RESEARCH ARTICLE

Direct Interaction of CD40 on Tumor Cells with CD40L on T Cells Increases the Proliferation of Tumor Cells by Enhancing TGF-β Production and Th17 Differentiation

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Abstract

It has recently been reported that the CD40-CD40 ligand (CD40L) interaction is important in Th17 development. In addition, transforming growth factor—beta (TGF-β) promotes tumorigenesis as an immunosuppressive cytokine and is crucial in the development of Th17 cells. This study investigated the role of CD40 in breast cancer cells and its role in immunosuppressive function and tumor progression. CD40 was highly expressed in the breast cancer cell line MDA-MB231, and its stimulation with CD40 antibodies caused the up-regulation of TGF-β. Direct CD40-CD40L interaction between MDA-MB231 cells and activated T cells also increased TGF-β production and induced the production of IL-17, which accelerated the proliferation of MDA-MB231 cells through the activation of STAT3. Taken together, the direct CD40-CD40L interaction of breast tumor cells and activated T cells increases TGF-β production and the differentiation of Th17 cells, which promotes the proliferation of breast cancer cells.

Introduction

CD40 is a 50 kDa type I transmembrane protein that belongs to the tumor necrosis factor (TNF)-receptor superfamily [1]. It is physiologically expressed in various cell types including dendritic cells, B cells and monocytes/macrophages [2]. Especially, CD40 in B cells has a critical role in the development and proliferation of B cells by enhancing interleukin (IL)-2, IL-4, and IL-5 and the production of other chemokines. [3–6]. CD40 also induces the differentiation of B cells into antibody-secreting plasma cells [7, 8]. The ligand for CD40 is CD154 (CD40L) which is predominantly expressed in activated T cells and has a crucial role in the regulation of B cell proliferation [9, 10]. Recently, it was reported that CD40-CD40L cross-talk is important in Th17 development [11]. In addition to B cells, CD40 is also expressed in several kinds of
tumor cells, such as breast cancer, ovarian cancer, colon cancer, and melanoma [10, 12]; however, its specific roles are still largely unknown.

Transforming growth factor-beta (TGF-β) is a pleiotropic cytokine that controls multiple cellular responses including the induction of cell growth inhibition, differentiation, cellular senescence, wound healing and apoptosis [13]. TGF-β cannot bind to its receptors in its latent form which is regulated by latency-associated protein and latent-TGF-β binding protein [14]. In vitro, these factors can be inactivated by extreme pH or heat, or by several proteases [15]. However, in vivo mechanisms for the activation of TGF-β are less clear; however, several models have been proposed including proteolysis by transglutaminase, conformational change of latency-associated proteins through physical interaction with thrombospondin, and modulation by αvβ6 integrins in epithelial cells [16–18]. TGF-β has dual roles in the progression and metastasis of cancer [19]. In human cancers, TGF-β promotes tumorigenesis through both decreased TGF-β signaling during early tumorigenesis and increased TGF-β signaling in advanced, progressive disease [13, 20]. TGF-β is a potent suppressor of proliferation in normal epithelial cells, notably breast; however, it converts to a promoter during cancer development [21]. In particular, TGF-β signaling has important roles during breast cancer progression and metastasis in various mouse models [19, 22, 23], and the level of TGF-β was increased in cancer patients [24, 25]. TGF-β has a role in the differentiation of CD4⁺CD25⁺ regulatory T cells which potently suppress both in vitro and in vivo effector T cell function and maintain Foxp3 expression [26–28], and it is also essential in the induction of Th17 cells [29, 30].

This study investigated the role of CD40 in the production of TGF-β in breast cancer cells, and the results show that the production of TGF-β induced by the CD40-CD40L interaction, results in the enhanced immunosuppressive function of breast cancer cells and could thereby contribute to tumor progression.

**Materials and Methods**

**Cells**

The human breast cancer cell lines, MDA-MB231 and HS-578T were purchased from American Type Culture Collection (Manassas, VA, USA). Cells were maintained in continuous log phase of growth at 37°C in a humidified atmosphere containing 5% CO₂ with RPMI 1640 medium supplemented with 2 mM L-glutamine, 100 units/ml penicillin, 100 μg/ml streptomycin (Welgene, Daegu, Korea), and 10% heat-inactivated fetal bovine serum (FBS, Hyclone, Utah, USA).

