

CHAPTER 9

PLANT DEFENCE COMPOUNDS AGAINST *BOTRYTIS* INFECTION

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Abstract. Plants possess a range of tools for combating a *Botrytis* infection. This chapter will describe three types of pre-formed and induced plant defence compounds and discuss their effectiveness in restricting *Botrytis* infection. Case studies are presented on several types of secondary metabolites: stilbenes including resveratrol, saponins including α -tomatin, cucurbitacins, proanthocyanidins and tulipalin A. Evidence is presented suggesting that *Botrytis* species have evolved mechanisms to counteract some of these defence responses. Secondly, we discuss the role of structural barriers and cell wall modification in preventing penetration. Finally the contribution of PR proteins to resistance is discussed.

1. Introduction

In response to pathogens, plants are generally able to mount a spectrum of defence responses, often coinciding with an oxidative burst involving active oxygen species (AOS) that commonly confers resistance to a wide range of (biotrophic) pathogens. The oxidative burst is however not completely effective against *Botrytis* and there is evidence that the pathogen actually benefits from it (Chapter 8). Nevertheless plants possess a range of tools for combating a *Botrytis* infection. This chapter will describe three types of pre-formed and induced plant defence compounds, namely secondary metabolites, structural barriers and antifungal pathogenesis-related (PR) proteins, and discuss their role and effectiveness in restricting *Botrytis* infection. In addition, mechanisms are discussed that suggest that *Botrytis* species have developed strategies to counteract some of these defence responses. Most of the information comes from research on *B. cinerea* but we will also discuss research on other, host-specialised *Botrytis* species where available and appropriate. The terms ‘compatible interaction’ and ‘incompatible interaction’ will be used for interactions

with a host and a non-host plant species, respectively. This is especially relevant for *Botrytis* species with a narrow host range, but in some cases also for *B. cinerea*.

2. Antimicrobial secondary metabolites

There are two types of antimicrobial metabolites: phytoalexins and phytoanticipins (VanEtten et al., 1994). Phytoanticipins are preformed, while phytoalexins are induced by pathogen infection. Excellent reviews on the phytoalexins identified in plants that are hosts for *Botrytis* spp. have been published by Mansfield (1980) and Daniel and Purkayastha (1995). These reviews also discuss the evidence for specific enzymatic degradation of some of these plant defence molecules, for example the metabolism of wyerone derivatives by *B. cinerea* and *B. fabae* in *Vicia faba*. The following section will discuss case studies of secondary metabolites that display inhibitory activity to *Botrytis*. Their chemical structures are very diverse and their modes of action presumably distinct.

2.1. Resveratrol and other stilbenes

Trans-resveratrol (3,5,4'-trihydroxystilbene) is one of the simplest stilbenes. It is a product of the plant secondary phenolic metabolism by the action of resveratrol synthase on p-coumaroyl-CoA and malonyl-CoA. It occurs in unrelated groups of angiosperms (Morales et al., 2000) and has been studied mainly in grapevine where it is the most abundant stilbene (Creasy and Creasy, 1998). In this species, it constitutes one of the major components of wood (Langcake and Pryce, 1976) and acts as a phytoalexin in leaves (Langcake and Pryce, 1977a; Langcake, 1981; Jeandet et al., 1995a). It is also found in the fruit skin (Jeandet et al., 1991).

Besides *trans*-resveratrol, numerous other stilbenes have been characterized in grapevine. These include a 3-O- β -glucoside of resveratrol called piceid that is formed by the action of a glycosyl transferase on resveratrol (Waterhouse and Lamuela Raventos, 1994; Romero-Perez et al., 1999) and a dimethylated derivative of resveratrol (3,5-dimethoxy-4'-hydroxystilbene) named pterostilbene of which the biosynthetic pathway remains to be resolved (Langcake et al., 1979; Pezet and Pont, 1988a). Piceids are a water-soluble form of resveratrol that can be reconverted into resveratrol by plant glycosidases (Ayran et al., 1987). Stressed or *Botrytis*-infected leaves also accumulate oligomers of resveratrol termed viniferins, the most abundant of which being *trans*- ϵ -viniferin, a resveratrol dehydrodimer (Langcake and Pryce, 1977b; Langcake, 1981) believed to result from the oxidative dimerisation of resveratrol by a plant peroxidase (Langcake and Pryce, 1977c; Langcake, 1981; Morales et al., 1997, 2000) or fungal laccase (Pezet, 1998; Schouten et al., 2002). *Cis* isomers of resveratrol, piceids and viniferins have also been detected in mature fruit and wine (Romero-Perez et al., 1999). They result from the isomerisation of *trans*-resveratrol by UV irradiation (Roggero and Garcia-Parrilla, 1995).

