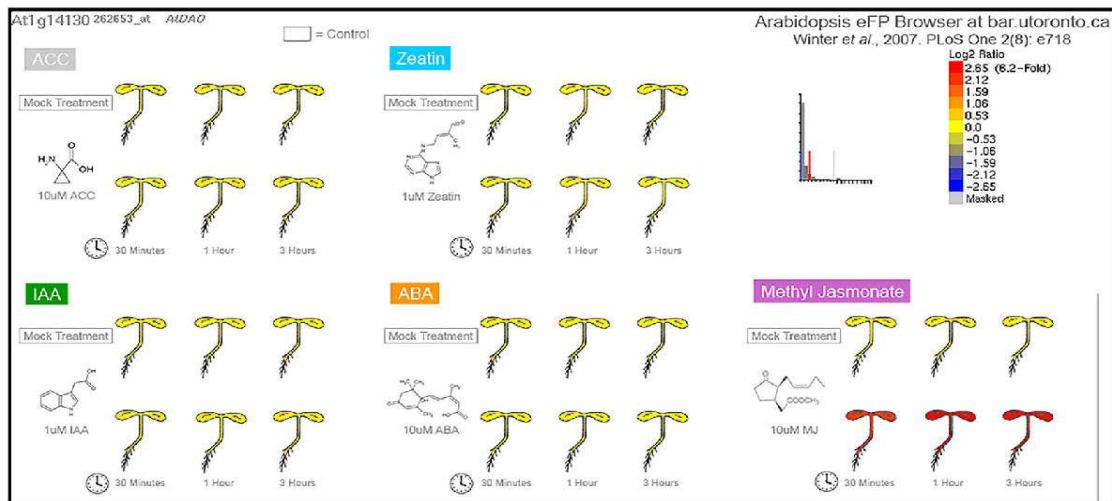


Supplementary Information

A



B

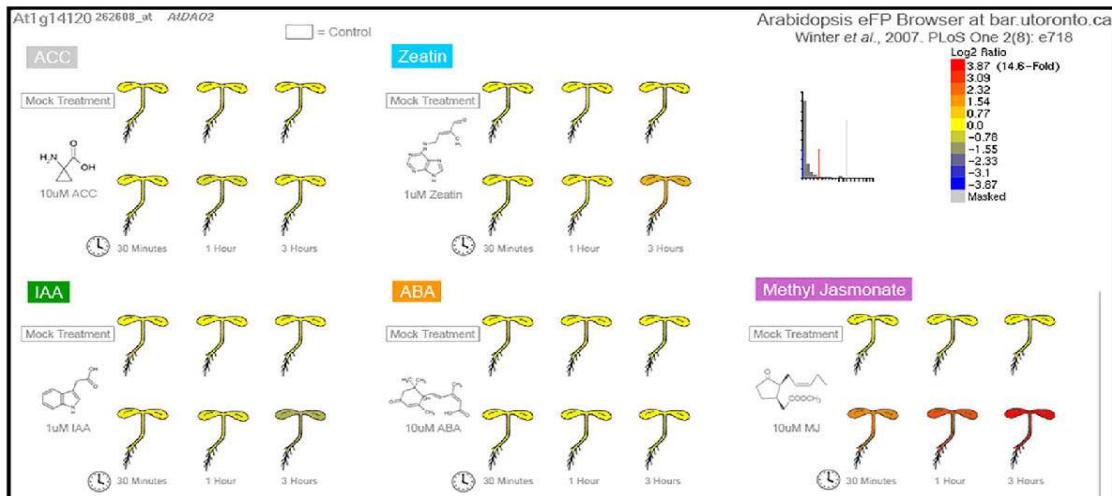
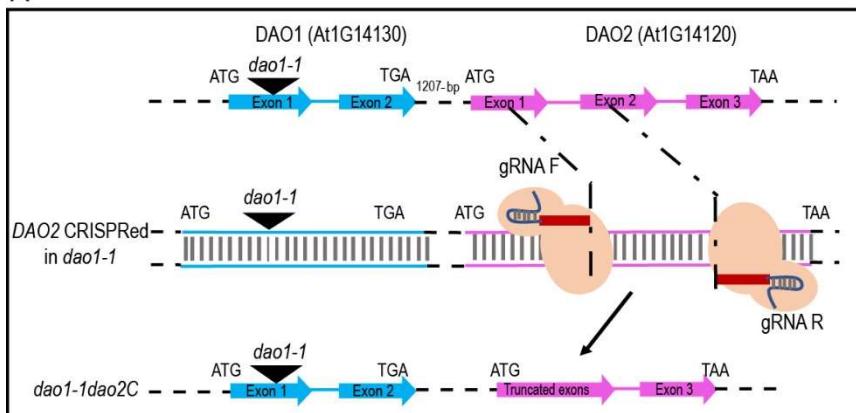


Figure 1. Relative expression data showing that MeJA specifically induces the expression of (A) DAO1 and (B) DAO2. Data retrieved from the publicly-available dataset at Arabidopsis eFP browser (<http://bar.utoronto.ca>).

