



UNIVERSITI PUTRA MALAYSIA

***HAEMOGLOBIN ADANA IN ALPHA THALASSAEMIA INTERMEDIA IN
THE MALAYSIAN POPULATION***

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THE MALAYSIAN POPULATION**

By

LEE TZE YAN

**Thesis Submitted to the School of Graduate Studies, Universiti Putra Malaysia, in
Fulfillment of the Requirements for the Degree of Doctor of Philosophy**

June 2017

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Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfillment of the requirement for the degree of Doctor of Philosophy

HAEMOGLOBIN ADANA IN ALPHA THALASSAEMIA INTERMEDIA IN THE MALAYSIAN POPULATION

By

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June 2017

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Alpha thalassaemia is an autosomal blood disorder due to a quantitative reduction or total absence of alpha globin chain synthesis caused by alpha globin gene mutations. Alpha thalassaemia intermedia or more commonly known as HbH disease is a form of thalassaemia with intermediate severity. A total of 3 out of 4 alpha globin genes are non-functional. There are two main types of HbH disease, the deletional and non-deletional HbH disease. There are limited studies in Malaysia on alpha thalassaemia intermedia particularly the severe form of non-deletional alpha thalassaemia like Hb Adana. Hence, it is imperative to identify alpha thalassaemia intermedia with its spectrum, origin of mutations and genotype-phenotype correlation. We hypothesized that Hb Adana is an important severe alpha thalassaemia and its molecular basis needs to be further elucidated for better understanding. Genotype-phenotype correlation was done on the 320 alpha thalassaemia intermedia cases collected. Three main groups from the pool of alpha thalassaemia cases were categorized into deletional HbH, non-deletional HbH, non-deletional alpha thalassaemia. The most common alpha thalassaemia genotypes found in this study is the $-\alpha^{3.7}/--^{SEA}$ which forms more than 50% (55.3%) of the cases. The non deletional HbH cases have a lower Hb, RBC, MCHC and HbA₂ mean value than deletional HbH. However, the RDW, MCV and MCH of the non-deletional HbH cases are higher than the deletional HbH. A rapid detection method to screen α -thalassaemia variants was also evaluated by using the new technology of digital droplet PCR (ddPCR). The design of this assay encompasses the detection of triplications, common and rare deletional alpha thalassaemia by utilising this new technology. This quantitative method was found to be able to measure and determine the copy number changes therefore could simultaneously detect deletions, duplications and triplications of the alpha globin gene cluster. Hence, ddPCR is an alternative method for rapid detection of alpha thalassaemia variants in Malaysia. The ddPCR was also able to detect Hb Adana (Cd59) which is a SNP mutation. Both the wild type and heterozygous Hb Adana mutation was successfully characterised using the ddPCR method. All Hb Adana cases were then subjected to direct sequencing and PCR RFLP to determine the position of Cd59 mutation on the alpha globin gene. All the 36 cases of Hb Adana have the Cd59 position on $\alpha 2$ globin gene. Majority of the Hb Adana samples found were of the Malay ethnicity

in the alpha thalassaemia intermedia pool. Further investigations on the haplotype patterns revealed that most of the Hb Adana cases belongs to the Haplotype I pattern (53.8%) in Malaysia. This suggests that the Hb Adana pool in Malaysia has a distinctive pattern and there is a high possibility that they might well be from the same genetic pool and quite similar to the one from Indonesia. In conclusion, Hb Adana with mutation at $\alpha 2$ globin gene and its distinctive haplotype pattern is found predominantly in the Malay population in Malaysia.



Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Doktor Falsafah

HEMOGLOBIN ADANA DALAM ALPHA TALASEMIA INTERMEDIA DALAM POPULASI MALAYSIA

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Alpha talasemia adalah penyakit darah autosomal yang disebabkan oleh pengurangan kuantitatif atau tidak adanya sintesis rantai alpha globin disebabkan oleh mutasi gen alpha globin. Alpha talasemia intermedia atau lebih dikenali sebagai penyakit HbH merupakan sejenis talasemia yang mempunyai keparahan yang sederhana. Sebanyak 3 daripada 4 alpha globin gen untuk penyakit ini tidak berfungsi. Terdapat dua jenis penyakit HbH iaitu penyakit HbH delesi dan bukan delesi. Kajian tentang alpha talasemia di Malaysia adalah terhad terutamanya yang merupakan alpha talasemia bukan delesi yang berbentuk parah. Oleh itu, adalah penting untuk mengenalpasti alpha talasemia intermedia dengan spektrum, sumber mutasi dan korelasi genotip fenotip. Kami membuat hipotesis bahawa Hb Adana merupakan sejenis alpha talasemia parah dan asas molekularnya perlu dikaji secara mendalam untuk pemahaman selanjutnya. Korelasi genotip dan fenotip telah dilakukan ke atas 320 kes alpha talasemia intermedia yang terkumpul. Tiga golongan utama daripada kumpulan kes alpha talasemia telah terbahagi kepada HbH delesi, HbH bukan delesi dan alpha talasemia bukan delesi. Genotip alpha tal assemia yang paling biasa dijumpai adalah jenis $-\alpha^{3.7}/--SEA$ yang merangkumi lebih daripada 50% (55.3%) kes. Kes-kes HbH bukan delesi mempunyai nilai min Hb, RBC, MCHC dan HbA₂ yang lebih rendah berbanding daripada HbH delesi. Namun begitu, nilai RDW, MCV dan MCH untuk kes HbH bukan delesi adalah lebih tinggi daripada HbH delesi. Satu kaedah cepat pengesanan variasi alpha talasemia telah dikaji dengan menggunakan teknologi terkini “digital droplet PCR (ddPCR)”. Reka-bentuk kaedah ini merangkumi pengesanan alpha talasemia delesi yang berbentuk triplikasi, biasa dan yang jarang berlaku dengan penggunaan teknologi baharu ini. Kaedah kuantitatif ini juga didapati dapat mengukur dan menentukan perubahan “copy number” yang sekaligus dapat mengesan delesi, duplikasi dan triplikasi untuk kluster gen alpha globin. Oleh itu, ddPCR merupakan kaedah alternatif untuk mengesan dengan pantas variasi alpha talasemia di Malaysia. ddPCR juga mampu mengesan Hb Adana (Cd59) yang merupakan sejenis mutasi titik. Kedua-dua jenis liar dan mutasi heterozigot telah berjaya dicirikan melalui kaedah ddPCR ini. Kesemua kes Hb Adana kemudian telah dilakukan teknik jujukan dan PCR RFLP untuk menentukan posisi mutasi Cd59 di gen alpha globin. Kesemua 36 kes Hb Adana mempunyai posisi

mutasi Cd59 di gen globin $\alpha 2$. Terdapat 32 daripada 36 kes Hb Adana merupakan kaum Melayu yang mencadangkan bahawa kumpulan gen kes Hb Adana di Malaysia hampir sama dengan kes yang terdapat di Indonesia. Kajian susulan ke atas corak haplotip mendedahkan kebanyakan kes Hb Adana mempunyai corak Haplotip I (53.8%) di Malaysia. Ini mencadangkan bahawa kumpulan Hb Adana yang terdapat di Malaysia mempunyai corak yang tersendiri and besar kemungkinannya kumpulan ini tergolong dalam kumpulan genetik yang sama. Konklusinya, Hb Adana berinteraksi dengan mutasi-mutasi alpha talasemia yang lain untuk membentuk alpha thalassemia intermedia di Malaysia.



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I certify that a Thesis Examination Committee has met on 15 June 2017 to conduct the final examination of Lee Tze Yan on his thesis entitled "Haemoglobin Adana in Alpha Thalassaemia Intermedia in the Malaysian Population" 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 Doctor of Philosophy.