**Isolation of T cells from human peripheral blood**

Heparinized peripheral blood was collected from healthy volunteers under protocol approved by an Institutional Review Board (IRB) of Seoul National University Hospital (SNUH) (IRB# 0902-022-271). Human T cells were enriched from peripheral blood by using RosetteSep (Stem Cell Technologies, Vancouver, Canada). Briefly, 40 ml of blood obtained from normal healthy volunteer was mixed with 2 ml of RosetteSep cocktail consisted of mouse IgG1 antibodies to human lineage antigens (CD16, CD19, CD36 and CD56) and incubated at room temperature for 30 min with gentle mixing. After dilution with an equal volume of phosphate buffered saline (PBS), T cells were isolated by density gradient centrifugation using pre-warmed Ficoll-Paque (GE healthcare lifesciences, Uppsala, Sweden) at 600 g for 20 min. The interface was harvested, centrifuged at 2,000 rpm for 10 min, and then pellet was suspended to RPMI 1640 medium contained 10% FBS. Otherwise, peripheral blood was mixed with an equal volume of PBS, and loaded onto pre-warmed Ficoll-Paque. After centrifuging at 600 g for 20 min, a buffy coat containing PBMC was harvested and washed with PBS twice. The red blood
cells (RBCs) were lysed with RBC lysis buffer (Sigma, St. Louis, MO, USA) in a 37°C water bath for 5 min with shaking, and the mononuclear cells were washed and counted. Human T cells among the isolated mononuclear cells were separated by using the Pan T Cell Isolation Kit (Miltenyi Biotec, Germany) with autoMACS Pro Separator (Miltenyi Biotec, Germany) according to the manufacturer’s instruction. In brief, determined cells were suspended with buffer and mixed with biotin-antibody cocktail (10 μl/10⁷ cells) for 5 min at 4°C. After washing, cells were mixed with anti-biotin microbeads (20 μl/10⁷ cells) for 10 min at 4°C. Washed cells were applied to the autoMACS separator, and negatively selected T cells were counted. We confirmed more than 95% of purified T cells were CD3+ cells by flow cytometry analysis, after staining with PE-conjugated anti-CD3 antibody (eBioscience, San Diego, CA, USA).

**Activation of T cells**

CD4 expression on activated T cells was reduced by stimulation with phorbol 12-myristate 13-acetate (PMA)/ionomycin reduces, but not by phytohemagglutinin (PHA) [31, 32]. However, PHA alone cannot effectively induces CD40L, but in combination with PMA showed CD40L expression comparable to those seen with a combination of CD3 mAb and PMA [33]. Purified T cells (2×10⁶/ml) were activated by of 5 μg/ml of PHA (Life Technologies, Grand Island, NY) for 69 hrs, and then activated with 10 ng/ml of PMA (Sigma, St.Louis, MO, USA) and 1 μg/ml of ionomycin for another 3 hrs. Activated T cells were analyzed by flow cytometry after staining with FITC-conjugated anti-CD69 or CD25 antibodies (BD Pharmingen, San Diego, CA, USA).

**Flow cytometry analysis**

MDA-MB231 cells were stained with PE-conjugated anti-human CD40 antibody (BD Pharmingen, San Diego, CA, USA), and activated T cells were stained with FITC-conjugated anti-CD25 antibody or PE-conjugated anti-CD40L antibody (BD Pharmingen, San Diego, CA, USA) for 30 min on ice. After washing with buffer containing 0.5% bovine serum albumin (BSA) in PBS, stained cells were analyzed by FACS Calibur (BD Bioscience, San Jose, CA, USA). To determine the Th17 differentiation by the ligation of CD40L on activated T cells with CD40 expressing MDA-MB231 cells or anti-CD40L agonistic antibody (2 μg/ml), intracellular IL-17 staining with Alexa Fluor 647-conjugated anti-IL-17 antibody and surface CD4 staining with FITC-conjugated anti-CD4 antibody was performed by BD Cytofix/Cytoperm Fixation/Permeabilization Solution Kit (BD Pharmingen, San Diego, CA, USA), and RORγt staining with PE-conjugated anti-RORγt antibody was performed by using a transcription factor staining buffer set (eBioscience, San Diego, CA, USA) according to the manufacturer’s instructions. Flow cytometry was performed with FACS Calibur, and analyzed with FlowJo software (Ashland, OR, USA).

**[^3H]-Thymidine incorporation assay**

MDA-MB231 cells growing at logarithmic growth phase were trypsinized and resuspended in growth media at 2.5×10⁴ cells/ml. A hundred microliter of the cell suspension was dispensed onto a 96-well plate. Quadruplicate samples were incubated in the presence or absence of agonistic anti-CD40 antibody (2, 4, 8, and 16 μg/ml) for 24 hrs. After addition of 1 μCi/mL of[^3H]-thymidin to each well, 96-well plate was further incubated for 18 hrs at 37°C in a 5% CO₂ atmosphere. Then, cells were harvested onto glass fiber filters using a cell harvester (Inotech Biosystems International, Dietikon, Switzerland). Glass fiber filters were dried and sealed into polythene bags with scintillation fluid (Betaplate Scint; PerkinElmer, Boston, MA, USA).
The incorporation of [3H]-thymidine counted on a MicroBeta Trilux1450 (PerkinElmer, Boston, MA, USA).

