



Ammonium aggravates salt stress in plants by entrapping them in a chloride over-accumulation state in an NRT1.1-dependent manner

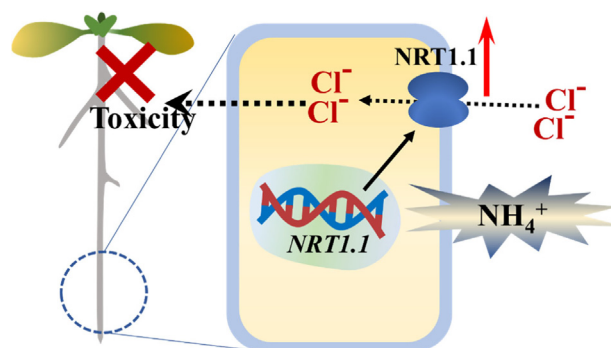
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HIGHLIGHTS

- The plants are hypersensitive to salt stress in NH_4^+ medium due to Cl^- uptake.
- Excess of Cl^- in plants grown in NH_4^+ medium is associated with the Cl^- uptake by NRT1.1.
- The impact of Cl^- toxicity in flooded-coastal areas requires serious consideration.

GRAPHICAL ABSTRACT



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ABSTRACT

Global climate change has exacerbated flooding in coastal areas affected by soil salinization. Ammonium (NH_4^+) is the predominant form of nitrogen in flooded soils, but the role played by NH_4^+ in the plant response to salt stress has not been fully clarified. We investigated the responses of *Arabidopsis thaliana*, *Oryza sativa*, and *Nicotiana benthamiana* plants fed with NH_4^+ . All species were hypersensitive to NaCl stress and accumulated more Cl^- and less Na^+ than those fed with NO_3^- . Further investigation of *A. thaliana* indicated that salt hypersensitivity induced by the presence of NH_4^+ was abolished by removing the Cl^- but was not affected by the removal of Na^+ , suggesting that excess accumulation of Cl^- rather than Na^+ is involved in NH_4^+ -conferred salt hypersensitivity. The expression of nitrate transporter NRT1.1 protein was also up-regulated by NH_4^+ treatment, which increased root Cl^- uptake due to the Cl^- uptake activity of NRT1.1 and the absence of uptake competition from NO_3^- . Knockout of NRT1.1 in plants decreased their root Cl^- uptake and retracted the NH_4^+ -conferred salt hypersensitivity. Our findings revealed that NH_4^+ -aggravated salt stress in plants is associated with Cl^- over-accumulation through the up-regulation of NRT1.1-mediated Cl^- uptake. These findings suggest the significant impact of Cl^- toxicity in flooded coastal areas, an issue of ecological significance.

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1. Introduction

Soil salinization negatively influences plant growth and crop productivity in coastal regions (Deinlein et al., 2014; Li et al., 2014). Grattan and Grieve (1992) proposed that salinity-driven root inhibition

was associated with inhibition of nutrient uptake via ion competition (e.g., Na^+/K^+ , $\text{Na}^+/\text{Ca}^{2+}$, $\text{Na}^+/\text{Mg}^{2+}$, and $\text{Cl}^-/\text{NO}_3^-$) during transportation across cell membranes. Following root uptake of Na^+ and Cl^- , the accumulation of these two ions in plant tissues may exacerbate nutritional imbalances via competition with other nutritional ions. Because of the negative effect of salt stress on plant nutrition, the nutritional conditions of the growth environment may exert a compounding effect on the severity of salt stress in plants (Bazihizina et al., 2019; Geilfus,

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2019). However, to date, the role of nutrients in the growth medium in the severity of salt stress for plants remains unclear.

Nitrogen is the essential nutrient required in the greatest quantity by plants and is one of the most limiting factors in crop production (Daniel-Vedele et al., 2010). Nitrogen is taken up by the roots primarily in the forms of ammonium (NH_4^+) and nitrate (NO_3^-). The presence of NO_3^- in the growth medium has been shown to enhance both the root uptake and xylem loading of Na^+ , and as a result, NO_3^- is considered to be a detrimental factor for plant growth under saline conditions (Álvarez-Aragón and Rodríguez-Navarro, 2017). Flooding in coastal areas is increasing due to rising sea levels and the increased frequency of prolonged rainfall events and storm surges, all due to global climate change (Wahl et al., 2015). Flooded soil environments tend to be oxygen depleted, causing nitrogen to remain in the form of NH_4^+ due to intense microbial denitrification (Benckiser et al., 2016). In light of these issues, the response of plants to NaCl stress in growth media with NH_4^+ as the nitrogen source has been addressed in several studies. For example, pea (*Pisum sativum* L.) (Frechilla et al., 2001), poplar (*Populus simonii*) (Meng et al., 2016), and wheat (*Triticum aestivum* L.) (Lewis et al., 1989) plants all exhibited increased sensitivity to salt stress when they were grown with NH_4^+ instead of NO_3^- . However, some species such as smooth cordgrass (*Spartina alterniflora*) (Hessini et al., 2013), nigella (*Nigella sativa* L.) (Bensalem et al., 2020) and Carrizo citrange (*Citrus sinensis* L.) (Fernández-Crespo et al., 2012) exhibit decreased sensitivity to salt stress when grown in NH_4^+ medium, as compared to those grown in NO_3^- medium. These findings indicate that the effect of NH_4^+ on salt sensitivity varies between plant species, underscoring the need for further clarification of the mechanisms underlying NH_4^+ -mediated salt sensitivity in plants. It should be noted that in flooded environments, the conversion of NO_3^- to NH_4^+ due to microbial denitrification may minimize the uptake competition between NO_3^- and Cl^- or improve the synergistic uptake of NH_4^+ and Cl^- , both of which would likely increase Cl^- acquisition by plants. Although Cl^- is a micronutrient for plants, excess accumulation can be toxic (Geilfus, 2018). In plants such as grapes (*Vitis* spp. L.) (Fort et al., 2013; Tregeagle et al., 2010), Carrizo citrange (Brumós et al., 2009), avocado (*Persea americana* L.) (White and Broadley, 2001), soybean (*Glycine max* L.) (Cox et al., 2018), and faba bean (*Vicia faba* L.) (Franzisky et al., 2019), Cl^- toxicity has been shown to inhibit transpiration and photosynthesis and to decrease biomass production. These findings strongly indicate that in an NH_4^+ -enriched growth medium, plant sensitivity to salt stress could be associated with an excess accumulation of Cl^- .

We investigated the above assumption by monitoring the plant responses to different supplied forms of nitrogen. Our results showed that NH_4^+ -enriched growth medium aggravated salt stress in *A. thaliana*, *N. benthamiana*, and *Oryza sativa* plants. Further investigation of *A. thaliana* revealed that the excess accumulation of Cl^- (rather than Na^+) was responsible for NH_4^+ -conferred salt hypersensitivity, which was found to be dependent on the up-regulation of root Cl^- uptake by nitrate transporter AtNPF6.3/NRT1.1. Our findings demonstrate the importance of considering the negative impacts of Cl^- toxicity on crop production in frequently flooded coastal regions, as well as in other soil environments where NH_4^+ is the predominant form of plant-available nitrogen.

2. Materials and methods

2.1. Plant materials

The *A. thaliana* ecotypes Columbia-0 (Col-0) and Landsberg erecta (Ler) were used in the present study. The single mutants *chl1-5*, *nrt1.1-1* (salk_097431), *nrt1.2* (cs859605), *nrt2.1* (salk_141712), *nrt2.2* (salk_043543), *nrt2.5* (GK 213H10), and *nlp7* (cs868891); the double-mutant *nrt2.1 nrt2.2* (cs859604); and the transgenic plants *pNRT1.1::NRT1.1-GUS* (cs6513) and *pNRT1.1::NRT1.1-GFP* all had a Col-0

background. The mutants *chl1-6* (cs6154) and *nrt2.4* (cs27332) had a Ler background. Seeds of *chl1-5* and *pNRT1.1::NRT1.1-GFP* were kindly provided by Dr. Philippe Nacry (Biochimie et Physiologie Moléculaire des Plantes, Montpellier, France). The insertions in the above lines were verified in our previous studies (Fang et al., 2016).

