

10. Isolation, Characterization and Proliferation of Cancer Cells from Breast Cancer Patients

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ORIGINAL PAPER

Isolation, Characterization and Proliferation of Cancer Cells from Breast Cancer Patients

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ABSTRACT

Introduction: The most cases of cancer death, which are in the first rank among cancers suffered by women is breast cancer. The breast cancer therapy for patients has been developed, but still not optimal, so it is necessary to understand the mechanism of therapy in model cell of breast cancer. **Aim:** This study aim to develop an isolation technique of breast cancer cell from patients as a cancer cell model. **Material and Methods:** Breast cancer cell isolation is performed by enzymatic methods using the collagen I and hyaluronidase. Then, breast cancer cells were characterized using flow cytometry based on the CD44/CD24 expression where MDA-MB468 and MCF-7 breast cancer cell lines were used as positive controls. Estrogen receptor (ER), progesterone receptor (PR), p53, HER2, and Ki67 expression were assessed using an immunohistochemistry assay. **Result and Discussion:** The morphology of cancer cells was fibroblast like cells the day 7th after isolation. Isolated breast cancer cells expressed 95.33±0.47% of CD44⁺/CD24⁺ and human epidermal growth factor receptor 2 (HER2) low expressions. Isolation of breast cancer cells can use In-house enzymatic protocol. Isolated breast cancer showed the same expression as MDA-MB468 (CD44⁺/CD24⁺) and HER2 compared to MCF-7 cell lines (CD44/CD24⁺). **Conclusion:** These cells belonged to a basal type of breast carcinoma and expressed CD44⁺/CD24⁺, then isolated BCCs can be used as model cancer cells for further research. **Keywords:** breast cancer, cell lines, CD44 antigen, CD24 antigen, enzymatic assay.

1. INTRODUCTION

Breast cancer is the first rank among cancers suffered by woman causing death. It often occurs recurrence after treatment (1). In the Asia-Pacific early a quarter of the population of women were diagnosed with breast cancer (24%), with the largest percentage occurring in China (46%), Japan (14%), and Indonesia (12%). The type of cancer in Indonesia is dominated by breast cancer (30%) and cervical cancer (24%) (2).

Breast cancer classification was based on different parameter, i.e.: tumor grade, lymph node status, histological type, the presence of estrogen receptor (ER) and human epidermal growth factor receptor 2 (HER2). Molecular profiling proved that breast cancer could be classified into: normal, basal, luminal A, luminal B, and HER2 based on ERα, progesterone receptor (PR) and HER2 expression in immunohistochemical assay (3). MDA-MB453 is HER2⁺ type which expresses high Ki67

(4), while MDA-MB-468 is a basal type with ER⁻, PR⁻ and HER2⁺ expression and MCF7 is luminal A type with of ER⁺, PR⁺ and HER2⁺ expression (5).

Breast cancer treatment depends on the stadium of breast cancer. Therefore, the selection of therapy at each stadium of the cancer is different (6). Surgery and chemotherapy are types of cancer therapy, which is effective to treat breast cancer (7). But, surgical therapy is costly and high risk of infection during surgery, as well as patient complaints regarding cosmetics (8). Chemotherapy can destroy normal cells and cause side effects, including: neutropenia, anemia, thrombocytopenia, alopecia, mucositis (8). The cell line is the most important factor in studying therapeutics target, carcinogenesis, and signal pathways for breast cancer (9).

2. AIM

The cell line of study is based on its biological aspects. Hence, research needs to be done on breast cancer iso-

lation. Therefore, it is essential to have isolated breast cancer cells from breast cancer patient as a cancer cell model in order to determine the effective cancer therapy.