Co-culture of MDA-MD231 cells and T cells
To investigate the requirement of interaction between CD40 and CD40L for IL-1β, IL-6, IL-21 and TGF-β production from MDA-MB231 cells and IL-17 differentiation from activated T cells, MDA-MB231 cells (1.5 x 10^5/well) were incubated with activated T cells (AT) and resting T cells (RT) in the same well on 6-well plate for 24 hrs at the ratio of 1:5, 1:10 and 1:20. And it was confirmed by the interference of interaction between CD40 and CD40L with anti-CD40 neutralizing antibody (2 μg/ml). It was also confirmed by the co-culture of MDA-MB231 cells and activated T cells by using of a Transwell culture system (Corning Life Sciences, Tewksbury, MA, USA). Briefly, MDA-MB231 cells (1.5 x 10^5/well) were cultured in the lower chamber (25 mm) of the Transwell system and activated T cells (7.5 x 10^5/well) were loaded on the upper chamber with a polycarbonate membrane bottom (0.4 μm pore size). Culture supernatants were harvested, centrifuged, and frozen in aliquots for further experiments.

Measurement of cytokine levels by ELISA
MDA-MB231 cells were stimulated with anti-CD40 agonistic antibody (2 μg/ml) or isotype (2 μg/ml). And then culture supernatants were harvested and stored at -70°C until use. The concentration of TGF-β (Biosource International, Inc. Camarillo, CA, USA), IL-1β, IL-6 (R&D system, Minneapolis, MN, USA) and IL-21 (eBioscience, San Diego, CA, USA) was measured with commercially available Enzyme Linked Immunosorbent assay (ELISA) kits, according to the manufacturers’ instruction. The final concentration of cytokines was normalized with cell numbers.

Reverse Transcriptase-Polymerase Chain Reaction (RT-PCR) and Real-time PCR
MDA-MB231 cells were seeded on a 24-well plate (1.5 x 10^5/well) and incubated overnight. Then, anti-CD40 agonistic antibody (2 μg/ml) and isotype antibody (Sigma, St. Louis, MO, USA) were treated to MDA-MB231 cells for 3, 6, 9 and 12 hrs. Total RNA was extracted from CD40-stimulated MDA-MB-231 cells by using Trizol (Invitrogen life technologies, Carlsbad, CA, USA), according to the manufacturers’ instruction, and RNAs were quantified with NanoDrop (Thermo scientific, Wilmington, DE, USA). Total RNA (1 μg) was transcribed to cDNA by avian myeloblastosis virus (AMV) reverse transcriptase (Promega, Madison, WI, USA). For conventional RT-PCR, cDNAs were amplified with flowing primers. TGF-β: Forward, 5’-GGG ACT ATC CAC GTG CAA GA-3’ and Reverse, 5’-CCT CCT TGG CGT AGT AGT CG-3’; β-actin: Forward, 5’-TCC TTA ATG TCA CGC ACG A-3’ and Reverse, 5’-GTG GGG CGC CCA GCC ACC A-3’. PCR products were separated on 1.5% agarose gel, stained with ethidium bromide and visualized under UV light. Quantitative Real-time RT-PCR analysis was performed by using SYBR Green Master mix (Fermentas, Hudson, NH, USA), and primers were used as followed. TGF-β: Forward, 5’- GGC GAT ACC TCA GCA ACC G-3’ and Reverse, 5’- CTA AGG CGA AAG CCC TCA AT-3’, GAPDH: GCC ACC CAG AAG ACT GTG GA-3’ and Reverse, 5’- CAG TGA GCT TCC GTG TCA GC-3’. Amplification conditions were: 95°C for 10 min, followed by 45 cycles of 95°C for 30s, 57°C for 30s and 72°C for 15s, and followed by an extension at 72°C for 5 min. For assessing the mRNA expression level, the Ct value for TGF-β was subtracted from the Ct value of GAPDH to yield a ΔCt value. The average ΔCt was calculated for the control group and subtracted from the ΔCt of all other samples (including the control
group). This resulted in a ΔCt value for all samples used to calculate the fold-induction of mRNA expression of target gene using the formula 2-ΔCt.

Western blot analysis
MDA-MB231 cells (1.5 x 10^5/well) were incubated with recombinant IL-17 or co-culture supernatants of MDA-MB231 cells and activated T cells for 15, 30, 60 min on 6-well plate. After washing with PBS, cells were homogenized with lysis buffer and total proteins were quantified by bicinchoninic acid (BCA) assay. Equal amounts of protein were resolved over 10% polyacrylamide gel and transferred to a nitrocellulose membrane. Followed blocking with 5% non-fat milk, membranes were incubated with anti-phospho-STAT3 antibody (1:200), anti-STAT3 antibody (1:200) (Cell signaling, Danvers, MA, USA) or β-actin (1:4000, Sigma, St. Louis, MO, USA) at 4°C for overnight. After incubating with horseradish peroxidase-conjugated anti-rabbit or anti-mouse IgG (1:1000, Cell signaling, Danvers, MA, USA), the immunoreactive proteins were visualized with the ECL detection system (Amersham Biosciences Corp., Piscataway, NJ, USA).

