# Inheritance of Resistance in Soybeans to Physiologic Races 5, 6, 7, 8, and 9 of *Phytophthora megasperma* var. *sojae*

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#### ABSTRACT

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The  $F_2$  populations from nine soybean crosses were tested with physiologic races 5, 6, 7, 8, and 9 of *Phytophthora megasperma* var. *sojae*. The gene *Rps*<sup>a</sup> in the soybean cultivar Mukden, which gives resistance to races 1 and 2 and susceptibility to races 3 and 4, gave susceptibility to races 5, 6, 7, 8, and 9. The gene *Rps*<sup>b</sup>, which conditions resistance to races 1, 3, and 4 but susceptibility to race 2 in PI 84637, gave resistance to races 5, 6, 7, 8, and 9. The gene *Rps*<sup>c</sup> in PI 54615-1, which gives resistance to races 1, 2, and

Additional key words: soybean disease, soybean genetics.

Recently, Mueller et al (3) reported results of a study of the inheritance of resistance to physiologic races 1, 2, 3, and 4 of *Phytophthora megasperma* Drechs. var. *sojae* A. A. Hildeb. in four soybean cultivars. The gene  $Rps^a$  in Mukden gave resistance to races 1 and 2. The gene  $Rps^b$  in PI 84637 gave resistance to races 1, 3, and 4. The gene  $Rps^c$  in PI 54615-1 gave resistance to races 1, 2, and 3. The gene  $Rps^a$ ,  $Rps^b$ , and  $Rps^c$  were at the same locus, whereas Rps3 was at a different locus and segregated independently of the other three genes. While that work was in progress, Haas and Buzzell (1) reported physiologic races 5 and 6 of the fungus and Laviolette and Athow (2) reported races 7, 8, and 9. The results reported here extend the knowledge of the genetics of resistance in the above four soybean cultivars to the five new physiologic races.

#### **MATERIALS AND METHODS**

Soybean cultivars Harosoy and Mukden, and plant introductions PI 54615-1, PI 84637, and PI 86972-1, and F<sub>2</sub> populations from crosses between them, were inoculated with races 5, 6, 7, 8, and 9 of P. megasperma var. sojae. Inoculum was prepared by growing the isolates of the physiologic races an oatmeal agar in petri dishes for 2-3 wk at 24 C. The same isolate of each race was used throughout the study. Inoculations were made by the hypocotyl method which consists of inserting a  $2 \times 2$  - mm piece of mycelium into a longitudinal slit in the hypocotyl and covering the wound with petrolatum to prevent desiccation of the inoculum and host tissues. Ten-day-old F<sub>2</sub> seedlings of each cross and appropriate parental checks were inoculated with each race and grown in the greenhouse at 24-27 C. Six days after inoculation the seedlings were classified as resistant (no external symptoms) or susceptible (dead). The data were analyzed by the chi-square test for goodness of fit.

3 and susceptibility to race 4, gave susceptibility to race 5 and resistance to races 6, 7, 8, and 9. These three genes are located at the same locus. The independent gene Rps3 in PI 86972-1, which conditions resistance to races 1, 2, 3, and 4, gave resistance to races 5, 8, and 9 but susceptibility to races 6 and 7. Lines with the genotypes  $Rps^{b} Rps^{b} Rps3 Rps3$  and  $Rps^{c} Rps^{c} Rps3 Rps3$  have been selected which are resistant to the nine reported physiologic races of the fungus.

## **RESULTS AND DISCUSSION**

The reactions of the parental cultivars to the nine physiologic races are shown in Table 1. Harosoy was susceptible to all nine races, Mukden was resistant to only races 1 and 2, PI 84637 was resistant to all but race 2, PI 54615-1 was susceptible to only races 4 and 5, and PI 86972-1 was resistant to all but races 6 and 7.

The segregation of the  $F_2$  populations of the nine soybean crosses to physiologic races 5, 6, 7, 8, and 9 is given in Table 2. The data from the Harosoy × Mukden cross are not included because both parents and the  $F_2$  population were susceptible to the five races. The  $F_2$  populations from the crosses of Harosoy or Mukden with PI 54615-1, PI 84637, and PI 86972-1 each segregated in a ratio of 3 resistant: 1 susceptible to those races to which the three PI's were resistant. When inoculated with races to which both parents were susceptible (race 5 with PI 54615-1, and races 6 and 7 with PI 86972-1), the  $F_2$  populations were uniformily susceptible except for a few apparent escapes (Table 2). These data indicate that the gene  $Rps^b$ in PI 84637 which was reported by Mueller et al (3) to confer resistance to races 1, 3, and 4, also gives resistance to races 5, 6, 7, 8, and 9. The gene  $Rps^c$  in PI 54615-1 reported to control resistance to races 1, 2, and 3, also controls resistance to races 6, 7, 8, and 9. The

TABLE 1. Reaction of parental soybean cultivars or plant introduction lines to physiologic races 1 through 9 of *Phytophthora megasperma* var. *sojae* 

| Cultivar   | Host reaction <sup>a</sup> to physiologic races |   |   |   |   |   |   |   |   |  |
|------------|---|---|---|---|---|---|---|---|---|--|
|            | 1   | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 |  |
| Harosoy    | S   | S | S | S | S | S | S | S | S |  |
| Mukden     | R   | R | S | S | S | S | S | S | S |  |
| PI 54615-1 | R   | R | R | S | S | R | R | R | R |  |
| PI 84637   | R   | S | R | R | R | R | R | R | R |  |
| PI 86972-1 | R   | R | R | R | R | S | S | R | R |  |

<sup>a</sup>Abbreviations: R = resistant, and S = susceptible.

