# Biochemical and molecular studies of early blight disease in tomato

Suchita J. Patel • R. B. Subramanian • Yachana S. Jha

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Abstract Tomato early blight occurs worldwide and it is prevalent wherever tomatoes are grown. Alternaria solani Sorauer, the causal agent, has been recognized as a serious foliar pathogen of tomato and there are very few cultivars which possess resistance against early blight. Alternaric acid is the major toxin of A. solani. In this study, alternaric acid and fungal culture filtrate were used as an elicitor in NDT-96 (tolerant) and GP-5 (susceptible) tomato varieties in order to study and compare their abilities to induce defense-related enzymes, viz., catalase, peroxidase,  $\beta$ -1,3 glucanase, phenylalanineammonia-lyase (PAL), chitinase and polyphenoloxidase (PPO) along with total phenols, and total soluble proteins. NDT-96 showed a rapid induction of all these pathogenesis-related enzymes except catalase along with total phenols as compared to GP-5 with both the treatments. Differential expression of total soluble proteins revealed higher protein content in NDT-96 as compared with GP-5. A 49.48 kDa protein was observed to be absent in GP-5. In addition, 25 microsatellite markers (SSR) were screened for polymorphisms among the above mentioned two tomato varieties. Of these, SSR 286 revealed a significant polymorphic band of 108 bp in NDT-96.

**Keywords** *Alternaria solani* · Pathogenesis-related enzymes · Simple sequence repeats · Systemic acquired resistance

## Introduction

Early blight, caused by Alternaria solani Sorauer, is one of the most common and destructive diseases of the cultivated tomato in areas of heavy dew, rainfall, and high relative humidity (Barksdale 1971; Nash and Gardner 1988). Alternaric acid is one of the major toxins found in the fungal culture filtrates (Brian et al. 1952); it causes chlorosis and necrosis and therefore plays a major role in early blight symptoms and defoliation (Pound and Stahmann 1951). Symptoms produced by an aqueous solution of crystalline alternaric acid when introduced into the plants are identical to symptoms produced by crude fungus filtrates. (Pound and Stahmann 1951) Control measures for these diseases include a 3- to 5-year crop rotation, routine applications of fungicides, and the use of disease-free transplants (Madden et al. 1978; Sherf and MacNab 1986).

Plants evoke a series of general defense reactions, including the production of phytoalexins and antimicrobial proteins, upon sensing invading microorganisms. (Radhajeyalakshmi *et al.* 2009). During the course of fungal attack, plants are induced to express a number of pathogenesis-related enzymes and proteins. Inoculation of plants with pathogens or treatment

S. J. Patel (⊠) • R. B. Subramanian • Y. S. Jha B. R. Doshi School of Biosciences, Satellite Campus, Sardar Patel University, Vallabh Vidhyanagar 388120, India e-mail: suchitapatel5@yahoo.com

with some chemical compounds can result in the establishment of systemic acquired resistance (SAR) (Ryals et al. 1996), which is accompanied by synthesis of pathogenesis-related proteins (PRproteins) (Linthorst 1991). Defense-related genes encode a variety of proteins including enzymes controlling secondary metabolism, PR-proteins and regulatory proteins that control the expression of other defense-related genes (Dixon et al. 1994). In early blight resistant lines a higher and more rapid induction of the PR-proteins, viz., chitinase and β-1, 3-glucanase (Lawrence et al. 1996, 2000), peroxidase (Fernandez et al. 1996), polyphenol oxidase (Thipyapong and Steffens 1997) and phenyalalanine ammonia lyase (Solorzano et al. 1996) are observed during the early infection process compared with those in susceptible lines (Lawrence et al. 1996, 2000). Catalase activity was observed to increase in a susceptible variety (Rani and Yasur 2009). Induction of defense proteins makes the plant resistant to pathogen invasion (Van Loon 1997), and has been correlated with defense against pathogen invasion in tomato (Bashan et al. 1985). Secondary plant metabolites correlated to early blight resistance include a higher total phenolic content (tannin, flavonol and phenol) in leaves and stems of early blight resistant varieties (Bhatia et al. 1972).