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LIST OF ABBREVIATIONS

ARMS	Amplification refractory mutations system
AS-PCR	Allele-specific PCR
ATMDS	Alpha thalassaemia and myelodysplastic syndrome
ATRX	Alpha thalassaemia mental retardation-X linked syndrome
Cd	Codon
CN	Copy number
CNV	Copy number variation
ddPCR	Digital droplet Polymerase Chain Reaction
DHPLC	Denaturing high-performance liquid chromatography
DNA	Deoxyribonucleic acid
DQ	Dosage Quotient
FAM	6-carboxyfluorescein
HbH	Haemoglobin H
HRM	High-resolution melting
IVS	Intervening sequence
LCR	Locus control region
MCH	Mean corpuscular haemoglobin
MCHC	Mean corpuscular haemoglobin concentration
MCV	Mean corpuscular volume
MLPA	Multiplex Ligation-dependent Probe Amplification
MSC-R2	Multispecies conserved sequences region 2
NGS	Next-Generation sequencing
OH	Hydroxyl group
PCR-RDB	Polymerase chain reaction reverse dot blot
RE	Restriction enzyme
RBC	Red blood cell
RDW	Red cell distribution width
RFLP	Restriction length polymorphism
SNP	Single nucleotide polymorphism
VIC	2'-chloro-7'-phenyl-1,4-dichloro-6-carboxyfluorescein

CHAPTER 1

INTRODUCTION

1.1 Structure of Haemoglobin

Haemoglobin is a tetrameric protein formed by 2 pairs of globin chains with an iron-containing haem molecule in each globin chain to transport oxygen in the human body (Lewis *et al.*, 2012; Bain *et al.*, 2010). HbA, a major haemoglobin in normal adults constitutes about 97% of the total haemoglobin and another component, HbA₂ accounts for a mere 2-3% of the total. Foetal haemoglobin (HbF) is the major haemoglobin in foetus and in most of the normal adults, however, a minute amount of HbF can still be present (Weatherall & Clegg, 2001).

The globin genes which encode the globin chains can be found in two clusters (Figure 1.1). They are located in chromosome 16 (alpha globin gene cluster) and chromosome 11 (beta globin gene cluster). Alpha globin gene cluster is made up of a few pseudogenes, a zeta (ζ) gene and two alpha (α) genes. On the other hand, beta globin gene cluster consists of an epsilon (ϵ) gene, two gamma (γ) genes, a delta (δ) gene and a beta (β) gene (Bain *et al.*, 2010). Both alpha and beta globin gene cluster are arranged according to their developmental expression (Thein, 2004).

