

Blue light specific and differential expression of a plastid σ factor, Sig5 in *Arabidopsis thaliana*

Yuichi Tsunoyama^a, Kazuya Morikawa^b, Takashi Shiina^c, Yoshinori Toyoshima^{b,*}

^aRadioisotope Research Center, Kyoto University, Kitashirakawa-oiwake-cho, Sakyo-ku, Kyoto 606-8502, Japan

^bGraduate School of Human and Environmental Studies, Kyoto University, Yoshida-nihonmatsu-cho, Sakyo-ku, Kyoto 606-8501, Japan

^cFaculty of Human and Environment, Kyoto Prefectural University, Shimogamo, Sakyo-ku, Kyoto 606-8522, Japan

Received 21 December 2001; revised 19 February 2002; accepted 22 February 2002

First published online 11 March 2002

Edited by Ulf-Ingo Flügge

Abstract The transcription of plastid gene *psbD* is under the control of the BLRP (blue-light-responsive promoter) recognized by plastid-encoded RNA polymerase, in which nuclear-encoded σ factors play a crucial role in the promoter recognition. We examined the effects of light on mRNA levels of six different *SIG* genes in *Arabidopsis* and found that blue light extensively induced the accumulation of *SIG5* transcripts, but red light did not. The blue light specificity was not observed in the accumulations of remaining five *SIG* genes. The blue light dependency of the *SIG5* expression well explains the light-dependent behavior of the *psbD* BLRP. © 2002 Federation of European Biochemical Societies. Published by Elsevier Science B.V. All rights reserved.

Key words: RNA polymerase; Blue light; σ factor; Sig5; *psbD*

1. Introduction

Expression of chloroplast *psbD* gene encoding the D2 subunit of the photosystem II reaction center is transcriptionally controlled by light. High-fluence blue light, but not red light, differentially activates *psbD* transcription from a blue light-responsive promoter (*psbD* BLRP) that is conserved among higher plants [1–3]. Even when the total plastid transcription declines as a consequence of chloroplast maturation, the *psbD* BLRP maintains high activity as to assist with maintaining the synthesis of D2, which is photo-damaged and undergoes rapid turnover in high light condition [1,4].

The *psbD* BLRP is recognized by the bacterial-type plastid-encoded RNA polymerase (PEP), but it is an unusual σ^{70} -type promoter that requires the ‘–10’ element for transcription, but not the ‘–35’ element. Instead, two well-conserved sequence elements, termed PGT box and AAG box, are located upstream of the ‘–35’ position [5–8]. In general, bacterial RNA polymerases are regulated by σ factors mediating the promoter recognition and by DNA-binding proteins functioning as activator or repressor. Since the plastome of higher plants encodes neither gene for σ factors nor DNA-binding protein, it is assumed that the blue light-dependent activation of the *psbD* BLRP is mediated by nuclear-encoded regulatory factors.

There are several proteins interacting with the AAG and PGT boxes of the *psbD* BLRP. However, it is reported that the PGT box is not necessary to activate the *psbD* transcription in vivo [9]. While, the AAG-box binding proteins (AGF) are constitutively expressed independent of light in matured chloroplasts [6,7]. Recently, PTF-1 that specifically binds to the ACC repeat in the AAG box was cloned in *Arabidopsis* [10]. Inactivation of PTF-1 by T-DNA insertion partially reduced the *psbD* BLRP activity, but did not eliminate the light-dependent transcription. These evidences suggest that AGF and PTF-1 act as positive regulators that generally enhance the *psbD* BLRP activity rather than the blue light-responsive factors that switch on the *psbD* BLRP activity.

The subunits of the core enzyme of PEP are encoded in plastid genome [11], while plastid σ factors are encoded in nuclear genome. In *Arabidopsis thaliana*, there are six putative plastid σ genes, which are homologous to bacterial group 1 and 2 σ factors [12–14]. We previously showed that in wheat chloroplasts, PEP changes its promoter preference depending on the light condition and the developmental stage of leaves possibly through the differential usage of the heterogeneous σ factors [6,15]. These findings might indicate that the blue light-induced differential expression of a particular σ factor is involved in the mechanism of the blue light-dependent activation of the *psbD* BLRP.

