



Identification Of Some Genetic Markers In Familial Breast Cancer In Egyptian Patients Thesis

Submitted for Partial Fulfillment of M.Sc. Degree In Clinical and Chemical Pathology

By
Sally Gharib Mohammed Abd Allah
M.B., B.Ch
Cairo University

Supervised by Professor. Dr. Mervat Al Ansary

Professor of Clinical and Chemical Pathology Faculty of Medicine - Cairo University

Doctor. Asaad El Gerzawy

Assistant Professor of Human Cytogenetics Human Cytogenetics Department Human Genetics and Genome Research Division National Research Center

Doctor. Iman Loay

Lecturer of Pathology National Cancer Institute Cairo University

Faculty of Medicine Cairo University 2012

Abstract

Background:

The Her- 2/neu gene encodes a receptor related to breast carcinogenesis and topoisomerase IIalpha (TOP2A) gene is located adjacent to Her-2/neu oncogene on chromosome 17 and encodes an enzyme plays a key role in DNA replication and is a target for multiple chemotherapeutic agents like anthracyclines.

The aim of the present study: was to evaluate cases of familial breast cancer (FBC) and compare them with sporadic cases (SBC) through hormonal status, immunostaining for Her-2/neu and TOP2A and Her-2/neu copy number alterations by FISH, and investigate the prevalence of alteration of TOP2A gene copy number in familial breast cancer.

Material and methods: 22 cases of invasive breast carcinomas: 12 cases with criteria that define familial breast cancer (FBC), and 10 cases of sporadic breast cancer (SBC) were involved in this study. Archival blocks of formalin fixed-paraffin embedded sections of these cases were used, these sections were subjected to immunostaining with Her-2/neu (Hercep test), ER (estrogen receptors) and PR (progesterone receptors), and Fluorescence in situ hybridization (FISH) was performed to evaluate TOP2A and Her-2/neu genes copy number alterations.

Results: revealed that the FBC tumors were more aggressive than SBCs regarding; pathological parameters (larger tumor size, and higher grade), and immunohistochemical markers (lower ER, and PR expression among FBCs), but the rate of Her-2/neu gene amplification and protein over-expression was equally likely between both groups. The rate of TOP2A gene amplification was 22.7%, co-amplification of Her-2/neu and TOP2A

genes was found in one case of the FBS patients (8.3%) and in two cases of the SBCs (20%). In contrast, Her-2/neu amplification alone was found in three patients of the FBCs (25%), compared to one case among the sporadic breast cancers (10%). Importantly, TOP2A amplification with normal Her-2/neu gene status was found in two of the FBCs (16.7%), but in none of the sporadic tumors (0%).

Our findings have potential therapeutic implications. Her-2/neu assessment is routinely used to select breast cancer patients for trastuzumab but also dose intensive anthracycline therapy. Our data suggest that FBCs also need to be tested for TOP2A amplification.

Conclusion: The clinical, pathological, and immunohistochemical characteristics of tumors should be of value in evaluating the genetic basis of breast cancer among families, as FBC tumors were more aggressive than SBCs. Identifying biological characteristics that can predict genetic background among breast cancer families and as a clinical consequence, we suggest that patients with suspected FBCs need more intensive therapy and careful follow up.

TOP2A gene amplification is more frequent in FBC than SBC and it may occur independent of Her-2/neu gene amplification, this finding together with the fact that TOP2A is the therapeutic target of anthracyclines suggests that this parameter should be determined on a routine basis in BC and especially in FBC.

Key Words:

Breast Cencar – TOP2A – Her-2/neu.

Acknowledgement

First of all, great thanks to **ALLAH** who is most merciful, for all the countless gifts I have been offered.

I would like to express my endless gratitude and appreciation to **Professor Dr. Mervat Mohammed Saad Al Ansary** Professor of Clinical and Chemical Pathology, Cairo University, for her continuous guidance, valuable suggestions and keen supervision throughout the work.

I would like to express my most sincere gratitude to **Professor Dr.**

Amal Mahmoud, Professor and Head of Human Genetics department,
National Research Center who has been there to guide me throughout my
studies. I encountered many challenges during my research and Dr. Amal
was always there to provide support and encouragement. I thank Dr. Amal
for sharing her enthusiasm and passion for science and research. I
especially thank her for the confidence she has instilled in me during my time
here that will help me in my future endeavors. Lastly, I thank her for sharing
some of her stories with me about her life outside of science, inspiration and
advice that will carry a long way.

I would like to express my sincere gratitude and deepest appreciation to **Professor. Sayeda Hammad**, Professor of Human Genetics, National Research Center, for her instructive supervision & encouragement, scientific and personal support throughout the course of this work.

My endless thanks to **Dr. Asaad El Gerzawy**, Assistant Professor of Human Genetics, National Research Center, for his valuable instructions, and support to get best out of this work.

