

**Investigating the potential use of unsaturated fatty acids as antifungal crop protective agents**

by  
Azadeh Yasari

A thesis submitted in partial fulfillment of the requirements for the degree of

Master of Science  
in  
Plant Science

Department of Agricultural, Food and Nutritional Science  
University of Alberta

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## ABSTRACT

Pathogenic fungi cause significant yield losses and quality reductions to major crops including wheat, canola, and barley. Toxic metabolites produced by some phytopathogenic fungi also pose significant risks to animal and human health. Extensive application of synthetic fungicides is not a sustainable solution since it poses risks to human, animal and environmental health. Unsaturated fatty acids may provide such an alternative because of their possible direct antifungal activity against phytopathogens as well as through the stimulation of plant defense pathways. The present study assessed the *in vitro* and *in vivo* efficacy of two hydroxy fatty acids; coriolic acid and ricinoleic acid, against the phytopathogens *Fusarium graminearum*, *Pyrenophora tritici-repentis*, *Pyrenophora teres* f. *teres*, *Sclerotinia sclerotiorum*, *Leptosphaeria maculans*, and *Aspergillus niger*. Antifungal activity was evaluated using the broth micro-dilution method to determine the minimum inhibitory concentration (MIC). Results indicated that both coriolic acid and ricinoleic acid possessed the strongest inhibitory activity against *L. maculans* with MIC of 0.73 g/L and 0.83 g/L, respectively, followed by *A. niger* with MIC of 0.78 g/L for coriolic and MIC of 0.88 g/L for ricinoleic acid. A weaker inhibitory activity of coriolic acid and ricinoleic acid was observed in terms of other fungal pathogens with MICs which varied greatly between species. An antifungal effect was observed for coriolic acid *in vivo* against pathogenic fungi of wheat and barley. This effect was not correlated to the *in vitro* activity because ricinoleic acid with equivalent *in vitro* antifungal activity showed no protective effect *in vivo*. Moreover, neither coriolic acid nor ricinoleic acid controlled fungal pathogens of canola. In conclusion, coriolic acid inhibits some phytopathogens *in vivo* and may have the potential to be an effective crop protection agent.

## **DEDICATION**

*This work is dedicated to my parents, Mr. Mohammadreza Yasari and Mrs. Farah Rasoulpour Hedayati for their love, sacrifice, and tolerance. It is also dedicated to my beloved husband, Mr. Aidin Foroutan Naddafi, for his support in every single moment of my life.*

## ACKNOWLEDGEMENTS

I would like to express my sincere gratitude to my supervisors Dr. Nat Kav and Dr. Michael Gänzle for giving me the opportunity to work with them, and for their great encouragement support, and guidance throughout my MSc program. I would also like to thank Dr. Stephen Ernest Strelkov for being my supervisory committee member and his great support and valuable suggestions throughout my research. Also, many thanks to Dr. Habibur Rahman for accepting to be my examiner and providing me his constructive comments.

My sincerest thanks to Nuanyi Liang for producing the coriolic acid and providing me with that as well as all her extended help in research and emotional support. I would also like to appreciate Dr. Jie Feng, Ms. Homa Askarian, Dr. Alireza Akhavan, Mr. Berriso Kebede, Dr. Reem Aboukhaddour and Mr. Klaus Strenzke for providing me with the fungal isolates, crops seeds and all their help and advices during my research work. Many thanks to Drs. Urmila Basu, Tiesen Cao, and Victor Manolii as well as my AFNS friends Ms. Homa Askarian, Dr. Swati Megha, Dr. Enid Perez Lara for their advices, constant encouragement and emotional support. I would also to extend my gratitude to Ms. Kelley Dunfield, Ms. Jody Forslund, Ms. Nikki-Karyssa Scott, Mr. Mehdi Farid and Ms. Robin Miles for their help during different steps of my research program and graduate study. I would also like to thank to all the past and current members of the university of Alberta Plant Pathology Lab and Food Microbiology Lab for their assistance, consideration, and encouragement to make a friendly atmosphere in the Labs.

I am also thankful to the Western Grains Research Foundation, Alberta Barley commission, Alberta Canola Producers Commission, Department of Agricultural, Food and Nutritional Science (AFNS), Faculty of Agriculture, Life and Environmental Sciences (ALES) for financial support and providing me with valuable scholarships and awards. The support of Alberta

Innovates - Technology Futures (in providing the ascospores of *Sclerotinia* fungus) is also acknowledged. Finally, I would like to express my gratitude to my beloved parents (Mohammadreza and Farah), and my dear husband (Aidin) for their unconditional love, sacrifice, and moral support throughout my study.

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## LIST OF ABBREVIATIONS

AOS	Allene oxide synthase
BLAST	Basic Local Alignment Search Tool
DAB	3,3'-Diaminobenzidine
FHB	Fusarium head blight
H <sub>2</sub> O <sub>2</sub>	Hydrogen peroxide
HSTs	Host-selective toxins
HUFAs	Hydroxy unsaturated fatty acids
ICS1	Isochorismate synthase 1
JA	Jasmonic acid
LPCB	Lacto Phenol Cotton Blue
MBSU	Molecular Biology Service Unit
MIC	Minimum Inhibitory Concentration
NBT	Nitroblue Tetrazolium
NFNB	Net form of net blotch
O <sub>2</sub> –	Superoxide
PAD4	Protein arginine deiminase 4
PCR	Polymerase chain reaction
PDA	Potato dextrose agar
PDB	Potato Dextrose Broth
PR	Pathogenesis-related
PR1	Pathogenesis-related protein 1
RCBD	Randomized complete block design

ROS	Reactive Oxygen Species
RT	Room temperature
SA	Salicylic acid
SDW	Sterile distilled water
SEM	Standard error of the mean
SFNB	Spot form of net blotch
TMV	Tobacco mosaic virus
UFAs	Unsaturated fatty acids
VSP2	Vegetative storage protein 2
WRKY70	WRKY transcription factor 70

# 1. GENERAL INTRODUCTION

Environmental factors (i.e. abiotic stresses) and challenges from disease causing organisms (i.e. biotic stresses) pose significant challenges to the growth and development of plants. Biotic stresses include diseases to plants caused by living organisms such as bacteria, fungi, viruses, nematodes, weeds and herbivorous insects (Fujita et al. 2006; Atkinson and Urwin, 2012) which negatively impact survival, crop productivity and food production worldwide (Agarwal et al. 2006). Amongst biotic stresses, pathogenic fungi pose some of the more serious threats for plants to cope with and are of major economic concerns to the agriculture industry (Montesinos et al. 2002; Strange and Scott 2005). For instance, pathogenic fungi cause severe yield losses and quality reductions to major crops including wheat (*Triticum aestivum* L.) (Shabeer and Bockus 1988; Savary et al. 2012), canola (*Brassica napus* L.) (Pageau et al. 2006; del Río et al. 2007; Hwang et al. 2016), and barley (*Hordeum vulgare* L.) (Jayasena et al. 2007; Murray and Brennan 2010). Therefore, continued research into the development of effective disease control strategies to combat this threat posed by fungal pathogens is of utmost relevance.

The importance of the aforementioned crops for the Canadian economy cannot be understated (Statistics Canada 2017a,b) as they collectively contribute around \$38.7 billion towards the economy per annum (Canola Council of Canada 2017a; The Canadian Wheat Alliance 2017; Barley Council of Canada 2017). In addition, their nutritional value for human health and wellness including the notable role of canola oil ingredients e.g.  $\alpha$ -linolenic acid (11% of oil content by weight) and linoleic acid (21% of the oil content by weight) (Przybylski et al. 2005; Gunstone 2011, Canola Council of Canada 2017b) in the prevention of cardiovascular

diseases further increase the value of some of the major Canadian crops (Ascherio et al. 1996; Hu et al. 1999; Djousse et al. 2001). Similarly, various health benefits of arabinoxylan oligosaccharides, which are the main component of dietary fiber in cereal grains such as wheat and barley, have also been demonstrated (Lu et al. 2000; Aune et al. 2011; Jonnalagadda et al. 2011; Broekaert et al. 2011; François et al. 2012). Therefore, the major Canadian crops have significant value from both economic and nutritional / health standpoints.

As alluded to earlier, severe diseases caused by pathogenic fungi have led to extensive yield losses in the production of wheat, canola and barley resulting in negative effects on the economy. In addition to crop and associated economic losses, toxic metabolites produced by some phytopathogenic fungi pose significant risks to animal and human health (Fajardo et al. 1995; Bottalico and Perrone 2002; Yoshida et al. 2008). Excessive application of synthetic fungicides also is not an ideal solution since it leads to insensitivity in pathogen populations (Brown et al. 2004; Gossen et al. 2014) and poses significant risks to human, animal, and environmental health (Alavanja et al. 2014). Therefore, the development of alternate crop protection strategies to mitigate the negative effects of fungal plant pathogens is of utmost necessity.

Studies have shown that unsaturated fatty acids (UFAs) and their derivatives (methylated or hydroxy) could have potential as environmentally-friendly fungicides, due to their direct antifungal activity against phytopathogens as well as through the stimulation of defense pathways in different plant species (Graner et al. 2003; Prost et al. 2005; Pohl et al. 2011). Numerous studies have been conducted on the application of UFAs and their derivatives against phytopathogens demonstrating their antifungal activity and potential as environmentally-friendly fungicides because of their molecular configuration and structure (Sjögren et al. 2003; Pohl et al.



2011; Black et al. 2013). However, the precise mechanism behind this antimicrobial activity and compound-target specificity is still unknown. Moreover, their potential as effective antifungal agents in the protection of important crops for the Canadian economy has not been investigated.

## 2. LITERATURE REVIEW

### 2.1. Wheat

Wheat is an important monocotyledonous crop belonging to the family Poaceae (Gramineae) and genus *Triticum* (NCBI 2017a). History showed that wheat was the first domesticated crop (Gill et al. 2004; Li et al. 2007); however, Canada has a relatively short history in terms of wheat cultivation (Agriculture and Agri-food Canada 2017). *T. aestivum* L. (bread or common wheat) and *T. turgidum* L. (durum or pasta wheat) are two important species in the *Triticum* genus, which are considered to be important food staples for human and animals (Paux et al. 2008; Shewry 2009; Zohary and Hopf 2012; Cooper 2015). Wheat grain is primarily composed of macronutrients including carbohydrates, proteins, lipids, fibre, as well as micronutrients (minerals and vitamins) (Šramková et al. 2009), supplying more than 20% of the world total dietary calories and proteins alone and making it one of the most important crops for global food security (Hawkesford et al. 2013; Shiferaw 2013). Industrial usage of wheat includes application of its fiber/gluten/starch in food additives, aquaculture feeding, adhesives and making plastics, and even production of ethanol (Day et al. 2006; FAOSTAT 2017).

Wheat is the second largest grain crop after corn, produced worldwide about 755 million tons in 2017 (USDA 2017a). Among all the wheat species, common wheat is the most widely grown wheat globally. The European Union, China, India, Russia, and the United States are the top five wheat producing countries around the world (USDA 2017b). Canada is the sixth largest producer of wheat with the seeded area of 9.12 million hectares and total production of 29.9 million tons in 2017 (Statistics 2017c). More than 90% of Canadian wheat crops are grown in the Prairie provinces Alberta, Saskatchewan and Manitoba (Aboukhaddour et al. 2013) with a

production of 9.98, 12.9, and 4.36 million tons in 2017, respectively (Statistics 2017c). Currently, Canada is the fourth largest exporter of wheat (21 million tons in 2017) in the world (USDA 2017b).

### **2.1.1. Important fungal diseases of wheat**

Fungal pathogens impact the development and production of the wheat crop. *Fusarium graminearum* Schwabe is the primary fungal agent of the devastating disease called fusarium head blight (FHB) or scab infecting wheat and other small grains worldwide (Parry et al. 1995). In 1982, scab reduced wheat production in the United States by about 4% in total, representing more than 2.72 million metric tons (Boosalis et al. 1982). Severe outbreaks of FHB also have been reported in Canada (Martin and Johnston 1982; Seaman 1982; Tekauz et al. 1986). However, other *Fusarium* species including *F. avenaceum*, *F. culmorum*, and *F. poae* can cause FHB as well as seedling blight and foot rot diseases in wheat and other cereals (Wong et al. 1992; Parry et al. 1995; McMullen et al. 1997; Schmale III and Bergstrom 2003).

*Puccinia graminis* f. sp. *tritici* Eriks & E. Henn. (causal agent of stem rust), *Puccinia triticina* Eriks. (= *Puccinia recondita* Rob. ex Desmaz. f. sp. *tritici*) (causal agent of leaf rust), and *Puccinia striiformis* Westend. f. sp. *tritici* (Pst) (causal agent of stripe or yellow rust) are also three obligate fungal parasites causing major losses (up to 100%) on susceptible cultivars of wheat around the world (Leonard and Szabo 2005; Chen 2005; Wegulo 2012a; Sharma et al. 2016). Powdery mildew, caused by biotrophic fungi *Blumeria graminis* f. sp. *tritici* Em., (Mwale et al. 2014) and Septoria tritici blotch caused by *Mycosphaerella graminicola* (anamorph: *Septoria tritici*) (Ponomarenko et al. 2011) are other important foliar pathogen of wheat worldwide. Tan spot caused by the fungus *Pyrenophora tritici-repentis* is another foliar disease which has the potential to reduce yield up to 50 % in all major wheat growing regions around the

world (Hosford 1982; Shabeer and Bockus 1988; Lamari and Strelkov 2010). In the subsequent section, the causal agent of tan spot disease in wheat, which was used for investigations in the research described in this thesis, is outlined.

## **2.1.2. *Pyrenophora tritici-repentis***

### **2.1.2.1. Taxonomy, epidemiology, and symptoms**

*Pyrenophora tritici-repentis* (Died.) Drechs. (Anamorph: *Drechslera tritici-repentis* (Died.) Shoem.) is an ascomycete fungus belonging to Pleosporaceae family. The pathogen can infect all classes of cereals such as wheat, rye, wild barley, oat as well as other grasses (Hosford 1971; Krupinsky 1992; Strelkov and Lamari et al. 2003; Wegulo et al. 2012b).

The disease caused by this fungus is recognized by two distinct symptoms on the leaves of susceptible wheat cultivars; this includes necrosis (tan color) and extensive chlorosis (yellow color). Initially, the lesions on leaves appear as tan/brown flecks, then they develop into oval- or diamond-shaped lesions with dark brown center and/or yellow circular border. In severe infection, the lesions may coalesce and cover most or the entire of the leaf surface resulting in their death (Kader 2010). These symptoms develop specifically and result from an interaction between the pathogen secreted host-selective toxins (HSTs) and the target receptors of a toxin-sensitive host plant (Strelkov and Lamari 2003; Singh et al. 2010; Aboukhaddour et al. 2011). HSTs lead to toxicity and disease induction in certain susceptible hosts (Scheffer and Briggs 1981). These compounds have no or little effect on resistant host genotypes (Scheffer and Livingston, 1984).

### **2.1.2.2. Importance of tan spot on wheat**

Tan spot is one of the major foliar diseases caused by *P. tritici-repentis* affecting wheat crops worldwide (Hosford 1982). The first severe outbreak of tan spot dates to 1974 in Canada

(Tekauz 1976). Severe disease epidemics have been also reported in South America, USA, and Australia (Kohli et al. 1992; Murray and Brown 1987; Schilder and Bergstrom 1995) as well as in some of the European countries (Cook and Yarham 1989; Leisova et al. 2008). Yield reductions due to tan spot may vary depending on the host growth stage at the time of infection. Yield loss of 13% during the seedling stage of infection, 35% during late growth stage and 48% from the presence of disease throughout the season were reported (Rees and Platz 1983). Not only *P. tritici-repentis* decrease wheat productivity, but it also affects the seed quality by causing red smudge (reddish discoloration of grain) and/or black smudge (blackening of the germ end) (Francl and Jordahl 1992; Fernandez et al. 2001). Such a reduction in grain yield associated with tan spot was reported in Germany ranging from 10-36% (Wolf and Hoffmann 1993). In Canada, the susceptibility of all registered durum wheat cultivars to red smudge has also been reported (Fernandez et al. 1997).

#### **2.1.2.3. Management of tan spot in wheat**

Genetically-resistant wheat cultivars are often considered as sustainable, most effective and most economical management methods for the control of tan spot (De Wolf et al. 1998; Gamba et al. 1998; Reide et al. 2003). However, in Canada, only a limited number of wheat cultivars have resistance to tan spot (Lamari et al. 2005). Tan spot disease severity can be reduced by tillage of the soil (Schuh 1990; Bockus and Claassen 1992; Stover et al. 1996). Crop rotation is also another effective cultural method for reduction of tan spot involving the cultivation of susceptible wheat crops with annual breaks in the same field (De Wolf et. 1998). A 3-year crop rotation between wheat crops can also decrease tan spot incidence (Rees and Platz 1979). Crops such as soybean, alfalfa, peas, and flax (Bailey et al. 1992; Hosford 1971), being non-hosts, are generally good choices for rotation with wheat crops.

Several fungi and bacteria have been regarded as biocontrol agents of *P. tritici-repentis* which ultimately result in yield growth (Gough and Ghazanfani 1982; Pfender et al. 1991). The antifungal metabolites produced by *Pseudomonas fluorescens* strain Pf-5, have shown the inhibitory function against both mycelial growth of *P. tritici-repentis in vitro* and ascocarp formation on wheat straw (Pfender et al. 1993). Six isolates of *Trichoderma harzianum* could reduce tan spot severity under field conditions either as seed treatments or foliar sprays at different growth stages (Perello et al. 2006).

Foliar application of fungicides is another effective approach for the management of wheat tan spot. So far, several fungicides from different classes have been applied to control tan spot (Colson et al. 2003; Wegulo et al. 2011; Patel et al. 2012; Harvey et al. 2015). Application of propiconazole found to decrease tan spot infection in wheat and increase grain yield (Entz et al. 1990). In Denmark, pyraclostrobin, picoxystrobin, propiconazole and prothioconazole were the most effective fungicides for tan spot controlling with efficacies 55-97% (Jorgensen and Olsen 2007). However, a reduced sensitivity to a range of strobilurin fungicides have been found for field isolates of *P. tritici-repentis* in Germany (Reimann and Deising 2005). In western Canada, numerous foliar fungicides are used to control tan spot disease which are listed in the “Crop Protection” e-book (Alberta Agriculture and Forestry 2017a).

## **2.2. Barley**

Barley (*Hordeum vulgare* L.) is an economically important cereal crop along with wheat which belongs to the family Poaceae (Gramineae) (NCBI 2017b). The first barley cultivar adopted for the Canadian climate with high yield was released in 1889 by the Ontario Agriculture College. During the 1950-1960s, desirable traits of Canadian six-row barley were identified (i.e. high malt extract levels, quick processing, and high levels of starch-degrading

enzymes), resulting in introduction of the cultivar “Harrington” as the first Canadian-bred two-row malting barley in the world market. Later, “AC Metcalfe” and “CDC Copeland” were developed as two-row malt cultivars and distinguished for their quality, leading to Canada’s reputation for premium malt barley (GoBarley 2017). Currently three different classes of barley are cultivated in Canada for general purposes (i.e. cosmetic industry), malting, and food barley (Schouest et al. 2012; Decloedt et al. 2015; Rodbotten et al. 2015; Rosser et al. 2016; Canadian Grain commission 2017a).

