# Chapter 2 Insights into Fruit Defense Mechanisms Against the Main Postharvest Pathogens of Apples and Oranges



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Abstract *Penicillium digitatum* and *P. expansum* are the most devastating pathogens of citrus and pome fruits respectively, causing significant economic losses during postharvest handling worldwide. To obtain new rational and environmentally friendly control alternatives, a better understanding of the fruit-pathogen interaction may be considered as a novel perspective for the control of postharvest diseases. The main objective of our studies was to gain insights into the fruit-pathogen interactions, specifically in oranges and apples defence responses against compatible and non-host pathogens. For such purposes, firstly a deep study of infection capacities of both pathogens in oranges and apples at different conditions was performed. Later, we characterized the effect of wound response in oranges and apples harvested at different maturity stages and temperatures, as well as we identified the possible compounds involved in the wound healing process. In addition, a visualization of lignin, suberin and callose in apple and oranges tissue was conducted using histochemical tests. Finally, in apples, a transcriptomic study in response to compatible and non-host pathogen was conducted; and the expression of several genes involved in the phenylpropanoid pathway were quantified in citrus fruit.

**Keywords** *P. expansum* · *P. digitatum* · Host-pathogen interaction · Lignification · Phenylpropanoid pathway · Transcriptomics analysis

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## Introduction

Oranges and apples are some of most produced fruits worldwide. The production of citrus in Mediterranean regions is 25,216 thousand tons in 2016, and Spain is the first producer with 6,882 thousand tons. Specifically, Spain contributed with 3,641 thousand tons of oranges and, therefore, remains the major contribution in Mediterranean Region with around 25% of total production (FAO 2017). In turn, as important apple producer, Spain contributed to the apple worldwide production with 473 thousand tons in 2018 (Prognosfruit 2018).

Quality deterioration of fresh fruits due to fungal pathogens is one of the main economic losses produced in postharvest handling worldwide. The most common cause of citrus and pome fruit decay is the development of rots caused by *Penicillium* spp. which are one of the most common and destructive genera in postharvest diseases, affecting a wide variety of fruit. In particular, *P. digitatum* and *P. expansum* are the most devastating postharvest pathogens of citrus and pome fruits, respectively. Although both are close related species, they show a different range of hosts. Whereas *P. digitatum* causes only green mould in citrus and could be considered as specialist species, *P. expansum* has the ability to infect a broader range of hosts (Fig. 2.1). These necrotrophic pathogens need wounds to enter tissues and start the infection process.

The traditional way to control these diseases is the usage of synthetic fungicides. However, there are important problems associated by their use, hence, actual trends are directed to find new rational and environmental friendly control alternatives focused on physical, chemical and biological strategies. These facts justify the need and the interest for more detailed studies of host-pathogen interaction to increase the knowledge on both virulence factors of pathogens and fruit defence mechanisms with the final aim to improve control strategies (Tian et al. 2016).



Fig. 2.1 The main postharvest pathogens of oranges and apples, *Penicillium digitatum* and *P. expansum*, respectively

At this regard, new perspectives to control postharvest diseases are needed and some of them are based on the knowledge of fruit-pathogen interaction. On one hand, the capacity of fungi to adapt to new conditions, overcoming plant defence mechanisms and becoming resistant to fungicides is huge. On the other hand, the defence machinery of the fruit is also complex and unknown, and evolved a multilevel series of structural and biochemical barriers that are both constitutive or preformed and inducible. Both components of host-pathogen interactions should be addressed from different perspectives, pathological, biochemical and molecular studies, to elucidate the factors that can modulate both compatible and non-host interactions.

The main proposal of this chapter is to approach the research line related to control postharvest diseases on pome and citrus fruit from new perspective, mainly focused on fruit-pathogen interactions in apple/oranges-*Penicillium* spp.

#### Fruit–Penicillium spp. Interaction

In many instances, the differences in the outcome of a host-pathogen interaction relies on a rapid and efficient deployment of defense responses, which will be modulated depending on whether the host-pathogen interaction involves a compatible or non-host interaction. For that purpose, *P. digitatum* and *P. expansum* were defined as compatible pathogens of oranges and apples, respectively, meanwhile *P. expansum* and *P. digitatum* were described as non-host pathogens in oranges and apples, respectively (Fig. 2.2).