CD40 siRNA transfection
CD40 siRNA and control siRNA were purchased from Santa Cruz Biotechnology (Palo alto, CA, USA). Cells in exponential phase of growth were plated in 6-well plates at 5 x 10^5 cells/well, grown for 24 hrs and then transfected with 20 nM of siRNA using oligofectamine and OPTI MEM I-reduced serum medium (Invitrogen Life Technologies, Carlsbad, CA, USA), according to the manufacturer’s protocol. The concentrations of siRNA were chosen based on dose-response studies. Transfection efficiency was examined by flow cytometry after staining with PE-conjugated anti-CD40 antibody at 72 hrs after transfection. Control cells were transfected with control siRNA with oligofectamine and serum-reduced medium (mock).

Statistics
Data were expressed as mean ± S.D. of each group in independent experiments. For comparison of three or more groups, data were analyzed by t-test or one-way analysis of variance (ANOVA) followed by Newman-Keuls multiple comparison tests. A value of P < 0.05 was considered statistically significant. Statistical tests were carried out using GraphPad InStat (GraphPad Software, San Diego, CA, USA).

Results
CD40 stimulation up-regulates TGF-β production in the malignant breast cancer cell line, MDA-MB231
First, the expression of CD40 was examined in the human breast cancer cell lines MDA-MB231 and Hs578T. MDA-MB231 cell is known to be more malignant than Hs578T cell. As shown in Fig 1A, CD40 is highly expressed on the surface of the MDA-MB231 cells; however, Hs578T cells hardly express CD40 on the surface. Because CD40 stimulation is known as a critical factor in the proliferation and differentiation of B cells [33], whether the proliferation of MDA-MB231 cells is also increased by stimulation of CD40 with anti-CD40 antibody was also investigated. The stimulation of CD40 in MDA-MB231 did not affect the proliferation of the MDA-MB231 cells (Fig 1B). Thus, we focused on the effect of CD40 stimulation in the production of immune modulatory molecules from tumor cells. TGF-β and IL-10 are the most well-known immunosuppressive molecules produced by tumor cells [34]. The transcriptional expression of TGF-β was determined by RT-PCR, and TGF-β mRNA
Fig 1. TGF-β production is up-regulated by CD40 stimulation in a breast cancer cell line, MDA-MB231. (A) The breast cancer cell lines, MDA-MB231 and Hs578T were collected at continuous log phase of growth. The expression of CD40 was examined by staining with PE-conjugated anti-human CD40 antibody (1 μg/ml) as described in Materials and Methods. Result is the representative of three independent experiments. (B) MDA-MB231 cells (2.5 x 10^3/well) were stimulated with the various concentrations of anti-CD40 agonistic antibody (2, 4, 8, and 16 μg/ml) and the same amount of isotype antibody for 1 hr on 96-well plate, and then cells were cultured for 24 hrs. After the addition of 1 μCi/mL of [3H]-thymidine, cells were culture for another 18 hrs. And then, the proliferation of cells by CD40 stimulation was measured as described in Materials and Methods. Data represents mean ± S.D. The assay was performed in quadruplicate and result is the representative of three independent experiments. There was no statistical significance among groups. (C and D) The expression of TGF-β mRNA transcript in MDA-MB231 cells by CD40 stimulation was investigated with semi-quantitative RT-PCR (C) and quantitative real-time PCR (D). Cells (1.5 x 10^5/well) were stimulated 2 μg/ml of anti-CD40 agonistic antibody for 3, 6, 9 and 12 hrs on 6-well plate. Cells were harvested and total RNA was extracted with Trizol, according to manufacturer’s instruction. After quantification, 1μg of RNA was converted to cDNA by reverse transcriptase. And then RT-PCR and real-time PCR were performed with specific primers for TGF-β as described in Materials and Methods. β-actin or GAPDH was used as a loading control. ***p < 0.001. (E) CD40 on MDA-MB231 cells (1.5 x 10^5/well) were stimulated with anti-CD40 agonistic antibody (2 and 4 μg/ml) and soluble CD40L (3 μg/ml) for 1 hr, and then cells were cultured for 24 hrs on 6-well plate. The amount of TGF-β in the culture supernatant was measured by ELISA, according to manufacturer’s instruction. Data represents mean ± S.D. Result is the representative of three independent experiments and each experiment was performed in triplicate. **p < 0.01 vs. untreated control. (F) Cells (1.5 x 10^5/well) were seeded on 6-well plate and then transfected with 20 nM of CD40 siRNA and control siRNA in the mixture of serum free media and oligofectamine as described in Materials and Methods. After 72 hrs, the down-regulation of CD40 expression on MDA-MB231 cells by CD40 siRNA transfection was confirmed by flow cytometry analysis. And then, cells (1.5 x 10^5/well) were stimulated with anti-CD40 agonistic antibody (2 μg/ml) and soluble CD40L (3 μg/ml) for 1 hr, and then cells were cultured for 24 hrs on 6-well plate. And then, the amount of TGF-β in the culture supernatant was measured by ELISA. Data represents mean ± S.D. Result is the representative of three independent experiments and each experiment was performed in triplicate. ***p < 0.001 vs. untreated control.
transcripts were increased by CD40 stimulation with the anti-CD40 antibody (Fig 1C). This result was also confirmed by real-time PCR (Fig 1D). Nine hours after stimulating CD40 in MDA-MB231 cells, TGF-β mRNA transcripts were increased and then decreased by the twelfth hour. Next, the production of TGF-β from MDA-MB231 cells was measured after treatment with anti-CD40 antibody or soluble CD40 ligand (sCD40L) (Fig 1E). MDA-MB231 cells secreted more TGF-β in response to CD40 stimulation. To confirm the TGF-β production from MDA-MB231 cells by CD40 stimulation on the surface, TGF-β production was measured from CD40 siRNA transfected MDA-MB231 cells. A decrease in CD40 expression on the surface of MDA-MB231 cells by CD40 siRNA transfection was confirmed (S1 Fig). As a result, TGF-β production was decreased from the CD40 siRNA transfected MDA-MB231 cells by CD40 stimulation (Fig 1F). Therefore, it seems that CD40 in MDA-MB231 triggers TGF-β production when it is stimulated with anti-CD40 Ab or CD40 ligand (CD40L).

