## RAPID COMMUNICATION

# **Rice** *PLASTOCHRON* genes regulate leaf maturation downstream of the gibberellin signal transduction pathway

Manaki Mimura · Yasuo Nagato · Jun-Ichi Itoh

Received: 29 February 2012/Accepted: 22 March 2012/Published online: 5 April 2012 © Springer-Verlag 2012

Abstract Rice PLASTOCHRON 1 (PLA1) and PLA2 genes regulate leaf maturation and plastochron, and their loss-of-function mutants exhibit small organs and rapid leaf emergence. They encode a cytochrome P450 protein CYP78A11 and an RNA-binding protein, respectively. Their homologs in Arabidopsis and maize are also associated with plant development/organ size. Despite the importance of PLA genes in plant development, their molecular functions remain unknown. Here, we investigated how PLA1 and PLA2 genes are related to phytohormones. We found that gibberellin (GA) is the major phytohormone that promotes *PLA1* and *PLA2* expression. GA induced PLA1 and PLA2 expression, and conversely the GA-inhibitor uniconazole suppressed PLA1 and PLA2 expression. In pla1-4 and pla2-1 seedlings, expression levels of GA biosynthesis genes and the signal transduction gene were similar to those in wild-type seedlings. GA treatment slightly down-regulated the GA biosynthesis gene GA20ox2 and up-regulated the GA-catabolizing gene GA2ox4, whereas the GA biosynthesis inhibitor uniconazole up-regulated GA20ox2 and down-regulated GA2ox4 both in wild-type and *pla* mutants, suggesting that the GA feedback mechanism is not impaired in *pla1* and *pla2*. To reveal how GA signal transduction affects the expression of PLA1 and PLA2, PLA expression in GA-signaling mutants was examined. In GA-insensitive mutant, gid1 and lesssensitive mutant, Slr1-d1, PLA1 and PLA2 expression was

**Electronic supplementary material** The online version of this article (doi:10.1007/s00425-012-1639-5) contains supplementary material, which is available to authorized users.

M. Mimura · Y. Nagato · J.-I. Itoh (⊠) Graduate School of Agricultural and Life Sciences, University of Tokyo, Tokyo 113-8657, Japan e-mail: ajunito@mail.ecc.u-tokyo.ac.jp down-regulated. On the other hand, the expression levels of *PLA1* and *PLA2* were highly enhanced in a GA-constitutive-active mutant, *slr1-1*, causing ectopic overexpression. These results indicate that both *PLA1* and *PLA2* act downstream of the GA signal transduction pathway to regulate leaf development.

**Keywords** Rice · *PLASTOCHRON 1* · *PLASTOCHRON 2* · Gibberellin · Leaf development

## Abbreviations

cid

# Introduction

Leaves are the main photosynthetic organ in plants. In addition, the number and arrangement of leaves greatly contributes to the establishment of plant shape. To understand the genetic mechanism underlying shoot formation, two aspects of leaf primordial formation must be considered: spatial (phyllotaxy) and temporal (plastochron) regulation. Given that the regular phyllotactic pattern has long fascinated plant scientists, a large number of studies have been conducted to investigate this phenomenon (for review, Steeves and Sussex 1989). However, rapid progress has been achieved only recently. In maize, the causal gene of the *abphyl1* mutant, which exhibits decussate phyllotaxy instead of 1/2 alternate, was isolated in 2004 (Giulini et al. 2004). The recent auxin-based model has been widely

accepted (Reinhardt et al. 2000, 2003; Jönsson et al. 2006; Smith et al. 2006).

In contrast, the molecular basis of plastochron regulation remains to be uncovered. *plastochron 1 (pla1)* is the first mutant that drastically alters plastochron, in which leaf primordia are formed approximately twofold faster than in the wild type (Itoh et al. 1998). Concomitantly, leaves of pla1 become short, suggesting that PLA1 regulates organ size. The PLA1 gene encodes a cytochrome P450 family protein (CYP78A11) (Miyoshi et al. 2004), but its substrate is unknown. An Arabidopsis homolog of PLA1, KLUH, was shown to regulate organ size (Anastasiou et al. 2007). Subsequently, the PLA2 and PLA3 genes, loss-of-function mutants of which exhibit similar phenotypes to that of pla1, were identified (Kawakatsu et al. 2006, 2009). These encode an RNA-binding protein and glutamate carboxypeptidase, respectively. Interestingly, PLA1 and PLA2 are expressed in young leaf primordia, but not in shoot meristems (Miyoshi et al. 2004; Kawakatsu et al. 2006). Therefore, based on analyses of the developmental processes of leaves, the primary functions of PLA1 and PLA2 are suppression of precocious leaf maturation, and that some non-cell autonomous signals move from leaf primordia through shoot meristems to suppress the formation of a new leaf primordium (Kawakatsu et al. 2006). Thus, PLA1 and PLA2 are key genes for elucidating leaf development. However, the regulation of PLA1 and PLA2 expression remains unknown.