An early study on the antifungal activity of stilbenes revealed that they rapidly inhibit the respiration of fungal cells, probably by acting as uncoupling agents and by forming protein-phenol complexes (Hart, 1981). Based on the structural similarity of hydroxystilbenes and aromatic hydrocarbons, it was inferred that their

mode of action may involve lipid peroxidation by blocking cytochrome c reductase and monooxygenases (Pezet and Pont, 1995). A study on structure-activity relationships of natural stilbenes and synthetic derivatives demonstrated a positive correlation between the biological activity of stilbenes and their hydrophobicity, their ability to form complexes with proteins and the electron-attractivity of their substituents (Pont and Pezet, 1990; Pezet and Pont, 1995). The hydroxystilbenes most active on fungal respiration were pterostilbene and ϵ -viniferin with respective EC₅₀ values of 20 $\mu\text{g/ml}$ (Pezet and Pont, 1988a) and 37 $\mu\text{g/ml}$ (Langcake and Pryce, 1977a).

Resveratrol displayed no immediate toxicity towards *B. cinerea* because of its hydrophilic character and inability to reach its targets in fungal cells (Pezet and Pont, 1995). Long-term incubation of *B. cinerea* with resveratrol, however, led to the inhibition of germination of conidia, as well as the elongation of germ tubes and hyphae (Adrian et al., 1997). Functional analysis of a resveratrol-inducible laccase gene, *Bclcc2*, recently clarified this discrepancy (Schouten et al., 2002). *Bclcc2* gene replacement mutants had an impaired ability to metabolise resveratrol and displayed improved growth characteristics on resveratrol containing media, suggesting that resveratrol is not toxic in itself, but the BcLCC2 protein is responsible for transforming resveratrol into a fungitoxic substance (member of the viniferins). Such a novel mechanism of activation of a phytoalexin presents obvious advantages for the plant. Plant secondary metabolites may be toxic to the plant itself and often have limited solubility. As resveratrol is one of the most soluble and least toxic stilbenes, it can be stored safely at high concentrations in vacuoles and cell walls (Morales et al., 2000) only to be activated upon contact with a microorganism. The activation of resveratrol into viniferins does not necessarily require the conversion by the pathogen. Increased ϵ -viniferin production also occurs in grapevine in the absence of a microbe, after UV irradiation (Langcake and Pryce, 1977b). The content ratios of resveratrol and ϵ -viniferin remained unchanged in several grapevine cultivars after UV irradiation or *B. cinerea* infection, even when their respective concentration increased (Douillet-Breuil et al., 1999; Adrian et al., 2000). Plant enzymes potentially involved in oxidation of resveratrol include peroxidases (Langcake and Pryce, 1977a, b). One constitutive basic grapevine peroxidase exhibits a high affinity for resveratrol (Morales et al., 1997). The peroxidase-mediated oxidation of resveratrol does not yield viniferins *in vitro*. However, the natural association of the benzene rings of resveratrol with cellulose fibres affects the stability of radical reaction intermediates such that peroxidase-mediated oxidation of resveratrol may generate viniferins *in planta* (Morales et al., 2000). Interestingly H₂O₂, a co-substrate of peroxidases, is generated during *B. cinerea* infection (Chapter 8).

Pterostilbene has the highest antifungal activity, but its concentration is less than 5 $\mu\text{g/g}$ in leaves (Douillet-Breuil et al., 1999) and fruit (Adrian et al., 2000) of various grapevine cultivars and is not enhanced by UV irradiation or *B. cinerea* infection. This raised doubts about the capacity of pterostilbene to protect plants against *B. cinerea* (Langcake, 1981; Douillet-Breuil et al., 1999; Adrian et al., 2000). The potency of pterostilbene increases in the presence of glycolic acid, an organic acid that accumulates to high concentrations in immature grape berries.

Pterostilbenes may thus act as a constitutive defence component in berries (Pezet and Pont, 1988b). In contrast, resveratrol acts in an inducible defence reaction against *B. cinerea*. A positive correlation was found between the resveratrol level in grapevine cultivars and their resistance to *B. cinerea* (Langcake and McCarthy, 1979). UV irradiation or *B. cinerea* challenge of leaves and fruit of resistant grapevine cultivars led to resveratrol concentrations up to 750 µg/g, exceeding levels needed for toxicity to *B. cinerea in vitro* (Jeandet et al., 1995a; Douillet-Breuil et al., 1999; Adrian et al., 2000). Local resveratrol concentrations may be more important than those found after grinding whole tissue. The interaction of grapevine leaves with an incompatible *B. cinerea* isolate led to a hypersensitive-like response during which stilbenes preferentially accumulated in the direct vicinity of the fungus (Derckel et al., 1999).

