Evaluation of Virulence of Phakopsora pachyrhizi and P. meibomiae Isolates

M. R. Bonde, S. E. Nester, C. N. Austin, C. L. Stone, and R. D. Frederick, USDA-ARS, Foreign Disease-Weed Science Research Unit, Fort Detrick, MD 21702-5023; G. L. Hartman, USDA-ARS and Dept. of Crop Sciences, University of Illinois, Urbana 61801; and M. R. Miles, USDA-ARS, University of Illinois, Urbana 61801

ABSTRACT

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Asian soybean rust (ASR), caused by Phakopsora pachyrhizi and recently discovered for the first time in continental United States, has been of concern to the U.S. agricultural industry for more than 30 years. Since little soybean rust resistance is known, and resistance is often difficult to detect or quantitate, we initiated a project to develop a better, more quantitative, method. The methodology determined the average numbers and diameters of uredinia in lesions that developed on leaves of inoculated plants 14 days after inoculation. It was used to compare virulence of P. pachyrhizi isolates from Asia and Australia and P. meibomiae from Puerto Rico and Brazil, collected as many as 30 years earlier, with isolates of P. pachyrhizi recently collected from Africa or South America. Susceptible reactions to P. pachyrhizi resulted in tan-colored lesions containing 1 to 14 uredinia varying greatly in size within individual lesions. In contrast, on these same genotypes at the same time of year, resistance to other P. pachyrhizi isolates was typified by 0 to 6 small uredinia in reddish-brown to dark-brown lesions. Using appropriate rust resistant and rust susceptible genotypes as standards, examination of uredinia 14 days after inoculation allowed quantitative comparisons of sporulation capacities, one measure of susceptibility or resistance to soybean rust. The study verified the presence and ability to detect all known major genes for resistance to soybean rust in the original sources of resistance. It demonstrated that soybean lines derived from the original PI sources, and presumed to possess the resistance genes, in actuality may lack the gene or express an intermediate reaction to the rust pathogen. We suggest that a determination of numbers and sizes of uredinia will detect both major gene and partial resistance to soybean rust.

Additional keywords: disease resistance, pathogen virulence, resistance screening, soybean rust histology

Asian soybean rust (ASR), incited by *Phakopsora pachyrhizi* Sydow, has been known in Asia since at least 1902 (9). In

Corresponding author: M. R. Bonde E-mail: morris.bonde@ars.usda.gov

Current address of C. N. Austin: 334 Plant Science Building, Dept. of Plant Pathology, Cornell University, Ithaca, NY 14853.

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1994, P. pachyrhizi was discovered on soybean (Glycine max (L.) Merrill) in Hawaii (11). From 1997 to 2001, P. pachyrhizi was reported for the first times from several central and southern African countries (14). Until 2001, only P. meibomiae, a minor rust pathogen on soybean, had been described in the Caribbean Islands and Central and South America, causing only minor yield losses. However, in 2001 ASR was discovered in Paraguay, the first time in the New World (21), and the following year it was discovered in Brazil (35) and Argentina (25). On 10 November 2004, ASR was reported for the first time to be present in the continental United States (26).

ASR is one of the most economically important soybean diseases in Asia, causing losses up to 80% in some fields (1). Liu (16) estimated the overall loss in Taiwan to be 20 to 30%. Yu et al. (36) reported that even with the use of chemical control in the Peoples Republic of China, yield losses were 15 to 20% in normal years, and 30 to 50% in years with severe rust epidemics. In the 1994 to 1995 growing season, soybean rust caused up to 80% losses in the northern districts of the state of Karnataka, India (24). An estimate of direct loss caused by rust in Brazil in 2003 is US\$2.0 billion (35).

In 1976, Vakili and Bromfield (32) discovered rust on soybean in the Limani Valley, Puerto Rico. Although they believed that the pathogen was *P. pachyrhizi*, they recognized that Puerto Rican isolates were considerably less virulent and produced fewer urediniospores on soybean than isolates from the Eastern Hemisphere.

Rust pathologists became aware that soybean rust actually was caused by at least two species when Ono et al. (22) in 1992 published a comprehensive paper on the morphology of the *Phakopsora* species on legumes. In their paper, which clarified the relationships of rust pathogens on legumes, they referred to the rust pathogen on soybeans in the New World as Phakopsora meibomiae (Arthur) Arthur, which had been known to be present on nonsoybean legumes since 1899 (22). Frederick et al. (5) in 2002 confirmed that two species caused soybean rust when they showed that isolates of P. meibomiae and P. pachyrhizi shared only 80% nucleotide sequence similarity within the ribosomal internal transcribed spacer region, and they developed polymerase chain reaction assays that differentiated the two species (5).

Bromfield (1) and Bromfield and Hartwig (2) described three infection types on soybean accessions inoculated with *P. pachyrhizi*: 1, RB, reddish-brown lesions with few uredinia and meager sporulation; 2, Tan, tan lesions with many uredinia and abundant sporulation; and, 3, Immune, lack of visible infection. All soybean accessions produced an RB reaction when inoculated with the organism from Puerto Rico, thought to be *P. pachyrhizi* and now known to be *P. meibomiae* (1,2).