Fruit defenses are complex and constitute a multilevel series of structural and biochemical barriers that are both constitutive and inducible. Therefore, a multidisciplinary approach including pathological, biochemical and molecular studies were conducted to unravel the underlying mechanisms related to fruit response against both postharvest pathogens. In particular, in this chapter we have focused our research in the following aspects: (1) infection capacity of each pathogen in oranges and apples; (2) characterization of fruit healing process; (3) identification and expression analysis of genes involved in fruit defence mechanisms.



Fig. 2.2 Compatible and non-host interaction in apple and oranges

### Infection Capacity of Both Pathogens on Apples and Oranges

The infection capacities of *P. digitatum* on oranges and of *P. expansum* on apples take into account several key factors such as i) maturity of fruit, ii) pathogen inoculum concentrations and iii) storage temperatures (Fig. 2.3).

Vilanova et al. (2012a, b) clearly defined the compatible and non-host interactions at both studied pathosystems. The main results showed that in compatible interactions the lowest growth rate of both pathogens was observed in the immature harvest fruit stored at both 20 °C and cold temperatures (4 °C and 0 °C for oranges and apples, respectively). In addition to that, the lower growth rate was accentuated at lower concentration of inoculum of pathogens as 10<sup>4</sup> conidia mL<sup>-1</sup>. Several authors have observed higher susceptibility of overripe apples compared to immature apples against *P. expansum* and *Botrytis cinerea*, which highlight the role of maturity in the susceptibility of fruit to fungal diseases (Torres et al. 2003; Neri et al. 2010; Su et al. 2011). Furthermore, in oranges no differences were observed between the growth rates of *P. digitatum* in oranges stored at both 20 °C and 4 °C, at the concentration of 10<sup>7</sup> conidia mL<sup>-1</sup> at any maturity stage. This fact could be explained because the inoculum concentration is too high that the maturity states of fruit do not affect the growth rate of this pathogen.

In apples and oranges, the storage temperature influenced the growth rate and visual symptoms of the decay of these pathogens and need shorter incubation times at 20 °C than at cold temperature (Vilanova et al. 2012a, b). These results coincide with those obtained by Baert et al. (2007a, b) in apples inoculated with different strains of *P. expansum*. Moreover, in *in vitro* studies in PDA medium, Buron-Moles et al. (2012) showed a decrease in the growth rate of *P. digitatum* and *P. expansum* at cold temperature (4 °C and 0 °C, respectively) compared to 25 °C.

Regarding inoculum concentration, in general no significant differences were observed at any concentration of the pathogen tested (from  $10^4$  to  $10^7$  conidia mL<sup>-1</sup>). Immature apples inoculated with *P. expansum* and stored at 20 °C, however, showed



a different growth rate dependent on inoculum concentration, in contrast, these differences were not observed in oranges (Vilanova et al. 2012a, b). In a similar work, Morales et al. (2008) observed differences in the growth rate of *P. expansum* at different inoculum concentrations in apples stored at 20 °C.

On the other hand, considering the non-host interaction, e.g. P. expansumoranges and P. digitatum-apples, these studies demonstrated that they became compatible interaction depending on the conditions and, therefore, they cannot strictly considered non-host pathogens. Previous studies conducted by Macarisin et al. (2007) have already shown that P. expansum could germinate and develop decay in citrus fruit but only at the point of inoculation. These authors only observed green mould in apples after treatment of wounds with different organic acids (citrus, ascorbic and oxalic) or with the enzyme catalase (CAT) before inoculation, pointing out the role of these substances in the suppression of the production of  $H_2O_2$ . Vilanova et al. (2012b) showed the ability of P. expansum to infect oranges by itself, without any pre-treatment, at 20 °C and 4 °C. However, incidence and severity of these infected oranges were higher at 4 °C than at 20 °C (Fig. 2.4a). This behaviour can be explained because wound healing, as fruit defence mechanisms, is slower at 4 °C than at 20 °C, together with the fact that *P. expansum* is a pathogen welladapted to low temperatures. Studies carried out by Ismail and Brown (1975) already pointed out that healing of "Valencia" oranges at 5 °C was slower than at 30 °C. Later, Brown and Barmore (1983) showed that both phenolic and lignification substances could be the responsible of resistance of curing oranges against *P. digitatum* infection. In turn, the formation of lignin increases at 20 °C more than at 2.5 °C (Mulas et al. 1996).

