

Expression Site of Protoporphyrinogen Oxidase Influences on Herbicide Resistance in Transgenic Rice

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형질전환 벼에서 Protoporphyrinogen Oxidase의 발현 위치가 제초제 저항성에 미치는 영향

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ABSTRACT The effect of Protox expression site on herbicidal resistance was investigated in wild-type and transgenic rice plants imposed by peroxidizing herbicide oxyfluorfen. The transgenic rice systems involved the plastidal expression of *Arabidopsis* protoporphyrinogen oxidase (Protox; AP line) and the dual expression of *Myxococcus xanthus* Protox in chloroplasts and mitochondria (TTS line). The oxyfluorfen-treated TTS4 line showed the lower levels of cellular leakage and malonyldialdehyde and the sustained capacity of 5-aminolevulinic acid synthesis, compared to the oxyfluorfen-treated AP and wild-type lines. During oxyfluorfen action, the TTS4 line had greater herbicide resistance than the AP1 line, indicating that the dual expression of *M. xanthus* Protox in chloroplasts and mitochondria prevented the accumulation of photodynamic protoporphyrin IX more effectively than the expression of *Arabidopsis* Protox only in chloroplasts. These results suggest that the ectopic expression of Protox in mitochondria greatly contributes to the herbicidal resistance in rice plants.

Key words: 5-aminolevulinic acid; herbicide resistance; oxidative stress; oxyfluorfen; protoporphyrinogen oxidase.

INTRODUCTION

Tetrapyrrole biosynthesis is important in plants because it provides many essential molecules that are involved in light harvesting, energy transfer,

signal transduction, detoxification, and systemic acquired resistance (Grimm 1998; Molina *et al.* 1999; Smith *et al.* 1993). The porphyrin ring system is derived from 5-aminolevulinic acid (ALA). ALA is formed in plants from the five-carbon skeleton of

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glutamate in unit of the C5 pathway (Beale 1978; von Wettstein *et al.* 1995). The last common step in the tetrapyrrole pathway to heme and chlorophyll is the oxidation of protoporphyrinogen IX (Proto IX) to protoporphyrin IX (Proto IX), which is catalyzed by protoporphyrinogen oxidase (Protox, EC 1.3.3.4) (Jacobs and Jacobs 1987; Matringe *et al.* 1992). Plants express two Protox isozymes : one is associated with the chloroplast envelope and thylakoid membranes (Matringe *et al.* 1992), and the other is localized to mitochondria (Smith *et al.* 1993).

There is increased interest in the mechanisms of plant chlorophyll synthesis owing to recent studies describing the light-dependent action of photobleaching diphenylether herbicides including oxyfluorfen and acifluorfen (Jacobs and Jacobs 1993; Jung and Back 2005; Kouji *et al.* 1988; Lee *et al.* 2003; Lydon and Duke 1988). They are photodynamically active and competitively block the substrate-binding region of Protox. The acute toxicity of oxyfluorfen is a result of the accumulation of Proto IX, which is oxidized in the cytoplasm to Proto IX by a nonspecific plasma membrane-bound peroxidase (Jacobs *et al.* 1991; Lee and Duke 1993). Proto IX in the cytoplasm is not metabolized by the porphyrin biosynthetic pathway because of Mg-chelatase and Fe-chelatase, which use Proto IX as a substrate, are located in chloroplasts and mitochondria. Cytoplasmic Proto (gen) IX, a potent photosensitizer, absorbs light that is used in detrimental reactions in which energy or electrons are subsequently transferred to oxygen, resulting in the formation of highly reactive oxygen, rapid lipid peroxidation, and cell death.

In the transgenic plants overexpressing Protox gene in either chloroplasts or mitochondria, the action of Protox-inhibiting herbicides was decreased by preventing the accumulation of Proto IX, decreasing the light-dependent phytotoxicity (Ha *et al.* 2004;

Jung *et al.* 2006; Lermontova and Grimm 2000; Warabi *et al.* 2001). This study investigated whether the expression site of Protox tightly correlates to the peroxidizing-herbicide resistance in the two transgenic rice systems involving the expression of *Arabidopsis* Protox in chloroplasts (AP line) (Ha *et al.* 2004) and the dual expression of *Myxococcus xanthus* Protox in chloroplasts and mitochondria (TTS line) (Jung *et al.* 2006). The relation between the levels of cellular leakage and oxidative status was monitored in the two transgenic and WT lines imposed by oxyfluorfen compared with controls. The regulation of ALA level upon the herbicidal treatment was also compared to the wild-type and transgenic rice plants.

