

**Supplementary Table S1** List of anthocyanins identified in the methanol extracts of *A. thaliana* rosettes.

Peak	Rt (min) <sup>a</sup>	UV, λmax (nm)	MS (m/z)	MS <sup>2</sup> fragments (m/z)	Assignment
1	19,7	527	1137	535,889	A8
2	20,8	520	1258	1095,449	A10
3	21,8	540	1344	1095,535	A11
4	22,8	535	1344*	1095,535	A11*
5	24,1	530	1095	933,449	A7
6	25	525	975	727,535,287	A5
7	26	530	1182	1137,933,535,491	A9
8	26,8	535	1182*	1137,933,535,491	A9*

\*Asterisk indicates a tautomer.

A8 – Cyanidin 3-O-[2"-O-(xylosyl) 6"-O-(p-O-(glucosyl) p-coumaroyl) glucoside] 5-O-[6'''-O-(malonyl) glucoside].

A10 – Cyanidin 3-O-[2"-O-(2'''-O-(sinapoyl) xylosyl) 6"-O-(p-O-(glucosyl) p-coumaroyl) glucoside] 5-O-glucoside.

A11. Cyanidin 3-O-[2"-O-(6'''-O-(sinapoyl) xylosyl) 6"-O-(p-O-(glucosyl)-p-coumaroyl) glucoside] 5-O-(6'''-O-malonyl) glucoside.

A7 – Cyanidin 3-O-[2"-O-(2'''-O-(sinapoyl) xylosyl) 6"-O-(p-coumaroyl) glucoside] 5-O-glucoside.

A5 – Cyanidin 3-O-[2"-O-(xylosyl)-6"-O-(p-coumaroyl) glucoside] 5-O-malonylglucoside.

A9 – Cyanidin 3-O-[2"-O-(2'''-O-(sinapoyl) xylosyl) 6"-O-(p-O-coumaroyl) glucoside] 5-O-[6'''-O-(malonyl) glucoside].

**Supplementary Table S2** *In vitro*-synthesized single-stranded RNA oligonucleotides.

	RNA oligonucleotide name	5' - 3' RNA oligonucleotides	Modification
1	ssCHS-s	5'-GUGACUGGAACUCCUCUUCU	5' phosphate
2	ssCHS-a	5'-AAGAGGGAGUUCCAGUCACUU	2'-O-methyl at 3' end
3	NPTII (R3-s-Me)	5'-AAUGGCCGUUUUCUGGAUUC	5' phosphate
4	NPTII (R3-a-Me)	5'-AUCCAGAAAAGCGGCCAUUU	2'-O-methyl at 3' end

**Supplementary Table S3** Primers used in RT-PCR and qRT-PCRs.

Gene name (ID number)	Primer name	Primers, 5'-3'
Primers for cloning full-length cDNA coding sequences, 5'-3'		
<i>AtCHS</i> (AT5G13930.1)	AtCHS-Nach-S AtCHS-konA	5'ATGGTGTGGCTGGCTTCTT 5'TTAGAGAGGAACGCTGTGCAAG
<i>AtMybL2</i> (AT1G71030.1)	AtMybL2-s1 AtMybL2-kon	5'ATGAACAAAACCCGCCTTC 5'TCATCGGAATAGAAGAAGC
<i>AtANAC032</i> (AT1G77450.1)	ANAC032-s1 ANAC032-kon	5'ATGATGAAATCTGGGCTG 5'TCAGAAAGTTCCCTGCCTA
Specific primers for dsRNA design, 5'-3'		
<i>AtCHS</i> (AT5G13930.1)	AtCHS-RNAs2 AtCHS- RNA-a2	5'TAATACGACTCACTATAAGGGAGAGCTTCTGGTCTCC GTCCTTCC 5'TAATACGACTCACTATAAGGGAGATTAGAGAGGAACG CTGTGCAAG
<i>AtMybL2</i> (AT1G71030.1)	MybL2-dsRNA-s1 MybL2-dsRNA-a1	5'TAATACGACTCACTATAAGGGAGAACAAAAACCC GCCTTC 5'TAATACGACTCACTATAAGGGAGATCATCGGAATAGA AGAAGCGTT

