Genetic Linkage Between Mex2, a Specific Resistance Gene to Anthracnose, and Anp, a Gene Involved in Pod Anthocyanin Accumulation in Bean

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ABSTRACT

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Breeding tests in bean (*Phaseolus vulgaris*) with Mex2, a single dominant resistance gene to bean anthracnose, suggested a linkage between this gene and pod anthocyanin accumulation. The cross between an inbred, purple-pod, anthracnose-resistant cultivar with Mex2 (AFN) and an inbred, green-pod, susceptible cultivar (La Victoire) was performed to study the segregation of the two characters and to recover green pod progeny resistant to Mex2, because purple pods are commercially unacceptable. Pod anthocyanin accumulation proved, at least in this cross, to be under the control of a single, dominant Mendelian gene, which has been designated Anp (anthocyanins in the pod). Among 548 plants of F_2 progenies, 13 recombinant phenotypes were identified. The genetic distance between Mex2 and Anp was estimated by maximum likelihood analysis to be 2.3 ± 1.3 cM (95% CI).

In bean (Phaseolus vulgaris L.), resistance to anthracnose caused by Colletotrichum lindemuthianum (Sacc. & Magnus) Lams.-Scrib. is controlled by a series of simple dominant genes (11). In Europe, because of limited pathogen variability, a single resistance gene (ARE) introduced into most inbred lines since 1960 is still effective against this disease (1). Other single dominant genes for specific resistance have been characterized in Latin America, protecting against Colletotrichum races present there (11). Compared to the ARE gene, which provides resistance against the socalled "ancestral" races of Colletotrichum $(\alpha, \beta, \gamma, \delta, \text{ and } \epsilon \text{ types})$ (3), these genes provide protection against additional races that normally attack ARE, such as α Brazil 54, "Ebnett" strain of kappa, Yotta, and λ mutant (6). Breeding programs are currently underway to introduce some of these genes into bean in combination with the ARE gene to increase protection against possible introductions of new Colletotrichum races (6). Bannerot et al (2) described a new resistance gene, Mex2, which provides very efficient resistance against a wide range of Colletotrichum races, and incorporated it into the P. vulgaris cultivar AFN. The combination of ARE and Mex2 resistances should provide very effective control against anthracnose, at least in Europe.

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During the breeding program of AFN, cosegregation of the Mex2 gene and the accumulation of anthocyanin pigments in the pod has been observed. Purple color, attributable to anthocyanins, has often been correlated with, or involved in, disease resistance in plants (4,7,8, 13,17). However, in many cases, resistance and pigment accumulation have been dissociated by genetic analysis. Linkages between genes involved in specific resistance and genes for anthocyanin metabolism have been reported for several host-pathogen interactions, such as tobacco-TMV (Tm2 gene) (12) and bean-BCMV (*I* gene) (15).

In this paper, we report a segregation analysis of *Mex2* and the purple pod phenotype. The results show that anthocyanin accumulation in the pod is determined by a single locus (*Anp*) and that *Anp* is linked but not identical to *Mex2*.

MATERIALS AND METHODS

Plant material. The inbred susceptible (La Victoire) and resistant (AFN) P. vulgaris genotypes were obtained from H. Bannerot and G. Fouilloux (INRA, Versailles, France). Upon exposure to light, La Victoire pods remain green, whereas under the same conditions, AFN pods show a purple-mottled phenotype. On the other hand, both La Victoire and AFN display anthocyanin pigmentation in different plant organs—flowers, cotyledons, and hypocotyls are violet, whereas seed coats are black in La Victoire and mottled black-gray in AFN. Plants were grown in a greenhouse under controlled conditions (8-hr dark/16-hr light photoperiod under fluorescent bulbs [250 μ E·m⁻²·s⁻¹ at the plant level], temperature at 24 C, and relative humidity

at 70%) in pots filled with vermiculite and watered with the nutrient solution of de Bildering and Lourtioux (5).

Five independent crosses between La Victoire (green pods, susceptible) and AFN (purple pods, resistant) were performed. Because of the autogamous behavior of P. vulgaris plants (selfing rate >98%), all crosses were performed manually. Young flowers in which the ovules are mature but pollen is not were handpollinated by delicately opening floral buds and fertilizing them with pollen obtained from mature flowers. La Victoire was used as a female parent to avoid artificial data in the progeny as a result of selfing. Genetic analysis was performed on F₁, F₂, and F₃ progenies. The purple and resistant (Mex2) F₁ plants were allowed to self to obtain F₂ pro-

Fungal material. C. lindemuthianum (race α Brazil 54) was supplied by F. Legrand and J. Tailler (Laboratoire Cryptogamie, Orsay, France), and was grown on malt (20 g/L) agar (20 g/L) plates at 23 C in the darkness. Conidia were obtained from 10-day-old cultures by gentle scraping with a spatula, suspended in sterile water, filtered through a sintered glass filter (20–40 μ m), and used directly for inoculation (10⁷ conidia per milliliter).

