

Ph D Thesis Award

Studies on cultural, morphological, pathogenic and molecular variability of *Alternaria brassicae*, the causal agent of blight disease of rapeseed-mustard

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Abstract

In India, rapeseed-mustard is one of the most edible oilseed crops after groundnut and contributes around 26.1 per cent of the total oilseed production. Among Brassica oil seed crops, Indian mustard [Brassica juncea (L.) Czern & Coss.] is commonly cultivated in most of the Indian states. Alternaria blight disease caused by A. brassicae has been reported from all the continents of the world causing up to 47 per cent yield losses with no proven source of resistance. The present investigation was carried out to study the cultural, morphological, pathogenic and molecular variability among A. brassiace isolates collected from different Brassica spp. (Pantnagar) and from different geographical locations of India (Karnal, Punjab, Bihar, Jammu and Kangra) to exploit them in breeding programme during screening for resistance sources and selection of host differentials. In the present studies A. brassicae isolates (20 nos.) showed the existence of genetic diversity. Significant variation in cultural, morphological, pathogenic and molecular variability was observed in A. brassiace isolates irrespective to geographical locations and Brassica spp. Maximum radial growth (82.0 mm) was in AB-B. juncea Pantnagar isolate, while minimum in AB- B. caulorapa (49.7 mm) on PDA. Variations were also observed in colony colour, appearance, margin and zonation number. Substantial variations were found in spore morphology in respect to conidial length, width and number of septa. Average conidial length and width were varied from 29.0-6.6 x 185.3-28.2 µm. Maximum spore length and width was in AB-B. carinata isolate (185.3 x 25.6 µm), while minimum in AB-B. caulorapa (29.0 x 6.6 µm). Number of horizontal and vertical septa ranged between 3.50-14.75 and 0.75-5.0 respectively. All the isolates were pathogenic to Brassica spp. AB-B. juncea Pantnagar isolate was most virulent, while AB-S. alba least virulent. RAPD (18 no.) and ISSR (4 no.) primers generated a total of 310 reproducible and scorable polymorphic bands ranged from 100 to 1350 bp which displayed genetic polymorphisms among the A. brassicae isolates. On the basis of cultural, morphological, pathogenic and molecular variability, the A. brassiace isolates has been categorized into 5 major groups. Group I: AB-R. sativus, AB-B. pekinensis, AB-B. oleracea, AB-B. botrytis and AB-B. caulorapa; Group II: AB-B. juncea Pantnagar, AB-Yellow sarson, AB-Toria, AB-B. nigra, AB-R. oleifera, AB-B. juncea Karnal, AB-B. juncea Punjab and AB-B. juncea Bihar; Group III: AB-B. napus, AB-B. carinata, AB-E. sativa, AB-S. alba; Group IV: AB-B. juncea Kangara and AB-B. juncea Jammu; Group V: AB-B. rugosa. The different Brassica spp. (08 nos.) used in the present investigation to differentiate different A. brassicae isolates (09 nos.) as host differentials has been categorized into 4 major groups on the basis of differential phenotypic disease reactions. Group I: Varuna, PT-303, PYS-6 and B. nigra; Group II: PBN-9501 and Kiran; Group III: E. sativa; Group IV: S. alba. Among different methods of culture preservation of A. brassicae isolates, 10 per cent glycerol at -20°C was best in terms of sporulation (66.7%) followed by PDA at 4°C (65.7%) and could be used for the preservation of A. brassicae isolates at least two years.

Key words: Alternaria brassicae, cultural, morphological, pathogenic and molecular variability

Introduction

In India, rapeseed-mustard is an important group of edible oilseed crops that contributes around 26.1 per cent of the total oilseed production. Indian mustard (*B. juncea*) is one of the major oilseed crops cultivated in India. Among various diseases Alternaria blight caused by *A. brassicae* has been reported from all the continents of the world. It is one of the most important diseases of mustard causing up to 47 per cent yield loss (Meena *et al.*, 2010) with no proven source of resistance till date. Studies on pathogen variability have to be the foundation for development of pre-breeding populations as strategic defense mechanism.

Severity of Alternaria blight on oilseed Brassica differs among regions and also Brassica crops within a region might be due to existence of variability among the isolates of A. brassicae. The existence of pathogen diversity among A. brassicae isolates has already been reported by earlier workers. Different isolates of A. brassicae showed variable response on different Brassica spp./genotypes. Significant tolerance was observed in B.alba, EC-399299, PAB 9511 (B. juncea), Eruca sativa, B. carinata and B. napus. Variation in tolerance and susceptibility on same host depending on aggressiveness of isolates revealed the existence of variability among A. brassicae isolates. PCRbased marker techniques viz. RAPD and ISSR have been used extensively for genetic variation in populations of A. brassicae isolates.

Since mid 90's, Alternaria blight disease of crucifers is being managed by the use of fungicides. Due to deleterious effects of fungicides, the focus now lays on the development of resistant varieties which is one of the most eco-friendly, economic, and effective method for the management of the disease. Sources of resistance has been reported but their utility and effectiveness in breeding for disease resistance cultivars is limited due to lack of information on the occurrence and distribution of pathotypes and suitable genotypes screening technique using different races of *A. brassicae*.

In view of the above facts the present investigation on study of cultural, morphological, pathogenic and molecular variability was undertaken to find out variability among *A.brassicae* isolates collected from different *Brassica* spp. and from different growing locations of India so that the isolates could be classified in different groups and further to be exploited for the testing of resistant sources in *Brassica* spp. Grouping of different *Brassica* spp. as host differentials may also be helpful to differentiate different *A. brassicae* isolates.

Materials and Methods Collection, isolation and purification

A. brassicae isolates (20 no.) collected from different Brassica spp. (Pantnagar) during the crop season of 2013-14 and from different growing locations of India (Karnal, Punjab, Bihar, Jammu and Kangra) were cultured on PDA and further purification was done using single spore isolation. The culture was preserved in the refrigerator (4°C) for further studies (Table 1).

