



UNIVERSITI PUTRA MALAYSIA

**GENE POLYMORPHISMS OF ANGIOTENSIN-CONVERTING ENZYME,
ANGIOTENSIN TYPE 1 RECEPTOR AND α -ADDUCIN ASSOCIATED WITH
RENIN ANGIOTENSIN-ALDOSTERONE SYSTEM IN MALAYSIAN
END-STAGE RENAL DISEASE PATIENTS**

AISYAH BINTI ALI

FPSK(m) 2012 36



CC
©

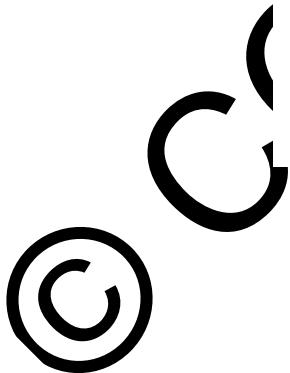
Thesis Submitted to the School of Graduate Studies,
Universiti Putra Malaysia, in Fulfilment of the
Requirements for the Degree of Master of Science

June 2012

COPYRIGHT

All material contained within the thesis, including without limitation text, logos, icons, photographs and all other artwork, is copyright material of Universiti Putra Malaysia unless otherwise stated. Use may be made of any material contained within the thesis for non-commercial purposes from the copyright holder. Commercial use of material may only be made with the express, prior, written permission of Universiti Putra Malaysia.

Copyright © Universiti Putra Malaysia



DEDICATION

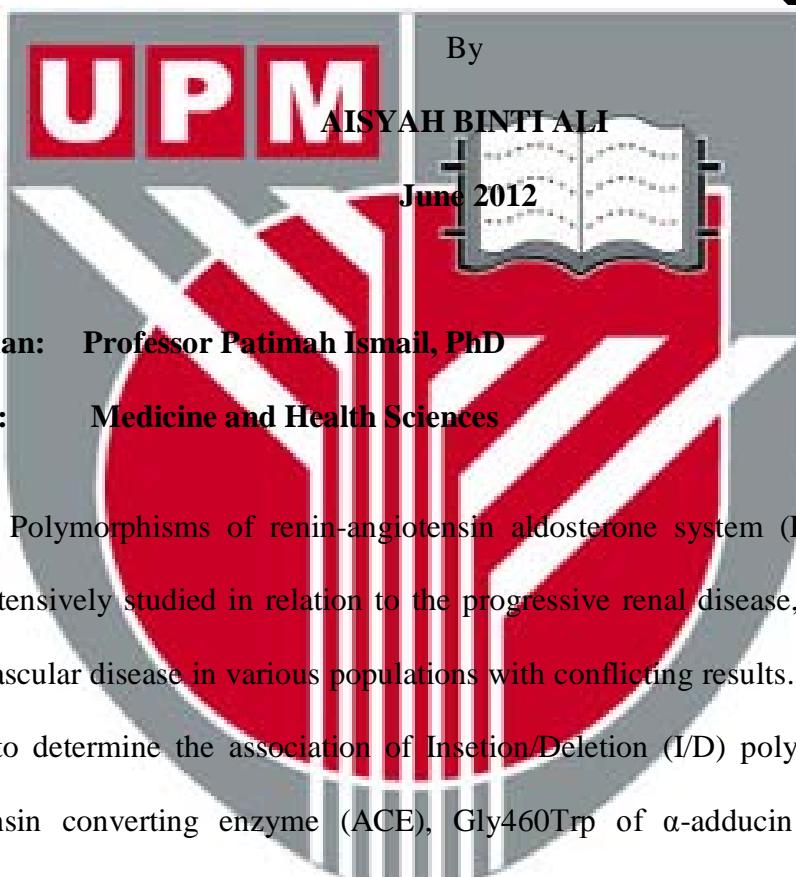


UPM

CC

Abstract of thesis presented to the Senate of Universiti Putra Malaysia
in fulfilment of the requirement for the degree of Master of Science

**GENE POLYMORPHISMS OF ANGIOTENSIN-CONVERTING ENZYME,
ANGIOTENSIN TYPE 1 RECEPTOR AND α -ADDUCIN ASSOCIATED WITH
RENIN ANGIOTENSIN-ALDOSTERONE SYSTEM IN MALAYSIAN
END-STAGE RENAL DISEASE PATIENTS**