Raising resveratrol levels in crop plants is an attractive option because of the potential of this substance to protect plants from the attack of pathogens and to improve human health (Chiou, 2002) by preventing and curing cancers (Savouret and Quesne, 2002) and guarding against vascular diseases (Hung et al., 2000). In plant species that do not naturally produce stilbenes, the production of resveratrol was achieved by genetic engineering. Introduction of the grapevine stilbene synthase gene *Vst1* into tobacco (Hain et al., 1993) and barley (Leckband and Lorz, 1998) resulted in resveratrol accumulation and enhanced protection against grey mould. In transgenic kiwifruit, however, the presence of high endogenous levels of glycosyl transferase activity led to the preferential accumulation of piceid over resveratrol and no protection to *B. cinerea* infection was observed (Kobayashi et al., 2000). In plants that naturally have the genetic potential to produce resveratrol, the expression of stilbene synthase is low in unstressed leaves. It can be induced in this organ by a variety of biotic and abiotic elicitors such as AlCl₃ (Adrian et al., 1996; Jeandet et al., 1998), UV, paraquat, wounding, H₂O₂, salicylic acid, jasmonic acid, ethylene (Grimmig et al., 1997; Adrian et al., 2000; Lippmann et al., 2000; Chung et al., 2003), methyl jasmonate (Larronde et al., 2003), systemic acquired resistance elicitors (Busam et al., 1997), abscisic acid (Ban et al., 2000) and ozone (Grimmig et al., 1997). The exogenous application of *B. cinerea* cell wall fragments (Liswidowati et al., 1991) or a live soil-borne biocontrol *Bacillus* species (Paul et al., 1998) have also been reported to lead to increased accumulation of resveratrol in grapevine leaves.

Fruit, such as grape berries, behave differently from leaves by expressing significant levels of resveratrol constitutively. One concern is that resveratrol levels decrease during fruit ripening due to lower expression of stilbene synthase (Bais et al., 2000) and increased substrate competition for this enzyme with chalcone synthase, whose activity is stimulated during fruit ripening to provide precursors for anthocyanin accumulation (Jeandet et al., 1995b). Nevertheless, grapes keep their capacity to accumulate higher levels of resveratrol after UV irradiation at any stage of ripening (Adrian et al., 2000) and even after harvest (Cantos et al., 2001). Another concern with resveratrol accumulation in grape is that it only occurs in the skin, implying that resveratrol can only participate in protection of the berry from outside intruders when the fruit skin is intact. Cultivation practices have also been shown to influence resveratrol accumulation in grapevine leaves and fruit. A

negative correlation was found between the resveratrol level and the extent of nitrogen fertilization (Fregoni et al., 2000; Bavaresco et al., 2001) or fungicide spray application (Magee et al., 2002) while increased potassium fertilization correlated with higher levels of resveratrol (Fregoni et al., 2000). The potential of grapevine cultivars to accumulate resveratrol was positively correlated with their tolerance to *Botrytis* (Douillet-Breuil et al., 1999), leading to the suggestion that resveratrol production capacity could be used as a criterion in the selection of resistant varieties (Pool et al., 1981; Creasy and Coffee, 1988; Sbaghi et al., 1995).

The ability of *B. cinerea* to cope with stilbenes should not be underestimated. Several studies did not corroborate a relationship between resveratrol accumulation and disease incidence in grapevine (Magee et al., 2002; Keller et al., 2003). Increasing conidia concentration in synthetic media containing resveratrol led to increased resveratrol catabolism and survival of conidia (Hoos and Blaich, 1990) and a correlation between resveratrol detoxification capacity of *B. cinerea* isolates and their virulence on grapevine leaves has been proposed (Sbaghi et al., 1996). Detoxification of resveratrol and the related pterostilbene has been attributed to their oxidative dimerisation by *B. cinerea* laccases (Hoos and Blaich, 1990; Pezet et al., 1991; Sbaghi et al., 1996; Adrian et al., 1998; Breuil et al., 1999). Application of cucurbitacins led to reduced production of laccases (Sect. 2.3) and correlated with increased resistance of several hosts, even those that do not accumulate stilbenes, such as carrot and cucumber (Bar-Nun and Mayer, 1990; Viterbo et al., 1993a). This suggested that *B. cinerea* laccases are part of a more general 'attack' machinery designed to detoxify phenolic defences from many host plants (Staples and Mayer, 1995). However, the product of one specific laccase gene, BcLCC2, has been implicated in the opposite effect by activating resveratrol into a fungitoxic substance (Schouten et al., 2002). Laccase production by *B. cinerea* increases during late stages of infection or development (Roudet et al., 1992; Manteau et al., 2003) in parallel with a decrease in resveratrol concentration in its host (Adrian et al., 2000; Montero et al., 2003) and can be stimulated *in vitro* by host-derived substances such as phenolics, pectins (Viterbo et al., 1993a) and ambient pH (Manteau et al., 2003). Proanthocyanidins in fruit act as competitive laccase inhibitors (Pezet et al., 1992).

