

## **Supplementary material**

**Overexpression of calmodulin-like (*ShCML44*) stress-responsive gene from *Solanum habrochaites*, enhances tolerance to multiple abiotic stresses**

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**Table S1.** List of primer sequences.

Gene	Forward primer	Reverse primer
<b>For gene amplification and qRT-PCR</b>		
<i>ShCML</i>	TCATGTCTCCGATCAACTCAAT	TTGGTAGATTGTAAGCGGGATT
$\beta$ -actin	ATGGCAGACGGAGAGGATATTCA	AGACGGAGAATGGCATGTGG
<i>EF1a</i>	CGTGGTTATGTTGCCCTAAA	ACAGCAATGTGGGAAGTGTG
<b>For promoter amplification</b>		
PCML	TCGGAGGCAGAAATTAAGGAA	TGACCGCAAATAATGATGCT
<b>For transgenic plants identification</b>		
<i>ShCML</i>	TCATGTCTCCGATCAACTCAAT	TTGGTAGATTGTAAGCGGGATT
CaMV35S	ACGCACAATCCCCTATCCTTC	
<b>For qRT-PCR</b>		
<i>CML</i>	ATGTCTCCGATCAACTCAAT	TACCAAGTAGCAACTCTAGC
<i>SOD</i>	TTTCTGGGAAGCAAATCGTC	TTGGGCAATGAAGAAGAAC
<i>Peroxidase</i>	CTTGCCCTAACATGCTCTCACC	GCATCACAAACCCCTGAACAAA
<i>GST</i>	GCAAGCCCATTGTGAGTCT	TGCTGACCCCTTATCATCG
<i>LOX</i>	TCATTTCCCCTGGCAAGTA	TGGTGCATTGGATCTTCCT
<i>PRI</i>	GGATTGACACCAGCAGAAC	CCATGTAACCCAATGCCTA
<i>PR2</i>	ATGGGCTGAAGGATCAGTTG	GACACTCTTCTTGAACTTGTTGG
<i>Catalase</i>	TCCCAGTTAACATGCTCCAAG	CTCAGCAGGACGACAAGGAT
<i>sucrose synthase</i>	GTGGAGACCGAAGAAAGGAA	GGGAAGAAATCCATCGGAAC
$\beta$ -amylase	GCATTCTGTGCCAACTTATCTT	GATTCACACTCCGTGCTTTC
<i>Hsp90-1</i>	GCACTTCTCTGTTGAAGGTCA	ATGAACACACGGCGAACATA
<i>ERD1</i>	CAGAACTGGCCTCAAGCAC	AGCATGGCAGATTCAAGATCC
<i>ERD2</i>	ATTTTGAGTGGCGAGGGTAA	TTGGGATAGTCGTGTTCTGG

## **Supplementary Figures legend**

**Figure S1.** Alignment of *ShCML44* nucleotide sequence with sequences from *S. lycopersicum* (*SlCML44*) and *S. pennellii* (*SpCML44*). Multiple sequence alignments were conducted using ClustalW2.

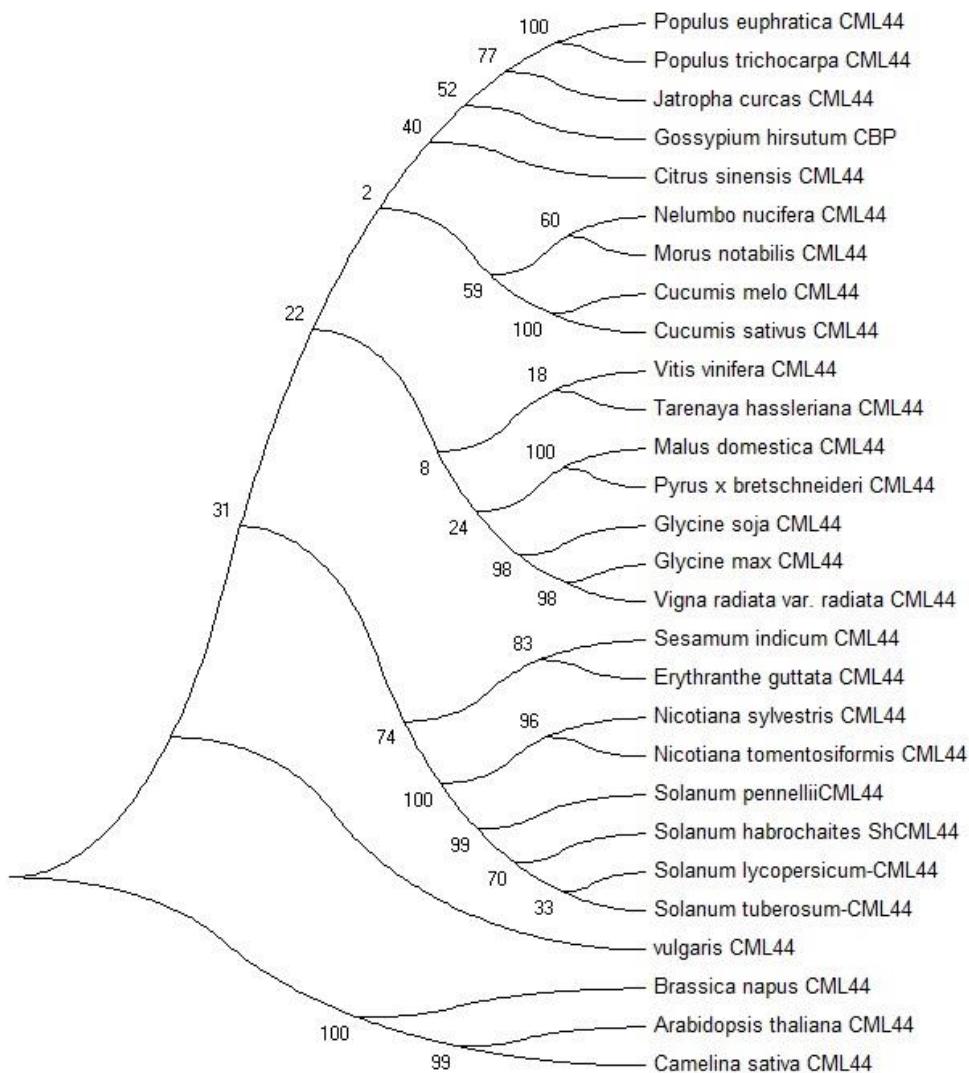
**Figure S2.** Phylogenetic analysis of *Solanum habrochaites* (*ShCML44*), *S. lycopersicum* (*SlCML44*) and *S. pennellii* (*SpCML44*) proteins with other species CML proteins.