2.2. Plant culture

The plants were grown in agar medium in sterile Petri dishes. The seeds were surface-sterilized using 25% NaClO and sown on basal medium containing 1% (w/v) sucrose, 0.8% (w/v) agar, and 2.5 mM 2-(N-morpholino) ethanesulfonic acid (MES). Rice (*Oryza sativa* L. ssp. *japonica*) seedlings were grown in a hydroponic medium. The nutrient composition of the basal medium was as follows: 2 mM KNO_3 , 1 mM CaCl_2 , 500 μM NaH_2PO_4 , 500 μM MgSO_4 , 25 μM Fe-EDTA, 10 μM H_3BO_3 , 0.5 μM ZnSO_4 , 0.5 μM MnSO_4 , 0.1 μM $(\text{NH}_4)_6\text{Mo}_7\text{O}_{24}$, and 0.1 μM CuSO_4 , with an overall pH of 6.0. After vernalization at 4 °C for 2 days, the Petri dishes were moved to a controlled-environment growth chamber with a daily cycle of 12 h daylight at 24 °C and 12 h darkness at 22 °C. The 4-day-old seedlings were treated according to experimental requirements. For the treatment with NH_4^+ as the sole nitrogen source, KNO_3 was replaced by 1 mM $(\text{NH}_4)_2\text{SO}_4$ and 1 mM K_2SO_4 . To assay the equivalent stoichiometric Na^+ sources supplied in different forms (NaCl , Na_2SO_4 , or $\text{Na}_2\text{C}_4\text{H}_6\text{O}_4$), the CaCl_2 in the medium was replaced with CaSO_4 to avoid interference with the Cl^- from CaCl_2 . In order to avoid acidification of the medium induced by the uptake of NH_4^+ , all growth media were buffered at pH 6.0 by the addition of 2.5 mM MES.

2.3. Root measurement

Seedlings were grown in media with various nitrogen sources and various doses of NaCl for 4 days and then photographed using a digital camera. The lengths of the primary roots in the images were analyzed using the ImageJ software (<http://imagej.nih.gov/ij/>).

2.4. Measurement of Na^+ and Cl^- concentrations

Following 4 days of treatment with or without 25 mM NaCl, the seedlings were harvested and separated into roots and shoots. The samples were subsequently rinsed with ultrapure water and oven-dried at 75 °C for 2 days; their dry weights were recorded, and they were then ground to a fine powder. For Na^+ measurement, the dried samples were wet-digested as described by He et al. (2017). The digests were diluted with ultrapure water, and the Na^+ concentrations were analyzed using MP-AES (Agilent Technologies, USA). For Cl^- measurement, the ground samples were incubated overnight in an acidic solution (0.1 M HNO_3 , 10% acetic acid). Following centrifugation at 15,000 $\times g$ for 5 min, the supernatant was filtered (0.22 μm pore size), and the Cl^- concentration was determined using a modified silver titration method as described by Zhang et al. (2015).

2.5. Analysis of green fluorescent protein (GFP) and histochemical staining

After 2 days of treatment with the various nitrogen sources (with or without 25 mM NaCl), we observed the expression of GFP in the roots of *pNRT1.1::NRT1.1-GFP* plants using an epifluorescence microscope (Ni-U, Nikon, excitation 488 nm, emission 525–550 nm) and imaged using a cooled charge-coupled device camera attached to the microscope. The histochemical β -glucuronidase (GUS) staining in the roots of *pNRT1.1::NRT1.1-GUS* plants was performed as described by Fang et al. (2016). The plasma membrane integrity of the root tissues was detected after 4 days of treatment by staining with Evans blue solution (0.25%, w/v) (Guan et al., 2018). The products of GUS staining and Evans blue staining of the roots were observed using a microscope.

2.6. RNA extraction and qPCR analysis

The root tissue samples of *A. thaliana* plants were ground in liquid nitrogen, and the total RNA was isolated using TRIzol reagent according to the manufacturer's instructions. Complementary DNA was synthesized using the reverse transcription kit HiScript II Q RT SuperMix for qPCR (Vazyme, Biotech) according to the manufacturer's instructions. The mRNA levels of *NRT1.1* were detected using ChamQ Universal SYBR qPCR Master Mix (Vazyme, Biotech) in a 25 μ L reaction system with pairs of *NRT1.1*-specific primers: 5'-GCACATTGGCATTAGGCTTT-3' (forward) and 5'-CTCAATCCCACTCAGCTA-3' (reverse). The qPCR analysis was performed using the ABI StepOnePlus Real-Time PCR system. Two pairs of housekeeping gene primers were used as the control in the qPCR analysis: 5'-ACCCTAACGGGAAAGACGA-3' (*UBQ10*-forward) and 5'-GGAGCCTGAGAACAAGATGAA-3' (*UBQ10*-reverse), and GCTGTCCTTATCATGACTCCACC (*EF1 α* -forward) and TCATACCAGTCTCAACACGTC (*EF1 α* -reverse). For each sample, relative transcript abundances, calculated by the comparative Ct (threshold cycle) method and normalized to the geometric mean of the expression of the two housekeeping genes and relative to a control (NO_3^- treatment), is given by $2^{-\Delta\Delta C_t}$ (Vandesompele et al., 2002).

2.7. Measurement of Cl^- uptake

The net Cl^- fluxes of the roots were measured in the maturation, elongation, and meristematic zones using a non-invasive microelectrode ion flux measurement system (SIET IPA-2, Applicable Electronics, Inc., Forestdale, MA, USA). The Cl^- flux in each section of the roots was measured for 20 min using a microelectrode filled with chloride ionophore I-cocktail A (Sigma-Aldrich, code 99408) (Lee et al., 2013), which exhibited high sensitivity for Cl^- over NO_3^- , SO_4^{2-} , HPO_4^{2-} , and HCO_3^- (Kondo et al., 1989). The presence of other anions, therefore, did not substantially interfere with the Nernstian response of the electrode to Cl^- in our study. The microelectrodes were calibrated in 1, 5, and 10 mM KCl prior to the Cl^- flux measurement. Only electrodes with a Nernstian slope above -56 mV per decade and a correlation of 0.999 or higher were used. To compare the effects of different nitrogen sources on the net Cl^- fluxes, the 4-day-old seedlings were pre-treated with either NO_3^- or NH_4^+ media with 25 mM NaCl for 2 days, after which the net Cl^- fluxes of each seedling were measured in a medium identical to that used in for pre-treatment. To assay the induction of *NRT1.1*-mediated Cl^- uptake by NH_4^+ , the seedlings were pre-treated in either NO_3^- or NH_4^+ medium with or without NaCl for 2 days. The net Cl^- fluxes of each seedling were then measured in NH_4^+ media with 25 mM NaCl.

2.8. Statistics analyses

Data were analyzed by one-way and two-way ANOVA with post-hoc Tukey HSD test. A P -value < 0.05 was considered significant. Statistical analyses were performed using GraphPad Prism, version 8 (GraphPad Software, La Jolla, CA).

3. Results

3.1. *A. thaliana* plants are hypersensitive to NaCl stress in NH_4^+ medium

To quantify the sensitivity of plants to NaCl stress in response to different forms of nitrogen, we monitored the elongation of the primary roots of *A. thaliana* Col-0 plants grown in salty medium supplied with either NO_3^- or NH_4^+ . After 4 days of 25 mM NaCl stress, the root elongation of plants grown with NO_3^- as the nitrogen source was unaffected (Fig. 1a, b), but the plants supplied with NH_4^+ as their sole nitrogen source exhibited mild NaCl stress and dramatically restrained root elongation (reduction of approximately 40%) as compared to the NaCl-free treatment (Fig. 1a–c). Two-way ANOVA revealed a significant interaction between NaCl stress and nitrogen form ($P = 0.0011$). Similar

results were observed in *Oryza sativa* and *N. benthamiana* seedlings treated with 50 mM and 25 mM NaCl, respectively (Fig. S1). However, when the Col-0 WT *A. thaliana* plants were exposed to 50 mM NaCl for 4 days, the NaCl-induced inhibition of root elongation in the NH_4^+ treatment was similar to that observed in the NO_3^- treatment (two-way ANOVA, $P = 0.6483$, Fig. S2). To control for this effect, we used the 25 mM dose of NaCl in the remaining evaluations.

