



# A vacuolar H<sup>+</sup>-pyrophosphatase differential activation and energy coupling integrate the responses of weeds and crops to drought stress



Josimara Barcelos Venancio <sup>a,b,1</sup>, Michelle Guedes Catunda <sup>c,1</sup>, Juarez Ogliari <sup>d</sup>, Janaína Aparecida Hottz Rima <sup>a,b</sup>, Anna Lvovna Okorokova-Facanha <sup>e</sup>, Lev Alexandrovitch Okorokov <sup>e</sup>, Arnaldo Rocha Facanha <sup>a,b,\*</sup>

<sup>a</sup> Laboratório de Biologia Celular e Tecidual, Centro de Biociências e Biotecnologia, Universidade Estadual do Norte Fluminense Darcy Ribeiro, Campos dos Goytacazes, RJ, Brazil

<sup>b</sup> Núcleo de Desenvolvimento de Insumos Biológicos para Agricultura (NUDIBA), Universidade Estadual do Norte Fluminense Darcy Ribeiro, Campos dos Goytacazes, RJ, Brazil

<sup>c</sup> Instituto Federal Fluminense, Macaé, RJ, Brazil

<sup>d</sup> Instituto Federal Fluminense, Bom Jesus do Itabapoana, RJ, Brazil

<sup>e</sup> Laboratório de Fisiologia e Bioquímica de Microorganismos, Centro de Biociências e Biotecnologia, Universidade Estadual do Norte Fluminense Darcy Ribeiro, Campos dos Goytacazes, RJ, Brazil

## ARTICLE INFO

### Article history:

Received 3 July 2013

Received in revised form 4 December 2013

Accepted 15 December 2013

Available online 21 December 2013

### Keywords:

Membrane-bound pyrophosphatase

V-type ATPase

Cytoplasmic pyrophosphate

Metabolic energy

Electrochemical gradient

Phosphate pool dynamics

## ABSTRACT

**Background:** *Cyperus rotundus* L. is a C4 weed of large vegetative and reproductive vigor endowed with competitive advantages over most crop species mainly under adverse environmental conditions. Vacuole functions are critical for the mechanisms of drought resistance, and here the modulation of the primary system of vacuolar ion transport is investigated during a transient water stress imposed to this weed and to C4 crop species (*Zea mays* L.).

**Methods:** The vacuolar H<sup>+</sup> pumps, the H<sup>+</sup>-ATPase and H<sup>+</sup>-PPase, expression, activities and the energy coupling were spectrophotometrically investigated as key elements in the differential drought-resistance mechanisms developed by weeds and crops.

**Results:** In *C. rotundus* tonoplasts, ATP hydrolysis was more sensitive to drought than its coupled H<sup>+</sup> transport, which was in turn at least 3-folds faster than that mediated by the H<sup>+</sup>-PPase. Its PP<sub>i</sub> hydrolysis was only slightly affected by severe water deficit, contrasting with the disruption induced in the PP<sub>i</sub>-dependent H<sup>+</sup>-gradient. This effect was antagonized by plant rehydration as the H<sup>+</sup>-PPase activity was highly stimulated, reassuming a coupled PP<sub>i</sub>-driven H<sup>+</sup> pumping. Maize tonoplasts exhibited 2–4 times lower hydrolytic activities than that of *C. rotundus*, but were able to overactivate specifically PP<sub>i</sub>-dependent H<sup>+</sup> pumping in response to stress relief, resulting in an enhanced H<sup>+</sup>-pumps coupling efficiency.

**Conclusion:** These results together with immunoanalysis revealed profiles consistent with pre- and post-translational changes occurring on the tonoplast H<sup>+</sup>-pumps, which differ between weeds and crops upon water deficit.

**General significance:** The evidences highlight an unusual modulation of the H<sup>+</sup>-PPase energy coupling as a key biochemical change related to environmental stresses adaptive capacity of plants.

© 2014 Elsevier B.V. All rights reserved.

## 1. Introduction

Purple nutsedge (*Cyperus rotundus* L.) has been considered by many authors the world's worst weed. A befitting designation for a species spread throughout in more countries (at least 92) than any other

weed, and which infests at least 52 different crops worldwide [1]. It grows in all types of soils and can survive the highest temperatures known in agriculture [2]. While it prefers moist soil, established nutsedge plants will thrive even under severe dry conditions. From the physiological point of view, these characteristics have been correlated with its high C4 metabolism efficiency, usually associated with the more invasive weeds. Indeed, C4 weeds have clear competitive advantage over C3 crop species under many field situations, especially in dry situations where stomatal conductance is low [3], and under high light intensities and temperatures [4]. Moreover, while weed and crop species differing in photosynthetic pathway (C<sub>4</sub> vs C<sub>3</sub>) are likely to respond differently to these factors, it is not clear at what extent weeds and crops sharing the same photosynthetic metabolism (C<sub>4</sub> vs C<sub>4</sub>) can differ in their stress tolerance-mechanisms [5]. Actually, nutsedge plants can also suppress other C<sub>4</sub> species growing in its vicinity, and efforts to establish the biochemical bases of such success

**Abbreviations:** ACMA, 9-amino-6-chloro-2-methoxyacridine; DTT, dl-dithiothreitol; EDTA, ethylenediamine tetraacetic acid; PMSF, phenylmethanesulfonyl fluoride; PVP-40, polyvinylpyrrolidone-40

<sup>\*</sup> In memoriam: Prof. Claudio A. Retamal, a deeply kind and generous colleague and an unforgettable friend.

<sup>\*</sup> Corresponding author at: Laboratório de Biologia Celular e Tecidual, Centro de Biociências e Biotecnologia, Universidade Estadual do Norte Fluminense Darcy Ribeiro, Campos dos Goytacazes, RJ, Brazil.

E-mail address: [arnaldo@uenf.br](mailto:arnaldo@uenf.br) (A.R. Facanha).

