

Cadmium Affects the Glutathione/Glutaredoxin System in Germinating Pea Seeds

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Abstract The aim of this work was to investigate the effects of cadmium (Cd) on thiol and especially glutathione (GSH)-dependent reactions (glutathione content, glutaredoxin (Grx) content and activity, “glutathione” peroxidase (Gpx) activity, and glutathione reductase (GR) activity) in germinating pea seeds. Under Cd stress conditions, the overall activity as well as more specifically the expression of Grx C4 and Grx S12 increased. On the contrary, when incubated with Cd ions in vitro, the disulfide reductase activity of both isoforms was drastically inhibited. In the case of Grx C4, this correlated with the formation of protein dimers of 28 kDa as evidenced by electrophoresis analysis. Oxidative stress also affected the GSH status, since Cd treatment provoked (1) a pronounced stimulation in Gpx (a thioredoxin-dependent enzyme in plants) expression and (2) a drastic decrease in GR activity. These results are discussed in relation with the known contribution of Grx system to the thiol status during the germination of Cd-poisoned pea seeds.

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Abbreviations

Cd	Cadmium
DTT	Dithiothreitol
DW	Dry weight
FW	Fresh weight
GPX	Glutathione peroxidase
GR	Glutathione reductase
Grx	Glutaredoxin
GSH	Glutathione
HED	Hydroxyethyl disulfide
β -MET	β -Mercaptoethanol

Introduction

Seeds are easy to grow and adapt to environmental stress [1], while the germination process is very sensitive to adverse situations, in particular heavy metal pollution [2]. Cadmium (Cd) is frequently accumulated by agriculturally important crops with a significant potential to impair animal and human health [1]. The inhibitory effects of Cd on various processes in adult pea and germinating seeds have been documented by several workers [3–8]. Bioassays reflect a toxicological damage at both the biochemical and physiological levels.

Proteins are the primary effector molecules of all living systems, and therefore, virtually, any response to environmental changes, physiological, or pathological conditions will be reflected by alterations in protein activity, location, and concentration [9]. Moreover, under heavy metal stress, the cellular thiol-disulfide redox status is important for the control of protein functions [10]. Glutaredoxin/glutathione/glutathione reductase (Grx/GSH/GR) is one of the major redox systems involved in maintaining the thiol reduced state [11–16]. This reducing system is predicted to be localized in multiple subcellular compartments as there are overall 31 Grx genes in *Arabidopsis thaliana* [14, 17]. There are 14 bicysteinic Grxs (nine in the cytosol, two in chloroplasts, and three secreted) and 17 monocysteinic (12 in cytosol, four in chloroplasts, and one in mitochondria). They represent a large family of GSH-dependent oxidoreductases that catalyze the GSH-dependent reduction of disulfides or glutathione-mixed disulfide by a dithiol or a monothiol mechanism through a CxxC or CxxS active site [14, 15, 17]. Using NADPH as an electron donor, GR catalyzes the conversion of oxidized glutathione to its reduced form which functions to reduce cellular disulfide bonds, often in conjunction with Grxs. Then, they are required for a number of antioxidant and metabolic enzymes that form a disulfide as part of their catalytic cycle [15].

The most documented functions of Grxs are their involvement in the oxidative stress response [15, 16, 18], notably in plants subjected to abiotic stress, as Cd [9, 19]. The goal of this study was to evaluate the effect of Cd addition into imbibition medium of pea seeds on Grx/GSH/GR system in post-mitochondrial fractions obtained from the reserve tissues (cotyledons) and growing embryonic axes. We show here that Cd influences the content and activity of several partners of this redox regulatory network in seeds.

