

Detection of Radiation-Induced DNA Damage in Breast Cancer Patients by Using Gamma H2AX Biomarker: A Possible Correlation with Their Body Mass Index

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ABSTRACT

Radiotherapy is one of the most important options for treating breast cancer in humans. The development of biomarkers to monitor radiosensitivity is scarce. The aim of this study is to investigate the γ H2AX levels in the human blood samples 0.5 h after radiotherapy compared to the levels before radiotherapy in breast cancer patients in relation to their respective body mass index (BMI). Blood plasma samples were collected from a total of 20 breast cancer patients before and after radiotherapy to measure γ H2AX levels with an antibody against γ H2AX based on enzyme-linked immunosorbent assay technique. The median BMI of the patients was 30 kg/m². γ H2AX was differentially expressed in breast cancer patients before radiotherapy. γ H2AX levels significantly increased in 14 patients after radiotherapy (*P* = 0.006), whereas γ H2AX levels decreased in three patients after radiotherapy, and three patients were excluded. There was no correlation between γ H2AX values after radiotherapy and BMI (*P* = 0.5, *r* = 0.1). Our results suggest that γ H2AX can be used by ELISA technique to measure γ H2AX in the blood plasma of breast cancer patients undergoing radiotherapy and can be considered a biomarker of radiosensitivity.

Key words: Body mass index, DNA damage, H2AX, radiotherapy

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Introduction

Breast cancer is considered the second most common cancer; among women, breast cancer is the leading cause of death worldwide.^[1] Ionizing radiation is increasingly used in therapy and diagnosis; radiotherapy is used to treat 50%–60% of cancers^[2] and is the main treatment option for all stages of breast cancer.^[3] However, biological markers to measure radiosensitivity are important to monitor individual response to radiotherapy and evaluate its benefit. Several assays, including double-strand break (DSB) repair, chromosomal aberrations, and radiation-induced apoptosis

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Address for correspondence: Dr. Alkhansa S. Mahmoud, Department of Radiobiology, Sudan Atomic Energy Commission, Khartoum 11111, Sudan. E-mail: alkhansa.salih@gmail.com in *ex vivo*-irradiated blood lymphocytes, have been described as predictors of radiosensitivity.^[4] Biomarkers of radiosensitivity may also be useful in normal tissue toxicity. In addition, several factors are known to enhance tumor response to radiation exposure, including total dose, fractionation, tumor potential doubling time, hypoxia, and intrinsic radiosensitivity.^[5] DNA DSBs are one of the most important types of DNA damage. Ionizing radiation causes DNA damage,^[6] and DNA DSBs are in turn responsible for the activation of three phosphatidylinositol 3-kinase-like kinases: ataxia telangiectasia-mutated (ATM), ataxia telangiectasia and Rad3-related, and DNA-dependent protein kinase (DNA-PK),

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which catalyze the phosphorylation of H2AX at serine 139 in the C-terminus named YH2AX.^[7] The phosphorylated H2AX plays a functional and structural role in DNA damage recognition and repair. Therefore, yH2AX may serve as a sensitive marker for detecting DSBs, which in turn may indicate genomic instability and potentially contribute to cancer development and progression.^[8] In breast cancer, yH2AX has been associated with triple-negative breast cancer,^[9] breast cancer gene (BRCA) 1, and p53 mutations.^[10] Meanwhile, yH2AX is a sensitive indicator of DNA DSBs and is considered an important regulator of DNA damage and repair.[11] The commonly used techniques to measure yH2AX level are immunostaining (analyzed by fluorescence microscopy), western blot, flow cytometry, and enzyme-linked immunosorbent assay (ELISA).^[12] The aim of this study is to measure the YH2AX concentration in the blood plasma of breast cancer patients before and after radiotherapy 0.5 h after sample collection by ELISA and then to investigate a possible relationship between YH2AX concentration and body mass index (BMI).

Materials and Methods

Study population

A total of 20 breast cancer patients who received chemotherapy and/or surgery before being subjected to radiotherapy were included. Patients were randomly selected to participate in this study according to whether they had undergone radiotherapy. Blood samples were collected before and after radiotherapy. All patients were informed of their concerns through individual written form. In this regard, the study was approved by the Ministry of Health in Khartoum, Sudan.