In the NH_4^+ treatments, the arrest in root elongation owing to mild NaCl stress was correlated with a clear decrease in the size of the meristem, as well as with the number of cortical cells in the meristem zone (Figs. 1d, S3). Furthermore, treatment with mild levels of NaCl and with NH_4^+ medium resulted in the destruction of the cell structure in the elongation zone (Fig. 1d). The loss of membrane integrity and the resulting cell death in the primary roots were also monitored by staining with Evans blue. As shown in Fig. 1e, the roots showed minimal staining with Evans blue in the NO_3^- growth medium, with or without the 25 mM NaCl addition. However, in the NH_4^+ growth medium, mild NaCl treatment resulted in heavy staining of the cells in the elongation zone. This result was inconsistent with the finding of the destructed cell structure in the elongation zone.

3.2. Excess accumulation of Cl^- rather than Na^+ induces salt hypersensitivity for plants in NH_4^+ conditions

Following NaCl treatment, the Na^+ levels in the roots and shoots of the plants grown with NH_4^+ were lower than those of the plants grown with NO_3^- (Fig. 2a, b). However, the Cl^- levels in both the roots and shoots of the plants in the NH_4^+ treatment were significantly higher than those of the plants in the NO_3^- treatment. This was particularly evident in the roots, where Cl^- levels were increased by more than 100% in the NH_4^+ treatment as compared with that in the NO_3^- treatment (Fig. 2c, d). We then used 12.5 mM Na_2SO_4 or 12.5 mM $(\text{Na})_2\text{C}_4\text{H}_6\text{O}_4$ (sodium succinate) instead of 25 mM NaCl to remove the Cl^- from NaCl. As shown in Fig. 2e and f, the Na_2SO_4 and $(\text{Na})_2\text{C}_4\text{H}_6\text{O}_4$ treatments had similar effects on the root elongation of seedlings to that of the NaCl and NO_3^- treatment but had clearly superior effects on the root elongation as compared to the NaCl and NH_4^+ treatment. These results indicate that the NaCl hypersensitivity of the plants grown with NH_4^+ could be abolished by removing Cl^- , supporting the hypothesis that the excess accumulation of Cl^- rather than Na^+ induces NaCl hypersensitivity in plants grown under NH_4^+ conditions. We next removed Na^+ from NaCl in the growth medium by replacing 25 mM NaCl with 25 mM KCl. This treatment had little effect on the root elongation in the NO_3^- treatment, but it significantly restrained root elongation in the NH_4^+ treatment (Fig. 2g, h). Regardless, the KCl-induced effect could be mitigated by replacing KCl with K_2SO_4 (Fig. S4), suggesting that the reduced root elongation observed in the 25 mM KCl and NH_4^+ treatment was a result of Cl^- toxicity rather than an excess of K^+ ions.

3.3. NaCl hypersensitivity of plants in NH_4^+ medium is dependent on *NRT1.1*

The inhibition of root growth by NaCl stress in NO_3^- treatment was less than that observed in the NH_4^+ treatment, indicating that the nitrate transporters (*NRTs*) involved in root NO_3^- acquisition may improve plant tolerance against mild NaCl stress in NH_4^+ -enriched environments. Six *NRTs* (*AtNPF6.3/NRT1.1*, *AtNPF4.6/NRT1.2*, *NRT2.1*, *NRT2.2*, *NRT2.4*, and *NRT2.5*) have been shown to be involved in root NO_3^- uptake in *A. thaliana* plants (Wang et al., 2018). In the NO_3^- condition, all of the tested mutants showed root growth similar to that in their corresponding WT plants, regardless of the 25 mM NaCl addition (Fig. 3a–c). Unexpectedly, the *NRT1.1*-knockout mutants *nrt1.1-1* and *chl1-5* in the NH_4^+ medium showed ~60% and ~69% greater elongation of the primary root, respectively, than Col-0 plants when grown in NaCl medium (Fig. 3a, c). The NaCl-induced inhibition of primary root growth in the third *NRT1.1*-null mutant (*chl1-6*) was also more severe than that in the corresponding ecotype (Ler) in the NH_4^+ medium (Fig. 3b, c). These results