In this work, we examined the effect of light on the accumulations of the transcripts from the six *SIG* genes in *A. thaliana* and compared them with the blue light-dependent behavior of the *psbD* BLRP. The results suggest that Sig5 is differentially expressed by blue light as to recognize the *psbD* BLRP and initiate the transcription from the promoter with the help of blue light-non-specific high-fluence irradiation.

2. Materials and methods

2.1. Plant materials, growth conditions and light treatment

A. thaliana, ecotype Columbia, was grown on vermiculites for 4 weeks at 22°C under continuous white light (10–20 $\mu\text{mol m}^{-2} \text{s}^{-1}$). Plants employed for Northern blot analysis were exposed to white or blue and/or red light after dark-adaptation for 16 h. LED panels (LED-B or LED-R, Eyela) were used as light sources for blue (470 nm, FWHM (full width at half maximum) 30 nm) or red (660 nm, FWHM 20 nm) light and their photon-fluence rates were measured using a quantum photometer (LI-250, Li-Cor). White light was provided by a fluorescence lamp (FPL27AX, Mitsubishi).

2.2. Northern blot analysis

Total RNAs were extracted from rosette leaves of *Arabidopsis* by RNeasy Plant Mini kit (Qiagen). cDNA fragments (1030–1509 of

*Corresponding author. Fax: (81)-75-753 7540.

E-mail address: ytoyoshi@ip.media.kyoto-u.ac.jp (Y. Toyoshima).

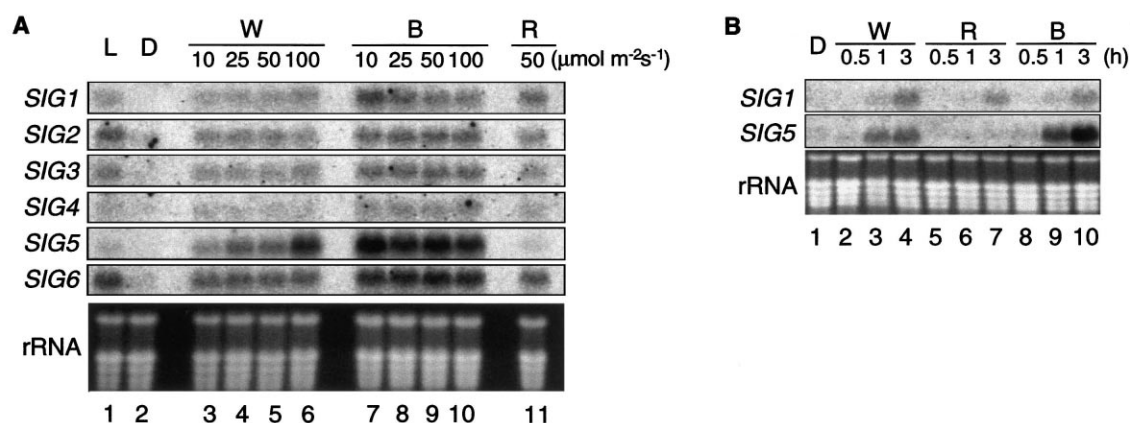


Fig. 1. Light-dependent accumulation of *SIG* transcripts in *Arabidopsis*. A: Effects of light quality and intensity on the expression of *SIG* genes. Light-grown plants (L) were dark-adapted for 16 h (D) and re-illuminated with white (W), blue (B) or red (R) light for 3 h at indicated fluences ($\mu\text{mol m}^{-2} \text{s}^{-1}$). Total cellular RNA (10 μg) was subjected to RNA gel-blot hybridization with six *SIG* probes. The ethidium bromide-stained rRNAs serve as gel-loading references. B: Time course of the induction of *SIG1* and *SIG5* transcripts by light. Dark-adapted plants (D) were re-illuminated with white (15 $\mu\text{mol m}^{-2} \text{s}^{-1}$), red (50 $\mu\text{mol m}^{-2} \text{s}^{-1}$) or blue (50 $\mu\text{mol m}^{-2} \text{s}^{-1}$) light for 0.5, 1 and 3 h. The amount of total cellular RNA per lane was 7 μg .