Sample preparation and analysis

Venous blood samples were collected in EDTA from breast cancer patients before and after radiotherapy. Samples were centrifuged at 2500 rpm to separate plasma for analysis. ELISA was used to quantitatively measure γ H2AX (Human Gamma H2AX ELISA Kit; Cat. No. SG-14818, Sino Geneclon Biotech Co., Ltd). The γ H2AX antibody was used based on the principle of ELISA technique. Then, the optical density at 450 nm was determined, and the concentrations were estimated using the standard curve for quantification of γ H2AX.

Body mass index

BMI was calculated as weight in kilograms divided by the square of height in meters (kg/m²) and classified as categories of normal weight (BMI <25), overweight (\geq 25), and obese (\geq 30) breast cancer patients.

Statistical analysis

Statistical analysis was performed with GraphPad Prism 8 software (GraphPad Software Inc., San Diego, California, USA). Correlation coefficient was used for the association between γ H2AX levels and BMI. Paired *t*-test was used to determine the significance between groups. Significance was considered at the level of P < 0.05.

Results

A total of 20 breast cancer patients with a median age of 44 years were considered for the present study. The median BMI was

30 kg/m²; therefore, mainly obese patients participated in the present study. Patients were treated with an absorbed dose of 40.5 Gy at 2.7 Gy/fraction. The γ H2AX values for all patients before and after radiotherapy are explained in Table 1. Of 20 patients, three were excluded because of very low γ H2AX concentrations. In 14 patients, γ H2AX levels were increased by 0.5 h after radiotherapy (*P* = 0.006), as shown in Figure 1. Meanwhile, no correlation was found between BMI and γ H2AX levels (*P* = 0.5, *r* = 0.1), as shown in Figure 2.

Discussion

yH2AX is a biomarker for detecting DNA double-strand damage. In this study, the level of YH2AX was measured in the blood samples from breast cancer patients who had undergone radiotherapy. Our results showed that YH2AX is expressed in breast cancer patients. In this regard, we found that another study confirmed this finding.^[8] In these findings, increased $\gamma \mathrm{H2AX}$ level seems to be associated with high BMI in this study because most of the patients showed high BMI; however, several studies found that obesity is associated with breast cancer.[13-15] In another study, an increase in YH2AX levels was observed in breast cancer with a BMI of <25.^[16] Obesity is a condition characterized by an increase in the mass of body adipose tissue and is associated with disturbances in lipid and glucose metabolism, chronic inflammation, and oxidative stress.^[17] Obesity is associated with DNA damage accumulation,^[18] and low body weight has been associated with a reduction in the extent of DNA damage.^[19] In addition, obesity can induce DNA damage and inhibit DNA repair mechanisms. The accumulation of DNA damage leading to the activation of various proteins can induce adipocyte differentiation, inflammation, and cell metabolism disorders.^[20] A relationship between BMI and DNA damage has been established.^[13] yH2AX levels increased in obese patients after 0.5 h of radiotherapy, indicating a strong DNA damage response leading to increased YH2AX levels after radiotherapy. After irradiation, histone H2AX is rapidly phosphorylated by ATM and/or DNA-PK at or near DNA DSB to form γ H2AX.^[21] Several studies reported that yH2AX is a good biomarker for



Figure 1: Mean levels of breast cancer patients showed increased γ H2AX levels postradiotherapy (*P* = 0.006)



Serial number of patients	BMI (kg/m²)	Plasma γH2AX preradiotherapy (pg/ml)	Plasma γH2AX postradiotherapy (pg/ml)	Status of γH2AX at 0.5 h
1	30	597	1692	+
2	26	1126	1710	+
3	31	1656	2356	+
4	30	2454	2975	+
5	28	516	1225	+
6	32	1925	2212	+
7	28	1025	1396	+
8	30	1396	1512	+
9	20	1037	3935	+
10	27	2030	220	-
11	21	1185	1378	+
12	34	1288	2894	+
13	34	929	4509	+
14	37	9045	4005	-
15	26	4050	3915	-
16	29	148	705	+
17	31	3379	8700	+

