Pathogen profile **Pantoea ananatis: an unconventional plant pathogen**

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SUMMARY

Pantoea ananatis causes disease symptoms in a wide range of economically important agricultural crops and forest tree species worldwide. It is regarded as an emerging pathogen based on the increasing number of reports of diseases occurring on previously unrecorded hosts in different parts of the world. Its unconventional nature lies in the fact that, unlike the majority of plant pathogenic microbes, *P. ananatis* is capable of infecting humans and occurs in diverse ecological niches, such as part of a bacterial community contaminating aviation jet fuel tanks and contributing to growth promotion in potato and pepper.

Taxonomy: Bacteria; *Gammaproteobacteria*; family *Entero-bacteriaceae*; genus *Pantoea*.

Microbiological properties: Gram-negative; facultatively anaerobic; most strains are motile and produce a yellow pigment in culture; indole positive.

Biology: Pantoea ananatis is a common epiphyte; it also occurs endophytically in hosts where it has been reported to cause disease symptoms and in hosts where no such symptoms have been described. Some strains are ice-nucleating, a feature which has been used as a biological control mechanism against some insect pests of agricultural crops and by the food industry.

Disease symptoms: *Pantoea ananatis* infects both monocotyledonous and dicotyledonous plants. The symptoms are diverse depending on the host infected, and include leaf blotches and spots, die-back, and stalk, fruit and bulb rot.

Biological control agent: *Pantoea ananatis* has both antifungal and antibacterial properties. These characteristics have the potential of being exploited by biological control specialists.

INTRODUCTION

The appearance of new and re-emerging plant diseases is having a significant constraint on agricultural productivity worldwide (Bandyopadhyay and Frederiksen, 1999; Bright, 1998; Palm, 1999).

* Correspondence: Tel.: +27-12-420 3934; Fax: +27 12 420 3960; E-mail: teresa.coutinho@ fabi.up.ac.za The extent of this threat has increased dramatically over the past two decades as a result of increased movement of agricultural products between countries. The threat of these emerging diseases is, however, not restricted to cultivated plants, as native and wild plants are also at risk (Anderson *et al.*, 2004). This is of considerable importance in terms of biodiversity conservation.

An emerging plant pathogen is considered to be one which has shown an increase in incidence, host and geographical range (Anderson et al., 2004). Since its initial discovery on pineapple, causing fruitlet rot in the Philippines in 1928 (Serrano, 1928), Pantoea ananatis has been found to cause a wide range of symptoms on both monocotyledonous and dicotyledonous plants. Its occurrence in these hosts leads to sporadic disease outbreaks, resulting in severe economic losses. From 1983 to date, the host range of *P. ananatis* has increased to eight, and the pathogen has now been reported to cause plant disease symptoms in at least 11 countries. The increase in geographical range is best illustrated by the distribution of palea browning of rice. This disease was initially reported from Japan (Tabei et al., 1988), but now also occurs in Korea (Kim et al., 1989), the Philippines (Xie, 1996 cited by Xie, 2001), China (Xie, 2001) and, more recently, Italy (Cortesi and Pizzatti, 2007).

Pantoea ananatis is a ubiquitous bacterium which, in itself, is not unique as a number of other bacterial plant pathogens share this characteristic. However, *P. ananatis*, when it is not associated with plants as an epiphyte, endophyte, pathogen or symbiont, also occupies diverse and unusual ecological niches. Here it may function as a saprophyte. In the case of humans, *P. ananatis* has also been reported to cause bacteraemia (De Baere *et al.*, 2004). With the exception of perhaps only *Pantoea agglomerans*, no other plant pathogen behaves in such an unconventional manner. An explanation of how *P. ananatis* has become so broadly adapted to these different habitats will probably only be resolved once species-specific genes have been identified.

