## Luciano Vilela Paiva

## List of Publications by Year

 in descending orderSource: https:/|exaly.com/author-pdf/3629717/publications.pdf
Version: 2024-02-01


[^0]Characterization of a Putative Serk-Like Ortholog in Embryogenic Cell Suspension Cultures of Coffea
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arabica L.. Plant Molecular Biology Reporter, 2014, 32, 176-184.
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A putative BABY BOOM-like gene (CaBBM) is expressed in embryogenic calli and embryogenic cell
3 suspension culture of Coffea arabica L. In Vitro Cellular and Developmental Biology - Plant, 2015, 51,
$2.1 \quad 25$
93-101.
$4 \quad$ Validation of reference genes for qPCR analysis of Coffea arabica L. somatic embryogenesis-related tissues. Plant Cell, Tissue and Organ Culture, 2017, 128, 663-678.
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5 In Silico and Quantitative Analyses of MADS-Box Genes in Coffea arabica. Plant Molecular Biology
Reporter, 2010, 28, 460-472.
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In Silico and Quantitative Analyses of the Putative FLC-like Homologue in Coffee (Coffea arabica L.).
Plant Molecular Biology Reporter, 2012, 30, 29-35.
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Gene expression and morphological characterization of cell suspensions of Coffea arabica L. cv.
$7 \quad$ CatiguÃ $\tilde{j}_{i}$ MG2 in different cultivation stages. Acta Physiologiae Plantarum, 2015, 37, 1.
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8 Genome-wide analysis, transcription factor network approach and gene expression profile of CH 3 genes over early somatic embryogenesis in Coffea spp. BMC Genomics, 2019, 20, 812.

In silico and in vivo analysis of ABI3 and VAL2 genes during somatic embryogenesis of Coffea arabica:
9 competence acquisition and developmental marker genes. Plant Cell, Tissue and Organ Culture, 2019, 137, 599-611.

10 Gene expression in two contrasting hybrid clones of Eucalyptus camaldulensis $\times$ Eucalyptus
urophylla grown under water deficit conditions. Journal of Plant Physiology, 2018, 229, 122-131.
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11 Gene Expression Profile Analysis is Directly Affected by the Selected Reference Gene: The Case of Leaf-Cutting Atta Sexdens. Insects, 2018, 9, 18.

Analysis of gene co-expression networks of phosphate starvation and aluminium toxicity responses in
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Populus spp.. PLoS ONE, 2019, 14, e0223217.
Comprehensive characterization of the ALMT and MATE families on Populus trichocarpa and gene
13 co-expression network analysis of its members during aluminium toxicity and phosphate starvation
$2.2 \quad 5$
stresses. 3 Biotech, 2020, 10, 525.

Transcriptional analysis of WUSCHEL-related HOMEOBOX (WOX) genes in Coffea arabica L.. Biologia
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(Poland), 2020, 75, 1483-1495.

15 The SAUR gene family in coffee: genome-wide identification and gene expression analysis during somatic embryogenesis. Molecular Biology Reports, 2022, 49, 1973-1984.
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[^0]:    1 Proteomic analysis of coffee grains exposed to different drying process. Food Chemistry, 2017, 221, 1874-1882.

