

Table S1. Primers for genes

№	Gene Bank ID	Gene Description	Gene	Primer 5'-3'	
				forward	reverse
1	ALN42232.1	Histidine-containing phosphotransfer 1	<i>HPT1</i>	GCTCAAGTATAGGAGCGCGG	CCAGCTTGTTTTTACGAGGT
2	EF083399.1	Jasmonate-Zim domain 1	<i>JAZa</i>	GGTGAACGTGTATGATGATATTCC	CGTTGCAGAGAATGCTTCCTC
3	AY289600.1	Auxin-induced protein 1 (IAA1)	<i>AUX/IAA</i>	GCCACCTGTCAAAGATTTCAG	TGAGGTCCACCTTTCTGAGA
4	CBB44933.1	Actin 1	<i>ACT1</i>	TTAGCAACTGGGATGACATGGA	CCTGAATGGCAACATACATAGCA
5	KY914544.1	auxin response factor 16	<i>ARF16</i>	TATATACCGTGGCACACCGC	ACACACAGCTCCCCATTGAG
6	A9NWW4	Steroid 5- $\alpha$ -reductase DET2	<i>BR-<math>\alpha</math>-RED</i>	CAACAGAGCTCTCAAGGCGA	GAAGAGTAGGGCGCTCATCC
7	MH017214.1	Brassinosteroid mediated signaling pathway gene	<i>BRZ2</i>	GCGTCTATGAGCCCATCCTC	CGGCGAGATTTTTCAGCCAG
8	JQ240296.1	Monoterpene synthase like TPS-mono1	<i>mono-TERP</i>	GGAGTATCCACCAGTCCCCT	TCCAATGCCTTTCTCGGTCC
9	JQ240308.1	$\alpha$ -terpineol synthase TPS- $\alpha$ -terp	<i><math>\alpha</math>-TERP</i>	AGCATCGAACGTTTGGGAGT	AAGAGTTCGAAGGCCCAGTG
10	MA_10426264g0020	Chalcone synthase	<i>CHS</i>	GGCATTGAGGAAGGCTCAGAGA	GGCACCTCCACCACAACCAT
11	FN433184.1	Auxin response factor 3/4	<i>ARF3/4</i>	CGTGGCTCCTTCCTCAACAT	TGGTCGGCGTGTAGATTGAC
12	L26923.1	Glyceraldehyde-phosphate dehydrogenase	<i>GAPDH</i>	GACCCAGATGTGCAGGTAGC	GTCAACAAACACGCCAGGTC