A direct CD40-CD40L interaction between MDA-MB231 cells and activated T cells increases TGF-β production

Next, the effect of activated T cells on TGF-β production was investigated because CD40L is known to be expressed in activated T cells [1]. The expression of CD40L and CD25 in the activated T cells was confirmed by flow cytometry (Fig 2A, insert). Then, MDA-MB231 cells were co-cultured with the activated T cells for 24 hrs, and the TGF-β level in the culture supernatant was examined by ELISA. TGF-β production was remarkably increased after co-culture with activated T cells but not with resting T cells (Fig 2A). However, the production of TGF-β did not increase when the direct contact between the MDA-MB231 cells and the activated T cells were inhibited by the inserts of the Trans-well system (Fig 2B). This result suggests that the production of TGF-β is mediated by direct contacts between MDA-MB231 cells and activated T cells, not by soluble factors. We also performed a neutralizing experiment to clarify that the increased TGF-β production is mediated by the interaction between the CD40 on the MDA-MB231 cells and the CD40L on the activated T cells. TGF-β production was inhibited when the activated T cells were pre-incubated with anti-CD40L neutralizing antibody prior to co-culture with the MDA-MB231 cells. In addition, the production of TGF-β was blocked when MDA-MB231 cells were pre-incubated with anti-CD40 neutralizing antibody (Fig 2C). Therefore, a direct interaction between MDA-MB231 cells and activated T cells induces the production of TGF-β.

A direct interaction between CD40 on MDA-MB231 cells and activated T cells induces Th17 differentiation

TGF-β is a critical factor in Th17 cell differentiation [35]. Because a direct interaction between CD40 on MDA-MB231 cells and CD40L on activated T cells increased TGF-β production, IL-17 production in activated T cells was examined. When activated T cells were directly co-cultured with MDA-MB231 cells, there was a large increase in IL-17-producing cells shown in Fig 3A. However, this increase was attenuated by indirect interaction with Transwell or by direct interaction between the activated T cells and MDA-MB231 cells treated with anti-CD40 neutralizing antibodies (Fig 3A). The cytokines IL-1β, IL-6 and IL-21 are also important in Th17 cell differentiation in humans [36]. Thus, the production of cytokines IL-1β, IL-6 and IL-21 secreted into the media was assessed with the direct or indirect co-culture of MDA-MB231 cells and activated T cells. There was an increased level of IL-1β, IL-6 and IL-21 in the direct co-culture compared to the indirect co-culture (Fig 3B–3D). This result suggests that Th17 differentiation is stimulated by cytokines induced by the direct interaction between activated T cells and MDA-MB231 cells. Additionally, the CD40-CD40L interaction has a key role in this process.
Retinoic acid receptor-related orphan nuclear receptor gamma t (RORγt), an orphan nuclear receptor, regulates IL-17 transcription and Th17 differentiation [37]. Therefore, we investigated whether CD40-CD40L interaction increases RORγt expression in activated T cells. As seen in Fig 3E, RORγt expression is increased when CD40L on activated T cells interacts with the anti-CD40 antibody, regardless of the presence of cytokines for Th17 differentiation such as TGF-β, IL-1β, IL-6, and IL-21. As shown in Fig 4, the increase in IL-17 producing cells through the direct interaction of the MDA-MB231 cells and the activated T cells was not completely but partially attenuated by treatment with one or more neutralizing antibodies against IL-1β, IL-6, IL-21 or TGF-β (Fig 4B and 4C). In addition, Th17 differentiation was not completely prevented.
CD40 Induces Th17 Differentiation and Tumor Proliferation

