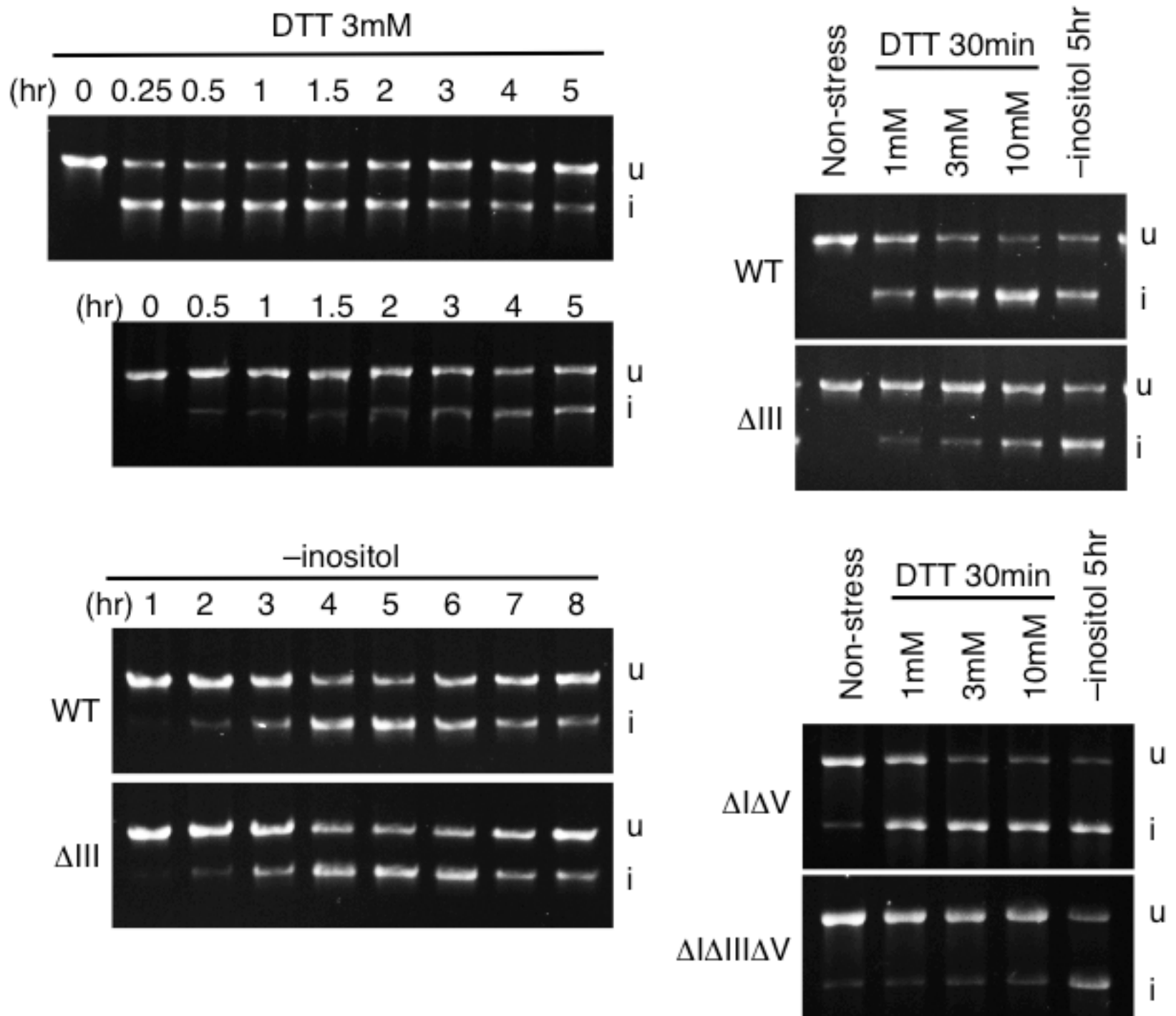


Figure S1

A Wild-type and ΔIII Ire1



B bZip-Ire1

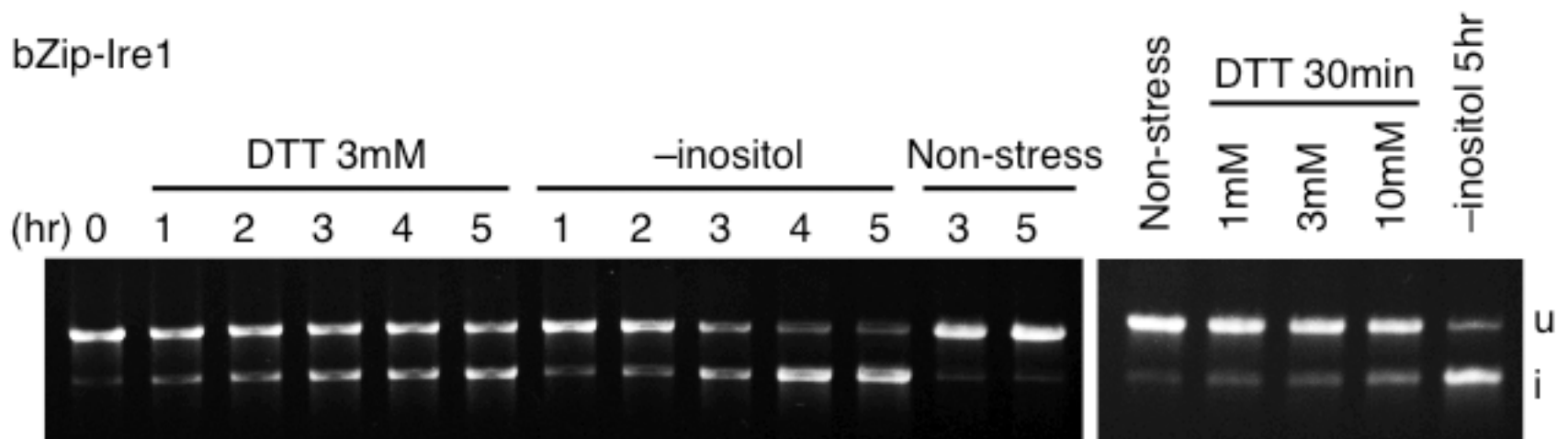


Figure S1 Raw-data examples for monitoring the *HAC1* mRNA splicing.

The *ire1Δ* cells KMY1516 carrying the *IRE1* gene (WT) or its mutants (on plasmid pRS313-IRE1 or its mutants) were stressed by