*pla1* and *pla2* show several phenotypes likely related to phytohormones, such as small leaf size, dwarfism and enlarged SAM. In addition, the phytohormone (CK, abscisic acid and IAA) contents of *pla* mutants differed from those of the wild type (Kawakatsu et al. 2009). These mutant phenotypes suggest that *PLA* genes have some relationship with phytohormones. However, the position of *PLA* genes in the phytohormone-related pathway remains unclear.

Several phytohormones are involved in the regulation of leaf development/growth. Auxin has pleiotropic functions on plant development, including leaf growth (for review, Teale et al. 2006). Rice tryptophan deficient dwarf 1 (tdd1) mutant exhibits low auxin content and dwarfism with small leaves (Sazuka et al. 2009). A gain-of-function mutant of rice, OsIAA3, which inhibits auxin signaling, shows an auxin-insensitive phenotype, and produces shorter leaves than the wild type, resulting in dwarfism (Nakamura et al. 2006a). Thus, auxin biosynthesis and signaling is important for normal leaf development and morphological processes in rice. Cytokinine (CK) is also profoundly associated with leaf development. For example, ABPYL1, which encodes the A-type response regulator, regulates phyllotaxy in maize (Jackson and Hake 1999; Giulini et al. 2004). Overexpression of a type-A response regulator caused dwarfism in rice (Hirose et al. 2007). Another phytohormone, brassinosteroid (BR), has a role in regulating plant growth. Loss-of-function mutants of BR biosynthetic and signaling genes frequently exhibit dwarfism and a reduced organ size. Rice D2 and D11 are BR biosynthesis genes, and regulate grain size and other traits (Hong et al. 2003; Tanabe et al. 2005). A BR-insensitive mutant, d61, also exhibits dwarfism, and a severe d61 allele, d61-4, exhibited rolled and twisted leaves (Yamamuro et al. 2000; Nakamura et al. 2006b).

Gibberellin (GA) is the most well-known phytohormone that affects plant height and organ (leaf) size. Semi-dwarf GA mutants were used in the wheat and rice green revolution (Hedden 2003). In rice, many dwarf mutants are associated with GA biosynthesis or signaling. For example, SEMIDWARF 1 (SD1), which encodes GA20 oxidase, was utilized in the rice green revolution (Ashikari et al. 2002; Sasaki et al. 2002). The d18 dwarf mutant of GA3ox2 has an extremely dwarf stature with small leaves (Itoh et al. 2001), and a prolonged juvenile phase (Tanaka et al. 2011). Other GA-deficient and GA-insensitive mutants commonly exhibit small leaves and a dwarf stature. In contrast, GA promotes the juvenile-adult phase change (Evans and Poehtig 1995; Teifer et al. 1997; Schwartz et al. 2008). In wild-type rice, plastochron is short in the juvenile compared with the adult phase (Itoh et al. 2005). Thus, GA seems to be associated with PLA functions, whose mutants show a short plastochron.

We examined the sensitivities of pla1 and pla2 to several phytohormones, and revealed that GA is the major influencer of *PLA* function. Using GA-related genes and mutants thereof, we determined that *PLA1* and *PLA2* function downstream of GA signal transduction.

## Materials and methods

## Plant materials

We used *pla1-4* and *pla2-1* mutants, which show the most severe phenotypes among their alleles, and share a cv Taichung 65 genetic background (Kawakatsu et al. 2006). We also used a GA-biosynthetic dwarf mutant, *d18-h*, which encodes GA<sub>3</sub> oxidase2 and has a low auxin content (Itoh et al. 2001). Three GA signaling mutants, *gibberellin insensitive dwarf 1 (gid1), slender rice 1-1 (slr1-1)* and *Slr1-d1* were used (Ikeda et al. 2001; Asano et al. 2009). *GID1* encodes a GA receptor, thus its loss-of-function mutant is GA insensitive (Ueguchi-Tanaka et al. 2005). *SLR1* encodes the DELLA protein and plays an important role in GA signal transduction (Ikeda et al. 2001; Itoh et al. 2002; Gomi et al. 2004). *slr1-1* forms a highly-elongated plant due to constitutive activation of GA signaling, while

*Slr1-d1* shows dwarf phenotype and is a dominant allele of *SLR1* (Ikeda et al. 2001; Asano et al. 2009).

## Application of phytohormones

Wild-type and mutant seeds were sterilized with 1 % NaClO for 40 min, and washed four times in sterile distilled water. The seeds were then placed on the Murashige and Skoog (1962) medium containing various concentrations of 2,4-D, kinetin, GA<sub>3</sub> 24-epiBL or uniconazole. Plants were grown in a growth chamber under continuous light at 28 °C. After 10 or 14 days, plant height and second leaf sheath length were measured for more than five plants of each treatment.

## Clearing of leaf sheath and measurement of cell size

To measure cell size, leaf sheaths were fixed with FAA (formalin: acetic acid: 50 % ethanol, 1:1:18) for 24 h at 4 °C. They were then dehydrated in a graded ethanol series and cleared in chloral hydrate at 96 °C in a heat block. We measured epidermal cell sizes on the adaxial side of cleared leaf sheaths under a light microscope. Measurements were performed on at least 100 cells per sample. Significant differences were analyzed by Student's *t* test.

