

Supplementary material

Table S1. Primers of genes analyzed by quantitative real-time reverse-transcription polymerase chain reaction.

Function	Gene	Primer Sequences	GenBank or TC TIGR* Accession Number
<i>Reference genes</i>	<i>EF1</i> (EF1- α elongation factor)	5'-GAACTGGGTGCTTGTAGGC-3' 5'-AACCAAAATATCCGGAGTAAAGA-3'	GU585871
	<i>60SRP</i> (60S ribosomal protein L18)	5'-ATCTACCTAAGCTCCTAGTC-3' 5'-CAATCTTGTCTCCTCTTCCT-3'	XM_002270599
	<i>39SRP</i> (39S ribosomal protein L41)	5'-AACCAAAATATCCGGAGTAAAGA-3' 5'-GACTGACTTCAAGCTTAAACC-3'	XM_002285709
<i>Phenylpropanoid metabolism</i>	<i>PAL</i> (Phenylalanine ammonia lyase)	5'-TCCTCCCGAAAACAGCTG-3' 5'-TCCTCCAATGCCCAAATCA-3'	X75967
	<i>STS</i> (Stilbene synthase)	5'-AGGAAGCAGCATTGAAGGCTC-3' 5'-TGCACCAAGGCATTCTACACC-3'	FJ851185
	<i>CHIT4C</i> (Chitinase class V)	5'-TCGAATGGGATGGTGAAA -3' 5'-TCCCCTGTCGAAACACCAAG -3'	NM_001281244
	<i>GLUC</i> (β -1,3 glucanase)	5'-TCAATGGCTGCAATGGTGC-3' 5'-CGGTCGATGTTGCGAGATTAA-3'	DQ267748
<i>Defense protein</i>	<i>PR1</i> (pathogenesis-related protein 1)	5'- GGAGTC CATTAGCACT CCTTG -3' 5'- CATAATTCTGGCGTAGGCAG -3'	XM_002273752
	<i>PR10</i> (pathogenesis-related protein 10)	5'- CGTTAAGGGCGGCAAAGAG -3' 5'- GCATCAGGGTGTGCCAAGA -3'	DQ396809
<i>Detoxification and Stress tolerance</i>	<i>GST1</i> (Glutathione s-transferase 1)	5'- TGCATGGAGGAGGAGTTCGT -3' 5'- CAAGGCTATATCCCCATTTCTT -3'	NM_001281248
	<i>PME25</i> (pectin esterase inhibitor)	5'-TGTCGGAGCTGAATGTGGA-3' 5'-CGATCCCGCTAACGAGAGTT-3'	XM_002273391.4
	<i>MSR</i> (Peptide methionine sulfoxide reductase))	5'-GCATTTGGCGTGTAGATGG-3' 5'-GCTTCAGTGTGGCCTGTTCT -3'	XM_010662104.2
<i>Arsenite recovery genes</i>	<i>WRKY</i> (transcription factors)	5'-GGGCAGAAGGACATTCTGGA -3' 5'-AGGGATCTCATCCGAACGC -3'	XM_002272684.3
	<i>HYD2</i> (ABA 8' hydroxylase 2)	5'-GAAGCTTGTCTCCGAGCCT -3' 5'-TATTGATCGACCGGCCTCC -3'	NM_001281052.1

* see <http://www.jcvi.org/cms/research/projects/tdb/overview/>

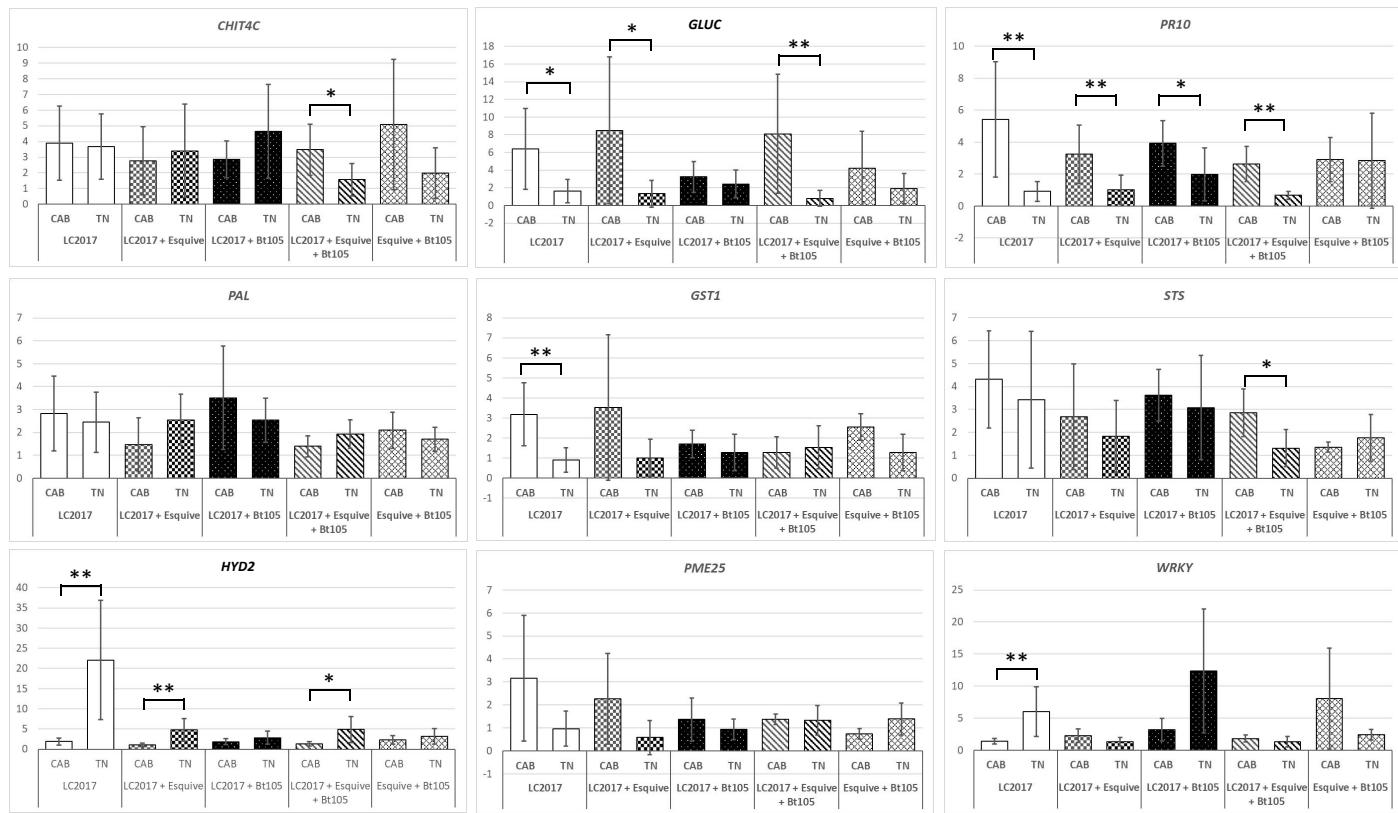


Figure S1: Comparison of the 9 targeted genes with induction values >2 recorded at T_0+5d on the different condition of treatments (LC2017 alone, LC2017 and Esquive) between Cabernet Sauvignon (CAB) and Touriga Nacional (TN) not infected or inoculated with *L. theobromae* (Bt105). The asterisk indicates a statistical significance (Mann-Whitney U test) between the medians of the two considered condition. * $= p<0.05$; ** $= p<0.01$