DOWN SYNDROME: DEVELOPMENT OF A NON-INVASIVE PRENATAL DNA SCREENING TEST USING SUPEROXIDE DISMUTASE 1 GENE IN MATERNAL BLOOD AND DETECTION OF CYSTATHIONINE p-SYNTHASE GENE MUTATIONS

THILAKAVATHY A/P KARUPPIAH

FPSK(P) 2004 7
DOWN SYNDROME: DEVELOPMENT OF A NON-INVASIVE PRENATAL DNA SCREENING TEST USING SUPEROXIDE DISMUTASE 1 GENE IN MATERNAL BLOOD AND DETECTION OF CYSTATHIONINE β-SYNTHASE GENE MUTATIONS

By

THILAKAVATHY A/P KARUPPIAH

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

February 2004
DEDICATION

This thesis is dedicated to my spiritual master,

HIS DIVINE GRACE YOGA JNANA SIVIDHAROM SRI RAJAYOGA GURU,

without whom none of this would have been even possible.
Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfilment of the requirements for the degree of Doctor of Philosophy

DOWN SYNDROME: DEVELOPMENT OF A NON-INVASIVE PRENATAL DNA SCREENING TEST USING SUPEROXIDE DISMUTASE 1 GENE IN MATERNAL BLOOD AND DETECTION OF CYSTATHIONINE β-SYNTHASE GENE MUTATIONS

By

THILAKAVATHY A/P KARUPPIAH

February 2004

Chairman: Associate Professor Rozita Rosli, Ph.D.

Faculty: Medicine and Health Sciences

Down syndrome or Trisomy 21, is the most commonly occurring genetic disorder that stems from the failure of chromosome 21 to segregate normally during meiosis, resulting in an individual carrying an extra copy of chromosome 21. The main aims of this study were to develop a relatively non-invasive prenatal DNA screening method using maternal blood and to detect mutations on cystathionine β-synthase (CBS) gene, a folate pathway gene located on chromosome 21. As an initial step, the presence of foetal cells and DNA in the maternal blood was firstly determined by foetal haemoglobin (HbF) staining and polymerase chain reaction (PCR). It was found that the ratio of the nucleated foetal cell to maternal cell increased from 2 in $10^6$ to 3 in $10^6$ and 5 in $10^6$ at the first, second and third trimester, respectively. By using Y chromosome specific primers, DNA from male foetuses could be detected as early as 6 weeks of gestation in 200 µl maternal blood obtained from fingertip. This is in line with the current technology in non-invasive screening methods of foetal aneuploidies which is focused on detecting Y chromosomal sequences which is impossible to be used
for female foetus pregnancies. Therefore, the superoxide dismutase 1 (SOD1) gene sequence, which is located on the Down Syndrome Critical Region, was used to overcome this situation by using real-time quantitative PCR. The level of SOD1 sequences in maternal blood was found to be significantly elevated in the third trimester normal pregnancies (mean = 11728 copies/μl) when compared to the second trimester (mean = 5705.6 copies/μl), p<0.005 and non-pregnant normal women (mean = 3580.2 copies/μl), p<0.0001. Down syndrome pregnancies have the greatest elevation compared to all the three trimesters of normal singleton pregnancies and twin pregnancies, p<0.05. The traditional approach of prenatal chromosomal diagnosis using amniotic fluid was found to be cumbersome and time-consuming compared to the newly developed method. The mutation detection on CBS gene was carried out using DNA sequencer and denaturing high performance liquid chromatography (DHPLC). This study revealed that the Down syndrome patients have four mutations, which are in intron 1 (A9231C), exon 10 (C20628T) and exon 17 (T27796C and C27817T). The Down syndrome children were found to have the same genotype as their mothers. The number of mothers and children having the substitutions in the CBS gene was twice the number of mothers and children with normal genotype, suggesting that the mothers who have these substitutions are at higher risk of having a child with Down syndrome. In conclusion, non-invasive prenatal diagnosis at first trimester using Y chromosomal sequence is feasible for diagnosis of foetal-derived paternally-inherited polymorphism/mutations or genes. Quantitative analysis using gene associated with a disorder has a potentially significant advantage over the invasive techniques currently used widely for prenatal diagnosis. Finally, the discovery of the mutations in the CBS
gene of Down syndrome patients and mothers will help contribute to new knowledge and the future studies on the folate pathway genes mutation and occurrence of Down syndrome. It may also suggest an opportunity to improve public health strategies for the primary prevention of Down syndrome.
Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Doktor Falsafah

**SINDROM DOWN: PENGHASILAN UJIAN PENYARINGAN PRENATAL DNA YANG TIDAK MERBAHAYA MENGGUNAKAN GEN SUPEROXADA DISMUTASE 1 DALAM DARAH IBU MENGANDUNG DAN PENGESANAN MUTASI GEN SISTATHIONIN β-SINTASE**

Oleh

**THILAKAVATHY A/P KARUPPIAH**

Februari 2004

Pengerusi: Profesor Madya Rozita Rosli, Ph.D.

