

Morphological, Pathological, Biochemical and Molecular Characterization of *Ralstonia solanacearum* Isolates in Bangladesh

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One thirty samples (fifty-five potato tubers, twenty-seven potato stems, three chili stems, twenty-eight soil samples, five weed samples, three banana leaves, and nine water samples) were examined and one hundred six (106) Bangladeshi isolates of Ralstonia solanacearum were isolated and identified. Isolation was made on selective media (Tetrazolium chloride media) and R. solanacearum was identified based on morphological, pathological and biochemical properties and polymerase chain reaction (PCR) by using the species-specific primers. Studies showed that 81.54% (106) samples were positive on tetrazolium chloride solid medium. Among them 90 isolates were virulent and rest of them were avirulent. Fifty isolates were selected for chemical characterization based on hypersensitivity test. R. solanacearum is gram negative, aerobic facultative bacteria on the basis of chemical characterization. Fifty tested isolates expressed as race 3 while in biovar test forty-eight showed as biovar III and the rest two showed as biovar I. In nine tested isolates from the three districts a species-specific band of 280 bp was amplified in PCR that confirmed the identity of R. solanacerum.

Keywords

Brown Rot of Potato, Latent Infection, Ralstonia solanacearum, Biovar, Race

1. Introduction

R. solanacearum [1] formerly called *Pseudomonas solanacearum* [2] is the most destructive, damaging soil-borne pathogen [3] [4] [5]. Complex species [6] *Ral-stonia solanacearum* is diversified and widely distributed with a wide host range

including 44 families and more than 200 plant species [7] [8]. It is very destructive pathogen of potatoes in temperate, subtropical and tropical regions throughout the world [9] [10].

R. solanacearum is gram-negative, rod-shaped bacterium measuring $0.5 - 0.7 \times 1.5 - 2.0 \ \mu\text{m}$ in size. This pathogen is severed at 24°C - 35°C temperature but optimum at 28°C - 32°C in aerobic conditions [11] [12]. *Ralstonia solanacearum* caused bacterial wilt, southern wilt and brown rot of potato world wide. Wilting and chlorosis of the leaves located at the tip of the branches are the first visible symptoms of *R. solanacearum* infection on potato (*Solanum tuberosum*), stunting and petiole epinasty may also occur due to the blocking of the vessels [13]. Infection of potato tubers with *R. solanacearum* may become latent under conducive environmental conditions [13].

R. solanacearum is historically sub divided into five races based on host range and five biovars based on different abilities to produce acid from a panel of a carbohydrate [7]. Race 1 hosts all solanaceous crops and many other plants which are available in Asia, Australia, America, Bangladesh, China, India, Japan, Nepal, Pakistan, Sri Lanka etc. belonged to biovar III, IV, and I; race 2 hosts triploid bananas, other *Musa* spp. are available in Caribbean, Brazil, Philippines which is belonged to biovar I; race 3 hosts only potato and tomato which is available worldwide except US and Canada belonged to biovar II or IIA; race 4 hosts ginger and unknown hosts available in Australia, China, Hawaii, India, Japan, Mauritius, South Asia and India belonged to biovar III and IV; and race 5 hosts mulberry tree in China which belongs to biovar V. However, typical race 3 strains are sometimes referred to as biovar IIA and new race 3 strains from the Amazon basin have been placed in a new biovar designed as IIT or NII and their relation to races is unclear [14] [15]. Based on the geographical distribution and interaction with climatic conditions, R. solanacearum was established in all agro-ecological zones of Bangladesh [16]. Ralstonia solanacearum caused the brown rot of potato found in Panchagar, Nilphamari, Rangpur, Lalmonirhat, Bogura, Joypurhat, Jashore and Rajshahi districts of Bangladesh [16]. The variation in *R. solanacea*rum isolates of potato which was observed different growing region of Bangladesh that belonging to race 3 and biovar III [17] [18] [19].

Therefore, the objectives of this study were to 1) isolate and identify *R. sola-nacearum* from potato, chili, banana, soil and water in different potato growing areas of Bangladesh by morphological, pathological, biochemical and molecular tools and 2) to determine race and biovar of *R. Solanacearum* present in Bangladesh.

2. Materials and Methods

2.1. Collection of Samples

The sample collection was carried out in Munshiganj, Narayangonj and Nilphamari districts under nine upazila viz. Munshiganj Sadar, Tongibari, Sirajdikhan, Narayganj Sadar, Sonargaon, Rupgonj, Domar, Nilphamari Sadar and Dimla from the field, farmer storage and cold storage during May 2017 to November 2019 in Bangladesh. Samples were collected randomly from potato tubers, potato stems, chili stems, banana leaves, weed, soil and irrigation water. Symptoms of the disease were studied by visual observation as per standard procedure [10]. In order to increase the likelihood for detection sampling was made at least three replications [20]. Collected samples were put in polyethylene bags immediately after collection to protect them from drying. Labelling of each sample was done with the sample's location, date and sample identification number. Samples were carried to laboratory within 24 - 26 hours after collection for investigation.

2.2. Observation of Visual Symptoms

First samples were cleaned and then the collected samples were stored in the refrigerator between 4°C and 10°C [15] [20]. The infected tuber and plant samples were checked for oozing out which is the simple method by using sterilized knife and a tube containing sterilized distilled water [21] [22]. All collected samples identification of the disease was finally confirmed through isolation, different biochemical tests and molecular characterization.

2.3. Isolation of Ralstonia solanacearum

Ralstonia solanacearum were isolated by dilution plate method. The collected samples from potato tuber and stem washed with sterilized distilled water containing 1% Clorox and then cut several small pieces. The cut surface was sterilized by dipping them in 5% sodium hypochlorite solution for 2 - 3 minutes. It was then washed three times with sterile water. After surface sterilization, the cut pieces were kept in a test tube containing 3 - 4 ml of sterile water and kept for 30 minutes for bacterial streaming and getting stock. 1 ml of this stock solution was transferred with the help of sterile pipette into the second test tube containing 9 ml sterile water and shaken thoroughly for 10⁻¹ dilution. Similarly, final dilution was made up to 10⁻⁴. Dried soil samples (1 g) were dissolved in 10 ml of sterilized distilled water. The samples were agitated for 20 minutes and a dilution plate method 10⁻⁴ was then carried out. Collected water samples (1 ml) were dissolved in 10 ml of sterilized distilled water. The samples were shake for 20 minutes and dilution plate method 10^{-4} was accomplished. Weed and other crop (chilli) collar and root regions were segmented, rinsed, vortexed and dilution plate method 10⁻⁴ were brought off. Then 0.1 ml solution directly placed on sterilized TZC [23] plates, incubated at growth chamber. 2, 3, 5-triphenyl tetrazolium chloride (TZC) selected media (1 L) contained casamino acids (1.0 g), peptone (10 g), glucose (5.0 g) and Agar (15 g). In TZC media 5 ml of 1% 2, 3, 5-triphenyltetrazolium chloride was added to sterilized medium before pouring in to the plate [24]. The medium was cooled at 50°C - 55°C, poured into the petri plates and stored at 4°C. The media was prepared following [14] [15] [22] [25] and was used to streak the bacterial suspension by using sterilized wire

loop. Inoculated plates were incubated at 28°C - 30°C for 2 - 3 days in growth chamber. Plates should be incubated in an inverted position because water condensation causes colonies to flow into each other, thereby limiting separation. Separately growing colonies can then be picked and sub-cultured into fresh media to obtain pure cultures collected.

