IMPACT OF DIFFERENT CADMIUM CONCENTRATIONS ON TWO PISUM SATIVUM L. GENOTYPES

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Abstract

This study was conducted to evaluate the effect of different concentrations of cadmium (Cd) (0, 100, 200, and 400 µM Cd) on two genotypes (AG-10 and AP-3) of *Pisum sativum*. Cd stress reduced the length, fresh weight, and dry weight of roots and shoots in both genotypes. Chlorophyll 'a' and 'b' and carotenoid content was also decreased and greater decreases were observed for higher concentrations of Cd. As Cd stress increased, levels of proline, glycine betaine, and soluble proteins were increased and sugar content was decreased. The application of Cd also caused an increase in hydrogen peroxide, lipid peroxidation, and electrolyte leakage in both genotypes. The activities of defensive antioxidant enzymes like superoxide dismutase, catalase, ascorbate peroxidase, glutathione reductase, and glutathione *S*-transferase were also increased with increasing Cd concentrations; however, the increase in AG-10 was greater than in AP-3. The accumulation of phenols was decreased with Cd stress in both pea genotypes, with the greatest decreases were observed in AP-3. In conclusion, Cd caused a marked reduction in growth, biomass yield, and pigment content in both pea genotypes. However, enhanced accumulation of osmolytes and antioxidant enzyme activities alleviated the adverse effects of Cd. AG-10 was more tolerant genotype than AP-3, as AG-10 showed less Cd induced damage and higher antioxidant enzyme activities.

Key words: Cadmium; Pisum sativum; Growth; Lipid peroxidation; Osmolytes; Antioxidants.

Introduction

Soil is polluted with different kinds of heavy metal elements, one of which is cadmium (Cd). Non-ferrous metal industries, fossil fuel combustion, and waste incineration primarily contribute to soil pollution (Ahmad et al., 2016). Cd is highly toxic and has detrimental effects on plant and animal life (Ahmad et al., 2011; Ahmad, 2012). After entering the human body through the food chain, Cd causes various health issues such as kidney problems, pulmonary dysfunction, neurotoxicity, and endocrine disruption (Godt et al., 2006). Plant roots easily absorb and transport Cd to other parts of the plant, causing toxicity there (Talukdar, 2012). Cd primarily affects plants by inhibiting seed germination, reducing plant growth (Siddiqui et al., 2012), reducing chlorophyll synthesis (Mobin & Khan, 2007), changing Calvin cycle enzyme activities (Mobin & Khan, 2007; Shamsi et al., 2008), and closing stomata (Perfus-Barbeoch et al., 2002). Cd also affects the synthesis of carbohydrates, proline, and protein (Siddiqui et al., 2012; El-Beltagi & Mohamed, 2013; Mondal et al., 2013) and the uptake of mineral elements (Sandalio et al., 2001; Siddiqui et al., 2012). In addition, excessive Cd exposure leads to the generation of reactive oxygen species (ROS), causing oxidative stress in plants (Ahmad et al., 2010; Ahmad et al., 2015). ROS are very reactive and have the capacity to oxidize organic molecules such as lipids, proteins, and nucleic acids (Ahmad et al., 2010; Ahmad et al., 2015). However, plants are equipped with antioxidants, which act as a defense mechanism against ROS (Ahmad et al., 2010; Ahmad et al., 2015). Plants contain both enzymatic and non-enzymatic antioxidants. Enzymatic antioxidants comprise superoxide dismutase (SOD), catalase (CAT), ascorbate peroxidase (APX), glutathione reductase (GR),

glutathione S-transferase (GST), and peroxidase (POD), whereas non-enzymatic antioxidants include ascorbic acid (AsA) and glutathione (GSH) (Ahmad *et al.*, 2010; Ahmad *et al.*, 2015).

Legumes are susceptible to Cd stress, which leads to global reductions in plant biomass and crop production (Shamsi *et al.*, 2008). Pea (*Pisum sativum*) is an important leguminous crop, serving as the main protein source for the growing population of humans worldwide. Thus, this study was conducted to determine which variety of pea is most tolerant to Cd pollution in soil.