Pantoea ananatis is not only important in the abovementioned roles. Its ice nucleation activity has been exploited by both the food industry and biological control specialists of insects. Extracellular ice nucleators from *P. ananatis* have been tested and applied in the freezing of foods in order to obtain the desired texture (Zasypkin and Lee, 1999), and in the freeze-drying of foods (Watanabe and Arai, 1994). Ice-nucleating strains of *P. ananatis* also markedly reduce the cold hardiness of mulberry pyralid larvae (Watanabe and Sato, 1999), and thus these strains have the potential to act as biological control agents of insect pests. In addition, the accomplishment of researchers in Switzer-land to genetically modify rice to produce 'yellow rice' owes its success to *P. ananatis*. In this situation, the phytoene desaturase from *P. ananatis* (a strain identified as *Erwinia uredovora*) was used to introduce the β -carotene biosynthesis pathway into rice (Beyer *et al.*, 2002).

In this review, we focus on the taxonomy, detection and identification of *P. ananatis*. We also discuss its role in different ecological niches with a focus on its association with plants as an epiphyte, endophyte and plant pathogen. The molecular basis for its role in these different niches is, however, not well understood.

TAXONOMY

Until recently all phytopathogenic *Enterobacteriaceae* belonged to a single genus *Erwinia*. The genus was proposed by Winslow *et al.* (1920) for all plant-associated, Gram-negative, non-sporeforming, peritrichous, fermentative, rod-shaped bacteria. Dye (1968, 1969a,b,c) divided the genus into four 'natural' clusters, namely the amylovora, carotovora, herbicola and 'atypical' *Erwinia* groups. The 'herbicola' group, of interest in this review, consisted of *Erwinia* strains that usually produced a yellow pigment in culture and related non-pigmented clinical isolates often named *Enterobacter agglomerans*. This group is referred to in the literature as the *Erwinia herbicola–Enterobacter agglomerans* complex. Many researchers have subsequently shown that this complex is heterogeneous and contains bacterial strains with different phenotypes and genotypes (Brenner *et al.*, 1984; Mergaert *et al.*, 1983, 1984; Verdonck *et al.*, 1987).

In 1989, Gavini *et al.* proposed the genus *Pantoea* (Gavini *et al.*, 1989). The type strains of *E. herbicola*, *Ent. agglomerans* and *E. milletiae* were found to belong to the same DNA hybridization group (Beji *et al.*, 1988), and thus the combined species, *P. agglomerans*, was proposed for these bacteria (Gavini *et al.*, 1989). Other species included in this genus are *P. dispera* (Gavini *et al.*, 1989), *P. punctata*, *P. citrea*, *P. terrea* (Kageyama *et al.*, 1992), *P. stewartii*, *P. stewartii* ssp. *indologenes*, *P. stewartii* ssp. *stewartii* and *P. ananatis* (Mergaert *et al.*, 1993). *Pantoea ananatis* was first described by Serrano (1928) as *Erwinia ananas*. Mergaert *et al.* (1993) proposed the name *P. ananas*, which was corrected to '*ananatis*' by Trüper and De'Clari (1997).

Pantoea ananatis and P. uredovora are listed in Bergey's Manual of Determinative Bacteriology (Holt, 1977) as differing in their ability to reduce nitrate to nitrite. Both species also differ substantially from each other in host range. Pantoea ananatis was initially described as a pathogen that caused fruitlet rot of pineapple (Serrano, 1928), whereas P. uredovora was described as a pathogen of the basidiomycete rust fungus, Puccinia graminis (Pon et al., 1954). On the basis of a high level of genotypic relatedness, Mergaert et al. (1993) synonymized these two species. They found that seven strains received as either one of these two species, and including their type strains, exhibited between 76% and 100% DNA binding and constituted a single DNA hybridization group. It has previously been shown that the phenotypic differentiation of these strains is extremely difficult. However, this synonymy was supported by numerical analysis (Dye, 1981; Mergaert et al., 1984; Verdonck et al., 1987). Furthermore, Waleron et al. (2002) have recently shown that the two species are in the same *recA* polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP) group, and Brady et al. (2007) have shown that these strains also form part of the same fluorescent amplified fragment length polymorphism (F-AFLP) cluster. Despite this research, the synonymy of these two species is not widely accepted, and the name E. uredovora is still commonly used in the literature (Beyer et al., 2002).