Cultural variability

Eight different media (Potato dextrose, Radish root dextrose, Cauliflower dextrose, Carrot potato dextrose, Mustard seed dextrose, Maize seed dextrose, Brassica leaf extract and Modified V-8 juice agar) were tested to determine their effect on growth of *A. brassicae* isolates. The observations on radial growth (mm), colony color, shape, margin, zonation number and appearance of each isolate on different media were recorded 10 DAI at $24\pm1^{\circ}$ C.

Morphological variability

Ten conidia of each isolates were examined at 10X and 40~X under advance light microscope. The conidial length, width and beak (μ m) were measured and number of horizontal and vertical septa and number of cells were recorded.

Pathogenic variability

The experiment was conducted in plastic pots under glasshouse during two consecutive seasons 2014-15 and 2015-2016. Forty days old plants (*B. juncea* var. Varuna) inoculated with conidial suspension (4x10⁴conidia/ml) of each *A. brassicae* isolates separately by single drop inoculation method (10µl/leaf) were kept in moist chamber for 48 hrs under glasshouse. The observations on spot and periphery

Table 1: Altenaria brassicae isolates collected from different Brassica spp. and from different geographical locations

A. brassicae isolate	Host	Location	Latitude & Longitude	Plant part
AB- B. juncea Pantnagar	B. juncea cv. Varuna	Pantnagar, CRC	29°N, 79.8°E	leaf
AB-B. napus	B. napus cv. PBN-9501	Pantnagar, CRC	29°N, 79.8°E	leaf
AB- B. carinata	B. carinata cv. Kiran	Pantnagar, CRC	29°N, 79.8°E	leaf
AB-Toria	B. rapa var. toria cv. PT-303	Pantnagar, CRC	29°N, 79.8°E	leaf
AB- Yellow sarson	B. rapa var. yellow sarson cv. PYS-6	Pantnagar, CRC	29°N, 79.8°E	leaf
AB- E. sativa	Eruca sativa (local)	Pantnagar, CRC	29°N, 79.8°E	leaf
AB- S. alba	Sinapis alba (local)	Pantnagar, CRC	29°N, 79.8°E	leaf
AB- R. oleifera	Raphanus sativus var. oleifera (local)	Pantnagar, CRC	29°N, 79.8°E	leaf
AB-B. nigra	B. nigra (local)	Pantnagar, CRC	29°N, 79.8°E	leaf
AB-B. rugosa	Brassica rugosa (local)	Pantnagar, CRC	29°N, 79.8°E	leaf
AB-R. sativus	Raphanus sativus	Pantnagar, VRC#	29°N, 79.8°E	leaf
AB-B. pekinensis	Brassica rapa var. pekinensis	Pantnagar, VRC	29°N, 79.8°E	leaf
AB- B. oleracea	Brassica oleracea var. capitata	Pantnagar, VRC	29°N, 79.8°E	leaf
AB- B. botrytis	Brassica oleracea var. botrytis	Pantnagar, VRC	29°N, 79.8°E	leaf
AB- B. caulorapa	Brassica caulorapa	Pantnagar, VRC	29°N, 79.8°E	leaf
AB-B. juncea Karnal	Brassica juncea	Karnal	$28^{0}N,77^{0}E$	leaf
AB- B. juncea Punjab	Brassica juncea	Punjab	31.1°N, 75.3°E	leaf
AB- B. juncea Bihar	Brassica juncea	Bihar	25°N, 85.3°E	leaf
AB-B. juncea Jammu	Brassica juncea	Jammu	32.4°N, 74.5°E	leaf
AB- B. juncea Kangara	Brassica juncea	Kangara	32°N, 76.18°E	leaf

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color, presence or absence of concentric rings, central point, yellow halo region and disease index (0-9 scale) were recorded at 50 DAI.

Molecular variability

DNA of twenty A. brassicae isolates was isolated by adapting Cetyl Trimethyl Ammonium Bromide (CTAB) extraction method (Doyle and Doyle, 1990). Molecular variability among A. brassicae isolates were studied by using RAPD (20 no.) and ISSR primers (05 no.) manufactured by Eurofins mwg operon (Eurofins genomics Pvt. ltd, Bangalore, India).

In RAPD, the polymerase chain reaction (PCR) was performed in 20 µl volumes containing 1 µl DNA (50 $ng/\mu l$), 1 μl primer (50 $ng/\mu l$), 2.0 μl of 10X PCR buffer, 1.2 µl of 25 mM MgCl₂, 0.75 µl Taq polymerase (1 U) and 1.0 µl dNTPs mix (10mM). Forty PCR amplification cycles were carried out in PCR machine [Eppendorf, Germany; model: Mastercycler(R) family] by denaturation at 94°C for 1 min, annealing at 37°C for 1 min and extension at 72°C for 1 min. Each PCR amplification reaction was preceded by an initial denaturation at 94°C for 5 min. followed by final extension at 72°C for 7 min. The amplified products were separated by electrophoresis in 1.5% (w/v) agarose (Genei, Bangalore) gel with 1X TBE buffer, stained with ethidium bromide (0.5 µg/ml) at 90 V for 3.0 to 3.5 h and photographed using gel documentation system (Alpha Innotech, USA Alpha Innotech, USA; model: AlphaImagerTM 3400). The sizes of the amplification product were estimated using 100 bp to 3.0 kb ladder ($\hat{O} \times 174$ DNA/ BsuRI (Hae III), Fermentas.

