

### **UNIVERSITI PUTRA MALAYSIA**

### EFFECTS OF GRACILARIA CHANGII EXTRACT ONAPOPTOSIS AND GENE EXPRESSION OF MCF-7AND MDA-MB-23 1 BREAST CANCER CELL LINES

### NAFEZAH ABDUL HAMID

FPSK(M) 2006 10

# 06 AUG 2008

# EFFECTS OF *GRACILARIA CHANGII* EXTRACT ON APOPTOSIS AND GENE EXPRESSION OF MCF-7 AND MDA-MB-231 BREAST CANCER CELL LINES

# NAZEFAH ABDUL HAMID

# MASTER OF SCIENCE UNIVERSITI PUTRA MALAYSIA

2006



# EFFECTS OF *GRACILARIA CHANGII* EXTRACT ON APOPTOSIS AND GENE EXPRESSION OF MCF-7 AND MDA-MB-231 BREAST CANCER CELL LINES

# NAZEFAH ABDUL HAMID

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

May 2006



#### DEDICATION

This thesis is dedicated to my loving family, who has been supporting me through thick and thin. Without them, none of this would have been possible.



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

#### EFFECTS OF *GRACILARIA CHANGII* EXTRACT ON APOPTOSIS AND GENE EXPRESSION OF MCF-7 AND MDA-MB-231 BREAST CANCER CELL LINES

By

#### NAZEFAH ABDUL HAMID

#### May 2006

#### Chairman: Associate Professor Rozita Rosli, PhD

#### Faculty : Medicine and Health Sciences

Cancer is a large group of diseases characterized by uncontrolled growth and spread of abnormal cells. Hundreds of research studies have demonstrated significant benefit of the use of natural products in the treatment of cancer and scientists believe examining new natural products will continue to turn up even more useful drugs to treat cancer. Marine organisms are a rich source for natural products and many compounds that are derived from these have generated interest for their cytotoxicities. *Gracilaria changii* is a type of red seaweed which comes from the family *Rhodophyta*. It is a relatively abundant seaweed in Malaysia can be found in the mangrove areas. In this study, the chemotherapeutic potential of *Gracilaria changii* in selected reproductive cancer cell lines was evaluated together with tamoxifen, a commercially used drug in cancer treatment. Exposure of breast, ovarian and cervical cancer cell lines, to a range of *Gracilaria changii* extracts demonstrated growth inhibition in some of these cancer cells in a dose-dependent manner. The *Gracilaria changii* extracts received from Kolej

iii

Universiti Sains dan Teknologi Malaysia (KUSTEM) were methanol, buthanol and diethyl ether extracts. The methanol extract gave the most promising IC50 values, as determined by the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) reduction assay and the results are as follows: MCF-7 (7.8 µg/ml), MDA-MB-231 (25  $\mu$ g/ml) HeLa (70.26  $\mu$ g/ml) and Caov-3 (90.46  $\mu$ g/ml). Since the results for the breast cancer cell lines were significant compared to the ovarian and cervical cancer cell lines, they were chosen for further analysis. The normal breast cell line, MCF-10A was also tested and the IC<sub>50</sub> value was found to be > 1000  $\mu$ g/ml, indicating that the methanol extract was not cytotoxic to normal cells. AOPI staining was used to study the morphology of the cells treated with the extract. Apoptotic features that included membrane blebbing and nucleus condensation were evident in MCF-7 and MDA-MB-231 cancer cells. Subsequently, the TUNEL assay was conducted to determine and quantitate the apoptotic cells within a cell population. The results suggest that the methanol extract was better of inducing cell death by stimulating apoptosis than tamoxifen. This is based on the significantly higher percentage of apoptotic cells in the G.changii methanol extract treated cancer cells as compared to tamoxifen. For MCF-7 and MDA-MB-231 cell lines, p was <0.01 when compared with control (24 hours) and Pof <0.001 when compared with control for 48 hours. In addition, gene expression analysis was performed using the microarray technology. This technology which allows the simultaneous analysis of a large number of nucleic acid hybridization experiments and was carried out to determine the gene expression profile. Preliminary work on micorarray was conducted using MCF-7 cell line only, due to time constraints and limited funding. Upon treatment with the methanol extract on MCF-7, several suppressed genes were



