

UNIVERSITI PUTRA MALAYSIA

GENETIC DIVERSITY OF SELECTED AROMATIC PLANTS IN AYER HITAM FOREST RESERVE, SELANGOR

MOHAMAD AZREN PUTRA BIN MAT DESA

FH 2016 53

GENETIC DIVERSITY OF SELECTED AROMATIC PLANTS IN AYER HITAM FOREST RESERVE, SELANGOR



FACULTY OF FORESTRY
UNIVERSITI PUTRA MALAYSIA
2016

GENETIC DIVERSITY OF SELECTED AROMATIC PLANTS IN AYER HITAM FOREST RESERVE, SELANGOR

Ву

MOHAMAD AZREN PUTRA BIN MAT DESA

A Project Report Submitted in Partial Fulfillment of the Requirements

for the Degree of Bachelor of Forestry Science in the

Faculty of Forestry

Universiti Putra Malaysia

DEDICATION

This is dedicated to my family and friends.

My deepest gratitude to my beloved parents; Mr. Mat Desa @ Mohd Wari Bin Haron and Mrs. Laili Binti Mohd Hashim, and to my family members; Nor Lailinda Hanum Binti Mat Desa, Mohd Azizi Bin Mat Desa, Nor Lailian MaimonHayu Binti Mat Desa, Nor Lily Maria Hadijah Binti Mat Desa and Mohamad Azrai Fitri Bin Mat Desa for their supports and loves. Not to be forget, my late sister Nor Lailenney Marlina Hanim Sara Binti Mat Desa, our love and pray always be with you.

I also dedicate this dissertation to my supervisor, Associate Professor Dr. Rozi Mohamed and my Forest Biotechnology's Colleagues.

Last but not least, I dedicate this dissertation to Dr. Razak for his wise and knowledge in evaluating this documentation of the project.

May God Bless All of Us

ABSTRACT

Malaysia is rich in aromatic plant species but they have not been fully explored. During an exploration in October 2015 at Ayer Hitam Forest Reserve (AHFR), Selangor, several plant species were identified from their ability to produce strong aroma. They included Trema angustifolia (Cenderai/Mengkirai), Cinnamomum iners (Medang Teja), Actinodaphne sphaerocarpa (Medang Payung), Syzygium polyanthum (Serai Kayu), Prismatomeris glabra (Tongkat Haji Samad), Monocarpia marginalis (Mempisang) and Chromolaena odorata (Kapal Terbang). Morphology of each species was described and DNA samples were isolated to provide for genetic information. DNA samples were extracted using FavorPrep Plant Genomic DNA Extraction Mini Kit. The DNA psbA-trnH region was amplified in PCR and sequenced. Sequence analysis and searches data at GeneBank's database revealed that many of Malaysian aromatic plant species have not been sequenced. This is the first report on genetic diversity of aromatic plants from AHFR. More efforts should be taken to document these plant species for potential use in producing aromatic-essential products.

ABSTRAK

Malaysia kaya dengan kepelbagaian spesis tumbuhan aromatik, namun masih belum diteroka sepenuhnya. Dalam satu ekspedisi pada Oktober 2015 di Hutan Simpan Ayer Hitam (HSAH), Selangor, beberapa spesis tumbuhan telah dikenalpasti berupaya menghasilkan aroma yang kuat. Spesies-spesies tersebut adalah Trema angustifolia (Cenderai/Mengkirai), Cinnamomum iners (Medang Teja), Actinodaphne sphaerocarpa (Medang Payung), Syzygium polyanthum (Serai Kayu), Prismatomeris glabra (Tongkat Haji Samad), Monocarpia marginalis (Mempisang) dan Chromolaena odorata (Kapal Terbang). Ciri-ciri morfologi kesemua species direkod dan sampel DNA digunakan bagi merekod maklumat genetik. Sampe DNA diekstrak dengan menggunakan FavorPrep Plant Genomic DNA Extraction Mini Kit. Struktur DNA psbA-trnH dikuatkan melalui PCR dan disusun. Analisis turutan DNA dan hasil carian data di pangkalan data GeneBank menunjukkan bahawa masih banyak spesis tumbuhan aromatik yang belum disusun dan direkod. Ini merupakan laporan pertama tentang kepelbagaian genetik spesis tumbuhan aromatik di HSAH. Tindakan lanjut dalam pendokumentasian spesis-spesis ini perlu diambil bagi kegunaan masa hadapan terutama dalam pembangunan produk berasaskan tumbuhan aromatik.

