

Effects of salt stress on low molecular antioxidants and redox state of plastoquinone and P700 in *Arabidopsis thaliana* (glycophyte) and *Eutrema salsugineum* (halophyte)

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Abstract

The effects of NaCl treatment were analysed in two species of considerably different resistance. In glycophyte, the content of ascorbate decreased but lipophilic antioxidants (α -tocopherol, plastochromanol, and hydroxy-plastochromanol) increased due to 150 mM NaCl. In halophyte, 300 mM NaCl caused a significant increase in hydrophilic antioxidants (ascorbate, total glutathione) but not in the lipophilic antioxidants. The redox states of plastoquinone (PQ) and P700 were also differently modulated by salinity in both species, as illustrated by an increased oxidation of these components in glycophyte. The presented data suggest that *E. salsugineum* was able to avoid a harmful singlet oxygen production at PSII, which might be, at least in part, attributed to the induction of the ascorbate-glutathione cycle. Another important cue of a high salinity resistance of this species might be the ability to sustain a highly reduced states of PQ pool and P700 under stress, which however, drastically affect the NADPH yield.

Additional key words: chloroplast; oxidative stress; photosystems; salinity.

Introduction

Salinity (in the general sense) causes disturbance in natural balance in plant cells, as well as, in the redox and energy status which affect cellular metabolism and result in less efficient photosynthesis (Takahashi and Murata 2008, Suzuki *et al.* 2012). Chloroplasts are particularly subjected to salt stress because of their sensitivity to external changes (Pfalz *et al.* 2012). Uptake of excessive sodium ions into cells disturbs the osmotic balance and water availability, which results in stomatal closure and decrease in the CO₂/O₂ ratio (Munns and Tester 2008). As a consequence, overreduction of photosynthetic electron transport (PET) occurs which leads to unbalanced ATP/NADPH ratio (Lokhande *et al.* 2011). Excess of electrons is transferred to molecular oxygen that is reduced to highly active species, namely, superoxide anion radical, H₂O₂, and

hydroxyl radical, which are produced mostly on the acceptor site of PSI (Mehler reaction), at PSII (at the Q_B site or *via* cytochrome *b*₅₅₉) or at the plastoquinone pool (PQ-pool) (Pospíšil 2009, Khorobrykh *et al.* 2015). Furthermore, energy transfer from triplet state of P₆₈₀ to molecular oxygen produces singlet oxygen in PSII. Although the leakage of electrons contributes to reactive oxygen species (ROS) production, it allows to control the redox poise of electron carriers (Noctor and Foyer 1998). In addition, a blockage of CO₂ flow to stomata also influences the Calvin cycle through depletion of 3-phosphoglyceric acid and NADP⁺ availability. The shortage of the latter contributes to ROS production, especially of H₂O₂, which is known to inhibit synthesis of D1 protein (Takahashi and Murata 2006, Murata *et al.* 2007).

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Abbreviations: NPQ – nonphotochemical quenching; PC-8 – plastochromanol-8; PET – photosynthetic electron transport; PQ – plastoquinone; ROS – reactive oxygen species; Toc – tocopherol; Y_I – the quantum efficiency of PSI, Y_{ND} – donor-side limitation to PSI; Y_{NA} – acceptor-side limitation to PSI.

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In order to minimize these negative effects and restore a disturbed homeostasis, plants activate short and long-term mechanisms in order to adjust the photosynthetic apparatus to new demands and activate acclimation responses (Scheibe *et al.* 2005, Dietzel *et al.* 2008). All these mechanisms, such as energy dissipation pathways of the PET chain (*i.e.* nonphotochemical quenching, state transitions, cyclic and pseudocyclic electron transport) or pathways involving cell compartments (*i.e.* malate valve, photorespiration or alternative oxidase pathway), allow to optimize electron flow and a required rate of the production of reducing equivalents for metabolic processes during stress conditions (Scheibe *et al.* 2005). Redox imbalance and subsequently ROS production is also sensed by the antioxidant systems. The enzymatic responses are mostly generated through the water-water cycle enzymes (superoxide dismutase, SOD, ascorbate peroxidase, APX, monodehydroascorbate reductase, MDHAR, dehydroascorbate reductase, DHAR, and glutathione reductase, GR) and redox regulatory enzymes and compounds, such as NADPH-dependent thioredoxin reductase (NTRC), 2-Cys peroxiredoxin (2-Cys Prx), thioredoxins or ferredoxin-thioredoxin reductase (Foyer and Noctor 2009). The second important target group are low molecular antioxidants, such as ascorbate, glutathione, prenyllipids, flavonoids, and carotenoids (Apel and Hirt 2004). Ascorbate and glutathione are redox buffers that have multiple functions in plants and serve as powerful antioxidants reacting directly with singlet oxygen, superoxide, and hydroxyl radical or as a reducing substrate for H₂O₂ detoxification (Foyer and Noctor 2011). In the latter case, it prevents overreduction of PET by regulating electron flow and production of photosynthetic assimilatory power and decrease the risk of singlet oxygen formation (Noctor and Foyer 1998). Ascorbate plays also an important role as a secondary antioxidant by reducing the oxidized form of α -tocopherol, thereby limiting lipid peroxidation in chloroplasts (Munné-Bosch 2005). Among prenyllipids antioxidants, the best known are tocopherols, plastochromanol, and plastoquinol (Munné-Bosch and Alegre 2002, Nowicka and Kruk 2010, Kruk *et al.* 2016). These lipophilic compounds are able to prevent lipid peroxidation or deactivate singlet oxygen (Munné-Bosch 2005, Gruszka *et al.* 2008, Nowicka and Kruk 2010).

