

## Expression of Recombinant *Alcohol Dehydrogenase* in *Escherichia coli* Strain BL21 (DE3) and *In Planta Agrobacterium* Transformation of Tomato Seeds

Mastura Sani<sup>1\*</sup> and Hairul Azman Roslan<sup>2</sup>

<sup>1</sup>Food Technology, School of Engineering and Technology, University College of Technology Sarawak, Sarawak, Malaysia

<sup>2</sup>Faculty of Resource Science and Technology, Universiti Malaysia Sarawak, Sarawak, Malaysia

Received: 26 December 2019, Revised: 13 March 2020, Accepted: 24 April 2020

### Abstract

Alcohol dehydrogenase is an enzyme that is involved in various roles in plant such as in plant development, growth and plant responses to abiotic and biotic stresses. A recombinant *alcohol dehydrogenase 1 (Adh1)* cDNA (*r-msAdh1*) from *Metroxylon sagu* has been previously isolated, containing 20 nucleotides derived from *Elaeis guineensis* at the 5'-end, with a molecular weight of 1.14 kb. The objective of this study is to determine the function of *r-msAdh1* via analyses in prokaryotic and eukaryotic hosts. For expression in prokaryotic system, pET-41a(+) with a 8x His tag at the C terminal was used for *r-msAdh1* protein purification and expression was achieved using IPTG for four to six hours in *Escherichia coli* strain BL21 (DE3) incubated at low temperature. The induced BL21 strain produced a small amount of soluble *r-msAdh1* protein while large amount was present as insoluble aggregates. Subsequently, the *r-msAdh1* cDNA was transformed into tomato seeds (*Solanum lycopersicum* cv. MT1) via *Agrobacterium*-mediated *in planta* transformation. The integration of *r-msAdh1* cDNA and the selectable marker were detected in transformed seedlings, T<sub>0</sub>, using polymerase chain reaction technique. The transformation efficiency was determined to be 33% for *r-msAdh1* cDNA and 46% for the selectable marker. For stability analysis of the transgene, eleven T<sub>1</sub> generation randomly selected from the transgenic T<sub>0</sub> were analyzed for the presence of the cDNA, and all seedlings were found to contain the full length of *r-msAdh1* cDNA. However, out of eleven T<sub>1</sub> transgenic lines produced, only four seedlings were used for expression analysis using the reverse transcriptase PCR (RT-PCR). Two transgenic lines, T<sub>1</sub>9 and T<sub>1</sub>11, were determined to contain *r-msAdh1* cDNA and verified by nucleotide sequencing. Although only a small number of T<sub>1</sub> transgenic seedlings was obtained, this study shows that tomato seeds could be used as a target tissue for *Agrobacterium*-mediated *in planta* transformation primarily because the protocol is easy, rapid and cheaper compared to tissue culture-based method.

**Keywords:** *Alcohol dehydrogenase*, *Metroxylon sagu*, BL21 (DE3), *Agrobacterium tumefaciens*, tomato seeds

DOI.....

\*Corresponding author: Tel.: +60 84 36 75 50, Fax: +60 84 36 73 01  
Email: masturasani88@gmail.com