

In Situ Evaluation of Estrogen Receptor Dimers in Breast Carcinoma Cells: Visualization of Protein-Protein Interactions

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The estrogen receptor (ER) functions as a dimer and is involved in several different biological functions. However ER dimeric proteins have not been identified by in situ methodologies. Structured illumination microscopy (SIM) has been recently developed. which enabled the localization of protein and protein interaction. Therefore, in this study, we firstly demonstrated that ERs formed both homodimers and heterodimers in breast carcinoma cell lines using Nikon's SIM (N-SIM). ERα/α homodimers were detected in the nuclei of both ERα-positive MCF-7 and T-47D cells; 23.0% and 13.4% of ERα proteins formed ERa/a homodimers, respectively. ERa/B heterodimers were also detected in MCF-7 and T-47D. Approximately 6.6% of both ER α and ER β 1 proteins formed ER α/β 1 heterodimers in MCF-7. In addition, 18.1% and 22.4% of ERa and ERB proteins formed ERa/β2 heterodimers and ERa/β5 heterodimers in MCF-7, respectively. In addition, by using proximity ligation assay (PLA) in MCF-7, estradiol-induced ER α/α homodimers and ER α/β 1 heterodimers were both detected after 15 to 45 min of treatment and at 15 min, respectively. The percentage of total ER proteins could also be determined using N-SIM. By using both methods, it has become possible to evaluate precise localization and ratio of ER dimers among different cell types.

Key words: structured illumination microscopy, proximity ligation assay, estrogen receptor dimer, protein-protein interaction, breast cancer

I. Introduction

Estrogen plays a physiologically important role in various organs [2, 26]. It is also well-known that estrogen exerts cell proliferative effects on breast carcinomas, and the effects of estrogen are principally mediated by specific receptor, estrogen receptor (ER) [21, 22]. ER has two isoforms, ER α and ER β , which form either homo (α/α , β/β) or heterodimers (α/β), each harboring different biological functions [4]. In breast carcinoma cells, it is well known

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that an ER β signal disrupts estrogen-induced cell proliferation via an ER α signaling pathway [14]. This ER dimerization pattern has also been known to exert different functions on intracellular estrogen signaling pathways. The ER α/α homodimer was reported to stimulate estrogeninduced cell proliferation, whereas the ER β/β homodimer exerts inhibitory effects on estrogen-dependent cell proliferation in breast carcinoma cells [14]. In addition, ER α/β heterodimers was also known to inhibit cell proliferation because ER β inhibits ER α -mediated cell proliferation through the co-expression of ER α and ER β in estrogen responsive cells [14]. Five different splice variants of ER β isoforms, designated ER β 1–5, has been so far reported [16]. ER β 1 and ER β 2 are also known as wild-type ER β and ER β cx, respectively. High levels of ER β 1, ER β 2, and ER β 5

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expression were reported in normal mammary gland and breast carcinoma cell lines [16]. In addition, nuclear ER β 2 and ER β 5, but not ER β 1, were reported to promote survival in breast cancer patients [13, 24]. However, Chantzi *et al.* reported that ER β 2 status was associated with adverse clinical outcome in ER α -negative breast cancer patients [3]. Ogawa *et al.* also reported that ER β 2 preferentially formed a heterodimer with ER α , rather than with ER β 1, thereby inhibiting DNA binding by ER α [17]. In addition, ER β 2, ER β 4, and ER β 5 do not necessarily form homodimers with themselves, but it readily form a heterodimer with ER β 1 [15]. Therefore, it has become pivotal to evaluate the status of ER dimers in breast cancer patients in order to obtain a better understanding of the potential effects of estrogen on carcinoma cell proliferation.