**Table S2.** Gene expression of Scots pine saplings

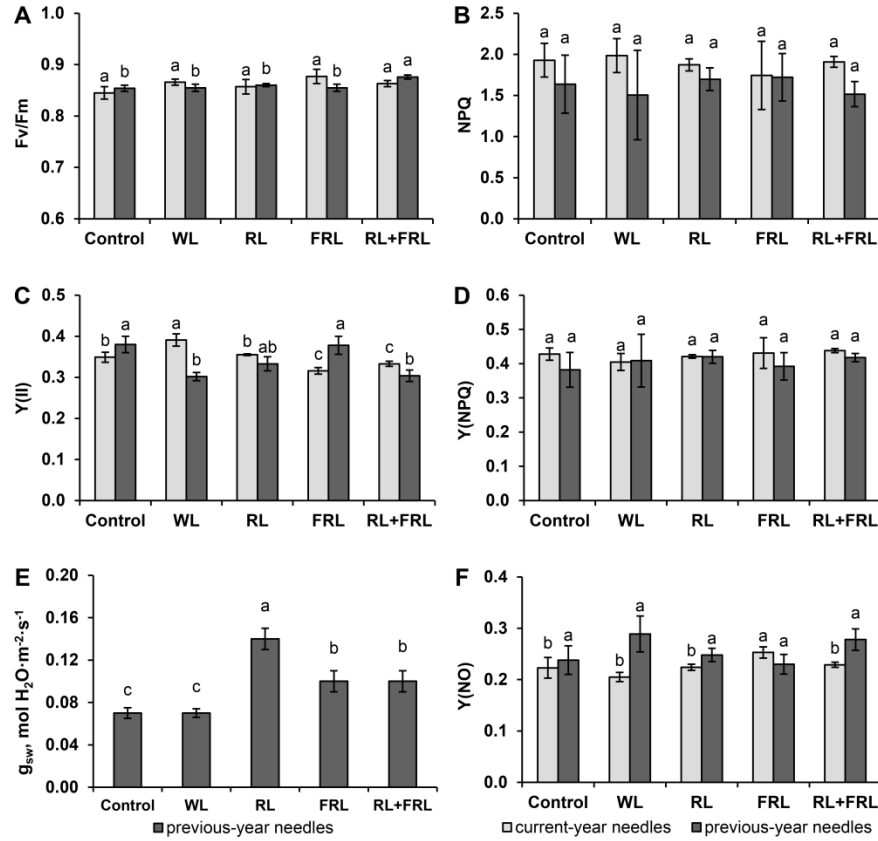
Genes	Control	Additional light			
		WL	RL	FRL	RL+FRL
current-year needles					
ARF16	1.00 ± 0.00 <sup>d</sup>	0.90 ± 0.05 <sup>d</sup>	4.72 ± 0.10 <sup>c</sup>	4.98 ± 0.12 <sup>b</sup>	9.01 ± 0.09 <sup>a</sup>
AUX/IAA	1.00 ± 0.00 <sup>c</sup>	0.66 ± 0.02 <sup>d</sup>	4.07 ± 0.46 <sup>b</sup>	0.61 ± 0.02 <sup>d</sup>	8.12 ± 0.10 <sup>a</sup>
JAZa	1.00 ± 0.00 <sup>d</sup>	1.35 ± 0.08 <sup>c</sup>	6.21 ± 0.55 <sup>b</sup>	0.10 ± 0.03 <sup>e</sup>	8.74 ± 0.50 <sup>a</sup>
HPT1	1.00 ± 0.00 <sup>b</sup>	1.32 ± 0.10 <sup>ab</sup>	0.88 ± 0.05 <sup>b</sup>	1.08 ± 0.08 <sup>b</sup>	1.50 ± 0.13 <sup>a</sup>
BR-a-RED	1.00 ± 0.00 <sup>c</sup>	1.66 ± 0.13 <sup>c</sup>	5.56 ± 0.71 <sup>b</sup>	5.55 ± 0.42 <sup>b</sup>	8.26 ± 0.62 <sup>a</sup>
BRZ2	1.00 ± 0.00 <sup>b</sup>	0.48 ± 0.19 <sup>b</sup>	0.92 ± 0.26 <sup>b</sup>	0.71 ± 0.21 <sup>b</sup>	5.19 ± 0.14 <sup>a</sup>
mono-TERP	1.00 ± 0.00 <sup>d</sup>	3.11 ± 0.22 <sup>c</sup>	4.89 ± 0.28 <sup>b</sup>	1.47 ± 0.12 <sup>d</sup>	8.56 ± 0.47 <sup>a</sup>
α-TERP	1.00 ± 0.00 <sup>c</sup>	0.62 ± 0.08 <sup>c</sup>	0.47 ± 0.06 <sup>c</sup>	3.64 ± 0.27 <sup>b</sup>	8.28 ± 0.42 <sup>a</sup>
CHS	1.00 ± 0.00 <sup>c</sup>	0.23 ± 0.02 <sup>d</sup>	1.62 ± 0.13 <sup>b</sup>	0.28 ± 0.02 <sup>d</sup>	8.61 ± 0.56 <sup>a</sup>
cambium of previous-year shoot					
ARF16	1.00 ± 0.00 <sup>b</sup>	0.80 ± 0.33 <sup>b</sup>	0.68 ± 0.30 <sup>b</sup>	1.95 ± 0.35 <sup>b</sup>	5.03 ± 0.09 <sup>a</sup>
AUX/IAA	1.00 ± 0.00 <sup>b</sup>	1.82 ± 0.37 <sup>b</sup>	2.29 ± 0.31 <sup>ab</sup>	2.42 ± 0.48 <sup>ab</sup>	3.40 ± 0.47 <sup>a</sup>
JAZa	1.00 ± 0.00 <sup>d</sup>	4.65 ± 0.38 <sup>b</sup>	2.68 ± 0.41 <sup>c</sup>	0.99 ± 0.18 <sup>d</sup>	8.33 ± 0.60 <sup>a</sup>
HPT1	1.00 ± 0.00 <sup>a</sup>	1.03 ± 0.17 <sup>a</sup>	1.25 ± 0.14 <sup>a</sup>	1.17 ± 0.31 <sup>a</sup>	1.55 ± 0.19 <sup>a</sup>
BR-a-RED	1.00 ± 0.00 <sup>c</sup>	0.61 ± 0.04 <sup>e</sup>	0.81 ± 0.06 <sup>d</sup>	2.75 ± 0.42 <sup>b</sup>	7.09 ± 0.79 <sup>a</sup>