Bromfield and Hartwig (2), and Hartwig and Bromfield (8) used *P. pachyrhizi* isolates India 73-1, Taiwan 72-1, Australia 72-1, and Philippines 77-1 to characterize rust reaction phenotypes and screen for resistant soybean lines. Accessions PI 200492 and PI 462312 gave immune and resistant (reddish brown hypersensitive) reactions, respectively, to India 73-1, but were susceptible (with tan colored lesions) to Taiwan 72-1 (8). PI 230970 was resistant to all isolates (1,8), but later was found to be susceptible to a new isolate,

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Taiwan 80-2 (1,7). Subsequently, a soybean introduction from central China, PI 459025, was identified as being resistant to Taiwan 80-2 (1,7). Four independently inherited genes for resistance to *P. pachyrhizi* (*Rpp 1-4*) were determined to be present in PI 200492 (*Rpp 1*), PI 230970 (*Rpp 2*), PI 462312 (cv. Ankur) (*Rpp 3*), and PI 459025 (*Rpp 4*), respectively (1,2,7,8).

Screening soybean germ plasm for rust resistance, and developing rust resistant cultivars, has been ongoing in Asia for several decades (3,6,17,18,20,24,27–31). For example, in the People's Republic of China (PRC), cultivars bred for resistance to *P. pachyrhizi* include You dou 84-87, Zan Bian #1, and Liu-dou #1 (3,20). 'September Yellow' has been widely promoted as rust tolerant in Datian county, Fujian province, PRC (30), and 'Baihua Jiuyuehuang' was part of the integrated control strategy for soybean rust in PRC (15).

Initially, the objective of this study was to characterize reaction phenotypes based on symptom expression as described by Bromfield and Hartwig (2) and relate this to pathogen development within plant leaves. It was our hypothesis that numbers of uredinia per lesion and average uredinia diameters are reflections of the growth of the pathogen in host tissue, and as such could be used as quantitative measures of susceptibility or resistance to soybean rust,

similar to measurements of spore producing capacity with wheat rust pathogens (23). During our study, P. pachyrhizi was discovered in South America for the first time, causing alarm throughout the U.S. soybean industry (19). As a result, we extended our study to compare virulence of Asian and Australian isolates collected as much as three decades earlier with newly collected isolates from Africa, South America, or Hawaii. These comparisons should provide insight into whether or not P. pachyrhizi had significantly changed in virulence, and the data from this study will supply valuable information with which to better screen for rust resistance. With the recent spread of P. pachyrhizi to the southeastern United States, it is important that rust resistant cultivars be developed and made available to U.S. growers as soon as possible. Results of studies with U.S. isolates will be included in a future publication.

MATERIALS AND METHODS

Pathogen isolates. Each isolate was a population of urediniospores harvested from numerous pustules on field-collected soybean leaves from the country and year indicated by the respective isolate designation (Table 1). Urediniospores of each isolate of *P. pachyrhizi* were increased on the soybean cultivar Wayne or Williams, and those of *P. meibomiae* on either red

kidney bean (*Phaseolus vulgaris* L.) or lima bean (*Phaseolus lunatus* L.), in isolation cubicles or greenhouses. Subsamples of all isolates were stored in liquid nitrogen (-196°C) refrigerators at the USDA, ARS, Foreign Disease-Weed Science Research Unit (FDWSRU) Biological Safety Level 3 (BSL3-P) Containment Facility (12). Harvesting urediniospores from infected leaves during spore increases was by a vacuum-type harvester (4), beginning 10 to 14 days after inoculation and continuing at weekly intervals.

Plant accessions and propagation. The soybean cultivars Williams, Centennial, and Wayne were used in the study as highly susceptible controls. Also used were the rust resistant accessions Tainung #4 (Nung-shih H 11 × PI 200492), Plant Introduction (PI) accessions 200492 (Komata), 230970, 459025B, and 462312 (Ankur), and the putative rust resistant lines D86-8286 (Rpp 3) (10) and D91-5987 (Rpp 4) (10). Plants were grown two per 10.2-cm-diameter clay pot. During the course of the study, plants were fertilized as needed with 14-14-14 Osmocote (Scotts-Sierra Horticultural Products Co., Marysville, OH).

Inoculation of plants. One day prior to inoculation, urediniospores were removed from liquid nitrogen storage, heat shocked while dry for 5 min at 40°C, and then hydrated for 16 h at 100% relative humidity

Table 1. Reaction types on commercial cultivars, breeding lines, and plant introduction accessions 14 days after inoculation with isolates of *Phakopsora* pachyrhizi or *P. meibomiae*^a

