

Petal cell shape and flower-pollinator interaction in *Nicotiana*

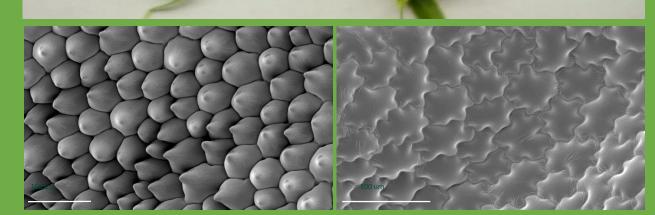
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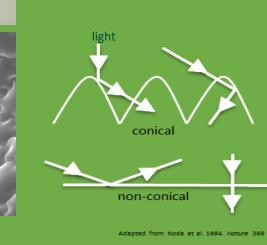


Petal epidermal cell shape plays a crucial role in plant-pollinator interaction. Conical cells are known to enhance petal colouration, regulate petal temperature and wettability, and increase grip for pollinators¹. This research looks to understand the developmental and genetic mechanisms that regulate petal cell shape in *Nicotiana* by comparing sister species with contrasting characteristics (conical vs. non-conical). It also explores the impact petal cell shape has on the interaction between flowers and pollinators in the genus.

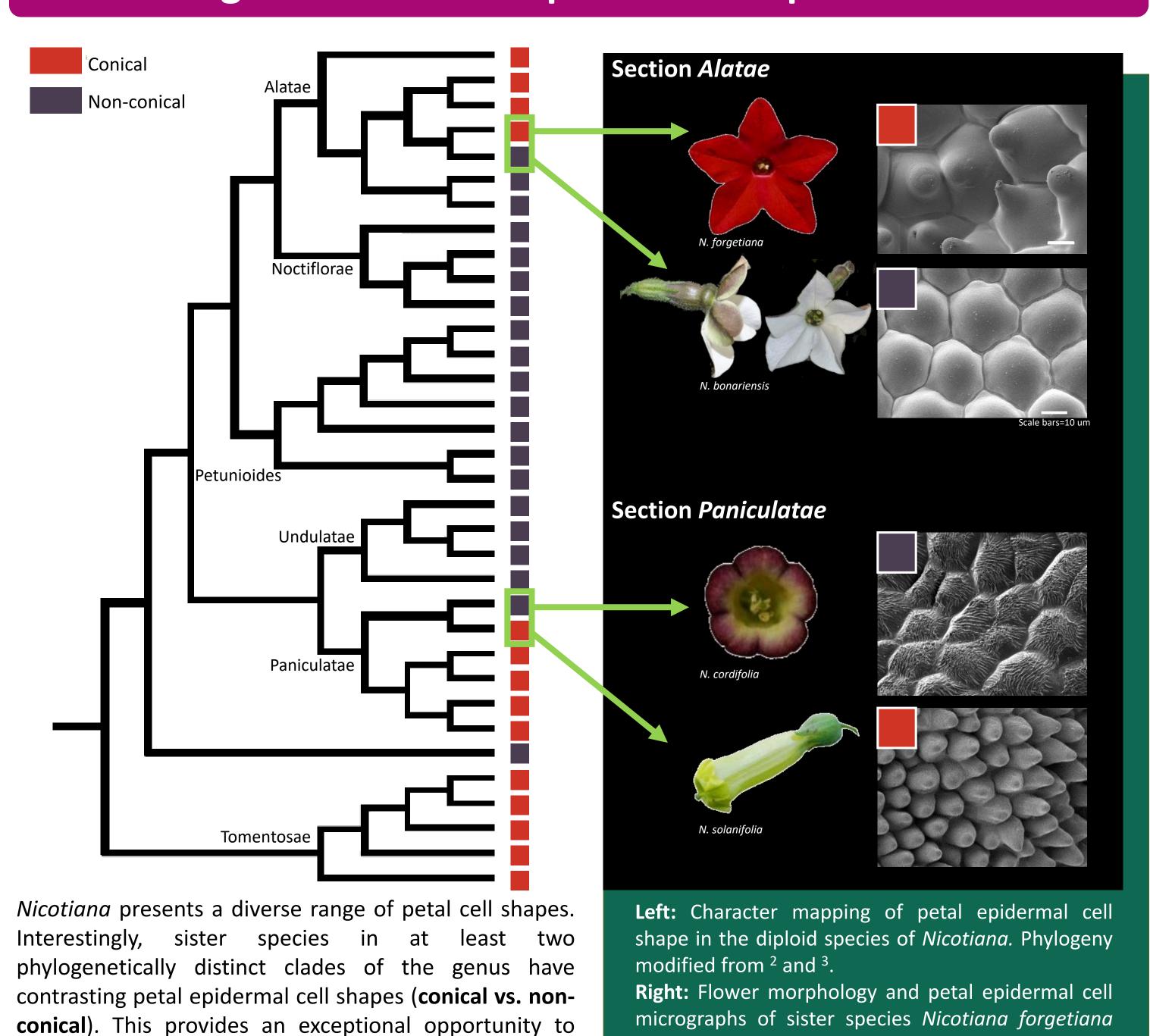


cells in wild type Antirrhinum majus (left) present enhanced colouration and provide increased grip for small pollinators (e.g. bees) compared to petals with nonconical cells in mutants (right)