Fig 3. CD40 ligand stimulation on the activated T cells induces Th17 differentiation. (A) Human T cells (2.5x10^6/ml) purified from human PBMCs were activated as described in Materials and Methods. After increase of CD40L expression on activated T cells was confirmed by flow cytometry analysis, they were co-cultured with MDA-MB231 cells in the same well on 6-well plate at the ratio of 5:1 for 24 hrs. Th17 differentiation was confirmed by staining of intracellular IL-17 in CD4+ T cells with Alexa Fluor 647-conjugated anti-human IL-17A antibody (1 μg/ml) and FITC-conjugated anti-human CD4 antibody (1 μg/ml). To confirm the role of CD40 on Th17 differentiation, the interaction between CD40 and CD40L was interfered with the addition of anti-CD40 neutralizing antibody (2 μg/ml) for 1 hr. Result is the representative of three independent experiments. (B-D) Culture supernatant from the experiment described in Fig 3A were collected and the amount of IL-1β (B), IL-6 (C) and IL-21 (D) were measured by ELISA, according to manufacturer’s instruction. Data represents mean ± S.D. Result is the representative of three independent experiments and each experiment was performed in triplicate. *p < 0.05. (E) Human T cells (2.5x10^6/ml) purified from human PBMCs were activated as described in Materials and Methods. After increase of CD40L expression on activated T cells was confirmed by flow cytometry analysis, activated T cells were stimulated by anti-CD40L agonistic antibody (1 μg/ml) or isotype (1 μg/ml) for 24 and 36 hrs. The expression of ROR gamma t (RORγt), the orphan nuclear receptor for IL-17 transcription and Th17 differentiation was determined by intracellular flow cytometry analysis, after staining with PE-conjugated anti- RORγt antibody (1 μg/ml) and PE-conjugated Rat IgG (1 μg/ml) as described in Materials and Methods. Result is the representative of three independent experiments.

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by antibodies against TGF-β, IL-1β, IL-6, and IL-21 (Fig 4D). It means that Th17 differentiation could be only induced by CD40-CD40L interaction.

**IL-17 induces proliferation of MDA-MB231 cells by activation of STAT3**

Tumor growth is suppressed in IL-17 knockout mice and IL-17/IFN-γ double-knockout mice [38], and a mechanism for this knockout is that IL-17 induces the production of IL-6 by tumor cells, which in turn promotes tumor growth through a STAT3-dependent pathway. Because Th17 cells were increased by the CD40-CD40L interaction between the MDA-MB231 cells and activated T cells, the proliferation of MDA-MB231 cells was examined next. MDA-MB231 cell proliferation was highly increased when MDA-MB231 cells were incubated with the direct co-culture supernatant but not with the indirect co-culture supernatant from the MDA-MB231 cells and activated T cells. However, this increase in proliferation was attenuated by neutralizing IL-17 with anti-IL-17 antibody in the direct co-culture supernatant (Fig 5A). It was also
Fig 5. IL-17 production via direct interaction of CD40 and CD40L increases STAT3 activation and the proliferation of MDA-MD231 cells. (A) MDA-MB231 cells and activated T cells were directly co-cultured at the ratio of 1:5 for 24 hrs in the presence of anti-IL-17 neutralizing antibody or anti-CD40 neutralizing antibody (2 μg/ml/each) on 96-well plate, and then cells were cultured for 24 hrs. After the addition of 1 μCi/mL of [3H]-thymidine, cells were culture for another 18 hrs. And the proliferation of cells was measured as described in Materials and Methods. Data represents mean ± S.D. The assay was performed in quadruplicate and result is the representative of three independent experiments. (B) MDA-MB231 cells were cultured in the presence of recombinant IL-17 (rIL-17, 50 ng/ml) for 15, 30 and 60 min. In addition, cells were cultured with direct co-culture supernatant of MDA-MB231 cells and activated T cells in the presence or absence of anti-IL-17 neutralizing antibody (nAb). And then, the activation of STAT3 was examined by western blot as described in Materials and Methods. Lane 1: Control, Lane 2: rIL-17 (50 ng/ml), Lane 3: Direct co-culture supernatant of MDA-MB231 cells and activated T cells, Lane 4: Direct co-culture supernatant of MDA-MB231 cells and activated T cells with anti-IL-17 nAb, Lane 5: Direct co-culture supernatant of MDA-MB231 cells and activated T cells with anti-CD40 nAb. (C) CPM
attenuated by incubating the cells with supernatant obtained from the direct co-culture of activated T cells and MDA-MB231 cells pre-treated with anti-CD40 neutralizing antibodies. Because IL-17 can promote tumor growth through the STAT3 signaling pathway [38], we investigated whether STAT3 is involved in IL-17-mediated MDA-MB231 cell proliferation. When MDA-MB231 cells were treated with recombinant IL-17 (rIL-17), phosphorylated STAT3 increased up until 30 min after treatment and then decreased at 60 min to the same extent as the MDA-MB231 cells that were incubated with the direct co-culture supernatant of the MDA-MB231 cells and activated T cells (Fig 5B). However, the activation of STAT3 was inhibited when the MDA-MB231 cells were incubated with direct co-culture supernatant pre-treated with anti-IL-17 neutralizing antibody. Moreover, the increased proliferation of the MDA-MB231 cells by the direct co-culture supernatant was attenuated by treatment with AG490, a STAT3 inhibitor (Fig 5C). In addition, the direct co-culture supernatant of Hs578T cells and activated T cells did not induce the proliferation of MDA-MB231 cells.