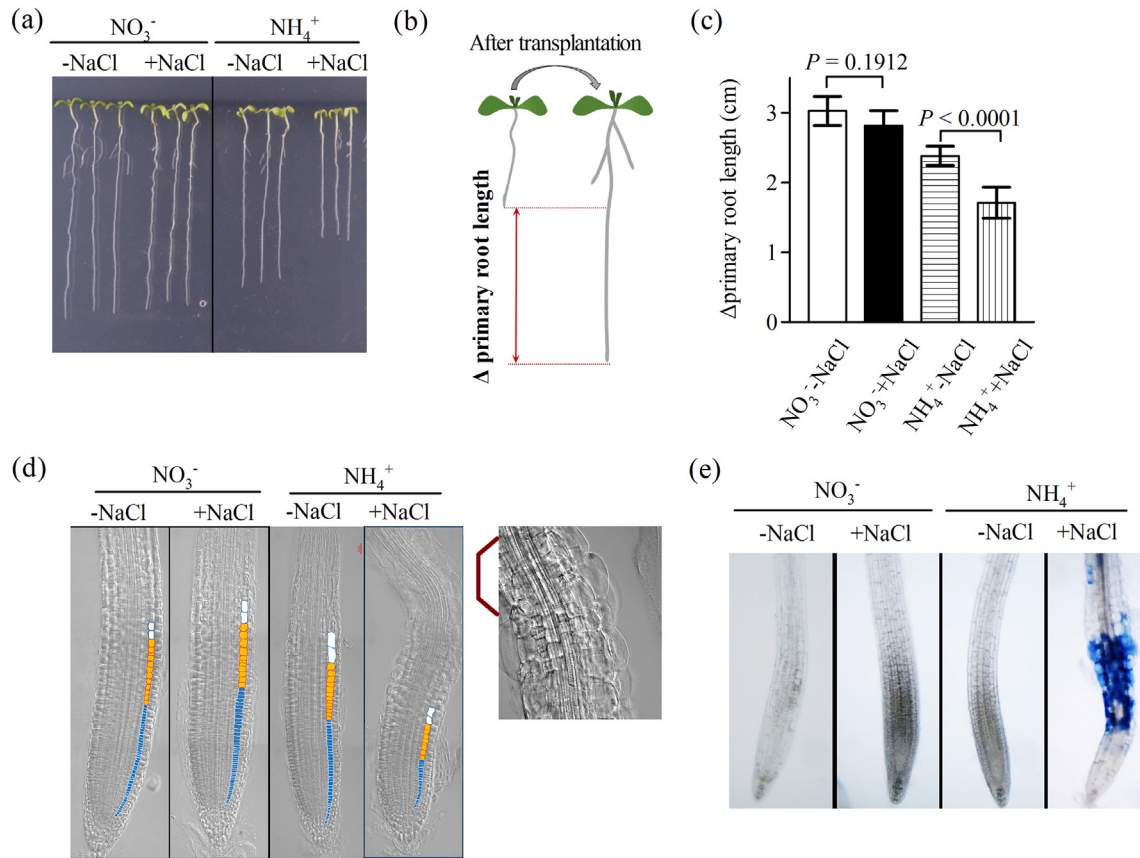


Fig. 1. Phenotype of Col-0 seedlings in response to 25 mM NaCl stress in treatments with different forms of nitrogen. (a) Representative images of the effects of different nitrogen sources on NaCl stress in Col-0 seedlings. (b) Scheme depicting the Δ primary root length of seedlings. (c) The Δ primary root length of the seedlings transferred to the treatment medium with different combinations of NaCl and forms of nitrogen. Bars and error bars indicate means \pm standard deviations ($n = 6-15$ biological replicates). (d) Differential interference contrast images of the root tips. The cortical cell files within the meristem are marked in blue, and the same files within the transition zone are marked in orange. The first two cortical cells in the elongation zone are marked in white. The image on the right depicts the destroyed cell structure in the elongation zone of the roots subjected to the NH_4^+ treatment with NaCl. (e) Detection of cell death by histochemical staining with Evans blue. The 4-day-old Col-0 seedlings were transplanted to basal agar medium with either 2 mM NO_3^- or 2 mM NH_4^+ as the sole nitrogen source and with (+NaCl) or without (-NaCl) 25 mM NaCl. The measurements were taken after 4 days of treatment. Statistical analysis: one-way ANOVA with post-hoc Tukey HSD test; P -values < 0.05 indicate significant differences between the -NaCl and +NaCl treatments within the same nitrogen condition.

suggested that NRT1.1 negatively regulated the plants' tolerance to NaCl stress in the NH_4^+ treatment. However, in contrast to the phenotypes observed in the *nrt1.1* mutants, none of the other five NRT-null single mutants (i.e., the *nrt1.2*, *nrt2.1*, *nrt2.2*, *nrt2.4*, and *nrt2.5* knockout mutants) showed improved tolerance to mild NaCl stress in the NH_4^+ medium as compared with their WT plants (Fig. 3c). We also examined NaCl sensitivity in an *nrt2.1 nrt2.2* double-mutant and found that this double-mutant also had a tolerance to mild NaCl stress similar to that of the Col-0 WT plants in the NH_4^+ medium (Fig. S5). These results indicate that NRT1.1 negatively mediates NH_4^+ -induced NaCl hypersensitivity in a relatively specific manner.

We then examined the root growth of the *pNRT1.1::NRT1.1-GFP* transgenic plants in the *chl1-5* background. The results indicated that the improved elongation of the primary roots observed in the *chl1-5* mutants grown in the NH_4^+ medium (with NaCl) could be restrained by complementation with *pNRT1.1::NRT1.1-GFP* (Fig. 3d). Furthermore, Evans blue staining was absent in the elongation zone of *chl1-5* roots, but the roots were heavily re-stained with the complementation of *pNRT1.1::NRT1.1-GFP* (Fig. 3e).

3.4. Excess accumulation of Cl^- in plants grown in NH_4^+ medium is associated with the Cl^- uptake activity of NRT1.1

In the NH_4^+ treatment with NaCl, the Cl^- levels in the roots and shoots of *nrt1.1* mutants were demonstrably lower than those of Col-0

plants (Fig. 4a, b). In contrast to the results observed in Col-0, the NH_4^+ condition did not increase Cl^- levels in the roots and shoots of *nrt1.1* mutants compared with the NO_3^- treatment. Furthermore, we observed that the difference in the root growth of Col-0 and *nrt1.1* mutants under NaCl stress in the NH_4^+ medium was abolished by replacing Cl^- with SO_4^{2-} or $\text{C}_4\text{H}_6\text{O}_4^{2-}$ (Figs. 4c, S6). Additionally, in the NH_4^+ medium containing NaCl, the root growth restrained by the complementation of *pNRT1.1::NRT1.1-GFP* in the *chl1-5* background could be restored by removing Cl^- (replacing the NaCl with Na_2SO_4) (Fig. S7).

A recent study revealed that NRT1.1 has Cl^- permeability in the *Xenopus* oocyte system (Wen et al., 2017). In light of this, we compared the Cl^- uptake between Col-0 and *chl1-5* using a non-invasive technique. The transmembrane Cl^- fluxes of the roots were measured in the meristematic, elongation, and maturation zones of the roots grown in a medium with 25 mM NaCl (Fig. 5a, b). In Col-0 plants, the net Cl^- influx in the elongation zone was greater than that of the other two root zones in both the NO_3^- and NH_4^+ treatments, particularly so in the latter (Fig. 5b). This finding is consistent with the previous findings indicating that the elongation zone was the most sensitive site to mild NaCl stress in the NH_4^+ treatment (Fig. 1e). As expected, in the NH_4^+ treatment, the loss of the NRT1.1 function in the *chl1-5* mutant led to a significant decrease in net Cl^- influx in the elongation and maturation zones but only a slight decrease in the meristematic zone, as compared to that in Col-0 plants (Fig. 5a, b). These results further demonstrated the Cl^- uptake activity of AtNPF6.3/NRT1.1 in plants.