3. MATERIAL AND METHOD

Sample Preparation

The sample was the tissue from breast cancer patients 52 years old with ductal carcinoma mammae sinistra grade III and invasive to lymphovascular. The informed consent used the guidelines approved by the Institutional Ethics Committee collaboration between Maranatha Christian University, Bandung, Indonesia and Immanuel Hospital Bandung, Bandung, Indonesia. Tissue was then washed using phosphate buffer saline (PBS 1x, Gibco, 1740576) preserved in the transfer medium consist of PBS 1x + 1% Abam (Gibco, 1772653) + 1% Amphotericin B (Gibco, 15290026) + 0.1% Gentamicin (Gibco, 15750060) before being isolated (10).

1st Cancer Cell Isolation

Medium preparation: Dulbecco's Modified Eagle Medium (DMEM, Gibco 11995065) : F-12K (Sigma, I6634) (1:1), 5% Fetal Bovine Serum (FBS, Gibco, 10270106), 1% Antibiotic-antimycotic (Gibco, 1772653), 1% Amphotericin B (Gibco, 15290026), 0.1% Gentamicin (Gibco, 15750060) were dissolved then filtered using Syringe Filter PES 0.22µm (TPP, 99722). Enzyme preparation: Collagenase Type I (Gibco, 10114532) 4 mg/ml, hyaluronidase (Sigma, 515397) 1 mg/ml, trypsin 0.1% (Gibco, 25200072) were dissolved in basal medium that consists of DMEM: F-12K (1:1), then filtered using syringe filter 0.22 µm PES (TPP, 99722).

Cells Isolation: Prepare a surgical device, PBS 1x and sterile petri dish. The tumor tissue was washed with PBS 1x on a petri dish and then cut it into small pieces using surgical scalpel. The sample fragments were washed again with PBS 1x (Gibco, 1740576), inserted into a 50 ml falcon tube (TPP, 91050) containing the mixed enzyme solution and then incubated 16 hours, 300 rpm, 37°C, 5% CO₂. After incubation sample was filtered using 100 µm strainer (Corning, 431752) then centrifuged at 1600 rpm for 10 min. Supernatant was discarded and the pellet was washed with PBS 1x and centrifuged at 1600 rpm for 5 min. The pellet was then resuspended with complete growth medium. The cell suspension was next incubated at 37°C, 5% CO₂ (10).

Breast Cancer Cells Characterization

Breast Cancer Cells (BCCs) at P3 were characterized using flow cytometry (MACSQuant, Analyzer 10) for positive and negative markers of breast cancer cells at density 1x10⁵ cells/tube. The cells were then stained with CD44 and CD24 according to manufacturer protocol (BioLegend kit, 338804/311118). The experiments and measurements of surface marker were performed in triplicate (11).

Proliferation Assay

BCC was cultured in medium consisting of DMEM: F-12K (1:1), 5% FBS, 1% Antibiotic-antimycotic, 1% Amphotericin B, 0.1% Gentamicin and the medium was replaced every 3 days. Briefly 3 x 10⁵ cells were seeded on T25 flask and cultured in complete medium until reaching confluence 80% around 3 days for proliferation assay. Briefly, cultured cells were detached using trypsin, then incubated for 1-3 min at 37°C, complete medium was added to neutralize trypsin and the detached cells were centrifuged (MPW-2000) at 1600

rpm for 5 min at room temperature. The cell pellet was resuspended with trypan blue solution (Sigma, 25200072) and diluted in 1:1 dilution. Then, cells were counted using a haemocytometer (Neubauer, 17849). Population Doubling (PD) was counted at every passage with the formula:

$$PD = [\log_{10}(NH) - \log_{10}(NI)] / \log_{10} 2$$

NI is the inoculum cell number and NH is the cell harvest number. Then, to determine cumulative PD data by added the PD at the previous passage.

The PD time (PDT) was determined by the formula (11).

$$PD \text{ time} = t (\text{time}) / PD (\text{in days})$$

Immunohistochemistry (IHC) Analysis

Breast cancer cells were processed using anti-S727p-STAT3 (BD Biosciences), anti-CA-IX and anti-EGFR (clone 31G7, Zymed Technologies, Invitrogen, Burlington, ON) protocols. The Idetect Ultra HRP Detection System (ID Labs Inc, London, Ontario) was used to visualize. Counterstaining was performed with Gill modified hematoxylin (Harleco®) (12).