Materials and Methods

Pisum sativum seeds were sterilized (using NaOCl 5%), placed in Petri dishes lined with blotting paper for germination, and provided with distilled water. Healthy germinated seedlings were transferred to pots filled with reconstituted sand that was supplemented with vermicompost. Seedlings were thinned to 10 per pot and grown for another 10 days. Seedlings were irrigated with full-strength (200 mL) Hoagland's nutrient solution every other day. After 10 days, a modified Hoagland solution supplemented with different concentrations of $CdCl_2$ (0, 100, 200 and 400 μ M) was provided for another 20 days. Pots were maintained in a greenhouse with the following conditions: day/night temperatures of 26°C/16°C, relative humidity of 70%-75%, and photoperiod of 18 h light/6 h night. Plants were harvested at 30 days of age for further analysis.

Plant length and dry weight: A manual scale was used to measure length and plant tissue was oven-dried overnight at 70°C to determine dry weight.

Photosynthetic pigments: To extract photosynthetic pigments, fresh leaf tissues were macerated in 80% acetone. Homogenates were centrifuged at 3000xg for 20 min and absorbance was measured at 480, 645, and 663 nm using a spectrophotometer (Beckman 640 D, USA) (Arnon, 1949).

Proline, glycine betaine, protein, and sugar content: Proline was measured using the method described by Bates *et al.*, (1973). To extract proline, leaf tissue was first homogenized in sulfosalicylic acid. The resulting supernatant was mixed with ninhydrin acid and incubated in boiling water for one hour. After cooling on ice, proline was separated using toluene and optical density was recorded at 520 nm using a spectrophotometer (Beckman 640 D, USA).

Glycine betaine was quantified using the method described by Grieve & Grattan (1983). Dry samples were extracted in distilled water. Extracted samples were mixed with a KI–I₂ (potassium iodide and iodine solution) reagent and the periodide crystals that formed were measured at 365 nm using a spectrophotometer (Beckman 640 D, USA). A graded concentration of glycine betaine was used for the standard curve.

Protein was estimated using the method described by Bradford (1976) and absorbance was recorded at 595 nm. Bovine serum albumin was used as the standard. Sugar was measured using the method described by Dey (1990) and absorbance was recorded at 485 nm using a spectrophotometer (Beckman 640 D, USA).

Determination of hydrogen peroxide (H₂O₂), lipid peroxidation (MDA), and electrolyte leakage: Electrolyte leakage was determined by incubating fresh leaf tissue at different temperatures and measuring electrical conductivities (Dionisio-Sese & Tobita, 1998). Leakage was calculated using the following formula: Electrolyte leakage (%) = (EC1-EC0)/(EC2-EC0) × 100

For lipid peroxidation, fresh tissue was macerated in trichloroacetic acid. Malondialdehyde content was measured after a known volume of supernatant was reacted with thiobarbituric acid (Madhava Rao & Sresty, 2000).

To determine hydrogen peroxide (H_2O_2) content, fresh tissue was extracted in trichloroacetic acid and optical density was measured at 390 nm (Velikova *et al.*, 2000).

Assay of antioxidant enzyme activities: Antioxidant enzymes were extracted by homogenizing 1 g fresh tissue in ice-cold 100 mm potassium phosphate buffer (pH 7.0) supplemented with Polyvinylpyrrolidone (PVP, 1%). Samples were centrifuged at 12000xg for 15 min at 4°C and the resulting supernatants were used for enzyme assays.

To determine superoxide dismutase activity, inhibition of the photochemical reduction of nitroblue tetrazolium (NBT) was measured at 560 nm. The assay mixture was exposed to fluorescent light for 15 min and read against unexposed samples (Dhindsa & Matowe, 1981).

The catalase assay was conducted by monitoring the decrease in optical density at 240 nm. The assay mixture contained a phosphate buffer (pH 6.0), Ethylenediamine tetraacetic acid (EDTA), and enzyme. H_2O_2 was used to initiate the reaction (Aebi, 1984).

Ascorbate peroxidase activity was determined in a reaction mixture containing a phosphate buffer (pH 7.5), EDTA, ascorbate, enzyme, and H_2O_2 , and changes in absorbance were recorded at 290 nm (Nakano & Asada, 1981).