SIG1 (AB019942), 1048–1719 of *SIG2* (AB019943), 1049–1716 of *SIG3* (AB019944), 739–1260 of *SIG4* (AB021119), 1096–1554 of *SIG5* (AB021120), 1158–1644 of *SIG6* (AB029916)) were amplified by PCR and used as probes for Northern analyses. The CDR probe (+78 to +1002 of the *psbD* translational start codon) was designed to detect the transcripts produced from all of multiple promoters in the *psbD/C* operon, and the UTR probe (–1085 to –276) to detect specifically the transcripts from the *psbD* BLRP. DNA fragments were labeled with [α - ^{32}P]dCTP using RTG DNA labeling beads (Amersham Pharmacia). 10 μg of total RNAs were separated on 1.0% agarose-formaldehyde denaturing gels, transferred onto a Hybond N+ membrane (Amersham Pharmacia), and hybridized at 60°C for 18 h with the labeled DNA probes. Final wash conditions were 0.1 \times SSC, 0.1% SDS at 60°C for 30 min.

3. Results

3.1. The effect of light on the transcript accumulation of six *SIG* genes in *Arabidopsis*

Light-dependent accumulation of the transcripts of *SIG* genes was reported in various plants including *Arabidopsis* [12–14,16–21], but the effects of the light quality and strength on the expression of each *SIG* gene remain unknown. Here we examined the effect of white, blue (470 nm) or red (660 nm) light on the accumulation of the transcripts of *SIG1* through *SIG6* genes in *Arabidopsis* by Northern blot analyses. As shown in Fig. 1A, the transcripts of all *SIG* genes were accumulated in rosette leaves of the plants grown under continuous white light for 4 weeks (Fig. 1A, L), although extents of their accumulation were dependent on the genes as *SIG6* > *SIG2* > *SIG3* > *SIG1* > *SIG5* = *SIG4*. The transcripts of all genes observed in the light-grown leaves virtually disappeared after dark-adaptation for 16 h (Fig. 1A, D), and recovered almost to the original levels by exposure to white light of 10 $\mu\text{mol m}^{-2} \text{s}^{-1}$ (Fig. 1A, W) within 3 h except for the *SIG5* transcript. In the case of *SIG5*, accumulation of the transcript significantly exceeded the original level. Increase of the white light fluence to 100 $\mu\text{mol m}^{-2} \text{s}^{-1}$ enhanced the accumulation of the *SIG5* transcript but scarcely affected to other genes. A dramatic light quality effect was observed in the accumulation of *SIG5* transcript. Blue light of 10–100 $\mu\text{mol m}^{-2} \text{s}^{-1}$ extensively induced the *SIG5* transcript, but red light of 50 $\mu\text{mol m}^{-2} \text{s}^{-1}$ never induced the *SIG5* tran-

scripts. The *SIG6* transcripts were induced by both of blue and red lights, although the effect of blue light slightly exceeded that of red light. In the cases of the other *SIG* genes, blue and red lights of 50 $\mu\text{mol m}^{-2} \text{s}^{-1}$ showed almost equal effects on the induction of their transcripts. Change of the blue light fluence from 10 to 100 $\mu\text{mol m}^{-2} \text{s}^{-1}$ did not make a significant difference in the induction of transcripts from all *SIG* genes. Fig. 1B compares the time courses of the accumulation of the *SIG1* and *SIG5* transcripts induced by white, blue and red lights. The transcript of *SIG1* was observed after 3 h illumination but not 1 h irrespective of light quality, while the transcript of *SIG5* appeared after 1 h illumination with white or blue light.

3.2. Comparison of the effect of blue light on the accumulation of the *SIG5* transcript and the *psbD* BLRP activity

It would be interesting to compare the effect of blue light on the induction of the *SIG5* transcript and that on the *psbD* BLRP activity. As shown in Fig. 2, the *psbD* CDR probe (see Section 2) detected four bands (4.5, 3.7, 2.8 and 2.6 kb) in the leaves illuminated with blue light of 100 $\mu\text{mol m}^{-2} \text{s}^{-1}$ (lane 12). Among them, the 4.5 and 3.7 kb transcripts were