Materials and Methods

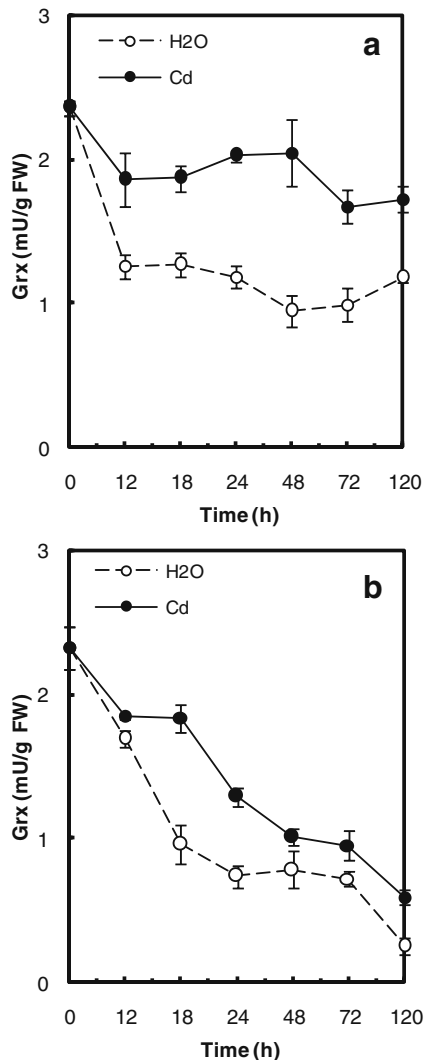
Germination and Cadmium Treatment

The germination conditions and Cd treatment of pea (*Pisum sativum* L. cv. douce province) seeds were as previously described [8].

Protein Extraction and Solubilization

Fresh tissues were ground with a mortar and pestle in a homogenization medium (pH 8.0) consisting of 50 mmol L⁻¹ Tris-HCl, 0.4 mol L⁻¹ sucrose, 1 mmol L⁻¹ EDTA, 5 mmol L⁻¹ ascorbic acid, and 1 mmol L⁻¹ MgCl₂ (1:2, w/v). The homogenate was

Fig. 1 Grx activity in cotyledons (a) and embryonic axes (b) of pea seeds during germination after imbibition with H₂O or 5 mmol L⁻¹ Cd. Data are the mean of six independent measurements \pm SE. Each measurement was performed in an extract obtained from several germinating seeds



filtered through two layers of miracloth and centrifuged at $3,000\times g$ for 10 min. The resulting supernatant was centrifuged again at $20,000\times g$ for 20 min. The pellet was washed with the homogenization medium and centrifuged in the same way. The mitochondrial pellet was then discarded for other uses [8]. The combination of supernatants was regarded as post-mitochondrial fraction whose quality was routinely evaluated [20] using cytochrome *c* oxidase test (data not shown). All operations were performed at 4°C .

Enzyme Activities and Thiol Determination

Grx activity was evaluated using the hydroxyethyl disulfide (HED) reduction test [17]. The activities of GR [21] and Gpx [22] were measured spectrophotometrically. Total protein thiols were assayed according to the method of Ellman [23]. Proteins were quantified using bovine serum albumin as a standard [24].

Electrophoresis and Western Blot Analysis

Sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE) was performed with 15% (*w/v*) polyacrylamide gels by the procedure of Laemmli [25]. Proteins were then transferred to nitrocellulose sheets as in Towbin et al. [26]. Immunodetection was performed with poplar Grx C4, Grx S12, and GPX 3 antibodies.

