Light quality-dependent regulation of photoprotection and antioxidant properties in rice seedlings grown under different light-emitting diodes

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Abstract

We examined light quality-dependent regulation of photoprotection and antioxidant properties in rice seedlings. Seedlings grown under red or green light-emitting diodes (LEDs) showed declines in quantum yield of electron transport through PSII, maximum relative electron transport rates, and photochemical quenching compared to the white and blue LED, but no difference in maximum photochemical efficiency of PSII was found. Blue LED treatment resulted in greater increases in not only nonphotochemical quenching, but also zeaxanthin formation, expressions of β-carotene hydroxylase and violaxanthin de-epoxidase, and anthocyanin compared to other LED treatments. By contrast, green and red LEDs increased activities of superoxide dismutase and ascorbate peroxidase as well as expression levels of catalase (CAT)α and CATβ to white and blue LEDs. Green and red LEDs also greatly upregulated expressions of phytochrome (PHY) A, PHYB, PHYC, and cryptochrome 2 compared to white and blue LEDs, whereas blue LED upregulated CRY1. Our results demonstrate that specific wavelengths of different LEDs differentially influence photosynthetic performance, photoprotection, and antioxidant properties, possibly through regulating expressions of phytochrome and cryptochrome genes in a light-quality-dependent manner.

Keywords: antioxidant enzyme; chlorophyll fluorescence quenching; xanthophyll.

Introduction

Light is not only an essential energy source for plants but also an important signal influencing plant development (Wu et al. 2007, Tran and Jung 2017, Yu et al. 2017). When more light is absorbed than utilized for photosynthesis, the excess excitation energy should be dissipated otherwise it is passed to molecular oxygen and reactive oxygen species (ROS) are generated. Therefore, photosynthesis could not exist without efficient photoprotective mechanisms against potential damage induced by unutilized light energy (Demmig-Adams et al. 2012, Wang et al. 2018). Excitation...
energy in excess can be removed by de-excitation of excited singlet chlorophyll (Chl*) (dissipation) and de-excitation of reactive forms of oxygen (oxidation) (Demmig-Adams et al. 2012).

Under excess light conditions, light-harvesting complexes (LHCs) perform photoprotective quenching functions to prevent harmful chemical species such as triplet chlorophyll (Chl*) and singlet oxygen (O₂) from forming and damaging the photosynthetic apparatus (Demmig-Adams et al. 2012). Chl a fluorescence is used as an indicator of photosynthetic energy conversion, and a decrease in fluorescence as a result of light energy being converted into chemical energy is called photochemical quenching (qP) (Murchie and Lawson 2013, Foyer et al. 2017). The qP indicates not only the capacity of open PSII reaction centers and the excited energy used in photosynthesis but also reflects the stress induced by excess energy excitation on PSII. Nonphotochemical quenching (NPQ) is a process in which excess absorbed light energy is dissipated into heat, and it is a photoprotection mechanism against high-light stress (Ruban 2016).

Carotenoids (Car) have been known as the nonenzymatic antioxidants to protect the photosynthetic apparatus, and photoprotection by the dissipation of excess light energy is aided by the xanthophyll cycle carotenoids (Gilmore and Yamamoto 1991, Demmig-Adams et al. 2012, Jahns and Holzwarth 2012). Violaxanthin de-epoxidase (VDE) converts violaxanthin to zeaxanthin via the intermediate antheraxanthin, thus sustaining dissipation of excess excitation energy as heat at PSII (Jahns and Holzwarth 2012). The other photoprotective pigments, anthocyanins, function as a light attenuator to protect the photochemical apparatus against excess light (Chalker-Scott 1999, Nakabayashi et al. 2014). While thermal dissipation of energy is useful in preventing the formation of ROS, additional photoprotective mechanisms exist to detoxify ROS. Antioxidation of harmful ROS generated under excess light acts as a sink for unutilized electrons (Demmig-Adams et al. 2012). ROS are generated when electrons not utilized in photosynthesis are donated to O₂, thus creating superoxide (O₂•−) that can be converted to hydrogen peroxide (H₂O₂) by superoxide dismutase (SOD). Ascorbate peroxidase (APX), peroxidase (POD), and catalase (CAT) subsequently detoxify H₂O₂ (Demmig-Adams et al. 2012, Foyer and Noctor 2013).

In higher plants, specific wavelengths of light play important roles in the regulation of growth, development of photosynthetic apparatus, Chl formation, and stomatal opening (Wu et al. 2007, Yu et al. 2017). Phytochromes are red/far-red light receptors that regulate photomorphogenesis, including de-etiolation and plant shape formation (Wang and Deng 2003). Cryptochromes are the blue-light photoreceptors that mediate inhibition of hypocotyl elongation, enhancement of cotyledon expansion, anthocyanin accumulation, and stomatal opening (Zhang et al. 2018). Also, cryptochromes regulate blue light-dependent biosynthesis of ROS (Consentino et al. 2015, El-Elawi et al. 2017) and are considered as key regulators of a range of plant stress responses, such as high light, heat, drought, and salinity (Ma et al. 2016, D’Amico-Damiano and Carvalho 2018). The green light has discrete effects on plant biology (Folta and Maruhnich 2007), and the mechanisms that sense this light quality are now being elucidated.

Light-emitting diodes (LEDs) are characterized by their narrow spectrum and emerging as a promising tool for greenhouse crop production (Mitchell et al. 2012). Many studies have shown the effects of light intensities on photoinhibition of PSII (Jung et al. 2000, Hakala et al. 2005, Kouřil et al. 2013) and light quality on antioxidant capacity (Wu 2016, Yu et al. 2017). However, the effects of light quality on photoprotection and enzymatic antioxidant systems, particularly with the involvement of photoreceptors in these processes, are not well characterized yet. In our previous study, rice seedlings grown under white and blue LEDs were stronger, as indicated by greater leaf area and shoot biomass, than those of green and red LEDs, showing that light qualities influence the growth characteristics of seedlings (Tran and Jung 2017).

