The gene encoding *Arabidopsis* Acyl-CoA-Binding Protein 3 is pathogen-inducible and subject to circadian regulation

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Supplementary Figure legends

Figure S1 Nuecleotide sequence of the *ACBP3* 5'-flanking region. Nucleotide numbers begin at the 5'-end of the *ACBP3* flanking region. Upward and downward arrows indicate, respectively, the positions of the primers used corresponding to the boundaries of the truncated promoter fragments (5'-end deletions). The TATA-box is in bold and underlined, the transcriptional start site (nucleotide A) is in bold and marked as +1. The translation initiation codon is in bold italic letters. The other predicted *cis*-elements are named above nucleotide sequences and are shown in gray-filled boxes.

Table S1 Oligonucleotide primers used in this study for PCR walking, sequence analysis, EMSAs and DNase I footprinting.

Supplemental Figure S1

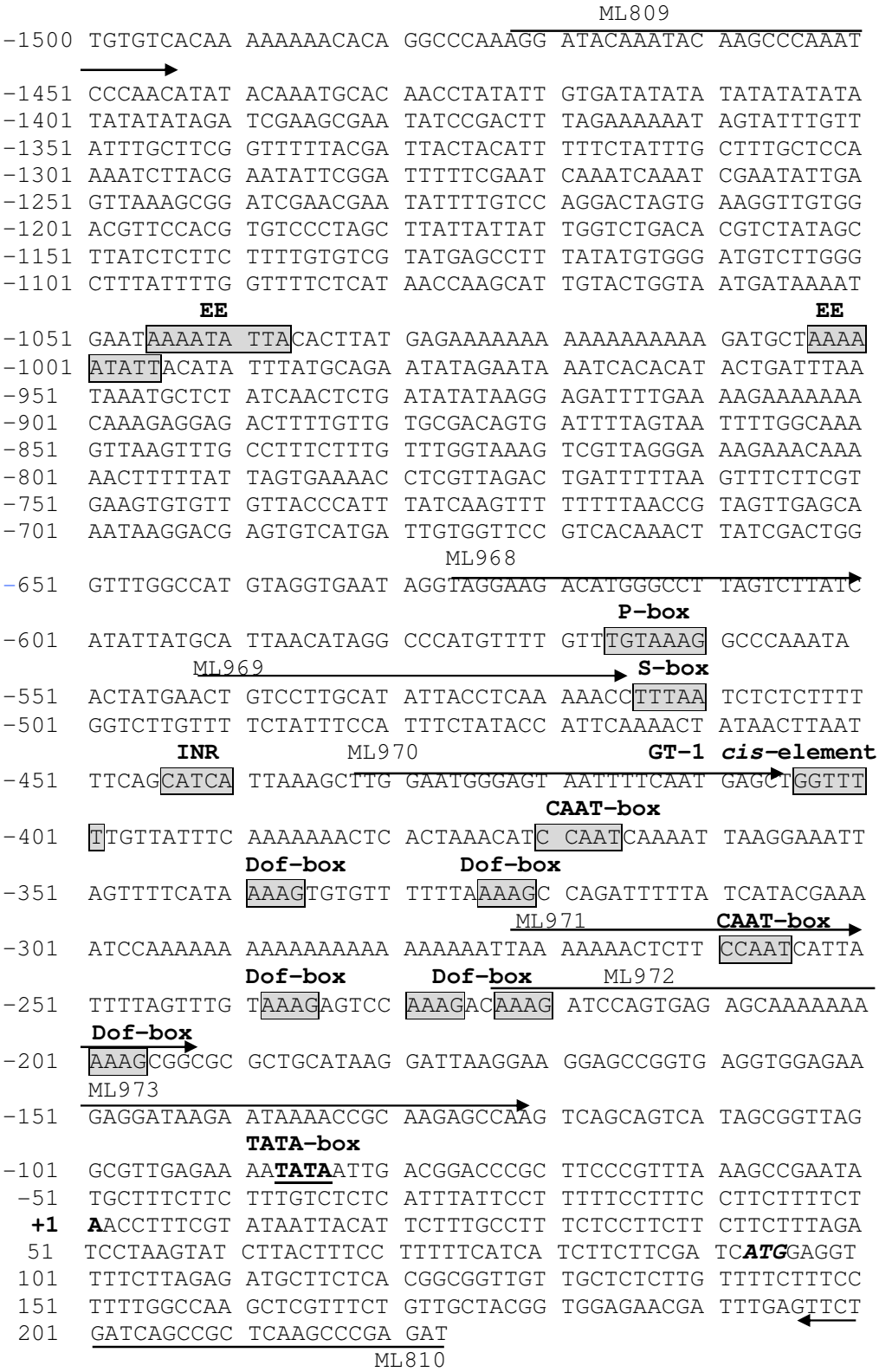

Table S1 Oligonucleotide primers used in this study for PCR walking, sequence analysis, EMSAs and DNase I footprinting

Primer	Sequence (from 5' to 3')	Nucleotide location	Orientation	Description
ML809	AAGGATCCAAATACAAGCCCAAATCCCAAC	-1475 to -1446 in *ACBP3* 5'-flanking region	Forward	For construction of pAT436, added *Bam*HI restriction site underlined
ML968	TAGGATCCCATGGGCCTTAGTCTTATC	-629 to -600 in *ACBP3* 5'-flanking region	Forward	For construction of pAT437, added *Bam*HI restriction site underlined
ML969	CTGGATCCGCATATTACCTCAAAAACC	-543 to -517 in *ACBP3* 5'-flanking region	Forward	For construction of pAT438, added *Bam*HI restriction site underlined
ML970	TTGGATCCGGAGTAATTTTCAATGAGC	-434 to -407 in *ACBP3* 5'-flanking region	Forward	For construction of pAT439, added *Bam*HI restriction site underlined
ML971	TAGGATCCCTCTTCCAATCATTA	-274 to -252 in *ACBP3* 5'-flanking region	Forward	For construction of pAT440, added *Bam*HI restriction site underlined
ML972	AAGGATCCAGTGAGAGCAAAAAAAAAAGCGGC	-226 to -194 in *ACBP3* 5'-flanking region	Forward	For construction of pAT441, added *Bam*HI restriction site underlined