<sup>1</sup> Contributions of these authors merit co-first authorship.

in dominating natural and agricultural habitats have mainly been focused on its ability to produce allelopathic substances [6].

On the other hand, stressful levels of environmental factors such as water and nutrient availability influence weed/crop interactions directly and also may interfere with weed control [5]. Weeds are highly efficient in water and nutrient acquisition, and thus are very competitive affecting even the most efficient crops developed for the agriculture. Drought is one of the most important environmental stresses limiting the productivity of crops around the world, and the comprehension of the different drought tolerance mechanisms is very important to develop new strategies to cope with this problem [7,8].

Drought-tolerant plants maintain their turgor at low water potentials by increasing solute molecules in the cell [9], a phenomenon tightly regulated in order to maintain the cell homeostasis. Indeed, in many cell types, any perturbations of the osmotic balance rapidly activate mechanisms involved in cell volume recovery, and the first responses involve changes in ion fluxes across the plasma membrane and vacuoles, which rapidly result in water flow across the cell membranes and induce changes in cell volume, and vice versa [10,11]. The vacuolar  $H^+$ -ATPase (V-ATPase; EC 3.6.1.3) plays important roles in controlling the cytoplasmic pH and ion fluxes in plant cell, which are essential for cellular processes including plant adaptation to stressful growth conditions [12]. However, a second proton pump, a vacuolar  $H^+$ -pyrophosphatase (V-PPase; EC 3.6.1.1), coexists with V-ATPase in the tonoplast (vacuolar membrane). Several evidences indicate that these enzyme functions like a backup to V-ATPase at limited energetic conditions [e.g., 13,14]. Furthermore, transgenic overexpression of the gene encoding the V-PPase improves drought and salt tolerance in different species, for example, in *Arabidopsis* [15], tobacco [16], maize [17] and cotton [18,19].

In order to provide new insights on the physiological and biochemical roles of the tonoplast  $H^+$  pumps in the development of different drought tolerance mechanisms, we compared the capacities of one of the most invasive weeds (*C. rotundus* L.) and a globally important  $C_4$  crop species (*Zea mays* L.) to cope with water deficit. A differential activation and energy coupling were found for the V-PPase, which seems to be related to the differential adaptive responses of weeds and crops to drought stress at the same time that it provides further support to recent findings that have motivated a reconsideration of the current knowledge of the role of V-PPase in plant physiology [20,21].

## 2. Materials and methods

### 2.1. Plant material

*Cyperus rotundus* L. and *Zea mays* L. (variety UENF 506-6) were cultivated in greenhouse, in pots with a substrate mixture of 65% sand and 35% commercial soil. Cultivated (15-days-old) plants were irrigated daily and 20 days after germination drought was imposed by withholding irrigation for about 11 days, plant substrate was monitored for its water content, and samples processed for analysis when the substrate water reached 30 and 70% of deficit, and also 24 h after re-watering. The substrate water content was measured by the Gravimetric Technique, in which a soil sample is collected, accurately weighted, completely dried out in an oven, re-weighted and the soil moisture percentage is calculated from the weight loss. Ten pots were maintained upon uninterrupted irrigation as controls.

### 2.2. Preparation of tonoplast-enriched membrane vesicles

Tonoplast vesicles were isolated from *C. rotundus* pseudostems or maize mesocotyls using differential centrifugation essentially as described by Giannini and Briskin [22], with minor modifications. About 50 g of mesocotyl was homogenized using a mortar and pestle in 2 mL/g (fresh weight) of ice-cold buffer containing 10% (v/v) glycerol, 0.5% (v/v) PVP (PVP-40, 40 kD), 5 mM EDTA, 0.13% (w/v) BSA, and

0.1 M Tris-HCl buffer, pH 7.6. Just prior to use, 150 mM KCl, 3.3 mM DTT, and 1 mM PMSF were added to the buffer. The homogenate was strained through four layers of cheesecloth and centrifuged at 10,000  $\times g$  for 10 min. The supernatant was centrifuged once more at 10,000  $\times g$  for 10 min and then at 100,000  $\times g$  for 30 min. The pellet was resuspended in a small volume of ice-cold buffer containing 10 mM Tris-HCl, pH 7.6, 10% (v/v) glycerol, 1 mM DTT, and 1 mM EDTA. The suspension containing the mesocotyl vesicles was layered over a 25/46% (w/w) discontinuous sucrose gradient that contained, in addition to sucrose, 10 mM Tris-HCl buffer, pH 7.6, 1 mM DTT, and 1 mM EDTA. After centrifugation at 100,000  $\times g$  for 2 h in a swinging bucket, the membrane vesicles were collected at the interface between 25 or 46% sucrose. All preparative steps were performed at 4 °C. The vesicles were frozen under liquid  $N_2$  and stored at -70 °C until use. Protein concentrations were determined by the Bradford's method [23].

### 2.3. Substrate hydrolysis assay

ATPase and PPase activity was determined colorimetrically by measuring the release of  $P_i$  as described by Fiske and Subbarow [24]. The reaction media contained 50 mM Tris-HCl, pH 6.5 or 7.0, 3 mM  $MgSO_4$ , 100 mM KCl, 1 mM ATP or 0.1 mM  $PP_i$ . The reaction was started by addition of tonoplast protein (40  $\mu g/mL$ ) and stopped with ice-cold 5% (w/v) trichloroacetic acid after 30 min of incubation at 25 °C. The  $H^+$ -ATPase activity was measured with and without 100 mM  $KNO_3$  or 10 nM Bafilomycin and the nitrate/Bafilomycin-sensitive activity was attributed to the V-ATPase [25]. The PPase activity stimulated by 100 mM KCl was considered as the V-PPase [26]. V-PPase activity was calculated as half of the rate of  $P_i$  released from  $PP_i$  ( $\mu moles PP_i$  hydrolyzed per minute).

