Corrigendum

The *Arabidopsis* homologs of trithorax (ATX1) and enhancer of zeste (CLF) establish ‘bivalent chromatin marks’ at the silent AGAMOUS locus

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The authors wish to draw the attention to two irregularities in Figures 2a and 4. Both concern errors in duplicating images of ‘empty’ lanes illustrating absence of DNA bands.

We wish to introduce the following corrections to Figure 2a and its legend, underlined text:

Figure 2. H3K27me2 and H3K27me3 profiles of three Arabidopsis genes. (a) ChIP assays with antibodies specific against di- and tri-methylated H3K27; superscripts ‘t’ and ‘d’ stand for the tri-, or di-methylated isoforms, respectively. The H3K27me2 and H3K27me3 levels for AG in clf are from the 5′-end of AG, shown in Figure 3a, below. (I) Input sample is 15% of the immunoprecipitated DNA; (-) is negative control without antibody. Quantified bands from three independently performed ChIP assays are shown as graphs. Changes in H3K27me3 levels of wild type and clf SUP and AP1/nucleosomes are insignificant (P > 0.5).

Figure 4. ChIP assays with antiATX1 antibodies. Bands corresponding to the 5′-end AG-region amplified after a ChIP with antiATX1 antibodies; downstream region nucleosomes were only weakly represented in the precipitated fraction.
The lower panel of Figure 4 was mistakenly duplicated from Figure 3a, lower left panel. The original gels are no longer available, we therefore wish to remove this panel from the figure and the sentence related to it from the Legend. A correction is required in the following sentence on page 6294, underlined text: “ChIP analysis with antiATX1 antibodies did not reveal presence of ATX1 at AP1 nucleosomes (not shown).”

We regret these omissions and apologize to the readers for the inconvenience caused. The results and conclusions remain valid.