Pathogen Isolate ^b	Williams ^c	Wayne ^c	Komata ^d <i>Rpp 1</i>	230970 <i>Rpp 2</i>	Ankur ^e <i>Rpp 3</i>	459025B ^f <i>Rpp 4</i>	Tainung#4 ^g <i>Rpp1</i> +	D86-8286 ^h <i>Rpp 3</i>	D91-5987 ⁱ <i>Rpp 4</i>
P. pachyrhizi									
Taiwan 72-1	Tan	Tan	Tan	RB	Tan/RB	Inter	Tan	RB	Tan
Taiwan 80-1	Tan	Tan	Tan	RB	Tan	Inter	Tan	RB	Tan
Taiwan 80-2	Tan	Tan	Tan	Tan	Tan	RB	ND^{j}	TAN	Tan
Philippines 77-1	Tan	Tan	Tan	RB	Tan	RB	Inter	TAN	Tan
Australia 79-1	Tan	Tan	Immune	RB	RB	Tan ^k	Immune	TAN	Tan
India 73-1	Tan	Tan	Immune	Inter	RB	RB	Immune	Tan/RB	Tan
Hawaii 94-1	Tan	Tan	Immune	RB	Imm/RB ¹	RB	Immune	Tan/RB	Tan
Thailand 01-1	Tan	Tan	Tan	Inter	Tan	RB	Tan	RB	Tan
Zimbabwe 01-1	Tan	Tan	Tan	Inter	Tan	Inter	Tan	Tan	Tan
South Africa 01-1	Tan	Tan	Tan	RB	RB	Inter	RB	Inter	Tan
Brazil 01-1	Tan	Tan	Tan	Inter	Mixed	RB	Mixed	Tan	Tan
Paraguay 01-2b	Tan	Tan	Tan	RB	Mixed	RB	RB	Tan	Tan
P. meibomiae									
Brazil 82-1	RB	RB	RB	Immune	Immune	RB	Immune	Immune	RB
Puerto Rico 76	RB	RB	RB	RB	RB	RB	RB	RB	RB

^a Soybean plants at the two- to four-trifoliolate leaf stage were atomized with a suspension of 1 to 2×10^4 urediniospores per ml of 0.01% Tween 20 in distilled water and incubated overnight in a dew chamber at 20°C. At 14 days after inoculation, leaflets were examined at ×10 magnification to determine reaction types. Reaction types were: 1, Tan, tan-colored lesions with 1 to 14 uredinia per lesion; 2, RB, reddish brown hypersensitive lesions with 0 to 6 uredinia per lesion; and 3, Immune, no visible symptoms. Tan/RB indicates that some plants produced a Tan reaction and some an RB reaction either within or between experiments. Mixed indicates Tan and RB reactions were evident on the same individual plants. Intermediate (Inter) indicates a reaction type somewhere between Tan and RB in the single experiment (experiment 3) where tested.

^b Pathogen isolates are designated based on country and year of origin.

^c U.S. commercial cultivars.

^d Cultivar Komata (PI 200492) with *Rpp 1* gene for resistance to soybean rust.

^e Cultivar Ankur (PI 462312), grown in India in the 1970s, with the Rpp 3 gene for resistance to soybean rust.

^f Selection from original PI 459025 with *Rpp 4* gene for soybean rust resistance.

^g Cultivar Tainung #4, with the *Rpp 1* gene, and believed to have an additional gene for resistance (18).

^h Breeding line selection reputed to have *Rpp 3* gene for rust resistance.

ⁱ Breeding line selection reputed to have the *Rpp 4* gene for rust resistance. The cross was initially between the cultivar Hardee and PI 459025. An advanced F3 line was crossed with the variety Lamar and advanced to the F5 generation. Results from this study indicate the gene is not present.

 j ND = not done.

^k Although scored as Tan, the low spore producing area per lesion suggests an RB reaction.

¹ In experiment 2, plants reacted with an immune reaction, and in experiment 3 an RB reaction.

over water in a glass desiccator at room temperature. Hydrated urediniospores were suspended in sterile distilled water containing one drop of Tween 20 per 100 ml to a final concentration of 1×10^4 or 2×10^4 per ml. Two and one-half milliliters of the urediniospore suspension was sprayed onto two plants at the two- to fourtrifoliolate-leaf stage in each of one or two 10.2-cm-diameter clay pots per soybean accession. All plants were maintained overnight (approximately 16 h) in dew chambers at 20° C. After the dew treatment, plants were removed from the chambers, allowed to dry, and then placed in a greenhouse maintained at 20 to 25° C for disease development. Experiment 1 was conducted from 19 September to 3 October 1995, ex-

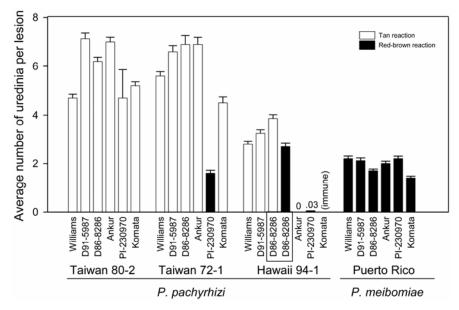


Fig. 1. Average number of uredinia per lesion (n = 100) in experiment 2 on cv. Williams, two reputed rust resistant breeding lines, and each of three PI accessions each containing a reputed singe major gene for rust resistance. Accessions were inoculated separately with isolates of *Phakopsora pachythizi* from Taiwan or Hawaii, or *P. meibomiae* from Puerto Rico. PI 200492, PI 230970, and PI 462312 contain the *Rpp 1*, *Rpp 2*, and *Rpp 3* genes, respectively, for rust resistance. Bracket signifies that not all plants of the accession inoculated with isolate Hawaii 94-1 were scored as having the same reaction type. Hawaii 94-1 produced a Tan reaction on on D86-8286 plant and an RB reaction on three other plants. Error bars represent standard errors of the means.