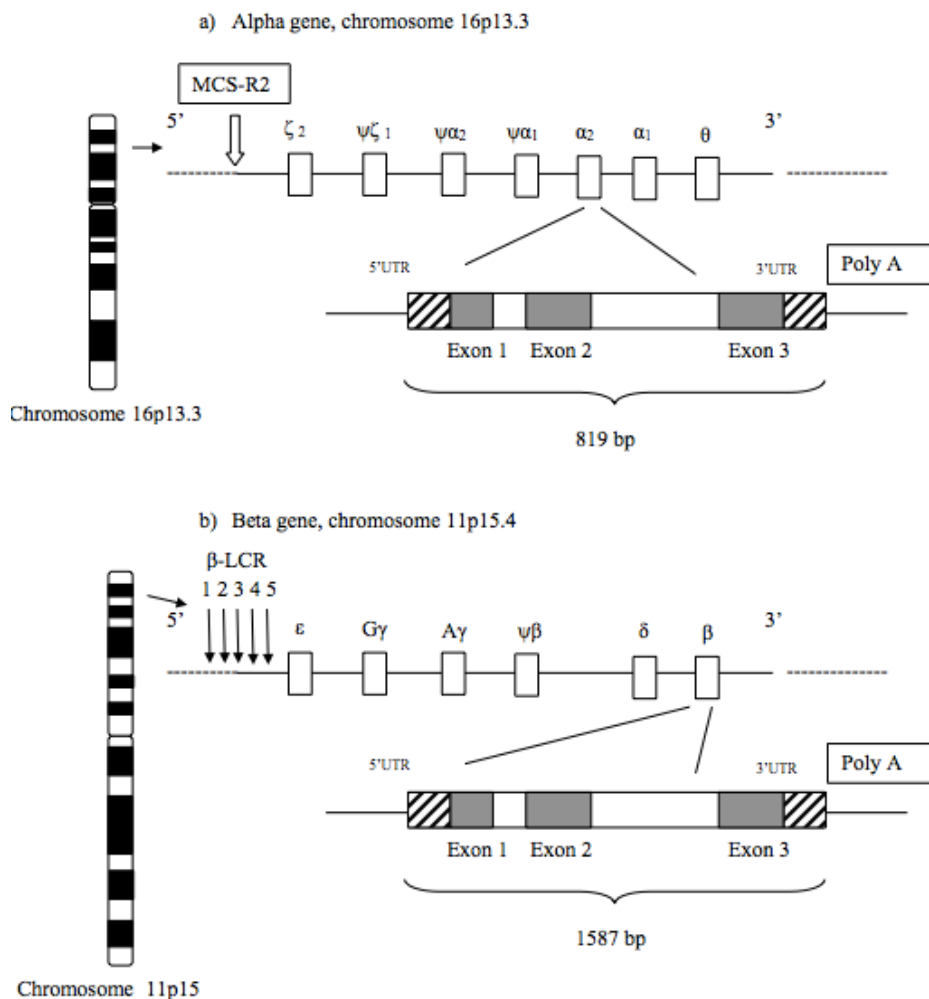


Figure 1.1: Maps of the α -like and β -like globin gene clusters. (a) Alpha gene locus on chromosome 16p13.3. Down-point vertical arrow represents the important gene expression locus, MSC-R2 (upstream hypersensitive site). (b) Beta locus on chromosome 11p15.4. Five hypersensitive sites are indicated by 5 vertical down point arrows, representing the gene regulatory sequences in the β -globin cluster, also known as locus control region (LCR) (Modified from Clark and Thein, 2004).

1.2 Haemoglobinopathies

Normal adult human haemoglobin is a paired complex of two α -globin and two β -globin chains (Bain *et al.*, 2010). The other two minor haemoglobins, Hb A₂ and Hb F are made up of 2 α and 2 δ chains and 2 α and 2 γ chains respectively. In embryo, pre-alpha chains (ζ globin chains) contribute to embryonic haemoglobin and for fetal development, β -like globin chains (ϵ and γ) are involved (Nemati *et al.*, 2010). There are basically two types of haemoglobin abnormalities (Figure 1.2). One is due to the synthesis of an abnormal haemoglobin (variant haemoglobin) while the other is caused

by the reduction or absence of normal alpha (α) and/or beta (β) chains production (thalassaemia syndromes) (Hoffbrand *et al.*, 2006). Variant haemoglobin or structural haemoglobinopathies is the term used to describe structurally abnormal haemoglobins which includes haemoglobin E, haemoglobin C and haemoglobin S (Bain *et al.*, 2006; Hoffbrand *et al.*, 2006). In Malaysia, thalassaemia is a public health problem among the ethnic Malay and Chinese populations whereas ethnic Indians form a minority percentage (Weatherall & Clegg, 2001).

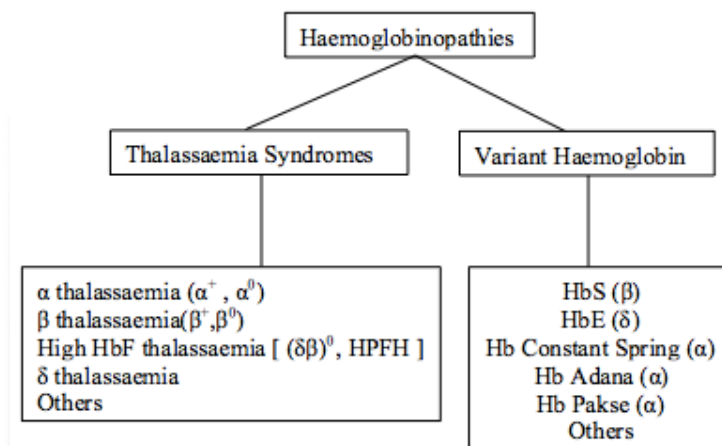


Figure 1.2: The 2 different types of haemoglobinopathies. Thalassaemia syndromes involve the reduction or absence of alpha or beta globin chains production while variant haemoglobin is the synthesis of an abnormal haemoglobin.