To determine glutathione reductase activity, the assay mixture consisting of phosphate buffer (pH 7.0), EDTA, oxidized glutathione, nicotinamide adenine dinucleotide phosphate (NADPH), and enzyme was observed for 3 min to measure changes in optical densities at 340 nm (Foster & Hess, 1980).

The activity of glutathione S-transferase was measured using the method described by Hasanuzzaman & Fujita (2013). Changes in optical densities were recorded at 340 nm using a spectrophotometer (Beckman 640 D, USA).

Determination of total phenolics: Phenols were extracted with ethanol and measured after reacting with a Folin–Ciocalteu reagent (Chun *et al.*, 2003). The Gallic acid standard was used for calculations.

Statistical analysis

Data were analyzed using analysis of variance. Duncan's multiple range test was used to determine the least significant difference (LSD) for a completely randomized design at 0.05% significance. Values represent the mean (\pm SE) of three replicates.

Results and Discussion

Growth and Biomass Yield: In this study, Cd treatment resulted in reduced growth of *Pisum sativum* cultivars. Lengths, fresh weights, and dry weights of roots and shoots were lower when Cd treatment was applied (Fig. 1A-F). "At 400 μ M, the length, fresh weight, and dry weight of cultivar AG-10 was 13.48, 37.11, 30.94% (shoot) and 12.28, 35.79, 9.7% (root) lower than the control, respectively. The length, fresh weight, and dry weight of cultivar AP-3 was 31.43, 37.11, 43.33% (shoot) and 20.53, 62.03, 43.37% (root) lower than the control, respectively."

Cd alters growth, activates leaf necrosis, and disrupts cell division and elongation (Liu et al., 2004). Cd stress has been shown to reduce growth in Lycopersicon esculentum (Jing et al., 2005; Ahmad et al., 2018), Brassica juncea (Ahmad et al., 2011; Per et al., 2016a, 2016b), Hordeum vulgare (Juknys et al., 2012) and Festuca arundinacea (Huang et al., 2017). Differences were observed between cultivars, as AG-10 exhibited a greater tolerance to increasing Cd concentrations. Cd can potentially disrupt the activities of key metabolic enzymes that control growth (Amaya-Carpio et al., 2009). In addition, growth reductions are a result of the irreparable effect of Cd on membrane leakage and proton pump activity (Karcz & Kurtyka, 2007). Cd stress hinders the hydraulic conductivity and extensibility of the cell wall, which damages the morphology of the plant (Ehlert et al., 2009).



Fig. 1. Effect of different concentrations of Cd on (A) shoot length, (B) root length, (C) shoot FW, (D) root FW, (E) shoot DW, and (F) root DW in two genotypes of *Pisum sativum* L. Data presented are the means \pm SE (n = 5). Different letters indicate significant difference at $p \le 0.05$.



Fig. 2. Effect of Cd toxicity on (A) chlorophyll 'a', (B) chlorophyll 'b', and (C) carotenoid content in two genotypes of *Pisum sativum* L. Data presented are the means \pm SE (n = 5). Different letters indicate significant difference at $p \le 0.05$.

Pigment content: Treatment of *Pisum sativum* cultivars with Cd led to a considerable reduction in chlorophyll synthesis, with greater reductions was observed when a high level of Cd (400 μ M) was applied. The observed reductions in Chl a, Chl b, and carotenoids were 36.08, 24.69, 12.29% (AG-10) and 38.61, 21.18, 19.15% (AP-3), respectively, at 400 M μ Cd (Fig. 2A-C).

Similarly, declines in pigment synthesis due to Cd application have been previously reported for *Helianthus annus* (Rivelli *et al.*, 2012), *Abelmoschus esculentus* (Mangal *et al.*, 2013) and *Gossypium hirsutum* (Liu *et al.*, 2014). Heavy metals (including Cd) have the potential to impede the biosynthesis of chlorophyll pigments (Hsu & Kao, 2003), potentially due to the enhanced activity of chlorophyll-degrading enzymes like chlorophyllase (Singh & Jain, 1981). Ahmad *et al.*, (2011) and Khan *et al.*, (2015) also observed chlorophyll degradation in mustard and wheat respectively due to Cd stress.