the indicated stimuli, and total RNA samples were subjected to RT-PCR to amplify the *HAC1* products (*HAC1^u* (u) and *HAC1ⁱ* (i)), which were then fractionated by 2% agarose-gel electrophoresis and visualized by EtBr staining.

Figure S2

ire1 Δ cells

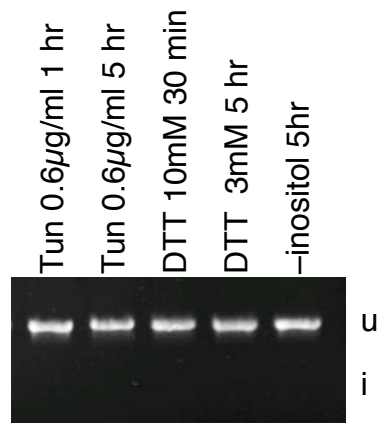
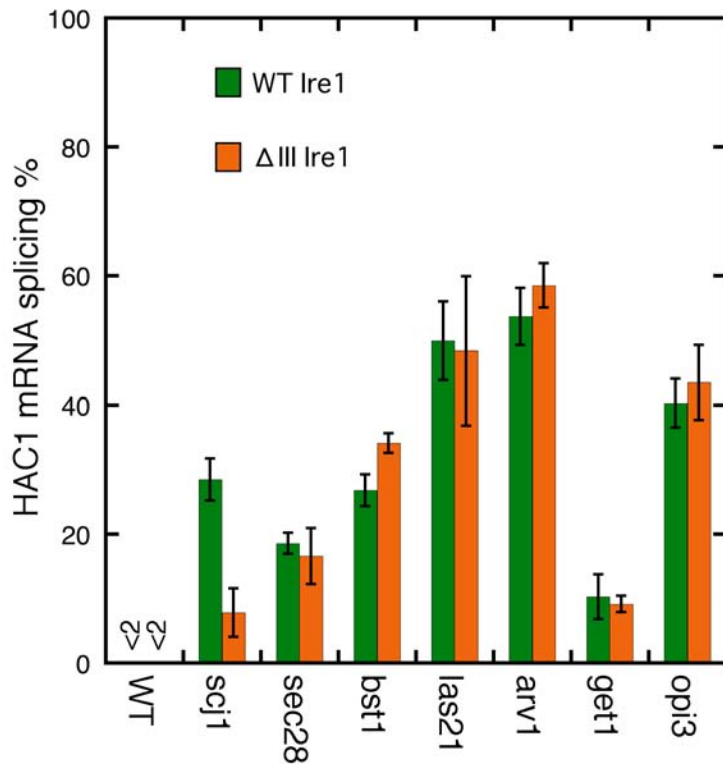