ER dimeric proteins have largely been identified by bioluminescence resonance energy transfer (BRET)/förster resonance energy transfer (FRET) analyses but application of in situ methodologies has not been reported, which has made it rather impossible to localize homo- or heterodimers in individual estrogen-responsive cells [1, 20, 27]. We have previously reported the detection of ERa homodimers and ER α/β heterodimers in breast cancer cells and tissues by using a proximity ligation assay (PLA) [10]. PLA was developed in order to visualize protein-protein interactions [25]. In addition, a specific immunohistochemical staining method using dual binders was reported as a tool for visualization of protein-protein interactions [28]. Recently, structured illumination microscopy (SIM) has been developed [7, 23]. Owing to the diffraction limit of light, the resolution of conventional fluorescence microscopy is rather limited to ~200 nm [29, 32]. In submicrometer-sized spines, virtually all the proteins could therefore appear as more or less co-localized, which prevented a detailed analysis of spatial relationships [29, 32]. The dimerization of some membrane receptors, such as epidermal growth factor receptor induced by its ligand, was evaluated by superresolution microscopic analysis [31]. However, the study regarding the detection of nuclear receptor dimers using super-resolution microscopic N-SIM (Nikon's structured illumination microscopy) analysis has not been reported to the best of our knowledge. Therefore, in this study, we firstly demonstrated the levels of the ER α/α homodimer and ER α/β isoform heterodimers formation in breast carcinoma cell lines using both N-SIM analysis and PLA. We also evaluated the effects of estradiol (E2) treatment on the status of ER α/α homodimer and ER α/β isoform heterodimers in breast carcinoma cell lines.

II. Materials and Methods

Cell lines and cell culture

The human breast carcinoma cell lines MCF-7, T-47D, and MDA-MB-231 were purchased from the American Type Cell Culture (ATCC; Manassas, VA). Cells were cultured in RPMI-1640 medium (Sigma-Aldrich, St Louis, MO) supplemented with 10% fetal bovine serum (FBS; Nichirei Biosciences, Tokyo, Japan) and 100 μ g/mL penicillin/streptomycin (Invitrogen, CA, USA). Cells were incubated in a humidified (37°C, 5% CO₂) incubator and passaged upon reaching 80% confluence.

Effects of estradiol on ERa expression in MCF-7 cells

MCF-7 cells were seeded at a density of 1.0×10^5 cells per mL and were quantified after culturing for 48 hr in phenol-red-free RPMI-1640 (Sigma-Aldrich) supplemented with 10% dextran-coated charcoal-treated FBS for estrogen free experiments. Next, estradiol (E2; Wako Pure Chemical Industries, Osaka, Japan) dissolved in DMSO was added to this culture media.

Immunoblot analysis

Cells were washed with phosphate buffered saline (PBS; Sigma-Aldrich) and proteins were extracted using Mammalian Protein Extraction Reagent (Pierce Biotechnology, Rockford, IL, USA) supplemented with Halt Protease Inhibitor Cocktail (Pierce Biotechnology). The total protein concentration was measured using the Protein Assay Rapid Kit (Wako Pure Chemical Industries). Proteins were separated by SDS-PAGE (10% acrylamide gel) and transferred to a Hybond-P PVDF membrane (GE Healthcare, Chalfont St Giles, UK). Membranes were then blocked with 4% skim milk (Wako) for 1 hr at room temperature, incubated with a primary antibody overnight at 4°C, washed with tris buffered saline, and incubated with a secondary antibody for 1 hr at room temperature. Antibody-protein complexes on the blots were detected using ECL Plus western blotting detection reagents (GE Healthcare), and the chemiluminescence signal was visualized using a ChemiDoc XRS+ System (BIO RAD, Tokyo, Japan). We examined the expression level of ER α , ER β 1, ER β 2, ER β 5, and ER β proteins in MCF-7, T-47D, and MDA-MB-231 cells. By immunoblotting analysis, a higher level of ER α protein was detected in MCF-7 cells compared to that in T-47D cells (Fig. 1A). In MDA-MB-231 cells, ERa was not detected (Fig. 1A). The immunoreactivity of the ER β antibody (68-4) that recognizes a common epitope of ER β isotypes was higher in T-47D cells compared to that in MCF-7 and MDA-MB-231 cells (Fig. 1B). Low levels of ERB1 and ER_{β5} were detected in MDA-MB-231 cells. Otherwise, ERβ2 levels were high in MDA-MB-231 cells (Fig. 1B).