Indeed, words do fail me when I come to express my profound gratitude and deep appreciation to **Dr. Iman Loay**, Lecturer of Pathology, National Cancer Institute, Cairo University for her everlasting support, honest encouragement and for offering me much of her time and effort.

I would like to thank **Dr. Fatma Abu Al Kasem**, Lecturer of Medical oncology, National Cancer Institute, Cairo University for providing me the clinical data needed to achieve this work.

Last, but not least, I would like to thank **my family**, especially my mom for their actual help, unconditioned love, and support received in many ways.

Sally Gharib

List of Contents

		Page
Ø	List of Abbreviations	
Ø	List of Tables	
Ø	List of Figures	
Ø	Introduction	1
Ø	Aim of the Work	4
Ø	Review of Literature	5
<u>Ch</u>	apter 1: PATHOLOGY OF BREAST CANCER	5
I. N	ormal Histological structure of the breast	5
II. Incidence of breast cancer		6
III. Risk factors of breast cancer		7
IV. Prognostic and predictive factors of breast cancer		14
V. Prognostic factors of breast cancer		16
Chapter 2: GENETICS OF BC		23
I. Criteria of FBC		24
II. The risk of FBC versus SBC		25
III.	Genes involved in breast carcinogenesis	26
IV.	Genes involved in familial breast cancer	30
<u>Ch</u>	apter 3: Her-2/neu GENE IN BC	46
I. T	he role of Her-2/neu in the development of breast cancer	46
II.	Her-2/ neu in carcinogenesis	50
III.	Mechanism of Her-2/neu amplification	51
IV.	The role of Her-2/neu in breast cancer	55
V. F	Ier-2/neu gene status in FBC versus SBC	61
VI.	Her-2/neu testing	62

<u>Chapt</u>	ter4: TOP2A GENE IN BC	69
I. Top	ology of DNA	69
II. Cla	sses of topoisomerases	71
III. DI	NA Topoisomerase II genes	72
IV. Str	ructural characterization	73
V. Me	echanism of action	75
VI. Tis	ssue distribution	77
VII. Subcellular localization of topoisomerases VIII. Cell cycle changes in protein level IX. Regulation of cell cycle changes in protein level X. Functions of topoisomerases XI. Topoisomerase II as a drug target		77
		78
		79
		80
		80
XII. Clinical significance of TOP2A		82
\geq S	ubjects and Methods	91
≥ R	Results	115
> Γ	Discussion	153
\geq S	ummary and Conclusion	171
≥ R	Recommendations	175
≥ R	References	176
\triangleright A	Arabic Summary	238

LIST OF ABREVIATIONS

(ER) α: Estrogen receptor alpha

(ER) β: Estrogen receptor beta

ACMG: American College of Medical Genetics.

ACOG: American College of Obstetricians and Gynecologists.

AJCC: American Joint Committee of Cancer.

ASCO: American society of clinical oncology

ATM: Ataxia telangiectasia mutation

BFB: Breakage Fusion Bridge

BRCA1: Breast Cancer susceptibility gene 1

BRCA2: Breast Cancer susceptibility gene 2

BRCT: Breast cancer gene 1(BRCA1) carboxyl terminal domain

CAP: College of American pathologist

CD: Cathpesin D

CGH: Comparative genomic hybridization

CHEK2: Cell cycle checkpoint kinase 2.

CI: Confidence interval.

CIN: Chromosomal instability.

CIS: Carcinoma in situ

DCIS: Duct carcinoma in situ.

DFS: Disease free survival.

DNA: Deoxy Nucleic Acid.

EGFR: Epidermal growth factor receptor.

ELISA: Enzyme linked immunosorbent assay

ER: Estrogen receptors

FBC: Familial breast cancer.

FDA: Food and drug administration in USA

FFPE: Formalin fixed paraffin embedded.

FGF: Fibroblast growth factor.

FISH: Fluorescent In Situ Hybridization

FISH: Fluorescnt insitu hybridization.

Her-2/neu: Human epidermal growth factor receptor 2.

HRT: Hormone replacement therapy

HSR: Homogenously stained regions

IHC: Immunohistochemistry

Kb: Kilo base

kDa: Kilo Dalton

LFS: Li- Fraumeni syndrome

LN: Lymph node.

LOH: Loss of heterozygosity

MIs: Mitotic indices

mRNA: Messenger ribonucleic acid.

NCCN: National Comprehensive Cancer Network.

NCI: National Cancer Institute.

OS: Overall Survival

PCBs: Polychlorinated biphenyls

PCR: Polymerase chain reaction

PGF: Platelet derived growth factor

PR: Progesterone receptors

PR-A: Progesterone receptor alpha.

PR-B: Progesterone receptor beta.

Rb1: Retinoblastoma gene.

RFS: Recurrence free survival

ROMA: Representational Oligonucleotide Microarray Analysis

SBC: Sporadic breast cancer.

SPF: S phase fraction.