Figure S2 *ire1* Δ cells do not exhibit splicing of the *HAC1* mRNA.

The *ire1* Δ cells KMY1516 were stressed by the indicated stimuli, and total RNA samples were subjected to RT-PCR to amplify the *HAC1* products (*HAC1*^u (u) and *HAC1*ⁱ (i)), which were then fractionated by 2% agarose-gel electrophoresis and visualized by EtBr staining.

Figure S3

A



B

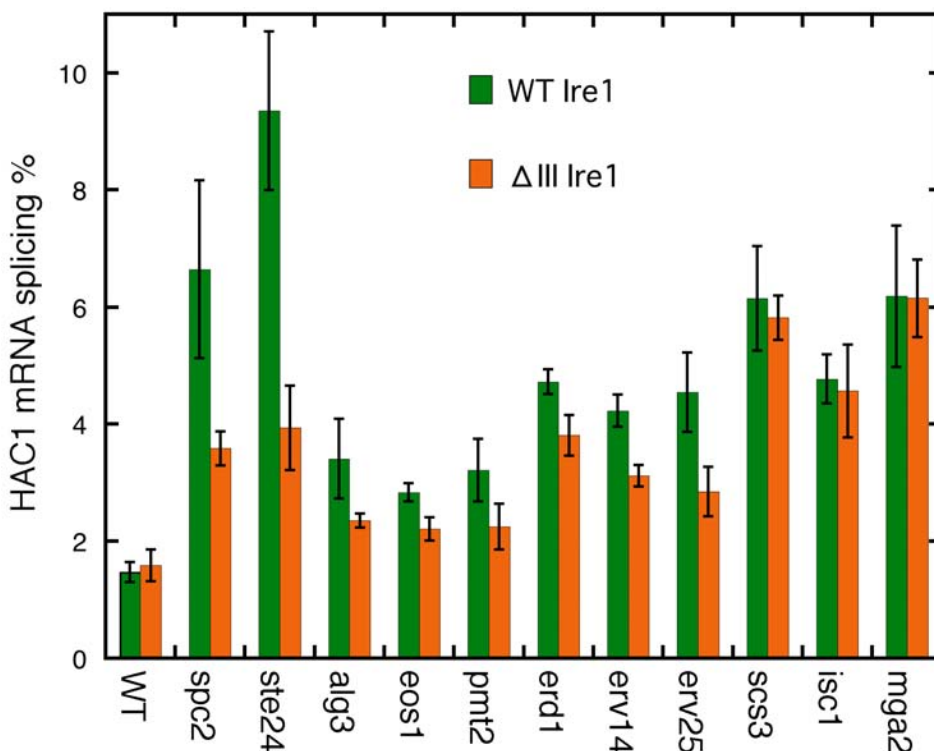


Figure S3 *HAC1* mRNA splicing induced by gene deletions.

