

## Corrigendum

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

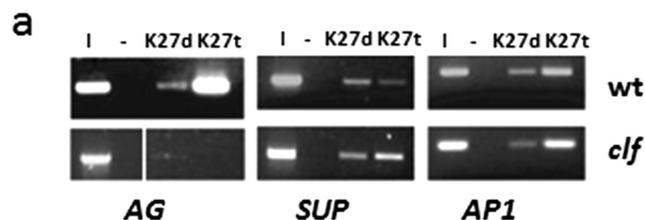
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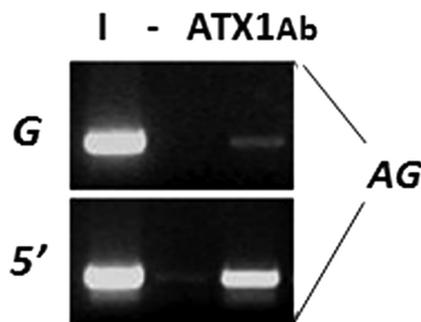
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The authors wish to draw the attention to two irregularities in Figures 2a and 4. Both concern errors in duplicating images of ‘empty’ lanes illustrating absence of DNA bands.

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. White line on AG panel indicates the two lanes were not adjacent to the input lane. (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 anti-ATX1 antibodies. Bands corresponding to the 5′-end *AG*-region amplified after a ChIP with anti-ATX1 antibodies; downstream region nucleosomes were only weakly represented in the precipitated fraction.

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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 *API*nucleosomes (not shown)”.

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