



UNIVERSITI PUTRA MALAYSIA

**HAEMATOLOGY, GENETICS AND MOLECULAR EPIDEMIOLOGY OF
DELETIONAL ALPHA THALASSAEMIA IN MALAYSIA**

ALIZA BINTI MOHD YACOB

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DELETIONAL ALPHA THALASSAEMIA IN MALAYSIA**



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By

ALIZA BINTI MOHD YACOB

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Chairman: Professor Elizabeth George, M.D

Faculty: Medicine and Health Sciences

Alpha thalassaemia is the most common autosomal recessive single gene disorder in Southeast Asia, encountered in increasing numbers all over the world and heterogenous. Deletions of all four alpha thalassaemia genes (--) result in fetuses mostly die before or shortly after birth known as Hb Barts hydrops foetalis syndrome. It also caused serious maternal complications in pregnancies in which without medical care half were estimated to die. Deletions of 3 alpha thalassaemia genes (-/- α) is known as HbH disease with moderate anaemia, usually was similarly seen in the Malays and Chinese. In contrast, antenatal diagnosis for alpha thalassaemia reported Hb Barts hydrops foetalis (--) mostly in the Chinese. Thalassaemia studies among blood donors of 91.3% Malays found out that 30% were anaemic with all donors had a negative H-inclusion test, which is usually positive with double deletions (-/ $\alpha\alpha$). Malaysia is a multiethnic country with mix marriages and 4.5% of Chinese-Malaysian is carriers of α^0

thalassaemia (--). Therefore the purpose of this study was to determine the current deletional alpha thalassaemia status and deletional alpha thalassaemia burden in our population. Changes in deletional alpha thalassaemia gene frequency may change the deletional alpha thalassaemia burden. In this study haematology, genetics and molecular epidemiology of deletional alpha thalassaemia were looked at and carrier detection of common deletions was carried out. This will identify phenotypic characteristics, type of deletions and genotypes present, determine prevalence, estimate disease burden of deletional alpha thalassaemia and determine foetal genotype from pregnancies at risk of deletional Hb Barts hydrops foetalis in the country.

Standard haematology protocol, DNA studies and statistical analysis of qualitative and quantitative were applied. A cross sectional study was carried out on 405 samples. These were 238 EDTA blood from blood donors from Universiti Putra Malaysia, 15 DNA of 5 Hb Barts hydrops foetalis (---) and 10 alpha thalassaemia spouses (-/-αα) from Universiti Malaya Medical Centre, 72 EDTA blood from HbH disease (-/-α) families from Institute for Medical Research and 80 DNA from blood donors from National Blood Bank. Blood count/blood film, HPLC, haemoglobin electrophoresis and multiplex PCR for detecting the 5 most common gene deletions were carried out. These are the common --^{SEA} in Southeast Asia, the ethnic origins --^{THAI} and --^{FIL}, the world common single deletions -α^{3.7} and -α^{4.2}. Multiplex PCR using validated primers was developed using samples from families of HbH disease, detected the --^{SEA}, --^{THAI}, --^{FIL} and -α^{3.7} giving 5 genotypes. This was compared with conventional method using samples from National Blood Bank and gave 100% accuracy, sensitivity and specificity