MATERIALS AND METHODS

Plant materials and growth conditions

The T₄ generations of homozygous rice lines (AP1) (Ha *et al.* 2004) expressing *Arabidopsis* Protox in chloroplasts and the T₃ generations of homozygous rice line (TTS4) expressing *Myxococcus xanthus* Protox in chloroplasts and mitochondria (Jung *et al.* 2006) were used for physiological experiments. Seeds of wild-type and transgenic lines were planted, and seedlings were potted in paddy soil and grown in the greenhouse, which was kept at 28 to 30°C. Parts of leaves from 3-week-old rice plants were taken for experiments.

Herbicide treatment

Technical-grade oxyfluorfen for conductivity measurement was generously provided by Kyungnong (Gyeongju, Korea). Commercially available oxyfluorfen (Goal[®]) was used for the foliar application test. Three-week-old rice plants were sprayed with 20 µM oxyfluorfen, placed in darkness for 12 h to

allow absorbance, and then illuminated for 30 h. The treated leaf tissues were used for determination of malonyldialdehyde (MDA) and ALA analyses.

Cellular leakage

The rice leaf tissues were treated with oxyfluorfen as described previously (Kenyon *et al.* 1985; Lee *et al.* 1995) by cutting mature leaves to 4-mm squares (0.1 g fresh wt.) and then placing them in a 6-cm diameter polystyrene Petri dish containing 5 mL of 1% sucrose, 1 mM 2-(N-morpholino)ethanesulfonic acid (MES, pH 6.5) with or without the herbicide dissolved in acetone. The MES buffer was used to reduce the oxyfluorfen-caused change in the pH of bathing solution. The controls contained the same amount of the solvent without the herbicide. The final concentration of acetone in all dishes was 1% (v/v). The tissues were incubated with 20 μM of oxyfluorfen in a growth chamber at 25°C in darkness for 12 h and then exposed to continuous light at 250 $\mu\text{mol m}^{-2} \text{s}^{-1}$ photosynthetically active radiation (PAR) for 24 h. Cellular leakage was determined periodically by detecting the electroconductivity of the bathing medium using a conductivity meter (Cole-Parmer Instruments) as previously described (Lee *et al.* 1995). Because of differences in the background conductivity of different treatment solutions, the results were expressed as changes in conductivity upon exposure to light.

Determination of malonyldialdehyde content

For the measurements of oxidative stress in leaves, the thiobarbituric acid (TBA) test, which determines MDA as an end product of lipid peroxidation (Buege and Aust 1978), was used. Leaves (0.1 g) were homogenized in 2 mL of a solution of 0.5% TBA in 20% trichloroacetic acid. The homogenate was centrifuged at 20,000 \times g for 15 min. and the supernatants were collected. The supernatants were heated in

boiling water for 25 min., and the reaction was stopped by placing the reaction tubes in an ice bath. Then the samples were centrifuged at 20,000 \times g for 15 min., and the absorbance of the supernatant at 532 nm was recorded for each sample and corrected for nonspecific turbidity at 600 nm. The MDA concentration was calculated using a molar extinction coefficient of 156 $\text{mM}^{-1} \text{cm}^{-1}$ (Buege and Aust 1978).

ALA-synthesizing capacity

ALA-synthesizing capacity was measured as described by Papenbrock *et al.* (1999). Leaf disks were harvested for each sample, incubated in 20 mM phosphate buffer containing 40 mM levulinic acid in the light for 6 h, and frozen in liquid nitrogen. Samples were homogenized, resuspended in 1 mL of 20 mM $\text{K}_2\text{HPO}_4/\text{KH}_2\text{PO}_4$ (pH 6.9), and centrifuged at 10,000 \times g. The 500- μL supernatant was mixed with 100 μL ethylacetoacetate, boiled for 10 min., and cooled for 5 min.. An equal volume of modified Ehrlich's reagent was added and the absorption of the chromophore was determined at 553 nm with the spectrophotometer.

RESULTS AND DISCUSSION

Comparison of herbicidal resistance in transgenic rice plants expressing either *Arabidopsis* or *M. xanthus* Protox

In the previous study, the transgenic rice overexpressing *M. xanthus* Protox (TTS) showed greatly increased Protox activity in both chloroplasts and mitochondria in comparison with wild-type, which suggested that the *M. xanthus* Protox was dually targeted into both organelles (Jung *et al.* 2006). However, the introduction of *Arabidopsis* Protox (AP) into rice led to the ectopic expression of Protox only in chloroplasts (Ha *et al.* 2004). The

overexpression of Protox in the two transgenic rice lines resulted in greater herbicidal resistance to peroxidizing herbicide oxyfluorfen in comparison to wild-type line. In the present study, the wild-type and transgenic lines, AP1 and TTS4, were exposed to oxyfluorfen treatment to assess the effect of differential Protox expression site on herbicidal resistance.

