



PROGRAM AND ABSTRACT BOOK

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Biomolecular and Biotechnology

Drug Development and Nutraceutical

Genetic Resources and Uses

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Program and Abstract Book

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A blue microphone with a perforated grille is the central focus, positioned on a stage. The background is a blurred conference room with round tables covered in white and red cloths, and rows of white chairs. The lighting is warm and ambient, typical of an indoor event space.

**INVITED
SPEAKERS**

Transcriptional and post-transcriptional regulation of genes involved in the production and scavenging of reactive oxygen species and antioxidant biosynthesis in *Hevea Brasiliensis* Laticifers

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Hevea brasiliensis is the main source of natural rubber accounting for 42 % of the worldwide rubber consumption. Natural rubber is synthesized in rubber particles of latex cells, which are differentiated from the vascular cambia and localized in the secondary phloem. Latex is collected by tapping the soft bark of rubber trees. Ethephon, an ethylene releaser, is applied on bark to stimulate latex flow and regeneration between two tappings. Above a certain threshold, environmental and harvesting stresses are known to induce an oxidative stress triggering Tapping Panel Dryness (TPD) [1]. TPD is a physiological syndrome affecting latex production through the agglutination of rubber particles. Four hundred and seven genes from thirty gene families related to reactive oxygen species (ROS) production and scavenging, and antioxidants biosynthesis genes were identified in the *Hevea* genome sequence [2]. Based on a transcriptome analysis [3], 161 ROS-related genes were found expressed in latex cells. Small RNA and degradome analysis revealed 13 genes targeted by 11 microRNAs and 15 genes targeted by 16 phased siRNA in latex. These post-transcriptional regulations dramatically affect their gene expression profile. HbRBOH2 was identified as the main source gene of ROS in latex, while HbCuZnSOD4 might be the most important ROS scavenging enzyme for ROS detoxification in latex. Overexpression of genes encoding a superoxide dismutase (HbCuZnSOD) and an enzyme of the glutathione biosynthetic pathway (EcGSH1) was successfully obtained in transgenic rubber plants. These latter showed an increase in plant growth and their tolerance to abiotic stress [4, 5]. This study revealed the crucial role of antioxidant in *Hevea brasiliensis* laticifers and suggests to seek genetic variability for antioxidant capacity in order to improve rubber tree for the tolerance to abiotic stress and TPD occurrence.

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