EXCESS COPPER INDUCES STRUCTURAL CHANGES IN CULTURED PHOTOSYNTHETIC SOYBEAN CELLS

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ABSTRACT

Soybean cell suspensions have the capacity to develop tolerance to excess copper, constituting a convenient system for studies on the mechanisms of copper tolerance. The functional cell organization changes observed in these cell cultures after both a short-term (stressed cells) and a long-term (acclimated cells) 10 μM CuSO_4 exposure are reported by using a structural, cytochemical and microanalytical approach. Cells grown in the presence of 10 μM CuSO_4 shared some main structural features with non-treated cells such as: *i*) a large cytoplasmic vacuole; *ii*) chloroplasts along the thin layer of cytoplasm; *iii*) nucleus in a peripheral location exhibiting circular-shaped nucleolus and a decondensed chromatin pattern; *iv*) presence of Cajal bodies in the cell nuclei. In addition, cells exposed to 10 μM CuSO_4 exhibited important differences compared with non-treated cells: *i*) chloroplasts displayed rounded shape and smaller size with a denser structured internal membranes specially in copper acclimated cells; *ii*) no starch granules were found within chloroplasts; *iii*) cytoplasmic vacuole was larger specially after a long-term copper exposure; *iv*) the levels of citrate and malate increased. Extracellular dark deposits attached at the outer surface of the cell wall with high copper content were only observed during a short-term copper exposure as a differential feature. Structural cell modifications, mainly affecting chloroplasts, accompanied the short-term copper induced response and were maintained as a stable character during the adaptive period to excess copper. Vacuolar changes rather accompanied the long-term copper response. The results indicate that the first response of soybean cells to excess copper avoids its entrance to the cell by its immobilization in the cell wall, and after an adaptive period copper acclimation may be mainly due to vacuolar sequestration.

Keywords: acclimation, copper, microscopical structural analysis, organic acids, soybean cell culture, tolerance

Abbreviations: Chl, chlorophyll; DAPI, 4'-6-diamidino-2-phenylindole; EDX, energy dispersive X-ray; LTSME; low temperature scanning electron microscopy.

INTRODUCTION

Copper (Cu) is an essential micronutrient for plants. Copper acts as a structural element in certain metalloproteins, many of which are involved in electron transport in mitochondria and chloroplasts, and oxidative stress response (Raven et al. 1999; Gratão et al. 2005). It also participates in cell wall metabolism, hormone signalling, protein trafficking machinery, transcription signalling and iron mobilization (for reviews see Yruela 2005; Pilon et al. 2006). Nevertheless, it can also be a toxic element at a tissue concentration slightly higher ($20\text{--}30\ \mu\text{g g}^{-1}$ dry weight) than the normal one ($6\text{--}12\ \mu\text{g g}^{-1}$ dry weight) (Baker and Senef 1995; Marschner 1995). Although normally copper binds to proteins, it may be released and become free to catalyze the formation of highly reactive hydroxyl radicals *via* Fenton-type reactions. Thus, copper has capacity to initiate oxidative damage and interfere with important cellular processes such as photosynthesis, pigment synthesis, plasma membrane permeability and other metabolic mechanisms causing a strong inhibition of plant development (Küpper et al. 2003; Bertrand and Poirier 2005; Yruela 2005). Since copper is both an essential cofactor and a toxic element its uptake and cellular concentrations must be strictly regulated. Thus, plants use different strategies to appropriately regulate copper homeostasis as a function of copper environmental level (Pilon et al. 2006; Krämer and Clemens 2006).

At present, there is an increasing concern about copper toxicity in agriculture and health since copper is considered a major heavy metal contaminant that results from fertilizers accumulation, the application of pig and poultry slurries rich in copper, fungicides, industrial and urban activities, mining or metal processing (Kabata-Pendias and Pendias 2001; Pilon-Smits and Pilon 2002). Copper concentration in non-contaminated soils and natural waters is *ca.*