2.4. Designation of Collected Isolates

The collected isolates were designated based on their location and source of collection [26]. For example, an isolate designated by PMMuTa₅CS₀₁ represents that this isolate was collected from potato (P), district Munshiganj (M), Upazila Munshiganj Sadar (Mu). Ta CS stands for Tamanna Cold Storage from where potato samples were collected. The sample collection number five is represented as Ta₅. CS₀₁ represents cold storage isolate number 01. The details of the collected isolates were as follows: PMMuTa₅CS₀₁ = Potato tuber, Munshiganj, Munshiganj Sadar, Tamanna Cold Storage, Sample No.01; PMMuFa₉ S₀₆ = Potato tuber, Munshiganj, Munshiganj Sadar, Farmer Storage, Sample No. 06; PMSiFi₃CS₀₈ = Potato tuber, Munshiganj, Sirajdikhan, Five Star Cold Storage, Sample No. 08; $PMSiFa_2S_{12} = Potato tuber$, Munshiganj, Sirajdikhan, Farmer Storage, Sample No. 12; PMTTo₅CS₁₆ = Potato tuber, Munshiganj, Tongibari, Tongibari Cold Storage, Sample No. 16; PMTFa $_{35}S_{21}$ = Potato tuber, Munshiganj, Tongibari, Farmer Storage, Sample No. 21; $PN_{ij}DSh_{43}CS_{24} = Potato tuber, Nilphamari, Do$ mar, Shawon Cold Storage, Sample No. 24; PN_{il}DFa₇S₂₈ = Potato tuber, Nilphamari, Domar, Farmer Storage, Sample No. 28; PN_{il}NiAn₁₉CS₂₉ = Potato tuber, Nilphamari, Nilphamari Sadar, Ankur Cold Storage, Sample No. 29; PN_{il}NiFa₂S₃₂ = Potato tuber, Nilphamari, Nilphamari Sadar, Farmer Storage, Sample No. 32; PN_{il}DiMk₄CS₃₃ = Potato tuber, Nilphamari, Dimla, Mukta Cold Storage, Sample No. 33; PN_{il}DiFa₂₄CS₃₆ = Potato tuber, Nilphamari, Dimla, Farmer Storage, Sample No. 36; PNNaHa₂CS₃₈ = Potato tuber, Narayangonj, Narayangonj Sadar, Hazi Rahamotolla Cold Storage, Sample No. 36; $PNNaFa_1S_{43} = Potato$ tuber, Narayangonj, Narayangonj Sadar, Farmer Storage, Sample No. 43; PNSSa₈CS₄₅ = Potato tuber, Narayangonj, Sonargaon, Sahin Cold Storage, Sample No. 45; PNSFa₂₀S₄₈ = Potato tuber, Narayangonj, Sonargaon, Farmer Storage, Sample No. 48; PNRuMa₃CS₅₀ = Potato tuber, Narayangonj, Rupgonj, Ma Cold Storage, Sample No. 50; PNRuFa₄S₅₄ = Potato tuber, Narayangonj, Rupgonj, Farmer Storage, Sample No. 54; P₁MMuF₆S₀₁ = Potato plant, Munshiganj, Munshiganj Sadar, Field, Sample No. 01; P₁MSiF₃₃S₀₄ = Potato plant, Munshiganj, Sirajdikhan, Field, Sample No. 04; P₁MTF₉S₀₇ = Potato plant, Munshiganj, Tongibari, Field, Sample No. 07; P₁N_{il}DF₁₁S₁₀ = Potato plant, Nilphamari, Domar, Field, Sample No. 10; $P_1N_{i1}NiF_{19}S_{13}$ = Potato plant, Nilphamari, Nilphamari Sadar, Field Sample No. 13; P₁N_{il}DiF₇S₁₆ = Potato plant, Nilphamari, Dimla, Field, Sample No. 16; $P_1NNaF_3S_{19}$ = Potato plant, Narayangonj, Narayangonj Sadar, Field, Sample No. 19; P₁NSF₄S_{22 =} Potato plant, Narayangonj, Sonargaon, Field, Sample No. 22; $P_1NRuF_3S_{25}$ = Potato plant, Narayangonj, Rupgonj, Field, Sample No. 25; CNRF₁₄S₀₁ = Chilli, Narayangonj, Rupgonj, Field, Sample No. 01; SoMMuF₁S₀₁ =

Soil, Munshiganj, Munshiganj Sadar, Field, Sample No. 01; SoMSiF₁S₀₄ = Soil, Munshiganj, Sirajdikhan, Field, Sample No. 04; SoMToF₁S₀₇ = Soil, Munshiganj, Tongibari, Field, Sample No. 07; SoN_{il}DF₁S₁₁ = Soil, Nilphamari, Domar, Field, Sample No. 11; SoN_{il}NiF₁S₁₄ = Soil, Nilphamari, Nilphamari Sadar, Field, Sample No. 14; SoN_{il}DiF₁S₁₇ = Soil, Nilphamari, Dimla, Field, Sample No. 17; SoN-NaF₁S₂₀ = Soil, Narayangonj, Narayangonj Sadar, Field, Sample No. 20; SoNSF₁S₂₃ = Soil, Narayangonj, Sonargaon, Field, Sample No. 23; SoNRuF₁S₂₆ = Soil, Narayangonj, Rupgonj, Field, Sample No. 26; WMSiF₁₃ S₀₁ = Weed, Munshigonj, Sirajdikhan, Field, Sample No. 01; WN_{il}NiF₁₆ S₀₃ = Weed, Nilphamari, Nilphamari Sadar, Field, Sample No. 03; WNRuFa₃₆ S₀₅ = Weed, Narayangonj, Rupgonj, Field, Sample No. 05; BNRF₂S₀₁ = Banana, Narayangonj, Rupgonj, Field, Sample No. 01; WaMSiF₁S₀₁ = Water, Munshigonj, Sirajdikhan, Field, Sample No. 01; WaN_{il}NiF₁₁S₀₄ = Water, Nilphamari, Nilphamari Sadar, Field, Sample No. 04; WaNRuF₁S₀₇ = Water, Narayangonj, Rupgonj, Field, Sample No. 04; WaNRuF₁S₀₇ = Water, Narayangonj, Rupgonj, Field, Sample No. 05; ONR