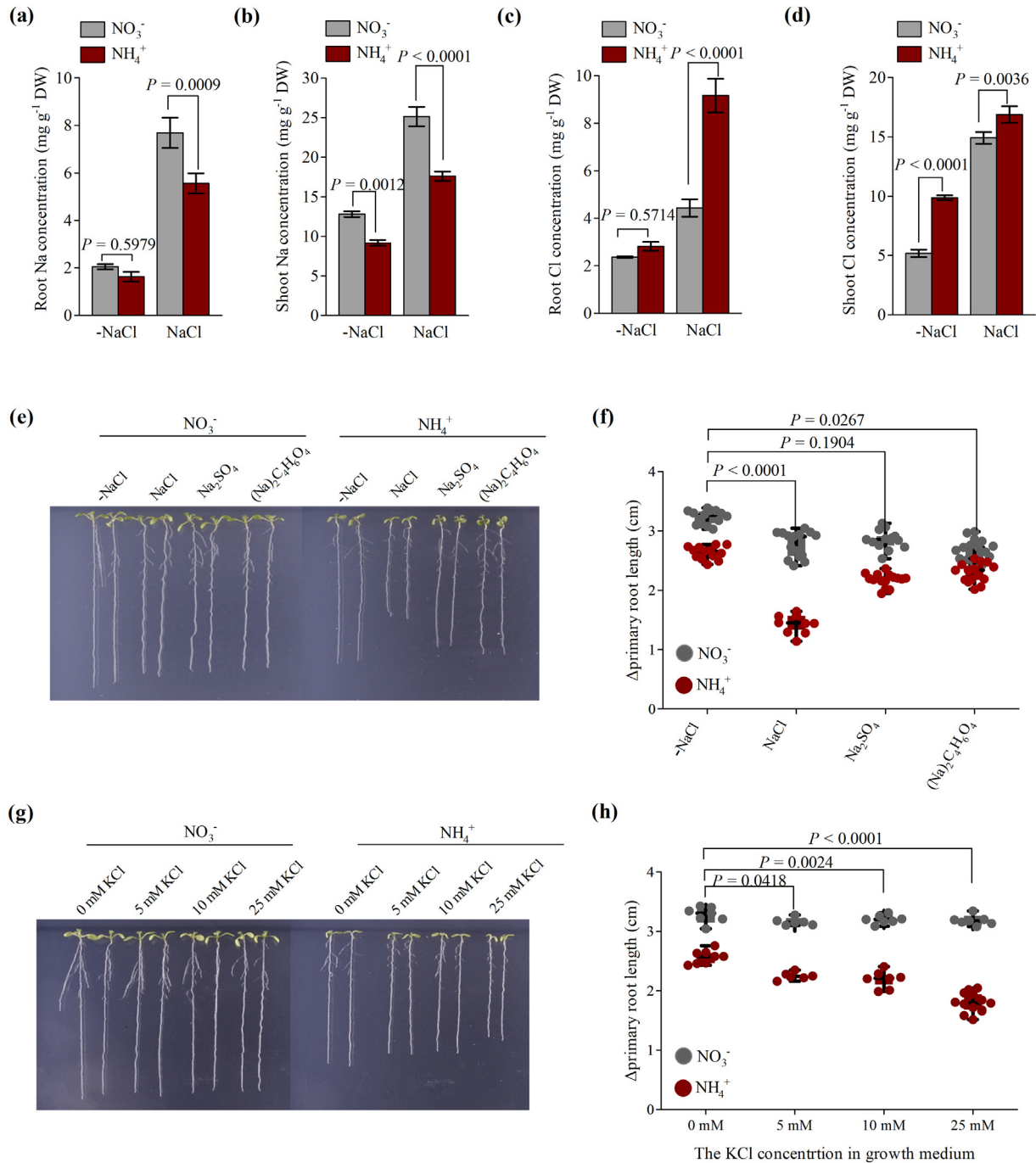


Fig. 2. Role of Cl^- in NH_4^+ -conferred salt hypersensitivity in Col-0 seedlings. (a–d) The Na and Cl concentrations in the roots and shoots of seedlings harvested after 4 days of treatment. The treatments were the same as those described in Fig. 1. Statistical analysis: one-way ANOVA with post-hoc Tukey HSD test; P -values < 0.05 indicate significant differences between the NO_3^- and NH_4^+ treatments with the same NaCl condition. Bars and error bars indicate means \pm standard deviations ($n = 5$ biological replicates). DW, dry weight. (e) Representative images of the effects of different Na^+ sources on Col-0 seedlings. (f) The Δ primary root length in response to different Na^+ sources. (g) Representative images of KCl dose-dependent responses of Col-0 seedlings. (h) The KCl dose responses of Δ primary root length. The 4-day-old Col-0 seedlings were transplanted to basal agar medium with either 2 mM NO_3^- or 2 mM NH_4^+ as the sole nitrogen source and containing either 0 mM NaCl (-NaCl), 25 mM NaCl (+NaCl), 12.5 mM Na_2SO_4 , or 12.5 mM $(\text{Na})_2\text{C}_4\text{H}_6\text{O}_4$ (e, f), or various doses of KCl as indicated in the Figures (g, h). In (e, f), the CaCl_2 in the basal agar medium was replaced with CaSO_4 to avoid interference with the Cl^- ions from CaCl_2 . Seedlings were analyzed after 4 days of treatment. Statistical analysis: two-way ANOVA with post-hoc Tukey HSD test; P -values < 0.05 indicate significant interactions between the nitrogen form and Na^+ source or KCl dose. For each box plot, the dots represent each specific datum, the interior line indicates the mean, and the edges represent the estimates of the first and third quartiles ($n = 6$ –15 biological replicates).

No statistically significant differences were observed between Col-0 and *chl1-5* in terms of the net Cl^- influx for all three root zones in the NO_3^- treatments. In addition, Col-0 seedlings showed lower net Cl^- influxes in the elongation and maturation zones (particularly in the former zone) in the NO_3^- treatment, as compared to those in the NH_4^+ treatment (Fig. 5a). These results may be due to competition between Cl^- and NO_3^- during their uptake by NRT1.1.

Accordingly, we examined the effect of additional NO_3^- on the net Cl^- influx of the elongation zone in an NH_4^+ medium containing 25 mM NaCl. As anticipated, the addition of 10 mM NO_3^- rapidly decreased the net Cl^- influx in Col-0 plants but had little effect on the net Cl^- influx in the *chl1-5* mutant (Fig. 5c). This swift inhibitory effect indicates competition between Cl^- and NO_3^- during their uptake by NRT1.1.

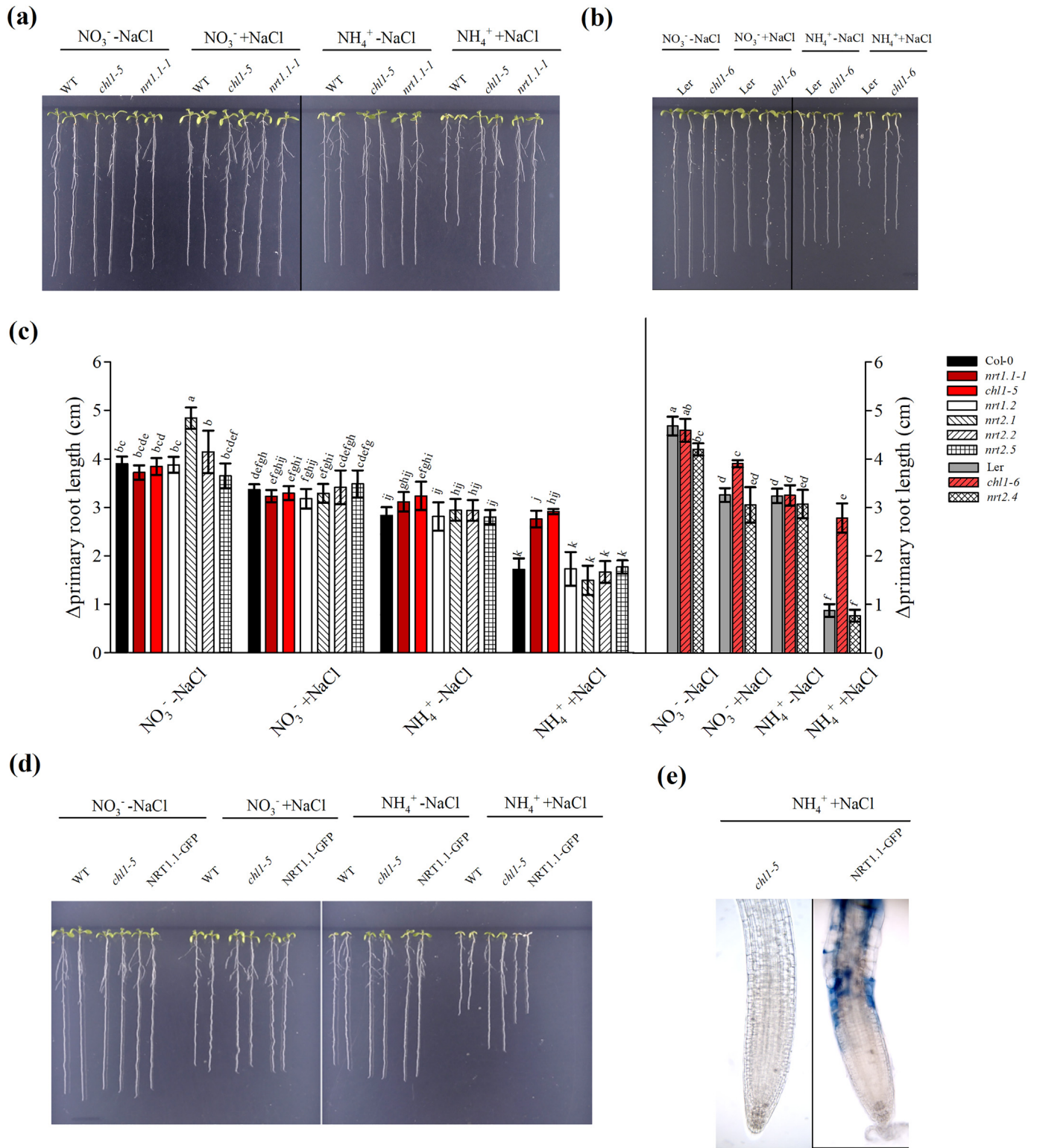


Fig. 3. Roles of nitrate uptake transporters (NRTs) in NH_4^+ -conferred salt hypersensitivity. (a, b) Phenotypic comparison of *nrt1.1* mutants and their corresponding wild-type seedlings in response to mild NaCl stress in different nitrogen conditions. (c) The Δ primary root length of the wild-type and NRT-null mutant seedlings grown in media with different combinations of NaCl and forms of nitrogen. (d) Appearance of Col-0, *chl1-5* mutant, and *NRT1.1-GFP* transgenic seedlings grown in media with different combinations of NaCl and nitrogen forms. (e) Cell death in the *chl1-5* and *NRT1.1-GFP* transgenic seedling roots, as evaluated by Evans blue staining. The 4-day-old seedlings were treated as described in Fig. 1. The measurements were performed 4 days after seedling transfer. Statistical analysis: one-way ANOVA analysis with post-hoc Tukey HSD test. Different letters indicate significant differences within the same treatment (P -values < 0.05). Bars and error bars indicate means \pm standard deviations ($n = 8-15$ biological replicates).

