

BRIEF COMMUNICATION

ABA or cadmium induced phytochelatin synthesis in potato tubers

A. STROIŃSKI*, T. CHADZINIKOLAU, K. GIŻEWSKA and M. ZIELEZIŃSKA

*Plant Physiology Department, University of Life Sciences, Wołyńska 35, PL-60637 Poznań, Poland***Abstract**

Short-term treatment of potato (*Solanum tuberosum* L.) tuber discs with CdCl₂ solution elevated both the *StPCS1* transcript level, phytochelatin synthase (PCS) activity and contents of phytochelatin (PC) and abscisic acid (ABA). Similar effects but less marked were noticed after treatment of tuber tissues with ABA solution. Cd-treatment increased also cysteine (CYS) content but did not change glutathione content. If ABA treatment preceded Cd-treatment, the elevation of CYS and PC contents were limited. The data suggest the participation of ABA in the regulation of PCS.

Additional key words: phytochelatin, phytochelatin synthase, *Solanum tuberosum*

The cadmium treatment may lead to plant growth inhibition, chlorosis and deficiencies of some essential elements (Stiborová *et al.* 1987, Stroiński *et al.* 1990, Karcz and Konopka 2007, Maksimović *et al.* 2007). Cd can intensify some harmful processes in plants, namely oxidative stress (Hendry *et al.* 1992, Stroiński and Kozłowska 1997, Sanita di Toppi and Gabrielli 1999, Skórzyńska-Polit *et al.* 2003) or inactivation of macromolecules and cellular structure (Stroiński *et al.* 1990, Jiang *et al.* 2009). The resistance of plant cells to Cd depends, among other things, on the content of phytochelatin (PC), which chelate cadmium ions. These peptides are synthesized from glutathione and related thiols by PC synthases (PCS; γ -glutamyl-cysteine dipeptidyl transpeptidases; EC 2.3.2.15) in the presence of heavy metal ions (Grill *et al.* 1989, Maitani *et al.* 1996, Clemens 2006). PCS genes were characterized in *Arabidopsis thaliana* (*AtPCS1*, *AtPCS2*), in wheat (*TaPCS1*), in *Brassica juncea* (*BjPCS1*), in rice (*OsPCS1*) (Cobbett and Goldsbrough 2002) and in potato (*StPCS1*) (Nuc *et al.* 2003). *AtPCS1* did not exhibit transcriptional regulation by Cd in *Arabidopsis*, although its activity was increased in the presence of Cd (Ha *et al.* 1999, Vatamaniuk *et al.* 2000). In other plants the biosynthesis of some phyto-chelatin synthases showed transcriptional and/or post-transcriptional regulation by Cd (Clemens *et al.* 1999, Stroiński and Zielezińska 2001, Cobbett and Goldsbrough 2002).

Until now there have been scarce reports characte-

rizing hormonal participation in plant response to Cd. Our initial experiment revealed that Cd infiltrating potato tuber tissues significantly changed ABA content in those tissues (Chadzinikolau and Stroiński 2003). Cd treatment resulted in an increase in ABA content in leaves of rice (Hsu and Kao 2003, 2005, Hsu *et al.* 2006) and in roots of *Typha latifolia* and *Phragmites australis* (Fediuc *et al.* 2005). ABA mediates plant responses to environmental stress, such as high salinity, drought, low temperature and mechanical wounding (Giraudat *et al.* 1994, Xiong *et al.* 2002, Shinozaki *et al.* 2003). Usually under these stress conditions ABA accumulation was observed and this effect was associated with growth inhibition (Cheng *et al.* 2002, Eckardt 2002). The present paper describes the effect of cadmium and exogenous ABA on ABA, cysteine (CYS), glutathione (GSH) and PC content and on *PCS1* gene expression and PCS activity in potato tuber tissues.

