

A novel isoprene synthase from the monocot tree *Copernicia prunifera* (Arecaceae) confers enhanced drought tolerance in transgenic Arabidopsis.

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Supplementary Table S1: Primers used in this study.

Primer name	Sequence (5' - 3')	Purpose
EgISPS_GW-Rev1	GGAAACCCGTGCTCTCAAGGAGTC	GenomeWalker
EgISPS_GW-Rev2	GTCATTTGAACATCAAGTCTGCCCTC	GenomeWalker
ISPS_For	CCAAACAATAGAACAGTAGCTCAAGC	full-length cDNA cloning
ISPS_Rev	CCCTTCTACACATTGCTGACC	full-length cDNA cloning
CprISPS_For	CACCATGGCATTCTCTACATGCTTCG	pENTR cloning
CprISPS_Rev	CTACACATTGCTGACCACCAACCT	pENTR cloning
pK7WG2_For	GACAATCTGATCCAAGCTCAAGC	Screening for transgene
CprISPS_RT_For	CAGCAATCAAATTTGGACGG	Semi-quantitative RT-PCR
CprISPS_RT_Rev	CTACACATTGCTGACCACCA	Semi-quantitative RT-PCR
DREB2A_RT_For	CTACAAAGCCTCAACTACGGAATAC	qRT-PCR
DREB2A_RT_Rev	AAACTCGGATAGAGAAATCACAGTC	qRT-PCR
COR15A_RT_For	GATA CATTGGTAAAGAAGCTGAGA	qRT-PCR
COR15A_RT_Rev	ACATGAAGAGAGAGGATATGGATCA	qRT-PCR
RAB18_RT_For	GGAAGAAGGAAATAACACAAAAAGAT	qRT-PCR
RAB18_RT_Rev	GCGTTACAAACCTCATTATTTTA	qRT-PCR
AtEF1α_RT_For	TGAGCACCGCTCTTCTGCTTTCA	qRT-PCR; Semi-quantitative RT-PCR
AtEF1α_RT_Rev	GGTGGTGGCATCCATCTGTTACA	qRT-PCR; Semi-quantitative RT-PCR
AtGAPDH_RT_For	TTGGTGACAACAGGTCAAGCA	qRT-PCR
AtGAPDH_RT_Rev	AAACTTGTCGCTCAATGCAAT	qRT-PCR

Supplementary Figure S1: Expression levels (semi-quantitative RT-PCR) of the *CpriSPS* transgene in the two Arabidopsis transgenic lines used in this study for all experiments. The gene *EF1 α* from Arabidopsis (*AtEF1 α*) was used as normalization control.