1.3 Alpha thalassaemia

Thalassaemia is the commonest monogenic disorder in human beings worldwide (Weatherall & Clegg, 2001). There is a double copy of the α -globin gene per haploid genome which gives rise to the annotation $\alpha\alpha/\alpha\alpha$. α^0 -thalassaemia with deletion of both α -globin genes will result in the absence of α -globin chain production whereas a reduction of α -globin chain production is caused by deletion of a single gene ($-\alpha/\alpha\alpha$) resulting in α^+ thalassaemia. Minor or trait (carriers) for α^0 -thalassaemia ($--/\alpha\alpha$), homozygous alpha thalassaemia ($-\alpha/-\alpha$) or α^+ -thalassaemia ($-\alpha/\alpha\alpha$) are normally asymptomatic or have mild anaemia. Compound heterozygous α^0 and α^+ -thalassaemia ($--/-\alpha$) will cause HbH disease or α -thalassaemia intermedia. The major form, homozygous α^0 -thalassaemia ($--/--$), results in intrauterine death and hydrops foetalis in the absence of intrauterine blood transfusions (Weatherall, 1997; Weatherall, 1999).

1.4 Haemoglobin Adana (Hb Adana)

Hb Adana [HBA2: c179G>A (or HBA1); p.Gly60Asp] is a common non-deletional α -thalassaemia variant found among the Malays (Ahmad *et al.*, 2013). This highly unstable variant is located at Cd 59 position either on $\alpha 1$ globin gene or $\alpha 2$ globin gene. Haemoglobin subtyping is unable to detect this highly unstable α -haemoglobin variant

since Hb Adana carriers have normal red blood cell indices and are therefore generally asymptomatic. (Azma et. al., 2014). Cases of Hb Adana were reported in the Malaysian Malays, Chinese and indigenous populations (Ahmad et al., 2013). Furthermore, the first reported case of Hb Adana in Malaysia was in a 52-year-old patient with α -thalassaemia intermedia (George et al, 2009). Confirmation of Hb Adana is a diagnostic predicament as this highly unstable haemoglobin variant has no product when observed during routine haematology tests. Hence, patients may remain undiagnosed until DNA studies are carried out. In Malaysia, the reliance on routine haematology studies and Hb subtype analysis to detect Hb Adana may not be sufficiently accurate in populations with different globin gene mutations (Tan J.A.M.A, et al., 2016). The interaction between Haemoglobin (Hb) Adana (HBA2:c.179>A) with deletional and nondeletional α -thalassaemia mutations will produce HbH disorders with varying clinical manifestations from asymptomatic to severe anaemia with significant hepatosplenomegaly.

1.5 Main challenges in Hb Adana

Similar to beta thalassaemia, screening, diagnosis and the understanding of the genotype-phenotype interactions of alpha thalassaemia often come with many challenges. For alpha thalassaemia intermedia, there are limited studies in Malaysia which focus on this category as most of the research are concentrated on the management and prevention of beta thalassaemia major. Besides that, the molecular basis of severe forms of alpha thalassaemia is not well-studied and a good example will be Hb Adana. Therefore, it is imperative to identify alpha thalassaemia intermedia at an early age with its spectrum of mutations, origin of mutations and genotype-phenotype correlation. It would be better if we could apply and evaluate rapid and newer techniques to detect some of the alpha thalassaemia mutations so that screening of alpha thalassaemia can be done faster and will then lead to accurate definitive diagnosis. Our hypothesis of this study is that Hb Adana is an important severe alpha thalassaemia when interacting with the other alpha mutations and its molecular basis needs to be further elucidated for better understanding.

1.6 Objectives

The general objective of this study is:

1. To determine the genotypes and its ethnic distribution of Hb Adana of alpha thalassaemia intermedia in Malaysia

The specific objectives of this study are:

- 1) To correlate the genotypes with phenotypes of alpha thalassaemia intermedia
- 2) To apply a rapid and newer technique to identify deletional alpha thalassaemia mutations and to screen for Hb Adana
- 3) To identify the position of Hb Adana mutation on the alpha globin gene in Malaysia
- 4) To determine the haplotypes associated with Hb Adana

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