Three laccase genes were described so far (Schouten et al., 2002) and limited information is available on their distribution among *B. cinerea* populations and their physiological role during pathogenesis. This role may prove to be complex since the incubation of *B. cinerea* with resveratrol has been shown to lead to the production of an array of oxidised derivatives, including the fungitoxic ϵ -viniferin (Langcake and Pryce, 1977a; Keller et al., 2003) and six other dimerisation products: δ -viniferin (Breuil et al., 1998; Pezet et al., 2003a), leachinol F, pallidol and restrytisols A-C (Cichewicz et al., 2000). Some information is available on the toxicity of these individual, pure metabolites towards *B. cinerea*, but one should consider that such *in vitro* tests may not reflect the chemical environment that the pathogen encounters during infection. Laccase activity on resveratrol yields radical quinone intermediates that may react with very distinct (phenolic and other) compounds, giving rise to heterogeneous products. The metabolism of resveratrol *in planta* therefore depends on the temporal and spatial distribution both of the fungal laccase(s), the plant peroxidase(s), their substrate(s), as well as on the enzyme kinetics for each substrate.

2.2. α -Tomatine and saponins

α -Tomatine is a secondary metabolite produced in tomato leaves and unripe fruit (Friedman, 2002). It is a potent antifungal (e.g. Sandrock and VanEtten, 1998) and insecticidal (e.g. Kowalski et al., 2000) compound that interacts with sterols in membranes (Keukens et al., 1995). Already in the 1970s it was reported that α -tomatine inhibits mycelial growth of *B. cinerea* while not affecting germination of conidia (Verhoeff and Liem, 1975). It was proposed that its fungistatic action was responsible for maintaining *B. cinerea* infections in a quiescent state. The reduction in α -tomatine content in ripening fruit (Friedman, 2002) supposedly relieves fungistasis and permits fungal outgrowth. Furthermore, Verhoeff and Liem (1975) reported that *B. cinerea* was able to convert α -tomatine to tomatidine by sugar hydrolysis, thereby actively detoxifying the compound and facilitating fungal outgrowth into tomato tissue prior to the natural drop in α -tomatine levels.

Supporting evidence for the important role of α -tomatine in conferring resistance towards *B. cinerea* came from work of Quidde et al. (1998). Among a set of 13 isolates, one field isolate was identified that lacked α -tomatine degrading ("tomatinase") activity. This isolate, designated M3 and originating from grape, was able to form primary lesions on tomato but these lesions never expanded. Virulence of isolate M3 on *Phaseolus vulgaris* was not affected (Quidde et al., 1998). Contrary to the report of Verhoeff and Liem (1975), the α -tomatine degradation product was identified as β -tomatine (Quidde et al., 1998). On the basis of this conversion it was proposed that the tomatinase should possess xylosidase activity. Quidde et al. (1999) cloned a *B. cinerea* gene homologous to the tomatinase gene *tom1* from *Septoria lycopersici*. The gene, designated *sap1*, encodes a β -glucosidase active on saponins. Gene replacement mutants deficient in *sap1* lost the ability to degrade avenacin (a phytoanticipin from roots of oat, a non-host species for *B. cinerea*), yet they remained able to detoxify tomatin, digitonin and avenacoside (a phytoanticipin from leaves of oat). *sap1*-deficient mutants remained able to infect tomato, suggesting that the *sap1* gene does not encode the true tomatinase. Thus *B. cinerea* can produce at least three distinct saponin-specific glycosidases (Quidde et al., 1999).

2.3. Cucurbitacins

Cucurbitacins are bitter-tasting triterpenoids, produced by the Cucurbitaceae, that are toxic to insects and mammals. There are at least six forms of cucurbitacins, one of which antagonises insect steroid responses (Dinan et al., 1997) while another form disrupts the cytoskeleton in carcinoma cells (Duncan et al., 1996). Application of cucurbitacin I to host tissue prior to inoculation prevented infection by *B. cinerea* (Bar-Nun and Mayer, 1990). When cucurbitacin was added to a *B. cinerea* liquid culture, laccase secretion into the medium was reduced (Bar-Nun and Mayer, 1989) while other enzymes were not affected (Viterbo et al., 1993a). Some cucurbitacin forms were more effective than others (Viterbo et al., 1993b). Radio-labelled cucurbitacin was taken up by *B. cinerea* mycelium suggesting that it acted intracellularly, reducing laccase activity (Viterbo et al., 1994), supposedly as a consequence of repression at the mRNA level (Gonen et al., 1996). These gene

expression studies were however performed with a probe corresponding to the *Bclcc1* gene, whereas the laccase gene that is induced by gallic acid is in fact *Bclcc2* (Schouten et al., 2002). The two genes show poor cross-hybridisation (A. Schouten, unpubl.). Neither of these laccase genes is important for virulence on a range of hosts (Schouten et al., 2002). It remains to be resolved whether the reduction of *B. cinerea* infection by application of cucurbitacins (Bar-Nun and Mayer, 1990) is caused indirectly by activation of defence or directly by inhibiting virulence factors.

