Supplementary Materials: Overexpression of S-Adenosyl-L-Methionine Synthetase 2 from Sugar beet M14 Increased *Arabidopsis* Tolerance to Salt and Oxidative Stress

Chunquan Ma, Yuguang Wang, Dan Gu, Jingdong Nan, Sixue Chen and Haiying Li

ATCTAC TCTCTCACCTCTTCG TC CAGGARATG GAGACC TTCC TATTCACCTCC GAG TC T H E T F L F T S E S ł H E T F L F T S E S GTGAAC GAG GGAC AC CC TGA CAAG CTCTGTGTGATCAGG TG TC TGATG CA GTGC TC GAC GC T 61 11 121 V N E G H P D K L C D Q V S D A V L D A TGCCTTGCCCAGGACCCTGAGAGCAAGGTTGCTTG TGAGAC TTGTACTAAGACAAACATG C L A Q D P E S K V A C E T C T K T N H GTTR TG GTC TT TG GTGRGRAT TR CRAC CRAG GCC GRGG TR GRC TATGARARGA TT GTC CG T 31 31 181 51 241 71 301 91 V H V F G E I T T K A E V D Y E K I V R GACACATEC CG TTCCATTEG CTTCAC GTCTGATGATE TE GE TC TTGATEC TEACAAATEC D T C R S I G F T S D D V G L D A D K C AAGG TC TTG GTCAATAT TGAGCAG CAGAGC CCAGACA TTGC ACAGG GTGTCCATGGTCAC E P D 27 Ι Ô Ô s A Ô CTCACCAAGCGCCCTGAGGAGATTGGTGCTGGTGATCAAGGCCACATGTTTGGCTATGCC 361 111 421 L T K R P E E I G A G D Q G H H F G Y A AC TGATGAGAC CCCG GAGCTTA TGCC TC TRAGC CACG TTCTTG CCACCAAGC TTGGTGC T 421 431 451 541 171 601 T D E T P E L H P L S H V L A T K L G A CGCCTCACTGAGGTGCGCAAGAATGGGACCTGTGCATGGCTGAGACCTGATGGTAAGACT E υ RKNG C A τσ L P D CARG TTACTGT TO AG TA CTA CARTGA CARTGGTGC CA TG GTAC CTG TTCG GG TTCACAC T Q V T V E Y Y N D N G A H V P V R V H T GTCCTTATATCCACCCAACACGATGAGACTGTCTCCAATGATGAAATTGCTGCTGTCTT 191 661 211 721 V L I S T Q H D E T V S N D E I A A D L ARGGAG CATGTCA TCARACC CG TCATCC CTGAGAAGTATCTTGATGAGAAGACCATC TTC K E H V I K P V I P E K Y L D E K T I F CACCTTAATCCATCTGGCCGATTTGTTATTGGTGGACCTCATGGAGATGCTGGTCTCACT H L N P S G R F V I G G P H G D A G L T GG TC GCAAGATCATTATCGACACA TACG GT GG TG GG TG GG GG GG C TC ATG GT GG TG GT GC TT TC 231 781 251 D т G TO THE GOARA GACC CAAC CARGE TO GACA GRAGTEG THE CT TA TA TOS TO AGAC AS SCT SCT 841 271 901 S G K D P T K V D R S G A Y I V R Q A A ARGAGCATAGTTS CCAATGG GC TC GC AGG GC CATTGTTC AGG TG TC ATATGCCATT K S I V A N G L A R R A I V Q V S Y A I GG TG TC CCTGAGC CTTTGTC CG TG TTTG TTGACAC CTATGG CACTG GAAA GA TACCC GAT G V P E P L S V F V D T Y G T G K I P D 291 961 311 311 G V P E P L S V F V D T Y G T G K I P D 1021 ARAGAGATTCTTRAGATTGTCAAGGAGACCTTTGACTTCAGGCCAGGAATGATGTCTATC 331 K E I L K I V K E T F D F R P G H H S I 331 K E I L K I V K E T F D F R P G H H S I 1081 ARCCTTGATCTCAAGAGAGGTG GCAA TG GCAGATTCCAGAAGACAGCTGCATAC GGG CAC 1001 IN L D L K R G G N G R F Q K T A A Y G N 1141 ITTS GAAGAGA TGAC CCAGACTTCAC CTGG GAG GAC CCC TGAAGTGG GAGAAA 371 F G R D D P D F T W E V V K P L K W E K 1201 ATCCCTGCG TRATGC CCACTARATGC TACTGCTGC TCAAGTAC GCG GAACCCCATGCA TGC I P A * TGATCC CCTTTTTTTATGTC: TGATCT TTTTCCA TTGAATTTTTTCCAA TTTC GA TTC GA T TCATGTACTGA TTGAAA TGAAA TG GA TTAC TGT TTGC TG TTTTGGTAC TTGAAAGTAGC C 391 1261 1321 TRAAGC GTTGC TS AA TC GTC GA CC TS CAGG CATGC AAGC TS GA TSA TT TS GA TS CTACTA 1381 ARCC CTATTAR TTGARGATR TTGTAC TR GRACARA TO CR TO CR TTT TR TO TRGT TO TCA T TC GRATCTRGC GRARARARARARARARARARARARARA 1441

