

Case Report

Running Title: Clinicopathological Characteristics of Patients with *BRCA* Mutation BC

Received: August 25, 2021; Accepted: July 12, 2022

Clinicopathological Characteristics of Patients with *BRCA* Mutation BC in North Sumatera: Case Report

Dedy Hermansyah[♦], MD, Gracia Pricilia, MD, Yolanda Simamora, MD, Denny Rifsal
Siregar, MD

*Department of Surgery, Faculty of Medicine, Universitas Sumatera Utara, Medan,
Indonesia*

♦Corresponding Author

Dedy Hermansyah, MD
Department of Surgery, Faculty of Medicine,
Universitas Sumatera Utara, Medan, Indonesia
Tel: +6281396743472
Email: escape744@gmail.com

Abstract

The most common cancer in women is breast cancer (BC) with an incidence of 24.2%. BC in younger patients will in general be more forceful, prompting more awful results and a requirement for more forceful treatment which may bring about a higher probability of long-haul treatment-related harmfulness and novel psychosocial issues. Furthermore, family inclination to breast cancer as *BRCA1* and *BRCA2* mutations is more prevalent in this age group. There were a total of five ladies who had tumor pathology testing with negative results. All intrusive BC examples were regularly assessed for estrogen receptor, progesterone receptor (PR), and human epidermal growth-factor receptor-2 (HER2)/neu status utilizing immuno-histochemistry. Cases with HER2/neu staining of 1+, 2+ or 3+ on immuno-histochemistry examination were additionally assessed by fluorescent in situ hybridization for the enhancement of the HER2/neu quality. In this examination, we distinguished clinicopathological attributes of patients with BC. We partitioned into two gatherings, *BRCA* positive change and *BRCA* negative transformation. Roughly 5%-10% instances of BC have a positive family ancestry and about 20%-40% BC development were in acquired variations. Our study revealed that 20% of cases included individuals who had a family history of *BRCA* mutation. Male relatives with breast cancer, earlier age at onset, a greater prevalence of reciprocal breast disease, and a connection to various malignancies in the ovary, colon, prostate, pancreas, and endometrial are only a few of the clear clinical characteristics of *BRCA1/2*-related BC.

Keywords: Breast neoplasms, Genes, *BRCA1*, *BRCA2*, Immunohistochemistry

Introduction

The most common cancer in women is breast cancer or carcinoma (BC) with an incidence of 24.2%. This type of cancer is the second-highest prevalence of incidence in the world with an incidence rate of 11.6%. Approximately 2,089 million new cases of BC were found in 2018. On the Asian continent, for example, BC is the main malignancy of women, which occurs in around 22.4% of the population with 911,014 new cases and 137,514 cases of which come from Southeast Asia.¹ As the phenomena of this case is discovered at an advanced stage, the prevalence of new BC cases is rising quickly in emerging nations. The most recent phenomenon is examined by data from GLOBOCAN 2020, which reveals that BC is the most prevalent cancer in women with an estimated 58,256 new cases or 30.9 percent, and that this disease will be the leading cause of death for women in Indonesia in 2020 with an incidence rate of 22,692 cases or 12.56 percent.²

BC in younger age populations is generally described as 'more-severe', as a requirement of longer yet aggressive approach will be applied which may bring about a higher probability of long treatment-related harmfulness and novel psychosocial issues. Furthermore, familial inclination to BC, as *BRCA1* and

BRCA2 transformations, is more normal in this age. Regarding high penetrance rates among such transformation carriers, it is critical to distinguish patients who may require extra danger decrease intercessions like two-sided mastectomies and oophorectomies. BC were divided into those with high articulation of the estrogen receptor (ER) quality (luminal A and luminal B subtypes) and those that do not express ER in light of the sub-atomic profiling of malignancies. Inside the ER-negative gathering, tumors that overexpress the *HER2/neu* oncogene are named the *HER2/neu*-positive subtype.³⁻⁵ ER negative tumors that express qualities found in basal epithelial cells and can be stained with antibodies to keratin 5/6 were distinguished as basal-like tumors. A larger part of these basal-like tumors is accepted to comprise of BC subtype in which does not express ER, progesterone receptor (PR), or *HER2/neu* (i.e., triple-negative breast cancer or TNBC). A few examinations have exhibited that *BRCA1*-change carriers are bound to be determined to have triple-negative BC than noncarriers.⁶

