

UNIVERSITI TEKNOLOGI MARA

**CLONING OF HUMAN *GAMMA GLUTAMYL
TRANSFERASE (GGT)* GENE**

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TABLE OF CONTENTS

	Page
TITLE PAGE	
APPROVAL FORM	
ACKNOWLEDGEMENT	ii
TABLE OF CONTENTS	iii
LIST OF TABLES	vi
LIST OF FIGURES	vii
LIST OF EQUATION	viii
LIST OF ABBREVIATION	ix
ABSTRACT	xi
CHAPTER 1: INTRODUCTION	1
CHAPTER 2: LITERATURE REVIEW	4
2.1 The γ -glutamyl Cycle	4
2.2 <i>GGT</i> Structure	6
2.3 <i>GGT</i> Localization	7
2.4 Clinical Impact	
2.4.1 Tumors growth and <i>GGT</i> levels	7
2.4.2 Alcohol abuse and <i>GGT</i> levels	9
2.4.3 <i>GGT</i> Deficiency	9
2.5 Polymerase Chain Reaction (PCR)	10
2.5.1 Principles of PCR	11
2.6 Principle of Gel Electrophoresis	14
2.7 TA Cloning	15
2.8 Transformation	16
2.9 DNA Sequencing	17
2.9.1 The Sequencing Reaction	17
2.9.2 Separation of Molecule	19

ABSTRACT

The γ -glutamyl transferase (*GGT*) plays an important role in the process of catalyzing the degradation of glutathione. *GGT* is widely used as a marker in preneoplastic lesions in the liver during chemical carcinogenesis. In this study, primers were designed without the restriction sites to investigate the success of cloning. The aim of this study was to clone *GGT* gene in plasmid by means of TA cloning. Methods used in this study were purification of human liver cDNA, PCR for gene amplification, TOPO TA cloning, transformation into *E.coli* and DNA sequencing. PCR was conducted under different annealing temperatures using Takara PCR Thermal Cycler Dice™ Gradient model to determine the optimal annealing temperature. TOPO TA cloning process with high cloning efficiency was completed in 5 minutes. Transformation was done using the One Shot® Chemical Competent *E.coli*. DNA sequencing was also performed using the plasmid samples from transformation process to validate the insert. Plasmid extraction yielded 168 ng/μl of liver cDNA. The PCR based method obtained an optimal annealing temperature for *GGT* amplification, which was at 47°C. Transformation was a success by using *E.coli* as a host where a 10 ml of LB broth was used and incubated at 37°C for 16 hours. Cloning of the *GGT* gene is important as it opens ways for further studies to look into its association with cancer and help in development of medication that prevents the progression of tumors. There is no specific treatment for patients with *GGT* deficiency and thus, researchers may develop protein-based drugs for them. *GGT* cloning can help in the establishment of gene collection in DNA libraries.

CHAPTER 1

INTRODUCTION

The human genome consists of high, medium, and low copy of repetitive DNA and single-copy DNA. The human *gamma glutamyl transferase (GGT)* cDNA was assigned to human chromosome 22 using *in situ* hybridization and at least four copies were identified in man (Collins *et. al.*, 1997). As reported by Sakamuro *et. al.*, (1988), *GGT* consists of 2 peptide chains, heavy and light, each composed of 351 and 189 amino acids, respectively. The peptide chains contain the catalytic site of *GGT*, which are encoded on a single mRNA (Courtay *et. al.*, 1994). *GGT* is a membrane-bound glycoprotein located on the outer surface of the cell membrane.

Breakdown of glutathione is catalyzed by *GGT*, which takes place by transfer of gamma glutamyl moiety of glutathione to a number of acceptors. Suggestion was made that transpeptidase mediates amino acid transport by the reaction of the gamma glutamyl cycle. This cycle is one of the pathways that functions in amino acid transport. Gamma glutamyl amino acids are formed in or on the cell membrane by *GGT* where the gamma glutamyl amino acids are then translocated into the cell before free amino acids are released (Allison & Meister, 1981).