20-30 mg kg⁻¹ and 2 µg kg⁻¹, respectively but in contaminated soils and waters can reach levels one hundred times higher (Fernandes and Henriques 1991). Some plant species or ecotypes are sensitive to metals, whereas others are tolerant showing little growth inhibition or damage even if they grow in severely polluted soils and accumulate high concentrations of these elements in their tissues. It was demonstrated that the plant *Silene vulgaris* exhibits tolerance to different metals (Zn, Cd or Cu) (Schat et al. 1997; Van Hoof et al. 2001). Different response to excess copper were observed in four Australian tree species, *Acacia holosericea*, *Eucalyptus crebra*, *Eucalyptus camaldulensis* and *Melaleuca leucadendra* (Reichman et al. 2006). Nowadays, mechanisms of heavy metal tolerance in plants are of interest and numerous works have been conducted on this subject. In general, control of metal uptake, excretion of accumulated metals and intracellular immobilization of those are the main mechanisms that operate alone or in concert to protect plants from toxic effects (Ernst et al. 1992; Clemens 2001; Hall and Williams 2003; Kramer and Clemens 2006). However, in some cases the mechanisms underlying the tolerance to metals are not fully understood and in particular those mechanisms related to copper tolerance. In soybean seedlings treated with excess copper it has been reported that the enhancement in peroxidase and laccase activities was accompanied by a rapid increase of lignin contents in the copper treated tissues (Lin et al. 2005). The increase in phenolic compounds, related enzymes and lignin was observed in root suspension cultures of *Panax ginseng* in response to copper stress (Ali et al. 2006). Intraspecific and interspecific differences in sensitivity to copper do occur between different plant species. On the other hand, with regard to mechanisms allowing a copper tolerance, a question of

interest is whether this tolerance is constitutive in each species or depends on previous long-term exposure to metal.

Plant cell cultures are widely used as suitable model system to analyse metabolic signalling pathways, developmental process and cell stress response and adaptation, among many other studies on plant physiology (Roitsch and Sinha, 2002; Allan et al. 2006; Zuppini et al. 2006). Plant cell cultures can be a useful tool for studying the characteristics and mechanisms of heavy metal tolerance in plants at cellular level. In this respect, only a few studies concerning heavy metal tolerance in plant cell suspensions exist. Selection and characterization of various cell cultures from *Nicotiana* tolerant to aluminium, cadmium, copper and zinc have been reported (Kishinami and Widholm 1986, 1987; Gori et al. 1998). Cadmium stress has been studied in sugar cane callus cultures (Fornazier et al. 2002). However, a very little information exist from cell suspensions of other species. Since the cell culture used in this work comes from mesophyll cells, the results presented here may provide information to advance in our knowledge of physiological aspects of leaf cells in plants. In a previous paper we reported that soybean cell suspensions have capacity to develop tolerance up to 50 μM CuSO_4 after a long-time copper exposure (Bernal et al. 2006). In this work we investigated more in detail the structural cellular changes induced by excess copper on soybean cell suspensions. The comparison between a short-term and a long-term copper exposure is presented.

MATERIALS AND METHODS

Cell culture conditions.- Photosynthetic cell cultures from soybean (*Glycine max* var. Corsoy) SB-P line were grown as described by Rogers et al. (1987) with some modifications. Cell suspensions were grown in liquid cultures photomixotrophically (KN¹ medium) under continuous low light ($30 \pm 5 \mu\text{E m}^{-2} \text{s}^{-1}$). The KN¹ medium contained thiamine (0.1 mg/L), kinetin (0.2 mg/L), naphthalene acetic acid (1 mg/L) and sucrose (1% w/v) as organic components. It has been reported that this kind of cultures grow optimally with $65\text{--}75 \mu\text{E m}^{-2} \text{s}^{-1}$ (Alfonso et al. 1996) and that high light cause photoinhibitory effects in these cell suspensions (personal communication; Alfonso et al. 1996). On the other hand, it is known that copper is a potential toxic element and its toxicity enhances with light. Thus, we decided to assay a low light regime ($30 \pm 5 \mu\text{E m}^{-2} \text{s}^{-1}$) following the light conditions used in previous works studying the copper stress effect on cell cultures (Gori et al. 1998, Bernal et al. 2006). To assay the copper effect on cell growth the media were supplemented with $10 \mu\text{M CuSO}_4 \cdot 5\text{H}_2\text{O}$. Control medium corresponded to $0.1 \mu\text{M CuSO}_4 \cdot 5\text{H}_2\text{O}$. Cell suspensions were sub-cultured each 23 days of growth (one transfer) by using fresh media containing the appropriated copper concentrations. For experiments cells were collected after the first 21 days of growth in the presence of $0.1 \mu\text{M CuSO}_4$ (non-treated cells) and $10 \mu\text{M CuSO}_4$ (Cu-stressed cells), and after 22 transfers in the presence of $10 \mu\text{M CuSO}_4$ (Cu-acclimated cells). For microscopical structural analysis, soybean cells were grown in 1.5% (w/v) agar plates with KN¹ medium at 24 °C and atmosphere with 5% CO₂. Cells cultured in these conditions were easier to handle during the fixation and sectioning procedures than liquid suspensions.