In contrast, Vilanova et al. (2012a) demonstrated that *P. digitatum* showed some visual symptoms of decay in apples but only restricted at the site of infection (Fig. 2.4b). Only overmatured apples were decayed by the highest concentration of *P. digitatum* (10<sup>7</sup> conidia mL<sup>-1</sup>), but exceptionally (Fig. 2.5). In addition, Buron-Moles et al. (2012) corroborated these results visualizing the infection process in apples using *P. digitatum* tagged with a green fluorescent protein (GFP). Likewise,





**Fig. 2.5** Green mould caused by *P. digitatum* on (**a**) immature, (**b**) matures, and (**c**) overmatured apples at inoculum concentration of  $10^7$  conidia mL<sup>-1</sup> and storage temperature of 20 °C

Fig. 2.6 Orange and apple inoculated with *P. expansum* and *P. digitatum*, respectively, with only visible reaction around the inoculation site



Louw and Korsten (2014) have also revealed that *P. digitatum* can infect different varieties of apples and pears. In contrast, at 0 °C, no signs of infection by *P. digitatum* were observed at any studied conditions. The overall results may explain by the ecophysiological difference between *P. digitatum* and *P. expansum* (Plaza et al. 2003; Buron-Moles et al. 2012).

A relevant fact in these non-host interactions were detected: when P. expansum and P. digitatum were not able to infect oranges and apples, an important reaction in the skin and pulp was observed (Fig. 2.6). In addition, this reaction increased proportionally to pathogen concentration and decreased as maturity advanced. This reaction could be associated to a rapid localized necrosis of non-host resistance (Mysore and Ryu 2004). Macarisin et al. (2007) in lemons also observed that approximately 4-5 days after inoculation with the non-host pathogen P. expansum appeared the hypersensitive response (HR), with dead cells and lignified tissue surrounding the wounds. HR usually appears in incompatible interactions (Mysore and Ryu 2004) and is recognized by brown necrotric areas resulting from a localized collapse of tissues as well as dead cells at the point of inoculation. In studies in coffee leaves (Silva et al. 2002) and melon (Romero et al. 2008), it was observed rapid HR in incompatible interactions. Whilst this response has been described to prevent the development of biotrophic fungi (Lamb and Dixon 1997; Lu and Higgins 1999), it may not prevent the consequent development of necrotrophic fungi (Mayer et al. 2001). In turn, some authors suggest that it may even stimulate the development of necrotrophic fungi (Govrin and Levine 2000).

In summary, our studies demonstrated that the inoculation of a compatible pathogen (*P. digitatum*-oranges and *P. expansum*-apples) at different inoculum concentrations always showed decay development. However, in oranges and apples, the most important differences in rot dynamics among harvests were found at the lowest inoculum concentration analysed ( $10^4$  conidia mL<sup>-1</sup>). Surprisingly, depending on the combination of factors (maturity stage, inoculum concentration and storage temperature), the non-host interaction (*P. expansum*-oranges and *P. digitatum*-apples) became compatible.

## **Orange and Apple Defence Mechanisms**

Lack of penetration by fungi into non-host tissue can be associated with the deposition of substances or compounds in the host epidermal cell walls. Based on above results, a study of the host defence responses to abiotic (wound) and biotic (pathogen) stresses was conducted following a pathological, biochemical and molecular approach to elucidate the defence reactions underlying in these interaction (Fig. 2.7).

## Wound Response Process

*P. digitatum* and *P. expansum* are considered as wound fungi, i.e., they need a wound on the fruit epidermis in order to start the infection (Kavanagh and Wood 1967; Spotts et al. 1998). Hence, good handling practices during harvest to avoid wounds in fruit are important to reduce postharvest losses. Beside that, a better knowledge on healing process of apples and oranges is needed.





**Fig. 2.8** Healing process in (**a**) oranges wounded and inoculated at different times with *P. digitatum* and (**b**) apples wounded and inoculated at different times with *P. expansum* and incubated at 20 °C and 85% RH

It is known that different treatments (curing, light UV) or inducing substances increased the resistance of fruit against post-harvest pathogens (Droby et al. 1993; Droby et al. 1999; Ballester et al. 2010, 2011; Shao et al. 2010). Vilanova et al. (2013, 2014a) studied the healing process *per se* of apples and oranges at different maturity stages when were wounded and infected after different times by *P. expansum* and *P. digitatum*. The healing process in both oranges and apples has an important effect limiting infection ability of both host pathogens when the fruit were stored at 20 °C (Fig. 2.8).

