



# Correction to: High throughput transcriptome analysis of coffee reveals prehaustorial resistance in response to *Hemileia vastatrix* infection

Juan Carlos Florez<sup>1</sup> · Luciana Souto Mofatto<sup>2</sup> · Rejane do Livramento Freitas-Lopes<sup>1</sup> · Sávio Siqueira Ferreira<sup>1</sup> · Eunize Maciel Zambolim<sup>1</sup> · Marcelo Falsarella Carazzolle<sup>2</sup> · Laércio Zambolim<sup>3</sup> · Eveline Teixeira Caixeta<sup>1,4</sup>

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## Correction to:

**Plant Molecular Biology (2017) 95(6):607–623**  
<https://doi.org/10.1007/s11103-017-0676-7>

All the transcriptome sequencing data mentioned in the original article is publicly available at the National Center of Biotechnology Information (NCBI).

## Links to bioprojects:

- PRJNA353233: Coffea arabica x Coffea canephora Raw RNAseq reads (Híbrido de timor)  
<https://www.ncbi.nlm.nih.gov/bioproject/PRJNA353233>

- PRJNA353185: Coffea arabica cv caturra raw RNAseq reads (Caturra)  
<https://www.ncbi.nlm.nih.gov/bioproject/PRJNA353185>

## Links to SRAs:

- SRP103086: Coffea arabica x Coffea canephora Raw RNAseq reads (Híbrido de timor)  
<https://www.ncbi.nlm.nih.gov/sra/?term=SRP103086>
- SRP103087: Coffea arabica cv caturra raw RNAseq reads (Caturra)  
<https://www.ncbi.nlm.nih.gov/sra/?term=SRP103087>

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Juan Carlos Florez and Luciana Souto Mofatto have contributed equally to this work.

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The original article can be found online at <https://doi.org/10.1007/s11103-017-0676-7>.

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✉ Eveline Teixeira Caixeta  
eveline.caixeta@embrapa.br

<sup>1</sup> Instituto de Biotecnologia Aplicada à Agropecuária (BIOAGRO), BioCafé, Universidade Federal de Viçosa, Campus Universitário, Avenida P.H. Rolfs, s/n, Viçosa, MG, Brazil

<sup>2</sup> Laboratório de Genômica e Expressão, Departamento de Genética, Evolução e Bioagentes, Instituto de Biologia, Universidade Estadual de Campinas, Cidade Universitária Zeferino Vaz, Distrito de Barão Geraldo, Campinas, SP 13083-970, Brazil

<sup>3</sup> Departamento de Fitopatologia, Universidade Federal de Viçosa, Campus Universitário, Avenida P.H. Rolfs, s/n, Viçosa, MG, Brazil

<sup>4</sup> Embrapa Café, Instituto de Biotecnologia Aplicada à Agropecuária (BIOAGRO), Universidade Federal de Viçosa, Campus Universitário, Avenida P.H. Rolfs, s/n, Viçosa, MG, Brazil

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