#### In situ hybridization

Ten-day-old shoot apices of wild-type plants treated with  $GA_3$  or uniconazole and of *slr1-1* plants were fixed with paraformaldehyde in 0.1 M sodium phosphate buffer, dehydrated through a series of butanol extractions, and embedded in paraplast plus. Microtome sections (8 mm thick) were applied to glass slides coated with APS (Matsunami Glasses, Japan). A digoxigenin-labeled antisense probe of *PLA1* was prepared as described previously (Miyoshi et al. 2004). Hybridization and immunological detection with alkaline phosphatase were preformed as described by Kouchi and Hata (1993).

#### Real-time PCR

Total RNA was extracted from shoot apices using TRIZOL reagent (Invitrogen). After RNase-free DNase I treatment, 1  $\mu$ g of RNA was used for RT-PCR using High-capacity cDNA Reverse Transcription Kits (Applied Biosystems, USA). To quantify *PLA1*, *PLA2*, *GA20ox2*, *GA3ox2*, *GA2ox4* and *SLR1* expression, real-time PCR was performed using SYBERGREEN (Applied Biosystems, USA) or the TaqMan Fast Universal PCR Master Mix, FAM-labeled TaqMan probes (Applied Biosystems, USA), and the StepOnePlus real-time PCR system (Applied Biosystems, USA). Each gene expression value is the average of

three independent real-time PCR assays. Expression levels were normalized to that of an internal control, *ACT1*. The primers and probes for each gene are listed in Supplementary Table S1.

## Results

Since the *pla1* and *pla2* mutants exhibit dwarf phenotypes and short plastochron (rapid leaf emergence), it is hypothesized that *pla* phenotypes are related to phyotohormones. Of the many *pla1* and *pla2* alleles, we used *pla1-4* and *pla2-1*, both of which are strong alleles with a common genetic background of cv. Taichung 65.

Gibberellin is the major phytohormone associated with *PLA1* and *PLA2* functions

First, we observed the responses of *pla1* and *pla2* seedlings when several phytohormones, which are known to affect plant growth and leaf development, were added to the culture media. Both pla1-4 and pla2-1 mutants showed similar responses to auxin (2,4-D) as did the wild type. That is, 2,4-D inhibited the growth of wild-type, pla1-4 and pla2-1 seedlings (Fig. 1a). CK (kinetin) application caused similar responses in wild-type and *pla* seedlings (Fig. 1b). Higher kinetin concentrations caused more severe growth inhibition in wild-type, *pla1-4* and *pla2-1* seedlings. BR is also known to affect plant height and leaf development. However, both *pla1-4* and *pla2-1* seedlings responded to external BR (24-epiBL) similarly to wild-type seedlings (Fig. 1c). Thus auxin, CK and BR are not related to PLA function. In contrast, GA3 application induced rapid growth of wild-type plants (Fig. 1d, e). In pla1-4 and pla2-1, however, growth induction was restricted (Fig. 1d, e). Accordingly, GA is the major phytohormone associated with PLA1 and PLA2 functions, while pla1-4 and pla2-1 seem to be less sensitive to GA than wild-type plants.

Responses of *pla1* and *pla2* to gibberellin application in cell size and leaf elongation

We examined responses to GA in cells and tissues of *pla1-4* and *pla2-1* plants. GA promotes cell elongation and tissue/organ elongation. We measured the length of more than 100 cells on the adaxial surface of the second leaf sheath. In wild-type plants, GA<sub>3</sub> application elongated leaf sheath cells by approximately 18 %, but by at most 6 % in *pla1-4*, and no elongation was observed in *pla2-1* plants (Fig. 2a, b). Leaf sheaths elongate in response to GA<sub>3</sub> application. In wild-type plants, elongation of the 2nd leaf sheath increased with GA<sub>3</sub> concentration, being *circa* threefold longer at  $10^{-5}$  than at 0 M, whereas *pla1-4* and

Fig. 1 Effect of several phytohormones on the growth of *pla1* and *pla2* seedlings. Wild-type, pla1-4 and pla2-1 seeds were inoculated on culture media containing phytohormones. Plants were grown for 14 days in 2,4-D (a) and kinetin (b) treatments, and for 10 days in 24-epiBL (BR) (c) and GA<sub>3</sub> (d) treatments. Data in **a**–**d** represent mean  $\pm$  SE. e Seedlings of wild type, pla1-4 and *pla2-1* grown for 10 days with or without GA<sub>3</sub> treatment. Bar 2 cm



pla2-1 showed a twofold or less elongation at the same GA<sub>3</sub> concentrations (Fig. 2c). In particular, a low GA<sub>3</sub> concentration did not promote pla1-4 and pla2-1 leaf elongation.

These results indicate that pla1-4 and pla2-1 are insensitive to GA in many traits, suggesting that *PLA1* and *PLA2* act downstream of GA signal transduction.