In addition, this effect was most outstanding in the immature and mature fruit than in overmatured. Regarding immature and commercial oranges, it was observed a significantly decrease in incidence and severity of wounds inoculated with *P. digitatum* 7 and 10 d after wounding, while on apples, a similar reductions in incidence and severity of blue mould it was achieved only 3 d after wounding (Vilanova et al. 2013, 2014a). Somehow, the skin of the oranges may be the responsible of this higher susceptibility to rot compared to apples. Brown et al. (1978) showed a reduction in the percentage of infected oranges by *P. digitatum* 3 d after wounding, whereas Su et al. (2011) and Shao et al. (2010) demonstrated a decrease in incidence of decayed apples by *B. cinerea* and *P. expansum*, respectively 96 h after wounding. In pears, Spotts et al. (1998) found greater resistance of fruit to *P. expansum* infection 2 d after wounding. Likewise, the greatest resistance was in green peppers inoculated with *Colletotrichum acutatum* only 1 h after wounding (Kim et al. 2008).

The studies on wounding process on oranges and apples demonstrated that it depends on temperature (Vilanova et al. 2013, 2014a). Both temperature and relative humidity are two of the most important factors affecting the process of healing

(Brown 1989). The temperature must be high enough to accelerate the development of metabolic reactions involved in the process of healing and, in turn, relative humidity should be sufficient to avoid drying or killing of tissues near damaged cells. Generally, temperatures above 10 °C and relative humidity higher than 85% are necessary to start the healing process in citrus fruit (Brown 1989). Moreover, the incidence and severity of wounded oranges inoculated with P. digitatum and stored at 4 °C for 30 d were lower when healing time increased. However, 45 d after inoculation, in commercial and overmatured oranges, there were no differences in the incidence between the different days of healing (Vilanova et al. 2013). This indicates that low temperatures make the healing process not effective enough to prevent infection with P. digitatum. In apples stored at 0 °C, the healing process was even less effective preventing infections of P. expansum than that observed in oranges. Lakshiminarayana et al. (1987) observed that apples inoculated with P. expansum or B. cinerea at 4 d after wounding and stored at 5 °C presented a substantial resistance to infection, although these authors considered that this healing time was scarce time to modify the cell wall to stop infection.

In contrast to these results, overmatured fruit showed few differences at different healing times, hence, once again it was clearly demonstrated the key role of maturity in susceptibility of oranges and apples to postharvest pathogen infection. Similar results were observed by Su et al. (2011) in overmatured apples inoculated with *B. cinerea* 96 h after wounding wounds: apples showed a higher incidence and severity of grey mould that immature or commercial apples. However, they did not observe differences in incidence of decay among harvest time when fruits were inoculated just after wounding.

Fruit maturity is a process that involves significant changes in the biochemical level such as alteration of the cell membrane (Cantu et al. 2008a). Consequently, maturity can modify the susceptibility of fruit by decreasing their defence responses and making them more susceptible to pathogen attack. As described above, overripe can trigger that the non-host *P. expansum* may even infect and develop decay in oranges. Some authors (Stange et al. 2002) defined that citrus skin is an inappropriate and even toxic environment for germination and growth of non-host moulds. In our study related to wound response, however, observed that *P. expansum* was able to infect and even develop rot at different times of healing in commercial and overmatured oranges at 20 °C, and at any maturity stage at 4 °C (Vilanova et al. 2013). Similar results were also observed in apples at 20 °C; *P. digitatum* was able to infect overmatured apples at 20 °C, even those inoculated 10 d after wounding, but not at 0 °C.

