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# Impact of Secondary Metabolites and Related Enzymes in Flax Resistance and or Susceptibility to Powdery Mildew

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Abstract: Changes in various physiological defenses including secondary metabolites, proline, total soluble protein and antioxidant enzymes were investigated in leaves and stems of eighteen flax lines either resistant or susceptible to powdery mildew. The total alkaloids content in flax stems significantly increased in resistant lines meanwhile they decreased in susceptible ones and susceptible parent when the both compared with resistant parent. Stems of resistant flax lines contained a higher content of total phenols than susceptible ones upon pathogen challenge. The activity of peroxidase, polyphenyl oxidase, catalase enzymes as well as proline content were significantly increased in powdery mildew infected leaves of flax lines as compared with either resistant or susceptible parents. This increase was much pronounced in tolerant lines than resistant lines. Total soluble protein content in flax leaves increased in resistant lines and decreased in susceptible lines when comparing both with resistant parent.

Key words: Flax · Antioxidant enzymes · Phenol · Proline · Secondary metabolites

## INTRODUCTION

Flax (*Linum usitatissimum* L.) is considered as the major bast fiber crop in Egypt and ranks just after cotton in relation to cultivated area and its economic importance [1]. Flax is cultivated in Egypt for both fiber and seed yields.

Powdery mildew (PM) is foliar disease attacks flax and is caused by Odium lini Skoric. The fungus attacks all aboveground parts of flax cultivars. In Egypt, it occurs wherever flax is cultivated when humidity and temperature conditions are favorable. Early infections may cause severe defoliation of flax plant and reduce yield and quality [1, 2]. The symptoms are characterized by a white powdery mass of mycelia that starts as small spots and rapidly spreads to cover the entire leaf surface [3]. Currently this disease is considered the most common, conspicuous, widespread and easily recognized as foliar disease of flax in Egypt [4]. Over the last decade, the importance of this disease has increased probably due to the appearance and rapid distribution of new races capable to attack the present resistant cultivars [1].

Plants responds to pathogen attack or elicitor treatments by activating a wide variety of protective mechanisms designed to prevent pathogen replication and spreading [5]. The defense mechanisms including production of reactive oxygen species (ROS) [6]; alterations in the cell wall constitution; accumulation of secondary metabolites [7-10]; activation and/or synthesis of defense peptides and proteins [11].

The involvement of phenols as secondary plant metabolites in plant disease resistance is based on to large extent on their cytotoxicty. This is associated with their oxidation products [12]. Phenolics present in healthy, uninfected plant tissues, as preformed antimicrobial compounds, that inhibit the growth of fungi may include simple phenols, phenolic acids, flavonols, some isoflavones and alkaloids [13, 14]. Phenolics seem to disease inhibit development through different mechanisms involving the inhibition of extracellular fungal enzymes (cellulases, pectinases, laccase and xylanase), inhibition of fungal oxidative phosphorylation, nutrient deprivation (metal complexation, protein insolubilisation), the inhibition of both spore germination and mycelial growth of different pathogenic fungi and antioxidant activity in plant tissues [15-17].

Corresponding Author: Heba I. Mohamed, Lecturer of Plant Physiology, Biological and Geological Sciences, Faculty of Education, Ain shams university, El Makres St. Roxy, Cairo, Egypt. (ARE) Post code: 1575, Mob: +0123754036, E-mail: hebaibrahim79@yahoo.com. The production of reactive oxygen species (ROS) is one of the earliest cellular responses following successful pathogen recognition. The generation of ROS such as the superoxide anion ( $O^{2-}$ ) and hydrogen peroxide ( $H_2O_2$ ) are a common event associated with normal plant biochemical processes [18] and also causes oxidative damage through actions such as lipid peroxidation with membrane destruction, protein inactivation and DNA mutation [19].  $H_2O_2$  generation have direct antimicrobial activity inhibiting germination of spores of many fungal pathogens and participation in the formation of phenoxylradicals during phenol-polymerization within the plant cell wall [20].

Various antioxidant enzymes such as peroxidase (POX), superoxide dismutase (SOD), catalase (CAT) and ascorbate peroxidase (APX) participate in ROS metabolism during the pathogen attack. POX may be some of the elements of the defense systems that are stimulated in plants in response to pathogen infection like *Fusarium oxysporum* [21].