### 2.4. Proton pumping assay

ATP- and  $PP_i$ -dependent  $H^+$  transport across membranes was measured as the initial rate of fluorescence quenching of 9-amino-6-chloro-2-methoxyacridine (ACMA) at 25 °C using a fluorimeter (model F-3010, Hitachi, Tokyo) using a protocol adapted by Facanha and de Meis [27]. The excitation wavelength was set at 415 nm and the emission wavelength was set at 485 nm. The reaction medium contained 10 mM Tris-HCl, pH 7.0, 2  $\mu M$  ACMA, 5 mM  $MgCl_2$ , 100 mM KCl, 1 mM ATP or 0.1 mM  $PP_i$ . The reaction was initiated by the addition of 50  $\mu g/mL$  of membrane vesicles. The addition of 20 mM  $NH_4Cl$  was used to show a recovery of the fluorescence that indicated a collapse of the preliminarily formed  $H^+$  gradient.

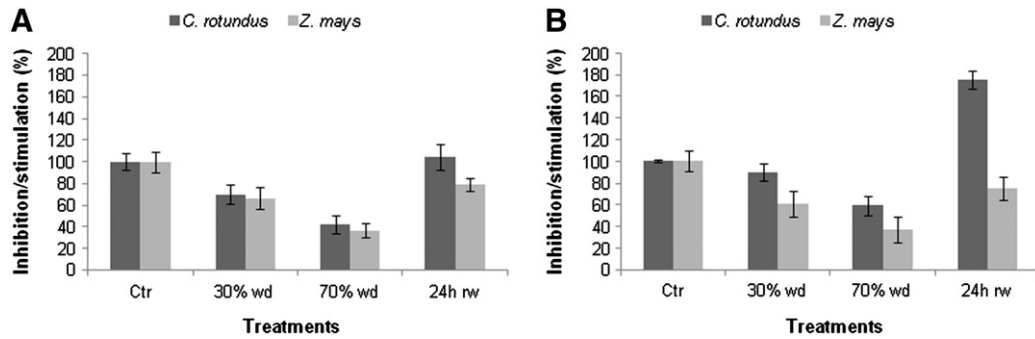
### 2.5. Western blot assays

Tonoplasts were isolated from *C. rotundus* pseudostems or maize mesocotyls. A sample of membrane proteins (30  $\mu g$ ), separated in a 12% SDS-PAGE was immunoblotted with antibodies against the subunit B of V- $H^+$ -ATPase or against  $H^+$ -PPase (Agrisera, Vännäs, SWEDEN). Cross-reacting proteins were revealed using peroxidase-conjugated secondary antibody (GE Healthcare Bio-Sciences Corp. Piscataway, NJ, USA) and the relative immunoresponse was estimated densitometrically as described by Retamal et al. [28].

## 3. Results

### 3.1. Changes in ATP and $PP_i$ hydrolysis

*C. rotundus* and *Z. mays* plants were submitted to drought stress and tonoplast vesicles were isolated by cell fractionation when the soil water reached 30 and 70% of deficit, and also after 24 h when the plants had been re-hydrated. Analysis of *C. rotundus* tonoplasts revealed that the ATP hydrolysis decreased progressively reaching 70% of inhibition upon the most severe stress (70% wd, Fig. 1A), while the PPase sustained high levels of  $PP_i$  hydrolysis being inhibited by only 30%



**Fig. 1.** V-ATPase (A) and V-PP<sub>i</sub>ase (B) hydrolytic activity in tonoplasts isolated from *C. rotundus* under increasing water deficit (wd) and 24 h after re-watering (rw). The reaction was started by addition of 1 mM ATP or 0.1 mM PP<sub>i</sub> at pH 7.0, and stopped after 30 min at 25 °C by addition of 5% TCA. Percentages were calculated from the specific enzymatic activities showed in Table 1. Values are means  $\pm$  SE of eight independent experiments.

under high dehydration (70% wd, Fig. 1B). Moreover, the PP<sub>i</sub> hydrolysis enhanced more than 2-fold in response to the relief of stress (24 h rw, Fig. 1B). Maize seedlings submitted to similar treatments were more sensitive to drought, with both ATP and PP<sub>i</sub> hydrolysis decreasing progressively during the stress, and just recovering the control levels 24 h after rewatering (Fig. 1). Furthermore, the rates of ATP and PP<sub>i</sub> hydrolysis were about 3-fold higher in tonoplast vesicles from *C. rotundus* plants (Table 1).

### 3.2. Proton pumping activities

Proton transport mediated by vacuolar H<sup>+</sup> pumps was analyzed using tonoplast vesicles isolated when the soil water reached 30 and 70% of deficit and 24 h after rewatering (Fig. 2). The H<sup>+</sup> pumping activities decreased along the stress, but this effect was more pronounced for the H<sup>+</sup>-PP<sub>i</sub>ase. The tonoplast H<sup>+</sup> gradient generated by *C. rotundus* V-ATPase was inhibited by ~60% upon the most severe stress (wd 70%, Fig. 2A), while no PP<sub>i</sub>-dependent H<sup>+</sup> pumping could be detected under this condition (Fig. 2B). Conversely, after re-watering, the initial rate of the PP<sub>i</sub>-dependent H<sup>+</sup> transport activity was stimulated by ~50% when compared to the controls (Fig. 2B), while the V-ATPase activity was only barely affected (Fig. 2A). The H<sup>+</sup> gradient generated by the maize V-ATPase was also only slightly modified by drought stress (Fig. 2C), while the maize V-PP<sub>i</sub>ase showed a comparable profile to that of *C. rotundus* with a complete inhibition of H<sup>+</sup> pumping during severe stress (wd 70%) and an overactivation of PP<sub>i</sub>ase after re-watering (Fig. 2D). However, for the maize V-PP<sub>i</sub>ase the activation occurred only in the H<sup>+</sup> gradient formation, resulting in more than 6-fold increase in the pump coupling efficiency (Table 2). On the other hand, for the *C. rotundus* pump it was observed in a higher stimulation of the PP<sub>i</sub> hydrolysis over that found in its related H<sup>+</sup> pumping capacity (Figs. 1B and 2D).