Barley is the fourth largest crop among grains behind maize, wheat, and rice; its total production in the world was 141.8 million tons in 2017 (USDA 2017c). The European Union, Russia, Ukraine, Australia, and Canada are, respectively, the top five producers of barley around the world (USDA 2017c). In 2017, Canada had a seeded area of 2.33 million hectares produced 7.9 million tons barley (Statistics 2017c), of which 7.5 million tons was produced in the Prairie provinces Alberta, Saskatchewan and Manitoba (Statistics 2017c). In the Prairie provinces, 60.8% of the total seeded area were allocated to malting barley and 31.8% assigned for barley used for general purposes. A relatively small percentage (1.3%) of seeded area was also devoted to food (Canadian Grain Commission 2017b). In 2017, Canada was ranked as the world’s sixth exporter, with 1.5 million tons, after European union, Australia, Russia, Ukraine, and Argentina (USDA 2017d).

### **2.2.1. Important fungal diseases of barley**

Pathogenic fungi pose a threat to barley crops and reduce their productivity. *Puccinia hordei* (causal agent of leaf rust), *Puccinia graminis* f.sp. *tritici* Eriks. & E. Henn. (causal agent of stem rust), *Puccinia striiformis* Westend. f. sp. *hordei* (causal agent of stripe rust), *Pyrenophora teres* Drechs. (causal agent of net blotch), *Blumeria graminis* (DC.) Golovin: Speer

f. sp. *hordei* Em. (causal agent of powdery mildew), and *Rhynchosporium secalis* (Oudem.) J.J. Davis (causal agent of scald) are the major fungi causing foliar diseases in barley with severe yield losses (Czembor 2001; Sun et al. 2006; Xi et al. 2008; Fetch et al. 2011; Safar Ali Safavi et al. 2012; Akhavan et al. 2017). In 1963, severe losses associated by FHB in barley threatened some of the population with starvation in South Korea (Vestal 1964). Loose smut (caused by *Ustilago nuda* (Jensen) Kellerman & Swingle.), covered smut (caused by *Ustilago hordei* (Pers.) Lagerh.) are other severe fungal diseases of barley affecting its seed, head and root with yield reductions (Menzies et al. 2014; Tekauz 2000). In the subsequent section, the pathosystem of necrotrophic fungi *Pyrenophora teres*, which is used in this research, is described.

## **2.2.2. Pyrenophora teres**

### **2.2.2.1. Taxonomy, epidemiology, and symptoms**

*Pyrenophora teres* Drechs. (Anamorph: *Drechslera teres* (Sacc) Shoem.) is an ascomycete fungus which belongs to the family Pleosporaceae (Liu et al. 2011). *Hordeum vulgare* and *H. vulgare* ssp. *spontaneum* are the primary hosts of this pathogen. However, other wild *Hordeum* species, as well as gramineous species from the genera such as *Bromus*, *Avena*, *Triticum*, and *Elymus* are other hosts of this fungus (Shipton et al. 1973; Liu et al. 2011).

*P. teres* has two forms: *P. teres* f. *teres* (*Ptt*) (causal agent of net form of net blotch, NFNB) and *P. teres* f. *maculata* (*Ptm*), (causal agent of spot form of net blotch, SFNB), on barley (Rau et al. 2007; Liu et al. 2011). These two forms are microscopically identical, morphologically similar but genetically different; however, they can be differentiated based on morphological characteristics and the polymerase chain reaction (PCR) based molecular markers (Leisova et al. 2005; Keiper et al. 2008; Lu et al. 2010; Bogacki et al. 2010).



NFNB and SFNB symptoms appear on all upper parts of the barley including leaves, stems, leaf sheaths, and kernels (Liu et al 2011). The pathogen causing NFNB directly penetrates the leaves and initially results small circular to elliptical pin-point lesions. As the symptoms develop in both horizontal and vertical directions, they cause distinct dark-brown net-like patterns with horizontal and vertical reticulations (Steffenson 1997; Liu et al. 2011). The pathogen causing SFNB results dark-brown and circular to elliptical lesions encircled by a chlorotic zone of varying width, depending on the isolates virulence and host resistance (McLean et al. 2009).

#### **2.2.2.2. Importance of net blotch on barley**

*P. teres* is an economically important foliar disease of barley throughout the world causing a loss of grain yield and the quality (Tekauz 1990; Steffenson 1997; McLean et al. 2009; Liu et al. 2011). NFNB and SFNB can typically reduce the productivity of barley cultivars up to 10-40% and 44%, respectively. However, they have the potential for causing 100% yield loss of susceptible cultivars under conducive environmental conditions for disease development (Mathre 1997; Steffenson et al. 1991; Jayasena et al. 2007; Murray and Brennan 2010). Each form of net blotch has different importance in barley-growing areas worldwide depending on susceptibility of the barley cultivars to local pathotypes of *P. teres*, availability of cultural control measures, and the climatic conditions (Steffenson 1997). In France, SFNB was found to be the predominant form of the disease in many regions (Arabi et al. 1992; while in Norway, both forms are present and no strong evidence of predominance of any of these two could be found (Wonneberger et al. 2017). In western Canada, the more prevalent form of *P. teres* seems to be the *Ptt*, which comprises 82% of the total isolates in the collection coming from the Prairie provinces (Alberta, Saskatchewan and Manitoba) (Tekauz 1990). However, several studies showed an increase in the

incidence of *Ptm* over time (Tekauz 1990; van den Berg and Rossnagel 1991; Liu and Friesen 2010), indicating changes in resistance in barley cultivars to *Ptt* and *Ptm* in this region, and/or even in climatic conditions (Tekauz 1990; Louw et al. 1996).

### **2.2.2.3. Management of net blotch in barley**

Crop rotation is one of the management strategies available for the control of net blotch of barley in western Canada (Tekauz 2003). Monoculture of barley resulted in increased net blotch severity compared with a non-host crop rotation (Krupinsky et al. 2004; Turkington et al. 2005; Turkington et al. 2012). However, the high demand of barley for livestock feed and market factors make crop rotation as short-term strategy for controlling net blotch severity and sustaining barley productivity (Turkington 2005). Burning stubble and conventional tillage are considered as other cultural control methods in UK (Jordan and Allen 1984). However, several studies showed that there is no significant difference in the adoption of tillage system either conventional tillage or conservation tillage, in the control of net blotch severity (Mathre 1997; Bailey et al. 2000; Martin et al 2001; Turkington et al. 2006).

The biocontrol activity of two *Pseudomonas fluorescens* strains (MKB100 and MKB156) was observed in foliar application against net blotch symptom development in detached leaf experiments in glasshouse conditions and small-scale field trials (Khan et al. 2010). *Pseudomonas chlororaphis* MA 342 was also found to be an effective and consistent biocontrol agent against the fungus *P. teres* (Hökeberg 1998; Tombolini et al. 1999).

Deployment of resistant cultivars is the most effective and environmentally-friendly means of disease management which is also cost-efficient for growers by reduction or elimination of the need for fungicide applications and cultural control practices. In Canada, the incidence of SFNB has been reduced by 61% with cultivation of resistant 6-rowed cultivars

relative to the susceptible 2-rowed barley cultivars (Tekauz 1976; Tekauz 1990). In another study performed by Turkington et al. (2006), susceptible barley cultivars showed 5-12 times greater potential of severe infection by NFNB compared to resistant cultivars. Currently, most of the commercial barley cultivars, especially the malting types, are susceptible to net blotch (Alberta Agriculture and Forestry 2017c), and very few NFNB-resistant cultivars are available (Alberta Agriculture and Forestry 2017a).

Foliar treatment with fungicides is another method of control by growers where the ecofriendly methods are inadequate (Mathre 1997; Tekauz 2003; Turkington et al. 2011; Turkington et al. 2015). Several classes of fungicides have been identified as effective chemicals in controlling SFNB and NFNB; this includes pyraclostrobin, propiconazole, azoxystrobin, epoxiconazole, and trifloxystrobin (Jayasena et al. 2002). The effectiveness of these fungicides for disease control is dependent on many factors such as their concentration, mode of action, active ingredient, and timing and numbering of applications (Khan et al. 1989; van den Berg and Rossnagel 1990). A list of foliar fungicides for the control of barley net blotch in western Canada is provided in the “Crop Protection” e-book (Alberta Agriculture and Forestry 2017a). However, the frequent application of fungicides on genetically diverse populations of NFNB and SFNB poses a high risk for resistance development, so finding an economically viable alternative to current fungicides in crop protection is a necessity.

### **2.3. Canola**

Canola (*Brassica napus* L) is an oilseed crop developed from rapeseed and belongs to the Brassicaceae (mustard) family (Casséus 2009). The elimination of undesirable characteristics of rapeseed including high content of erucic acid (C22:1) from seed oil and high level of sulphur compounds glucosinolates from seed meal led to the development of Canada’s new version of

rapeseed, called “Canola”, in the early 1970s (Canola Council of Canada 2017a; Busch et al. 1994). Canola is an abbreviated name from "Canadian Oil Low Acid" referring to the cultivars of rapeseed which produce seed oils with less than 2% erucic acid and seed solid components with less than 30  $\mu\text{mol}$  of glucosinolates per gram in the meal (Casséus 2009; Rempel et al. 2014; Canola Council of Canada 2017a). By the 1980's, the production of rapeseed almost replaced by canola in Canada (Casséus 2009).

*Brassica* species including *B. napus*, *B. rapa*, and *B. juncea* are cultivated for canola grade oil and meal. Canola oil, after extraction and refinement, can be utilized as a cooking oil, deep-frying oil, salad seasoning, and in margarine preparation (Rempel et al. 2014). Canola meal is a nutritive source of vitamins B and E as well as proteins which can be used in aquaculture and livestock feed industries (Rempel et al. 2014). Other usage of canola oil is involved in preparation of many non-edible products, such as biodiesel, cosmetics, detergents, inks, lacquers, lubricants, pharmaceuticals, and plastics (Canola Council of Canada 2017c).

The total worldwide production of brassica oilseeds, as one of the world’s most important crops, was 72.06 million tons in 2017 (USDA 2017e). This crop is widely grown in many countries throughout the world where Canada and China were the two leading countries producing by 19.90 and 13.10 million tons in 2017, respectively (USDA 2017e). Ninety-nine percent of the Canada’s canola is produced in the prairie provinces Saskatchewan, Alberta, and Manitoba while British Columbia, Ontario, Quebec and New Brunswick producing the rest 1% (Statistics Canada 2017c). Canola has been considered as one of the major cash crops in Canada, which contributes more than \$26 billion annually. Canada exports 90% of its production to other countries, such as the United States, Japan, Mexico, and China (Rempel et al. 2014; Canola Council of Canada 2017a).

### **2.3.1. Important diseases of canola**

There are various sources of fungal pathogens of canola which are transferred from the soil matrix to canola seed and residues, as well as from other susceptible host plants (Canola Council of Canada 2017d). Clubroot is an important soil-borne disease of canola, caused by *Plasmodiophora brassicae* Woronin, resulting in yield loss of 30-100% in severely infected canola crops (Strelkov et al. 2007; Hwang et al. 2011). *Rhizoctonia solani* Kühn, *Pythium* spp., and *Fusarium* spp. are the other soil-borne pathogenic fungi of canola, infecting roots and young seedlings and causing seedling blight, damping off, foot rot, and brown girdling root rot diseases. White rust or staghead (*Albugo candida* Pers.), alternaria black spot (*Alternaria brassicae* Berk. and *Alternaria raphani* Groves and Skolko), white leaf spot and gray stem (*Pseudocercospora capsellae* Ellis and Everh.), and blackleg (*Leptosphaeria maculans* (Desm.) Ces. & De Not.) are known as stubble- or residue-borne diseases of canola (Martens et al. 1988; Guo et al. 2005). The two destructive fungal pathogens, which are studied in this thesis research, *L. maculans* and *S. sclerotiorum*, (West et al. 2001; Canola Council of Canada 2017d) are described in the following sections.

### **2.3.2. Leptosphaeria maculans**

#### **2.3.2.1. Taxonomy, epidemiology, and symptoms**

*Leptosphaeria maculans* (Desm.) Ces. & de Not. (anamorph *Phoma lingam* Tode ex Fr.) is an ascomycete fungus which belongs to Leptosphaeriaceae family (Rouxel and Balesdent 2005). Members of this order are mostly saprophytes with an ability to live on dead plant materials, while some can live as parasites on living plants (Kaczmarek and Jedryczka 2011). *L. maculans* affects host species mainly of the genus *Brassica* (Rouxel and Balesdent 2005), including cabbage, cauliflower, broccoli, mustard, rapeseed, and canola.

The fungus gradually grows from lesions of leaves via the veins, colonizes in the petiole, and moves systemically through vascular tissue to the stem base without causing any macroscopic symptoms (West et al. 2001; Hayward et al. 2012). Inside the plant, blockage of the xylem vessels by the fungus restricts water and nutrient transmission which contributes to premature ripening, lodging and ultimately yield reduction (Kaczmarek and Jedryczka 2011). At the end of the growing season, the fungus becomes necrotrophic when it completely expands to the plant's crown and roots (Hayward et al. 2012) and causes crown cankers and stem lesions (West et al. 2001). On stems, lesions are usually found at the base with a distinct dark brown or purple margin (Hammond et al. 1985) and have the potential to girdle and even separate the stem in severe infections (West et al. 2001). Seeds also may become dry and pods fracture easily at harvest time, resulting in seed loss (Davies 1986) and denoting that canola is susceptible to blackleg infection from the seedling to the pod-set stages.

#### **2.3.2.2. Importance of blackleg in canola**

Blackleg is one of the most economically important diseases of canola worldwide which cause extensive yield losses in Europe, Australia, and Canada (Gugel and Petrie 1992; Chen and Fernando 2006). It was not considered as a great economic concern until the 1950s, when severe epidemics occurred in France (1950), Australia (1972), and England (1977), with reported yield losses from 50% to 96% (Gugel and Petrie 1992; Toscano- Underwood et al. 2001). In 1998 and 1999, Australia also suffered from significant losses of \$18.6 and \$ 49.4 million in the oilseed industry as a result of this disease, respectively (Khangura and Barbetti 2001). In western Canada, *L. maculans* was firstly identified on canola stubble in central Saskatchewan in 1975. The disease incidence had increased by ten-fold between 1978 and 1981 (Juska et al. 1997) until it was found in 65% of the Saskatchewan's fields in 1986 (Jespersion 1989); then it dramatically

spread to Ontario (1986), Manitoba (1987) and Alberta (1988) (Gugel and Petrie 1992). In 1997, the Canola Council of Canada estimated the annual loss due to blackleg disease about 50 million CAD annually (Juska et al. 1997).

### **2.3.2.3. Management of blackleg in canola**

Crop rotation and the cropping of blackleg resistant cultivars are of utmost importance in cultural control methods of blackleg disease management (Kharbanda and Tewari 1996; Gout et al. 2006). Short rotations or continuous cropping of canola cultivars prevents the complete decomposition of crop residues and readily breaks down the crop qualitative resistance, leading to development of *L. maculans* inoculum on infected canola stubble and debris (Huang et al. 2009; Marcroft et al. 2012; Kutcher et al. 2013). Breakdown of resistance has been reported in France (Rouxel et al. 2003) and Australia (Li et al, 2003). In addition to crop rotation, weed control is also considered as an effective method for blackleg management since *L. maculans* can survive on volunteer canola and other weed species (Kutcher et al. 2011). Other methods of cultural control of blackleg disease in canola are time adjustment of seeding in the season prior to maturation of pseudothecia and peak periods of ascospore discharge (Aubertot et al. 2004), tilling and burning the crop residue (Kharbanda and Tewari 1996; Guo et al. 2005), as well as flooding canola basal stems infected with *L. maculans* (Peluola et al. 2013). In Canada, all registered cultivars of *B. napus* have moderate to high resistance against blackleg disease (with unclear resistance type or specific resistance genes) (Kutcher et al. 2011). The heterothallic nature of *L. maculans* may increase genetic diversity, result in more virulent genotypes of the fungus population, and ultimately defeat the resistance (Aubertot et al. 2006).

Biological control procedures have been found to be useful for the management of blackleg in canola. For example, the Bird's nest fungi, *Cyathus striatus* and *Cyathus olla*, reduce

the stubble food for *L. maculans* survival and available inoculum for further spread (Shinners and Tewari, 1997). Two bacterial strains, *Bacillus endophyticus* and *Bacillus amyloliquefaciens* have been recognized as potential biocontrol agents against *L. maculans* (Danielsson et al. 2007). The bacterial isolate of *Serratia plymuthica* HRO-C48 and fungal isolate of *Gliocladium catenulatum* J1446 found to decrease the disease severity of *L. maculans* on infected cotyledons of *B. napus* by 44% and 52%, respectively (Hammoudi et al. 2012). The antifungal activity of peptides produced by the bacterium *Paenibacillus polymyxa* Prazmowski against *L. maculans* was also reported (Kharbanda et al. 2003).

In western Canada, synthetic fungicides are generally applied on canola both as seed and foliar treatments for the reduction of the incidence of blackleg disease. Seed treatment destroys seed borne inoculum, prevents the spread of blackleg into un-infected areas, and protects the young seedling from airborne ascospores with their systemic activity (Gugel and Petrie 1992). Foliar treatment reduces blackleg disease symptoms on leaves and results in increased yield (Kutcher et al. 2011); however, mixed successes have been reported regarding the efficacy of foliar fungicides in blackleg control (Gugel and Petrie 1992; Khangura and Barbetti 2004). In Alberta, there are currently six registered products for seed treatments and eight registered foliar fungicides for management of blackleg in canola (Alberta Agriculture and Forestry 2017a).

### **2.3.3. Sclerotinia sclerotiorum**

#### **2.3.3.1. Taxonomy, epidemiology, and symptoms**

*Sclerotinia sclerotiorum* (Lib.) de Bary is an ascomycete fungus which belongs to the Sclerotiniaceae family (Willetts and Wong 1980; Bolton et al. 2006). *S. sclerotiorum* is a non-specific and necrotrophic pathogen which is known as the causal agent of disease in over 400 plant species from 75 different families (Boland and Hall 1994; Bolton et al. 2006; Attanayake et



al. 2013). The initial symptoms of infected tissues usually show as water-soaked lesions then they rapidly enlarge and develop into necrotic tissues with fluffy white mycelium (Boland and Hall 1994; Bolton et al. 2006). Later, the old lesions become bleached, shredded, and shattered when dry (Bolton et al. 2006).

### **2.3.3.2. Importance of Sclerotinia stem rot on canola**

Sclerotinia stem rot is one the most destructive diseases of canola in many countries around the world, with the potential to cause severe yield losses and oil quality reduction (Bardin & Huang 2001; Bolton et al. 2006; Gao et al. 2014). In China, this fungal disease could reduce canola yield ranging from 10% to 80% as well as the oil quality (Wang et al. 2014). In 2008, 36% of the canola crops cultivated in UK was damaged by sclerotinia stem rot leading to economic loss of £20 million (Young and Werner 2012). Annual incidence of *S. sclerotiorum* in the United States averaged around 13.6% (Bradley and Lamey 2005; Lamey 2003), which could contribute to an economic loss of \$94 million between 1991 and 2002 (Lamey 2003). In western Canada, yield losses ranging from 5 to 10% due to Sclerotinia stem rot are typical in canola fields (Fang and Platford 1995; Platford 1996), although they may reach up to 94% in drastically infected fields (McLaren et al. 2005).

