Evaluating Bacterial Wilt-resistant Tomato Genotypes Using a Regional Approach

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Additional index words. Lycopersicon esculentum, Pseudomonas solanacearum, Physalis ixocarpa, resistance, tomatillo

Abstract. Several procedures for evaluating the resistance of tomato (Lycopersicon esculentum Mill.) to bacterial wilt were used to account for diversity in strains of Pseudomonas solanacearum Smith and to approximate resistance under field conditions. Five strains of P. solanacearum from Florida and one from North Carolina were inoculated onto 19 tomato genotypes and one tomatillo (Physalis ixocarpa Brot.) genotype using a stem-puncture technique. Genotypes were also transplanted as seedlings into naturally infested soil. Resistance was evaluated by comparing the response of each genotype to the susceptible cultivars Bonny Best and Sunny. With the stem-puncture technique, the mean incidence of disease ranged from 30% with the strain from North Carolina to 94% with a strain from northern Florida. Significant differences in the resistance of genotypes and pathogenicity of strains were observed. However, no interaction between strain and genotype was observed. Using naturally infested soil, the mean incidence of disease was 51% and significant differences in the resistance of genotypes were also observed. Hawaii 7997, Hawaii 7998, and CRA 66 had the lowest incidence of disease, regardless of inoculation method. The results indicate that assessing pathogen diversity and using a combination of resistance screening techniques can facilitate the evaluation of many genotypes, account for potential regional variability in the pathogen, and differentiate levels of field resistance to tomato bacterial wilt.

Bacterial tomato wilt, caused by *Pseudomonas solanacearum*, is a limiting factor for tomato production in many tropical, subtropical, and warm temperate regions of the world (Hayward, 1991). In Florida, bacterial tomato wilt was first reported in 1897 (Rolfs, 1898) and has caused sporadic losses in the southern region of the state (Sonoda et al., 1979). In the 2000-ha Quincy tomato production area in northern Florida, recent crop losses due to bacterial wilt have been especially severe and in some cases have forced growers to abandon entire fields.

Crop rotation is not a viable control tactic because the bacterium is indigenous to soils of Florida and Georgia (Jaworski and Morton, 1964; Sonoda, 1978) and can persist indefinitely in infested fields. Soil fumigation does not provide acceptable levels of season-long disease control (Enfinger et al., 1979). Developing resistant cultivars is the most logical solution for suppressing bacterial wilt epidemics. Although the emphasis of bacterial wilt research in Florida has been on developing resistant cultivars, a cultivar having resistance to *P. solanacearum* and good horticultural characteristics has been difficult to obtain (Scott et al., 1993; Sonoda et al., 1979).

Several bacterial wilt-tolerant genotypes have been released (Anais, 1986; Henderson and Jenkins, 1972; Jaworski et al., 1987), but the resistance often breaks down outside the specific region in which they were developed (Scott et al., 1993). The inability of resistant genotypes to perform consistently in different locations

Received for publication 2 Apr. 1992. Accepted for publication 25 June 1993. Florida Agricultural Experiment Station Journal series R-03038. We thank Susan Lee for technical assistance, Robert Stall for access to his collection of *P. solanacearum* strains, and the Gadsden County Tomato Growers Association for financial assistance. The cost of publishing this paper was defrayed in part by the payment of page charges. Under postal regulations, this paper therefore must be hereby marked *advertisement* solely to indicate this fact.

has been attributed to environmental influences such as temperature (Krausz and Thurston, 1975; Mew and Ho, 1977; Sonoda, 1978) and variability among *P. solanacearum* strains (McLaughlin and Sequeira, 1989; Prior et al., 1990). One way to overcome this problem is to develop a screening process that accounts for the various factors affecting resistance.

The purpose of this study was to evaluate the potential for reducing the incidence of tomato bacterial wilt in the Quincy tomato production area using resistant genotypes. Regional diversity in the pathogen was assessed and several screening techniques were used to account for the potential variability of local *P. solanacearum* strains, minimize variation in results due to fluctuating environmental conditions, and to approximate infections under natural field conditions.

Materials and Methods

Experiments were conducted in a temperature-controlled green-house, with days at 32 to 37C and nights at 16 to 21C, to minimize variation in resistance expression due to temperature.

Genotypes. Nineteen tomato genotypes (Table 1) and one tomatillo genotype (Lot 8-33487; Rogers NK Seed Co., Naples, Fla.) were selected based on previous reports (published and unpublished) of bacterial wilt resistance. To reduce variability among replications, reactions of the selected genotypes were compared to those of the susceptible tomato cultivars Bonny Best and Sunny in each replication.