Table 1: Individual γ H2AX levels expressed in a study population of breast cancer patients with normal weight, overweight, and obese

+: Increase and -: Decrease, BMI: Body mass index



Figure 2: Correlation between body mass index and γ H2AX (P = 0.5, r = 0.1)

radiosensitivity of human tumor cells.^[22-24] Phosphorylation of Υ H2AX is induced by DNA DSBs. Since tumor cells usually exhibit deficiencies in DNA damage response, it has been suggested that constitutive expression of histone γ H2AX may indicate DNA damage response dysfunction and genomic instability.^[25] Several studies have found that radiosensitivity increases with genetic defects in DNA repair^[26] and that the DNA damage marker γ H2AX can be used to assess the radiation exposure of biological samples.^[27] Thus, γ H2AX can be used as a sensitive biomarker for DNA damage caused by ionizing radiation^[28] and can be used as a therapeutic marker against cancer.^[29]

Our data are limited by sample size, and further studies are recommended to support our findings. YH2AX can be measured in the bloodstream, and there are several techniques to measure its concentration, including ELISA. The development of new biomarkers to monitor radiosensitivity could be useful to optimize treatments and improve patient care. The initial effect of ionizing radiation or its early detection, e.g., after 0.5 h, is important for studying the DNA damage response, and this could be revealed by γ H2AX. However, the changes in γ H2AX concentrations after radiotherapy in breast cancer patients seem to enhance this histone variant by ionizing radiation.

Conclusion

Individual γ H2AX values in breast cancer patients with normal and high BMI increased significantly after radiotherapy. The decrease in γ H2AX levels in some patients might indicate a low DNA damage response. On the other hand, no correlation was found between the increase in γ H2AX and BMI in our data. The results show that γ H2AX can be measured in the blood plasma of breast cancer patients and can be considered as a biomarker for radiosensitivity.

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Conflicts of interest

There are no conflicts of interest.

References

1. Ferlay J, Soerjomataram I, Dikshit R, Eser S, Mathers C, Rebelo M, *et al.* Cancer incidence and mortality worldwide: sources, methods and major patterns in GLOBOCAN 2012. Int J Cancer

2015;136:359-86.

- Rosenblatt E, Izewska J, Anacak Y, Pynda Y, Scalliet P, Boniol M, et al. Radiotherapy capacity in European countries: An analysis of the Directory of Radiotherapy Centres (DIRAC) database. Lancet Oncol 2013;14:e79-86.
- Wang W. Radiotherapy in the management of early breast cancer. J Med Radiat Sci 2013;60:40-6.
- Chua ML, Rothkamm K. Biomarkers of radiation exposure: Can they predict normal tissue radiosensitivity? Clin Oncol (R Coll Radiol) 2013;25:610-6.
- Forker LJ, Choudhury A, Kiltie AE. Biomarkers of tumour radiosensitivity and predicting benefit from radiotherapy. Clin Oncol (R Coll Radiol) 2015;27:561-9.
- Jeggo PA, Löbrich M. DNA double-strand breaks: Their cellular and clinical impact? Oncogene 2007;26:7717-9.
- Hanasoge S, Ljungman M. H2AX phosphorylation after UV irradiation is triggered by DNA repair intermediates and is mediated by the ATR kinase. Carcinogenesis 2007;28:2298-304.
- Varvara PV, Karaolanis G, Valavanis C, Stanc G, Tzaida O, Trihia H, et al. Gamma-H2AX: A potential biomarker in breast cancer. Tumour Biol 2019;41:1-7.
- Nagelkerke A, van Kuijk SJ, Sweep FC, Nagtegaal ID, Hoogerbrugge N, Martens JW, *et al.* Constitutive expression of γ-H2AX has prognostic relevance in triple negative breast cancer. Radiother Oncol 2011;101:39-45.
- Deniz M, Kaufmann J, Stahl A, Gundelach T, Janni W, Hoffmann I, et al. In vitro model for DNA double-strand break repair analysis in breast cancer reveals cell type-specific associations with age and prognosis. FASEB J 2016;30:3786-99.
- Novotna E, Tichy A, Foltanova K, Vavrova J. Role of γ-H2AX in DNA damage response and its possible clinical applications. Mil Med 2011;80:169-77.
- Pouliliou S, Koukourakis MI. Gamma histone 2AX (γ-H2AX) as a predictive tool in radiation oncology. Biomarkers 2014;19:167-80.
- Włodarczyk M, Jabłonowska-Lietz B, Olejarz W, Nowicka G. Anthropometric and dietary factors as predictors of DNA damage in obese women. Nutrients 2018;10:578.
- Bukhari SA, Rajoka MI, Ibrahim Z, Jalal F, Rana SM, Nagra SA. Oxidative stress elevated DNA damage and homocysteine level in normal pregnant women in a segment of Pakistani population. Mol Biol Rep 2011;38:2703-10.
- James FR, Wootton S, Jackson A, Wiseman M, Copson ER, Cutress RI. Obesity in breast cancer – What is the risk factor? Eur J Cancer 2015;51:705-20.