for --^{SEA} . --^{FIL} , --^{THAI} and $-\alpha^{3.7}$ were also detected. In blood donors from Universiti Putra Malaysia, $\text{MCV} < 80 \text{ fL}$ was identified in 16.5% (with exclusion of Indians), deletional α -thalassaemia prevalence was 7.4% (3.5% double and 3.9% single deletions), equally distributed among Malays and Chinese. The most common double deletion detected was --^{SEA} and $-\alpha^{3.7}$ for single followed by $-\alpha^{4.2}$. The prevalence was 5.3% in the Malays with the most common was $-\alpha^{3.7}$ (4.3%), 8.5% in the Chinese with 5.4% --^{SEA} indicating an increase from 4.5% α^0 thalassaemia ($--/-$) and 0.4% in others. All the --^{SEA} were with $\text{MCV} \leq 68 \text{ fL}$ and $\text{MCH} < 22 \text{ pg}$. The projected number of pregnancies each year at risk of deletional Hb Barts hydrops foetalis syndrome and HbH disease are 30 and 120 in Malays, 250 and 150 in Chinese, 640 of both and on average 600 and 669 respectively. A typical α -thalassaemia phenotype was observed in HbH disease, and carriers of α^0 and α^+ thalassaemia. Real-time quantitative PCR (TaqMan[®]) with validated primers was optimized for --^{SEA} . Fetal genotype values of $\alpha^{\text{SEA}}\text{-BR}_N$ were 0.6 to 1.6 for $(\text{--}^{\text{SEA}}/\text{--}^{\text{SEA}})$, 0.2 to 0.5 for $(\text{--}^{\text{SEA}}/\alpha\alpha)$ and 0 for $(\alpha\alpha/\alpha\alpha)$ giving distinctly different values. The methods used were simple, reliable and useful in monitoring allele status. Identified carriers can be counseled and DNA from pregnancies at risk for $(\text{--}--)$ can be analysed. These will help in increasing awareness among carriers, bringing down deletional α -thalassaemia births and eventually lower the disease burden in Malaysia.

Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai
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HEMATOLOGI, GENETIK DAN EPIDEMIOLOGI MOLEKUL ALPHA THALASEMIA DELESI DI MALAYSIA

Oleh

ALIZA BINTI MOHD YACOB

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Pengerusi: Profesor Elizabeth George, M.D

Fakulti: Perubatan dan Sains Kesihatan

Alfa talasemia adalah kecacatan gen tunggal autosomal resesif paling lazim berlaku di Asia Tenggara, bilangannya semakin meningkat di seluruh dunia dan heterogenus. Delesi kesemua empat gen alfa talasemia (--) mengakibatkan fetus kebanyakannya meninggal dunia sebelum atau sejurus selepas dilahirkan dan dikenali sebagai sindrom Hb Barts hydrops foetalis. Ia juga menyebabkan komplikasi maternal yang serius pada kandungan di mana tanpa penjagaan perubatan separuh dijangka meninggal dunia. Delesi 3 gen alfa talasemia (-/- α) dikenali sebagai penyakit HbH dengan anemia sederhana, selalunya dilihat samarata pada kedua-dua orang Melayu dan Cina. Walaubagaimanapun, diagnosis antenatal untuk penyakit alfa talasaemia melapurkan sindrom Hb Barts hydrops foetalis (--) kebanyakannya pada orang Cina. Kajian talasemia dikalangan penderma darah yang 91.3% Melayu mendapat 30% adalah anemik dan kesemua penderma adalah negatif ujian H-inklusi, yang selalunya positif

dengan 2 delesi ($--/\alpha\alpha$). Malaysia adalah negara berbilang kaum dengan perkahwinan campur dan 4.5% daripada orang Cina-Malaysia adalah pembawa α^0 talasemia ($--$). Oleh sebab itu tujuan kajian ini adalah untuk menentukan status terkini penyakit alfa talasemia delesi dan beban penyakit alfa talasemia delesi dalam populasi kita. Perubahan pada frekuensi gen penyakit alfa talasemia delesi boleh mengubah beban penyakit alfa talasemia delesi. Dalam kajian ini hematologi, genetik dan epidemiologi molekul penyakit alfa talasemia delesi dilihat dan pengesanan pembawa delesi lazim dijalankan. Ini akan mengenalpasti sifat fenotip, jenis delesi dan genotip yang hadir, menentukan prevalen, menganggar beban penyakit alfa talasemia delesi dan menentukan genotip fetus pada kandungan berisiko sindrom Hb Barts hydrops foetalis delesi di negara ini.

