## **Biological Control of Oomycetous Plant Pathogens: A Review**

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## Abstract

Oomycetes are generally known as water molds, and include diverse plant pathogenic organisms. In this review, we summarized plant diseases mainly caused by oomycetes and highlighted ongoing trends in controlling and managing these pathogens using eco-friendly ways.

Key words: antagonistic microorganisms, biological control, oomycet

#### Introduction

Oomycetes, commonly known as water molds (Winter 1880) are detrimental plant pathogens infecting a wide range of host plants such as native weeds, ornamental plants, and trees (Erwin & Ribeiro 1996, Margulis & Schwartz 2000, van West *et al.* 2003, Sanogo & Ji 2012). The pathogenicity of oomycetes is rendered by their spore production, development of infecting structures, and dispersal of spores (Endo & Colt 1974, Kramer *et al.* 1997). In molecular aspects, effector proteins recognized by signature amino acid motifs RxLR (arginine, any amino acid, leucine, arginine), and dEER (a string of acidic amino acids followed by arginine) are known to facilitate the oomycetes virulence in host plant (Kale & Tyler 2011, Tyler 2011).

Oomycetes are being controlled by numerous approaches which include clean nursery stock, use of resistant varieties, chemical, physical, and systemic fungicides. Biological control agents (BCAs) are also used to suppress oomycetes and their related diseases (Pal & Gardener 2006, Lee *et al.* 2005, Sang & Kim 2012). Aside from these, however, various *Pythium*-and *Phytophthora*-causing diseases exhibited the

resistance to BCAs such as propamocarb, mefenoxam, and metalaxyl, no longer (Titone *et al.* 2009, Moorman & Kim 2004, Parra *et al.* 2001). Therefore, development of more advanced and efficient biological control is of utmost necessity for future success to control oomycetes. This mini review summarized major diseases caused by oomycetous pathogens, efficient BCAs against oomycetous diseases, and their relevant mechanisms.

#### Major diseases caused by oomycetes

The plant pathogenic oomycetes contains many taxa and exhibit remarkably diverse lifestyles ranging from obligate biotroph to necrotroph (Agrios 2011). General life cycle of these pathogens can be exemplified by *Phytophthora capsici* (Fig 1a). Few representative disease symptoms caused by them are shown in (Fig 1b). The diseases caused by major genera such as *Phytophthora, Pythium, Peronospora, Albugo,* and *Aphanomyces* are summarized in (Table 1). Species of *Pythium, Phytophthora, Aphanomyces* and *Rhizoctonia,* etc. are known to cause damping-off disease (Agrios 2011). *Albugo candida* causes white rust on *Erysimum crassicaule* (Mirzaee *et al.* 2009). Soil borne *Phytophthora* and *Pythium* spp. are also widespread and cause major losses on crops of soybeans (Schmitthenner 1985) and avocados (Cohen 1981, Darvas *et al.* 1984). In addition, *Phytophthora* and *Pythium* spp. were responsible for many pre- and post-harvest problems on fruits and vegetables, including brown rot of citrus (Cohen 1981a, b, 1982, Gutter 1983), and black pod of cocoa (McGregor 1983, 1984). Recently, new diseases are emerging caused by these oomycetes; for example, severe rotting and blight of seedlings of soybean (Tomioka *et al.* 2013), root rot disease of legumes (Gaulin *et al.* 2007), etc. New species were also reported in many crops: *Pythium solare* (wilt and death of adult plants of

Phaseolus vulgaris) (de Cock et al. 2008), Pythium myriotylum (root and crown necrosis) (Serrano et al. 2008), Phytophthora bisheria (raspberry, rose, and strawberry diseases) (Abad et al. 2008), Pustula sp. (sunflower white rust) (Rost & Thines 2012), Pythium echinogynum (severe "damping-off pathogen" to tomato and cucumber seedlings) (Balghouthi et al. 2013), etc. Some other oomycetes such as Phytophthora gallica (Jung & Nechwatal 2008), Pythium indigoferae, and Pythium irregulare (Souli et al. 2011) caused diseases in oak and apple trees, respectively.



