

Isolation of cadmium-induced DNA sequence in microalga *Nannochloropsis oculata*

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(Received: March 22, 2007; Revised received: November 03, 2007; Accepted: December 26, 2007)

Abstract: The gene induced in response to Cd exposure in microalga *Nannochloropsis oculata* was examined in this investigation. In order to isolate this gene by using nested PCR, degenerated primers were designed on the basis of highly conserved regions of the amino acid sequences of various phytochelatin synthase. The size of the nested PCR amplification product from *Nannochloropsis oculata* by using two degenerated primers was measured to be 266 bp. The 266 bp DNA fragment was sequenced, and it might be induced by Cd exposure in microalga *Nannochloropsis oculata*.

Key words: Cadmium, DNA, *Nannochloropsis oculata*

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Introduction

There has been considerable concern about the increasing levels of heavy metals like cadmium (Cd) in the environment due to the potential toxic effects on living organism (Pinto *et al.*, 2003; Akinola and Ekiyoyo, 2006; Phetsombat *et al.*, 2006). The photosynthetic process is very sensitive to Cd; it reduces growth, metabolism and low biomass accumulation (Payne and Price, 1999; Sanita and Gabbrielli, 1999). Cd also produces oxidative stresses in the cell possibly by generating free radicals and reactive oxygen species as we have reported previously (Lee *et al.*, 2000; Lee, 2002; Lee and Shin, 2003). One protective strategy against metal excess is the expression of high-affinity binding sites to suppress uncontrolled binding of metal ions to functional groups (Ahner and Morel, 1995; Pistocchi *et al.*, 2000). Phytochelatin (PC), glutathione-derived metal binding peptides, are heavy metal binding peptides that play an important role in sequestration and detoxification of heavy metals in various organisms. A few years ago the genes encoding phytochelatin synthase (PCS) were cloned from plants, fungi and nematodes. Expression of Arabidopsis PCS in Indian Mustard enhanced tolerance for Cd and Zn (Gasic and Korban, 2006). The tolerance of zebrafish to heavy metal toxicity was also enhanced by the expression of plant PCS (Konishi *et al.*, 2006).

PCS was more strongly activated by Cd²⁺ than by Zn²⁺ in marine green alga, *Dunaliella tertiolecta* (Tsuji *et al.*, 2003). Proline-mediated tolerance to Cd through increased phytochelatin synthesis and sequestration of Cd in transgenic microalgae was also reported (Siripornadulsil *et al.*, 2002).

However, the information about cadmium-induced genes in marine microalgae is not sufficient in spite of the importance for environmental genomics. In this investigation, cadmium-induced DNA sequence was isolated and characterized from *Nannochloropsis oculata* by using nested PCR with degenerated primers for PCS.

Materials and Methods

Algal culture and cadmium treatment: A unialgal strain of *Nannochloropsis oculata* was obtained from the Korea Marine Microalgae Culture Collection (KMCC) and cultured in the f/2 medium of Guillard and Ryther (1962). Erlenmeyer flasks containing 100 ml medium were inoculated with 25 ml of a mother culture. All the media for microalgal growth were made with electrolyzed water. The alga was grown at room temperature with shaking at 100 rpm under a 12:12 light-dark cycle of 50 $\mu\text{mol photon m}^{-2} \text{s}^{-1}$ illumination. For cadmium treatment, a 10-day old culture was further cultured for 4 days in similar medium, but with 10 $\mu\text{M CdCl}_2$.

Algal DNA extraction: Algal material treated with 10 $\mu\text{M CdCl}_2$ for 4 days was powdered in liquid nitrogen. This powdered was further homogenized in a solution of 50 mM Tris-HCl (pH 8.0), 5 mM EDTA, and 50 mM NaCl. Lysozyme was added and the solution was incubated at 55°C for 30 min. After incubation, proteinase K (20 mg/ml) and 10% SDS were added, and the mixture was incubated at 55°C for 30 mins. This solution was chilled on ice and then extracted with equal volume of phenol-chloroform-isoamylalcohol. The organic extraction step was repeated, and the supernatant was added to an equal volume of 4M ammonium acetate. The DNA was precipitated by adding 2 volumes of isopropanol, and collected by centrifugation for 10 min at room temperature (Sambrook and Russell, 2001).

Nested PCR with degenerated oligonucleotides: Degenerated oligonucleotides designed on the basis of the highly conserved regions of the amino acid sequences of various phytochelatin synthases were used for nested PCR. PCR was performed in a total volume of 20 μl containing 200 $\mu\text{M dNTPs}$, 2 mM MgCl_2 , 1 unit of Taq polymerase, 1X PCR buffer, 1 μl of DNA template and 1 pmol of sp-1 primer (garccngartaytygyg) and asp-2 primer (naccartgngnggrga). Amplification was carried out with 30 cycles of 94°C for 1 min, 55°C for 1 min and 72°C for 1 min using a PROGENE thermal cycler. The primary PCR product was used as



a template for a secondary PCR with a nested sp-2 primer (nnnnatgctngaytytg) and asp-1 primer (nnnnggrctnggrtgnc) in a reaction mixture. A secondary amplification was also carried out with 30 cycles of 94°C for 1 min, 55°C for 1 min and 72°C for 1 min using the same thermal cycler. The PCR product was subcloned and characterized by DNA sequencing (Sambrook and Russell, 2001).

