1050 Brief Communication

Ancient asymmetries in the evolution of flowers Pilar Cubas*, Enrico Coen⁺ and José Miguel Martínez Zapater*

Dorsoventral asymmetry in flowers is thought to have evolved many times independently as a specialized adaptation to animal pollinators [1, 2]. To understand how such a complex trait could have arisen repeatedly, we have compared the expression of a gene controlling dorsoventral asymmetry in Antirrhinum with its counterpart in Arabidopsis, a distantly related species with radially symmetrical flowers. We found that the Arabidopsis gene is expressed asymmetrically in floral meristems, even though they are destined to form symmetrical flowers. This suggests that, although the flowers of the common ancestor were probably radially symmetrical, they may have had an incipient asymmetry, evident at the level of early gene activity, which could have been recruited many times during evolution to generate asymmetric flowers.

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There are many examples in which complex morphological traits have arisen repeatedly during evolution [3, 4, 5]. However, the molecular mechanisms underlying such events are unclear in most cases. One possibility is that entirely different mechanisms arose for each event. Alternatively, the ancestor could have been evolutionarily poised, such that slight modifications of the same mechanism allowed the trait to be generated repeatedly. We have tested these possibilities in the case of floral asymmetry evolution.

The *cycloidea* (*cyc*) gene plays a key role in establishing dorsoventral asymmetry in *Antirrhinum* (snapdragon) flowers (Figure 1a) [6]. The *cyc* gene is expressed at a very early stage in the dorsal (adaxial) region of floral meri-

stems, where it affects growth rate and primordium initiation [6]. Expression continues through to later stages in dorsal primordia to affect the asymmetry, size and shape, and cell types of petals and stamens. CYC is a member of the TCP family of DNA-binding proteins, many examples of which have now been identified in the genome of Arabidopsis [7]. Of these, TCP1 is the most similar to CYC and represents a likely ortholog. We analyzed the expression of TCP1 in Arabidopsis to determine how a cyc ortholog might be expressed in a species with radially symmetrical flowers. Like cyc, TCP1 was expressed at very early stages of development, stages 1 and 2 [8], in the dorsal part of flower meristems (Figure 1c). However, in contrast to cyc, TCP1 was not expressed from stage 3 onward (Figure 1c). The transience of TCP1 expression may account, at least in part, for the lack of dorsoventral asymmetry detected at the morphological level in the adult Arabidopsis flowers. TCP1 was also expressed in the dorsal (adaxial) region of all axillary shoot meristems (Figure 1d), in a similar pattern to that in floral meristems. This shows that common prepatterns are present in both types of axillary meristems and makes TCP1 a molecular marker for axillary versus apical meristems. This also shows that, similar to lateral organs [9], axillary meristems have an underlying dorsoventral asymmetry. In contrast to the situation in flowers, TCP1 expression in axillary shoot meristems was maintained during later stages of development (Figure 1e).

These results indicate that dorsoventral asymmetry in *cyc/TCP1* gene expression predates the divergence of *Antirrhinum* and *Arabidopsis*. Thus, the common ancestor of these species, which presumably had radially symmetrical flowers, would have had asymmetric expression of the ancestral *cyc/TCP1* gene in its axillary flower and/or shoot meristems. This incipient asymmetry could then have been recruited many times independently, by changes in timing of expression and interactions with target genes, so that it became manifest at the morphological level as the dorsoventral asymmetry of the flower.

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Figure 1

Asymmetrical expression of TCP1 during Arabidopsis development. (a) Antirrhinum flower. (b) Arabidopsis flower. (c-e) Sections of Arabidopsis tissue probed with digoxigenin-labeled TCP1 antisense RNA according to the protocol described in [10]. (c) TCP1 mRNA accumulates in the dorsal (adaxial) region of stage 2 floral meristems. This expression may be responsible for the delayed initiation of the dorsal sepal primordium. TCP1 is not expressed in stage 5 flower buds, when sepals, petals, and stamens are being formed. (d) TCP1 is transcribed in the dorsal part of shoot axillary meristems. Note that this early expression pattern is similar to that in (c); (e) TCP1 expression is maintained in the dorsal part of lateral shoots during their elongation. Numbers indicate stages according to Smyth [8]. Cl, cauline leaf; Sam, shoot apical meristem; Axm, shoot axillary meristem; Fm, flower meristem; Ms, main shoot; Ls, lateral shoot.



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