2.5. Hypersensitive Reaction

Hypersensitive reaction was performed on the leaf of *Nicotiana tabacum* [27] [28]. At 30 - 40 days aged tobacco plants were selected for inoculation. Single colony of *R. solanacearum* showing virulent, fluidal, irregular and creamy white with pink at the centre was selected for each isolate and multiplied in a CPG (without adding triphenyl tetrazolium chloride) medium for 48h prior to inoculation. Bacterial suspensions were prepared in distilled sterilized water and 10^8 cfu/mL, cell density was adjusted through benchtop spectrophotometer at 620 nm. 1ml plastic syringe with 25 mm gauge needles was used to inject cell suspension into the abaxial side of the mature tobacco leaves. Alternating interveinal regions were infiltrated with bacterial suspension. Each injected area was labelled with appropriate letters. A fine mist of water was lightly sprayed over the whole plants and placed at room temperature ($25^{\circ}C - 27^{\circ}C$). Plants were covered with clear plastic bags. Data were recorded 1, 3 and 7 days after inoculations [28].

2.6. Biochemical Test

Hypersensative reaction positive isolates were studied according to specific six biochemical tests for *Ralstonia solanacearum*.

2.6.1. Gram Staining

Young pure cultures (24 hrs) were used for gram-staining. A loop of the bacterium was spreaded on a glass slide and fixed by heating on a very low flame. Aqueous crystal violet solution (0.5%) was spread over the smear for 30 seconds and then washed with running tape water for one minute. It was then flooded with iodine for one minute, rinsed in tape water and decolorized with 95% ethanol until colourless runoff. After washing the specimen was counter-stained with safranin for approximately 10 seconds, washed with water, dried and observed microscopically at 10X, 40X and 100X using oil imersion [29].

2.6.2. Potassium Hydroxide (KOH) Test

Isolates of *R. solanacearum* were aseptically taken from 24 - 36 hrs old culture plates with a sterile tooth pick. It was then placed on glass slide in a drop of 3% KOH solution, stirred for 10 seconds and observed for the formation of slime threads [30].

2.6.3. Catalase Oxidase Test

Catalase test was performed with isolates to check their liveliness. Young agar cultures (18 - 24 hrs) and 3% hydrogen peroxide (H_2O_2) were used to observe production of gas bubbles. A loop full of bacterial culture was mixed with a drop of H_2O_2 on a glass slide and observed for the production of gas bubbles with naked eye and under a dissecting magnification of 25X [29].

2.6.4. Kovac's Oxidase Test

Oxidase reagent (1% tetramethyl-p-phenyl diamine dihydrochloride) solution (100 ml) was prepared and kept in rubber–stopper dark bottle. A drop of reagent was added to a piece of filter paper (whatman no. 1) placed within a glass petri dish. Small quantities of the inoculum were rubbed on the filter paper impregnated with 1% (w/v) oxidase reagent solution. Bacteria were then noted for the development of purple colour in 10 - 60 seconds indicating the oxidation of the reagent [25].

2.6.5. Pectolytic Test

Potato tubers were disinfected with 99% ethanol, cut up into slices of about 7 - 8 mm thick, and then placed on moistened sterile filter paper in sterile petri dishes. Bacterial cell suspension was pipetted into a depression cut in the potato slices. One potato slice pipetted with sterile water was treated as control. Development of rot on the slices was examined 24 - 48 h after incubation at 25°C. Examination was done 5 days after inoculation. Two slices were inoculated for each isolate [25] [31] [32].

2.6.6. Temperature Sensitivity Test

The ability of bacterial isolates to grow at different selected temperature. Bacterial suspensions from different isolates were prepared from 24 hrs TZC cultures and 2 - 3 drops of suspension were plated on TZC medium which was placed at $4^{\circ}C \pm 2^{\circ}C$, $28^{\circ}C \pm 2^{\circ}C$, $37^{\circ}C \pm 2^{\circ}C$ and $40^{\circ}C \pm 2^{\circ}C$ temperatures [25].

2.7. PCR for Molecular Confirmation of R. solanacearum Isolates

The isolates of *R. solanacearum* collected from Munshingonj, Narayngonj and Nilphamari districts were tested for molecular detection. Three isolates from each district were selected and confirmed by polymerase chain reaction (PCR) technique using 1 kb DNA ladder (Promega, USA, dispensed Lot # 0000339931) on agarose gel. A pair of universal species-specific primer pair AU759f (5'-GTC

GCC GTC AAC TCA CTT TCC-3') and AU760r (5'-GTC GCC GTC AGC AAT GCG GAA TCG-3') were used [33]. A well separated bacterial colony on TZC medium were cultured on CPG broth (Peptone 10 g; glucose, 5 g; Casamino acid, 1 g; distilled water 1 l; pH 7.0 - 7.1) and incubated at 28°C for 48 - 72 h with vigorous shaking prior to the DNA extraction process in a shaking incubator. Genomic DNA extracted by Gene JET Genomic DNA Purification Kit from Thermo Fisher Scientific (www.thermoscientific.com/onebio). PCR amplification was carried out in a total volume of 25 µl using 12 µl of PCR Master Mix (Thermo fisher), 1 μ l of forward primer, 1 μ l of rivers primer, 5 μ l of genomic DNA and 6 µl Milli Q water. The following cycling program was used in a thermal cycler: an initial denaturation step at 95°C for 5 mins, followed by 35 cycles of 95°C for 30 s, 57°C for 45 s and 72°C for 45 s, followed by a final extension step of 72°C for 5 mins. After completion of cycling program, reaction was held at 4°C. Bacterial suspensions of *R. solanacearum* at approximately 10° CFU/ml and sterile water were used as the positive and negative control, respectively. PCR product of expected 280 bp was visualized under UV light after electrophoresis in 1.5% agarose and staining in ethidium bromide solution (1 µg·ml⁻¹) and photographed.

2.8. Race Determination

The races of *R. solanacearum* were determined by pathogenicity test on wide host range [34] [35]. Soil was prepared by integration of farmyard manure, sand and clay in 1:1:1 ratio. This mixture was sterilized by 37% formalin in 1:9 ratio (1 part formalin and 9 parts soil). Soil mixture was covered with polyethylene sheet and kept for 4 days. After that the polyethylene sheet was removed and treated soil was exposed in air for additional two days. Plastic tray was filled by sterilized soil. Potato (Solanum tuberosum), tomato (Solanum lycopersicum) and brinjal (Solanum melongena) seeds were surface sterilized, rinsed with sterile water and sown in separate tray. Seedlings bearing 4/5 leaves were used for performing pathogenicity tests through artificial inoculation and race assinged [36]. The seedlings were inoculated with R. solanacearum by wounded root dippingin bacterial suspension for 30 mins [25] [28] [37] [38]. Plants were not watered a day before inoculation to reduce moisture in the pots. Roots of potato, tomato and brinjal seedlings were slightly injured by inserting a scalpel in the pots in order to facilitate bacterial infection. About 10 ml of bacterial suspension (10⁸ cfu/mL) of each isolate was used for the study. Inoculated plants were regularly watered and kept at temperature range from 28°C - 30°C and 90% RH [39]. The inoculated plants were then kept in observation until the symptoms were developed.