## Induction of PLA gene expression by gibberellin

The above results suggest that *PLA1* and *PLA2* gene expression is associated with GA signaling. Thus, we examined the effect of GA on *PLA1* and *PLA2* gene expression. Ten-day-old seedlings were treated with 10  $\mu$ M GA<sub>3</sub> and *PLA* gene expression was monitored for 24 h by real-time PCR. *PLA1* and *PLA2* expression increased as early as 3 h after treatment, and a high level of expression was maintained for 24 h (Fig. 3a). To investigate the long-term effect of GA, wild-type seeds were inoculated and

grown for 8 days on culture media containing 10  $\mu$ M GA<sub>3</sub>. *PLA1* and *PLA2* expression was maintained at a high level for 8 days (Fig. 3b). Next, we examined the effect of uniconazole, a GA biosynthesis inhibitor. Uniconazole treatment markedly suppressed *PLA1* and *PLA2* gene expression (Fig. 3b). Therefore, GA regulates the expressions of both *PLA1* and *PLA2*.

Since GA and uniconazole treatments up- and downregulated *PLA1* expression, respectively, the *PLA1* expression pattern was investigated using in situ hybridization. *PLA1* is expressed in the basal and abaxial regions of leaf primordia, but not in shoot meristems (Fig. 3c, Miyoshi et al. 2004). When treated with GA<sub>3</sub>, strong and ectopic *PLA1* expression was detected in the adaxial region of leaf primordia and in the tip of shoot meristems in addition to the normal expression domain (Fig. 3d). The expression pattern was not unaffected by uniconazole treatment, but hybridization signals were weakened (Fig. 3e).

Fig. 2 Response of *pla1* and pla2 to gibberellin application. a Adaxial surface of cleared 2nd leaf sheath of wild type, pla1-4 and *pla2-1* treated with GA<sub>3</sub>. Bars 200 µm. b Effects of GA<sub>3</sub> treatment on cell length in 2nd leaf sheath. In WT, GA<sub>3</sub> treatment significantly increased cell length. Double asterisks significantly longer at 1 % level than in -GA (*t* test). **c** Effects of GA<sub>3</sub> treatment on the length of 2nd leaf sheath. Double asterisks significant at 1 % level compared with 0 M (t test). Data in **b**, **c** represent mean  $\pm$  SE

Fig. 3 Induction of PLA1 and PLA2 expression by gibberellin. a, b Real-time PCR assays. a Short-term induction of PLA1 and PLA2 expression by 10 µM GA<sub>3</sub>. Each expression level is represented relative to that at 0 h. b Effects of GA<sub>3</sub> and uniconazol treatments for 8 days on PLA 1 and PLA2 expressions. Expression level is represented relative to that in the control. Data in **a**, **b** represent mean  $\pm$  SE. **c**–**e** In situ hybridization of *PLA1* in wild-type shoot apex (c), wildtype shoot apex treated with 10 µM GA<sub>3</sub> for eight days (d) and wild-type shoot apex treated with 1 µM uniconazole for 8 days (e). Arrows indicate ectopic expression. Bars 100 µm



Expression of GA-related genes in pla mutants

Given that GA regulates *PLA1* and *PLA2* expression, we next examined the expression of GA-related genes in *pla1* and *pla2* mutants. The expression of the GA biosynthesis genes, *GA20ox2* and *GA3ox2*, and the GA-catabolizing

gene, *GA20x4*, did not largely differ among wild-type, *pla1-4* and *pla2-1* plants, although *GA30x2* expression was somewhat increased in *pla1-4* and *pla2-1* compared with the wild type (Fig. 4a). Similarly, *SLR1* (a gene involved in GA signal transduction) expression was comparable in wild-type, *pla1-4* and *pla2-1* plants (Fig. 4a). These data



**Fig. 4** Expression of gibberellin-related genes in *pla1* and *pla2* seedlings. **a–c** Real-time PCR assays. **a** Expression of GA-biosynthetic (*GA20ox2* and *GA30x2*), catabolizing (*GA20x4*) and signaling gene (*SLR1*) genes in wild type, *pla1-4* and *pla2-1*. Expression level in *pla* mutants is represented relative to that in wild type. **b**, **c** Effect of GA<sub>3</sub> and uniconazole treatments on *GA200x2* (**b**) and *GA20x4* 

(c) expression in wild type, pla1-4 and pla2-1. Each expression level is represented relative to that in wild-type control. In **b** and **c**, expression level of *GA20ox2/GA2ox4* in the control (non-treatment) does not significantly differ among wild type, pla1-4 and pla2-1. Data in **a**-**c** represent mean  $\pm$  SE

suggest that *PLA1* and *PLA2* do not affect GA biosynthesis or signal transduction.