Cell blocks

Cells were suspended in 500 μ L of PBS and then centrifuged at 1500 rpm for 3 min. Pellets obtained were fixed for 3 min in 10% neutral-buffered formalin, centrifuged at 1500 rpm for 3 min and then washed in PBS. Fibrinogen dissolved in PBS was added to the cells and then rapidly mixed with thrombin. The clot was gently removed from the tube, transferred to a tissue cassette, and moved into ethanol and xylene, and finally embedded in paraffin.



Fig. 1. Immunoblotting analysis of ER proteins in breast cancer cells. (A) Expression of ER α proteins. β -actin functions as a control. (B) Expression of ER β proteins. β -actin functions as a control.

Immunofluorescence

Paraffin sections (3-µm thick) were dewaxed with xylene and ethanol. Antigen retrieval was performed by heating the slides in an autoclave at 121°C for 5 min in citrate buffer, pH 6.0. Sections were washed with PBS and incubated for 30 min at room temperature with blocking solution. These sections were further incubated overnight in a moist chamber at 4°C with primary antibodies. The characteristics of the primary antibodies employed in this study were summarized in Table 1. The sections were subsequently incubated with fluorescence-labeled secondary antibodies (Alexa Fluor 488 anti-rabbit and Alexa Fluor 594 anti-mouse; Invitrogen) for 1 hr at room temperature. The reacted slides were then mounted with mounting medium with DAPI. The dimeric proteins were detected by super-resolution imaging using structured illumination microscopy (Nikon, Tokyo, Japan). The areas of the yellow, green, and red signals were analyzed by Lumina Vision (Mitani Corp, Japan). We then determined the "% of ER dimer (i.e., ER α/α and ER α/β)" according to the following equation: (Yellow area/Yellow + Green + Red areas) \times 100 (%).

In situ proximity ligation assay (PLA)

In this study, we employed *in situ* PLA to detect ER dimerization according to our previous study [10].

We used the Duolink *in situ* PLA kit from Olink Bioscience (Olink Bioscience, Uppsala, Sweden). The cells grown on cover slides were fixed in 4% paraformaldehyde and permeabilized using 0.1% Triton-X-100. Cells were then incubated with blocking solution and incubated overnight with primary antibodies at 4°C. The characteristics of the primary antibodies used in this study were summarized in Table 1. The cells were subsequently incubated with PLA PLUS and MINUS probes for mouse and rabbit and incubated with ligation-ligase solution, and subsequently with amplification-polymerase solution according to the manufacturer's instructions. The slides were mounted with mounting medium with DAPI. The number of ER dimers in breast carcinoma cell lines was quantified as the area of the dots in the nuclei using image analysis (Lumina Vision, Mitani Corp, Japan).

III. Results

Imaging of ERa/a homodimers in breast carcinoma cells using N-SIM

 $ER\alpha/\alpha$ homodimer was identified as double-stained ERα proteins using two different anti-ERα antibodies, such as 6F11 (red) and SP-1 (green), and analyzed by using super-resolution microscopy (i.e., N-SIM). Both green and red fluorescent signals were detected in the nuclei of MCF-7 and T-47D cells (Fig. 2A, B). The ERa/a homodimer was detected as yellow signal spots, resulting from overlapping green and red fluorescent signals (Fig. 2A, B). Results of our study revealed that 23.0% and 13.4% of ER α/α homodimers were detected in MCF-7 and T-47D cells, respectively (Fig. 2C). ER α/α homodimers were also detected in both MCF-7 and T-47D cells using the PLA method as well as N-SIM (Fig. 2D). The PLA score in MCF-7 cells tended to be higher than that in T-47D cells (Fig. 2F). The ratio of ER α/α homodimers in MCF-7 and T-47D cells were significantly higher than that in MDA-MB-231 cells using both N-SIM (p = 0.0003, p = 0.0120) and PLA (p = 0.0012, p = 0.0236) (Fig. 2C, F). In ER α negative MDA-MB-231 cells, the ER signal could not be detected by either analysis (Fig. 2B–E).

Detection of ERa/ β heterodimers in MCF-7 and T-47D cell lines

Using N-SIM, ER α/β isotype heterodimers were detected in ER α -positive MCF-7 and T-47D cells (Fig. 3A).