ISOLATION, DETECTION AND IDENTIFICATION

Most general media, such as blood agar, nutrient agar, tryptic soy agar or media specifically designed for the isolation of the *Enterobacteriaceae*, for example MacConkey and Hektoen agar, can be used for the isolation of *P. ananatis* (Grimont and Grimont, 2005). From diseased plants, the bacterium is usually isolated using nutrient agar (Bruton *et al.*, 1991; Coutinho *et al.*, 2002; Gitaitis and Gay, 1997; Schaad, 2001), but other media, such as yeast extract–dextrose–calcium carbonate agar (Azad *et al.*, 2000) and King's Medium B (Cother *et al.*, 2004), have also been employed.

Currently, there are only a few semi-selective media for the isolation of *P. ananatis*. Hasegawa *et al.* (2003) developed NSCV-In medium for the selective isolation of pathogenic *P. ananatis* from diseased rice plants. PA 20 was developed for the isolation of *P. ananatis* from onion seed (Goszczynska *et al.*, 2006a,b), and was reported to inhibit the growth of most of the common saprophytes associated with the seed. Similar to the use of general isolation media to obtain *P. ananatis*, further tests to confirm the identity and pathogenicity of isolates are still required when using selective media.

The tentative identification of isolates as belonging to the genus *Pantoea* is usually performed using commercial identification systems (Azad *et al.*, 2000; Cortesi and Pizzatti, 2007) or sequencing of the 16S rRNA gene (Coutinho *et al.*, 2002). For this group of bacteria, 16S rRNA is, however, too highly conserved to differentiate reliably between closely related species (Stackebrand and Goebels, 1994). API 20E or Biolog systems also have problems in identifying specific *Pantoea* species accurately as they share many phenotypic characteristics. The production of indole by *P. ananatis* is often used to distinguish between this

species and *P. agglomerans* strains, but this characteristic is shared by strains of *P. stewartii* ssp. *indologenes* (Grimont and Grimont, 2005). The identity of *P. ananatis* strains is therefore often confirmed by means of other techniques, such as whole-cell fatty acid methyl ester profiles (Azad *et al.*, 2000; Cother *et al.*, 2004; Gitaitis and Gay, 1997; Schwartz and Otto, 2000), PCR-RFLP analysis of the *recA* gene (Waleron *et al.*, 2002), DNA–DNA hybridization (Coutinho *et al.*, 2002) or F-AFLP (Brady *et al.*, 2007).

At present, there is no recommended method for the direct detection of *P. ananatis* in plant material. Walcott *et al.* (2002) designed species-specific primers for the 16S–23S rDNA internal transcribed spacer region, but the best primer pair detected both *P. ananatis* and *P. stewartii* ssp. *stewartii*. These researchers also tried to overcome the interference of saprophytes on onion seed by enriching samples for *P. ananatis* by means of polyclonal immunomagnetic bead capturing before proceeding with PCR. Cortesi and Pizzatti (2007) used another primer set targeting the same region, but no data on the assay's specificity are available. An oligonucleotide probe for fluorescent *in situ* hybridization was also not species specific and detected both *P. ananatis* and *P. aqglomerans* (Nakanishi *et al.*, 2006).

Currently, the most promising approach for species assignment is the use of multilocus sequence analysis (MLSA) (Gevers *et al.*, 2005). Using sequence data from four housekeeping genes, Brady *et al.* (2008) showed that MLSA could clearly differentiate between all the current *Pantoea* spp. A high level of congruence was observed between the *gyrB* sequence data and DNA–DNA hybridization values, and it was recommended that accurate identification of *P. ananatis* strains could be achieved with comparison of either the *gyrB* (Fig. 1) or *rpoB* gene sequences (Brady *et al.*, 2008).