Peroxidases (POXs, EC 1.11.1.7; donor: H<sub>2</sub>O<sub>2</sub> oxidoreductase) are oxido-reductive enzymes that participate in the wall-building processes such as oxidation of phenols, suberization and lignifications of host plant cells during the defense reaction against pathogenic agents [22]. One of the important physiological roles of POXs is the synthesis of cell-wall polymers (lignin and suberin), which constitute physical barriers for both biotic and abiotic stresses [23, 24], which might confer the plant with high rigidity. In sorghum, POXs are involved in resistance to fungal infection [25]. Enhanced POX activity has been correlated with resistance in rice [26, 27], wheat [28], barley [29] and sugarcane [30] following the inoculation with phytopathogens.

Polyphenol oxidases (PPOs, EC1.14.18.1; monophenol, 3,4-L-dihydroxyphenylalanine: oxygen oxidoreductase) play an important role in plant defense via the oxidation of endogenous phenolic compounds into o-quinones, which are toxic to invading pathogens and pests [13]. PPO activity in plants increases upon fungal infections [25].

Accumulation of proline occurs in response to many abiotic stresses including drought, salinity and frost as well as biotic stresses such as pathogen infection [31, 32]

The aim of the present study was to determine the changes in secondary metabolites of flax leaves and stems in response to powdery mildew infection.

#### MATERIALS AND METHODS

**Plant Material:** Flax cvs., i.e Wildon (P2) as tolerant parent and Cortland (P3) as sensitive ones were crossed,  $F_1$  was self pollinated to generate  $F_2$  seeds.  $F_2$  individuals were subjected to PM infection; disease incidence and severity were rated as follows:

Disease Incidence; plants showing even a single spot of infection.

Disease severity; Percentage of number of infected leaves out of the total number of leaves on the plant. Individual plants showing disease severity 0-10% were selected as resistant, while plants showing 90-100% disease severity were selected as susceptible. Flowers of selected plants were bagged and left for self pollination; seeds of each individual selected plant were harvested separately and planted at the next generation. The same selection criteria were used for five successive seasons; as a result, several lines with constant response to infection were generated (either resistant or sensitive). F<sub>7</sub> lines were used for this study. Twelve resistant lines and five susceptible ones along with their respective tolerant and susceptible parent were used. The selected lines were grown in a pot trial in three replicates and left out doors during the winter seasons 2008-2009. Natural infection was occurred, samples were obtained at vegetative stage (65 days after germination) and the following parameters were tested.

Flax lines were considered resistant when showing disease severity 0-10%, while lines showing 90-100% disease severity were selected as highly susceptible.

**Determination of Soluble Protein:** Two grams of fresh sample were homogenized in cold phosphate buffer (0.05 M at pH 6.5). The homogenate was centrifuged at 1000 rpm for 10 minutes. The filtrate was completed to a known volume. The total protein content of plant tissue was estimated according to the method of [33] and was measured at 595 nm using spectrophotometer.

**Determination of Antioxidant Enzymes:** Peroxidase (EC 1.11.1.7) and Polyphenol oxidase (EC 1.14.18.1) was assayed following the method described by [34]. The colour intensity was read at 430 nm and the enzyme activity was expressed as the change in the optical density/ gram fresh weight/hour.

Catalase (EC 1.11.1.6) was assayed following the method of [34]. The colour intensity was read at 240 nm. Catalase activity was expressed as imole  $H_2O_2$  destroyed /gram fresh weight/hour.

**Determination of Total Proline:** The total proline of fresh leaves was estimated according to [35]. The absorbance was determined at 520 nm. The concentration of proline was determined as ug g<sup>-1</sup> fresh leaves.

**Determination of Total Phenols:** Levels of soluble phenols in stems of flax lines were determined in accordance with [36]. The absorbance of the developed blue colour was read at 725 nm. Tannic acid was used as standard and the amount of soluble phenols was expressed as mg tannic acid  $g^+$  dry weight.

**Determination of Alkaloids:** Alkaloids were measured according to the method described by [37]. The weight of alkaloids was then expressed as mg g<sup>-1</sup> dry weight.

**Statistical Analysis:** All data were subjected to statistical analysis, means were compared by Duncan's multiple range test using Mstat C commuter package.

