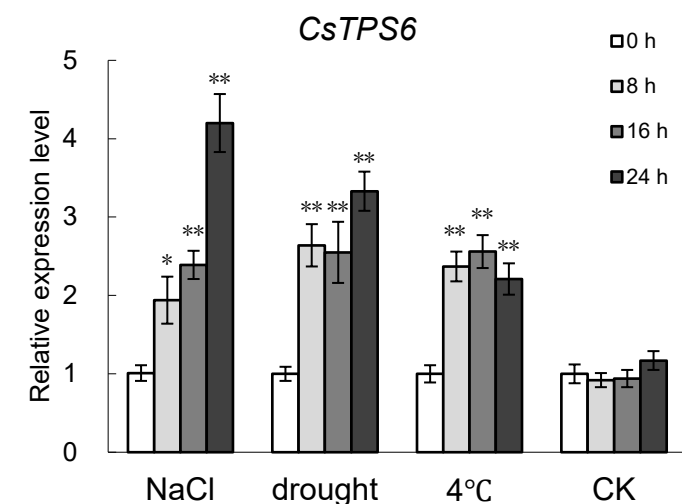
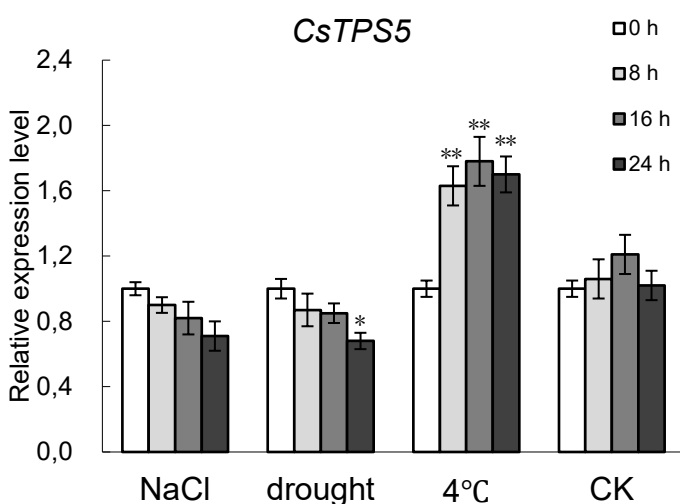
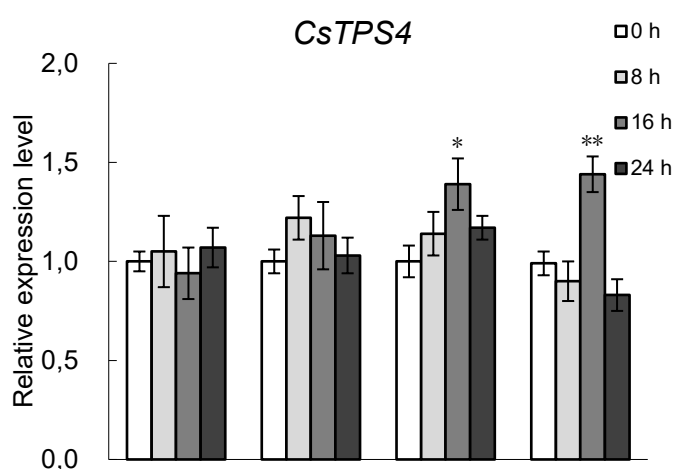
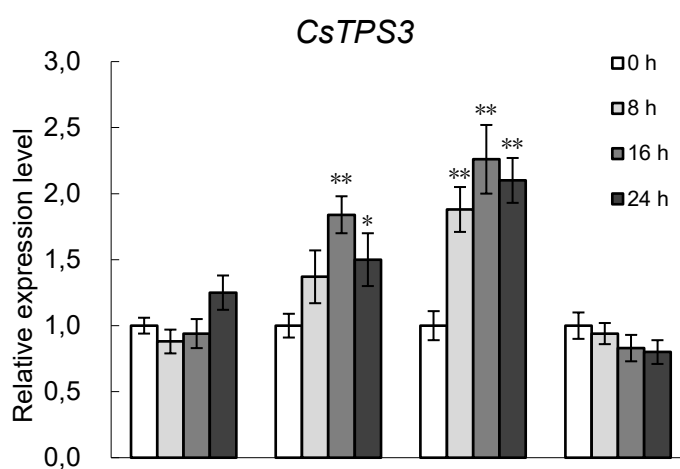
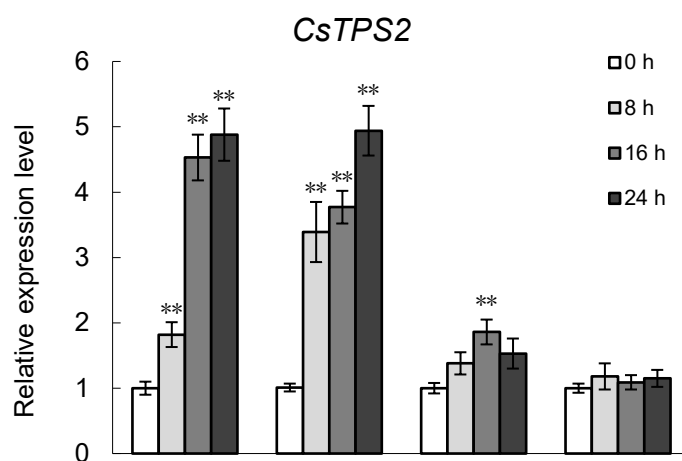
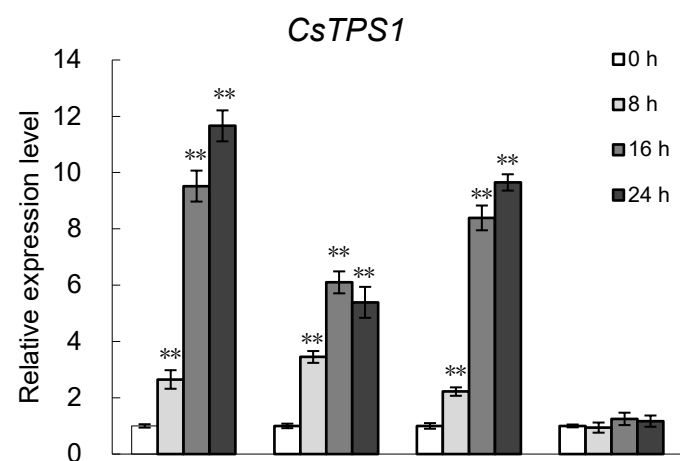
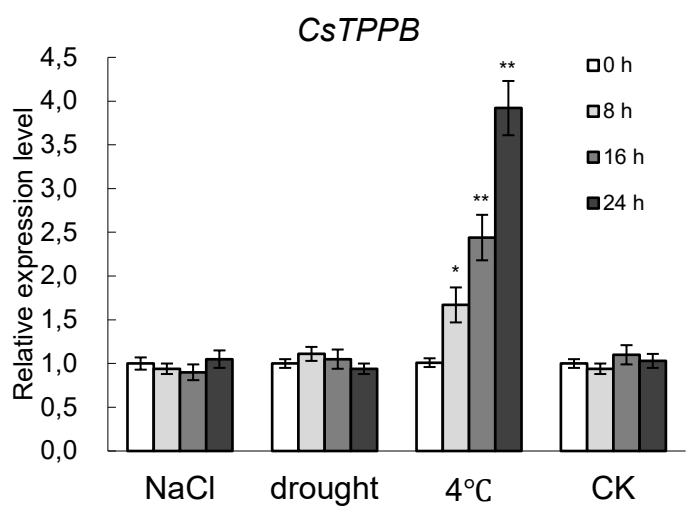
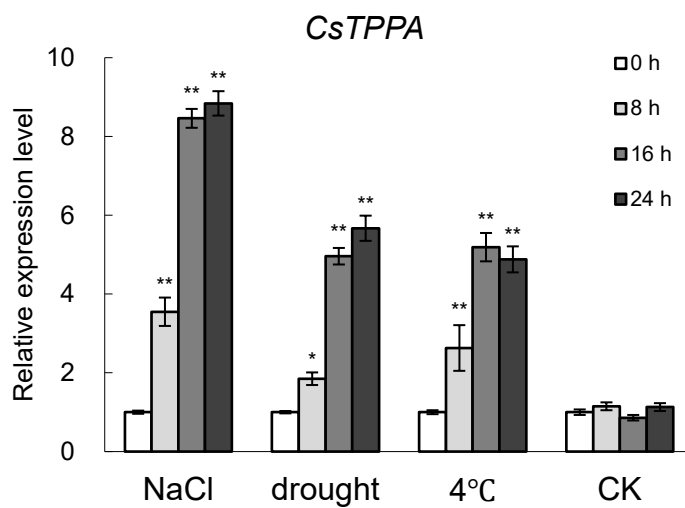
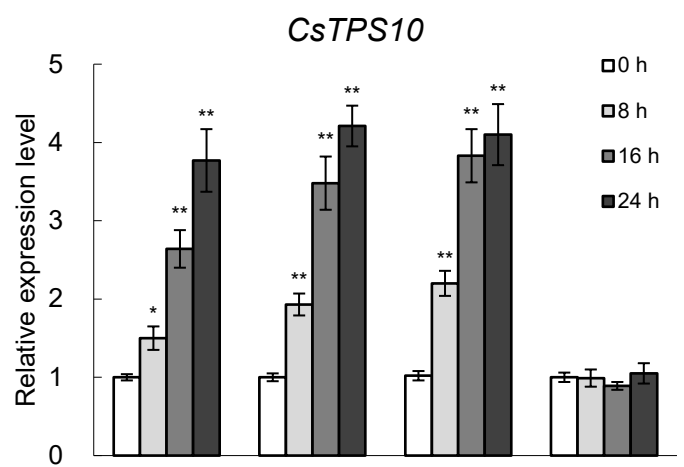
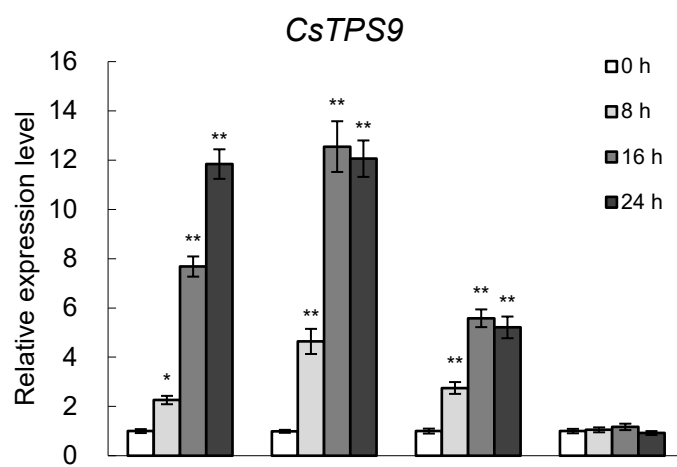
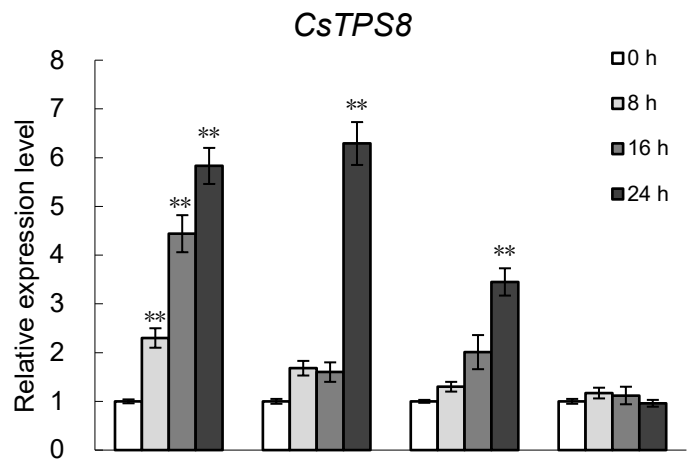