Characterization of P. solanacearum strains. Strains of P. solanacearum were isolated from tomato plants with symptoms of bacterial wilt from various locations in the Quincy production region. Strains were purified and stored in sterile tap water at room temperature between experiments. The ability of strains to oxidize various disaccharides and hexose alcohols was examined to deter-

Table 1. Bacterial wilt-resistant tomato genotypes used in screening tests.

Genotype	Location or institution ^z	Source of resistance
Capitan	Petoseed Co.	CRA 66
Caraibo	INRA, West Indies	CRA 66
CL 5915-93	AVRDC, Taiwan	?
CL 5915-153	AVRDC, Taiwan	?
CRA 66	INRA, West Indies	West Indies ecotype
FMX 192	Ferry Morse Seed Co.	?
GA 219	Univ. of Georgia	PI 126408 (Lycopersicon esculentum)
GA 1095	Univ. of Georgia	PI 196298 (L. esculentum)
GA 1405	Univ. of Georgia	PI 251323 (L. pimpinellifolium)
GA 1565	Univ. of Georgia	PI 263722 (L. esculentum)
Hawaii 7997	Univ. of Hawaii	PI 127805A (L. pimpinellifolium)
Hawaii 7998	Univ. of Hawaii	PI 127805A (L. pimpinellifolium)
Island Red	Yates Brothers, Trinidad	?
IHR 66	ICAR, India	?
PI 126408	Univ. of Florida	L. esculentum
Tomatillo	Rogers NK Seed Co.	Physalis ixocarpa
Venus	North Carolina State Univ. and Beltsville 3841 (<i>L.esculentum</i>)	PI 129080 (L. esculentum var cerasiforme)
XPH 5675	Asgrow Seed Co.	?
XPH 5677	Asgrow Seed Co.	?
84 BWR	ICAR, India	?

²Location or institution where resistance screening was performed. INRA = Institute National de la Recherche Agronomique; AVRDC = Asian Vegetable Research and Development Center; ICAR = International Center for Agricultural Research.

mine biovar grouping (Hayward, 1964). Race designation was determined by inoculating seedlings of 'Hicks' tobacco (*Nicotiana tabacum*), 'Bonny Best' tomato, 'Golden Summer' pepper (*Capsicum annuum*), and 'Mini Finger' eggplant (*Solanum melongena*) (Buddenhagen et al., 1962). Five strains from the Quincy production area and a sixth strain (K60) isolated from tomato in North Carolina (Kelman and Person, 1961) were used in inoculations. Seedlings were inoculated by the stem-puncture technique described below and observed for developing bacterial wilt symptoms for 3 weeks.

A quantitative measure of diversity among strains isolated from Quincy was obtained by comparing their fatty acid profiles to profiles of 20 *P. solanacearum* strains collected from various plant hosts and locations around the world (from R.E. Stall, Univ. of Florida, Gainesville). The collection of strains contained members from races 1, 2, and 3 and biovars 1, 2, and 3.

Strains were grown on trypticase soy broth agar for 24 ± 2 h at 28 ± 1 C. Cellular fatty acids were derivatized to fatty acid methyl esters (FAMEs) (Miller, 1982). Statistical analyses of FAME profile data were conducted using the MIDI Microbial Identification System, Newark, Del. version 3.60 (Sasser, 1990a). Cluster analysis of the FAME profiles was performed with the MIDI dendrogram software program. Strains within a Euclidian distance (ED) of 10 were considered to be the same species, while strains that link within six units were considered to be the same subspecies or biotype (Sasser, 1990b).

Stem-puncture inoculation technique. Seeds were sown in Metro-Mix 220 (W.W. Grace, Cambridge, Mass) in polystyrene planter flats containing 72 individual cells, each with a volume of 65.5 cm³. Seedlings were thinned to one per cell. Four-week-old seedlings were inoculated using the stem-puncture technique (Winsted and Kelman, 1952), which consists of forcing a sharp needle into the stem through a drop of bacterial suspension that has been placed in the axil of the second or third expanded leaf below the stem apex.

Inoculum was prepared from 2-day-old colonies growing on

nutrient agar with yeast extract at 27C. Bacterial suspensions were diluted in sterile tap water to 5.0×10^8 colony-forming units (cfu)/ml (OD_{600nm} = 0.775). Ten plants per genotype were inoculated. Plants were incubated in the greenhouse and observed for the presence of bacterial wilt symptoms after 18 days. The experiment was replicated three times.

Infested soil technique. Infested soil, a Carnegie loamy fine sand (Plinthic Kandiudult: Clayey, Kaolinitic, Thermic) with a pH of 5.3 and an organic content of 4.0%, was obtained from a tomato farm in Gadsden County, Fla., in Nov. 1991. Soil was collected from the base of plants with symptoms of bacterial wilt, passed through a 1-cm mesh screen to remove plant debris, and stored in plastic containers. The inoculum density of the soil was determined before each replication using a modified soil dilution plating technique (Chellemi et al., 1992).