TDLU: Terminal duct lobular unit

TKRs: Tyrosine kinase receptor

TLIs: Thymidine labeling indices

TOP2A: Topoisomerase II alpha.

TOP2A: TopoisomeraseII alpha.

uPA: Urokinase plasminogen activator

u-PA: urokinase plasminogen activator.

VEGF: Vascular endothelial growth factor

Vs: versus

List Of Errata

Chapter of results:

P.127, table 12; Menopausal status of FBC:

6 Cases premenopausal = 75%.

2 Cases post menopausal = 25%.

The same in P.119, table 16 and figure 30.

P130: table29: ER+&PR+ in FBC: 16.7%.

Chapter of discussion:

P.157: line 11; ASCO, 2003&2011.

P.157: line 16; premenopausal cases (75%) and two postmenopausal cases (25%).

P.166: line 10; lost word: these contributing results could be <u>also</u>

Chapter of summary and conclusion:

P.175: line 15: through hormonal receptor status, and

P176: line 14: (75% Vs 20%).

P.176: line 18: and lower PR expression.

LIST of TABLES

	Page
Table 1: Breast cancer is typically a disease which occurs with advancing age.	8
Table 2: Genes whose mutations are reported to increase risk of hereditary breast cancer.	24
Table 3: The most important genes involved in breast carcinogenesis.	26
Table4: Comparison of BRCA1 and BRCA2.	35
Table 5: Proposed mechanisms of Herceptin action.	61
Table 6: Different methods for Her-2/neu testing .	63
Table 7: Classes of topoisomerases and their criteria.	71
Table 8: Analyses of incremental anthracycline benefit according to topo IIa amplification from Table .	88
Table 9: Pertinent data of the monoclonal antibodies used to	96
immunocharacterize carcinoma of the breast. Table 10: Allred score.	100
Table 11: Her-2/neu score used to evaluate Hercep test.	101
Table 12: Age of patients in FBC group Vs SBC.	117
Table 13: Sex of the studied cases.	118
Table 14: Sex of patients in FBC group Vs SBC group.	118
Table 15: Menopausal status of the studied cases.	118
Table 16: Menopausal status of FBC Vs SBC group.	119
Table 17: Tumor size of the studied case.	120

Table 18: Tumor size of FBC group Vs SBC cases.	121
Table 19: Histological grade of the studied cases .	121
Table 20: Histological grade of the FBC group Vs SBC group	122
Table 21: Histological type of the studied cases.	123
Table 22: Lymph nodal status of the studied cases cancer.	124
Table 23: Lymph nodal status of FBC group Vs SBC group.	124
Table 24: Distribution of cases according to the LN categories.	124
Table 25: Summary of the clinicopathological parameters of the studied cases .	127
Table 26: Distribution of the cases according to ER staining.	128
Table 27: Distribution of the cases according to PR staining.	129
Table 28: Distribution of the studied cases according to the combined ER &	130
PR immunostaining . Table 29: Combined ER and PR immunostaining among FBC group Vs SBC	130
groups. Table 30: Distribution of the studied cases according Her -2/neu gene	132
expression. Table 31: Her-2/neu gene expression among FBC group Vs SBC cases.	133
Table 32: Summary of the immunohistochemical results in FBCs Vs SBCs.	134
Table 33: Rate of Her-2/neu gene amplification in the studied patients.	139
Table 34: The rate of Her -2/neu gene amplification in FBC group versus (Vs)	140
SBC group. Table 35: Rate of TOP2Agene amplification in the studied patients.	142
Table 36: Rate of TOP2Agene amplification in FBC group Vs SBC group.	142

Table 37: Relationship between IHC and FISH diagnostics of Her-2/neu.			144
Table 38: The relationship between Her	-2/neu and TOP	2A gene status	146
detected by FISH.			
Table 39: Her-2/neu and TOP2A genes status in FBCs Vs SBCs.			147

LIST of FIGURES

	Page
Figure 1: Anatomy of the breaste.	6
Figure 2: Familial and sporadic breast cancer	11
Figure 3: Genes involved in FBC breast cancer	26
Figure 4: BRCA1 gene exons , distribution and relative frequency of mutations	30
Figure 5: BRCA1 domain structure and sites of protein interaction	31
Figure 6: cellular functions regulated by BRCA1	32
Figure 7: BRCA2 gene exons and protein structure	33
Figure 8: functions of BRCA2 gene	34
Figure 9: DNA repair by BRCA1 and BRCA2 genes	34
Figure 10: Model of the function of p53 gene	42
Figure 11: Li-Fraumeni syndrome .	42
Figure 12: binding of a ligand to domains I and III	48
Figure 13: Summary of all signaling pathways induced by ErbB receptor	48
Figure 14: The ErbB2 receptor dimerising with ErbB3 receptor	49
Figure 15: The ErbB family of receptors	50
Figure16: The Breakage-Fusion-Bridge cycle	53
Figure 17: Cytogenetic ideogram showing Her-2/neu Amplicon	54