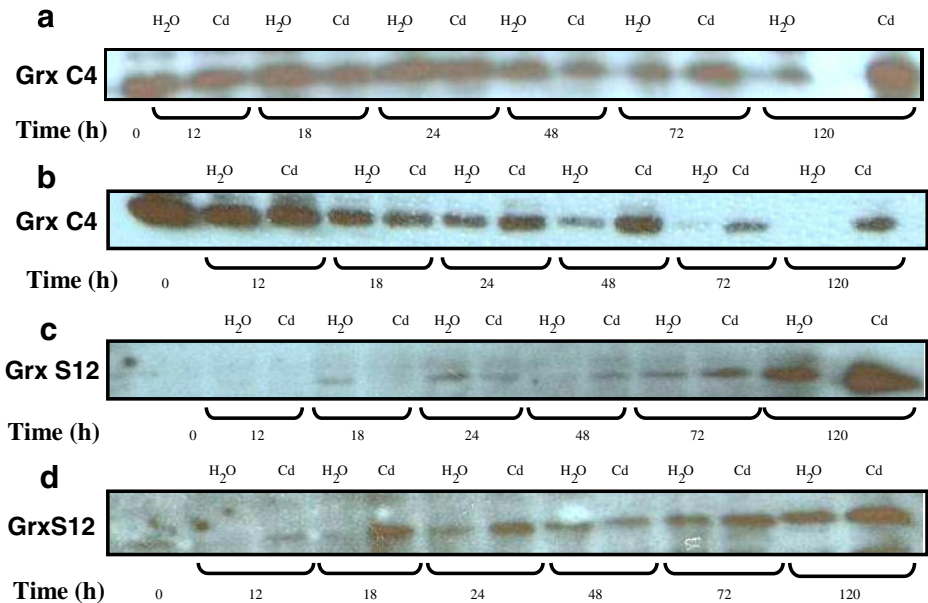


Fig. 2 Western blot analysis of Grx expression in cotyledons (**a**, **c**) and embryonic axes (**b**, **d**) of pea seeds during germination after imbibition with H₂O or 5 mmol L⁻¹ Cd. Analysis were performed after SDS-PAGE of post-mitochondrial proteins obtained from several germinating seeds (20 μg per track), transferred to nitrocellulose sheet and immunodetected with poplar Grx C4 and Grx S12 antibodies. The blot is representative of two experiments

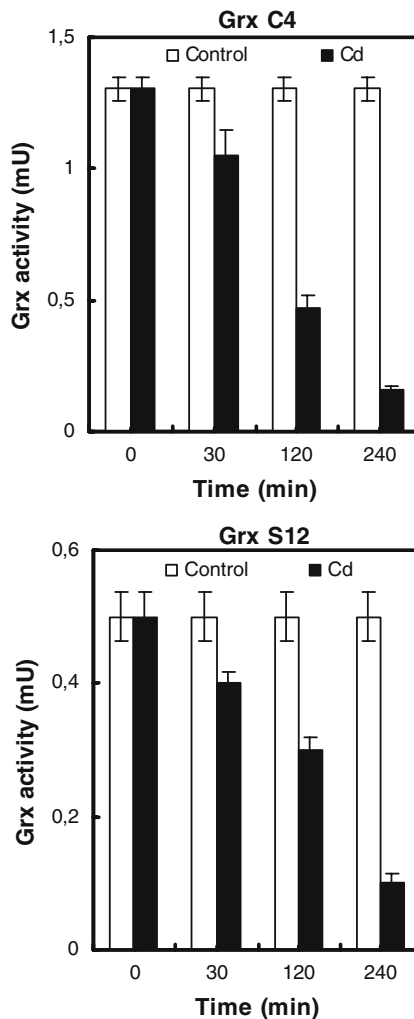
Effect of Cd Ions on Grx in Vitro

Proteins were incubated with appropriate amounts of CdCl_2 corresponding to real metal contents in tissues of Cd-treated germinating seeds [8]. Disulfide reductase activity of Grx [17] was performed after an incubation period of 30, 120, and 240 min at 4°C . Grx dimer and monomer were resolved by SDS-PAGE after incubation of protein with Cd ions in the absence or in the presence of 30 mmol L^{-1} β -mercaptoethanol (β -MET) and dithiothreitol (DTT).

Glutathione Assay

Non-protein thiols were extracted by homogenizing the post-mitochondrial fraction in 5% *w/v* sulfosalicylate ($\text{pH} < 1$) containing 6.3 mmol L^{-1} diethylene triamine pentaacetic acid. After centrifugation at $10,000\times g$ for 30 min at 4°C , the supernatants were used for total GSH determination [27].

Fig. 3 Effect of Cd ions on Grx activity in vitro. Poplar proteins ($5 \mu\text{g}$) were mixed with $5 \mu\text{g}$ Cd or equal volume of H_2O (*Control*) and incubated for different times before the enzyme assay. Disulfide reductase activity was performed as HED reduction test at 25°C . The test was performed in $500 \mu\text{L}$ of 100 mmol L^{-1} Tris-HCl, $\text{pH} 8$, 2 mmol L^{-1} EDTA, $150 \mu\text{mol L}^{-1}$ NADPH (Sigma, St. Louis, MO, USA), 1 mmol L^{-1} glutathione, 1 U GR (Sigma), and 7 mmol L^{-1} HED (Acros, Noisy-le-grand, France). The Grx isoforms were added after 2 min of incubation, allowing disulfide bridge formation between glutathione and HED. The oxidation of NADPH was measured at 340 nm . Data are the mean of six independent measurements \pm SE



Results and Discussion

The Grx system plays a fundamental role in controlling the redox status in animals and plants subjected to abiotic stress as Cd [19, 28]. Exposure of pea seeds to cadmium during germination results in the increase of the overall Grx activity: about 47% and 136% from