## **RESULTS AND DISCUSSION**

**Changes in Secondary Metabolites:** Table (1) showed that the total alkaloids and phenols content in flax stems increased significantly in resistant lines as compared with resistant parent. Also, they decreased significantly in susceptible lines and susceptible parent when both comparing with resistant parent. Plants have the ability to synthesize a large number of aromatic substances, most of which are phenols or their oxygen-substituted derivatives [38]. Most are secondary metabolites, which may be in the form of simple phenols and phenolic acids, flavonoids and alkaloids. These substances serve as plant defense mechanisms against predation by insects, herbivores and microorganisms [39]. Different studies showed that there are often large increases in phenolic synthesis in plants after attack by plant pathogens [40, 41].

In resistant plants, phenolic based defense responses are characterized by the early and rapid accumulation of phenolics at the infection site resulting in the effective isolation of the pathogen [42-44]. Results of many previous studies suggested that esterification of phenols to cell wall materials and the accumulation and deposition of phenols in and on cell walls are usually considered as an increase in resistance to fungal hydrolytic enzymes as well as a physical barrier against fungal penetration [45].

**Changes in Proline and Total Soluble Proteincontent:** Table (1) showed that proline markedly increased in resistant and susceptible lines as compared with their respective parents. This increase was much pronounced in tolerant than resistant lines. These results are in agreement with the records [46] that proline concentration peaked in frost- and snow mould resistant genotype 561 and were 25 times higher in comparison with the control

| Lines        | Disease reaction | Total phenol (mg tannic acid/g) | Alkaloids (mg/g)   | Total soluble protein (mg/ml) | Proline(µg / g fresh weight) |
|--------------|------------------|---------------------------------|--------------------|-------------------------------|------------------------------|
| 1            | R Parent         | 3.20 <sup>d-h</sup>             | 14.67 efg          | 39.11 <sup>d-g</sup>          | 1.94 <sup>fgh</sup>          |
| 2            | R                | 3.50 <sup>d-g</sup>             | 29.67 ª            | 42.33 bcd                     | 2.70 <sup>e</sup>            |
| 3            | R                | 3.36 <sup>d-h</sup>             | 21.00 bcd          | 45.11 <sup>b</sup>            | 5.67 <sup>b</sup>            |
| 4            | R                | 5.41 <sup>ab</sup>              | 21.67 bcd          | 44.33 <sup>b</sup>            | 6.46 <sup>a</sup>            |
| 5            | R                | 4.12 <sup>cde</sup>             | 18.00 cde          | 50.08 <sup>a</sup>            | 3.54 <sup>d</sup>            |
| 6            | R                | 4.93 <sup>abc</sup>             | 17.33 def          | 41.11 <sup>b-e</sup>          | 5.32 bc                      |
| 7            | R                | 4.32 bcd                        | 20.33 bcd          | 44.04 bc                      | 2.21 efg                     |
| 8            | R                | 3.53 <sup>d-g</sup>             | 22.33 <sup>b</sup> | 44.14 <sup>b</sup>            | 2.19 fg                      |
| 9            | R                | 5.53 <sup>f-i</sup>             | 22.33 <sup>b</sup> | 42.11 bcd                     | 2.44 ef                      |
| 10           | R                | 6.03 <sup>a</sup>               | 18.33 cde          | 39.88 <sup>c-f</sup>          | 2.20 efg                     |
| 11           | R                | 4.34 <sup>bcd</sup>             | 17.00 fgh          | 44.62 <sup>B</sup>            | 3.54 <sup>d</sup>            |
| 12           | R                | 3.66 def                        | 19.67 bcd          | 42.27 <sup>bcd</sup>          | 4.85 °                       |
| 13           | H.S Parent       | 2.88 <sup>e-i</sup>             | 13.67 fgh          | 21.10 <sup> i</sup>           | 1.47 <sup>h</sup>            |
| 14           | S                | 1.81 <sup>ij</sup>              | 11.00 ghi          | 35.40 <sup>gh</sup>           | 1.72 <sup>gh</sup>           |
| 15           | S                | 1.75 <sup>hij</sup>             | 13.67 fgh          | 35.82 <sup>fgh</sup>          | 1.46 <sup>h</sup>            |
| 16           | S                | 2.37 <sup>g-j</sup>             | 9.67 <sup> i</sup> | 36.78 fg                      | 2.28 ef                      |
| 17           | S                | 1.41 <sup>j</sup>               | 10.00 hi           | 37.72 <sup>efg</sup>          | 2.26 ef                      |
| 18           | S                | 2.59 <sup>f-j</sup>             | 10.00 hi           | 32.01 <sup>h</sup>            | 2.15 fg                      |
| L. S.D at 5% |                  | 1.26                            | 3.76               | 4.2030                        | 0.5087                       |

Different letters indicate a significant difference according to Duncan's multiple range test.

