A Distinct Role for Interleukin–6 as a Major Mediator of Cellular Adjustment to an Altered Culture Condition

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ABSTRACT
Tissue microenvironment adjusts biological properties of different cells by modulating signaling pathways and cell to cell interactions. This study showed that epithelial–mesenchymal transition (EMT)/mesenchymal–epithelial transition (MET) can be modulated by altering culture conditions. HPV E6/E7-transfected immortalized oral keratinocytes (IHOK) cultured in different media displayed reversible EMT/MET accompanied by changes in cell phenotype, proliferation, gene expression at transcriptional, and translational level, and migratory and invasive activities. Cholera toxin, a major supplement to culture medium, was responsible for inducing the morphological and biological changes of IHOK. Cholera toxin per se induced EMT by triggering the secretion of interleukin 6 (IL-6) from IHOK. We found IL-6 to be a central molecule that modulates the reversibility of EMT based not only on the mRNA level but also on the level of secretion. Taken together, our results demonstrate that IL-6, a cytokine whose transcription is activated by alterations in culture conditions, is a key molecule for regulating reversible EMT/MET. This study will contribute to understand one way of cellular adjustment for surviving in unfamiliar conditions. J. Cell. Biochem. 116: 2552–2562, 2015. © 2015 The Authors. Journal of Cellular Biochemistry Published by Wiley Periodicals, Inc. This is an open access article under the terms of the Creative Commons Attribution-NonCommercial-NoDerivs License, which permits use and distribution in any medium, provided the original work is properly cited, the use is non-commercial and no modifications or adaptations are made.

KEY WORDS: IL-6; CULTURE CONDITION; EPITHELIAL MESENCHYMAL TRANSITION (EMT); MESENCHYMAL EPITHELIAL TRANSITION (MET); CHOLERA TOXIN

Microenvironment is a crucial factor that regulates cellular growth and differentiation by modulating complex signaling pathways and intercellular interactions in different cell types [Abbott et al., 2008]. All viable organisms persistently respond to an altered environment to maintain homeostasis, and epithelial cells actively interact with their microenvironment to facilitate normal tissue homeostasis [Cousin et al., 2009]. Likewise, molecular crosstalk occurs between cancer cells and their microenvironment, which actively participates in cancer initiation and progression [Ghotra et al., 2009]. Culture medium represents in vitro microenvironment, and different culture conditions have triggered diverse phenotypic alterations of cells [Yu et al., 2009]. For human cancer cells, biochemical, and morphological changes have been induced by various culture conditions [Terasaki et al., 1986; Winkenwerder et al., 2003]. Epithelial–mesenchymal transition (EMT) is defined as a phenomenon in which epithelial cells are transformed into motile mesenchymal cells through a process known as mesenchymal metaplasia or the bidirectional differentiation of undifferentiated
stem cells [Battifora, 1976]. EMT is a well-known physiologic process in development [Lee et al., 2006]. Moreover, EMT plays a central role for invasion and metastasis of cancer cells [Niu et al., 2012]. Similarly, mesenchymal–epithelial transition (MET) occurs as a physiological process in embryonic development [Vainio and Lin, 2002], and MET enables migratory cancer cells to colonize and survive at remote sites for successful completion of metastasis [Brabletz et al., 2005]. EMT occurs at an early stage of carcinogenesis for migration and invasion, while MET occurs at a later stage for stabilization of invasive or migratory cells [Chaffer et al., 2006]. In this sense, EMT and MET are transient processes that occur sequentially to facilitate the formation of metastatic cancer [Brabletz et al., 2004].

Upon conducting an experiment in which culture medium was altered to assess the consequent changes in cultured cells, we made an interesting discovery: EMT/MET occurred reversibly in immortalized oral keratinocytes (IHOK) upon changing the culture medium in which the cells were grown. As EMT/MET is a coupled physiologic process that plays a pivotal role in development and cancer progression, the finding that EMT/MET can be reversibly induced by modifying microenvironment implies that developmental abnormalities or cancer progression can be controlled by making alterations to the environment. Hence, we attempted to search for key components and molecules in culture conditions that modulate EMT/MET. To achieve this goal, we closely examined reversible EMT that occurred in human papilloma virus (HPV) Er/E7-transfected IHOK when they were grown in different culture media.