Fig. 1a. Diagram depicting the life cycle of *Phytopthora capsici*.

Life cycle figure was provided by C.D. Smart, Cornell University, with some modifications.

Oospores; Reproduced by permission, from Gallup, C. A., Sullivan, M. J., and Shew, H. D. 2006. Black shank of tobacco. The Plant Health Instructor. DOI: 10.1094/PHI-I-2006-0717-01.

Photo courtesy Zoospores: Fred Brooks, University of Hawaii at Manoa, Bugwood.org.

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Fig. 1b. Representative photographs of some disease symptoms caused by oomycetes:

a) Symptoms of Phytopthora blight on pepper plant with characteristic wilting due to Phytopthora capsici leonian, b) Downy mildew on lettuce plant by Bremia lactucae, c) White rust on morning glory leaf with heavy sporulation of white rust caused by Albugo ipomoe panduratae, d) White rust on mustard with white rust pustules on the leaf underside due to Albugo candida, e) Downy mildew on soyabean caused by Peronospora manshurica, f) Damping off of tobacco with characteristic large and wet lesions caused by Pythium sp. pringsh., g) Damping off characterized by root rot external symptoms on mature beet, superficial scaring caused by Aphanomyces cochlioides, and h) Damping off of common bean caused by Pythium spp.

Photo courtesy; a, b, c, d; Gerald Holmes, Valent USA Corporation, Bugwood.org.

e; Clemson University, USDA Cooperative Extension Slides Series, Bugwood.org.

f; R. J. Reynolds, Tobacco Company Slide Set, Bugwood.org. g; Oliver T. Neher, The Amalgamated Sugar Company, Bugwood.org.

h; Howard F. Schwartz, Colorado State University, Bugwood.org.

Pathogen	Disease caused	References			
Phytopthora species	Root rot pathogen of soybean	Tyler 2007, Souli et al. 2011, Sang et al. 2013			
	Root and crown necrosis of bean	Abad et al. 2008			
	Damping off disease	Agrios 2011			
	Root rot on ginseng	Sang et al. 2006			
Pythium species	Damping-off, root-rot damping-off pathogen" to tomato and cucumber seedlings	Cohen 1981a, Cohen 1981b, Cohen 1982 Van West <i>et al.</i> 2003, Schmitthenner 1985, Cohen 1981a Balghouthi <i>et al.</i> 2013			
	Root rot disease of legumes	Cohen 1981b, Cohen 1982, Gutter 1983, de Cock <i>et al.</i> 2008, Serrano <i>et al.</i> 2008, Balghouthi <i>et al.</i> 2012			
Peronospora Bremia,		Souli et al. 2011			
Plasmopara	Various downy mildews	Agrios 2011			
Albugo species	White blister	Abbasi & Mohammadi, 2009			
	Sunflower white rust	Rost & Thines, 2012			
Aphanomyces	Damping off disease	Agrios 2011, Gaulin <i>et al</i> . 2007			

Table 1	<b>. Im</b>	portant	t plan	t pat	hogen	ic oomy	cetes	and c	liseases	caused	by	them
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# Biological control of oomycetes and mechanisms involved

Microorganisms from different sources such as rhizosphere and phylloshere can potentially reveal biological control effects against different plant pathogenic oomycetes. In mechanistic basis, these microorganisms control the target pathogens by antibiotic production, root colonization, nutrient competition, induced systemic resistance, plant growth promotion, mutualism, mycoparasitism, and predation. Some of the common bacteria, fungi, and actinomycetes against oomyceteous pathogens were summarized (Table 2). The most effective bacterial isolates were *Pseudomonas*, *Bacillus*, *Lysobacter*, *Enterbacter*, and *Paenibacillus*. Fungi such as *Trichoderma*, endophytic *Fusarium*, and *Ganoderma* spp. also controlled oomycetes. Moreover, about 9% of the total number of isolated bacteria identified as Firmicutes,  $\alpha$ -Proteobacteria,  $\gamma$ -Proteobacteria and Actinomycetes exhibited anti-oomycetic activity (Bibi *et al.* 2012).

