

CONSTRUCTION OF FULL-LENGTH pKT9FL CLONE
ENCODES FOR GLUTELIN FROM OIL PALM

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**CONSTRUCTION OF FULL-LENGTH pKT9FL CLONE
ENCODES FOR GLUTELIN FROM
OIL PALM**

By
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A research report submitted in partial fulfillment of
the requirements for the award of the degree of
Bachelor of Science (Biological Sciences)

**DEPARTMENT OF BIOLOGICAL SCIENCES
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RESEARCH REPORT VERIFICATION

Adalah ini diakui dan disahkan bahawa laporan penyelidikan bertajuk:
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DECLARATION

I hereby declare that this thesis entitled Construction of Full-Length pKT9FL Clone Encodes for Glutelin from Oil Palm is the result of my own research except as cited in the references.

Signature



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ABSTRACT

Seed storage protein is classified into three classes, which are globulin, prolamin and glutelin. The protein was utilized during post-germinative period of plant growth. Three different isoforms of glutelin genes were isolated from oil palm kernel (*Elaeis guineensis*, Tenera) in previous research. One of the isoforms consists of two separate fragments, the 3'-end-fragment (900 bp) isolated from cDNA library and the 5'-end fragment (820 bp) isolated by using 5'-RACE method. The full length of this clone is 1621 bp. Thus, this study was aimed to construct a complete full-length sequence of this oil palm glutelin cDNA by joining the two fragments at the overlapping sequence. The 5'-end fragment was first cloned into *Eco*RI and *Sma*I site of pUC19 vector. The 3'-end fragment was generated by PCR and joined with the 5'-end fragment at the *Hpa*I site located at nt. 894 of the glutelin gene. The positive recombinant clone was selected by colony PCR technique and named as pKT9FL. The orientation and the joining site of *Hpa*I in clone pKT9FL were verified by DNA sequencing. This complete sequence of clone pKT9FL consists of 1621 bp that encodes a polypeptide of 469 amino acid. Its 5'and 3'-untranslated region is 52 bp and 152 bp respectively.

ABSTRAK

Protein simpanan biji di klasifikasikan kepada tiga kelas iaitu globulin, prolamin dan glutelin. Protein simpanan biji digunakan ketika fasa selepas percambahan dalam kitaran perkembangan tumbuhan. Dalam kajian terdahulu, tiga isoform gen glutelin yang berbeza telah dipencarkan daripada isirong kelapa sawit (*Elaeis guineensis*, Tenera). Salah satu isoform, terdiri daripada dua serpihan yang berasingan iaitu bahagian 3'-akhiran (900bp) yang dipencarkan daripada perpustakaan cDNA isirong manakala bahagian 5'-akhiran dipencarkan daripada eksperimen 5'-RACE. Saiz lengkap jujukan klon itu ialah 1621 bp. Oleh itu, objektif utama penyelidikan ini adalah untuk membentuk jujukan lengkap klon cDNA gen glutelin kelapa sawit ini dengan menyambungkan kedua-dua serpihan pada kawasan bertindih. Bahagian 5'-akhiran diklonkan pada tapak *Eco*R1 dan *Sma*I pada vektor pUC19. Manakala, bahagian 3'-akhiran pula diamplifikasi dengan PCR dan disambungkan pada bahagian 5'-akhiran pada tapak *Hpa*I (nt. 894) dalam vektor pUC19. Klon rekombinan positif dipilih melalui teknik PCR koloni dan dinamakan pKT9FL. Orientasi jujukan dan tapak penyambungan *Hpa*I dalam klon pKT9FL telah dipastikan melalui penjuzzukan DNA. Jujukan lengkap klon ini ialah 1621 bp yang mengekodkan polipeptida sepanjang 469 amino asid. Saiz kawasan 5' dan 3' yang tidak mengekod ialah masing-masing 52 bp dan 152 bp.