#### Cell Wall Reinforcement

Wound healing process could be associated with a limited accumulation of phenolics and lignin-like compounds in cell wall thickening around wounds and, hence, be an effective protection against pathogen infections. In our studies, fruits inoculated with the non-host pathogen (oranges inoculated with P. expansum and apples inoculated with *P. digitatum*) showed a visible reaction around wounds when pathogen was not able to infect them (Fig. 2.6). To identify the substances involved in this reaction, a more in-depth study was conducted. Previous authors showed a direct relation of deposition of some substances and orange resistance against fungal infections (Brown and Barmore 1983; Baudoin and Eckert 1985). Therefore, different histochemical tests to detect lignin, suberin and callose and to characterize the accumulation of substances at the site of inoculation were used. Lignin was detected using Maüle and toluidine blue O reactions, suberin by Sudan IV test, and callose by Aniline blue test (Fig. 2.9). Both tests used to detect lignin showed a positive reaction at short response times (24 h and 48 h for apples and oranges, respectively) in both compatible and non-host interactions. These results suggest that lignin production is not exclusive for compatible pathogens. Remarkably, reaction of control fruits (fruits wounded) was not equal on oranges and on apples: wounded apples showed a positive reaction in lignin test at short time, while oranges always showed a negative reaction. Therefore, lignin production seems to be related with abiotic (wounds) and biotic (pathogen) stresses, and appears more important in apples than in oranges.

Moreover, lignification observed in both oranges and apples inoculated with the non-host pathogen at long-term of response (7 d) was stronger than at 24 h or 48 h. In addition, lignification was apparently more significant in immature fruit than in commercial fruit since non-positive reaction was observed in overmatured fruit.

Neither of the tests used in Vilanova et al. (2012a, b) to detect suberin and callose showed a positive reaction either in oranges or in apples. According to Lai et al. (2003) using nuclear magnetic resonance spectroscopy (NMR spectroscopy) the substances present in curing grapefruit were suberin and not lignin. Regarding the quantification of lignin, Vilanova et al. (2013, 2014a) showed that immature control (mock inoculation at different times after wounding) in oranges and apples increased lignin content along storage period. Su et al. (2011) also observed an increase in lignin content in immature "Gala" apples compared to overmatured ones. Valentines



**Fig. 2.9** Images of histochemical tests: (**a**) Maüle reaction, (**b**) Blue Toluidine and (**c**) Sudan IV, in immature apples inoculates inoculated with *P. digitatum* at the concentration of  $10^7$  conidia mL<sup>-1</sup> after 7 d at 20 °C and 85% RH. Pictures (**a**) and (**b**) show a positive reaction to lignin, and (**c**) test Sudan IV shows a negative reaction to suberin

et al. (2005) also found higher lignin content in apples which denoted a major resistant to *P. expansum* infection. Overall results suggest that lignin has a role in apple defence against *P. expansum*.

Immature oranges and apples inoculated with the non-host pathogen showed an important increase in lignin content 7 d and 72 h after inoculation, respectively. These results are in agreement with histochemical results obtained by Vilanova et al. (2012a, b), in which the most intense reaction was observed in immature fruits 7 d after inoculation. In the non-host interactions, lignification could begin few days after infection; however, its accumulation was more remarkable at long-term response. In addition, the highest accumulation of lignin was observed in immature fruit that usually are not infected.

Regarding the lignin content in compatible interaction, our studies showed unusually high absorbance values. These values obtained with the acetyl bromine method may be attributed to an increase in polysaccharides degradation and suggests that other factors than solubilisation of lignin were responsible for the increase in absorbance (Hatfield et al. 1999). When necrotrophic pathogens infect fruits, they secrete various cell wall degrading enzymes that increase the polysaccharides (Cantu et al. 2008b). Therefore, the acetyl bromide method applied to macerated tissues could produce erroneous absorbance values.

### Molecular Studies in Fruit-Penicillium Interactions

The availability of the complete genome sequence of fruit has greatly aided the development of "omic" approaches such as genomics, transcriptomics and proteomics. Specifically, in *Malus* x *domestica*, the information gained from those studies could be particularly useful in the field of postharvest pathology. In our studies, a transcriptomic analysis of apple to understand fruit disease resistance in response to compatible (*P. expansum*) and non-host (*P. digitatum*) pathogens was conducted (Vilanova et al. 2014b). Moreover, the expression of several genes involved in the phenylpropanoid pathway were quantified to define their role in citrus fruit response against pathogen and non-host pathogen (Vilanova et al. 2013).