Table 1. Antiboules employed in this study					
Antigen	Clone	Species	Description	Reference	Application
ERα	6F11	Mouse	Monoclonal	Leica (HE, GER)	IF, PLA
	SP-1	Rabbit	Monoclonal	Abcam (MA, USA)	IF, PLA
	polyclonal	Rabbit	Polyclonal	Santa Cruz	WB
ERβ	68-4	Rabbit	Monoclonal	Merck Millipore	WB
ERβ1	PPG5/10	Mouse	Monoclonal	AbDSerotec (OXF, UK)	IF, PLA
	7DN	Rabbit	Polyclonal	Invitrogen (CA, USA)	WB
ERβ2	57/3	Mouse	Monoclonal	AbDSerotec	IF, PLA, WB
ERβ5	5/25	Mouse	Monoclonal	AbDSerotec	IF, PLA, WB

Table 1. Antibodies employed in this study

PLA, ploximityligation assay; IF, immunofluorescence; WB, western blotting.



Fig. 2. Expression of ERα/α homodimers in breast carcinoma cells. (**A**) Detection of ERα homodimers using N-SIM in MCF-7 cells. Cells were doublestained for anti-ERα antibody clones 6F11 (Alexa Fluor 594: red) and SP-1 (Alexa Fluor 488: green). Homodimers are represented by the yellow areas, and nuclei are stained blue (DAPI). The areas of the yellow, green, and red signals respectively were analyzed as the yellow-green area by Lumina Vision (lower figures). Bar = 5 µm. (**B**) Detection of ERα homodimers using N-SIM in T-47D and MDA-MB-231 cells. Cells were doubleimmunostained for anti-ERα antibody clones 6F11 (Alexa Fluor 594: red) and SP-1 (Alexa Fluor 488: green). Bar = 5 µm. (**C**) The ratios of the ERα/α homodimer were quantified as the yellow area in the nuclei using Lumina Vision. *p = 0.0003 vs. MDA-MB-231, *p = 0.0120 vs. MDA-MB-231 for the ERα/α homodimer. (**D**) Detection of ERα homodimers was quantified as the area of the dots in the nuclei using Lumina Vision. *p < 0.0001 vs. MDA-MB-231, *p = 0.0012 vs. T-47D, *p = 0.0236 vs. MDA-MB-231 for the ERα/α homodimer.

The percentages of each dimer determined by N-SIM analysis were as follows (Fig. 3B). MCF-7: ER $\alpha/\beta1$, 6.6%; ER $\alpha/\beta2$, 18.1%; ER $\alpha/\beta5$, 22.4%. T-47D: ER $\alpha/\beta1$, 5.1%; ER $\alpha/\beta2$, 10.1%; ER $\alpha/\beta5$, 10.1%. The ratio of ER $\alpha/\beta1$ heterodimers in MCF-7 cells were significantly higher than that in MDA-MB-231 cells (p = 0.0206) and that of ER $\alpha/\beta2$ heterodimers in MCF-7 and T-47D cells were significantly higher than that in MDA-MB-231 cells (p < 0.0001, p = 0.0004). In addition, the ratio of ER $\alpha/\beta2$ heterodimers in MCF-7 cells was significantly higher than

that in T-47D cells (p = 0.0020). The ratio of ER α/β 5 heterodimers in MCF-7 and T-47D cells was also significantly higher compared to that in MDA-MB-231 (p < 0.0001, p = 0.0109). In addition, the ratio of ER α/β 5 heterodimers in MCF-7 cells was significantly higher than that in T-47D cells (p = 0.0034).

