

# **UNIVERSITI PUTRA MALAYSIA**

# BCL-2, HER1, HER2, HER3 AND HER4 EXPRESSION IN PRIMARY COLORECTAL ADENOCARCINOMA

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### BCL-2, HER1, HER2, HER3 AND HER4 EXPRESSION IN PRIMARY COLORECTAL ADENOCARCINOMA



### NURUL MAHIRAH BINTI AHMAD ZUBIR

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

November 2017

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

### BCL-2, HER1, HER2, HER3 AND HER4 EXPRESSION IN PRIMARY COLORECTAL ADENOCARCINOMA

By

#### NURUL MAHIRAH BINTI AHMAD ZUBIR

November 2017

### Chairman : Associate Professor Norhafizah Mohtarrudin, MBBS, MPath, AM Faculty : Medicine and Health Science

In Malaysia, CRC is the most frequently reported cancer among males and second in females with the prevalence of 16.4% and 10.7%, respectively. Most cases occur sporadically, only 5% of the cases are inherited. There is a 50% risk of recurrence for the CRC survivor. Hence biomarkers to improve the prognosis, treatment outcomes and risk assessment are in dire need. The potential biomarkers include BCL-2 and HERs family members (HER1, HER2, HER3, and HER4). BCL-2 is an anti-apoptotic protein and its expression has been associated with tumour grading, staging, and prognosis. It affects tumour genesis by inhibiting cell apoptosis. Presence of BCL-2 expression enhances cancer survival rate. HERs family members are known as a surface receptor. Its activation occurs via ligands binding, leading to downstream signalling that influences cell proliferation. Mutations in HERs family members lead to activation of a series of signalling cascade which has numerous effects on protein expression. HER1, HER2, HER3 and HER4 aberrations in signalling contributes to malignancies, gene amplification and enhanced transcription in the CRC. This study aims to determine the protein expression of BCL-2 and HERs family members in CRC cases, using immunohistochemistry (IHC) technique. A total of 94 FFPE samples were collected from the Hospital Serdang archives from year 2008 until year 2015. Data on demographic and clinicopathologic were retrieved from the database. Protein expressions were detected in the cytoplasm, membranous, or/and nucleus region. The expressions were scored with semi-quantitative scoring system and graded as 0 (absent), 1+ (weak), 2+ (moderate), or 3+ (strong). Association test was used to determine the pvalue (p < 0.05). BCL-2 expression was seen immunopositive in 26.6% cases. HERs family members showed immunopositivity in 31.9%, 13.8%, 42.6%, and 31.9% in HER1, HER2, HER3, and HER4 respectively. There was a significant association between HER1 and HER3 expression (p < 0.001), HER1 and HER4 expression (p =(0.019) and between HER3 and HER4 expression (p = 0.019). Significant association was seen between BCL-2 and HER1 expression (p = 0.022). HER1/HER3 co-expression was significantly associated with tumour grading (p = 0.030) and HER3/HER4 co-expression was significantly related with lymph node metastasis (p = 0.018). HER1 expression was associated with age (p = 0.018), Malay ethnicity (p = 0.038), tumour grading (p = 0.011), and diabetes mellitus (p = 0.044). A significant association was seen between HER2 and diabetes mellitus (p = 0.024). HER3 expression was significantly associated with gender (p = 0.027), tumour grading (p = 0.023), lymph nodes metastasis (p = 0.049), and smoking (p = 0.033). Likewise, HER4 expression was significantly associated with ethnicity (p < 0.001) and tumour grading (p = 0.021). This study shows that BCL2 and the HERs family members are overexpressed in CRC. The implications of these results for the CRC biomarkers are that, generally, overexpression of HERs family members are overexpression and generality. Results in co-expression among antibodies indicated relation and dependency with each other through downstream of pathways. BCL-2, HER1, HER2, HER3 and HER4 could be further developed for targeted therapy receptors and prognostic biomarkers for CRC.

Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Master Sains

### BCL-2, HER1, HER2, HER3 DAN HER4 EKSPRESI DALAM KOLOREKTAL ADENOKARSINOMA PRIMER

Oleh

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### Pengerusi : Profesor Madya Norhafizah Mohtarrudin, MBBS, MPath, AM Fakulti : Perubatan dan Sains Kesihatan

Di Malaysia, kolorektal kanser adenokarsinoma (CRC) merupakan kanser yang paling kerap dilaporkan di kalangan lelaki dan kedua di kalangan wanita dengan statistik masing-masing adalah 16.4% dan 10.7%. Kebanyakan kes yang berlaku adalah secara rawak dan hanya 5% adalah kes yang diwarisi secara genetik. Terdapat 50% risiko keberulang-laku bagi CRC pemandiri. Justeru itu penanda-penanda bio untuk menambah baiki prognosis, hasil rawatan dan penilaian risiko sangat diperlukan. Penanda aras berpotensi yang dikenal pasti termasuklah BCL-2 dan dari ahli keluarga HER (HER1, HER2, HER3, dan HER4). BCL-2 adalah dari protein anti-apoptosis dan ekspresinya berkait rapat dengan gred tumor, peringkat tumor dan prognosis. Ia menjejaskan genetik tumor dengan menghalang sel-sel apoptosis. Ekspresi BCL-2 meningkatkan kadar kelangsungan sel-sel kanser. Ahli keluarga HER dikenali sebagai reseptor permukaan. Pengaktifannya terjadi melalui ligan-ligan mengikat, yang membawa kepada isyarat hiliran yang mempengaruhi proliferasi sel. Mutasi di kalangan ahli keluarga HER membawa kepada pengaktifan satu siri lata isyarat yang mempunyai banyak kesan terhadap ekspresi protein. Penyimpangan isyarat HER1, HER2, HER3 dan HER4 menyumbang kepada kejadian kanser, amplifikasi gen dan peningkatan transkripsi dalam CRC. Kajian ini dilakukan untuk menentukan ekspresi protein BCL-2 dan ahli keluarga HER dalam kes-kes CRC dengan menggunakan teknik imunohistokimia (IHC). Sejumlah 94 sampel tisu yang diproses dalam bentuk tisu terendam formalin-parafin (FFPE) telah dikumpulkan dari arkib Hospital Serdang diantara tahun 2008 hingga tahun 2015. Data mengenai demografi dan klinikopatologi diambil daripada pangkalan data. Ekspresi protein dikesan pada kawasan sitoplasma, membran atau/dan nukleus. Ekspresi diberi skor berdasarkan sistem pemarkahan semi-kuantitatif dan dinilai sebagai 0 (tidak hadir), 1+ (lemah), 2+ (sederhana), atau 3+ (kuat). Ujian pertalian digunakan untuk menentukan nilai p (p < 0.05). Terdapat 26.6% kes imunopositif untuk BCL-2. Ahli keluarga HER menunjukkan imunopositifiti sebanyak 31.9% (HER1), 13.8% (HER2), 42.6% (HER3) dan 31.9% (HER4). Terdapat persamaan yang signifikan antara ekspresi HER1 dan HER3 (p < 0.001), ekspresi HER1 dan HER4 (p = 0.019) dan antara ekspresi HER3 dan HER4 (p = 0.019). Persamaan yang signifikan dilihat antara ekspresi BCL-2 dan HER1 (p = 0.022). HER1 / HER3 ekspresi bersama dikaitkan dengan peringkat tumor (p = 0.030) manakala ekspresi bersama HER3 / HER4 berkaitan dengan metastasis nodus limfa (p = 0.018). Ekspresi HER1 dikaitkan dengan umur (p = 0.018), etnik Melayu (p = 0.038), gred tumor (p = 0.011) dan kencing manis (p = 0.044). Pertalian yang ketara dilihat antara HER2 dan kencing manis (p = 0.024). Ekspresi HER3 secara signifikan dikaitkan dengan jantina (p = 0.027), gred tumor (p = 0.023), metastasis nodus limfa (p = 0.049) dan merokok (p = 0.033). Selain itu, ekspresi HER4 berkait dengan etnik (p < 0.001) dan gred tumor (p = 0.021). Kajian ini menunjukkan bahawa ekspresi BCL2 dan ahli keluarga HER wujud dalam CRC. Implikasi keputusan penanda-bio ini terhadap CRC adalah, secara umumnya, ekspresi ahli keluarga HER lebih cenderung kepada prognosis yang buruk. Walau bagaimanapun penanda BCL-2 gagal menunjukkan sebarang hubungkait dengan parameter demografi dan klinikopatologi. Ekspresi bersama antara antibodi menunjukkan hubungan dan pergantungan antara satu sama lain melalui tapak jalan. BCL-2, HER1, HER2, HER3 dan HER4 boleh dikaji dengan lebih mendalam sebagai reseptor sasaran terapi dan penanda bio prognostik untuk CRC.