### **2.3.3.3. Management of Sclerotinia stem rot in canola**

Tilling the soil is a potential method for management of Sclerotinia stem rot, causing the sclerotia to be buried and preventing their successful germination (Williams and Stelfox 1980). It also may be effective in increased parasitism of the buried sclerotia resulting in reduced viability of the sclerotia (Kurle et al. 2001). Altering the crop canopy phenology through avoiding dense-planting is another cultural control strategy against *S. sclerotiorum* reducing the disease dispersal via plant to plant contact (Tu and Zeng 1997). Crop rotation is focused on decreasing the amount

of inoculum present in the field (Twengström et al. 1998). However, the ineffectiveness of a 3- or 4-year crop rotation away from a susceptible host in reduction of viable sclerotia populations in the field was also reported (Williams and Stelfox 1980; Morrall and Dueck 1982). This ineffectiveness could be attributed to the scattering of *S. sclerotiorum* ascospores by wind (Suzio and Koayashi 1972; Williams and Stelfox 1979), causing the infection of host crops by externally produced inoculum (Morrall and Dueck 1982). Moreover, the vast host range of *S. sclerotiorum* and presence of susceptible weeds and volunteers may lead to inoculum preservation, even in the absence of a canola crop (Boland and Hall 1994).

More than 30 species of fungi and bacteria exist with antagonistic or mycoparasitic activity against *Sclerotinia* spp (Adams and Ayres 1979). Fungi belonging to the genera *Fusarium*, *Gliocladium*, *Hormodendrum*, *Mucor*, *Pencillium*, *Trichoderma*, *Verticillium*, *Conithyrium*, and *Ulocladium* were identified as biocontrol agents of *S. sclerotiorum* (Adams and Ayres 1979; McLaren et al. 1996; Li et al. 2003). A bacterial suspension of *Bacillus subtilis* could strongly inhibit the mycelial growth of *S. sclerotiorum* and sclerotia germination at concentrations of  $10^9$  and  $10^{11}$  CFU/ml. It also significantly reduced the disease severity and incidence in field trials by 50-70% was as effective as the fungicide treatments including tebuconazole and carbendazim used for canola stem rot control (Gao et al. 2014). In Canada, three registered bio-control products are Serenade Max (PCP# 28549, Bayer Crop Science), Serenade CPB (PCP# 30143, Bayer Crop Science), and Contans WG (PCP#29066, Bayer Crop Science) which are used on canola against *Sclerotinia* stem rot (Government of Saskatchewan 2016; Alberta Agriculture and Forestry 2017a).

Deployment of resistant cultivars is another control measure for the management of *Sclerotinia* stem rot. However, there are no commercial cultivars with complete resistance

against *Sclerotinia* stem rot in the Canadian market (Canola Council of Canada 2017e). Some tolerant cultivars such as ‘45S52’ from DuPont Pioneer were introduced to the market which can be cultivated by growers as another control measure, although they still can get damaged when the disease pressure is high (Pratt 2012).

In Canada and United States, application of fungicides at the flowering stage of canola is one of the conventional methods for controlling *Sclerotinia* stem rot, which often must be made prior to symptoms appearance (del Río et al. 2007; Koch et al. 2007; Turkington et al. 2011). In Alberta, several registered fungicides such as Acapela, Lance AG, Priaxor, Proline 480 SC, and Quadris are used for the management of this disease (Alberta Agriculture and Forestry 2017a). The application of fungicides in disease control may lead be successful, but it may not be guaranteed every year since over time the infection level by *S. sclerotiorum* changes. The extensive application of fungicides also causes fungicide insensitivity via increased selection pressure on pathogen populations. Such an example was confirmed in Canada when *S. sclerotiorum* populations developed resistance to the fungicide benomyl (Benlate), formerly used as chemical agent for control of canola and alfalfa diseases caused by *S. sclerotiorum* (Gossen et al. 2001).

#### **2.4. Antimicrobial activity of UFAs against phytopathogens**

Difficulties and challenges associated with pathogenic fungi such as severe yield losses, wide host range, lack of highly resistant cultivars, and increased virulence or resistance toward synthetic fungicides have motivated researchers to develop alternative or complementary fungicides.

UFAs and hydroxy unsaturated fatty acids (HUFAs) as organic compounds provide one such alternative. They include compounds with antifungal activity due to their specific structural

characteristics (Prost et al. 2005; Pohl et al. 2011; Cantrell et al. 2008). Coriolic acid (13-hydroxy-9(Z),11(E)-octadecadienoic acid), which was first extracted from the seed oil of *Coriaria nepalensis* (Tallent et al. 1966), could inhibit the spore germination and germ tube elongation of the rice blast fungus, *Pyricularia oryzae* (Namai et al. 1993). The accumulation of coriolic acid in Sasanishiki rice cultivar after infection with the rice blast fungus as anti-rice blast fungus compounds was also reported (Kato et al. 1993). The inhibitory activity of coriolic acid also was demonstrated for the *in vitro* growth of fungal pathogens such as *Cladosporium herbarum*, *Botrytis cinerea*, *Phytophthora infestans*, and *Phytophthora parasitica* var. *nicotianae* (Prost et al. 2005). Didehydrocoriolic acid has been shown to act as a self-defense compound against the rice blast disease (Ding et al. 2012). Application of coriolic acid by 0.15% in bread making could extend the mold-free shelf life of wheat bread from 2 to more than 6 days (Black et al. 2013). The growth-inhibiting activity of coriolic acid against food spoilage as well as fungal pathogens *Aspergillus niger* and *Penicillium roqueforti* was also observed (Liang et al. 2017).

Ricinoleic acid (12-hydroxy-cis-9-octadecenoic acid), constituting more than 80% of seed oil of castor plant (*Ricinus communis* L.) (Bafor et al. 1991), showed antifungal activity against pathogenic fungi *Aspergillus niger* and *Penicillium roqueforti* *in vitro* (Black et al. 2013; Chen et al. 2016; Liang et al. 2017). The methylated derivatives of ricinoleic acid such as methyl-12-(4-hydroxy-3-methoxyphenyl propenoate) octadec-9-en-1-oate and methyl -12-(4-hydroxy-3-methoxybenzoate) octadec-9-en-1-oate also possessed antifungal activity against fungal pathogens *Candida albicans*, *Candida rugosa*, *Saccharomyces cerevisiae*, *Rhizopus oryzae*, and *Aspergillus niger* *in vitro* experiments via agar well diffusion method. The antifungal activity of coriolic acid and ricinoleic acid accumulated in supernatant of *Pseudomonas aeruginosa* 42A2 against pathogenic fungi *Verticillium dhaliae*, *Macrophomina phaesolina*,

*Arthroderma uncinatum*, *Trycophyton mentagrophytes* and *Penicillium funiculosum* have been also reported (Martin-Arjol et al. 2010).

Other UFAs and their derivatives exert antimicrobial activity against diverse range of species. For example, 13(S)-hydroperoxy-9(Z),11(E),15(Z)-octadecatrienoic acid, 13(S)-hydroxy-9(Z),11(E),15(Z)-octadecatrienoic acid, and 5(Z)-etherolenic acid inhibited the mycelial growth of *Leptosphaeria maculans* and *Alternaria brassicae* at 1 mM concentration but their relative potency of inhibition was different (Graner et al. 2003). However, none of these UFAs were effective in inhibition of mycelial growth of *Sclerotinia sclerotiorum* at the same concentration, indicating species-specific mode of action (Graner et al. 2003). The growth-inhibiting activity of *threo*-12,13-dihydroxy-9(Z)-octadecenoic acid and *cis*-12,13-epoxy-9(Z)-octadecenoic acid was also reported against *Sclerotinia sclerotiorum* (Graner et al. 2003). The antiviral activity of oleic acid against *tobacco mosaic virus* (TMV) was demonstrated *in vivo* (Zhao et al. 2017). Methyl- and ethyl- oleate (3 g/L) showed antifungal activity against the powdery mildew of barley caused by *Blumeria graminis* f. sp. *hordei* (Choi et al. 2010). However, many of these UFAs are not stable chemically and tend to be degraded in the presence of biomolecules (Prost et al. 2011) or to be metabolized by several enzymes (Schaller et al. 2001; Chechetkin et al. 2004), explaining the transient nature of growth inhibitory effect and making intricate the antifungal functionality of the UFAs against the target fungi (Graner et al. 2003).

Apart from the direct antifungal activity of UFAs, they may also act simultaneously as signaling molecules to regulate plant self-defense pathways through their phytohormone mediators such as salicylic acid (SA) and jasmonic acid (JA) (as reviewed by Kunkel and Brooks 2002; and Kachroo and Kachroo 2009). Accumulation of oxylipins such as coriolic acid in rice plants led to enhanced resistance to rice blast fungal pathogen (Yara et al. 2008). Rice plants

treated with 2,6-dichloroisonicotinic acid (a synthetic resistance-inducer) and JA were efficiently protected from infection by the rice blast fungus being related to the induction of a set of pathogenesis-related (*PR*) genes (Schweizer et al. 1997). Increased expression of PR genes resulted in increased resistance to TMV also was illustrated in oleic acid-treated tobacco leaves (Zhao et al. 2017). Exogenous application of arachidonic acid (10  $\mu$ M) on wild-type leaves of *Arabidopsis* demonstrated an increase in basal levels of JA which was related to upregulation of JA biosynthesis pathways gene (Allene oxide synthase, *AOS*) and JA-responsive gene (Vegetative storage protein 2, *VSP2*) in response to arachidonic acid treatment (Savchenko et al. 2010). However, this treatment reduced the SA levels and downregulated the expression of the genes protein arginine deiminase 4 (*PAD4*), isochorismate synthase 1 (*ICS1*), WRKY transcription factor 70 (*WRKY70*), and pathogenesis-related protein 1 (*PR1*), indicating the antagonistic interactions of JA and SA pathways to facilitate the fine-tuning of defense responses to different plant pathogens (Savchenko et al. 2010).

Signaling in plant defense pathways leads to the formation of UFAs, as signaling molecules or antimicrobial agents, which ultimately contributes to pathogen defense (Kachroo and Kachroo 2009). To date, several studies have been conducted regarding the application of UFAs and HUFAs against phytopathogens illustrating their antifungal activity (Graner et al. 2003; Prost et al. 2005; Pohl et al. 2011) and their potential as environmentally-friendly fungicides because of their molecular configuration. However, the exogenous application of UFAs on major Canadian crops as protective agents against pathogenic fungi has not been reported yet. Therefore, in the present research work, the efficacy of two HUFAs, coriolic acid and ricinoleic acid will be assessed.

## 2.5. Research objectives

The first objective of this project was to determine the antifungal activity and efficacy of coriolic acid (13-hydroxy-9(Z),11(E)-octadecadienoic acid) and ricinoleic acid (12-hydroxy-cis-9-octadecenoic acid) against important fungal pathogens of canola (*Leptosphaeria maculans* and *Sclerotinia sclerotiorum*), wheat (*Fusarium graminearum* and *Pyrenophora tritici-repentis*), barley (*Pyrenophora teres* f. *teres*), and fruits (*Aspergillus niger*) *in vitro*. The second objective was to figure out the potential contribution of coriolic acid and ricinoleic acid in crop protection against the selected fungal pathogens of canola, wheat, and barley when they are applied exogenously on plants and on seeds (*in vivo*).

## 2.6. Research Hypotheses

The specific hypotheses of this study are as follows:

- Coriolic acid and ricinoleic acid could have significant inhibitory effects on mycelial growth of pathogenic fungi including *L. maculans*, *S. sclerotiorum*, *F. graminearum*, *P. tritici-repentis*, *P. teres* f. *teres*, and *A. niger* *in vitro*.
- Exogenous application of coriolic acid and ricinoleic acid on plants and/or seeds (*in vivo*) may inhibit the disease development and control plant infection.

### 3. MATERIALS AND METHODS

#### 3.1. Preparation of UFAs

Coriolic acid (13-hydroxy-9(*Z*),11(*E*)-octadecadienoic acid) was generously provided by the Food Microbiology lab (Nuanyi Liang, PhD Candidate) of the Department of Agricultural, Food and Nutritional Science, University of Alberta. It was produced either through enzymatic oxidation of linoleic acid or extracted from seed oil of *Coriaria nepalensis* Wall (XinTai Seed Production and Wholesale Company, Jiangsu, China), then purified by high speed counter current chromatography (Nanda and Yadav 2003; Liang et al. 2017). Ricinoleic acid (12-hydroxy-9-*cis*-octadecenoic) and oleic acid (9-*cis*-octadecenoic acid), each with purity > 99%, were also purchased from Nu-Check Prep, Inc. (Elysian, MN).

#### 3.2. Preparation of fungal isolates

The fungal isolates *Pyrenophora tritici-repentis* AB 7-2 (Aboukhaddour et al. 2013), *Pyrenophora teres* f. *teres* AB 06 and AB 34 (Akhavan et al. 2016; 2017), and *Sclerotinia sclerotiorum* SS-01 (Navabi et al. 2010) were used in this study. *Aspergillus niger* FUA5001 was kindly provided by Dr. M. Gänzle, Department of Agricultural, Food and Nutritional Science, University of Alberta. *Fusarium graminearum* G-1, *Leptosphaeria maculans* RL-60 were generously provided by Dr. S.E. Strelkov, Department of Agricultural, Food and Nutritional Science, University of Alberta. The identities of both *F. graminearum* and *L. maculans* were confirmed via DNA sequence analysis (Demeke et al. 2005; Liu et al. 2006). Briefly, DNA was extracted from actively growing mycelia on PDA plates using Promega DNA isolation kit (Promega Corporation, Madison, WI, USA) and quality and quantity of the isolated DNA were determined using NanoDrop 1000 spectrophotometer (Thermo scientific, Waltham, MA, USA).



Two specific primer pairs were also selected from the literature (Table 3-1) for these two fungi and used for PCR amplification reactions (Demeke et al. 2005; Liu et al. 2006). PCR amplification was performed in a 25 µl of reaction including 2.5 µl of 10X PCR buffer (150 mM Tris-HCl, 500 mM KC), 2 mM Mgcl<sub>2</sub>, 0.2 mM of dNTPs, 1 units of *Taq* DNA Polymerase, 0.2 µM each forward and reverse primer (Table 3-1) and 1 µl of mycelial DNA. The entire reaction was then incubated at 95 °C for 3 min for initial denaturation, followed by 30 cycles of 30 s denaturation at 95 °C, 30 s annealing at 62 °C for *F. graminearum* and 68°C for *L. maculans*, and 1 min extension at 72 °C, plus a final extension of 5 min at 72 °C in GeneAmp PCR System 9700 (Applied Biosystems, USA). PCR products were separated by electrophoresis in 2.0% (wt/vol) agarose gels stained with 0.6 µl of syber safe DNA gel stain (Invitrogen, USA). Then, the amplified products of expected size were isolated from agarose gels using a sterile scalpel blade on UV exposure box. DNA sequences were determined at the Molecular Biology Service Unit (MBSU), Department of Biological Sciences, University of Alberta. DNA sequences were confirmed using Basic Local Alignment Search Tool (BLAST; <https://blast.ncbi.nlm.nih.gov/Blast.cgi>).

Table 3-1. Specific primers used for identification of *Fusarium graminearum* and *Leptosphaeria maculans*

<b>Fungal pathogens</b>	<b>Primer sequences</b>	<b>References</b>
<i>Leptosphaeria maculans</i>	Lm-F: CTTGCCCCACCAATTGGATCCCCTA Lm-R: GCAAATGTGCTGCGCTCCAGG	Demeke et al. 2005
<i>Fusarium graminearum</i>	Fg16-F: CTCCGGATATGTTGCGTCAA Fg16-R: GGTAGGTATCCGACATGGCAA	Liu et al. 2006

### **3.3. Preparation of fungal inoculum**

The required growth conditions for sporulation of all fungal isolates is described in Table 3-2. When fungal isolates sporulated on their specific media, approximately 5 ml of sterile distilled water was added to sporulating cultures and gently scraped using a sterile inoculation loop. Under sterile conditions, the spore suspensions were collected and passed through four layers of cheesecloth to eliminate mycelial cells. Spores were counted using a Neubauer counting chamber (Fein-Optik, Jena, Germany) and spore concentrations were adjusted for subsequent experiments.

Table 3-2. Fungal pathogens, pathogenicity and their specific sporulation conditions

<b>Fungal isolates</b>	<b>Pathogenicity</b>	<b>Culture conditions for sporulation</b>
<i>Pyrenophora tritici-repentis</i> AB 7-2	Tan spot in wheat	The fungus was grown on V8-PDA medium (150 ml V8 juice, 3 g CaCO <sub>3</sub> and 10 g agar, 10 g PDA for 1 L) at room temperature (RT: 21 ± 2 °C) under continuous darkness until colonies reached a diameter up to 5 cm. Mycelia were covered by sterile distilled water (SDW) and gently streaked with the bottom of a sterilized glass tube. Subsequently, plates were incubated for 16-19 h under fluorescent light at RT. Then they were incubated at 14°C for 24 h (Aboukhaddour et al. 2013).
<i>Pyrenophora teres f. teres</i> AB 06 & AB 34	Net boltch of barley (NFNB)	The fungus was grown on V8-agar (100 ml V8 juice, 3 g CaCO <sub>3</sub> and 20 g agar, for 1 L) at RT under continuous darkness until colonies reach the diameter up to 5 cm. Mycelia were covered by sterile distilled water and gently streaked with the bottom of a sterilized glass tube. Subsequently, plates were incubated for 16-19 h under fluorescent light at RT. Then they were incubated at 15 °C for 24 h (Akhavan et al. 2017).
<i>Sclerotinia sclerotiorum</i> SS 01	Sclerotinia stem rot on canola	The fungus was grown on Difco potato dextrose agar (PDA; Dickinson and Company, Sparks, MD, USA) medium at RT for 3 days. Then sub-cultured on fresh PDA media for additional 3 days at RT to use for inoculation of canola seedlings (Joshi et al. 2016).
<i>Leptosphaeria maculans</i> RL-60	Blackleg on canola	The fungus was grown on V8-agar medium (150 ml V8 juice, 1.5 g CaCO <sub>3</sub> and 15 g agar for 1 L) at RT under continuous cool-white fluorescent light and incubated for 14 days (Feng et al. 2014).
<i>Fusarium graminearum</i> G-1	Fusarium head blight on wheat and barley	The fungus was grown on PDA plates sealed with parafilm at 25°C for 12 h light/darkness for 5 days. (Geddes et al. 2008).
<i>Aspergillus niger</i> FUA5001	Mold spoilage	The fungus was grown on PDA plates at 25 °C in darkness for 7 days (Magnusson and Schnurer 2001).

### 3.4. Minimum Inhibitory Concentration (MIC) Assay

Antifungal activity of fatty acids was determined using the broth micro-dilution method as described by Magnusson and Schnurer (2001) with some modifications. Pure fatty acids were dissolved in 100% ethanol at a concentration of 53.33 g/L. The fatty-acid stock solution (100 µl) was mixed with 100 µl of the specific liquid medium used to grow each fungus (described in Table 3-3) in a 96-well microtiter plate to make a series of 2-fold diluted of HUFAs at 20, 10, 5, 2.5, 1.25, 0.625, 0.312, 0.156, 0.078 and 0.039 g/L concentrations. Ethanol was evaporated by placing the microtiter plate without a lid in a laminar flow hood until 100 µl of 50% ethanol placed in an empty well of the same plate was totally evaporated. Subsequently, the wells were inoculated with 33.33 µl of spore suspensions of the aforementioned pathogenic fungi at concentration of  $10^4$  spores/ml. The well consisting of only the medium was considered as the negative control and the well including the medium plus spore suspension served as the positive control. The MIC of samples was defined as the lowest concentration to completely inhibit the visible growth of the fungal strains and determined visually one day after that the growth was visible in the positive-control wells. The MIC values were determined from the average of three independent experiments using replicate preparations of the conidiospores. The entire experiment was also repeated twice.