The necrotic phenotypes of herbicide-treated plants typically display leaf desiccation, veinal necrosis, and leaf deformation (Knörzer and Böger 1999). With 20 μM oxyfluorfen treatment, the necrosis of leaf squares (data not shown) in wild-type plants was associated with time-dependent increases in conductivity until 18 h after illumination (Fig. 1), an indication of cellular leakage, caused by an accumulation of Proto IX. Excited Proto IX molecules present in excess interact with O_2 and give rise, by triplet-triplet interchange, to highly toxic $^1\text{O}_2$, that in turn triggers membrane lipid

peroxidation, protein denaturation, and pigment bleaching (Dolphin 1994; Reinbothe *et al.* 1996). In the transgenic line, AP1, the conductivity began to drastically increase 12 h after illumination and then continuously increased until 18 h after the illumination (Fig. 1). In contrast, the TTS transgenic line, TTS4, showed only a minor increase in conductivity in response to oxyfluorfen treatment. Overall, the dual expression of *M. xanthus* Protox in chloroplasts and mitochondria in the TTS4 line resulted in a lesser increase in conductivity during oxyfluorfen action than that of the AP1 line, indicating the reduction in the leakage of photosensitizing Protoporphyrin IX from chloroplasts and mitochondria to the cytosolic membranes.

MDA content, which is a measure of lipid peroxidation, was assayed in rice plants. The levels of MDA were the same in the untreated leaf disks of wild-type and the two transgenic lines (Fig. 2). The MDA level in the wild-type line was greatly

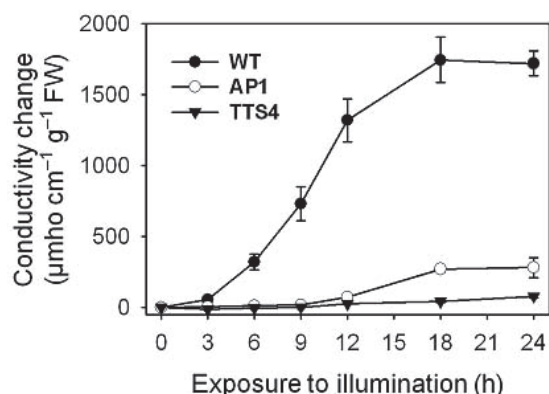


Fig. 1. Effect of oxyfluorfen on time-dependent change of cellular leakage from leaf squares of rice plants. Conductivity, which reflects electrolyte leakage, was measured in the bathing solution after leaf segments were treated with 20 μM oxyfluorfen. Tissues were exposed to continuous white light at 250 $\mu\text{mol m}^{-2} \text{s}^{-1}$ PAR for 24 h following a 12-h dark incubation. WT, wild type; AP1, transgenic line overexpressing *Arabidopsis* Protox; TTS4, transgenic line overexpressing *M. xanthus* Protox. The data represent the mean \pm S.E. of three replicates.

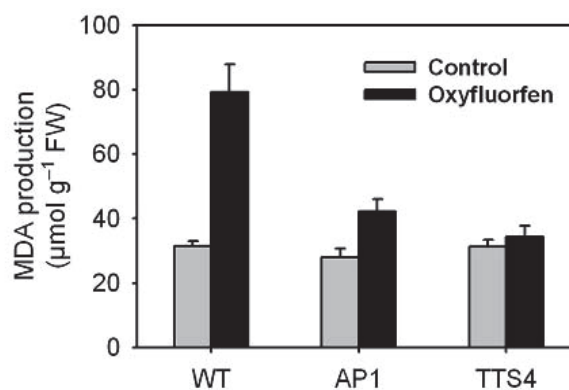


Fig. 2. Effect of foliar application of oxyfluorfen on MDA production from leaf tissues of rice plants. MDA production was measured in leaf tissues after treatments with oxyfluorfen. Three-week-old rice plants were sprayed with a 20 μM concentration of oxyfluorfen, placed in darkness for 12 h to allow absorbance, and then illuminated for 30 h. WT, wild type; AP1, transgenic line overexpressing *Arabidopsis* Protox; TTS4, transgenic line overexpressing *M. xanthus* Protox. The data represent the mean \pm S.E. of three replicates.

increased up to 150% in response to 20 μM of oxyfluorfen treatment, but it was not altered in the transgenic line, TTS4. The AP1 line showed 51% increase in MDA production relative to its control. Membrane disruption by nonenzymatic lipid peroxidation destroys cellular compartments, causes loss of solutes and dehydration, and finally leads to cell death (Mock *et al.* 1999). Due to the remarkable reactivity of all tetrapyrroles, living organisms have a substantial danger that uncontrolled chemical reactions may occur (Jung *et al.* 2006, 2008; Yao and Greenberg 2006). The dual expression of Protox in TTS plants efficiently prevented the oxyfluorfen-caused accumulation of Proto(gen) IX, thereby suffering less oxidative stress due to the reduced production of reactive oxygen species, compared to the wild-type and AP1 plants.