Pathogenicity test. The test was adapted from Tu (16). Isolated cotyledonary leaves were inoculated on the abaxial surface by applying a suspension of conidia with a paintbrush. Control leaves were treated with sterile water. Infected and control leaves were then incubated on Whatman No. 3 paper moistened with sterile water in petri dishes for a 10-hr dark/14-hr light photoperiod under fluorescent light (Philips P.L. 24W 183) with the temperature at 20 C and saturated humidity. Ten days after inoculation, the presence or absence of visible symptoms (+/-) was recorded. Statistical analysis. Theoretical and

Statistical analysis. Theoretical and observed F_2 segregations were compared by the chi-square test. Genetic distance was estimated by maximum likelihood analysis (10).

RESULTS AND DISCUSSION

To score for both resistance and pod color on the same F₂ plant, it was necessary to use a reliable, nondestructive in vitro pathogenicity test. This also per-

mitted later crossing or selfing of the same plant. For this reason, the detached leaf inoculation technique described by Tu (16) was ideal for segregation test. Preliminary experiments with hypocotyl inoculation in vivo demonstrated that detached leaves show the same differential specificity toward the Colletotrichum races as the whole plant (data not shown). Symptoms on susceptible leaves, as in planta, involve a general browning that spreads from the vascular bundles.

Resistance segregation analysis of F_2 progenies from the crosses between La Victoire and AFN was done by inoculation with *C. lindemuthianum* race α Brazil 54, a race that attacks La Victoire but not AFN. The observed segregation was an expected 3:1 (nonsignificant at $\alpha = 0.05$), confirming that the resistance is under the control of a single, dominant Mendelian locus present in AFN.

Plants of the F₂ progeny were also examined for pod color phenotype. In the field or greenhouse without additional light, the purple pod phenotype of AFN is barely visible. Therefore, the use of fluorescent bulbs was necessary to obtain a reproducible and intense purple-mottled phenotype on the exposed pods of AFN, whereas La Victoire pods maintained under the same conditions remained green. F₁ plants had purple pods, and the segregation ratio of pod color in 548 plants was 3:1 (nonsignificant at $\alpha = 0.05$), thus demonstrating that purple pod color of AFN is under the control of a single, dominant Mendelian locus. This locus has been designated Anp (anthocyanins in the pod). The Anp gene seems to be involved in a specific step of the anthocyanin metabolism in the pod, because anthocyanins, which are absent from La Victoire pods, are normally accumulated in the hypocotyls, cotyledons, flowers, and seed coats of both genotypes.

Among 548 F₂ plants, 396 were resistant with red pods, 139 susceptible with green pods, and 13 have been identified as harboring a recombinant phenotype, i.e., green pod-resistant (six plants) or purple pod-susceptible (seven plants). These recombinant phenotypes were confirmed by analysis of F₃ progeny from each plant (a minimum of 20 plants for each assay). Thus, the two genetic loci involved in anthracnose resistance and pod color, respectively, appeared to be

strongly linked but cannot be at the same locus. The genetic distance between Anp and Mex2 was estimated by the maximum likelihood test to be 2.3 ± 1.3 cM. This linkage supports the original observations of Bannerot and Fouilloux (personal communication) in the course of Mex2 breeding.

The green pod-resistant recombinants show the same level of resistance as AFN, whereas the purple pod-susceptible ones display the same symptoms as La Victoire. In this case, anthocyanins appear not to have fungistatic or fungitoxic effect, which is in contrast to previous reports concerning the effect of delphinidin on conidial germination of Fusarium oxysporum Schlechtend.:Fr. f. sp. pisi (J. C. Hall) W. C. Snyder & Hanna (9) or the biostatic activity of pelargonidin-3-glucoside from bean seeds (13).

Several linkages between specific resistance genes to plant disease and anthocyanin gene markers have been described. Two examples of this are the linkage between a gene controlling bean seed color and a dominant gene involved in hypersensitive response to bean common mosaic virus (15), and between a gene for tomato anthocyanin pigmentation and a resistance gene to tobacco mosaic virus (12). All of the observed linkages between anthocyanin color and specific resistance genes to pathogens are in the range of 1-4 cM.

The Mex2-Anp recombinants obtained will be used to introduce the Mex2 resistance gene in breeding programs to obtain Mex2 resistant plants with green pods, because purple pods are commercially unacceptable. The pod color character is currently being used as a tag to construct nearly isogenic lines differing for the presence or absence of Mex2 (18). These nearly isogenic lines will be suitable to unravel molecular aspects of Mex2 resistance specificity. RFLP markers will be searched for in the Anp-Mex2 area to allow Mex2 gene cloning by chromosome walking strategies (14).

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