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This thesis was submitted to the Senate of the Universiti Putra Malaysia and has been accepted as fulfilment of the requirement for the degree of Master of Science. The members of the Supervisory Committee were as follows:

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# LIST OF ABBREVIATIONS

| °C             | Degree Celsius                                 |
|----------------|--|
| $\mu_{\rm C}$  | Proportion Of Unexposed With Outcome           |
| μm             | Micrometre                                     |
| μ <sub>T</sub> | Proportion Of Exposed With Outcome             |
| 5-FU           | 5-Fluorouracil                                 |
| ABC            | Avidin-Biotin-Peroxidase Complex               |
| AKT            | Protein Kinase B                               |
| APAAP          | Alkaline Phosphatase-Anti Alkaline Phosphatase |
| ATP            | Adenosine Triphosphate                         |
| BAX            | Bcl-2 Associated X Protein                     |
| BCL-2          | B-Cell Lymphoma 2                              |
| CEA            | Carcinoembryonic Antigen                       |
| cm             | Centimetre                                     |
| CRC            | Colorectal Cancer                              |
| CT-scan        | Computed Tomography Scan                       |
| DAB            | 3,3'-Diaminobenzene                            |
| DM             | Diabetes Mellitus                              |
| DNA            | Deoxyribonucleic Acid                          |
| ECD            | Extracellular Domain                           |
| ELISA          | Enzyme-Linked Immunosorbent Assay              |
| ErbB           | Avian Erythroblastosis Oncogene B              |
| ERK            | Extracellular Signal-Regulated Kinase          |
| ERRP           | EGFR-Related Protein                           |

 $(\mathbf{C})$ 

- FDA Food and Drug Administration
- FFPE Formalin Fixed Paraffin Embedded
- FOBT Faecal Occult Blood Testing
- GIT Gastrointestinal Tract
- HB-EGF Heparin Binding Epidermal Growth Factor Like Growth
- HER Human Epidermal Growth Factor Receptor
- HER1 Human Epidermal Growth Factor Receptor-1
- HER2 Human Epidermal Growth Factor Receptor-2
- HER3 Human Epidermal Growth Factor Receptor-3
- HER4 Human Epidermal Growth Factor Receptor-4
- ICD Intracellular Domain
- IHC Immunohistochemistry
- JM Juxtamembrane
- kDa Kilo Dalton
- LN Lymph Node
- mAbs Monoclonal Antibodies
- MAPK Mitogen Activated Protein Kinase
- MEK MAPK/ERK Kinase
- MRI Magnetic Resonance Imaging
- NRG Neuregulin
- OS Overall Survival
- PAP Peroxide-Antiperoxidase
- pH Potential of Hydrogen
- PI3K Phosphatidylinositol 3-Kinase

## RNA Ribonucleic Acid

ROS Reactive Oxygen Species

RT Room temperature

SH2 Src homology 2

SH3 Src homology 3

G

- TKI Tyrosine Kinase Inhibitors
- TKR Tyrosine Kinase Receptor
- TM Transmembrane Domain
- WHO World Health Organization

### CHAPTER 1

### **INTRODUCTION**

#### 1.1 Study Background

Colorectal cancer (CRC) is an asymptomatic disease in its early development, and generally takes up five to ten years to be obvious. In the United Kingdom (UK), CRC is the second highest cause of death by cancer with 15,000 cases in 2014 (Cancer Research UK). The National Institute of Health (NIH) reported the United State spent more than \$216.6 billion for cancer management in 2009 alone. Globally, CRC is the third leading cause of cancer death (Garza-Treviño *et al.*, 2015) and is the third most frequently diagnosed cancer in both men and women (Sabounchi *et al.*, 2012). Among all the cancer survivors, there is an alarming risk of 50% relapse and it is difficult to achieve a total cancer-free condition (Bandrés *et al.*, 2007; Labianca *et al.*, 2010). The increasing incidence of CRC has been reported in Asian region, although it was previously considered as a low risk region (Labianca *et al.*, 2010). According to the Malaysian National Cancer Registry Report 2007-2011 (2016), CRC is the most common cancer in male Malaysian population and second in female population. Cumulatively, CRC is the second most reported cancer with 13,693 cases between the year 2007 to 2011 (Azizah *et al.*, 2016).