Table 3-3. Specific liquid media used in MIC for each fungal pathogen

<b>Fungal pathogens</b>	<b>Specific liquid media used for MIC</b>
<i>Pyrenophora tritici-repentis</i>	Potato Dextrose Broth (PDB)
<i>Pyrenophora teres</i> f. <i>teres</i>	PDB
<i>Leptosphaeria maculans</i>	V8-juice broth
<i>Sclerotinia Sclerotiorum</i>	PDB
<i>Aspergillus niger</i>	mMRS broth

### 3.5. Plant materials

*Brassica napus* cv. ‘Westar’ and DH12075 line used as susceptible genotypes for experiments with *L. maculans* and *S. sclerotiorum*, respectively (Sharma et al. 2010; Joshi et al. 2016). Their seeds were sown in plastic inserts (5 cm × 5 cm; one seed per insert) filled with Sunshine potting mix (W.R. Grace and Co., Fogelsville, PA, USA) and grown in controlled growth cabinets under 22 °C day/18 °C night and 16 hours photoperiod for three weeks. Seedlings were watered regularly and were fertilized by Peters ® NPK (20-20-20) solution on 2-week old seedlings at concentration of 200 ppm (Sharma et al. 2010). *Triticum aestivum* cv. ‘Katepwa’ and *Hordeum vulgare* cv. ‘Xena’ were treated as susceptible genotypes for *P. tritici-repentis* and *P. teres* f. *teres* experimentation, respectively. Their seeds were sown in plastic square pots (5 inches; 6 seeds per pot) filled with Sunshine potting mix (W.R. Grace and Co., Fogelsville, PA, USA) and grown in growth cabinets under 20 °C day/18 °C night and 16 hours photoperiod and watered regularly for 10-13 days to reach the 2-3 leaf stage (Aboukhaddour et al 2013; and Akhavan et al. 2016). No fertilizer was used for wheat and barley cultivars.

### 3.6. Foliar treatment and plant inoculation procedures

To evaluate the reaction of canola cv. ‘Westar’ to *L. maculans* after fatty acids treatment, 3-week old seedlings were placed in a misting chamber 24 hour prior to inoculation with a relative humidity of > 95%. Then seedlings were air-dried in greenhouse for about 2 hours and treated manually using a plastic-bottle sprayer with approximately 10 ml of coriolic acid and ricinoleic acid per seedling at concentrations of 0.12, 0.25, 0.5, 1, and 2 g/L. Four hours later, two true leaves of each seedlings were wounded at two sites of the mid-rib using a sterile needle gauge and inoculated with 10 µl of spore suspension with concentration of 10<sup>7</sup> spores/ml. To prevent the falling of inoculum droplets, the seedlings were left for about two hours at the same

place and then returned to the humidity chamber for 48 hours. After transferring plants to the previous growth cabinets, disease severity was measured 14 days post inoculation (Sharma et al. 2010). Briefly, all treated leaves and control leaves were excised from seedlings, gently wiped clean with a wet sponge to remove any debris that might interfere with the analysis, and then scanned using EPSON Perfection V19 Scanner (Epson America Inc, USA). Subsequently, all pictures were analyzed by APS Assess 2.0 software (The American Phytopathological Society 2008) and the mean disease severity was calculated as the percentage of lesion area divided by leaf area for each leaf.

Inoculation of the canola line DH12075 by *S. sclerotiorum* was done based on the methodology described by Joshi et al. (2016) with some modifications. Briefly, 21-day old seedlings were placed in the humidity chamber 24 hours prior to inoculation. The seedlings were air-dried in greenhouse for about 2 hours and then were treated manually using a plastic-bottle sprayer with approximately 10 ml of coriolic acid and ricinoleic acid per each one at five different concentrations including 0.12, 0.25, 0.5, 1, and 2 g/L. Four hours later, two true leaves of each seedlings were wounded at two sites of the mid-rib using a sterile needle gauge and inoculated by 3-day old mycelial plug (5 mm) of actively growing *S. sclerotiorum* excised from PDA plates. Similarly, leaves of negative control plants were inoculated with sterile PDA plugs without fungal cultures. After inoculation, the plants were left in the humidity chamber for 24 h and then transferred to the growth cabinets. Disease severity also was measured 48 hours post inoculation using APS Assess 2.0 software as described for ‘Westar’ cultivar (The American Phytopathological Society 2008).

Seedlings of the wheat cv. ‘Katepwa’ were treated manually using a plastic-bottle sprayer with approximately 8 ml of coriolic acid, ricinoleic acid, and oleic acid at five different

concentrations with two-fold dilutions including 0.12, 0.25, 0.5, 1, and 2 g/L four hours before inoculation. Inoculation with *P. tritici-repentis* was performed as described by Aboukhaddour et al. (2013) with some modifications. Briefly, seedlings in two- to three-leaf stage were sprayed by conidial suspension with concentration of  $4 \times 10^3$  spores/ml using a pressurized sprayer connected to an air-line. Immediately, the inoculated seedlings were covered with moist plastic bags for 24 hours to maintain the relative humidity at almost 100%. Subsequently, the bags were removed, plants were returned to the previous growth conditions, and disease severity was measured six days after inoculation using APS Assess 2.0 software as described for ‘Westar’ cultivar (The American Phytopathological Society 2008).

Seedlings of the barley cv. ‘Xena’ were also treated manually using a plastic-bottle sprayer with approximately 8 ml of coriolic acid, ricinoleic acid, and oleic acid at 0.12, 0.25, 0.5, 1, and 2 g/L concentrations four hours before inoculation. Infection by *P. teres* f. *teres* was performed as described by Akhavan et al. (2016) with some modifications. The seedlings at 3-leaf stage were inoculated by conidial suspension with concentration of  $1 \times 10^4$  spores/ml using an automatic sprayer connected to an air-line, then they were covered with moist plastic bags for 24 hours to preserve the relative humidity around 100%. After bag removal and plant transition to the growth cabinets (mentioned as above), the disease severity was measured six days post inoculation using APS Assess 2.0 software as described for ‘Westar’ cultivar (The American Phytopathological Society 2008).

In all plant experiments, sterile distilled water containing 0.05% Tween-20 (PH= 7.5) was used as the solvent of fatty acids at such low concentrations with two-fold ranging from 0.12 to 2 g/L (Bajpai et al. 2009). Treatment of seedlings with sterile distilled water containing 0.05% Tween-20 served as positive control when inoculated with specific fungal spore suspension and

served as negative control when un-inoculated. A fungicide, Bumper 418 EC (Propiconazole 418 g/L), also was used as a chemical control at a concentration of 1.5 ml/L which was applied on seedlings 36 hours before inoculation. Oleic acid also was applied on wheat and barley crops as another control of fatty acids without hydroxy group in its structure. Experiments were arranged in a randomized complete block design (RCBD) with three biological replicates for each treatment and were repeated twice for each crop. Six plastic inserts each with one seedling represented one biological replicate in case of 'Westar' cultivar and DH12075 line. A single pot with six seedlings represented one biological replicate in case of the 'Katepwa' and 'Xena' cultivars. In total, there were 18 replicates (6 technical replications  $\times$  3 biological replicates) for each treatment.

### **3.7. Seed germination assay**

The effects of coriolic acid, ricinoleic acid, and oleic acid was assessed on seed germination of 'Katepwa', 'Xena', 'Westar', and DH12075 cultivars/lines as described by Nwachukwu and Umechuruba (2001) with some modifications. Six seeds of each cultivar, as one biological replicate, were soaked in aqueous form of coriolic acid, ricinoleic acid and oleic acid at five different concentrations with two-fold dilution ranging from 0.12 g/L to 2 g/L for two hours. Seeds then were air-dried for two hours and placed on Petri-dishes covered with Whatman filter paper saturated with SDW. Petri-dishes were kept in an incubator set at of 20 °C and 16/8 hours of light/darkness to prepare suitable condition for seed germination. The Petri-dishes were watered regularly and monitored for 7 days for seed germination and seedling growth. A seed was considered to have germinated if both roots and shoots were present. Soaked-seeds in the fatty acid solvent (water-tween 0.05%) and in SDW served as the controls. The experiment was



designed as randomized complete block design (RCBD) and performed with three biological replicates over time for each treatment and repeated twice for each crop.

### **3.8. Seed treatment assay**

The efficacy of coriolic acid, ricinoleic acid, and oleic acid in crop protection as seed treatment compounds was assessed on ‘Katepwa’, ‘Westar’, and ‘DH12075’ cultivars/lines. Six seeds of each crop cultivar, as one biological replicate, were soaked in aqueous form of coriolic acid, ricinoleic acid, and oleic acid at five different concentrations ranging from 0.12 g/L to 2 g/L for two hours. Following that, seeds were sown in plastic inserts (5 cm x 5 cm; one seed per insert) filled with sterile Sunshine potting mix (W.R. Grace and Co., Fogelsville, PA, USA) and watered regularly. Seeds soaked in the solvent (water-tween 0.05%), without fatty acids, which were later inoculated with specific phytopathogens, served as the positive control. Seeds soaked in water-tween 0.05% served as a negative control and were not inoculated with the pathogen. Plant infection procedures by specific phytopathogens and disease severity measurements were performed as described previously in plant inoculations section. The experiment was designed as randomized complete block design (RCBD) and performed in three biological replicates for each treatment and repeated twice for each crop.

### **3.9. Histological and morphological studies**

The effect of coriolic acid, ricinoleic acid, and oleic acid treatment on clogging of stomatal pores was investigated using light microscopy. Two hours after spraying the 3-week old ‘Westar’ seedlings by two concentrations of these fatty acids (2 and 1 g/L), a small piece of leaf from each treatment including fatty acids-treated as well as controls was prepared (8 × 10 mm). The tissues were fixed in a solution consisting of formaldehyde solution (37%), acetic acid, and

ethanol (50%) overnight at RT (Yeung and Saxena 2005). After fixation, the tissues were sectioned paradermally (8  $\mu$ m thickness), affixed to glass slides, and stained as described by Sharma et al. (2010) with some modifications. Briefly, the affixed tissues were de-waxed in two series of toluene solutions each for 5 min, rehydrated to 50% ethanol and then stained with Lacto Phenol Cotton Blue (LPCB) for 20 minutes. Subsequently, the stained sections were washed 3 times with water (3 min each), counterstained with Safranin for 20 seconds, dehydrated in ethanol and toluene, and then mounted with DPX (Electron Microscopy Sciences, Hatfield, PA, USA) mounting medium. The sections were viewed using an Optronics digital camera interfaced to an Axio Scope.A1 - Zeiss Microscope (Carl Zeiss MicroImaging, Oberkochen, Germany).

The stomatal pores were also observed directly by a layer of leaf surface using an Axio Scope.A1 - Zeiss Microscope (Carl Zeiss MicroImaging, Oberkochen, Germany) for all treated and untreated leaves of the 'Westar' cultivar, then photographed with an Optronics digital camera. In addition, a relief map of the surface of all treated and untreated Westar leaves was prepared through painting each side of the midrib with a clear nail polish. When the nail polish thoroughly dried, the edge of the nail polish was carefully peeled off using a small piece of scotch tape. Then the nail polish was placed on a microscope slide and visualized under Axio Scope.A1 - Zeiss Microscope by an Optronics digital camera.

### **3.10. Determination of reactive oxygen species (ROS) levels**

Accumulation of two major types of ROS,  $O_2^-$  (superoxide) and  $H_2O_2$  (hydrogen peroxide) (O'Brien et al. 2012) was determined through histochemical staining of 'Westar' canola cultivars treated with coriolic acid, ricinoleic acid, and oleic acid at the two highest concentrations used in this study, 2 g/L and 1 g/L. Two different methods of fatty acids treatment were used for determination of ROS level including foliar and wounding treatments. Briefly, the

second leaf of 3-week old of 'Westar' seedlings were wounded at two sides of the mid-rib using a sterile needle gauge. Then 10 µl of coriolic acid, ricinoleic acid, and oleic acid at the mentioned concentrations was applied on each wounded site. While, in foliar treatment method, the 3-week 'Westar' seedlings were sprayed with 10 ml of coriolic acid, ricinoleic acid, and oleic acid at the aforementioned concentrations. Six hours later, the second leaf was excised from each wounded- and foliar-sprayed seedling (three seedlings for each treatment) and placed in staining falcon tubes separately. Untreated seedlings and treated ones by the solvent of the fatty acids which was water-tween 0.05%, served as the controls. Staining of the leaves with Nitroblue Tetrazolium (NBT) and 3,3'-Diaminobenzidine (DAB) was conducted as described by Kumar et al. (2014) Daudi and O' Brien (2012) with some modifications. Briefly, 16 ml of 1 M sodium phosphate monobasic solution ( $\text{NaH}_2\text{PO}_4$ ) and 84 ml of 1 M sodium phosphate dibasic solution ( $\text{Na}_2\text{HPO}_4$ ) were mixed together and then the volume increased to 2 L with SDW to prepare 50 mM sodium phosphate buffer (pH=7.5). Subsequently, 50 mg of DAB was dissolved in 45 ml of SDW in a bottle covered with aluminum foil. The pH was also adjusted to 3.8 while the solution was mixing properly on a magnetic stirrer. The total volume of the solution was also increased to 50 ml to get 1 mg/ml of DAB solution. For NBT staining, 100 mg of NBT was dissolved in 50 mM sodium phosphate buffer (pH=7.5) with volume of 50 ml to make a 0.2% NBT solution. The solution was blended on magnetic stirrer thoroughly in an amber bottle and prepared freshly before use. After preparation of DAB and NBT solutions freshly and incubation of treated and control leaves over-night in the staining falcon tubes covered with aluminum foil, the staining solutions were poured and replaced by bleaching solution (ethanol: acetic acid: glycerol= 3:1:1). After, the tubes were placed in boiling water bath at 90-95 °C for  $15 \pm 5$  min to bleach out the

chlorophyll of leaves, then the leaves were photographed on a white background under uniform lighting.

### **3.11. Statistical analysis**

Disease scores and seed germination data were analyzed using the MIXED procedure of SAS version 9.4 (Statistical Analysis system; SAS Institute Inc., Cary, NC, USA). The model for the experiment included different treatments (Negative control, Positive control, coriolic acid 0.12, coriolic acid 0.25, coriolic acid 0.5, coriolic acid 1, coriolic acid 2, oleic acid 0.12, oleic acid 0.25, oleic acid 0.5, oleic acid 1, oleic acid 2, ricinoleic acid 0.12, ricinoleic acid 0.25, ricinoleic acid 0.5, ricinoleic acid 1, ricinoleic acid 2 g/L, and Fungicide (Bumper 418 EC at concentration of 1.5 ml/L)) as the fixed independent variables and disease severity as the dependent variable. If the data for certain treatments were not normally distributed, the Box-Cox transformation within PROC TRANSREG was used to find the most appropriate data transformation. Differences between means were analyzed using a Tukey's Multiple Comparison Test with a 95% confidence level and were reported as a mean  $\pm$  the standard error of the mean (SEM). The cut of p-value of a significance was  $P < 0.05$ .

## 4. RESULTS

### 4.1. MICs of coriolic acid and ricinoleic acid against pathogenic fungi

The antifungal activity of coriolic acid and ricinoleic acid was determined using the broth micro-dilution method (Table 4-1). Coriolic acid and ricinoleic acid possessed the strongest inhibitory activity against *L. maculans* with MIC of 0.73 and 0.83 g/L, respectively, followed by *A. niger* with MIC values of 0.78 g/L for coriolic acid and 0.88 g/L for ricinoleic acid. A weaker inhibitory activity of coriolic acid was observed against the phytopathogens *S. sclerotiorum*, *P. teres* f. *teres* and *P. tritici-repentis* with MIC values of 2.92, 1.66, and 1.64 g/L, respectively. Ricinoleic acid also could inhibit the mycelial growth of *P. teres* f. *teres* and *P. tritici-repentis* at concentrations of 2.08 and 2.18 g/L, respectively. Amongst all pathogenic fungi used in this study, the weakest inhibitory activity of coriolic acid and ricinoleic acid were observed for *F. graminearum* with MIC of 7.50 and 8.33 g/L, respectively (Table 4-1).

Table 4-1. Minimum Inhibitory Concentrations (MICs) of coriolic acid and ricinoleic acid against phytopathogenic fungi

Fungal species	Mean MIC (g/L) $\pm$ SE <sup>a</sup>	
	Coriolic acid (13-OH C 18:2)	Ricinoleic acid (12-OH C 18:1)
<i>Leptosphaeia maculans</i>	0.73 $\pm$ 0.10	0.83 $\pm$ 0.10
<i>Aspergillus niger</i>	0.78 $\pm$ 0.10	0.88 $\pm$ 0.34
<i>Pyrenophora teres</i> f. <i>teres</i>	1.66 $\pm$ 0.20	2.08 $\pm$ 0.41
<i>Pyrenophora tritici-repentis</i>	1.64 $\pm$ 0.46	2.18 $\pm$ 0.32
<i>Sclerotinia sclerotiorum</i>	2.92 $\pm$ 1.09	ND <sup>b</sup>
<i>Fusarium graminearum</i>	7.50 $\pm$ 1.44	8.33 $\pm$ 1.66

<sup>a</sup> SE: standard error of the mean

<sup>b</sup> ND: not determined

## 4.2. Effects of foliar treatment of UFAs on plant-pathogen interaction

### 4.2.1. Dicotyledons

The effects of coriolic acid and ricinoleic acid were investigated through foliar spraying on whole plant systems. Two genotypes of canola including Westar and DH12075 infected by *L. maculans* and *S. sclerotiorum*, respectively, were tested for the inhibitory activity of coriolic acid and ricinoleic acid. There was no significant reduction on disease severity of blackleg caused by *L. maculans* in ‘Westar’ cultivar treated with five different concentrations of coriolic acid and ricinoleic acid including 0.12, 0.25, 0.5, 1, and 2 g/L compared with the disease severity of a positive control (3.99%) (Figure 4-1). Similar observations were also made for coriolic acid- and ricinoleic acid-treated DH12075 canola line inoculated with *S. sclerotiorum* in comparison with the disease severity of positive control 18.33% (Figure 4-2). The highest concentration of coriolic acid and ricinoleic acid (2 g/L) used in this study caused extensive necrotic lesions on the leaves of both genotypes of canola including ‘Westar’ and DH12075, which ultimately contributed to death of these cultivars after couple of days (Figure 4-3).