In plants, GA content is regulated by a feedback mechanism involving GA signal transduction; e.g., GA20ox2 expression is enhanced in GA-insensitive mutants such as gibberellin insensitive dwarf 1 (gid1), and downregulated in the GA-constitutive-active mutant slr1-1 (Yamaguchi 2008). Therefore, we investigated whether GA signal transduction is operating normally in *pla1* and *pla2* by determining the effect of GA and an inhibitor thereof on expression of the above genes. Application of GA slightly decreased the expression of GA20ox2 in wild-type, pla1-4 and pla2-1 plants (Fig. 4b). In contrast, uniconazole treatment markedly enhanced expression in wild type and *pla1-4*, and moderately enhanced it in pla2-1, plants (Fig. 4b). The opposite effect was detected for GA2ox4, which encodes a GA-catabolizing enzyme. GA treatment strongly enhanced GA2ox4 expression in wild-type and *pla1-4* plants (Fig. 4c). In *pla2-1* plants, GA also induced the expression, but to a limited extent (Fig. 4c). Uniconazole treatment suppressed GA2ox4 expression in wild-type, pla1-4 and *pla2-1* plants (Fig. 4c).

These results show that the feedback mechanism is operating normally in *pla1-4* and *pla2-1* mutants, although somewhat weakened in *pla2-1*. In addition, *PLA1* and *PLA2* may be positioned downstream of GA biosynthesis and signal transduction genes.

## PLA gene expression in GA-related mutants

To confirm the relationship between *PLA* genes and GArelated genes, we examined *PLA1* and *PLA2* expression in GA-related mutants. d18 is a dwarf mutant, whose wildtype gene encodes GA3 oxidase 2. *PLA1* and *PLA2* expression in d18 was slightly down-regulated compared with the wild type (Fig. 5a). However, this difference in *PLA* expression between the wild type and d18-h was not large compared with that between the wild-type and GA-signaling mutants below. In GA-insensitive mutant *gid1* and reduced GA-sensitivity mutant *Slr1-d1*, *PLA1* and *PLA2* expression was severely suppressed, whereas in the GA-constitutive-active mutant *slr1-1*, their expression was markedly increased (Fig. 5b).

To determine the effect of *PLA1* overexpression on the expression domain, we examined the *PLA1* expression pattern by in situ hybridization. Compared with that in the wild-type shoot apex, *PLA1* expression in *slr1-1* was expanded to the adaxial region and the upper regions of leaf primordia and shoot meristems, as well as the normal basal and abaxial regions of leaf primordia (Fig. 5c–e). This ectopic expression of *PLA1* coincided with that induced by GA treatment (Fig. 3d). Since *PLA1* and *PLA2* expression was strongly affected by GA-signaling genes, both *PLA1* and *PLA2* likely act downstream of the GA signal transduction pathway to regulate leaf development.

## Discussion

GA is associated with various developmental and physiological processes, such as seed germination, stem elongation, vegetative phase change, flowering and pollen maturation (Olszewski et al. 2002; Yamaguchi 2008). It is

Fig. 5 PLA1 and PLA2 gene expression in GA-related mutants. a. b Real-time PCR assays of PLA1 and PLA2 expression in GA-deficient mutant (d18-h) (a) and in inactive (gid1 and Slr1-d1) and constitutive active (*slr1-1*) mutants of GA signaling (b). Expression level in each mutant is represented relative to that in wild type. Data in a, b represent mean  $\pm$  SE. **c**–**e** in situ hybridization of PLA1 in wildtype shoot apex (c) and slr1-1 shoot apex (d, e). Arrows indicate ectopic expression. Bars 100 µm



1087



also proposed that GA functions in early leaf development (Olszewski et al. 2002; Yamaguchi 2008). A subset of class I KNOX genes represses GA-biosynthetic gene expression in the SAM by direct transcriptional regulation, resulting in prevention of SAM cells from entering a determinate state (Sakamoto et al. 2001) Once a leaf is initiated from the shoot apex, negative regulation of GA occurs in the leaf primordium, and GA contributes to leaf expansion and differentiation (Olszewski et al. 2002; Yamaguchi 2008). This model is widely accepted, and it is believed that GA is an important regulator of young leaf development. However, the downstream mechanism involved in GA-dependent leaf development remains unknown.

We showed that PLA1 and PLA2 expression was positively regulated by GA. Previous studies indicated that both PLA1 and PLA2 expression is restricted in the leaf primordia, but not in the SAM (Miyoshi et al. 2004; Kawakatsu et al. 2006). This is consistent with GA synthesis in leaf primordia but not in the SAM. GA-related genes (GA3ox2, GA20ox2, SLR1) are expressed in young leaf primordial, including P0 and P0 primordia (Kaneko et al. 2003). This expression domain overlaps with that of *PLA1* and PLA2, supporting the present finding that GA is involved in the regulation of PLA gene expression. The proposed primary function of PLA1 and PLA2 is precocious leaf maturation (Kawakatsu et al. 2006). Leaf maturation is a complex process involving organized expansion and differentiation of cells/tissues. Because GA also plays a role in this organized expansion and differentiation, it is thought that GA action in leaf primordia could be closely related to PLA1 and PLA2 functions. It is assumed that GA regulates leaf development through controlling PLA gene expression. Our in situ hybridization experiments revealed that expression of PLA1 was not only quantitatively enhanced, but was also ectopically expanded to the SAM in GA-signaling mutants and GA-applied plants. These indicate that GA also spatially regulates PLA1 and PLA2 expression.