Because cholera toxin stimulates the growth of mammary cells, it has been used as a culture supplement for epithelial biology and carcinogenesis studies [Stampfer, 1982]. Cholera toxin affects G-protein through the stimulation of adenylyl cyclase, which results in an increased level of intracellular cAMP [Evans et al., 1987]. In turn, cAMP regulates IL-6 as a transcription factor in a cell type-specific manner [Hershko et al., 2002]. IL-6, a multifunctional cytokine, has shown diverse effects depending on cell types. That is, IL-6 induced differentiation of neuronal or neuroglial cells in malignant glioma cells [Kimura et al., 2001; Shu et al., 2011], while IL-6 promoted cancer progression by inducing EMT in human breast or prostate cancer cell lines [Sullivan et al., 2009; Rojas et al., 2011].

From this study, we found that cholera toxin triggered the up-regulated secretion of IL-6, an E-cadherin repressor. In turn, IL-6 was responsible for the induction of reversible EMT, representing an adaptive process for IHOK cells to adjust to an altered culture condition. This study will contribute to understand one way of cellular adjustment for surviving in unfamiliar conditions, opening the possibility of controlling incurable human diseases through modulation of tissue microenvironment.

**MATERIALS AND METHODS**

**REAGENTS AND ANTIBODIES**

Antibodies against E-cadherin, snail, p21, CDK4, vimentin, and cyclin D1 were obtained from Cell Signaling Technology (Danvers, MA). Antibodies against p27 and CDK2 and Alexa Fluor 488-conjugated anti-rabbit IgG secondary antibody were obtained from Santa Cruz Biotechnology (Santa Cruz, CA). Anti-actin, FITC monoclonal anti-pan cytokeratin conjugate, and monoclonal anti-vimentin cy3 conjugate were purchased from Sigma (St. Louis, MO). Type I collagen was purchased from Nitta Gelatin Inc (Osaka, Japan).

**CELL CULTURE**

IHOK cell line was established and reported previously [Lee et al., 2005]. To analyze medium–dependent phenotypic changes, two kinds of keratinocyte culture media were used. IHOK cells were maintained in keratinocyte growth medium (KGM, Lonza, MD) with supplementary bullet kit (Lonza), which were then named as IHOK-KGM. As an alternative culture medium, EF-medium was used to culture IHOK, and the cells were named as IHOK-EF. After IHOK-EF cells were cultured for 120 days in EF-medium, they were cultured again in KGM medium for 120 days, and the cell line was subsequently named as IHOK-EF(KGM). IHOK-EF and IHOK-EF(KGM) required 120 days for adjustment before the aforementioned culture method could be applied. When IHOK cells started to proliferate without floating, 1 × 10^6 cells were plated and then transferred to a new dish once in 3–4 days. The medium was changed every other day. Lovo colorectal cancer cell line, MCF-7 human breast cancer cell line, and HT1080 human fibrosarcoma cells were obtained from American Type Culture Collection (ATCC, Manassas, VA). Lovo cells were maintained in minimum essential medium (MEM, Gibco, Carlsbad, CA). MCF-7 and HT1080 cells were maintained in Dulbecco’s Modified Eagles Medium (DMEM, Gibco BRL). YD-38 cells, an oral squamous cell carcinoma cell line established in our laboratory, were maintained in EF-medium. MCF-7 and Lovo cells with 1 × 10^6 cells/8 mL were seeded in 100 mm dishes. YD-38 and HT1080 cells with 5 × 10^5 cells/8 mL were seeded in 100 mm dishes, respectively. Each medium was changed every other day, and the cells were transferred to a new dish once in 3 days. For the control of mesenchymal cells, normal oral gingival fibroblasts were used, which were derived from mucosal disease-free patients who underwent wisdom tooth extraction. Informed consent was obtained from the patients for this study (IRB-2–2009-0002). Gingival fibroblasts were selected by a versene solution (0.2% EDTA in PBS) supplemented with 0.04% glucose from the explanted tissues, and the selected cells were maintained in F-medium. The detailed composition of each medium was presented in Supplementary Table S1.

