In Vitro Inhibitory Activity of Antimicrobial Peptides Cecropin, α-Thionin DB4, and γ-Thionin RsAFP1 Against Several Pathogens of Strawberry and Highbush Blueberry

F.A. Hammerschlag

Fruit Laboratory, Agricultural Research Service, U.S. Department of Agriculture, 10300 Baltimore Avenue, Beltsville, MD 20705

Abstract. As part of a program to develop transgenic highbush blueberry (Vaccinium corymbosum L.) and strawberry (Fragaria xannanassa Duchesne) cultivars with increased levels of disease resistance, we have investigated the feasibility of introducing genes for the antimicrobial peptides cecropin B and MB39, α-thionin DB4 (DB4) and γ-thionin RsAFP1 (RsAFP1) by testing the effects of these peptides on several important pathogens of these two crop species. A thin-layer plate bioassay was conducted with these peptides and the pathogenic Botrytis cinerea (Pers. ex. Fr.), Botryosphaeria dothidea (Mouq. ex. Fr.) Ces & de Not., Colletotrichum acutatum Simmonds, C. gloeosporioides (Penz.) Penz.et Sacc., C. fragariae Brooks, Monilinia vaccinii-corymbosi Reade (Honey), Phytophthora fragariae Hickman and Xanthomonas fragariae Kennedy and King. The minimum lethal concentration (µM) ranged from 0.13 for X. fragariae strains 10 and 128 to 72.8 for C. gloeosporioides isolate Akp1. For DB4, the minimum inhibition concentration (µM) ranged from 0.03 for X. fragariae strain 6 to 87.2 for B. cinerea isolate cc. For RsAFP1, the minimum inhibition concentration (µM) ranged from 0.13 for X. fragariae strain 6 to 61.4 for M. vaccinii-corymbosi isolate 9423-X-45. These results indicate that introducing genes for either cecropin, DB4 or RsAFP1 into strawberry may be useful for controlling bacterial angular leaf spot disease caused by X. fragariae.

Cultivated strawberry and highbush blueberry are hosts to a large and diverse number of pathogens and pests including fungi, bacteria, viruses, phytoplasma, nematodes, and arthropods. The pathogens that are considered major obstacles to maintaining or increasing strawberry production include P. fragariae, C. fragariae (Hancock et al., 1991; Maas et al., 1995; Roberts et al., 1995), and M. vaccinii-corymbosi, B. dothidea and C. gloeosporioides for blueberry (Luby et al., 1991). There is no effective chemical control for X. fragariae (Maas et al., 1995), and although pesticides are currently used to control the above mentioned fungal pathogens, the need to reduce production costs, coupled with concerns over the chemical impact on the environment and safety of food products, increased resistance of pests and pathogens to pesticides, failure of chemical companies to renew registrations, and phasing out of pesticides (Cornelissen and Melchers, 1993; Gilmartin, 1983; Wilhelm, 1999) suggest that introducing disease resistance into fruit crops may be a viable approach to controlling crop losses. Although genetic sources of resistance to disease exist for blueberry and strawberry, the many generations of hybridization and selection required to produce commercial quality fruit with adaptive traits obtained from noncommercial genotype are a severe impediment to crop improvement (Hancock et al., 1991; Luby et al., 1991). Over the past 10 years, some groundwork has been laid for utilizing molecular approaches to improve small fruit crops (Hokanson and Maas, 2001; Rowland and Hammerschlag, 2003), however, to date there are still only a few examples of using gene transfer to introduce horticulturally useful genes into strawberry (Gilpatrick et al., 1995; Mathews et al., 1995; Owen et al., 2002) and only preliminary transformation studies have been conducted on blueberry (Cao et al., 1998, 2003; Graham et al., 1996).