Sample processing for microscopical structural analysis.- Soybean cells were fixed overnight at 4°C in 4% (w/v) formaldehyde in phosphate buffered saline solution (PBS), pH 7.3. Then, they were washed in PBS and dehydrated through an acetone series (30%, 50%, 70% and 100% (v/v)) at 4°C. The samples were infiltrated and embedded in Historesin 8100 at 4°C. Semithin sections (1 µm thickness) were obtained and used for light microscopy observations. Toluidine-blue stained semithin sections were observed under bright field and phase contrast field for structural analysis in a Leitz microscope fitted with a digital camera Olympus DP10.

Morphometric analysis of cell, chloroplast and vacuole size.- Random sampling was carried out over toluidine-stained micrographs of the three cultures. The number of micrographs to be taken was determined using the progressive mean test, with a maximum confidence limit of $p \leq 0.05$. The cell, chloroplast and vacuolar area was measured in µm² using a square lattice composed by 40 x 57 squares of 5 x 5 mm (cell and vacuole) or 65 x 96 squares of 3 x 3 mm (chloroplast) each. For each culture, the mean area per cell was estimated and the results were compared among the cultures. Duncan's multiple range test ($p \leq 0.05$) was used to compare the data among the cultures.

Cytochemical stainings for starch and DNA.- Starch was detected by I₂KI staining (O'Brien and McCully 1981) on Historesin semithin sections and observed under bright field (Barany et al. 2005). DAPI staining for DNA was applied to semithin sections (Testillano et al. 1995) and observed under UV light in a Zeiss Axiophot epifluorescence microscope fitted with a CCD camera.

Copper determination.- Cells were harvested and washed twice with 3 mM EDTA and once with distilled H₂O to remove free cations. After washing, cells were filtered through a layer of Miracloth (Calbiochem, EMD Biosciences Inc, San Diego, CA, USA) and dried in a ventilated oven at 60 °C for 48 h. Dried samples were treated using a standard procedure (Abadía et al. 1985). Analyses were performed in an atomic absorption spectrometer (UNICAN 969).

Organic acids analysis.- Cells were harvested on Miracloth layer (Calbiochem, EMD Biosciences Inc, San Diego, CA, USA), washed three times with cold distilled water and then homogenized in 70% (v/v) ethanol in a Teflon homogenizer. The homogenate was centrifuged at 10,000 x g for 10 min at 4°C and the pellet was washed with 70% (v/v) ethanol three times. The supernatants were combined and dried under vacuum at 35°C in SPD Speed Vac model SPD111V (Thermo Savant, Holbrook, NY, USA). The residue was resuspended in 0.1% (v/v) formic acid and filtered with 0.2 µm nylon filter (Tecknokroma, Barcelona, Spain), taken to a final volume of 0.5 ml and stored at -80°C until analysis. Organic acids were analysed by HPLC-ESI/MS with a BioTOF® II (Bruker Daltonics, Billerica, MA, US) coaxial multipass time of flight mass spectrometer (MS (TOF)) equipped with an Apollo electrospray ionization source (ESI), and coupled to a Waters Alliance 2795 HPLC system (Waters Corp., Mildford, MA, US) with a Supelcogel H 250 x 4,6 mm column. The mobile phase was 0.1% (v/v) formic acid. Duncan's multiple range test ($p \leq 0.05$) was used to compare the data among the cultures.

Energy-dispersive X-ray microanalysis.- Cells were cry fixed using a high-pressure freezer (Leica EMPact; Leica Microsystems, Gladesville, Australia) and were then stored in liquid nitrogen until freeze-substitution. Frozen cells were freeze-substituted with diethyl ether containing 10% acrolein with freshly activated 5Å molecular sieve (Baker Analyzed) for 4 days at -90°C, 2 days at -70°C and 1 day at -30°C. Subsequently, samples were gradually brought to 4°C by raising the temperature for 1.5 h (Bidwell et al. 2004). The freeze-substituted cells were infiltrated with Araldite resin over 4 days. Araldite contains negligible levels of elements detectable by energy-dispersive spectrometry (Pålsgård et al. 1994). Cell sections were cut dry at 0.5 µm using an ultramicrotome (Leica Ultracut UCT) and mounted directly onto titanium grids (75 mesh) that were coated with formvar. Sections were stored in a desiccator under vacuum until analysis to prevent absorption atmospheric water and hence possible redistribution of ions.

