

Corrigendum: the comparison of expressed candidate secreted proteins from two arb...

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A corrigendum on

The Comparison of Expressed Candidate Secreted Proteins from Two
Arbuscular Mycorrhizal Fungi Unravels Common and Specific Molecular Tools
to Invade Different Host Plants

by Kamel, L., Tang, N., Malbreil, M., San Clemente, H., Le Marquer, M., Roux,
C., et al. (2017). *Front. Plant Sci.* 8: 124. doi: 10.3389/fpls.2017.00124

In the original article, there was an error in Table S1A concerning
Rhizophagus irregularis in planta RNAseq libraries.

The corrected Table S1A has been published as Supplementary material.

Accordingly, an addendum has been made to Materials and Methods section,
RNA Production and Sequencing, first paragraph:

“ Total RNA extraction and sequencing were performed according to
Tisserant et al. (2013) for *R. irregularis* and Tang et al. (2016) for *G. rosea* .
Apart from ERM of *G. rosea* where short paired- end sequencing reads were
obtained from Illumina Miseq1000 protocols (2 × 151 bp), all libraries were
obtained from Illumina Hiseq2000 protocols (2 × 101 bp). Library
constructions and sequencing were performed on the GeT-PlaGe facility
(Toulouse, France), according to standard Illumina protocols. Data are
available at NCBI GEO portal (GSE67906 and GSE67911) for *G. rosea* , and at
NCBI Sequence Read Archive (SRR1027885) and at NCBI GEO portal
(GSE67926) (—see details on libraries in Table S1A) for *R. irregularis* .
Number of reads per libraries, representativeness of fungal reads in

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symbiotic tissues and variability of data are presented in Tables S1A, S5A for *R. irregularis* and *G. rosea* respectively.”

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.

Conflict of Interest Statement

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.