Discussion

CD40 is mainly expressed on antigen presenting cells (APCs) including B cells, dendritic cells, and macrophages to activate immune responses after interaction with its ligand, CD40L, on activated T cells [39–41]. There are recent reports regarding CD40 expression on several kinds of tumor cells [42–45]; however, its role in tumor cells is not clearly known.

We found that CD40 is highly expressed in the breast cancer cell line MDA-MB231 and hardly expressed in the breast cancer cell line Hs578T (Fig 1). However, CD40L is not expressed in both of the cell lines (S2 Fig). This means that the possibility of an interaction between CD40 and CD40L in MDA-MB231 cells could be ignored. In contrast, CD40L is expressed in activated T cells (Fig 2A); however, CD40 is not expressed in activated T cells (S3 Fig). Therefore, an interaction between CD40 and CD40L in activated T cells could also be ignored. Based on our findings regarding CD40 and CD40L expression, the function of CD40 signaling in breast cancer cells was investigated with CD40 stimulating antibody, soluble CD40L and activated T cells.

First, we investigated the role of CD40 in the proliferation of MDA-MB231 cells because its original function in the immune system is that of a key mediator for the proliferation and differentiation of B cells [46, 47]. In contrast to our expectation, the proliferation of MDA-MB231 cells was not observed by stimulation of CD40 with anti-CD40 antibody (Fig 1B). There are several molecules that have different functions according to their origins [48, 49]. For example, IL-18 produced by immune cells has an anti-tumor activity; however, IL-18 produced by tumor cells has an important role in the immune escape process [50, 51]. Tumor cells secret TGF-β and IL-10, which are the most well-known immunosuppressive cytokines, to escape immune surveillance [30, 52]. For this reason, we next examined whether CD40 also has a role in the immune escape mechanism of breast cancer through the production of TGF-β. As shown in Fig 1C and 1D, CD40 stimulation increased the expression of TGF-β mRNA transcripts in MDA-MB231 cells. In addition, its secretion from MDA-MB231 cells is also increased by...
CD40 stimulation when treating the cells with its agonistic antibody and soluble ligand and by co-culturing with CD40L-expressing activated T cells (Fig 1E and 1F). Moreover, the increase in TGF-β production is completely blocked by pre-treatment with CD40 and CD40L neutralizing antibodies or by indirect co-culture of MDA-MB231 cells and activated T cells (Fig 2C). Therefore, our data not only show that CD40 has a role in the production of TGF-β from the tumor to suppress the immune system in the tumor microenvironment, but also has a novel mechanism in the increase of TGF-β production by CD40 stimulation.

Immune suppression by TGF-β is achieved by regulating cells in adaptive immunity components, such as T and B cells, as well as in innate immunity components, such as natural killer (NK) cells [42, 53, 54]. Additionally, it was recently reported that TGF-β suppresses immune responses by promoting regulatory T cell (Treg) induction and Th17 differentiation from conventional T cells [43, 44]. Because forkhead box protein 3 (Foxp3) is expressed when T cells become Treg, we examined whether Foxp3 is increased in T cells after stimulation of CD40L with anti-CD40L agonistic Ab or by co-culture with CD40-expressing MDA-MB231 cells. However, we could not detect an increase in Foxp3 expression (data not shown). Therefore, we next examined the role of the CD40-CD40L interaction in Th17 differentiation.