PANTOEA ANANATIS AS AN EPIPHYTE

Pantoea ananatis is a common epiphyte on host and non-host plants. Gitaitis et al. (2002) detected and cultured this bacterium as an epiphyte from 25 asymptomatic weed species, including crabgrass, sicklepod and yellow nutsedge, and from crop plants, such as Bermuda grass, cowpea and soybean. It has also been reported as an epiphyte on numerous economically important plant species, including rice (Watanabe et al., 1996), maize (Paccola-Meirelles et al., 2001), barley, buckwheat, uredospores of Ustilago smut of maize (Coplin and Kado, 2001), cotton lint (Chun and Perkins, 1997), mulberry (Takahashi et al., 1995), poplar trees (Zeng et al., 1999) and wheat heads (Legard et al., 1994). In many of these cases, the occurrence of *P. ananatis* on the plant surface has not been linked to a specific disease on the host from whence it was isolated as an epiphyte. However, these asymptomatic non-hosts could be providing a source of inoculum, causing disease outbreaks of susceptible hosts grown in their vicinity.

The occurrence of *P. ananatis* on plant surfaces may be beneficial to these plants. The bacterium has both antibacterial and antifungal activity in vitro and in vivo, thus protecting host plants against infection by other pathogenic fungi and bacteria. A strain, previously described as P. uredovora, was found to have in vitro antibacterial activity against Xanthomonas, Agrobacterium and all species belonging to the 'amylovora' group, including Erwinia amylovora (Vantomme et al., 1989). Isolates of P. ananatis from buckweed seed have been shown to have strong antifungal activity against Rhizopus spp. in vitro (limura and Hosono, 1998). In a study by Pajk (2004), it was reported that 'P. uredovora' reduced the infection of pome fruit trees by E. amylovora. A strain, CPA-3, of P. ananatis was discovered to rapidly colonize wounds of harvested pome fruit, protecting them against Penicillium expansum (Torres et al., 2005). Similarly, strains of *P. ananatis* have been found to protect tomato fruit against the grey mould fungus, Botrytis cinerea, by producing antifungal compounds (Enya et al., 2007). Rice blast caused by Pyricularia oryzae has also been controlled by a strain of *P. ananatis* transformed with a chitinolytic enzyme gene (*ChiA*) from Serratia marcescens (Simeya et al., 2003).

Pantoea ananatis is one of only a few species of bacteria shown to contain ice-nucleating strains (Abe et al., 1989; Obata et al., 1990). An ice nucleation active (INA) gene, inaA, has been sequenced and found to be similar to that of Pseudomonas species (Abe et al., 1989). This implies that frost formation on sensitive plants is induced at temperatures slightly higher than would normally occur when the bacterium is absent. Examples of frostsensitive plants are *Eucalyptus*, poplar, pea, pome and stone fruit trees (Lindow, 1983). Pantoea ananatis has been found to be the dominant INA bacterium on maize leaves in China (Sun et al. 2005), and has also been reported to cause frost injury to mulberry (Takahashi et al., 1995), tea (Goto et al., 1988), strawberry (Obata et al., 1990), apricot (Zhao et al., 2007) and citrus (Wang et al., 2008). The freezing injury caused by P. ananatis in poplar has resulted in an increased incidence of infection by the fungal pathogen Dothiorella gregaria in China (DePeng et al., 1999; Xiang et al., 2001; Zeng et al., 1999). The exact role of frost injury in outbreaks of many of the diseases caused by P. ananatis has not been elucidated clearly.

PANTOEA ANANATIS AS AN ENDOPHYTE

Although the term 'endophyte' is more commonly associated with fungi, bacteria are also capable of colonizing the interior of a plant. In this environment, they may be considered beneficial, neutral or existing as latent pathogens (Lodewyckx *et al.*, 2002). In recent years, *P. ananatis* has been isolated as an endophyte from a number of plants, including, for example, coffee (Nunes and de Melo, 2006) and ginseng (Cho *et al.*, 2007). Its role as an endophyte has mainly been found to be beneficial to the plant.

