loamy fine sandy soil, moderately to strongly calcareous | Mianwali             | MP-12, MP-13, MP-14, MP-15, MP-16 |
|                           |                                               | Bhakkar              | MP-17, MP-18, MP-19, MP-20, MP-21 |
|                           |                                               | Layyah               | MP-22, MP-23, MP-24, MP-25, MP-26, MP-27, MP-28 |
| Northern irrigated plain  | Sandy loam to clay loam                       | Faisalabad           | MP-29, MP-30, MP-31, MP-32, MP-33 |
|                           |                                               | Muzaffargarh         | MP-8, MP-9, MP-10, MP-11 |
| Barani (rainfed)          | Non-calcareous to moderately calcareous silt loams | Narowal              | MP-34, MP-35, MP-36 |
|                           |                                               | Sialkot              | MP-37, MP-38, MP-39, MP-40, MP-41 |
|                           |                                               | Chakwal              | MP-42, MP-43, MP-44, MP-45 |
| Wet mountains             | Silt loam to silty clays, non-calcareous to slightly calcareous | Islamabad            | MP-46, MP-47, MP-48, MP-49 |
|                           |                                               | Rawalpindi           | MP-1, MP-2, MP-3 |
| Western dry mountains     | Loamy, deep, and strongly calcareous          | Kohat                | MP-56, MP-57, MP-58, MP-59 |
|                           |                                               | Banu                 | MP-60, MP-61, MP-62, MP-63, MP-64, MP-65 |
| Sulaiman piedmont         | Loamy to clayey and strongly calcareous       | Dera Ghazi Khan      | MP-4, MP-5, MP-6, MP-7 |
|                           |                                               | Dera Ismail Khan     | MP-50, MP-51, MP-52, MP-53, MP-54, MP-55 |

The diseased samples were first packed in paper bags and then in 15 × 20 cm polyethylene bags, labelled, brought to the lab, and stored at 4°C until processed for identification.

#### 2.2. Isolation, Purification, and Identification of *M. phaseolina*.

The fungus was isolated from stem bark tissues of mungbean bearing fungal sclerotia and showing characteristic charcoal rot symptoms. The samples were cut into small pieces (5–10 mm long) and surface sterilized with 1% sodium hypochlorite for 2 minutes and then rinsed thrice in sterilized distilled water. The pieces were placed on Chloroneb Mercury Rose Bengal Agar (CMRA) medium [25] in Petri dishes and incubated in dark at 25 ± 1°C for 7 days. A small portion of the fastest growing colony of *M. phaseolina* was taken from the periphery of a 90 mm diameter Petri dish, spread onto Petri dishes containing glucose agar medium (glucose, 20 g; agar, 20 g; and water, 1 L), and incubated in the dark at 25 ± 1°C for 7 days. A small portion of the colony having sclerotia was taken up into a drop of sterilized water and agitated with a sterilized needle to separate the sclerotia from the mycelia. Sclerotia were then transferred to 90 mm diameter Petri dishes containing CMRA medium. Colonies appearing from single sclerotia were again transferred to CMRA medium in 90 mm Petri plates, incubated as mentioned above and identified as described [26].

#### 2.3. Storage of Pure Cultures of *M. phaseolina*.

The purified culture (5 mm disc) from each isolate growing on PDA was transferred to 10 mL culture tubes and incubated in the dark at 25 ± 1°C for 6 days, until the surface of PDA was covered with a dense sclerotial layer of the fungal culture. The culture tubes were labeled and stored at 4°C.

#### 2.4. Multiplication of *M. phaseolina*.

Ground sorghum seeds were water-soaked overnight, air-dried under room temperature, and placed in conical flasks. The mouth of each flask was plugged with cotton wool, wrapped in aluminum foil,
Table 2: Morphological variations among different isolates of *M. phaseolina*