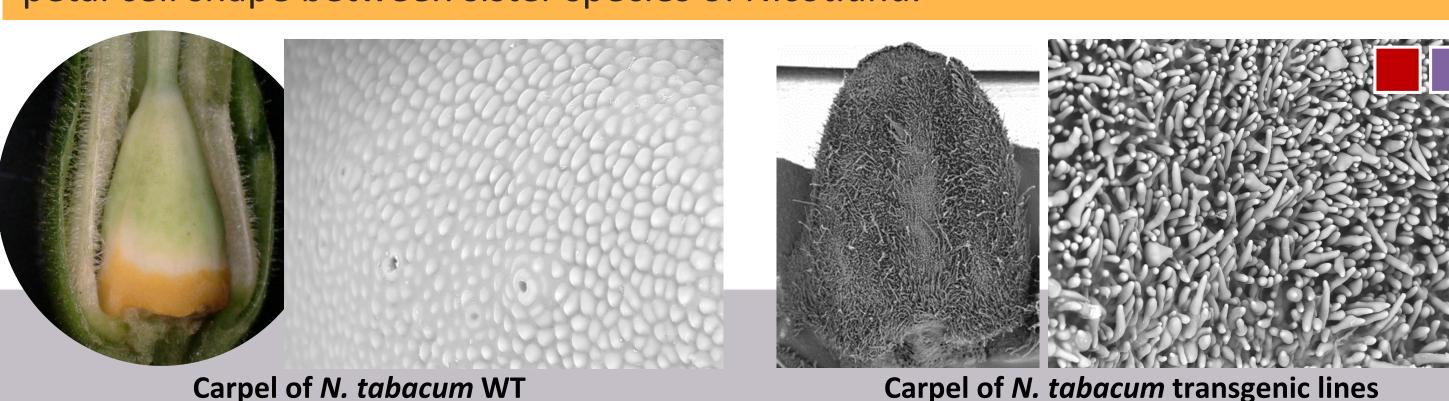


Convergent evolution of petal cell shape in Nicotiana



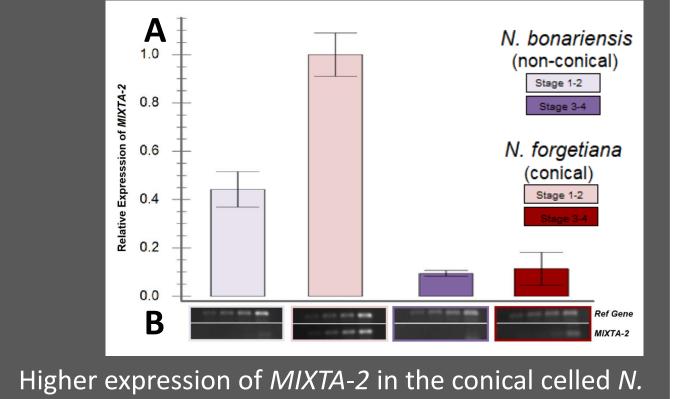
Molecular control of petal cell development

The R2R3 MYB subgroup 9 transcription factors are known to be involved in the regulation of epidermal cell outgrowths in plants. A candidate gene approach demonstrated that differences in gene expression rather than nucleotide substitutions in the gene sequences might be responsible for the differences in petal cell shape between sister species of Nicotiana.



Alternative versions of the candidate genes, those from the conical celled species as well as those from the nonconical celled species, were ectopically expressed in *Nicotiana tabacum* via *Agrobacterium tumefaciens* genetic transformation. All versions of the candidate genes, regardless of their origin (conical or non-conical celled species), had the ability to develop epidermal outgrowths, including conical cells.