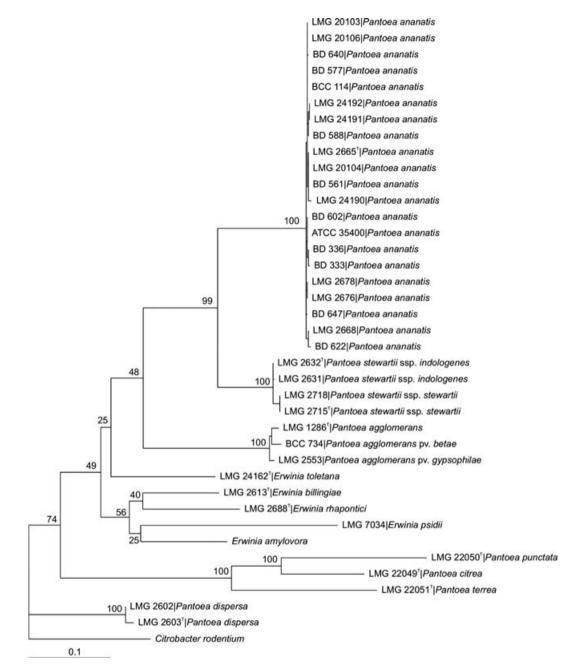


Fig. 1 Maximum likelihood tree based on the *gyrB* sequences of *Pantoea* species and their closest phylogenetic neighbours. Bootstrap values after 1000 replicates are expressed as percentages. *Citrobacter rodentium* was included as an outgroup. BD 333 and 336 were isolated from onion seed in South Africa, ATCC 35400 from honeydew melons in the USA, LMG 2676 and 2678 from *Puccinia graminis* in the USA and Zimbabwe, respectively, LMG 2668 and 2665 from pineapple in Hawaii and Brazil, respectively, LMG 20103, 20104, 20106 and BD 114 from *Eucalyptus* in South Africa, LMG 24190 from onion in the USA, and BD 561, 577, 588, 602, 622, 640, 647 and LMG 24191 and 24192 from maize in South Africa.

In pepper, *P. ananatis* promoted significant growth and induced systemic resistance against *Xanthomonas axonopodis* pv. *vesicatoria* (Kang *et al.*, 2007). Likewise, when shoot tips of papaya were inoculated with *P. ananatis* strains originally acquired as endophytic residents from within papaya tissue, they displayed significantly better root and shoot growth (Thomas *et al.*, 2007).

Strains found to occur endophytically in maize kernels have *in vitro* activity against *Lecanicillium aphanocladii* (Rijavec *et al.*, 2007).

There are few reports of *P. ananatis* behaving as a latent pathogen within susceptible host tissue. It has been reported as an endophyte in rice plants (Mano and Morisaki, 2008), and we

Table 1 Host range of Pantoea ananatis

Natural host	After artificial inoculation	Symptoms	Reference
Pineapple		Fruitlet rot	Serrano (1928)
	Sugarcane	Leaf streaks	Serrano (1928)
Sudangrass		Leaf blotches and streaks	Azad <i>et al</i> . (2000)
	Sorghum and oats	Leaf blotches and streaks	Azad <i>et al</i> . (2000)
Cantaloupe fruit		Brown spot	Bruton <i>et al</i> . (1991)
Honeydew melons		Brown spot	Ceponis <i>et al</i> . (1985)
Onions including giant onions		Leaf blight, seed stalk rot, bulb decay	Gitaitis and Gay (1997)
			Yumiko <i>et al</i> . (2005)
Eucalypts		Leaf blight, shoot tip die-back	Coutinho <i>et al</i> . (2002)
Maize		Necrotic spots and streaks	Paccola-Meirelles et al. (2001)
		Brown stalk rot	Goszczynska <i>et al</i> . (2007)
Rice		'Palea' browning	Tabei <i>et al</i> . (1988)
		Stalk rot	Cother <i>et al</i> . (2004)
Tomato		'Graywall'	Stall <i>et al</i> . (1969)
Puccinia graminis (causal agent of leaf rust of wheat)		Parasite of the rust	Pon <i>et al.</i> (1954)
King oyster mushroom		Soft rot	Kim <i>et al</i> . (2007)
	Cotton	Internal boll rot	Bell <i>et al.</i> (2007)
Watermelon			Walcott <i>et al</i> . (2003)

have also isolated *P. ananatis* as an endophyte from both susceptible and resistant *Eucalyptus grandis* \times *Eu. nitens* (GN) clones in South Africa (T. A. Coutinho and S. N. Venter, unpublished results). In susceptible clones, this pathogen causes leaf blight and dieback (Coutinho *et al.*, 2002). *Pantoea ananatis* also occurs endophytically in rice seed (Mano *et al.*, 2006; Okunishi *et al.*, 2005) and maize kernels (Rijavec *et al.*, 2007). Both are known hosts of this pathogen and its occurrence within seed may be of considerable epidemiological significance.

