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Supplemental Tables

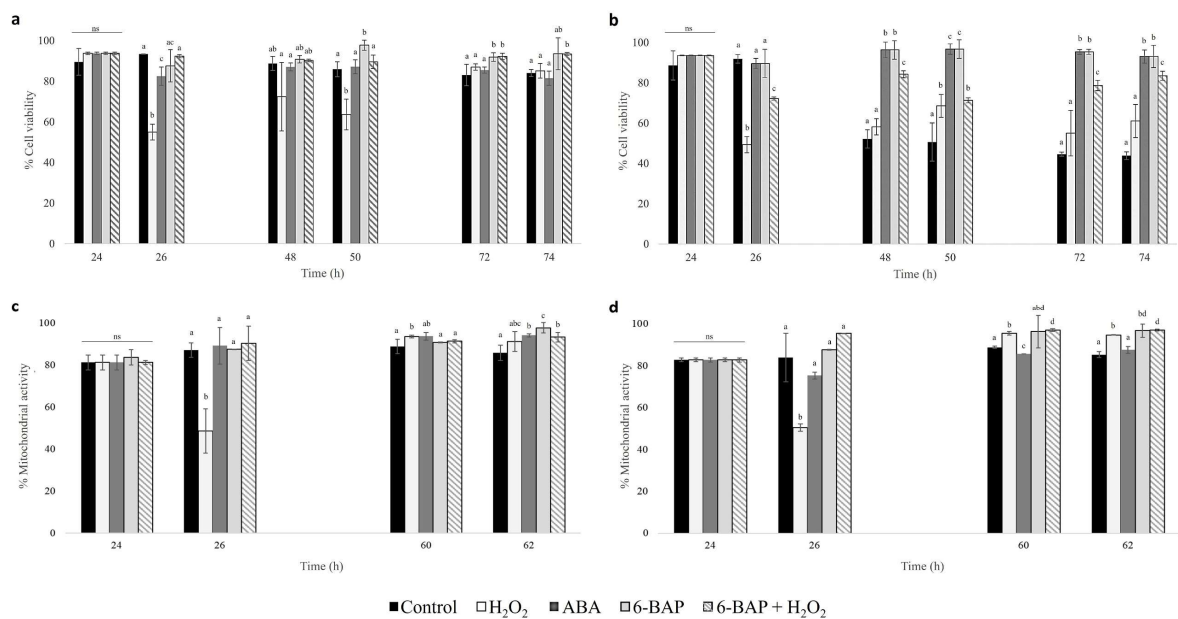
Supplemental Table S1. Mature transcript gene-specific primers used for RT-qPCR analysis in this study.

Primer name*	Sequence (5'-3')	Target gene cDNA	Source
mact-F	CCGCCCTCGTGATTGATAAC	<i>actin</i>	Casteblanco-Matiz et al. 2015
mact-R	TCACCAACGTAGGAGTCCTT	<i>actin</i>	Casteblanco-Matiz et al. 2015
mmhmgR-F	CATTGATGGAGGCTGTGAACG	<i>hmgR</i>	This study
mmhmgR-R	GTCGATCGATTGTGCTTCATGTG	<i>hmgR</i>	This study
mmidi-F	GAGTACGATGAGGAGCAGGTCAG	<i>idi</i>	This study
mmidi-R	GTCGAACACCGATCTGGTTC	<i>idi</i>	This study
mmcrtYB-F	AGACAGCGGAAGAATACCGAC	<i>crtYB</i>	This study
mmcrtYB-R	GAACAAACGATGGAAAGGAGGA	<i>crtYB</i>	This study
mmcrtR-F	CTGGGAAACAAGACCTACGA	<i>crtR</i>	This study
mmcrtR-R	GGAACCTCGGTTACGACAAA	<i>crtR</i>	This study
mmcrtS-F	ATGGCTCTTGCAGGGTTTGA	<i>crtS</i>	Casteblanco-Matiz et al. 2015
mmcrtS-R	TGCTCCATAAGCTCGATCCCAA	<i>crtS</i>	Casteblanco-Matiz et al. 2015