Experiments were carried out using tubers of potato (*Solanum tuberosum* L.) cv. Bzura (cultivar tolerant to cadmium; Stroiński *et al.* 1990). Discs (18 mm in diameter, 4 mm thick) were infiltrated with H₂O or with ABA solution (0.1 mM) under two vacuum impulses for 5 min, washed with distilled water and incubated for 24 h in a moist chamber at 20 °C in the dark. Next day, they were infiltrated in similar manner with CdCl₂ solution (1 mM), washed with distilled water, and finally incubated in a moist chamber at 20 °C in the dark for 24 h. Then the materials were immediately frozen in liquid nitrogen.

Received 11 June 2007, accepted 21 September 2008.

Abbreviations: ABA - abscisic acid; CYS - cysteine; GSH - glutathione; PC - phytochelatin; PCS - phytochelatin synthase.

Acknowledgements: This study was supported by KBN grant No. 6 P06A 014 20.

* Corresponding author; fax: (+48) 61 8487179, e-mail: astroins@o2.pl

Thiols (cysteine, glutathione and PC) were isolated and separated by a reverse-phase HPLC with post column derivatization as described earlier (Tukendorf and Rauser 1990, Stroiński and Zielezińska 1997). PCS activity was determined according to Grill *et al.* (1989). The amount of PC was calculated from a comparison to GSH standard. Total protein contents were measured according to Bradford (1976).

Total RNA was extracted from tuber tissues using the phenol-chloroform method (Branch *et al.* 1989). Poly (A) RNA was isolated from the total RNA with a *Dynabeads* mRNA isolation kit (*Dynal*, Hamburg, Germany) and used to prepare the cDNA. The first-strand cDNAs were synthesized by reverse transcription from 0.1 - 0.5 µg of mRNA isolated from *Solanum tuberosum* by using AMV reverse transcriptase (*Promega*, Madison, USA) and an oligo dT primer according to the manufacturer's instructions. Amounts of synthesized cDNA for the PCR reaction were measured spectrofluorometrically by using the OilGreen dye with oligonucleotide standard M13 (*Molecular Probes*, Carlsbad, USA). The PCR reaction was carried out by using 2× PCR Master Mix (*Fermentas*, Vilnius, Latvia), which contained a concentrated solution of 0.05 U mm⁻³ *Taq* DNA polymerase (recombinant). Specific primers were used for the phytochelatin synthase gene (*StPCS1*): 5'AAATGGAAAGGGCCTTGGAG 3' and 5' GACCCAGTGAGGGGGATA 3' (Nuc *et al.* 2003). PCR products were separated electrophoretically on a 1.1 % agarose gel with the addition of ethidium bromide.

The isolation and estimation of ABA by high-performance liquid chromatography (HPLC) were done according to Moore (1990) with ABA-methyl ester as an internal standard for estimating extraction efficiency. Amounts of ABA were calculated from its peak area, and peak area of internal standard using Chromatography Station program (*Data Apex*, Prague, Czech Republic).

To determine the soluble (Cd_s) and bound (Cd_B) cadmium, the potato tissues were homogenized in 200 mM phosphate buffer at pH 7.8 and the resultant homogenates were centrifuged at 20 000 g for 20 min. Supernatants were collected to estimate Cd_s concentrations. On the other hand, to determine Cd_B concentration, the precipitates were wet mineralized to ash by the use of acid mixture (36 % HCl + 20 % HClO₄

+ H₂O, 2:1:1, v/v). Cadmium was determined by means of flame atomic absorption spectrometry.

The data presented in the figures are means of three or more independent replications. The significance of differences between means was calculated using unpaired *t*-test.

Our previous studies indicated that a short-term treatment of potato (*Solanum tuberosum* L.) tuber discs with CdCl₂ induced an increase of phytochelatin synthase (PCS) activity in concentration and time dependent manner (Stroiński and Zielezińska 2001).