Moreover, carriers of *BRCA2* changes appear to have comparable pathologic attributes with noncarriers. Furthermore, previous analyses have often ignored critical clinical parameters that could

influence the total tumor improvement. Hence, as the consequence, this study aims to overview the clinicopathological characteristics of patients with *BRCA*-Positive and *BRCA*-Negative BC in North Sumatera region in Indonesia.

Case Presentation

Five women in total were included from the Division of Oncology in Department of Surgery of Universitas Sumatera Utara General Hospital with confirmed triple-negative breast cancer (TNBC) by immunohistopathological evaluations in the surgical pathology in our center. The ethical consideration of this study was approved by the ethical committee of the same institutions with the registered ethics code of 130/UN5.2.1.1.1.19/PPM/2022. The collected data include family history of cancer, cancer history, age, BC grade and stage, immunohistochemistry (IHC) testing, nodal metastasis, and sites of metastasis including lung, liver, brain, and bone.

Tumor pathology for the examples with BC was looked into by the pathologists. Data with respect to the histologic sort of BC; tumor grade utilizing the adjusted Black's atomic reviewing framework; and ER, PR, and HER2/neu status of BC tests were acquired from the patients' institutional

pathology reports. All intrusive BC examples were regularly assessed for ER, PR, and HER2/neu status utilizing IHC. Cases with HER2/neu staining of 1+, 2+ or 3+ on IHC examination were additionally assessed by fluorescent in situ hybridization for the enhancement of the HER2/neu quality. The clinicopathological attributes of patients with *BRCA*-Positive and *BRCA*-Negative BC appeared on the table 1.

Discussion

Similar to our own Indonesia, BC in developing countries is associated with unusual highlights that are often underrepresented in Western-culture of attention. In the first place, middle age at determination is at any rate 10 years younger than that of the West, a reality that should be contemplated when planning early-location programs and furthermore managing psychosocial outcomes and long treatment confusions in a younger age. Second, a huge level of patients presents with privately progressed or metastatic infection, even among the most youthful.⁷

We distinguished clinicopathological attributes of patients with BC. We divided into two groups: *BRCA* negative transformation and *BRCA* positive change. As we presumably all know, this illness has a strong genetic component

but may also develop in an atypical way. *BRCA* was a significant characteristic that signified as a propensity or risk factor in this circumstance. Roughly 5%-10% instances of BC have a positive family history and about 20%-40% BC development was in acquired variations. Our examination shows that one of case (20%) were happen in the patients that has family ancestry in *BRCA* transformation. This outcome is in accordance with an investigation of 207 families that foreseeing changes in the *BRCA* qualities utilizing family ancestry data, was expanded by including the ER and PR receptor status and pathologic evaluation of the tumor. Extra pathologic factors that may anticipate *BRCA1* change status incorporate Ki67 and epidermal development factor receptor. In young (age less than 54 years) ladies with BC development, significant degrees of Ki67 articulation anticipated an opportunity of having a *BRCA1* change as high as 75%. In light of these studies and mounting evidence that suggests *BRCA1* tumors exhibit unusual pathologic characteristics, doctors may want to consider combining pathology findings with family ancestry information when determining if a patient is at increased risk for genetic BC. This might be especially valuable when family ancestry data brings about a moderate worry for

innate malignancy. Notwithstanding, on the grounds that apparently *BRCA2*-related malignancies have pathology like that of non-*BRCA* carriers, it is right now indistinct whether pathologic outcomes might be utilized in foreseeing *BRCA2* change status. Extra examination ought to be led to decide how much accentuation ought to be set on tumor pathology and how this data can be remembered for effectively settled models.^{8,9}

