



BRIEF COMMUNICATION

Identification of the Thioredoxin-Like 2 Autoantibody as a Specific Biomarker for Triple-Negative Breast Cancer

Jee Min Chung^{1,2,3,*}, Yongsik Jung^{4,*}, Young Pil Kim⁵, Jinsue Song⁵, Soyeon Kim^{1,2,3}, Ji Young Kim⁴, Mira Kwon^{1,2,3}, Jung Hyun Yoon^{1,2,3}, Myo-Deok Kim⁵, Jun-kyoung Lee⁵, Da-Yoon Chung⁵, Seo Yun Lee^{1,2}, Jooseong Kang⁵, Ho Chul Kang^{1,2,3}

¹Genomic Instability Research Center and Departments of ²Physiology, ³Biomedical Sciences, and ⁴Surgery, Ajou University School of Medicine, Suwon; ⁵Department of Bio-Engineering, Life Science RD Center, Sinil Pharmaceutical Co., Seongnam, Korea

Triple-negative breast cancer (TNBC) has a higher risk of death within 5 years of being diagnosed than the other forms of breast cancer. It is the second leading cause of death due to cancer among women. Currently, however, no diagnostic blood-based biomarker exists to identify the early stages of TNBC. To address this point, we utilized a human protein microarray system to identify serum autoantibodies that showed different expression patterns between TNBC and normal serum samples, and identified five autoantibodies showing TNBC-specific expression.

Among them, we selected the thioredoxin-like 2 (TXNL2) autoantibody and evaluated its diagnostic relevance by dot blot analysis with the recombinant TXNL2 protein. We demonstrated that the TXNL2 autoantibody showed 2- to 6-fold higher expression in TNBC samples than in normal samples suggesting that serum TXNL2 autoantibodies are potential biomarkers for TNBC.

Key Words: Autoantibodies, Biomarkers, Breast neoplasms, Protein array analysis

Breast cancer has been divided into five subtypes based on their intrinsic molecular entities: normal-like, luminal A, luminal B, human epidermal growth factor receptor 2-enriched, and basal-like [1]. Among them, basal-like breast cancer, commonly known as triple-negative breast cancer (TNBC), has attracted much interest due to its poor prognosis and lack of effective targeted therapies [2]. Intensive studies are, therefore, currently underway to identify the underlying molecular mechanism that drives the growth and metastasis of TNBC [3]. In addition, as early diagnosis enables early treatment and

Correspondence to: Ho Chul Kang

Department of Physiology, Ajou University School of Medicine, 206 World cup-ro, Yeongtong-gu, Suwon 16499, Korea Tel: +82-31-219-5044, Fax: +82-31-219-5049 E-mail: hckang@ajou.ac.kr

*These authors contributed equally to this work.

This work was supported by the Technology Innovation Program (R0003804) funded by the Ministry of Trade, Industry & Energy, Republic of Korea and the Korea Health Technology R&D Project through the Korea Health Industry Development Institute, funded by the Ministry of Health & Welfare, Republic of Korea (HI16C0992). This work was also supported in part by the Brain Research Program through the National Research Foundation of Korea funded by the Ministry of Science, ICT & Future Planning (NRF-2015M3C7A1029038).

Received: January 4, 2018 Accepted: March 4, 2018

helps significantly delay disease progression and expand the overall lifetime of patients, the discovery of specific or selective biomarkers for early stage TNBC has become an attractive research area [4]. Nonetheless, there is still no specific biomarker that can provide a definitive diagnosis of TNBC. Current cancer screening is based on a combination of mammography and self-examination, but their prognostic value remains controversial [5]. No laboratory tests utilizing serum, saliva, or urine samples have been established to be effective in the primary diagnosis of breast cancer [5]. Thus, early stage biomarkers of TNBC to establish or develop precise early diagnostic methods that meet the criteria for an accurate, relatively noninvasive, reliable, and low-cost test remain to be discovered [6,7]. Recently, disease-specific autoantibodies derived from patient serum have become useful and cost-effective biomarkers, providing a potential gateway for diagnosing cancer [8]. Compared with other biomarkers such as DNA, RNA, or proteins, autoantibodies have several advantages in that they are structurally stable, specific, and present in large amounts in serum [7]. Thus, disease-specific autoantibodies are currently being investigated in a range of cancers to overcome the obstacles encountered for their early detection [8-10]. In an effort to identify a useful early stage autoantibody biomarker in the sera from TNBC samples, we utilized a hu88 Jee Min Chung, et al.

