Figure S1. Alignment of ZmDREB2.1/2A, ZmDREB2.9 and AtDREB2A isoforms (indicated according to the Table 1).
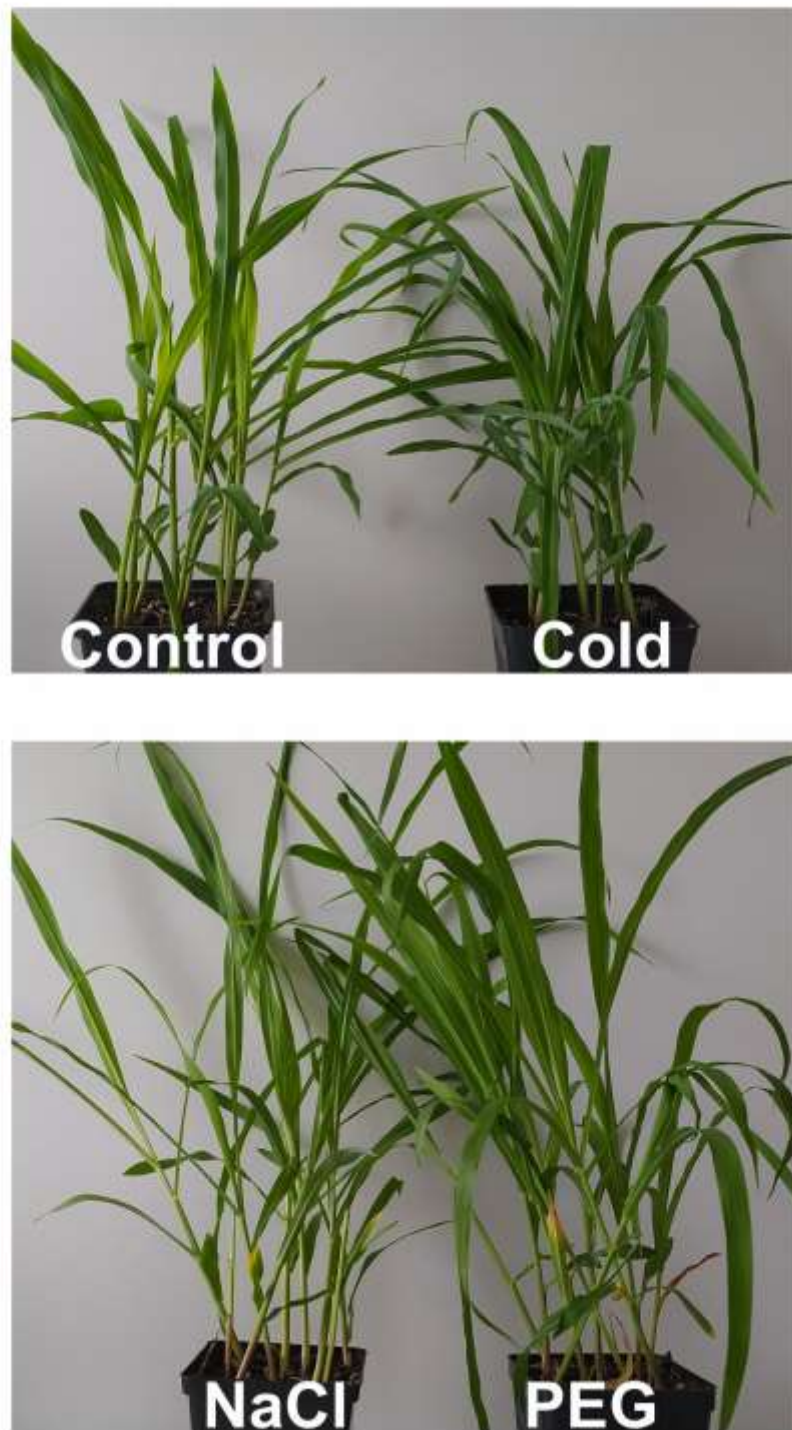
Figure S2. Photographs of maize seedlings 24 h after stresses (cold, salinity [NaCl], or drought [PEG]) compared to the untreated control. Seedlings after treatment with ABA are not shown, since outwardly, they did not differ from the control.