ML973	GAGGATCCGAATAAAACCGCAAGAGCCAAG	-151 to -122 in *ACBP3* 5'-flanking region	Forward	For construction of pAT442, added *Bam*HI restriction site underlined
ML810	ATCCCGGGCTTGAGCGGCTGATCAGAAC	+196 to +223 in *ACBP3* 5'-flanking region	Reverse	For construction of *ACBP3pro::GUS* deletion series, added *Sma*I restriction site underlined
GUS-3'	TCCACAGTTTTCGCGATCCA	+64 to +84 in pBI101.3	Reverse	For identification of transgenic lines with *ACBP3* 5'-flanking sequence-specific forward primers by PCR
ML1112	CATAAAAGTGTGTTTTTTAAAAGCCAGATT	-345 to -316 in *ACBP3* 5'-flanking region	Forward	For preparation of Gp-I-Dof-wt probe used in EMSAs, Dof-boxes underlined
ML1113	AATCTGGCTTTTAAAAAACACACTTTTATG	-345 to -316 in *ACBP3* 5'-flanking region	Reverse	For preparation of Gp-I-Dof-wt probe used in EMSAs, Dof-boxes underlined
ML1114	GTAAAGAGTCCAAAGACAAAGATCC	-242 to -218 in *ACBP3* 5'-flanking region	Forward	For preparation of Gp-II-Dof-wt probe used in EMSAs, Dof-boxes underlined
ML1115	CGGATCTTTGTCTTTGGACTCTTTA	-242 to -218 in *ACBP3* 5'-flanking region	Reverse	For preparation of Gp-II-Dof-wt probe used in EMSAs, Dof-boxes underlined
ML1116	AAAAAAAAAGCGGCGCGC	-206 to -190 in *ACBP3* 5'-flanking region	Forward	For preparation of Gp-III-Dof-wt probe used in EMSAs, Dof-box underlined
ML1117	GCGCGCCGCTTTTTTTT	-206 to -190 in *ACBP3* 5'-flanking region	Reverse	For preparation of Gp-III-Dof-wt probe used in EMSAs, Dof-box underlined
ML1160	CATACACGTGTGTTTTTTAAAAGCCAGATT	-345 to -316 in *ACBP3* 5'-flanking region	Forward	For preparation of Dof-(-341/-338)-mut probe used in EMSAs, mutated Dof-box (-341/-338) underlined
ML1161	AATCTGGCTTTTAAAAAACACACGTGTATG	-345 to -316 in *ACBP3* 5'-flanking region	Reverse	For preparation of Dof-(-341/-338)-mut probe used in EMSAs, mutated Dof-box (-341/-338) underlined
ML1162	CATAAAAGTGTGTTTTTTACACGCCAGATT	-345 to -316 in *ACBP3* 5'-flanking region	Forward	For preparation of Dof-(-326/-323)-mut probe used in EMSAs, mutated Dof-box (-326/-323) underlined