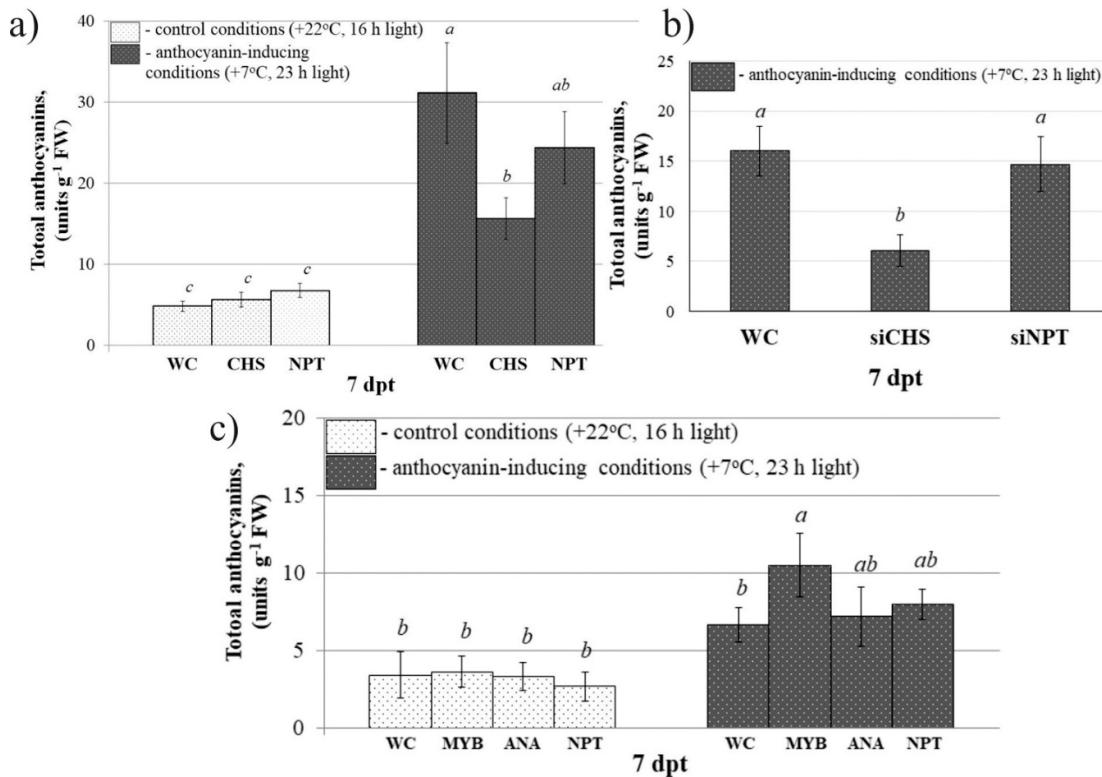
<i>AtANAC032</i> (AT1G77450.1)	ANA-dsRNA-s1  ANA-dsRNA-a1	5'TAATACGACTCACTATAAGGGAGAATGATGAAATCTG GGGCTGATTG  5'TAATACGACTCACTATAAGGGAGATCAGAAAGTTCCC TGCCTAACCC
<i>NPTII</i> (AY818371)	npt-T71-s  npt-T72-a	5'TAATACGACTCACTATAAGGGAGAATGTGGATTGAAC AAGATGGATTG  5'TAATACGACTCACTATAAGGGAGATCCACCATGATATT CGGCAAGCAG
Primers for cDNA check-up on DNA contamination, 5'-3'		
<i>AtGAPDH</i> (GenBank NM_111283)	AtGapdh-s  AtGapdh-a	5'CTG GAA TGT CTT TCC GTG TC 5'ATT CGT TGT CGT ACC ATG AC
Primers for PCR and real-time PCR, 5'-3'		
<i>AtCHS</i> (AT5G13930.1)	AtCHS-Nach-S  AtCHS-realA	5'ATGGTGATGGCTGGTGTGCTTCTT 5'-CACATGGTTCTCAGGGTTAGC
<i>AtMybL2</i> (AT1G71030.1)	AtMybL2-reals1  MybL2-3UTR-realA1	5'TTGCCTGACCTAACATTG 5'GCCGGTCCAATTCAAGGATTAAC
<i>AtANAC032</i> (AT1G77450.1)	ANAC032-reals1  AtANA-3UTR-real-a1	5'CGTTTAATTACGTAGATGC 5'CACTTCCACTAACTCTAATCGC
<i>NPTII</i> (GenBank AJ414108)	nptII-realS  nptII-realA	5'TTGCTGAAGAGCTGGCGGCGAAT 5'TCAGAAGAACTCGTCAAGAAGG
<i>AtGAPDH</i> (GenBank NM_111283)	AtGapdh-real-s  AtGapdh-real-a	5'TTG GTG ACA ACA GGT CAA GCA 5'AAA CTT GTC GCT CAA TGC AAT
<i>AtUBQ</i> (GenBank NM_001084884)	AtUBQ-realS  AtUBQ-realA	5'GGCCTTGATAATCCCTGATGAATAAG 5'AAAGAGATAACAGGAACGGAAACATACTAGT