Increased disease resistance to both fungal and bacterial pathogens has been achieved in a range of plants following the introduction of genes encoding the small antimicrobial proteins e-thionin (Carmona et al., 1993), γ-thionin (a plant defense) (Terras et al., 1995), and cecropin (Jaynes et al., 1993; Liu et al., 2001). The thionins, found in many plants, are small proteins, only ≈50 amino acid units in length, that display varying levels of activity against a broad range of plant pathogenic fungi and some bacterial pathogens (Boholm et al., 1988; Moreno et al., 1994; Terras et al., 1995). Cecropins, found in the giant silk moth (Hyalopha cecropia) (Boman and Hultmark, 1987), comprise a family of naturally occurring lytic peptides that exhibit both antibacterial and antifungal activity (Li and Gray, 2003; Mills and Hammerschlag, 1993; Owens and Huette, 1997). Because of their broad spectrum antimicrobial activity and because they exhibit little cytotoxicity to animal and plant cells (Bechinger, 1997; Sharma et al., 2000), the cecropins are particularly attractive candidates for developing transgene-induced resistance. In the present study, we report on the effect of these peptides on the growth of a range of pathogens of strawberry and blueberry as a prerequisite to using a transformation approach to generate increased levels of disease resistance in these crops.

Materials and Methods

Maintenance of plant pathogens. B. cinerea isolates cc and wk (obtained from W. Conway, USDA–ARS, Beltsville, Md.), B. dothidea isolates #3 and #5 (obtained from J. Polashock, USDA–ARS, Chatsworth, N.J.), M. vaccinii-corymbosi isolates 9520-X-7, 963-X-Snas, 9423-X-45 (obtained from J. Polashock, USDA–ARS, Chatsworth) were maintained on Difco potato dextrose agar (PDA) at 4 °C in the dark. During the bioassay period, 2 to 3-mm-diameter hyphal plugs were transferred weekly to PDA and incubated in the dark at 25 °C. Colletotrichum acutatum isolates cal c and Golf, C. fragariae isolate Fla 2, and C. gloeosporioides isolates Akp-1 and 162 (obtained as silica gel cultures from Barbara Smith, USDA–ARS, Poplarville, Miss.) were stored at 4 % C in the dark. During the bioassay period, fungal-coated silica gel was transferred to PDA and then 2-3-mm hyphal tips transferred weekly to fresh PDA plates. Petri plates were incubated at 25 °C under continuous cool white fluorescent light. About every 6 to 8 weeks these three Colletotrichum spp. were reisolated from silica gel cultures.

Phytophthora fragariae (obtained from John Maas, USDA–ARS, Beltsville) was maintained in the dark at 4 °C on petri plates containing a soil and rhizosphere microorganism isolation medium (dRSM) (Buyer, 1994). For bioassays, P. fragariae was transferred weekly to dRSM and maintained in the dark at 25 °C.

Xanthomonas fragariae strains 6, 10, 128, and ATCC33239 (obtained from John Hartung, USDA–ARS, Beltsville) were maintained at 4 °C on SPA medium (Hayward, 1960) consisting of (in g·L–1) 20 sucrose, 5 peptone, 0.5 KH2PO4, 0.25 MgSO4 • 7H2O, 15 Bactoagar. For bioassays, a loopful of bacteria was streaked onto SPA medium and incubated at 28 °C in the dark for 5 to 7 d to provide cells for inoculum.

Peptides. Cecropin B was purchased from Bachem California (Torrance, Calif.). Cecropin MB39 and thionins RsAFP1 and DB4 were obtained from Lowell Owens, USDA–ARS.
Beltsville, who synthesized these peptides as reported (Owens and Huette, 1997).

**Thin-agarose radial diffusion assay.** The method of Hultmark et al. (1983) as modified by Nordeen et al. (1992) was used to assay antimicrobial activity of selected peptides. Petri plates were prepared by adding 6 ml of the media specified above for each pathogen, but with 1% seaplaque agarose instead of agar and without MgSO₄•7H₂O for X. fragariae. To test peptides on fungal pathogens and P. fragariae, an upside-down plug of fungal mycelia (7 mm in diameter), taken from the edge of actively growing culture, was placed onto the center of the solidified agarose. Petri plates were wrapped with parafilm, incubated in the dark at 25°C until the radius of the fungal colony was grown out 1 cm from the original inoculum (0.5 cm for P. fragariae). Wells 2 mm in diameter were then positioned 1 cm from the periphery of the mycelia and two microliters of test protein (1 µg·µL⁻¹) dissolved in water was added to each well. For assays with X. fragariae, the bacterium was grown overnight in liquid SPA medium, adjusted to an optical density of 0.165 at 620 nm to give a standard inoculum of ≈4 × 10⁶ colony forming units (CFU/mL), diluted with sterile distilled water and added to cooled assay medium to give a final concentration of 1 × 10⁶ cfu/mL. Wells were then positioned 2 cm from the center of the petri plate and the test solutions added as above.