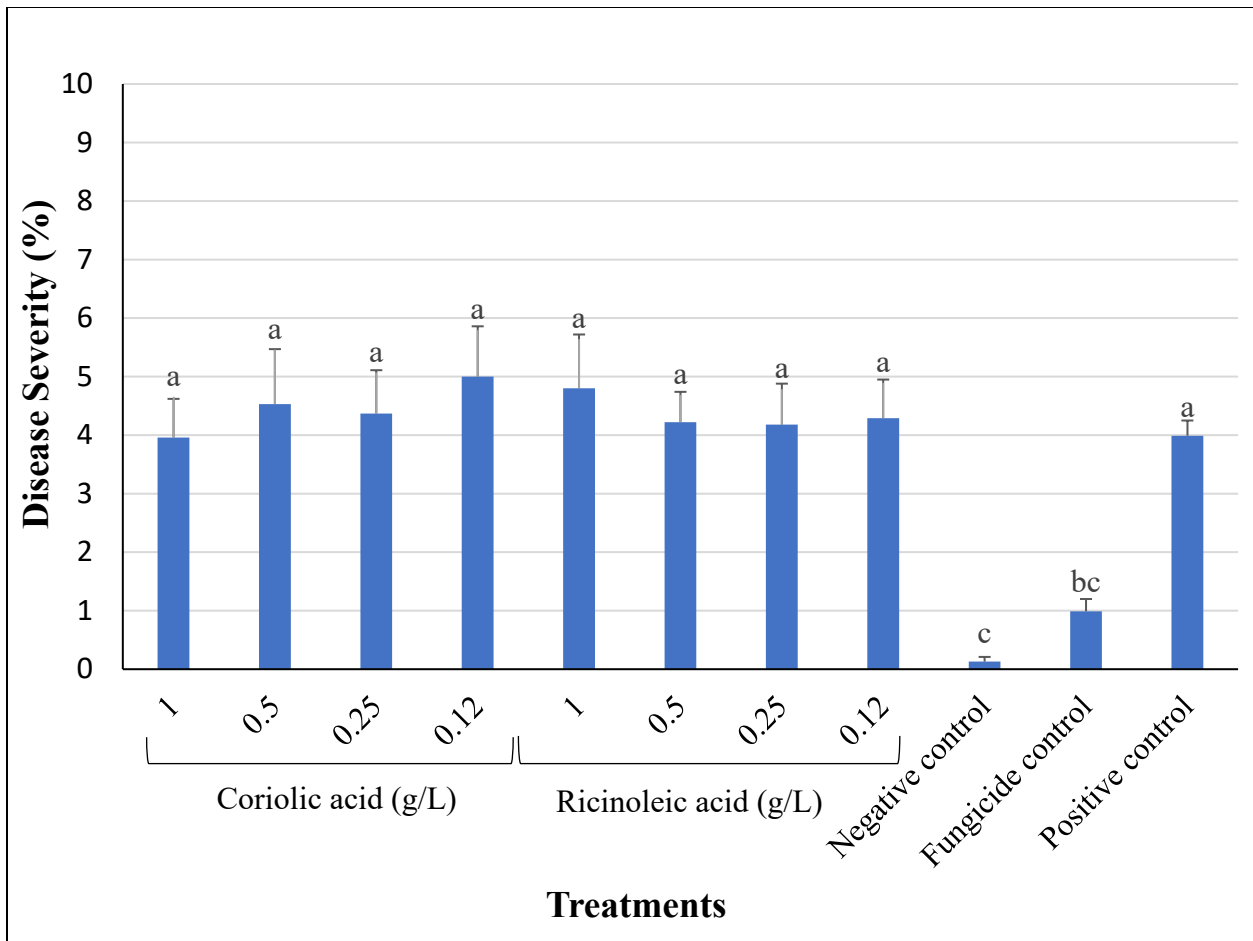


Figure 4-1. Effects of foliar treatment of coriolic acid and ricinoleic acid on disease severity induced by *Leptosphaeria maculans* in *Brassica napus* cv. ‘Westar’ under controlled-environment growth chambers conditions. Bars with the same lowercase letter indicate means that do not differ significantly ( $P < 0.05$ ) based on Tukey's Multiple Comparison Test. Untransformed means are presented  $\pm$  standard error of the means.



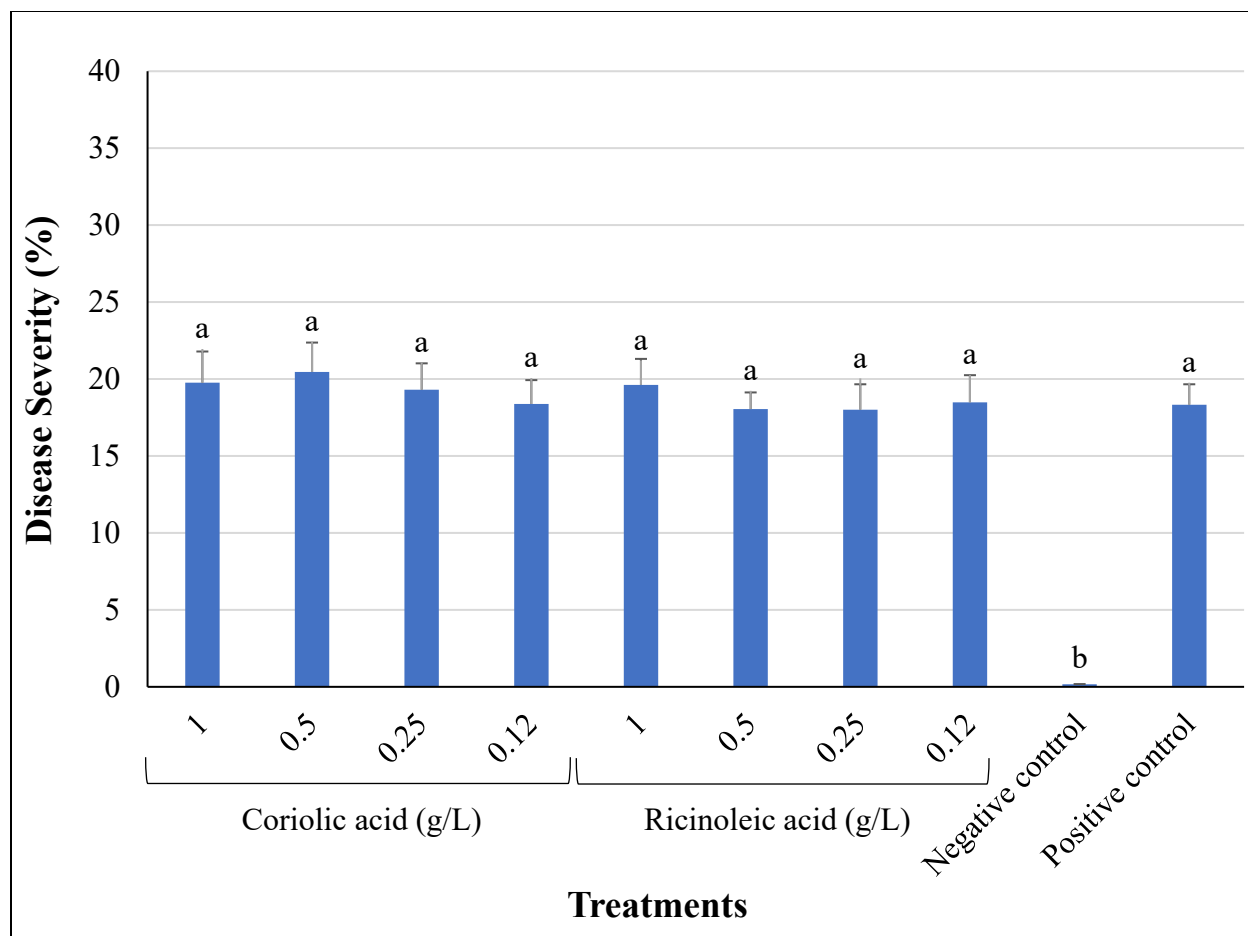


Figure 4-2. Effects of foliar treatment of coriolic acid and ricinoleic acid on disease severity induced by *Sclerotinia sclerotiorum* in *Brassica napus* line DH12075 under controlled-environment growth chambers conditions. Bars with the same lowercase letter indicate means that do not differ significantly ( $P < 0.05$ ) based on Tukey's Multiple Comparison Test. Untransformed means are presented  $\pm$  standard error of the means.

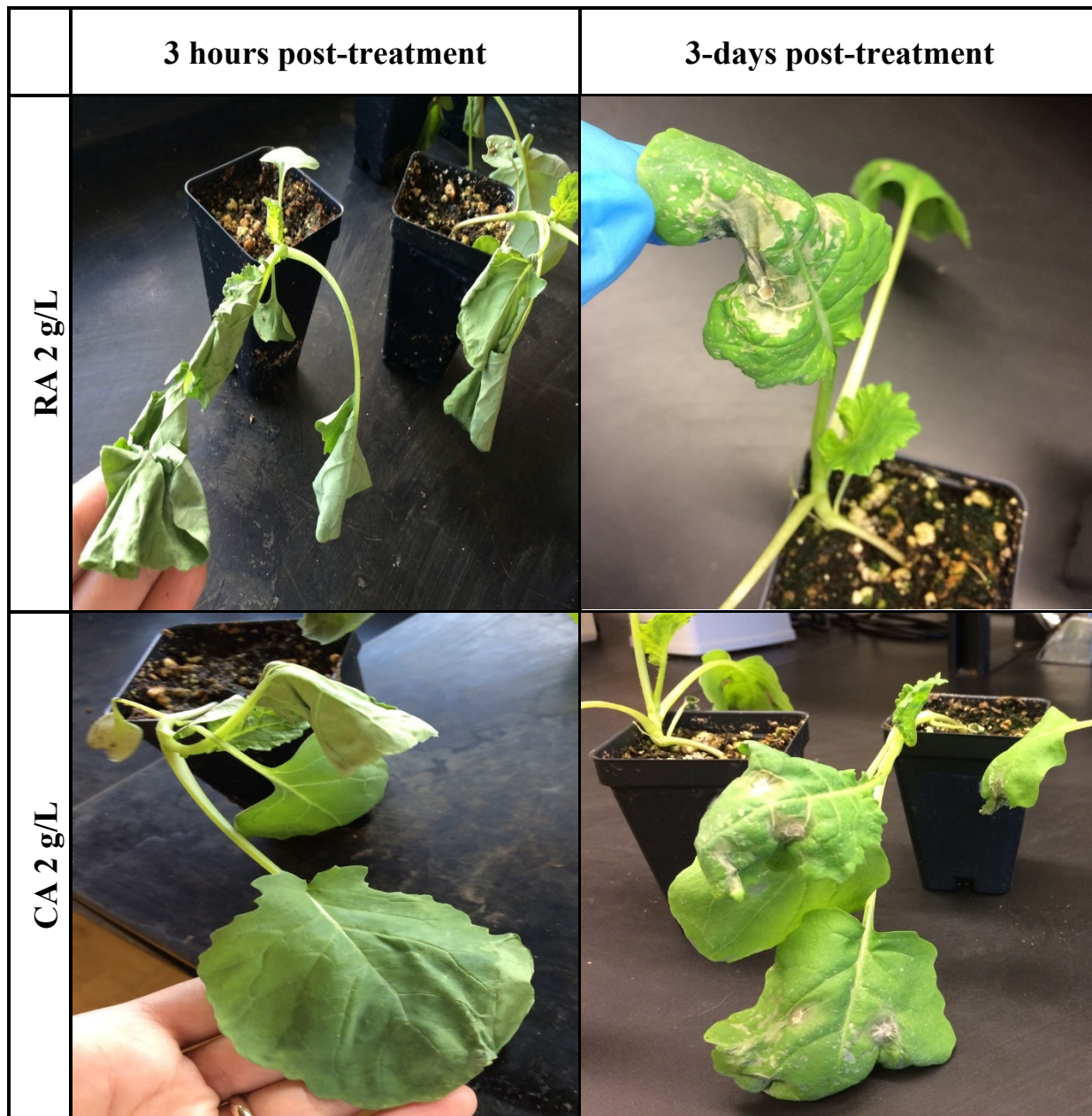


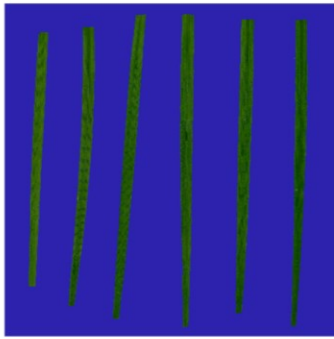
Figure 4-3. Effects of foliar treatment of coriolic acid and ricinoleic acid at 2 g/L concentration in three-week old *Brassica napus* cv. 'Westar' 3 hours and 3 days post treatment, which caused wilting and formation of necrotic lesions on the seedlings.

#### 4.2.2. Monocotyledons

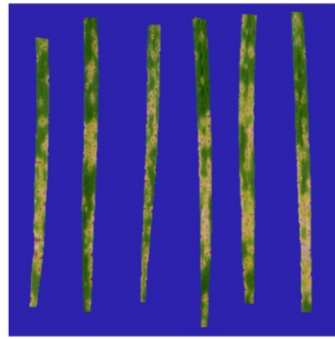
The effects of coriolic acid, ricinoleic acid, and oleic acid were investigated through foliar spraying on whole plant systems. Coriolic acid-treated wheat inoculated with *P. tritici-repentis* (isolate AB 7-2) resulted in a reduction of visual symptoms (Figure 4-4) and a significant ( $P < 0.05$ ) reduction in disease severity at concentrations of 2 (1.26%), 1 (3.16%), and 0.5 (12.30%) g/L when compared with the positive control with disease severity of 21.23% (Figure 4-5). Additionally, higher concentrations of coriolic acid (i.e. 2 and 1 g/L) showed significant ( $P < 0.05$ ) disease reduction in comparison with a concentration of 0.5 g/L. A statistically significant effect on disease severity was not observed in wheat seedlings treated with ricinoleic acid and oleic acid. These results suggest that, under the experimental conditions tested, coriolic acid afforded a higher degree of protection in wheat against this pathogen.

In barley, coriolic acid-treated seedlings at 0.5 g/L concentration showed a significant ( $P < 0.05$ ) reduction in disease symptoms caused by *P. teres* f. *teres* (isolate AB 34) (30.64%) compared with the positive control (59.00%) (Figure 4-6). Furthermore, coriolic acid at 2 and 1 g/L concentrations decreased the disease severity on barley seedlings infected by *P. teres* f. *teres* by 40.93% and 36.25%, respectively. No significant difference was observed amongst the five different concentrations of ricinoleic acid and oleic acid including 2, 1, 0.5, 0.25, and 0.12 g/L on the disease severity caused by *P. teres* f. *teres* in barley seedlings compared with the positive control (Figure 4-6). Similar to wheat, ricinoleic acid and oleic acid were not effective to control the disease severity caused by *P. teres* f. *teres* in barley seedlings. Although coriolic acid was effective to control the disease severity on barley but it was not as effective as on wheat. Formation of necrotic lesions on wheat and barley seedlings were in a similar manner with the

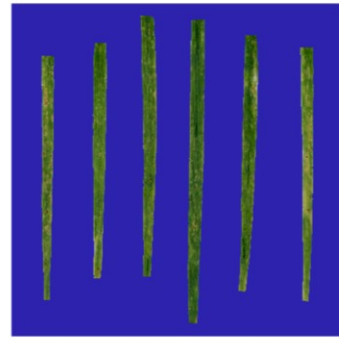
canola cultivars but their numbers were much fewer which did not kill these monocotyledonous crops.



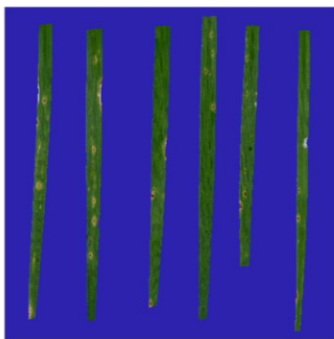
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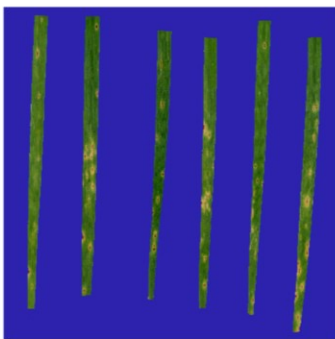
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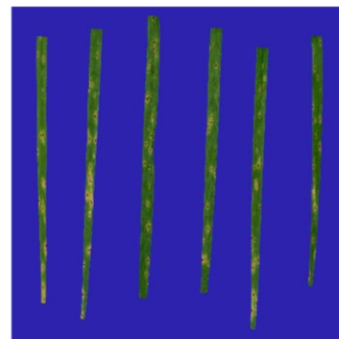
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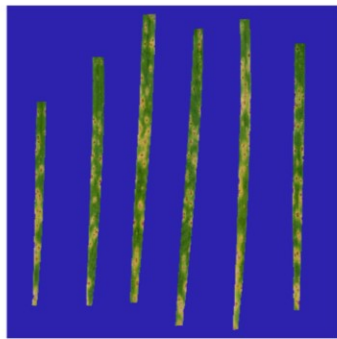
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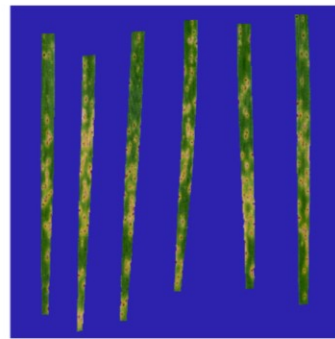
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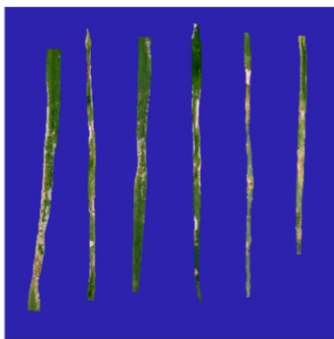
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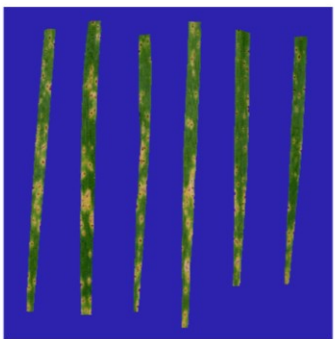
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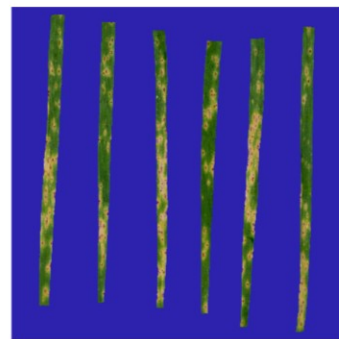
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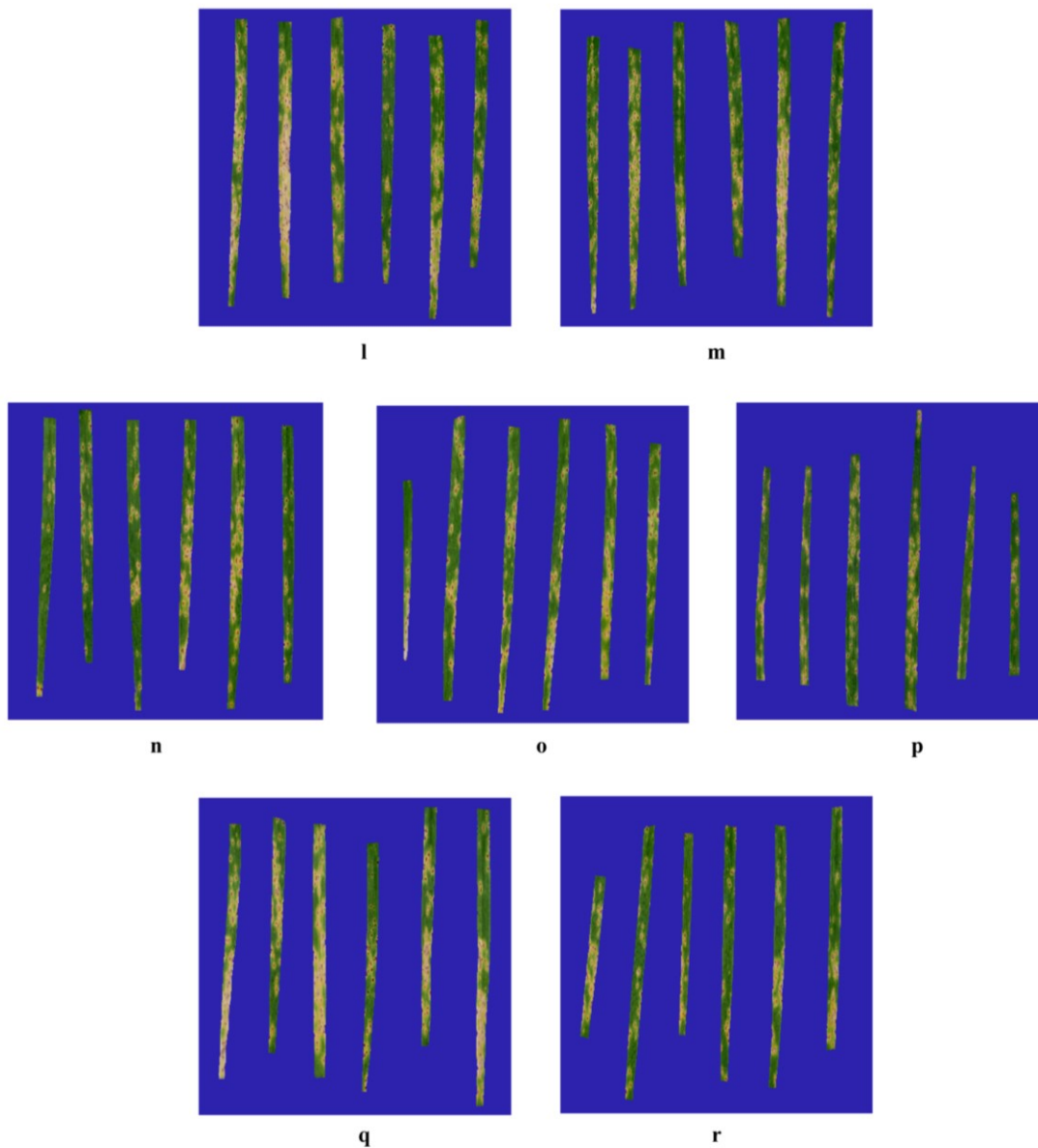