### 3.3. Western blot analysis of the tonoplast proton pumps

The expression of the vacuolar H<sup>+</sup> pumps in the tonoplast under drought stress and after re-watering was evaluated using western blot analysis. As shown in Fig. 3A and C, V-ATPase immunoreactivity in *C. rotundus* vacuolar membrane increased by ~20% at the initial stage of the stress (30% wd), and then, the enzyme level reduced as the soil

water deficit reached 70%. The protein content was recovered to the control levels after rehydrating (Fig. 3A and C). Interestingly, the level of H<sup>+</sup>-PP<sub>i</sub>ase in *C. rotundus* increased ~80% when submitted to initial stress (Fig. 3B and D), but the immunoreactivity was restored to the control level during severe stress (70% wd), and remained unaltered even after 24 h rehydration (Fig. 3B and D). In *Z. mays*, a lower sensitivity was observed at the V-ATPase level in response to stress and rehydration (Fig. 3A). Contrasting with the stimulatory effect found in *C. rotundus*, *Z. mays* H<sup>+</sup>-PP<sub>i</sub>ase immunoreactivity was reduced not only by 60% in the initial stress, but was also restored to the control level in the subsequent treatments (Fig. 3B).

## 4. Discussion

Plant distribution and abundance in natural environments and crop yield in agricultural systems are largely determined by water availability. Plants respond and adapt to water deficit at both the cellular and molecular levels, for instance by the accumulation of osmolytes and proteins specifically involved in stress tolerance. The plant vacuole is directly involved in this process as well as in ion signaling for drought responses [29]. In order to gauge whether the vacuolar H<sup>+</sup> pumps play a coordinated and active role on different mechanisms of plant drought adaptation, we compared the changes in the enzyme activities and expression occurring in *C. rotundus* L., a rustic stress-tolerant weed species, and in an agriculturally adapted crop variety (*Z. mays* L.).

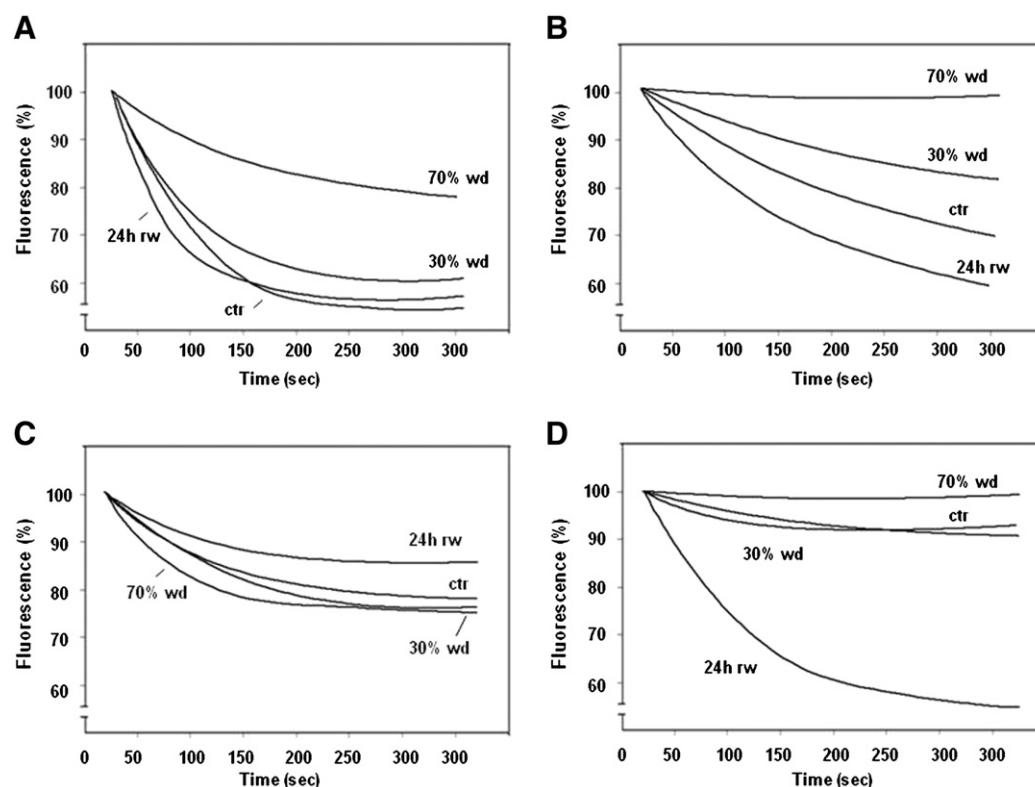
The ATP hydrolysis (Fig. 1A) as well as the ATP-dependent H<sup>+</sup> pumping activity (Fig. 2A) of *C. rotundus* V-ATPase were inhibited in the same range during the stress, and after rewatering both activities also recovered their control levels. A close profile was observed in immunodetection assay (Fig. 3A) suggesting that such behavior reflects a modulation of the pump expression in the vacuolar membrane. This hypothesis is in line with previous findings indicating that whereas salinity had a minor effect on the expression of V-ATPase genes in *Arabidopsis thaliana*, heat and drought stress led to alterations in transcript amount and isoforms of its subunits [30]. However, such regulation seems to be characteristic of species adapted to stress like *C. rotundus* and *A. thaliana* [31], rather than of more sensitive cultivated species, like maize and tomatoes. Actually, it was shown that the message level for the catalytic (70-kDa) subunit A of the V-ATPase did not vary in tomato under drought stress [32]. Here, also no significant change was found in immunodetection of the subunit B in maize tonoplasts, whereas for *C. rotundus*, a clear decrease occurred under severe stress and a rapid recovery upon rehydration was observed (Fig. 3A).