BRZ2	1.00 ± 0.00 <sup>b</sup>	0.47 ± 0.11 <sup>b</sup>	2.72 ± 0.28 <sup>a</sup>	2.23 ± 0.26 <sup>a</sup>	2.61 ± 0.28 <sup>a</sup>
mono-TERP	1.00 ± 0.00 <sup>a</sup>	0.55 ± 0.34 <sup>a</sup>	0.38 ± 0.16 <sup>a</sup>	0.44 ± 0.25 <sup>a</sup>	0.87 ± 0.29 <sup>a</sup>
α-TERP	1.00 ± 0.00 <sup>a</sup>	0.40 ± 0.04 <sup>a</sup>	0.20 ± 0.04 <sup>a</sup>	0.48 ± 0.20 <sup>a</sup>	0.56 ± 0.30 <sup>a</sup>
CHS	1.00 ± 0.00 <sup>a</sup>	0.85 ± 0.08 <sup>ab</sup>	0.56 ± 0.06 <sup>b</sup>	0.26 ± 0.05 <sup>b</sup>	0.47 ± 0.21 <sup>b</sup>
ARF 3/4	1.00 ± 0.00 <sup>d</sup>	2.17 ± 0.31 <sup>c</sup>	2.93 ± 0.28 <sup>c</sup>	4.41 ± 0.32 <sup>b</sup>	6.73 ± 0.52 <sup>a</sup>
young roots					
ARF16	1.00 ± 0.00 <sup>b</sup>	2.30 ± 0.18 <sup>b</sup>	2.19 ± 0.14 <sup>b</sup>	6.23 ± 0.50 <sup>a</sup>	7.92 ± 0.60 <sup>a</sup>
AUX/IAA	1.00 ± 0.00 <sup>c</sup>	1.12 ± 0.11 <sup>c</sup>	1.05 ± 0.14 <sup>c</sup>	8.65 ± 0.23 <sup>a</sup>	7.86 ± 0.43 <sup>b</sup>
JAZa	1.00 ± 0.00 <sup>a</sup>	1.31 ± 0.18 <sup>a</sup>	1.19 ± 0.10 <sup>a</sup>	1.04 ± 0.11 <sup>a</sup>	0.98 ± 0.07 <sup>a</sup>
HPT1	1.00 ± 0.00 <sup>c</sup>	1.69 ± 0.24 <sup>c</sup>	6.16 ± 0.61 <sup>b</sup>	4.63 ± 0.36 <sup>b</sup>	9.02 ± 0.29 <sup>a</sup>
BR-a-RED	1.00 ± 0.00 <sup>d</sup>	1.29 ± 0.10 <sup>d</sup>	2.16 ± 0.20 <sup>c</sup>	2.93 ± 0.19 <sup>b</sup>	6.14 ± 0.45 <sup>a</sup>
BRZ2	1.00 ± 0.00 <sup>d</sup>	1.65 ± 0.18 <sup>cd</sup>	2.18 ± 0.08 <sup>c</sup>	3.83 ± 0.28 <sup>b</sup>	8.20 ± 0.54 <sup>a</sup>
mono-TERP	1.00 ± 0.00 <sup>b</sup>	0.93 ± 0.14 <sup>b</sup>	1.29 ± 0.18 <sup>b</sup>	1.12 ± 0.21 <sup>b</sup>	3.12 ± 0.35 <sup>a</sup>
α-TERP	1.00 ± 0.00 <sup>c</sup>	0.32 ± 0.13 <sup>c</sup>	0.32 ± 0.10 <sup>c</sup>	4.11 ± 0.36 <sup>b</sup>	7.75 ± 0.27 <sup>a</sup>
CHS	1.00 ± 0.00 <sup>b</sup>	1.02 ± 0.26 <sup>b</sup>	1.26 ± 0.19 <sup>b</sup>	2.97 ± 0.12 <sup>a</sup>	3.46 ± 0.30 <sup>a</sup>
ARF 3/4	1.00 ± 0.00 <sup>d</sup>	0.40 ± 0.07 <sup>e</sup>	6.87 ± 0.68 <sup>b</sup>	8.98 ± 0.25 <sup>a</sup>	4.52 ± 0.38 <sup>c</sup>