Osmolytes

The effect of Cd on the accumulation of osmolytes, including proline, glycine betaine (GB), total protein, and soluble sugar, is shown in Fig. 3A-D. Significant increases in proline, GB, and protein accumulation were observed for both cultivars when higher Cd concentrations were applied. Relative to the controls, proline and GB increased up to 75.47% and 85.98%, respectively, in AG-10 and 68.09% and 77.18%, in AP-3 at 400 µM Cd stress (Fig. 3A-D). However, soluble protein content increased at 200 µM Cd stress by 47.11% and 2.48% in AG-10 and AP-3, respectively, but decreased at 400 µM Cd stress. Sugar content declined at all Cd concentrations, with the greatest declines observed at 400 µM Cd (AG-10: 33.84%, AP-3: 46.53%).

Accumulation of proline, GB, and sugars have been implicated in the maintenance of water balance and thus, cell osmolarity (Abdul Jaleel et al., 2007; Ahanger et al., 2014). Environmental stress conditions enhance the activity of proline-synthesizing enzymes and down-regulate its catabolism (Ahmad et al., 2010; Ahanger et al., 2014), resulting in proline accumulation. Similar to the results of this study, increased accumulation of proline in Cd-fed seedlings has been shown in Helianthus annuus (Zengin & Munzuroglu, 2006), Lycopersicon esculentum (Ahmad et al., 2018), and Brassica juncea (Irfan et al., 2014). Although the increased accumulation of osmolytes, including proline, GB, and sugars, does not interfere with metabolic pathways of cells, these osmolytes do replace water in key metabolic processes (Zhifang & Loescher, 2003). Accumulation of proline, GB, and sugars helps prevent stress-induced oxidative damage to plants by maintaining enzyme activity, membrane functionality, and by eliminating ROS (Ahanger & Agarwal, 2017). GB, which is exclusively found in chloroplasts, protects the thylakoid membranes, Rubisco activity, oxygen evolving complex, and, thus, the photosynthetic efficiency (Yokoi et al., 2002). GB accumulation stabilizes extrinsic PSII complex proteins at non-physiological conditions (Papageorgiou & Murata, 1995; Chen & Murata, 2011). Soluble sugars (such as disaccharides, oligosaccharides in the raffinose family, and fructans) and the enzymes involved in their regulation are believed to have a strong correlation with stress-induced ROS accumulation in plants (Keunen et al., 2013). Ahanger & Agarwal (2017) observed higher stress tolerance in Triticum aestivum L. as a result of sugar accumulation. Accumulated sugars regulate the interplay between ROS and other tolerance strategies and enhance growth and crop production (Sami et al., 2016).



Fig. 3. Effect of different concentrations of Cd on (A) proline, (B) glycine betaine, (C) total protein, and (D) soluble sugar content in two genotypes of *Pisum sativum* L. Data presented are the means \pm SE (n = 5). Different letters indicate significant difference at $p \le 0.05$.

H2O2, MDA, and EL: H2O2, MDA, and EL increased by 40.63%, 43.70%, and 27.92% (AG-10) and 39.30%, 42.14%, and 33.46% (AP-3), respectively, relative to control (Fig. 4A-C). AG-10 had lower reductions in H₂O₂, MDA, and EL compared to AP-3, indicating that AG-10 has a higher tolerance to Cd. Greater H₂O₂, MDA, and electrolyte leakage has been observed for Cd-stressed tomato (Ouariti et al., 1997), Chrysanthemum morifolium (Hossain et al., 2006), and mustard (Ahmad et al., 2011). Lipid membranes are sensitive to excess ROS and often trigger oxidation, causing the generation of peroxide radicals through a chain of reactions (Ahanger & Agarwal, 2017; Ahmad et al., 2018). This results in the loss of structural and functional integrity and leakage of essential elements and other cellular constituents (Djebali et al., 2005). Mahmood et al., (2007) reported reduced membrane stability leading to the leakage of essential ions in wheat, barley, and rice due to heavy metal stress. Heavy metal stress enhances lypoxygenase activity and production of peroxides and hydroxyl radicals, thereby deteriorating the structural and functional integrity of membranes and other cellular components (Macrí et al., 1994; Djebali et al., 2005). Increased generation of ROS are responsible for oxidative stress and growth reduction (Kuo & Kao, 2004).