Although ROS interact with many important cellular components, they function as signalling molecules (Noctor and Foyer 1998). Moreover, glutathione and ascorbate are not only essential to annihilate ROS but can also act as signal transducers initiating signalling cascades towards changes in nuclear gene expression (Baier and Dietz 2005, Noctor 2006, Scheibe *et al.* 2005). The main redox sensor in chloroplast is the PQ-pool (Pfannschmidt *et al.* 2009). This pool is composed of the fractions photosynthetically active (localized mainly in thylakoid membranes) and

inactive (present in plastoglobules). The inactive part, owing to its antioxidant properties, contributes to scavenging of singlet oxygen and superoxide anion radical (Szymańska and Kruk 2010). It has been shown that the amount and size of plastoglobules increase during salt stress (Hernandez *et al.* 1995). The redox state of the PQ-pool regulates many processes, such as state transitions, expression of PSII and PSI proteins, SOD, APX, and others (Pfannschmidt 2003, Ślesak *et al.* 2003, Chang *et al.* 2004, Puthiyaveetil *et al.* 2012).

One of many approaches to study salt defense mechanisms is the use of halophytic plants. Several studies pointed to a constitutive preparedness of these plants to salt conditions (Taji *et al.* 2004, Amtmann 2009, Dassanayake *et al.* 2011). In these plants, several protecting strategies are activated in order to stabilize photosystems, maintain the rate of photosynthesis, secure Calvin cycle enzymes, and modulate redox state of many components (Niewiadomska and Wiczarz 2015). Moreover, halophytes seem to avoid excessive ROS production by a more pronounced ability to activate and regulate antioxidant response. By contrast to glycophytes, this response is more intense and rapid, allowing to control ROS production at a certain threshold (Ellouzi *et al.* 2011, Ozgur *et al.* 2013, Bose *et al.* 2014). Likewise, in glycophytic *Arabidopsis thaliana*, a strong mobilization is observed at the level of stress hormones, transcription of stress genes, as well as photosynthetic performance, due to salinity treatment with 100–200 mM NaCl. Whilst, in glycophytic *Eutrema salsugineum*, a very weak change in these stress markers was found under a much higher salinity levels (Gong *et al.* 2005, Stepien and Johnson 2009, Wiczarz *et al.* 2015, Pilarska *et al.* 2016). Lokhande *et al.* (2011) suggested that changes in the redox and energetic status in halophytes accounts for their tolerance to salinity. However, in this type of analysis much attention is directed toward enzymatic response to salt treatments. Less emphasis is placed on low molecular antioxidants, although their role in a stress defence is crucial (Bose *et al.* 2014).

In our previous study, we have shown enhanced PSII electron transport rate in *Eutrema salsugineum* (*Thellungiella salsuginea*) which was accompanied by pronounced H₂O₂ production from the PQ-pool and increased chlororespiration (Wiczarz *et al.* 2015). A goal of this study was to verify hypothesis that a considerable difference between the halophyte and glycophyte in response to salinity stress, as previously reported, is related to the engagement of soluble nonenzymatic antioxidants. Considering the above-mentioned phenomenon of high H₂O₂ generation in chloroplasts of halophyte, we also aimed to characterize a lipophilic antioxidants, which are localised close to PSII and may modify the functioning of the overall photosynthetic electron transport.