Fakulti: Perubatan dan Sains Kesihatan

Sindrom Down atau trisomi 21, merupakan keabnormalan genetik yang paling biasa berlaku akibat kegagalan kromosom 21 untuk membahagi secara normal semasa meiosis, menyebabkan seseorang individu mempunyai tambahan satu kromosom 21. Objektif-objektif utama kajian ini adalah untuk menghasilkan sebuah teknik penyaringan prenatal DNA yang tidak merbahaya secara relatif dan juga mengesan mutasi-mutasi pada gene sistathionin β-sintase (CBS), satu gen yang terletak di tapak-jalan folat pada kromosom 21. Sebagai langkah pertama, kehadiran sel-sel dan DNA janin di dalam darah ibu hamil ditentukan dengan pewarnaan haemoglobin janin (HbF) dan tindakan rantai polymerase (PCR). Didapati bahawa, nisbah antara sel janin bernukleus dengan sel ibu hamil bertambah dari 2 dalam $10^6$ kepada 3 dalam $10^6$ dan 5 dalam $10^6$ masing-masing pada trimester pertama, kedua dan ketiga. Primer-primer yang spesifik bagi kromosom Y dapat mengesan DNA dari janin-janin lelaki seawal enam minggu gestasi dengan menggunakan 200 μl darah ibu hamil yang diambil dari hujung jari. Ini adalah sejajar dengan teknologi terkini dalam penyaringan janin
aneuploidi yang tidak berbahaya yang difokus pada pengesanan jujukan kromosom Y di mana ianya tidak mungkin dapat digunakan untuk kandungan janin perempuan. Oleh itu, jujukan gen superoxida dismutase 1 (SOD1), yang berlokal pada Kawasan Kritikal Sindrom Down, digunakan untuk mengatasi situasi ini melalui aplikasi kuantitatif masa-benar PCR. Tahap jujukan SOD1 dalam darah ibu hamil didapati meningkat secara ketara pada kehamilan normal trimester ketiga (min = 11728 salinan/μl) apabila dibandingkan dengan kehamilan normal trimester kedua (min = 5705.6 salinan/μl), $p<0.005$ dan wanita normal yang tidak hamil (min = 3580.2 salinan/μl), $p<0.0001$. Kehamilan sindrom Down mempunyai peningkatan yang paling tinggi berbanding dengan kehamilan tunggal kesemua trimester dan kehamilan kembar, $p<0.05$. Diagnosis prenatal kromosom tradisional yang menggunakan cecair amnion didapati sukar dan mengambil lebih masa berbanding dengan teknik baru yang telah dihasilkan. Penentuan mutasi pada gen CBS dilakukan dengan menggunakan penjujuk DNA dan "denaturing high performance liquid chromatography" (DHPLC). Kajian ini telah menunjukkan bahawa pesakit sindrom Down mempunyai empat mutasi, iaitu di dalam intron 1 (A9231C), exon 10 (C20628T) dan exon 17 (T27796C dan C27817T). Kanak-kanak sindrom Down didapati mempunyai genotaip yang sama dengan ibu mereka. Bilangan ibu dan anak yang mempunyai substitusi ini adalah dua kali ganda daripada ibu dan anak dengan genotaip normal, mencadangkan bahawa ibu yang mempunyai substitusi ini berisiko tinggi untuk mendapat anak sindrom Down. Sebagai kesimpulan, diagnosis prenatal yang tidak berbahaya pada trimester pertama menggunakan jujukan kromosom Y boleh digunakan untuk mendiagnosis janin yang membawa polimorfisma/mutasi-mutasi atau gen-gen warisan-bapa. Analisis kuantitatif
menggunakan gen yang berkaitan dengan penyakit mempunyai potensi yang tinggi berbanding dengan teknik-teknik merbahaya yang digunakan kini secara meluas untuk diagnosis prenatal. Penemuan mutasi-mutasi pada gen CBS dalam pesakit sindrom Down dan ibu mereka akan membantu menyumbangkan pengetahuan baru dan kajian masa hadapan ke atas mutasi gen-gen di tapak-jalan folat dan kejadian sindrom Down. Ia juga mungkin memberi peluang untuk memperbaiki strategi kesihatan awam bagi pencegahan awal sindrom Down.
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I owe my respectful gratitude to my spiritual master, His Divine Grace Yoga Jnana Sitthar Om Sri Rajayoga Guru, whose blessings have accompanied me every step of the way in finishing this thesis work and made the impossible possible.