Fungicide treatments are generally the most effective control measures, but are not economically feasible in all areas of the world and may not be effective under weather conditions favorable for epidemics (Herriot et al. 1986). Resistant cultivars are potentially the most economical control measure because they can extend the intervals between fungicide sprays while maintaining control of the disease (Keinath et al. 1996; Madden et al. 1978; Shtienberg et al. 1995). The development of the polymerase chain reaction (PCR) (Saiki et al. 1988) has led to the development of many techniques to detect polymorphism at the DNA level (Powell et al. 1995; Rafalski et al. 1996). Of these techniques, some require the use of sequence-specific primers, such as microsatellite markers (SSR) or microsatellites (Powell et al. 1996). The SSR are short (mostly 2-4 bp) tandem repeats of DNA sequences and are useful due to their high degree of polymorphism and co-dominant character of heredity (He et al. 2003). The use of microsatellite polymorphisms to study the genetic diversity and variability was described for a number of plant species especially in tomato (Wang *et al.* 2006). They are mostly used as markers for genomic mapping, variety identification and marker-assisted selection in tomato (He *et al.* 2003).

The objectives of this investigation were to: (1) screen the two selected tomato varieties, *viz.*, NDT-96 and GP-5, by *in vitro* and *in vivo* bioassays using alternaric acid as an elicitor; (2) study and compare the effect of alternaric acid and fungal culture filtrate (FCF) on accumulation of phenolics and activities of catalase (EC 1.11.1.6), peroxidase (EC 1.11.1.7), phenylalanine ammonia-lyase (PAL) (EC 4.3.1.5),  $\beta$ -1,3 glucanase (EC 3.2.1.39), polyphenol oxidase (PPO; EC 1.14.18.1) and chitinase (EC 3.2.1.14) in tomato varieties; (3) study the induction of total soluble proteins in tomato varieties; and (4) study polymorphism among NDT-96 and GP-5 through SSR markers.

### Materials and methods

Fungal culture and its maintenance Alternaria solani isolate was grown on potato-dextrose agar (PDA) plates and these plates were incubated at  $25^{\circ}C\pm1^{\circ}C$ . After sporulation these plates were maintained at  $4^{\circ}C$  until further use.

*FCF production and its analysis* An 8 mm mycelial mat was cut with a sterile cork borer from 1-monthold *A. solani*, grown on a PDA plate and transferred into a 250 ml Erlenmeyer flask containing 100 ml of autoclaved potato-dextrose broth (PDB). Flasks were incubated from 21 days at  $25^{\circ}C \pm 1^{\circ}C$  in static submerged condition. Thereafter, the broth was filtered using Whatman filter paper No. 1.

Isolation of alternaric acid For development of crystals, an 8-mm mycelial mat was cut with a sterile cork borer from 1-month-old *A. solani*, grown on a PDA plate and transferred into a 250 ml Erlenmeyer flask containing 100 ml of PDB. Flasks were incubated for 18 days at  $25^{\circ}\pm1^{\circ}$ C in a static submerged condition. Mycelial mat from an 18-dayold culture grown on PDB was removed and the broth was filtered using Whatman No. 1 filter paper. The pH of the filtrate was adjusted to between 3.0 and 3.5 by 1 N HCl and extracted with an equal volume of ethanol. The colorless solution obtained was dissolved by adding drop by drop boiling carbon

**Table 1** Microsatellite marker (SSR) primers used for thescreening of two tomato cultivars (F: forward primer; R:reverse primer)