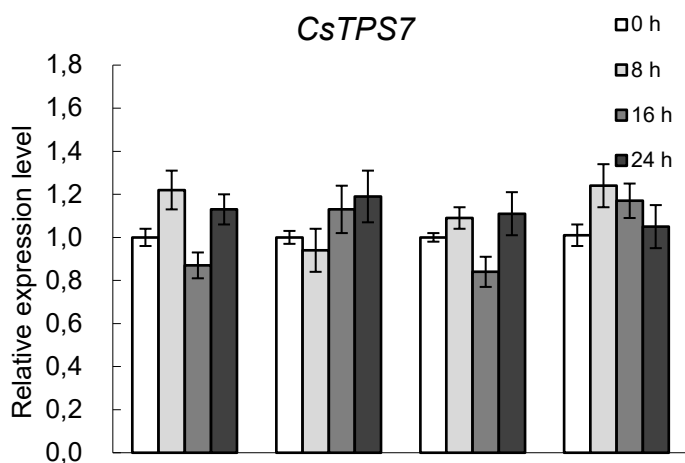


Table 1 Suppl. Specific primers for *CsTPS* and *CsTPP* genes.

Gene	Primer sequence (5'-3')	Product size [bp]
<i>CsTPS1</i>	F: CAAAGGAAGTGAGGGTC R: TCAAGTACATTCCACGAA	192
<i>CsTPS2</i>	F: ACGACCTTCTAAACAGCG R: CAGGCACATTTACACCAG	176
<i>CsTPS3</i>	F: TTCTTGCCTCATAACAGG R: GAGATATTGAAGCAGGTC	213
<i>CsTPS4</i>	F: TTTGGGCAACCAGGCTAT R: TGTCACCACAAGGCAATC	102
<i>CsTPS5</i>	F: GGGTTATGATCCCGTTGT R: ATTCCATCCCTAACTGCT	108