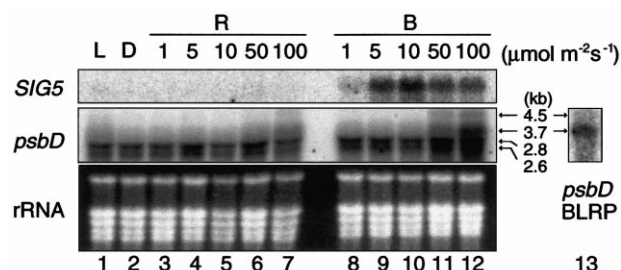


Fig. 2. Comparison of the light-dependent accumulation of *SIG5* and *psbD* transcripts. Light-grown plants (L) were dark-adapted for 16 h (D) and re-illuminated with red (R) or blue light (B) for 3 h at indicated fluences ($\mu\text{mol m}^{-2} \text{s}^{-1}$). Total cellular RNA (7 μg) was subjected to RNA gel-blot hybridization with the *SIG5* and *psbD* CDR probes (lanes 1–12). The *psbD* UTR probe was used to detect transcripts from the *psbD* BLRP in leaves illuminated 100 $\mu\text{mol m}^{-2} \text{s}^{-1}$ of blue light for 3 h after dark-adaptation (lane 13). The ethidium bromide-stained rRNAs serve as gel-loading references.

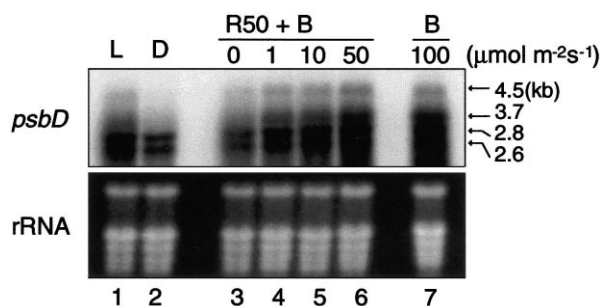


Fig. 3. The effect of blue light intensity on the *psbD* BLRP activity under the simultaneous illumination with an intensive red light ($50 \mu\text{mol m}^{-2} \text{s}^{-1}$). Light-grown plants (L) were dark-adapted for 16 h (D) and simultaneously re-illuminated with $50 \mu\text{mol m}^{-2} \text{s}^{-1}$ of red and blue light at indicated fluences ($\mu\text{mol m}^{-2} \text{s}^{-1}$) for 3 h following dark-adaptation. Total cellular RNA (7 μg) was subjected to RNA gel-blot hybridization with the *psbD* CDR probe. The ethidium bromide-stained rRNAs serve as gel loading references.

detected by the *psbD* UTR probe (lane 13), which was designed to detect specifically the transcripts from the *psbD* BLRP based on the 5'-ends mapping analysis of *psbD* transcripts in *Arabidopsis* [3], indicating that these two transcripts were originated from the *psbD* BLRP. These BLRP transcripts were induced by high-fluence of blue light (lanes 8–12), but not red light (lanes 3–7) irradiation. These results were in agreement with the previous findings [3,22]. These two transcripts strongly induced by $100 \mu\text{mol m}^{-2} \text{s}^{-1}$ blue light (lane 12) were reduced to less than one fifth when the blue light intensity was at $50 \mu\text{mol m}^{-2} \text{s}^{-1}$ (lane 11), and were undetectable at less than $10 \mu\text{mol m}^{-2} \text{s}^{-1}$ (lane 10). Contrary to the case of the *psbD* BLRP transcription, the *SIG5* transcript was detected at a low intensity such as $1 \mu\text{mol m}^{-2} \text{s}^{-1}$ of blue light (lane 8) and the transcript level increased with intensifying the light up to $5 \mu\text{mol m}^{-2} \text{s}^{-1}$ and saturated (lanes 9–12). This result seems to indicate that there is a great difference in the intensity requirement of blue light between the activation of the *psbD* BLRP and the expression of *SIG5*. As shown in Fig. 3, however, a weak blue light such as $1 \mu\text{mol m}^{-2} \text{s}^{-1}$ that induced the *SIG5* transcript was able to induce the accumulation of the 4.5 and 3.7 kb transcripts from the *psbD* BLRP to the original level (lane 1), when the leaves were simultaneously illuminated with $50 \mu\text{mol m}^{-2} \text{s}^{-1}$ red light. Simultaneous illumination with $50 \mu\text{mol m}^{-2} \text{s}^{-1}$ red light and $50 \mu\text{mol m}^{-2} \text{s}^{-1}$ blue light (lane 6) was able to enhance almost equivalently the accumulation of these two transcripts from the *psbD* BLRP to the illumination with $100 \mu\text{mol m}^{-2} \text{s}^{-1}$ blue light (lane 7). These evidences indicate that red light equally satisfies the requirement of high light intensity to the enhancement of the *psbD* BLRP activity, as long as a weak blue light is present.