Ser. No.	Name of marker	Sequence (5'-3')
1	SSR 45	F: TGTATCCTGGTGGACCAATG
		R: TCCAAGTATCAGGCACACCA
2	SSR 46	F: TGTATCCTGGTGGACCAATG
		R: TCCAAGTATCAGGCACACCA
3	SSR 52	F: TGATGGCAGCATCGTAGAAG
		R: GGTGCGAAGGGATTTACAGA
4	SSR 67	F: GCACGAGACCAAGCAGATTA
		R: GGGCCTTTCCTCCAGTAGAC
5	SSR 76	F: ACGGGTCGTCTTTGAAACAA
		R: CCACCGGATTCTTCTTCGTA
6	SSR 80	F: GGCAAATGTCAAAGGATTGG
		R: AGGGTCATGTTCTTGATTGTCA
7	SSR108	F: TGTGTTGGATGTTTGGCACT
		R: GCCATTGAAACTTGCAGAGA
8	SSR 136	F: GAAACCGCCTCTTTCACTTG
		R: CAGCAATGATTCCAGCGATA
9	SSR 181	F: CAATCGAAACCGACGATACA
		R: GGTAGATCTGGATCGAGGAGG
10	SSR 241	F: TCAACAGCATAGTGGAGGAGG
		R: TCCTCGGTAATTGATCCACC
11	SSR 276	F: CTCCGGCAAGAGTGAACATT
		R: CGACGGAGTACTTCGCATTT
12	SSR 285	F: AGTGGCTCTCACCTACTGCG
		R: CAATTCTCAGGCATGAAACG
13	SSR 286	F: AGCTATGGAGTTTCAGGACCA
		R: ATTCAGGTAGCATGGAACGC
14	SSR 304	F: TCCTCCGGTTGTTACTCCAC
		R: TTAGCACTTCCACCGATTCC
15	SSR 565	F: GAGGATGATGAGAACTCGCC
		R: TCAGAGGCTTCTGGGTCAGT
16	SSR 637	F: AATGTAACAACGTGTCATGATTC
		R: AAGTCACAAACTAAGTTAGGG
17	Tom 8–9	F: GCATTGATTGAACTTCATTCTCGTCC
	ATT7	R: ATTTTTGFCCACCAACTAACCG
18	Tom 31A-	F: AATGTAATGGTGATGCTCTTCC
	32A TA11	<b>R:</b> CTCGGTTTTAATTTTTGTGTCT
19	Tom 41–	F: GAAATCTGTTGAAGCCCTCTC
17	42 TCC6	<b>B:</b> GACTGTGATAGTAAGAATGAG
20	Tom 43–	F: GCAGGAGATAATAACAGAATAAT
	44 TCC6	<b>B:</b> GGTAGAAGCCCGAATATCATT
21	Tom 47–	F: CAAGTTGATTGCATTACCTATTG
	48 AT10	<b>B:</b> TACAACAACATTTCTTCTTCCTT
22	Tom 49–	F: AAGAAACTTTTTGAATGTTGC
	50 AT10	R: ATTACAATTTAGAGAGTCAAGG
23	Tom 57–	F: TCTAAGTGGATGACCATTAT
	58 CT8	<b>R:</b> GCAGTGATAGCAAATGAAAAC

Table 1	(continued)	
Ser. No.	Name of marker	Sequence (5'-3')
24	Tom 144	F: CTGTTTACTTCAAGAAGGCTG R: ACTTTAACTTTATTATTGCGACG
25	Tom 196	F: CCTCCAAATCCCAAAACTCT R: TGTTTCATCCACTATCACGA

tetrachloride (2 ml) using a glass dropper. Residual ethanol was removed by evaporation and crystalline alternaric acid appeared on cooling.

Plant materials The following two tomato (Solanum *lycopersicum* L., syn. *Lycopersicon esculentum* Mill.) varieties were included in our studies: NDT-96 and GP-5. Seeds of NDT-96 were obtained from Anand Agriculture University, India, and seeds of GP-5 were obtained from Junagadh Agriculture University, India. The surface of tomato seeds was sterilized in 70% ethanol for 2 min, and thoroughly washed with sterile distilled water (Jung et al. 2005). Thereafter, seeds of both varieties were treated with 1% H<sub>2</sub>O<sub>2</sub> (Jaskani et al. 2006) and allowed to germinate in petri plates at 28°C for 2 days and sown in plastic cups filled with cocopeat (made from coconut husks) initially and after one week the seedlings were transferred to plastic cups filled with soil. The plants were allowed to grow for 4 weeks and watered daily; they were maintained under natural conditions.

Phytotoxicity tests of culture filtrates and toxin on tomato plants

Detached leaf bioassay Tomato leaves were washed in tap water. The leaves were surface sterilized with a cotton swab dipped in 4% sodium hypochlorite solution for 10–15 s. They were then washed thoroughly in sterile water and placed in sterile petri plates containing Whatman No. 1 paper. Thereafter these leaves were treated with toxin and FCF as follows: (*a*) 100  $\mu$ l of crystals along with water was applied gently on the leaf surface using a micropipette and 100  $\mu$ l of Water was applied on control leaves; (*b*) 100  $\mu$ l of FCF was applied gently on the leaf surface using a micropipette and 100  $\mu$ l sterile PDB was applied on control leaves. These plates were allowed to incubate at 25°±1°C for 1 to 2 days and the

Table 2	Resistant	ce of NI	DT-96 a	nd GP-	5 to 2	Alterna	ria s	solani,
using a (	–9 scale	disease	index (	DI) and	two	types of	of as	ssay

Evaluation technique	Cultivar	DI for FCF treatment	DI for toxin treatment
Detached-leaf bioassay	NDT-96	2.6	2.5
,	GP-5	7.2	7.4
In vivo bioassay	NDT-96	3.5	3.6
	GP-5	8.2	8.3

All the values were significant at  $P \le 0.001$  according to Tukey's Multiple Comparison Test method

moisture in dishes was maintained throughout the experiment. Visible symptoms were recorded after 24 h. The experiment was carried out in triplicate.