In the present study, we examined the influence of light spectral quality on photosynthetic performance and energy dissipation by analyzing photosynthetic efficiency and fluorescence quenching in rice seedlings grown under different LED wavelengths. To assess photoprotection and antioxidant properties under different LEDs, we determined the activities and gene expression of ROS-scavenging enzymes as well as the contents of photoprotective pigments. We also analyzed expression profiles of phytochrome and cryptochrome genes to characterize the involvement of photoreceptors in the modulation of energy dissipation and antioxidation.

Materials and methods

Plant growth and light treatment conditions: Rice (Oryza sativa cv. Dongjin) seeds were germinated at 25°C under the dark conditions for 5 d before irradiation. Then, the rice seedlings were grown under various types of light-emitting diodes (Union LED Electronic Co., Siheung-si, Korea) for 10 d. Each LED treatment was conducted in separately controlled chambers to be free from spectral interference among treatments. Fifteen pots were placed in each LED chamber and seven plants were grown in each pot which was filled with commercial greenhouse compost. The LED array chambers were programmed to provide a 14-h-light/10-h-dark photoperiod at PPFD of approximately 150 μmol/m² s −1. The rice seedlings were grown under four different light sources with broad-spectrum white LED (420–680 nm) as a control, blue LED (460–490 nm), green LED (520–550 nm), and red LED (620–650 nm) conditions. The youngest, fully developed leaves of rice seedlings were sampled 4 h after illumination (11:00 h) on the last day of LED treatments.

Measurement of Chl a fluorescence: Chl a fluorescence was measured in vivo using a pulse amplitude modulation fluorometer (JUNIOR-PAM, Walz, Effeltrich, Germany) after dark adaptation for 30 min. The minimum fluorescence (F₀) at open PSII centers was determined by measuring
light, while the maximum fluorescence \((F_{\text{m}})\) at closed PSII centers was examined after an application of a 0.8-s pulse of saturating light \([3,500 \mu\text{mol}(\text{photon}) \text{m}^{-2} \text{s}^{-1}]\). Actinic light \([150 \mu\text{mol}(\text{photon}) \text{m}^{-2} \text{s}^{-1}]\), growth light source for each LED treatment was applied to measure the steady-state Chl fluorescence (\(F_s\)). In the light-adapted state, \(F_{\text{m}}\) was measured by applying a saturating pulse, while \(F_{\text{r}}\) was measured by switching off the actinic light for 2 s after the saturating pulse and applying far-red light. The ratio of \(F_{\text{r}}\) to \(F_{\text{m}}\), representing the activity of PSII, was used to assess the functional damage to the plants. The quantum yield of electron transport through PSII (\(\Phi_{\text{PSII}}\)) was calculated according to Genty et al. (1989). Coefficient of photochemical fluorescence quenching \((q)\) was used to estimate the fraction of open PSII centers and calculated as follows: \(q = (F_{\text{m}}' - F')/(F_{\text{m}}' - F_0)\), where \(F_{\text{r}}\), the minimal Chl fluorescence yield of a preilluminated sample; \(F_{\text{r}}\), the level of the fluorescence curve during treatments and shortly before a saturating pulse is applied; \(F_{\text{r}}\), maximal Chl fluorescence in the light-adapted state (Schreiber et al. 1986). NPQ was also quantified, as previously done by Bilger and Björkman (1990) according to the Stern-Volmer equation, \(NPQ = F/F_{\text{m}} - 1\).

The rapid light curve (RLC) was also determined with a gradual increase of irradiance in 12 steps with 10-s intervals ranging from 0 to 1,500 \(\mu\text{mol}(\text{photon}) \text{m}^{-2} \text{s}^{-1}\) using blue light of PAM instrument as actinic light for RLC analysis. The parameters of the RLCs, including maximum relative ETR \((\text{re} \text{ETR}_{\text{max}})\), the efficiency of electron transport \(\alpha\), the initial slope of the light curve), electron transport \((\alpha, \text{the initial slope of the light curve})\), and the light saturation \(I_{\text{sat}}\) \((\text{irradiance at the onset of light saturation})\), were calculated following the Platt’s formula (Jassby and Platt 1976, Harrison and Platt 1986).

**Protein extraction, SDS-PAGE, and immunoblot analysis:** Total protein was extracted by grinding frozen leaf material in a buffer containing 2% (w/v) SDS, 56 mM Na2CO3, 12% (w/v) sucrose, 56 mM diithiothreitol (DTT), and 2 mM EDTA, pH 8.0, followed by sonication. Total soluble proteins were collected by centrifugation at 12,000 \(\times g\) for 20 min and used for gel blot analysis. The proteins were separated on a 14% SDS-PAGE and electroblotted onto PVDF membranes. Immunodetection was performed using standard procedures (Roche, Basel, Switzerland). The polyclonal antibodies against light-harvesting chlorophyll-binding proteins of PSI (Lhca) and PSII (Lhcb) and \(\alpha\)-tubulin were purchased from Agrisera.