Figure 4-4. Effects of foliar treatment of unsaturated fatty acids on the disease severity induced by *Pyrenophora tritici-repentis* in *Triticum aestivum* cv. 'Katepwa'. [a] negative control, [b] positive control, [c] fungicide control, [d] coriolic acid 2 g/L, [e] coriolic acid 1 g/L, [f] coriolic acid 0.5 g/L, [g] coriolic acid 0.25 g/L, [h] coriolic acid 0.12 g/L, [i] ricinoleic acid 2 g/L, [j] ricinoleic acid 1 g/L, [k] ricinoleic acid 0.5 g/L, [l] ricinoleic acid 0.25 g/L, [m] ricinoleic acid 0.12 g/L, [n] oleic acid 2 g/L, [o] oleic acid 1 g/L, [p] oleic acid 0.5 g/L, [q] oleic acid 0.25 g/L, [r] oleic acid 0.12 g/L. Photos were taken 6 days after inoculation.

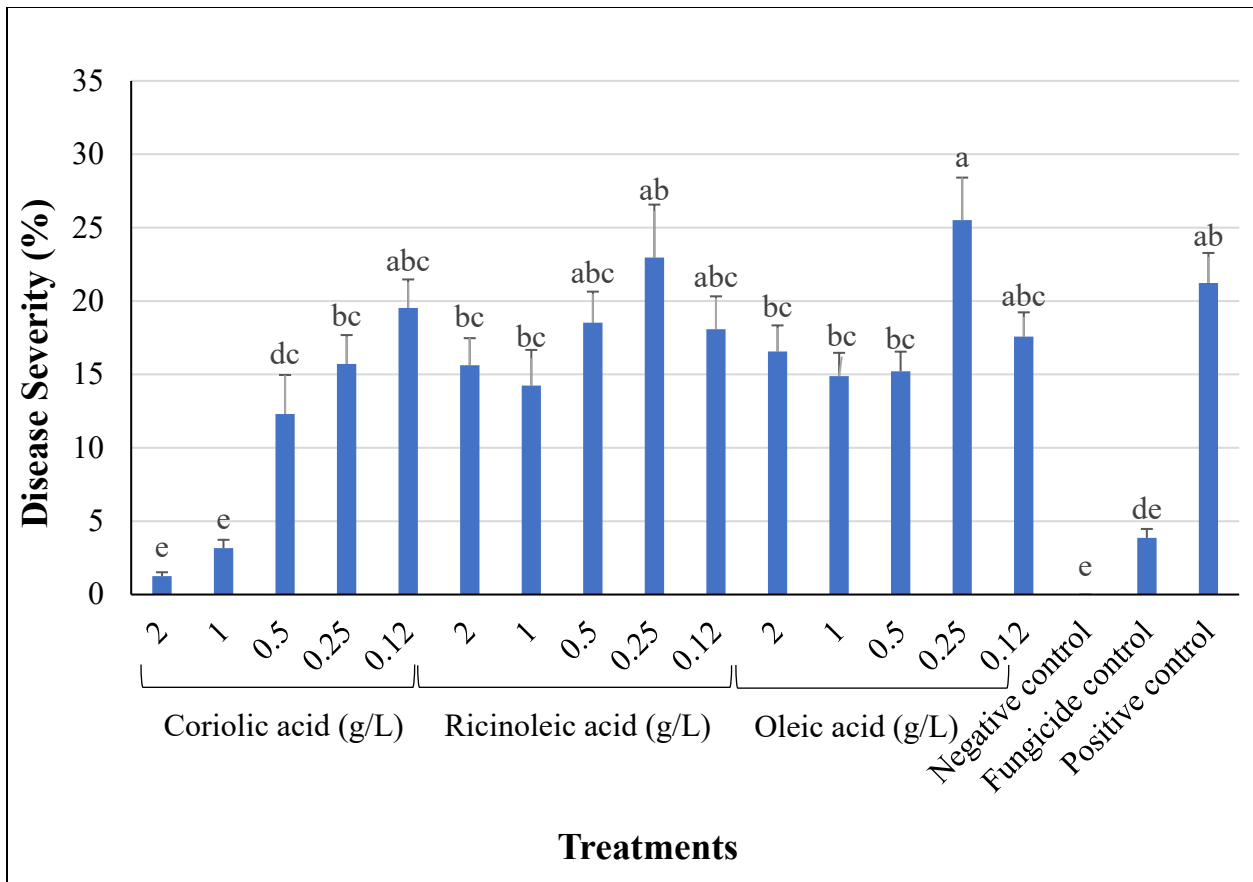


Figure 4-5. Effects of foliar treatment of coriolic acid, ricinoleic acid, and oleic acid on the disease severity induced by *Pyrenophora tritici-repentis* in *Triticum aestivum* cv. ‘Katepwa’ under controlled-environment growth chambers conditions. Bars with the same lowercase letter indicate means that do not differ significantly ( $P < 0.05$ ) based on Tukey's Multiple Comparison Test. Untransformed means are presented  $\pm$  standard error of the means.

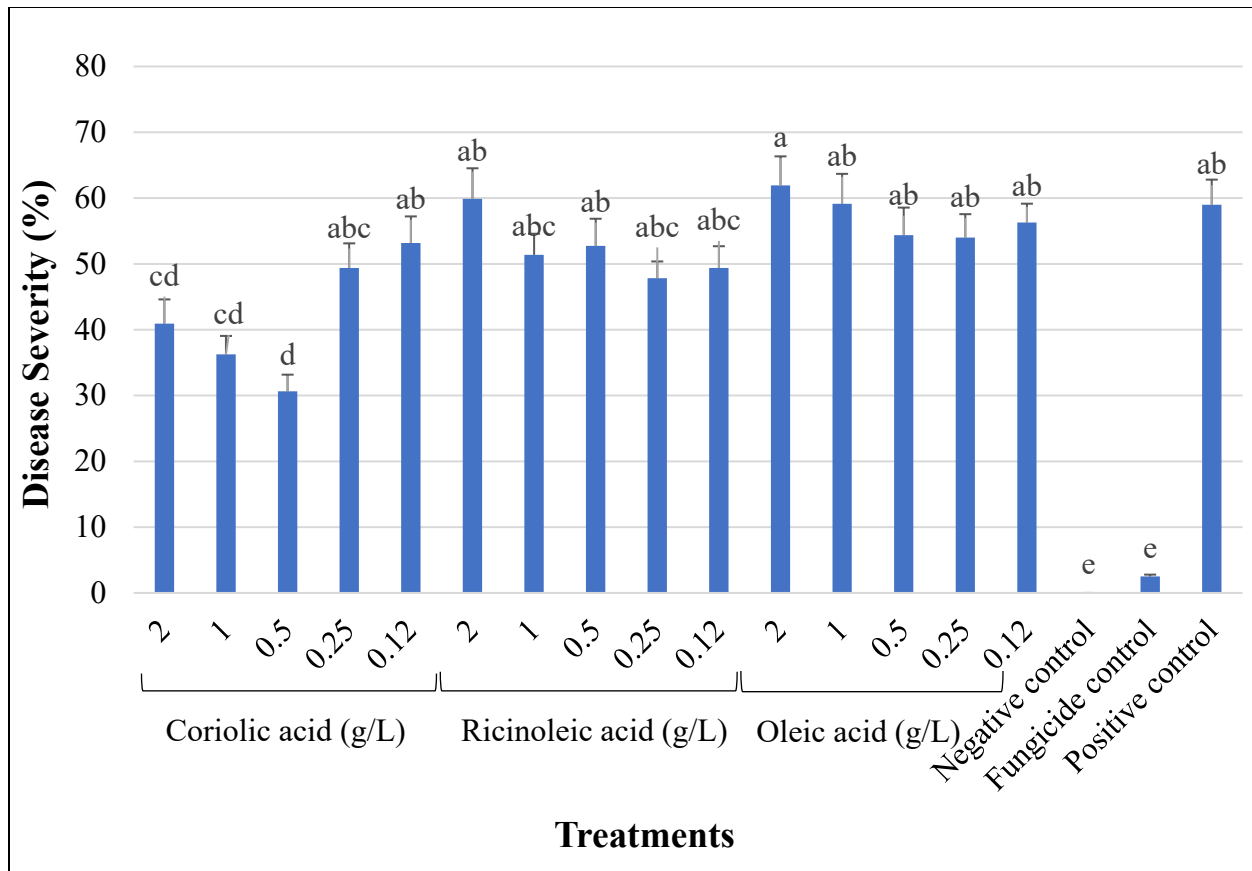


Figure 4-6. Effects of foliar treatment of coriolic acid, ricinoleic acid, and oleic acid on the disease severity induced by *Pyrenophora teres* f. *teres* in *Hordeum vulgare* cv. ‘Xena’ under controlled-environment growth chambers conditions. Bars with the same lowercase letter indicate means that do not differ significantly ( $P < 0.05$ ) based on Tukey's Multiple Comparison Test. Untransformed means are presented  $\pm$  standard error of the means.



### **4.3. Effects of coriolic acid, ricinoleic acid and oleic acid on seed germination**

The effects of coriolic acid, ricinoleic acid and oleic acid were assessed on seed germination of wheat, barley and canola genotypes (Westar and DH12075). The results indicated that coriolic acid, ricinoleic acid, and oleic acid did not have any inhibitory effect on seed germination of wheat and 'Westar' compared with the controls when their seeds were tested on moist Whatman paper (Table 4-2). However, in terms of canola line DH12075, coriolic acid at 2 g/L concentration showed lower seed germination compared to controls and other treatments. On barley, ricinoleic acid with 0.25 and 0.12 g/L concentrations had a better seed germination compared to other treatments, with the lowest germination rate for coriolic acid at 2 g/L concentration (Table 4-2).

Table 4-2. Effects of treatment of coriolic acid, ricinoleic acid, and oleic acid at concentrations with two-fold dilutions ranging from 0.12 to 2 g/L on seed germination of wheat, Barley, and Canola

Treatment	Canola-Westar (%)	Canola-DH12075 (%)	Wheat-Katepwa (%)	Barley-Xena (%)
Coriolic acid 2 g/L	94.44 <sup>a</sup> ± 5.55	88.88 <sup>b</sup> ± 5.55	88.88 <sup>b</sup> ± 5.55	27.77 <sup>c</sup> ± 5.55
Coriolic acid 1 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	88.88 <sup>b</sup> ± 5.55	50.18 <sup>abc</sup> ± 16.85
Coriolic acid 0.5 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	94.44 <sup>a</sup> ± 5.55	66.85 <sup>abc</sup> ± 9.78
Coriolic acid 0.25 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	88.88 <sup>b</sup> ± 5.55	61.11 <sup>abc</sup> ± 5.55
Coriolic acid 0.12 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	88.88 <sup>b</sup> ± 5.55	61.11 <sup>abc</sup> ± 5.55
Ricinoleic acid 2 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	83.33 <sup>a</sup> ± 9.62	38.88 <sup>bc</sup> ± 5.55
Ricinoleic acid 1 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	66.85 <sup>abc</sup> ± 9.78
Ricinoleic acid 0.5 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	55.55 <sup>abc</sup> ± 5.55
Ricinoleic acid 0.25 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	94.44 <sup>a</sup> ± 5.55	83.88 <sup>a</sup> ± 0.00
Ricinoleic acid 0.12 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	88.88 <sup>b</sup> ± 5.55	83.51 <sup>a</sup> ± 9.62
Oleic acid 2 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	83.33 <sup>a</sup> ± 9.62	55.55 <sup>abc</sup> ± 11.11
Oleic acid 1 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	83.33 <sup>a</sup> ± 9.62	78.14 <sup>ab</sup> ± 5.74
Oleic acid 0.5 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	88.88 <sup>b</sup> ± 5.55	55.55 <sup>abc</sup> ± 5.55
Oleic acid 0.25 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	94.44 <sup>a</sup> ± 5.55	72.40 <sup>ab</sup> ± 5.74
Oleic acid 0.12 g/L	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	94.44 <sup>a</sup> ± 5.55	66.66 <sup>abc</sup> ± 0.00
Solvent control	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	94.44 <sup>a</sup> ± 5.55	61.11 <sup>abc</sup> ± 5.55
SDW control	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	100 <sup>a</sup> ± 0.00	61.29 <sup>abc</sup> ± 14.83

Values show the mean percentage of germination for each treatment ± standard error of the mean which is followed by different letters, showing significant ( $P < 0.05$ ) difference between treatments based on Tukey's Multiple Comparison Test.

#### 4.4. Effects of seed treatment of UFAs on plant-pathogen interaction

The effect of seed treatment with coriolic acid, ricinoleic acid, and oleic acid was also investigated on the disease severity induced by *P. tritici-repentis*, *L. maculans*, and *S. sclerotiorum* in wheat cv. 'Katepwa', canola cv. 'Westar', and canola line DH12075, respectively. No significant reduction was observed on disease severity caused by *P. tritici-repentis* in wheat cv. 'Katepwa' (Figure 4-7), *L. maculans* in canola cv. 'Westar' (Figure 4-8), and *S. sclerotiorum* in canola line DH12075 (Figure 4-9) compared with the positive control, when their seeds were treated with five different concentrations of coriolic acid, ricinoleic acid, and oleic acid at two-fold dilutions ranging from 0.12 to 2 g/L.

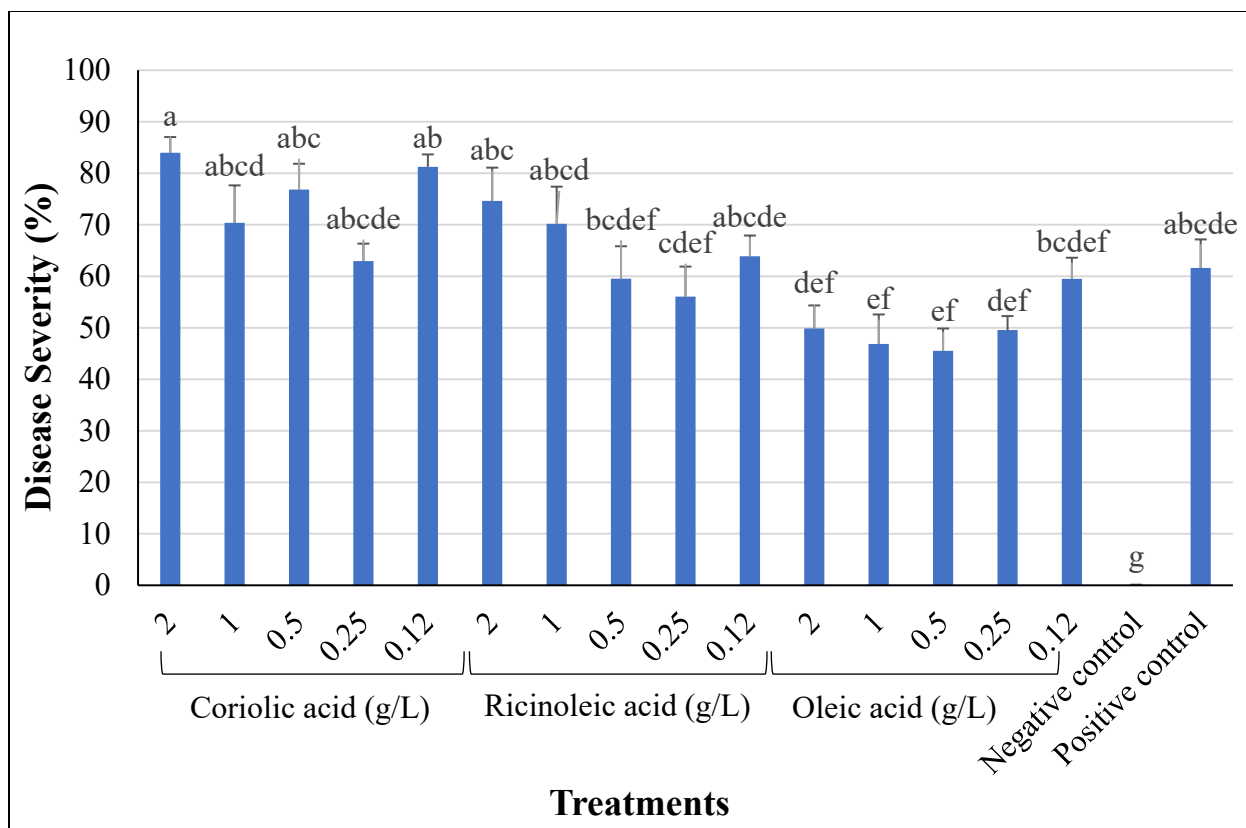


Figure 4-7. Effects of seed treatment of coriolic acid, ricinoleic acid, and oleic acid on the disease severity induced by *Pyrenophora tritici-repentis* in *Triticum aestivum* cv. ‘Katepwa’ under controlled-environment growth chambers conditions. Columns with the same lowercase letter indicate means that do not differ significantly ( $P < 0.05$ ) based on Tukey's Multiple Comparison Test. Untransformed means are presented  $\pm$  standard error of the means.