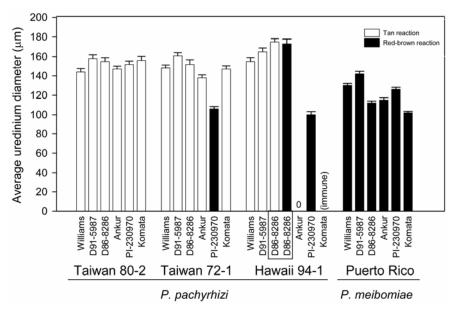


Fig. 2. Average uredinium diameter (μ m) (n = 100) for cv. Williams, two reputed rust resistant breeding lines, and each of three rust resistant PI accessions containing single genes for rust resistance inoculated separately with isolates of *Phakopsora pachyrhizi* from Taiwan or Hawaii, or *P. meibomiae* from Puerto Rico. PI 200492, PI 230970, and PI 462312 contain the *Rpp 1*, *Rpp 2*, and *Rpp 3* genes, respectively, for rust resistance. Bracket signifies that not all plants of the accession inoculated with isolate Hawaii 94-1 were scored as having the same reaction type. Hawaii 94-1 produced a Tan reaction on one D86-8286 plant and an RB reaction on three other plants. Error bars represent standard errors of the means.

periment 2 from 3 to 17 April 1996, and experiment 3 from 21 February to 8 March 2002. The second experiment was conducted at a time when environmental conditions apparently were more conducive for disease development, based on subsequent density of lesions, than the other times.

Determination of infection types. In each experiment, 14 days after inoculation, the two or four test plants per soybean accession/pathogen isolate were examined to determine symptoms at ×1 and ×15 magnification in the greenhouse and scored using the criteria of K. R. Bromfield (1) as having a Tan, RB, or Immune reaction phenotype. Tan-colored lesions (designated Tan reaction), the color primarily due to the large numbers of tan urediniospores, indicated a highly susceptible reaction, and reddishbrown lesions (designated RB reaction), a form of resistance (1). An immune reaction (designated Immune) indicated a lack of obvious symptoms (1). In some instances, an intermediate, weakly hypersensitive response with moderate sporulation (designated Intermediate) was produced. If two reaction types were expressed on the same plant, generally on the same leaflet, the reaction was designated as Mixed and assumed to be the result of a mixture of races in the inoculum.

Histological comparisons. At 14 days after inoculation, leaflets from the test plants of each accession inoculated with P. pachyrhizi or P. meibomiae, and previously examined for reaction phenotype, were excised from the plants, and leaf pieces approximately 1×1 cm placed into a solution of absolute ethanol/acetic acid (3:1, vol/vol) for at least 24 h to fix and remove plant pigments from the tissue. The leaf pieces were transferred to lactophenol for 24 h to further clear the tissue and make translucent, and stained in 0.1% cotton blue in lactophenol for 24 h. The stained leaf pieces were examined microscopically at ×100 and ×400 and measurements of uredinia made with an ocular micrometer as described below.

Experimental design. Leaf samples for each soybean accession/rust isolate were combined within an experiment. For experiment 1 (data not presented), in which specific parameters were tested to determine their potential for detecting resistance, all combinations of seven rust isolates and 10 soybean accessions were examined 14 days after inoculation. Data included uredinia diameter, number of uredinia per lesion, number of uredinia per mm² lesion area, and reaction type (i.e., Tan, RB, or Immune). Sample size for each parameter, other than reaction type, was 100, and measurements were obtained from randomly selected leaf pieces collected from the two or four inoculated plants, depending upon the number of plants in a particular experiment.

In experiment 2 (Table 1, Figs. 1 and 2), four rust isolates were inoculated onto six