| Isolates | Radial growth (mm) | Sclerotial size (μm) | Sclerotial weight (mg) |
|----------|--------------------|----------------------|------------------------|
| 1        | 2                  | 3                    | 4                      |
| MP-1     | 63.17 i-r          | 26.67 a-d            | 0.10 h-l               |
| MP-2     | 65.33 i-p          | 23.50 b-l            | 0.12 e-l               |
| MP-3     | 75.17 a-i          | 27.33 ab             | 0.10 h-l               |
| MP-4     | 71.67 f-m          | 19.67 g-o            | 0.13 d-l               |
| MP-5     | 72.33 e-m          | 20.67 f-o            | 0.15 a-k               |
| MP-6     | 78.00 a-h          | 20.33 f-o            | 0.16 a-h               |
| MP-7     | 87.17 a            | 24.17 a-j            | 0.15 a-k               |
| MP-8     | 32.00 t            | 20.83 f-o            | 0.11 g-l               |
| MP-9     | 52.83 q-s          | 24.67 a-h            | 0.13 d-l               |
| MP-10    | 83.00 a-f          | 23.00 b-m            | 0.13 d-l               |
| MP-11    | 54.67 o-s          | 19.67 g-o            | 0.15 a-k               |
| MP-12    | 64.33 i-r          | 18.17 l-o            | 0.16 a-h               |
| MP-13    | 68.33 h-n          | 21.17 e-o            | 0.15 a-k               |
| MP-14    | 84.67 a-e          | 19.33 h-o            | 0.13 d-l               |
| MP-15    | 74.50 b-k          | 19.00 l-o            | 0.14 c-l               |
| MP-16    | 68.00 h-n          | 21.83 e-o            | 0.14 b-l               |
| MP-17    | 82.17 a-g          | 24.50 a-i            | 0.18 a-f               |
| MP-18    | 74.83 a-j          | 26.83 a-c            | 0.19 a-d               |
| MP-19    | 85.33 a-c          | 21.17 e-o            | 0.19 a-d               |
| MP-20    | 62.50 j-r          | 29.00 a              | 0.21 a                 |
| MP-21    | 78.00 a-h          | 20.33 f-o            | 0.17 a-g               |
| MP-22    | 71.83 f-m          | 20.17 f-o            | 0.19 a-d               |
| MP-23    | 65.17 i-q          | 17.50 m-o            | 0.20 ab                |
| MP-24    | 64.00 i-r          | 22.50 b-o            | 0.20 a-c               |
| MP-25    | 66.33 h-o          | 22.67 b-n            | 0.18 a-f               |
| MP-26    | 86.67 ab           | 23.33 b-l            | 0.17 a-g               |
| MP-27    | 82.83 a-f          | 21.67 c-o            | 0.18 a-f               |
| MP-28    | 43.83 s            | 17.33 no             | 0.17 a-g               |
| MP-29    | 43.67 s            | 18.00 l-o            | 0.13 d-l               |
| MP-30    | 44.00 s            | 22.67 b-n            | 0.13 d-l               |
| MP-31    | 53.83 p-s          | 26.33 a-e            | 0.15 a-k               |
| MP-32    | 83.33 a-f          | 19.83 g-o            | 0.12 f-i               |
| MP-33    | 84.17 a-f          | 17.67 m-o            | 0.12 f-l               |
| MP-34    | 53.17 p-s          | 21.83 c-o            | 0.10 h-l               |
| MP-35    | 72.50 d-m          | 24.67 a-h            | 0.13 d-l               |
| MP-36    | 61.50 m-r          | 22.17 b-o            | 0.10 h-l               |
| MP-37    | 52.33 rs           | 19.33 h-o            | 0.13 d-l               |
| MP-38    | 73.00 c-m          | 17.50 m-o            | 0.09 k-l               |
| MP-39    | 67.33 h-n          | 17.00 o              | 0.09 i-l               |
| MP-40    | 74.17 b-l          | 25.00 a-g            | 0.10 h-l               |
| MP-41    | 71.83 f-m          | 23.50 b-l            | 0.12 e-l               |
| MP-42    | 73.17 c-m          | 24.33 a-j            | 0.13 d-l               |
| MP-43    | 81.33 a-g          | 18.83 j-o            | 0.12 e-l               |

Table 2: Continued.

| Isolates | Radial growth (mm) | Sclerotial size (μm) | Sclerotial weight (mg) |
|----------|--------------------|----------------------|------------------------|
| 1        | 2                  | 3                    | 4                      |
| MP-44    | 85.00 a-d          | 19.00 i-o            | 0.12 e-l               |
| MP-45    | 86.17 ab           | 22.50 b-o            | 0.13 d-l               |
| MP-46    | 84.83 a-e          | 17.50 m-o            | 0.08 l                 |
| MP-47    | 65.17 i-q          | 21.33 d-o            | 0.09 k-l               |
| MP-48    | 81.17 a-g          | 18.50 k-o            | 0.10 h-l               |
| MP-49    | 64.17 i-r          | 22.33 b-o            | 0.09 j-l               |
| MP-50    | 81.83 a-g          | 20.33 f-o            | 0.18 a-f               |
| MP-51    | 56.50 n-r          | 21.67 c-o            | 0.17 a-g               |
| MP-52    | 78.00 a-h          | 25.67 a-f            | 0.20 ab                |
| MP-53    | 67.17 h-n          | 23.50 b-1            | 0.18 a-f               |
| MP-54    | 78.50 a-h          | 23.50 b-1            | 0.19 a-d               |
| MP-55    | 63.83 i-r          | 24.17 a-j            | 0.20 a-c               |
| MP-56    | 62.17 k-r          | 27.50 ab             | 0.16 a-i               |
| MP-57    | 71.83 f-m          | 22.83 b-n            | 0.18 a-e               |
| MP-58    | 74.17 b-l          | 24.00 a-k            | 0.15 a-k               |
| MP-59    | 78.17 a-h          | 26.67 a-d            | 0.16 a-i               |
| MP-60    | 61.17 m-r          | 19.33 h-o            | 0.18 a-e               |
| MP-61    | 69.67 g-m          | 19.67 g-o            | 0.18 a-f               |
| MP-62    | 84.50 a-e          | 21.50 c-o            | 0.17 a-g               |
| MP-63    | 61.67 l-r          | 24.17 a-j            | 0.15 a-k               |
| MP-64    | 78.67 a-h          | 19.83 g-o            | 0.15 a-j               |
| MP-65    | 54.50 o-s          | 24.17 a-j            | 0.16 a-i               |

Values are means of five replicates in case of radial growth and sclerotial weight and ten replicates in case of sclerotial size. Values sharing common letters in each column do not differ significantly at $P < 0.05$ according to Fisher's protected least significant difference test.

and autoclaved at 15 psi (121°C) for 20 minutes. After cooling, the seeds in flasks were inoculated with 4 mm mycelial plugs from a 7-day old culture of *M. phaseolina* and incubated at 25 ± 1°C for 15 days. The flasks were shaken at alternate days for uniform colonization of the grains. The inoculum thus produced was used in pot assay.