mm: mature transcript

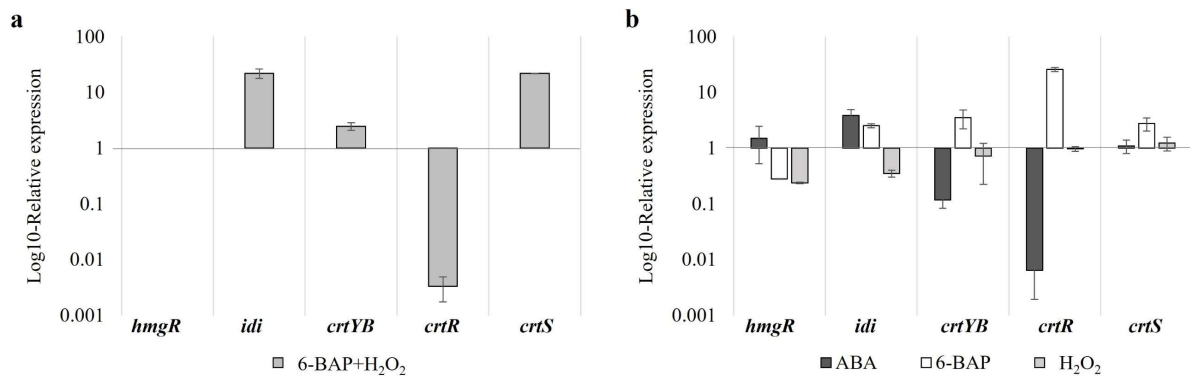
*F (forward) and R (reverse) in the primer name indicate the primer orientation.

22 Supplemental Figures



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24 **Supplemental Fig. S1.** Cell viability and mitochondrial activity of stimulated *X. dendrorhous* cultures determined by
25 flow cytometry. **a)** Cell viability of stimulated *X. dendrorhous* grown in a CD medium enriched with glutamate, **b)**
26 cell viability of stimulated *X. dendrorhous* grown in CD culture medium formulated with 0.1 g/L glutamate (control
27 concentration). Cell viability was evaluated times before stimulating agents addition: 24, 48 and 72 h, and,
28 subsequently, at evaluated times after stimulating agents addition at 26, 50 and 74 h. **c)** Mitochondrial activity in
29 stimulated *X. dendrorhous* grown in a CD medium enriched with glutamate, and **d)** mitochondrial activity for
30 stimulating agents in a CD culture medium formulated at 0.1 g/L of glutamate (control concentration). Mitochondrial
31 activity was evaluated times before stimulating agents addition: 24 and 60 h, and, subsequently, at evaluated times
32 after stimulating agents addition at 26 and 62 h. Lettering shows means that data are significantly different at evaluated
33 time (one-way statistical analysis; p -value ≤ 0.05) and ns (not significant) indicate p -value ≥ 0.05 .

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37 **Supplemental Fig. S2.** RT-qPCR expression analysis of *X. dendrorhous* ATCC 24202 genes involved in
 38 carotenoids/astaxanthin biosynthesis under stimulation conditions at 38 h fermentation. **a)** Relative expression levels
 39 of genes using 6-BAP and H₂O₂ combination treatment and **b)** relative expression levels using ABA, 6-BAP and H₂O₂
 40 individual treatments. Each transcriptional level was normalized to the expression of the housekeeping actin gene and
 41 expressed as a function of the control treatment (relative expression = 1). The values are the mean of four independent
 42 experiments, and the error bars correspond to the standard deviations. *hmgR*, hydroxymethylglutaryl-CoA reductase;
 43 *idi*, isopentenyl pyrophosphate isomerase; *crtYB*, phytoene-β-carotene synthase; *crtR*, cytochrome P450 reductase;
 44 *crtS*, astaxanthin synthase.