In contrast, expression of GA biosynthetic, and GA-catabolizing and signaling genes was not significantly altered in *pla1* and *pla2* mutants. In addition, feedback regulation of GA-biosynthetic and GA-catabolizing genes after GA application was normal in *pla1* and *pla2* mutants. The rice DELLA protein, SLR1, is a key component of GA signaling, and is a principal factor responsible for feedback regulation of GA biosynthesis (Itoh et al. 2008; Yamaguchi 2008). The normal SLR1 expression level in the pla mutants suggests that SLR1-dependent GA signal transduction is operating normally.

Many GA-related and GA responsive genes have been identified; for example, PHYTOCHROME INTERACTING FACTOR in skotomorphogenesis, (Feng et al. 2008; de Lucas et al. 2008) and  $\alpha$ -amylase genes in seed germination (Kaneko et al. 2002). In addition, several microarray experiments revealed many GA responsive genes (Yazaki et al. 2003; Yang et al. 2004; Jan and Komatsu 2006). In terms of leaf development, however, less is known about the downstream pathway of GA. Our analyses indicate that PLA1 and PLA2 are factors downstream of GA in leaves, and one action of GA is the PLA-dependent suppression of precocious leaf maturation.

Although the *pla1* and *pla2* leaf phenotypes were similar, PLA1 and PLA2 regulate leaf maturation process through independent genetic pathways. Indeed, a pla1 and pla2 double mutant showed a more severe phenotype than that of the single mutants (Kawakatsu et al. 2006). With regard to their molecular function, PLA1 encodes a member of the cytochrome P450 family thought to be involved in the biosynthetic pathway of an unknown substance (Mivoshi et al. 2004), and PLA2 encodes a MEI2-like RNA-binding protein that may interact with unidentified RNA molecules (Kawakatsu et al. 2006). Considering the independent expression regulation and molecular functions of PLA1 and PLA2, it would not be surprising if PLA1 and PLA2 have functionally diversified. The present results, however, show that both genes are regulated by GA and act downstream of GA signaling. The only difference between *pla1* and *pla2* was in the feedback regulation of GA-related genes. In pla1, the effect of GA<sub>3</sub> and uniconazole treatment on GA20ox2 and GA2ox4 expression was almost identical to those in the wild type. In contrast, pla2 showed a weaker response to these treatments. This indicates that *PLA2* is partially involved in the GA feedback mechanism, but PLA1 is not.

Acknowledgments We thank Dr. H. Kitano (Nagoya University) for kind gift of *d18-h*, *slr1-1*, *Slr1-d1* and *gid1* seeds. This work is supported in part by a Grant-in-Aid for Scientific Research from the Ministry of Education, Culture, Sports, Science and Technology of Japan (23658005 to Y.N.).

#### References

- Anastasiou E, Kenz S, Gerstung M, MacLean D, Timmer J, Fleck C, Lenhard M (2007) Control of plant organ size by *KLUH/ CYP78A5*-dependent intercellular signaling. Dev Cell 13:843–856
- Asano K, Hirano K, Ueguchi-Tanaka M, Angeles-Shim RB, Komura T, Satoh H, Kitano H, Matsuoka M, Ashikari M (2009) Isolation and characterization of dominant dwarf mutants, *Slr1-d*, in rice. Mol Genet Genomics 281:223–231
- Ashikari M, Sasaki A, Ueguchi-Tanaka M, Itoh H, Nishimura A, Datta S, Ishiyama K, Saito T, Kobayashi M, Khush GS, Kitano H, Matsuoka M (2002) Loss-of-function of a rice gibberellin biosynthetic gene, *GA20 oxidase* (*GA20ox-2*), led to the rice 'Green revolution'. Breed Sci 52:143–150
- de Lucas M, Davière JM, Rodríguez-Falcón M, Pontin M, Iglesias-Pedraz JM, Lorrain S, Fankhauser C, Blázquez MA, Titarenko E, Prat S (2008) A molecular framework for light and gibberellin control of cell elongation. Nature 451:480–484
- Evans MM, Poehtig RS (1995) Gibberellins promote vegetative phase change and reproductive maturity in maize. Plant Physiol 108:475–487
- Feng S, Martinez C, Gusmaroli G, Wang Y, Zhou J, Wang F, Chen L, Yu L, Iglesias-Pedraz JM, Kircher S, Schäfer E, Fu X, Fan LM, Deng XW (2008) Coordinated regulation of *Arabidopsis thaliana* development by light and gibberellins. Nature 451:475–479
- Giulini A, Wang J, Jackson D (2004) Control of phyllotaxy by the cytokinin-inducible response regulator homologue *ABPHYL1*. Nature 430:1031–1034