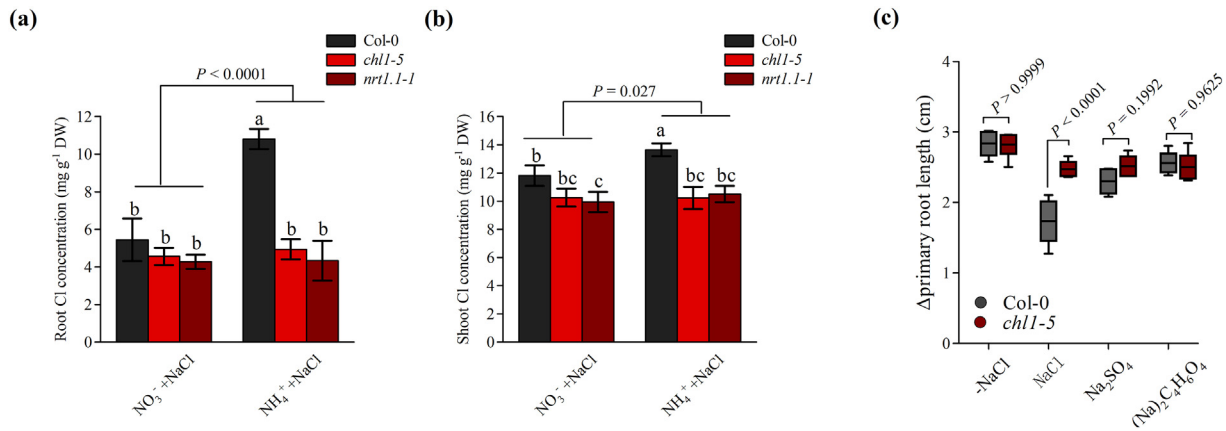


Fig. 4. Alleviation of NaCl toxicity in NH_4^+ -fed *nrt1.1* mutants, associated with a low accumulation of Cl^- . (a, b) The Cl^- concentrations in the roots and shoots of Col-0, *chl1-5*, and *nrt1.1-1* seedlings harvested after 4 days of treatment. The treatments were the same as those described in Fig. 1. Statistical analysis: one-way ANOVA analysis with post-hoc Tukey HSD; P -values < 0.05 indicate significant differences between Col-0 and *chl1-5* or *nrt1.1-1* mutants within the same treatment. Bars and error bars indicate means \pm standard deviations ($n = 5$ biological replicates). (c) The Δ primary root length of Col-0 and *chl1-5* mutants treated with 2 mM NH_4^+ and the equivalent stoichiometric Na^+ source. The CaCl_2 in the basal agar medium was replaced with CaSO_4 to avoid interference with the Cl^- ions from CaCl_2 . Statistical analysis: one-way ANOVA with post-hoc Tukey HSD test; P -values < 0.05 indicate significant differences between Col-0 and *chl1-5* or *nrt1.1-1* mutants within the same treatment. For each box plot, the interior line indicates the mean, and the edges represent the estimates of the first and third quartiles ($n = 6-15$ biological replicates).

3.5. NRT1.1 protein expression is increased in the NH_4^+ medium

Real-time quantitative PCR analysis showed that the form of nitrogen and the mild NaCl treatment had little effect on the expression of *NRT1.1* in the roots (Fig. 6a). We then analyzed the expression of *NRT1.1* protein in the roots. As shown in Fig. 6b, the *NRT1.1*-GFP protein expression (as indicated by GFP-associated fluorescence in the roots of *pNRT1.1::NRT1.1*-GFP transgenic plants) was minimally affected by mild NaCl stress in the NO_3^- medium. However, when the plants were grown in NH_4^+ medium, the expression of *NRT1.1*-GFP protein was demonstrably increased, and this increase occurred regardless of the addition of NaCl (Fig. 6b). We tested the rate of net Cl^- fluxes at the meristematic, elongation, and maturation zones of the roots following pre-culture with different nitrogen sources, with or without NaCl, for 2 days. The Cl^- flux test was conducted in the NH_4^+ medium with NaCl to avoid any resulting interference from uptake competition between Cl^- and NO_3^- (Fig. 6c). In the Col-0 plants, pre-culture with NH_4^+ increased the net Cl^- influx in the elongation and maturation zones in the same testing medium as compared to the pre-culture treatments with NO_3^- , although the net Cl^- influx in the meristematic zone did not differ significantly between the two pre-culture treatments with different nitrogen forms (Fig. 6d-f). Furthermore, the increase in the net Cl^- influx in Col-0 plants pre-cultured in NH_4^+ was not affected by the presence of NaCl in the pre-culture medium, confirming the above finding regarding the expression pattern of *NRT1.1* protein in response to different nitrogen sources (Fig. 6d-f). In the *NRT1.1*-null mutant *chl1-5*, however, no significant differences were observed in the net Cl^- influxes in all three measured zones between the two nitrogen-source pre-treatments, with or without the NaCl addition (Fig. 6d-f). We, therefore, posit that the increased Cl^- uptake by the roots in the NH_4^+ growth medium is associated with the improved expression of *NRT1.1* protein in the roots.

4. Discussion

Studies of salt stress in plants have largely focused on the effects of Na^+ , neglecting the effect of Cl^- (Li et al., 2016). In this study, we demonstrated that the excess accumulation of Cl^- , rather than Na^+ , was the key factor responsible for salt hypersensitivity in plants grown in NH_4^+ conditions. Cl^- toxicity may be a serious concern and should, therefore, be taken into account in frequently flooded coastal areas where NH_4^+ is the predominant form of soil nitrogen.

Although a transporter specifically responsible for root Cl^- uptake from the growth medium has not yet been identified, several transport proteins have been shown to be permeable to Cl^- and involved Cl^- homeostasis in plants over the past 10 years (Li et al., 2016; Li et al., 2017b). CLCc compartmentalizes Cl^- in the vacuoles (Jossier et al., 2010), while CCC (Colmenero-Flores et al., 2007), NPF2.4 (Li et al., 2016), SLAH1, and SLAH3 (Cubero-Font et al., 2016) facilitate Cl^- loading into the root xylem; NPF2.5 extrudes Cl^- from the root (Li et al., 2017a). Among these, NPF2.4 and NPF2.5 belong to the NITRATE EXCRETION TRANSPORTER (NAXT) subfamily (Segonzac et al., 2007; Tsay et al., 2007). The nitrate transporter *NRT1.1* was also shown to have Cl^- permeability in the *Xenopus* oocyte system (Wen et al., 2017). However, the permeability of such NO_3^- transporters to Cl^- in plant systems requires verification. In this study, we showed that this nitrate transporter exhibited Cl^- permeability in *A. thaliana* plants (Fig. 5a, b) and that this function was enhanced in the NH_4^+ conditions as compared to the NO_3^- conditions (Fig. 5a, b). This was likely the result of uptake competition between Cl^- and NO_3^- . Furthermore, increased uptake of the cation NH_4^+ should be accompanied by increased anion uptake to maintain the charge balance in the roots. In this context, the uptake synergy between Cl^- and NH_4^+ may be a driving force for NH_4^+ -improved Cl^- uptake by *NRT1.1*. Thus, enhanced Cl^- uptake by *NRT1.1* could be a mechanism for NH_4^+ -conferred salt hypersensitivity in plants. In this study, *nrt1.1* mutants showed greater tolerances to Cl^- exposure in the NH_4^+ medium, confirming this finding (Figs. 3a-b, 4c).

