

**UNIVERSITI TEKNOLOGI MARA**

**DEVELOPMENT OF PCR-BASED METHOD FOR  
DETECTION OF THE POLYMORPHISM OF N-  
METHYL-D-ASPARTATE RECEPTOR GENE**

**FARRAH IDAYU ABDUL MUTALIB**

**Dissertation submitted in partial fulfillment of the requirements for  
the Bachelor of Pharmacy (Hons)**

Faculty of Pharmacy

November 2009

## ACKNOWLEDGMENT

I am very grateful and thank a lot to Almighty Allah S.W.T. in giving me patience and strength to complete this final project.

Firstly, I would like to thank to my family who is very understanding and consistently support during doing this project.

I would like to take this opportunity to express my gratitude and appreciation to all pharmacy staff and lecturers especially to my supervisor, Professor Abu Bakar bin Abdul Majeed and also my co-supervisor, En. Mohd Nazif bin Hj. Samat@Darawi for their continuous guidance, invaluable advices, constructive comments and patience during the course of this project.

I wish to express my deepest appreciation to my project coordinator, Dr. Kalavaty for her guidance and information throughout this project.

Not forgotten to all laboratory staff at Life Sciences Research Laboratory, Faculty of Pharmacy for their technical assistance and valuable comments.

Last but not least, I like to extend my heartiest appreciation to all my friends who have contributed towards the success of this project.

Thank you very much.

## TABLE OF CONTENTS

	Page
TITLE PAGE	
APPROVAL SHEET	
ACKNOWLEDGEMENT	iii
TABLE OF CONTENTS	iv
LIST OF TABLES	vi
LIST OF FIGURES	vii
LIST OF ABBREVIATIONS	viii
ABSTRACT	ix
CHAPTER ONE (INTRODUCTION)	1
1.1 Background of study	1
1.2 Justification	2
1.2.1 Problem statement	2
1.2.2 Study output	3
1.3 Objectives of the study	3
1.4 Hypothesis	3
CHAPTER TWO (LITERATURE REVIEW)	4
2.1 Polymerase Chain Reaction	4
2.2 Genetic Variants	5
2.2.2 Single Nucleotide Polymorphism (SNP)	6
2.3 Allele-Specific PCR	7
2.4 N-Methyl-D-Aspartate (NMDA)	8
2.4.1 Neurotransmitter	8
2.4.2 The NR2 subunits	10
2.5 NMDA receptor and neurodegeneration	13
2.5.1 Diseases	15
CHAPTER THREE (METHODOLOGY)	17
3.1 Methodology	17
3.2 Primer design	17
3.2.1 Methods of primer design	18
3.2.2 Designing allele-specific primer and common primer	20
3.2.3 Design primers for sequencing	25
3.3 Samples	27
3.4 Methods for DNA extraction from blood using DNeasy Kit	27
3.4.1 Apparatus	28
3.4.2 Substances	28

## ABSTRACT

The purpose of this study was to design the allele-specific polymerase chain reaction for detection of the polymorphism of N-Methyl-D-Aspartate receptor gene and establish the frequencies of the selected genetic variants amongst Chinese ethnic in Malaysia. Single nucleotide polymorphism (SNP) of GRIN2B, rs1805247 was selected to be studied. The allele-specific primers were designed by using Oligo Explorer software and the unique sequence of the primers has been sent to First Base Laboratory Sdn. Bhd to be synthesised. The primers were Rcm\_2\_NMDAR, Fmt\_2\_NMDAR and Fwt\_2\_NMDAR. The most suitable annealing temperature for these primers was found to be 67°C where the intended band appeared and matched with DNA sequencing results. In this study, the most common genotype was G/A (55.8%) and A was the predominant allele (62.8%). It can be concluded that the method has been successfully developed because it matched with gold standard validation method. The assay would be useful for detection of the genetic polymorphism of NMDAR rs1805247 because variations that occurred in analysed samples are more than 1%.

# CHAPTER 1

## INTRODUCTION

### 1.1.1 Background of study

Polymerase chain reaction (PCR) is the key technique used in molecular genetics for amplifying deoxyribonucleic acid (DNA) sequences as template for replication in vitro by separating the DNA into two strands. By using PCR methods, we can amplify the DNA by more than a million times to produce more copies of the DNA piece. Besides, we can perform a wide array of genetic manipulations by using this method.

In this study, allele specific polymerase chain reaction (AS-PCR) was developed to genotype DNA sample for genetic polymorphism of N-Methyl-D-Aspartate (NMDAR2B) or glutamate receptor ionotropic N-Methyl-D-Aspartate 2B (GRIN2B). Allele specific primers were designed to detect the mutation in selected variant in GRIN2B. In this PCR method, only a single specific primer was used in combination with a common primer to amplify the target sequence. The 3' end of the primer designed should be complementary to the target DNA sequence.