In ISSR, the polymerase chain reaction (PCR) was performed in 20 µl volumes containing 1 µl DNA (50 ng/ μl), 1 μl primer (50 ng/ μl), 2.0 μl of 10X PCR buffer, 1.2 µl of 25 mM MgCl₂, 0.75 µl Taq polymerase (1 U) and 1.0 µl dNTPs mix (10mM). Forty PCR amplification cycles were carried out in PCR machine [Eppendorf, Germany; model: Mastercycler (R) family] by denaturation at 94°C for 1 min, annealing at 51°C for 1 min and extension at 72°C for 2 min. Each PCR amplification reaction was preceded by an initial denaturation at 94°C for

5 min followed by final extension at 72°C for 7 min. The amplified products were separated by electrophoresis in 1.5% (w/v) agarose (Genei, Bangalore) gel with 1X TBE buffer, stained with ethidium bromide (0.5 ig/ml) at 90 V for 3.0 to 3.5 h and photographed using gel documentation system (Alpha Innotech, USAAlpha Innotech, USA; model: Alpha Imager TM 3400). The sizes of the amplification product were estimated using 100 bp to 3.0 kb ladder ($\hat{O} \times 174$ DNA/ BsuRI (Hae III), Fermentas. All the reactions were repeated in at least two independent experiments. All the amplified bands were scored as present or absent for each DNA sample and further, to analyze the relatedness among the species, a similarity matrix was generated from the binary data using DICE similarities coefficient in SIMQUAL program of NTSYS-PC Package (Rohlf, 2000). Cluster analysis was done with the Unweighed Pair Group Method with Arithetic Mean (UPGMA) in the SAHN program of NTSYS-PC Package, and constructed dendrogram based on genetic distances.

Selection of host differentials

Eight different Brassica spp. viz. Varuna (B. juncea), PBN-9501(B. napus), Kiran (B. carinata), Sinapis alba (local), PT-303 (B. rapa var. toria), PYS-6 (B. rapa var. yellow sarson), Eruca sativa (local), and B. nigra (local) were cross inoculated with each nine A. brassicae isolates collected from different Brassica spp. B. juncea, B. napus, B. carinata, Toria, Yellow sarson, B. nigra, E. sativa, S. alba and R. oleifera by single drop inoculation method (10 ul/leaf) at 40 DAS. The plants were kept in glasshouse under suitable environmental conditions. The observations on phenotypic disease reactions and percent disease index on each *Brassica* spp. against each isolate were recorded at 50 DAI. Based on different phenotypic disease reactions, the different Brassica spp. will be categorized in different groups as host differentials.

Long term preservation methods of A. brassicae

One year old A. brassicae isolates preserved in PDA alone (4°C), 10 per cent glycerol (-20°C) and 5 per cent DMSO (-20°C) collected from Amity University, Noida were used for the study of their sporulation in vitro and pathogenecity in glasshouse in two successive years. All thirty five A. brassicae isolates were re-cultured on PDA. The observations on mycelial growth and sporulation were recorded (0 days after incubation (DAI). The pathogenic virulence was tested on Varuna cultivar grown in pots under glasshouse by inoculating with the conidial suspension of each A. brassicae isolates separately and kept in glasshouse under suitable environmental conditions. The observations on disease severity against each isolates were recorded using 0-9 scale and per cent disease index was calculated.

Statistical analysis

Data obtained on various traits under laboratory and glasshouse experiments were assessed using STPR software and R-studio software.

Results and Discussion Cultural variability

Twenty A. brassicae isolates showed variable cultural characteristics viz. radial growth, colony colour (olive green, whitish gray, grayish, dark green, olive gray, whitish black, dark gray and greenish), shape (circular and irregular), zonation number and appearance (compressed and fluffy) on PDA. On the basis of colony colour i.e. olive green, whitish gray, grayish, dark green, olive gray, whitish black, dark gray and greenish, the isolates were categorized into 8 groups viz. Group I (olive green): AB-B. carinata, AB-Toria, AB-R. oleifera, AB-B. juncea Karnal, AB-B. juncea Kangara and AB-B. botrytis; Group II (whitish gray): AB-Yellow sarson, AB-B. rugosa, AB-B. juncea Bihar and AB-B. juncea Jammu; Group III (grayish): AB-B. juncea Pantnagar, AB-B. juncea Punjab and AB-B. pekinensis; Group IV (dark green): AB-S. alba and AB-B. caulorapa; Group V (olive gray): AB-B. napus and AB-E. sativa; Group VI (whitish black) AB-B. nigra, Group VII (dark gray) AB-R. sativus; Group VIII (greenish) AB-B. oleracia (Table 2; Fig 1). The present results are in accordance with Kaur et al. (2007), Sharma et al. (2013), Pramila et al. (2014) and Saha et al. (2015) who reported cultural variability among different A. brassicae collected from different geographical regions of India on the basis of colony colour, shape, zonation number and appearance.

Table 2: Cultural characters of A. brassicae isolates on Potato dextrose agar medium

A. brassicae isolate	Colony colour	Growth	Shape	Margin	Zonation (No.)	Appearance
AB-B. juncea Pantnagar	grayish	fast	circular	smooth	2	fluffy
AB-B. napus	olive gray	medium	circular	smooth	2	fluffy
AB-B. carinata	olive green	medium	circular	smooth	2	compressed
AB-Toria	olive green	medium	irregular	smooth	3	compressed
AB-Yellow sarson	whitish gray	medium	irregular	smooth	4	fluffy
AB-E. sativa	olive gray	medium	irregular	smooth	2	compressed
AB- S. alba	dark green	medium	irregular	smooth	3	compressed
AB- R. oleifera	olive green	medium	circular	smooth	-	compressed
AB-B. nigra	whitish black	medium	circular	smooth	-	compressed
AB-B. rugosa	whitish gray	medium	irregular	smooth	-	compressed
AB-B. juncea Karnal	olive green	fast	circular	smooth	-	compressed
AB- <i>B. juncea</i> Punjab	grayish	medium	irregular	smooth	-	fluffy
AB-B. juncea Bihar	whitish gray	medium	circular	smooth	4	compressed
AB- <i>B. juncea</i> Jammu	whitish gray	medium	irregular	smooth	2	compressed
AB-B. juncea Kangara	olive green	medium	circular	smooth	-	compressed
AB-R. sativus	dark gray	medium	irregular	smooth	2	fluffy
AB-B. pekinensis	grayish	medium	irregular	smooth	-	fluffy
AB-B. oleracea	greenish	slow	circular	smooth	-	compressed
AB-B. botrytis	olive green	medium	irregular	smooth	2	compressed
AB-B. caulorapa	dark green	slow	irregular	smooth	2	compressed