identified including haplotype n1b mitochondrion complete genome, melanomaassociated antigen p97 isoform 1 and damage-specific DNA binding protein 2 (ddb2). The results showed that the three genes regulated by the methanol extract encode proteins that belongs to DNA repair, protection against membrane-lipid peroxidation and maternal inheritance family, which may play an important role for the cancer treatment. It was further confirmed using Reverse Transcription Polymerase Chain Reaction (RT-PCR). Therefore, the methanol extract of *Gracilaria changii* is a potential candidate to be developed as a chemotherapeutic agent in the treatment of estrogen receptor-positive breast cancers.



#### EFFECTS OF *GRACILARIA CHANGII* EXTRACT ON APOPTOSIS AND GENE EXPRESSION OF MCF-7 AND MDA-MB-231 BREAST CANCER CELL LINES

Oleh

#### NAZEFAH ABDUL HAMID

#### Mei 2006

#### Pengerusi : Profesor Madya Rozita Rosli, PhD

#### Fakulti : Perubatan dan Sains Kesihatan

Kanser merupakan satu kumpulan penyakit yang dikategorikan sebagai tumbesaran sel yang tidak dapat dikawal dan penyebaran sel-sel yang tidak normal. Sel-sel ini boleh membesar menjadi kelompok tisu yang dipanggil tisu malignan. Apabila kanser merebak, ia akan menyerang dan memusnahkan tisu normal dan juga akan merebak ke bahagian lain badan. Kanser yang merebak ke bahagian lain atau ke organ lain dipanggil kanser metastatik. Jika penyebaran kanser tidak dikawal, ia akan menyebabkan kematian. Kebanyakan kanser boleh dikawal atau dicegah jika dikesan awal dan dirawat dengan segera. Kebanyakan penyelidikan yang telah dijalankan menunjukkan kesan yang signifikan daripada penggunaan alam semulajadi dan saintis mempercayai bahawa ujianujian terhadap alam semulajadi yang baru akan membuahkan lebih banyak ubat antikanser. Organisma marin kaya dengan sumber alam semulajadi dan banyak sebatian daripadanya telah menarik minat terhadap kesan sitotoksik. *Gracilaria changii* merupakan rumpai laut merah daripada keluarga *Rhodophyta*. Ia merupakan rumpai laut



yang banyak terdapat di Malaysia dan boleh didapati di kawasan paya bakau. Di dalam kajian ini, potensi kemoterapeutik Gracilaria changii untuk kanser reproduktif terpilih dikaji bersama tamoxifen, drug antikanser komersil yang sekarang digunakan untuk rawatan kanser. Pendedahan sel-sel kanser payudara, ovari dan serviks kepada julat kepekatan ekstrak yang berbeza menghasilkan perencatan pertumbuhan bagi kedua jenis sel kanser dengan bergantung kepada nilai kepekatan. Ekstrak yang diterima daripada Kolej Universiti Sains dan Teknologi Malaysia (KUSTEM) merupakan ekstrak metanol, butanol dan dietil eter. Ekstrak methanol memberikan nilai IC<sub>50</sub> yang paling baik, didapati daripada asai penurunan 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) adalah seperti berikut: MCF-7 (7.8 ug/ml), MDA-MB-231 (25 ug/ml) HeLa (70.26 ug/ml) and Caov-3 (90.46 ug/ml). Oleh kerana sel selanjar payudara memberikan keputusan yang baik berbanding dengan sel-sel selanjar ovari dan servik, maka ia dipilih untuk analisis seterusnya. Sel selanjar normal, MCF-10A juga diuji dan nilai IC<sub>50</sub> lebih daripada 1000 ug/ml, menunjukkan ekstrak metanol tidak sitotoksik terhadap sel normal. Pewarnaan AOPI digunakan untuk melihat perubahan morfologi sel selepas rawatan dengan ekstrak tersebut. Ciri-ciri apoptosis seperti gelembung membran dan kondensasi nukleus telah didapati bagi sel kanser MCF-7 dan MDA-MB-231. Seterusnya, asai TUNEL dijalankan untuk menentukan apoptosis. Menariknya, ekstrak metanol didapati menyebabkan kematian sel dengan merangsang apoptosis lebih baik daripada tamoxifen berdasarkan peratusan sel-sel apoptotik yang menunjukkan perbezaan yang signifikan berbanding sel-sel kanser dengan rawatan ekstrak metanol. Sebagai tambahan, analisis pengekspresan gen dijalankan dengan menggunakan teknologi mikroarray. Teknologi mikroarray membenarkan analisis terhadap asid nukleik yang



