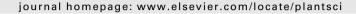
ELSEVIER

#### Contents lists available at ScienceDirect

### Plant Science





Cloning and expression of a plastid-encoded subunit, beta-carboxyltransferase gene (*accD*) and a nuclear-encoded subunit, biotin carboxylase of acetyl-CoA carboxylase from oil palm (*Elaeis guineensis* Jacq.)

Alisa Nakkaew<sup>a</sup>, Wilaiwan Chotigeat<sup>a</sup>, Theera Eksomtramage<sup>b</sup>, Amornrat Phongdara<sup>a,\*</sup>

- <sup>a</sup> Center for Genomics and Bioinformatics Research, Faculty of Science, Prince of Songkla University, Hat-Yai, Songkhla 90110, Thailand
- b Department of Plant Science, Faculty of Natural Resources, Prince of Songkla University, Hat-Yai, Songkhla 90110, Thailand

#### ARTICLE INFO

Article history:
Received 10 October 2006
Received in revised form 16 May 2008
Accepted 28 May 2008
Available online 11 June 2008

Keywords:
Oil palm
Elaeis guineensis Jacq.
Beta-carboxyltransferase
accD, acetyl-CoA carboxylase (ACCase)
accC, biotin carboxylase (BC)

#### ABSTRACT

Palm oil is the second largest traded oil or fat in the world market and palm is the most important crop grown mainly for its oil. Identified varieties of Elaeis guineensis Jacq., with a high oil content and produced through a selective breeding program, are desirable for improving the yield of oil, subsequently enhancing the economic feasibility of using oil palm in various applications, including bio-diesel. We have cloned the gene of biotin carboxylase (accD) from E. guineensis Jacq. This gene encodes a plastidcoded subunit of heteromeric acetyl-CoA carboxylase (ACCase). The cDNA of accD gene (accession number DQ004687) has an open reading frame of 1479 bp that encodes a putative protein of 492 amino acid residues (AAY86362) with a predicted molecular mass of 55.47 kDa. The heteromeric form of ACCase is important as it catalyzes the first committed step of fatty acid synthesis. There is evidence that collectively suggests that the expression of accD in plastids is crucial to the levels of heteromeric ACCase and in turn, to the amount of seed oil in plant. Here we support the hypothesis that the expression level of accD is correlated with the oil palm production by using a semi-quantitative reverse transcription polymerase chain reaction (RT-PCR) and quantitative real-time polymerase chain reaction (real-time PCR) analysis. Moreover, we observed the similar expression profile in nuclear-encoded subunit, biotin carboxylase (accC). This finding represents the genetic background of the expressed genes that correlate to high yield in plant and that ACCase can be used as a marker in the breeding program.

© 2008 Elsevier Ireland Ltd. All rights reserved.

#### 1. Introduction

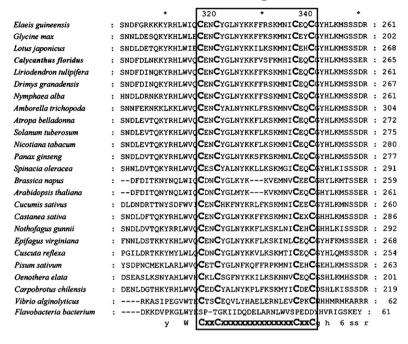
Oil palm (*Elaeis guineensis* Jacq.) is a monocotyledonous plant of the palm family (Arecaceae), with a long life cycle and no natural vegetative reproduction. Oil palm is a species of particular economic importance as it provides one of the most important sources of vegetable oil for use in a wide range of edible products. The global production of palm oil in 2003–2004 was 29.1 million tons, second only in importance to soybean [1]. Because of the oil crisis, uses of palm oil will increase in the future because palm oil has the potential for use in the production of bio-diesel. Oil palm is cultivated in the inter-tropical regions of Asia, Latin America, and Africa. The preferred plants for cultivation are *tenera* hybrids (bearing fruits with shells of intermediate thickness). These

E-mail address: pamornra@yahoo.com (A. Phongdara).

originate from crosses between dura (thick shell) and pisifera (thin shell) types [2,3]. Due to a long selection cycle, poorly characterized genotypes and a high heterogeneity prevalent among hybrids, the use of modern breeding strategies with DNA marker-assisted breeding [4-6] is recommended for the improvement of crop quantities and qualities. There has been considerable work done to establish a genetic map for this species, notably by using RFLP, microsatellite and transposon markers [4,7-11]. In order to speed up the process of crop improvement, one possible marker looks up a key enzyme involved in fatty acid biosynthesis. Biochemical studies have indicated that the acetyl-CoA carboxylase (ACCase) gene product may be involved in the control of the lipid accumulation process [12]. The plant has two distinct forms of this enzyme: the homomeric and the heteromeric forms. In Arabidopsis, a homomeric form of ACCase is encoded by two genes: ACC1 and ACC2 [13]. ACC1 is present in the cytosol and ACC2 is predicted by bioinformatics tool to be present in the plastid [14]. The genome information of four subunits of heteromeric form are accC for a biotin carboxylase (BC), accB for biotin carboxyl carrier protein (BCCP), accA for  $\alpha$ , and accD for  $\beta$  subunits of carboxyl

<sup>\*</sup> Corresponding author at: Center for Genomics and Bioinformatics Research, Faculty of Science, Prince of Songkla University, 15 Karnjanavanit Road, Hat-Yai, Songkhla 90110, Thailand. Tel.: +66 74 288384; fax: +66 74 288384.

#### Zinc-binding Domain



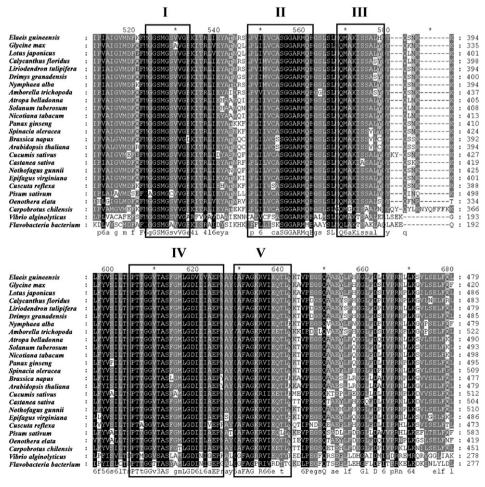


Fig. 1. Conserved zinc-binding domain (CX2CX15CX2C) and I–V motifs at the C-terminal amino acid sequence in plant accDs from oil palm (AAY86362; Elaeis guineensis) with that of other accD's. The species and corresponding accession number are as follows: E. guineensis (AAY86362), Glycine max (AAA80643), Lotus japonicus (BAB33205), Calycanthus floridus (NP\_862763), Liriodendron tulipifera (YP\_740211), Drimys granadensis (YP\_784395), Nymphaea alba (CAF28602), Amborella trichopoda (CAD45116), Atropa belladonna (NP\_783241), Solanum tuberosum (YP\_635648), Nicotiana tabacum (NP\_054508), Panax ginseng (YP\_086975), Spinacia oleracea (CAB88738), Brassica napus (CAA9747), Arabidopsis thaliana (NP\_051068), Cucumis sativus (ABI97426), Castanea sativa (AAS55872), Nothofagus gunnii (AAT79506), Epifagus virginiana (AAA65854),

## Download English Version:

# https://daneshyari.com/en/article/2018140

Download Persian Version:

https://daneshyari.com/article/2018140

<u>Daneshyari.com</u>