2.9. Biovar Test

Biovars of *R. solanacearum* was determined on the basis of carbon utilization in disaccharides and hexose alcohols [7] [35]. The disaccharides were cellobiose,

lactose, and maltose while the hexose alcohols were dulcitol, mannitol and sorbitol. Sterilized solutions (10% W/V) of these carbon sources were added to a basal medium (Table 1) before inoculation of *R. solanacearum* isolates.

The medium was brought to boil with constant stirring and raised the pH of the medium to 7.0 - 7.1 by drop wise addition of 1.0N sodium hydroxide. When the medium turned green, the medium was then autoclaved at 121°C, 15 psi for 20 mins.

Cellobiose, lactose, maltose, dulcitol, mannitol and sorbitol were taken 1 g from each disaccharide and hexose alcohols were dissolved in 10 ml distilled water separately. Solutions were filter-sterilized using 0.22 μ m filters.

After autoclaving, the basal medium was cooled at 65°C. 10 ml each of disaccharides and hexose alcohols solutions were mixed with 90 ml basal medium. For the control, 10 ml sterile water was mixed (without disaccharides and hexose alcohols) with 90 ml basal medium. The media containing 1% of each disaccharide and hexose alcohols was prepared to determine the biovars of isolates of *R. solanacearum* [40]. After melting the medium, it was then dispensed into the wells of microtiter plate.

Isolates of *R. solanacearum* were grown on CPG solid media to test the biovar expression on the basis of carbon utilization in disaccharides and hexose alcohols which were incubated in 28°C temperature [7] [15] [22] [35]. CPG media compounds were tabulated in Table 2.

The ingredients were autoclaved at 121°C for 15 mins and cool to about 40°C - 45°C before pouring. The homogenized medium was then poured into sterile petri dishes and let solidify for 30 mins. Isolates of *R. solanacearum* were then taken from pure culture to dilute @10 microliters/ml and spread plated and incubated at optimum temperature. Inocula for each isolate were prepared by adding a loopful of the bacteria from 24 - 48 h old cultures to 1ml sterile in a 2.0 ml centrifuge tube. One small drop of cloudy bacterial suspension was added in each well of a column. After inoculation microtiter plate incubated at 28°C. Reactions were observed and recorded after seven days of inoculation. A colour change to yellow (acid pH < 6) indicates oxidation of the carbon source [7] [41]. This usually occurs within 3 - 5 days; those biovars capable of oxidizing disaccharides take a few days longer to give a clear positive result. The inoculated tubes compared with a non-inoculated control tube.

3. Results

It was observed that fifty-five tuber samples showed positive in oozing and browning in vascular bundle (Table 3). In streaming test, it was observed that twenty-seven potato plant samples showed positive bacterial ooze streaming in clear water but three chilli samples showed negetive in streaming test (Table 3). Selected weed, soil, banana and water samples did not show any streaming and oozing test. Twenty-eight soil samples, five weed samples, three banana samples and nine water samples were also used to isolation of *Ralstonia solanacearum* (Table 3).

Components	Per 1000 ml distilled water liter						
Ammonium dihydrogen phosphate (NH ₄ H ₂ PO ₄)	1.0 g						
Potassium chloride (KCl)	0.2 g						
Magnesium sulphate (MgSO ₄ ·7 H_2 O)	0.2 g						
Peptone	1.0 g						
Bromothymol blue	0.03 g						
Agar	3.0 g						
Water	1.0 L						

Table 1. Components for preparing basal medium.

Table 2. Component of CPG medium.

Components	Per 1000 ml distilled water liter ⁻¹
Casamino acids (Difco)	1.0 g
Bacto-Peptone (Difco)	10.0 g
Glucose	10.0 g
Bacto-Agar (Difco)	18.0 g

Table 3. Host and sources of collected isolates of *R. solanacearum* from Bangladesh and their reaction response on TZC, virulent colony (VC) and hypersensitivity (HR) test.

Isolates	source	Host	TZC	vc	HR test	
PMMuTa ₅ CS ₀₁			+ve	+ve	+ve	
PMMuTa ₁₁ CS ₀₂			+ve	+ve	-ve	
PMMuTa ₁₇ CS ₀₃	Munshiganj, Munshiganj Sadar (Tamanna Cold Storage)	Solanum tuberosum	+ve	-ve	-	
PMMuTa ₁₉ CS ₀₄	(Tullullu Cold Storage)		+ve	+ve	+ve	
PMMuTa ₂₃ CS ₀₅			-ve	-ve	-	
PMMuFa ₉ S ₀₆	Munshiganj, Munshiganj Sadar		+ve	-ve	-	
PMMuFa ₁₅ S ₀₇	(Farmer Storage)	Solanum tuberosum	+ve	+ve	+ve	
PMSiFi ₃ CS ₀₈			+ve	+ve	+ve	
PMSiFi ₁₁ CS ₀₉	Munshiganj, Sirajdikhan		+ve	+ve	-ve	
PMSiFi ₂₁ CS ₁₀	(Five Star Cold Storage)	Solanum tuberosum	+ve	+ve	+ve	
PMSiFi ₂₄ CS ₁₁			+ve	-ve	-	
PMSiFa ₂ S ₁₂			+ve	+ve	+ve	
PMSiFa ₁₂ S ₁₃	Munshiganj, Sirajdikhan		+ve	-ve	-	
PMSiFa ₂₂ S ₁₄	(Farmer Storage)	Solanum tuberosum	+ve	-ve	-	
PMSiFa ₂₈ S ₁₅			+ve	+ve	+ve	
PMTTo ₅ CS ₁₆			+ve	+ve	-ve	
PMTTo7CS17			+ve	-ve	-	
PMTTo ₁₅ CS ₁₈	Munshiganj, Tongibari (Tongibari Cold Storage)	Solanum tuberosum	+ve	+ve	+ve	
PMTTo ₁₈ CS ₁₉	(Tongibari Cold Storage)		+ve	-ve	-	
PMTTo ₂₀ CS ₂₀			+ve	+ve	+ve	