Protokol hematologi piawai, kajian DNA dan analisis statistik kualitatif dan kuantitatif diaplikasikan. Kajian rentas dijalankan ke atas 405 sampel. Ini adalah 238 darah EDTA penderma darah dari Universiti Putra Malaysia, 15 DNA daripada 5 sindrom Hb Barts hydrops foetalis ($--/-$) dan 10 pasangan alpha talasaemia ($--/\alpha\alpha$) dari Pusat Perubatan Universiti Malaya, 72 darah EDTA ahli keluarga penyakit HbH ($--/\alpha$) dari Institut Penyelidikan Perubatan dan 80 DNA penderma darah dari Tabung Darah Negara. Kiraan darah/filem darah, HPLC, elektroforesis hemoglobin dan multiplex PCR untuk mengesan 5 delesi gen lazim dijalankan. In adalah delesi lazim $--^{SEA}$ di Asia Tenggara, mengikut asal etnik $--^{THAI}$, $--^{FIL}$, tunggal lazim di dunia $-\alpha^{3.7}$ dan $-\alpha^{4.2}$. PCR Multiplex menggunakan primer yang disahkan dibangunkan menguna sampel ahli keluarga pesakit

penyakit HbH, mengesan --^{SEA} , --^{THAI} , --^{FIL} dan $-\alpha^{3.7}$ dan memberi 5 genotip. Ini dibandingkan dengan kaedah konvensional menggunakan sampel dari Tabung Darah Negara dan memberi 100% betul, sensitif dan spesifik untuk --^{SEA} , --^{FIL} , --^{THAI} dan $-\alpha^{3.7}$ juga dikesan. Pada penderma darah Universiti Putra Malaysia, $\text{MCV} < 80 \text{ fL}$ dikenalpasti pada 16.5% (eksklusif orang India), prevalen α -talasemia delesi ialah 7.4% ($3.5\% \text{ } \text{--}^{\text{SEA}}$ dan 3.9% delesi tunggal) dan ditaburkan samarata diantara orang Melayu dan Cina. --^{SEA} adalah delesi kembar dan $-\alpha^{3.7}$ adalah delesi tunggal paling lazim dikesan diikuti $-\alpha^{4.2}$. Prevalen ialah 5.3% pada orang Melayu dengan $-\alpha^{3.7}$ (4.3%) paling lazim, 8.5% pada orang Cina dengan 5.4% --^{SEA} iaitu menunjukkan peningkatan daripada 4.5% α^0 talasemia ($--/$) dan 0.4% pada bangsa lain. Semua --^{SEA} mempunyai $\text{MCV} \leq 68 \text{ fL}$ dan $\text{MCH} < 22 \text{ pg}$. Unjuran kandungan setiap tahun berisiko sindrom Hb Barts hydrops foetalis delesi dan penyakit Hb H delesi ialah 30 dan 120 pada orang Melayu, 250 dan 150 pada orang Cina, 640 pada keduanya dan secara purata 600 dan 669. Fenotip tipikal α -talasemia diperhatikan dari penyakit HbH, dan pembawa α^0 dan α^+ talasaemia. Kuantitatif PCR Real-time (TaqMan) menggunakan primer yang disahkan dioptima untuk (--^{SEA}). Nilai genotip fetus $\alpha^{\text{SEA}}\text{-BR}_N$ ialah 0.6 ke 1.6 untuk ($\text{--}^{\text{SEA}}/\text{--}^{\text{SEA}}$), 0.2 ke 0.5 untuk ($\text{--}^{\text{SEA}}/\alpha\alpha$) dan 0 untuk ($\alpha\alpha/\alpha\alpha$) memberi perbezaan nilai yang jelas. Kaedah-kaedah ujian yang digunakan adalah ringkas, boleh diharap dan berguna untuk monitor status alel. Pembawa yang dikenalpasti dapat diberi kaunseling dan DNA dari kandungan berisiko ($--/-$) alfa talasemia boleh dianalisis. Ini membantu meningkatkan kesedaran dikalangan pembawa, menurunkan kelahiran alfa talasaemia delesi dan seterusnya mengurangkan beban penyakit ini di Malaysia.

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Elizabeth George, M.D

Professor

Faculty of Medicine and Health Science

Universiti Putra Malaysia

(Chairman)

Abdul Manaf Ali, PhD

Professor

Faculty of Agriculture and Biotechnology

Universiti Sultan Zainal Abidin Malaysia

(Member)

Zubaidah Zakaria, M.D

Ketua Pusat Penyelidikan Kanser

Ketua Unit Hematologi

Institut Penyelidikan Perubatan

(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 quotations 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.