Petri plates were wrapped with parafilm and incubated in the dark at 25°C with the exception of X. fragariae which was incubated at 28°C in the dark. Minimum lethal (bacteria) or minimum inhibitory (all other pathogens) concentrations were calculated from inhibition zone diameters measured when the pathogens reached the control well containing bovine serum albumin, according to the mathematical model of Hultmark et al. (1983), where minimum inhibitory or lethal concentration (Cₐ₅₀) = 0.468nα²d⁻¹n [n = mmole of peptide in well, α = agarase depth (0.11 cm), and d = zone diameter of inhibition (cm)]. After scoring, mycelia continued to grow at reduced rates into the inhibition zones, hence the term minimum inhibitory concentration is used to describe the fungal and P. fragariae response rather than minimum lethal concentration descriptive of the bacterial response. Each assay consisted of two replicate petri plates with one well per test solution per plate. Assays were conducted a minimum of three times.

**Results and Discussion**

Cecropin and thionins DB4 and RsAFP1 were all highly inhibitory (at sub-micromolar levels) to X. fragariae isolates, but not to any of the three Colletotrichum species, P. fragariae or to isolates of B. cinerea, B. dothidea and M. vaccinii-corymbosi (Table 1). The minimum lethal concentration of these peptides against X. fragariae ranged from 0.02 µM for cecropin B against strains 10 and 128 to 0.04 µM for DB4 against strains 10 and 128. The minimum inhibitory concentration of these peptides against the fungal pathogens ranged from 10.6 µM for cecropin MB39 against B. cinerea isolate wk to 87.2 µM for DB4 against C. acutatum isolate calc and B. cinerea isolate cc.

Cecropin B and its structural analog material MB39 (Boman and Hultmark, 1987; Owens and Huette, 1997) were selected for these studies over Shiva-1, another cecropin B structural analog with potent lytic activity (Jaynes et al., 1993), because previous studies showed cecropin B and Shiva-1 to be equally toxic to X. campestris pv. campestris (Jaynes et al., 1993), and because cecropin B showed cytotoxicity in the micromolar range against a range of bacteria and fungi (Akan and Earle, 2002; Mills and Hammerschlag, 1993; Owens and Huette, 1997). Recently, cecropin B was found to be significantly more effective than Shiva-1 in inhibiting growth of the bacterial pathogens Xylella fastidiosa and Agrobacterium tumefaciens (Li and Gray, 2003).

Cecropin MB39 was initially used for our studies, but cecropin B was substituted when our MB39 supply was exhausted. MB 39 differs from the latter by 1) substitution of Val for Met, to eliminate a translational start codon in the corresponding engineered gene, 2) the addition of a tripeptide to the N terminus, to create a signal-peptide cleavage site, and 3) the substitution of the amide group at the C terminus, Gly being the post-translational source of the amide group in cecropin (Owens and Huette, 1997). These two peptides have been shown to be equally active against a number of plant pathogens, but the MB39 form is potentially more useful for combating bacterial plant pathogens, because it is less susceptible to degradation by plant proteases in the intercellular fluid (Owens and Huette, 1997). The cecropin MB39 results suggest that cecropin MB39 could potentially be used to control X. fragariae. Other studies have demonstrated that MB39 transgenic tobacco (Nicotiana tabacum) (Huang et al., 1997) and apple (Malus x domestica) (Liu et al., 2001) exhibit increased levels of resistance to Pseudomonas syringae and Erwinia amylovora, respectively; however, its gene will need to be tailored to reduce the rate of degradation of the protein in strawberry.

Cecropin B was initially selected for testing against P. fragariae because it was shown to be toxic to P. infestans (Owens and Huette, 1997) as well as to a range of P. infestans genotypes (Owens, unpublished data); however, it was not effective in inhibiting growth of P. fragariae. These findings point out the difficulty of using this approach for disease control since the introduction of defense genes does not generally provide broad spectrum control (Stuiver and Custers, 2001), either because of 1) differential sensitivity of the same organism to the same gene product or 2) because of different defense systems within the different crops. Another example of the first was the differential response of the B. cinerea isolates and the Mollisinae vaccinii-corymbosi isolates to the same antimicrobial peptide in the present study (Table 1). An example of the second was encountered when the introduction of a chitinase gene into cucumber and carrot resulted in increased resistance only in carrot, even when the same pathogen was used to challenge the two transgenic crops (Punja and Raharjo, 1996).