man protein microarray system containing 16,368 human recombinant proteins. Human serum proteome analysis demonstrated that approximately 3,700 separate proteins exist in a physiological state. One issue with using human serum to identify biomarkers is the high background protein content, which causes noise signals under certain experimental conditions [11]. To minimize background signals in the protein microarray, the amount of serum was adjusted so that up to 200 µg of serum proteins was used, which greatly reduced the interference by noise signals (data not shown). To identify autoantibodies in TNBC samples, we used the HuProtTM human proteome microarray v2.0 (CDI Laboratories Inc., Mayaguez, USA), each containing 16,368 unique full-length human protein antigens (Figure 1A). In total, 20 human serum samples (10 TNBC and 10 normal) were subjected to protein microarray analysis. This study was approved by the Institutional Review Board of Ajou University Hospital (IRB No. AJIRB-BMR-KSP-15-362) and written informed consent was obtained from all participants. The result of protein microarray was analyzed

by GenePix analysis software (Molecular Devices, San Jose, USA) (Figure 1B). Briefly, microarray slides were blocked with filtrated dry milk and were then incubated with serum samples. After washing, the arrays were incubated with antihuman IgG (H+L) conjugated to AlexaFluor 546 (Cat. No. A-21445; Invitrogen, Carlsbad, USA). The arrays were then washed, dried, and immediately scanned with a GenePix 4000B Fluorescence Scanner (Molecular Devices). The GenePix analysis software determined that five autoantibodies, thioredoxinlike 2 (TXNL2), ADP-ribosylhydrolase like 1 (ADPRHL1), glycyl-TRNA synthetase (GARS), spinocerebellar ataxia type 3 protein (ATXN3) and solute carrier family 16 member 4 (SLC16A4), had a significantly higher expression in the TNBC samples than in the normal samples. The differential expression of these five autoantibody biomarkers is shown in Figure 1B. Among them, we selected the TXNL2 autoantibody biomarker not only because it showed the largest difference between the TNBC and normal samples but also because the TXNL2 autoantigen is known as a potential diagnostic in-

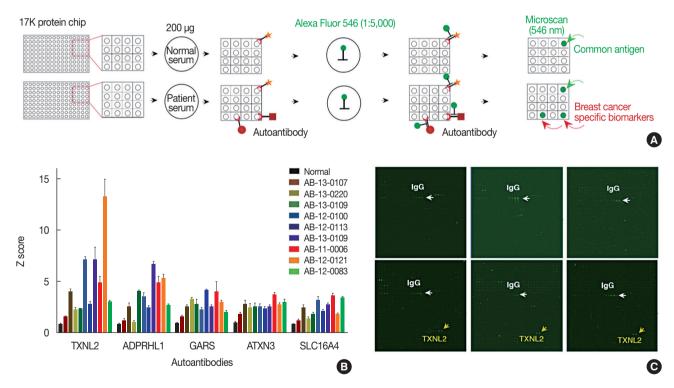


Figure 1. Identification of five autoantibodies for triple-negative breast cancer (TNBC) biomarkers using a 17K human protein microarray system. (A) To identify novel TNBC-specific biomarkers, normal or TNBC patient sera containing autoantibodies were applied on 17K protein microarray. TNBC-specific autoantibodies were detected by anti-human Alexa Flour 546 (Invitrogen) and scanned by GenePix 4000B (Molecular Devices). Signal intensity for each spot was obtained as the ratio of foreground to background signals and was normalized with the glutathione s-transferase signal intensity. The mean signal intensity of each protein on the chip was calculated. (B) Five autoantibodies, thioredoxin-like 2 (TXNL2), ADP-ribosylhydrolase like 1 (ADPRHL1), glycyl-TRNA synthetase (GARS), spinocerebellar ataxia type 3 protein (ATXN3), and solute carrier family 16 member 4 (SLC16A4), were determined by the GenePix analysis. The differential expression of these five autoantibody biomarkers were calculated by statistical analysis. (C) Highpower image of 17K protein microarray. Yellow arrow indicated TXNL2 autoantibody and white arrow was shown IgG as a positive control. TXNL2 autoantibody specifically and strongly interacted with the TXNL2 antigen.

dicator of breast cancer [12]. Figure 1C shows that the TXNL2 autoantibody specifically and strongly interacted with the TXNL2 antigen, suggesting that it is a highly specific biomarker for TNBC.

For independently verifying the TXNL2 autoantibody as a TNBC-specific biomarker, we attempted to purify the recombinant TXNL2 antigen from BL21-Gold(DE3)pLysS Competent Cells (Agilent, Santa Clara, USA). The purity of the TXNL2 protein is directly associated with the reliability of the TXNL2 autoantibody as a TNBC-specific biomarker. To obtain high-purity and sufficiently large yields of recombinant TXNL2, the full-length cDNA of TXNL2 was optimized for codon usage. Fortunately, we found that glutathione s-transferase (GST)-TXNL2 protein was successfully expressed and purified using a Glutathione Sepharose 4B (GE Healthcare, Little Chalfont, UK). Most GST-TXNL2 proteins were observed in the soluble fraction after sonication and migrated to ~63 kDa on an sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) as a single band (Figure 2A). Densitometric analysis by the ImageJ program (https://imagej.

