

Correction

Drevensek, S., Gousot, M., Duroc, Y., Christodoulidou, A., Steyaert, S., Schaefer, E., Duvernois, E., Grandjean, O., Vantard, M., Bouchez, D., and Pastuglia, M. (2012). The *Arabidopsis* TRM1–TON1 interaction reveals a recruitment network common to plant cortical microtubule arrays and eukaryotic centrosomes. *Plant Cell* **24**: 178–191.

In the original publication, panel O of Figure 7, which represents the overlay of panels M and N, was mistakenly produced as a double overlay of panel M. The corrected overlay image is presented here. The figure legend is unchanged from the original version. We apologize for this error, which occurred at original review stage and was not detected prior to publication.

The conclusions of our article are unaffected by this correction.

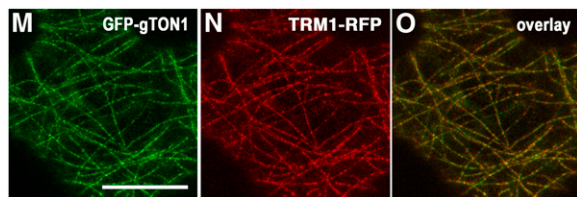


Figure 7. TRM1 Targets TON1 to Microtubules through the TRM1 M2 Motif.

(M) to (O) Coexpression of GFP-TON1 and TRM1-RFP at lower expression levels shows a punctate staining reminiscent of TRM1 and TON1 localization in *Arabidopsis*. To decrease expression levels of the TON1 fusion, we used the GFP-gTON1 construct. To decrease expression levels of the TRM1 fusion, agrobacteria carrying the TRM1-RFP construct were resuspended in infiltration buffer to an OD₆₀₀ of 0.05 (instead of 0.5).

All micrographs are projections of Z-stack confocal images. Bars = 20 μ m.

Editor's note: the corrected figure and accompanying text were reviewed by members of *The Plant Cell* editorial board.

Correction

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