**Supplementary Fig. S1** Spectrophotometric results of total anthocyanins in the rosettes of *A. thaliana* grown under the control (+22°C, 16 h light) and anthocyanin-inducing (+7°C, 23 h light) conditions for seven days. (a) total anthocyanins in *A. thaliana* treated with sterile water or synthetic *AtCHS*- or *NPTII*-dsRNAs; (b) total anthocyanins in *A. thaliana* treated with sterile water or synthetic *siCHS* or *siNPTII*; (c) total anthocyanins in *A. thaliana* treated with sterile water or synthetic *MYBL2*- or *ANAC032*-dsRNAs.

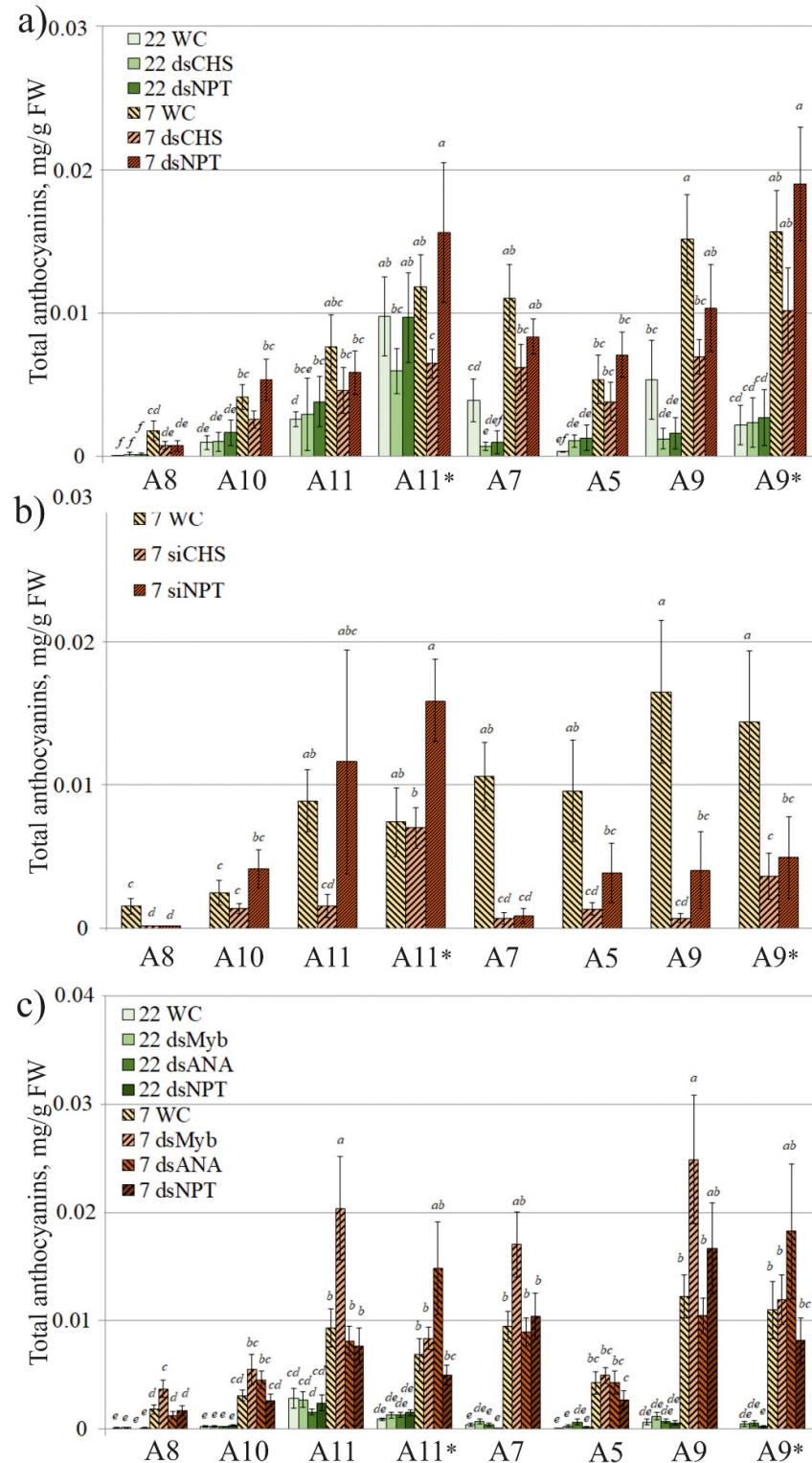
WC – *A. thaliana* treated with sterile water, CHS – *A. thaliana* treated with *AtCHS*-dsRNAs, NPT – *A. thaliana* treated with *NPTII*-dsRNAs; siCHS - *A. thaliana* treated with *AtCHS*-siRNAs; siNPTII - *A. thaliana* treated with *NPTII*-siRNAs; MYB – *A. thaliana* treated with *AtMYB*-dsRNAs; ANA – *A. thaliana* treated with *AtANAC032*-dsRNAs; dpt – days post treatment.