Four-week-old seedlings were transplanted into 700-cm³ pots containing infested soil. Ten pots, each containing one seedling, were prepared for each genotype. After transplanting, the pots were incubated in the greenhouse and observed for the presence of bacterial wilt symptoms for 42 days. At the end of each experiment, the bacterium was isolated from several symptomatic plants to confirm the presence of the pathogen. The experiment was repeated three times.

Data analysis. Disease incidence for each genotype was expressed as a percentage of the incidence on the susceptible control cultivars Sunny and Bonny Best. Data were checked for normality and, when appropriate, transformed using the inverse sign transformation. Data from the stem-puncture inoculations were analyzed as a 6×20 factorial design with combinations of six strains and 20 genotypes. An analysis of variance was conducted to determine the significance of main effects or interactions using SAS's General Linear Model Procedure (SAS Institute, 1988). Data from the infested soil inoculations were analyzed as a randomized complete-block design, with replications over time as the block. Analysis of variance was conducted to determine if there were significant differences in resistance among genotypes.

Results

All strains were characterized as race 1, producing symptoms of bacterial wilt in tomato, pepper, and eggplant. No symptoms were produced on tobacco. All of the strains failed to oxidize the disaccharides and hexose alcohols and were classified as belonging to biovar 1.

A dendrogram of fatty acid profiles linked the 30-member population of strains at <10 ED units (Fig. 1), a result indicating that all of the samples are likely to be the same species (Sasser, 1990b). Six clusters linked at >6 ED units, a result indicating that the population may represent several *P. solanacearum* subspecies or biotypes. Strains from the Quincy production area were found in four of the six clusters.

The stem-puncture technique indicated significant differences in strain pathogenicity and genotype resistance (Table2). However, there was no significant strain × genotype interaction. Strains from the Quincy production area were significantly more virulent than the strain from North Carolina (Table 3). Symptoms developed in at least 60% of the plants for all genotypes (Table 4). The genotypes Hawaii 7997, Hawaii 7998, and CRA 66 had significantly fewer diseased plants than eight other genotypes.

Inoculum density in the infested soil ranged from 1.6×10^2 to 1.7×10^3 cfu/g soil. Disease incidence ranged from 0 in Hawaii 7997 to 92% in XPH 5677, using the infested soil technique (Table 4). Differences in resistance were significant at P < 0.10 (Table 2). As in the stem-puncture technique, Hawaii 7997, Hawaii 7998, and CRA 66 had the lowest disease incidence.

Discussion

The *P. solanacearum* strains were characterized to assess the potential strain variability within the Quincy production region and to draw comparisons to strains from other hosts or regions throughout the world. The race 1, biovar 1 designation is typical of *P. solanacearum* strains in the southeastern United States (Hayward, 1991; Velupillai and Stall, 1984). However, these characterizations alone do not indicate the potential variability among strains. For example, McLaughlin and Sequeira (1989) found that large biochemical and pathogenic differences existed among strains of *P. solanacearum* isolated from a single field in Hastings, Fla. In this study, comparing bacterial fatty acid profiles of a collection of strains from worldwide locations comprising three different races and biovars provided corroborative evidence of diversity among the Quincy strains.

Without a clear understanding of variability in virulence to tomato within a production region, breeders will be unsure of the stability of their resistant cultivars and the adaptability of resistant cultivars developed in other geographic regions. Successful screening will depend on including strains representative of the population found in commercial fields. The stem-puncture technique was used because it helps evaluate many P. solanacearum strains more than would be possible in evaluations conducted in naturally infested soil. Other common resistance screening techniques for bacterial wilt that permit the evaluation of many bacterial strains include the root-injury, root-dip, and clipping techniques (McCarter, 1973; Winstead and Kelman, 1952). In this study, the stem-puncture technique demonstrated that no significantly different responses occurred between strain and genotype. In addition, the Quincy strains were more virulent than the North Carolina strain. Thus, we would expect the resistance of genotypes Hawaii 7997, Hawaii 7998, and CRA 66 to be stable in this region.

In the previously mentioned screening techniques, the plant

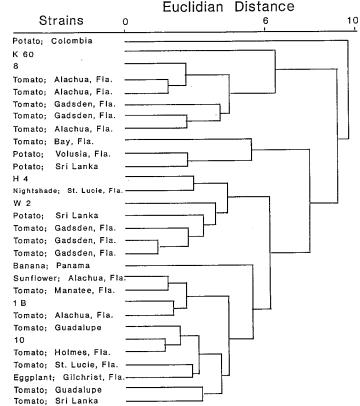


Fig. 1. Dendrogram derived from cluster analysis of fatty acids from 30 strains of *Pseudomonas solanacearum*. Strains are designated according to the plant host and county, state, or country from which they were isolated. Accession numbers are available from the authors. Strains H4, K60, W2, 1B, 8, and 10 were used in evaluating resistance to bacterial wilt. Strains within a Euclidian distance of ten are likely to be the same species, while strains that link within six units are likely to be the same subspecies or biotype.