Management of CRC depends on the grade and stage of the disease. Currently, surgery is the gold standard for CRC treatment, common for stage II and stage III. However, the routine chemotherapy cocktail use in CRC does not only eradicate cancerous cells, it also indirectly weaken nearby healthy cells (Caley & Jones, 2012). Antibody targeted therapy which selectively kills cancer cells has lesser side effects and could enhance patients recovery rate (Sudhakar, 2009). Molecular markers have been correlated with CRC prognosis and therapeutics approaches, in addition for use in risk assessment. Efficient biomarkers are needed for prognostication and measurement of competency of therapeutic regimens. The ability of cancer cells to avoid apoptosis and increase proliferation rate leads to cell cycle aberration. Concurrently, analysing two related biomarkers brings more information with the potential to serve as dual targeted antibodies (Newton *et al.*, 2012; Reimers *et al.*, 2013).

The B-cell lymphoma-2 (BCL-2) is a member of the anti-apoptotic proteins which are composed of more than 25 members. BCL-2 is overexpressed in various types of cancers including breast, ovaries, thyroid and colon. A previous in vitro study done on colorectal cell lines (CaCo2, Colo205, HT29 and SW480) demonstrated that BCL-2 was significantly overexpressed. Knockdown of BCL-2 hindered invasiveness and motility in colorectal cell lines, while overexpression of BCL-2 enhances cell migration in two-fold (Koehler *et al.*, 2013). Several studies have also indicated that BCL-2 expression is significantly associated with better prognosis in long term and improved survival (Han *et al.*, 2006; Kouraklis *et al.*, 2003; Meterissian *et al.*, 2001).

Multiple studies have suggested that the heterodimerization of HER family members plays a crucial role in tumourigenesis and in the development of resistance to therapy in CRC cases. HER or ErbB family members have been extensively studied in different types of cancer including breast, lung, skin and colon cancer for biomarkers purposes. Numerous studies have indicated that HER1, HER2, and HER3 properties are significantly associated with CRC conditions. The role of HER4 is still understudied, although its involvement in CRC carcinogenesis seems likely. HER1 expression has been found to be related to TNM stage III. Subsequently, HER1 knockdown interfered with SW480 colon cancer cells anchorage-independent cell growth ability (Li et al., 2014; Spano et al., 2005). Recent studies have suggested that HER2 activation correlated with lymph node and tumour differentiation (Half et al., 2004; Kountourakis et al., 2006). Further study in colon cancer cell line indicated that HER2 overwhelming signals increase cancer metastasis activity and lead to poor clinical outcome (Yu & Hung, 2000). An earlier study in gastric cancer determined that HER3 overexpression is strongly associated with advanced tumour progression and is a poor prognostic indicator (Hayashi et al., 2008). Another study in breast cancer revealed connection between HER3 overexpression and reduced survival rate (Witton et al., 2003).

HER4 cell lines study indicated that the disruption of HER4 signalling in poorly differentiated CRC cell lines inhibits cell survival mechanism, but HER4 overexpression enhance cell survival and growth. Excessive HER4 signalling is also documented as a factor for epithelial inflammation (Williams *et al.*, 2015). Recent studies have established that HER family members can interact and bond with each other to form an array of heterodimer complexes. Therefore, the activation of one receptor could modulate the activity of other HER family members. Co-expression could extend the range of downstream signalling events. Their interaction among each other might enable cells to specifically react to exogenous stimuli. Co-expression of HER2 and HER4 had a shorter overall survival in late stage and poor prognosis CRC cases (Lee *et al.*, 2002). A better survival in bladder cancer was found in HER3 and HER4 co-expression (Memon *et al.*, 2004). Co-expression of HER2 and HER3 also were highly associated in breast cancer (Witton *et al.*, 2003).