ER α // β heterodimers were detected in both MCF-7 and T-47D cells using the PLA method as well as N-SIM (Fig. 3C, D). The area of each dimer determined by PLA were as follows (Fig. 3D). MCF-7: ER α / β 1, 3.3 µm²;



Fig. 3. Expression of ERα/β heterodimers in breast carcinoma cells. (A) Detection of ERα/β heterodimers using N-SIM in breast carcinoma cells. Cells were double-stained for anti-ERα antibody (Alexa Fluor 488: green) and anti-ERβ antibody (Alexa Fluor 488: green). Heterodimers were represented by the yellow areas, and nuclei stained blue (DAPI). Bar = 5 µm. (B) The ratios of ERα/β heterodimers were quantified as the yellow areas in the nuclei using Lumina Vision. *p = 0.0206 vs. MDA-MB-231 for the ERα/β1 heterodimer; *p < 0.0001 vs. MDA-MB-231, *p = 0.0020 vs. T-47D, *p = 0.0004 vs. MDA-MB-231 for the ERα/β2 heterodimer; *p < 0.0001 vs. MDA-MB-231, *p = 0.0109 vs. MDA-MB-231 for the ERα/β5 heterodimer. (C) Detection of ERα/β heterodimers using PLA. Heterodimers were represented by the red dots (Texas red), and nuclei labeled blue (DAPI). Bar = 50 µm. (D) The number of ERα/β heterodimers was quantified as the area of the dots in the nuclei using Lumina Vision. *p = 0.0039 vs. MDA-MB-231 for the ERα/β1 heterodimer; *p = 0.0051 vs. MDA-MB-231, *p = 0.0337 vs. T-47D for the ERα/β5 heterodimer.</p>



Fig. 4. Expression of the ERα/α homodimer induced by estradiol in MCF-7 cells and time comparison. (**A**) Detection of ERα homodimers using N-SIM. Cells were double-stained for anti-ERα antibody clones 6F11 (Alexa Fluor 594: red) and SP-1 (Alexa Fluor 488: green). Homodimers were represented by the yellow areas, and the nuclei labeled blue (DAPI). Bar = 5 µm. (**B**) The ratios of ERα/α homodimers were quantified as the yellow areas in the nuclei using Lumina Vision. *p = 0.0121 vs. control for the ERα/α homodimer. (**C**) Immunoblotting analysis of ERα proteins in breast carcinoma cells. β-actin functions as a control. (**D**) Detection of ERα/α homodimers was quantified as the area of the dots in the nuclei using Lumina Vision. *p = 0.0252 vs. control for the ERα/α homodimers was quantified as the area of the ERα/α homodimer.

ER $\alpha/\beta2$, 4.2 µm²; ER $\alpha/\beta5$, 4.3 µm². T-47D: ER $\alpha/\beta1$, 2.5 µm²; ER $\alpha/\beta2$, 3.5 µm²; ER $\alpha/\beta5$, 1.1 µm². ER $\alpha/\beta1$ heterodimers in MCF-7 and T-47D cells were significantly higher than that in MDA-MB-231 cells (p = 0.0011, p = 0.0039). The number of ER $\alpha/\beta2$ heterodimers in MCF-7 and T-47D cells was higher than that in MDA-MB-231 cells. However, these differences did not reach statistical significance. The number of ER $\alpha/\beta5$ heterodimers in MCF-7 and T-47D cells was significantly higher than that in MDA-MB-231 cells (p = 0.0051, p = 0.0337). In ER α -negative MDA-MB-231 cells, ER α/β heterodimers were not detected (Fig. 3A–D).

Formation of estradiol-induced ERa/a homodimers in MCF-7 cells evaluated by N-SIM

E2-induced ER α/α homodimers were detected in MCF-7 cells (Fig. 4A, B). We added 10 nM of E2 to breast carcinoma cells and evaluated how this influenced the expression of ER α/α homodimers after 15, 45, and 90 min of incubation by employing N-SIM (Fig. 4A, B). Intranuclear ER α/α homodimers were detected in control and E2-treated MCF7 cells. The ratio of ER α/α homodimers in MCF-7 cells increased after 15 and 45 min of E2 treatment (Fig. 4A, B). In addition, the ratio of ER α/α homodimers after 15 min of E2 treatment was significantly higher than