Fig 1: Cultural characteristics of A. brassicae isolates on PDA medium

Table 3:	Radial	growth of <i>A</i> .	brassicae	isolates	on different	solid media

A. brassicae isolate	Radial growth (mm)								
	Potato dextrose	Radish dextrose	Cauli flower dextrose	Carrot Potato dextrose	Mustard seed leaf	Brassica dextrose seed	Maize extract dextrose	Modified V-8 juice	Mean
AB-B. juncea Pantnagar	82.0	72.7	71.7	70.3	74.3	64.3	68.7	60.7	76.9
AB-B. napus	68.0	61.3	63.3	42.0	65.0	48.3	45.0	47.0	64.3
AB-B. carinata	63.0	72.0	65.7	67.3	67.7	50.0	65.7	52.3	67.9
AB-Toria	80.0	73.3	67.3	54.7	71.0	69.0	57.7	65.3	76.34
AB-Yellow sarson	76.0	70.3	66.0	64.7	72.0	65.7	61.0	48.3	72.9
AB-E. sativa	63.3	68.3	62.0	62.7	60.3	63.3	50.3	49.7	66.1
AB- S. alba	62.3	44.7	48.0	59.7	57.7	62.7	46.7	59.7	52.6
AB-R. oleifera	68.0	57.7	57.3	60.7	52.7	60.3	52.7	51.0	62.3
AB-B. nigra	70.3	69.0	64.7	70.0	69.7	59.7	66.3	42.0	69.6
AB-B. rugosa	68.0	75.7	69.0	55.0	53.7	58.7	63.3	55.7	72.2
AB-B. juncea Karnal	78.7	72.7	69.7	60.3	64.0	65.7	64.0	62.3	75.4
AB-B. juncea Punjab	76.7	56.3	68.7	70.0	62.0	68.3	58.0	48.0	65.5
AB-B. juncea Bihar	70.7	62.7	54.7	69.0	68.3	59.7	42.0	62.7	66.3
AB-B. juncea Jammu	70.3	63.3	60.3	62.0	60.0	63.7	66.7	61.7	66.5
AB-B. juncea Kangara	66.0	67.7	42.0	53.3	55.0	47.7	57.7	59.7	66.9
AB-R. sativus	71.0	70.3	65.0	62.7	64.7	62.7	62.7	55.7	70.6
AB-B. pekinensis	67.7	66.7	43.3	64.7	68.3	60.0	58.7	61.7	67.1
AB-B. oleracea	53.3	66.3	64.0	61.7	62.7	68.0	46.3	62.0	60.5
AB-B. botrytis	69.3	62.3	66.3	62.0	55.0	62.7	47.7	53.0	65.5
AB-B. caulorapa	49.7	61.0	50.3	68.7	51.3	55.0	48.3	50.0	55.9
CD 5%	4.3	3.8	3.3	3.5	4.3	3.2	3.3	3.3	2.5
CV	3.7	3.5	3.3	3.4	4.2	3.2	3.5	3.6	3.8

Eight different nutrient media brought variation in the radial growth of twenty different A. brassicae isolates (Table 3). Significant maximum radial growth (82.0 mm) was observed in PDA followed by Brassica leaf extract (74.3 mm) and Radish root dextrose (72.7 mm), while significantly minimum growth in Maize seed dextrose (42.00 mm) medium.

On the basis of radial growth on different solid media the A. brassicae isolates were categorised into seven different groups Group I: AB-B. juncea Pantnagar (76.9 mm), AB-Toria (76.3 mm), AB-B. juncea Karnal (75.4 mm); Group II: AB-Yellow sarson (72.9 mm), AB-B. rugosa (72.2 mm) AB-R. sativus (70.6 mm); Group III: AB-B. nigra (69.6 mm), AB-B. carinata (67.9 mm), AB-B. pekinensis (67.1 mm); Group IV: AB-B. juncea Kangara (66.9 mm), AB-B. juncea Jammu (66.5 mm), AB-B. juncea Bihar (66.3 mm), AB-E. sativa (66.1 mm), AB-B. juncea Punjab (65.5µ mm), AB-B. botrytis (65.5 mm); Group V: AB-B. napus (64.3 mm), AB- R. sativus (local) (62.3 mm); Group VI: AB-B. oleracea (60.5 mm); Group VII: AB-B. caulorapa (55.9 mm), AB-S. alba (52.6 mm).

Earlier workers also reported variability on the basis of radial growth on different media. In the present findings PDA was found best in terms of radial growth of all the A. brassicae isolates followed by Radish dextrose agar and Brassica leaf extract agar media which are in accordance with the findings of Kumar and Singh (2003) and Singh et al. (2015). However, Selvamani et al. (2013) found Cauliflower Leaf Extract Agar as the best medium followed by Potato Dextrose Agar.

Morphological variability

The A. brassicae isolates purified through single spore isolation showed significant morphological variability in respect of conidial length, conidial width, conidial beak length, number of septa and number of cells (Table 4; Fig 2). Average conidial length