Statistical Analysis

Statistical analysis was conducted using SPSS software (version 20.0). Data were presented as Mean ± Standard Deviation. Significant differences among treatments were determined using the one-way Analysis of variance (ANOVA) and $p < 0.05$ were considered as statistically significant, along with Turkey honestly significant difference post hoc test and 95% confidence interval. Z score values were used for plotting the protein expression based on immunohistochemistry signal. Clustering methods based UPGMA statistical analysis were performed by R software ver.3.4.3.

4. RESULTS

Isolation of BCCs with enzymatic method

BCCs were successfully isolated using an enzymatic method and cultured until passage 9. The cells attached to the surface of the plastic dish a day after seeding (P0) and were maintained up to two weeks. The cells showed short fibroblastic-like morphology in growth medium and there were still blood cells at the beginning of P0 (Figure 1A). From the first passage onwards, particularly at P1, P2, P3, the cells in growth medium changed their shape into long fibroblast cells (Figure 1B-1D).

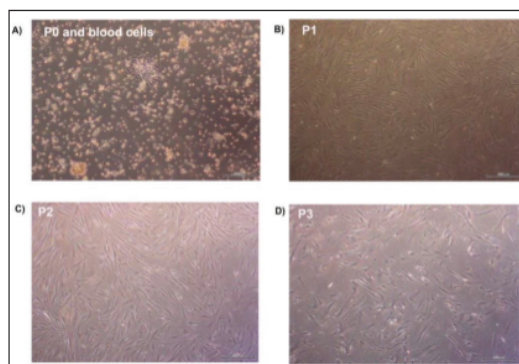


Figure 1. The morphology of breast cancer cells in the different passages. (A) BCCs at P0 and blood cells. (B) BCCs at P1. (C) BCCs at P2. (D) BCCs at P3. Magnification 40x. Scale Bars: 500µm.

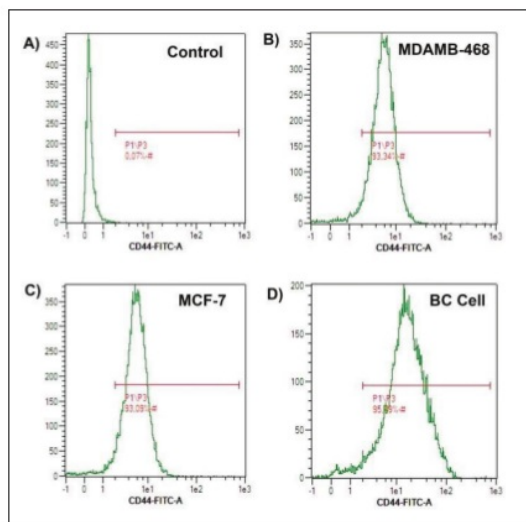


Figure 2. Density plot CD44 expressions in the unstained cell as a control (A), MDAMB-468 cell (B), MCF-7 cell (C), and breast cancer cell/BC cell (D)

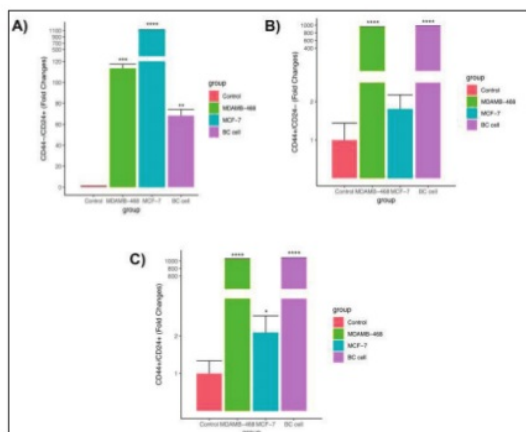


Figure 3. Expression of CD44+/CD24+ cell (A), CD44+/CD24- cell (B) and CD44+/CD24+ cell (C) characterization in breast cancer cell (BC cell) compared to MCF-7 cell lines and MDAMB-468 cell lines. Significance difference with p value $p < 0.05$