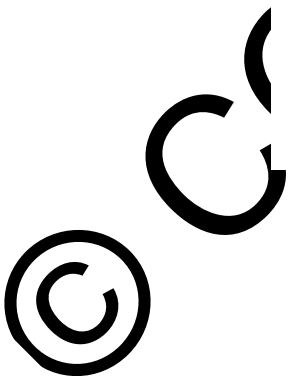
Chairman: Professor Patimah Ismail, PhD

Faculty: Medicine and Health Sciences

Genetic Polymorphisms of renin-angiotensin aldosterone system (RAAS) genes has been extensively studied in relation to the progressive renal disease, hypertension and cardiovascular disease in various populations with conflicting results. The present study sought to determine the association of Insertion/Deletion (I/D) polymorphisms of the angiotensin converting enzyme (ACE), Gly460Trp of α -adducin and A1166C of angiotensin type 1 receptor(AT1R) of RAAS genes in Malaysian end stage renal subjects. A total of 380 subjects consisted of 190 end stage renal disease (ESRD) patients and 190 unrelated healthy individuals were recruited in this study. Genotypes of RAAS gene polymorphisms were determined using mutagenically separated PCR and PCR-RFLP method. There was significant difference ($p<0.05$) found in age, systolic blood pressure (SBP), creatinine level, triglycerides and total cholesterol between the



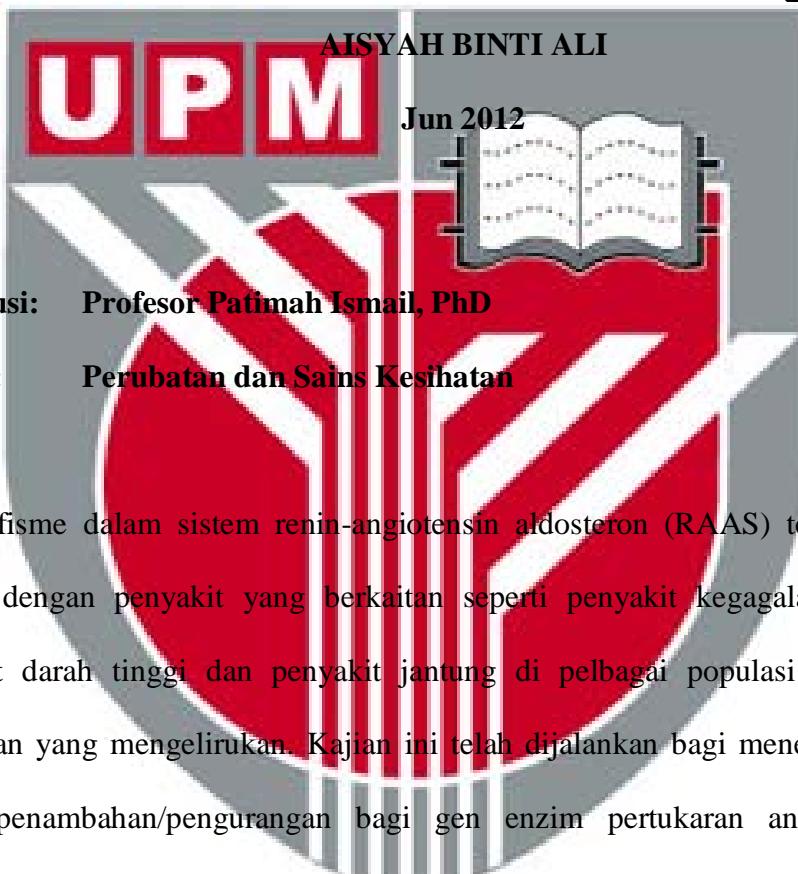
ESRD and control subjects. There was statistically significant differences ($p<0.05$) were found in I/D polymorphisms of ACE and Gly460Trp polymorphism of α -adducingene and no significant difference ($p>0.05$) was found in A1166C polymorphism of AT1R genes between the ESRD and control subjects. The findings of this study indicate that I/D polymorphisms of the ACE gene and Gly460Trp polymorphism of α -adducingene are a useful marker and are likely to play a major role in determining genetic susceptibility to Malaysian ESRD subjects.



Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia
Sebagai memenuhi keperluan untuk Ijazah Master Sains

**POLIMORFISME GEN BAGI ENZIM PERTUKARAN ANGIOTENSIN,
RESEPTOR JENIS 1 ANGIOTENSIN DAN . -ADDUCIN BERKAITAN
DENGAN SISTEM RENIN ANGIOTENSIN ALDOSTERON DIKALANGAN
PESAKIT BUAH PINGGANG PERINGKAT AKHIR DI MALAYSIA**

Oleh



Pengerusi: Profesor Patimah Ismail, PhD

Fakulti: Perubatan dan Sains Kesihatan

Polimorfisme dalam sistem renin-angiotensin aldosteron (RAAS) telah dikaji secara meluas dengan penyakit yang berkaitan seperti penyakit kegagalan buahpinggang, penyakit darah tinggi dan penyakit jantung di pelbagai populasi dengan pelbagai keputusan yang mengelirukan. Kajian ini telah dijalankan bagi menentukan hubungan antara penambahan/pengurangan bagi gen enzim pertukaran angiotensin (ACE), polimorfisme Gly460Trp bagi gen α -adducin dan polimorfisme A1166C bagi gen Angiotensin type 1 receptor (AT1R) di kalangan pesakit buah pinggang peringkat akhir di Malaysia. Seramai 380 orang telah terlibat dalam kajian ini dimana terdiri daripada 190 orang pesakit buah pinggang peringkat akhir (ESRD) and 190 orang yang sihat sebagai kawalan. Genotip polimorfisme bagi gen RAAS telah ditentukan menggunakan kaedah tindak balas rantaian polymerase polimorfisme panjang jalur terpotong (PCR-