- Barba M, Vici P, Pizzuti L, Di Lauro L, Sergi D, Di Benedetto A, et al. Body mass index modifies the relationship between γ-H2AX, a DNA damage biomarker, and pathological complete response in triplenegative breast cancer. BMC Cancer 2017;17:101.
- 17. Scherer PE, Hill JA. Obesity, diabetes, and cardiovascular diseases: A compendium. Circ Res 2016;118:1703-5.
- Zaki M, Basha W, El-Bassyouni HT, El-Toukhy S, Hussein T. Evaluation of DNA damage profile in obese women and its association to risk of metabolic syndrome, polycystic ovary syndrome and recurrent preeclampsia. Genes Dis 2018;5:367-73.
- Heilbronn LK, de Jonge L, Frisard MI, DeLany JP, Larson-Meyer DE, Rood J, et al. Effect of 6-month calorie restriction on biomarkers of longevity, metabolic adaptation, and oxidative stress in overweight individuals: A randomized controlled trial. JAMA 2006;295:1539-48.
- Włodarczyk M, Nowicka G. Obesity, DNA damage, and development of obesity-related diseases. Int J Mol Sci 2019;20:1146.
- Redon CE, Weyemi U, Parekh PR, Huang D, Burrell AS, Bonner WM. γ-H2AX and other histone post-translational modifications in the clinic. Biochim Biophys Acta 2012;1819:743-56.
- Zhao J, Guo Z, Pei S, Song L, Wang C, Ma J, et al. pATM and γH2AX are effective radiation biomarkers in assessing the radiosensitivity of ¹²C⁶⁺in human tumor cells. Cancer Cell Int 2017;17:49.
- Kawashima S, Kawaguchi N, Taniguchi K, Tashiro K, Komura K, Tanaka T, *et al.* γ-H2AX as a potential indicator of radiosensitivity in colorectal cancer cells. Oncol Lett 2020;20:2331-7.
- Lee Y, Wang Q, Shuryak I, Brenner DJ, Turner HC. Development of a high-throughput γ-H2AX assay based on imaging flow cytometry. Radiat Oncol 2019;14:150.
- Bonner WM, Redon CE, Dickey JS, Nakamura AJ, Sedelnikova OA, Solier S, *et al.* GammaH2AX and cancer. Nat Rev Cancer 2008;8:957-67.
- Yin X, Mason J, Lobachevsky PN, Munforte L, Selbie L, Ball DL, et al. Radiation therapy modulates DNA repair efficiency in peripheral blood mononuclear cells of patients with non-small cell lung cancer. Int J Radiat Oncol Biol Phys 2019;103:521-31.
- Rothkamm K, Horn S. Gamma-H2AX as protein biomarker for radiation exposure. Ann Ist Super Sanita 2009;45:265-71.
- Redon CE, Dickey JS, Bonner WM, Sedelnikova OA. γ-H2AX as a biomarker of DNA damage induced by ionizing radiation in human peripheral blood lymphocytes and artificial skin. Adv Space Res 2009;43:1171-8.
- Sedelnikova OA, Bonner WM. GammaH2AX in cancer cells: A potential biomarker for cancer diagnostics, prediction and recurrence. Cell Cycle 2006;5:2909-13.

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