**IMMUNOBLOTTING**

When the cells were grown overnight to 80% confluency, the cells (3 × 10^6 cells) were washed with cold PBS and lysed in Cell Lysis Buffer (Cell Signaling Technology). After lysis, protein complexes were boiled for 5 min in sodium dodecyl sulfate (SDS) sample buffer. Thirty microgram protein per sample was loaded on gel and separated by 10% SDS-PAGE. Then, proteins were transferred to polyvinylidene fluoride membranes, and the membranes were blocked in 5% milk in Tris-buffered saline containing tween 20 and then immunoblotted with appropriate antibodies.

**CELL PROLIFERATION ASSAY**

Proliferation of IHOKs was determined directly by counting cells. In brief, IHOKs cells (2 × 10^3) were seeded in 6-well plates. The number of cells per well was measured in a time-dependent manner. We detached the cells at indicated time points and centrifuged the cells with culture media. Supernatant medium was discarded, and the
cells were resuspended in a fresh medium. Ten microliter of culture medium-cell mixture was placed in hemacytometer, and the number of cells was counted using microscope. In the proliferation assay conducted for the normalization of invasion assay, $2 \times 10^4$ IHOK cells were seeded in 6-well plates.

**IMMUNOFLUORESCENCE**

IHOKs were seeded in a 4-well chamber slide. Cells were fixed in 4% formaldehyde in PBS for 15 min, permeabilized in 0.5% triton-X-100 solution, and rinsed three times in PBS for 5 min each. Then cells were labeled with antibodies against E-cadherin or snail (1:100), followed by incubation with Alexa Fluor 488-conjugated anti-rabbit IgG secondary antibodies. For cytokeratin and vimentin staining, cells were incubated with FITC monocular anti-pan cytokeratin conjugate (1:200) and monocular anti-vimentin cy3 conjugate (1:200), respectively. To detect DNA, cells were counterstained with DAPI, and the slides were mounted with mounting medium. Fluorescent images were captured and analyzed using LSM 510 laser scanning confocal microscope and software with 40× lens (Carl Zeiss Inc., Oberkochen, Germany).

**TRANSWELL MIGRATION ASSAY**

IHOKs were plated and cultured to 70% confluency. To evaluate the migratory activity of each cell line, we used each cell line’s own culture medium to fill the lower chamber, for the migratory potential was altered by changing the medium. In brief, $2 \times 10^4$ cells/well were resuspended either in F-medium without FBS or in KGM without the supplementary bullet kit. Then, the cells were seeded in the upper wells of 24-well transwell chambers (Coster, Cambridge, MA) for 48 hr. The cells’ own media were placed in the lower wells, and the cells migrated to the lower wells. After 48 hr incubation in $5\%$CO$_2$ incubator, cells were fixed in 10% formalin solution and stained in 0.025% crystal violet (Junsei, Japan). Then, the number of migratory cells was counted in five fields at 200× magnification. The mean for each chamber was calculated and was normalized by counting the number of IHOK cells that were separately incubated in the same condition.

**TRANSWELL INVASION ASSAY**

The transwell invasion assay was carried out by coating the upper chamber with Type I collagen, and the rest of the procedure was same as the procedure for transwell migration assay explained above. The data were normalized by counting the number of IHOK cells that were separately incubated in the same condition.

**GELATIN ZYMOGRAPHY**

Gelatinolytic activities of MMP-2 and MMP-9 were analyzed by gelatin zymography. IHOKs were seeded in 100 mm dish and incubated for 16 hr. Cells were then incubated in proper media for 48 hr. The conditioned media were centrifuged and quantified for protein content. The samples were loaded on a 10% SDS polyacrylamide gel copolymerized with 0.2% gelatin. After electrophoresis, the gel was renaturated in zymogram renaturation buffer (Bio–Red Laboratories) at room temperature for 2 hr, and then incubated in zymogram development buffer (Bio–Red Laboratories) for 16 hr at 37°C. The gels were stained with Coomassie blue and then destained.