ACKNOWLEDGEMENT

I would like to express my gratitude to Allah S.W.T for His blessing and guidance in giving me strength to complete this thesis entitled "Genetic Diversity of Selected Aromatic Plants in Ayer Hitam Forest Reserve, Selangor". Special thanks to my supervisor, Assoc. Prof. Dr. Rozi Mohamed for the endless support, guidance and advice for me to finish this project. I also want to thank to Dr. Razak Terhem for his wise and acknowledge in evaluating this project.

My special gratitude to my Forest Biotechnology Lab's colleagues; Mr. Lee Shiou Yih, Miss Siti Rahimah Jumaat, Miss Aimi Zafirah Adam and Miss Wong Mun Theng. Their helps and guidance are really being appreciated.

Special appreciation is given to my family members for supporting me and giving me the encouragement to finish my final year project. They are always by my side along the time I was preparing for this thesis.

Last but not least, to all my friends those have helped me; those have never given up giving moral supports to me, thank you very much. I really appreciate what you have given to me and I would never forget all your kindness.

APPROVAL SHEET

I certify that this research project report entitled Genetic Diversity of Selected Aromatic Plants in Ayer Hitam Forest Reserve, Selangor by Mohamad Azren Putra Bin Mat Desa has been examined and approved as a partial fulfillment of the requirements for the degree of Bachelor of Forestry Science in the Faculty of Forestry, Universiti Putra Malaysia.

Approved by:

U P M

Assoc. Prof. Dr. Rozi Mohamed Faculty of Forestry Universiti Putra Malaysia (Supervisor)

Prof. Dr. Mohamed Zakaria bin Hussin Dean Faculty of Forestry Universiti Putra Malaysia

Date: 27 June 2016

TABLE OF CONTENT

		Page
DED	ICATION	i
ABS ¹	TRACT	ii
ABS ¹	TRAK	iii
ACK	NOWLEDGEMENTS	iv
APP	ROVAL SHEET	V
TABI	LE OF CONTENTS	vi
LIST	OF TABLES	viii
LIST	OF FIGURES	ix
LIST	OF ABBREVITIONS	X
СНА	PTER	
1	INTRODUCTION	
	1.1 Background 1.2 Problem Statement	1
	1.3 Objectives	4
	1.4 Justification of Study	5
2	LITERATURE REVIEW	
2	2.1 General Background of Aromatic Plants	6
	2.2 Description of Ayer Hitam Forest Reserve, Selango	
	2.3 General Background of Lauraceae	8
	2.3.1 Description of Cinnamomum iners	8
	2.3.2 Description of Actinodaphne sphaerocarpa	9
	2.4 General Background of Annonaceae	9
	2.5 General Background of Myrtaceae	10
	2.6 General Background of Rubiaceae 2.7 General Background of Asteraceae	10 11
	2.8 General Background of Cannabaceae	12
	2.9 General Background of Genetic Diversity	12
0		
3	METHODOLOGY 3.1 Plants Identification and Selection	14
	3.1 Plants Identification and Selection 3.2 Plants Materials	14
	3.3 Plant DNA Extraction	15
	3.4 PCR Amplification	17
	3.5 Agarose Gel Electrophoresis	18
	3.6 DNA Sequencing and Alignment	19
	3.7 Phylogenetic Analyses	19
1	RESULTS	
4	4.1 Samples Documentation	20
	4.1 Samples Documentation 4.2 Quantification of Genomic DNA	20 29
	4.3 PCR Amplification Product	30
	4.4 DNA Sequences Analyses	34
	4.5 Phylogenetic Tree	36

5	DISCUSSION 5.1 Plants Identification and Selection 5.2 Phylogenetic Tree	39 39
6	CONCLUSION AND RECOMMENDATION 6.1 Conclusion 6.2 Recommendation	41 42
	REFERENCES	43
	APPENDICES Appendix A: Sequences Alignment Appendix B: TAE Buffer 50X Ingredients	46 55
	PUBLICATION OF THE PROJECT UNDERTAKING	56

LIST OF TABLE

Table		Page
3.1	List of selected species	15
3.2	PCR mixture components and their volumes	17
3.3	Non-coding cpDNA of psbA-trnH primer sequences	18
4.1	Morphological information of selected aromatic species	20
4.2	Mean ratio and concentration for seven selected aromatic plants	30
4.3	List of sequences used in the phylogenetic analysis	35
4.4	Sequences producing significant alignments	38