**Determination of carotenoids (Car):** For Car analysis, leaf tissues were ground in 100% acetone containing 10 mg of CaCO3. The extracts were centrifuged at 16,000 \(\times g\) for 10 min and the resulting supernatants were collected. The pigments were separated by HPLC as previously described by Gilmore and Yamamoto (1991) using Prominance HPLC system (Shimadzu, Kyoto, Japan) equipped with SPD-M20A Prominance Photodiode Array Detector (Shimadzu). A Spherisorb ODS-1 column (5-μm particle size, 250 mm × 4.6 mm inner diameter) was obtained from Waters (Milford, MA, USA). Solvent A (acetonitrile:methanol:HCl buffer 0.1 M, pH 8.0, 72:8:3, v/v/v) was run isocratically from 0 to 4 min followed by a 2.5-min linear gradient to 100% solvent B (methanol:hexane, 4:1, v/v) at flow rate of 2 ml min\(^{-1}\). HPLC runs were carried out at 445 nm for xanthophylls and \(\beta\)-carotene. Specific extinction coefficients (E1% 1 cm\(^{-1}\)) used for quantification were lutein (2,550 at 445 nm), zeaxanthin (2,540 at 450 nm), antheraxanthin (2,350 at 446 nm), violaxanthin (2,550 at 434 nm), neoxanthin (2,243 at 439 nm), and \(\beta\)-carotene (2,590 at 450 nm).

**Determination of anthocyanin:** Anthocyanins were extracted by homogenizing leaves [0.1 g of fresh mass (FM)] in a precooled mortar with 1 ml of acidified (1% HCl) methanol and maintained at 4°C for 4 h to avoid the degradation of Chl whose products may interfere with the absorption of anthocyanins at 530 nm (peak absorption of anthocyanin) and 657 nm (peak absorption of Chl degradation products) (Mancinelli et al. 1975). Particles were removed by centrifugation at 10,000 \(\times g\) for 30 min. Anthocyanins were determined by using \(A_{530} = 0.25 \Delta A_{530}\) to account for the contribution by Chl. Anthocyanin content was calculated as cyanidin-3-glucoside using 29,600 M\(^{-1}\) cm\(^{-1}\) as molecular extinction coefficient and 445 as molecular mass.

**Assays for antioxidant enzymes:** Leaves (0.25 g of FM) were macerated to a fine powder in a mortar under liquid \(\text{N}_2\). Soluble proteins were extracted by homogenizing the powder in 2 ml of 100 mM potassium phosphate buffer, pH 7.5, containing 2 mM EDTA, 1% PVP-40, and 1 mM phenylmethylsulfonyl fluoride. Insoluble material was removed by centrifugation at 15,000 \(\times g\) for 20 min at 4°C. As for maintenance of consistent catalase (CAT, EC 1.11.1.6) electrophoretic mobility was found to require the presence of DTT, an aliquot of each sample was made to 10 mM DTT to be used for the CAT zymogram. Equal amounts of protein were electrophoresed on 10% non-denaturing polyacrylamide gels at 4°C for 1.5 h at a constant current of 30 mA. Following separation of ascorbate peroxidase (APX, EC 1.11.1.11), gels were soaked in 50 mM potassium phosphate buffer, pH 7.0, containing 2 mM ascorbate for 30 min (Rao et al. 1996). The gels were incubated in the same buffer containing 4 mM ascorbate and 2 mM \(\text{H}_2\text{O}_2\) for 20 min and then soaked in 50 mM potassium phosphate buffer, pH 7.8, containing 28 mM tetramethyl ethylenediamine and 2.45 mM nitroblue tetrazolium (NBT) for 15 min. The CAT activity was detected by incubating the gels in 3.27 mM \(\text{H}_2\text{O}_2\) for 25 min, rinsing them in water, and staining them in a solution of 1% potassium ferricyanide and 1% ferric chloride for 4 min (Woodbury et al. 1971). The staining of peroxidase (POD, EC 1.11.1.7) isoforms was achieved by incubating gels in sodium citrate buffer (pH 5.0) containing 9.25 mM \(\beta\)-phenylenediamine and 3.92 mM \(\text{H}_2\text{O}_2\) for 15 min (Olson and Varner 1993). Gels were stained for superoxide dismutase (SOD, EC 1.15.1.1) isoforms by soaking in 50 mM potassium phosphate (pH 7.8) containing 2.5 mM NBT in darkness for 25 min, followed by soaking in 50 mM potassium phosphate (pH 7.8) containing 28 mM NBT and 28 \(\mu\)M riboflavin in
darkness for 30 min. The gels were then exposed to light for 30 min (Rao et al. 1996).

APX activity was measured spectrophotometrically (UV-2550, Shimadzu, Japan) by monitoring the decline in A_290 as ascorbate was oxidized (Chen and Asada 1989). The 3-ml reaction volume contained 100 mM potassium phosphate buffer (pH 7.5), 0.5 mM ascorbate, and 0.2 mM H_2O_2 at 25°C. Total CAT activity was performed spectrophotometrically in a 3 ml volume containing 50 mM potassium phosphate buffer, pH 7.0, containing 20 mM H_2O_2 by monitoring H_2O_2 destruction at 240 nm (Beers and Sizer 1952). POD activity was determined specifically with guaiacol at 470 nm (Egley et al. 1983). The reaction mixture contained 40 mM potassium phosphate buffer (pH 6.9), 1.5 mM guaiacol, and 6.5 mM H_2O_2 in a 3-ml volume. For APX, CAT, and POD, enzyme activities were expressed as μmol g–1(FM) min–1. SOD activity was determined using the xanthine oxidase-cytochrome c method at pH 10.0 (Baum and Scandalios 1981). The assay was performed at 25°C in a 3-ml volume containing 50 mM sodium carbonate buffer (pH 10.0), 0.1 mM EDTA, 15 μM ferricytochrome c, and 0.05 mM xanthine. SOD activity was expressed as units g–1(FM).