Name of BCAs	Target disease(s)/ pathogen (s)	Product name (if available)	Mode of action	Reference
Bacteria				
Bacillus licheniformis	Turfgrass diseases	Ecogurad <sup>TM</sup>		Nelson 2004
Bacillus				
lentimorbusWJ5a17				
mutants	Pythium root rot		Radiation	Lee et al. 2003
	Soybean seed, and root	TM		
Bacillus pumilus	rots (Pythium)	GB34 <sup>1M</sup>		Nelson 2004
<b>.</b>	Various foilar, and root	TT II I TM		
Bacillus subtilis	diseases	Kodiak <sup>1</sup>		Nelson 2004
	Seed, seedlings, and root	Avogreen <sup>TM</sup> ,		N.1. 2004
Burkholderia cepacia	rots (Pythium)	Deny <sup>m</sup>		Nelson 2004
Bacillus				
<i>amyloliquefaciens</i> , and	Pythium			Elazzay <i>et al.</i>
P seuaomonas aeruginosa	apnaniaermatum		Colonization and	2012
	Phytopthore blight of		colonization, and	
Chrysophactarium	Pepper Phytopthora		of HCN with	
waniuansa strain K 19C8	cansici		swarming effect	Kim et al. 2012
wangaense, stant 13900	Pythium seed rot and		swarming erreet	Nelson 1988 van
Enterobacter cloacae.	pre-emergence damping-		Competition for	Diik & Nelson
and Erwinia herbicola	off of cotton		nutrient (fatty acid)	2000
			Volatile organic	
			compound 2,4-di-ter-	
			butylpheonl inhibits	
			mycelial growth,	
			sporulation, zoospore	
Flavobacterium			formation, and	Sang & Kim,
johnsoniae strain GSE09	Phytopthora capsici		colonization	2012
Fluorescent				Mezaache et al.
Pseudomonad spp	Pythium ultimum,			2010
	Damping off,			
	Peronosporomycetes,			
Lysobacter sp strain SB-	Aphanomyces		Colonization, and	
K88	cochlioides		antibiosis	Islam <i>et al</i> . 2005
			Antibiosis;	
			production of	
			secondary metadolite	
Insohactor antibiotions	Phytopthora blight of		4- hudroxyphenylacetic	
HS174	nenner		acid	Ko et al 2009
11.712-1	r-ppor			110 01 01. 2007

Table 2. Biological control agent (BCA) and some commercial microbial inoculants for control of plant disease