Different letters within each row indicate significant differences ( $p < 0.05$ ), according to ANOVA, followed by Duncan's method (regular letters) or Kruskal–Wallis one-way ANOVA on ranks, followed by the Student–Newman–Keuls post hoc test (italic letters).

**Table S3.** Effect of light of different spectral compositions on antioxidant capacity, phenolic compound, total terpenoids and main photosynthetic pigments content in previous-year needles.

Previous-year needles	Control	WL	RL	FRL	RL+FRL
TEAC, μmol Trolox/g FW	555.1 ± 63.5 a	532.8 ± 46.9 a	490.1 ± 16.1 a	563.4 ± 38.6 a	469.2 ± 41.9 a
GAE, mg/g FW	22.2 ± 2.1 <i>a</i>	21.1 ± 0.8 <i>a</i>	21.8 ± 0.6 <i>a</i>	24.5 ± 2.1 <i>a</i>	20.1 ± 0.9 <i>a</i>
Flavonoids, mg catechin/g FW	11.3 ± 1.5 a	10.7 ± 0.5 a	10.2 ± 0.6 a	11.5 ± 0.7 a	9.8 ± 0.6 a
Catechins+proanthocyanidins, mg catechin/g FW	30.9 ± 3.9 a	33.9 ± 3.2 a	33.1 ± 1.9 a	35.0 ± 1.8 a	33.0 ± 2.3 a
Proanthocyanidins, mg cyaniding/g FW	11.1 ± 0.9 a	12.4 ± 0.8 a	11.7 ± 0.4 a	12.4 ± 0.3 a	11.5 ± 0.4 a
Terpenoids, mg/g DW	5.96 ± 0.36 a	6.45 ± 0.18 a	6.25 ± 0.47 a	7.12 ± 0.29 a	6.06 ± 0.27 a
Chl <i>a</i> , mg/g DW	2.66 ± 0.12 a	2.79 ± 0.24 a	2.96 ± 0.13 a	2.92 ± 0.29 a	2.78 ± 0.14 a
Chl <i>b</i> , mg/g DW	1.23 ± 0.06 a	1.27 ± 0.11 a	1.33 ± 0.08 a	1.55 ± 0.17 a	1.37 ± 0.09 a
Carotenoids, mg/g DW	0.54 ± 0.02 a	0.57 ± 0.05 a	0.61 ± 0.03 a	0.65 ± 0.06 a	0.58 ± 0.03 a
Chl <i>a</i> /Chl <i>b</i>	2.16 ± 0.02 a	2.19 ± 0.02 a	2.24 ± 0.04 a	1.89 ± 0.03 c	2.04 ± 0.04 b
Carotenoids/chlorophylls <i>a+b</i> ratio	0.139 ± 0.001 b	0.139 ± 0.001 b	0.143 ± 0.001 ab	0.145 ± 0.002 a	0.141 ± 0.002 ab

Different letters within each row indicate significant differences ( $p < 0.05$ ) according to ANOVA on ranks followed by Duncan's Multiple Range post hoc test (regular letters) or Kruskal–Wallis one-way ANOVA on ranks followed by Student–Newman–Keuls post hoc test (*italic letters*). TEAC, Trolox equivalent antioxidant capacity; GAE, gallic acid equivalents.



**Figure S1.** Effect of additional light on stomatal conductance ( $g_{sw}$ , mol  $H_2O$   $m^{-2}s^{-1}$ ) (E) and the main parameters of chlorophyll *a* fluorescence:  $F_v/F_m$  (the maximum quantum yield of PSII) (A);  $Y(II)$  (PSII effective quantum yield) (C); NPQ (non-photochemical fluorescence quenching) (B);  $Y(NO)$  (quantum yield of non-regulated non-photochemical energy dissipation in PSII) (F),  $Y(NPQ)$  (quantum yield of regulated non-photochemical energy dissipation in PSII) (D) in the previous-year and current-year needles of Scots pine.