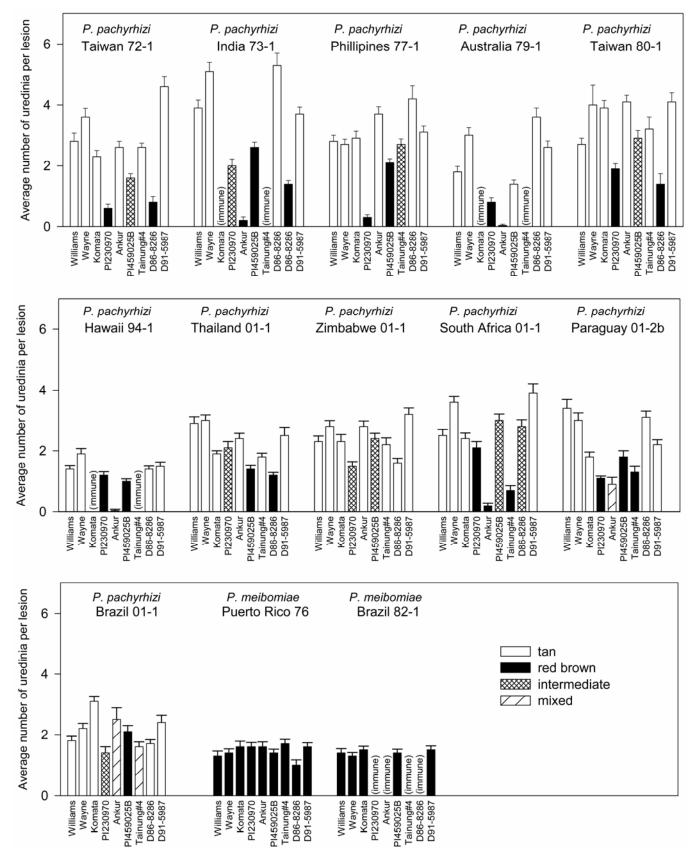


Fig. 3. Average number of uredinia per lesion (n = 25) 14 days after inoculation with 11 separate isolates of *Phakopsora pachyrhizi* or two of *P. meibomiae*. Cultivars Williams and Wayne are representative U.S. cultivars highly susceptible to Asian Soybean Rust. The Japanese cultivar Komata (PI 200492) and the Taiwanese cultivar Tainung #4 (PI 368039) contain the *Rpp 1* gene for soybean rust resistance, PI 230970 the *Rpp 2* gene, and Indian cultivar Ankur (PI 462312) the *Rpp 3* gene. Breeding line D86-8286 (10) is reputed to contain *Rpp 3* and D91-5987 *Rpp 4* (10). Note that the latter had a large number of ured-inia per lesion and always produced a Tan reaction.

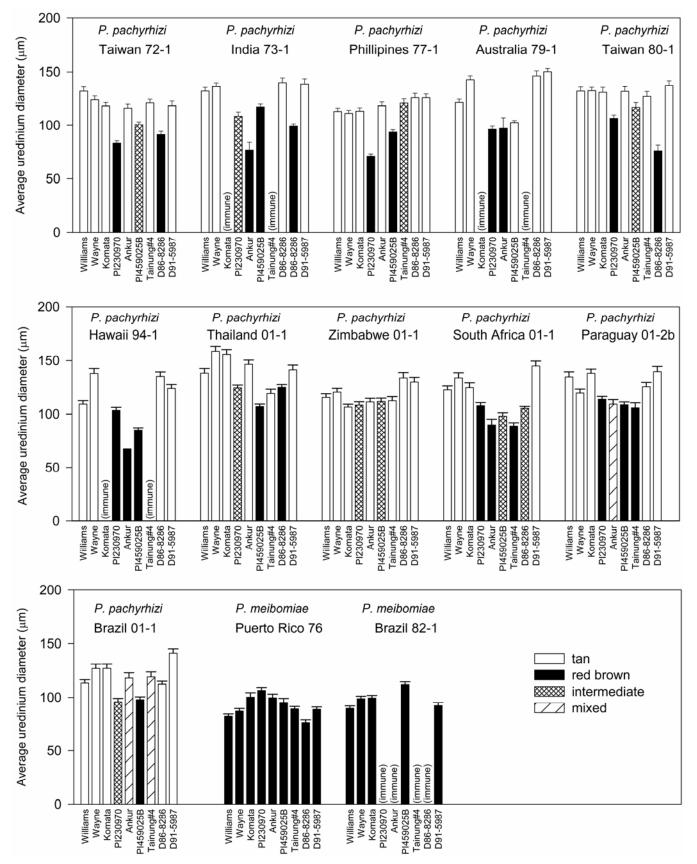


Fig. 4. Average uredinium diameter (n = 50) at 14 days after inoculating 11 isolates of *Phakopsora pachyrhizi* and two of *P. meibomiae* separately onto nine soybean accessions. Cultivars Williams and Wayne are representative U.S. cultivars highly susceptible to Asian Soybean Rust. The Japanese cultivar Komata (PI 200492) and the Taiwanese cultivar Tainung #4 (PI 368039) contain the *Rpp 1* gene for soybean rust resistance, PI 230970 the *Rpp 2* gene, and Indian cultivar Ankur (PI 462312) the *Rpp 3* gene. Breeding line D86-8286 (10) is reputed to contain *Rpp 3* and D91-5987 *Rpp 4* (10). Note that D91-5987 had relatively large uredinia diameters and lesions were Tan, regardless of pathogen isolate used for inoculum.

soybean accessions. Data included reaction type, uredinia diameter, and number of uredinia per lesion; sample size for the latter two each was 100.

In experiment 3 (Table 1, Figs. 3 and 4), 13 rust isolates, including five not available for use in previous experiments, were inoculated separately onto each of nine soybean accessions. Besides reaction type, data included uredinia diameter and number of uredinia per lesion. Sample size was 25 lesions to determine number of uredinia per lesion, and 50 uredinia to determine average uredinia diameter for each pathogen isolate/soybean accession.

In each experiment, the means and standard errors of the means were calculated for each parameter.

RESULTS

Lesion size. Lesion diameters, determined in experiment 1 for 70 pathogen isolate/soybean accession combinations, were not correlated with resistance or susceptibility. Often, RB reactions produced the largest lesions. Tan lesions frequently were difficult to measure. As a result, this parameter was not included in subsequent experiments.