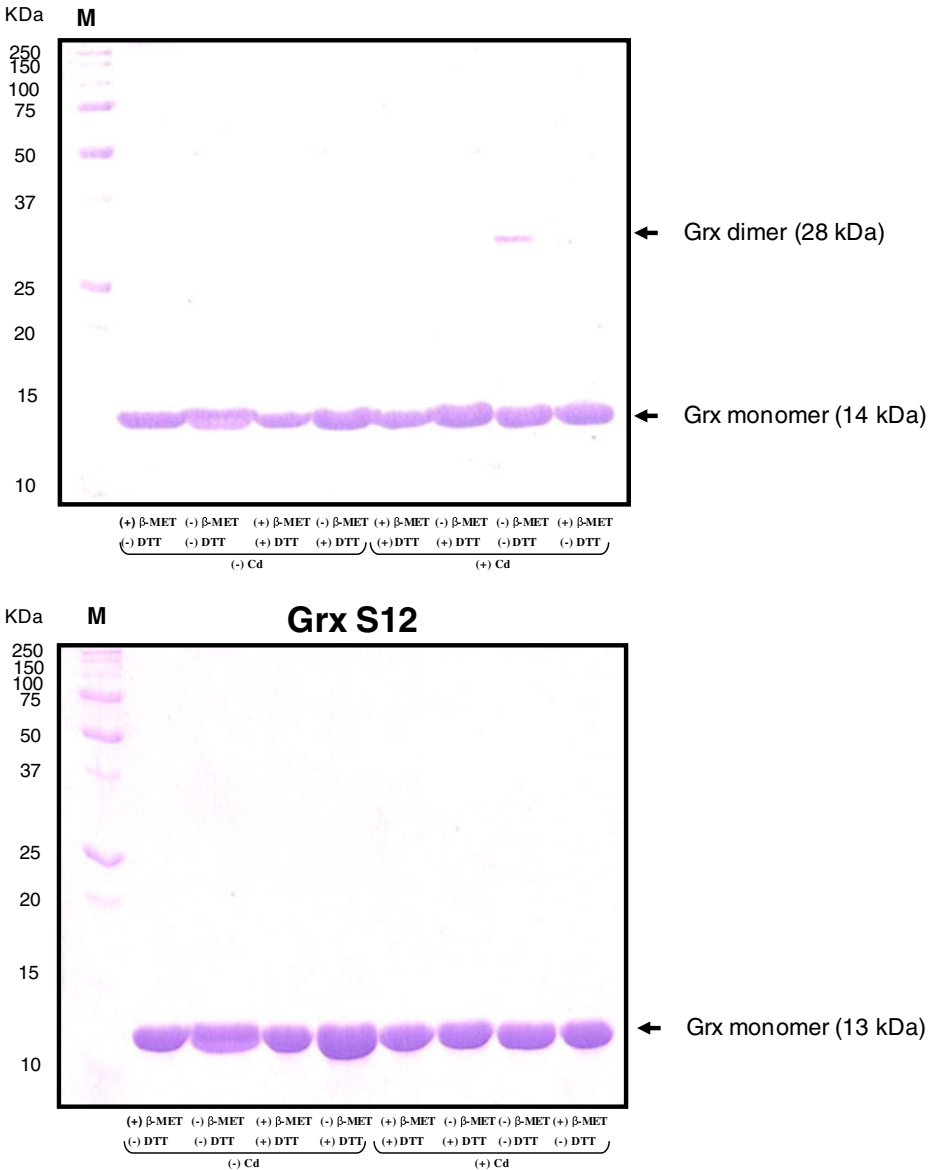
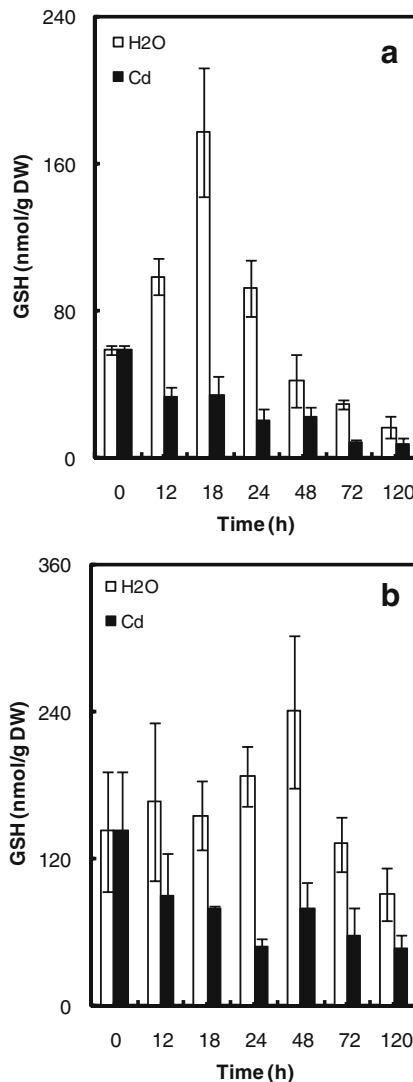