**IHOKS XENOGRAFT IN ZEBRAFISH AND MICROSCOPIC OBSERVATION**

Zebrafish were maintained in accordance with the standard guidelines [Nüsslein-Volhard and Dahm, 2002]. Care and treatment of zebrafish were conducted in accordance with the guidelines established by the Animal Care and Ethics Committee of the Gwangju Institute of Science and Technology, Republic of Korea. After staining of cells, zebrafish embryos were de-chlorionized using micro-forceps, anesthetized with 0.0016% tricaine, and positioned on their right side on a wet 1.0% agarose pad. IHOKs were detached from culture dishes using 0.05% trypsin-EDTA and washed twice with PBS at room temperature. Cells were stained with 2 μg/ml Dil diluted in PBS and washed four times. $1 \times 10^2$ cells / ml were suspended in 10% FBS, which were then injected into the center of yolk sac using an injector (PV820 pneumatic picopump, World Precision Instruments, FL) equipped with borosilicate glass capillaries (World Precision Instruments, FL). The number of embryos exhibiting cell dissemination from the injection site was counted by inverted microscopy (Leica DMIRB microscope 090-132.101, JH Technologies, Boston, MA), and the embryos were imaged at 4 days post injection (DPI) by upright microscopy (Leica DMRBE microscope 301-371.011, JH Technologies).

**MICROARRAY DATA ANALYSIS**

IHOK-KGM, IHOK-EF, and IHOK-EFKGM were cultured to 70% confluency. Total RNA was isolated using RNeasy kit (Qiagen) according to the manufacturer’s instructions. Each total RNA sample (200 ng) was labeled and amplified using Low Input Quick Amp labeling kit (Agilent technologies, CA). The Cy3-labeled aRNAs were resuspended in 50 μl of hybridization solution (Agilent technologies). After labeled aRNAs were placed on Agilent Sure Print G3 Human GE 8 × 60K array (Agilent technologies) and covered by a Gasket 8-plex slide (Agilent technologies), the arrays were analyzed using an Agilent scanner with associated software. Gene expression levels were calculated with Feature Extraction v10.7.3.1 (Agilent technologies). Relative signal intensity for each gene was generated using the Robust Multi-Array Average algorithm. The data were processed based on the median polish normalization method using Gene Spring GX 7.3.1 (Agilent technologies). The normalized and log-transformed intensity values were then analyzed using Gene Spring GX 7.3.1 (Agilent technologies).

**QUANTITATIVE REAL-TIME PCR**

Total RNA was extracted using RNeasy kit (Qiagen), and cDNA was synthesized from 1 μg of the total RNA by using Transcriptor First Strand cDNA synthesis Kit (Roche) according to the manufacturer’s protocol. Real-time PCR was carried out using SYBER green Master (Roche applied science, Switzerland), and the results were analyzed using the Light Cycler 480 software. The primers used for real-time PCR were presented in Supplementary Table 2.

**ENZYME LINKED IMMUNOSORBENT ASSAY (ELISA)**

Each IHOK cell line was cultured with proper culture medium for 48 hr in a 100 mm dish. Their conditioned media were harvested for the analysis of IL-6 secretion. Concentration of IL-6 was measured...
and exhibited elongated shape. After additional 60 days, the cells lost cell to cell contacts, scattered, and recovered to polygonal epithelial shape, the shape of IHOK-KGM, but were hardly detected in IHOK-EF. The expressions of vimentin and snail, which are decreased upon changing the culture medium in which IHOKs were grown: up or down-regulated genes in IHOK-EF returned to their original expression levels in IHOK-KGM cells, in 120 days (Fig. 1A).

To investigate whether spindle-shaped cells of IHOK-EF went through EMT, EMT marker proteins were examined. Expression of E-cadherin, an epithelial differentiation marker, was reduced by 58% in IHOK-EF compared with IHOK-KGM, and involucrin expression was not detected in IHOK-EF. The expressions of vimentin and snail, an E-cadherin repressor, were 7.9-fold and 2.65-fold higher, respectively, in IHOK-EF than in IHOK-KGM (P < 0.05) (Fig. 1B). In immunofluorescence analysis, E-cadherin and cytokeratin were clearly localized in IHOK-KGM, but were hardly detected in IHOK-EF (Fig. 1C and Supplementary Fig. S1). However, snail and vimentin were clearly localized in IHOK-EF (Fig. 1D and Supplementary Fig. S1). Interestingly, IHOK-EFKGM expressed E-cadherin and cytokeratin with reduced snail and vimentin expression (Fig. 1B, C, D, and Supplementary Fig. S1). Taken together, these results indicate that culture conditions can modulate reversible EMT/MET in IHOK.