In control materials used in the present experiments, we found small amounts of Cd, and low levels of the *StPCS1* transcript and PCS activity. Cadmium treatment of tuber tissues caused a considerable accumulation of Cd about thirty times and simultaneous increase in *StPCS1* transcript level and PCS activity. Cd stress also increased the contents of CYS and PC (Table 1, Fig. 1). Similar CYS changes caused by Cd treatment were observed in soybean seedlings (El-Shintinawy and El-Ansary 2000) and in maize roots (Nocito *et al.* 2002). At the same time, the GSH content did not change in the tuber tissues (Table 1). Previously Cd-induced decrease in GSH content was observed in roots of maize seedlings (Tukendorf and Rauser 1990) and in spinach roots (Tukendorf 1993).

Until now there have been no reports which would characterize the hormonal participation in the regulation of cadmium detoxification, particularly in the regulation of PCS activity. The ABA content increased approximately four times after Cd application. Similar results were found by Hsu and Kao (2003, 2005), Hsu *et al.* (2006) and Fediuc *et al.* (2005). The exogenous ABA treatment of tuber tissues increased the endogenous

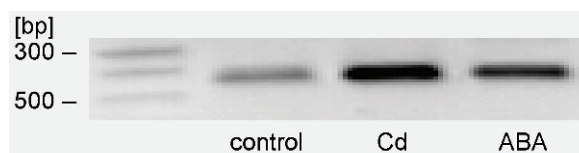


Fig. 1. Influence of exogenous ABA and Cd on expression of *StPCS1*. RT-PCR were conducted using 40 ng of total mRNA extracted from tissues exposed or not exposed to CdCl₂ (0.1 mM) for 48 h, or with ABA (0.1 mM) for 24 h. Specific primers to *StPCS1* cDNA for 352 bp product were used.

Table 1. Effect of exogenous ABA and Cd on Cd, ABA, CYS, GSH and PC content and on PCS activity in potato tuber tissues. Values are the means of three replicates ± SE. The values marked with different letters are significantly different at *P* = 0.05.

	Control	ABA	ABA + Cd	Cd
Soluble Cd content [µg g ⁻¹ (f.m.)]	0.90 ± 0.1 ^a	-	22.60 ± 1.5 ^b	21.30 ± 1.4 ^b
Bound Cd content [µg g ⁻¹ (f.m.)]	2.20 ± 0.2 ^a	-	30.60 ± 4.0 ^b	55.20 ± 0.6 ^c
ABA content [µg g ⁻¹ (f.m.)]	0.44 ± 0.0 ^a	0.88 ± 0.0 ^b	1.56 ± 0.1 ^c	1.72 ± 0.1 ^c
CYS content [nmol g ⁻¹ (f.m.)]	7.40 ± 1.0 ^a	5.00 ± 0.5 ^b	8.00 ± 1.4 ^a	12.30 ± 1.0 ^c
GSH content [nmol g ⁻¹ (f.m.)]	15.30 ± 1.1 ^a	15.70 ± 1.3 ^a	16.60 ± 1.0 ^a	19.40 ± 1.4 ^a
PC content [nmol g ⁻¹ (f.m.)]	6.70 ± 0.7 ^a	11.90 ± 1.5 ^b	26.00 ± 2.6 ^c	36.30 ± 3.5 ^c
PCS activity [nmol(PC) mg ⁻¹ (protein) h ⁻¹]	4.40 ± 1.1 ^a	8.70 ± 1.6 ^b	13.60 ± 3.6 ^c	16.20 ± 2.1 ^c

ABA content (Table 1) and it confirmed earlier observations of the positive feedback in ABA biosynthesis by ABA (Audran *et al.* 2001, Xiong and Zhu 2003). However in this experiment, ABA can be accumulated nonspecifically in plant tissue by passive transport.

Simultaneously, this ABA treatment induced significant increase in the *PCS1* transcript level, PCS activity and PC content (Table 1, Fig. 1) similarly as Cd treatment. Additionally, the experiments when ABA

treatment preceded Cd stress indicated that these effects were not additive (Table 1, Fig. 1). It seems that both factors can modulate PCS activity differently using signaling pathways possessing some common elements (Knight and Knight 2001).

These observations, for the first time, can suggest the participation of ABA in the regulation of phytochelatin synthase, the main step of the cadmium detoxification process.

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