Characteristic of Breast Cancer Cells

The flow cytometry analysis displayed the biological properties of BCCs, which was compared to MDA-MB468 and MCF-7 as cells control. BCCs showed a highest CD44⁺ cell intensity expression compared to the MCF-7 cell line, but had no significant difference with MDA-MB468. The intensity of CD44⁺ in most cells around 10^1 (log1) (Figure 2). However, MDA-MB468 and BCCs had a lower number of CD44⁺/CD24⁺ cells than MCF-7. On the other hand, CD44⁺/CD24⁺ and CD44⁺/CD24⁻ of BCCs and MDA-MB468 were higher than MCF-7 (Figure 3).

Population Doubling Time of Breast Cancer Cells

PDT and PD rate were used to evaluate the breast cancer cell proliferation capacity. PDT is the time needed by cell to divide into two cells. According to the results shown in Table 1, BCCs suggested high PDT at early P2 and P3 around 21.38 ± 1.34 and 18.10 ± 1.69 respectively compared to late pas-

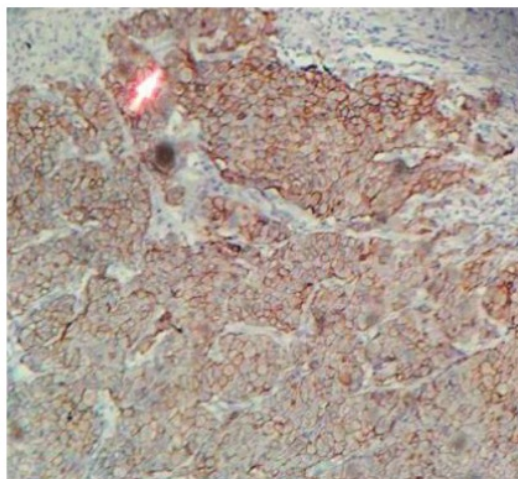


Figure 4. Immunoprofile of breast tissue from breast cancer patient. Immunohistochemistry result of HER2 expression in breast cancer.

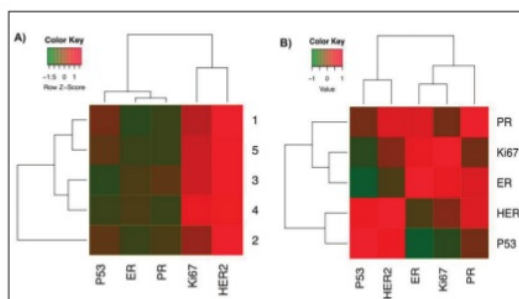


Figure 5. Gene Correlation Analysis (A) Heatmap protein expressions in breast cancer were analyzed by immunohistochemistry. (B) heatmap correlation graph by "Pearson" analysis in five surface protein expressions

sages (P7 and P8). It could be seen that BCCs reduced their PDT after every passaging (Table 1).

Immunohistochemistry (IHC) and Gene Correlation Analysis

IHC analysis was identified for immunoprofile of breast tissue from breast cancer patient (Figure 4). This procedure was performed to double check flow cytometry analysis of isolated BCCs in order to determine the molecular classification of cancer cell subtype. Five gene analysis (p53, ER, PR, Ki67, HER2) using an IHC staining method showed that HER2 was the strongest upregulated gene expression followed by Ki67 and other genes were down regulated (Figure 5A). In contrast, DeFazio-Eli et al. (2011), Holliday and Speirs (2011) reported that MDA-MB468 suggested very low level of HER2. In a gene correlation analysis using Pearson method, HER2 had a high positive correlation with p53, PR, ER and low positive correlation with Ki67 (Figure 5B).