Table 4: Morphological variability among different A. brassicae isolates

A. brassicae isolate	Conidial length (µm)	Conidial beak length (µm)	Conidial width (µm)	Horizontal septa (No.)	Vertical septa (No.)	Number of cells (No.)
AB-B. juncea Pantnagar	61.0	17.6	9.1	5.3	1.5	9.5
AB-B. napus	58.5	20.1	10.0	6.8	2.0	9.3
AB-B. carinata	185.3	93.4	25.6	14.8	5.0	17.3
AB-Toria	107.8	31.9	15.1	8.8	4.0	11.3
AB-Yellow sarson	59.7	12.0	10.1	7.3	1.5	10.5
AB-E. sativa	91.4	31.8	11.7	8.3	2.8	10.3
AB- S. alba	175.8	78.2	22.8	12.5	5.0	16.5
AB- R. oleifera	51.8	15.0	10.6	6.3	1.8	9.5
AB-B. nigra	50.6	16.0	10.1	7.8	1.5	9.5
AB-B. rugosa	38.9	11.0	8.1	6.3	1.5	8.8
AB-B. juncea Karnal	131.8	41.2	28.2	10.0	4.8	13.5
AB-B. juncea Punjab	62.5	19.9	12.3	6.5	2.0	9.3
AB-B. juncea Bihar	57.9	18.4	10.0	8.0	2.3	11.5
AB-B. juncea Jammu	44.3	14.7	9.9	5.8	0.8	7.8
AB-B. juncea Kangara	45.4	15.6	8.3	6.3	1.0	8.5
AB-R. sativus	44.7	16.0	9.4	7.0	1.5	10.0
AB-B. pekinensis	39.1	10.8	8.7	5.5	0.8	8.5
AB-B. oleracea	30.9	8.7	8.8	5.8	1.3	8.0
AB-B. botrytis	40.5	5.5	9.0	6.0	0.8	8.8
AB-B. caulorapa	29.0	7.1	6.6	3.5	0.8	5.8
CD 5%	5.7	2.5	2.0	1.6	1.0	2.0
CV	5.8	7.3	11.4	15.5	34.4	13.7

and width in A. brassicae isolates varied from 29.0 to 182.3µm x 6.6 to 28.2µm. significantly maximum conidial length was in AB-B. carinata isolate (182.3µm) and minimum in AB-B. caulorapa (29.0µm). Maximum conidial width was in AB-B. juncea Karnal isolate (28.2µm), while minimum in AB-B. caulorapa (6.6 µm). Average conidial beak length varied from 5.5 to 93.4 µm. Maximum conidial beak length was in AB-B. carinata isolate (93.4µm), while minimum in AB-B. botrytis (5.5µm). Average number of horizontal septa varied from 3.5 to 14.8 no. Maximum horizontal septum was in AB-B. carinata isolate (14.8 no.), while minimum in AB-B. caulorapa (3.5 no.). Vertical septa varied from 0.75 to 5.0 no. Maximum vertical septa were in AB-B. carinata isolate (5.0 no.), while minimum in AB-B. caulorapa (0.75 no.). The largest conidia and highest number of septa was observed in AB-B. carinata isolate followed by AB-S. alba, AB-B. juncea Karnal and AB-Toria isolates, while smallest conidia and lowest number of septa was in AB-B. caulorapa isolate. Earlier workers have also characterized different A. brassicae isolates on the basis of morphological characters viz. conidial length, conidial width, number of horizontal and vertical septa, beak length (Mehta et al., 2003; Kaur et al., 2007; Meena et al., 2012; Selvamani et al., 2013; Saha et al., 2015). They reported conidia length from 51.4-481.2 µm, width from 6.9-285.9µm, beak length 16.3-266.9 µm, number of horizontal and vertical septa from 3.0-9.0 and 1-2 respectively.

In the present studies on the basis of conidial length A. brassicae isolates were categorized into 4 different groups viz. Group I (200-150 µm): AB-B. carinata (182.3µm), AB-S. alba (175.8µm); Group II (150-100 μm): AB-B. juncea Karnal (131.8μm), AB-Toria (107.8μm); Group III (100-50 μm): AB-E. sativa (91.4μm), AB- B. juncea Punjab (62.5μm), AB-B. juncea Pantnagar (61.0µm), AB-Yellow sarson (59.7µm), AB-B. napus (58.5µm), AB-B. juncea Bihar (57.9µm), AB-R. oleifera (51.8µm), AB-B. nigra (50.6μm); Group IV (< 50): AB-R. sativus (44.7µm), AB-B. juncea Jammu (44.3µm), AB-B. juncea Kangara (45.4µm), AB-B. pekinensis (39.1µm), AB-B. rugosa (38.90µm), AB-B. botrytis (40.5µm), AB- B. oleracea (30.9µm) and AB-B. caulorapa (29.0µm).

Pathogenic variability

The qualitative characters viz. spot colour, periphery colour, concentric rings, central point and yellow halo were recorded on leaves at 40 days after inoculation (DAI). The A. brassicae isolates showed variability in spot colour i.e. brown and gray in periphery colour i.e. brown, gray and green; presence or absence of concentric rings; presence or absence of central point and colour (black, brown, gray); presence or absence of yellow halo. The dendogram constructed (using statistical software R studio) for pathogenic variation among A. brassicae isolates (Fig. 3) was categorized in 5 groups. Group I: AB-B. caulorapa; Group II: AB-B. rugosa, AB-E. sativa, AB-S. alba; Group III: AB-B. juncea Kangara, AB-B. juncea Jammu; Group IV: AB-B. juncea Punjab, AB-R. oleifera, AB-B. nigra, AB-B. juncea Bihar, AB-B. juncea Karnal, AB-B. juncea Pantnagar, AB-Yellow sarson, AB-Toria; Group V: AB-B. carinata, AB-B. pekinensis AB-B. botrytis, AB-R. sativus, AB-B. napus and AB-B. oleracea. The quantitative character viz. per cent disease index was recorded during 2014-15 and 2015-16, 50 DAI revealed significantly maximum per cent disease index and

AUDPC in Varuna with AB-B. juncea Pantnagar isolate (49.3 %, 44.8 % & 313.4, 278.3), followed by AB-B. juncea Karnal (42.6 %, 37.0% & 262.5, 220.8), AB-Toria (41.9%, 37.8% & 259.3, 226.4) and AB-Yellow sarson (39.6 %, 34.1 % & 244.9, 202.1), respectively and were at par with each other. Minimum per cent disease index and AUDPC was observed with AB-S. alba (18.9 %, 17.0% & 102.3, 81.5) followed by AB-E. sativa (20.4%, 18.89 %& 113.4, 93.5), respectively (Fig 4 & 5). In the present studies AB-B. juncea Pantnagar isolate was found most virulent, while AB-S. alba least virulent.

Studies on pathogenic variability are essential for the development of pre-breeding populations as strategic defense mechanism. The pathogenic variability among A. brassicae isolates on host genotypes, on the basis of qualitative and quantitative characters were also reported by earlier workers (Kolte et al., 1991; Kaur et al., 2007; Khan et al., 2007; Goyal et al., 2013; Pramila et al., 2014 and Saha et al., 2015).

