Epigallocatechin gallate inhibits the growth and promotes the apoptosis of bladder cancer cells

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Abstract. Epigallocatechin gallate (EGCG) has been revealed to inhibit the proliferation and induce the apoptosis of several types of tumor, in addition to inhibiting DNA methyltransferase activity, leading to CpG demethylation. Tissue factor pathway inhibitor 2 (TFPI-2) expression is downregulated in bladder cancer. The present study revealed that this downregulation was partly due to hypermethylation of the TFPI-2 gene promoter, which was decreased by EGCG treatment. In addition, the present study demonstrated that EGCG could inhibit the viability and invasion, and induce the apoptosis, of bladder cancer T24 cells. Furthermore, western blot analysis and reverse transcription-quantitative polymerase chain reaction analyses demonstrated that EGCG could upregulate the expression of TFPI-2. These results suggest that EGCG inhibits the growth and induces the apoptosis of bladder cancer cells through restoring TFPI-2 expression. Thus, EGCG is a potential therapeutic candidate for the treatment of bladder cancer.

Introduction

Tissue factor pathway inhibitor 2 (TFPI-2) is a recently identified serine proteinase inhibitor, which inhibits a number of proteinases, including matrix metalloproteinase (MMP), trypsin, and fibrinogenase (1). MMP, the key enzyme that degenerates extracellular matrix, can be inhibited by TFPI-2, thus reducing tumor invasion and metastasis. TFPI-2 has been reported to have this effect in ovarian cancer, pancreatic cancer, prostate cancer and glioma (2-5). It has also been reported that the overexpression of TFPI-2 in U-251 glioma cells induces apoptosis (5). However, the mechanism by which TFPI-2 is downregulated in various cancers remains

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undetermined. It has been reported that the downregulation of TFPI-2 in pancreatic cancer, glioma and hepatic cancer could result from CpG hypermethylation of promoters (6-8). Using a methylation inhibitor such as 5-Aza-2'-deoxycytidine (5-Aza-dC) could induce the expression of TFPI-2 in hepatocellular carcinoma cells (9).

Cancer is subject to genetic and epigenetic modifications. The methylation of CpG islands in tumor suppressors is currently thought to be the primary mechanism of epigenetic modifications in cancer. DNA methylation refers to the methylation of the fifth carbon on the pyridine ring of cytosine by DNA methyltransferase (DNMT) (10). Therapeutic methods of targeting CpG island methylation primarily focus on the inhibition of DNMT using drugs like 5-Aza-dC. A recent report demonstrated that epigallocatechin gallate (EGCG) can inhibit DNMT with minimal cytotoxicity and potent efficacy in solid tumors (11). EGCG is the main component of catechin in tea and conjugates with DNMT1 via four hydrogen bonds. However, no proof of health benefits of EGCG have been acknowledged in Europe or the US (12,13). This may be due to the poor bioactivity of EGCG when taken orally. However, in experiments using purified EGCG, there has been promising results showing its antitumor activity (14).

Bladder cancer is the eighth most common cancer, the ninth most common cause of cancer-associated mortality and was once the most commonly diagnosed urological cancer in China prior to the era of prostate-specific antigen screening (15). However, bladder cancer remains difficult to treat. The resection of a bladder tumors that do not invade the muscle is followed by a 50% recurrence rate and ~50% of muscle-invading bladder tumors metastasize (14). Novel treatments based on the underlying molecular mechanisms of bladder cancer pathogenesis are urgently required. The present study aimed to determine whether TFPI-2 is hypermethylated in bladder cancer and whether the DNMT inhibitor EGCG could inhibit the progression of bladder cancer.

Materials and methods

Methylation-specific polymerase chain reaction (PCR) analysis. A total of 24 fresh tissue samples from patients diagnosed with bladder cancer were included in the present study. Normal and cancerous bladder tissue samples were obtained via transurethral resections or radical cystoprostatectomies, which were performed at Huashan Hospital (Shanghai, China). All patients provided written informed consent and the Ethics Committee of Huashan Hospital judged the present study to be exempt from ethical approval. Probes were constructed using PyroMark Assay Design software (version 2.0; Qiagen, Inc., Valencia, CA, USA) as follows: TFPI-2 forward, 5'-TGGTTTT TTAAAGTGTTGGGGATTA-3' and reverse, 5'-ATATAACCA CCAATAAAAAATTAAAACATT-3' with 5' biotinylation; and TFPI-2 probe, 5'-AGTGTTGGGATTATAGG-3' without 5' biotinylation. Genomic DNA was extracted using the Qiagen DNeasy tissue kit (Qiagen GmbH; Hilden, Germany) according to the manufacturer's instructions and was subsequently dissolved using sodium bisulfite mixed with 800 μ l of RNase-free water. The mixture was placed at room temperature (15-25°C) and was then subject to PCR conversion (95°C for 5 min, 60°C for 25 min, 95°C for 5 min, 60°C for 85 min, 95°C for 5 min, 60°C for 175 min and 20°C for holding). Freshly prepared Buffer BL was added and samples were transported to EpiTect spin columns. Following centrifugation at 16,000 x g for 1 min at room temperature, 500 μ l of Buffer BW was added to the columns, which were subsequently treated with Buffer BD and left incubated at 15-25°C for 15 min. Following further centrifugation at 16,000 x g for 1 min at room temperature, columns were treated with Buffer BW. When all remnant liquid was removed via several centrifuges, columns were incubated at 56°C for 5 min. Purification was finalized with Buffer EB treatment for the collection of purified DNA. Methylation PCR was conducted using the 50 µl system. Reactions were run at 95°C for 3 min; 94°C for 30 sec, 54°C for 30 sec, and 72°C for 1 min for 40 cycles; followed by 72°C for 7 min. Pyrosequencing was performed by adding conjugated beads into PCR tubes. The beads were aspired using a vacuum pump and were immersed into 70% ethanol, 0.2 M NaOH and rinsing buffer for 5 sec each. Beads and PCR products were subsequently treated with 40 μ l of annealing buffer and subjected to 85°C for 2 min prior to cooling down to room temperature. Quantity of substrate, enzyme, and dNTP (Qiagen, Inc.) added was obtained using Pyrose quenching software (PyroMark Q96 ID; Qiagen, Inc.). Pyro Q-CpG software (version 1.0) was used for automatic analysis of the methylation status of TFPI-2.