Unstained sections were analysed at room temperature in a carbon holder by energy-dispersive X-ray microanalysis using a transmission electron microscope (model H-800-MT, Hitachi) and X-ray EDX Quantum detector (model 3600, Kevex). The detector was interfaced to a signal processing unit (Röntec, Ltd, Normanton, UK). The electron microscope was operated at 100 kV in STEM mode with a spot size 10-15 nm and a beam current 15 µA. The spectral data from individual cells and compartments in each section were collected using selected area analysis with an acquisition time of 300 s. Analyses were carried out by Quantax 1.5 program (Röntec Ltd, Normanton, UK). Spectra were normalized at titanium peak used as an internal standard, which came from the grid material.

Low Temperature Scanning Electron Microscopy (LTSEM).- Samples were examined by using the Low Temperature Scanning Electron Microscopy LTSEM technique (De los Ríos et al. 2004). Cells were mechanically fixed onto the specimen holder of a cryotransfer system (Oxford CT1500), plunged into subcooled liquid nitrogen, and then transferred to a preparation unit via an air lock transfer device. The frozen specimens were cryofractured and transferred directly via a second air lock to the microscope cold stage, where they were etched for 2 min at -90°C. After ice sublimation, the etched surfaces were sputter coated with gold in the preparation unit. Samples were subsequently transferred onto the cold stage of the scanning electron microscope chamber. Fractured surfaces were observed with a DSM 960 Zeiss scanning electron microscope at -135°C.

RESULTS

The main features of the cell structure in the soybean photosynthetic cell cultures grown in control medium (non-treated cells) and in the presence of 10 μM CuSO_4 after either the first 21 days of copper treatment (Cu-stressed cells), or 22 transfers exposed to excess copper (Cu-acclimated cells) were analyzed (for details see Materials and Methods). Cells accumulated copper during the first 21 days of growth in the presence of 10 μM CuSO_4 and the copper concentration in acclimated cells even slightly increased during the 10 μM CuSO_4 treatment (Fig. 1). Semithin sections were observed at light microscopy after different staining and cytochemical procedures (Figs. 2-4): *i*) toluidine blue for general structural analysis; *ii*) iodide-based staining to reveal starch and *iii*) DAPI for DNA. The observations revealed common and differential features in the structural organization of cells from different cultures assayed. Scanning electron microscopy observations are also shown in Fig. 5.

Cells of the three types of *in vitro* cultures studied, with different copper concentrations and times (non-treated, Cu-stressed and Cu-acclimated), shared some main structural features and exhibited several differences, which characterized their structural organization. Non-treated cells (Fig. 2, 5a), Cu-stressed cells (Fig. 3, 5b) and Cu-acclimated cells (Fig. 4, 5c) showed rounded shape, most of them with a large cytoplasmic vacuole occupying a high proportion (ca. 50-60%) of the cellular volume; the rest of the cytoplasm appeared as a thin peripheral layer containing numerous chloroplasts. Cells grown under the three culture conditions assayed were found either isolated or forming spherical structures with two cells; the two-cell structures were divided by a straight cell wall located in a lateral position, probably originated by the asymmetric division of individual isolated cells (Figs. 2a, b, f-h; 3a; 4a, b). The

nucleus was observed in a peripheral location exhibiting a patent and circular-shaped nucleolus (Figs. 2b, f-h; 3b, c; 4b, c), and a pattern of chromatin mostly decondensed, with several small patches of heterochromatin in linear arrays along to the nuclear envelope, as clearly seen by DAPI staining (Fig. 2c). Nuclear bodies were frequently observed in the cell nuclei of the photosynthetic cultures as small rounded structures of medium contrast in toluidine-blue stained preparations (arrows in Figs. 2f, h, and open arrow in Fig. 3b). Their size, localization and presence were similar to that of Cajal bodies, ribonucleoprotein structures found in most eukaryotic cells and also described in plants (Risueño and Medina, 1986; Beven et al. 1995). They are typical structures of metabolically-active and proliferating plant cells (Beven et al. 1995, Testillano et al. 2005, Seguí-Simarro et al. 2006). Morphometric analysis revealed a similar cellular size in non-treated, Cu-stressed and Cu-acclimated cells, with no statistically significant differences (Table 1).