In 2009, Kwong et al. announced the clinicopathological qualities of Chinese patients with *BRCA*-related BC development. Among 226 high-hazard Hong Kong Chinese ladies, 28 ladies (12.4%) conveyed *BRCA* transformations (*BRCA1* change, 11 patients; *BRCA2* changes, 17 patients), and 55.6% of these carriers were determined to have BC before age 40 contrasted and 36.0% of non-carriers ($P = 0.05$). A family history of breast and ovarian cancer, high-grade tumors, and TNBCs is a given for *BRCA* change carriers. TNBC was essentially more prevalent in *BRCA1* carriers (67.7%) than in *BRCA2* carriers (35.3%) and non-carriers (25.6%). Negative cancer found in the emergency department was mostly linked to *BRCA1* mutations, especially in individuals under the age of 40.¹⁰

Period of beginning had significant role as one of clinical qualities in innate

tumors particularly patients in youthful beginning. In innate BC development, the patients endure this carcinoma in the youthful beginning. As in our reports, the patient determined to have BC development with *BRCA* transformation and positive family ancestry were in younger age. This outcome is in accordance with meta-examinations indicate that *BRCA1* and *BRCA2* carriers have a 57%–65% and 45%–49% likelihood of creating BC over lifetime, individually. Patients with a familial history of breast or ovarian cancer, a personal history of BC at a younger age, or triple-negative aggregate have *BRCA1/2* germline alterations that are more common (for *BRCA1* as it were). As of late, Lang et al. revealed the pervasiveness of *BRCA* transformation and highlights of *BRCA*-related BC in Chinese patients using cutting edge sequencing on 2,991 BC patients and 1,043 solid people as controls. *BRCA* transformations were available in 9.1% (232/2,560) of patients with in any event one danger factor for genetic BC development contrasted with 3.5% (15/431) in irregular patients and 0.38% (4/1,043) in sound controls. Family background of breast/ovarian malignant growth, younger age, negative HER2, high Ki-67 list, and high tumor grade were related with *BRCA* transformations.

Though *BRCA2*-changed BCs were guaranteed to be ER- or PR-positive, *BRCA1* carriers were guaranteed to be ER- or PR-negative compared to *BRCA1* non-carriers. Patients with *BRCA1* mutations also presented a higher stage at the hour of conclusion, while *BRCA2* mutation carriers had more precise lymph nodes. There were no distinctions in infection free endurance among *BRCA1* carriers, *BRCA2* carriers, and non-carriers. In any case, among non-TNBC patients, *BRCA2* transformation carriers showed diminished disease-free survival contrasted with *BRCA2* change non-carriers (risk proportion, 1.892; 95% certainty stretch, 1.132 to 3.161; $P = 0.013$).^{11,12}

As in the site of metastasis, it was appeared in *BRCA*-negative patients that there are two cases with lung metastasis, and every last one of the cerebrum and bone metastasis. These results were also consistent with a study that was shown on television, which found that 63 (15.1%) individuals had metastatic disease, with 13 (20.6%) having bone-only metastasis and 50 (79.4%) having instinctive metastases to the liver, lung, or brain. A significant number of the 354 patients with nonmetastatic sickness had poor clinical and neurotic highlights, including 237 (66.9%) with positive axillary lymph hubs, 182 (51.4%) with grade III, 100

(28.2%) with T3 or T4 infection, and 172 (48.6%) with positive lymphovascular attack. Both estrogen receptors (ERs) and progesterone receptors (PRs) were positive in 284 (68.1%) patients, while 52 (12.5%) others had ER-or PR-positive sickness and 80 (19.2%) had chemical receptor-negative infection. Not all patients were tried for HER2; notwithstanding, 122 (31.5%) of the 387 those tried patients were HER2-positive on immunohistochemical staining or fluorescence in situ hybridization.^{13,14}