On the other hand, the H<sup>+</sup>-PP<sub>i</sub>ase hydrolytic activity of *C. rotundus* was barely affected by drought treatment (Fig. 1B), while its H<sup>+</sup> pumping was severely inhibited at this stress level (Fig. 2B). Upon 70% of water deficit no PP<sub>i</sub>-dependent H<sup>+</sup> pumping was detected, but at the same condition (Fig. 2B) the PP<sub>i</sub> hydrolysis was inhibited only by ~30% (Fig. 1B). This may indicate that the H<sup>+</sup>-PP<sub>i</sub>ase is assuming an uncoupled functional conformation, since a possible membrane leakage

**Table 1**

Comparison between the vacuolar proton pumps hydrolytic activities ( $\mu\text{mol} \cdot \text{mg}^{-1} \cdot \text{min}^{-1}$ ) of *C. rotundus* and *Z. mays* under water deficit (wd) and 24 h after rewatering (rw). Values are means  $\pm$  SE of eight independent experiments.

Treatments	ATP hydrolysis		PP <sub>i</sub> hydrolysis	
	<i>C. rotundus</i>	<i>Z. mays</i>	<i>C. rotundus</i>	<i>Z. mays</i>
Control	1.73 $\pm$ 0.03	0.64 $\pm$ 0.014	1.31 $\pm$ 0.23	0.56 $\pm$ 0.009
30% wd	1.10 $\pm$ 0.03	0.41 $\pm$ 0.009	1.04 $\pm$ 0.06	0.31 $\pm$ 0.007
70% wd	0.70 $\pm$ 0.09	0.26 $\pm$ 0.008	0.74 $\pm$ 0.06	0.17 $\pm$ 0.004
24 h rw	1.74 $\pm$ 0.11	0.50 $\pm$ 0.008	1.75 $\pm$ 0.12	0.45 $\pm$ 0.015



**Fig. 2.** Proton pumping activities of V-ATPase (A) and V-PPase (B) of tonoplast vesicles isolated from *C. rotundus* and V-ATPase (C) and V-PPase (D) from *Z. mays* seedlings irrigated daily (control, ctr), upon 30% or 70% of soil water deficit (30wd, 70wd), and 24 h after re-watering (rw). Proton translocation across membrane vesicles was monitored by the fluorescence quenching of ACMA in the presence of 50  $\mu$ g protein of membrane vesicles; the reaction was started by addition of 1 mM ATP or 0.1 mM  $PP_i$ . The data are representative of three independent experiments.

could be ruled out taking into account that a significant ATP-dependent  $H^+$  gradient was formed in the same tonoplast preparation (Fig. 2A). This possibility is reinforced by the data obtained 24 h after rewatering, where the  $PP_i$  hydrolysis enhanced  $\sim 110\%$ , while the initial velocity of  $PP_i$ -dependent  $H^+$  pumping increased only by  $\sim 60\%$ .

It seems likely that the  $H^+$ - $PP_i$ ases could operate mainly as phosphate-recycling enzymes during severe drought stress, by undergoing an unusual and still unexplored uncoupling regulation. This possibility is especially interesting from the energetic viewpoint, because  $H^+$ - $PP_i$ ases use a simple, low-cost substrate pyrophosphate that is generated as a byproduct of several biosynthetic processes for macromolecules, such as protein, RNA and polysaccharides. Moreover, a single gene encoding a protein of  $\sim 80$  kDa encodes the V- $PP_i$ ase and, theoretically, under metabolic stress, it should be easier and energetically safer to trigger its up-regulation than for the multimeric V-ATPase ( $\sim 700$  kDa). This is in line with the notion that for plants to react to their constantly changing environments and at the same time maintain

optimal metabolic conditions, the expression, activity and interplay of the  $H^+$  pumps have to be tightly regulated [33,34]. In this context, the present study also provides new insights on how transgenic plants that overexpress the  $H^+$ - $PP_i$ ases acquire a higher tolerance to  $P_i$  starvation [35], drought and salt stress [15,36–38] and capacity to nitrogen use efficiency [39]. Further studies are clearly required to better investigate the role of  $H^+$ - $PP_i$ ases as  $P_i$ -recycling enzyme active in  $P_i$  signaling, providing a complementary view on recent findings on phosphorus deficiency resulting in increased expression of the *AVP1*  $H^+$ - $PP_i$ ases and subsequent enhancement of the P-type plasma membrane  $H^+$ -ATPases [35].