banyak secara serentak bertujuan untuk menentukan profil pengekspresan gen. Sebagai permulaan, kajian mikroarray hanya dijalankan ke atas sel selanjar MCF-7 sahaja, disebabkan factor masa dan geran yang terhad.Daripada rawatan ekstrak methanol terhadap MCF-7, didapati tiga gen yang telah ditindas iaitu haplotype n1b mitochondrion complete genome, melanoma-associated antigen p97 isoform 1 dan damage-specific DNA binding protein 2. Keputusan kajian menunjukan gen tersebut terdiri daripada gen yang mengkodkan protein kumpulan pembaikkan DNA, perlindungan terhadap peroksidasi membrane-lipid dan warisan daripada ibubapa, yang mungkin memainkan peranan yang penting terhadap rawatan kanser. Seterusnya ia disahkan dengan menggunakan Reverse Transcription Polymerase Chain Reaction (RT-PCR). Oleh itu, ekstrak methanol *Gracilaria changii* didapati mempunyai potensi untuk dikembangkan sebagai agen kemoteraputik untuk rawatan kanser yang bergantung kepada hormone.



#### ACKNOWLEDGEMENTS

I am giving all my thanks to Allah S.W.T whose blessings have accompanied me with the energy, time and ideas in every step of the way in finishing this work and made it a possible task.

This project would have never been initiated and completed without the persistent help from so many people. Greatest debt of gratitude therefore is owed to my supervisor, Assoc.Prof. Dr. Rozita Rosli whose inspiration, gratitude, advice and unwavering support made the completion of this project a reality. I thank her again for trusting me so much in this project and all the help she provided during the accomplishment of this project. I am also grateful to my co-supervisor, En. Muhammad Nazil Salleh for the comments, suggestions and attention in this project.

Very special thanks to K. Nurma for her invaluable contribution in knowledge and guidance as well as tirelessly accompanying me throughout the course of my project. Recognition also goes to members of the Molecular Genetics lab, Wong, King Hwa, Syaban, Lama, Nasir, K. Za, Chan, Chin, Radha and K. Shariza Sr. Not forgetting Molecular Microbiology's lab members, Matun, Mas, Farah, Lai and Hana for their support and helping hands.

Loving thanks to all my collegues, Yati, Zet, Fatan, Arie, Ajantha, Nirmala, Ahmad, Pike See, Yunus and Im, who assisted me with their invaluable advice and encouragement.



Special thanks to Thila, K. Shahriza and K. Sue, who kept on providing encouragement and help along the journey.

My heartiest gratitude and sincere thanks to my family for their love, encouragement and support in completing this project. On a personal note, I would like to express my gratitude to my beloved husband, Nizam Nordin for his support and concern all this while. Words cannot describe their patience and kindness in helping and guiding me in every part of this project.

Last but not least, I would like to thank to everyone who have helped me directly or indirectly, towards the completion of this research project.



# 06 AUG 2008

## TABLE OF CONTENTS

DEDICATION	ii
ABSTRACT	iii
ABSTRAK	vi
ACKNOWLEDGEMENTS	ix
APPROVAL	xi
DECLARATION	xiii
LIST OF TABLES	xvii
LIST OF FIGURES	xviii
LIST OF ABBREVIATIONS	XX