R means resistant

S means susceptible

| Lines        | Disease reaction | Polypheny oxidase activity/g fresh weight/h) | Peroxidase (activity/g fresh weight/h) | Catalase(µM H2O2/ g fresh weight/h) |
|--------------|------------------|--|--|-------------------------------------|
| 1            | R Parent         | 15.26 <sup>d</sup>                           | 9.14 <sup>fg</sup>                     | 26.20 <sup> i</sup>                 |
| 2            | R                | 18.11 <sup>g</sup>                           | 10.16 fg                               | 27.56 <sup>g</sup>                  |
| 3            | R                | 19.01 <sup>hij</sup>                         | 22.08 bcd                              | 28.55 def                           |
| 4            | R                | 15.78 <sup>ij</sup>                          | 25.24 ª                                | 28.68 <sup>cdef</sup>               |
| 5            | R                | 18.03 <sup>f</sup>                           | 21.44 <sup>cd</sup>                    | 27.48 <sup>g</sup>                  |
| 5            | R                | 18.70 <sup>g</sup>                           | 20.16 <sup>d</sup>                     | 28.88 <sup>cde</sup>                |
| 7            | R                | 19.04 <sup>g</sup>                           | 11.63 <sup>fg</sup>                    | 30.96 ª                             |
| 3            | R                | 17.05 °                                      | 23.00 abc                              | 29.56 <sup>ь</sup>                  |
| )            | R                | 19.76 <sup>k</sup>                           | 24.56 <sup>ab</sup>                    | 27.28 <sup>g</sup>                  |
| 0            | R                | 16.63 <sup>hi</sup>                          | 20.84 <sup>cd</sup>                    | 27.36 <sup>g</sup>                  |
| 11           | R                | 18.54 j                                      | 9.82 <sup>fg</sup>                     | 27.39 <sup>g</sup>                  |
| 2            | R                | 21.04 <sup>b</sup>                           | 15.66 °                                | 29.03 bcd                           |
| 3            | H.S Parent       | 12.47 °                                      | 4.61 <sup>i</sup>                      | 25.48 <sup>j</sup>                  |
| 4            | S                | 12.91 <sup>j</sup>                           | 4.62 <sup>i</sup>                      | 26.71 <sup>hi</sup>                 |
| 5            | S                | 12.03 <sup>a</sup>                           | 5.86 <sup>hi</sup>                     | 27.08 <sup>gh</sup>                 |
| 6            | S                | 13.94 °                                      | 6.05 <sup>hi</sup>                     | 28.19 <sup>f</sup>                  |
| 17           | S                | 14.70 <sup>g</sup>                           | 8.03 <sup>gh</sup>                     | 28.44 <sup>ef</sup>                 |
| 8            | S                | 13.34 <sup>hi</sup>                          | 6.51 <sup>hi</sup>                     | 29.23 <sup>bc</sup>                 |
| L. S.D at 5% |                  | 0.091  | 2.16                                   | 0.550                               |

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Different letters indicate a significant difference according to Duncan's multiple range test.

S means susceptible

plants and 5 times higher than that of the susceptible genotype 621. Also, proline index the indicator of predisposition in tomato plants to *Phytophtora nicotianae* infection as influenced by abiotic stresses [47]. In addition proline concentration in foliar tissues of pepper plants inoculated with *Verticillium dahliae* increased significantly between days 21 and 28 after inoculation, while they did not change in the leaves of control plants [48]. In cotton, higher proline levels were detected in leaves from inoculated plants with *Verticillium* than in those from controls [49].

The amino acid proline may act as a potent scavenger of ROS and this property of proline might prevent the induction of programmed cell death by ROS [50]. Proline may also function as a protein compatible hydrotrope [51] and as a hydroxyl radical scavenger [52]. In any case the higher proline accumulation in diseased tissue might be related to pathological disorder [53].