Molecular variability **RAPD** markers

Fig 4: Percent disease index and AUDPC on B. juncea cv. Varuna 2014-15

Fig 3: Dendogram on the basis of pathogenic qualitative characteristics of A. brassicae isolates (20 no.) on B. juncea var. Varuna

Fig 5: Percent disease index and AUDPC on B. juncea cv. Varuna 2015-16

Twenty random decamer oligonucleotide primers were screened initially, of these two primers did not give any amplification. Eighteen primers therefore used in molecular diversity analysis of A. brassicae isolates. A total of 259 reproducible and scorable polymorphic bands ranging from 100-1350 bp were generated for all twenty A. candida isolates with 18 RAPD primers. The number of polymorphic bands produced were maximum (21) with the primers LC 106 and LC 79, while minimum (7) with primer LC 94.

The UPGMA (un-weighted pair group method with arithmetic mean) dendrogram was constructed using Jaccard's similarity coefficient of RAPD marker data of 18 polymorphic primers generated for twenty A. brassicae isolates employing the program NTSYS 2.11 (Table 5, Fig 6, Plate 1). RAPD marker based dendrogram categorised twenty A. brassiace isolates into two major groups i.e. A and B. Group A and Group B were differentiated to each other at similarity coefficient of 0.59. In Group A isolates were categorized into Cluster I, consisted of 5 isolates AB-R. sativus, AB-B. pekinensis, AB-B. oleracea, AB-B. botrytis and AB-B. caulorapa. Intra cluster similarity coefficient between pairs of isolates within Cluster I varied from 0.87-0.76. In Group B isolates were categorized into Cluster II, III, IV and V on the basis of similarity coefficient. Cluster II consisted of only one isolate AB-B. rugosa and differentiated from remaining of the clusters of Group B at similarity coefficient of 0.70. Cluster III

Plate 1: RAPD profile of A. brassicae isolates with primer LC 80

consisted of 9 isolates viz. AB-E. sativa, AB-S. alba, AB-R. sativus (local), AB-B. nigra, AB- B. juncea Karnal, AB-B. juncea Punjab, AB-B. juncea Jammu, AB-B. juncea Bihar and AB-B. juncea Kangara and exhibited similarity coefficient of 0.70 with Cluster IV. Similarity coefficient between isolates of Cluster III varied from 0.76 to 0.91. Cluster IV consisted of only one isolate i.e. AB-Yellow sarson and established similarity coefficient of 0.7 with Cluster V. Cluster V consisted of 4 isolates viz. AB-B. juncea Pantnagar, AB-B. napus, AB-B. carinata and AB-Toria and pair wised similarity coefficient varied from 0.8 - 0.7.

ISSR markers

Five ISSR primers were screened initially, of these four ISSR primers gave distinct and reproducible marker profile against A. brassicae isolates. A total of 51 reproducible and scorable polymorphic bands ranging from 100-950 bp were generated with four ISSR primers for genetic polymorphisms among the isolates. The number of polymorphic bands were

Table 5: Similarity coefficient for A. brassicae isolates using RAPD primers

```
1 000
            0.733 1.000
3.
            0.791 0.810 1.000
            0.768 0.764 0.822 1.000
4
5.
            0.710\ 0.729\ 0.718\ 0.764\ 1.000
            0.741 0.698 0.764 0.810 0.768 1.000
            0.756 0.706 0.764 0.841 0.745 0.861 1.000
            0.745 0.702 0.752 0.814 0.733 0.810 0.880 1.000
            0.710\ 0.667\ 0.718\ 0.803\ 0.729\ 0.837\ 0.876\ 0.818\ 1.000
10.
            0.675\ 0.648\ 0.667\ 0.776\ 0.710\ 0.710\ 0.733\ 0.752\ 0.733\ 1.000
11.
            0.714\ 0.671\ 0.745\ 0.806\ 0.733\ 0.826\ 0.872\ 0.830\ 0.849\ 0.729\ 1.000
            0.772\ 0.656\ 0.706\ 0.776\ 0.756\ 0.795\ 0.857\ 0.837\ 0.826\ 0.729\ 0.891\ 1.000
12.
            0.749 \ 0.667 \ 0.710 \ 0.764 \ 0.760 \ 0.791 \ 0.783 \ 0.803 \ 0.768 \ 0.749 \ 0.833 \ 0.826 \ 1.000
13
14.
            0.749\ 0.683\ 0.733\ 0.818\ 0.768\ 0.783\ 0.837\ 0.833\ 0.806\ 0.779\ 0.864\ 0.872\ 0.837\ 1.000
            0.679\ 0.644\ 0.687\ 0.741\ 0.729\ 0.737\ 0.776\ 0.741\ 0.729\ 0.694\ 0.741\ 0.787\ 0.752\ 0.799\ 1.000
15.
16.
            0.583 \ 0.571 \ 0.606 \ 0.613 \ 0.648 \ 0.586 \ 0.610 \ 0.613 \ 0.563 \ 0.637 \ 0.621 \ 0.621 \ 0.640 \ 0.648 \ 0.602 \ 1.000
            0.548 0.552 0.571 0.579 0.613 0.536 0.559 0.571 0.521 0.617 0.579 0.571 0.613 0.613 0.567 0.864 1.000
17.
18
            0.552 0.540 0.559 0.575 0.610 0.563 0.579 0.613 0.525 0.644 0.583 0.590 0.625 0.633 0.579 0.791 0.772 1.000
19.
            0.555\ 0.544\ 0.548\ 0.555\ 0.613\ 0.528\ 0.536\ 0.586\ 0.498\ 0.602\ 0.540\ 0.548\ 0.621\ 0.598\ 0.583\ 0.756\ 0.722\ 0.872\ 1.000
            0.598 \ 0.579 \ 0.606 \ 0.621 \ 0.640 \ 0.579 \ 0.610 \ 0.637 \ 0.579 \ 0.652 \ 0.629 \ 0.621 \ 0.671 \ 0.656 \ 0.586 \ 0.776 \ 0.764 \ 0.845 \ 0.810 \ 1.000
                                                             8
                                                                         10
                                                                                       12
                                                                                              13 14
                                                                                                            15
                                                                                                                          17
                                                                                 11
                                                                                                                  16
                                                                                                                                18
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Plate 2: ISSR profile of *A. brassicae* isolates with primer ISSR 2

dendogram categorised twenty A. brassiace isolates into two major groups i.e. A and B. Group A and B were differentiated at similarity coefficient of 0.67. Group A isolates were categorized into Cluster I, consisted of 5 isolates viz. AB-R. sativus, AB-B. pekinensis, AB-B. oleracea, AB-B. botrytis and AB-B. caulorapa. Intra cluster similarity coefficient between pairs of isolates within Cluster I varied from 0.90-0.79. Group B isolates were categorized into Cluster II, III, IV and V. Cluster II consisted only one isolate i.e. AB-B. napus and differentiated from remaining of the clusters of Group B at similarity coefficient of 0.74. Cluster III again had only isolate i.e. AB-R. oleifera, and exhibited similarity coefficient of 0.75 with Cluster IV. Similarity coefficient of Cluster III varied from 0.75 to 0.98. Cluster IV consisted of six isolates viz. AB-S. alba, AB-B. juncea Jammu, AB-B. rugosa, AB- B.

Fig 6: Dendogram of *A. brassicae* isolates using RAPD primers

maximum (15) with ISSR 2 and ISSR 10, while minimum (9) with the ISSR 12 primers.

The ISSR markers were also used to determine level of genetic diversity among twenty *A. brassicae* isolates. The UPGMA dendrogram was constructed using Jaccard's similarity coefficient of ISSR marker and data were generated using four ISSR primers (Table 6, Fig 7, Plate 2). ISSR marker based

Fig 7: Dendogram of *A. brassicae* isolates using ISSR primers

Table 6: Similarity coefficient for A. brassicae isolates using ISSR primers

