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# Corrigendum: A promoter toolbox for tissue-specific expression supporting translational research in cassava (*Manihot esculenta*)

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## KEYWORDS

cassava, biotechnology, promoter, storage root, parenchyma, phloem, xylem, tissue

## A Corrigendum on

## A promoter toolbox for tissue-specific expression supporting translational research in cassava (*Manihot esculenta*)

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### Error in Figure Legend

In the published article, there was an error in the legend for Figure 2 and the legend for Figure S2 as published. In the Figure 2 legend, subfigure F<sub>2</sub> – F<sub>6</sub> were falsely labeled subfigure H<sub>2</sub> – H<sub>6</sub>. In the Figure S2 legend, the subfigures C-F were incorrectly labeled and the labeling for subfigures G-H was missing. The corrected legend appears below.

#### FIGURE 2

Representative GUS staining pattern of at least three events from *pAtCAB1*, *pStLS1*, *pAtRBCS3B*, *pMeGBSS1*, *pStSSS3*, and *pStSTP1* promoter-reporter plants. *pAtCAB1::GUS* = **A**<sub>1</sub>) Source leaf (Inlay = Close-up), **B**<sub>1</sub>) Sink leaf, **C**<sub>1</sub>) Emerging leaves, **D**<sub>1</sub>) Petiole cross-section, **E**<sub>1</sub>) Upper stem cross-section, **F**<sub>1</sub>) Lower stem cross-section, **G**<sub>1</sub>) Storage root cross-section, **H**<sub>1</sub>) Fibrous roots, **I**<sub>1</sub>) GUS expression levels of four *pAtCAB1::GUS* lines relative to three *pCaMV35S::GUS* lines in %. *pStLS1::GUS* = **A**<sub>2</sub>) Source leaf, **B**<sub>2</sub>) Sink leaf, **C**<sub>2</sub>) Petiole cross-section, **D**<sub>2</sub>) Upper stem cross-section, **E**<sub>2</sub>) Storage root cross-section, **F**<sub>2</sub>) Fibrous roots. *pAtRBCS3B::GUS* = **A**<sub>3</sub>) Source leaf, **B**<sub>3</sub>) Sink leaf, **C**<sub>3</sub>) Petiole cross-section, **D**<sub>3</sub>) Upper stem cross-section, **E**<sub>3</sub>) Storage root cross-section, **F**<sub>3</sub>) Fibrous roots. *pMeGBSS1::GUS* = **A**<sub>4</sub>) Source leaf, **B**<sub>4</sub>) Sink leaf, **C**<sub>4</sub>) Petiole cross-section, **D**<sub>4</sub>) Upper stem cross-section, **E**<sub>4</sub>) Storage root cross-section, **F**<sub>4</sub>) Fibrous roots. *pStSSS3::GUS* = **A**<sub>5</sub>) Source leaf, **B**<sub>5</sub>) Sink leaf, **C**<sub>5</sub>) Petiole cross-section, **D**<sub>5</sub>) Upper stem cross-section, **E**<sub>5</sub>)

Storage root cross-section, **F<sub>5</sub>**) Fibrous roots. *pStSTP1::GUS* = **A<sub>6</sub>**) Source leaf, **B<sub>6</sub>**) Sink leaf, **C<sub>6</sub>**) Petiole cross-section, **D<sub>6</sub>**) Upper stem cross-section, **E<sub>6</sub>**) Storage root cross-section, **F<sub>6</sub>**) Fibrous roots. Plants were either grown on the field at NCHU experimental station Taichung, Taiwan or in a greenhouse in Erlangen, Germany. Tissues from approximately 3-month-old cassava plants were used.

#### FIGURE S2

Representative GUS staining pattern of three *pCaMV35S::GUS* promoter-reporter lines. **A)** Source leaf, **B)** Sink leaf, **C)** Emerging leaves, **D)** Petiole cross-section, **E)** Upper stem cross-section, **F)** Lower stem cross-section, **G)** Storage root cross-section, **H)** Fibrous roots.

The authors apologize for these errors 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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