Data presented in Table (1) also showed that, total soluble protein content in flax leaves increased significantly in resistant lines as compared with resistant parent. In this regard, an opposite feature was observed in susceptible lines comparing with resistant parent. Plant pathogens such as bacteria and fungi elicit the synthesis of host proteins which help in restricting the multiplication and spread of pathogens in the healthy tissue [54]. Also, [55] showed that the presence of pathogenesis-related (PR) proteins, such as chitinase,  $\beta$ -1,3-glucanase or thaumatin in the tissue infected by pathogens positively correlates with the plant resistance to micro-organisms. In addition to, in cucumber, the resistance was associated with the induction of pathogenesis-related proteins (i.e. chitinases) [56] and POX [57] in extracellular spaces of the host tissues following treatment.

**Changes in Antioxidant Enzymes:** It is well known that organisms protect themselves against oxidatives stress by the synthesis of various antioxidant enzymes. Results in Table (2) showed that infection with powdery mildew significantly increased peroxidase, polyphenyl oxidase and catalase activities in leaves of flax lines as compared with resistant and susceptible parents.

POX activity was noted to increase significantly in both resistant and susceptible lines as compared with their parent. These results are in agreement with other studies [58], which reported that increase in peroxidase activity enhance lignifications in response to chocolate spot infection which may restrict the fungal penetration. These findings indicate to a positive relationship between resistance and peroxidase activity. Peroxidase also produces free radicals and hydrogen peroxide which are toxic to many microorganisms [59]. Another supportive suggestion was brought by [13] who found that, a significant increase in POX specific activity in both resistant (Wang-shuibai) and susceptible (Falat) wheat

R means resistant

cultivars following the inoculation with *F. graminearum* conidia. Also, an increase in peroxidase activity is considered as a preliminary indicator for resistance of broad beans to chocolate spot disease [60]. These compounds act as barriers against pathogen invasion.

POXs are usually associated with induced resistance [61] and they are also implicated in several plant defense mechanisms such as lignin synthesis, oxidative cross linking of different plant cell wall components or generation of reactive oxygen species [62].

Data also showed that PPO markedly increased in resistant and susceptible lines as compared with resistant and susceptible parents. While in susceptible lines, PPO significantly decreased as compared to resistant parent. These results are in agreement with findings that PPO may confer resistance to some fungal species such as R. sorghicola [25]. PPO specific activity significantly increased in wheat heads of resistant and susceptible cultivars following the inoculation with F. graminearum conidia [13]. Similar results have been obtained in plant-pathogenic fungal interactions such as cabbage/F. oxvsporum, onion/Botrytis, sunflower/Sclerotinia sclerotiorum, soybean/Phytophthora megasperma and bean /Rhizoctonia [63]. In addition, pretreatment of Chick pea with rhizobium increased significantly the levels of peroxidase and polyphenyl oxidase and total phenolics was recorded [64]. These increases were the most effective reducing Fusarium wilt development. POX and PPO are important in the defense mechanism against pathogens, through its role in the oxidation of phenolic compounds to quinines, causing increasing in antimicrobial activity. Therefore, its may be directly involved in stopping pathogen development [65, 66]; accelerating the cellular death of cells close to the infection site, preventing the advance of infection and/or by generating a toxic environment which will inhibit the growth of the pathogen inside the cells [67].

Catalase activity was increased markedly in flax leaves of resistant and susceptible lines (Table 2). This increase was much pronounced in tolerant lines than resistant lines. This increase in catalase activity may provide its protection from oxidative damage by rapid removal of  $H_2O_2$ . These results were in agreement with those recorded that, the activity of antioxidant enzymes in leaves under *Fusarium oxysporum* infection increased and might be affective in scavenging mechanism to remove  $H_2O_2$  and  $O_2^-$  produced in leaves [68]. Also, in banana plants, it was reported that, levels of  $H_2O_2$  and other enzyme activities were increased to the levels of tolerance or susceptibility to *Fusarium* wilt disease [69]. The obtained results in the present study lead to conclude that the total alkaloids, phenols and total soluble protein content in flax stems significantly increased in resistant lines but decreased in susceptible ones and susceptible parent when the both compared with resistant parent. The activity of peroxidase, polyphenyl oxidase, catalase enzymes as well as proline content were significantly increased in powdery mildew infected leaves of flax lines as compared with either resistant or susceptible parents.

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