A



B

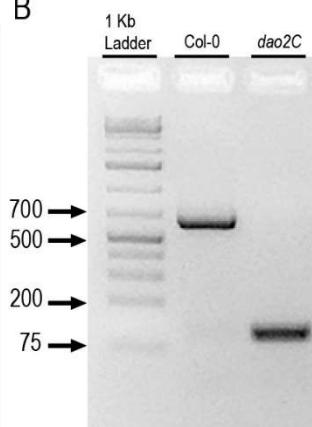


Figure 2. Illustration showing the CRISPR-Cas9 strategy adopted. (A) Two guide RNAs were designed to target a relatively large DNA fragment from the *DAO2* gene in a *dao1-1* loss of function mutant background. (B) Agarose gel showing 647 bp fragment in the *DAO2* wild type and homozygote deletion of ~500 bp resulting in ~100 bp fragment in the *dao2C* allele.

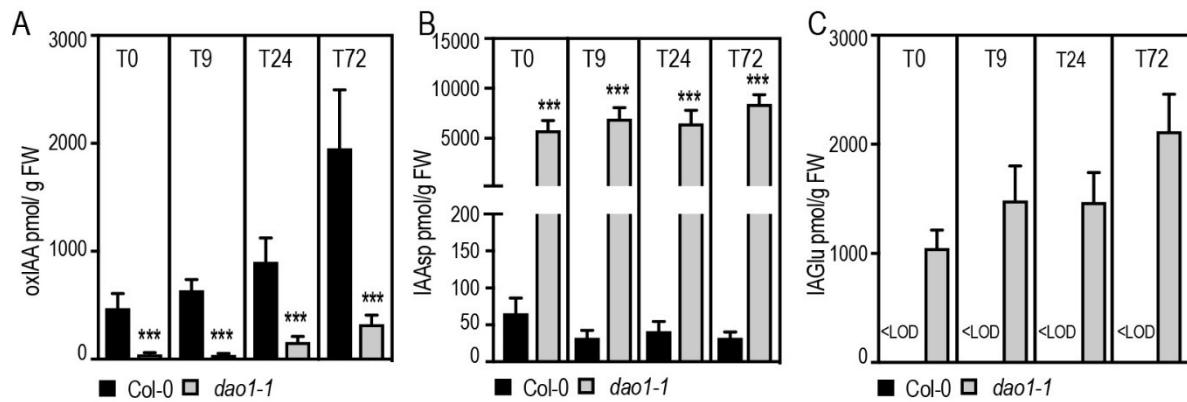


Figure 3. The *dao1-1* mutant accumulates significantly less OxIAA and more amino acid conjugates (IAAsp and IAGlu). (A-C) Endogenous hormone contents. (A) OxIAA, (B) indole-3-acetyl-L-aspartic acid (IAAsp) and (C) indole-3-acetyl glutamic acid (IAGlu) were quantified in the hypocotyls of wild-type and *dao1-1* mutant seedlings grown in the dark until the hypocotyl reached ~ 6 mm long (T0) and after their transfer to the light for 9 h (T9), 24 h (T24) or 72 h (T72). Error bars indicate \pm SD of six biological replicates. Asterisks indicate statistically significant difference in the mutant lines versus the wild type (Col-0) in an ANOVA analysis (*, **, and *** correspond to P-values of 0.05 > p > 0.01, 0.01 > p > 0.001, and p < 0.001, respectively). <LOD means under the limit of detection.

Table 1: list of primers used in qRT-PCR and genotyping

Gene name	Gene number	Forward primer	Reverse primer
<i>GH3.3</i>	At1g77850	ACAATTCCGCTCCACAGTTTC	ACGAGTTCTTGCTCTCCAA
<i>GH3.5</i>	At4g27260	GTCCTCGAGGACTGCTGCTT	ATGTCCCTGGCTAACAAATC
<i>GH3.6</i>	At5g54510	CCTTGTTCGGTTTGATGCTT	CGTGTACCAGTTCAAGCAGA
<i>OPR3</i>	At2G06050	TGGTTGGCATGCTCAATAAG	GCCTTCCAGACTCTGTTGC
<i>AOC2</i>	At3G25770	GGTGCCTACGGACAGGTCAAGC	GCGGTACCGGTGTTCCGGTG
<i>TIP41</i>	At4g34270	GCTCATCGGTACGCTCTTT	TCCATCAGTCAGAGGCTTCC
<i>dao2C</i>	At1g14120	TTGCTTGACTAGAGAAAAGCCTT	TTGGCTTGGCCATCCTCTCA
U626-ID		TGTCCCAGGATTAGAATGATTAGGC	AGCCCTTCTCTTCGATCCATCAAC