RFLP), tindak balas rantaian polimerase mutagenic (MS-PCR) dan tindak balas rantaian polimerase Hot-Start (Hot-Start PCR). Perbezaan signifikan telah dijumpai dalam umur, tekanan darah sistol, kretinin, trigliseride, dan jumlah kolesterol apabila dibandingkan antara pesakit buah pinggang peringkat akhir dengan yang normal. Perbezaan signifikan ($p<0.05$) telah dijumpai dalam polimorfisme penambahan/pengurangan bagi gen enzim pertukaran angiotensin dan polimorfisme Gly460Trp bagi gen α -adducin gene apabila dibandingkan antara pesakit buah pinggang peringkat akhir dengan yang normal. Tiada perbezaan signifikan ($p>0.05$) dijumpai dalam polimorfisme A1166C bagi gen AT1R apabila dibandingkan dengan pesakit buah pinggang dengan yang normal. Penemuan dalam kajian ini telah menunjukkan bahawa polimorfisme penambahan/pengurangan bagi gen enzim pertukaran angiotensin dan polimorfisme Gly460Trp bagi gen α -adducin merupakan penanda yang berguna dan memainkan peranan besar dalam menentukan kestabilan genetic terhadap pesakit buah pinggang terakhir di Malaysia.



ACKNOWLEDGEMENTS

First and foremost, I would like to extend my deep gratitude towards my dearest Supervisor, Professor Dr. Patimah Ismail for her generous guidance, kindness, helpful and valuable support to complete my dissertation. I would like to express my heartfelt appreciation to my co-supervisors; Dr. Srikumar Chakravarti and Dr Christopher Lim Thiam Seong for their helpfulness, wise guidance and incessant support throughout the study.

Apart from that, I would like to express my full gratitude to all my friends; Dr R. Vasudevan, Rusni, Mimi, and GRG group members who greatly helped me throughout the project and for their great support, encouragement, having fun and entertainment and for their fruitful advices and sweet full memories.

I would like to express my deepest gratitude to my greatest parents and family for their endless encouragement, patience and sacrifices, which had helped me in all my undertakings and the completion of the project.

I would like to thank to all the staff in the Department of Biomedical Sciences and others for their kind co-operation. I would like to acknowledge the Ministry of Science, Technology and Environment (MOSTE), RUGS project number 91104, for their full funding to complete this project.

I certify that a Thesis Examination Committee has met on 25 June 2012 to conduct the final examination of Aisyah binti Ali on her thesis entitled "Gene Polymorphisms of Angiotensin-Converting Enzyme, Angiotensin Type 1 Receptor and α -Adducin Associated with Renin Angiotensin-Aldosterone System in Malaysian End-Stage Renal Disease Patients" in accordance with the Universities and University Colleges Act 1971 and the Constitution of the Universiti Putra Malaysia [P.U.(A) 106] 15 March 1998. The Committee recommends that the student be awarded the Master of Science.

Members of the Thesis Examination Committee were as follows:

Fauziah binti Othman, PhD

Professor

Faculty of Medicine and Health Sciences

Universiti Putra Malaysia

(Chairman)

Cheah Yoke Kqueen, PhD

Associate Professor

Faculty of Medicine and Health Sciences

Universiti Putra Malaysia

(Internal Examiner)

Abdah binti Md Akim, PhD

Senior Lecturer

Faculty of Medicine and Health Sciences

Universiti Putra Malaysia

(Internal Examiner)

Ammu K. Radhakrishnan, PhD

Professor

Faculty of Medicine and Health Sciences

International Medical University

Kuala Lumpur

(External Examiner)



SEOW HENG FONG, PhD

Professor and Deputy Dean

School of Graduate Studies

Universiti Putra Malaysia

Date: 23 January 2013

This thesis was submitted to the Senate of Universiti Putra Malaysia and has been accepted as fulfillment of the requirement for the degree of Masters of Science. The members of the Supervisory Committee were as follows:

Patimah Ismail, PhD

Professor

Faculty of Medicine and Health Sciences

Universiti Putra Malaysia

(Chairman)

Dr Srikumar Chakravarthi, MD

Senior Lecturer

Department of Pathology

International Medical University

(Member)

Dr Christopher Lim Thiam Seong, FAMS

Senior Lecturer

Faculty of Medicine and Health Sciences

Universiti Putra Malaysia

(Member)

BUJANG BIN KIM HUAT, PhD

Professor and Dean

School of Graduate Studies

Universiti Putra Malaysia

Date:



DECLARATION

I declare that the thesis is my original work except for quotation and citations which have been duly acknowledged. I also declare that it has not been previously, and is not concurrently, submitted for any other degree at Universiti Putra Malaysia or at any other institution.