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I would like to express my heartfelt thanks to my beloved mother, brother and sister for their endless encouragement, patience, care and sacrifices that had helped me in my undertakings and to complete this research study successfully.

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I certify that an Examination Committee met on 18th February 2004 to conduct the final examination of Thilakavathy a/p Karuppiah on her Doctor of Philosophy thesis entitled "Down Syndrome: Development of a Non-Invasive Screening Test for Prenatal Diagnosis Using Superoxide Dismutase 1 Gene in Maternal Blood and Detection of Cystathionine β-Synthase Gene Mutations" in accordance with Universiti Pertanian Malaysia (Higher Degree) Act 1980 and Universiti Pertanian Malaysia (Higher Degree) Regulations 1981. The Committee recommends that the candidate be awarded the relevant degree. Members of the Examination Committee are as follows:

**Elizabeth George, FRCPE**  
Professor  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Chairman)

**Rozita Rosli, Ph.D.**  
Associate Professor  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Member)

**Jammal Ahmad Essa, MRCOG**  
Associate Professor  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Member)

**Tan Boon Chong, MRCOG**  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Member)

**TP Baskaran, MOG**  
Department of Obstetrics & Gynaecology  
Hospital Kuala Lumpur  
(Member)

**Abdul Rahman Abdul Jamal, MRCP**  
Professor  
Faculty of Medicine  
Hospital Universiti Kebangsaan Malaysia  
(Independent Examiner)

---

MAD NASIR SHAMSUDIN, Ph.D.  
Professor/Deputy Dean  
School of Graduate Studies  
Universiti Putra Malaysia

Date: 14 APR 2004
This thesis submitted to the Senate of Universiti Putra Malaysia and was accepted as fulfilment of the requirements for the degree of Doctor of Philosophy. The members of the Supervisory Committee are as follows:

**Rozita Rosli, Ph.D.**
Associate Professor  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Chairman)

**Jammal Ahmad Essa, MRCOG**  
Associate Professor  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Member)

**Tan Boon Chong, MRCOG**  
Faculty of Medicine and Health Sciences  
Universiti Putra Malaysia  
(Member)

**TP Baskaran, MOG**  
Hospital Kuala Lumpur  
(Member)

---

**AINI IDERIS, Ph.D.**
Professor/Dean  
School of Graduate Studies  
Universiti Putra Malaysia

Date: 05 JUL 2004
DECLARATION

I hereby declare that the thesis is based on my original work except for quotations and citations which have been duly acknowledged. I also declare that it has not been previously or concurrently submitted for any other degree at UPM or other institutions.

THILAKAVATHY A/P KARUPPIAH

Date: 15/4/04
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M = 50 bp marker; 1-3 = foetal cells from women with male foetus at 38, 38, 39 weeks gestation, respectively; 4 = male positive control

Annealing temperature optimization of polymerase chain reaction amplification of *SRY* gene from cord blood, maternal blood and amniotic fluid of a male pregnancy. 

M = 50 bp DNA marker; 1-7 = DNA from cord blood (lane 1: 54°C, 2: 54.8°C, 3: 56.3°C, 4: 57.1°C, 5: 58.0°C, 6: 59.3°C, 7: 59.9°C); 8-14 = DNA from maternal blood (lane 8: 54°C, 9: 54.8°C, 10: 56.3°C, 11: 57.1°C, 12: 58.0°C, 13: 59.3°C, 14: 59.9°C); 15-21 = DNA from amniotic fluid (lane 15: 54°C, 16: 54.8°C, 17: 56.3°C, 18: 57.1°C, 19: 58.0°C, 20: 59.3°C, 21: 59.9°C). Black arrow shows the amplification of *SRY* gene in maternal blood at 58°C