2.5 Determination of Morphological Variability. Morphological variability among 65 isolates of *M. phaseolina* was studied on the basis of the following parameters.

2.5.1 Radial Growth. For studying variability in radial growth, the isolates were grown on Potato Dextrose Agar [25]. Fifteen milliliters of autoclaved PDA was poured in 90 mm diameter Petri plates, allowed to solidify, and inoculated in the center with a 5 mm plug from the actively growing culture of each isolate of the fungus. The plates were incubated at 25 ± 1°C for 7 days. Each isolate was replicated five times. After the stipulated period, the growth of each isolate was measured in terms of colony diameter and their means were computed. On the basis of radial growth, the isolates were categorized as fast (>80 mm), medium (61–80 mm), and slow (<61 mm) growing.
2.5.2. Sclerotial Size. For measuring sclerotial size, slides from 7-day-old pure cultures of *M. phaseolina* isolates were prepared and examined under a microscope ocular micrometer. Sizes of ten randomly selected sclerotia were measured and their means were calculated. The isolates were classified as large (>25 μm), medium (21–25 μm), and small (<21 μm) sized.

2.5.3. Sclerotial Weight. In order to measure the dry weight of sclerotia, each isolate of the fungus was cultured in 100 mL sterilized Potato Dextrose Broth in 250 mL Erlenmeyer flasks of sclerotia, each isolate of the fungus was cultured in 100 mL sterilized Potato Dextrose Broth in 250 mL Erlenmeyer flasks with five replicates. The flasks were incubated at 25 ± 1°C for 15 days. The sclerotia were filtered through Whatman filter paper no. 41, wrapped in aluminium foil, and oven-dried at 45°C for 24 hrs. The sclerotia of each isolate were weighed using electric balance and grouped as high (>0.15 mg), medium (0.11–0.15 mg), and low (<0.11 mg) weight.

2.6. Determination of Pathogenic Variability. The pathogenicity of 65 isolates was studied on three cultivars of mungbean (NM-92, NM-51, and AEM-96) in the glasshouse in a split plot design with cultivars as main plots and the isolates as subplots. Each treatment was repeated thrice. Seeds were disinfected by immersing in 2.5% NaOCl for 5 min, rinsed in sterilized water, and air-dried.

Ten seeds of each of the three cultivars of mungbean were sown in pots containing 2 kg soil infested with each isolate of *M. phaseolina* @ 2 g/kg soil. Pots without inoculum served as controls. The pots were placed in a glasshouse at 30 ± 2°C. Disease severity caused by each isolate on each cultivar was assessed after 20 days of emergence using the disease rating scale developed by Abawi and Pastor-Corrales [6].

2.7. Statistical Analysis. Data were subjected to analysis of variance (ANOVA) using GenStat package 2009 (12th edition) version 12.1.0.3278 (http://www.vsni.co.uk). The differences among means were compared by Fisher's protected least significant difference test at *P* ≤ 0.05. Euclidean distances were used to construct a dendrogram by unweighted paired group method with arithmetic averages (UPGMA) using Statistica version 6.1.

3. Results

3.1. Morphological Variability among *M. phaseolina* Isolates. Significant variations were observed in the morphological parameters among 65 isolates of *M. phaseolina* collected from six agroecological zones of Pakistan.

3.1.1. Radial Growth. Significant differences among 65 isolates of *M. phaseolina* collected from different districts were observed on the basis of radial growth (*F* = 11.75; *df* = 64, 130; *P* < 0.001). The individual average radial growths of 65 isolates of *M. phaseolina* ranged from 32.00 to 87.17 mm observed 7 days after incubation. Maximum colony diameters of 87.17 and 86.67 mm were observed in case of isolate MP-7 (Dera Ghazi Khan) and MP-26 (Layyah) proving to be the fast growing, while isolates MP-8, MP-29, and MP-30 showed the minimum radial growths and were rated as slow growing. The individual radial growths of all the isolates are shown in Table 2 (Column 2). Sixteen isolates showed radial growths above 80 mm and were rated as fast growing while the growth of 11 isolates was found below 61 mm and were categorized as slow growing. The rest of the isolates showed growth between 61 and 80 mm and hence were classified as medium growing (Table 3).

3.1.2. Sclerotial Size. Significant variations were also observed among these isolates regarding the size of their sclerotia (*F* = 3.53; *df* = 64, 130; *P* < 0.001). Maximum sclerotial size was observed in case of isolates MP-20 and MP-3 showing 29.00 and 27.33 μm diameter, respectively, while the isolates MP-39 and MP-28 were found to be the smallest in size. The individual average sclerotial sizes of isolates ranged from 17.00 to 29.00 μm which are given in Table 2 (Column 3). The size of sclerotia of 9 isolates was above 25 μm and were classified as large size while 26 isolates with sclerotial size less than 21 μm were rated as small sized. The remaining 30 isolates ranged between 21 and 25 μm in sclerotial size and were categorized as medium sized (Table 4).

3.1.3. Sclerotial Weight. Sclerotial weight was another parameter considered for variability. The analysis of variance showed significant variability in sclerotial weight among the isolates (*F* = 6.07; *df* = 64, 130; *P* < 0.001). Analyzing the data of sclerotial weight revealed that isolates MP-20, MP-23, MP-24, and MP-52 produced maximum sclerota giving maximum weight above 0.20 mg. The lowest sclerotial production was found in isolate MP-46 with average weight of 0.08 mg. The mean individual sclerotial weight of each isolate is given in Table 2 (Column 4). Thirty five isolates were ranked as heavy weight as these produced sclerota more than 15 mg while 12 isolates produced sclerota weighing less than 11 mg and were graded as low weight. The rest of isolates being weighed between 11 to 15 mg were grouped as medium weight (Table 5).