```
1.000
            0.784 1.000
3.
            0.882 0.784 1.000
            0.862 0.725 0.862 1.000
4.
            0.862 0.764 0.862 0.921 1.000
            0.843 0.823 0.882 0.862 0.862 1.000
            0.745  0.686  0.745  0.882  0.803  0.823  1.000
            0.705 0.725 0.745 0.764 0.725 0.784 0.803 1.000
            0.843 0.705 0.882 0.901 0.901 0.843 0.823 0.705 1.000
10.
            0.803 0.745 0.843 0.862 0.862 0.882 0.862 0.745 0.921 1.000
11
            0.823 0.764 0.862 0.882 0.882 0.901 0.882 0.764 0.941 0.980 1.000
12.
            0.725  0.627  0.725  0.823  0.745  0.764  0.823  0.705  0.803  0.843
                                                                               0.862 1.000
13.
            0.882\ 0.745\ 0.921\ 0.941\ 0.941\ 0.882\ 0.823\ 0.745\ 0.960\ 0.921
                                                                               0.941 0.803 1.000
            0.784\ \ 0.686\ \ 0.784\ \ 0.843\ \ 0.803\ \ 0.823\ \ 0.921\ \ 0.764\ \ 0.862\ \ 0.901
                                                                               0.921 0.862 0.862 1.000
14.
15.
            0.764 0.705 0.803 0.862 0.823 0.843 0.862 0.745 0.882 0.921
                                                                               0.941 0.882 0.882 0.901 1.000
            0.667 0.529 0.705 0.764 0.725 0.667 0.725 0.647 0.745 0.705
16.
                                                                               0.725 0.627 0.784 0.725 0.705 1.000
17
            0.667 0.529 0.705 0.725 0.686 0.667 0.686 0.607 0.705 0.667
                                                                               0.686 0.588 0.745 0.686 0.667 0.764 1.000
            0.705 \ \ 0.568 \ \ 0.784 \ \ 0.764 \ \ 0.725 \ \ 0.705 \ \ \ 0.725 \ \ 0.725 \ \ 0.745 \ \ 0.705
                                                                               0.725\ 0.667\ 0.784\ 0.725\ 0.705\ 0.803\ 0.803
                                                                                                                              1.000
18.
19.
            0.588 0.450 0.667 0.686 0.647 0.588 0.647 0.607
                                                                 0.667 0.627
                                                                               0.647\ 0.549\ 0.705\ 0.647\ 0.627\ 0.764\ 0.764
                                                                                                                              0.843 1.000
            0.607 \ \ 0.470 \ \ 0.686 \ \ 0.705 \ \ 0.667 \ \ 0.607 \ \ 0.667 \ \ 0.627 \ \ 0.686 \ \ 0.647
                                                                               0.667 0.568 0.725 0.667 0.647 0.823 0.823 0.862 0.901 1.000
20.
                     2
                           3
                                  4
                                         5
                                               6
                                                      7
                                                             8
                                                                    9
                                                                         10
                                                                                11
                                                                                       12.
                                                                                              13
                                                                                                     14
                                                                                                            15
                                                                                                                  16
                                                                                                                         17
                                                                                                                                18
                                                                                                                                       19
                                                                                                                                             20
```

juncea Karnal, AB-B. juncea Kangara and AB-B. juncea Punjab and established similarity coefficient of 0.84 with Cluster V. Cluster V consisted of seven isolates viz. AB-B. juncea Pantnagar, AB-B. carinata, AB-Toria, AB-B. nigra, AB-B. juncea Bihar, AB-Yellow sarson and AB-E. sativa pair wised similarity coefficient varied from 0.96-0.86.

In the present study, molecular profile based on dendogram generated by 18 RAPD and 4 ISSR markers revealed genetic diversity among different A. brassicae isolates as both RAPD and ISSR primers successfully differentiated A. brassicae isolates into different clusters. Goyal et al. (2013), Pramila et al. (2014) and Saha et al. (2015) also studied genetic variability among A. brassicae isolates RAPtD primers. Sharma et al. (2013) characterized 32 Indian isolates using ITS primers between 550- 600 bp. Kumar et al. (2014) performed DNA fingerprinting using ISSR primers to study genetic diversity among 32 isolates of A. brassicae isolates.