The main differences observed among the cellular types of the three cultures affected chloroplasts, starch granules, vacuoles and extracellular deposits. Chloroplasts of non-treated cells were ellipsoids (Fig. 2a, b, f-h) and most of them contained large starch deposits, as revealed by the iodide-based cytochemistry (Fig. 2d, e); the starch granules were clearly observed at LTSEM occupying large volumes in the chloroplast (Fig. 5a). In contrast, chloroplasts of Cu-stressed and Cu-acclimated cells lost the ellipsoidal shape and displayed rounded shape (Figs. 3b, c; 4b, c; 5b, c). The chloroplast size in these cells was on average 2-fold and 3-fold smaller, respectively, compared with those of non-treated cells (Table 1). Moreover, chloroplasts of Cu-acclimated cells appeared more numerous with denser structured internal membrane compared with non-treated cells (Fig. 4b,c). These observations are in agreement with those

previously reported by Bernal et al. (2006). No starch granules could be found after iodide cytochemistry in copper treated cells (Figs. 3d, e; 4d, e). The cells of Cu-acclimated cultures showed larger cytoplasmic vacuoles than the non-treated cells and Cu-stressed cells, exhibiting a vacuole size 34% larger (Figs. 4a-c; 5c; Table 1). In Cu-stressed cells, numerous dark and small rounded deposits were observed attached at the outer surface of the cell wall (Fig. 3a, b, c, f-h). The cells of the other cultures did not show similar extracellular features (Figs. 2, 4). EDX microanalyses revealed a high copper content in these attached dark deposits (Fig. 6b). Since low molecular weight metal complexes could be organic acid-metal chelates (Kishinami and Widholm 1987), the levels of citrate and malate were determined in the three types of *in vitro* cultures studied (Table 2). In Cu-acclimated cells, both citrate and malate content increased to levels 2-3 fold and 3-4 fold higher, respectively, compared with non-treated cells. Interestingly, in Cu-stressed cells citrate increased to levels 3-4 fold those of non-treated cells and remained or slightly decreased in Cu-acclimated cells.

DISCUSSION

In the present work we compare the effect of 10 μM CuSO_4 on the functional cellular structure of soybean cell cultures after different copper exposure times. This study provided information about which Cu-induced effects on soybean cell structure accompany a short-term and a long-term response associated to adaptive processes. Cells accumulated high copper concentration when they were grown in medium supplemented with 10 μM CuSO_4 with no toxicity symptoms, in agreement with previous results (Bernal et al. 2006). Tolerance to increased concentrations of copper has been induced in the laboratory in several strains of the algae *Scenedesmus* and *Chlorella* (Küpper et al. 2003). The green alga *Scenedesmus quadricauda* was capable of developing a reversible (*i.e.* not genetically fixed) high resistance to copper. The results reported here show that the general cell organization pattern of the non-treated soybean cultures is maintained by the cells after the copper treatments. Moreover, cell features typical of proliferating cells were found not only in the non-treated cultures but also in both short- and long-term treated cultures such as nuclear Cajal-like bodies (Beven et al. 1995, Testillano et al. 2005, Seguí-Simarro et al. 2006) or straight thin cell walls dividing spherical two-cell structures (Satpute et al. 2005). The results also showed that the copper exposure induced changes in specific subcellular structures. The main target of copper effect was the chloroplast compartment. Indeed, smaller chloroplasts with a rounded shape were observed in soybean cells grown in the presence of 10 μM CuSO_4 compared with non-treated ones. This modification was observed either in Cu-stressed or Cu-acclimated cells indicating that this event is a fast Cu-induced response in soybean cell cultures. This finding contrasts with certain observations reported in the literature. For instance, no

significant alterations in chloroplast size and shape were found in 100 μM Cu-tolerant cultures of *Nicotiana tabacum* (Gori et al. 1998) and mesophyll cells from *Triticum durum* seeds exposed to 10-50 μM Cu (Ciscato et al. 1997; Quartacci et al. 2000). Chloroplasts of seven-week-old *Arabidopsis thaliana* plants exposed to 50 μM Cu during 2-14 days showed similar size that the control ones but rather circular than ellipsoidal shape (Wójcik and Tukiendorf 2003). Different chloroplast response has been found by exposure to other metals, i.e., cadmium (Cd) stress increased the chloroplast size in *Myriophyllum spicatum* (Stoyanova and Tchakalova 1997) and *Raphanus sativus* (radish) (Vitória et al. 2004). Our results also showed that chloroplasts in Cu-acclimated cells were more numerous in agreement with Bernal et al. (2006). This higher chloroplast number could maintain a constant plastid to mesophyll cell volume, which could compensate for the reduced chloroplast size. Recently, it has been reported that *Arabidopsis thaliana* mutants altered in the accumulation and replication of chloroplasts (*arc* mutants) present reduced chloroplast number and increased plastid size due to a compensatory mechanism (Austin II and Webber 2005).

