Table 1 Suppl. The primers used for the real-time quantitative PCR.

Gene name	Sequences (5′-3′) forward	reverse
Actin	GGAACTTGAGAAGGAGCCTAAG	CAACACCAACAGCAACAGTCT
SlNAC35	AGGAATTAATCATGTATTATCT	TCCAATAACCAGACACAGCT
SlCuZnSOD	TCTTCACCACAACCAGCACT	CAGTAAGGGGTTTAGGGGTAGT
SlCAT	TCCTTGTCGTCCTGCTGAG	TTGATGTATCTGTCTTGCCTGTC
SlCBF1	ATCATCTTCTTCTTCATCGTCATC	TCCTTGTCTTCTTATTTGGTTCTC
SlCBF2	ATGACGTGGCGGCTATAGCAT	AGTCACTACTCTCTTGCACAT
SlCOR518	ACTATTGTAGTCATGCGGTT	ACGCGATGGAGAAACGACAT
SlCOR413IM1	ATTGATGTAAGTGGAGAGAT	AACTGCAGTAGGAGCTTGT

Table 2 Suppl. Main putative *cis*-acting elements and their positions in the 2 000 bp upstream the *SlNAC35* transcription initiation site. The data have been obtained with *PlantCARE* (http://bioinformatics.psb.ugent.be/webtools/plantcare/html/). Indicated positions are relative to the transcription initiation site.

Cis-element	Consensus sequence	Responsiveness	Position
DRE	TACCGACAT	chilling and drought stress	-1972
CAT-box	GCCACT	meristem expression	-1168
MBS	TAACTG	drought	-1039
TC-rich repeats	ATTTTCTCCA	defense and stress	-981
GARE-motif	AAACAGA	gibberellin	-656
TCA-element	CCATCTTTTT	salicylic acid	-1613, -1640
CGTCA-motif	CGTCA	methyl jasmonate	-1141
HSE	AAAAAATTTC	heat stress	-1131, -1263, -1486
MRE	AACCTAA	irradiance	-304