SAPROPHYTIC NATURE OF P. ANANATIS

Pantoea ananatis has been isolated from a diverse range of environments as a saprophyte. These include rivers (Morohoshi et al., 2007), soil (Lai and Hsu, 1974), aviation fuel tanks (Rauch et al., 2006), sorghum fermentation (Mohammed et al., 1991), from the rhizosphere of soft rush (Juncus effusus) (Halda-Alija, 2003) and on ticks (Murrell et al., 2003), and is a common inhabitant of the gut microflora of brown plant hoppers (*Nilaparvata lugens*) (Watanabe et al., 1996), mulberry pyralid (Glyphodes pyloalis) (Takahashi et al., 1995), cotton fleahoppers (Pseudatomoscelis seriatus) (Bell et al., 2007) and tobacco thrips (Frankliniella fusca) (Gitaitis et al., 2003; Wells et al., 2002), Pantoea ananatis has also been isolated from Pinus elliottii roots colonized by ectomycorrhizal fungi in Australia (Izumi et al., 2008). Many of these studies which show that *P. ananatis* is present as a saprophyte involved the analyses of bacterial communities either occurring in or contaminating a specific location. The identification of the inhabitants of these sites was mostly based on partial sequencing of the 16S rRNA gene and determining the similarity of sequences to those in GENBANK. This introduces an element of caution into the interpretation of these results, and one cannot state with absolute confidence that *P. ananatis* does indeed reside in these habitats.

PANTOEA ANANATIS AS A PATHOGEN

Host range and symptom expression

Pantoea ananatis infects both monocotyledonous and dicotyledonous plants (Table 1). This bacterium has also been deposited in numerous culture collections, where its role, as an epiphyte or pathogen, is not clearly indicated. These hosts include Cattleya sp. (LMG 2807), Musa sp. (LMG 2628), Cassia pectuta (ICMP 12183) and sugarcane (ICMP 10132). The symptoms caused by P. ananatis are diverse depending on the host infected, and include blotches and spots (Fig. 2A), die-back, fruit, boll and bulb rot (Fig. 2B), and decay (Fig. 2C). In the case of infection of honeydew melons and cantaloupes, the brown spot symptoms occur only after harvesting (Bruton et al., 1991; Wells et al., 1987). The disease apparently originates in the form of field infections that remain guiescent until the fruit ripens. Following the infection of young Eucalyptus trees or seedlings/cuttings with P. ananatis, the shoots wilt and die-back occurs (Coutinho et al., 2002). The pathogen appears to spread from the petioles into the main leaf veins, and from these parts into adjacent leaf tissue. In cases in which monocotyledonous plants are infected, symptoms are in the form of blotches and streaks, forming parallel to the main leaf vein (Azad et al., 2000; Paccola-Meirelles et al., 2001). Outbreaks of diseases caused by P. ananatis are usually sporadic, possibly as a result of its opportunistic nature.

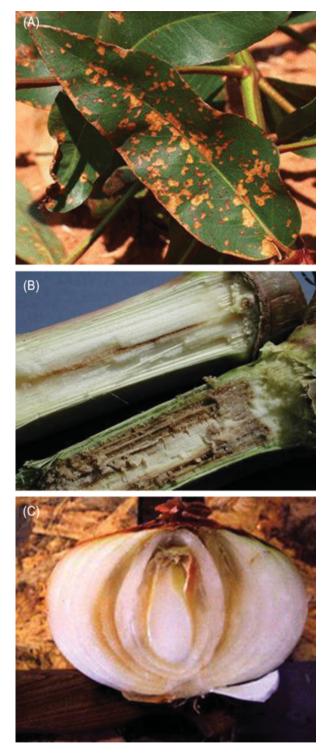


Fig. 2 Disease symptoms caused by *Pantoea ananatis*. (A) Bacterial blight of *Eucalyptus*. (B) Brown stalk rot of maize (photograph courtesy of Dr Teresa Goszczynska, Agricultural Research Council-Plant Protection Research Institute, Pretoria, South Africa). (C) Centre rot of onion (photograph courtesy of Professor Ron Gitaitis, University of Georgia, Tifton, GA, USA).