RNA extraction and qRT-PCR: Total RNA was prepared from leaf tissues using TRizol Reagent (Invitrogen, Waltham, MA, USA), and 5 μg of RNA from each sample was used for the reverse transcription reaction (ImProm-II™ Reverse Transcription System, Promega, WI, USA). Subsequently, 50 ng of cDNA was used for qRT-PCR analysis. The qRT-PCR analysis was carried out using the StepOnePlus Real-Time PCR System (Applied Biosystems, Waltham, MA, USA) using Power SYBR Green PCR Master Mix (Applied Biosystems) and gene-specific primers (Table 1S, supplement). The qRT-PCR program consisted of 2 min at 50°C, 10 min at 95°C, and 40 cycles of 15 s at 95°C, and 1 min at 60°C. A melting curve analysis was performed after every PCR reaction to confirm the accuracy of each amplified product. All reactions were set up in triplicate. The sample of white LED treatment was used as the calibrator, with the expression level of the sample set to 1. Actin was used as an internal control.

Statistical analysis: Data were expressed as means ± SE. Differences were analyzed by Duncan's multiple range test. p-values<0.05 were considered to be significant. Correlation analysis was also performed using the SPSS software (SPSS Inc., Chicago, IL, USA).

Results

Photosynthetic efficiency and photoprotective quenching mechanisms: The 5-d-old etiolated rice seedlings were exposed to white, blue, green, or red LED light in cycles of 14-h light and 10-h dark for 10 d. In all seedlings grown under different LEDs, the youngest, fully developed leaves did not develop any necrosis. Although seedlings were grown under a moderate PPFD light intensity of 150 μmol( photon) m−2 s−1, necrotic spots, which are characteristics of photodamage, often appeared on older leaves of seedlings grown under green or red LED (data not shown). Seedlings grown under green and red LEDs also exhibited reductions in surface area of the youngest, fully developed leaves and dry shoot biomass compared to those of white and blue LEDs, while seedlings grown under blue LED exhibited the shortest shoot height and a significant increase in leaf width (Tran and Jung 2017). We supposed that a narrow spectrum of green and red LEDs may impose a lower PSII excitation pressure on whole seedlings compared to blue and broad-spectrum white LEDs, reducing shoot biomass. Thus, we examined how seedlings treated with different wavelengths of LEDs develop differential photoprotective mechanisms to optimize their growth.

For rapid estimation of photosynthetic characteristics and quenching mechanisms under different LED light qualities, we measured Chl fluorescence parameters in the youngest, fully developed leaves. There was no difference in F_v/F_m, a trait positively correlated with the vitality of PSII, of seedlings grown under monochrome blue, green, or red LED compared to broad-spectrum white LED (Fig. 1A). Seedlings grown in green or red LED showed declines in Φ_PSII, which is the quantum yield of electron transport through PSII, compared to white and blue LEDs (Fig. 1B).

Leaf photosynthetic capacity was also assessed by the RLC approach. The relative electron transfer rate of PSII (rETR) rapidly increased with increasing light intensity (Fig. 1E). The rETR in red and green LED treatments reached a steady state faster, while white and blue LED treatments reached the steady state at higher light levels. The maximum electron transport rate of PSII, rETR_max, and the light-saturation point, I_s, in seedlings under white and blue LED treatments were significantly greater than those of green and red LEDs (Table 1). The initial slope of the light response, α, was lower in green and red LEDs than that of white and blue LEDs, with the lowest value in green LED.

The qP is used to indicate the onset of photoinhibition because a linear relationship was observed between qP and photoinhibition of PSII (van Wijk and van Hasselt 1993). Seedlings grown in red and green LEDs showed lower qP values as compared to white and blue LEDs (Fig. 1C). Nonradiative energy dissipation through NPQ was higher by 44% in blue LED treatment, compared to that of white LED treatment (Fig. 1D). In comparison to white LED, NPQ levels were lower by 60 and 54% in seedlings under green and red LEDs, respectively (Fig. 1D). Next, we evaluated the organization of the photosynthetic apparatus in seedlings under respective light irradiations of LEDs by quantitatively analyzing Lhca and Lhcb proteins of PSI and PSII, respectively. Expression levels of Lhca and Lhcb proteins, which are major components of light-harvesting antennae, were not different after all LED treatments (Fig. 1F).

Photoprotective pigments and expression of genes involved in zeaxanthin biosynthesis: To assess the effects of light quality on photoprotective pigments, we
examined the contents of carotenoids and anthocyanins. The contents of lutein and β-carotene, which are potent ROS scavengers, in seedlings treated with white and red LEDs were higher than those of blue and green LEDs (Table 2). In seedlings grown under blue LED, contents of zeaxanthin, an important component of the xanthophyll cycle, increased by 65% compared to white LED and were higher than those of green and red LEDs by 253 and 283%, respectively (Table 2). Contents of antheraxanthin, violaxanthin, and neoxanthin as well as the sum of violaxanthin, antheraxanthin, and zeaxanthin (VAZ) were not significantly different between various LED treatments (Table 2). The de-epoxidation state of xanthophyll cycle pigments, which is expressed as \([Z + 0.5 A/VAZ]\), exhibited higher values in white and blue LEDs compared to those in green and red LEDs. In comparison to white LED, anthocyanin contents were the highest in seedlings grown under blue LED and lowest in seedlings grown under white and green LEDs (Table 2).

We used qRT-PCR to analyze the effect of different light wavelengths on the expression of genes involved in zeaxanthin formation. In comparison to white LED treatment, the expression level of \(BCH\) encoding β-carotene hydroxylase (BCH), which catalyzes the hydroxylation reaction from β-carotene to zeaxanthin, was about 2-fold upregulated in blue LED treatment (Fig. 2A). Seedlings grown under blue LED also upregulated expression of \(VDE\) encoding VDE, which catalyzes the conversion from violaxanthin to zeaxanthin, compared to white LED (Fig. 2B). Expression levels of \(BCH\) and \(VDE\) in seedlings treated with green and red LEDs were similar to those of white LED.