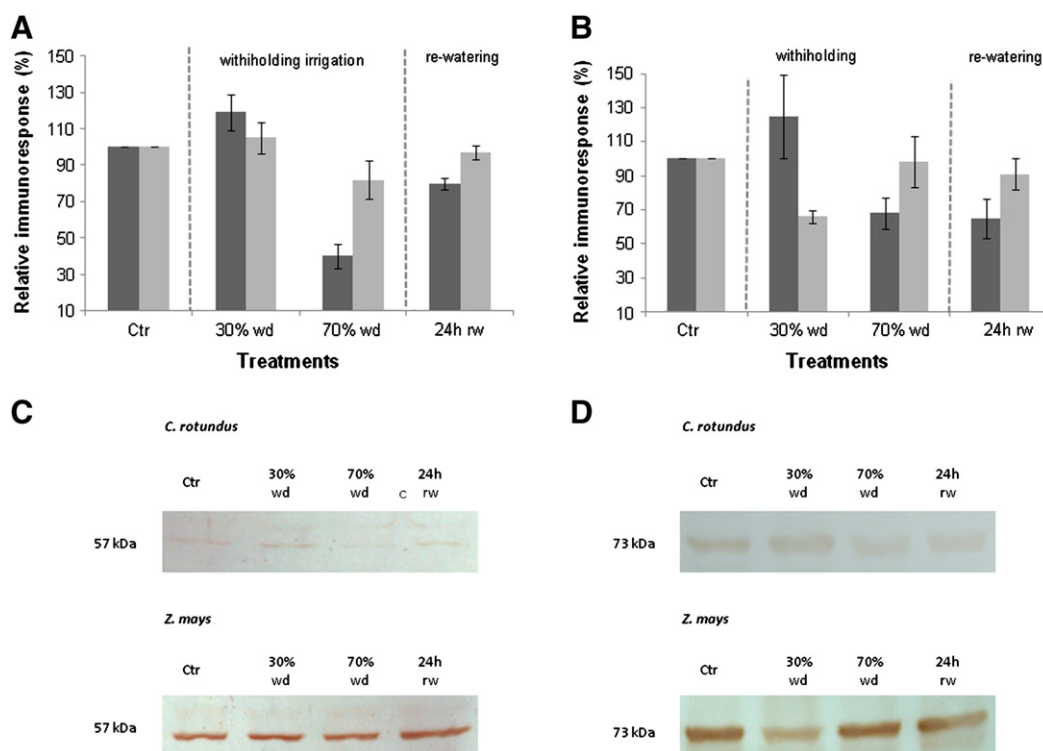
By overactivating their  $H^+$ - $PP_i$ ases, in the first drought stages and/or after the relief of stress, tolerant plant cells will decrease the ATP demand for maintenance of vacuolar electrochemical gradient, at the same time that the expression/activity of some uncoupled pumps will enhance  $P_i$  availability for the higher ATP synthesis and consume cytosolic  $PP_i$  required to overcome the constraints of transient stress exposures. Pyrophosphate is formed in a variety of biosynthetic reactions and needs to be promptly hydrolyzed in order to keep the overexpression of a myriad of stress responsive genes. This is also in line with a new emerging concept that in specific physiological conditions the role of  $H^+$ - $PP_i$ ase as a proton pump seems to be negligible [20,21]. These authors have provided very compelling evidences that during early seedling development, the major role of *Arabidopsis*  $H^+$ - $PP_i$ ase is the removal of the inhibitory metabolite  $PP_i$  rather than vacuole acidification. Previously, Hirono et al. [40] have also elegantly demonstrated, by random and site-directed mutagenesis, that discrete changes in specific amino acid residues dramatically affect the efficiency of  $H^+$  translocation and energy coupling of the  $H^+$ - $PP_i$ ase of *Streptomyces coelicolor*. Thus, it is tempting to speculate that there could exist physiological ways to modulate the functionality of related residues in the plant enzyme, in response to several environmental stresses.

**Table 2**

Comparison between the vacuolar proton pumps coupling ratio of *C. rotundus* and *Z. mays* under water deficit (wd) and 24 h after rewatering (rw). The  $H^+$  pumps' coupling was calculated by the difference between the initial rates of  $H^+$  pumping activity and the respective substrate hydrolysis. Values are means  $\pm$  SE of three independent experiments.

Treatments	Coupling ratio			
	$(V_0 H^+ \text{ pumping}/ATP \text{ hydrolysis})$		$(V_0 H^+ \text{ pumping}/PP_i \text{ hydrolysis})$	
	<i>C. rotundus</i>	<i>Z. mays</i>	<i>C. rotundus</i>	<i>Z. mays</i>
Control	21.6 $\pm$ 2.2	29.1 $\pm$ 1.7	11.6 $\pm$ 1.2	10.6 $\pm$ 1.4
30% wd	28.9 $\pm$ 3.7	43.4 $\pm$ 7.8	7.54 $\pm$ 1.8	34.8 $\pm$ 2.1
70% wd	19.4 $\pm$ 1.8	104.0 $\pm$ 15.9	–	–
24 h rw	22.2 $\pm$ 2.9	23.8 $\pm$ 1.4	13.3 $\pm$ 1.2	70.4 $\pm$ 6.2





**Fig. 3.** Immunoblot analysis of V-ATPase subunit B and H<sup>+</sup>-PPase under increasing water deficit (wd) and after re-watering (rw). Membranes were isolated from *C. rotundus* (dark bars) and *Z. mays* (light bars) plants as described in the [Material and methods](#) section, and submitted to immunoblot analysis. Western blots were used for quantification of the relative levels of the subunit B of V-ATPase (A and C) and H<sup>+</sup>-PPase (B and D). The immunoresponse of control plants was considered as 100% in each case. The data are representative of three independent experiments.

In regard to the issue of biochemical adaptive differences developed between weeds and crops, it is worth noting the much lower hydrolytic activities of maize proton pumps in comparison to that of the weed. It seems likely that a high performance of the proton pumps can be one of the features developed by plants evolving natural environments, with the tonoplast electrochemical gradient being mainly sustained by a resistant V-ATPase and the H<sup>+</sup>-PPase assuming an uncoupled conformation towards warrant of the Pi supply and the PPi cytoplasmic control. On the other hand, cultivated crops were selected presenting high performance in agricultural fields based at least in part on the stability of their V-ATPase and on the capacity to strongly enhance the H<sup>+</sup>-PPase coupling, which assumes the main role in the vacuolar energization as soon as a stress condition is overcome. Taken together these results argue in favor of a reconsideration of the biochemical and physiological roles of the vacuolar proton pumps, with emphasis on the impressive H<sup>+</sup>-PPase coupling plasticity and the respective relevance to the energy balance upon stress, not only for the vacuolar system but also for the cellular biosynthetic metabolism as a whole.

In conclusion, the lack of correlation between the PP<sub>i</sub> hydrolysis, H<sup>+</sup> pumping activity data and protein expression in the tonoplast suggests a complex regulation of the H<sup>+</sup>-PPases, which argues against a unique role as backup system to V-ATPases in maintaining the vacuolar H<sup>+</sup> gradient. From the physiological aspect, this study provides new perspectives to address the essential question of why H<sup>+</sup>-PPase coexists with H<sup>+</sup>-ATPase in the same vacuolar membrane. This work also proposes a new energetic adaptation characteristic of invasive weeds that enable them to disperse rapidly even into dry areas and outcompete crops and native vegetation.