*In vivo bioassay* Tomato plants were grown and maintained in the similar way as described above for about 4 weeks and were used for *in vivo* bioassay. The tomato plants were treated with toxin and FCF as follows: (*a*) 0.5 ml of crystal solution along with water was taken in an injection syringe and this was

Fig. 1 The enzymatic activity of (a) Catalase (FCF) and (b) Catalase (toxin) was investigated in susceptible (GP-5) and tolerant (NDT-96) tomato genotypes following infection with alternaric acid toxin and fungal culture filtrate (FCF) of Alternaria solani. GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96Control and NI = NDT-96 Infected. PAL = phenylalanine ammonia-lyase; PPO = polyphenol oxidase. The error bar shows the standard error for the means of three replicates

injected in the petiole region of the plant; sterile water was injected in the control plants; (b) 0.5 ml of FCF was taken in an injection syringe and was injected in the petiole region of the plant; sterile PDB was injected in the control plants. The surface of the leaf was slightly pricked with a sterilized syringe in order to enhance the rate of infection. The plants were monitored regularly for the development of visible symptoms. The experiment was carried out in triplicate.

*Disease assessment* Disease severity was rated as described previously (Gaube *et al.* 2004), by visual assessment based on a 0–9 scale. A disease index (DI), obtained from the average of each leaf value, was used to assess plant resistance. Differences between cultivars were tested using Tukey's Multiple Comparison Test method. The software used was GraphPad Prism Version 3.

Induction of biochemical defense mechanisms To study the induction of catalase, peroxidase,  $\beta$ -1,3-glucanase, PAL, chitinase, PPO and phenolics,



one-month-old tomato plants were given the treatments by toxin and FCF as described above for *in vivo* bioassay. Four sets of tomato plants (GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96 Control, and NI = NDT-96 Infected) were treated similarly as above. Whole plants from four sets were harvested at various time intervals (0, 6, 12, 24, 48, 120 and 168 h) after treatment and leaf samples were collected and used for enzyme analysis. The experiment was carried out in triplicate.

Catalase (CAT) assay CAT activity was measured using the modified Barber (1980) method. The enzyme activity was expressed as  $\mu$ g/min/ml.

Peroxidase (POX) assay

POX activity was measured using the modified Sumner and Gjessing (1943) method. The enzyme activity was measured at 430 nm and expressed as change in units/min/g/ml. *PAL assay* PAL activity was measured using the Dickerson *et al.* (1984) method. The enzyme activity was measured at 290 nm and expressed as nmol transcinnamic acid/min/g.

 $\beta$ -1,3 glucanase assay  $\beta$ -1,3 glucanase activity was measured using the Pan *et al.* (1991) method. The enzyme activity was measured at 500 nm and expressed as units/ml/min.

*PPO assay* PPO activity was measured using the Siriphanich and Kader (1985) method. The change in O.D. was recorded at 30-s intervals up to 3 min at 495 nm. The enzyme activity was expressed as changes in units/min/mg total protein.

Chitinase assay 1% colloidal chitin (pH 6.6) was prepared by Rani and Yasur's method (2009). A 1% solution was prepared from this precipitate and stored at 4°C until use. Chitinase activity was measured using the Reissig *et al.* (1959) method. Enzyme activity was expressed as  $\mu$ mol GlcNAc/h/mg protein.

Fig. 2 The enzymatic activity of (a) Peroxidase (FCF) and (b) Peroxidase (toxin) was investigated in susceptible (GP-5) and tolerant (NDT-96) tomato genotypes following infection with alternaric acid toxin and fungal culture filtrate (FCF) of Alternaria solani. GC = GP-5 Control. GI = GP-5Infected, NC = NDT-96Control and NI = NDT-96 Infected. PAL = phenylalanine ammonia-lyase; PPO = polyphenol oxidase. The error bar shows the standard error for the means of three replicates



*Estimation of total phenolic content* Total phenolic content was measured using the Zieslin and Ben-Zaken (1993) method. The content of the total soluble phenols was calculated based on a standard curve obtained from a Folin Ciocalteau reaction with phenol and expressed as  $\mu g/g$  f.wt.

## Data analyses

Data were subjected to analysis of variance (ANOVA). Multiple mean comparisons were performed using Tukey's Multiple Comparison Test and comparisons between treatments were made using unpaired *t*-test in GraphPad Prism Version 3. One-way ANOVA and unpaired *t*-test were performed between both controls and between both infected tomato varieties.