Selection of host differentials

Based on phenotypic diseases reactions against nine

different A. candida isolates, the Brassica genotypes were categorized into 4 different groups as host differentials viz., Group I: Varuna, PT-303, PYS-6 and B. nigra which showed susceptible disease reaction (S) i.e. coalescing larger brown spots with a distinct margin yellow halo, 26-50 per cent leaf area covered by the lesions, against all the isolates; Group II: PBN-9501 and Kiran which showed moderately resistant reactions (MR) i.e. small roundish slightly sporulating small brown necrotic spots, 11-25 per cent leaf area covered by lesions; Group III: E. sativa which showed resistant disease reaction (R) i.e. pinpoint size or small brown necrotic spots, less than 5-10 per cent leaf area covered by lesions: Group IV: S. alba which showed disease resistant reaction (R) i.e. pinpoint size or small brown necrotic spots, less than 5-10 per cent leaf area covered by lesions against AB-B. juncea Pantnagar, AB-Toria, AB-Yellow sarson and AB-S. alba isolates, while showed immune response against AB-B. napus, AB-B. carinata, AB-B. nigra, AB-E. sativa and AB-R. oleifera isolates (Table 7). Earlier workers (Sangwan and Mehta, 2007 and Meena et al., 2012) used seventeen and six different Brassica genotypes respectively for the

Table 7: Grouping of Brassica spp. based on phenotypic disease reactions against different A. brassicae isolates

Brassica spp.				A. b	rassicae is	solate			
•	AB-B. juncea	AB-B.	AB-B. carinata	AB- Toria	AB- Yellow	AB-E. sativa	AB- S. alba	AB- R. oleifera	AB-B. nigra
	Pantnagar				sarson				
Group I									
Varuna	S	S	S	S	S	S	S	S	S
Toria	S	S	S	S	S	S	S	S	S
Yellow sarson	S	S	S	S	S	S	S	S	S
B. nigra	S	S	S	S	S	S	S	S	S
Group II									
B. napus	MR	MR	MR	MR	MR	MR	MR	MR	MR
B. carinata	MR	MR	MR	MR	MR	MR	MR	MR	MR
Group III									
E. sativa	R	R	R	R	R	R	R	R	R
Group IV									
S. alba	R	-	-	R	R	-	R	-	-

⁽⁻⁾ no reaction

differentiation of different A. brassicae isolates on the basis of phenotypic disease reactions.

Long term preservation methods of A. brassicae culture

Mycelial growth was observed in all of the A. brassicae cultures (56 nos.) preserved in PDA (4°C), 10 per cent glycerol (-20°C) 5 per cent DMSO (-20°C). However, sporulation was observed only in thirty five A. brassicae cultures. On PDA (4°C) out of 35 isolates only 23 isolates sporulate well; On 10 per cent glycerol (-20°C) out of 9 isolates only 6 isolates sporulate well; On 5 per cent DMSO (-20°C) out of 12 isolates only 7 isolates sporulate well. The above findings revealed that culture preservation in 10 per cent glycerol at -20°C was the best in terms of sporulation (66.7%) followed by PDA at 4°C (65.7%) and could be used for preservation and storage of A. brassicae isolates at least for 2 years (Table 8). The present findings are in accordance with the work of Aneja et al. (2014) who reported that A. brassicae isolates, could be maintained by preserving in PDA at (4°C); 10% glycerol and 5% DMSO (-20°C) at least 6 months of storage.

Conclusion

In present studies significant variation in cultural, morphological, pathogenic and molecular variability

was observed in A. brassiace isolates irrespective to geographical locations and Brassica spp. On the basis of cultural, morphological, pathogenic and molecular variability, the A. brassiace isolates are categorized into 5 major groups. Group I: comprised 5 isolates viz. AB-R. sativus, AB-B. pekinensis, AB-B. oleracea, AB-B. botrytis and AB-B. caulorapa; Group II: comprised 8 isolates viz. AB-B. juncea Pantnagar, AB-Yellow sarson, AB-Toria, AB-B. nigra, AB-R. oleifera, AB-B. juncea Karnal, AB-B. juncea Punjab and AB-B. juncea Bihar; Group III: comprised 4 isolates viz. AB-B. napus, AB-B. carinata, AB-E. sativa, AB-S. alba; Group IV: comprised 2 isolates viz. AB-B. juncea Kangara and AB-B. juncea Jammu; Group V: comprised 1 isolate *i.e.* AB-B. rugosa. These A. brassicae isolates could be used during screening of resistant sources in Brassica genotypes against Alternaria blight disease.

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Table 8: Effect of different preservation methods on the growth and sporulation of A. brassicae isolates

Observation		A. brassicae isolates				
		Methods of preservation				
	PDA(4°C)	10% glycerol (-20°C)	5% DMS	O (-20°C)		
Mycelia growth with sporulation	ABc-D01, ABc-D02, ABo-D03, ABo-P03, ABo-P04, ABc-P08, ABc-P10, ABc-P12, ABc-L06, ABc-L08, ABc-L11, ABc-L12, ABc-L13, ABc-L14, ABc-L20, ABc-L01, ABc-L04, ABc-L09, ABc-L16, ABc-L18, ABc-Kg02, ABc-B01, ABc-B03	ABc-D01, ABc-D02, ABo-D03, ABo-P03, ABo-P04, ABc-P08		, ABc-D02, ABo-D03, B, ABo-P04, ABc-P10,		
Mycelia growth without sporulation	ABa-Kn02, ABa-B02, ABo-D04, ABo-D05, ABc-D06, ABc-D07, ABo-P02, ABo-P09, ABc-P11, ABc-L02, ABc-L03, ABc-L05	ABo-D04, ABo-D05, ABo-P02		1, ABo-D05, ABo-P02, 9, ABc-P11		
Sporulation perce	ntage	65.7%	66.7%	58.3%		

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