Continued					
PMTFa ₃₅ S ₂₁			+ve	+ve	+ve
PMTFa ₃₆ S ₂₂	Munshiganj, Tongibari (Farmer)	Solanum tuberosum	+ve	-ve	-
PMTFa ₄₁ S ₂₃			+ve	+ve	+ve
PN _{il} DSh ₄₃ CS ₂₄			-ve	-ve	-
PN _{il} DSh ₄₆ CS ₂₅	Nilphamari, Domar		+ve	+ve	+ve
PN _{il} DSh ₄₇ CS ₂₆	(Shawon Cold Storage)	Solanum tuberosum	+ve	+ve	-ve
PN _{il} DSh ₅₀ CS ₂₇			+ve	+ve	+ve
PN _{il} DFa ₇ S ₂₈	Nilphamari, Domar (Farmer)	Solanum tuberosum	+ve	+ve	-ve
PN _{il} NiAn ₁₉ CS ₂₉			+ve	-ve	_
PN _{il} NiAn ₃₁ CS ₃₀	Nilphamari, Nilphamari Sadar	Solanum tuberosum	+ve	+ve	+ve
PN _{il} NiAn ₃₁ CS ₃₁	(Ankur Cold Storage)		+ve	-ve	_
PN _{il} NiFa ₂ S ₃₂	Nilphamari, Nilphamary Sadar (Farmer Storage)	Solanum tuberosum	+ve	+ve	-ve
PN _{il} DiMk ₄ CS ₃₃			+ve	+ve	+ve
PN _{il} DiMk ₁₄ CS ₃₄	Nilphamari, Dimla (Mukta Cold Storage)	Solanum tuberosum	+ve	-ve	-
PN _{il} DiMk ₁₇ CS ₃₅	Storage)		+ve	+ve	+ve
PN _{il} DiFa ₂₄ CS ₃₆	Nilphamari, Dimla		+ve	+ve	-ve
PN _{il} DiFa ₂₉ CS ₃₇	(Farmer Storage)	Solanum tuberosum	+ve	+ve	+ve
PNNaHa ₂ CS ₃₈			+ve	+ve	+ve
PNNaHa ₁₃ CS ₃₉			-ve	-ve	-
PNNaHa ₁₅ CS ₄₀	Narayangonj, Narayangonj Sadar (Hazi Rahamotolla Cold Storage)	Solanum tuberosum	+ve	+ve	+ve
PNNaHa ₁₆ CS ₄₁	(Trazi Kanamotona Cold Storage)		+ve	+ve	-ve
PNNaHa ₂₁ CS ₄₂			+ve	-ve	-
PNNaFa ₁ S ₄₃	Narayangonj, Narayangonj Sadar		+ve	+ve	-ve
PNNaFa ₁₁ S ₄₄	(Farmer Storage)	Solanum tuberosum	+ve	+ve	+ve
PNSSa ₈ CS ₄₅			+ve	+ve	-ve
PNSSa ₉ CS ₄₆	Narayangonj, Sonargaon	Solanum tuberosum	+ve	+ve	+ve
PNSSa ₁₅ CS ₄₇	(Sahin Cold Storage)		+ve	-ve	_
PNSFa ₂₀ S ₄₈	Narayangonj, Sonargaon		+ve	+ve	+ve
PNSFa ₂₅ S ₄₉	(Farmer Storage)	Solanum tuberosum	+ve	+ve	-ve
PNRuMa ₃ CS ₅₀			+ve	+ve	+ve
PNRuMa7CS51	Narayangonj, Rupgonj		+ve	+ve	+ve
PNRuMa ₁₃ CS ₅₂	(Ma Cold Storage)	Solanum tuberosum	+ve	+ve	-ve
PNRuMa ₁₉ CS ₅₃			+ve	+ve	+ve
PNRuFa ₄ S ₅₄	Narayangonj, Rupgonj		+ve	+ve	-ve
PNRuFa ₄ S ₅₅	(Farmer Storage)	Solanum tuberosum	-ve	-ve	_
$P_1MMuF_6S_{01}$			+ve	+ve	+ve
$P_lMMuF_{14}S_{02} \\$	Munshiganj, Munshiganj Sadar (Field)	Solanum tuberosum	+ve	+ve	+ve
	(Field)				

Continued					
P ₁ MSiF ₃₃ S ₀₄			+ve	+ve	+ve
$P_{l}MSiF_{35}S_{05}$	Munshiganj, Sirajdikhan (Field)	Solanum tuberosum	+ve	+ve	+ve
$P_1MSiF_{39}S_{06}$			+ve	+ve	-ve
P ₁ MTF ₉ S ₀₇			-ve	-ve	-ve
$P_l MTF_{20}S_{08}$	Munshiganj, Tongibari (Field)	Solanum tuberosum	+ve	+ve	+ve
P1MTF29S09			+ve	+ve	-ve
$P_l N_{il} DF_{11} S_{10}$			+ve	+ve	+ve
$P_1N_{il}DF_{21}S_{11}$	Nilphamari, Domar (Field)	Solanum tuberosum	-ve	-ve	-
$P_l N_{il} DF_{23} S_{12}$			+ve	+ve	+ve
P _l N _{il} NiF ₁₉ S ₁₃			+ve	+ve	+ve
$P_l N_{il} Ni F_{21} S_{14}$	Nilphamari, Nilphamari Sadar (Field)	Solanum tuberosum	-ve	-ve	-
$P_l N_{il} Ni F_{31} S_{15} \\$	(Tett)		+ve	+ve	+ve
$P_1N_{il}DiF_7S_{16}$			+ve	+ve	-ve
$P_l N_{il} Di F_{17} S_{17}$	Nilphamari, Dimla (Field)	Solanum tuberosum	-ve	-ve	-
$P_l N_{il} Di F_{25} S_{18}$			+ve	+ve	+ve
P ₁ NNaF ₃ S ₁₉			+ve	+ve	+ve
$P_1NNaF_{13}S_{20}$	Narayangonj, Narayangonj Sadar (Field)	Solanum tuberosum	-ve	-ve	-
$P_1NNaF_{30}S_{21}$	(Fleid)		+ve	+ve	+ve
P ₁ NSF ₄ S ₂₂			+ve	+ve	+ve
$P_1NSF_{14}S_{23}$	Narayangonj, Sonargaon (Field)	Solanum tuberosum	-ve	-ve	-
$P_1NSF_{25}S_{24}$			+ve	+ve	+ve
P ₁ NRuF ₃ S ₂₅			+ve	+ve	+ve
$P_1NRuF_{13}S_{26}$	Narayangonj, Rupgonj (Field)	Solanum tuberosum	-ve	-ve	-
$P_1NRuF_{22}S_{27}$			+ve	+ve	+ve
CNRF ₁₄ S ₀₁			-ve	-ve	-
CNRF ₂₁ S ₀₂	Narayangonj, Rupgonj (Field)	Capsicum annum	-ve	-ve	-
CNRF ₄₁ S ₀₃			-ve	-ve	-
SoMMuF ₁ S ₀₁			+ve	+ve	+ve
SoM MuF ₂ S ₀₂	Munshiganj, Munshiganj Sadar (Field)	Soil	+ve	+ve	+ve
SoM MuF ₃ S ₀₃	(Field)		+ve	-ve	-
SoMSiF ₁ S ₀₄			+ve	+ve	+ve
SoMSiF ₂ S ₀₅	Munshiganj, Sirajdikhan (Field)	Soil	+ve	+ve	+ve
SoMSiF ₃ S ₀₆			+ve	-ve	-
SoMToF ₁ S ₀₇			+ve	+ve	+ve
SoMToF ₂ S ₀₈			-ve	-ve	-
SoMToF ₂₁ S ₀₉	Munshiganj, Tongibari (Field)	Soil	-ve	-ve	-
SoMToF ₃₀ S ₁₀			+ve	+ve	+ve