Materials and methods

Plant material and growth conditions: *Arabidopsis thaliana* (Col-0) and *Eutrema salsugineum* (*Thellungiella salsuginea*) ecotype Shandong were used for all experiments. Seeds were obtained from the *Nottingham Arabidopsis Stock Centre*, UK. Growth conditions were as those described earlier (Wiciaz *et al.* 2015). After development of leaf rosette, four-week-old *A. thaliana* and *E. salsugineum* plants were daily irrigated with NaCl solutions (0.15 M for *A. thaliana* and 0.3 M for *E. salsugineum*), while watered plants served as controls. These two NaCl concentrations were chosen on basis of earlier reports describing stress responses of this two species under a broad salinity range (Vinocur and Altman 2005, Gong *et al.* 2005, Stepień and Johnson 2009). They revealed that a much stronger salinity treatment is needed to evoke stress effects in halophyte than in glycophyte. Chosen concentrations also correspond with our earlier characteristic of changes in chloroplast metabolism in these two species (Wiciaz *et al.* 2014, Pilarska *et al.* 2016). After 7 d of NaCl treatment, the plants were collected and used for further experiments.

Ascorbate and total glutathione in leaves: The ascorbate content was determined according to Takahama and Oniki (1992). Fresh leaves (2–3) were ground in a mortar with 10 ml of 6% (w/v) metaphosphoric acid solution with addition of 0.5 mM diethylenetriaminepentaacetic acid (DTPA) and at the proportion of 500 μ l of metaphosphoric acid solution for 50 mg of leaf fresh mass (FW). The homogenate was centrifuged at 4°C for 4 min at $10,000 \times g$. The supernatant was transferred to an Eppendorf tube and carefully neutralised with 2.5 M potassium carbonate. Ascorbate concentration in the probes was determined by test stripes (*Merckoquant*, *Merck KGaA*, Darmstadt, Germany) and reflectometers (*RQflex*, *Merck KGaA*, Darmstadt, Germany) according to the producer's instruction, and compared with ascorbate standards.

Total glutathione content was measured according to Luwe *et al.* (1993). The samples were prepared in the same way as for ascorbate measurements and the supernatants were used in a further procedure. The reaction mixture consisted of 100 mM potassium phosphate buffer, pH 7.5, containing 6.3 mM ethylenediaminetetraacetic acid (EDTA), 0.3 mM NADPH, and 6 mM 5,5'-dithiobis-2-nitrobenzoic acid (DTNB). The reaction was initiated by the addition of glutathione reductase (type III, 0.1 U μ l⁻¹, *Sigma-Aldrich*, Poznań, Poland) into the cuvette (1 ml) and the increase of absorbance at 412 nm (*DU-640*, *Beckman Instruments Inc.*, Brea, USA) was measured and compared with GSH standard. Control rates in the absence of the extract were subtracted.

Prenylipids content and the redox state of total plastoquinone: The concentration of tocopherols (α -Toc, PC-8), the total (oxidized plus reduced) plastoquinone

(PQ) and its redox state in leaves was determined according to Kruk and Karpinski (2006). In general, three to four leaves were homogenized in a cold mortar with 750 μ l of cold ethyl acetate, then next 750 μ l of ethyl acetate was added, and homogenization continued. The extract (400 μ l) was transferred to an Eppendorf tube and immediately evaporated to dryness under stream of nitrogen. Afterwards, the extract was dissolved in 200 μ l of methanol/hexane (340/20, v/v) solvent, shortly centrifuged, and 100 μ l of the extract was analysed by HPLC system, composed of *PU-980* pump and UV-VIS detector system *UV-970* (*Jasco Inc.*, Tokyo, Japan), and fluorescence detector *RF10-AXL* (*Shimadzu*, Kyoto, Japan). The concentration of PQ redox forms, as well as tocopherols, was determined from the area of respective peaks and were compared to standards. The measurements were performed using reversed-phase C₁₈ column (*Nucleosil 100*, 250 \times 4 mm, 5 μ m, *Teknokroma*, Barcelona, Spain) with the flow rate of 1.5 ml min⁻¹ and absorption detection at 255 nm for prenylquinones and fluorescence detection at λ_{ex} = 290 nm, λ_{em} = 330 nm for chromanols. PC-OH was determined as described previously (Szymańska *et al.* 2014).

Chlorophyll (Chl): The total Chl content was determined spectrophotometrically (*DU-640*, *Beckman Instruments Inc.*, Brea, USA) and calculated according to Lichtenthaler (1987). Ethyl acetate extract (400 μ l) was evaporated to dryness under stream of nitrogen and dissolved in 2 ml of methanol.