Reaction phenotypes. Reaction phenotypes (Fig. 5) in our study generally were consistent for each of the PI accessions inoculated with the specific isolates used previously to describe the four known *Rpp* genes (1,2,7,8; Table 1). Furthermore, P. pachyrhizi isolates Taiwan 72-1, India 73-1, Australia 79-1, and Taiwan 80-2; and P. meibomiae isolate Puerto Rico 76 produced for us essentially the same reaction on each of the four PI accessions as they did for Bromfield and Hartwig (2), Hartwig (7), and Hartwig and Bromfield (8). The only exceptions were in a few instances in which we interpreted a reaction to be intermediate (between Tan and RB), instead of RB as reported previously by Hartwig and Bromfield (8).

Our results with soybean line D86-8286, however, were inconsistent among experiments, and sometimes within an experiment (Table 1). For example, in experiment 3, one D86-8286 plant inoculated with India 73-1 produced obvious Tan lesions and one clearly RB lesions (Figs. 3 and 4). Likewise, in experiment 2, one D86-8286 plant produced a Tan reaction while three plants gave an RB reaction when inoculated with Hawaii 94-1 (Figs. 1 and 2). It was difficult to distinguish between Tan and RB on these plants, and the histological measurements indicated little or no difference among plants.

Although reaction phenotypes usually grouped into either Tan or RB, each pathogen isolate produced its own characteristic Tan or RB reaction. For example, Hawaii 94-1 in experiment 2 inoculated onto susceptible soybean accessions, lacking major genes for rust resistance, produced a Tan reaction with an average of 2.7 to 3.8 uredinia per lesion (Fig. 1), and the uredinia varied greatly in size within individual lesions (e.g., Fig. 6A). Isolate Hawaii 94-1 produced uredinia with mean diameters of 155, 165, and 176 µm on Williams, D91-5987, and D86-8286, respectively (Fig. 2).

Hawaii 94-1 on PI 230970 (with the *Rpp* 2 gene) produced a strong RB reaction (Fig. 6C). No uredinia were evident in at least 97% of the lesions, and those that were present were small, averaging only 100 μ m in diameter (Fig. 2). Uredinia often stained poorly with cotton blue (Fig. 6C), and as expected, sporulation was sparse.

Soybean accessions PI 200492 (Komata) and PI 462312 (Ankur), containing the *Rpp 1* and *Rpp 3* genes, respectively, in experiment 2 were immune to Hawaii 94-1 (Figs. 1 and 2). However, in experiment 3, PI 462312 inoculated with Hawaii 94-1 produced an RB reaction with few uredinia (Fig. 3).

Relationship between number of uredinia and uredinia diameters to Tan, RB, and Immune reactions. Tan reactions usually had higher average numbers of uredinia per lesion (Figs. 1 and 3) and larger average uredinia diameters (Figs. 2 and 4) than typical RB reactions. In experiment 2, conducted when symptoms were well expressed and differences between resistant and susceptible reactions were great, Tan lesions had 1 to 14 (averages varied from 2.8 to 7.0) uredinia per lesion, and RB lesions, excluding those on D86-8286, had 0 to 6 (averages 0.0 to 1.7) uredinia per lesion (Fig. 1). Many RB lesions were devoid of uredinia. Although average diameters of uredinia usually were larger in Tan reactions than in RB reactions, uredinia diameters did not separate pathogen isolate/soybean accession responses to the degree as numbers of uredinia per lesion.

Comparison of P. pachyrhizi isolates. In contrast to older isolates collected in Asia and Australia, isolates of P. pachyrhizi acquired in 2001 from Africa or South America never caused an Immune reaction on any of the soybean accessions tested. When RB reactions were observed, the number of uredinia per lesion usually was not greatly less than in Tan reactions on the same soybean accession (Fig. 3). In a few instances, an Intermediate reaction was observed in which the reaction was neither RB nor Tan, but intermediate between RB and Tan (Figs. 3 and 4). Based on the lower frequency of RB reactions and complete absence of Immune reactions on soybean accessions containing the known major genes for rust resistance, the newer isolates from Africa and South America were more virulent and/or aggressive than the older isolates. The most virulent was Zimbabwe 01-1 (Table 1, Figs. 3 and 4), in which no RB or Immune reactions were observed.

Reactions of soybean to *P. meibomiae*. *P. meibomiae* isolates Puerto Rico 76 and Brazil 82-1 produced red-brown to darkbrown lesions with few uredinia, or an immune reaction (Table 1). Even during the most conducive time of year for disease, lesions had an average of only 1.4 to 2.2 uredinia per lesion (Fig. 1). Uredinia produced by these two *P. meibomiae* isolates were small, with average diameters ranging from 101 to 142 μ m when inoculated onto the various soybean accessions (Fig. 2 and 6E).

DISCUSSION

Knowledge of how a pathogen develops within a plant can be very valuable when evaluating disease resistance. In this paper, we examined the relationships of numbers of uredinia per lesion and average uredinia diameters that developed during expression of the Tan, RB, and Immune reaction phenotypes previously described by Bromfield (1) and Bromfield and Hartwig (2). It was particularly significant that each soybean accession used by them behaved the same, or nearly the same, for us. Because we used the same pathogen isolates, stored in liquid nitrogen refrigerators, as Bromfield (1) and Bromfield and Hartwig (2), we were able to make conclusions otherwise not possible.

