

Phylogenetic analysis of *Tecticornia* using the external transcribed spacer (ETS) nuclear marker: insights into the evolution of C₄ photosynthesis

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BACKGROUND

Tecticornia is a genus of succulent halophytes in the family Chenopodiaceae. The genus currently contains two C₄ species and 39 C₃ species. Many species are endemic to Western Australia, where they play a key ecological role in salt-affected areas (Fig. 1). Photosynthesis is carried out in fused leaves and stems, known as vegetative articles (Fig. 2). High salt tolerance and unique C₄ anatomy (Kranz-Tecticornoid) make it a valuable genus for studying the steps in the evolution of C₄ photosynthesis.

Previous phylogenetic studies, using the internal transcribed spacer (ITS) nuclear marker, and chloroplastic markers, found difficulties resolving relationships among the species.

To study the steps of C₄ evolution in this genus, the C₃ species most closely related to the C₄ species must be identified. These species will be used in further molecular and anatomical studies.



Figure 1. *Tecticornia* plants on a salt lake (Cowcowing Lakes, Western Australia)



Figure 2. *Tecticornia indica* subsp. *bidens* (a C₄ species).

AIMS

- To resolve the phylogeny of *Tecticornia* using the external transcribed spacer (ETS) nuclear marker.
- To determine the number of origins of C₄ photosynthesis in *Tecticornia*.
- To identify C₃ species most closely related to C₄ species.

RESULTS

Sequences were obtained from 68 herbarium specimens, including hybrids, subspecies, and putative new species. Combining ETS and ITS sequence information produced a phylogeny with greater resolution than previous phylogenetic studies. (Fig. 3).

Paralogous ETS sequences were amplified in many species, including all C₄ species. This produced conflicting topology in early ITS and ETS trees, and some species are represented only by the ITS sequence. The presence of paralogues in *Tecticornia* species is possibly due to recent hybridisation events.



Figure 3. Phylogeny of *Tecticornia* constructed using ETS and internal transcribed spacer (ITS) sequences. The tree was obtained through Bayesian inference. Bayesian posterior probabilities (> 0.7) appear above the branches. C₄ species are highlighted in orange. Supported clades are in bold.

CONCLUSIONS AND FURTHER WORK

- Combining ITS and ETS marker sequences has resolved individual clades of species, but the backbone of the phylogeny remains unresolved.
- The C₄ species *T. indica* is not monophyletic. The phylogeny supports two origins of C₄ photosynthesis in *Tecticornia*.
- Strategies to obtain missing orthologue sequences are underway.

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