Proliferation of IHOKs was determined directly by counting cells. Proliferation of IHOK-EF reduced to 80% and to 40% compared with IHOK-KGM after 3 days and 7 days, respectively (*P < 0.05 by t-test). Both expressions of p21 and p27 increased and cyclin D1 expression reduced in IHOK-EF compared with IHOK-KGM (Supplementary Fig. S2A, B).

IHOK-EF CELLS HAVE HIGH MIGRATORY AND INVASIVE ACTIVITIES

To determine whether IHOK-EF had high migratory and invasive activities, we carried out transwell migration and invasion assays. To exclude proliferation-dependent effects of IHOK cells, the data were normalized by counting the number of IHOK cells that were separately incubated in the same condition used for migration and invasion assays. Briefly, we used 2 × 10^4 cells as was the case for migration and invasion assays. After 2 days, the 0.37% of IHOK-KGM (total number 3.5 × 10^5), 4.72% of IHOK-EF (total number 2.75 × 10^5), and 1.2% of IHOK-EFKGM (total number 3.0 × 10^5) migrated, respectively (P < 0.05) (Fig. 2A). To confirm that the in vitro migratory activity of IHOKs is correlated with in vivo dissemination of xenografted IHOKs, we injected three groups of cells including IHOK-KGM, IHOK-EF, and IHOK-EFKGM into zebrafish embryos. IHOK-EF showed 1.3-fold higher cell dissemination than IHOK-KGM (P < 0.05). Cell dissemination was reduced by 13% in IHOK-EFKGM compared with IHOK-EF (P < 0.05) (Fig. 2B). For invasive activity, the same number of cells with migratory activity was seeded. After 2 days, the 2.37% of IHOK-KGM, 8.0% of IHOK-EF, 2.43% of IHOK-EFKGM invaded, respectively (P < 0.05) (Fig. 2C). Gelatinolytic activity of MMP-2 in IHOK-EF was up-regulated and 12-fold higher than that in IHOK-KGM, but the activity in IHOK-EFKGM was reduced to the level of IHOK-KGM (Fig. 2D). Taken together, these results indicate that IHOK-EF acquired remarkable migratory and invasive activities, which were reversible, due to the changes in culture conditions.

GENE EXPRESSIONS IN IHOKS WERE ALTERED BY DIFFERENT CULTURE CONDITIONS

To identify whether gene expressions in IHOKs were altered by different culture conditions, we analyzed gene expression profiles by microarray. In the three types of IHOKs, mRNA levels were examined using Agilent Whole Human Genome Array which included 22602 human genes. Among the genes that were up-regulated or down-regulated in IHOK-EF compared with IHOK-KGM, those with > twofold changes were selected. Among the arrayed human genes, 13.6% (3080 genes) were altered in IHOK-EF, including 1518up-regulated genes and 1562 down-regulated genes. Figure 3 displays a reversible change in gene expression upon changing the culture medium in which IHOKs were grown: up or down-regulated genes in IHOK-EF returned to their original expression levels in IHOK-EFKGM. Next, we selected E-cadherin repressors, which were considered as principal mediators of property changes in IHOK, from the microarray data, and compared their expressions in three IHOK cell lines. As can be seen in Supplementary Table 3, five known E-cadherin repressors including SNAI1, IL-6, LOXL2, ZEB2, and TWIST1 showed higher expressions in IHOK-EF than in IHOK-KGM, and these expressions were reduced in IHOK-EFKGM compared with IHOK-EF.