2.4. Proanthocyanidins

Proanthocyanidins (condensed tannins) are polymeric flavonoids that result from the condensation of two or more derivatives of flavan-3,4-diol. The term was first coined by Weinges et al. (1969) to designate these colourless tannins that form intensely coloured anthocyanidins upon heating with acid. Proanthocyanidins are widely distributed in the plant kingdom and are constitutive components in a number of discrete tissues in most plant organs. In leaves, they are mostly present in vascular tissue and in fruit they preferentially accumulate in the epidermis and seeds (Porter and Schwartz, 1962; Bachmann and Blaich, 1979; Hills et al., 1981; Jersch et al., 1989; Prieur et al., 1994; Prusky, 1996; Souquet et al., 1996). The chemical structure and composition of proanthocyanidins vary among plant species, organs and also with the stage of organ development. In grape berries, the mean degree of polymerisation of the proanthocyanidins, the proportion of epigallocatechin extension subunits and the level of anthocyanin association increase during ripening (Hills et al., 1981; Kennedy et al., 2001).

Because of the potential association of proanthocyanidins with sugars (including pectins), flavonoids and proteins (Kennedy et al., 2001), reports on their toxicity towards *B. cinerea* (conidia germination and germ tube elongation) have yielded opposite results depending on the state of purity of the proanthocyanidins (Hills et al., 1981; Jersch et al., 1989; Hebert et al., 2002). Nevertheless, these studies agree in suggesting that plant proanthocyanidins maintain *B. cinerea* in a quiescent stage, leading to delayed development of symptoms. The transition from quiescence into expansion is triggered during host senescence or ripening and occurs at a less senescent or ripe stage in susceptible varieties (reviewed by Elad, 1997). For example, grapes develop latent infections as early as the blooming stage (McClellan and Hewitt, 1973) while lesions are increasingly likely to start spreading only when the fruit ripens (Hills et al., 1981). The development of quiescent *B. cinerea* infections on grape did not correlate with conidia density (Coertze and Holz, 1997), the stage of ripening (Hills et al., 1981) or the level of resistance of the cultivar to grey mould (Pezet et al., 2003b). Even though the production of phytoalexins after pathogen entry is one way to keep *B. cinerea* quiescent (Sect. 2.1), some fruit, such as strawberries, do not mount a phytoalexin response (Jersch et al., 1989), implying that fungal quiescence can be maintained by additional mechanisms. These may involve the inhibition of fungal pathogenicity factors by proanthocyanidins (Prusky, 1996). *B. cinerea* virulence factors include a range of secreted enzymes (Chapter 7) and proanthocyanidins can complex with proteins (Haslam, 1966, 1974) resulting in non-specific inhibition of the activity of fungal polygalacturonases and

cellulases (Porter and Schwartz, 1962; Hills et al., 1981; Jersch et al., 1989). Proanthocyanidins additionally act as competitive inhibitors of *B. cinerea* laccase, thereby preventing detoxification of the phytoalexin pterostilbene. Pezet et al. (1992) reported an EC₅₀ value of 12 µg/ml for the inhibition of *B. cinerea* laccase by grape skin proanthocyanidins. Their levels in grape skin vary from 50 to 250 µg/g dry weight (Hills et al., 1981; Pezet and Pont, 1992) and 14-50 µg/g fresh weight in strawberry (Jersch et al., 1989) depending on the cultivar or the stage of ripening.

Despite the fact that proanthocyanidins do not accumulate in all plant tissues, it is suggested that endogenous proanthocyanidins contribute to maintaining *B. cinerea* in a quiescent state. More tolerant grape and strawberry cultivars accumulate larger quantities of proanthocyanidins (Hebert et al., 2002; Pezet et al., 2003b). Proanthocyanidin content decreases in grape during fruit ripening (Hills et al., 1981) while they remain constant in strawberries (Jersch et al., 1989). However, the ripening-related modifications of proanthocyanidins led to a decrease in their capacity to inhibit *B. cinerea* macerating enzymes (Hills et al., 1981; Jersch et al., 1989; Pezet et al., 1992; Pezet and Pont, 1992). This phenomenon parallels the loss of resistance of these fruit during ripening (Hills et al., 1981; Jersch et al., 1989) and occurs to a larger extent in susceptible cultivars (Pezet et al., 2003b). Finally, the addition of proanthocyanidin at 0.1% at the point of inoculation blocked the development of spreading lesions of *B. cinerea* in susceptible grapevine varieties (Hills et al., 1981).

Prolonging the quiescence of *B. cinerea* infections by increasing the proanthocyanidin content would reduce losses to grey mould, especially after harvest. However, proanthocyanidin levels are constitutive and are not known to be subject to modulation by external elicitors. Moreover, knowledge is lacking on the genes and enzymes involved in the subtle modifications of proanthocyanidin structure that affect their biological activity. The use of proanthocyanidin content as an indicator of grey mould resistance for the selection of cultivars with improved shelf-life has been suggested for grape (Pezet et al., 2003b) and strawberry (Jersch et al., 1989; Hebert et al., 2002). However, proanthocyanidins negatively affect the taste and colour (via anthocyanin binding) of fresh produce and plant-derived food products (Noble, 1990; Gawel, 1998). Attempts to modify their composition may result in a modification of the perception of these products by consumers.

