

Table S1 Primer sequences used for qRT-PCR.

gene ID	Genen Name	Forward primer (5' -3' )	Reverse primer (5' -3' )
Gh_A13G2023	CNP 60A-1	CGAGGGTGCTAGTAACTGAT	TCCAGTAACATCCTCCGCAA
Gh_D13G2425	CNP 60A-2	CGAAAGTAGTCAATGATGGG	CCATCACCTGCAGAGTCATT
Gh_A05G1083	CNP 60B-1	TGGCTATAATGCTGCAACTGG	GATGTCGACAACCACGGCAT
Gh_D05G1261	CNP 60B-2	CTAGTAGATGTGGCTGCTGT	TTGTTTTCCGAAGTGTACCT
Gh_A01G0845	CAT1	TGCATTTTGCCCTGCCATTGTG	TGTGCCTCTGGGTATCAGCGTA
Gh_D06G1170	POD	TCAAGAGGCTGTCAATCTGGCG	TGGCCAGTGTTCGATTGCTGT
Gh_D13G1062	SODCC	ACTCAAGAGGGAGATGGCCAA	AATGGAGTGTGGCCAGAGAGA
Actin	—	ATCCTCCGTCTTGACCTTG	TGTCCGTCAGGCAACTCAT

Table S2 Effect of exogenous silicon on stomatal characteristics of Z9807 and Z0102 under saline and non-saline conditions

Genotype	Treatment	Length ( $\mu\text{m}$ )	Width ( $\mu\text{m}$ )	Surface ( $\mu\text{m}^2$ )	ESD (No. $\text{mm}^{-2}$ )
Z9807	CK	14.33 <sup>c</sup>	6.41 <sup>a</sup>	14.65 <sup>b</sup>	397.7 <sup>a</sup>
	CKSi	15.04 <sup>b</sup>	6.41 <sup>a</sup>	15.70 <sup>b</sup>	386.4 <sup>a</sup>
	NaCl	13.07 <sup>d</sup>	6.51 <sup>a</sup>	4.06 <sup>e</sup>	181.8 <sup>d</sup>
	NaClSi	14.94 <sup>b</sup>	6.35 <sup>a</sup>	12.14 <sup>c</sup>	340.9 <sup>b</sup>
Z0102	CK	14.25 <sup>c</sup>	6.31 <sup>a</sup>	15.02 <sup>b</sup>	431.8 <sup>a</sup>
	CKSi	15.99 <sup>a</sup>	6.38 <sup>a</sup>	17.18 <sup>a</sup>	375.0 <sup>a</sup>
	NaCl	13.11 <sup>d</sup>	6.51 <sup>a</sup>	2.99 <sup>e</sup>	159.1 <sup>d</sup>
	NaClSi	15.22 <sup>b</sup>	6.31 <sup>a</sup>	7.01 <sup>d</sup>	250.0 <sup>c</sup>

ESD, Effective stomata density (number of opening stomata  $\text{mm}^{-2}$ ). Data represents means of three replicates. Different letters within same testing trait indicates significant differences between treatments ( $p \leq 0.05$ ).