5. DISCUSSION

There are two kinds of breast cancer, namely ductal and lobular carcinoma that attack the milk duct and the milk producing gland respectively (13). Breast cancer therapy, which has existed is still not optimal. Surgery therapy was costly and has a risk of infection during surgery (8). It also often

occurs recurrence after treatment (1). Chemotherapy also has not shown optimal results, because the chemicals also target normal cells in the body, so that they can harm cells in the body. This causes new issues alternative therapy is needed to determine the mechanism and causes of breast cancer. Therefore, to understand the mechanism in breast cancer cell, model cell is needed. Even though utilization of cell lines models cell for most the basic research, they are already engineered for some reasons which may not clinically related to the population of breast cancer.

The morphology of isolated BCCs at P0 show different shape compared to BCCs at P1 onwards. This is caused by the adaptation ability of isolated BCCs in a new environment. Since isolated BCCs have not only similar biological properties to MDA-MB468 but also its morphology like fibroblast cells (14). Isolated BCCs seem to keep with a good state in fibroblast like morphology at each passage.

This study use MCF-7 and MDA-MB468 as a comparison because both cell lines are breast cancer cell lines, MCF-7 is the luminal type with ER⁺ and PR⁺ and MDA-MB468 is the basal type with ER⁻, PR⁻ and HER2⁻ breast cancer cell line (5). IHC analysis suggests that isolated BCCs are the basal type of breast cancer cell. This finding highlights the importance of classifying isolated BCCs based on molecular classification using IHC assay and surface marker expression using flow cytometry analysis into consideration of clinical cancer therapy.

The isolated BCCs express low HER2 expression and high Ki67 expression based on IHC analysis. It is in line with research from DeFazio-Eli et al. that MDA-MB468 cell line has also very low levels of HER 2. Also, Holliday and Speirs (3) mention that MDA-MB468 is a basal type because this cell line has ER⁻, PR⁻ and HER2⁻ and high Ki67 characteristic around 20%. Ki67 expression ranging under 50%, while cancer cell lines ranging from 20%-100% is one explanation of striking differences between cell lines and isolated primary tumor cell (9). The similarities between basal and triple-negative phenotypes cannot guarantee the type of breast cancer. In line with recent studies, there is still no convincing that triple negative is a feature of basal type cancer (15). Therefore, our finding can be assumed that isolated BCCs have the same characteristics as MDA-MB468 according to IHC assay and flow cytometry analysis.

According to the results shown in Table 1, breast cancer cell proliferation in the second passage need 21 days to doubling their population. While in the third and fourth passage need 18 day and 13 day to doubling their population respectively. In earlier passages, the cell takes time to proliferate because it adapts to the new environment, however, in the next passage the cell has adapted well and high proliferate. Furthermore, doubling time in cancer cells is faster than other cells isolated from human. It is because cell cancer is a malignant cell that has a fast proliferation time than normal cells. It is supported by Liu et al. (16) she states that malignant cells have faster doubling times compared to other normal cells. The doubling time for MDA-MB468 (ATCC) should be 30-40 hours (17), while in this study PDT is 5,73 days in passage 8. This indicates that the isolated cells had a lower proliferation rate compared to MDA-MB468.

6. CONCLUSION

This study concludes that BCCs could express CD44⁺/CD24⁻, then expected act as model breast cancer cells effectively.

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- **Author's contribution:** This work was carried out in collaboration between all authors. W.W., supervised the laboratory work, contributed to the experimental design and the protocol (writing of the manuscript). Y.H., D. R. L., D. K. J. and T. L. W. carried out laboratory work, contributed to the experimental design and the protocol. R. R., F. S. P., A. A., A. A., and Z. K. carried out laboratory work, the experimental design and contributed to the protocol. and M. S. performed the statistical analysis and contributed to the protocol. All authors read and approved the final manuscript.
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