

Corrections

Milo J. Aukerman and Hajime Sakai. (2003). Regulation of Flowering Time and Floral Organ Identity by a MicroRNA and Its *APETALA2*-Like Target Genes. *Plant Cell* **15**, 2730–2741.

In Table 1 on page 2732, the second and fifth entries under “Floral Phenotype” are incorrect. The corrected table is printed below.

Table 1. Flowering Times and Floral Phenotypes of Lines Used in This Study

Genotype	Rosette Leaf No. (Average) ^a	SD	Floral Phenotype
Wild type	11.4	1.2	Wild type
Wild type, short days	36.7	5.2	Wild type
<i>eat-D</i>	3.1	0.8	<i>ap2</i>
<i>35S-EAT</i>	2.0	0.2	<i>ap2</i> plus additional ^b
<i>35S-EAT</i> , short days	6.1	1.2	<i>ap2</i> plus additional ^b
<i>35S-eatdel</i>	11.1 ^c	1.1	Wild type
<i>35S-miR172a-1</i>	2.1	0.3	<i>ap2</i> plus additional ^b
<i>toe1-1D</i>	22.5	2.1	Wild type
<i>35S-TOE1</i>	28.6	3.6	Wild type
<i>toe1-2</i>	8.7	0.6	Wild type
<i>toe2-1</i>	10.2 ^c	1.4	Wild type
<i>toe1-2 toe2-1</i>	6.0	0.8	Wild type

Flowering time was determined by counting the number of rosette leaves, and floral phenotypes were observed visually. All plants were in the Col-0 genetic background and were grown in long-day conditions (16 h of light and 8 h of dark), except as indicated (short days [8 h of light and 16 h of dark]).

^a Average values from at least 10 plants per line.

^b See text and Figure 1 for details.

^c No statistically significant difference compared with the wild type. All other lines were significantly different from the wild type (Student's *t* test, $P < 0.0001$).