Nonphotochemical quenching and PSI parameters: Nonphotochemical quenching (NPQ) and PSI parameters were determined with *Dual PAM* (*Heinz Walz GmbH*, Effeltrich, Germany). Due to technical limitations, measurements were done on cut-off leaves dark-adapted for 15–20 min. To reach a steady state of electron transport an illumination with red actinic irradiation of 126 μ mol(quantum) m⁻² s⁻¹ for 5 min was used. NPQ and PSI parameters were calculated according to Bilger and Björkman (1990) and Klughammer and Schreiber (1994), respectively.

NADP-MDH activity and NADPH in leaves: The activity of chloroplast NADP-dependent malate dehydrogenase (NADP-MDH; EC 1.1.1.82) was assayed according to Scheibe and Stitt (1988) with minor modifications. The powdered leaves were homogenized in 0.1 M Tris-HCl buffer, pH 8.0, containing 1 mM EDTA, 5 mM DTT, and 0.5% PVP-40. Homogenates were centrifuged for 5 min at $13,300 \times g$ and the supernatant was used for measurements. Extracts of soluble proteins were incubated for 15 min in 0.1 M Tris-HCl buffer, pH 8, containing 1 mM EDTA, 20 mM DTT, and 0.2 mM NADPH. Decrease of absorbance at 340 nm (*DU-640*, *Beckman*

Instruments Inc., Brea, USA) was followed for 7 min after starting the reaction by addition of 1 mM oxaloacetate. Activity of NADP-MDH was expressed in nmol of consumed NADPH per mg protein per min.

The concentration of NADPH was measured by *Amplite Colorimetric NADP/NADPH Assay Kit* (AAT Bioquest Inc., Sunnyvale, USA) according to manufacturer's instructions. Fifty mg of leaves were ground in a mortar with 1 ml of PBS buffer, centrifuged in a cold benchtop for 5 min at $10,000 \times g$ and stored on ice. The

supernatant was incubated for 45 min in the reaction mixture at room temperature, then absorbance was measured at 575 nm (*DU-640*, Beckman Instruments Inc., Brea, USA) and compared with NADPH standard.

Statistical analysis: Significances between the means were analysed with the *Student's t*-test. Significant differences are marked at $P \leq 0.05$. Calculations were made with *MS Excel 2013*.

Results

Effect of moderate salinity on low molecular antioxidants: To have an insight into the antioxidant response to mild salinity treatment, we compared low molecular antioxidants, such as ascorbate, glutathione, α -Toc, and PC-8. The analysis of hydrophilic antioxidants (Fig. 1A,B) involved in the ascorbate-glutathione cycle revealed that in *E. salsugineum*, a significant increase of ascorbate and total glutathione concentration were found after salinity. In

turn, concentration of ascorbate decreased in *A. thaliana*, while that of glutathione remained unaffected. In the case of hydrophobic antioxidants, an opposite effect was observed (Fig. 1C–E). In leaves of *E. salsugineum*, the concentrations of α -Toc and PC-8 were not affected significantly after salt treatment. In contrast, in *A. thaliana* the content of these compounds increased by 50 and 60% after NaCl treatment, respectively. In the control