SoN _{il} DF ₁ S ₁₁			+ve	+ve	_
$SoN_{il}DF_2S_{12}$	Nilphamari, Domar (Field)	Soil	+ve	+ve	
$SoN_{il}DF_3S_{13}$			+ve	+ve	
SoN _{il} NiF ₁ S ₁₄			+ve	+ve	
$SoN_{il}NiF_2S_{15}$	Nilphamari, Nilphamari Sadar	Soil	-ve	-ve	
SoN _{il} NiF ₃ S ₁₆	(Field)	001	+ve	+ve	+
SoN _{il} DiF ₁ S ₁₇			+ve	+ve	
$SoN_{il}DiF_2S_{18}$	Nilphamari, Dimla (Field)	Soil	+ve	+ve	
$SoN_{il}DiF_{2}S_{18}$ $SoN_{il}DiF_{3}S_{19}$	Niphaniari, Dinna (Field)	501	-ve	-ve	'
SoNNaF ₁ S ₂₀			-ve +ve		
	Narayangonj, Narayangonj Sadar	Soil		+ve	
SoNNaF ₂ S ₂₁	(Field)	3011	+ve	+ve	
SoNNaF ₃ S ₂₂			+ve	+ve	
SoNSF ₁ S ₂₃			+ve	+ve	
SoNSF ₂ S ₂₄	Narayangonj, Sonargaon (Field)	Soil	+ve	+ve	
SoNSF ₃ S ₂₅			+ve	+ve	+
SoNRuF ₁ S ₂₆			-ve	-ve	
SoNRuF ₂ S ₂₇	Narayangonj, Rupgonj (Field)	Soil	+ve	+ve	+
SoNRuF ₃ S ₂₈			+ve	+ve	-
WMSiF ₁₃ S ₀₁	Munshigonj, Sirajdikhan (Field)		+ve	+ve	-
WMSiF ₃₃ S ₀₂			+ve	+ve	-
WN _{il} NiF ₁₆ S ₀₃	Nilphamari, Nilphamari Sadar	Cynodon dactylon	-ve	-ve	
WN _{il} NiF ₂₃ S ₀₄	(Field)		-ve	-ve	
WNRuF ₃₆ S ₀₅	Narayangonj, Rupgonj (Field)		+ve	+ve	_
$BNRF_2S_{01}$			+ve	+ve	-
$BNRF_{13}S_{02}$	Narayangonj, Rupgonj (Field)	M <i>usa spp</i>	-ve	-ve	
BNRF ₂₂ S ₀₃			+ve	+ve	-
$WaMSiF_1S_{01}$			-ve	-ve	
WaMSiF ₁₂ S ₀₂	Munshigonj, Sirajdikhan (Irrigation Water-Field)	Water	+ve	+ve	+
WaMSiF ₃₀ S ₀₃	,		+ve	+ve	_
WaN _{il} NiF ₁₁ S ₀₄			+ve	+ve	_
WaRNiF ₂₃ S ₀₅	Nilphamari, Nilphamery Sadar (Irrigation Water-Field)	Water	+ve	+ve	_
WaRNiF ₃₄ S ₀₆	(-ve	-ve	
WaNRuF ₁ S ₀₇			-ve	-ve	
WaNRuF ₂ S ₀₈	Narayangonj, Rupgonj (Irrigation Water-Field)	Water	+ve	+ve	_
WaNRuF ₃ S ₀₉	water-rieid)		+ve	+ve	

HR = hypersensitive reaction; VC = virulent colony; +ve = positive reaction; -ve = negative reaction; - = not applicable in HR test.

Finally, one hundred and thirty isolates of *R. solanacearum* were isolated from potato tuber, potato stems, chili stems, banana leaves, weed, soil and irrigation water. Above them one hundred and six isolates were positive on TZC solid medium. Fluidal pinkish red centered colonies, typical of *R. solanacearum* were observed on TZC solid medium. Among them ninety isolates were virulent and sixteen isolates were avirulent (**Table 3** and **Plate 1**). All virulent isolates were

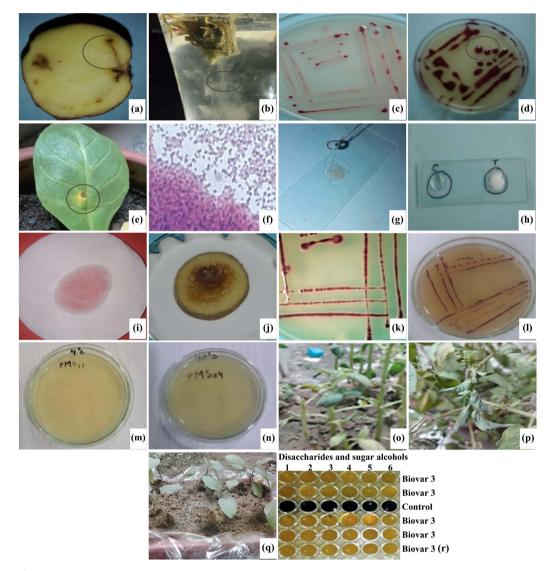


Plate 1. (a) Brown discoloration of vascular tissue; (b) Bacterial ooze from cut stem; (c) Virulent colonies on TZC medium; (d) Avirulent colonies on TZC medium (uniformly round and dark red); (e) Death cell showed on *R. solanacearum* inoculated leaf of tobacco due to hypersensitive reaction; (f) Gram staining test; (g) Catalase test; (h) KOH test; (i) Kovacs oxidase test; (j) Pectolytic test; (k) *R. solanacearum* at 28° C (Typical growth); (l) *R. solanacearum* at 37° C (Typical growth); (m) No potential growth of *R. solanacearum* at 4° C; (n) No potential growth of *R. solanacearum* at 40° C; (o) Pathogenicity test positive on potato seedlings after artificial inoculation of *R. solanacearum*; (q) Pathogenicity test negetive on brinjal seedlings after artificial inoculation of *R. solanacearum* and (r) biovar (s); (s) test of *R. solanacearum* showing a positive result (yellow colour) and a negative result (green colour) (1 = Cellobiose: 2 = Lactose; 3 = Maltose; 4 = Dulcitol; 5 = Mannitol; 6 = Sorbitol).

tested for hypersensitive response on tobacco plants (**Table 3** and **Plate 1**). Depending upon the appearance of hypersensitive reaction, fifty-seven isolates producing symptoms within 2 - 5 days after bacterial inoculation on tobacco leaves. On the basis of all tests performed which were positive for hypersensative reaction. Fifty isolates out of fifty-seven isolates showed positive reaction in gram staining test, KOH test, catalase test, kovac's oxidase test and pectolytic test (**Table 5** and **Plate 1**). In temperature sensitivity test, the test isolates showed positive and typical growth at 28°C and 37°C but did not show any potential growth at 4°C and 40°C (**Table 5** and **Plate 1**).

