

**Transcriptome profiling of ripening nectarine (*Prunus persica* L. Batsch) fruit treated with 1-MCP. Fiorenza Ziliotto, Maura Begheldo, Angela Rasori, Claudio Bonghi, and Pietro Tonutti**

**SUPPLEMENTARY DATA**

**Table S1.** Primers (L: left and R: right) used for qRT-PCR analyses.

<b>Gene</b>	<b>Primer L/R</b>
<i>AUX/IAA</i> <i>Ctg_768</i>	5-CTGGATGTTAGTTGGAGACGTCCCTTGG-3 5-GTGGCAGAACCTGAGACGTCCTTGGAG -3
<i>β carotene hydroxylase</i> <i>Ctg_711</i>	5-GCACGAGTCTCACCACCGACCCAGAG-3 5-TCCAAGACCAGCACCAAAACATAGGC-3
<i>Catalase</i> <i>Ctg_1024</i>	5-GATCCTGTTCGTCATGCTGAGCGCTAC-3 5-GCGCCCAAGATCTGTATCTCTCTCCA-3
<i>ERF-like</i> <i>Ctg_2757</i>	5-CGCATACATGGCTCCAAGGCTTTGCTC-3 5-CTGGGACAAAGGGAAGGCCGTCAC-3
<i>ETR2</i> <i>Ctg_4109</i>	5-CTCATGAGGGAGAAGTTGGCCGAGC -3 5-GCCTCCTCATCCCATCACTCATTACC-3
<i>TPPA</i> <i>Ctg_4621</i>	5-CCGCAATGTAGATGAGAAGAGTTGGCC-3 5- GTGACGGCCTTCCCTTTGTCCCAG -3