

Engineering Salt Tolerance in Plants

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Aims

- Ultimately, to develop a new approach of engineering salt tolerance in plants by modifying their DNA
- To successfully transfer a particular segment of DNA (a gene) from a salttolerant plant to a salt-sensitive plant
- To grow the newly transformed plants and compare them, at both the genetic level and physiological level, to the untransformed plants to see if the transformation was successful

Introduction & Background

It is estimated that **food production must increase by 70%** in the next 40 years^[1] in order to adequately sustain the projected human population.

Materials & Methods

- Wild type *A. thaliana* transformed with P5CS1 gene (from *T. salsuginea*) by inoculating with bacteria containing the plasmid (circular DNA)
- P5CS1 is an important gene, in both A. thaliana and T. salsuginea (Fig. 1), involved in salt-tolerance by regulating production of proline (a solute which regulates water loss/uptake)
- Plasmid also contained gene for kanamycin (antibiotic) resistance
- Plants grown; seeds harvested
- Next generation plated on agar containing kanamycin
- Seedlings grown in tubes; transformation confirmed via DNA isolation and extraction, amplification via PCR and

Results & Discussion

- The recipient plant was shown to have taken up and expressed the P5CS1 gene
- The expression of the transgene should have increased the capacity to resist saltstress in the transformed plant
- The fact that the transformed seedlings grew and survived on agar containing kanamycin (Fig. 3) was an early clue that the transformation was successful





Fig. 3A Ten-day old transgenic *A. thaliana*

Fig. 3B Five-week old transgenic *A. thaliana*

By the year 2050, there will be over **2.3 billion more mouths to feed**^[2] than there are today.

However, **\$27 billion dollars**^[3] worth of crops are lost each year due to the adverse effects of salinity (high salt levels).

This is a result of over 20% of global farmland being contaminated with high levels of salt. **This is an area the size of France** and is set to increase even further in the forthcoming decades.

While some plants are naturally resistant to salt-stress and can grow under high levels of salinity, others, including many crop plants, **die quickly when exposed to high salt levels**.

Extensive work has been carried out over the past 20 years to help understand how and why these plants are able to not only survive, but **thrive in highly saline environments.**

The key mechanisms behind salt-tolerance are now clear, and it is also clear that these mechanisms are expressed (but regulated differently) by both salt-tolerant and saltsensitive plants.

What is not clear, however, are the reasons behind the differential expressions in different plants.



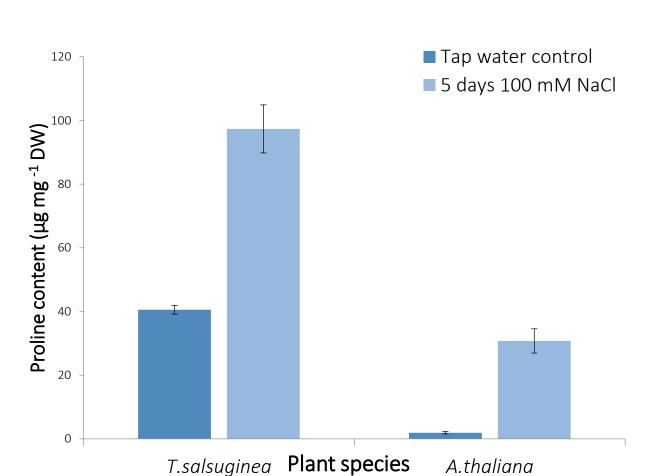
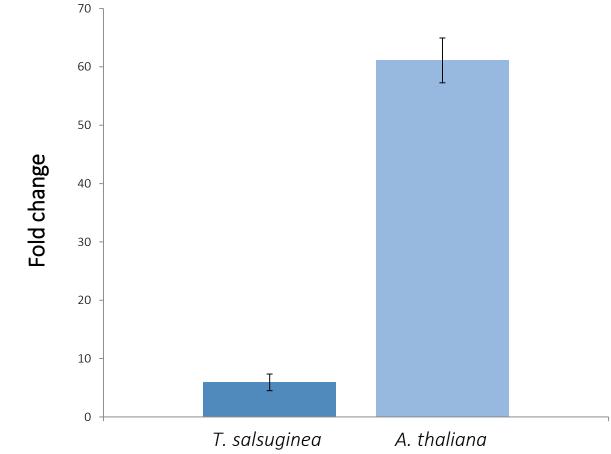


Fig. 1 Proline content in shoots of *A. thaliana* and *T. salsuginea* after either 5 days of tap water or salt-treatment with 100 mM NaCl. Both plants produced proline in response to salt-stress, but *T. salsuginea* produced more and could therefore tolerate salt better.



seedlings on an agar plate containing kanamycin. seedlings growing in a glass tube on agar containing 50 µg ml⁻¹ kanamycin.

Although we did not have time to test the transformed plants' capacity to resist saltstress (they took too long to mature), it is well established that proline accumulation is an important mechanism for resisting saltstress^[4] and plants expressing the P5CS1 gene have an increased capacity to do so.^[5]

The next step would be to engage these methods on crop plants that are saltsensitive. This would probably be more difficult than the work performed here: the plants used were closely related, and so successful transfer of genetic material between the two is, in comparison, easier to achieve.

Nonetheless, the results achieved here are a positive step in the right direction in terms of engineering salt-tolerance in plants.

Conclusions

- The P5CS1 gene was successfully transferred from a salt-tolerant plant to a salt-sensitive plant
- The gene was shown to have been

In this project, salt-sensitive plants (*Arabidopsis thaliana*), which had been transformed with DNA from salt-tolerant plants (*Thellungiella salsuginea*), were grown and studied in order to determine if salt tolerance can be engineered in salt-intolerant plants.

Plant species

Fig. 2 Relative quantification of *P5CS1* transcript fold change in *T. salsuginea* and *A. thaliana* after 5 days treatment with 100 mM NaCl. Each point is the fold change comparatively to unstressed control plants. Despite a bigger fold-change in proline, *A. thaliana* does not produce enough initially to be able to tolerate salt-stress.

expressed in the recipient plant through a variety of molecular and biochemical techniques, allowing comparison to the donor plant

 A new approach of engineering salt tolerance in plants by modifying their DNA was developed

References

1. Food and Agriculture Organisation (2009) Global agriculture towards 2050

http://www.fao.org/fileadmin/templates/wsfs/docs/Issues_papers/HLEF2050_Global_Agriculture.pdf

2. United Nations Department of Economic and Social Affairs (2015) World Population Prospects: Key findings and advance tables https://esa.un.org/unpd/wpp/Publications/Files/Key_Findings_WPP_2015.pdf

3. Qadir M et al. (2014) Economics of salt-induced land degradation and restoration. Natural Resources Forum 38: 282-295

4. Huang Z et al. (2013) Salt Stress Encourages Proline Accumulation by Regulating Proline Biosynthesis and Degradation in Jerusalem Artichoke Plantlets <u>http://dx.doi.org/10.1371/journal.pone.0062085</u>

5. Ibragimova SM et al. (2015) Evaluation of Salt Tolerance of Transgenic Tobacco Plants Bearing with P5CS1 Gene of *Arabidopsis thaliana*. Genetika 51: 1368-1375

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