4. Discussion

Phylogenetic analysis revealed that higher plant σ factors so far known were similar to the bacterial principal σ factors (group 1 or 2 σ factors) and formed a monophyletic group [14]. Among plant σ factors, *Arabidopsis* Sig5 appears isolated from other σ factors in a phylogenetic tree constructed by analyzing different plant σ factors [14] and intron sites of *SIG5* are distinct among *Arabidopsis* *SIG* genes, possibly indicating an unique function of Sig5 in plastid transcription. In this

work, we found that all of *Arabidopsis* σ factors are expressed light-dependently in matured leaves, but the expression of *SIG5* is significantly different from the others in the selectivity for blue light and the rapid response to the light. These evidences may suggest Sig5 to be a special σ factor, which contributes to a blue light-dependent and gene-specific transcription in plastids.

The accumulation of 4.5 and 3.7 kb *psbD* transcripts derived from the *psbD* BLRP are notably induced in *Arabidopsis* matured leaves by the illumination with high-fluence blue light ($100 \mu\text{mol m}^{-2} \text{s}^{-1}$) but not with monochromatic red light, in agreement with the previous work [1]. Inhibitor experiments had suggested that blue light regulated the expression of nuclear factors that comprise the pathway activating the *psbD* BLRP [23–25], but the key factor induced by blue light to activate the *psbD* BLRP has not been identified yet. The blue light-specific and differential response of the *SIG5* transcription seems to indicate that *SIG5* is the nuclear-encoded key factor, which mediates blue light signal from cytoplasm/nuclei to chloroplasts and functions as a molecular switch to activate the *psbD* BLRP.

The strength of monochromatic blue light required for activating the *psbD* BLRP was higher than that for inducing the transcription of *SIG5* by one order of magnitude. It has been a common understanding that the *psbD* BLRP requires high-fluence of blue or UV-A light to be activated, although the observation indicating that far-red light given together with high-fluence of blue light enhanced the *psbD* transcription in a synergistic manner in barley was reported [26]. In this work, however, we revealed that the requirement of high light intensity for the activation of the *psbD* BLRP was not confined to blue light, but red light was almost equally effective under simultaneous illumination with low-fluence of blue light such as $1 \mu\text{mol m}^{-2} \text{s}^{-1}$, which can induce the *SIG5* transcription. These evidences strongly suggest that the effect of blue light and that of high light intensity on the activation of the *psbD* BLRP originate from the different molecular events mediated by at least two photo-sensory pathways. Comparing the blue light-dependent behaviors of the accumulation of the *SIG5* transcript and that of the transcripts from the *psbD* BLRP, it seems quite probable that blue light-dependent expression of Sig5 is essential for the *psbD* BLRP to be recognized by PEP. However, the accommodation of Sig5 to PEP is not enough to initiate the transcription from the *psbD* BLRP effectively. Other unknown events induced by the high-fluence light are also required. In vitro transcription analyses in wheat indicated that the deletion of two enhancer regions of the *psbD* BLRP resulted in a severe reduction of the transcription activity, although the light-responsive transcription was still observed [6,23]. High-fluence light might induce the full activity of PEP on the *psbD* BLRP transcription via the modifications of enhancer proteins and/or PEP. Thus, the blue light-induction of *SIG5* expression observed in this work well explains the light-responsive activation of the *psbD* BLRP, although the direct evidence has not been obtained. It will be interesting to examine whether over-expression of Sig5 results in the continuous activation of the *psbD* BLRP.

Blue light responses are mediated by at least two different types of photoreceptors, cryptochromes (cry1 and cry2) and phototropins (nph1 and npl1). Since the activity of the *psbD* BLRP was repressed in a *CRY1/CRY2* double mutant [27], it is likely that both cry1 and cry2 are involved in the blue light-

specific pathway, regulating the *psbD* BLRP activity possibly through the activation of *SIG5* gene. On the other hand, the BLRP activity was also repressed in phytochrome (*phyA*) mutants [27]. PhyA signaling pathway might be involved in the high-fluence light requirement of the *psbD* BLRP activation. Alternatively, the high-fluence response of the *psbD* BLRP might be related to the highly reduced redox state in plastids generated under the high light intensity.

Acknowledgements: We thank Y. Isozumi for providing us the facilities of the Radioisotope Research Center. This work was supported by Grants-in-Aid for Scientific Research in Priority Areas (number 12025216) and Grants-in-Aid for Scientific Research from the Ministry of Education, Culture, Sports, Science and Technology of Japan (Monbukagakusho).