The effect of oxyfluorfen on ALA-synthesizing capacity

The rate of synthesis of the porphyrin precursors is tightly controlled to reflect the requirements of the plant for chlorophyll in the light. ALA is a key precursor in the early step of porphyrin biosynthesis such as chlorophyll and heme, and its formation may be the rate limiting step (von Wettstein *et al.* 1995). The ALA-synthesizing capacity was significantly decreased in the oxyfluorfen-treated wild-type and AP1 transgenic lines, as compared to controls, with a greater decrease in the wild-type line (Fig. 3). The accumulated levels of Protogen IX caused by oxyfluorfen treatment were toxic enough to cause the deregulation of the precursor ALA in the wild-type and AP1 transgenic lines. The Protogen IX accumulation due to Protox inhibition may cause the feedback inhibition of ALA synthesis for decreasing the new synthesis of Protogen IX in the oxyfluorfen-treated plants. However, the transgenic line TTS4 did not alter the ALA-synthesizing capacity

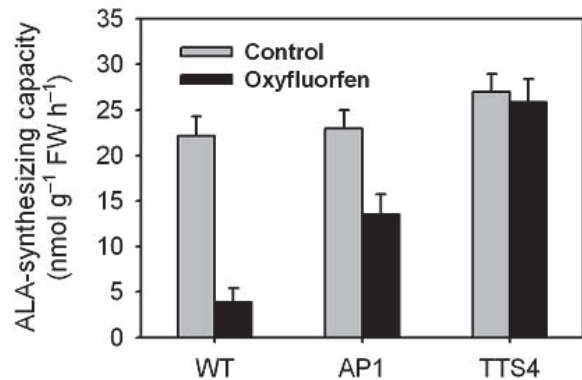


Fig. 3. Effect of oxyfluorfen on the ALA-synthesizing capacity in WT and transgenic plants. Three-week-old rice plants were sprayed with a 20 μM concentration of oxyfluorfen, placed in darkness for 12 h to allow absorbance, and then illuminated for 30 h. WT, wild type; AP1, transgenic line overexpressing *Arabidopsis* Protox; TTS4, transgenic line overexpressing *M. xanthus* Protox. The data represent the mean \pm S.E. of three replicates.

in response to oxyfluorfen treatment (Fig. 3), demonstrating that the ALA synthesis in TTS line was not disturbed by oxyfluorfen treatment.

In conclusion, this study provides evidence of a relation of Protox expression site with herbicidal resistance in rice plants. Both wild-type and AP1 lines greatly increased the levels of conductivity and MDA in response to oxyfluorfen (Figs. 1 and 2), which may be due to a drastic accumulation of Proto IX in the oxyfluorfen-treated tissues. However, the transgenic line TTS4 did not show any sign of oxidative stress with oxyfluorfen treatment. The herbicidal resistance in TTS4 may be because of the dual expression of *M. xanthus* Protox in chloroplasts and mitochondria, which greatly reduces the leakage of photosensitizing Protogen IX from chloroplasts and mitochondria to the cytosolic membranes. This result also suggests that the increased expression level of Protox in mitochondria greatly contributes to the herbicidal resistance in rice plants. The elucidation of a critical role of Protox activity in herbicidal resistance from this study is particularly

important because it may provide the basis for the construction of new, porphyrin-based, photobleaching herbicides to be exploited in modern weed control.

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요 약

과산화제 제초제 oxyfluorfen이 처리된 비형질전환 벼와 형질전환 벼에서 Protox 발현 위치가 제초제 저항성에 미치는 영향을 비교하였다. *Arabidopsis* protoporphyrinogen oxidase(Protox; AP 계통)를 색소체에만 발현하는 형질전환 벼와 *Myxococcus xanthus* Protox 유전자를 색소체와 미토콘드리아에 모두 발현하는 형질전환 벼(TTS 계통)가 형질전환 시스템으로 사용되었다. Oxyfluorfen이 처리된 TTS4 계통은 AP 계통이나 비형질전환 벼에 비해 낮은 수준의 세포질 누출 및 malonyldialdehyde를 보여주었고, 높은 5-aminolevulinic acid 합성 능력을 유지하였다. Oxyfluorfen 작용 동안, TTS4 계통은 AP1 계통보다 높은 제초제 저항성을 보여주었는데, 이는 색소체만에서의 *Arabidopsis* Protox의 발현에 비해 색소체와 미토콘드리아에서의 *M. xanthus* Protox의 쌍발현이 광역학적인 protoporphyrin IX의 축적을 더 효율적으로 억제하였기 때문일 것이다. 이 결과들은 미토콘드리아 내 Protox의 발현이 Protox 저해형 제초제에 대한 식물의 저항성에 크게 기여함을 의미한다.

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