

Erratum to: Epigenetic regulation of *MdMYB1* is associated with paper bagging-induced red pigmentation of apples

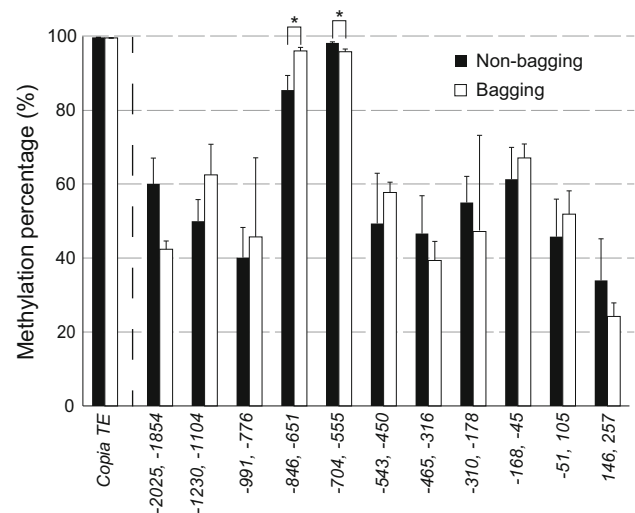
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In the online version of the original article, Fig. 7 was incorrect. The correct version is given below. The original has been corrected.

Fig. 7 DNA methylation levels in the 5' upstream region of *MdMYB1-2/-3* alleles in non-bagged and bagged 'Mutsu' fruits (Fig. 3a). The methylation percentage was estimated using an assay combining *McrBC* digestion and qPCR. The X axis indicates the position from ATG translation start sites for qPCR amplification. Digestion efficiency was evaluated by a highly methylated *Copia*-type retrotransposon, *Ppcrt1*. The statistically different (two-tail Student's *t* test) methylation regions are marked with an asterisk ($P < 0.05$); Error bars show SE. Digestion reactions were performed in triplicate with independently extracted genomic DNA (three biological replications), and qPCR was performed with three technical replicates per biological replicate



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