Fig. 5. Expression of ER dimers induced by estradiol in MCF-7 cells and time comparison. (**A**) Detection of ERα/β1 heterodimers using PLA. Homodimers were represented by the red dots (Texas red), and nuclei stained blue (DAPI). Bar = 50 μm. (**B**) The number of ERα/β1 heterodimers was quantified as the area of the dots in the nuclei using Lumina Vision. *p < 0.0001 vs. control, *p = 0.0030 vs. control, *p = 0.0030 vs. control for the ERα/β1 heterodimer. (**C**) Detection of ERα/β2 heterodimers using PLA. Homodimers were represented by the red dots (Texas red), and nuclei labeled blue (DAPI). Bar = 50 μm. (**D**) The number of ERα/β2 heterodimers was quantified as the area of the dots in the nuclei using Lumina Vision. (**E**) Detection of ERα/β5 heterodimers using PLA. Homodimers was quantified as the area of the dots in the nuclei using Lumina Vision. (**E**) Detection of ERα/β5 heterodimers using PLA. Homodimers were represented by the red dots (Texas red), and nuclei labeled blue (DAPI). Bar = 50 μm. (**B**) The number of ERα/β2 heterodimers was quantified as the area of the dots in the nuclei using Lumina Vision. (**F**) Detection of ERα/β5 heterodimers using PLA. Homodimers were represented by the red dots (Texas red), and nuclei labeled blue (DAPI). Bar = 50 μm. (**F**) The number of ERα/β5 heterodimers was quantified as the area of the dots in the nuclei using Lumina Vision. *p = 0.0040 vs. control, *p = 0.0040 vs. control for the ERα/β1 heterodimer.

that obtained in the control (p = 0.0121). Similar results were also obtained by PLA analysis (Fig. 4D, E). The PLA signal increased after 15 and 45 min of E2 treatment (p = 0.0252) (Fig. 4D, E). Results of immunoblotting analysis revealed that the ER α protein level decreased after 15, 45, and 90 min of E2 treatment (Fig. 4C).

Visualization of the effects of E2 treatment upon ERa/ β heterodimer formation in MCF-7 cells using PLA

We also evaluated the formation of ER α/β heterodimers using PLA. We added 10 nM of E2 to breast carcinoma cells and used PLA to examine the status of ER α/β isotype heterodimers after 15, 45, and 90 min of treatment (Fig. 5).

The number of E2-induced ER α/β 1 heterodimers increased after 15 min of E2 treatment (Fig. 5A, B). There were no changes in the number of either ER α/β 2 or ER α/β 5 heterodimers in MCF-7 cells treated with E2 (Fig. 5C–F). None of the ER α/β subtype heterodimers could be detected after 45 or 90 min of treatment (Fig. 5A–F). The number of ER α/β 1 heterodimers after 15 min of E2 treatment was significantly higher than that in the control (p < 0.0001). In addition, the number of ER $\alpha/\beta 1$ and ER $\alpha/\beta 5$ heterodimers was significantly less after 45 and 90 min of treatment compared to that in the control (ER $\alpha/\beta 1$; 45 min, p = 0.0030; 90 min, p = 0.0030) (ER $\alpha/\beta 5$; 45 min, p = 0.0040; 90 min, p = 0.0040) (Fig. 5B, D, F).

IV. Discussion

This is the first study to detect ER dimeric proteins using N-SIM. Both ER homo- and heterodimers were detected in ERs double-positive MCF-7 and T-47D cells. In ER α -/ER β + MDA-MB-231 cells, only ER α/β heterodimers were detected and immunoblotting analysis also revealed that a higher level of ER α protein was detected in MCF-7 cells compared to that in T-47D cells. In both N-SIM analysis and PLA, the level of ER α/α homodimer in MCF-7 cells was higher than that in T-47D cells. In addition, no ER α/α homodimer signal was detected by either N-SIM analysis or PLA in ER α -negative MDA-MB-231 cells. In addition, the E2-treatment assay in MCF-7 cells demonstrated that the peak PLA signal for the ER α/α homodimer occurred after 45 min of treatment, as shown in both our previous report and this study [10]. The ER α/α homodimer patterns obtained by N-SIM analysis was similar to that in E2treated MCF-7 cells by PLA. Results of our present study also indicate that the detection sensitivity of the ER dimer by PLA was equal to that detected by FRET [10]. Therefore, these findings did indicate that N-SIM could be a useful analytical tool, comparable to PLA, in detecting ER α/α homodimer in breast carcinoma cells.

