



Correction to: Detection of QTL for apple fruit acidity and sweetness using sensorial evaluation in multiple pedigreed full-sib families

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In the original publication, the positions of the SNP markers from the 50K SNP array are given according to GDDH13 v1.0 version of the apple reference genome and not according to GDDH13 v1.1 version as stated in the original publication.

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