### Activity and gene expression of ROS-scavenging enzymes

We monitored changes in isozyme profiles and activities of ROS-scavenging enzymes in response to different LED wavelengths. Total SOD activities were higher in seedlings grown under green and red LEDs compared to white and red LEDs, with a greater increase in green LED (Fig. 3A). These were mainly due to the elevated activity of SOD isozymes 2 and 3 in seedlings grown under green LED, and isoform 3 in the case of red LED (Fig. 3E). Although the total CAT activities did not differ between different LEDs (Fig. 3B), the staining activities of CAT isozymes 1 and 2 greatly increased in green LED (Fig. 3D). These were mainly due to the elevated activity of SOD isozymes 2 and 3 in seedlings grown under green and red LED treatments and decreased in blue LED treatment (Fig. 3F) as compared to white LED. The total POD activity was lower in blue LED, but higher in red LED than that of white LED (Fig. 3C). In a comparison of isoforms, seedlings grown under blue LED exhibited lower activities of all POD isozymes compared to white LED (Fig. 3G). POD isozyme 2 decreased in green LED treatment compared to white LED, whereas POD isozymes 3 and 4 increased in green and red LED treatments. Marked increases in both APX-staining activities and total APX activities were observed in seedlings grown under green or red LED, with a greater increase in green LED (Fig. 3D, H).
Table 2. Effects of different wavelengths of LEDs on levels of carotenoid biosynthetic intermediates and anthocyanin. The plants were subjected to the same treatments as in Fig. 1, and treatment notations are the same as in Fig. 1. Total RNAs were purified from plants and reverse transcribed. The resulting cDNAs were used as templates for qRT-PCR using Actin as an internal control. The data represent the mean ± SE of nine replicates from three independent experiments. Means denoted by the same letter did not differ significantly at P<0.05 according to Duncan's multiple range test.

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Lutein [nmol g(^{-1}) (FM)]</th>
<th>β-carotene [nmol g(^{-1}) (FM)]</th>
<th>Z - zeaxanthin [nmol g(^{-1}) (FM)]</th>
<th>A - antheraxanthin [nmol g(^{-1}) (FM)]</th>
<th>V - violaxanthin [nmol g(^{-1}) (FM)]</th>
<th>V + A + Z [nmol g(^{-1}) (FM)]</th>
<th>(Z + 0.5 A)/V [nmol g(^{-1}) (FM)]</th>
<th>Anthocyanin [µg g(^{-1}) (FM)]</th>
</tr>
</thead>
<tbody>
<tr>
<td>White</td>
<td>368 ± 27(a)</td>
<td>3.8 ± 1.5(a)</td>
<td>201 ± 12(a)</td>
<td>156 ± 4(a)</td>
<td>157 ± 3(a)</td>
<td>0.075 ± 0.007(a)</td>
<td>31.5 ± 1.5(a)</td>
<td>3.5 ± 0.3(b)</td>
</tr>
<tr>
<td>Blue</td>
<td>128 ± 1.0(b)</td>
<td>5.2 ± 0.5(b)</td>
<td>183 ± 9(b)</td>
<td>84 ± 3(c)</td>
<td>115 ± 7(b)</td>
<td>0.036 ± 0.007(b)</td>
<td>3.8 ± 0.7(c)</td>
<td>4.2 ± 1.7(c)</td>
</tr>
<tr>
<td>Green</td>
<td>134 ± 7(a)</td>
<td>3.8 ± 0.7(a)</td>
<td>149 ± 4(a)</td>
<td>83 ± 3(c)</td>
<td>149 ± 2(c)</td>
<td>0.036 ± 0.007(b)</td>
<td>3.8 ± 1.1(c)</td>
<td>4.2 ± 1.7(c)</td>
</tr>
<tr>
<td>Red</td>
<td>203 ± 15(b)</td>
<td>3.5 ± 0.3(b)</td>
<td>156 ± 4(a)</td>
<td>103 ± 21(a)</td>
<td>157 ± 3(a)</td>
<td>0.065 ± 0.008(a)</td>
<td>3.5 ± 0.3(b)</td>
<td>4.2 ± 1.7(c)</td>
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</table>

Fig. 2. Expression of genes involved in zeaxanthin formation in rice seedlings grown under different wavelengths of LEDs. (A) β-carotene hydroxylase (BCH). (B) Violaxanthin de-epoxidase (VDE). The plants were subjected to the same treatments as in Fig. 1. Treatment notations are the same as in Fig. 1. Total RNAs were purified from plants and reverse transcribed. The resulting cDNAs were used as templates for qRT-PCR using Actin as an internal control. The white LED (control) was used for normalization, with the expression level of the sample set to 1. The data represent the mean ± SE of nine replicates from three independent experiments. Means denoted by the same letter did not differ significantly at P<0.05 according to Duncan's multiple range test.

We next examined whether the transcript levels of genes encoding H\(_2\)O\(_2\)-scavenging enzymes, CAT and APX, directly correspond to their enzyme activities under different wavelengths of LEDs. Expression of CATa exhibited 2.2- and 3.1-fold upregulation in seedlings treated with green and red LEDs, respectively, compared to white LED treatment, but downregulation in blue LED (Fig. 4A). Seedlings grown under blue, green, or red LEDs exhibited increases in transcript levels of CATb compared to white LED, with greater increases in green and red LEDs (Fig. 4B). However, the transcript level of CATc was highest in seedlings treated with blue LED compared to other LEDs (Fig. 4C). In comparison to white LED, transcript levels of APXa increased only in green LED, whereas transcript levels of APXb increased in blue, green, and red LEDs (Fig. 4D,E). CAT activity showed positive correlations to CATa and CATb with Pearson's correlation coefficients of 0.897 and 0.537, respectively, but a negative correlation to CATc with Pearson's correlation coefficient of −0.712 (Table 3). APX activity showed positive correlations to APXa and APXb with Pearson's correlation coefficients of 0.763 and 0.705, respectively.