IL-6 MODULATES REVERSIBLE EMT IN IHOK CELLS

To validate the microarray expression data for E-cadherin repressors, we performed real-time RT-PCR. Real-time RT-PCR data showed that the mRNA expression of IL-6 was changed by the largest amount upon changing the culture medium, unlike the microarray data (Fig. 4A). Among the five E-cadherin repressors, IL-6 showed 55-fold higher mRNA expression in IHOK-EF than in IHOK-KGM, but the expression was markedly reduced in IHOK-EFKGM. Accordingly, we assumed IL-6 to be a major contributor to the induction of reversible EMT in IHOK. Next, we examined the secretion level of IL-6 in the conditioned media of the three IHOKs. Secretion level of IL-6 was the highest in IHOK-EF, being 39.3-fold higher than that in IHOK-KGM.
Fig. 1. Modulation of IHOK phenotype by culture conditions. (A) a–d, IHOK-KGM was maintained in KGM medium for 120 days, which was used as a control IHOK cell line. e–h, IHOK-KGM was cultured in EF-medium for 120 days (IHOK-EF). i–l, IHOK-EF was re-cultured in KGM medium, which was the original medium for the control cells, for 120 days (IHOK-EFKGM). (B) Western blot was performed to measure the protein expression levels of E-cadherin and involucrin, the epithelial markers, and vimentin and snail, the mesenchymal markers. (C), (D) Cells were labeled with antibody against E-cadherin (Bar indicates 20 μm.) or antibody against snail (Bar indicates 10 μm.) (1:100), followed by incubation with Alexa Fluor 488-conjugated anti-rabbit IgG secondary antibody.
Fig. 2. Migratory and invasion activities of IHOKs grown in different culture conditions. (A) Each cell line was seeded on a transwell chamber, and the number of cells that migrated to the other side of the chamber was counted after 48 hr. In the lower chamber, each culture medium was applied as a chemoattractant. Each assay was performed in triplicate (*\( P < 0.05 \)). (B) Transplanted embryos were incubated in E3 media without methylene blue at 31°C. Four days later, the number of embryos exhibiting cell dissemination was counted (24 larvae/group). Representative fluorescent images (bottom) and merged DIC: fluorescent images (top). Blue arrow indicates cell mass disseminated from the injection site (yolk sac). Bar = 200 μm. Data are presented as average ± SD of six independent assays. Student’s t-test was used for comparison between experimental groups (# \( P < 0.01 \), * \( P < 0.05 \)). (C) For invasion assay, each cell line was seeded on a transwell chamber that was pre-coated with collagen type I. After 48h, the number of cells that invaded the collagen and moved to the other side of the chamber was counted (*\( P < 0.05 \)). (D) Each cell line was incubated in its culture medium for 48h. The conditioned media were collected and subjected to gelatin zymography. Gelatin zymography was performed using electrophoresis gels that contained 0.2% gelatin (*\( P < 0.05 \)).
IHOK-KGM (Fig. 4B). To investigate whether IL-6 induces reversible EMT in IHOK, we treated IHOK-KGM with IL-6. In 11 days, E-cadherin expression was markedly decreased, while expressions of both snail and vimentin were increased. The down-regulated expression of E-cadherin in IL-6-treated IHOK-KGM was reversed when IL-6 treatment ceased, accompanied by down-regulation of snail and vimentin (Fig. 4C and D). To evaluate whether IL-6 is capable of modulating reversible EMT in cancer cell lines, we treated MCF7 breast cancer cells, LOVO colon cancer cells, and YD-38 oral squamous cell carcinoma cells with IL-6. As shown in supplementary Figure S3, IL-6 induced reversible changes in the E-cadherin expression of LoVo and YD-38 cancer cells. MCF7 breast cancer cells exhibited a decrease in E-cadherin expression 2 hr after adding IL-6, but the expression was restored to the constitutional level after 24 hr.