After copper exposure, starch accumulation in chloroplasts was especially affected either in Cu-stressed or in Cu-acclimated cells. No accumulation of starch was observed in those cells. This finding contrasts with the significant increase in starch observed in seven-week-old *Arabidopsis thaliana* and cucumber plants exposed to excess copper (100 μM and 20 $\mu\text{g g}^{-1}$, respectively) (Wójcik and Tukiendorf 2003; Alaoui-Sossé et al. 2004). Similarly, cadmium or nickel stress was accompanied by an increase in carbohydrate contents in rice seedlings (Moya et al. 1993). Such discrepancies could be due to differences in starch metabolism between plants and cell cultures. The

relationship between starch accumulation in leaves and chloroplasts from stressed plants and inhibition of photosynthesis is well documented (Foyer 1988; Moya et al. 1993). Starch accumulation may result from a fall in the rate of utilisation of assimilates in the sink organs. Decrease in chlorophyll and sugars, and increase in starch was observed in the medicinal plant *Phyllanthus amarus* exposed to cadmium stress (Rai et al. 2005). On the contrary, the absence of starch accumulation may be attributed to a faster metabolism of assimilates. In this respect, previous results (Bernal et al. 2006) demonstrated that Cu-acclimated soybean cell suspensions have a higher growth rate as consequence of additional cell division. Starch consuming could be also required for an enhanced antioxidant activity. It has been reported that the induction of glucose-6-phosphate dehydrogenase (G6PDH) could contribute to antioxidant activity by providing more NADPH, which is required for the detoxification of ROS and peroxides (Kuo and Tang 1998; Ali et al. 2006). In this respect we observed no significantly differences in the ascorbate peroxidase (APX) and total superoxide dismutase (SOD) enzyme activities between copper-treated and non-treated cells although Cu/Zn SOD activity was stimulated in cells exposed to excess copper (unpublished data). On the other hand, the photosynthetic oxygen-evolution activity, which generates ROS and peroxides, was stimulated in Cu-acclimated cells in comparison with non-treated cells (Bernal et al. 2006) but this was not the case in Cu-stressed cells. Thus, a clear direct correlation between Cu-induced effect on starch reduction, antioxidant activity and photosynthesis cannot be established in these cell cultures. Interestingly, a correlation between the size of chloroplasts and the content of starch granules could be established. It is worth mentioning that starch accumulation strongly decreased in both Cu-stressed and Cu-acclimated

cells and this event accompanied the reduction of chloroplast size in those cells. These phenomena were independent of the copper exposure time and they were rather a short-term response than a long-term response as consequence of the excess copper adaptive period. Recently, it has been probed that manipulation of *FtsZ* expression levels, a key structural component of the chloroplast division machinery, can result in a modified size distribution of starch granules (De Parter et al. 2006). Further investigations are necessary to understand starch metabolism in chloroplasts of soybean cells and its relation with chloroplast division and structure.

The strategies for avoiding heavy metal toxicity are diverse. A first barrier against copper stress can be the immobilization of copper by means of cell wall and extracellular compounds such as organic acids (citrate, malate, oxalate), carbohydrates, proteins or peptides enriched in cysteine or histidyl groups. However, the importance of these mechanisms may vary in accordance with the concentration of metal supplied, the plant species involved and the exposure time. Dark deposits attached at the outer surface of the cell wall containing high level of copper were observed in Cu-stressed cells. Similar deposits have been observed in plants grown under metal stress conditions (Vitória et al. 2006). Interestingly, those dark deposits were not observed in Cu-acclimated cells indicating that their presence is really a short-term response, which disappears with a longer time exposure. Extracellular chelation by organic acids such as citrate and malate is a mechanism of metal tolerance (Rauser 1999). The secretion of organic acids is reasonably well established as a mechanism for aluminium tolerance (Kochian et al. 2004, Mariano et al. 2005), but there is very little number of studies in support of organic acid mechanism for copper tolerance. In this work we show that the accumulation of higher levels of citrate