Of particular interest is that symptoms caused by *P. ananatis* on the same host may differ from country to country. On maize in Brazil, the symptoms described are necrotic or white leaf spots and streaks (Bomfeti *et al.*, 2008; Paccola-Meirelles *et al.*, 2001), whereas, in South Africa, infection results in stalk rot (Goszczynska *et al.*, 2007). Similarly, on rice in Japan and elsewhere in the world, the pathogen infects the developing seed causing palea browning, whereas, in Australia, the symptom caused by *P. ananatis* is stem necrosis (Cother *et al.*, 2004).

Epidemiology

The epidemiology of plant diseases caused by *P. ananatis* on different hosts is relatively unknown. What has been established is that the pathogen enters its host through flowers (Hasegawa *et al.*, 2003; Serrano, 1928) and/or wounds created by feeding insects (Gitaitis *et al.*, 2003; Watanabe *et al.*, 1996; Wells *et al.*, 2002), mechanical injury (Serrano, 1928) and plant to plant contact during high winds (Azad *et al.*, 2000; Cother *et al.*, 2004). The development of brown hopper burn symptoms on rice was found to be accelerated when *P. ananatis* was present on the leaf surfaces. Gitaitis *et al.* (2003) were also able to show that tobacco thrips vector *P. ananatis* in onion fields. Although *P. ananatis* was isolated from Miridae feeding on infected plant tissue in a field outbreak of blight and die-back of eucalypts in South Africa, its exact role in disease outbreaks is currently unknown (J. Roux and T. A. Coutinho, unpublished results).

Pantoea ananatis is both seed-borne and seed-transmitted in onions (Goszczynska et al., 2006a; Walcott et al., 2002), sudangrass (Azad et al., 2000) and rice (Azegami et al., 1983; Tabei et al., 1988). Together with a number of other bacterial species, this bacterium was also found in buckwheat seed (limura and Hosono, 1996) and in maize kernels (Rijavec et al., 2007). The recent appearances of bacterial blight and die-back on *Eucalyptus* in countries that have purchased seed from South Africa suggest seed transmission (Dr. G. Nakabonge, FABI, University of Pretoria, Pretoria, South Africa, unpublished data). Similarly, outbreaks of centre rot of onions may be a result of the introduction of infested seed into new environments/countries (Gitaitis et al., 2004; Goszczynska et al., 2006a).

Environmental factors influence the severity of the diseases caused by *P. ananatis* on its different hosts. In the case of maize and *Eucalyptus*, high humidity and moderate temperature conditions (between 20 and 25 °C) increase the incidence and severity of the disease (Coutinho *et al.*, 2002; Paccola-Meirelles *et al.*, 2001). In contrast, on sudangrass, infection was found to be worst at a temperature of 32 °C and a high relative humidity (Azad *et al.*, 2000). This is similar to the situation with onions, where *P. ananatis* was found to be active at bulb formation when moisture was high and temperatures ranged from 28 to 35 °C (Schwartz *et al.*, 2003).

Economic importance

As a plant pathogen, P. ananatis can lead to serious economic losses. In the case of sudangrass, blighted foliage can reach 50% or more of the total leaf area (Azad et al., 2000). Losses of up to 100% have been recorded in the case of onions (Gitaitis and Gay, 1997). Severe infections of maize resulted in leaf senescence and a sharp decrease in grain size and weight (Pinto, 1995 cited by Paccola-Meirelles et al., 2001). Where an outbreak of P. ananatis occurs in a Eucalyptus plantation, trees either fail to survive or are multistemmed (Coutinho et al., 2002). What is of particular concern is the fact that this pathogen is able to infect a wide range of Eucalyptus species, hybrids and clones. If left unchecked, the disease on its numerous hosts has the potential to reach epidemic proportions under favourable environmental conditions. In addition to being a primary plant pathogen in the field, P. ananatis is also responsible for post-harvest losses of cantaloupe fruit (Bruton et al., 1991), honeydew melons (Wells et al., 1987) and onions (Gitaitis et al., 2003).