net/) indicated that GST-TXNL2 comprised 42% of the total proteins. As shown in Figure 2B, following affinity chromatography, the fractions containing GST-TXNL2 were further separated by using gel filtration with Superdex 75 prep grade resin (GE Healthcare) that displays a high resolution for proteins in the molecular weight range of 3,000 to 70,000 Da. Finally, we found that the final eluted fractions contained highly pure GST-TXNL2 (Figure 2C). The molecular weight and homogeneity of the sample were checked using a TSKgel-S3000SW_{XL} (Tosoh Bioscience, Tokyo, Japan) (Figure 2D). Taken together, these data suggest that GST-TXNL2 protein might be a usable antigen for testing of the TXNL2 autoantibody as a TNBC-specific biomarker.

To validate the specificity of the TXNL2 autoantibody for TNBC, we employed a quantitative dot blot system. The dot blot method is the best technique for rapidly assessing the quantity of a target antigen across many samples at once. It is also a popular method for screening antibodies for target specificity. The feasibility of the dot blot analysis was verified by analyzing the GST protein and the specificity of a rabbit

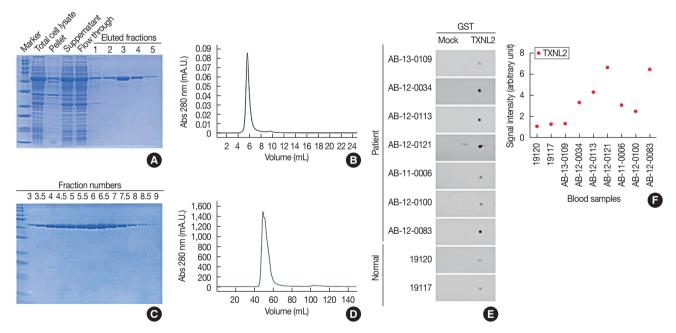


Figure 2. Thioredoxin-like 2 (TXNL2) autoantibody is a potential biomarker for diagnosing triple-negative breast cancer (TNBC). (A) Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) analysis of the recombinant glutathione s-transferase (GST)-tagged TXNL2 protein. Total cell lysate and each subcellular fractions of the isopropyl β-D-1-thiogalactopyranoside-induced *Escherichia coli* BL21 (DE3) expressing GST-tagged TXNL2 protein was loaded in SDS-PAGE. The soluble fraction containing GST-tagged TXNL2 protein was applied in a Glutathione Sepharose 4B resin (GE Healthcare). Subsequently resin conjugated TXNL2 was detached by elution buffer containing reduced glutathione. (B) Superdex 75 (GE Healthcare) gel-filtration fractionation profile for the further purification of the target protein. (C) SDS-PAGE profile of GST-tagged TXNL2 protein eluted from gel filtration. (D) A high-performance liquid chromatography chromatogram of recombinant GST-tagged TXNL2 protein. (E) Dot blot analysis of TXNL2 for detection of TXNL2 autoantibody in TNBC serum sample. Purified GST or GST tagged TXNL2 protein was spotted on nitrocellulose membrane and applied with normal or TNBC patient sera. GST was used as a negative control. Autoantibody against TXNL2 was detected by anti-human horseradish peroxidase conjugated antibody and developed using an enhanced chemiluminescent substrate. (F) Graph shows that amount of TXNL2 autoantibody derived from human sera. TXNL2 autoantibody was expressed 2- to 6-fold higher in TNBC serum than in normal. Densitometric analysis was performed using the ImageJ program (https://imagej.net/).

90 Jee Min Chung, et al.

anti-GST antibody was evaluated (data not shown). For the dose-response curve, we serially diluted GST proteins from 0.1 µg to 2 µg under nondenaturing condition. Dot blot analysis yielded a linear curve between 0.2 µg and 1.8 µg of protein with a coefficient of determination (R2) of 0.999 when simple linear regression analysis was performed (data not shown). Beyond 2 µg, the signal intensity started to plateau. Based on the control experiment, we loaded 1.5 µg GST or GST-TXNL2 proteins onto the nitrocellulose membrane; we then incubated each membrane with normal or TNBC sera, which were used in the protein microarray experiment, as shown in Figure 2E. Dot blot data revealed that the amount of the TXNL2 autoantibody derived from each serum sample showed similar expression levels to those obtained using the protein microarray, indicating that the protein microarray data are reliable for detecting autoantibodies from sera (Figure 2F). In addition, we found that the TXNL2 autoantibody was more highly expressed in TNBC samples than in normal samples. Taken together, these results suggest that the TXNL2 autoantibody is a strong serum biomarker to detect TNBC.