Quantification of the relative expression of candidate genes across developmental stages of petals, using real-time quantitative PCR (A) and semiquantitative PCR (B), revealed significant differences between the conical celled and the non-conical celled species of Nicotiana.



forgetiana, compared to the non-conical celled N. bonariensis, suggests this gene might be involved in conical cell development in this species.

Development of petal epidermal cells in *Nicotiana*



study the evolution of petal epidermal cell shape and

explore convergent evolution of this trait in the genus.

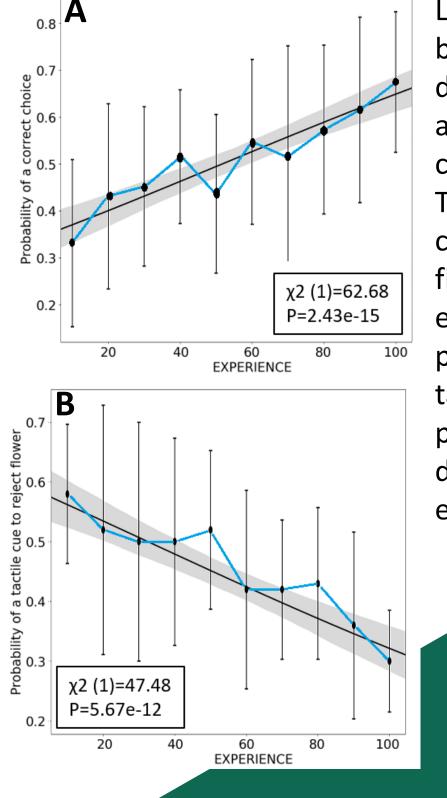
development are reached floral similar stages of development in *N. forgetiana* (conical) and N. bonariensis (non-conical). Both species start with flat cells (stage 1). The cells expand and become more rounded (stage 2-3). By stage 4 the cells of both species are dome shaped. The differences in final cell shape of these two species do not become apparent until late in floral development (stage 5).

and N. bonariensis (Section Alatae) and N. solanifolia

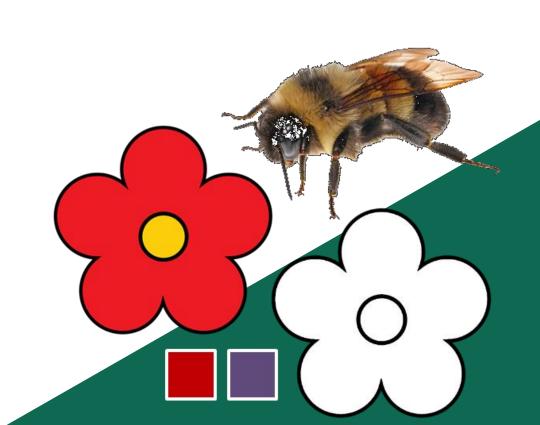
and N. cordifolia (Section Paniculatae).

Flower morphology and cryo-SEM characterization of N. forgetiana (A) and N. bonariensis (B). Numbers indicate successive developmental stages. Scale bars for morphology series = 10 mm; scale bars for SEM series = 10 μm

Exploring the function of petal cell shape in pollination

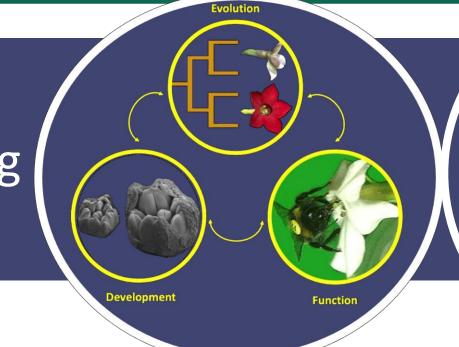


Learning curves for 20 bees choosing from six discs, three with conical and three with nonconical celled texture. The probability of a bee choosing a rewarding flower (A) increases with experience, while the probability of a bee using tactile clues to reject a punishing flower (B) decreases with experience.



Differential conditioning experiments with artificial flowers indicate that the bumblebees can discriminate flowers with conical from flowers with nonconical surfaces, on a red and on a white background, using visual cues alone as well as tactile cues alone.

Nicotiana (Solanaceae) as a system to further understand the EVOLUTION, DEVELOPMENT and **FUNCTION** of petal cell shape in flowering plants and its implications in pollination systems, by combining tools of molecular biology, morphology and pollinator behaviour experiments.





References: ¹Whitney et al., 2009. Curr. Biol. 19: 948–53; ²KNAPP, S. et al. 2004. *Taxon* 53: 73–82; ³CLARKSON, J.J. et al. 2010. *Mol. Phylogenet. Evol.* 55: 99–112.