**Figure S3.** *ShCML44* overexpression plants response towards drought tolerance in tomato. Comparison of plant height (a), number of leaves per plant (b), leaf length (c) leaf width (d), of transgenic and wild-type plants under normal and drought stress conditions.

**Figure S1**

S1CML ATGTCTCCGATCAACTCAATTAAATTGTCAAAAATCTTCTCAAAGCTTGACAAGAATGGT  
SpCML ATGTCTCCGATCAACTCAATTAAATTGTCAAAAGATCTTCTCAAAGCTTGACAAGAATGGT  
ShCML ATGTCTCCGATCAACTCAATTAAATTGTCAAAAATCTTCTCAAAGCTTGACAAGAATGGT  
\*\*\*\*\*  
  
S1CML GATGGCCTTGTGTCTTGATGAGTTAAAGGGATTCTTGATACAATAGGAATTATTGCA  
SpCML GATGGCCTTGTGTCTTGATGAGTTAAAGGGATTCTTGATACAATAGGAATTATTGCA  
ShCML GATGGCCTTGTGTCTTGATGAGTTAAAGGGATTCTTGATACAATAGGAATTATTGCA  
\*\*\*\*\*  
  
S1CML AGCCAAGAGGAGCTAGAGTTGCTACTTGGTAAAACAAGCCTAGACTCCATTGATTTC  
SpCML AGCCAAGAGGAGCTAGAGTTGCTACTTGGTAAAACAAGCCTAGACTTCATTGATTTC  
ShCML AGCCAAGAGGAGCTAGAGTTGCTACTTGGTAAAACAAGCCTAGACTCCATTGATTTC  
\*\*\*\*\*  
  
S1CML TTCTTCTATGATGCTATCACAAAGGCAAATATAAAAGGTAGCAATTATAAGCACGAG  
SpCML TTCTTCTATGATGCTATCACAAAGGCAAATATAAAAGGTAGCAATTATAAGCACGAG  
ShCML TTCTTCTATGATGCTATCACAAAGGCAAATATAAAAGGTAGCAATTATAAGACACGAG  
\*\*\*\*\*  
  
S1CML GATCGAGAAAATGTTTCTTGGAAAATGACCTACGTAAAGTGTAGAGTATTGATT  
SpCML GATCGAGAAAATGTTTCTTGGAAAATGACCTACGTAAAGTGTAGAGTATTGATT  
ShCML GATCGAGAAAATGTTTCTTGGAAAATGACCTACGTAAAGTGTAGAGTATTGATT  
\*\*\*\*\*  
  
S1CML AACGAGGACGGATTCATATGTTGTGAGGAGCTGCAAAGAGCATTGTCAGATTAGGATTG  
SpCML AACGAGGACGGATTCATATGTTGTGAGGAGCTGCAAAGAGCATTGTCAGATTAGGATTG  
ShCML AACGAGGACGGATTCATATGTTGTGAGGAGCTGCAAAGAGCATTGTCAGATTAGGATTG  
\*\*\*\*\*  
  
S1CML TGGGATGAAACATGTGGGAAAGATTGTAAGAGTATGATCAATGTTTATGACAAAATTAA  
SpCML TGGGATGAAACATGTGGGAAAGATTGTAAGAGTATGATCAATGTTTATGACAAAATTAA  
ShCML TGGGATGAAACATGTGGGAAAGATTGTAAGAGTATGATCAATGTTTATGACAAAATTAA  
\*\*\*\*\*  
  
S1CML GATGGAAAACCTGATTATGAGGAGTTAAAGACATGATGTTGATAATTAA  
SpCML GATGGAAAACCTGATTATGAGGAGTTAAAGACATGATGTTGATAATTAA  
ShCML GATGGAAAACCTGATTATGAGGAGTTAAAGACATGATGTTGATAATTAA  
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**Figure S2**



**Figure S3**