# Transcriptomic Approach in Apples

The first transcriptomics study to elucidate the genes involved in the fruit development of "Fuji" apples was conducted by Lee et al. (2007) using a microarray containing 6,253 cDNAs. Few years later, Soria-Guerra et al. (2011) developed a microarray containing approximately 40,000 probes, including positive and negative controls, from 34 libraries of cDNA constructed from vegetative and reproductive apple tissues at different developmental stages, different genotypes and different types of biotic and abiotic stresses. This microarray was used and validated in

different studies: in one of them the aim was to characterize the gene expression along fruit development of different varieties of apple (Soria-Guerra et al. 2011); in another one the goal was to study the response of apples after the application of an abscision-promoting treatment (Zhu et al. 2011); in the last ones the objectives were to investigate the response of apples to *Erwinia amylovora* (Bocsanczy et al. 2009; Sarowar et al. 2011). The microarray was also used in our study (Vilanova et al. 2014b) to perform a global analysis of genes involved in response of immature apples to a compatible pathogen (*P. expansum*) and a non-host pathogen (*P. digitatum*) at 20 °C (Fig. 2.10). The results of this study indicated a different pattern of apples in response to *P. expansum* and *P. digitatum* infection. Functional annotation based on gene ontology terms (GO) showed that biological processes of apples significantly modified in response to *P. expansum* were related to defence responses and oxidative stress, meanwhile to *P. digitatum* were related to phenylpropanoid pathway. Our microarray study identified 3,770 differentially expressed probes that could be used to further study of apple-*Penicillium* spp.

One of the first defence responses detected in a host-pathogen interaction is the oxidative burst that is characterized by a significant increase in ROS levels (Levine et al. 1994; Low and Merida 1996). Plant nicotinamide adenine dinucleotide phosphate oxidases, also called "Respiratory burst oxidases homologues (Rbo)" was described as an important source of ROS species in majority of plant–pathogen interactions (Torres et al. 2006). ROS metabolism is controlled by Rbos, and other enzymes including superoxide dismutase (SOD), catalase (CAT), and ascorbate peroxidase (APX) (De Gara et al. 2003). SOD activity plays a role in the dismutation of superoxide radicals, while CAT and APX activity contribute to the elimination of hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) (Levine et al. 1994). Our microarray data showed



Fig. 2.10 General scheme to conduct the global study in apple in response to *P. expansum* (pathogen) and *P. digitatum* (non-host pathogen)

an induction of the expression of the Rbo gene in apples inoculated with P. digitatum compared to apples inoculated with P. expansum. First, this fact pointed out that an increase in ROS content could be related to defence response of apples against P. digitatum. On the other hand, apples inoculated with P. expansum showed a greater increase in the expression of genes encoding ROS-detoxifying enzymes such as SOD, APX and POX (peroxidase) compared to apples inoculated with P. digitatum. Nevertheless, there was no increase in the expression of the Rbo gene when the apples were inoculated by *P. expansum*, indicating that the highest level of Rbo expression may be earlier than 24 h. Torres et al. (2003) showed that the largest production of H<sub>2</sub>O<sub>2</sub> in apples inoculated with P. expansum awas detected approximately 6 h after inoculation. This ability of P. expansum to prevent the oxidative burst, suppressing the production of  $H_2O_2$  in apples 24 h after inoculation, may be closely related to its pathogenicity. These results are in agreement to those obtained by Macarisin et al. (2007) in oranges where they observed that P. digitatum (compatible pathogen) suppressed the oxidative burst on host cell, meanwhile P. expansum (non-host pathogen) triggered an accumulation of  $H_2O_2$ . The relationship between ROS levels and detoxifying enzyme levels is considered to be a key factor in determining the  $H_2O_2$  and  $O_2^-$  tissue-levels (Mittler et al. 1999). Ballester et al. (2006) suggested an imbalance among the activities of enzymes that generate and degrade H<sub>2</sub>O<sub>2</sub> to promote the infection process, as has been reporter for the necrotrophic fungi B. cinerea and Sclerotinia sclerotiorum (Govrin and Levine 2000). Pathogenicity of both moulds seems to be dependent on ROS levels generated during the infection because they may benefit from cell necrosis caused by a hypersensitive response (HR) (Govrin and Levine 2000). Additionally, the bcsod1 gene encoding for a Cu-Zn SOD has been described in B. cinerea as a virulence factor (Rolke et al. 2004). Other study also shows that the virulence factor of Monilinia fructicola MfCUT1 is induced in the presence of ROS generated during infection process (Chiu et al. 2013).