Analysis of total soluble proteins Four sets of tomato plants (GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96 Control, and NI = NDT-96 Infected) were treated with toxin in a similar way as described

Fig. 3 The enzymatic activity of (a)  $\beta$ -1,3-glucanase (FCF) and (b)  $\beta$ -1,3-glucanase (toxin) was investigated in susceptible (GP-5) and tolerant (NDT-96) tomato genotypes following infection with alternaric acid toxin and fungal culture filtrate (FCF) of Alternaria solani. GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96 Control and NI = NDT-96 Infected. PAL = phenylalanine ammonia-lyase; PPO = polyphenol oxidase. The error bar shows the standard error for the means of three replicates

above. Whole plants from four sets were harvested at various time intervals (0, 12, 24, 48, 120 and 168 h) after treatment and leaf samples were collected and used for total soluble protein analysis. Tomato leaves (1 g) were homogenized in 10 ml of 0.1 M phosphate buffer (pH 7.0). The mixture was centrifuged at 10,000g for 20 min. The supernatant was used as a sample to analyze soluble proteins. The concentration of proteins was estimated by the method of Lowry et al. (1951) using bovine serum albumin as the standard. For one-dimensional separation, 12% linear sodium dodecyl sulfate (SDS)-polyacrylamide slab gels (1 mm thickness) overlaid with stacking gel (Laemmli 1970) were used. The electrophoresis was carried out at 50 V. Once tracking dye reached the anode, the run was stopped and the gels were carefully removed and placed in 5 volumes of Colloidal Blue Stain by the modified method of Neuhoff et al. (1988) for 1 h or overnight. The gels were washed with distilled water and destained using deionized water. The electropherograms were photographed and analyzed using the gel documentation



system (AlphaEase FC 4.0, Alpha Innotech, Minneapolis, MN, USA).

DNA isolation and marker analysis For the SSR analysis, DNA was isolated from young and fresh leaves of NDT-96 and GP-5 according to Doyle and Doyle's method (1990). A total of 25 SSR primers were surveyed (Table 1). Primers for SSR 45 to SSR 637 and Tom 144-Tom 198 were obtained from the Sol Genomics Network at www.sgn.cornell. edu. Primers for Tom 8-9 to Tom 57-58 were obtained from the literature (Rajput et al. 2006). PCR was conducted in 12-µl volumes containing the following: 1 µl DNA, 50 ng µl<sup>-1</sup>; 0.75 µl, 10x assay buffer; 1.5 µl, 2.5 mM dNTPs; 0.5 µl, 15 mM MgCl<sub>2</sub>; 0.25 µl, 3 Units Taq DNA polymerase (Bangalore Genei, Karnataka, India); 1 µl, 10 pmol  $\mu$ <sup>-1</sup> each of forward and reverse primers and 7  $\mu$ l of sterile water. The PCR profile starts with 95°C for 1 min followed by 35 cycles of denaturation at 95°C for 30 s, annealing at 54°C for 1 min, extension at 72°C for 2 min. A final extension at

Fig. 4 The enzymatic activity of (a) PAL (FCF) and (b) PAL (toxin) was investigated in susceptible (GP-5) and tolerant (NDT-96) tomato genotypes following infection with alternaric acid toxin and fungal culture filtrate (FCF) of Alternaria solani. GC = P-5 Control, GI = GP-5 Infected, NC = NDT-96 Control and NI = NDT-96 Infected. PAL = phenylalanineammonia-lyase; PPO = polyphenol oxidase. The error bar shows the standard error for the means of three replicates

72°C for 5 min was included. The PCR products were electrophoresed in a 3% agarose gel with ethidium bromide at 50 V for 2–3 h. The gel was then observed on a UV transilluminator. DNA samples were sent for sequencing to AN Xplorigen Technologies Pvt. Ltd. (New Delhi, India). The sequence obtained was submitted to GenBank. The GenBank accession number after acceptance of sequence is GU930288.

## Results

Phytotoxicity tests of culture filtrates and toxin on tomato plants

Treatment with toxin and FCF showed a similar pattern of infection. Cultivar NDT-96 showed tolerance to *A. solani*, but GP-5 showed clear susceptibility to *A. solani*. Significant differences in resistance were identified between the two cultivars (Table 2).



