



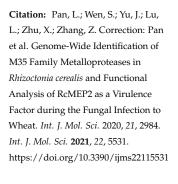
Correction

Correction: Pan et al. Genome-Wide Identification of M35 Family Metalloproteases in *Rhizoctonia cerealis* and Functional Analysis of RcMEP2 as a Virulence Factor during the Fungal Infection to Wheat. *Int. J. Mol. Sci.* 2020, 21, 2984

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In the original article, there was a mistake in Figure 8 as published. The picture of His-TF-RcMEP2 at 60 min was confused by L.J.P. because the sample container marker was blurred [1]. Thus, Figure 8 should be replaced with the following figure (Figure 1).



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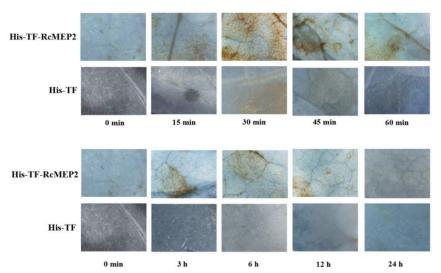


Figure 1. H_2O_2 accumulation triggered by RcMEP2 in the infiltrated *N. benthamiana* leaves. H_2O_2 accumulation (as indicated by diaminobenzidine staining) appeared obviously in the veins and stomata of His-TF-RcMEP2 infiltrated leaves but not obviously in leaves infiltrated with 5 μ M His-TF-tag solution (CK). These stains were observed and photographed under a stereomicroscope. Bar, 2 mm.

The correction does not change the scientific conclusions of the article in any way. The authors apologize for any confusion this error may have caused to the readers.

Conflicts of Interest: The authors declare no conflict of interest.

Reference

 Pan, L.; Wen, S.; Yu, J.; Lu, L.; Zhu, X.; Zhang, Z. Genome-Wide Identification of M35 Family Metalloproteases in *Rhizoctonia cerealis* and Functional Analysis of RcMEP2 as a Virulence Factor during the Fungal Infection to Wheat. *Int. J. Mol. Sci.* 2020, 21, 2984. [CrossRef] [PubMed]