Table S1 Oligonucleotides used for experiments

Experiment	Name	Sequence[a]
cis-analysis	DREB1Cpro_mini_F_HindIII	5'-CCCAAGCTTATATCTTCCGGCCAATATAAAC-3'
	DREB1Cpro_mini_R_XhoI	5'-CCGCTCGAGTGATCAGAAGAGTACTCTGTTTC-3'
	DREB1Cpro_-394_F_HindIII	5'-CCCAAGCTTTTTGAAAATATCTCGTTTC-3'
	DREB1Cpro_-256_F_HindIII	5'-CCCAAGCTTTCATACAATAAGAAG-3'
	DREB1Cpro_-190_F_HindIII	5'-CCCAAGCTTCCAAGATGGGTCAAAGGAC-3'
	DREB1Cpro_-113_F_HindIII	5'-CCCAAGCTTAGCTGTTTCTTATCC-3'
	DREB1Cpro_-47_R_HindIII	5'-CCCAAGCTTGGATATTTGTGGGGTC-3'
	M1_F_HindIII	5'-CCCAAGCTTAGCTGTTTCTTATCCttGTGGC-3'
	M2_F_HindIII	5'-CCCAAGCTTAGCTGTTTCTTATCCACGTGGCATTtACA-3'
	M3_F_HindIII	5'-CCCAAGCTTAGCTGTTTCTTATCCACGTGGCATTCACAttGACAG-3'
	M4_R_HindIII	5'-CCCAAGCTTGGATATTTGTGGGGTCGAAaaCGGAG-3'
EMSA	1A_WT_F_HindIII	5'-CCCAAGCTTCGCTGTGTATAGTTTACG-3'
	1A_WT_R_HindIII	5'-CCCAAGCTTAGAGGCAAAAGAAAAGG-3'
	1B_WT_F_HindIII	5'-CCCAAGCTTCGCTATGTACTATACACG-3'
	1B_WT_R_HindIII	5'-CCCAAGCTTAACGGATATGTGGGGAG-3'
transgenic plant	PIF7pro_F_XbaI	5'-GCTCTAGAATATCTCTTTGAGAAATC-3'
	PIF7pro_F_HindIII	5'-CCCAAGCTTATATCTCTTTGAGAAATC-3'
	PIF7pro_R_SmaI	5'-TCCCCCGGGGTGTTACTTAGGCCGCACG-3'
	PIF7_F_XbaI	5'-GCTCTAGAATGTCGAATTATGGAGTTAAAG-3'
	PIF7_R_SmaI	5'-TCCCCCGGGCTAATCTCTTTTCTCATG-3'
	PIF7_R-3_SmaI	5'-TCCCCCGGGATCTCTTTTCTCATGATTCG-3'
	PIF7_fusion_F	5'-CGGCCTAAGTAACACATGTCGAATTATGGAG-3'
	PIF7_fusion_R	5'-CTCCATAATTCGACATGTGTTACTTAGGCCG-3'
transient expression assay	DBD-PIF7_F_SpeI	5'-GGGACTAGTGGACCGATGGTTCGCTGTTC-3'
	DBD-PIF7_R_SpeI	5'-GGGACTAGTCTAATCTCTTTTCTCATG-3'
	DBD-PIF7dN89_F_SpeI	5'-GGGACTAGTATGGACCGATGGTTCGCTG-3'
	DBD-PIF7dN328_F_SpeI	5'-GGGACTAGTAAGAGCCAACTTGCCACAAC-3'
	DBD-PIF7dC88_R_SpeI	5'-GGGACTAGTTTATTCTTGAGGATAACCGC-3'
	DBD-PIF7dC223_R_SpeI	5'-GGGACTAGTTTATCTTAGGCTCATGAAC-3'
	DBD-PIF7dC327_R_SpeI	5'-GGGACTAGTTTATGTCTGTGCGAAAAAGG-3'