3.1.4. Cluster Analysis Based on Radial Growth, Sclerotial Size and Sclerotial Weight. The cluster analysis of 65 isolates of 14 districts on an average basis of three morphological parameters (radial growth, sclerotial size, and sclerotial weight) is shown in Figure 1. In the dendrogram, three main clusters were distinguished, at a linkage distance of around 50%. The first cluster comprised 8 isolates of Faisalabad and Muzaffar Garh districts; the second cluster comprised 13 isolates of Chakwal, Bhakkar, and Dera Ghazi Khan districts, and the third cluster consisted of 44 isolates of the remaining districts. The isolates belonging to the districts Chakwal, Bhakkar, and D.G. Khan showed optimum growth performance, while the isolates belonging to M. Garh and Faisalabad exhibited poor growth performance. The isolates in the third group were found intermediate in their growth performance.

3.2. Pathogenic Variability among *M. phaseolina* Isolates. Highly significant differences were observed among isolates, varieties, and their interactions. Significant variations in
Table 3: Categorization of *M. phaseolina* isolates on the basis of radial growth.

| S. No. | Category                  | Number | Isolates                          |
|--------|---------------------------|--------|-----------------------------------|
| 1      | Fast growing (>80 mm)     | 16     | MP-7, MP-10, MP-14, MP-17, MP-19, MP-26, MP-27, MP-32, MP-33, MP-43, MP-44, MP-45, MP-46, MP-48, MP-50, MP-62 |
| 2      | Medium growing (61–80 mm) | 38     | MP-1, MP-2, MP-3, MP-4, MP-5, MP-6, MP-12, MP-13, MP-15, MP-16, MP-18, MP-20, MP-21, MP-22, MP-23, MP-24, MP-25, MP-35, MP-36, MP-38, MP-39, MP-40, MP-41, MP-42, MP-47, MP-49, MP-52, MP-53, MP-54, MP-55, MP-56, MP-57, MP-58, MP-59, MP-60, MP-61, MP-63, MP-64 |
| 3      | Slow growing (<61 mm)     | 11     | MP-8, MP-9, MP-11, MP-28, MP-29, MP-30, MP-31, MP-34, MP-37, MP-51, MP-65 |

Table 4: Categorization of *M. phaseolina* isolates on the basis of size of sclerotia.

| S. No. | Category                  | Number | Isolates                          |
|--------|---------------------------|--------|-----------------------------------|
| 1      | Large sized (>25 μm)      | 9      | MP-1, MP-3, MP-18, MP-20, MP-31, MP-40, MP-52, MP-56, MP-59 |
| 2      | Medium sized (21–25 μm)   | 30     | MP-2, MP-7, MP-9, MP-10, MP-13, MP-16, MP-17, MP-19, MP-24, MP-25, MP-26, MP-27, MP-30, MP-34, MP-35, MP-36, MP-41, MP-42, MP-45, MP-47, MP-49, MP-51, MP-53, MP-54, MP-55, MP-57, MP-58, MP-62, MP-63, MP-65 |
| 3      | Small sized (<21 μm)      | 26     | MP-4, MP-5, MP-6, MP-8, MP-11, MP-12, MP-14, MP-15, MP-21, MP-22, MP-23, MP-28, MP-29, MP-32, MP-33, MP-37, MP-38, MP-39, MP-43, MP-44, MP-46, MP-48, MP-50, MP-60, MP-61, MP-64 |

Table 5: Categorization of *M. phaseolina* isolates on the basis of weight of sclerotia.

| S. No. | Category                  | Number | Isolates                          |
|--------|---------------------------|--------|-----------------------------------|
| 1      | Heavy weight (>0.15 mg)   | 35     | MP-5, MP-6, MP-7, MP-11, MP-12, MP-13, MP-17, MP-18, MP-19, MP-20, MP-21, MP-22, MP-23, MP-24, MP-25, MP-26, MP-27, MP-28, MP-31, MP-50, MP-51, MP-52, MP-53, MP-54, MP-55, MP-56, MP-57, MP-58, MP-59, MP-60, MP-61, MP-62, MP-63, MP-64, MP-65 |
| 2      | Medium weight (0.11–0.15 mg) | 18 | MP-2, MP-4, MP-9, MP-10, MP-14, MP-15, MP-16, MP-29, MP-30, MP-32, MP-33, MP-35, MP-37, MP-41, MP-42, MP-43, MP-44, MP-45 |
| 3      | Low weight (<0.11 mg)     | 12     | MP-1, MP-3, MP-8, MP-34, MP-36, MP-38, MP-39, MP-40, MP-46, MP-47, MP-48, MP-49 |

Figure 1: Dendrogram derived from cluster analysis (UPGMA) showing relationship among the 65 isolates of *M. phaseolina* on the basis of morphological characters collected from 14 districts of Punjab and KPK provinces.
pathogenicity were found among 65 isolates of the fungus ($F = 34.31; \text{df} = 64, 388; P < 0.001$) when tested against three mungbean cultivars which also varied in response to the isolates ($F = 52.049; \text{df} = 2, 388; P < 0.001$). Six isolates, namely, MP-7, MP-13, MP-18, MP-48, MP-56, and MP-64, were found highly virulent against NM-92 with mean disease severity scoring of 7.3. Five isolates, namely, MP-8, MP-10, MP-26, MP-35, and MP-60, were found to be the least virulent in their reaction with average disease score ranging between 2.3 and 3.7 showing that the cultivar is resistant against these isolates (Table 6, Column 2).

