

Supplementary File S1

Genome-wide expression analysis and Metabolite Profiling Elucidate Transcriptional Regulation of Flavonoid Biosynthesis and Modulation under Abiotic Stresses in Banana

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>GSMUA_Achr6T08170_001_(MaCHS1)_Genomic

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>GSMUA_Achr4P16830_001_(MaCHI1)_Protein

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>GSMUA_Achr2T02770_001_(MaF3H1)_Genomic

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>GSMUA_Achr7P16780_001 (MaF3H2)_Protein

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>GSMUA_Achr3P07030_001_(MaFLS1)_Protein

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>GSMUA_Achr10T25240_001_(MaFLS3)_Genomic

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>GSMUA_Achr3P30350_001_(MaF3'H)_Protein

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>GSMUA_AchrUn_randomP12120_001_(MaF3'5'H7)_Protein

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>GSMUA_Achr3P31110_001_(MaDFR1)_Protein

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>GSMUA Achr4T10640 001 (MaDFR2) Modified CDS

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>GSMUA_Achr4P10640_001_(MaDFR2)_Protein

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