Results of immunoblotting analysis revealed that the expression levels of ER β 1 and ER β 2 in MCF-7 cells were very similar to those in T-47D cells but the ERa protein levels in MCF-7 cells were markedly higher than those in T-47D cells. However in dimeric signaling analysis using both N-SIM and PLA, there were no differences in ER α / ER β 1 and ER α /ER β 2 heterodimer formation between MCF-7 and T-47D cells. Therefore, the formation of ER α / ER β 1 or ER α /ER β 2 heterodimers could be determined more by the expression levels of ER β 1 or ER β 2 than that of ERa. The ERa/ β 1 heterodimer signal revealed by PLA increased after 15 min of E2 treatment in MCF-7 cells. However, E2 treatment had no effects upon the levels of $ER\alpha/\beta^2$ or $ER\alpha/\beta^5$ heterodimers. A possible reason for this discrepancy might be that ER_{β2} formed heterodimers with ER α or ER β 1 without ligand binding [17, 18]. Ligandindependent dimer formation of ERB5 has not been reported but ERB5 was reported to bind to E2 with moderate affinity compared to that of ER β 1 [15]. In our present study, PLA signals from ERa/ERB isoform heterodimers disappeared after 45 or 90 min of E2 treatment. One possible reason for this is that E2 treatment decreased the protein levels of ERa. However, in both N-SIM analysis and PLA, the formation of the ER α /ER α homodimers remained at a level equal to or more than that of the intact MCF-7 level because ERB1 is well known to bind to E2 with lower affinity than ERa [11]. Therefore, these findings did suggest that the ER α/α homodimer could be more predominantly induced by E2 compared to that of the ER α/β heterodimer in breast carcinoma cell lines.

As for ER α/β heterodimer, all ER β isoforms inhibited the transcriptional activities of $ER\alpha$ for an estrogen response element-containing promoter but it is also true that the various ERB isoforms had different degrees of these effects [19]. Nuclear ER β 2 and ER β 5, but not ER β 1, were reported to be significantly correlated with good overall survival in breast cancer patients [13, 24]. However, in both ER α -negative and HER2-positive breast cancer, ER β 2 status was indeed associated with adverse clinical outcome of the patients [30]. In addition, ER β 2 preferentially formed a heterodimer with ER α rather than with ER β 1 in ER-transfected COS-7 cells [17]. These findings all indicated that the effects of ER on breast cancer could be different depending on the dimer patterns of ER α and ER β . However, further examination, such as an analysis of the dimer patterns of ER α and ER β (including its variant forms) in breast cancer patients are required for clarification of the significance of ER dimmers in biological behavior of ER positive breast cancer patients.

In this study, we examined the dimeric protein pattern of ER by both N-SIM and PLA in breast carcinoma cells. Both N-SIM analysis and PLA are generally considered powerful laboratory methods for the detection of nuclear receptor dimers such as ER α and ER β in breast carcinoma cells. Results of our previously reported study revealed that PLA was suitable for a large-scale study using tissue samples as in immunochemical analysis [10]. However, N-SIM analysis has been reported to have high sensitivity, specificity, and resolution compared to light microscopy [8, 9]. Raloxifen were reported to translocate ERa into the nucleoli in breast cancer cell line [6]. Therefore, using N-SIM technique, it is possible to detect the patterns of intranuclear distribution of ER dimers stimulated by its agonists. In addition, N-SIM analysis could visualize proteinprotein interactions of more than three proteins using multiple staining, whereas PLA could only visualize protein-protein interactions of two proteins. Therefore, in addition to detecting the ER dimers, we could distinctively identify the complex patterns of ER dimer with cofactor proteins using N-SIM. Furthermore, N-SIM analysis also could indeed evaluate fine structures and function in single living cell [5, 12, 31]. This could allow evaluation of dynamic changes in ER α and ER β dimer formation using N-SIM analysis.

V. Disclosure/Duality of Interest

All authors declare that they have no competing interests.

VI. Acknowledgments

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