2.5. Non-host resistance

2.5.1. Phytoanticipins of tulip as mediators of *Botrytis* non-host resistance

Tulip bulbs and pistils contain high concentrations of fungitoxic compounds, identified as lactones and termed tulipalins (Bergman et al., 1967; Beijersbergen, 1969). Tulipalin A is found in bulbs, whereas aerial plant parts contain mainly tulipalin B. Tulipalins A and B are stored in a glycosylated form, named tuliposides A and B. Tuliposides are less toxic to fungi than the corresponding aglycosylated lactones and are lactonised into tulipalins when the pH exceeds 5. Plant enzymes may enhance the conversion rate of tuliposides into tulipalin A (Beijersbergen and Lemmers, 1972). The lactone itself is relatively stable *in vitro* at pH values between 5.4 and 6.5 (Beijersbergen, 1969). It was suggested that precursor tuliposides, stored

in acidic vacuoles, are released upon disruption of the vacuolar membranes during cell collapse mediated by pathogens (Schönbeck and Schlösser, 1976). The lactones themselves spontaneously hydrolyse at pH values above 7.5 into the corresponding non-toxic butyric acids (Beijersbergen, 1969).

All *Botrytis* species tested so far are sensitive to pure tulipalin A, except for the tulip-specific pathogen *B. tulipae* (Schönbeck and Schroeder, 1972). Partial or complete fungistasis occurs *in vitro* at 3-5 μM . Higher doses (7-10 μM , more than 30 μM for *B. tulipae*) are lethal to *Botrytis* spp. Mycelium exposed to lethal tulipalin A concentrations does not resume growth when transferred to fresh agar without tulipalin. Microscopic observation of tulipalin A-treated mycelium showed a disappearance of hyphal cell contents and altered autofluorescence of hyphal walls as compared to vital cultures. Conidia are at least three times more sensitive to tulipalin A than mycelium (P. van Baarlen and M. Staats, unpubl.). *B. tulipae* is more tolerant to tulipalin A than other *Botrytis* species. Infection of tulips by *B. tulipae* is associated with a conversion of tuliposides into the corresponding, inactive hydroxylic acids. In contrast, infection by *B. cinerea* leads to a conversion into the active lactones (Schönbeck and Schroeder, 1972). This suggests that *B. tulipae* contains a factor that can hydrolyse the tuliposides or the lactones into the corresponding hydroxylic acids. Indeed, a total protein extract from *B. tulipae* mycelium was able to neutralise tulipalin A; incubation of tulipalin A with this protein extract resulted in loss of toxicity towards sensitive *B. cinerea* isolates (P. van Baarlen and M. Staats, unpubl.). Lactone detoxification was also reported in *Fusarium oxysporum* isolates that produce a lactonohydrolase (Shimizu et al., 1992). It is tempting to speculate that *B. tulipae* possesses an enzyme with similar activity. Such an enzyme might act as a host specificity determinant for *B. tulipae*.

2.5.2. Other monocot secondary metabolites involved in non-host resistance

The genus *Allium* contains various compounds associated with resistance to fungal disease. Some are constitutive inhibitors, such as the phenolic compound catechol that is present in the outer bulb layers of pigmented *Allium cepa* (onion) cultivars where it confers resistance to *Colletotrichum* (Link and Walker, 1933). Onion also produces a class of cyclopentane phytoalexins upon pathogen infection, designated tsibulins (Dmitriev et al., 1990), which accumulate in bulb scales at infection sites during incompatible interactions with *B. cinerea*. Tsibulins inhibited spore germination and germ tube elongation of *B. cinerea* *in vitro*. The ED_{50} values were lower than the actual phytoalexin content in bulb scale spots, where *B. cinerea* lesion formation was restricted (Dmitriev et al., 1990). Only little accumulation of cyclopentane phytoalexins was observed in onion bulb scales during infection by the specialised pathogen *B. allii* (Dmitriev et al., 1990). *B. allii* seems to either suppress tsibulin accumulation, analogous to the interaction of *B. narcissicola* and its host narcissus (discussed below) or actively degrade tsibulins, as discussed for other antifungal compounds above. It would be of interest to investigate whether *B. allii* is able to break down tsibulin cyclopentanes.

3. Tolerance of *Botrytis* to antifungal metabolites

In many cases *Botrytis* species have adapted to antifungal compounds produced by their host plants by developing mechanisms to counteract their toxicity. In fact the ability to counteract toxicity of phytoalexins and phytoanticipins is often crucial for successful host colonization. As a first line of defence to toxic compounds, fungi possess ATP-binding cassette transporters (ATR) and Major Facilitator (MFS) proteins that are able to excrete a spectrum of chemically unrelated toxic metabolites from the cytoplasm. ATR and MFS proteins serve as membrane pumps with a broad substrate range that expel chemically heterogeneous antifungal compounds at the expense of ATP or proton extrusion (reviewed by de Waard, 1997). These proteins may also be involved in fungicide resistance (Chapter 12). *B. cinerea* possesses a large family of functional transporter genes, some of which confer protection to antibiotics (Schoonbeek et al., 2002) and plant defence metabolites including resveratrol (Schoonbeek et al., 2001). The degree of tolerance that ATR and MFS transporters confer towards these toxicants is in several cases small, only rarely do they provide full resistance. This low degree of tolerance may however be biologically significant as it allows the pathogen time to activate true enzymatic detoxification mechanisms. Such true detoxification mechanisms are specific for each individual antifungal compound, as discussed in Sect. 2.