#### **1.2 Problem statement**

CRC is one of the most common cancers with high recurrence and mortality rates. Currently, biomarkers available for CRC are less sensitive and specific for prognostic and therapeutic purposes. Treatment resistance is also an issue in CRC. One of the factors contributing to treatment resistance and relapse issues is the aberrant apoptotic signalling. High anti-apoptotic signals from BCL-2, for example, can reduce chemotherapy efficiency and enhance cancer cell survival. This situation is likely to cause poor prognosis and relapse in patients. Targeted cancer therapy promises less side effects and better quality of life. Blockage of HER family members contributes to the decrease of cancer cell activities including delaying tumour growth, inducing tumour shrinkage as well as reducing metastasis rate. Correlation of the molecular alterations with demographic and clinicopathologic data could also provide important information on the aetiology of CRC.

### 1.3 Objectives

General objective:

To determine protein expression of BCL-2, HER1, HER2, HER3, and HER4 in CRC patients using immunohistochemistry (IHC) technique.

Specific objectives:

- 1. To determine the demographic and clinicopathologic distribution of CRC cases.
- 2. To determine the distribution of BCL-2 and HERs (HER1, HER2, HER3, and HER4) protein expression in CRC cases.
- 3. To determine the association among HERs (HER1 and HER2, HER1 and HER3, HER1 and HER4, HER2 and HER3, HER2 and HER3, HER2 and HER3 and HER4) expression in CRC cases.
- 4. To determine the association between BCL-2 and HERs (BCL-2 and HER1, BCL-2 and HER2, BCL-2 and HER3, and BCL-2 and HER4) expression in CRC cases.
- 5. To determine the association of BCL-2 and HERs (HER1, HER2, HER3, and HER4) protein expression with the demographic and clinicopathologic parameters.

### 1.4 Research hypothesis

- 1. BCL-2, HER1, HER2, HER3, and HER4 overexpression are detected in CRC cases.
- 2. There are significant associations among HERs (HER1 and HER2, HER1 and HER3, HER1 and HER4, HER2 and HER3, HER2 and HER4, and HER3 and HER4) expression in CRC cases.
- 3. There are significant associations between BCL-2 and HERs (BCL-2 and HER1, BCL-2 and HER2, BCL-2 and HER3, and BCL-2 and HER4) expression in CRC cases.
- 4. There are significant associations of BCL-2 and HERs (HER1, HER2, HER3, and HER4) protein expression with the demographic and clinicopathologic parameters.

### 1.5 Significance of study

- 1. This study provides baseline data on CRC cases in relations to BCL-2, HER1, HER2, HER3, and HER4 protein expressions.
- Findings of this study could suggest the potential use of biomarkers for therapeutic purposes such as in BCL-2 (Venetoclax, (Cang *et al.*, 2015)), HER1 (Cetuximab (Pozzi *et al.*, 2016)), and HER2 (Neratinib (Kavuri *et al.*, 2015)). Including in some dual targeted therapy drugs in HER1/HER3 (MEHD7945A (Huang *et al.*, 2013)) and HER1/HER2 (Lapatinib (Kim *et al.*, 2009)).
- 3. BCL-2, HER1, HER2, HER3, and HER4 antibodies could be used as IHC biomarkers to determine suitable patients with proper therapeutic approaches.



Clinicopathologic 1. Grading (Ueno *et al.*, 2008) 2. TNM staging (Srinivasan *et al.*, 2000) 3. LN metastasis (Kountourakis *et al.*, 2006) 4. Dukes staging (Kountourakis *et al.*, 2006) 5. Tumour location (Wray *et al.*, 2009)

### Demographic

Age
 (Labianca *et al.*, 2010)
 Gender
 (Sung *et al.*, 2005)
 Ethnicity
 (Rashid *et al.*, 2009)

### COLORECTAL CANCER (CRC)

# **Other Risk Factor**

Smoking
 (Limsui *et al.*, 2010)
 Alcoholic drink
 (Derry *et al.*, 2013)
 Family history
 (Guttmacher *et al.*, 2003)
 Diabetes mellitus
 (Sun & Yu, 2012)

## **Proteins Expression**

- 1. BCL-2 (Poincloux et al., 2009)
- 2. HER1 (Rego et al., 2010)
- 3. HER2 (Seo et al., 2014)
- 4. HER3 (Seo et al., 2015)
- 5. HER4 (Williams *et al.*, 2015)

Figure 1.1 : Variables associated with CRC cases

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