Expression profiles of phytochrome and cryptochrome genes: To assess the possible role of photoreceptors in photoprotection and antioxidant properties under different wavelengths of LEDs, we monitored the expression of genes encoding the apoproteins of photoreceptors and cryptochromes, which are red/far-red light and blue light photoreceptors, respectively. In comparison to broad-spectrum white LED, seedlings grown under green or red LED greatly upregulated PHYA, PHYB, and PHYC genes encoding phytochrome A, B, and C, with greater increases of PHYA and PHYB in red LED (Fig. 5A–C). Transcript levels of PHYA, PHYB, and PHYC in seedlings treated with blue LED were not significantly different from those of white LED.
In seedlings grown under blue LED, the expression level of *CRY1*, a gene encoding cryptochrome 1, increased by 78% compared to white LED (Fig. 5D), whereas the expression level of *CRY2* was similar to that of white LED. Expression levels of *CRY1* did not differ significantly between white, green, and red LED treatments, whereas expression levels of *CRY2* were upregulated by 85 and 53% in green and red LED treatments, respectively, compared to white LED (Fig. 5D,E). These results demonstrate that different wavelengths of LEDs greatly influence expression profiles of phytochrome and cryptochrome genes.

**Correlation between antioxidant enzyme activity and expression of photoreceptor genes:** Pearson’s correlation coefficient was applied to evaluate the relationship between antioxidant enzyme activity and expression of photoreceptor genes. The enzyme activities of SOD, CAT, POD, and APX showed positive correlations to *PHYA*, *PHYB*, and *PHYC* (Table 3). Among these, total APX activity showed a significant positive correlation to *PHYC* with Pearson’s correlation coefficient of 0.946. Negative correlations between *CRY1* vs. enzyme activities of SOD, CAT, POD, and APX were observed, with a significant coefficient of –0.931 between *CRY1* and POD activity. By contrast, *CRY2* showed a significant positive correlation to SOD and APX activities.

Expression levels of *PHYA*, *PHYB*, and *PHYC* showed positive correlations to *CATa*, *CATb*, and *APXb*, with significant correlations to *CATa* and *CATb*, whereas negative correlations between *CATc* vs. expression levels of *PHYA*, *PHYB*, and *PHYC* were found (Table 3). Negative correlations between *CRY1* vs. expression levels of *CATa* and *CATb* were observed, whereas *CRY1* showed a significant positive correlation to *CATc* with a coefficient of 0.995. *CRY2* showed positive correlations to *CATa*, *CATb*, *APXa*, and *APXb* with Pearson’s correlation coefficients of 0.748, 0.969, 0.833, and 0.728, respectively. Overall, the activities of SOD, CAT, POD, and APX as well as the expression levels of *CATa*, *CATb*, and *APXb* showed positive correlations to *PHYA*, *PHYB*, and *PHYC*. On the other hand, *CRY1* and *CRY2* exhibited negative and positive correlations, respectively, to the activities and expression of some antioxidant enzymes.

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**Fig. 3.** Isozyme profiles and activities of ROS-scavenging enzymes in rice seedlings grown under different wavelengths of LEDs. (A,E) Superoxide dismutase (SOD); (B,F) catalase (CAT); (C,G) peroxidase (POD); (D,H) ascorbate peroxidase (APX). The plants were subjected to the same treatments as in Fig. 1. Treatment notations are the same as in Fig. 1. The numbers indicate each isozyme of antioxidant enzymes in order of detected bands from the top. The data represent the mean ± SE of nine replicates from three independent experiments. Means denoted by the same letter did not differ significantly at P<0.05 according to Duncan’s multiple range test.

**Fig. 4.** Effects of different wavelengths of LEDs on the expression of genes encoding the H2O2-scavenging enzymes, catalase (CAT) (A–C) and ascorbate peroxidase (APX) (D,E). The plants were subjected to the same treatments as in Fig. 1. Treatment notations are the same as in Fig. 1. Total RNAs were purified from plants and reverse transcribed. The resultant cDNAs were used as templates for qRT-PCR using Actin as an internal control. The white LED (control) was used for normalization, with the expression level of the sample set to 1. The data represent the mean ± SE of nine replicates from three independent experiments. Means denoted by the same letter did not differ significantly at P<0.05 according to Duncan’s multiple range test.
Discussion

Specific wavelengths of different LEDs greatly influence not only growth characteristics (Tran and Jung 2017), but also photoprotection and antioxidation mechanisms in rice seedlings. When rice seedlings were exposed to different wavelengths of LEDs, levels of Fv/Fm, as well as Lhca and Lhcb proteins in blue, red, and green LEDs were the same as in white LED (Fig. 1A,F), indicating no noticeable damage of photosynthetic apparatus. However, seedlings grown in red or green LED showed low efficiency of photochemical utilization of absorbed light energy, as indicated by the decrease of $\Phi_{PSII}$, rETR$_{max}$, $\alpha$, and $I_h$, compared to seedlings grown in white or blue LED (Fig. 1, Table 1). These results imply the negative effects of monochromatic green and red LED on the photosynthetic performance. Since Chls have light absorption peaks at blue and red wavelengths, light energy absorbed by Chls is likely to be associated with photodamage to PSII (Zavafer et al. 2015). In comparison to white and blue LEDs, Chl contents of green and red LEDs were lower, while Chl $a/b$ higher in blue, red, and green LEDs than that in white LED, indicating that structure of photosynthetic apparatus is altered depending on the light quality, as indicated previously by Tran and Jung (2017). The wavelength of blue light is known as particularly effective photoinhibition (He et al. 2015). However, rice seedlings treated with blue LED showed higher $\Phi_{PSII}$, RLC parameters, and $q_P$ compared to monochromatic green and red LEDs (Fig. 1), which may explain the greatest shoot biomass among different monochrome LEDs (Tran and Jung 2017).