Intriguingly, loss of the *NRT1.1* function also decreased the plant's Na^+ levels (Álvarez-Aragón and Rodríguez-Navarro, 2017), which should have improved the plant's tolerance to salt stress. This finding disagrees with our conclusion that excess Cl^- , rather than Na^+ , was responsible for the *NRT1.1*-conferred salt hypersensitivity of plants grown in NH_4^+ medium (Figs. 4a-b, S8). This discrepancy may be due to the variance in growth conditions between the different experiments because only the presence of NO_3^- ensured a lower Na^+ level in *nrt1.1* mutants than in the WT plants; this difference was abolished when NO_3^- was removed from the growth medium (Álvarez-Aragón and Rodríguez-Navarro, 2017). Therefore, the mechanism of *NRT1.1*-intensified salt stress in plants is likely to be associated with the form of nitrogen present in the growth medium; in short, *NRT1.1* aggravates Na^+ accumulation in plants grown in NO_3^- conditions but entraps plants in a Cl^- -excess state in NH_4^+ conditions. In light of this finding and the disparate stress effects of Na^+ and Cl^- , the effect of different forms of nitrogen on the plant response to salt stress may be associated with the level

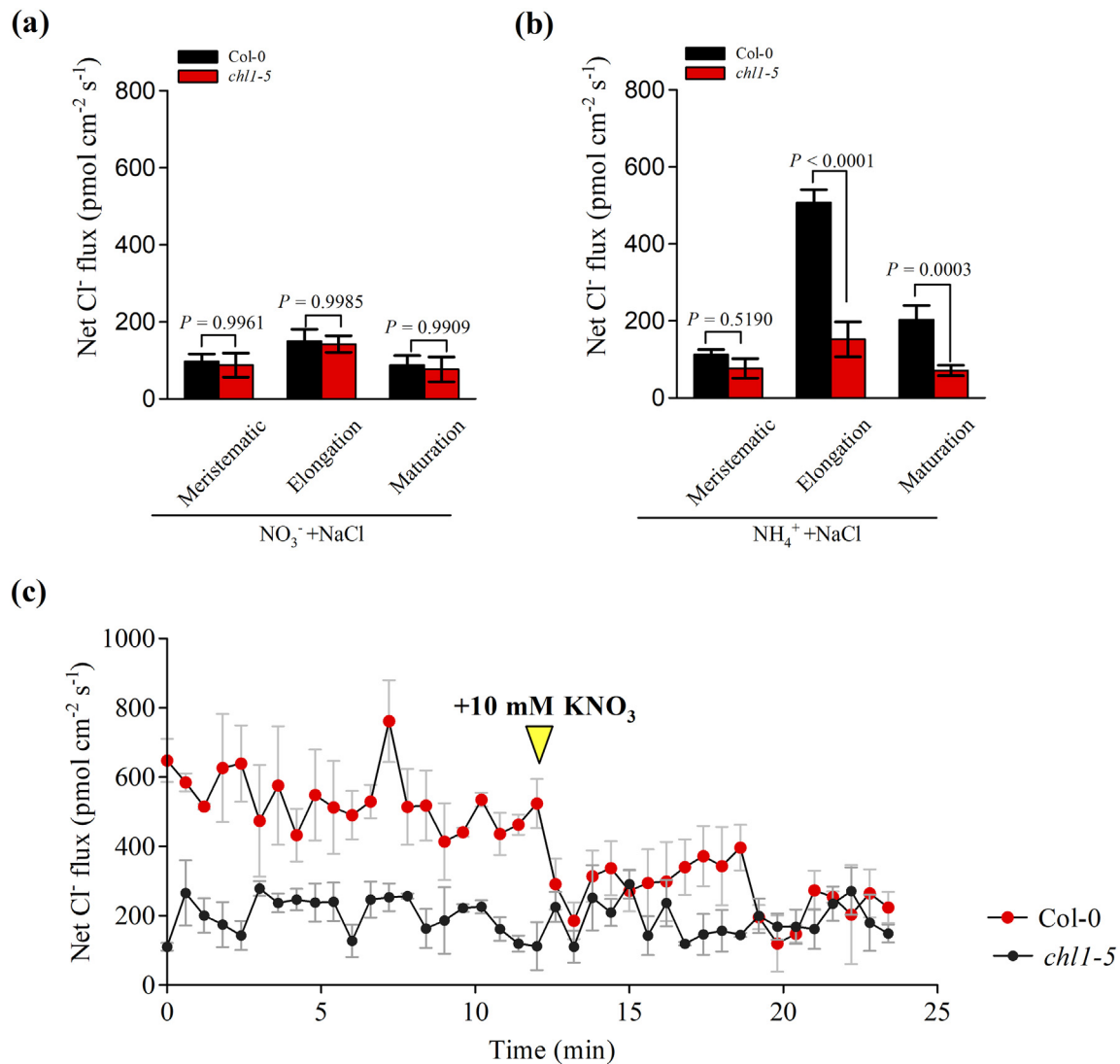


Fig. 5. Comparison of the net Cl⁻ influx between Col-0 and *chl1-5* seedlings. (a, b) Net Cl⁻ flux in roots of Col-0 and *chl1-5* mutants. The Col-0 and *chl1-5* seedlings were pre-cultured for 2 days in basal agar medium with either 2 mM NO₃⁻ or 2 mM NH₄⁺ as the sole nitrogen source and 25 mM NaCl. The net Cl⁻ fluxes in the different root zones were measured in a testing solution that had an identical ion composition to that of the medium used for the corresponding seedling treatments. Statistical analysis: one-way ANOVA with post-hoc Tukey HSD test; P-values < 0.05 indicate significant differences in the net Cl⁻ fluxes between Col-0 and *chl1-5* mutants within the same treatment. Bars and error bars indicate means ± standard deviations (n = 4 biological replicates). (c) The effect of additional NO₃⁻ on the net Cl⁻ flux in the elongation zone. The Col-0 and *chl1-5* seedlings were pre-cultured in basal agar medium with 2 mM NH₄⁺ for 2 days. Subsequently, the net Cl⁻ flux in the elongation zone of the roots was measured in the testing solution with a medium containing 2 mM NH₄⁺ and 25 mM NaCl for the first 12 min and then further measured after the addition of 10 mM NO₃⁻ (added in the form of KNO₃).

of salt stress. This is supported by the observation that no significant differences in root growth were observed between the NH₄⁺ and NO₃⁻ treatments when the NaCl stress was raised to 50 mM; Miranda et al. (2017) showed that 75 mM NaCl resulted in a higher Na⁺ accumulation in plants grown with NO₃⁻ and exhibited higher sensitivity to the salt stress than the plants grown with NH₄⁺. It is important to note, as mentioned previously, that the effect of the form of nitrogen on salt sensitivity also varies between plant species (Ullrich, 2002). For example, smooth cordgrass (*Spartina alterniflora*) plants prefer NH₄⁺ as a nitrogen source and exhibit better growth in response to salt stress when they were grown with NH₄⁺ instead of NO₃⁻ (Hessini et al., 2013; Hessini et al., 2009). This indicates that the salt sensitivity of a plant may also be affected by the nitrogen preference of the plant.