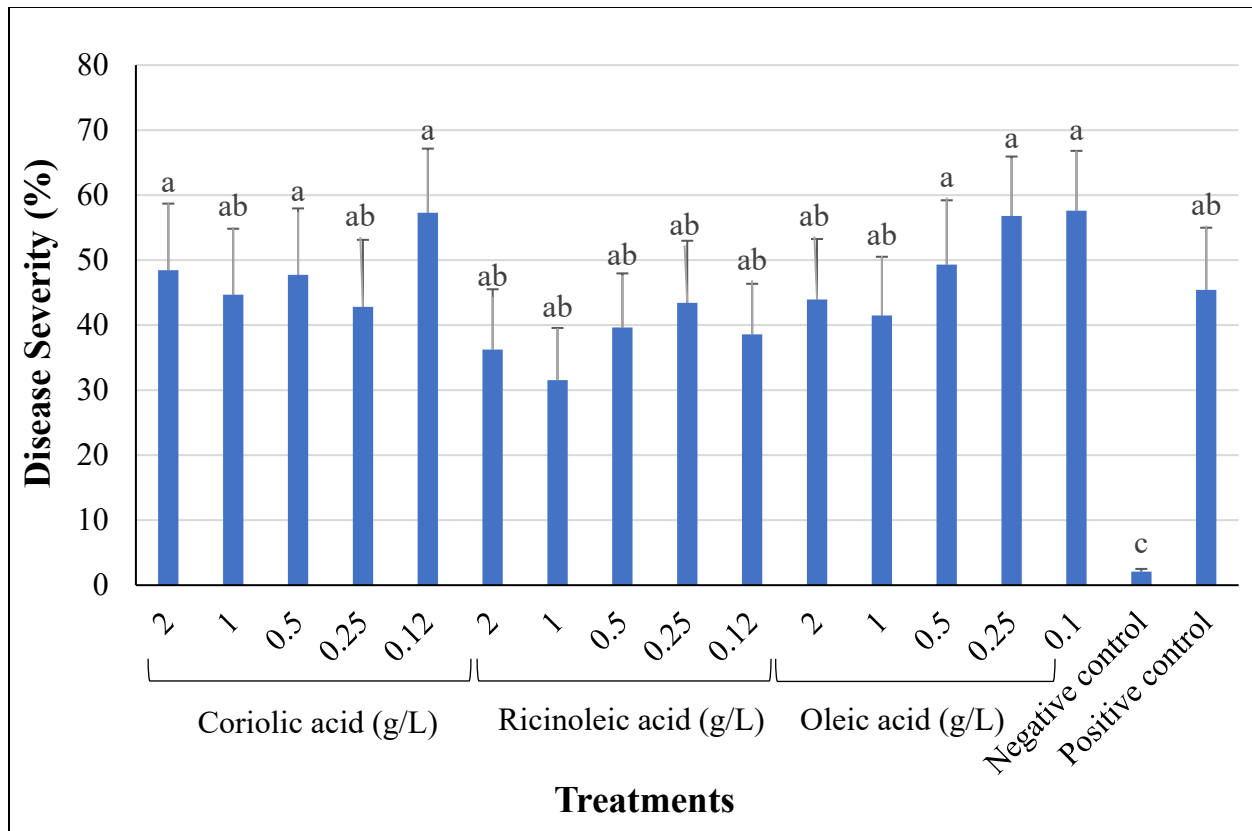


Figure 4-8. Effects of seed treatment of coriolic acid, ricinoleic acid, and oleic acid on the disease severity induced by *Leptosphaeria maculans* in *Brassica napus* cv. 'Westar' under controlled-environment growth chambers conditions. Columns with the same lowercase letter indicate means that do not differ significantly ( $P < 0.05$ ) based on Tukey's Multiple Comparison Test. Untransformed means are presented  $\pm$  standard error of the means.

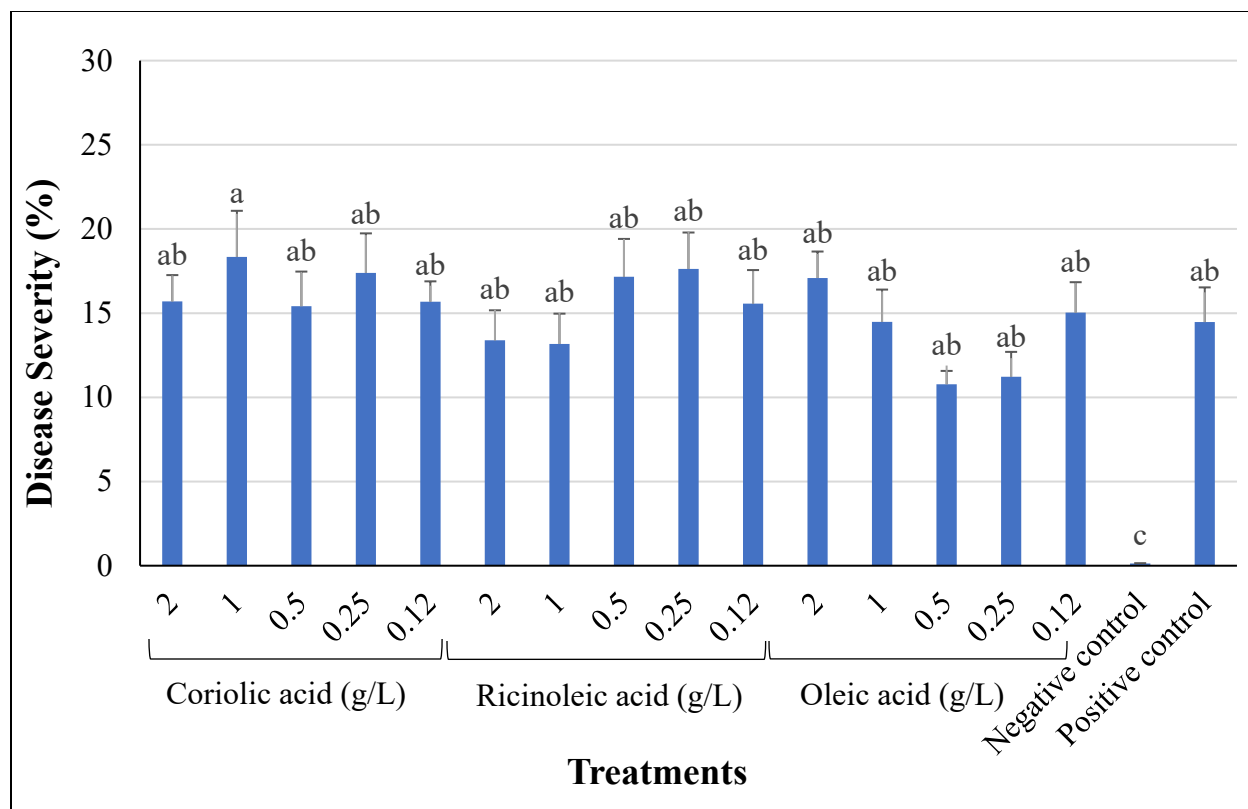
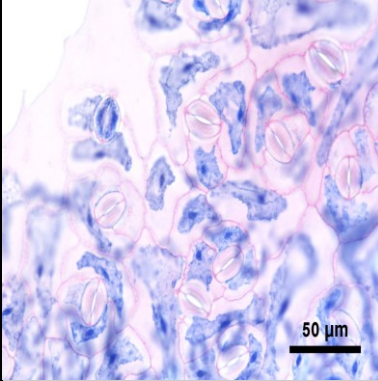
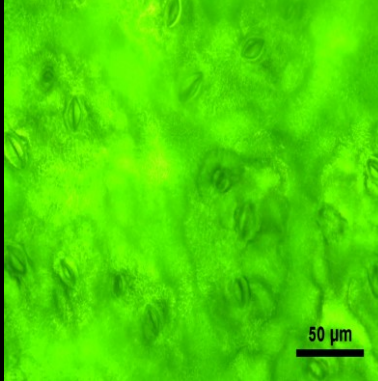
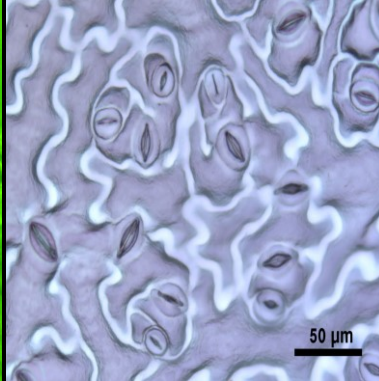
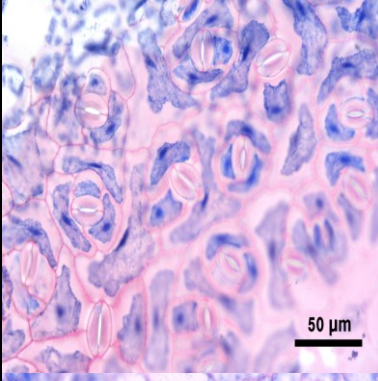
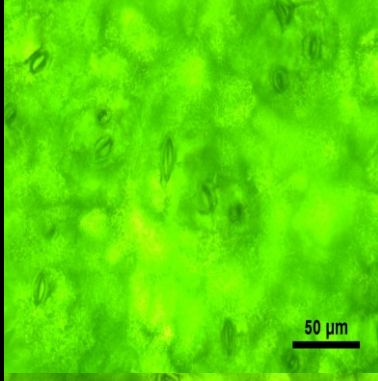
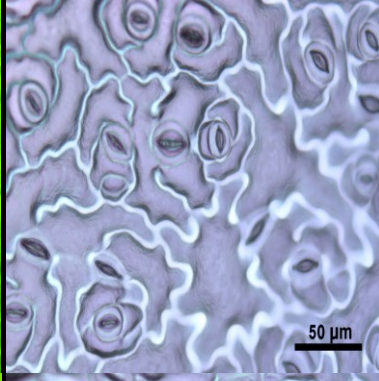
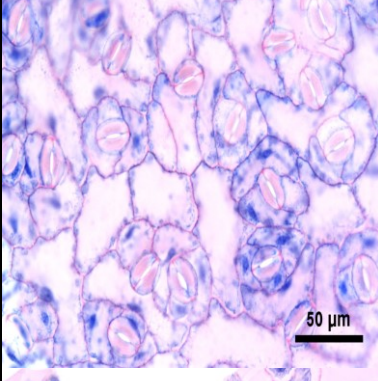
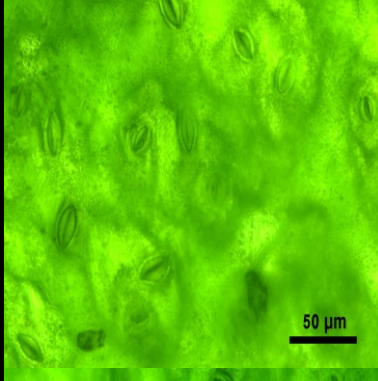
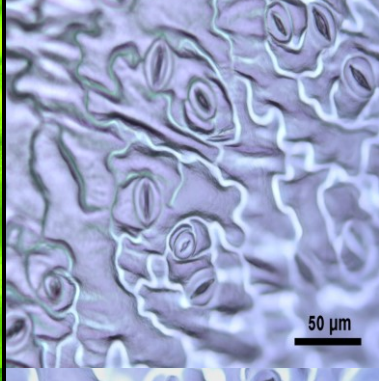
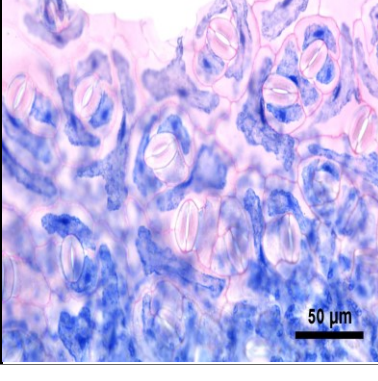
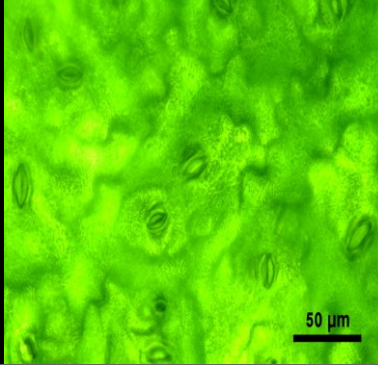
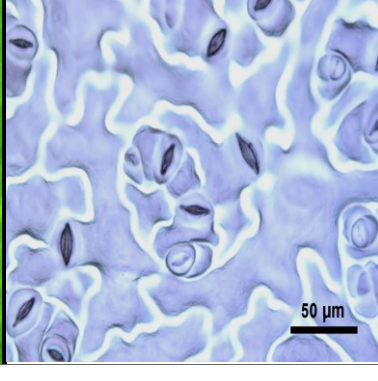


Figure 4-9. Effects of seed treatment of coriolic acid, ricinoleic acid, and oleic acid on the disease severity induced by *S. sclerotiorum* in *Brassica napus* line DH12075 under controlled-environment growth chambers conditions. Columns with the same lowercase letter indicate means that do not differ significantly ( $P < 0.05$ ) based on Tukey's Multiple Comparison Test. Untransformed means are presented  $\pm$  standard error of the means.

#### **4.5. Histological and morphological studies**

Spraying canola seedlings with coriolic acid and ricinoleic acid at 2 and 1 g/L concentrations led to formation of necrotic lesions on the leaves, followed by wilting of the plants. It was hypothesized that such high concentrations may clogged the stomatal pores and inhibit transpiration, which ultimately contributes to the senescence and death of the canola seedlings. Therefore, the effects of coriolic acid, ricinoleic acid, and oleic acid treatments at 2 and 1 g/L concentrations were investigated via light microscopy method as well as direct observation of leaves under the microscope to understand whether there are any lipid layer levels on stomatal pores and evidence of clogging. The results revealed no evidence of oily layers on stomatal pores between all the UFAs-treated leaves as well as control leaves (Figure 4-10).

	Light microscopy by LPCB & safranin staining (A)	Direct observation of leaf surface (B)	Direct observation of relief map of leaf surface (C)
Coriolic acid 2 g/L			
Coriolic acid 1g/L			
Ricinoleic acid 2 g/L			
Ricinoleic acid 1 g/L			



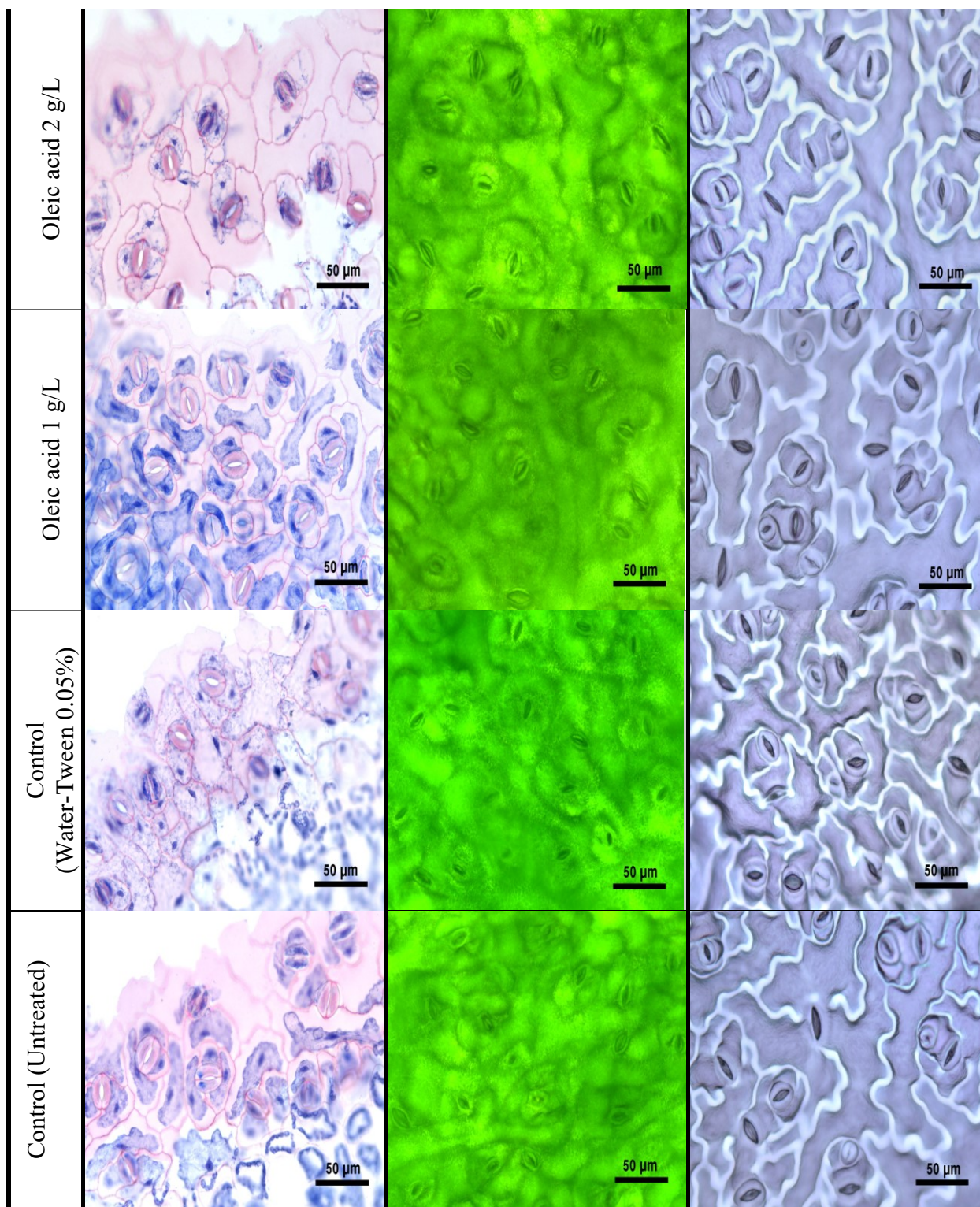












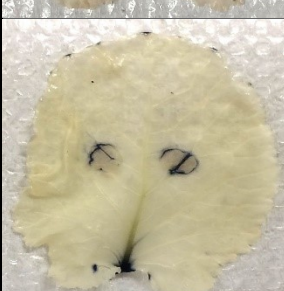









Figure 4-10. Effects of high concentrations (2 and 1 g/L) of coriolic acid, ricinoleic acid, and oleic acid on stomatal pores of *Brassica napus* cv. 'Westar' using light microscopy via Lacto Phenol Cotton Blue and safranin staining (A), direct observation of leaf surface (B), and relief map (C), via optronics digital camera interfaced to Axio Scope.A1 - Zeiss microscope with magnification of 40× (Scale bar = 50 μm).

#### 4.6. Effects of coriolic acid, ricinoleic acid, and oleic acid on the ROS levels

To understand whether coriolic acid, ricinoleic acid and oleic acid triggered the oxidative stress through disturbance of a steady-state level of ROS in canola plants, accumulation of  $\text{H}_2\text{O}_2$  and  $\text{O}_2^-$  was detected in the canola cv. 'Westar' after treatment with concentrations of 1 and 2 g/L of coriolic acid, ricinoleic acid and oleic acid using histochemical staining. The histochemical staining of  $\text{O}_2^-$  with NBT indicated that the oleic acid-treated leaves stained heavily as a result of  $\text{O}_2^-$  accumulation (Figure 4-11A). However, compared with controls, no staining of  $\text{O}_2^-$  was observed for coriolic acid- and ricinoleic acid-treated leaves using both foliar- and wounding-treatment procedures (Figure 4-11A).  $\text{H}_2\text{O}_2$  accumulation was also detected using histochemical staining with DAB. DAB reacts with  $\text{H}_2\text{O}_2$  in the presence of peroxidases to produce a brown polymerization product. Treatment of leaves with coriolic acid, ricinoleic acid and oleic acid at 2 g/L concentration induced intense staining as compared with the controls, indicating the accumulation of  $\text{H}_2\text{O}_2$  in 2 g/L UFA-treated leaves (Figure 4-11B). In contrast, coriolic acid-, ricinoleic acid-, and oleic acid-treated leaves at 1 g/L concentration had similar staining patterns as controls, suggesting that all these treatments probably had similar accumulation of  $\text{H}_2\text{O}_2$  in their leaves (Figure 4-11B).