and malate takes place in Cu-acclimated cells. Similar levels of citrate and malate in copper tolerant *Nicotiana plumbaginifolia* cells have been reported (Kishinami and Widholm 1987). It is worth mentioning that citrate increased in Cu-stressed cells and remained or slightly decreased in Cu-acclimated ones. This fact could indicate that citrate synthesis is preferentially stimulated during the first time of copper exposure in these soybean cells, being one of the fastest responses to copper exposure. Two organic acids exudation responses differing in time have been observed in roots of aluminium resistant plants (for review see Mariano et al. 2005). In the former response organic acids release is rapidly activated after aluminium exposure and the rate of release remains constant with time. In this case it has been suggested that aluminium activates a constitutive mechanism of organic acids transport in the plasma membrane and the activation of genes is not necessary. Aluminium can activate anion channels, which have been proposed as the mediators of organic acids transport across the cell membranes. In the second one there is a delay in the organic acids release after the addition of aluminium and this release increases with time. In this case the activation of genes related to the metabolism and membrane transport of organic acids might be required. According to that, the rapid induction of citrate in Cu-stressed cells might correspond to a mechanism similar to that taking place in the former response. On the other hand, the slower induction of malate observed in Cu-acclimated cells might rather correspond to the second mechanism mentioned above. In this point it is clear that further experiments varying time of copper exposure are necessary to define and understand the mechanisms involved in organic acid respond to excess copper in these cells.

Vacuole was also affected by copper treatments. Accumulation of metals in vacuoles is known as a detoxification mechanism to maintain lower metal concentrations in other subcellular compartments (chloroplast, cytoplasm, nucleus). The results showed that the cytoplasmic vacuole area increased in Cu-acclimated cells in comparison with non-treated and Cu-stressed ones. This fact can be associated with an accumulation of copper within the vacuole. Indeed, previous results showed a higher level of copper in vacuoles of Cu-acclimated cells compared with those of non-treated ones (Bernal et al. 2006). Furthermore, the analyses did not reveal the copper accumulation in vacuoles of Cu-stressed cells (data not shown). Vacuole modifications can be associated to dynamic conversion between lytic and storage functions (Murphy et al. 2005). Our observations could indicate that in the acclimated state the detoxification relies on compartmentation through active copper pumping into the vacuole rather than on complexation by ligands, or exclusion by an active efflux or reduced uptake. The results are in agreement with a picture where the first response of soybean cells to excess copper exposure avoids its entrance to the cell by its immobilization in the cell wall, probably through citrate copper chelates among other mechanisms. Then, after an adaptive long-term exposure, the Cu-acclimation would be mainly due to vacuolar sequestration. Further research is underway to determine whether the copper tolerance developed by the soybean cell cultures is really a stable character. As preliminary result we found that the suppression of excess copper in the growth medium of Cu-acclimated cells during four subsequent transfers did not alter the copper content within the cells and their oxygen evolution-activity, which was different to that of non-treated one (data not shown).

In summary our results indicate that the structural cell modifications, mainly affecting chloroplasts and starch, accompany a short-term Cu-induced response and they are maintained as a stable character during an adaptive period to those conditions, whereas vacuole changes rather accompany a long-term response.

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FIGURE LEGENDS

FIGURE 1: Cu content in soybean photosynthetic cells during 21 days of growth. Control (non-treated) cells (●); Cu-stressed cells (▲); Cu-acclimated cells (■). For details see Materials and Methods.

FIGURE 2: Structural organization of soybean photosynthetic cells of non-treated cultures. Histoiresin semithin sections after toluidine blue staining (a, b, f-h), DAPI staining for DNA (c) and I₂IK staining for starch (d, e). **a)** Panoramic view of the cells in the culture, isolated and in two-cell groups. **b)** Cell structure observed at higher magnification, cells show large vacuoles (v), perypheric nucleus (N) with one nucleolus each, cytoplasm (ct) with ellipsoid chloroplasts, many of them containing clear starch deposits (arrows). **c)** Chromatin pattern showed by DAPI staining: nuclei exhibit an homogeneous medium-bright fluorescence and several brighter small spots aligned at the nuclear periphery, corresponding to condensed chromatin patches. The nucleoli appear as dark areas (arrows). **d, e)** Starch grains revealed as dark inclusions by iodide-base cytochemistry, observed under bright field (d) and phase contrast (e). **f-h)** Several two-cell structures showing different asymmetric position of the dividing cell wall and some details of the cell organization. In some nuclei, Cajal-like bodies (arrows) can be observed. Bars represent 10 μm.