As shown in Fig 1E and 1F, MDA-MB231 cells produced TGF-β by stimulating CD40 with anti-CD40 agonistic antibody or sCD40L, and by co-culture with activated T cells. TGF-β has multiple cellular responses including the induction of cell growth inhibition, differentiation, wound healing and apoptosis [45]. TGF-β signaling can act as a tumor suppressor or tumor promoter depending on the tumor type and the stage of tumor progression [55]. Furthermore, TGF-β has a crucial role in Th17 cell lineage commitment [29]. It is known that TGF-β and either IL-6 or IL-21 are essential factors in the induction of Th17 differentiation [43]. Furthermore, we showed that the production of IL-1β, IL-6 and IL-21 is increased by direct co-culture of MBA-MB231 cells with activated T cells (Fig 3B–3D). This result suggests that the optimal condition for Th17 differentiation could be induced by the interaction between CD40 on the surface of the MDA-MB231 cells and CD40L on the surface of the activated T cells. In fact, we observed an increased population of IL-17-producing CD4+ T cells (Fig 3A). Our study coincides with a report that CD40-CD40L cross-talk is important in Th17 development [11].

As seen in Fig 1B, there was no effect from CD40 stimulation on the proliferation of MDA-MB231 cells in vitro. The proliferation of MDA-MB231 cells did not change even though TGF-β production was increased by CD40 stimulation. However, we found that the production and secretion of IL-17 were increased through the CD40-CD40L interaction. It is still controversial whether IL-17 has a tumor-suppressing effect or tumor-promoting effect [56]. In our study, IL-17 increased the proliferation of MDA-MB231 cells through the activation of STAT3. IL-17-mediated proliferation of MDA-MB231 cells was inhibited by the treatment with a STAT3 inhibitor AG490 and anti-IL-17 neutralizing antibody (Fig 5). CD40L is expressed in many cells including mast cells, macrophages, basophils, NK cells, B cells, smooth muscle cells, endothelial cells, and epithelial cells [44]. Based on the activated T cells, it seems that these cells also get a signal through CD40L found on the surface of the cells by interacting with CD40 on the surface of the MDA-MB231 cells. CD40 has an important role in generating T cell responses against viruses and bacteria through the interaction with CD40L on T cells [57, 58]. In particular, the role of CD40 in generating T cell responses provides the possibility of eliciting effective anti-tumor immune responses because CD40 on APC can deliver co-stimulatory signaling for the activation of CD8+ cells directly without the activation of CD4+ helper T cells [59, 60]. In fact, it was reported that efficient cytotoxic T lymphocytes responded against tumors when administering CD4 knock-out mice with CD40 activating monoclonal antibody [61]. That is, the ligation of CD40 on B cells up-regulates their co-stimulatory activity, and these cells might have a role in the generation of cytotoxic T lymphocyte responses against
tumors. However, the cytotoxicity of activated T cells against MDA-MB231 cells was not observed because unlike B cells, MDA-MB231 cells do not express co-stimulatory molecules on their surface. Therefore, the significance of activated T cells delivering their signals through CD40L will be further investigated. In addition, in vivo administration of CD40 stimulating antibody for the induction of cytotoxic T lymphocyte responses against tumors should be considered carefully because it can suppress immune responses through an increase in TGF-β production from tumors based on the results of this study.

In summary, an interaction between CD40 on MDA-MB231 breast cancer cells and CD40L on activated T cells increases the production of Th17 differentiating cytokines including TGF-β, IL-1β, IL-6, and IL-21. Moreover, the increased IL-17 promotes the proliferation of the breast cancer line MDA-MB231 through the activation of STAT3. Therefore, it seems that CD40 on tumor cells is closely related to the immune escape of tumors. Thus, by modulating the interaction between CD40 on tumor cells and CD40L on activated T cells, the efficiency of cancer immunotherapy could be increased by regulating the TGF-β production in a tumor microenvironment.

Supporting Information

S1 Fig. CD40 expression is down-regulated on MDA-MB231 cells by CD40 siRNA transfection. MDA-MB231 cells (1.5 x 10^5/well) were seeded on 6-well plate and then transfected with 20 nM of CD40 siRNA and control siRNA in the mixture of serum free media and oligofectamine as described in Materials and Methods. After 72 hrs, the down-regulation of CD40 expression on MDA-MB231 cells by CD40 siRNA transfection was confirmed by flow cytometry analysis. Result is the representative of three independent experiments. (TIF)

S2 Fig. CD40L is not expressed on MDA-MB231 and Hs578T cells. Both breast cancer cell lines, MDA-MB231 and Hs578T were collected at continuous log phase of growth. The expression of CD40L was examine by staining with PE-conjugated anti-human CD40L antibody (1 μg/ml), as described in Materials and Methods. Result is the representative of three independent experiments. (TIF)

S3 Fig. CD40 expression is not induced on activated T cells. Human T cells (2.5x10^6/ml) purified from human PBMCs were activated as described in Materials and Methods. And then CD40 expression on activated T cells was examine by staining with PE-conjugated anti-human CD40 antibody (1 μg/ml). Result is the representative of three independent experiments. (TIF)

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Author Contributions

Conceived and designed the experiments: JSK. Performed the experiments: HK YK SB JMK. Analyzed the data: YH WJL JSK. Contributed reagents/materials/analysis tools: Jiwon Choi JMH MJ Jiyea Choi. Wrote the paper: HK JSK.
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