Total RNA samples obtained from the strains employed in Figure 8 were analyzed by RT-PCR to evaluate splicing efficiency of the *HAC1* mRNA. In A, the RT-PCR products were conventionally fractionated by EtBr-containing gels, while in order to detect weak *HAC1*ⁱ signal, we employed a fluorescent dye-labeled PCR primer in B. Error bars represent the standard deviations from three independent transformants.

Figure S4

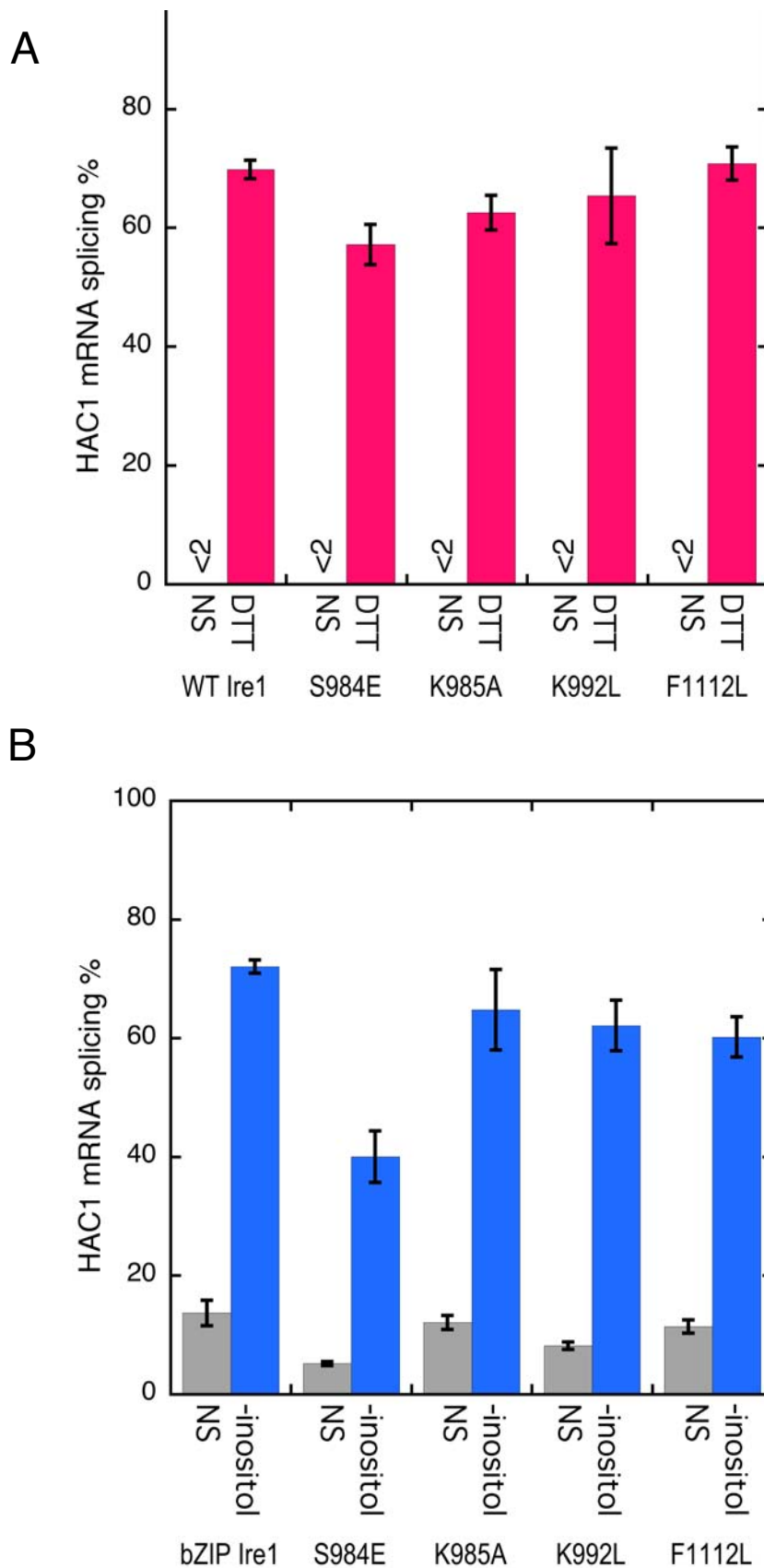


Figure S4 Activity of wild-type and bZIP-Ire1 carrying the quercetin-non-responding mutations.