The $q_P$, that is the relative reduction state of QA, reflects the fraction of open PSII reaction centers (Murchie and Lawson 2013). The high $q_P$ value in white and blue LEDs (Fig. 1C) indicates that more absorbed light energy was consumed by photochemical processes in seedlings grown under white or blue light. By contrast, the low $q_P$ value in seedlings grown under green or red LED implies a decrease in the proportion of available excitation energy used for photochemistry. NPQ, a protective process, increased by 44% in blue LED treatment compared to white LED treatment (Fig. 1D). The significant increase of NPQ in blue LED indicates the increased thermal dissipation of excess excitation energy in PSII, which is the first line of defense in photoprotection. In seedlings under green and red LEDs, NPQ levels significantly decreased compared to white LED (Fig. 1D), whereas the red alga Pyropia haitanensis under red light showed the highest NPQ compared to white, blue, and green light (Wu 2016).

Zeaxanthin enhances thermal dissipation of excess absorbed light energy, which is measured as NPQ (Demmig-Adams et al. 2012, Jahns and Holzwarth 2012). Zeaxanthin is synthesized by either conversion of the xanthophylls $\alpha$, $\alpha'$, and $\beta$, or by de-epoxidation of the xanthophylls $\alpha$, $\alpha'$, and $\beta$. This conversion is thought to be a means of transferring excitation energy to non-photochemical processes in green plants (Zavafer et al. 2015). In this study, the xanthophylls are synthesized by either conversion of the xanthophylls $\alpha$, $\alpha'$, and $\beta$, or by de-epoxidation of the xanthophylls $\alpha$, $\alpha'$, and $\beta$. This conversion is thought to be a means of transferring excitation energy to non-photochemical processes in green plants (Zavafer et al. 2015).

Table 3. Pearson’s correlation coefficients of antioxidant enzyme activity vs. the expression of photoreceptor genes. ’’’’ correlation is significant at $P<0.05$ and $P<0.01$, respectively. APX – ascorbate peroxidase; CAT – catalase; CRY – cryptochrome; PHY – phytochrome; POD – peroxidase; SOD – superoxide dismutase.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>PHYA</th>
<th>PHYB</th>
<th>PHYC</th>
<th>CRY1</th>
<th>CRY2</th>
<th>CAT</th>
<th>APX</th>
</tr>
</thead>
<tbody>
<tr>
<td>SOD</td>
<td>0.845</td>
<td>0.768</td>
<td>0.872</td>
<td>$-0.521$</td>
<td>0.998***</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>CAT</td>
<td>0.781</td>
<td>0.824</td>
<td>0.768</td>
<td>$-0.792$</td>
<td>0.408</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>POD</td>
<td>0.723</td>
<td>0.729</td>
<td>0.769</td>
<td>$-0.931^*$</td>
<td>0.436</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>APX</td>
<td>0.904</td>
<td>0.832</td>
<td>0.946*</td>
<td>$-0.688$</td>
<td>0.989***</td>
<td>-</td>
<td>-</td>
</tr>
<tr>
<td>CATa</td>
<td>0.944*</td>
<td>0.933*</td>
<td>0.963**</td>
<td>$-0.873$</td>
<td>0.748</td>
<td>0.897</td>
<td>-</td>
</tr>
<tr>
<td>CATb</td>
<td>0.951*</td>
<td>0.908</td>
<td>0.944*</td>
<td>$-0.567$</td>
<td>0.969***</td>
<td>0.537</td>
<td>-</td>
</tr>
<tr>
<td>CATc</td>
<td>$-0.634$</td>
<td>$-0.574$</td>
<td>$-0.755$</td>
<td>0.995**</td>
<td>$-0.565$</td>
<td>$-0.712$</td>
<td>-</td>
</tr>
<tr>
<td>APXa</td>
<td>0.446</td>
<td>0.327</td>
<td>0.514</td>
<td>$-0.220$</td>
<td>0.833</td>
<td>-</td>
<td>0.763</td>
</tr>
<tr>
<td>APXb</td>
<td>0.875</td>
<td>0.904</td>
<td>0.773</td>
<td>$-0.257$</td>
<td>0.728</td>
<td>-</td>
<td>0.705</td>
</tr>
</tbody>
</table>
from β-carotene or de-epoxidation from violaxanthin (Demming-Adams et al. 2012, Jahns and Holzwarth 2012). In addition to low β-carotene content, expression levels of BCH and VDE were greater in blue LED treatment than those of white, green, and red LED treatments (Table 2, Fig. 2), showing that the great accumulation of zeaxanthin can be explained by transcriptional regulation of zeaxanthin biosynthetic genes, with more conversion from β-carotene. Anthocyanin is another pigment that has some protection for the chloroplasts against excess radiation (Chalker-Scott 1999, Zhang et al. 2016). Among different LEDs, the highest anthocyanin contents in seedlings grown under blue LED (Table 1) were similar to the findings that anthocyanin contents in grape and strawberry treated with blue wavelength were higher than those with red wavelength (Kondo et al. 2014, Zhang et al. 2018). Our results demonstrate that rice seedlings grown under different light qualities produce differential contents of photoprotective pigments, particularly with high accumulations of zeaxanthin and anthocyanin under blue LED. Although anthocyanins are not involved in the plasticized energy dissipation processes mediated by NPQ, the accumulation of anthocyanins may provide an additional means of photoprotection by acting as a light screen.