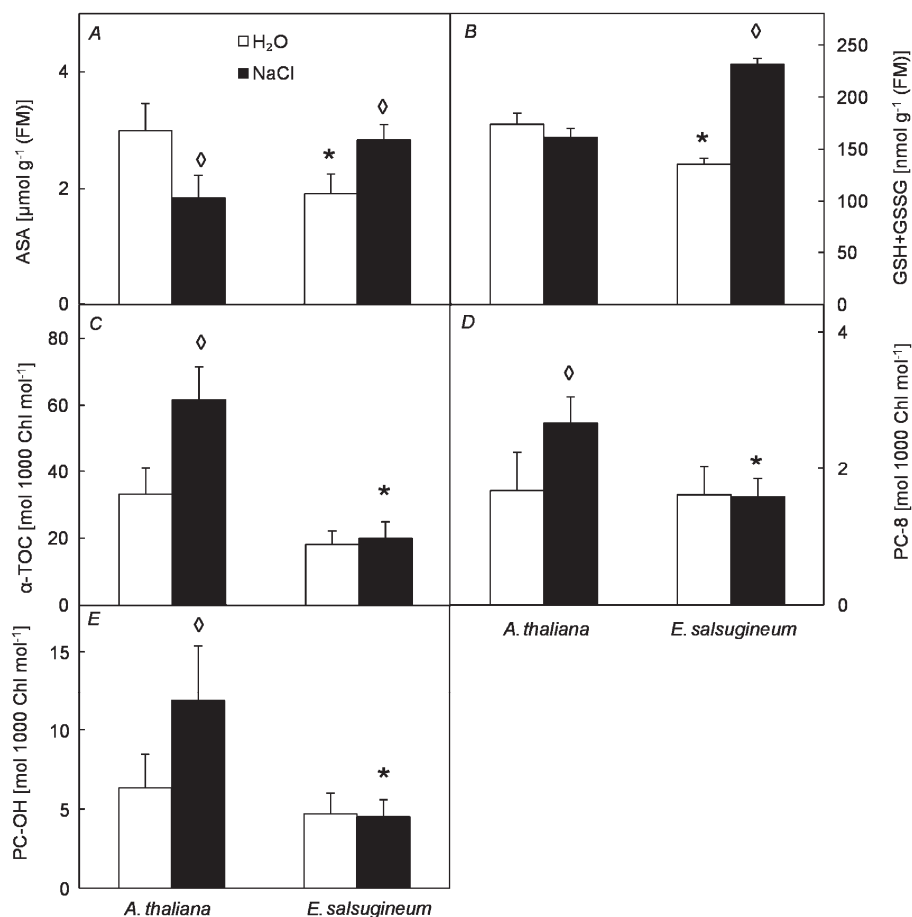


Fig. 1. The content of ascorbate (A), total glutathione content (B), the content of α -tocopherol (α -Toc; C), plastochromanol (PC-8; D), and hydroxyplastochromanol (PC-OH; E) in leaves of *Arabidopsis thaliana* and *Eutrema salsugineum* plants irrigated with water and with NaCl solution. Salinity of 0.15 and 0.3 M NaCl, was used for *A. thaliana* and *E. salsugineum*, respectively. Data represent means \pm SD, $n = 6$ (A,B), 3–4 (C,D,E). \diamond – a significant difference between control and salinity-treated plants; * – a significant difference from *A. thaliana* from the same treatment.

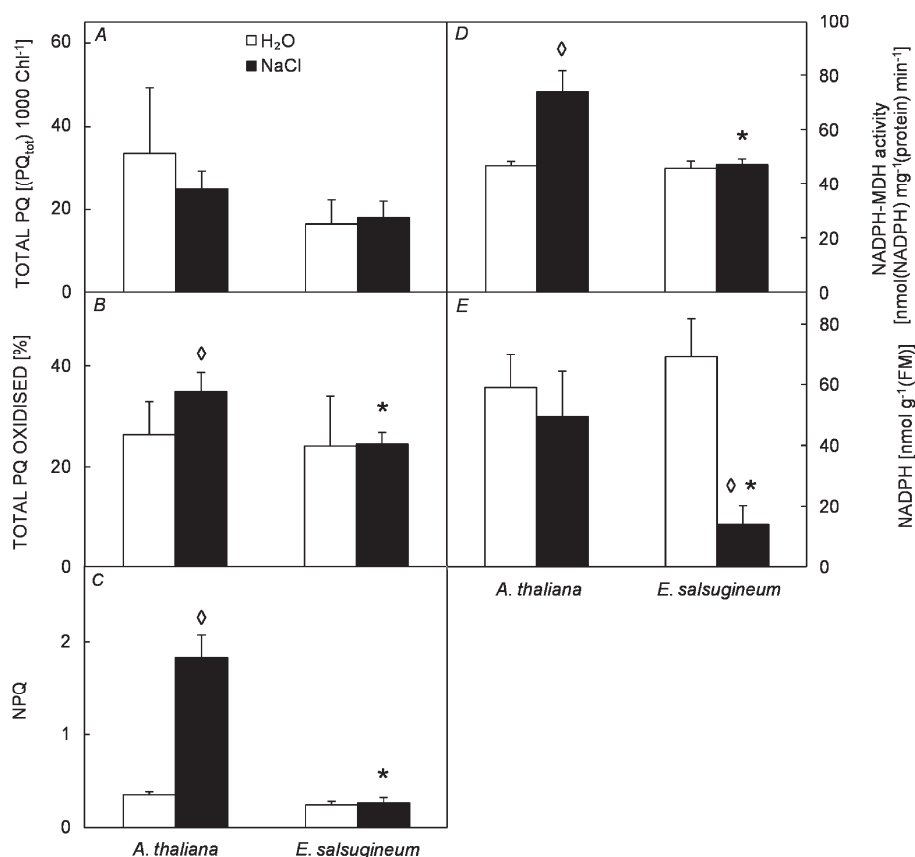


Fig. 2. Effects of salinity on the content of total foliar PQ (A), the redox state of total foliar PQ (oxidized and reduced, B), NPQ parameter (C), the activity of NADP-MDH (D), and the content of NADPH (E) in leaves of *Arabidopsis thaliana* and *Eutrema salsugineum* plants irrigated with water and with NaCl solution. Salinity of 0.15 and 0.3 M NaCl, was used for *A. thaliana* and *E. salsugineum*, respectively. Data represent means \pm SD, $n = 3-5$ (A, B), 6–7 (C), 5 (D), 3–6 (E). ♦ – a significant difference between control and salinity-treated plants; * – a significant difference from *A. thaliana* from the same treatment

Arabidopsis plants, the content of α -Toc was twofold compared to that in *Eutrema*, and this difference increased to threefold under salt conditions. The content of PC-8 was the same in control plants of both species. The content of PC-OH was similar in control plants of both species and increased only in *A. thaliana* after salinity.