Among fifty isolates, nine isolates selected from three districts produced the bands of 280 bp size (**Table 4** and **Figure 1**) specific for *R. solanacearum* with species specific primers which mean that, those isolate(s) were belonged to *R. solanacearum*.

In race determination test fifty isolates showed wilting symptoms on potato and tomato seedlings except brinjal seedlings (**Table 5** and **Plate 1**). Fifty isolates were expressed as race 3. Forty-eight isolates were expressed as biovar III after 7 - 8 days and the remaining two isolates from soil and water were expressed as biovar I on the basis of utilization of different sugars (**Table 5** and **Plate 1**).

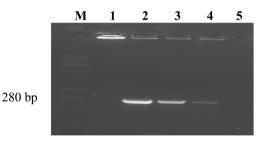


Figure 1. Gel electrophoresis of amplified 280 bp DNA fragments from *R. solanacearum* with species specific primers. M = 100 bp DNA ladder, 1 = negative control, 2 to 4 = representative three strains of *R. solanacearum* from three districts and 5 = blank control.

Isolates	Locations of sampling	Districts	Source of infection	Band size	Inference
PM MuTa ₅ CS ₀₁	Munshiganj Sadar (Tamanna Cold Storage)		Solanum tuberosum	280 bp	R. solanacearum
PM MuFa ₁₅ S ₀₇	Munshiganj Sadar (Farmer Storage)	Munshigonj	Solanum tuberosum	280 bp	R. solanacearum
PlMSiF ₃₃ S ₀₄	Sirajdikhan (Field)		Solanum tuberosum	280 bp	R. solanacearum
PNSSa ₉ CS ₄₆	Sonargaon (Sahin Cold Storage)		Solanum tuberosum	280 bp	R. solanacearum
PNSFa ₂₀ S ₄₈	Sonargaon (Farmer Storage)	Narayangonj	Solanum tuberosum	280 bp	R. solanacearum
$P_1NNaF_3S_{19}$	Narayangonj sadar (Field)		Solanum tuberosum	280 bp	R. solanacearum
PN _{il} DSh ₄₆ CS ₂₅	Domar (Shawon Cold Storage)		Solanum tuberosum	280 bp	R. solanacearum
$P_l N_{il} Di Fa_{25} S_{18} \\$	Dimla (Farmer)	Nilphamari	Solanum tuberosum	280 bp	R. solanacearum
$P_l N_{il} DF_{11} S_{10} \\$	Domar (Field)		Solanum tuberosum	280 bp	R. solanacearum

	gu		ų	÷			l'empo ensitiv							Н	exose alcoh	ols	I	Disaccarid	es		
Isolates	Gram Staining	KOH test	Catalase oxidase test	Kovacs oxidase test	Pectolytic test	4°C	28°C	37°C	40°C	Potato	Tomato	Brinjal	Race	Dulcitol	Sorbitol	Mannitol	Cellobiose	Maltose	Lactose	Biovar	Inference
PMuTa ₅ CS ₀₁	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMMuTa ₁₉ CS ₀₄	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMMuFa ₁₅ S ₀₇	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMSiFi ₃ CS ₀₈	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMSiFi ₂₁ CS ₁₀	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMSiFa ₂ S ₁₂	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$PMSiFa_{28}S_{15}$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMTTo ₁₅ CS ₁₈	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMTTo ₂₀ CS ₂₀	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMTFa ₃₅ S ₂₁	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PMTFa ₄₁ S ₂₃	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$PN_{il}DSh_{46}CS_{25} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PN _{il} DSh ₅₀ CS ₂₇	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PN _{il} NiAn ₃₁ CS ₃₀	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PN _{il} DiMk ₄ CS ₃₃	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PN _{il} DiMk ₁₇ CS ₃₅	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PN _{il} DiFa ₂₉ CS ₃₇	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PNNaHa ₂ CS ₃₈	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PNNaHa ₁₅ CS ₄₀	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PNNaFa ₁₁ S ₄₄	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PNSSa ₉ CS ₄₆	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$PNSFa_{20}S_{48}$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PNRuMa ₃ CS ₅₀	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PNRuMa ₇ CS ₅₁	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
PNRuMa ₁₉ CS ₅₃	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_{l}MMuF_{6}S_{01} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_{l}MMuF_{14}S_{02} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_1MSiF_{33}S_{04}$	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_lMSiF_{35}S_{05}$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_l MTF_9 S_{07}$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_l MTF_{20}S_{08}$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_l N_{il} DF_{11} S_{10} \\$	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_l N_{il} DF_{23} S_{12} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum

 Table 5. Result of different biochemical test, race determinationand biovar test for characterization of different isolates of *R. sola-nacearum* collected from different locations of Bangladesh.

Con	tin	116	А
COIL	CTTT	uc	u

$P_l N_{il} Ni An_{19} S_{13} \\$	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_l N_{il} Ni An_{31} S_{15} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_lN_{il}DiFa_{25}S_{18} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_1NNaF_3S_{19}$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_1NNaF_{30}S_{21} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_1 NSF_4 S_{22} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_l NSF_{25}S_{24}$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_1NRuF_3S_{25}\\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$P_l NRuF_{22}S_{27}$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
SoMSiF ₁ S ₀₄	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$SoMSiF_2S_{05}$	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$SoN_{il}NiF_1S_{14} \\$	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$SoN_{il}NiF_3S_{16} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$SoN_{il}DiF_2S_{18} \\$	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	ve	ve	ve	ve	ve	ve	1*	R. solanacearum
$SoNNaF_1S_{20}$	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
SoNRuF ₂ S ₂₇	-ve	+ve	+ve	+ve	+++ve	-ve	+ve	+ve	-ve	+++ve	+++ve	-ve	3**	++++ve	++++ve	++++ve	++++ve	++++ve	++++ve	3*	R. solanacearum
$WaMSiF_{12}S_{02} \\$	-ve	+ve	+ve	+ve	++ve	-ve	+ve	+ve	-ve	++ve	+++ve	-ve	3**	ve	ve	ve	ve	ve	ve	1*	R. solanacearum

"+ve" means positive reaction; "-ve" means negative reaction; "+++ve" means strongly positive reaction; and "++ve" means weakly positive reaction; "++++ve" means positive reaction (Yellow color produced due to change in pH); "--ve" means negative reaction (Green color remained as pH did not change); 3** = Race3; 3* = Biovar III; 1* = Biovar 1.