## Acknowledgements

This study was supported by the Conselho Nacional de Desenvolvimento Científico e Tecnológico (CNPq) and the Fundação

Carlos Chagas Filho de Amparo à Pesquisa no Estado Rio de Janeiro (FAPERJ). JBV and JAH are fellows of Graduate Program of Plant Production at the Universidade Estadual do Norte Fluminense Darcy Ribeiro.

## References

- [1] L.G. Holm, D.L. Plucknett, J.V. Pancho, J.P. Herberger, *Cyperus rotundus* L, The World's Worst Weeds: Distribution and Biology, University Press of Hawaii, Honolulu, 1977, pp. 8–24.
- [2] K. Ueki, Studies on the control of nutsedge (*Cyperus rotundus*): on the germination of the tuber, Proceedings of the 2nd Asian-Pacific Weed Control Interchange, Asian-Pacific Weed Science Society, Los Banos, Philippines, 1969, pp. 355–370.
- [3] O. Björkman, Adaptive and genetic aspects of C4 photosynthesis, in: R.H. Burris, C.C. Black (Eds.), CO<sub>2</sub> Metabolism and Plant Productivity, Univ. Park Press, Baltimore, 1976, pp. 287–309.
- [4] G.E. Edwards, S.C. Huber, The C4 pathway, in: M.D. Hatch, N.K. Boardman (Eds.), The Biochemistry of Plants, Photosynthesis, Academic Press, New York, 1981, pp. 237–281.
- [5] D.T. Patterson, Effects of environmental stress on weed/crop interactions, Weed Sci. 43 (1995) 483–490.
- [6] R. Sharma, R. Gupta, *Cyperus rotundus* extract inhibits acetylcholinesterase activity from animal and plants as well as inhibits germination and seedling growth in wheat and tomato, Life Sci. 80 (2007) 2389–2392.
- [7] H.J. Bohnert, D.E. Nelson, R.G. Jensen, Adaptations to environmental stress, Plant Cell 7 (1995) 1099–1111.
- [8] D. Bartels, R. Sunkar, Drought and salt tolerance in plants, Crit. Rev. Plant Sci. 24 (2005) 23–58.
- [9] E.A. Bray, J. Bailey-Serres, E. Weretilnyk, Responses to abiotic stresses, in: W. Gruissem, B. Buchanan, R. Jones (Eds.), Biochemistry and Molecular Biology of Plants, American Society of Plant Physiologists, Rockville, 2000, pp. 1158–1249.
- [10] S.N. Shabala, R.R. Lew, Turgor regulation in osmotically stressed *Arabidopsis* epidermal root cells. Direct support for the role of inorganic ion uptake as revealed by concurrent flux and cell turgor measurements, Plant Physiol. 129 (2002) 290–299.
- [11] L. Zonia, T. Munnik, Life under pressure: hydrostatic pressure in cell growth and function, Trends Plant Sci. 12 (2007) 90–97.
- [12] K.J. Dietz, N. Tavakoli, C. Kluge, T. Mimura, S.S. Sharma, G.C. Harris, A.N. Chardonens, D. Goldack, Significance of the V-type ATPase for the adaptation to stressful growth conditions and its regulation on the molecular and biochemical level, J. Exp. Bot. 52 (2001) 1969–1980.
- [13] R.A. Leigh, A.J. Pope, I.R. Jennings, D. Sanders, Kinetics of the vacuolar H<sup>+</sup>-pyrophosphatase. The roles of magnesium, pyrophosphate, and their complexes as substrates, activators, and inhibitors, Plant Physiol. 100 (1992) 1698–1705.