Results and Discussion

Both essential and non-essential transition metal ions can be toxic to cells. The physiological range for essential metals between deficiency and toxicity is therefore extremely narrow and a tightly controlled metal homeostasis network to adjust to fluctuations in micronutrient availability is necessary for all organisms (Clemens, 2006). One protective strategy against metal excess is the production of high affinity binding sites to inhibit uncontrolled binding of metal ions to physiologically important functional groups. Phytochelatins are heavy metal-binding peptides that play important roles in the detoxification of toxic heavy metals. Phytochelatins also regulate intracellular concentrations of essential metals in eukaryotes, including

Table - 1: Degenerated primers used in the nested PCR

Primer	Sequence (5'→3')	Product length (bp)
PCS sp-1	garccngartaytygg	423
PCS asp-2	naccartgngnggrga	
PCS sp-2	nnnnatgctngaytytg	237
PCS asp-1	nnnnggrctnggrtgnc	

higher plants, fungi, and microalgae. Recently, PC synthase (PCS) genes in higher plants and fission yeast have been identified and characterized (Clemens, 2006), enabling molecular biological investigation to clarify the mechanisms underlying PC synthesis. Moreover, recent routine database searches have unexpectedly identified genes that are similar to plant PCS genes in the genomes of worms and some prokaryotes (Hirata *et al.*, 2005). In response to heavy metal exposure, the constitutively expressed phytochelatin synthase enzyme (PCS) is activated leading to synthesis of PCs in the cytosol (Picault *et al.*, 2006). Moreover, recent attempts to enhance metal accumulation and tolerance in plant reported that

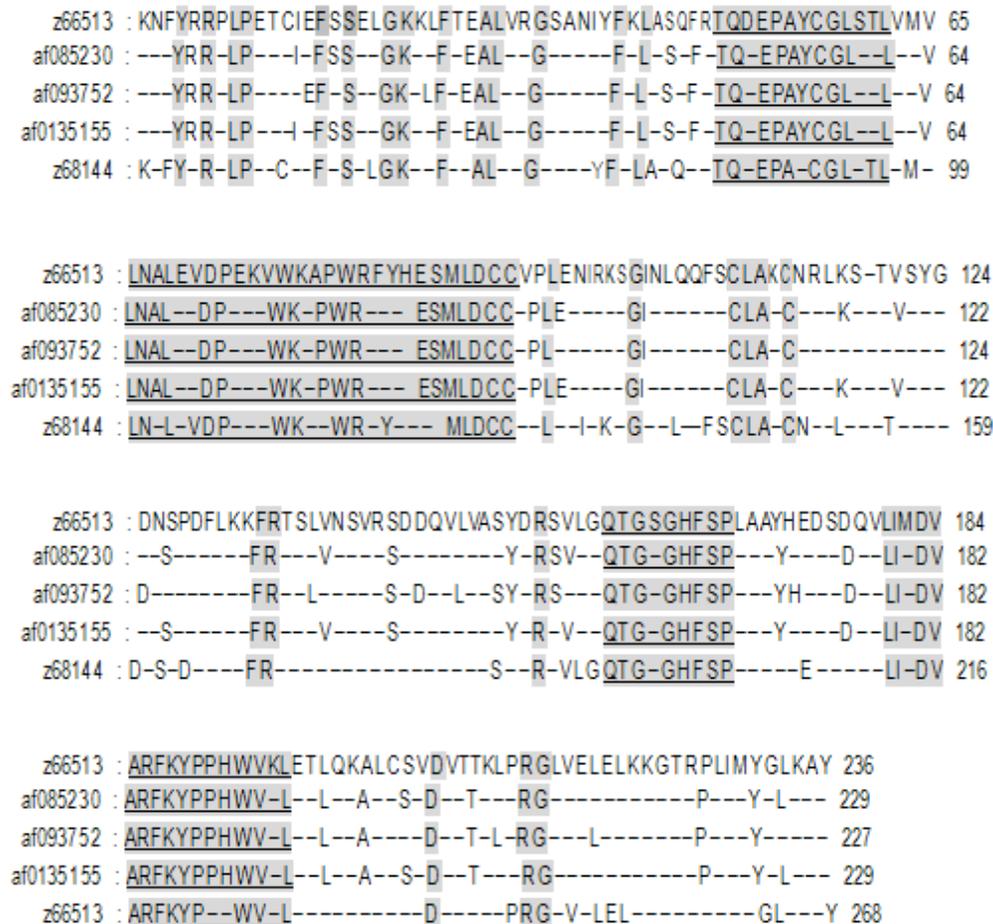


Fig. 1: Amino acid sequence alignment of putative phytochelatin synthase. Degenerate primers underlined were designed based on the highly conserved regions