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1 = male positive control; 2-6 = DNA from pregnant women with male foetus at 6, 7, 8, 9, 10 weeks gestation, respectively; M = 50 bp DNA marker; 8-12 = DNA from pregnant women with female foetus at 6-10 weeks gestation, respectively; 13 = non template control

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40. The SRY gene sequence amplified to determine the presence of the foetal cell and DNA in the maternal blood

41. The SOD1 gene sequence used for the non-invasive prenatal DNA screening test

42. Purine-purine and pyrimidine-pyrimidine base pairing with a water molecule. (From http://www.bi.umist.ac/users/mjfasjw/2MMB/mutants/diagrams.asp)
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<td>CBS</td>
<td>Cystathionine beta-synthase</td>
</tr>
<tr>
<td>CT</td>
<td>Threshold cycle</td>
</tr>
<tr>
<td>dATP</td>
<td>Deoxyadenosine triphosphate</td>
</tr>
<tr>
<td>dCTP</td>
<td>Deoxycytosine triphosphate</td>
</tr>
<tr>
<td>dGTP</td>
<td>Deoxyguanidine triphosphate</td>
</tr>
<tr>
<td>DHPLC</td>
<td>Denaturing high performance liquid chromatography</td>
</tr>
<tr>
<td>DNA</td>
<td>Deoxyribonucleic acid</td>
</tr>
<tr>
<td>dUTP</td>
<td>Deoxyuridine triphosphate</td>
</tr>
<tr>
<td>E-type</td>
<td>Epitheloid cell</td>
</tr>
<tr>
<td>F-type</td>
<td>Fibroblast like cell</td>
</tr>
<tr>
<td>GAPDH</td>
<td>Glyceraldehydes-3-phosphate dehydrogenase</td>
</tr>
<tr>
<td>HbF</td>
<td>Foetal haemoglobin</td>
</tr>
<tr>
<td>hCG</td>
<td>Human chorionic gonadotropin</td>
</tr>
<tr>
<td>ME-THF</td>
<td>5-methyltetrahydrofolate</td>
</tr>
<tr>
<td>MgCl₂</td>
<td>Magnesium chloride</td>
</tr>
<tr>
<td>MTHFR</td>
<td>5,10-methylene-tetrahydrofolate reductase</td>
</tr>
<tr>
<td>MTRR</td>
<td>Methionine synthase reductase</td>
</tr>
<tr>
<td>Abbreviation</td>
<td>Definition</td>
</tr>
<tr>
<td>--------------</td>
<td>------------</td>
</tr>
<tr>
<td>NaCl</td>
<td>Sodium chloride</td>
</tr>
<tr>
<td>NF</td>
<td>Nuclear factor</td>
</tr>
<tr>
<td>PCR</td>
<td>Polymerase chain reaction</td>
</tr>
<tr>
<td>R_N</td>
<td>Fluorescence intensity</td>
</tr>
<tr>
<td>SAM</td>
<td>S-adenosylmethionine</td>
</tr>
<tr>
<td>SNPs</td>
<td>Single nucleotide polymorphisms</td>
</tr>
<tr>
<td>SOD1</td>
<td>Superoxide dismutase 1</td>
</tr>
<tr>
<td>SRY</td>
<td>Specific Region of Y</td>
</tr>
<tr>
<td>UE3</td>
<td>Unconjugated estriol</td>
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CHAPTER I
INTRODUCTION

Chromosomal abnormalities are the most frequent genetic disorders seen in both live born babies and miscarriages. Down syndrome is a chromosomal abnormality, which manifests itself in a set of common physical and mental characteristics. This abnormality is due to the presence of an extra chromosome (chromosome 21). Hence, the scientific name, trisomy 21.

Most people are aware that the chance of having a baby with Down syndrome is greater in older women. But Down syndrome can occur at any maternal age. In fact, 75-80% of babies with Down syndrome are born in younger women simply because that age group has more babies (Benke et al., 1995). Studies have proven that inadequate folate status at the time of conception increases the risk of Down syndrome. Women with genetic mutations which interfere with the body's ability to absorb folic acid, are at higher risk for having children with Down syndrome (James et al., 1999).

Down syndrome is a major reason for prenatal diagnosis. Prenatal diagnosis employs a variety of techniques to determine the health and condition of an unborn foetus. Such diagnosis is usually performed by means of karyotyping and depends on analysis at 11-18 weeks of gestation. Karyotyping is not always possible, especially when the number of cells obtained is limited, where cell culture fails (in 1-2% of patients), or when the