- [14] G.D. Carystinos, H.R. MacDonald, A.F. Monroy, R.S. Dhindsa, R.J. Poole, Vacuolar  $H^+$ -translocating pyrophosphatase is induced by anoxia or chilling in seedlings of rice, *Plant Physiol.* 108 (1995) 641–649.
- [15] F. Brini, M. Hanin, I. Mezghani, G.A. Berkowitz, K. Masmoudi, Overexpression of wheat  $Na^+/H^+$  antiporter *TNAX1* and  $H^+$ -pyrophosphatase *TVP1* improve salt- and drought-stress tolerance in *Arabidopsis thaliana* plants, *J. Exp. Bot.* 58 (2007) 301–308.
- [16] F. Gao, Q. Gao, X.G. Duan, G.D. Yue, A.F. Yang, J.R. Zhang, Cloning of an  $H^+$ -PPase gene from *Thellungiella halophila* and its heterologous expression to improve tobacco salt tolerance, *J. Exp. Bot.* 57 (2006) 3259–3270.
- [17] B. Li, A. Wei, C. Song, N. Li, J.R. Zhang, Heterologous expression of the *TsVP* gene improves the drought resistance of maize, *Plant Biotechnol. J.* 6 (2008) 146–159.
- [18] S.L. Lv, L.J. Lian, P.L. Tao, Z.X. Li, K.W. Zhang, J.R. Zhang, Overexpression of *Thellungiella halophila*  $H^+$ -PPase (*TsVP*) in cotton enhances drought stress resistance of plants, *Planta* 229 (2009) 899–910.
- [19] V. Pasapula, G. Shen, S. Kuppu, J. Paez-Valencia, M. Mendoza, P. Hou, J. Chen, X. Qiu, L. Zhu, X. Zhang, D. Auld, E. Blumwald, H. Zhang, R. Gaxiola, P. Payton, Expression of an *Arabidopsis* vacuolar  $H^+$ -pyrophosphatase gene (*AVP1*) in cotton improves drought- and salt tolerance and increases fibre yield in the field condition, *Plant Biotechnol. J.* 9 (2011) 88–99.
- [20] A. Ferjani, S. Segami, G. Horiguchi, Y. Muto, M. Maeshima, H. Tsukayad, Keep an eye on *PPi*: the vacuolar-type  $H^+$ -pyrophosphatase regulates postgerminative development in *Arabidopsis*, *Plant Cell* 23 (2011) 2895–2908.
- [21] A. Ferjani, S. Segami, G. Horiguchi, A. Sakata, M. Maeshima, H. Tsukaya, Regulation of pyrophosphate levels by  $H^+$ -PPase is central for proper resumption of early plant development, *Plant Signal. Behav.* 7 (2012) 38–42.
- [22] J.L. Giannini, D.P. Briskin, Proton transport in plasma membrane and tonoplast vesicles from red beet (*Beta vulgaris* L.) storage tissue, *Plant Physiol.* 84 (1987) 613–618.
- [23] M.M. Bradford, A rapid and sensitive method for the quantitation of microgram quantities of protein utilizing the principle of the protein-dye binding, *Anal. Biochem.* 72 (1976) 248–254.
- [24] C.F. Fiske, Y. Subbarow, The colorimetric determination of phosphorus, *J. Biol. Chem.* 66 (1925) 375–400.
- [25] S.D. O'Neill, A.B. Bennet, R.M. Spanswick, Characterization of a  $NO_3^-$  sensitive  $H^+$ -ATPase from corn roots, *Plant Physiol.* 72 (1983) 837–846.
- [26] Y. Wang, R.A. Leigh, K.H. Kaestner, H. Sze, Electrogenic  $H^+$ -pumping pyrophosphatase in tonoplast vesicles of oat roots, *Plant Physiol.* 81 (1986) 497–502.
- [27] A.R. Facanha, L. de Meis, Reversibility of  $H^+$ -ATPase and  $H^+$ -pyrophosphatase in tonoplast vesicles from maize coleoptiles and seeds, *Plant Physiol.* 116 (1998) 1487–1495.
- [28] C.A. Retamal, P. Thiebaut, E.W. Alves, Protein purification from polyacrylamide gels by sonication extraction, *Anal. Biochem.* 268 (1999) 15–20.
- [29] H. Knight, A.J. Trewavas, M.R. Knight, Calcium signaling in *Arabidopsis thaliana* responding to drought and salinity, *Plant J.* 12 (1997) 1067–1078.
- [30] M. Hanitzsch, D. Schnitzer, T. Seidel, D. Gollack, K.J. Dietz, Transcript level regulation of the vacuolar  $H^+$ -ATPase subunit isoforms *VHA-a*, *VHA-E* and *VHA-G* in *Arabidopsis thaliana*, *Mol. Membr. Biol.* 24 (2007) 507–518.
- [31] N. Vartanian, L. Marcotte, J. Ciraudat, Drought rhizogenesis in *Arabidopsis thaliana*: differential responses of hormonal mutants, *Plant Physiol.* 104 (1994) 761–767.
- [32] M.L. Binzel, J.R. Dunlap, Absciscic acid does not mediate NaCl-induced accumulation of 70 kDa subunit tonoplast  $H^+$ -ATPase message in tomato, *Planta* 197 (1995) 563–568.
- [33] R.A. Gaxiola, M.G. Palmgren, K. Schumacher, Plant proton pumps, *FEBS Lett.* 581 (2007) 2204–2214.
- [34] D.B. Zandonadi, L.P. Canellas, A.R. Facanha, Indolacetic and humic acids induce lateral root development through a concerted plasmalemma and tonoplast  $H^+$  pumps activation, *Planta* 225 (2007) 1583–1595.
- [35] H. Yang, J. Knapp, P. Koirala, D. Rajagopal, W.A. Peer, L.K. Silbart, A. Murphy, R.A. Gaxiola, Enhanced phosphorus nutrition in monocots and dicots overexpressing a phosphorus-responsive type I  $H^+$ -pyrophosphatase, *Plant Biotechnol. J.* 5 (2007) 735–745.
- [36] R.A. Gaxiola, J. Li, S. Undurraga, L.M. Dang, G.J. Allen, S.L. Alper, Drought- and salt-tolerant plants result from overexpression of the *AVP1*  $H^+$ -pump, *Proc. Natl. Acad. Sci. USA* 98 (2001) 11444–11449.
- [37] S. Park, J. Li, J.K. Pittman, G.A. Berkowitz, H. Yang, S. Undurraga, J. Morris, K.D. Hirschi, R.A. Gaxiola, Up-regulation of an  $H^+$ -pyrophosphatase ( $H^+$ -PPase) as a strategy to engineer drought-resistant crop plants, *Proc. Natl. Acad. Sci. USA* 102 (2005) 18830–18835.
- [38] S. Guo, H. Yin, X. Zhang, F. Zhao, P. Li, S. Chen, Y. Zhao, H. Zhang, Molecular cloning and characterization of a vacuolar  $H^+$ -pyrophosphatase gene, *SsVP*, from the halophyte *Suaeda salsa* and its overexpression increases salt and drought tolerance of *Arabidopsis*, *Plant Mol. Biol.* 60 (2006) 41–50.
- [39] J. Paez-Valencia, J. Lares-Sanchez, E. Marsh, L.T. Dorneles, D. Sanchez, M.P. Santos, A. Winter, S. Murphy, J. Cox, M. Trzaska, J. Metler, A. Kozic, A.R. Facanha, D. Schachtman, C. Sanchez, R.A. Gaxiola, Enhanced  $H^+$ -PPase activity improves nitrogen use efficiency in Romaine lettuce (*Lactuca sativa* cv. conquistador), *Plant Physiol.* 161 (2013) 1557–1569.
- [40] M. Hirono, Y. Nakanishi, M. Maeshima, Essential amino acid residues in the central transmembrane domains and loops for energy coupling of *Streptomyces coelicolor* A3(2)  $H^+$ -pyrophosphatase, *Biochim. Biophys. Acta* 1767 (2007) 930–939.