ALIZA BINTI MOHD YACOB

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TABLE OF CONTENTS

	Page
ABSTRACT	ii
ABSTRAK	v
ACKNOWLEDGEMENTS	viii
APPROVAL	ix
DECLARATION	xi
LIST OF TABLES	xii
LIST OF FIGURES	xiii
LIST OF ABBREVIATIONS AND SYMBOLS	xvi
CHAPTER	
1 INTRODUCTION	1
2 LITERATURE REVIEW	
2.1 The haemoglobin molecule	8
2.2 Molecular genetics of the human α -globin gene cluster	12
2.3 Epidemiology of α -thalassaemia	16
2.4 Clinical description and laboratory detection of α -thalassaemia	18

3 MATERIALS AND METHODS

3.1	Blood sample for analysis	28
3.1.1	Subgroup samples	28
3.1.2	Comparison samples	30
3.1.3	Control samples	30
3.1.4	Inclusion and Exclusion Criteria	31
3.2	BHES protocol, DNA testing and Statistical analysis	32
3.3	BHES protocol	35
3.3.1	Blood cell count	35
3.3.2	Blood film preparation	37
3.3.3	Blood film staining	38
3.3.4	H-inclusion test	39
3.3.5	VARIANT™ hemoglobin test	40
3.3.6	Haemoglobin electrophoresis	44
3.4	DNA extraction	46
3.4.1	DNA yield and purity estimation	47
3.5	Multiplex PCR	49
3.5.1	Preparation of primers	50
3.5.2	Preparation of multiplex PCR reactions	53
3.5.3	Optimization and development of multiplex assays	54
3.5.4	Separation and identification of multiplex PCR products	56
3.6	Conventional routine method	57
3.7	Real-Time Quantitative PCR	57
3.7.1	Preparation of primers and TaqMan® probes	60
3.7.2	Preparation of human β-actin	62
3.7.3	Preparation of standards	63
3.7.4	Preparation of real time Q-PCR reactions	64

3.7.5	Minimizing DNA contaminants	70
3.7.6	Analysis of Data	71
3.8	DNA Sequencing	73
3.9	Statistical Analysis	74

4 RESULTS

4.1	Identification of rare and common α -thalassaemia deletions and α -globin gene triplication	77
4.1.1	Heterogeneity of α -thalassaemia	77
4.1.2	Genotypes of α -thalassaemia	79
4.1.3	Sensitive detection for rare mutations	81
4.1.4	Correct detection	81
4.2	Haematology phenotypic characteristics	82
4.3	Comparison between test method and routine method	83
4.3.1	Performance of test method	83
4.3.2	Evaluation of test method	84
4.4	Detection of five common α -thalassaemia deletions in blood donors	85
4.4.1	Presumptive identification of thalassaemia trait	85
4.4.2	Double and single α -thalassaemia deletions	87
4.4.3	Reliability of α -deletions detected	89
4.4.4	Correlating conventional findings to DNA findings	92
4.5	Prevalence of common α -thalassemia deletions in blood donors	94
4.5.1	Ethnicity	94
4.5.2	Allele frequency	97
4.5.3	The predicted future health burden	99

4.6	Quantitative analysis of α^{SEA} by real-time PCR for prenatal diagnosis	101
4.6.1	Raw Data Plot	101
4.6.2	Multicomponent Plot	101
4.6.3	Quality of Standard Curve	102
4.6.4	Absolute quantification	105
4.6.5	Normalization of α^{SEA} -BR and α -GCR	106
4.6.6	Local DNA Sequence Alignment	110
5	DISCUSSION	
5.1	Correct approach in the detection of alleles of 5 common deletional α -thalassaaemia in Malaysia	114
5.2	Screening for common deletional α -thalassaaemia carriers in blood donors	118
5.3	Prevalence of common α -thalassemia deletions in blood donors	124
5.4	Quantitative analysis of deletional α -thalassaaemia for prenatal diagnosis of Hb Barts hydrops foetalis by real-time PCR	130
6	SUMMARY, CONCLUSION AND RECOMMENDATIONS FOR FUTURE RESEARCH	134
REFERENCES		140
APPENDICES		149
BIODATA OF STUDENT		159
LIST OF PUBLICATIONS		160