The clinical stages of BC were logically shown in *BRCA*-with three cases in the IV stage, yet one case in III stage. On the other hand, one patient with II stage had *BRCA*+ alongside them. In 2014, Yu et al. looked examined the characteristics of BC from 55,387 irregular breast tumors from the Korean BC Registry and 181 *BRCA1/2* transformation carriers' cases (80 patients with *BRCA1* alteration and 101 patients with *BRCA2* transformation). In this report, middle patient age was fundamentally lower in the *BRCA1* and *BRCA2* change groups than in the registry group (37 years and 41 years versus 48 years; $P < 0.001$ for both). Tumor size was not diverse between *BRCA1* and *BRCA2* gatherings and the registry group. The extent of patients with axillary hub metastasis was not essentially unique

between *BRCA1* and library gatherings; nonetheless, axillary nodal association was available more regularly in the *BRCA2* group than in the registry group (45.5% versus 33.5%, $P = 0.002$).¹²

The *BRCA1* and *BRCA2* populations showed no significant relationships between tumor size and axillary nodal association. When compared to the library group's tumors, *BRCA1* tumors performed better (64.3 percent vs 27.5 percent, $P < 0.001$) In comparison to the registry group, the *BRCA1* group had a larger percentage of chemical receptor-negative tumors and a lower percentage of HER2-overexpressing tumors. TNBCs were more pervasive in the *BRCA1* group than in the registry group (61.3% versus 12.4%, $P < 0.001$). Moreover, chemical receptor articulation was not altogether unique between the *BRCA2* group and registry group. The recurrence of DCIS was lower in the *BRCA1* (3.7%) and *BRCA2* (5%) groups than in the registry group (10.3%).¹⁵

Conclusion

In this case report, we found one patient who had a *BRCA 1/2* mutation with a triple negative type of BC grade 3 and had a family history of BC. According to the existing theory, BC related to *BRCA 1/2* has aggressive

characteristics, high grade, and is often found in the triple negative type.

Informed Consent

Before beginning any medical or legal processes, we got the participants' informed consents; in other words, patients have given their consent to participate in formal scientific research ever since they registered at our facility. The verbal agreement from all patients regarding the data documentation, discussion, and possible publication of the cases had been obtained since we explained the main objective of this report is solely on scientific purpose.

Conflict of Interest

None declared.

References

1. World Health Organization. Global Health Estimates 2020: Deaths by Cause, Age, Sex, by Country and by Region, 2000-2019 [Internet]. WHO. 2020. Available from: <http://who.int/data/gho/data/themes/mortality-and-global-health-estimates/gh-leading-causes-of-death>
2. Sung H, Ferlay J, Siegel RL, Laversanne M, Soerjomataram I, Jemal A, et al. Global Cancer statistics 2020: GLOBOCAN estimates of incidence and mortality worldwide for 36 cancers in 185 countries. *CA Cancer J Clin.* 2021;71(3):209-49. doi:10.3322/caac.21660.
3. Fackenthal JD, Olopade OI. BC risk associated with *BRCA1* and *BRCA2* in diverse populations. *Nat Rev Cancer.* 2007;7(12):937-48. doi:10.1038/nrc2054.
4. Ripperger T, Gadzicki D, Meindl A, Schelegelberger B. BC susceptibility: current knowledge and implications for genetic counselling. *Eur J Hum Genet.* 2009;17(6):722-31. doi:10.1038/ejhg.2008.212.
5. Torre LA, Bray F, Siegel RL, Ferlay J, Lortet-Tiulent J, Jemal A. Global cancer statistics, 2012. *CA Cancer J Clin.* 2015;65(2):87-108. doi:10.3322/caac.21262.
6. Ford D, Easton DF, Stratton M, Narod S, Goldgar D, Devilee P, et al. Genetic heterogeneity and penetrance analysis of the *BRCA1* and *BRCA2* genes in breast cancer families. *Am J Hum Genet.* 1998;62(3):676-89. doi:10.1086/301749.
7. Antoniou A, Pharoah PDP, Narod S, Risch HA, Eyfjord JE, Hopper JL, et al. Average risks of breast and ovarian cancer associated with *BRCA1* or *BRCA2* mutations detected in case series unselected for family history: A combined analysis of 22 studies. *Am J Hum Genet.* 2003;72(5):1117-30.