- Gomi K, Sasaki A, Itoh H, Ueguchi-Tanaka M, Ashikari M, Kitano H, Matsuoka M (2004) GID2, an F-box subunit of the SCF E3 complex, specifically interacts with phosphorylated SLR1 protein and regulates the gibberellin-dependent degradation of SLR1 in rice. Plant J 37:626–634
- Hedden P (2003) The genes of the Green Revolution. Trends Genet 19:5–9
- Hirose N, Makita N, Kojima M, Kamada-Nobusada T, Sakakibara H (2007) Overexpression of a type-A response regulator alters rice morphology and cytokinin metabolism. Plant Cell Physiol 48:523–539
- Hong Z, Ueguchi-Tanaka M, Umemura K, Uozu S, Fujioka S, Takatsuto S, Yoshida S, Ashikari M, Kitano H, Matsuoka M (2003) A rice brassinosteroid-deficient mutant, ebisu *dwarf* (*d2*), is caused by a loss of function of a new member of cytochrome P450. Plant Cell 15:2900–2910
- Ikeda A, Ueguchi-Tanaka M, Sonoda Y, Kitano H, Koshioka M, Futsuhara Y, Matsuoka M, Yamaguchi J (2001) Slender rice, a constitutive gibberellin response mutant, is caused by a null mutation of the SLR1 gene, an ortholog of the height-regulating gene GAI/RGA/RHT/D8. Plant Cell 13:999–1010
- Itoh JI, Hasegawa A, Kitano H, Nagato Y (1998) A recessive heterochronic mutation, *plastochron1*, shortens the plastochron and elongates the vegetative phase in rice. Plant Cell 10:1511–1522
- Itoh H, Ueguchi-Tanaka M, Sentoku N, Kitano H, Matsuoka M, Kobayashi M (2001) Cloning and functional analysis of two gibberellin 3 beta-hydroxylase genes that are differently expressed during the growth of rice. Proc Natl Acad Sci USA 98:8909–8914
- Itoh H, Ueguchi-Tanaka M, Sato Y, Ashikari M, Matsuoka M (2002) The gibberellin signaling pathway is regulated by the appearance and disappearance of SLENDER RICE1 in nuclei. Plant Cell 14:57–70
- Itoh J, Nonomura K, Ikeda K, Yamaki S, Inukai Y, Yamagishi H, Kitano H, Nagato Y (2005) Rice plant development: from zygote to spikelet. Plant Cell Physiol 46:23–47
- Itoh H, Ueguchi-Tanaka M, Matsuoka M (2008) Molecular biology of gibberellins signaling in higher plants. Int Rev Cell Mol Biol 268:191–221
- Jackson D, Hake S (1999) Control of phyllotaxy in maize by the *abphyl1* gene. Development 126:315–323
- Jan A, Komatsu S (2006) Functional characterization of gibberellinregulated genes in rice using microarray system. Genomics Proteomics Bioinformatics 4:137–144
- Jönsson H, Heisler MG, Shapiro BE, Meyerowitz EM, Mjolsness E (2006) An auxin-drive polarized transport model for phyllotaxis. Proc Natl Acad Sci USA 103:1633–1638
- Kaneko M, Itoh H, Ueguchi-Tanaka M, Ashikari M, Matsuoka M (2002) The alpha-amylase induction in endosperm during rice seed germination is caused by gibberellins synthesized in epithelium. Plant Physiol 128:1264–1270
- Kaneko M, Itoh H, Inukai Y, Sakamoto T, Ueguchi-Tanaka M, Ashikari M, Matsuoka M (2003) Where do gibberellin biosynthesis and gibberellin signaling occur in rice plants? Plant J 35:104–115
- Kawakatsu T, Itoh J, Miyoshi K, Kurata N, Alvarez N, Veit B, Nagato Y (2006) *PLASTOCHRON2* regulates leaf initiation and maturation in rice. Plant Cell 18:612–625
- Kawakatsu T, Taramino G, Itoh JI, Allen J, Sato Y, Hong SK, Nagasawa N, Kojima M, Kusaba M, Sakakibara H, Sakai H, Nagato Y (2009) *PLASTOCHRON3/GOLIATH* encodes a glutamate carboxypeptidase required for proper development in rice. Plant J 58:1028–1040
- Kouchi H, Hata S (1993) Isolation and characterization of novel nodulin cDNAs representing genes expressed at early stages of soybean nodule development. Mol Gen Genet 238:106–119

