

RESEARCH ARTICLE

Overexpression of annexin gene AnnSp2, enhances drought and salt tolerance through modulation of ABA synthesis and scavenging ROS in tomato

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Fig. S1 *Cis*-acting element analysis of AnnSp2 promoter. ABRE (TACGTG), *cis*-acting element involved in abscisic acid responsiveness. ATCT-motif (AATCTAATCT), part of a conserved DNA module involved in light responsiveness. BOX 4 (ATTAAT), part of a conserved DNA module involved in light responsiveness. GCN4-motif (TGTGTCA), *cis*-acting regulatory element involved in endosperm expression. G-BOX (CACGTC/TACGTG), *cis*-acting regulatory element involved in light responsiveness. HSE (AAAAAAATTTC), *cis*-acting element involved in heat stress responsiveness. SP1 (CC(G/A)CCC), light responsive element. TCA-element(GAGAAGAATA), *cis*-acting element involved in salicylic acid responsiveness. AT1-motif (ATTAATTTCACA/ AATTATTTTTATT), part of a light responsive module. AE-box (AGAACAT), part of a module for light response. GT1-motif (GGTTAA), light responsive element. TATC-box (TATCCC), gibberellin-responsive element. MBS (TAACTG/CAACTG), MYB binding site involved in drought inducibility. ARE (TGGTTT), *cis*-acting regulatory element essential for the anaerobic induction.

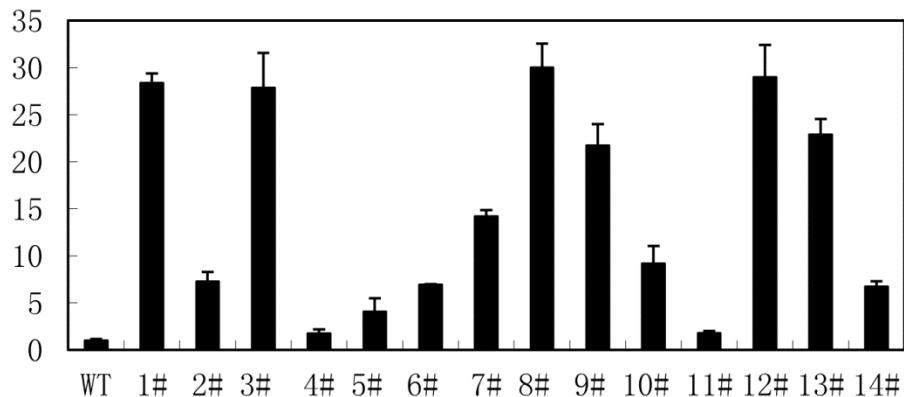
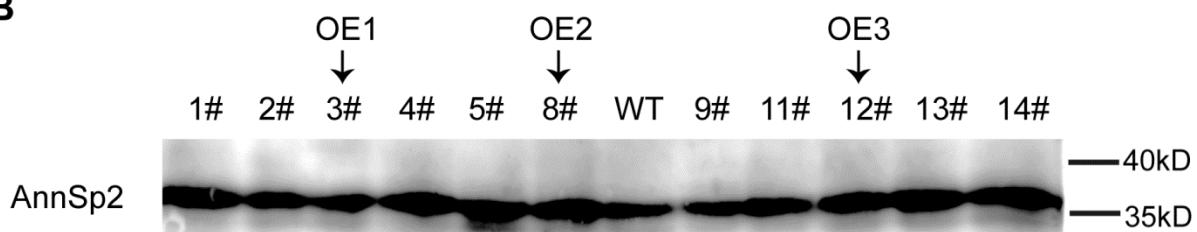
A**B**

Fig. S2 A Identification of transgenic plants and expression of AnnSp2. (A) q-PCR analysis of AnnSp2 expression in wild-type (WT) and T2 OE plants. (B) **Protein accumulation** in AnnSp2. Western blot analysis of AnnSp2 transgenic plants and in wild-type (WT) plants.

Table S1 Details of the primers used in this study

Primer name	Forward primer sequence	Reverse primer sequence	Purpose
AnnSp2	TTCCCACGTATATTCTTGAGATTG	CACACCATAACATCATAAAAGCAAA	Gene amplification
QAnnSp2	GATTCTTATGCTGCTGCTTATGGA	CAGTCAGACGTTGGTAGCCTCA	Quantitative real-time-PCR primer
AnnSP2	CCG <u>CTCGAG</u> ATGTCTAGTCTAAAGTTCCA	TGCT <u>CTAGAT</u> CAAGCATCTCCGTGCCAATC	Subcellular localization primer
CaMV 35S	TTGAATCCTGTTGCCGGTCT	GGAAGGGTCTTGCAGAAGGATAG	Transgenic plants confirmation
AREB	TTGCTGGTGGAAATGTAAGTGC	GGAATGTAACATCCTTGAGTATCG	qRT-PCR
SPERD	AGGCATCAAGTCATCACTCTGGT	GAGTAATGTGAGTAAGAACCAACG	qRT-PCR
DREB	AGTTGGGGAAAATGGGTGTC	CGAGGCAATGAGTCAATTAGGT	qRT-PCR
NCED	AGTTCCCGATTGGTATTTC	TGGAGTCTGGTGGTGTCA	qRT-PCR
P5CS	GCGATTCCCTGGAACTGTTGG	AGCATGGCCAAGAACAGGAA	qRT-PCR
β-actin	ACCTTCAATGTCCCTGCTATG	CTCCACCTTCAGAAACGCAAC	β-actin

Table S2 The genetic segregation analysis of over expression AnnSp2 plants in T1 generation by spraying kanamycin test

Number	Gene Name	T1 segregation		Single locus with 3:1 segregation*
		Green	yellow	
1	AnnSp2 -6	18	3	No
2	AnnSp2 -14	32	7	No
3	AnnSp2 -18	29	10	Yes
4	AnnSp2 -20	33	16	No
5	AnnSp2 -26	44	12	No
6	AnnSp2 -27	21	5	No
7	AnnSp2 -29	27	13	No
8	AnnSp2 -31	36	12	Yes
9	AnnSp2 -36	15	6	Yes
10	AnnSp2 -38	15	9	No
11	AnnSp2 -42	14	2	No
12	AnnSp2 -44	12	4	Yes
13	AnnSp2 -50	23	16	No
14	AnnSp2 -56	17	4	No

* The analysis based on chi square test