4. Structural barriers and cell wall modifications

Structural barriers can be mounted during resistance responses and non-specific wound responses (Heath, 2000, 2002) by means of incorporating and cross-linking phenolic compounds at penetration sites. Cell wall modifications may directly pose physical barriers for fungi and they may interfere with degradation of wall components that function as nutrient sources for *Botrytis* species. One form of plant cell wall modification associated with disease resistance is their lignification. The phenylpropanoid pathway generates coumaryl Co-A and cinnamyl Co-A esters that serve as precursors for diverse compounds, including lignin (Dixon and Paiva, 1995). Wall modification occurs in the incompatible interaction of *B. cinerea* with a non-host, wheat. Challenging wheat leaves with *B. cinerea* conidia results in a localised, sharp increase of sinapyl alcohol dehydrogenase, phenylalanine ammonia lyase and peroxidase expression and progressive lignification (Mitchell et al., 1994). The phenylpropanoid pathway also generates phytoalexins (discussed above). The simultaneous formation of structural barriers and phytoalexin production blocks the infection and lesion spread of *B. cinerea* during incompatible interactions with carrot, narcissus and wheat (Garrod et al., 1982; O'Neill and Mansfield, 1982; Mitchell et al., 1994). The relative contribution of wall modifications and phytoalexin production to the effective inhibition of infection is often unclear.

Other wall modifications are established by a class of phenolic amides, mainly hydroxycinnamic acid derivatives and tyramine (McLusky et al., 1999). The synthesis and incorporation in cell walls of amides is associated with the inhibition of outgrowth of *B. allii* germ tubes within modified infection sites (Stewart and Mansfield, 1985). The amides that are produced in onion cells are not fungitoxic to

B. allii and *B. cinerea* conidia and germ tubes up to concentrations of 1 mM (McLusky et al., 1999). The fact that amide synthesis and their incorporation into cell walls were associated with attempted penetrations of *B. allii*, a compatible pathogen of onions, suggests that this defence mechanism provides a general protection against *Botrytis*. The efficiency of the resistance mechanism may depend on spore concentration, H₂O₂ availability and occurrence of other stress factors.

The cell wall modification response (Heath, 2000) has been studied in lily and tulip cultivars in compatible and incompatible interactions, with *B. cinerea*, *B. elliptica* or *B. tulipae*. Cell wall modification was observed during incompatible interactions but not during compatible interactions (P. van Baarlen and J. van Kan, unpubl.). Autofluorescence of cell walls at penetration sites occurred within 18 h upon inoculation of tulip and lily leaves with dry conidia. Fluorescent vital staining of plant cells showed that in an incompatible interaction, the non-host plant cells were not killed upon expression of this resistance response. Germ tubes did not penetrate the modified walls any further. Primary lesion formation was not observed in incompatible interactions, suggesting that wall modifications can effectively restrict *Botrytis* infection. A more extensive study of wall modifications during compatible and incompatible plant-*Botrytis* interactions seems worthwhile, although it will remain difficult to distinguish the contribution to resistance of cell wall modification from that of the concomitant phytoalexin production.

Several methods by which *B. cinerea* could interfere with cell wall resistance responses may be envisaged. The pathogen may actively suppress resistance responses in a compatible interaction. Preliminary microscopy studies have shown that lily cells, at the sites of penetration by *B. elliptica*, did not show the characteristic local yellow cell wall autofluorescence observed in incompatible interactions. Also in the compatible interaction of narcissus and *B. narcissicola*, cell wall modifications were mostly absent (O'Neill and Mansfield, 1982). Both phytoalexin production and wall modification depend upon the phenylpropanoid phenolic acid pathway. It is conceivable that in a compatible plant-*Botrytis* interaction, the cell wall modifications and phytoalexin production are suppressed by (a) secreted fungal factor(s). One study has shown that during a compatible interaction of the cowpea rust fungus (*Uromyces vignae*) with its host, cell wall-associated defence responses are suppressed (Heath, 2002). In this pathosystem, suppression of defence is established through interference with wall-membrane communication via secreted fungal peptides that interfere with integrin proteins that mediate wall-membrane adhesion (Mellersh and Heath, 2001). We are, however, not aware of literature reporting active suppression of defence by *Botrytis*.

5. Pathogenesis-related proteins

Pathogenesis-related proteins (PR proteins) represent a large array of proteins coded by the host plant that are co-ordinately expressed under pathological or related situations. They have been characterised in over 70 plant species and 13 plant families including mono- and di-cotyledonous plants. They are extremely diverse in terms of enzymatic and biological activity and have been grouped into 13 protein families based on primary structure and serological relationships (Van Loon, 1999).