4. Discussion

Samples were collected from different sources viz. potato tuber, potato stem, chili stem, banana leaves, soil, weeds and irrigation water from major potato growing nine upazila under three districts of Bangladesh. Potato samples were collected from field, farmer storage and cold storage and other samples were collected from positive location under streaming and oozing test to investigate the presence of R. solanacearum. The causal agent (Ralstonia solanacearum) isolated from infected organism which has been described all around the world, tuber [36] [42], plant [15] [36] [42], soil [37] [43], banana [2], asymptomatic weeds [44] [45], chilli [46] and water [15] [47]. In potato tuber, no oozing was found but browning of the vascular bundle region of seed tuber were observed. Cross cut of plant samples showed bacterial ooze streaming in clear water. Cutting a diseased tuber will reveal browning and necrosis of the vascular ring and in adjacent tissues. A creamy fluid exudate usually appears spontaneously from the vascular ring at the cut surface. Bacterial ooze can emerge from the eyes and stem-end attachment of whole tubers, to which soil adheres. If cut stem or tuber vascular tissue is placed in water, threads of bacterial ooze exude [13].

TZC medium [23] is widely used and best studied media to characterize R. solanacearum. Virulent isolates showed the typical colony characters which are

mucous and pink-centered on TZC solid media and smooth margins are normally avirulent [24] [29] [30]. A total of one hundred thirty samples were used for isolation on TZC solid medium out of which 94% (*i.e.* 106) found positive for *R. solanacearum* presence.

Among the isolates fifty-seven isolates were able to cause rapid death of local cell of tissue between veins of tobacco leaves. *R. solanacearum* was able to produce HR in tobacco leaves. These results clearly indicated that like other plant pathogenic bacteria, *R. solanacearum* isolates possess Hrp type III secretion system which is responsible for inducing HR on tobacco leaves [28].

Incase of biochemical tests, fifty isolates showed positive gram reaction, KOH reaction, catalase, kovac'soxidase and pectolytic tests and showed similar reaction to temperature sensitivity test. So, isolates of R. solanacearum were gram negative and straight or curved rod shaped which is the characteristic feature of any plant pathogenic bacteria [29]. Gram negative bacteria have fragile cell walls which are bounded by an outer membrane that is why gram negative bacteria produce slime threads in a 3% KOH. Those slime threads were actually DNA. That is why this test is also called lytic release of DNA. But Gram-positive bacteria by contrast possess a thicker, more rigid cell wall which resists the disruptive effect of KOH [30]. Entire gram-negative bacteria produce gas bubbles when these were mixed with a drop of 3% H₂O₂ on glass slide. Production of gas bubbles gives a clue for presence of aerobic and facultative bacteria [29]. Kovacs oxidase test was used for differentiation between aerobic and anaerobic bacteria [48]. Fifty Isolates of *R. solanacearum* produced dark purple color within 10 - 60 seconds. It indicates those isolates were aerobic bacteria [25] [49]. Pectolytic test was physiological test. In pectolytic test, fifty isolates of *R. solanacearum* showed positive reaction producing soft rot in inoculated potatoes [25] [29] [32] [50]. In temperature sensitivity test all test isolates showed positive and typical growth at 28°C and 37°C but did not show any potential growth at 4°C and 40°C. R. solanacearum growth on TZC media at 28°C and 37°C not growth at 41°C [51]. R. solanacearum is able to survive even in colder environments for about 3 years and it is severe in temperature ranges of 24°C - 35°C with optimum of 27°C [52]. Nine isolates out of three districts produced the bands 280 bp specific for *R*. solanacearum. Many scientists in world wide also used 759/760 primers for detecting the *R. solanacearum* in potato [53] [54] [55].

Fifty isolates showed wilting symptoms on potato and tomato seedlings except brinjal seedlings except some chlorosis on leaves. At last wilting and collapse of whole plants can lead to rapid death. When the cut end of infected stem kept in a glass of water the milky or cloudy threads like streaming signifies the presence of *R. solanacearum* [19] [56]. So, it was observed that all (fifty) tested isolates expressed as race 3. However, in the present study race 1, race 2, race 4 and race 5 were not detected since race 1 is infect all solanaceous and a wide range of plants (solanaceous and nonsolanaceous weeds, diploid bananas, groundnut, olive, ginger, straw berry, geranium, eucalyptus, some other plants) which were

available in Asia, Australia and America while race 3 infects only potato and tomato worldwide [36], race 2 is restricted to triploid banana and heliconia; race 4 infects ginger; and race 5 is pathogenic on mulberry [40] [57] [58].

In case of biovar test, forty eight out of fifty isolates of R. solanacearum had shown to oxidize the different carbon sources from both disaccharides and sugar alcohols and changed the green color of the medium into yellow which was indicating the oxidation of both carbons so expressing those as biovar III. But, the remaining two isolates from soil and water neither changed the color for disaccharides nor for sugar alcohols which meant that those were not able to utilize either of the carbon sources and they were biovar I. R. solanacearum has been grouped into five biovars on the basis of utilizing or oxidizing three hexoses (mannitol, dulcitol and sorbitol) and three disaccharides (lactose, maltose and cellobiose) [31] [35]. Biovar I oxidize none of disaccharide sugar and hexose alcohol, biovar II oxidizes only disaccharide sugars, Biovar III oxidizes both of disaccharide sugars and hexose alcohols, biovar IV oxidizes only hexose alcohols whereas biovar V disaccharide sugar and oxidizes only mannitol hexose alcohol not sorbito and dulcitol [31] [35]. Bangladeshi R. solanacearum isolates of potato belonging to race 3 biovar III [17] [18]. It is clearly revealed that all groups of R. solanacearum isolates oxidized disaccharides (sucrose, lactose, and maltose) and sugar alcohols (manitol, sorbitol and dulcitol) within 3 - 5 days and confirmed biovar as III.

5. Conclusions

Among 106 collected isolates from different sources, 81.54% isolates showed positive reaction on TZC test, 50 isolates showed race 3; biovar III except one from each of Munshigonj and Nilphamari which showed as biovar I; and nine isolates out of three districts showed 280 bp DNA fragment amplification through the species-specific primer of Bangladeshi isolates.

The most prevalent and frequent isolate, race 3 biovar III were observed in Bangladesh at Munshigonj, Narayangonj and Nilphamari districts that are major potato growing areas and sources of seed potatoes for all over Bangladesh. It is urgently important to confirm the presence of the most disastrous *R. solanaceaum* isolate, race 3 biovar III isolates in other regions and prevent further spread to other hosts or regions of Bangladesh.

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Conflicts of Interest

The authors declare no conflicts of interest regarding the publication of this paper.

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