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Lysobacter enzymogenes 3.1 T8	Pythium root rot in cucumber, <i>Pythium</i> aphanidermatum		Colonization; root of cucumber plant Production of organic acids such as propionic or lactic	Folman 2003,
Lactic acid bacterial strains	Pythium ultimum		pH, accumulation of H <sub>2</sub> O <sub>2</sub> ; antimicrobial compounds Effective selection procedure,	Lutz et al. 2012
Novosphingobium capsulatum strain YJR107	Phytopthora blight of pepper, <i>Phytopthora capsici</i>		colonization, and inhibition of mycelial growth	Sang <i>et al</i> . 2013
CCR80), and Chryseobacterium indologenes ISE14	Phytopthora blight of pepper, <i>Phytopthora capsici</i>		Antagonism, inhibition of mycelial growth Antagonism;	Sang <i>et al.</i> 2008
Pseudomonas fluorescens EBL 20-PF	<i>Pythium</i> <i>aphanidermatum</i> Pythium damping off, and Aphanomyces root		inhibition of mycelial growth; Induce systemic resistance	Muthukumar <i>et al.</i> 2011
Pseudomonas cepacia	rot of peas		Seed colonization 2,4- diacetylphloroglucino	Parke <i>et al.</i> 1991
Pseudomonas fluorescens F113	Pythium ultimum damping-off of sugar beet		l; a natural phenol renders antiphtyopathogenic action Antibiosis,	Dunne et al. 1998
Pseudomonas flourescens CV69 and V11	Cucumber root, and crown rot by <i>Phytopthora dreschsleri</i> <i>Pythium ultimum</i> or <i>Pythium</i>		siderophores, and indole-3-acetic acid (IAA)	Maleki <i>et al.</i> 2010, Maleki <i>et</i> <i>al.</i> 2011
Pseudomonas marginalis Pseudomonas	aphanidermatum	AtEze <sup>TM</sup> ,	growth	Gravel <i>et al</i> . 2005
chlororaphis Pseudomonas aeruginosa	Pythium damping off of	Cedomon	Antibiosis due to siderophere mediated	Buysens <i>et al.</i>
strain 7NSK2 Pseudomonasv sp	tomato Rhizoctoina, and Pythium root of wheat		Antagonism,	1996, Williams & Asher1996
Paenibacillus polymyxa GBR-462	Phytopthora capsici Phytopthora		muotion of mycelium growth, and zoospore formation	Kim et al. 2009
Paenibacillus polymyxa	palmivora, and Pythium aphanidermatum		Biofilm formation, and niche exclusion Production of	Timmusk <i>et al.</i> 2009
Paenibacillus lentimorbus WJ15	Phytopthora capsici,and Pythium ultimum		antifungal metabolites	Lee et al. 2008

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Paenibacillus sp	Damping off (Pythium) Pythium ultimum, and		Indirect; Inducing plant systemic resistance by plant growth promotion	Li <i>et al</i> . 2007
Rhizobacteria	Aphanomyces cochlioides on sugar-beet seedlings		Colonization, and antifungal metabolites	Mavrodi <i>et al.</i> 2012
Rhizobium	seedings			2012
leguminosarum Jordan	Pythium damping-off of			
bv. Viceae Serratia plymuthica	pea, and sugar beet		Colonization Inhibition of mycelium growth, and zoospore	Bardin <i>et al</i> . 2004
A21–4,	Phytopthora capsici		formation	Shen <i>et al</i> . 2002 Dunne <i>et al</i> 1997, Dunne
Stenotrophomonas maltophilia W81	<i>Pythiumultimum</i> damping -off of sugar beet		Production of extracellular protease	<i>et al.</i> 1998
<b>Fungi</b> Chaetomium cupreum/C. globosum	Disease caused by Phytophthora	Ketomium (R) $^{TM}$		Nelson 2004
Chaetomium globosum Clonostachys rosea	Phytopthora infestans Pythium tracheiphilum Pythium ultimum, Phytophthora		PGP Antioomycete	Ramarethinam <i>et</i> <i>al.</i> 2008 Moller <i>et al.</i> 2003
Fusarium oxysporum EF119	infestans, and Phytophthora capsici		Bikaverin, and fusaric acid Direct; antibiosis and mycoparasitism	Kim <i>et al</i> . 2007, Son <i>et al</i> . 2007
<i>Fusarium oxysporum</i> Strain Fo47	Pythium ultimum	Soil	indirect; inducing systemic resistance	Benhamou <i>et al.</i> 2002
Glicocladium virens	Seed seedlings, and root rots ( <i>Pythium</i> )	Guard <sup>TM</sup> , Gliomix <sup>TM</sup>	Inhibition of	Nelson 2004
Ganoderma appalantum	Sclerospora graminicola		sporangia, zoospore release, and zoospore motility, <i>Phoma</i> impairs <i>P.</i> <i>parasitica</i> mycelium	Sudisha & Shetty 2009
Phoma nov.sp	Phytopthora parasiticia		growth, and prevents <i>P. parasitica</i> infection of the leaf Induction of defense-	French patent application (FR 1051767)
Trichoderma viride	Pythium		related enzymes, and	Muthukumar et
(TVA) Trichoderma	aphanidermatum Pythium damping-off		phenolic compound	<i>al.</i> 2011 Yang <i>et al.</i> 2004
Actinomycetes	r yunun damping-on			1 ang et ut. 2004
A stin on lan soo amn an dat	Dudium		Antibiosis; cell wall	El Tombily et al
us. Micromonospora	r yınıum aphanidermatum		and inducing	2010. El-Tarabily
<i>chalcea</i> , and	damping off disease of		systemic resistance in	<i>et al.</i> 2009, El-
Streptomyces spiralis	cucumber		cucumber plant	Tarabily 2006
Actinoplanes spp	Phytophthora megasperma f. sp. Glycinea Wilte good and meta-ta-ta-	Muoostar TM	Antagonism	Filonow & Lockwood 1985
sirepiomyces	withs, seed, and root rols	wrycostop		11015011 2004

