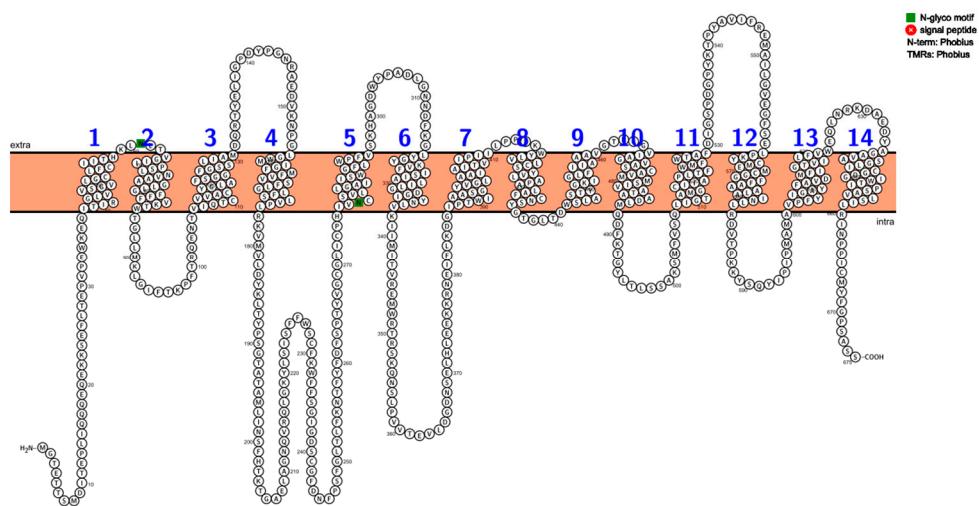
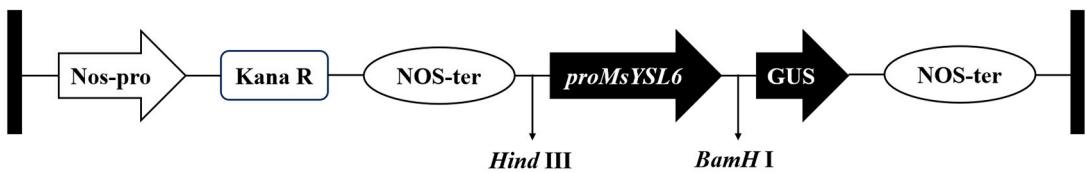


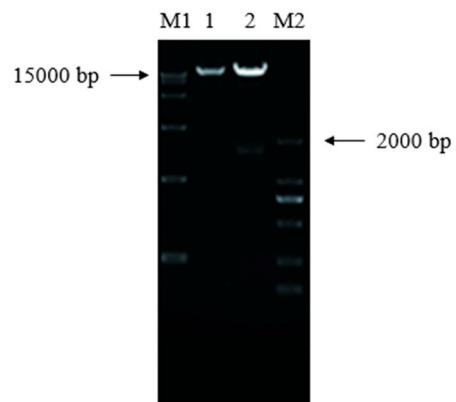
**Figure S1.** The cloning of *MsYSL6* from alfalfa and the transformation into tobacco. (A) The full-length of *MsYSL6* amplified by PCR from complementary DNA (cDNA) of alfalfa. M: DL 2000; 1: PCR product (*MsYSL6* CDS with 2, 028 bp). (B) Construction of pBI121-*MsYSL6* vector. LB: T-DNA left border; RB: T-DNA right border; p35S: cauliflower mosaic virus promoter; Npt II: Neomycin phosphotransferase II gene. (C) The transcript levels of *MsYSL6* in transgenic tobacco. Different letters represent significant differences ( $p < 0.05$ ). (D) The *MsYSL6* transgenic tobacco plant.