	NBT staining (A)		DAB Staining (B)	
	Wounding treatment	Foliar treatment	Wounding treatment	Foliar treatment
Coriolic acid 2 g/L				
Coriolic acid 1g/L				
Ricinoleic acid 2 g/L				
Ricinoleic acid 1 g/L				
Oleic acid 2 g/L				

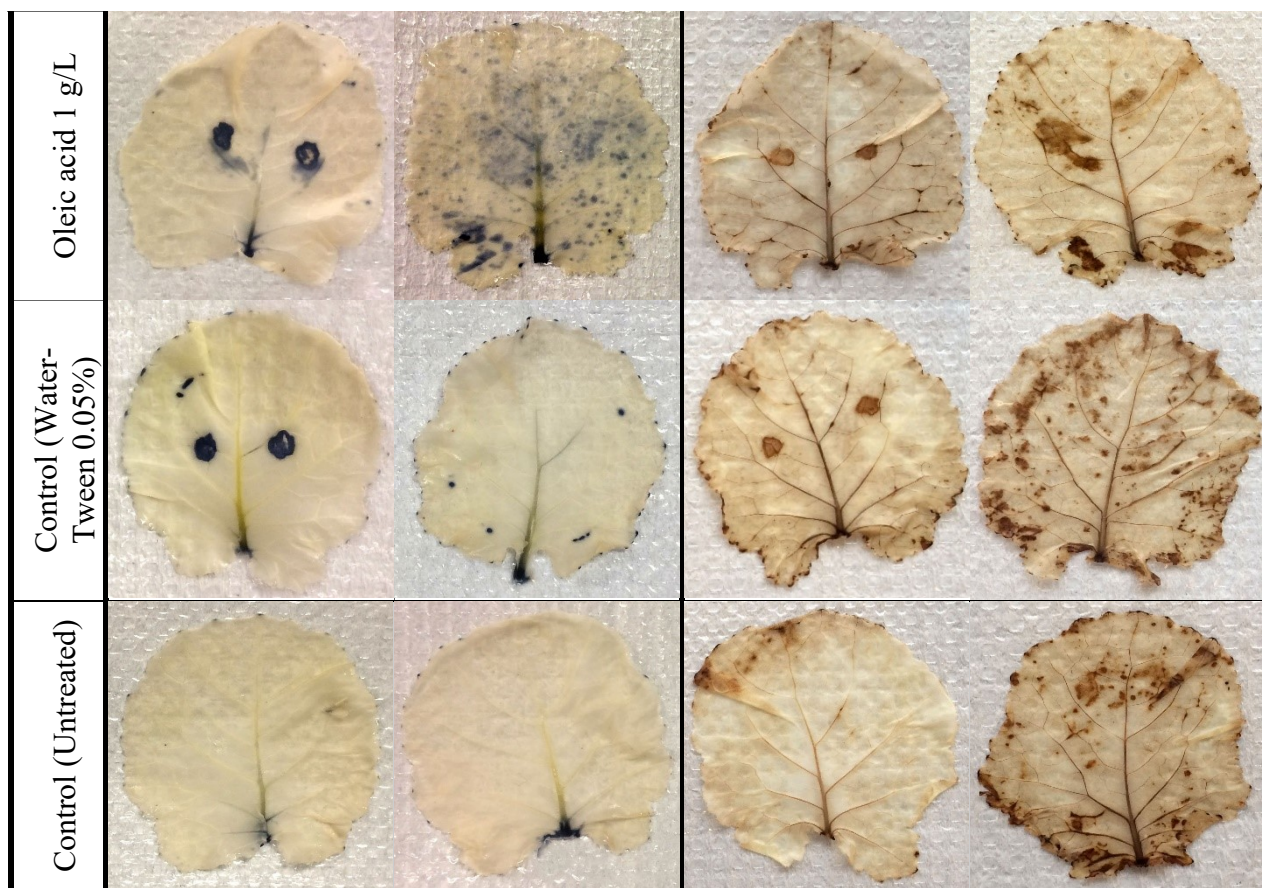


Figure 4-11. Effects of coriolic acid, ricinoleic acid and oleic acid at 2 and 1 g/L concentrations on the accumulation of  $O_2$  and  $H_2O_2$  in the leaves of *Brassica napus* cv. 'Westar' using Nitroblue Tetrazolium (A) and 3,3'-Diaminobenzidine (B) staining procedures.

## 5. DISCUSSION

Results described in this thesis provide evidence that HUFAs such as coriolic acid are potent inhibitors of plant pathogenic fungi *in vitro* and *in vivo*. It appeared that coriolic acid and ricinoleic acid exhibited large antifungal activity in active concentrations *in vitro*, affecting different pathogenic fungi. The lowest MICs of coriolic acid and ricinoleic acid were observed for *L. maculans* respectively by 0.73 and 0.83 g/L, indicating that coriolic acid and ricinoleic acid at lower concentrations than that had no effect on *L. maculans* mycelial growth. Our observation is similar with the results obtained by Graner et al. (2003), where the mycelial growth of *L. maculans* could not be inhibited by 0.29 g/L concentration of coriolic acid. *A. niger* was the second most sensitive fungi to coriolic acid and ricinoleic acid with MICs of 0.78 and 0.88 g/L, respectively, being in general agreement with the results obtained in previous studies (Chen et al. 2016; Liang et al. 2017). Two closely-related pathogens, *P. tritici-repentis* and *P. teres* f. *teres*, showed a similar range of MICs for coriolic acid and ricinoleic acid. Although there is no published research in the literature indicating the antifungal activity of coriolic acid and ricinoleic acid against these two species, there are studies showing the antifungal activity of other related UFAs such as linoleic acid and linolenic acid against *P. avenae* (Walters et al. 2004). Coriolic acid at concentrations lower than 2.92 g/L had no inhibitory activity against *S. sclerotiorum*. Similarly, a Graner et al. (2003) study on inhibitory effect of different concentrations of coriolic acid against *S. sclerotiorum* indicated that coriolic acid at 0.29 g/L or lower did not affect the growth rate of *S. sclerotiorum* mycelium, considering 0.29 g/L was the highest concentration they used in their experiment while the highest concentration that we used was 20 g/L to determine MIC. The differences in the MICs of coriolic acid and ricinoleic acid against *F. graminearum* were as high as ten-fold as compared with MICs against *L. maculans*,

indicating antifungal activity to be specific for different fungi (Graner et al. 2003; Prost et al. 2005; Pohl et al. 2011).

An explanation for differences in antifungal activities may be related to the diversity in the chemical composition of the membranes of different fungi such as sterol content (Avis and Belanger 2001). The general mechanism of antifungal UFAs with cell membranes has been described as detergent-like properties, in which antifungal UFAs directly insert into the phospholipid bilayer of the fungal membranes, disturb the membrane physically, and release intracellular electrolytes and proteins due to increased fungal membrane fluidity, ultimately leading to cytoplasmic disintegration of fungal cells (Avis and Belanger 2001; Pohl et al. 2011). Fungal sterols such as ergosterol, can moderate such stress-induced elevations in the membrane fluidity (Avis and Belanger 2001; Pohl et al. 2011); therefore, differences in sterol content of fungal membranes may lead to variations in fungal sensitivity. However, the antifungal activity of UFAs may also be linked to other features of fungal physiology. For example, laetiseric acid ((R)8-hydroxy-*cis*-9*cis*-12-octadecadienoic acid), an isomer of coriolic acid, acts as a premature-sexual-inducer ( $\psi$ ) Ba factor, regulating the sexual development of the ascomycetous fungus *Aspergillus nidulans* (Mazur et al. 1991), while showing antifungal properties against phycomycetous fungi (Bowers et al. 1986). The association of laetiseric acid in regulation of conidia formation and production of fumonisins by *Fusarium verticillioides* was also reported (Scala et al. 2014). Consequently, these diverse roles of HUFAs in fungal physiology and ecology may illustrate their species-specific modes of action.

Our observation from the *in vivo* studies illustrated that coriolic acid exhibited the most control activity against tan spot of wheat and subsequently on net blotch of barley. Both ricinoleic acid and oleic acid did not control the disease severity of tan spot and net blotch on

wheat and barley cultivars, respectively. The protective and curative applications of oleic acid at 0.33 and 1 g/L concentrations also could not control the powdery mildew caused by *Podosphaera xanthii* on cucumber plants, while its methyl and ethyl esters at the same concentrations could effectively control the disease severity (Choi et al. 2010). Methyl- and ethyl-oleate also showed control activity against the powdery mildew of barley at 3 g/L concentration (Choi et al. 2010). However, another *in vivo* study illustrated that oleic acid at 0.5 or 1 g/L concentrations exerted antiviral activity against tobacco mosaic virus in tobacco seedlings, attributed to the activated expression levels of defense-related genes such as PR-1 and PR-5 (Zhao et al. 2017).

The difference in antifungal activity of coriolic acid, ricinoleic acid, and oleic acid against the tested pathogenic fungi of wheat and barley may also be explained by their structural differences, such as the presence and location of hydroxy groups, providing the specific structure-function relationships of UFAs in inhibition of pathogenic fungi (Prost et al. 2005; Pohl et al. 2011; Black et al. 2013; Liang et al. 2017). The 12,13,17-trihydroxy-9-octadecenoic acid did not inhibit the growth of rice blast fungus (*Pyricularia oryzae*) in rice crops (Hou and Forman 2000), while 9,12,13-trihydroxy-10-octadecenoic acid shown to be active against the fungus (Kato et al. 1985), indicating that the position of hydroxy groups plays a key role in growth inhibition of rice blast fungus. Furthermore, the presence and number of carbon-carbon double bonds in long-chain UFAs contributes to the higher levels of antifungal activity of UFAs (Prost et al. 2005; Black et al. 2013). Thus, our results can support the conclusion that HUFAs with more C=C bonds also show higher levels of antifungal activity against pathogenic fungi *in vivo*, although their antifungal activity were observed to be similar in *in vitro* experiments.

Control of tan spot of wheat and net blotch of barley by coriolic acid at 0.5 g/L concentration or higher, may relate to coriolic acid effects on germination of their conidiospores, germ-tubes growth, or development of the infection structure, appressorium. To the best of knowledge, there is no published article showing the effects of coriolic acid on fungal infection development, but it has been previously indicated that coriolic acid can reduce spore germination in *L. maculans* (Garner et al. 2003) and inhibit mycelial growth in *Aspergillus spp*, *Penicillium roqueforti*, *Mucor plumbeus*, *Cladosporium herbarium*, *Botrytis cinerea*, *Phytophthora infestans* and *Fusarium oxysporum* (Prost et al. 2005; Pohl et al. 2011; Black et al. 2013; Liang et al. 2017).

The effects of coriolic acid and ricinoleic acid on disease severity caused by *L. maculans* and *S. sclerotiorum* on canola cultivars were not as promising as the results obtained for monocotyledons crops. None of these fatty acids at 1 g/L or lower concentrations were effective against the pathogenic fungi affecting canola cultivars. Canola cultivars treated with coriolic acid and ricinoleic acid at the highest concentration (2 g/L) showed soaking and wilting symptoms which ultimately caused necrotic leaf tissues and dead seedlings. Although little information about the phytotoxicity of long chain fatty acids has been documented, the phytotoxic activity of stearic acid (C18:0), myristic acid (C14:0), and decanoic acid (C 12:0), extracted from root exudates of barnyard grass (*Echinochloa crusgalli* (L.) Beauv.), at 100 ppm concentrations has been reported on germination and growth rate of alfalfa, lettuce, monochoria, Indian jointvetch, and sesame (Xuan et al. 2006). Foliar treatment of middle-chain fatty acids including caproic acid (C6), caprylic acid (C8), peralgonic acid (C9), capric acid (C10), undecanoic acid (C11), lauric acid (C12), and myristic acid (C14) at 0.1 M concentration also resulted in severe to low levels of damage on crabgrass (*Digitaria ciliaris* Koel.) by degradation of cell membranes and



thylakoid membranes of the treated leaves (Fukuda et al. 2004; Lederer et al. 2004). It was also indicated that the observed variations in phytotoxicity is related to differences in hydrophobicity of the applied fatty acids which increases by increasing in the number of carbon-chain (Fukuda et al. 2004). According to these observations and the fact that amphiphilic monocarboxylic acids can disturb membrane-mediated processes by their partition into the bilayer of cell membranes (Gruber and low 1988), it can be suggested that coriolic acid and ricinoleic acid, as amphiphilic molecules with 18 carbons in their alkyl chains and high levels of hydrophobicity, possibly have the potential to damage the cell and thylakoid membranes of canola cultivars especially when uses at such high concentration (2 g/L). Although this phytotoxicity effect can be specific to some species, as indicated by this study, the monocotyledons crops were affected much lesser compared to dicotyledons ones.

The difference in functional doses of HUFAs to inhibit the disease severity caused by various pathogenic fungi illustrates that the inhibitory effects are not only dose-dependent but also fungus and/or host specific. However, the role of UFAs as signaling molecules to regulate plant defense pathways in the induction of resistance responses to the diseases cannot be ignored. Oleic acid treatment in tobacco leaves induced resistance responses to TMV, resulting in increased activity of defense related enzymes (phenylalanine ammonia lyase) and increased expression levels of defense related genes (PR-1a and PR-5) (Zhao et al. 2017). The exogenous application of UFAs such as arachidonic acid, docosahexaenoic acid, linolenic acid, eicosapentaenoic acid, oleic acid and linoleic acid on potato plants also induced systemic resistance against infection of potato by *Phytophthora infestans* (Cohen et al. 1991). Consequently, it can be speculated that coriolic acid treatments on monocotyledons crops probably triggered the defense mechanisms such as SA-dependent and/or JA-dependent

signaling pathways, resulting in induced resistance against subsequent challenge by the pathogenic fungi. Although further research is needed to understand the biology behind this defense mechanism.

Under various abiotic and biotic stresses, generation of ROS superoxide, hydrogen peroxide, hydroxyl radicals, and peroxy radicals are increased, thereby imposing a secondary oxidative burst in plant cells which can damage cellular structures and macromolecules and ultimately lead to cell death (Bhattacharjee 2005; Sewelam et al. 2016). In the present study to establish a link between formation of necrotic tissues on the canola cv. Westar treated by coriolic acid and ricinoleic acid at 2 and 1 g/L concentrations and enhanced ROS levels, we detected the accumulation of  $\text{H}_2\text{O}_2$  and  $\text{O}_2^-$  in the treated leaves. The histochemical staining of coriolic acid- and ricinoleic acid-treated leaves showed no apparent accumulation of  $\text{O}_2^-$  compared with control leaves at both concentrations; however, increased staining suggested that the accumulation of  $\text{H}_2\text{O}_2$  was probably higher in coriolic acid- and ricinoleic acid-treated leaves at 2 g/L concentration six hours post-treatment. It can be speculated that coriolic acid and ricinoleic acid at a 2 g/L concentration are able to trigger the enhanced production of  $\text{H}_2\text{O}_2$  in foliar tissues of ‘Westar’ as a result of the oxidative burst through degradation of membrane lipids which eventually lead to oxidative damage and plant cell death (Bhattacharjee 2005; Gill 2010; Sharma 2012). This enhancement in accumulation of  $\text{H}_2\text{O}_2$  may relate to changes in the activity of enzymatic antioxidant defense systems developed in plants against oxidative stress damage or non-enzymatic systems (Mittler 2004; Gill 2010), which will require further investigations to reveal. In contrast, increased staining for  $\text{O}_2^-$  observed in oleic acid-treated leaves at 2 and 1 g/L concentrations compared with controls suggests that structural differences of fatty acids may play a role in production of various ROS indicators such as free radicals ( $\text{O}_2^-$ ) or non-radical

molecules (H<sub>2</sub>O<sub>2</sub>). Although there is no literature regarding the exogenous application of coriolic acid, ricinoleic acid, and oleic acid on ROS accumulation in plant cells, several studies on mammalian cells illustrated the activity of long chain fatty acids including oleic acid in stimulation of the oxidative burst and enhancement of ROS levels (Inoguchi et al. 2000; Cury-Boaventura and Curi 2005; John Aitken et al. 2006; Hatanaka et al. 2013).

Long-chain fatty acids are known to be germination-inhibiting compounds and their inhibitory activity increases by increasing the number of carbon atoms present in the alkyl chains (Marambe et al. 1993; Edney and Rizvi 1996). In this study, however, ricinoleic and oleic acid had no effect on seed germination of the tested cultivars compared with controls, supporting the results obtained by Ferrarese et al. (1998) in which different concentrations of oleic acid had no effect on seed germination of *B. napus* L. cv Iciola 41. Likewise, among the long-chain fatty acids in animal-waste compost, saturated fatty acids are more potent germination-inhibitors of sorghum seeds than the UFAs (Marambe et al. 1993). In the case of coriolic acid, we observed a significant ( $p < 0.05$ ) germination-inhibiting effect at 2 g/L concentration for DH12075 but not at other concentrations. Currently, to our knowledge, there are no reports of coriolic acid effects on seed germination rate of monocotyledons and dicotyledons. Therefore, this inhibitory effect may be attributed to the threshold effect of fatty acids on seed germination (Buller et al. 1976) or changes in the rate of physiological processes of treated seeds such as water uptake and  $\alpha$ -amylase activity (Marambe et al. 1992; Marambe et al. 1993). The diversity in germination rate of treated seeds by coriolic acid 2 g/L between crops may also be explained by the species-specific differences in the responses of crop species to the germination-inhibiting compounds (Marambe et al. 1991). The results of seed treatment assays shown the ineffectiveness of coriolic acid, ricinoleic acid, and oleic acid as seed treatment compounds against the tested

pathogenic fungi, which suggests the chemical instability and biodegradability of UFAs because of the high activity of double bonds (Pohl et al. 2011; Liu et al. 2014), as well as intricacy of antifungal functionality of the UFAs against the target fungi.

## **6. CONCLUSION AND FUTURE PROSPECTS**

In conclusion, a comparison amongst the antifungal activity of coriolic acid, ricinoleic acid, and oleic acid provided insights into the structure-function relationships of UFAs as well as their host-pathogen interactions, meaning that the antifungal activity of UFAs is dependent on their chemical structure as well as the type of fungus and its host. Until now, a limited number of fatty acids were developed as commercial crop protective compounds against fungal diseases. As shown by the present study, UFAs such as coriolic acid possess superior activity against pathogenic fungi of monocotyledons both *in vitro* and *in vivo*. This suggests the potential of UFAs as environmentally-friendly crop protective agents for commercial application to control fungal diseases of plants, reduce environmental pollution in agricultural ecosystems and keep human and animal health safe. As the future line of works, evaluating the antifungal activity of coriolic acid against a wide range of fungal pathogens is valuable. Conducting seed treatment experiments to assess the efficacy of coriolic acid against seed-borne and/or soil-borne fungal pathogens affecting the seed or seedling viability is also worthwhile in management of diseases caused by them. Furthermore, it has been demonstrated that the antifungal activity, both *in vitro* and *in vivo*, does not explain what is happening in the plant; thus, the molecular mechanism behind this antifungal activity should be investigated further to understand how these fatty acids can stimulate plant defense signaling pathways. Investigating the metabolic profiling between coriolic acid treated plants and untreated ones is also important for identification of accumulated and/or diminished metabolites inside the plants which are involved in resistance induction. These

results will hopefully provide some useful insights into future control strategies of fungal diseases.

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