Similarly nine isolates, namely, MP-5, MP-16, MP-17, MP-22, MP-33, MP-37, MP-46, and MP-63, were detected to be virulent against NM-51 with an average disease score ranging from 7.0 to 7.7, while six isolates, namely, MP-11, MP-30, MP-31, MP-38, MP-53, and MP-60, exhibited least pathogenic reaction against NM-51 with disease severity ranging from 2.0 to 3.0 and the remaining isolates proved to be intermediate in their pathogenicity (Table 6, Column 3). Seventeen isolates appeared to be highly virulent towards AEM-96 as these gave disease scores above 7, while six isolates, namely, MP-8, MP-25, MP-30, MP-31, MP-35, MP-36, and MP-41, with an average disease score ranged up to 3 proved to be least virulent. The rest of the isolates were found to be moderately virulent (Table 6, Column 4).

The cluster analysis on the basis of pathogenicity is shown in Figure 2. Sixty five isolates were categorized into five clusters on the basis of pathogenicity against three mungbean cultivars. Ten fungal isolates placed in clusters 2 and 3 appeared to be least virulent, whereas eight isolates of diverse origin in cluster 5 proved to be highly virulent for their virulence against mungbean cultivars. The remaining isolates falling under clusters 1 and 4 were regarded as moderately virulent.

4. Discussion

Macrophomina phaseolina, a soil- as well as seed-borne fungus, induces charcoal rot in different crops including mungbean. In the present studies, 65 isolates of M. phaseolina belonging to different regions of Punjab and Khyber Pakhtunkhwa provinces of Pakistan showed variations in different morphological traits such as radial growth, sclerotial size, and weight as well as in pathogenicity. The variations in morphology might be due to differences in temperature, moisture, soil types, and other edaphic factors of various districts of Punjab and KPK. Morphological variability has also been reported by many workers in terms of growth, color, pycnidium production, and chlorate sensitivity among different isolates of M. phaseolina on different hosts [11, 27–31] which corroborated our findings. Similarly, variations in morphology and pathogenicity among M. phaseolina isolates taken from different hosts as well as from different parts of the same host have also been observed by Beas-Fernández et al. [32]. However, in the present studies, no relationship was found among the morphological characters and pathogenicity of the isolates. Among the highly virulent isolates of M. phaseolina, namely, MP-7, MP-13, MP-18, MP-48, MP-56, and MP-64, against mungbean, not all the isolates were fast growing (radial growth $> 80$ mm) or large sized ($> 25$ $\mu$m) or have high weight ($> 0.15$ mg). Of these highly virulent isolates, two (MP-7 and MP-48) were the fast growing, and the remaining 4 were medium growing. Similarly, isolates MP-18 and MP-26 produced large sized, MP-7 and MP-13 medium sized, and MP-48 and MP-64 small sized sclerotia. Likewise, MP-7, MP-13, MP-18, and MP-64 were high and MP-48 was low weight. Similar pattern was observed in moderately and least virulent isolates. Confirmatory and contradictory findings in this regard have also been reported by others. A close linkage between virulence and growth was reported by
Table 6: Differential response of selected mungbean cultivars against various isolates of *M. phaseolina*.