Fig. 4 Effect of Cd binding on Grx. Poplar proteins ($5 \mu\text{g}$ per lane) were mixed with $5 \mu\text{g}$ Cd, 30 mmol L^{-1} β-MET, and DTT (+) or equal volume of H_2O (-) and incubated for 240 min before being subjected to 15% SDS-PAGE. M, molecular weight markers are indicated in kilodaltons. Grx dimer and monomer were stained with Coomassie blue. Experiments were performed in duplicate

controls in cotyledons and embryonic axes, respectively, after 5 days of treatment (Fig. 1). Moreover, the expression of some isoforms, namely, Grx C4 and Grx S12, was induced following the exposure to Cd (Fig. 2).

However, heavy metals are known to bind thiols with high affinity [29, 30]. Although Cd has been implicated as a pro-oxidant, it differs from other metal ions like Cu and Fe because it does not participate in redox cycling [31]. This can implicate a direct binding of Cd ion to critical cellular components such as proteins with vicinal dithiols that are expected to be particularly sensitive to inactivation. Consequently, the active site of reduced Grxs is a possible target of Cd fixation. To test further the supposed mechanism of inactivation, we have analyzed the effect of pre-treatment with Cd ions on the activity of (a) bicysteinic Grx C4 and (b) monocysteinic Grx S12

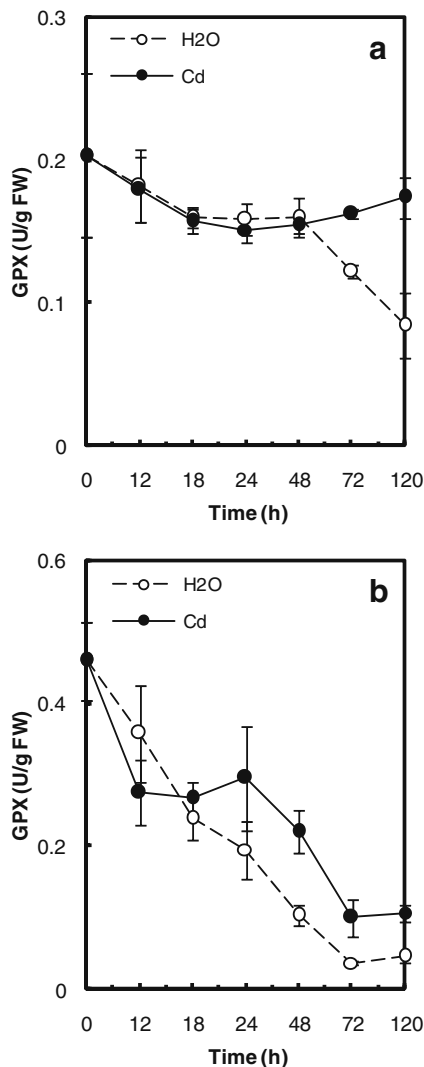
Fig. 5 Content of total glutathione in cotyledons (a) and embryonic axes (b) of pea seeds during germination after imbibition with H₂O or 5 mmol L⁻¹ Cd. Data are the mean of six independent measurements \pm SE. Each measurement was performed in an extract obtained from several germinating seeds



where the active site contains one thiol (Cys) and one hydroxyl (Ser) [15, 17]. The disulfide reductase activity, performed as HED reduction test, of both isoforms was markedly diminished in a time-dependent manner (Fig. 3). The inhibition of the wild-type Grx C4 activity may be due, at least in part, to the formation of dimer. In fact, the electrophoretic analysis showed that the reduced form of Grx C4 (monomer; 14 kDa) can be converted to an oxidized state in the presence of cadmium (dimer; 28 kDa) (Fig. 4). The reduced state was restored by the addition of either β -MET or DTT. These experiments indicate that Grxs are able to bind Cd, presumably at the level of their active site thiols. However, Grx S12 remained in the monomeric state (13 kDa) even in the presence of the heavy metal (Fig. 4).