LIST OF FIGURE

Figure		Page
3.1	DNA extraction's protocol	15
4.1	Specimen voucher of Prismatomeris glabra	22
4.2	Specimen voucher of Cinnamomum iners	23
4.3	Specimen voucher of Syzygium polyanthum	24
4.4	Specimen voucher of Actinodaphne sphaerocarpa	25
4.5	Specimen voucher of <i>Trema angustifolia</i>	26
4.6	Specimen voucher of Chromolaena odorata	27
4.7	Specimen voucher of Monocarpia marginalis	28
4.8	The agarose gel of PCR product (Sample 1)	31
4.9	The agarose gel of PCR product (Sample 2-5)	32
4.10	The ag <mark>arose gel of PCR product (Sample 6</mark> and 7)	33
4.11	The phylogenetic tree of selective aromatic plants from Ayer Hitam Forest Reserve, Selangor with selected aromatic plant species available in Malaysia	37

LIST OF ABBREVITION

AHFR Ayer Hitam Forest Reserve

BLAST Basic Loci Alignment Tool

bp Base pair

cpDNA Chloroplast DNA

DNA Deoxyribonucleic acid

MAPs Medicinal and Aromatic Plants

MEGA Molecular Evolutionary Genetics Analysis

NCBI National Centre for Biotechnology Information

NJ Neighbor-Joining

PCR Polymerase Chain Reaction

RNase Ribonuclease

SISFEC Sultan Idris Shah Forestry Education Centre

sp. Species

TAE Tris-acetate-EDTA

UCMP University of California Museum of Paleontology

UPM Universiti Putra Malaysia

USA United State of America

UV Ultraviolet

CHAPTER ONE

INTRODUCTION

1.1 General Background

Plants have various types of uses. Due to the variety uses of plants, they are playing big role as economic contributor in most of the countries that relaying on agriculture-based products. Other than as source of food, plants are also used as sources for non-food industrial products. Plants are the provider of textiles and fabric materials, which we make clothes from them. The raw materials especially woods are used world widely such as for constructions, buildings, crafts and furniture. Plants are also used in the production of biomass for energy resource. In other hand, many plant species are cultivated for their secondary metabolites, which useful for production of fine chemicals and other specialty products (Lubbe & Verpoorte, 2011).

Plants that used primarily for their medicinal or aromatic properties in pharmacy and perfumery are defined as medicinal and aromatic plants (MAPs) (Planta Europa, 2010). These including trees, herbs, shrubs, creepers, climbers and epiphytes. Medicinal and aromatic plants (MAPs) can be used for the production of essential oils, pharmaceutical, herbal health products, dyes and colorants, cosmetic and personal care products, plant protection products and intermediate products produced from the primary products (Lubbe & Verpoorte, 2011).

In Malaysia, the development of aromatic plants is still unclear. But it is known that most of the aromatic plants are used by the locals as traditional medicines, ornamental plants and source of foods. There is no record that the aromatic plants being planted in large scale for trading, except for *Aquilaria sp.* The *Aquilaria sp.* are being planted for their agarwood, which economically give great income for the people.

This study focused on seven selected species of aromatic trees in Ayer Hitam Forest Reserve, Puchong, Selangor. The selected species are *Trema angustifolia* (Cenderai/Mengkirai), *Cinnamomum iners* (Medang Teja), *Actinodaphne sphaerocarpa* (Medang Payung), *Syzygium polyanthum* (Serai Kayu), *Prismatomeris glabra* (Tongkat Haji Samad), *Monocarpia marginalis* (Mempisang) and *Chromolaena odorata* (Kapal Terbang). Most of the species come from different families, with few sharing the same family classification. This forest reserve is house for 430 tree species from 230 genus and 72 families (Faridah Hanum, 1999). The most diverse families are from Euphobiacea (39 species), Myrtaceae (26 species) and Lauraceae (23 species).

1.2 Problem Statement

AHFR encompasses varies plants species from different genus and families. Out of them, there are a few species that can be categorized as aromatic plants. Yet, there is no specific documentation about aromatic plants in AHFR. Plant lists and documentation are usually done via categorizing and classifying the plants by their species, genus and families. This is giving some difficulties to identify aromatic plants species in AHFR. In term of laborious, the lack of expertise in this molecular study by our locals, especially on aromatic plants can cause its less in genetic information.