Induction of biochemical defense mechanisms

*CAT assay* By comparing the treatment of tomato leaves with FCF and with toxin, a similar pattern of induction of defense-related enzymes was observed. Studies of induction of defense mechanism revealed that a high accumulation of CAT was observed in FCF-and toxin-treated GP-5. The levels of CAT increased by 85.97 (t=42.01, P<0.0001) at 168 h (Fig. 1a) with FCF treatment and 84.97 (t=19.31, P<0.0001) (Fig. 1b) with toxin treatment in susceptible cultivar GP-5 at 168 h in response to the said treatments.

*POX assay* A significant increase in POX activity was observed in FCF- and toxin-treated NDT-96. The levels of POX increased by 7.867 (t=4.667, P< 0.0001) (Fig. 2a) with FCF treatment and by 6.867 (t=1.909, P<0.0001) (Fig. 2b) with toxin treatment in NDT-96 at 168 h.

*PAL assay* A significant increase in PAL activity was recorded in FCF- and toxin-treated NDT-96 plants.

The levels of PAL increased by 93.58 (t=39.48, P< 0.0001) (Fig. 3a) with FCF treatment and by 93.97 (t=37.98, P<0.0001) (Fig. 3b) with toxin treatment in NDT-96 at 168 h.

 $\beta$ -1,3 glucanase assay Treatments of tomato plants with FCF and toxin elicited an increase in  $\beta$ -1,3glucanase activity. A significant increase in  $\beta$ -1,3glucanase activity was found: 0.6167 (*t*=46.67, *P*< 0.0001) in NDT-96 at 168 h with FCF treatment (Fig. 4a) and 0.5967 (*t*=42.43, *P*<0.0001) at 168 h with toxin treatment (Fig. 4b).

*PPO assay* Upon pathogen infection the activities of PPO increased 3.967 (t=1.909, P<0.0001) at 168 h in NDT-96 with FCF treatment (Fig. 5a), and increased 4.167 (t=1.697, P<0.0001) at 168 h (Fig. 5b) in NDT-96 with toxin treatment.

*Chitinase assay* Chitinase activity increased rapidly in tomato upon treatment with FCF and with toxin. It increased significantly 7.867 (t=10.18, P<0.0001) at



Fig. 5 The enzymatic activity of (a) Chitinase (FCF) and (b) Chitinase (toxin) was investigated in susceptible (GP-5) and tolerant (NDT-96) tomato genotypes following infection with alternaric acid toxin and fungal culture filtrate (FCF) of Alternaria solani. GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96Control and NI = NDT-96 Infected. PAL = phenylalanine ammonia-lyase; PPO = polyphenol oxidase. The error bar shows the standard error for the means of three replicates

168 h in NDT-96 with FCF treatment (Fig. 6a) and 7.967 (t=6.152, P<0.0001) with toxin treatment (Fig. 6b) in NDT-96. A significantly higher enzyme activity was detected at all sampling days.

*Estimation of total phenolic content* Phenolic content increased significantly in FCF-and toxin-treated NDT-96. The levels of total phenol increased 94.97 (t= 88.47, P<0.0001) at 168 h in NDT-96 with FCF treatment (Fig. 7a) and 92.87 (t=72.13, P<0.0001) with toxin treatment (Fig. 7b) in NDT-96.

*Analysis of total soluble proteins* (Figs. 8 and 9) SDS gel (12%) of total soluble protein showed differential expression of PR-proteins in leaves of the tolerant (NDT-96) and susceptible (GP-5) tomato cultivars inoculated with alternaric acid toxin at different periods of inoculation. A specific band of 49.48 kDa was observed to be absent in GP-5 at all time intervals.

*Molecular marker analysis* (Fig. 10) A 3% PCR agarose gel showed the presence of 108 bp band present in NDT-96. This particular band was absent in GP-5.

# Discussion

Reliable and repeatable techniques for large-scale screening are necessary to identify host plant resistance. Techniques have been developed for early blight resistance screening under field, glasshouse, and laboratory conditions. In our studies, in the laboratory as well as under field conditions, FCF and fungal toxin have been used in screening for resistance. Locke (1948) used detached leaflet assays for evaluation of early resistance as a means to circumvent the influence of growth habit, which may affect the reaction of plants in the field or glasshouse. In field tests, large populations can be assessed under normal growing conditions during the whole life cycle of the plants. By both in vitro and in vivo assays it can be proved that NDT-96 is a tolerant variety and GP-5 is a susceptible variety.

The qualitative estimations of catalase, peroxidase,  $\beta$ -1,3 glucanase, phenylalanine-ammonia-lyase, chitinase and polyphenol-oxidase enzymes indicated the role of these oxidative and hydrolytic enzymes in plant responses towards the pathogen infection.