Figure S1. Sequence analysis of a cDNA encoding a *BvM14-SAMS2* isolated from the monosomic addition line M14 roots. Nucleotide and deduced amino acid sequence of *BvM14-SAMS2*.

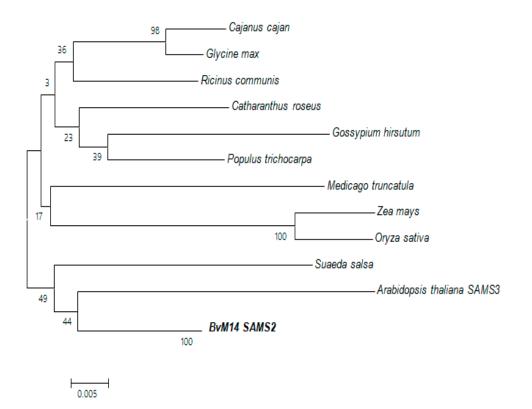


Figure S2. Phylogenetic tree of *BvM14-SAMS2* and *SAMS* gene members. The name of the gene (*BvM14-SAM2*) isolated in this study is in bold.

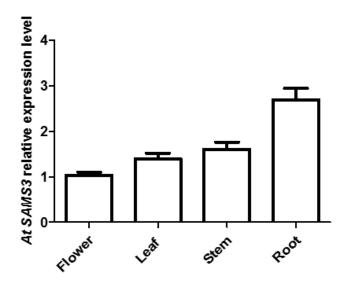


Figure S3. Tissue specific expression of AtSAMS3 gene in Arabidopsis plants

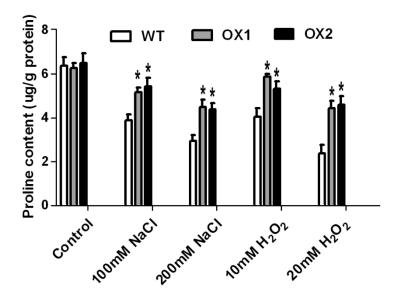


Figure S4. Effects of salt and H₂O₂ stress on proline content in wild type (WT) and *BvM14-SAMS2*overexpressed seedlings in wild type Arabidopsis (OX).

Primer pair 1								
	Sec	quence (5'->3')		Length	Tm	GC%	Self complementarity	Self 3' complementarit
Forward primer	GT	GTCTGATGATGTGGGTCTTGATGCT			62.88	48.00	3.00	0.00
Reverse primer	GAGTCTTACCATCAGGTCTCAGCCA			25	63.60	52.00	3.00	1.00
Products on target t	emplat	es						
>XM_010681173.2 PF	REDICT	ED: Beta vulgaris subsp. vulgaris S-ade	nosylmethic	onine synthase	2 (LOC1	04894834), mRNA	
product length	= 275							
Forward primer	1	GTCTGATGATGTGGGTCTTGATGCT	25					
Template	375	•••••	399					
Reverse primer	1	GAGTCTTACCATCAGGTCTCAGCCA	25					
Template	649		625					
	FDIOT	ED: Beta vulgaris subsp. vulgaris S-ade			4 // 0.04			
> <u>XM_010676210.2</u> PF	EDICI	ED. Beta vulgans subsp. vulgans S-ade	nosyimetric	onine synthase	T (LUCT	04890646), INRNA	
product length	= 275							
Forward primer	1	GTCTGATGATGTGGGTCTTGATGCT	25					
Template	394	AC.CC	418					
Reverse primer	1	GAGTCTTACCATCAGGTCTCAGCCA	25					
		.gg						

Figure S5. Primer-BLAST result for *BvM14- SAMS2* in sugar beet genome.