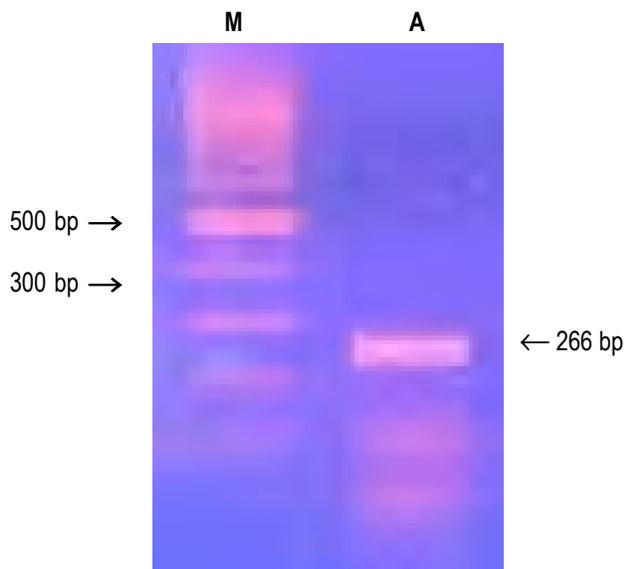


Fig. 2: PCR amplification of a PCS gene from *Nannochloropsis oculata* using degenerated primers. Lane M: 100 bp DNA ladder, Lane A: Nested PCR products from *Nannochloropsis oculata*

PCS over-expression in transgenic plants paradoxically induced cadmium hypersensitivity (Picault *et al.*, 2006).

In this study, our goal is to isolate the gene induced in response to Cd exposure in marine microalga *Nannochloropsis oculata*. Cd-induced oxidative stress and the involvement of antioxidant enzymes in *Nannochloropsis oculata* following 10 μ M Cd treatment were reported in our previous investigation (Lee and Shin, 2003). Increasing concentrations of Cd produced growth inhibition. The reduction was about 15% with 10 μ M Cd and 50% growth with 20 μ M Cd. The markedly elevated levels of various antioxidant enzymes, named ascorbate peroxidase, guaiacol peroxidase and catalase following 10 μ M Cd treatment indicated the protective role of the enzymes against Cd induced oxidative stress. Moreover, the oxidative stress might result in the activations of various stress-related signals at gene levels. Therefore, in order to isolate Cd stress-related gene, degenerated primers were designed on the basis of highly conserved regions of the amino acid sequences from various phytochelatin synthases. Amino acid sequence alignment of various putative phytochelatin synthases was shown in Fig. 1. Table. 1 indicates the degenerated oligonucleotide primers designed on the basis of the highly conserved regions of the amino acid sequences from various phytochelatin synthases. The size of the nested-PCR amplification product from *Nannochloropsis oculata* using two sets of degenerated primers was measured to be 266 bp as shown in Fig. 2.

Fig. 3 shows the nucleotide sequences of the nested PCR product. This sequence indicates the Cd-induced DNA fragment in marine alga *Nannochloropsis oculata*. Further molecular biological studies in *Nannochloropsis oculata* will allow testing of this DNA

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1 ATCGGGGCTC GGATGGCCCC CAAGCACAGT CAGTTTGCCA
  GACTCACAGA TTTTCTAGT
61 TGCCAACGGT TCCATTCTTG GGGATACATT ATTTTCTAGT
  ACAAATGAGA ATTTAGCAAT
121 GGTAAGTGGT TATTCAGGAT TTCCTCAGTC AATAAAGGGG
  GCTGCAGATC TCCATACAAG
181 CTCTATATAC GATCATCATG TGAAGTCAAG TGACGGGGAT
  GTTGTTGGC TTAAGACAGA
241 TAAATTCGGC AGCAATCAAG CATTCA

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Fig. 3: Nucleotide sequences of the nested PCR product from *Nannochloropsis oculata*

sequence for toxic Cd sequestration and detoxification. More detailed experiments will be needed to fine out and isolate full length Cd-induced gene in *Nannochloropsis oculata*. The information on the Cd-induced DNA sequence obtained can be utilized as indicators for environmental stresses. For understanding the roles of the Cd-induced gene in the alga, experiments using transgenic alga with induction or suppression of the gene will be needed. The induction or repression of genes under the cadmium stress should be helpful in understanding the mode of action of marine organisms to defend themselves against cadmium toxicity.

Acknowledgments

The nucleotide sequence reported in this paper has been submitted to the NCBI data bank with accession number BU709656. This work was supported by Korea Ministry of Environment as "The Eco-technopia 21 project".

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