Mechanisms of anti-oomycetic activity are mainly due to colonization, antibiotic production, hyphal lysis, sporangium abortion, oospore parasitism and siderophore production (Buysens et al. 1996, Broadbent et al. 1971, Drapeau et al. 1973, Honor & Tsao 1973, Broadbent & Baker 1974, Wynn & Epton 1979). Colonization of bacteria (e.g. Enterobacter cloacae) resulted in a competitive exclusion of nutrients from Pythium, Phytopthora capsici, and Phytophthora cactorum (Nelson 1988, Sang et al. 2007, Sang et al. 2006). Various antibiotics and lytic enzymes produced by microorganisms revealed antagonism against oomycetes (Dunne et al. 1997, Lee et al. 2003, Lee et al. 2008, Timmusk et al. 2009, Muthukumar et al. 2011). Recently, Streptomyces producing chitinase, β-1, 3-glucanase, lipase and protease showed direct lysis of Phytophthora capsici hyphae (Nguyen et al. 2012). In addition, Pseudomonas ûuorescens and Serratia plymuthi showed the antagonisms to Pythium aphanidermatum and Phytophthora capsici (Muthukumar et al. 2011, Shen et al. 2002). The compounds originated from Streptomyces koyangensis and Ganoderma appalantum restricted the growth of oomycetes (Lee et al. 2005, Sudisha & Shetty 2009).

On the other hand, various BCAs were suggested to control oomycetes by modulating the induced systemic resistance (ISR) of host plants either directly or through volatile organic compounds produced by them (Benhamou *et al.* 2002, Sang & Kim 2012). Most of the BCAs reported are able to suppress more than one pathogen; however, some of them were pathogen specific and even some were host-specific showing selective influence of BCAs (Maurhofer *et al.* 1994, Van Dijk & Nelson 2000, Mavrodi *et al.* 2012, Sang *et al.* 2013). Combination treatment of bacteria-fungi or bacteria-bacteria are also effective to control oomycetes (Dunne *et al.* 1998, Muthukumar *et al.* 2011).

#### **Future perspectives**

Development of anti-oomycetic BCAs is very important and utmost necessity for managing oomycetic diseases as it is considered as an alternative or a supplemental way of reducing the use of chemicals in agriculture (De weger *et al.* 1995, Gerhardson 2002). More researches should be carried out to elucidate the mechanism involved in the microorganism-pathogen interaction and to identify the novel efficient BCAs in future to establish sustainable BCAs against oomycetous diseases. Finally, we can conclude that different biological control approaches summarized in this review can shed light on future directions in developing and choosing different biological control agents against oomycetes.

## Acknowledgements

This work was carried out with the support of Cooperative Research Program for Agriculture Science & Technology Development (PJ009411) RDA, Korea, the National Research Foundation of Korea (NRF) grant funded by the Korean government (MEST) (No2011-0020202), and the research funds of Chonbuk National University.

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