Clinically, autoantibodies are currently being used to diagnose various immune diseases. However, they are also found in various cancers and neurological diseases, indicating that a diagnostic method using autoantibodies can be applied to a wide range of diseases beyond immunological diseases. That said, autoantibodies against cancer-related antigens may be detected in the sera of patients with various types of cancer, and there are still few ways to identify disease-specific autoantibodies from patient blood. Of these methods, protein microarrays have the advantage of being able to identify autoantibodies efficiently by using only a small amount of blood. For this reason, many research groups are now trying to identify autoantibodies as specific biomarkers for a number of diseases [13].

Recently, it has been reported that TXNL2 is also a TNBCspecific marker that is highly expressed in breast cancer patients [12]. Moreover, the same study determined that TXNL2 protein is directly linked to the growth and metastasis of breast cancer cells and is associated with decreased overall patient survival, suggesting that abnormal expression of TXNL2 is directly or indirectly linked to TNBC initiation or progression [12]. TXNL2 has a unique protein structure consisting of a Trx homology region in the N terminus, followed by two tandem repeats of the Grx domains, which play a role in reactive oxygen species (ROS) detoxification and cell viability [12]. Remarkably, the deletion of TXNL2 gene in mice leads to embryonic lethality [14], indicating its role in protecting cells against oxidative stress during embryogenesis. In addition, TXNL2 is a critical factor in regulating ROS levels and nuclear factor κB activity for controlling tumor development and metastasis in primary human breast cancers, demonstrating that it has essential roles in cancer cells [12]. In the current study, we identified the TXNL2 autoantibody as a TNBC-specific biomarker. Although the disease-specific linkage between autoantibodies and autoantigens is not clear, our data suggests that abnormal expression or dysfunctions of TXNL2 might be closely related to the generation of the TXNL2 autoantibody in TNBC.

CONFLICT OF INTEREST

The authors declare that they have no competing interests.

REFERENCES

- Perou CM, Sørlie T, Eisen MB, van de Rijn M, Jeffrey SS, Rees CA, et al. Molecular portraits of human breast tumours. Nature 2000;406:747-52.
- Kalimutho M, Parsons K, Mittal D, López JA, Srihari S, Khanna KK. Targeted therapies for triple-negative breast cancer: combating a stubborn disease. Trends Pharmacol Sci 2015;36:822-46.
- Collignon J, Lousberg L, Schroeder H, Jerusalem G. Triple-negative breast cancer: treatment challenges and solutions. Breast Cancer (Dove Med Press) 2016;8:93-107.
- Fleisher B, Clarke C, Ait-Oudhia S. Current advances in biomarkers for targeted therapy in triple-negative breast cancer. Breast Cancer (Dove Med Press). 2016;8:183-97.
- Wahba HA, El-Hadaad HA. Current approaches in treatment of triplenegative breast cancer. Cancer Biol Med 2015;12:106-16.
- Ramachandran N, Srivastava S, Labaer J. Applications of protein microarrays for biomarker discovery. Proteomics Clin Appl 2008;2:1444-59.
- Conrad K, Roggenbuck D, Reinhold D, Sack U. Autoantibody diagnostics in clinical practice. Autoimmun Rev 2012;11:207-11.
- Xu YW, Peng YH, Chen B, Wu ZY, Wu JY, Shen JH, et al. Autoantibodies as potential biomarkers for the early detection of esophageal squamous cell carcinoma. Am J Gastroenterol 2014;109:36-45.
- Zaenker P, Ziman MR. Serologic autoantibodies as diagnostic cancer biomarkers: a review. Cancer Epidemiol Biomarkers Prev 2013;22: 2161-81
- Macdonald IK, Parsy-Kowalska CB, Chapman CJ. Autoantibodies: opportunities for early cancer detection. Trends Cancer 2017;3:198-213.
- 11. Pieper R, Gatlin CL, Makusky AJ, Russo PS, Schatz CR, Miller SS, et al. The human serum proteome: display of nearly 3700 chromatographically separated protein spots on two-dimensional electrophoresis gels and identification of 325 distinct proteins. Proteomics 2003;3:1345-64.
- 12. Qu Y, Wang J, Ray PS, Guo H, Huang J, Shin-Sim M, et al. Thioredoxin-like 2 regulates human cancer cell growth and metastasis via redox homeostasis and NF-kappaB signaling. J Clin Invest 2011;121:212-25.
- Zhu H, Cox E, Qian J. Functional protein microarray as molecular decathlete: a versatile player in clinical proteomics. Proteomics Clin Appl 2012;6:548-62.
- Cha H, Kim JM, Oh JG, Jeong MH, Park CS, Park J, et al. PICOT is a critical regulator of cardiac hypertrophy and cardiomyocyte contractility. J Mol Cell Cardiol 2008;45:796-803.