The microarray data also indicated a change in the gene expression of heat shock proteins (HSPs). HSPs are one class of chaperones produced in response not only to high temperatures if not to stress, to exposure to heavy metals, pathogens attack and disturbance in intracellular calcium levels (Vierling 1991). In our study, we observed an induction in the *HSP101* gene expression in apples inoculated with *P. expansum* and in the *HSP70* gene expression in apples inoculated with *P. expansum* and in the *HSP70* gene expression in apples inoculated with *P. expansum* and *P. digitatum* (Vilanova et al. 2014b). These results indicated that *HSP101* might be more specific for pathogen compatible although *HSP70* may be related to the pathogen attack. Little is known about the role of HSPs in response to pathogens because most of the HSP studies have been focused on plant abiotic stresses. In fruits, for example, the expression HSP has been characterized in response to heat treatments used for control post-harvest diseases (Lauxmann et al. 2012; Pavez et al. 2013). Specifically, Yun et al. (2013) and Pavoncello et al. (2001) observed an induction of genes encoding HSP in heat-treated citrus fruit related with an increase of resistance to *P. italicum* and *P. digitatum*.

A differential expression in other genes related to defence-related proteins was observed, as genes encode for pathogenesis-related proteins (PR-proteins), and

other related to synthesis of callose (Vilanova et al. 2014b). The PR proteins are divided into 17 families, including glucanases (GLU), chitinases (CHI), taumatins (TAU), peroxidase (POX) and defensins (DEFL) (van Loon et al. 2006; van Loon and van Strien 1999). The results showed that the inoculation of apples with P. expansum induced more PR proteins than P. digitatum, including two CHI, one endoglucanase (EGL), a TAU and a DEFL proteins. The outcomes indicated that induction of genes encoding PR proteins are related to fruit defence against pathogenic fungi. Concerning fruit, there is scarce information on induction of PR genes in response to pathogens. Most of the studies are based on the induction of PR proteins in response to PR proteins biological control and abiotic stress conditions. Ballester et al. (2010) correlated an increase in CHI and GLU enzymes activity with upregulation of genes encoding these enzymes when oranges are firstly cured and after that inoculated with *P. digitatum*. On the other hand, in oranges, Hershkovitz et al. (2012) detected an induction of the CHI and in apples, Ouaglia et al. (2011) detected an induction of the CHI, GLU and TAU when the fruits were treated with biocontrol agents. Buron-Moles et al. (2014) in proteomic studies, revealed a correlation between an increase in the amount of TAU protein and wound response of apples.

An induction of the expression of  $\beta$ -glucosidase13 and  $\beta$ -glucosidase40 genes in response to inoculation by *P. digitatum* and *P. expansum*, respectively, was also observed (Vilanova et al. 2014b). Sánchez-Torres and González-Candelas (2003) found that the glycosidase gene was specifically expressed in apple tissue infected by *P. expansum*. The first evidence of the expression of the  $\beta$ -glucosidase gene in plants was reported by Simmons et al. (2001), that correlated upregulation of the *rhm1* corn mutant to resistance to *Bipolaris maydis*.

# Gene Expression of Genes Involved in the Phenylpropanoid Pathway

Because of the involvement of lignin in defence processes of oranges and apples against biotic and abiotic stresses, a gene expression study of several phenylpropanoid pathway-related genes was conducted for apples (Vilanova et al. 2014b) and oranges (Vilanova et al. 2013).

#### Apples

Microarray study revealed changes in the expression of genes involved in phenylpropanoid pathway, as the phenylalanine ammonium liase (*PAL1* and *PAL2*), caffeic acid O-methyltransferase (*COMT*) and cinnamoyl-CoA reductase (*CCR*) genes, in apple inoculated with *P. digitatum* (non-host pathogen). Transcription levels of some genes (*PAL1*, *PAL2*, *COMT2* and *POX64*) were evaluated in apples at immature and commercial harvest, inoculated with *P. expansum*, *P. digitatum* and water (control) at 8, 24 and 48 h post inoculation (hpi) using the quantitative PCR (qPCR) technique.

