

Genome-wide identification and characterization of the late embryogenesis abundant (LEA) protein-encoding gene family related to water deficit response in *Theobroma cacao*

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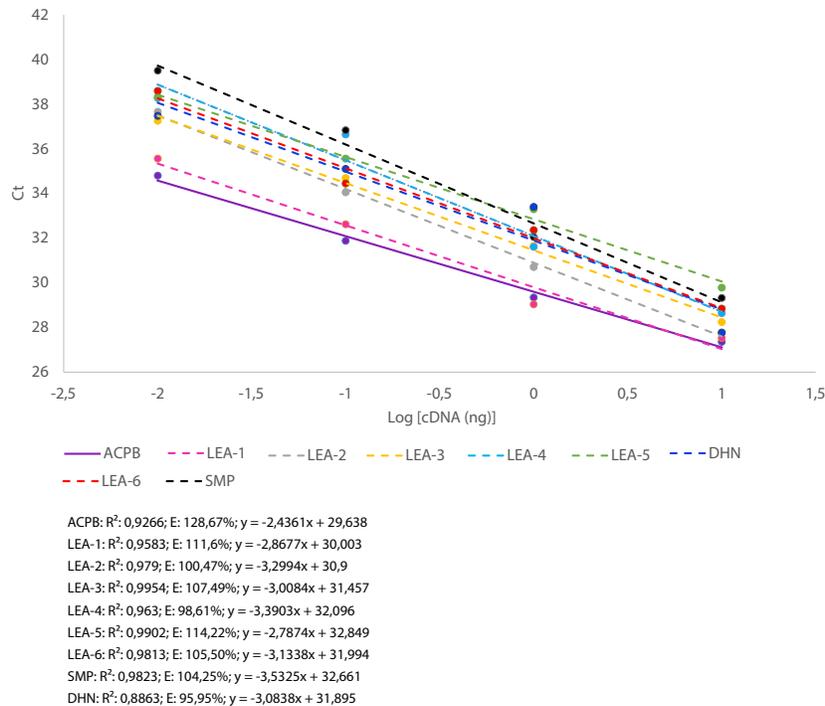
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Online Resource 4 Standard curves of all RT-qPCR targeted genes. The plotted values correspond to the averages resulting from the serial dilutions 15.00, 1.50, 0.15 and 0.015 ng of cDNA, (n = 3). For each gene, the values of R^2 , amplification efficiency (E) and equation of the linear regression are presented, where Y axe corresponds to the Ct value obtained and X axe corresponds to Log [cDNA] in nanograms.