### CHAPTER

1	INTE	RODUCTION	
	1.1	Cancer	1
	1.2	Significance of study	2
	1.3	Objectives	3
2	LITE	ERATURE REVIEW	
	2.1	Natural products in cancer treatment	4
	2.2	Gracilaria changii	7
	2.3	Cancer Cases in Malaysia	10
	2	.4 Breast cancer	
		2.4.1 Overview	11
		2.4.2 Epidemiology of Breast Cancer	12
		2.4.3 Treatment of Breast Cancer	14
		2.4.4 Tamoxifen as treatment in breast cancer	16
		2.4.5 Side effects of tamoxifen	17
	2.5	Breast Cancer Cell Lines	
		2.5.1 MCF-7	18
		2.5.2 MDA-MB-231	19
		2.5.3 MCF-10A	19
	2.6	Apoptosis	
		2.6.1 Overview	20
		2.6.2 Apoptosis and Cancer Chemotherapy	25
	2.7	Microarray	
		2.7.1 Overview	26
		2.7.2 Microarray and breast cancer	28



Page

2.8	Rever	Reverse Transcription Polymerase Chain Reaction (RTPCR)		
		2.8.1 Overview	31	
3	МАТ	ERIALS AND METHODS		
-	3.1	General outline of the study	35	
	3.2	Gracilaria changii extracts	37	
	3.3	Cancer cell lines	37	
	3.4	Cell Culture		
		3.4.1 Preparation of media	37	
		3.4.2 PBS-EDTA solution for washing the cell lines	39	
		3.4.3 Trypsinizing solution	39	
	3.5	Cell culture protocols		
		3.5.1 Thawing cryopreserved cells	40	
		3.5.2 Media renewal	40	
		3.5.3 Subculture of adherent cells	41	
		3.5.4 Cell plating	42	
		3.5.5 Cryopreservation	43	
	3.6	MTT Cytotoxicity Assay	43	
	3.7	Morphological Studies Using Phase Contrast Microscope	45	
	3.8	Acridine Orange/Propididum Iodide (AOPI) Staining	45	
	3.9	Detection of Apoptosis Using DeadEnd <sup>tm</sup> Fluorimetric TUNEL System	46	
	3.10	Statistical Analysis	48	
	3.11	Microarray	48	
		3.11.1 Spotted slide		
		Array fabrication	50	
1		Resuspension of oligonucleotides	50	
		Array printing	51	
		Post treatment	51	
		Slide Quality Control	52	
		3.11.2 Commercially available slide		
		MWG slide	52	
		Extraction of Total RNA	53	
		RNA Quantification		
		RNA Quantification by	54	
		spectrophotometer		
		Electrophoresis of RNA	54	
		cDNA Synthesis	54	
		Double stranded DNA Purification	55	
		T7 Transcription of cRNA and	56	
		Labeling of cRNA		
		Fragmentation of labeled cRNA	56	
		Hybridization	57	
		Washing	57	
		Scanning	58	



	Data Analysis	58
3.11	Reverse Transciption Polymerase Chain Reaction (RTPCR)	
	3.11.1 Primer design	59
	3.11.2 Reverse Transcription	59
	3.11.3 Polymerase Chain Reaction (PCR)	59
	3.11.4 Agarose Gel Electrophoresis	61
	3.11.5 Agarose Gel Formula	61
	3.11.6 Electrophoresis	61

#### **RESULTS**

4.1	Cytotoxicity assay	63
4.2	Morphological studies using phase contrast	69
	inverted microscope	
4.3	Cell Viability and Apoptosis Assay using AO/PI Staining	77
4.4	TUNEL assay	85
4.5	Microarray	97
DISC	CUSSION	
5.1	Effect of methanol extract of Gracilaria changii on	118
	breast cancer cell lines and induction of apoptosis	
5.2	Місгоантау	124
CON	CLUSION	
6.1	Conclusion	132
6.2	Achievement of the projected objectives	133
6.3	Limitations of the study	134
6.4	Future studies	134

