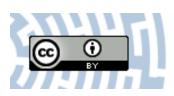


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Corrigendum: To Be or Not to Be Expressed: The First Evidence of a Nucleolar Dominance Tissue-Specificity in *Brachypodium hybridum*

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Keywords: nucleolar dominance, 35S rDNA, secondary constriction, *Brachypodium*, allopolyploidy, 3D-FISH, rRNA gene expression

A Corrigendum on

To Be or Not to Be Expressed: The First Evidence of a Nucleolar Dominance Tissue-Specificity in *Brachypodium hybridum*

by Borowska-Zuchowska, N., Robaszkiewicz, E., Mykhailyk, S., Wartini, J., Pinski, A., Kovarik, A., and Hasterok, R. (2021). Front. Plant Sci. 12:768347. doi: 10.3389/fpls.2021.768347

In the original article, there was a mistake in **Figure 3B** as published. The gel lines were marked in an incorrect order. The corrected **Figure 3B** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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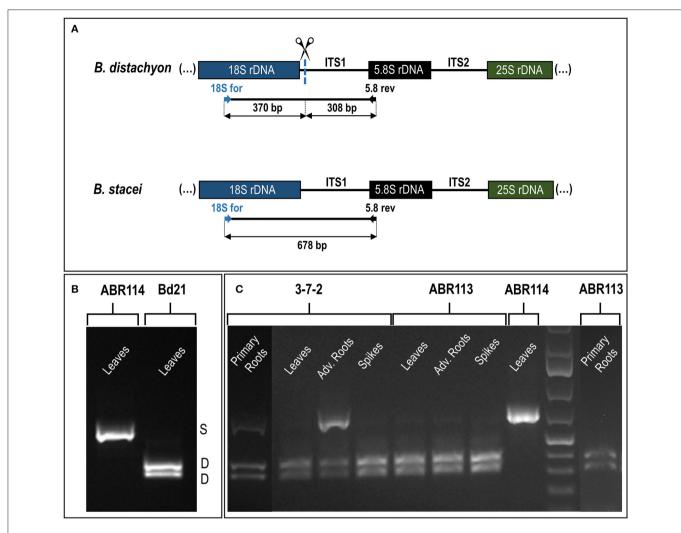


FIGURE 3 | The 35S rDNA expression analysis in the different tissues of *B. hybridum* (genotypes 3-7-2 and ABR113) and *B. stacei* (genotype ABR114; control) and *B. distachyon* (genotype Bd21; control) using the RT-PCR CAPS method. (A) The *Mlul* restriction profile of the *B. distachyon*-like and *B. stacei*-like ITS1 PCR products. The expected sizes of the bands after *Mlul* digestion are presented. (B) The *Mlul* restriction profiles of ITS1 amplification products that were obtained from the leaves cDNAs of *B. distachyon* and *B. stacei*. (C) The *Mlul* restriction profiles of ITS1 amplification products that were obtained from the primary roots, leaves, adventitious roots, and immature spikes cDNAs. Adv. Roots, adventitious roots.