All share several physicochemical properties such as solubility in acidic buffers, resistance to proteolysis, a molecular mass less than 50 kDa and a lack of quaternary structure and glycosylation (Stintzi et al., 1993). They primarily accumulate in plant cell walls and vacuoles. *B. cinerea* infection leads to PR protein induction in many plants (Van Loon, 1985). Fruit tissues differ significantly from other plant organs by accumulating unusually high concentrations of a limited set of PR-like proteins that share sequence similarities with known PR proteins but accumulate with fruit development (most often during fruit ripening) and not after stimulation with pathogenesis-derived signals such as salicylic acid or wounding (Derckel et al., 1998). Grapes, for example, mainly accumulate one thaumatin-like protein and one chitinase which make up 80% of the total soluble protein content of the fruit at harvest (Derckel et al., 1998; Salzman et al., 1998; Waters et al., 1998).

Members of several PR protein families display some toxicity towards *B. cinerea in vitro*. For some of them, this may be caused by their potential to degrade chitin and β -glucan fragments of *B. cinerea* cell walls (Gomez-Miranda et al., 1981; Punja and Zhang, 1993; Simmons, 1994). However, fungitoxicity varies greatly among members of one PR protein family, just like their specific enzyme activity which may differ by up to 250-fold towards a given substrate (Kauffmann et al., 1987; Stintzi et al., 1993). A grape PR-like protein (chitinase) has one of the highest botryticidal activities. It inhibits germination of conidia with an EC_{50} value of 7.5 μ g/ml (Derckel et al., 1998) and it restricts the elongation of hyphae (Salzman et al., 1998). The chitinase is present at levels up to 26 μ g/g in fruit of resistant cultivars (Derckel et al., 1998; Salzman et al., 1998). The grape thaumatin-like protein, on the other hand, exhibited no toxicity towards *B. cinerea in vitro* (Salzman et al., 1998). In the presence of 1M glucose (a physiological concentration for a ripe grape), however, the toxicity of the grape chitinase was potentiated by as much as 70% and the grape thaumatin-like protein became equally toxic as the chitinase at similar concentrations (Salzman et al., 1998).

Despite their anti-microbial activity *in vitro*, there is little evidence to support a potential role of PR proteins in effective plant disease resistance to *B. cinerea*. The homologous or heterologous expression of PR proteins in transgenic plants, or the infiltration of single PR proteins in leaves, has rarely led to any significant level of protection against *B. cinerea* (Lawton et al., 1993; Datta et al., 1999; Neuhaus, 1999; Van Loon, 1997). Only the coordinated expression of members of different PR protein families yielded some (though limited) degree of tolerance due to synergistic effects (Zhu et al., 1994; Jach et al., 1995; Datta et al., 1999). Additionally, treatments that stimulate the coordinate expression of a large array of PR proteins in tomato leaves such as ethylene treatment (Diaz et al., 2002) or *B. cinerea* primary lesion formation (Benito et al., 1998) did not prevent the development of *B. cinerea* spreading lesions. To the contrary, a higher degree of susceptibility (higher fungal growth as estimated by the level of fungal actin mRNA) correlated with higher levels of PR proteins mRNAs (Diaz et al., 2002). In any case, the commercial potential of plants exhibiting higher levels of PR proteins will be hampered by the fact that PR proteins are associated with several undesirable effects such as the formation of haze in grape juices (Waters et al., 1996) and allergic reactions (Breiteneder and Ebner, 2000; Hoffmann-Sommergruber, 2000).

6. Conclusions

There are interesting perspectives for altering secondary metabolism in plants to optimise the contents of antifungal compounds, either by transgenes or by classical breeding. The example of enhanced *Botrytis* resistance in transgenic tobacco producing resveratrol (Hain et al., 1993) already provides evidence for its feasibility. Now that the field of plant pathology and breeding has entered the genomics era, it will be increasingly feasible to modulate the levels of secondary metabolites, such as phenylpropanoids, to enhance defence (Dixon et al., 2002). The information on the *Botrytis* genome will be a useful tool. When plants can be equipped with antifungal metabolites for which *Botrytis* has no detoxifying enzymes, it is likely that these metabolites will be more effective. Conversely, when *Botrytis* possesses enzymes that may convert non-toxic metabolites into antifungal compounds (Schouten et al., 2002) one can perhaps make use of these enzymes. When considering the options for enhancing plant resistance to *Botrytis* by modulation of secondary metabolites, it should be taken into serious account that it is often not required to achieve complete resistance as conferred by the classical R-genes. Especially in the case of post-harvest problems, it will be sufficient to delay the disease outbreak and attenuate disease out-growth. We recommend the design of strategies that extend the latent infection phase. Chapter 20 deals in more detail with novel approaches to reduce the damage inflicted by *Botrytis* diseases.

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