Antioxidants: The activity of antioxidant enzymes SOD, CAT, APX, GR and GST was increased significantly at all levels of Cd stress. Relative to the control, SOD, CAT,

APX, GR and GST activities increased by 12.9%, 15.52%, 16.70%, 20.29% and 8.82% (AG-10) and by 13.83%, 14.95%, 15.94%, 16.95% and 6.25% (AP-3), respectively, at 100 μ M Cd (Fig. 5A-E). At a higher concentration of Cd (400 μ M), antioxidant activity was increased in both genotypes; however, the greatest increase was observed in AG-10.

Within the antioxidative protective system, SOD is a key component that actively eliminates toxic superoxide radicals, leading to greater protection of cellular structures and processes such as the electron transport chain. Our results support the findings of Milone et al., (2003), Shan et al., (2012), Irfan et al., (2014), Per et al., (2016b), and Ahmad et al., (2018). Greater SOD activity brings modulation in Haber-Weiss reaction substrates. superoxides, and H₂O₂, leading to reductions in the generation of toxic hydroxyl (OH-) radicals. SOD generated H₂O₂ is scavenged either by CAT or APX and GR via the ascorbate-glutathione cycle, where AsA and GSH are the key catalytic components (Ahmad et al., 2010; Ahanger et al., 2017). Greater activity of CAT, GR, and GST mediates the efficient removal of toxic H₂O₂ to maintain the functional and structural integrity of cells, ultimately leading to improved stress adaptation (Ahanger & Agarwal, 2017). John et al., (2009) also showed a gradual increase in the activities of antioxidant enzymes in mustard with increasing Cd concentrations in soil.



Fig. 4. Enhanced levels of (A) hydrogen peroxide, (B) malondialdehyde, and (C) electrolyte leakage in two genotypes of *Pisum sativum* L. subjected to different concentrations of Cd. Data presented are the means \pm SE (n = 5). Different letters indicate significant difference at $p \le 0.05$.

Phenols: Cultivar AP-3 showed greater declines in the accumulation of phenols with increasing Cd concentrations (Fig. 5F). Phenol content was decreased by 22.91% (AG-10) and 39.14% (AP-3) at 400 μ M Cd relative to the controls. This also indicates that AG-10 is more tolerant to Cd than AG-3.

Higher synthesis of phenols and phenol derivatives has been shown to confer stress tolerance in crop plants (Tomar & Agarwal, 2013; Ahanger & Agarwal, 2017). Plant species maintaining higher phenol content are known to have higher peroxidase enzyme activities, leading to better growth (Zupan et al., 2014). Polyphenols have a strong capacity to donate hydrogen ions, and can, therefore, be among the best antioxidant molecules to eliminate ROS (Hernández et al., 2009). Multiple studies have shown that increased induction of several genes under stress conditions (e.g., drought, metal toxicity, nutrient deprivation, and wounding) causes increase in phenol levels (Dixon & Paiva, 1995; Winkel-Shirley, 2002; Ahanger & Agarwal, 2017). Griesser et al., (2015) demonstrated that the high photosynthetic efficiency of Vitis vinifera was due to the enhanced accumulation of wide variety of polyphenolic compounds, which led to the maintenance of a higher water potential.

Conclusions

The results from this study showed that Cd toxicity decreased plant growth and photosynthetic activity through decreased chlorophyll content in two pea genotypes. Cd leads to increase in ROS, which are dangerous for biomolecules. However, osmolytes (such as proline, GB, and protein) help pea plants to withstand Cd stress. Antioxidant enzyme activities also increased in order to maintain the redox balance under Cd stress. The genotype AG-10 showed smaller reductions in growth, biomass yield, and pigment content under Cd stress than AP-3. AG-10 also showed higher accumulations of proline and GB and enhanced antioxidant enzyme activities than AP-3, indicating that the AG-10 genotype is more tolerant to Cd than AP-3.

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Fig. 5. Effect of different concentrations of Cd on (A) superoxide dismutase, (B) catalase, (C) ascorbate peroxidase, (D) glutathione reductase, (E) glutathione S-transferase, and (F) phenol content in two genotypes of *Pisum sativum* L. Data presented are the means \pm SE (n = 5). Different letters indicate significant difference at $p \le 0.05$.

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