References

- [1] Christopher, D.A. and Mullet, J.E. (1994) *Plant Physiol.* 104, 1119–1129.
- [2] Christopher, D.A., Kim, M. and Mullet, J.E. (1992) *Plant Cell* 4, 785–798.
- [3] Hoffer, P.H. and Christopher, D.A. (1997) *Plant Physiol.* 115, 213–222.
- [4] Stern, D.B., Higgs, D.C. and Yang, J. (1997) *Trends Plant Sci.* 2, 308–315.
- [5] Allison, L.A. and Maliga, P. (1995) *EMBO J.* 14, 3721–3730.
- [6] Nakahira, Y., Baba, K., Yoneda, A., Shiina, T. and Toyoshima, Y. (1998) *Plant Physiol.* 118, 1077–1088.
- [7] Kim, M. and Mullet, J.E. (1995) *Plant Cell* 7, 1445–1457.
- [8] Kim, M., Thum, K.E., Morishige, D.T. and Mullet, J.E. (1999) *J. Biol. Chem.* 274, 4684–4692.
- [9] Thum, K.E., Kim, M., Morishige, D.T., Eibl, C., Koop, H.U. and Mullet, J.E. (2001) *Plant Mol. Biol.* 47, 353–366.
- [10] Baba, K., Nakano, T., Yamagishi, K. and Yoshida, M. (2001) *Plant Physiol.* 125, 595–603.
- [11] Hu, J. and Bogorad, L. (1990) *Proc. Natl. Acad. Sci. USA* 87, 1531–1535.
- [12] Tanaka, K., Tozawa, Y., Mochizuki, M., Shinozaki, K., Nagatani, A., Wakasa, K. and Takahashi, H. (1997) *FEBS Lett.* 413, 309–313.
- [13] Isono, K., Shimizu, M., Yoshimoto, K., Niwa, Y., Satoh, K., Yokota, A. and Kobayashi, H. (1997) *Proc. Natl. Acad. Sci. USA* 94, 14948–14953.
- [14] Fujiwara, M., Nagashima, A., Kanamaru, K., Tanaka, K. and Takahashi, H. (2000) *FEBS Lett.* 481, 47–52.
- [15] Satoh, J., Baba, K., Nakahira, Y., Tsunoyama, Y., Shiina, T. and Toyoshima, Y. (1999) *Plant J.* 18, 407–415.
- [16] Tozawa, Y., Tanaka, K., Takahashi, H. and Wakasa, K. (1998) *Nucleic Acids Res.* 26, 415–419.
- [17] Kestermann, M., Neukirchen, S., Kloppstech, K. and Link, G. (1998) *Nucleic Acids Res.* 26, 2747–2753.
- [18] Morikawa, K., Itoh, S., Tsunoyama, Y., Nakahira, Y., Shiina, T. and Toyoshima, Y. (1999) *FEBS Lett.* 451, 275–278.
- [19] Tan, S. and Troxler, R.F. (1999) *Proc. Natl. Acad. Sci. USA* 96, 5316–5321.
- [20] Lahiri, S., Yao, J., McCumbers, C. and Allison, L.A. (1999) *Mol. Cell Biol. Res. Commun.* 1, 14–20.
- [21] Oikawa, K., Fujiwara, M., Nakazato, E., Tanaka, K. and Takahashi, H. (2000) *Gene* 261, 221–228.
- [22] Yao, W.B., Meng, B.Y., Tanaka, M. and Sugiura, M. (1989) *Nucleic Acids Res.* 17, 9583–9591.
- [23] Satoh, J., Baba, K., Nakahira, Y., Shiina, T. and Toyoshima, Y. (1997) *Plant Mol. Biol.* 33, 267–278.
- [24] Gamble, P.E. and Mullet, J.E. (1989) *EMBO J.* 8, 2785–2794.
- [25] Christopher, D.A., Li, X., Kim, M. and Mullet, J.E. (1997) *Plant Physiol.* 113, 1271–1282.
- [26] Christopher, D.A. (1996) *Photosynth. Res.* 47, 239–251.
- [27] Thum, K.E., Kim, M., Christopher, D.A. and Mullet, J.E. (2001) *Plant Cell* 13, 2747–2760.