Species identification by DNA sequencing method is useful and advantageous. This method serves as a supporting aid to the conventional plant identification through morphology characteristics. This method does not require the whole plants for species identification, but just needs a piece of dried or fresh tissue. Through this method, we can distinguish a group of aromatic plants in a short time by performing both botanical observation and molecular study. The results are useful information for conservation and economic purpose in the future. Aromatic plants contain extractives which give big contributions to economic development. This molecular study is important to establish the relatedness between the selected aromatic plants species.

1.3 Objectives

The objectives of this study are:

- To identify and select aromatic plants species in the AHFR via present publications and site-visits.
- II. To collect selected aromatic plants species samples and document their morphological characteristics.
- III. To study the phylogenetic relationship of the selected aromatic plants species by using PCR analysis.

1.4 Justification of Study

The results of this study might help to improve and develop method in species identification and classification based on morphological and molecular information. Through this study, the molecular information can help in species identification and classification. Commonly, species identification and classification are based on morphological information such as leaves, flowers, fruits and bark's scale. Yet, the identification of species based on morphological information may be wrong as its need broad knowledge in botany's field. Thus, the DNA sequences database can be reference as supporting information in plant identification.

Phylogenetic information is important and can be used for varies of purposes. It can be used for analysis of morphological and several kinds of molecular data. Furthermore, this information can be used in making estimation of evolutionary relationships and the relatedness among the selected aromatic plants species. However, the accurate estimates of phylogenetic relationships between the selected species are still lacks. This is a limitation that must be faced by ecology community from apply the evolutionary information within their field and studies. Therefore, this study is quite useful and has its own priority due to its importunateness in supplying molecular information.

REFERENCES

American Botanical Council (2013). Retrieved from http://abc.herbalgram.org/on 3rd October, 2015.

Anderberg, A. A. (1991). Taxonomy and phylogeny of the tribe Gnaphalieae (Asteraceae). *Opera Botanica*, *104*, 1-195.

Bawon, P., & Yaman, A. R. (2007). Ayer Hitam Forest Reserve: Multimedia, Super Corridor, Community Heritage. Faculty of Forestry, Universiti Putra Malaysia.

Bremer, K., & Anderberg, A. A. (1994). *Asteraceae: cladistics and classification*. Portland (Oregon). *Timber Press*.

Briggs, B. G., & Johnson, L. A. S. (1979). Evolution in the Myrtaceae-evidence from inflorescence structure. In *Proceedings on Linnean Society of New South Wales*, *102*(4), 157-256.

Frankham, R., Briscoe, D. A., & Ballou, J. D. (2002). *Introduction to conservation genetics. Cambridge University Press.*

Frankham, R. (2005). Genetics and extinction. *Biological conservation*, *126*(2), 131-140.

Fries, R. E., & Pulle, A. (1940). Annonaceae. Flora of Suriname, 2(2), 341-83.

Gittleman, J. L., Anderson, C. G., Kot, M., & Luh, H. K. (1996). Phylogenetic lability and rates of evolution: a comparison of behavioral, morphological and life history traits. *Phylogenies and the comparative method in animal behavior*, 166-205.

Govaerts, R., Sobral, M., Ashton, P., Barrie, F., Holst, B. K., Landrum, L. L., & Soares-Silva, L. H. (2008). *World checklist of Myrtaceae*. Royal Botanic Gardens.

Grudzinskaya, I. A. (1988). On the taxonomy of the family Cannabaceae. Botanichnyi Zhurnal Society for the Scientific Study of Reading, 73, 589-93.

Hall T.A. (1999). BioEdit: A user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. *Nucleic Acids Symposium Series*, 41: 95-98.

Hanum, I. F., Ibrahim, A. Z., Khamis, S., Nazre, M., Lepun, P., Rusea, G., & Latiff, A. (2001). An annotated checklist of higher plants in Ayer Hitam Forest Reserve, Puchong, Selangor. *Pertanika Journal of Tropical Agricultural Science*, *24*(1), 63-78.

Heath T.A., Hedtke S.M., & Hillis D.M. (2008). Taxon sampling and accuracy of phylogenetic analyses. *Journal of Systematics and Evolutions*, *46*(3), 239-257.

Ibrahim, F. H. (1999). Plant diversity and conservation value of Ayer Hitam forest, selangor, peninsular Malaysia. *Pertanika Journal of Tropical Agricultural Science*, 22(2), 73-83.