One anthocyanin unit equals one absorbance unit [A530 - (1/4A657)] in 1 mL of extraction solution.

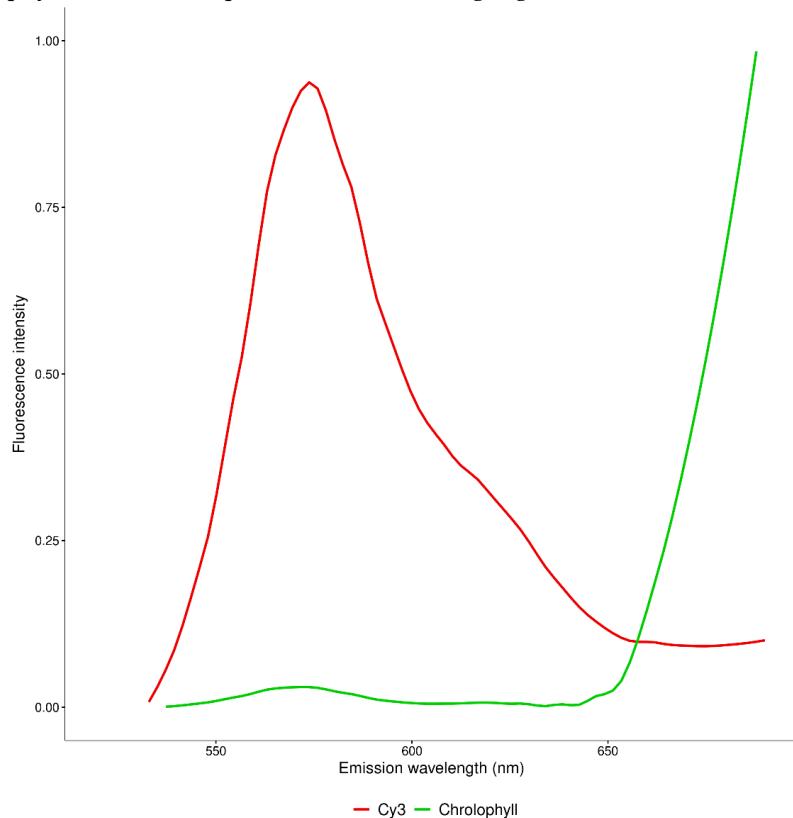
The data are presented as the mean ± SE (four independent experiments). Means followed by the same letter were not different using Student's *t* test. *P* ≤ 0.05 was considered statistically significant.



**Supplementary Fig. S2** HPLC profile of anthocyanins in the leaves of *Arabidopsis thaliana* grown under the control (+22°C, 16 h light) and anthocyanin-inducing (+7°C, 23 h light) conditions. WC – *A. thaliana* treated with sterile water, dsCHS – *A. thaliana* treated with *AtCHS*-dsRNAs, dsNPT – *A. thaliana* treated with *NPTII*-dsRNA. The data are presented as the mean ± SE (three independent experiments). Means followed by the same letter were not different using Student's t test.  $p < 0.05$  was considered statistically significant.  
\* – Asterisk indicates a tautomer.



**Supplementary Fig. S3** Emission spectrum of Cy3 and chlorophyll obtained in the  $\lambda$ -scanning mode of Zeiss LSM 780 laser scanning microscope. Cy3 fluorescence (peak at  $\approx$  570 nm–580 nm) is highlighted in green and chlorophyll fluorescence (peak at  $\approx$  690 nm) is highlighted in red.



### Supplementary Video S1

The dsRNAs were applied on the foliar surface of four-week-old *Arabidopsis thaliana* by spreading with individual soft brushes (natural pony hair) sterilized by autoclaving. All leaves of one rosette for each type of condition were treated on both the adaxial (upper) and abaxial (lower) sides. The leaf surface of each individual plant was treated with 0.35  $\mu\text{g}/\mu\text{L}$  of dsRNA or water (100  $\mu\text{L}$  per individual plant). For each dsRNA treatment, 35  $\mu\text{g}$  of the dsRNA were diluted in 100  $\mu\text{L}$  of nuclease-free water.