Fig. 6 The enzymatic activity of (a) PPO (FCF) and (b) PPO (toxin) was investigated in susceptible (GP-5) and tolerant (NDT-96) tomato genotypes following infection with alternaric acid toxin and fungal culture filtrate (FCF) of Alternaria solani. GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96 Control and NI = NDT-96 Infected. PAL = phenylalanine ammonia-lyase; PPO = polyphenol oxidase. The error bar shows the standard error for the means of three replicates



Fig. 7 The enzymatic activity of (a) Total phenol (FCF) and (b) Total phenol (toxin) was investigated in susceptible (GP-5) and tolerant (NDT-96) tomato genotypes following infection with alternaric acid toxin and fungal culture filtrate (FCF) of Alternaria solani. GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96 Control and NI = NDT-96 Infected. PAL = phenylalanineammonia-lyase; PPO = polyphenol oxidase. The error bar shows the standard error for the means of three replicates





**Fig. 8** Polyacrylamide gel electrophoresis (12%) of total soluble protein showing differential expression of PR-proteins in leaves of the tolerant (NDT-96) and susceptible (GP-5) tomato varieties inoculated with alternaric acid toxin at different times. Loaded samples were adjusted to a constant amount of protein (15  $\mu$ g). M = marker (kDa), GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96 Control and NI = NDT-96 Infected. Figures 8 and 9 show the time differential expression of total soluble proteins from 0 h to 168 h

Certain quantitative differences in oxidative enzymes in pathogen-infected as well as uninfected plants had occurred. These changes were also prominent between the two varieties of the tomato plants. CAT is an oxygen-scavenging enzyme that removes toxic substrates during development, which are otherwise lethal. Large increases in foliar catalase activity were observed in Algerian-Susceptible but not in Algerian-Resistant barley (Hordeum vulgare L.) leaves inoculated with Blumeria graminis (Vanacker et al. 1998). However, catalases were earlier reported to act as antagonist to the enzyme peroxidase so this could be the reason for the decreased activity of the enzyme in NDT-96. They were also known to be inhibited by the salicylic acid release in the plants due to SAR (Chen et al. 1993).

Peroxidase is involved in the production of reactive oxygen species, which are directly toxic to the pathogen or indirectly reduce the spread of the pathogen by increasing the cross-linkage and lignifi-



Fig. 9 Polyacrylamide gel electrophoresis (12%) of total soluble protein showing differential expression of PR-proteins in leaves of the tolerant (NDT-96) and susceptible (GP-5) tomato varieties inoculated with alternaric acid toxin at different times. Loaded samples were adjusted to a constant amount of protein (15  $\mu$ g). M = marker (kDa), GC = GP-5 Control, GI = GP-5 Infected, NC = NDT-96 Control and NI = NDT-96 Infected. Figures 8 and 9 show the time differential expression of total soluble proteins from 0 h to 168 h

cation of the plant cell walls (Hammond-Kosack and Jones 1996). POX is required for the final polymerization of phenolic derivatives into lignin and also involved in suberization or wound healing (Ward *et al.* 1991). Increased POX activity was observed in a number of resistant interactions involving plantpathogenic fungal and bacterial interactions (Deborah *et al.* 2001; Flott *et al.* 1989; Reimers *et al.* 1992). The results of the present study indicated that POX activity significantly increased in NDT-96 after treatment when compared to GP-5, with both the treatments.

Phenylalanine ammonia-lyase is the key enzyme in the synthesis of the secondary, endogenous signaling molecule salicylic acid, which in turn activates the expression of a variety of PR genes (Mauch-Mani and Slusarenko 1996). Therefore, these results suggest that increased accumulation of PAL with a pathogen could re-establish the notion that in response to invasion by a pathogen, PAL is the first enzyme of phenylpropanoid metabolism and plays a significant role in the regulation of biosynthesis of phenols in plants. Induction of PAL enzyme activity is correlated with increased resistance to pathogenic infection (Bell *et al.* 1984). It was demonstrated that treatment with PAL inhibitors suppressed resistance and increased susceptibility in several host-pathogen interactions. Further, it was demonstrated that reduction of phenyl-propanoid metabolism through inhibition of PAL activity in transgenic tobacco also rendered tissues more susceptible to *Cercospora nicotianae* (Maher *et al.* 1994). The results of the present study indicated that PAL activity increased significantly from 0 h after treatment and reached the maximum 168 h after treatment in NDT-96 as compared with GP-5 plants.