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TABLE 2. Segregation of the F2 populations of nine soybean crosses to physiologic races 5, 6, 7, 8, and 9 of Phytophthora megasperma var. sojae

|                             | Plants resistant (R) or susceptible (S) to race |         |         |         |         |         |                  |         |                  |         |  |
|-----------------------------|---|---------|---------|---------|---------|---------|------------------|---------|------------------|---------|--|
| Parentage                   | 5   |         | 6       |         | 7       |         | 8                |         | 9                |         |  |
|                             | (No. R)   | (No. S) | (No. R) | (No. S) | (No. K) | (No. S) | (No. R)          | (No. S) | (No. R)          | (No. S) |  |
| Harosoy $\times$ PI 84637   | 159   | 50      | 150     | 55      | 119     | 43      | 130              | 38      | 152              | 36      |  |
| Harosoy $\times$ PI 54615-1 | 0   | 208     | 182     | 73      | 91      | 29      | 115              | 42      | 102              | 38      |  |
| Harosoy × PI 86972-1        | 164   | 58      | 1       | 203     | 0       | 170     | 159              | 57      | 127              | 50      |  |
| Mukden $\times$ PI 54615-1  | 1   | 222     | 105     | 27      | 122     | 37      | 91               | 27      | 95               | 35      |  |
| PI 54615-1 × PI 84637       | 157   | 56      | 203     | 0       | 150     | 0       | 127              | 0       | 114              | 0       |  |
| Mukden × PI 84637           | 160   | 50      | 155     | 44      | 151     | 47      | 148              | 53      | 158              | 52      |  |
| Mukden $\times$ PI 86972-1  | 156   | 44      | 4       | 216     | 0       | 194     | 122              | 47      | 151              | 47      |  |
| PI 86972-1 × PI 54615-1     | 157   | 56      | 156     | 50      | 68      | 25      | 146 <sup>a</sup> | 10      | 101 <sup>a</sup> | 8       |  |
| PI 86972-1 × PI 84637       | 194 <sup>a</sup>                                | 10      | 163     | 47      | 133     | 49      | 162 <sup>a</sup> | 8       | 150 <sup>a</sup> | 12      |  |

<sup>a</sup>All segregation was in a ratio of 3 resistant: 1 susceptible (P = > 0.05) except these which segregated in a ratio of 15 resistant: 1 susceptible (P = > 0.30).

gene Rps3 in PI 86972-1, reported to give resistance to races 1, 2, 3, and 4, also gives resistance to races 5, 8, and 9.

The F<sub>2</sub> population from the cross PI 54615-1 × PI 84637 was resistant to races 6, 7, 8, and 9, to which both parents were resistant, and segregated in a ratio of 3 resistant:1 susceptible to race 5, to which PI 84637 was resistant and PI 54615-1 was susceptible. This indicates that  $Rps^b$  and  $Rps^c$  are allelomorphic for resistance to races 6, 7, 8, and 9, in addition to races 1 and 3 previously reported (3).

The  $F_2$  populations from crosses of PI 54615-1 and PI 84637 with PI 86972-1 each segregated in a ratio of 15 resistant: 1 susceptible to those races to which both parents were resistance. The  $F_2$ populations from these crosses segregated 3 resistant:1 susceptible to those races to which only one parent was susceptible. These data confirm that the gene *Rps3* in PI 86972-1 which controls resistance to races 5, 8, and 9, in addition to races 1, 2, 3, and 4 previously reported (3), is not at the same locus as the allelomorphic series *Rps<sup>a</sup> Rps<sup>b</sup> Rps<sup>c</sup>* represented by the soybean cultivars Mukden, PI 84637, and PI 54615-1, respectively.

The gene  $Rps^a$  gives resistance to only races 1 and 2, but the resistance conferred by  $Rps^b$ ,  $Rps^c$ , and Rps3 has been extended as

follows:  $Rps^{b}$  gives resistance to races 1, 3, 4, 5, 6, 7, 8, and 9;  $Rps^{c}$  gives resistance to races 1, 2, 3, 6, 7, 8, and 9; and Rps3 gives resistance to races 1, 2, 3, 4, 5, 8, and 9. The significance of these findings is the selection of lines with the genotypes  $Rps^{b} Rps^{b} Rps3$  Rps3 and  $Rps^{c} Rps^{c} Rps3 Rps3$  which are resistant to the nine reported physiologic races of the fungus. Lines with the former genotype have two genes for resistance to all but races 2, 6, and 7, whereas lines with the latter genotype have two genes for resistance to all but races 4, 5, 6, and 7.

### LITERATURE CITED

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