Salinity-evoked changes in the foliar plastoquinone and NPQ

Change of the redox state of PQ pool is typically observed under stress, however, studies on both of the redox state of the PQ-pool and total PQ (photosynthetically active and inactive) during salinity are rather scarce. Therefore, we analysed the content of total foliar PQ (oxidized and reduced) and its redox state in both species after salinity (Fig. 2A,B). Under control conditions, the total PQ content was similar in both species and maintained mostly in a reduced state (72.7% for *A. thaliana* and 75.9% for *E. salsugineum*). The salinity caused a significant oxidation of total PQ in *A. thaliana* (to 65.4%), and at the same time the total PQ amount in relation to Chl did not change considerably. In turn, in *E. salsugineum*, salinity did not affect neither the amount of total PQ nor its redox state which remained mostly reduced (75.5%).

In the light of a crucial importance of PQ involvement in many physiological processes, we investigated NPQ as a feasible factor that could influence PQ redox state during mild salinity. In order to estimate if NPQ contributes to the PQ-pool oxidation under conditions of mild salinity, we measured this process before and after salinity treatments (Fig. 2C). Under well-watered conditions, NPQ was fairly similar in *A. thaliana* and *E. salsugineum* plants. However, the harmful effect of moderate salinity on the photosynthetic capacity of *A. thaliana* was illustrated by the significant increase in NPQ. In contrast, treatment of *E. salsugineum* with high salinity did not affect the NPQ value that remained as low as in the control plants.

Donor and acceptor site of photosystem I: Salinity-induced stomatal closure reduces an availability of CO₂ which is followed by an overaccumulation of NADPH allowing to increase ROS formation, as shown in salt-treated leaves of *Rumex* (Chen *et al.* 2004). This situation negatively affected PSI due to the limited pool of electron acceptors. In control condition, the efficiency of PSI (Y_I) was higher in *E. salsugineum* than that in *A. thaliana* (Table 1). This was associated by a very low value of

Table 1. The effect of salinity on the quantum efficiency of PSI (Y_I), donor-side limitation to PSI (Y_{ND}), and acceptor-side limitation to PSI (Y_{NA}) measured in leaves of *Arabidopsis thaliana* and *Eutrema salsugineum* plants irrigated with water and with NaCl solution. Salinity of 0.15 and 0.3 M NaCl, was used for *A. thaliana* and *E. salsugineum*, respectively. Data represent means \pm SD ($n = 7-8$). “Diamond” indicates a significant difference between control and salinity-treated plants. “Asterisk” indicates a significant difference from *A. thaliana* from the same treatment.

Parameter	Control <i>A. thaliana</i>	<i>E. salsugineum</i>	Salinity <i>A. thaliana</i>	<i>E. salsugineum</i>
Y_I	0.739 \pm 0.081	0.940 \pm 0.040*	0.442 \pm 0.067 [◇]	0.941 \pm 0.030*
Y_{ND}	0.121 \pm 0.061	0.004 \pm 0.002*	0.453 \pm 0.078 [◇]	0.003 \pm 0.000*
Y_{NA}	0.141 \pm 0.004	0.058 \pm 0.035	0.115 \pm 0.019	0.060 \pm 0.033

donor- and acceptor-side limitations to PSI represented by Y_{ND} and Y_{NA} , respectively. A noteworthy decrease in Y_I was noted in salinity-treated *A. thaliana*, in concert with a strongly increased Y_{ND} and only slightly decreased Y_{NA} . Whereas, in salinity-treated *E. salsugineum*, efficiency of PSI remained at high level and a primary electron donor to PSI (P700) was kept at highly reduced state.

To get an insight how the redox state of the PSI acceptors is affected under the conditions of moderate salinity, we compared the enzyme activity of chloroplastic NADP-MDH and the content of NADPH in both species,

before and after salinity treatment (Fig. 2D,E). In control plants, the enzyme activity was similar both in *A. thaliana* and *E. salsugineum*, whereas after salinity the activity of this enzyme was stimulated only in *A. thaliana*. In turn, a different pattern of responses were observed in the concentration of NADPH under salinity, where NaCl treatment considerably decreased the content of NADPH in *E. salsugineum* and did not affect its amount in *A. thaliana*. These results pointed to the clear change in the maintenance of NADPH under salinity stress in the two species.