- Miyoshi K, Ahn BO, Kawakatsu T, Ito Y, Itoh J, Nagato Y, Kurata N (2004) *PLASTOCHRON1*, a timekeeper of leaf initiation in rice, encodes cytochrome P450. Proc Natl Acad Sci USA 101: 875–880
- Murashige T, Skoog F (1962) A revised medium for rapid growth and bioassays with tobacco tissue cultures. Physiol Plant 15:473–497
- Nakamura A, Fujioka S, Sunohara H, Kamiya N, Hong Z, Inukai Y, Miura K, Takatsuto S, Yoshida S, Ueguchi-Tanaka M, Hasegawa Y, Kitano H, Matsuoka M (2006a) The role of *OsBR11* and its homologous genes, *OsBRL1* and *OsBRL3*, in rice. Plant Physiol 140:580–590
- Nakamura A, Umemura I, Gomi K, Hasegawa Y, Kitano H, Sazuka T, Matsuoka M (2006b) Production and characterization of auxininsensitive rice by overexpression of a mutagenized rice IAA protein. Plant J 46:297–306
- Olszewski N, Sun TP, Gubler F (2002) Gibberellin signaling: biosynthesis, catabolism, and response pathways. Plant Cell 14:61–80
- Reinhardt D, Mandel T, Kuhlemeier C (2000) Auxin regulates the initiation and radial position of plant lateral organs. Plant Cell 12:507–518
- Reinhardt D, Pesce ER, Stieger P, Mandel T, Baltensperger K, Bennett M, Traas J, Friml J, Kuhlemeier C (2003) Regulation of phyllotaxis by polar auxin transport. Nature 426:255–260
- Sakamoto T, Kamiya N, Ueguchi-Tanaka M, Iwahori S, Matsuoka M (2001) *KNOX* homeodomain protein directly suppresses the expression of a gibberellin biosynthetic gene in the tobacco shoot apical meristem. Genes Dev 15:581–590
- Sasaki A, Ashikari M, Ueguchi-Tanaka M, Itoh H, Nishimura A, Swapan D, Ishiyama K, Saito T, Kobayashi M, Khush GS, Matsuoka M (2002) A mutant gibberellin-synthesis gene in rice. Nature 416:701–702
- Sazuka T, Kamiy N, Nishimura T, Ohmae K, Sato Y, Imamura K, Nagato Y, Koshiba T, Nagamura Y, Ashikari M, Kitano H, Matsuoka M (2009) A rice *tryptophan deficient dwarf* mutant, *tdd1*, contains a reduced level of indole acetic acid and develops abnormal flowers and organless embryos. Plant J 60:227–241
- Schwartz S, Grande H, Bujdosso N, Saedler H, Huijser P (2008) The microRNA regulating? SEP-box genes SPL9 and SPL15 control shoot maturation in Arabidopsis. Plant Mol Biol 67:183–195
- Smith RS, Guyomarc'h S, Mandel T, Reinhardt D, Kuhlemeier C, Prusinkiewicz P (2006) A plausible model of phyllotaxis. Proc Natl Acad Sci USA 103:1301–1306

- Steeves TA, Sussex IM (1989) Patterns in plant development. Cambridge University Press, Cambridge
- Tanabe S, Ashikari M, Fujioka S, Takatsuto S, Yoshida S, Yano M, Yoshimura A, Kitano H, Matsuoka M, Fujisawa Y, Kato H, Iwasaki Y (2005) A novel cytochrome P450 is implicated in brassinosteroid biosynthesis via the characterization of a rice dwarf mutant, dwarf11, with reduced seed length. Plant Cell 17:776–790
- Tanaka N, Itoh H, Sentoku N, Kojima M, Sakakibara H, Izawa T, Itoh JI, Nagato Y (2011) The COP1 ortholog PPS regulates the juvenile-adult and vegetative-reproductive phase changes in rice. Plant Cell 23:2143–2154
- Teale WD, Paponov IA, Palme K (2006) Auxin in action: signalling, transport and the control of plant growth and development. Nat Rev Mol Cell Biol 7:847–859
- Teifer A, Bullman KM, Poethig RS (1997) Phase change and the regulation of trichome distribution in *Arabidopsis thaliana*. Development 124:1889–1898
- Ueguchi-Tanaka M, Ashikari M, Nakajima M, Itoh H, Katoh E, Kobayashi M, Chow TY, Hsing YI, Kitano H, Yamaguchi I, Matsuoka M (2005) *GIBBERELLIN INSENSITIVE DWARF1* encodes a soluble receptor for gibberellin. Nature 437:693–698
- Yamaguchi S (2008) Gibberellin metabolism and its regulation. Annu Rev Plant Biol 59:225–251
- Yamamuro C, Ihara Y, Wu X, Noguchi T, Fujioka S, Takatsuto S, Ashikari M, Kitano H, Matsuoka M (2000) Loss of function of a rice *brassinosteroid insensitive1* homolog prevents internode elongation and bending of the lamina joint. Plant Cell 12:1591–1606
- Yang GX, Jan A, Shen SH, Yazaki J, Ishikawa M, Shimatani Z, Kishimoto N, Kikuchi S, Matsumoto H, Komatsu S (2004) Microarray analysis of brassinosteroids- and gibberellin-regulated gene expression in rice seedlings. Mol Genet Genomics 271:468–478
- Yazaki J, Kishimoto N, Nagata Y, Ishikawa M, Fujii F, Hashimoto A, Shimbo K, Shimatani Z, Kojima K, Suzuki K, Yamamoto M, Honda S, Endo A, Yoshida Y, Sato Y, Takeuchi K, Toyoshima K, Miyamoto C, Wu J, Sasaki T, Sakata K, Yamamoto K, Iba K, Oda T, Otomo Y, Murakami K, Matsubara K, Kawai J, Carninci P, Hayashizaki Y, Kikuchi S (2003) Genomics approach to abscisic acid- and gibberellin-responsive genes in rice. DNA Res 10:249–261