| Isolates | NM-92 | NM-51 | AEM-96 |
|----------|-------|-------|--------|
| 1        | 2     | 3     | 4      |
| MP-1     | 5.7 defg | 5.7 defg | 5.7 defg |
| MP-2     | 7.0 abc | 4.7 ghij | 7.3 ab  |
| MP-3     | 5.7 defg | 5.3 efgh | 7.0 abc |
| MP-4     | 6.7 abcd | 7.0 abc | 7.3 ab  |
| MP-5     | 6.0 cdef | 7.7 a   | 7.3 ab  |
| MP-6     | 6.3 bcde | 5.7 defg | 6.0 cdef |
| MP-7     | 7.3 ab  | 6.7 abcd | 7.0 abc |
| MP-8     | 2.7 mn  | 5.7 defg | 3.0 lmn |
| MP-9     | 4.3 hijk | 5.3 efgh | 4.7 ghij |
| MP-10    | 2.3 mn  | 5.7 defg | 5.0 fgh |
| MP-11    | 5.7 defg | 2.7 mn  | 4.3 hijk |
| MP-12    | 6.0 cdef | 4.7 ghij | 5.7 defg |
| MP-13    | 7.3 ab  | 6.0 cdef | 4.3 hijk |
| MP-14    | 4.7 ghij | 7.0 abc  | 5.3 efgh |
| MP-15    | 4.3 hijk | 5.0 fgh  | 7.0 abc  |
| MP-16    | 6.3 bcde | 7.3 ab  | 5.7 defg |
| MP-17    | 7.0 abc | 7.3 ab  | 6.0 cdef |
| MP-18    | 7.3 ab  | 5.7 defg | 5.7 defg |
| MP-19    | 5.7 defg | 6.0 cdef | 4.3 hijk |
| MP-20    | 4.3 hijk | 6.3 bcde | 7.3 ab  |
| MP-21    | 4.7 ghij | 7.0 abc  | 5.3 efgh |
| MP-22    | 6.0 cdef | 7.3 ab  | 4.7 ghij |
| MP-23    | 5.3 efgh | 5.7 defg | 7.7 a   |
| MP-24    | 4.7 ghij | 5.0 fgh  | 5.7 defg |
| MP-25    | 5.7 defg | 5.3 efgh | 2.7 mn  |
| MP-26    | 2.7 mn  | 6.7 abcd | 7.3 ab  |
| MP-27    | 3.3 klm | 7.0 abc  | 7.7 a   |
| MP-28    | 5.3 efgh | 4.3 hijk | 5.7 defg |
| MP-29    | 6.0 cdef | 4.0 ijk  | 7.0 abc  |
| MP-30    | 5.3 efgh | 2.3 mn  | 3.0 lmn |
| MP-31    | 4.3 hijk | 2.0 n    | 3.3 klm |
| MP-32    | 2.3 mn  | 4.7 ghij | 4.7 ghij |
| MP-33    | 5.7 defg | 7.3 ab  | 6.7 abcd |
| MP-34    | 5.3 efgh | 4.3 hijk | 7.3 ab  |
| MP-35    | 3.3 klm | 4.7 ghij | 2.3 mn  |
| MP-36    | 5.3 efgh | 4.3 hijk | 2.7 mn  |
| MP-37    | 5.7 defg | 7.3 ab  | 7.0 abc  |
| MP-38    | 4.3 hijk | 3.0 lmn  | 5.3 efgh |
| MP-39    | 6.0 cdef | 5.7 defg | 6.7 abcd |
| MP-40    | 2.7 mn  | 4.0 ijk  | 5.7 defg |
| MP-41    | 7.0 abc | 5.3 efgh | 3.3 klm |
| MP-42    | 5.7 defg | 7.0 abc  | 7.3 ab  |
| MP-43    | 6.7 abcd | 5.7 defg | 7.0 abc  |
| MP-44    | 6.0 cdef | 7.7 a    | 7.7 a   |
| MP-45    | 4.7 ghij | 7.0 abc  | 5.7 defg |
| MP-46    | 5.7 defg | 4.3 hijk | 2.7 mn  |
| MP-47    | 5.3 efgh | 4.7 ghij | 5.7 defg |

Table 6: Continued.

| Isolates | NM-92 | NM-51 | AEM-96 |
|----------|-------|-------|--------|
| 1        | 2     | 3     | 4      |
| MP-48    | 7.3 ab  | 6.0 cdef | 7.0 abc |
| MP-49    | 5.3 efgh | 4.0 ijk | 5.7 defg |
| MP-50    | 5.7 defg | 5.7 defg | 7.3 ab  |
| MP-51    | 5.3 efgh | 7.0 abc | 7.7 a   |
| MP-52    | 4.7 ghij | 4.3 hijk | 5.3 efgh |
| MP-53    | 5.7 defg | 2.7 mn  | 7.3 ab  |
| MP-54    | 4.3 hijk | 4.0 ijk  | 5.7 defg |
| MP-55    | 7.0 abc  | 5.3 efgh | 7.3 ab  |
| MP-56    | 7.3 ab  | 7.3 ab  | 6.0 cdef |
| MP-57    | 5.7 defg | 5.3 efgh | 7.7 a   |
| MP-58    | 4.7 ghij | 5.7 defg | 7.0 abc  |
| MP-59    | 5.7 defg | 4.3 hijk | 5.3 efgh |
| MP-60    | 3.7 jkl | 2.3 mn  | 6.7 abcd |
| MP-61    | 4.3 hijk | 4.7 ghij | 7.3 ab  |
| MP-62    | 5.3 efgh | 5.7 defg | 5.7 defg |
| MP-63    | 6.7 abcd | 7.3 ab  | 4.7 ghij |
| MP-64    | 7.3 ab  | 5.3 efgh | 7.7 a   |
| MP-65    | 5.7 defg | 4.0 ijk  | 5.3 efgh |

Values are means of three replicates.

Values sharing common letters in each column do not differ significantly at \( P < 0.05 \) according to Fisher's protected least significant difference test.

Rayner [33]. Purkayastha et al. [34] also found relationship between morphological variations and pathogenicity. On the other hand, Dhingra and Sinclair [11] and Beas-Fernández et al. [32] reported that pathogenicity has no relation with size and weight of sclerotia.

The pathogenic fungus, *M. phaseolina*, has a broad host range and exits in two asexual forms which maintain its survival better [11, 13, 35]. Some workers also related variability to the phenomena of host specialization in *M. phaseolina*. Su et al. [15] found host specialization in maize on the basis of pathogenic, genetic, and physiological differences. Similarly, Cloud and Rupe [35] analyzed host specialization in soybean. This mechanism takes long time to establish within a specific host. Mihail and Taylor [13] suggested that, due to heterogenic nature of *M. phaseolina*, categorization into distinct subgroups based upon pathogenicity and morphology could not take place. Pathogenesis along with genetic diversity plays a specific role in host-plant resistance. Isolates having morphological similarity are not necessarily identical genetically, they might have some differences. The variable genetic pattern contributes to variation in morphology and pathogenesis, which has been confirmed by using different molecular tools [14, 16, 21, 22, 36–38]. As the pathogen has no sexual phase, genetic diversity is produced either by fusion of vegetative cells or by parasexual recombination between nuclear genes [39]. In nature genetic variability improves survival of a fungus [37].