CHOLERA TOXIN INDUCES EMT BY UPREGULATING THE PRODUCTION OF IL-6

In addition to investigating IL-6 for its involvement in reversible EMT, we also compared the supplements between KGM and F-medium, for changing the culture medium resulted in reversible EMT of IHOKs. Among the supplements to these two media, cholera toxin, and Triiodothyronine (T3) were included only in EF-medium. For this reason, we cultured IHOK-KGM with F-medium, which had the same composition as EF-medium except the five supplements including cholera toxin and T3, plus either cholera toxin or T3. As each group of cells was cultured for 120 days, the cells treated with cholera toxin lost cell to cell adhesion and showed spindle-like cell morphology compared with the non-treated control cells cultured only in F-medium. However, the cells treated with T3 showed no morphological changes compared with the control cells (Fig. 5A). To examine whether the alteration of cholera toxin-treated IHOK was correlated with the changes that occurred in IHOK-EF, we measured the expression of EMT-associated proteins and cell cycle-regulatory proteins, as the altered expression of these proteins was characteristic of IHOK-EF. In the cells treated with cholera toxin, expression of E-cadherin was down-regulated and expressions of vimentin and snail were up-regulated compared with the control cells, suggesting that cholera toxin contributed to the induction of EMT in IHOK. In addition, expression of cyclin D1 was down-regulated, while expression of p21, a CDK inhibitor, was up-regulated compared with the control cells (Fig. 5B). These results were compatible with the results of IHOK-EF shown in the supplementary figure S2A and B. Then, we looked into the relationship between IL-6 and the EMT of IHOK induced by cholera toxin. First, we measured the secretion level of IL-6 from the cells treated with cholera toxin or T3. The secretion level of IL-6 from the cells treated with cholera toxin was 3.6-fold higher than that from the control cells (P < 0.05) (Fig. 5C). Next, we analyzed whether a neutralizing antibody against IL-6 can block EMT induced by cholera toxin. As IL-6 concentration was markedly reduced (Fig. 5D), Cholera toxin-induced EMT was gradually reversed as could be seen by increased E-cadherin and reduced snail expressions (Fig. 5E). Therefore, these results suggest that cholera toxin-mediated increase in the secretion of IL-6 contributes to the induction of EMT in IHOK.

DISCUSSION

This study showed that the changes of culture conditions induce reversible EMT in IHOKs, not only in morphological changes and proliferation but also in migratory and invasive activities (Fig. 1 and Fig. 2). Reversibility of EMT/MET has been documented with respect to a variety of culture conditions. For instance, hypoxia can induce EMT, while reoxygenation triggers MET to reversibly modulate EMT/MET in cancer cells [Jo et al., 2009]. In addition, estrogen promotes reversible EMT-like transition and collective motility in MCF-7 breast cancer cells [Planas-Silva and Waltz, 2007], and high concentrations of glucose induce reversible EMT in human peritoneal mesothelial cells [Yu et al., 2009].

Our data showed that cholera toxin, the major difference between the supplements to KGM and EF-medium (Supplementary Table 1), induced EMT in IHOK cells (Fig. 5). With regard to morphological changes, cholera toxin induced morphological alteration in a cell type specific manner: In ovarian cancer cells, it induces to polygonal shape through MET process [Han et al., 1999]; In malignant glial cells, it induces neuroglial or neuronal differentiation [Kumura et al., 2001]; In other types of cells including Chinese hamster ovary cells, it induces EMT-like morphological changes [Rieber et al., 1975; Morinaga et al., 2001]. According to our results, cholera toxin is likely to trigger EMT in IHOK cells.

Cholera toxin has been used as a culture supplement, based on the knowledge that it stimulates the growth of mammary cells [Stampfer, 1982]. However, the conflicting data were reported with regard to cell growth. Cholera toxin enhanced the growth of
Fig. 4. IL-6-mediated induction of reversible EMT in IHOKs. (A) mRNA expressions of SNAI1, IL-6, LOXL2, ZEB2, and TWIST1 were analyzed by real-time PCR in IHOK-KGM, EF, and EFKGM cells. Real-time PCR was carried out using SYBER green I Master, and the results were normalized to the housekeeping gene GAPDH. (B) IHOKs were incubated in each culture medium for 48h for detection of IL-6. IL-6 concentration was measured in each conditioned medium by using IL-6 ELISA Kit (*P < 0.05). (C) IHOK-KGM was cultured in KGM medium treated with 10 ng/ml IL-6. Fresh medium containing IL-6 was added every other day. Then, the cells were cultured again in original KGM medium that lacked IL-6 for 11 days. (D) Immunofluorescence analysis confirmed the changes in the expression of E-cadherin, vimentin, and snail in IHOK-KGM that was treated with IL-6. Bars indicate 20 μm in the panels of E-cadherin and vimentin, and bar indicates 10 μm in the panels of snail.
human normal keratinocytes [Green, 1978] and type II pneumocytes [Uhal et al., 1998], whereas cholera toxin inhibited the proliferation of ovarian [Shaw et al., 2002] and lung cancer cells [Viallet et al., 1990]. On the other hand, accumulating data reported the relationship between cholera toxin and cell cycle regulatory proteins. Cholera toxin induces cell cycle arrest by blocking expression of cell cycle regulatory proteins such as cyclin D1 and CDK2 in malignant glioma cells [Li et al., 2007]. In addition, cAMP, which was elevated by cholera toxin, has been associated with cell cycle regulatory proteins. In cAMP-induced inhibition of cell cycle progression, cyclin D1 has been major target for cAMP [L’Allemain et al., 1997] and p27, upregulated by cAMP, promotes growth arrest [Kato et al., 1994]. Collectively, the role of cholera toxin with regard to cell cycle may depend on the biological status of the involved cells. In our study, we found that cholera toxin induced EMT and growth inhibition in IHOK cells (Fig. 5).