It became apparent in our study that the RB reaction (2) actually is a spectrum of reaction types. RB lesions, in addition to being reddish brown, were sometimes dark chocolate brown, or even light brown. Whereas RB lesions generally had sparse sporulation, there were a few instances in which RB lesions sporulated prolifically. In the susceptible reactions, Tan lesions also varied. Fully susceptible Tan lesions usually produced more numerous and larger uredinia than RB lesions. The greatest differences between RB and Tan lesions occurred during March to mid-November. This was observed in other soybean rust studies also being conducted in our greenhouses (M. R. Miles, G. L. Hartman, R. D. Frederick, and M. R. Bonde, unpublished data).

Whereas the reaction phenotypes on the original sources were easy to score in this study, it was very difficult to ascertain whether some lines derived from them had RB or Tan lesions. For example, in experiment 2, not only were the D86-8286 plants inoculated with Hawaii 94-1 difficult to assign as Tan or RB, the numbers of uredinia per lesion and average uredinia diameters within the lesions were very similar (Figs. 1 and 2). This supports our belief that soybean lines derived from the original sources of *Rpp 1-4* sometimes produce intermediate reactions.

Average number of uredinia per lesion (Figs. 1 and 3) was a better indicator of reaction phenotype than average uredinia diameter; however, we believe that a combination of the two parameters might be a better and more prudent measure of resistance or susceptibility. This concept is

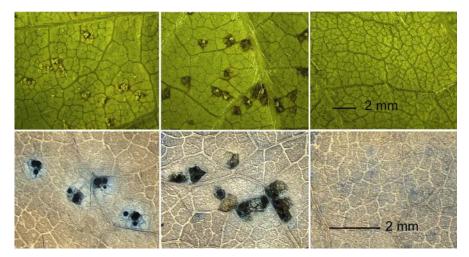


Fig. 5. Typical symptoms of Tan, RB, and Immune reactions 15 days after inoculation (top row); and leaf tissue after removal of pigments and staining with cotton blue (bottom row). Left is Taiwan 80-1 on PI 200492 (Tan reaction), middle Taiwan 80-1 on PI 230970 (RB reaction), and right India 73-1 on PI 200492 (Immune reaction). PI 200492 and PI 230970 contain the *Rpp 1* and *Rpp 2* genes, respectively, for rust resistance. The top row of illustrations is the lower leaf surface of the fresh leaf, and the bottom row the lower leaf surface of the same combination of pathogen isolate/soybean accession fixed, cleared, and stained with cotton blue.

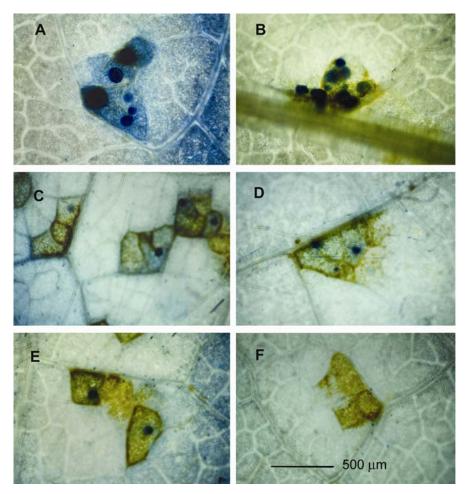


Fig. 6. Typical reactions within leaves of soybean cultivars or lines exhibiting susceptible or resistant phenotypes following fixing, clearing, and staining with cotton blue. **A**, Hawaii 94-1, Tan reaction on cultivar Williams; **B**, Taiwan 80-2, Tan reaction on PI 230970; **C**, Hawaii 94-1, RB reaction on PI 230970; **D**, Taiwan 72-1, RB reaction on D86-8286; **E**, Puerto Rico 76, RB reaction on PI 230970; and **F**, Brazil 82-1, RB reaction on PI 230970. Note that the susceptible reactions (**A and B**) resulted in both large and small uredinia. Resistant RB reactions were typified by either a few small uredinia or complete lack of uredinia.

being pursued in more detail in our current research.

Leaf pieces collected from greenhousegrown soybean plants 14 days after inoculation and stained with cotton blue produced results that were relatively easy to interpret, particularly when resistant and susceptible standards were included within the same experiment. For example, under highly favorable conditions for disease, a susceptible reaction was typified by a mean of 3 to 7 uredinia per lesion, and uredinia varied greatly in size within individual lesions (Fig. 6A and B). In contrast, in the same experiment, resistance was typified by a mean of only 0 to 2 small uredinia per lesion (Fig. 6C and D). A particularly useful criterion was that highly susceptible reactions had a wide range of uredinia sizes within the same lesion, whereas resistant reactions had only small uredinia (Fig. 6C and D), or in some instances no uredinia.

We examined inoculated soybean plants at 14 days after inoculation. It is possible that determination of uredinia numbers and diameters at times other than 14 days might have given somewhat different results, but it is unlikely that trends or main conclusions would differ. Numbers and sizes of uredinia are quantitative measurements, and as such probably would detect partial resistance (23). This hypothesis is being pursued in our research program.