**Figure S2.** The predicted transmembrane domains of MsYSL6 protein.



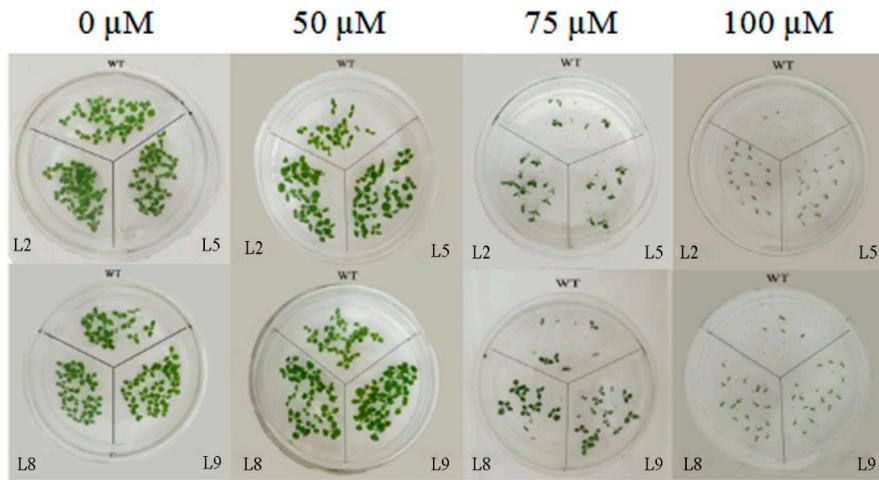
**Figure S3.** Construction of pBI121-*MsYSL6pro*::GUS.



**Figure S4.** Identification of pBI121-*MsYSL6pro*::GUS. M1, Marker DL15000; M2, Marker DL2000; 1, pBI121-*proMsYSL6*::GUS; 2, Recombinant plasmid digested by BamH I, Hind III.



**Figure S5.** The pBI121-*MsYSL6pro*::GUS induced alfalfa hairy roots.



**Figure S6.** The germination of *MsYSL6OE* tobacco under Cd stress. The seeds of four *MsYSL6OE* lines (L2, L5, L8, and L9) and WT tobacco were germinated on a half-strength MS solid medium containing 0, 50, 75, 100  $\mu\text{M}$   $\text{CdCl}_2$ .

**Table S1.** The germination rates of *MsYSL6Ex* lines and WT under Cd stress.

Line	Cd			
	Control	50 $\mu\text{M}$	75 $\mu\text{M}$	100 $\mu\text{M}$
WT	95.00±0.05	75.00±0.05	45.00±0.05	26.67±0.06
L2	96.67±0.03	83.33±0.03*	66.67±0.03*	53.33±0.03*
L5	95.00±0.05	83.33±0.03*	68.33±0.03*	55.00±0.00*
L8	96.67±0.03	83.33±0.03*	68.33±0.03*	55.00±0.05*
L9	96.67±0.03	83.33±0.03*	66.67±0.03*	50.00±0.05*

Note: \* $p < 0.05$ .

**Table S2.** The primer sequence.

Primer Comments	Primer Sequence
MsYSL6 cloning	F: ATGGGTACAGAAACA R: TCAGCTGCTTGCAGAA
qRT-PCR of <i>MsYSL6</i>	F: CTCAGTCACAACGGGAAGG R: TGGAACAGCCACAGCAAT
<i>MsYSL6pro</i> fragment cloning	F: CGGGATCCGCTATCTATTATGTGCTATCT R: CCAAGCTTGTGAGGTGAATCTGAG
<i>MsNAS</i> gene cloning	F: CGGGATCCTATAGAGATGATAACCCACCCG R: GAAGATCTTGGATTGAATTGAAGTAATTAA

qRT-PCR of <i>MsNAS</i>	F: AAGATTGTGGCTTCGGAT R: TTCACTCCTACGCATACCA
<i>MsNAS</i> pro fragment cloning	F: GGGATCC TATAGAGATGATAACCACCCG R: GAAGATCTTGGATTGAATTGAAGTAATTAA
<i>MsACTIN</i>	F: ACGAGCGTTTCAGATG R: ACCTCCGATCCAGACA
<i>NtGAPDH</i>	F: TAAGGGTGGTCCAAGAAGGT R: AGCAAGAGGAGCAAGGCACCT