#### **REFERENCES BIODATA OF AUTHOR**



### LIST OF TABLES

Table		Page
1	Differential features and significance of necrosis and apoptosis, including morphological, biochemical and physiological features.	23
2	Temperature and time of PCR parameters that were optimized for the three genes.	60
3	MTT assay results on the five cell lines. Data represents the mean $IC_{50}$ value and standard error (SEM) for each of the cell lines tested with the extracts and tamoxifen.	65
4	Chemotherapeutic significance values for anticancer screening according to NCI standard for crude extract.	66
5	The concentration and purity total of RNA extracted from MCF-7. Concentration of RNA extracted is expressed as $\mu$ g/ml.	102
6	List of genes in the block on the commercial slide which were analyzed using QuantArray software.	105
7	Genes regulated by methanol extract in MCF-7. For fold changes, (-) indicates suppression. Significant differences were determine according to cut-off values of two-fold or greater.	108 d
8	Gene-specific primers designed for RT-PCR. The primers were designed using Primer Premier 5.0 software.	114



xvii

#### LIST OF FIGURES

Figur	'e	Page
1	Gracilaria changii harvested from Morib, Selangor.	9
2	Ten most frequent cancers in females, Peninsular Malaysia 2003.	11
3	Structure of tamoxifen.	16
4	The general pipeline for the study outlining methods used in the study.	36
5	Pipeline for microarray.	49
6	Effect of increasing concentrations of methanol extract and tamoxifen on the cell viability.	67
7	The effect of methanol extract and tamoxifen on MCF-10A	68
8	Changes in morphology of MCF-7 cells after treatment with methanol extract and tamoxifen, 24 hours.	71
9	Morphological changes (inverted microscopy) in control and treated cells, 48 hours (MCF-7).	72
10	Morphological changes of MCF-7 cells treated with methanol extract and tamoxifen at 72 hours.	73
11	Morphological analysis of MDA-MB-231 cells after 24 hours exposure to methanol extract and tamoxifen.	74
12	Methanol-induced morphologic changes characteristic of apoptosis in MDA-MB-231 cell lines for 48 hours.	75
13	Changes in morphology of MDA-MB-231 cells after treatment with methanol extract and tamoxifen (72 hours).	76
14	Fluorescent microscopy analysis of the morphological changes in MCF-7 cell lines.	79
15	Fluorescence images of MCF-7 treated for 48 hours.	80
16	MCF-7 cell line treated for 72 hours.	81
17	Morphological changes of MDA-MB-231 undergoing apoptosis. (24 hours)	82



18	MDA-MB-231 cell line treated for 48 hours.	83
19	Fluorescent images MDA-MB-231 treated for 72 hours.	84
20	TUNEL assay performed on MCF-7 at day 1 (24 hours).	87
21	TUNEL assay performed on MCF- 7 at day 2 (48 hours).	88
22	Effect of methanol extract on MDA-MB-231 cells at 24 hours.	89
23	Fluorescent TUNEL labeling of methanol treated MDA-MB-231 cells at 48 hours.	90
24	TUNEL assays demonstrate the differential induction of apoptosis in MCF-10A at 24 hours treatment.	91
25	TUNEL assay performed on MCF-10A at day 2 (48 hours).	92
26	Percentage of apoptosis induced by methanol extract and tamoxifen on MCF-7.	94
27	Percentage of MDA-MB-231 cells stained positive for apoptosis using TUNEL assay.	95
28	Quantification of apoptosis effects of methanol extract on MCF-10A for 24 and 48 hour.	96
29	Slide design which contains 3 replicates of 380 genes.	98
30	A slide scanned without using any dye.	99
31	Spotcheck slide scanned using Cy3 channel.	101
32	A representative of the quality of RNA from untreated and untreated MCF-7.	102
33	Image scanned from the microarray slide after hybridization.	104
34	Scatter plot of fluorescence intensity values from MWG slide hybridized with untreated and treated MCF-7.	111
35	Histogram of Cy3 and Cy5 ratio according to each of the genes.	112
36	Histogram of M log <sub>2</sub> vs. A log <sub>2</sub>	113
37	RT-PCR analysis of ddb2 expression in MCF-7.	115
38	RT-PCR analysis of mfi2 expression in MCF-7.	116
39	RT-PCR analysis of haplotype nlb expression in MCF-7.	117