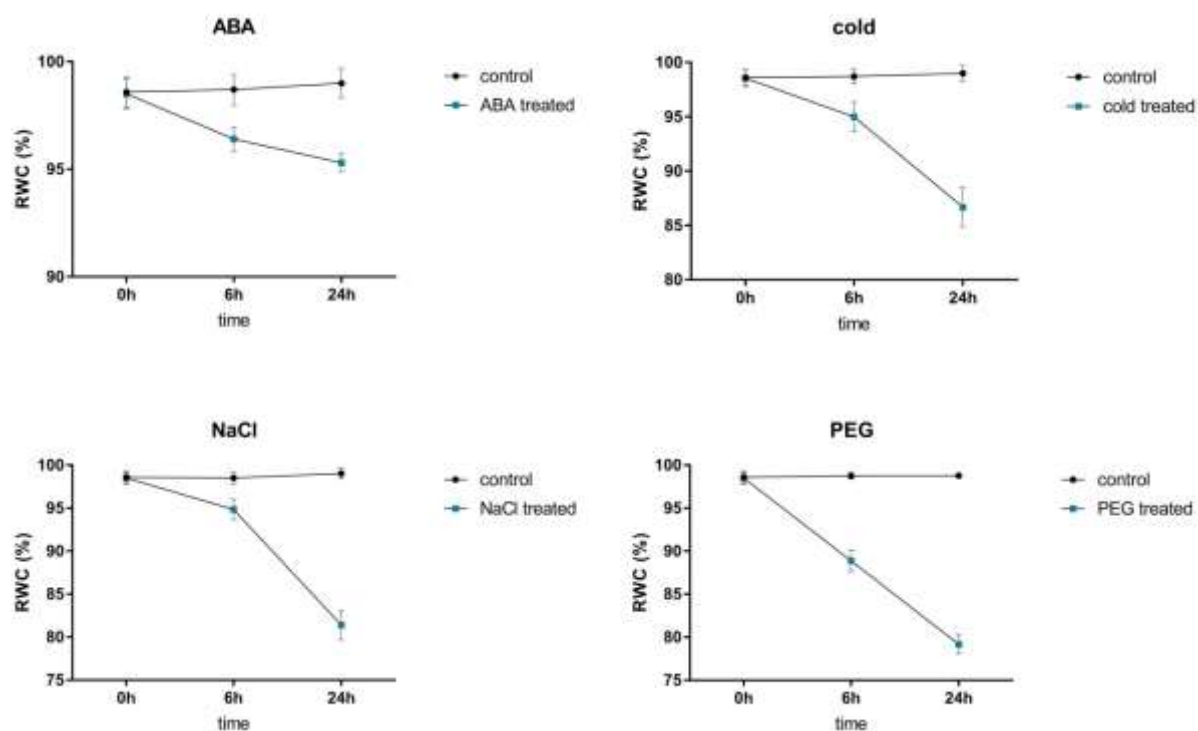
Figure S3. Relative water content in the leaf of corn seedlings 6 and 24 hours after stresses (ABA treatment, cold, salinity [NaCl], drought [PEG]) compared to the untreated control.

Table S1. List of primers for *ZmDREB2.1/2A–2.9* gene expression analysis.

Genes	Primer sequences (5′→3′)
ZmDREB2.1/2A (X2; X3)	GTTCCATCAGAACAGTAGCAGG CAGATTAGAGGCCAGCAGTCG
ZmDREB2.1/2A (X1; iso1)	GTTCCATCAGAACAGTAGGAG CAGATTAGAGGCCAGCAGTCG
ZmDREB2.2	GGACGACGACCCTCATGAC GCCATCGAACCCCTCATGC
ZmDREB2.3	AAGAAGGGCTCCGGCTCGT TACAGGAGCCGCAGGTAGG
ZmDREB2.4	CCTGGACCTGGACCTGGA GAACTCCCTGAGGTCGAGC
ZmDREB2.5	CCCCAGATCGACCTCAGC AGCTGGCTTTAGCAGTAGTCC
ZmDREB2.6	CAGGGAGTTCTTGCAACAGATC GCCGCTGACGAAGCAAGACA
ZmDREB2.7	GATGTTCCAGACACCTGCACT CCATCGCGGTGGTAGTGTC
ZmDREB2.8	GTGATGATGGCTGCCGTCC TCCTGATGCAGAGGAGCCG
ZmDREB2.9-S (X1)	GACGATGAATTGCCCTGTTGG CTCTTTCCACCACCTGATCG
ZmDREB2.9-L (iso1)	ATGGAGGAGCTGGGAGACG CTCTTTCCACCACCTGATCG