

**Nucleotide sequence of a full length cDNA clone of light harvesting chlorophyll a/b binding protein gene from green dark-grown pine (*Pinus thunbergii*) seedling**

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Light has been shown to induce the expression of gene encoding light harvesting chlorophyll a/b binding protein(LHCP)(1). Dark-grown seedling of coniferous plant is green. We have constructed cDNA library and cloned full length cDNA (pDDLHC2176) for LHCP from dark-grown pine (*Pinus thunbergii*) seedlings in pUC8, which was identified by cross-hybridization with duckweed genomic clones(2) and complete sequencing. The cDNA included 801 bp of the open reading frame, and 49 and 128 nucleotides of 5'- and 3'-untranslated sequences, respectively. From the putative cleavage site(arrow head), the transit and mature polypeptides have been estimated to be composed of 37 and 229 amino acids, respectively. This result also shows the expression of the gene in the dark-grown pine seedling.

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-49 CGCATTACCTGACTTTGGTCTGAGCATTMTTCTCGTCGTCACGTTCCACC
1 ATGGCAACAGCTTCAGCCATCCAAAGCTCAAGCTTGGCAGGCCAGACCCCTCCTAAGGCCG
M A T A S A I Q S S S L A G Q T L L R P
61 CAGCAGAATGAGCTCGTCAAGAAAGTGGGCACGGCGCAGGCTCGAATCACCATGCGAAGA
Q Q N E L V K K V G T A Q A R I T M R R
121 ACCGTAAGGAGCGCCCCGAGAGCATTGGTATGGACCTGACCGCCCAAGTACCTAGGC
T V R S A P E S I W Y G P D R P K Y L G
181 CCCTCTCGGAAGGACCGCTCATATCTCACCGGAGAATTTCCCGCGACTACGGGTGG
P F S E G T P S Y L T G E F P G D Y G W
241 GACACTGCGCCGCTCTCGGCGGATCCAGAGACCTTCGCAAAAAACAGAGAGCTGGAGGTG
D T A A V S A D P E T F A K N R E L E V
301 ATCCAATGAGATGGCCATGTTGGGAGCGCTCGGCTGCGTTTCCCGGAGCTGTGGCC
I H C R W A M L G A L G C V F P E L L A
361 AAAAAATGGTTGAAAATTTGGGGAAGCTGTGTGGTTCAAGGCGGGGCGAGATATTCTCA
K N G L K F G E A V W F K A G A Q I F S
421 GAGGAGGCCTTGACTACGCTGGGAACCCCAACCTGATCCACGCGCAGAGCATTCTAGCC
E G G L D Y A G N P N L I H A Q S I L A
481 ATCTGGGCTGCCAGGTTGTTCTCATGGGATTGATTGAAGGATACAGAGTGGGAGGAGG
I W A C Q V V L M G L I E G Y R V G G G
541 ACCCTTGGAGAGGGTTGGACCCCTGTTACCAGGGGTGCTTCGACCCACTGGGGCTG
T L G E G L D P L L P G G A F D P L G L
601 GCOGACGACCCGAGGCCTGCGCGGAGCTGAAGGTGAAAGAGATTAAGAACGGTCCGGCTG
A D D P E A C A E L K V K E I K N G R L
661 GCCATGTTCTCCAATGTTCCGTTTCTTCGTTGAGGCAATCGTGACCGGGAAGGCCCCATT
A M F S M F G F F V Q A I V T G K G P I
721 GAAAATCTCTACGACCACCTGGCCGACCCCGTTGCCAACAAATGCGCTGGCCCTACGCCACC
E N L Y D H L A D P V A N N A W A Y A T
781 AATTTCTGTTCTGGCAAGTGAAGGTGACGGAAAAATAAAGAGGCCTGTGATCTGTGCATC
N F V P G K -
841 AATCATTTGACAGCCTTAGTGTAAATAAAATATGTTCTTTCAGCTGGATGTATTTGTGG
901 TGATCTTCGTTAATAAAATATTTTCTTTC

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