Innis, M. A., Gelfand, D. H., Sninsky, J. J., & White, T. J. (2012). *PCR protocols: a guide to methods and applications. Academic press.*

Inoue, M., & Craker, L. E. (2014). Medicinal and Aromatic Plants—Uses and Functions. In *Horticulture: Plants for People and Places, Volume 2* (pp. 645-669). *Springer Netherlands*.

Jantan, I. B., & Goh, S. H. (1992). Essential oils of Cinnamomum species from Peninsular Malaysia. *Journal of Essential Oil Research*, *4*(2), 161-171.

Kim S.C., & Crawford D.J. (1997). The use of noncoding region of chloroplast DNA in phylogenetic studies of the subtribe Sonchinese (Asteraceae: Lactuceae). *Plant System Evolution*, *215*, 85-99.

King, R. M., & Robinson, H. (1987). The genera of the Eupatorieae (Asteraceae). *Monographs in systematic botany*, 22.

Lee, S. Y., & Mohamed, R. (2016). Rediscovery of Aquilaria rostrata (Thymelaeaceae), a species thought to be extinct, and notes on Aquilaria conservation in Peninsular Malaysia. *Blumea*, *61*, 13-19.

Lubbe, A., & Verpoorte, R. (2011). Cultivation of medicinal and aromatic plants for specialty industrial materials. *Industrial Crops and Products*, *34*(1), 785-801.

Ma, X. Y., Xie, C. X., Liu, C., Song, J. Y., Yao, H., Luo, K., ... & Chen, S. L. (2010). Species identification of medicinal pteridophytes by a DNA barcode marker, the chloroplast psbA-trnH intergenic region. *Biological and Pharmaceutical Bulletin*, 33(11), 1919-1924.

Nor Aini. A.S. 1999. A Preliminary Study of Genetic of Selected Species of a Lowland Forest at Ayer Hitam Forest Reserve, Selangor. *Pertanika Journal Tropical Agriculture Science*, 22(2), 111-116.

Planta Europa (2013). Retrieved from http://www.plantaeuropa.net/ on 1st October, 2015.

Richter, H. G. (1981). Wood and bark anatomy of Lauraceae. I. Aniba Aublet. *IAWA Journal*, 2(2-3), 79-87.

Robbrecht, E. (1988). Tropical woody Rubiaceae. *Opera Botanica Belgica*, 1(272), 599-602.

Robbrecht, E. (1988). *Tropical woody Rubiaceae. Characteristic features and progressions. Contributions to a new subfamilial classification.* Nationale Plantentuin van België.

Rohwer, J. G. (1993). Lauraceae (pp. 366-391). Springer Berlin Heidelberg.

Sarah, A. R., Nuradnilaila, H., Haron, N. W., & Azani, M. (2015). A Phytosociological Study on the Community of Palaquium gutta (Hook. f.) Baill.(Sapotaceae) at Ayer Hitam Forest Reserve, Selangor, Malaysia. *Sains Malaysiana*, *44*(4), 491-496.

Schä, H., Bernard, L., Courties, C., Lebaron, P., Servais, P., Pukall, R., & Muyzer, G. (2001). Microbial community dynamics in Mediterranean nutrient-enriched seawater mesocosms: changes in the genetic diversity of bacterial populations. *FEMS Microbiology Ecology*, *34*(3), 243-253.

The Plant List (2014). Version 3.0. Retrieved from http://www.theplantlist.org/on 25th September, 2015.

Toro, M. A., & Caballero, A. (2005). Characterization and conservation of genetic diversity in subdivided populations. *Philosophical Transactions of the Royal Society of London B: Biological Sciences*, *360*(1459), 1367-1378.

van der Werff, H., & Richter, V. D. H. (1996). Toward an improved classification of Lauraceae. *Annals of the Missouri Botanical Garden*, 409-418.

Verdcourt, B. (1958). Remarks on the classification of the Rubiaceae. *Bulletin du Jardin botanique de l'Etat, Bruxelles/Bulletin van den Rijksplantentuin, Brussel*, 209-290.

Van Heusden, E. C. H. (1992). Flowers of Annonaceae: morphology, classification, and evolution. *Blumea. Supplement*, 7(1), 1-218.

Walsh, N. G., & Entwisle, T. J. (1996). Flora of Victoria. Volume 3: Dicotyledons Winteraceae to Myrtaceae. Inkata Press.

Yang, M. Q., van Velzen, R., Bakker, F. T., Sattarian, A., Li, D. Z., & Yi, T. S. (2013). Molecular phylogenetics and character evolution of Cannabaceae. *Taxon*, 62(3), 473-485.