### LIST OF ABBREVIATIONS

μΙ	Microliter
μΜ	Micromolar
ATCC	American Type Culture Collection
cDNA	complementary deoxyribonucleic acid
cRNA	complementary ribonucleic acid
DMSO	Dimethylsulfoxide
dNTP	Deoxyribonucleotide triphosphate
dsDNA	Double stranded deoxyribonucleic acid
EDTA	Ethylenediaminetetra-acetic acid
mg	Milligram
ml	Milliliter
mm	Millimeter
MTT	3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide
mV	Millivolt
PBS	Phosphate buffer saline
pmol	picomole
RNA	Ribonucleic acid
RT-PCR	Reverse Transcription Polymerase Chain Reaction
SDS	Sodium Deodecyl Sulphate



#### **CHAPTER 1**

#### **INTRODUCTION**

#### 1.1 Cancer and natural product

Cancer is a collective term that covers hundreds of different diseases characterised by invasive and uncontrolled cell growth. It is a chronic disease that can consist of specific stages, from genetic predisposition through various forms of premalignant and malignant degeneration of cells to disease progression. Breast cancer is one of the most common cancers diagnosed in women worldwide and is a leading cause of cancer-related deaths (Greenlee *et al.*, 2000). In Malaysia, as of 2003, it accounted for 31 % of newly diagnosed female cases, and was the commonest cancer in all ethnic groups and all age groups in females from the age of 15 years (Lim and Halimah, 2004).

Mortality that results from the common forms of cancer is still unacceptably high. Natural or semisynthetic compounds may be used to block, reverse, or prevent the development of invasive cancers. Cellular carcinogenesis forms the biological basis for the identification of preventive products, the assessment of their activity, and ultimately the success or failure of a therapy (Reddy *et al.*, 2003).

Ideally, chemotherapeutic drugs should specifically target only neoplastic cells and should decrease tumor burden by inducing cytotoxic and/or cytostatic effects with minimal "collateral damage" to normal cells. (Ricky *et al.*, 2002). Many pharmaceutical



agents have been discovered by screening natural products from plants, animals, marine organisms and microorganisms. Despite major scientific and technological progress in combinatorial chemistry, drugs derived from natural products still make an enormous contribution to drug discovery today (Rocha *et al.*, 2001).

Epidemiological data indicated that ubiquitous consumption of seaweeds in Japan may be a possible protective factor against some types of tumor (Okai *et al.*, 1994). Therefore in this study, *Gracilaria changii*, from the family of *Rhodophyta* or red seaweed was chosen. It is indigenous agarophytic seaweed in Malaysia (Phang, 1994). *Gracilaria changii* when used as food provides substantial amounts of fiber, minerals, lipids and protein (Norziah *et al.*, 2000). At present, this seaweed is only consumed in certain coastal areas especially along the east coast of Peninsula Malaysia and in East Malaysia.

#### 1.2 Significance of study

In recent years, improved diagnostic tools have made it possible to detect breast cancers at early, even pre-invasive stages leading to a significant decrease in breast cancer mortality rates over the past decades. Breast cancer has been the major killer in women all over the world, and the number is increasing year by year. The use of natural product in cancer treatment has shown good results, and extensive research is being carried out. In Japan, the cancer mortality rate is among the lowest in the world, and this has been associated with having seaweed in their dietary intake. Thus, this study is hoped to



highlight on our own seaweed as a potential anticancer agent. The molecular studies of *Gracilaria changii* on breast cancer may contribute insight on drug targets and formulation against novel apoptosis pathway.

#### 1.3 Objectives

There has been no previous study reported on the effect of *Gracilaria* changii on cancer cells. Thus, the objectives of the study are:

- to determine whether *Gracilaria changii* extract is effective in inhibiting the proliferation of selected breast cancer cell lines (MCF-7, MDA-MB-231), cervical cancer (HeLa) and ovarian cancer cell lines (Caov-3).
- to select the cell lines most effective against the extracts and comparing this with a normal cell line.
- 3) to evaluate apoptosis inducing ability of the *Gracilaria changii* extract by morphology through the apoptotic features of the cells and to confirm through quantification using TUNEL assay
- 4) to determine the gene expression profile following treatment of the extract on the cancer cells using microarray technology and to validate the expression using RT-PCR.