The *ire1Δ* strain KMY1516 transformed with the wild-type IRE1 (A) or the bZIP-Ire1 (B) plasmid carrying the indicated mutations (pRS313-IRE1 or its derivatives) were cultured under non-stress conditions (NS) or stressed by DTT treatment (A; 3mM for 30 min) or inositol depletion (B; 5 hr) before RT-PCR amplification of the *HAC1* products from total RNA samples. Error bars represent the standard deviations from three independent transformants.

Table S1 Oligonucleotide primers to generate CPY and CPY* plasmids.

Sequence	Target	Direction	Attached restriction site
atatatata gcggccgc ATAGCTTCAAAAATGTTCTACTCC	TEF1 promoter	Forward	Not I
atatata actagt AAACTTAGATTAGATTGGCTATGCT	TEF1 promoter	Reverse	Spe I
atatat gcggccgc ACGGATTAGAAAGCCGCCGAGC	GAL1 promoter	Forward	Not I
atata actagt GGTTTCTCTCCCTTGACCGTTAAAGTATAGAGG	GAL1 promoter	Reverse	Spe I
gtt taactagt AAATGAAGCATTCACCAGTTTACTATGTGG	CPY or CPY* (PRC1 or prc1-1)	Forward	Spe I
tt taagaatt CTAAGGAAACCACCGTGGATCCAT	CPY or CPY* (PRC1 or prc1-1)	Reverse	Eco RI
ggttt agaatt cAAAGGTGAAGAATTATTCACTGGTGTG	GFP	Forward	Eco RI
tt aaactcga GTATTGTATCAATTCCATACCATG	GFP	Reverse	Xho I

The hybridizing sequences appear as capital letters.

The introduced restriction sites are highlighted by red letters.

Table S2 Antibodies used in this study.

Description	Working concentration	Purpose	Source
12CA5 mouse monoclonal IgG to the HA epitope	1 μ g/ml 2 μ g/ml	Western blotting	Roche
Rabbit anti-yeast BiP antiserum	4 μ g/ml	Immunoprecipitation	Higashio et al. (2000) (a)
	1: 1000 dilution	Western blotting	
	1: 500 dilution	Immunofluorescent staining	
Chicken anti-GFP IgY	1: 100 dilution	Immunofluorescent staining	Ave Labs
Rabbit anti-GFP IgG	1: 1000 dilution	Western blotting	MBL International Corporation
HRP-conjugated goat antibody to mouse Igs	1: 150 dilution	Immunoprecipitation	Dako
	1: 1000 dilution	Western blotting	
HRP-conjugated donkey antibody to rabbit IgG	1: 1000 dilution	Western blotting	GE Healthcare
FITC-conjugated goat antibody to mouse Igs	1: 100 dilution	Immunofluorescent staining	Cappel
Cy5-conjugated donkey antibody to mouse IgG	1: 100 dilution	Immunofluorescent staining	Jackson ImmunoResearch Laboratories
FITC-conjugated goat antibody to rabbit IgG	1: 100 dilution	Immunofluorescent staining	Cappel
FITC-conjugated donkey antibody to chicken IgY	1: 50 dilution	Immunofluorescent staining	Jackson ImmunoResearch Laboratories
Cy3-conjugated donkey antibody to rabbit IgG	1: 200 dilution	Immunofluorescent staining	Jackson ImmunoResearch Laboratories

(a) Higashio, H., Y. Kimata, T. Kiriyama, A. Hirata, and K. Kohno (2000) *J. Biol. Chem.* 275: 17900–17908.