Although potentially very valuable, microscopic examination of leaf tissue should not, as yet, replace examination of intact plants in a rust resistance-screening program. In our rust-screening program at Fort Detrick, we are now using histological measurements to support our observations in the greenhouse. Smaller and fewer uredinia are manifestations of at least one form of disease resistance. Although we did not critically compare spore production, observations in our greenhouse suggest, and logic dictates, that decreases in numbers and sizes of uredinia can greatly decrease spore production.

The differences in numbers and sizes of uredinia in a susceptible, typically Tan, reaction, compared with a resistant, RB, reaction can be rationalized. After initial pathogen establishment in a susceptible reaction, the hyphae rapidly spread within the leaf tissue, producing new uredinia over an extended time period. This progression of pathogen development results in uredinia varying widely in maturities and, hence, size within individual lesions. Large uredinia often are present at the center, surrounded by smaller uredinia, sometimes in concentric circles. In contrast, pathogen development is restricted in a resistant reaction, resulting in fewer and smaller uredinia (e.g., Fig. 6C and D), and in some instances no uredinia. Indeed, resistance to soybean rust often can be detected merely by a quick visual examination of the stained leaf tissue.

Soybean cultivars that display extreme reductions in the number and size of uredinia following inoculation with specific isolates of *P. pachyrhizi* probably result from the presence of a major resistance gene. More subtle differences may result from partial resistance, and detection likely will require more critical measurements. Although not part of this study, resistance to soybean rust could also be a result of reduced numbers of pathogen penetrations, resulting in fewer points of initial establishment in the host, leading to fewer lesions.

The Rpp 1-4 genes are believed to be race specific (1,2,7,8). In Taiwan, three physiological races of P. pachyrhizi were identified by inoculating five differential soybean cultivars with 50 singleurediniospore isolates (34). In another study using 10 differential hosts, 42 rust isolates were differentiated into nine races (34). The investigators suggested that the predominate races in Taiwan possess multiple genes for virulence (34). Other rust isolates were reported to cause disease on all known suspected sources of specific resistance (31). Tschanz et al. (31) pointed out that the occurrence of multiple virulence genes in P. pachyrhizi was not what one would expect since no soybean lines were known to have more than one specific resistance gene for soybean rust. The presence of multiple virulence genes in P. pachyrhizi could make disease control strategies difficult (6).

Historically, genes conferring specific resistance to soybean rust rapidly became ineffective. For example, soybean cv. Komata, which contains the *Rpp 1* gene, was found to be resistant in germ plasm evaluations in Australia in 1961 to 1963; however, by 1966, Tan-type lesions (susceptible reaction phenotype) were observed in field trials. By the mid 1970s, cv. Komata was considered no longer useful as a source of rust resistance (13). Similarly, soybean cv. Ankur, which contains the *Rpp 3* gene resistant in India in the early 1970s (29), was totally susceptible to rust by the 1980s (1).

Results from our study indicate races may also exist within *P. meibomiae*. Whereas Puerto Rico 76 infected all soybean accessions, producing an RB reaction, isolate Brazil 82-1 didn't produce symptoms (Immune reaction phenotype) on accessions with either *Rpp 1* or *Rpp 2* (Table 1). This indicates that these two *Rpp* genes also recognize avirulence genes in *P. meibomiae*, which might suggest that these avirulence genes either have coevolved or have been transferred between these two *Phakopsora* species.

There is evidence from our study that the new isolates are somewhat more virulent, and/or more aggressive, than the older isolates. In only two instances in experiment 3 were the spore producing areas per lesion, calculated by multiplying average number of uredinia per lesion × average cross-sectional area per uredinium, less than $5 \times 10^3 \,\mu\text{m}^2$. This equated to only 4% of the total number of rust reactions producing low-level sporulation. In contrast, in 14 instances with old isolates, equal to 25%, the spore producing area per lesion was less than $5 \times 10^3 \,\mu\text{m}$.

With the recent discovery of P. pachyrhizi in the continental United States, it is important that efforts to screen for rust resistance be increased, and newly discovered resistance genes be incorporated into U.S. cultivars as quickly as possible. Since single gene, specific resistance appears to be of limited value, horizontal or ratereducing resistance is desirable. Evidence for such resistance in soybean exists (31,33). Tolerance to soybean rust also has been identified (6,27), and this trait should be incorporated into elite soybean cultivars in each of the maturity groups grown in the United States. Determining the number of uredinia per lesion and average uredinia diameters provides a direct and quantitative method to measure levels of rust resistance, and should be useful for identifying both specific and rate-reducing resistance. With modification, the methodology could be made more rapid, thus accommodating the testing of large numbers of soybean accessions.

This study confirms the presence of the Rpp genes and our ability to recognize reaction phenotypes in the original PIs. Evidence from this study suggests that D91-5987, derived from a cross between PI 459025, containing the Rpp 4 rust resistance gene, and the cultivar Hardee, lacks resistance to ASR. Other soybean lines, presumably containing rust resistance genes, in the past also gave indications of being either intermediate in reaction or completely susceptible. We found intermediate reactions difficult to score based on the current RB/Tan designations, and therefore believe that in many instances reactions must be quantitatively measured. The methodology described here, using histology, can be used to quantitate a resistance reaction and assist in correctly identifying rust resistance during development of resistant varieties. Histological measurements potentially will detect both specific and partial resistance.

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