Table 2. Analysis of variance (ANOVA) for data (transformed) from both resistance screening techniques.

df	Mean square	P > F
Stem-punc	ture technique	
5	4.41	< 0.01
18	0.43	< 0.01
90	0.14	0.38
Infested-s	soil technique	
2	0.17	0.52
18	0.44	0.08
	Stem-punc 5 18 90 Infested-s	Stem-puncture technique 5 4.41 18 0.43 90 0.14 Infested-soil technique 2 2 0.17

Table 3. Pathogenic aggressiveness of six strains of *Pseudomonas solan-acearum*.

Strain	Disease incidence ^z
10	94% ^y a ^x
W2	91% a
8	89% a
1B	89% a
H4	88% a
K60	30% b

^zMean incidence of disease on 19 tomato genotypes and one tomatillo genotype inoculated using the stem-puncture technique.

yWeighted mean derived from transformed data.

^xMean separation ($P \le 0.05$) by LSD.

Table 4. Susceptibility of 19 tomato genotypes and one tomatillo genotype using the stem puncture and infested soil inoculation techniques.

	Disease incidence ^z		
Genotype	Stem puncture	Infested soil	
XPH5675	98% ^y a ^x	92% a	
XPH 5677	98% a	47% a-d	
Island Red	94% ab	23% a-d	
FMX 192	93% a-c	67% a-c	
Venus	92% a–d	91% a	
Tomatillo	91% a–d	73% ab	
GA 219	87% a–e	23% a-d	
GA 1405	85% a–e	73% ab	
IHR 66	83% b-f	42% a-d	
Caraibo	81% b-f	75% ab	
84 BWR	80% b-f	78% ab	
CL 5915-93	79% b–f	62% abc	
GA 1095	78% c-f	45% a-d	
CL 5915-153	78% c-f	84% a	
Capitan	76% d-f	91% a	
GA 1565	75% d-f	51% a-d	
PI 126408	69% ef	23% a-d	
Hawaii 7997	67% f	0 d	
CRA 66	65% f	7% b-d	
Hawaii 7998	63% f	2% cd	

^zDisease incidence expressed as a percentage of the disease incidence on the susceptible cultivars Sunny and Bonny Best. Disease incidence for the stem-puncture technique is a mean of the six strains used in this study. ^yWeighted mean derived from transformed data.

must respond to the presence of *P. solanacearum* after some type of stress has been applied. The response of healthy plants to a bacterial challenge is not considered. The high concentrations of the bacterial suspension used in these techniques (10⁷ to 10⁹ cfu/ml) are not representative of soil populations encountered in the fields (Granada and Sequeira, 1983; Hara and Ono, 1984; Khan et al., 1988). Such high concentrations may further overwhelm the already weakened defense mechanisms of the plant. Therefore, these inoculation techniques, when used as a screening method, do not reflect the level of resistance that may be expressed under field conditions.

Breeding programs in Florida have traditionally relied on field trials to differentiate resistance levels (Locascio et al., 1988; Sonoda and Augustine, 1978). Variable inoculum densities and uncontrolled environmental conditions may severely affect the outcome of field trials (Mew and Ho, 1977; Sonoda, 1978) and lead to variable results. The infested-soil technique was used because it permits the assessment of field resistance by allowing the infection process to take place under natural conditions with realistic doses of naturally produced inoculum. By conducting the technique under controlled conditions, the influence of temperature or moisture stress on disease is minimized. In this study, the high probability value associated with replication (Table 2) is a good indication that greenhouse conditions did not significantly affect results between treatments.

By combining several resistance screening techniques with the analysis of bacterial fatty acid profiles, a standardized method for quantifying diversity of *P. solanacearum* strains and differentiating levels of field resistance was obtained. From this, we conclude that, although *P. solanacearum* strains in the Quincy production region of northern Florida are biochemically diverse, based on fatty acid profiles, there is no significant specific interaction

between strains and the 20 sources of resistance evaluated. Furthermore, the genotypes CRA 66, Hawaii 7998, and especially Hawaii 7997, seem to have a high level of field resistance to bacterial wilt. These observations agree with those by Sonoda et al. (1979) for the southeastern Florida tomato production area, but differ from those of Scott et al. (1993) for the southwestern Florida tomato production area, where H7998 tended to be more susceptible.

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^xMean separation ($P \le 0.05$) by LSD.

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