It is quite evident that variability in morphology, physiology, genetics, pathogenicity, and so forth is imperative for the fungus to have better adaptation in response to
diversified environmental behavior. It also leads to host-plant resistance, development of resistant varieties of different crops against disease, and implementation of new disease controlling strategies [14, 40].

5. Conclusions

The determination of variability among *M. phaseolina* isolates is fundamental to guide the development of appropriate strategies for disease management according to different agroecological zones. As there are no reports about the determination of morphological and pathogenic variability, the present studies for the first time provide information on the variability of *M. phaseolina* in major mungbean growing areas of Pakistan. These results will be useful in developing integrated strategies for the management of charcoal root rot and breeding programs for pulses and other crops.

Conflict of Interests

The authors declare that there is no conflict of interests.

References

[1] Anonymous, Agricultural Statistics of Pakistan, Ministry of Food, *Agriculture and Live Stock*, Agriculture and Livestock Division, Islamabad, Pakistan, 2011.

[2] M. Bashir and B. A. Malik, “Diseases of major pulse crops in Pakistan—a review,” *Tropical Pest Management*, vol. 34, no. 3, pp. 309–314, 1988.

[3] Y. Y. Charles, “Mungbean diseases and control,” in *Proceedings of the 1st International Mungbean Symposium*, AVRDC, 1978.

[4] J. A. Hoes, “*Macrophomina phaseolina* causal agent of charcoal rot of sunflower and other crops,” Agricultural Canada Research Statistics, Modern Manitoba, 1985.

[5] C. Cottingham, “Numbers and distribution of sclerotia of *Macrophomina phaseolina* in soils of South Carolina,” *Plant Disease Reporter*, vol. 65, no. 4, pp. 355–356, 1981.

[6] G. S. Abawi and M. A. Pastor-Corrales, *Root Rots of Beans in Latin America and Africa: Diagnosis, Research Methodologies and Management Strategies*, CIAT, Cali, Colombia, 1990.

[7] O. D. Dhingra and D. Chagas, “Effect of soil temperature, moisture and nitrogen on competitive saprophytic ability of *Macrophomina phaseolina*,” *Transactions of British Mycological Society*, vol. 77, no. 1, pp. 15–20, 1981.

[8] J. B. Sinclair, *Compendium of Soybean Diseases*, American Phytopathological Society, Saint Paul, Minn, USA, 2nd edition, 1982.

[9] H. Wheeler, *Plant Pathogenesis*, Academic press, London, UK, 1975.

[10] G. C. Papavizas, “Some factors affecting survival of sclerotia of *Macrophomina phaseolina* in soil,” *Soil Biology and Biochemistry*, vol. 9, no. 5, pp. 337–341, 1977.

[11] O. D. Dhingra and J. B. Sinclair, *Biology and Pathology of Macrophomina phaseolina*, Universidade Federal de Vicsna, Minas Gerais, Brazil, 1978.

[12] N. Mayek-Pérez, C. López-Castañeda, and J. A. Acosta-Gallegos, “Variación en características culturales en vitro de aislamientos de *Macrophomina phaseolina* y su virulencia en frijol,” *Agrociencia*, vol. 31, no. 2, pp. 187–195, 1997.

[13] J. D. Mihail and S. J. Taylor, “Interpreting variability among isolates of *Macrophomina phaseolina* in pathogenicity, pyridinium production, and chlorate utilization,” *Canadian Journal of Botany*, vol. 73, no. 10, pp. 1596–1603, 1995.

[14] N. Mayek-Pérez, C. López-Castañeda, M. González-Chavira et al., “Variability of Mexican isolates of *Macrophomina phaseolina* on basis of pathogenesis and AFLP genotype,” *Physiological and Molecular Plant Pathology*, vol. 59, no. 5, pp. 257–264, 2001.

[15] G. Su, S.-O. Suh, R. W. Schneider, and J. S. Russin, “Host specialization in the charcoal rot fungus, *Macrophomina phaseolina*,” *Phytopathology*, vol. 91, no. 2, pp. 120–126, 2001.

[16] M. C. Reyes-Franco, S. Hernández-Delgado, R. Beas-Fernández, M. Medina-Fernández, J. Simpson, and N. Mayek-Pérez, “Pathogenic and genetic variability within *Macrophomina phaseolina* from Mexico and other countries,” *Journal of Phytopathology*, vol. 154, no. 7–8, pp. 447–453, 2006.

[17] S. S. Aboshosha, S. I. Atta Alla, A. E. El-Korany, and E. El-Aargawy, “Characterization of *Macrophomina phaseolina* isolates affecting sunflower growth in El-Behera governorate, Egypt,” *International Journal of Agriculture and Biology*, vol. 9, no. 6, pp. 807–815, 2007.

[18] T. E. Chase, Y. Jiang, and J. D. Mihail, “Molecular variability in *Macrophomina phaseolina*,” *Phytopathology*, vol. 84, no. 10, pp. 1149, 1994.

[19] G. Vandemark, O. Martinez, V. Pecina, and M. J. Alvarado, “Assessment of genetic relationships among isolates of *Macrophomina phaseolina* using a simplified AFLP technique and two different methods of analysis,” *Mycologia*, vol. 92, no. 4, pp. 656–664, 2000.

[20] V. Pecina-Quintero, O. Martinez-de la Vega, M. J. Alvarado-Balleza, G. J. Vandemark, and H. Williams-Alanis, “Comparación de dos sistemas de marcadores moleculares en el análisis de las relaciones genéticas de *Macrophomina phaseolina*,” *Revista Mexicana de Fitopatología*, vol. 19, pp. 128–139, 2001.