Discussion

There are numerous data on the correlation between an increased antioxidant capacity and salt tolerance (for a recent review see: Ozgur *et al.* 2013, Bose *et al.* 2014, Niewiadomska and Wiczarz 2015). Within the antioxidative response, an activation of the ascorbate-glutathione cycle plays a major role, as documented for several salt-tolerant glycophytic and halophytic species: *Pisum sativum* (Hernandez *et al.* 1995), *Lycopersicon pennelli* (Mittova *et al.* 2003), *Oryza sativa* (Vaidyanathan *et al.* 2003), *Suaeda salsa* (Cai-Hong *et al.* 2005), *Cakile maritima* (Amor *et al.* 2006), and *Thellungiella parvula* (Uzilday *et al.* 2015). Here documented increase in the contents of ascorbate and glutathione in *E. salsugineum* due to salinity stress is in agreement with this view. As suggested by Bose *et al.* (2014), in halophytes, an efficient operation of this cycle is crucial for decreasing the amounts of H_2O_2 “signatures” required for signalling purposes. In contrast, in *A. thaliana* plants, lack of positive responses to salinity from ascorbate and glutathione may suggest a weak involvement of the ascorbate-glutathione cycle.

However, an opposite salinity stress response was related to the lipophilic antioxidants (α -Toc and PC-8) and PC-OH, which increased only in *A. thaliana*. Since PC-OH, the hydroxyl derivative of plastochromanol, has been assumed as an indicator of singlet oxygen-mediated oxidative damage (Szymańska *et al.* 2014), we argued that *A. thaliana* plants activated an α -Toc turnover under a mild salinity in order to counteract the toxicity of 1O_2 . Interestingly, in *E. salsugineum*, the content of α -Toc was rather low, which is in contrast to the data, documenting a

constitutively higher amounts of α -Toc in salinity-resistant variety of *Gossypium hirsutum* and halophytic *Cakile maritima* (Gossett *et al.* 1994, Ellouzi *et al.* 2011). This may suggest a very efficient PSII protection in *E. salsugineum* independent from that provided by α -Toc. At least in part, this enhanced protection may be given by cycling of electrons around PSII in concert with an enhanced H_2O_2 production, as it was shown before (Stepien and Johnson 2009, Wiczarz *et al.* 2015).

One of the typical effects of salinity stress is a declined photochemical efficiency of PSII. A high excitation pressure on PSII reduces the PQ-pool and disturbs linear electron transport which results most likely from acidification of the thylakoid lumen which in turn activates the xanthophyll cycle (Sacksteder and Kramer 2000). This cycle is involved in a harmless dissipation of excess of absorbed light energy from PSII (Baker 2008). An activation of the xanthophyll cycle, visualized by NPQ parameter, in turn lowers the excitation pressure on PSII and minimizes 1O_2 production within the PSII antenna, and prevents overreduction of the PQ-pool (Müller *et al.* 2001). However, under high light stress, enhanced singlet oxygen production may influence redox state of the PQ-pool by its fast oxidation (Kruk and Szymańska 2012). Our results support the latter view, since NPQ increased after salinity in *A. thaliana* and at the same time, a significant oxidation of total PQ was observed. If we also take into account that plastoquinol is an effective singlet oxygen scavenger (Kruk and Trebst 2008) and it is oxidized to PQ and H_2O_2 in this reaction (Khorobrykh *et al.* 2015), it could be

assumed that production of singlet oxygen in *A. thaliana* additionally contributes to PQ-pool oxidation.

Halophytic plants seem to use other strategies than NPQ to avoid PSII photoinhibition under salinity. This has been earlier documented for a halophytic C₃-CAM intermediate *Mesembryanthemum crystallinum* (Niewiadomska *et al.* 2011). Here documented, lack of NPQ increase, as well as of PQ pool oxidation in salinity-treated *E. salsugineum* additionally confirmed the aforementioned results on a minimal singlet oxygen production at PSII. Our data are in agreement with the results of Degl'Innocenti *et al.* (2009), where a comparison of glycophytic *Hordeum vulgare* with a more salinity-resistant *H. maritimum* revealed a higher PQ reduction in the latter species both in control plants and after salinity treatment. As documented earlier using EPR method, chloroplasts of *E. salsugineum* are also capable for a decreased formation of the toxic O₂⁻ in relation to less toxic H₂O₂ in comparison to *A. thaliana* (Pilarska *et al.* 2016). We hypothesize that a highly reduced state of PQ in *E. salsugineum* already under control conditions may create a basis for a high H₂O₂ formation, which in turn activates defence mechanisms to prevent the production of a more toxic forms of ROS (O₂⁻ and ¹O₂) and to prevent lipid peroxidation. Such a strategy would explain a constitutively enhanced expression of stress genes, repeatedly reported for this species (Taji *et al.* 2004, Gong *et al.* 2005, Amtmann 2009).