In our study, we focused on the search for a modulator of EMT. Previous literature has indicated that E-cadherin repressors play a major role in the induction of EMT by repressing E-cadherin transcription [Mejbjerg et al., 2007]. Thus, we selected five E-cadherin repressors as potential candidates for EMT modulation among genes whose expressions were significantly altered upon changing culture conditions: SNAI1, IL-6, LOXL2, ZEB2, and TWIST1. In particular, we focused on IL-6, for the expression of IL-6 was the highest among the five E-cadherin repressors in IHOK-EF (Fig. 4). We confirmed that IL-6 significantly contributed to the
reversible induction of EMT in IHOK by treating neutralizing antibody of IL-6 in the media.

We tested whether the reversible EMT by IL-6 could be induced in other human cancer cell lines such as oral squamous cancer cells, breast, and colon cancer cells. Notably, E-cadherin expression was responsive to IL-6 treatment in these cell lines, however, the responsive time point of each cell line was different. Moreover, we could not find further correlation between IL-6 treatment and the expression of mesenchymal marker proteins and cell cycle-associated proteins (data not shown). From these results, we considered that IL-6 is a molecule crucial for the initiation of EMT. However, the responsiveness by IL-6 treatment may be variable according to the cell types or biological conditions of the involved cells such as proliferation rate, the expression levels of endogenous IL-6 and its receptors and other molecular factors.

Apart from the EMT-inducing function of IL-6 as an E-cadherin repressor, IL-6 has also been known to be directly involved in tumorigenesis. With regard to this function of IL-6, accumulating evidence suggests that IL-6 promotes cancer progression by inducing invasion [Sansone et al., 2007; Wu et al., 2012]. However, IL-6 can also act as a cancer-protective factor in oncogene-induced cellular senescence, for it promotes the elimination of early neoplastic cells, similar to the transforming growth factorβ that performs dual functions in carcinogenesis [Kuilman et al., 2008]. Thus, the cytokastic function of IL-6 is likely to be context-dependent as suggested from these data. Likewise, the function of IL-6 as a molecular modulator of cellular adjustment to an altered environment depends on the context: EMT triggered by IL-6 might provide better survival opportunity for cells that are in the process of adapting to a new, unfamiliar environment.

In this study, we showed that cholera toxin induced EMT in IHOK by up-regulating IL-6 secretion. Neutralization of IL-6 in cholera toxin–treated IHOKs made the cells undergo a process of MET at the protein expression level, confirming the role of IL-6 for inducing EMT in IHOKs. However, we cannot exclude the possibility that other unknown components contained in the basal medium of KGM might be related to this reversible property by enhancing synergistic or antagonistic effects, for the composition of basal KGM medium could not be informed. Additionally, even if cholera toxin did regulate cell proliferation and EMT/MET in IHOKs, the effects of cholera toxin might be concentration-specific, for the growth of cells was uninhibited and EMT not induced when the cells were treated with 10-fold higher concentration of cholera toxin (data not shown) compared to the cholera toxin concentration in EF-medium.

In summary, our study demonstrated that pleiotropic cytokine IL-6, whose production is regulated by changes in culture conditions, modulates EMT/MET in an autocrine manner to facilitate adjustment of cells to an altered environment. This result will contribute to understand one way of cellular adjustment for surviving in unfamiliar conditions, opening up the possibility of controlling incurable human diseases through modulation of tissue microenvironment.

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