At 24 hpi, higher transcript abundance of *PAL1*, *PAL2* and *POX64* genes was observed for both apples inoculated by the non-host pathogen (*P. digitatum*) than other treatments. In contrast, at 48 hpi the highest increase in gene expression was found in *PAL1*, *PAL2*, *COMT2* and *POX64* in apples inoculated with *P. expansum*. Ballester et al. (2013) found that some genes related to the phenylpropanoid pathway showed their maximum expression at 48 hpi and decreased at 72 hpi. These variances in the expression along the time may be due to differences in inoculum concentrations. Furthermore, these results are in agreement with that observed in histochemical studies: in a short-term response both fungi showed a positive reaction to lignin, while in a long-term response only a positive reaction was obtained for the non-host pathogen, because the other samples were completely rotten (Vilanova et al. 2012a). Overall results support the hypothesis that apple attempt to defend against both pathogens, although host pathogen is finally able to overcome apple defences and develop infection.

Regarding apple maturity stages, microarray study indicated that *PAL1*, *PAL2* and *POX64* genes showed a greater abundance in commercial than in immature apples. Both PAL and POX belong to a family of enzymes that are involved in various biochemical functions in plants as growth, development and senescence. High PAL activity has also been associated with the maturity stage of apples (Wang et al. 2000) and strawberries (Cheng and Breen 1991; Villarreal et al. 2010) due to the accumulation of anthocyanins and other phenolic compounds. An increase in POX activity during ripening has been observed in apples (Torres et al. 2003) and plums (Singh et al. 2012). Therefore, the observed increase in the expression of *PAL1*, *PAL2* and *POX64* genes may be more related to fruit ripening rather than defence responses.

#### Oranges

In previous studies, Ballester et al. (2011, 2013) analysed the gene expression of several genes involved in the phenylpropanoid pathway and the modifications in phenolic compounds associated with citrus resistance against *P. digitatum*. In Vilanova et al. (2013), some of these genes were analysed in immature oranges infected by *P. digitatum* but also by *P. expansum*, because the response of citrus fruit to non-host pathogen remained unexplored.

In oranges, the expression of *PAL1*, *COMT1*, *POX1*, cinnamyl alcohol dehydrogenase (*CAD*) and sinapyl alcohol dehydrogenase (*SAD*) genes was analyzed after inoculation with *P. digitatum*, *P. expansum* and water (control), 24 and 48 hpi using the semi-quantitative PCR technique (Vilanova et al. 2013). Gene expression analysis showed different behaviour depending on the time after inoculation. At 24 hpi, oranges inoculated with the compatible pathogen (*P. digitatum*) showed a higher expression of the *PAL1* and *COMT1* genes compared to oranges inoculated with *P. expansum* and control, although the *POX1* gene showed greater expression in oranges inoculated with both moulds than control. At 48 h, however, oranges inoculated with *P. digitatum* displayed an important decrease in expression of *PAL1*, *COMT1*, *POX1* and *SAD* genes, and oranges inoculated with *P. expansum* and control the gene expression remained constant. This decrease in expression seems to indicate that the compatible pathogen is able to suppress the gene expression related to the phenylpropanoid pathway and therefore start the infection.

#### **Concluding Remarks**

Our studies suggest that maturity stage is an important factor to be considered on pathological studies. Therefore, it would be interesting to continue exploring the relationship between maturity factors and fruit susceptibility in response to pathogens. Besides, the increase of resistance or susceptibility of apples and oranges to host and non-host pathogens depend on it and may vary based on the climacteric or non-climacteric respiration pattern of fruit.

Depending on the combination of certain conditions as maturity stage, inoculum concentration and storage temperature, the non-host pathogen become compatible pathogen in these fruit –pathogen interactions. Furthermore, wounded fruit stored at cold temperatures are, in general, more susceptible to pathogen attack. Lignification is a key factor in fruit defence to both biotic and abiotic stresses together with other multiple reactions. This process was apparently more important in immature than in commercial and over-mature fruit.

Finally, the global study conducted on apples give us an overall picture of metabolic pathways to may activate in response to host and non-host pathogens to continue to work in this research line with the aim of unravelling the fruit defence mechanisms.

Overall knowledge obtained over the time-course of our studies can help us to establish the basis towards the improvement of postharvest control strategies in apples and oranges.

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