	ICE1_F+1_SpeI	5'-GGGACTAGTGATGGGTCTTGACGGAAAC-3'
	ICE1_R-3_SpeI	5'-GGGACTAGTGATCATACCAGCATACCC-3'
	MYB15_F+1_SpeI	5'-GGGACTAGTGATGGGAAGAGCTCCATG-3'
	MYB15_R-3_SpeI	5'-GGGACTAGTGAGCCCGGCTAAGAGATC-3'
	CAMTA3_F+1_SpeI	5'-GGGACTAGTGATGGCGGAAGCAAGACG-3'
	CAMTA3_R-3_SpeI	5'-GGGACTAGTACTGGTCCACAAAG-3'
	TOC1_F+1_XbaI	5'-GCTCTAGAGATGGATTTGAACGGTGAG-3'
	TOC1_R-3_XbaI	5'-GCTCTAGAAGTTCCCAAAGCATCATCC-3'
	PRR3_F+1_XbaI	5'-GCTCTAGAGATGTGTTTTAATAACATTG-3'
	PRR3_R-3_XbaI	5'-GCTCTAGAATTGTCTTCACTTCCTG-3'
	PRR5_F+1_XbaI	5'-GCTCTAGAGATGACTAGTAGCGAGGAAG-3'
	PRR5_R-3_XbaI	5'-GCTCTAGATGGAGCTTGTGTGGATTGG-3'
	PRR7_F+1_XbaI	5'-GCTCTAGAGATGAATGCTAATGAGGAG-3'
	PRR7_R-3_XbaI	5'-GCTCTAGAGCTATCCTCAATGTTTTTTATG-3'
	PRR9_F+1_XbaI	5'-GCTCTAGAGATGGGGGAGATTGTGG-3'
	PRR9_R-3_XbaI	5'-GCTCTAGATGATTTTGTAGACGCGTC-3'
	VP16AD_F_BamHI	5'-CGGGATCCGGCACCCAAGAAGAAGCGGAAG-3'
	VP16AD_R_BamHI	5'-CGGGATCCGGGAATTCCATGGTGATCC-3'
	PIF7_mAPB_F	5'-GTTAAAgctCTCACATGGGAAAATgctCAAC-3'
	PIF7_mAPB_R	5'-GTTGagcATTTTCCCATGTGAGagcTTTAAC-3'
	PIF4_F_XbaI	5'-GCTCTAGAATGGAACACCAAGGTTGGAG-3'
	PIF4_R-3_SmaI	5'-TCCCCCGGGGTGGTCCAAACGAGAACCGTC-3'
	PhyB_F_XbaI	5'-GCTCTAGAATGGTTTCCGGAGTCGGGGGTAG-3'
	PhyB_R-3_SmaI	5'-TCCCCCGGGAATATGGCATCATCAGCATC-3'
	H2B_F_XbaI	5'-GCTCTAGAATGGCACCAAGAGCCGAGAAG-3'
	H2B_R-3_SmaI	5'-TCCCCCGGGAGAGCTAGTAAACTTGGTAAC-3'
RT-PCR	PIF7_634F	5'-TTGAAACAGCTACAAGCACAAGTACAG-3'
	PIF7_713R	5'-GGTAGTTGCGGAATCATCATTTG-3'
	PIF7_1084R	5'-GATTCGAAGAACTTGAAGGCATG-3'
	DREB1A_436F	5'-CGCTGACTCGGCTTGGA-3'
	DREB1A_543R	5'-GCATCACACATCTCATCCTGAAAC-3'
	DREB1B_504F	5'-AGTCAACATGCGCCAAGGAT-3'
	DREB1B_606R	5'-ATGTCCAGGCCATGATTCG-3'
	DREB1C_562F	5'-CGACGGATGCTCATGGTCTT-3'
	DREB1C_655R	5'-TCTTCATCCATATAAAACGCATCTTG-3'

[a] Bases with underline shows cutting sites of restriction enzymes. Bases with small characters indicate mutations.