<i>CsTPS6</i>	F: TTGAAAGATGGGTTAGGA R: TCAGGTGGGAGAAAAGTGGGA	143
<i>CsTPS7</i>	F: AGCTAAGCGTCGACTGGAT R: AGCATTCAAAGACCCCTAC	124
<i>CsTPS8</i>	F: GCCTCAAACCTCCATCTC R: TTCCCTCTGCCACTAACC	108
<i>CsTPS9</i>	F: CCAGTTACCGATACGAGC R: CACTTCGAGACAACCCAC	139
<i>CsTPS10</i>	F: TCATAAGGTCAAGCGGAACA R: ACAGGAACCGAAATCAGG	185
<i>CsTPPA</i>	F: AGATAACCCTGACTGTGC R: GTCCATCCCATGACTACC	163
<i>CsTPPB</i>	F: CGGCAATCATAAGTGGAA R: TTAGCAGGCTGGAAGAGC	163
<i>CsTPPC</i>	F: TATTTATTATGCTGGGAGTC R: GCATGGCAGGCAAGAACC	140
<i>CsTPPD</i>	F: AATAATCCTGCTTCTGTCT R: CTCGATCTGGGTCTTCTA	178
<i>CsTPPE</i>	F: GCCCAGCAAGTAAAGTCA R: TTTGCGAATCCAAGTGA	143
<i>CsTPPF</i>	F: ATTGGGATAAAGGTCGTG R: TCTGAGGGAGTATGAAGC	197
<i>ACTIN</i>	F: TTGCTGGTCGTGATCTTACTG R: GTCTCCATCTCCTGCTCAAAG	118





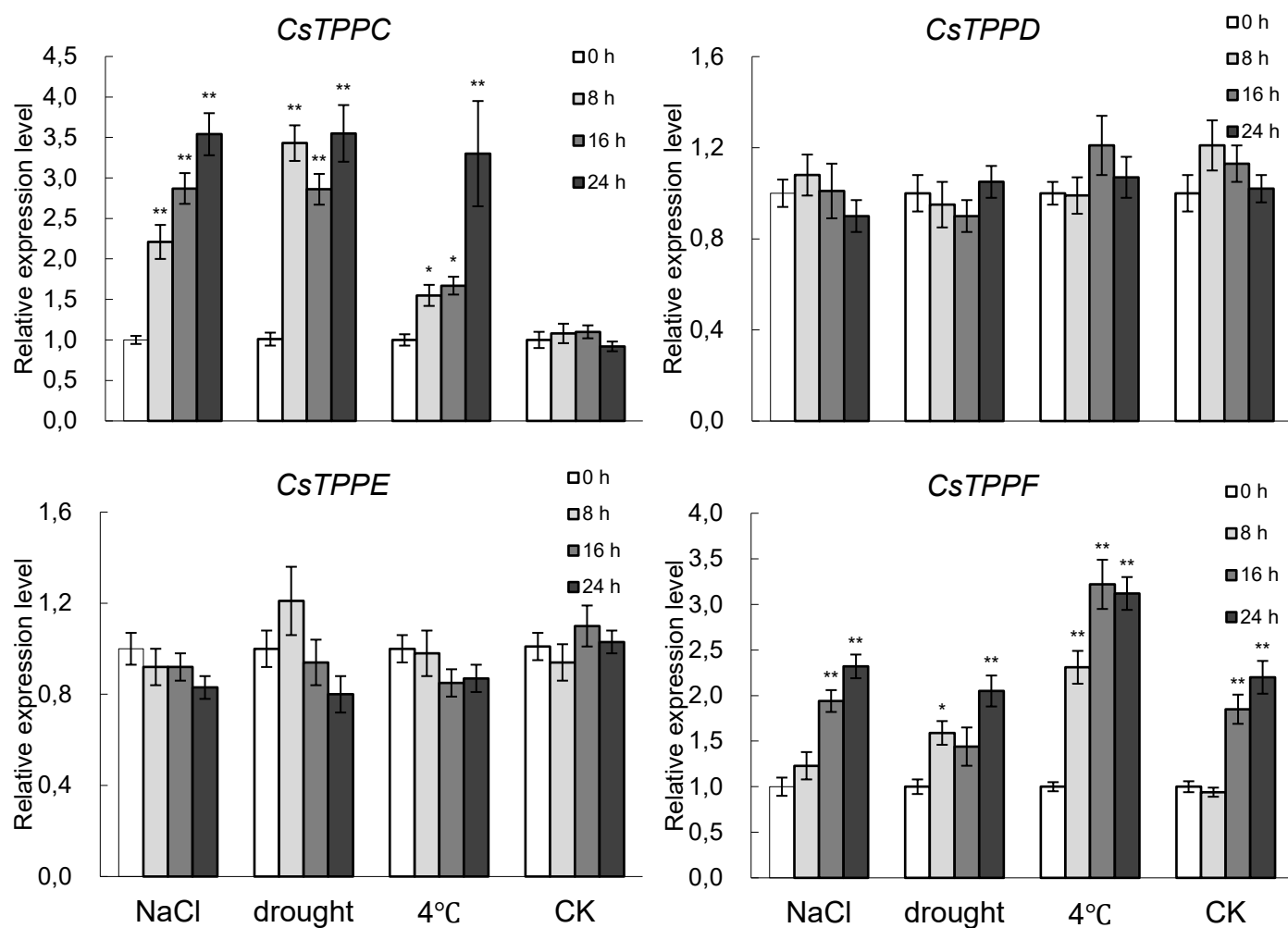


Fig. 1 Suppl. Expression patterns of TPS and TPP family genes in *Cannabis* under salt, drought, and low temperature stresses. The x-axis represents the different stress treatments (NaCl, PEG-6000, 4 °C) and control (CK). The y-axis represents the relative expression. The columns from left to right of each treatment represent the sampling time points of 0, 8, 16, and 24 h, respectively. Means $\pm$ SEs, * and ** - significantly different from 0 h at $P \leq 0.05$ and $P \leq 0.01$, respectively.