Thus, in spite of the negative effect of a direct interaction between Cd ion and Grx, it seems that heavy metal stress might enhance the protein expression and,

Fig. 6 GPX activity in cotyledons (a) and embryonic axes (b) of pea seeds during germination after imbibition with H₂O or 5 mmol L⁻¹ Cd. Data are the mean of six independent measurements \pm SE. Each measurement was performed in an extract obtained from several germinating seeds



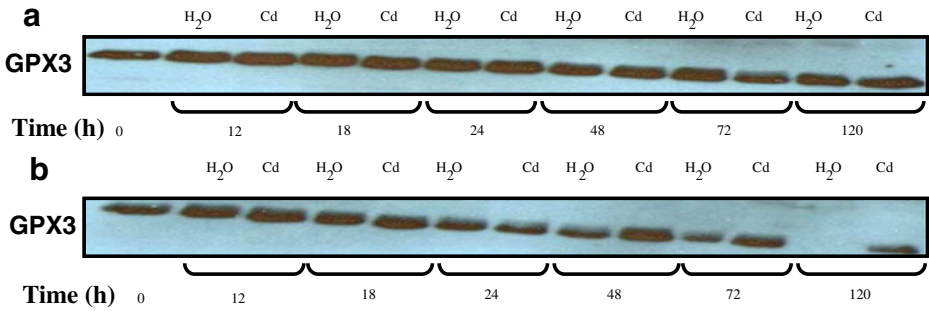
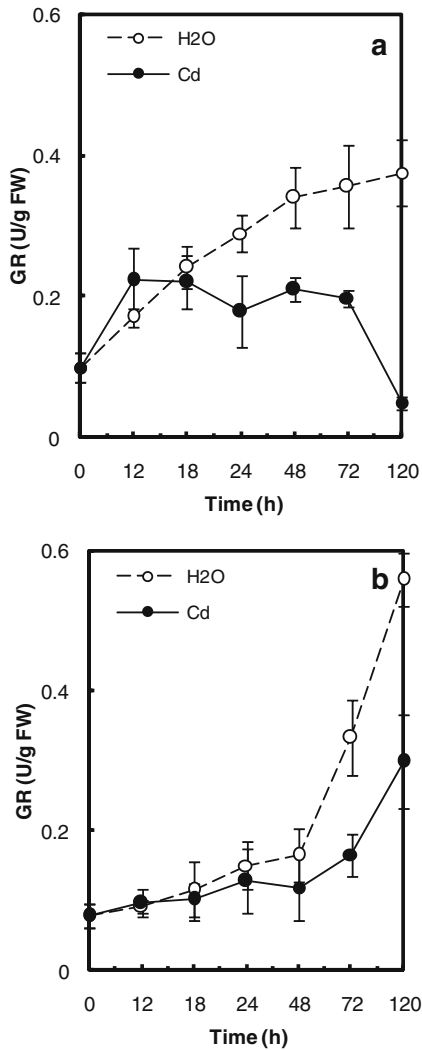


Fig. 7 Western analysis of GPX expression in cotyledons (a) and embryonic axes (b) of pea seeds during germination after imbibition with H₂O or 5 mmol L⁻¹ Cd. Analysis were performed after SDS-PAGE of post mitochondrial proteins obtained from several germinating seeds (20 µg per track), transferred to nitrocellulose sheet and immunodetected with poplar GPX3 antibody. The blot is representative of two experiments

Fig. 8 GR activity in cotyledons (a) and embryonic axes (b) of pea seeds during germination after imbibition with H₂O or 5 mmol L⁻¹ Cd. Data are the mean of six independent measurements ± SE. Each measurement was performed in an extract obtained from several germinating seeds