FIGURE 3: Structural organization of soybean photosynthetic cells of Cu-stressed cultures. Histoiresin semithin sections after toluidine blue staining (a, b, c, f-h), and I₂IK staining for starch (d, e). **a)** Panoramic view of the cells in the culture, isolated and in two-cell groups, showing high vacuolation. **b, c)** Cell

structure observed at higher magnification: the chloroplasts (arrows) appear dense and rounded, with no clear areas inside. The nuclei exhibit small dark nucleoli each and some of them Cajal-like bodies (open arrow). Extracellular dark deposits (arrowheads) are observed in contact with the outer face of the cell walls in many cells. **d, e)** Iodide-based cytochemistry revealed no presence of starch deposits in the chloroplasts, observed under bright field (d) and phase contrast (e). **f-h)** Extracellular deposits of different sizes and shapes (rounded and ellipsoid) are found on the cell walls. Bars in a, b, c, d, e represent 10 μm ; bars in f, g, h represent 5 μm .

FIGURE 4: Structural organization of soybean photosynthetic cells of Cu-acclimated cultures. Historesin semithin sections after toluidine blue staining (a, b, c), and I_2/IK staining for starch (d, e). **a)** Panoramic view of the culture, cells appear isolated and in two-cell groups with a cell wall in asymmetric position. **b, c)** Higher magnification of the cell organization details: the cytoplasmic vacuole (v) appears very large in most cells, the peripheral layer of cytoplasm is thin and contains rounded and dense chloroplasts (arrows), with no clear areas inside. **d, e)** Iodide-based cytochemistry for starch does not provide any contrast under bright field (d) revealing no presence of starch deposits in the chloroplasts which are observed under phase contrast (e). Bars represent 10 μm .

FIGURE 5: Low Temperature Scanning Electron micrographs of soybean photosynthetic cells of non-treated (a), Cu-stressed (b) and Cu-acclimated (c) cultures. Etching of frozen native samples. The aqueous content of the vacuole (v) appears forming a reticulum due to the freezing. In the peripheral

layer of cytoplasm (ct) the chloroplasts are observed as spherical structures, in non-treated cultures, most of them contain inner large structures, starch grains (arrows in a). Bars represent 5 μm .

FIGURE 6.- Energy dispersive X-ray spectra of extracellular dark deposits attached at the cell wall in Cu-stressed soybean photosynthetic cells. a) cell wall of non-treated cells; **b)** extracellular dark deposits. Typically, 6 cells of each type were analysed. Y-axis scale for b) was 4-fold.

TABLE 1. Morphometric analysis of cell, chloroplast and vacuole size from soybean cell cultures grown in control media (Non-treated), media supplemented with 10 μ M CuSO₄ during 21 days of growth (Cu-stressed) and after 22 transfers (Cu-acclimated).

Cell cultures	Cell ¹	Chloroplast ¹	Vacuole ¹
Non-treated	209.6 ^a $\pm$ 34.8	7.5 ^a $\pm$ 1.2	103.8 ^a $\pm$ 22.2
Cu-stressed	185.4 ^a $\pm$ 26.2	3.7 ^b $\pm$ 0.7	111.4 ^a $\pm$ 17.4
Cu-acclimated	220.6 ^a $\pm$ 38.0	2.3 ^c $\pm$ 0.5	138.0 ^b $\pm$ 25.4

¹ area in μm^2

^{a,b,c} For the same column, different letters indicate significant differences at $p \leq 0.05$. Data were obtained on average from 15 images for each condition.

TABLE 2. Organic acid content of cells from soybean cell cultures grown in control media (Non-treated), media supplemented with 10 μ M CuSO₄ during 21 days of growth (Cu-stressed) and after 22 transfers (Cu-acclimated).

Cell cultures	Citrate ¹	Malate ¹
Non-treated	2.4 ^a $\pm$ 0.2	0.3 ^a $\pm$ 0.1
Cu-stressed	7.6 ^b $\pm$ 0.3	n.d.
Cu-acclimated	5.9 ^c $\pm$ 0.2	1.1 ^b $\pm$ 0.3

¹ μ moles g⁻¹ fresh weight

^{a,b,c} For the same column, different letters indicate significant differences at $p \leq 0.05$. Data were obtained on average from 15 images for each condition.

Fig. 1

Fig. 2

Fig. 3

Fig. 4

Fig. 6

Fig. 5