[21] A. M. R. Almeida, R. V. Abdelnoor, C. A. A. Arias et al., “Genotypic diversity among Brazilian isolates of *Macrophomina phaseolina* revealed by RAPD,” *Fitopatologia Brasileira*, vol. 28, no. 3, pp. 279–285, 2003.

[22] T. Jana, T. R. Sharma, R. D. Prasad, and D. K. Arora, “Molecular characterization of *Macrophomina phaseolina* and Fusarium species by a single primer RAPD technique,” *Microbiological Research*, vol. 158, no. 3, pp. 249–257, 2003.

[23] T. Jana, T. R. Sharma, and N. K. Singh, “SSR-based detection of genetic variability in the charcoal root rot pathogen *Macrophomina phaseolina*,” *Mycological Research*, vol. 109, no. 1, pp. 81–86, 2005.

[24] T. K. Jana, N. K. Singh, K. R. Koundal, and T. R. Sharma, “Genetic differentiation of charcoal root pathogen, *Macrophomina phaseolina*, into specific groups using URP-PCR,” *Canadian Journal of Microbiology*, vol. 51, no. 2, pp. 159–164, 2005.

[25] W. A. Meyer, J. B. Sinclair, and M. N. Khare, “Biology of *Macrophomina phaseolina* in soil studies with selective media,” *Phytopathology*, vol. 63, no. 5, pp. 613–620, 1973.

[26] H. L. Barnett and B. B. Hunter, *Illustrated Genera of Imperfect Fungi*, Burgess Publishing Company, Minneapolis, Minn, USA, 1972.

[27] O. D. Dhingra and J. B. Sinclair, “Location of *Macrophomina phaseolina* on soybean plants related to cultural characteristics and virulence,” *Phytopathology*, vol. 63, no. 7, pp. 934–936, 1973.

[28] C. A. S. Pearson, J. F. Leslie, and F. W. Schwenk, “Variable chlorate resistance in *Macrophomina phaseolina* from corn,
soybean, and soil," *Phytopathology*, vol. 76, no. 6, pp. 646–649, 1986.

[29] T. Adam, *Contribution à la connaissance des maladies du niébé (Vigna unguiculata (L.) Walp.) au Niger avec mention spéciale au Macrophomina phaseolina (Tassi) Goid [Thèse de doctorat]*, Université de Renne I, 1986.

[30] M. Atiq, A. Shabeer, and I. Ahmed, “Pathogenic and cultural variation in *Macrophomina phaseolina*, the cause of charcoal rot in sunflower,” *Sarhad Journal of Agriculture*, vol. 2, pp. 253–255, 2001.

[31] A. Riaz, S. H. Khan, S. M. Iqbal, and M. Shoaib, “Pathogenic variability among *Macrophomina phaseolina* (Tassi) Goid, isolates and identification of sources of resistance in mash against charcoal rot,” *Pakistan Journal of Phytopathology*, vol. 19, no. 1, pp. 44–46, 2007.

[32] R. Beas-Fernández, A. de Santiago-de Santiago, S. Hernández-Delgado, and N. Mayek-Pérez, “Characterization of Mexican and non-Mexican isolates of *Macrophomina phaseolina* based on morphological characteristics, pathogenicity on bean seeds and endoglucanase genes,” *Journal of Plant Pathology*, vol. 88, no. 1, pp. 53–60, 2006.

[33] A. D. M. Rayner, “The challenge of the individualistic mycelium,” *Mycologia*, vol. 83, no. 1, pp. 48–71, 1991.

[34] S. Purkayastha, B. Kaur, N. Dilbaghi, and A. Chaudhury, “Cultural and pathogenic variation in the charcoal rot pathogen from cluster bean,” *Annals of Agri Bio Research*, vol. 9, no. 2, pp. 217–221, 2004.

[35] G. L. Cloud and J. C. Rupe, “Preferential host selection of isolates of *Macrophomina phaseolina*,” *Phytopathology*, vol. 78, no. 12, p. 1563, 1988.

[36] M. Fuhlbohm, “Genotypic diversity among Australian isolates of *Macrophomina phaseolina*,” in *Proceedings of the Biennial Australian Plant Pathology Society Conference*, Lincoln University, Lincoln, New Zealand, 1997.

[37] Rajkumar, F. Bashasab, and M. S. Kuruvinashetti, “Genetic variability of sorghum charcoal rot pathogen (*Macrophomina phaseolina*) assessed by random DNA markers,” *Plant Pathology*, vol. 23, pp. 45–50, 2007.

[38] N. Allaghebandzadeh, S. Rezaee, B. Mahmoudi, and H. Z. Zadeh, “Pathogenic and genotypic analysis among Iranian isolates of *Macrophomina phaseolina*,” *Plant Pathology*, vol. 98, p. 11, 2008.

[39] M. J. Carlile, “Genetic exchange and gene flow: their promotion and prevention,” in *Evolutionary Biology of the Fungi*, A. D. M. Rayner, C. M. Brasier, and D. Moore, Eds., pp. 203–214, Cambridge University Press, Cambridge, UK, 1986.

[40] S. Purkayastha, B. Kaur, N. Dilbaghi, and A. Chaudhury, “Characterization of *Macrophomina phaseolina*, the charcoal rot pathogen of cluster bean, using conventional techniques and PCR-based molecular markers,” *Plant Pathology*, vol. 55, no. 1, pp. 106–116, 2006.