ML1163	AATCTGGCGTGTAAAAAACACACTTTTATG	-345 to -316 in *ACBP3* 5'-flanking region	Reverse	For preparation of Dof-(-326/-323)-mut probe used in EMSAs, mutated Dof-box (-326/-323) underlined
ML1198	CAATGAGCTGGTTTTTGTTATTTC	-415 to -392 in *ACBP3* 5'-flanking region	Forward	For preparation of GT-1-(-406/-401)-wt probe used in EMSAs, GT-1 *cis*-element (-406/-401) underlined
ML1199	GAAATAACAAAAACCAGCTCATTG	-415 to -392 in *ACBP3* 5'-flanking region	Reverse	For preparation of GT-1-(-406/-401)-wt probe used in EMSAs, GT-1 *cis*-element (-406/-401) underlined
ML1200	CAATGAGCTCCTTTTTGTTATTTC	-415 to -392 in *ACBP3* 5'-flanking region	Forward	For preparation of GT-1-(-406/-401)-mut probe used in EMSAs, mutated GT-1 *cis*-element (-406/-401) underlined
ML1201	GAAATAACAAAAAGGAGCTCATTG	-415 to -392 in *ACBP3* 5'-flanking region	Reverse	For preparation of GT-1-(-406/-401)-mut probe used in EMSAs, mutated GT-1 *cis*-element (-406/-401) underlined
ML1451	CCTCAAAAACCTTTAATCTCTC	-527 to -506 in *ACBP3* 5'-flanking region	Forward	For preparation of S-box-(-516/-512)-wt probe used in EMSAs, S-box (-516/-512) underlined
ML1452	GAGAGATTAAAGGTTTTTGAGG	-527 to -506 in *ACBP3* 5'-flanking region	Reverse	For preparation of S-box-(-516/-512)-wt probe used in EMSAs, S-box (-516/-512) underlined
ML1453	CCTCAAAAACCCAGTCTCTCTC	-527 to -506 in *ACBP3* 5'-flanking region	Forward	For preparation of S-box-(-516/-512)-mut probe used in EMSAs, mutated S-box (-516/-512) underlined
ML1454	GAGAGAGACTGGGTTTTTGAGG	-527 to -506 in *ACBP3* 5'-flanking region	Reverse	For preparation of S-box-(-516/-512)-mut probe used in EMSAs, mutated S-box (-516/-512) underlined
ML1171	TCAGCATCATTAAAGCTTGG	-450 to -431 in *ACBP3* 5'-flanking region	Forward	5'-end labeled with 6-FAM for preparation of DNase I footprinting probe
ML1172	GAATAGGAGAAGAGGTGGAG	-162 to -143 in *ACBP3* 5'-flanking region	Reverse	5'-end labeled with NED for preparation of DNase I footprinting probe