 $\beta$ -1,3-glucanases (PR-2 family) and chitinases (PR-3 family) degrade the fungal cell wall and cause lysis of fungal cells. Chitin and glucan oligomers released during degradation of the fungal cell wall act as an elicitor that elicits various defense mechanisms in plants (Frindlender et al. 1993). Induction of defense proteins makes the plant resistant to pathogen invasion (Van Loon 1997), and has been correlated with defense against pathogen invasion in tomato (Bashan et al. 1985). Chitinases and glucanases probably slow fungal ingress in the plant, as indicated by their inhibition of the in vitro growth of A. solani (Lawrence et al. 1996). Enzyme preparations from resistant lines also induced the in vitro release of elicitors of the hypersensitive response from A. solani, whereas enzymes from susceptible lines did not (Lawrence et al. 2000). The results of the present study indicated that  $\beta$ -1,3-glucanase activity increases in NDT-96 plants. A significant increase in chitinase activity was observed at 168 h in NDT-96 plants.

Polyphenol oxidase is systemically up-regulated in response to *A. solani* infection and is detected in

**Fig. 10** Polymerase chain reaction gel showing the presence of marker linked to resistance in the tolerant tomato variety. M=100 bp DNA ladder, T = Tolerant variety (NDT-96) and S = Susceptible variety (GP-5)



leaves of upper nodes but not of lower nodes (Thipyapong and Steffens 1997). This induction pattern coincides with the observation of temporary resistance of young leaves to A. solani infection (Johanson and Thurston 1990). PPOs catalyze the oxygen-dependent oxidation of o-dihydroxyphenols to *o*-quinones, which are more toxic to pathogens than the former. Direct toxicity of quinones against pathogens has also been proposed (Mayer et al. 1965). In addition, many studies have shown that PPO is induced in response to mechanical wounding, fungal and bacterial infection, and by treatment with signaling molecules such as jasmonic acid/methyl jasmonate (MeJA), systemin and salicylic acid (Constabel et al. 2000; Stewart et al. 2001). Systemic induction of PPO expression in response to wounding and pathogens might provide an additional line of defense to protect plants against further attack by pathogen and insects (Stout et al. 1999; Thipyapong et al. 1995). In the present study, PPO activity increased at 168 h in NDT-96.

The phenolic compounds may contribute to enhancing the mechanical strength of host cell walls and may also inhibit the fungal growth, as phenolics are fungitoxic in nature. Altering the level of phenolic compounds in plants has been demonstrated to change disease susceptibility (Yao et al. 1995). Increase in phenolic content in plants has been correlated with increased resistance to pathogens (Velazhahan and Vidhyasekaran 1994). Carrasco et al. (1978) showed that treatment of susceptible tomato plants with precursors of phenolic compounds such as quinic acid and phenylalanine increased their phenolic content and their resistance to Fusarium oxysporum. However, in the present study, the accumulation of phenolics was higher in the tolerant cultivar than in the susceptible cultivar.

Expressed defense-related genes such as those encoding pathogenesis-related proteins are used as markers for the establishment of SAR (Du and Klessig 1997). PR-proteins are host-coded proteins induced by different types of pathogens and abiotic agents. Synthesis and accumulation of PR proteins have been reported to play an important role in plant disease resistance (Van Loon 1997). In our study, 49.48 kDa protein has been induced by *A. solani* in leaves of a tolerant cultivar, but not of a susceptible cultivar. SSRs provide highly informative markers because they are co-dominant (unlike RAPDs and AFLPs) and generally highly polymorphic (Daugrois *et al.* 1991; Devi and Reddy 2002). Furthermore, the ease and speed of genetic analysis based on SSRs enhance the ability to make a greater number of SSRs available to the scientific community, at least for most of the species of economic value, such as the tomato.

Thus, it can be suggested that there is a correlation between constitutive induced levels of these enzymes and plant resistance. On application of each of these treatments there was an elicitation of array of signals, which helps in turn to increase the levels of the biochemicals that play a role in plant immunity. The induction level of all these enzymes except catalase and total phenols was observed to increase in variety NDT-96 more than in their respective controls. However, a considerable increase of catalase enzyme had been found in variety GP-5. Both FCF and toxin treatments showed a similar pattern of induction of these enzymes as well as total phenols. These findings strongly suggest that alternaric acid induces SAR and is associated with resistance. The result of differential expression of total soluble proteins indicates the possibility of involvement of this particular protein in the defense of tomato against early blight. The presence of the 108 bp band in the tolerant variety may be used as a marker for genomic mapping, variety identification and marker-assisted selection in tomato.

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