ROS depend on not only nonenzymatic antioxidants including Car but also enzymatic antioxidants (Suzuki et al. 2012, Noctor et al. 2014). Increased activities of SOD in seedlings grown under green or red LED (Fig. 3A,E) indicated the increased conversion of O₂⁻ into H₂O₂. Upregulations of CATa and CATb in seedlings treated with green and red LEDs were correlated with the increased staining activities of CAT, but not with total CAT activities (Figs. 3, 4). Red light also significantly increased CAT and SOD activities compared to other light sources in P. haitanensis (Wu 2016). In comparison to white, green, and red LEDs, staining and total activities of POD decreased under blue LED treatment (Fig. 3C,G). APXa was upregulated only in green LED, whereas APxb was upregulated in blue, green, and red LEDs compared to white LED (Fig. 4D,E). It is assumed that both transcriptional upregulation of APX and the direct effect of specific light wavelengths on APX enzyme can be the basis for the marked increase in APX activity under green and red LEDs (Fig. 3D,H). Among different LED lights, blue LED light caused low demand for antioxidant enzymes, i.e., the low activities of POD and APX, which is likely due to sufficient energy dissipation through zeaxanthin and NPQ. By contrast, the lack of dissipative processes in green and red LED treatments may necessitate increased demand for SOD and APX.

Photoreceptor-mediated light signaling plays a pivotal role in photomorphogenesis (Neff and Chory 1998, Wang and Deng 2003, Zhang et al. 2018), but its role in photoprotection especially against excess excitation energy needs further clarification. Rice seedlings grown under green or red LED significantly upregulated PHYA, PHYB, and PHYC compared to broad-spectrum white LED (Fig. 5A–C). Exposure of etiolated Avena seedlings to red-light pulse also showed a higher phytochrome transcript level than continuous white light (Colbert et al. 1985). The green light-mediated upregulations of the three PHYs (Fig. 5A–C) support the idea that phytochromes can absorb some green light (Folta and Maruhnich 2007). The high expression levels of the three PHY genes were positively correlated with the increased activities of SOD, CAT, and APX (Figs. 3, 5). Moreover, activities of SOD, CAT, POD, and APX as well as expressions of CATa, CATb, and APxb showed positive correlations to PHYA, PHYB, and PHYC (Table 3). These results suggest a possible role of phytochromes in light quality-dependent regulation of antioxidant enzymes. Sharma et al. (1976) has already observed in maize that the transformation of Pr into Pfr by absorption of red light increased POD activity. In phyA, phyB, phyAB, and cry1phyAB1 mutants, deficiencies in phytochromes and cryptochrome 1 affected the levels of antioxidant enzymes in response to UV stress or high light (Rusaczonek et al. 2015, Kreslaski et al. 2020). The expression patterns of PHYs in blue LED treatment were similar to those of white LED treatment (Fig. 5A–C) indicated that besides red and far-red light, phytochromes have been shown to act as blue-light photoreceptors (Mohr 1994, Neff and Chory 1998).

To assess the involvement of cryptochromes in differential photoprotection under different wavelengths of LEDs, we examined expression profiles of CRYs. In comparison to the white LED, the expression level of CRY1 significantly increased in seedlings grown under blue LED, which was followed by a higher content of anthocyanins (Table 1, Fig. 5D). CRY1 protein contents also increased severalfold on irradiation of Brassica napus seedlings with blue light (Chatterjee et al. 2006). On the other hand, expression levels of CRY2 were upregulated in green and red LED treatments, compared to white LED (Fig. 5E). The upregulation of CRY2 in red LED treatment could be explained by the fact that CRY is a functionally important blue and red light-activated flavoprotein (Beel et al. 2012). Besides, cryptochromes are known as green-light receptors (Folta and Maruhnich 2007), and the green-absorbing state of CRY1 and CRY2 reverses blue light-induced responses (Banerjee et al. 2007, Bouly et al. 2007) such as the degradation of CRY2 (Li et al. 2011). Photolabile CRY2 is rapidly degraded in blue light, while CRY1 is a relatively stable photoreceptor (Ahmad et al. 1998). Among different LED treatments, the highest expression level of CRY2 in green LED treatment (Fig. 5E) may be related to green light-induced prevention of CRY2 degradation. Also, CRY1 and CRY2 exhibited negative and positive correlations, respectively, to the activities and expression of some antioxidant enzymes (Table 3). However, the correlations were weaker than those of PHYs. Kreslaski et al (2020) suggests that phytochromes together with CRY1 control the expression of APX genes under high-intensity light.

Conclusions: Our results demonstrate that specific wavelengths of different LEDs greatly influence photosynthetic performance, photoprotection, and antioxidant properties in rice seedlings. Seedlings grown under blue LED ensured sufficient levels of photoprotection by not only high NPQ,
but also great increases in zeaxanthin, expressions of \textit{BCH} and \textit{VDE}, and anthocyanin. Also, blue light-induced upregulation of \textit{CRY1} indicates that \textit{CRY1} is involved in energy dissipation mechanisms, thereby optimizing photosynthetic efficiency and photoprotection of PSII. By contrast, in seedlings grown under green and red LEDs, low levels of NPQ, zeaxanthin, and anthocyanin may necessitate increased activities of SOD and APX. The activities and expressions of antioxidant enzymes showed positive correlations to the expression of \textit{PHY} genes, suggesting a possible role of phytochromes in the regulation of antioxidant enzymes. Our study provides more insight into light quality-dependent regulation of photoprotection and antioxidation mechanisms and their relationship with the expressions of phytochrome and cryptochrome genes.

References


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