Pathogenesis

Pantoea ananatis has been reported to produce indole-3-acetic acid (Enva et al., 2007; Halda-Alija, 2003; Mano and Morisaki, 2008; Sessitsch et al., 2004), which could play a role in pathogenesis. This plant growth promoter affects plants at very low concentrations and promotes cell wall loosening during cell elongation (Brandl and Lindow, 1998). The major virulence factors of *P. ananatis* are, however, currently unknown. The genome of a virulent strain of *P. ananatis* from eucalypts, as well as the type strain from pineapple, have recently been sequenced using 454 pyrosequencing (De Maayer et al., 2008) and solexa technology, respectively, and detailed information will soon become available. Initial comparative genomics revealed the absence of the Type II, Type III and Type IV secretion systems in *P. ananatis*. These secretion systems are located on pathogenicity islands in a broad range of animal- and other plant-associated bacteria. The genome of *P. ananatis* does, however, contain a cluster of genes with high homology to members of the novel Type VI secretion system. This system plays a role in diseases caused by several human and animal-as well as plant-pathogenic Gram-negative bacteria, but whose function has yet to be elucidated fully (Mattinen et al., 2008; Pukatzki et al., 2006).

Pantoea ananatis produces quorum sensing-related signal molecules. Yoshida *et al.* (2006) revealed that, when this bacterium inhabited wheat heads, it produced at least two *N*-acyl-L-homoserine lactones (AHLs): *N*-hexanoyl-L-homoserine lactone (C6-HSL) and *N*-(3-oxohexanoyl)-L-homoserine lactone (3-oxo-C6-HSL). Pomini *et al.* (2006) reported that *P. ananatis* produced three AHLs with the major substance identified being C6-HSL. Morohoshi *et al.* (2007) identified the LuxRI homologue, EanRI,

and C6-HSL and 3-oxo-C6-HSL in a strain isolated from the Shirakwa river in Japan. They were able to show the involvement of the quorum-sensing system in the regulation of exopolysaccharide biosynthesis, biofilm formation and the infection of onion leaves.

Control

The control of diseases caused by P. ananatis is usually achieved through the deployment of resistant/tolerant cultivars/clones. Blight and die-back of eucalypts in South Africa are currently controlled by the use of resistant clones (T.A. Coutinho, unpublished data). They are selected in a cutting production nursery where the incidence of disease is high. Similarly, in a study by Paccola-Meirelles et al. (2002), it was discovered that, by artificially inoculating maize lines, it was possible to select genotypes resistant to P. ananatis. The control of white spot disease of maize in Brazil has also been achieved by applying the fungicide Mancozeb in the initial phases of disease development (Bomfeti et al., 2007). The use of mulch and irrigation systems has been investigated for the control of centre rot of onion (Gitaitis et al., 2004). Irrigation type had no effect on the incidence and severity of disease. However, the use of straw mulch or bare ground was found to delay symptom development by 7-14 days compared with the use of black plastic. Avoidance and eradication of the initial inoculum are probably the most appropriate management strategies that can be recommended against diseases caused by P. ananatis and most other phytopathogenic bacteria.

CONCLUSIONS

• The conclusive identification of strains of *P. ananatis* has been difficult in the past. However, the use of gene sequences, such as *gyrB* or *rpoB*, now provides plant pathologists with a tool to rapidly and reliably identify this species. In addition, this will allow bacteriologists to clarify the ecological role of this bacterium in the natural environment.

• The association of *P. ananatis* with plants as a pathogen has been known since 1928. In the past 20 years, new reports of *P. ananatis* on previously unreported hosts have highlighted the re-emergence of this bacterium as a potentially economically important plant pathogen.

• Little is known of how *P. ananatis* induces diseases in its hosts. However, now that the genomes of virulent strains from pineapple and eucalypts have been sequenced, information pertaining to pathogenicity and host specificity will become available. This approach will facilitate the development of novel approaches to pathogen control in the future.

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