The two thionins, DB4 and RsAFP1, were selected for this study because they were shown to be active against a range of fungi (Bolhmann et al., 1988; Terras et al., 1992, 1995). Although none of the fungi tested were highly sensitive to either of these two peptides, differential sensitivity of isolates of M. vaccinii-corymbosi and B. cinerea to each peptide was observed (Table 1). Both DB4 and RsAFP1 were as toxic to X. fragariae as cecropin B (Table 1). This is in contrast to Terras et al. (1992) who found most bacteria insensitive to defenses. However, several bacteria such as Bacillus subtilis (Osborn et al., 1995), and Pseudomonas solanacearum and Clavibacter michiganisis (Moreno et al., 1994) have been shown to be sensitive to plant defenses. Of significance is that both the thionins and cecropin MB39 were equally effective against the four X. fragariae representing four genotypic strain groups as defined by repetitive element PCR-based assays (Pooler et al., 1996), and thus, these peptides are good candidates for insertion into

| Table 1. In vitro inhibitory or lethal activity of cecropins B and MB39 and thionins DB4 and RsAFP1 against several pathogens of blueberry and strawberry.³ |
|---------------------------------|-----------------|-----------------|-----------------|
| **Plant pathogen**              | **Isolate/strain** | **Minimum inhibitory or lethal conc (µM)** |
| **Botrytis cinerea**            | cc               | 28.8 ± 9         | 87.2 ± 0         | 59.8 ± 14 |
| **Botryosphaeria dothidea**     | wk               | 10.6 ± 2         | 38.4 ± 24        | 16.8 ± 5 |
| **Colletotrichum acutatum**     | cal c            | 16.7 ± 1         | 21.4 ± 4         | 29.7 ± 3 |
| **C. gloeosporioides**          | Goff             | 2.1 ± 0.5        | 22.0 ± 4         | 21.3 ± 5 |
| **C. fragariae**                | Flx 2            | 31.8 ± 7         | 62.8 ± 25        | 23.1 ± 5 |
| **M. vaccinii-corymbosi**       | 9423-x-45        | 41.9 ± 12        | 61.4 ± 12        | 55.8 ± 16 |
| **Phytophthora fragariae**      | 9520-x-27        | 21.5 ± 12        | 37.3 ± 20        | 18.1 ± 10 |
| **Xanthomonas fragariae**       | 963-x-Sns        | 54.1 ± 25        | 16.4 ± 2         | 18.1 ± 4 |

³Values are the means and standard errors of two replicate plates from a minimum of three replicate experiments.
strawberry to control bacterial angular leaf spot disease. Bacterial angular leaf spot disease of strawberry has become increasingly important to strawberry fruit and plant production in Canada, the United States, as well as in other countries (Maas et al., 1995). Under certain conditions it can cause considerable crop loss (Roberts et al., 1995). The seriousness of the disease has caused many countries to close their borders to importation of plants from regions which have this disease problem (Calzolari, 1994). Because the mode of action of all three peptides is different, a strategy to pyramid these transgenes might be a viable approach to generating a more durable and broad-spectrum resistance in strawberry. Gene pyramiding via genetic engineering for enhanced disease resistance has been reported (Jach et al., 1995; Maqbool et al., 2001).

**Conclusions**

This study suggests a strategy for control of bacterial angular leaf spot disease by introducing the antimicrobial peptides cecropin, DB4 and RsAFP1 into strawberry. Although these peptides were not highly active against the important fungal pathogens of strawberry and blueberry, these are but a few of the wide range of antifungal plant proteins that have been identified, which include ribosome-inactivating proteins (Leah et al., 1991), chitin-binding proteins of the PR-4 and hevein types (Hejgaard et al., 1992; van Parijs et al., 1991), thaumatin-like proteins (Hejgaard et al., 1991; Vigers et al., 1991), lipid transfer proteins (Monila et al., 1993), and other cysteine-rich proteins of low molecular weight including the β-thionins (Garcia-Olmedo et al., 1992).

**Literature Cited**