Stress-driven increase in NPQ is associated with an increase in ΔpH across the thylakoid membranes. This, in turn, inhibits cytochrome *b6f* complex leading to the limited electron donation to PSI (Johnson 2011). This was visualized by a highly increased donor-side limitation in salt-treated *A. thaliana* plants. In contrast, *E. salsugineum* plants maintained an extremely low Y_{ND}, indicative for a very effective electron donation to PSI (Klughammer and Schreiber 1994) both in control and under salinity, as well as, a highly active PSI. An increased PSI efficiency of this species is in line with our earlier determinations made with oxygen electrode (Wiczarz *et al.* 2015). Extremely low Y_{ND} detected in *E. salsugineum* seems to be closely linked to a more reduced PQ pool. A decreased Y_{ND} due to salinity was documented previously with halophytic *M. crystallinum* (Niewiadomska *et al.* 2011). Because our measurements were done with excised leaves, an effect of desiccation on PSII and PSI parameters might also be taken into account. Typically, an increase in NPQ occurs due to stomatal closure. In respect to PSI performance, Peeva *et al.* (2012) described an acceleration of P700⁺ dark reduction caused by the activity of NADPH dehydrogenase (NDH) in excised leaves of barley. This is, however, unlikely not to affect PSI in a light-adapted state. Hence, a clear differences in NPQ and in PSI parameters persistent in excised leaves of the two species rather

illustrated a specific performance of *E. salsugineum*. On this basis, we speculate that ability to preserve an efficient electron donation to PSI under stress is one of the key traits for a high salinity resistance. It is tempting to speculate that a stimulation of cycling electron transport at PSI might be responsible for this phenomenon, and verification of this hypothesis is under way.

A clear discrepancy in the functioning of PSI between the two species became less evident as far as the acceptor site of PSI is concerned. However, in *E. salsugineum*, a tendency to lower Y_{NA} was detected. Under salinity-dependent stomatal closure and lowered CO₂ supply, a restriction occurs in usage of NADPH by Calvin cycle. In this situation, several strategies of processing of an excessive reducing power might be involved. As visualized in our study, at similar levels of FNR protein (Fig. 1S, *supplement available online*) in *A. thaliana* and *E. salsugineum*, different strategies of NADPH management were activated under salinity. In *A. thaliana*, a salinity-dependent increase in the activity of NADPH-MDH may indicate an intensified export of reducing equivalents from chloroplasts *via* so-called malate valve. This safety valve takes advantage of excess NADPH to convert oxaloacetate to malate which is then transported to cytosol (Scheibe 2004). The activity of this enzyme is regulated by increasing reduction of thioredoxin pool and switches off when NADPH is consumed for assimilatory processes (Scheibe *et al.* 2005). In contrast, in *E. salsugineum*, a strong salinity-dependent decline in the content of NADPH may suggest either its limited production or intensified consumption. One explanation of this phenomenon might be an activation of other efficient electron sinks from reduced ferredoxin. A competition for electrons from PSI has been clearly shown with *Arabidopsis nadp-mdh* mutant, in which a set of compensatory mechanisms, such as NTRC/2-Cys Prx system, proline biosynthesis or photorespiration, were activated in order to avoid excess of reducing equivalents (Hebbelmann *et al.* 2012). In agreement with this view, in *E. salsugineum*, an activation of alternative sinks for NADPH, such as adenosine-5'-phosphosulfate reductase involved in sulfate assimilation (Gong *et al.* 2005), thioredoxin CDSP32 (M'rah *et al.* 2007), peroxiredoxins BAS1 and PrxQ (Gao *et al.* 2009, Pilarska *et al.* 2016), and proline biosynthesis (Ghars *et al.* 2012) may be suggested. Another possibility to explain a low NADPH production might be an activation the cyclic electron flow around PSI.

In conclusion, the new features of high salinity resistance of *E. salsugineum* were resolved in this study: an activation of the ascorbate-glutathione cycle, avoidance of singlet oxygen-mediated damage at PSII, and in the same time, the ability to sustain a highly reduced states of PQ pool and P700. These are, however, associated with a very low NADPH yield.

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