It should be noted that the expression of NRT1.1 protein in the roots was increased in the NH₄⁺ treatment, which improved the Cl⁻ uptake (Fig. 6b). Up-regulation of NRT1.1 should be detrimental for plants coping with salt stress in an NH₄⁺-dominated soil environment; therefore, this may represent another mechanism for the NH₄⁺-conferred salt

hypersensitivity in plants. It is also noteworthy that NRT1.1 has been shown to negatively regulate plant tolerance to NH₄⁺ stress (Hachiya et al., 2011; Jian et al., 2018). This finding was further confirmed in the present study, wherein we demonstrated that the NRT1.1-null mutants *chl1-5* and *nrt1.1-1* had better root growth than the Col-0 plants in the NH₄⁺ treatments without NaCl (Fig. 3a). Because 2 mM Cl⁻ (added in form of 1 mM CaCl₂) was also present in the NH₄⁺ growth medium in our study, we examined whether NRT1.1-mediated Cl⁻ uptake playing a role in NH₄⁺ toxicity to plants. The replacement of CaCl₂ with CaSO₄ in the NH₄⁺ growth medium did not improve the growth of WT plants over that of NRT1.1-null mutants, indicating that Cl⁻ uptake by NRT1.1 should be not a cause of the growth restraint of plants grown in the NH₄⁺ medium and containing low levels of Cl⁻. In the present study, the two-way ANOVA indicated that NRT1.1 conferred sensitivity to NaCl stress more significantly in Ler than in Col-0 (Fig. S10), indicating that the genetic background of a genotype or species may affect the degree of Cl toxicity conferred by NRT1.1. Studies have shown that sensitivity to high Cl⁻ concentrations varies greatly between plant species;

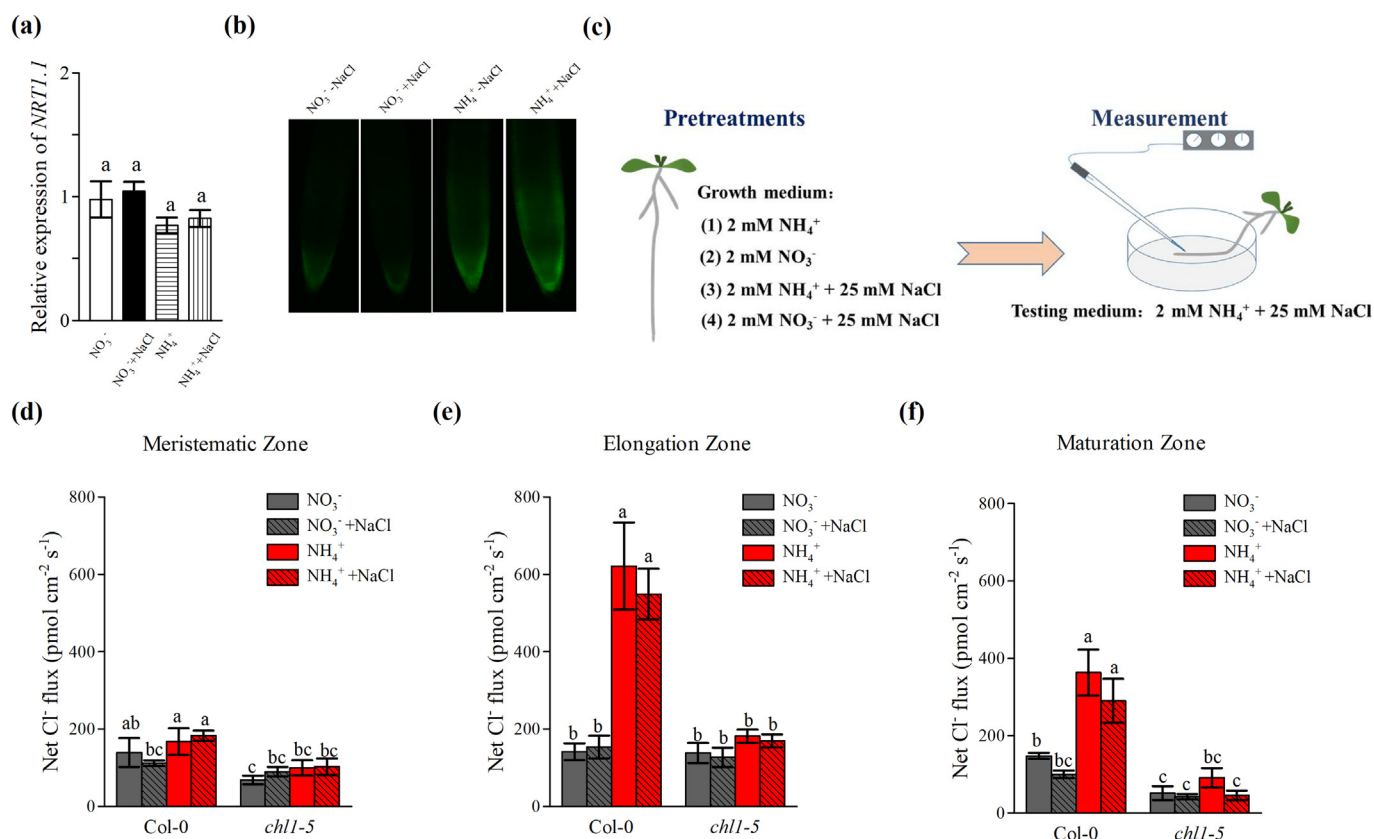


Fig. 6. Effect of NH_4^+ on the induction of NRT1.1-mediated Cl^- uptake. (a) Expression of the *NRT1.1* gene in the roots of the Col-0 seedlings. (b) The expression of NRT1.1-GFP in *pNRT1.1::NRT1.1-GFP* plants. The 4-day-old seedlings were cultured in basal agar medium with either 2 mM NO_3^- or 2 mM NH_4^+ as the sole nitrogen source, with or without 25 mM NaCl. The measurements were taken after 2 days of treatment. Bars and error bars indicate means \pm standard deviations ($n = 4$ biological replicates). (c) Scheme depicting the pre-treatment of seedlings and the measurement of net Cl^- flux. (d–f) Net Cl^- flux in the meristematic, elongation, and maturation zones of roots. The 4-day-old Col-0 and *chl1-5* seedlings were pre-treated in the basal agar medium with either 2 mM NO_3^- or 2 mM NH_4^+ as the sole nitrogen sources and with or without 25 mM NaCl, as depicted in Figure (c). The net Cl^- fluxes in the different root zones were measured 2 days after treatment with 2 mM NH_4^+ and 25 mM NaCl. Statistical analysis: two-way ANOVA with post-hoc Tukey HSD test. Different letters indicate significant differences (P -values < 0.05). Bars and error bars indicate means \pm standard deviations ($n = 4$ biological replicates).

for example, the toxicity threshold of Cl^- concentrations accumulated in plant tissues is 4–7 mg g^{-1} for citrus (*Citrus sinensis*) (Bell et al., 1997), 7–8 mg g^{-1} for rice (*Oryza sativa*) (Yin et al., 1989), about 12 mg g^{-1} for potato (*Solanum tuberosum*) (Corbett and Gausman, 1960), 25 mg g^{-1} for cotton (*Gossypium raimondii*) (Tan and Shen, 1993), and 33 mg g^{-1} for corn (*Zea mays*) (Parker et al., 1985). In this study, under the 25 mM NaCl treatment, the *A. thaliana* plants grown with NH_4^+ had about 9 mg g^{-1} Cl^- in their roots and showed evidence of toxicity (indicated by the inhibition of root growth) (Fig. 2c). *A. thaliana* likely has a lower toxic threshold for Cl^- concentration than that of potato, cotton, and corn. The above findings suggested that screening crop cultivars with low efficacy of NRT1.1-conferred Cl^- toxicity or a high toxic threshold of Cl^- concentration may help to reduce the risk of decreasing crop yields in both coastal and non-coastal areas with frequent flooding and saline soils.

5. Conclusion

The NH_4^+ -enrichment of the growth environment aggravates salt stress in *A. thaliana* plants as a result of excess accumulation of Cl^- , attributed to the increased Cl^- acquisition by NRT1.1 and lack of competition from NO_3^- .

CRedit authorship contribution statement

Xing Xing Liu: Investigation, Formal analysis, Writing – original draft. **Ya Xin Zhu:** Methodology. **Xian Zhi Fang:** Methodology. **Jia**

Yuan Ye: Methodology. **Wen Xin Du:** Resources. **Qing Yang Zhu:** Resources. **Xian Yong Lin:** Writing – review & editing. **Chong Wei Jin:** Conceptualization, Writing – original draft.

Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.scitotenv.2020.141244>.

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