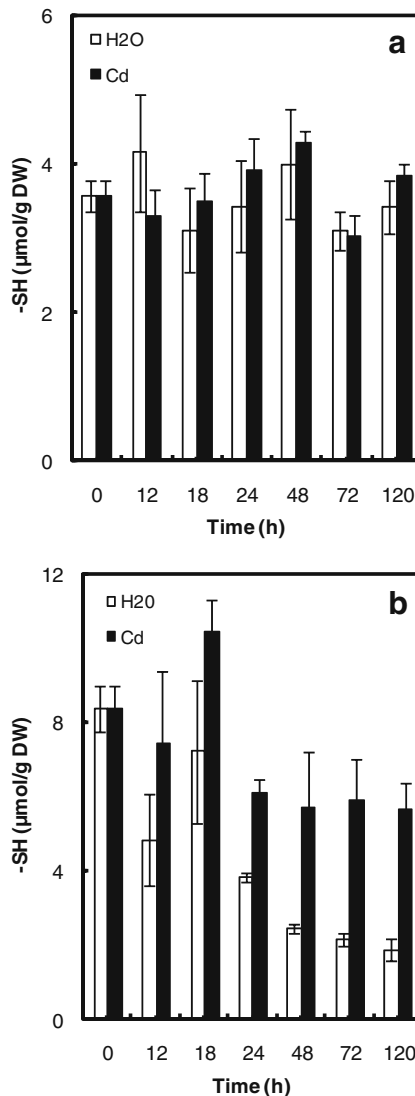


consequently, the absolute activity. The stimulatory effect should compensate the vulnerability of Grx to dimerization or any other mechanism of chemical inactivation imposed by Cd.

Under normal conditions, the intra-cellular redox state is predominately reducing, but processes like oxidative stress can shift the redox balance toward an oxidizing state [15, 16, 18, 32]. In adult pea, we have demonstrated that Cd causes an oxidative stress [4, 5]. In this study, we bring evidence for relationships between the GSH-dependent redox status, Grx, and Cd treatment during pea seed germination.

The level of GSH decreased at least twofold in Cd-treated germinating seeds as compared to controls (Fig. 5). Peroxidation is one of the expected consequences of oxidative stress in pea [4, 5]. Cadmium treatment stimulated the enzymatic activity of GPX

Fig. 9 Content of total protein thiol in cotyledons (a) and embryonic axes (b) of pea seeds during germination after imbibition with H₂O or 5 mmol L⁻¹ Cd. Data are the mean of six independent measurements \pm SE. Each measurement was performed in an extract obtained from several germinating seeds



after 48 h of exposure (Fig. 6). GPX 3 isoform was immunodetected with high expression levels in dry seed tissues and was markedly decreased during H₂O imbibition but remained significantly high at least in the Cd-treated embryos (Fig. 7b). Moreover, Cd stress provoked a drastic decrease in GR activity: about 87% and 47% from cotyledon and embryo controls after 5 days, respectively (Fig. 8).

The formation of disulfides and mixed disulfides between GSH and other thiols such as protein can be expected to occur under oxidative stress conditions [15]. A similar phenomenon might be also responsible for the decline in the GSH status in Cd-treated germinating pea seeds, since protein–glutathionyl–mixed disulfides formation is a mechanism of regulation and implicates Grx as a key player in this cellular process [15].

We can exclude a possible role for peroxidases in the GSH depletion (Figs. 6 and 7). Indeed plant peroxidases have a thiol-oxidase function [33], but so-called plant “glutathione” peroxidases are in fact thioredoxin- and not glutathione-dependent enzymes [22]. Anyway, under Cd stress, an increase in the GPX activity has been reported during germination [9]. Moreover, the resulting loss in the activity of enzymes such as GR (Fig. 8) leads to an increase in the generation of an intracellular oxidative stress. These observations are in agreement with the results obtained by Noriega et al. [34] and Yannarelli et al. [35]. It has been also reported that redox perturbations, especially in GSH levels, act as signal in stimulating stress response [10, 36–40].

On the other hand, the protein thiols seem to be protected against the Cd-imposed oxidation. In fact, SH-protein content remained statistically unchanged in the cotyledons and enhanced markedly in the embryos of Cd-treated seeds, as compared to the respective controls (Fig. 9). This can attest to the possibility of the cellular thiol response of germinating pea seed against Cd toxicity.

Conclusion

All data led us to underlie the relationships between thiol management and some Grx thiol-dependent responses during the germination of Cd-treated pea seeds. Despite the loss in GSH pool and GR activity due to the generation of an intracellular oxidative stress, a protective action of high Grx expression and activity on thiols is possible to improve the redox status. However, Grxs are often involved in concert with thioredoxins whose gene expression and activity can be also regulated by heavy metals. For these reasons together, we have started an investigation to elucidate how the thioredoxin system might contribute to control the redox homeostasis in germinating pea seeds exposed to cadmium.

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