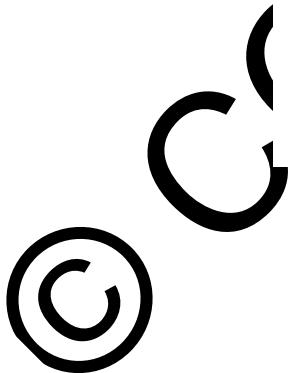


TABLE OF CONTENTS

	Page
DEDICATION	ii
ABSTRACT	iii
ABSTRAK	iv
ACKNOWLEDGEMENT	viii
APPROVAL	ix
DECLARATION	x
LIST OF TABLES	xiv
LIST OF FIGURES	xv
LIST OF ABBREVIATIONS	xvi
CHAPTER	
1.0 INTRODUCTION	
1.1 Background	1
1.2 Problem Statements	3
1.3 Significant of Study	4
1.4 Hypothesis	5
1.5 Objective	5
1.5.1 Main Objective	5
1.5.2 Specific Objectives	6
2.0 LITERATURE REVIEW	
2.1 End Stage Renal Disease (ESRD)	7
2.1.1 Prevalence of ESRD	8
2.1.2 Risk factors of ESRD	9
2.1.2.1 Environmental Factor	9
2.1.2.2 Genetic factors of ESRD	11
2.1.3 Pathophysiology of ESRD	12
2.2 Renin Angiotensin-Aldosterone System (RAAS)	13
2.2.1 Mechanism of RAAS	14
2.2.2 Relationship between RAAS and ESRD	15
2.3 Genetic Polymorphism	17



2.4	Angiotensin Converting Enzyme (ACE) Gene	18
2.5	Angiotensin II Type 1 Receptor (AT1R) Gene	19
2.6	Alpha-Adducin Gene	20
2.7	Genetic Association Analysis	21
	2.7.1 Techniques to Detect Genetic Polymorphism	21
	2.7.1.1 Hot Start PCR	22
	2.7.1.2 Mutagenic Separated PCR	22
	2.7.2 Restriction Fragment Length Polymorphism (RFLP)	23
2.8	Agarose Gel Electrophoresis	24
2.9	DNA Sequencing	24
2.10	Statistical Package for the Social Sciences (SPSS)	25
3.0	METHODOLOGY	
3.1	Study Design	26
3.2	Ethical Approval	26
3.3	Sample Size	27
3.4	Sampling	28
	3.4.1 Case Subjects	28
	3.4.2 Control Subject	29
	3.4.3 Questionnaire	30
	3.4.4 Sampling Method	30
3.5	Genomic DNA Extraction	30
	3.5.1 Quantification of Genomic DNA	31
	3.5.2 Purification of Impurity DNA	31
	3.5.3 Agarose Gel Electrophoresis for Genomic DNA	32
3.6	Polymerase Chain Reaction (PCR)	32
3.7	Optimization of PCR	32
3.8	Confirmation of PCR Product	33
3.9	Determination of ACE genotyping	33
	3.9.1 Conventional PCR	33
3.10	Determination of AT1R genotyping	36



3.10.1	Hot-Start PCR-RFLP	36
3.11	Determination of Alpha-Adducin genotyping	37
3.11.1	Mutagenic Separated PCR	37
3.12	Agarose Gel Electrophoresis	38
3.13	Staining and Visualization	39
3.14	Purification of PCR products and DNA sequencing	40
3.15	Statistical Analysis	40
4.0	RESULT	
4.1	Clinical Characteristic of All subjects	42
4.2	Insertion/deletion (I/D) polymorphism of ACE Gene	49
4.2.1	PCR Amplification	49
4.2.2	Genotype and Allele Frequencies Analysis	50
4.3	A1166C polymorphism of AT1R Gene	52
4.3.1	PCR Amplification	52
4.3.2	Genotype and Allele Frequencies Analysis	54
4.4	Gly460Trp polymorphism of α -Adducin Gene	56
4.4.1	PCR Amplification	56
4.4.2	Genotype and Allele Frequencies Analysis	57
5.0	DISCUSSION	
5.1	Clinical Characteristic of All subjects	59
5.2	Genotype and Allele Frequencies Analysis	61
6.0	CONCLUSION AND FUTURE RECOMMENDATIONS	65
REFERENCES		66
APPENDICES		75
BIODATA OF STUDENT		95
LIST OF PUBLICATION		96