doi:10.1086/375033.

8. Daly MB, Axilbund JE, Buys S, Crawford B, Farrell CD, Friedman S, et al. Genetic / familial assessment : Breast and ovarian. *Genet Couns.* 2010;8(5):562-94.

9. Gonzalez-Angulo AM, Timms KM, Liu S, Chen H, Litton JK, Potter J, et al. Incidence and outcome of *BRCA* mutations in unselected patients with triple receptor-negative breast cancer. *Clin Cancer Res.* 2011;17(5):1082-9. doi:10.1158/1078-0432.CCR-10-2560.

10. Kwong A, Wong LP, Wong HN, Law FBF, Ng EKO, Tang YH, et al. Clinical and pathological characteristics of Chinese patients with *BRCA* related breast cancer. *Hugo J.* 2009;3(1):63-76. doi:10.1007/s11568-010-9136-z.

11. Shin HR, Carlos MC, Varghese C. Cancer control in the Asia Pacific region: Current status and concerns. *Jpn J Clin Oncol.* 2012;42(10):867-81. doi:10.1093/jjco/hys077.

12. Lang GT, Shi JX, Hu X, Zhang CH, Shan L, Song CG, et al. The spectrum of *BRCA* mutations and

characteristics of *BRCA*-associated breast cancers in China: Screening of 2,991 patients and 1,043 controls by next-generation sequencing. *Int J Cancer.* 2017;141(1):129-42. doi:10.1002/ijc.30692.

13. Kim H, Choi DH. Distribution of *BRCA1* and *BRCA2* mutations in Asian patients with breast cancer. *J Breast Cancer.* 2013;16(4):357-65. doi:10.4048/jbc.2013.16.4.357.

14. Byung HS, Beom SK, Jeong KK, Hee JK, Soo JH, Jung SL, et al. Changing patterns in the clinical characteristics of Korean patients with breast cancer during the last 15 years. *Arch Surg.* 2006;141(2):155-60. doi:10.1001/archsurg.141.2.155.

15. Robson M, Im SA, Senkus E, Xu B, Domchek SM, Masuda N, et al. Olaparib for metastatic breast cancer in patients with a germline *BRCA* mutation. *N Engl J Med.* 2017;377(6):523-33. doi:10.1056/NEJMoa1706450. Erratum in: *N Engl J Med.* 2017;377(17):1700.

Table 1. Sample characteristics

Characteristics	<i>BRCA</i>	
	+	-
Family History	1	4
History of Cancer	-	5
Age		
<30	-	-
30-40	1 (20)	3 (60)
41-50	-	1 (20)
>50	-	-
Grade		
1	-	-
2	-	2 (40)
3	1	2 (40)
Stadium		
1	-	-
2	1 (20)	-
3	-	1 (20)
4	-	3 (100)
IHC		
ER		
Positive	-	2
Negative	1	2
PR		
Positive	-	1
Negative	1	3
HER2		
Positive	-	1
Negative	1	3
Nodal Metastasis		
Positive	-	3
Negative	1	1
Site of Metastasis		
Lung	-	2
Brain	-	1
Liver	-	-
Bone	-	1

BRCA: Breast cancer gene; ER: Estrogen receptor; HER2: Human epidermal growth factor receptor-2; IHC: Immunohistochemistry; PR: Progesterone receptor