Reverse transcription-quantitative PCR (RT-qPCR). For RT-qPCR, total RNA was extracted from T24 cells or normal bladder tissue with TRIzol (Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA). Tissue was ground and treated with 1 ml of TRIzol reagent. Samples were subsequently treated with chloroform and subject to centrifugation at 16,000 x g for 15 min at 4°C. Supernatants were extracted and treated with isopropanol. Following centrifugation at 16,000 x g for 10 min at 4°C, supernatant was dispersed and RNA was dissolved in ethanol. Following centrifugation, products was maintained in diethyl pyrocarbonate water. The RNA concentration was determined using a spectrophotometer. Retro-transcription was conducted in the system with 5X All-In-One RT MasterMix, extracted RNA, and water. Reactions were run at 37°C for 15 min and 85°C for 5 sec and RNA was maintained at -20°C. The primers used for qPCR were as follows: TFPI-2 forward, 5'-CTTGCGACGATGCTT GCTG-3' and reverse, 5'-ACACCCACCGGAAAAGAA TTT-3'; and GAPDH forward, 5'-ATGGGGAAGGTGAAGG TCG-3' and reverse, 5'-GGGGTCATTGATGGCAACA ATA-3'. PCR reactions were run in the system containing SYBR-Green, primers, RoxDye (II), RT products, and water. Cycling condition used were as follows: 95°C for 30 sec; 95°C for 3 sec and 60°C for 30 sec for a total of 40 cycles. For each sample, the average threshold cycle value was normalized to GAPDH and the relative expression was quantified according to the $2^{-\Delta\Delta Cq}$ method (16).

Western blotting. An established protocol was used for western blotting (17-20). Total protein was extracted from T24 cells or normal bladder tissue. Briefly, to prepare cellular protein, plates were placed on ice and treated with pre-chilled proteinase inhibitor consisting of phenylmethylsulfonyl fluoride and phosphatase and were subsequently treated with protein extraction kit (Sangon Biotech Co., Ltd., Shanghai, China). Cells were scraped and debris were filtered using centrifuge. Tissue chunks were sliced and homogenated using the same agents. Samples were treated with lysate and proteins were prepared similarly to cells. Protein concentration was determined via the bicinchoninic acid (BCA) assay (Sangon Biotech Co., Ltd.). A total of 50 µg protein were separated by SDS-PAGE on a 10% gel and subsequently transferred to a polyvinylidene difluoride membrane at a constant voltage. Following blocking using 5% bovine serum albumin (Gibco; Thermo Fisher Scientific, Inc.) at room temperature overnight, TFPI-2 primary antibodies (cat. no. YT5159; ImmunoWay Biotechnology Company, Suzhou, Shina) were diluted to 1:5,000 with Tris-buffered saline/Tween-20 and incubated with the membranes for 90 min at room temperature. Following washing using tris buffered saline with Tween-20 three times, the membrane was incubated with horseradish peroxidase-labeled goat anti-rabbit IgG secondary antibody diluted at 1:1,000 (cat. no. 111-035-003; Jackson ImmunoResearch Laboratories, Inc., West Grove, PA, USA). Protein bands were then detected using chemiluminescence and autoradiography (ChemiDoc XRS; Bio-Rad Laboratories, Inc., Hercules, CA, USA). Densitometry was determined using the Quantity One software version 4.6.2 (Bio-Rad Laboratories, Inc., Hercules, CA, USA).

Cell culture. Human T24 bladder urothelial cancer cells were purchased from the Cell Bank of the Chinese Academy of Sciences (Shanghai, China). T24 cells were cultured in RPMI-1640 (Gibco; Thermo Fisher Scientific, Inc.) supplemented with 10% fetal bovine serum (Gibco; Thermo Fisher Scientific, Inc.) at 37°C in an atmosphere containing 5% CO₂. Cells were subsequently treated with different concentrations of EGCG (10 and 20 μ M) or PBS buffer (control) for 6 days.

MTT assay. Briefly, 500 cells/well were seeded into 96-well plates. Cells were incubated at 37°C and treated with MTT and dimethyl sulfoxide for 4 h and 10 min at 37°C, respectively. Treated (EGCG vs. PBS as control) cells were analyzed on a plate reader and quantified by measuring the absorbance at 490 nm.

Apoptosis assay. For flow cytometry, the Annexin V-fluorescein isothiocyanate (FITC) and propidium iodide (PI) (BD) method was employed using an Annexin V/FITC kit (BD Biosciences,



Figure 1. Methylation of the TFPI-2 gene in bladder cancer is decreased by EGCG. Methylation-specific polymerase chain reaction analysis revealed (A) a significantly higher methylation level of TFPI-2 in bladder cancer compared with normal bladder tissue, and (B) significantly decreased methylation of TFPI-2 in T24 bladder cancer cells treated with EGCG. **P<0.001. TFPI-2, tissue factor pathway inhibitor 2; EGCG, epigallocatechin gallate.



Figure 2. EGCG restores the expression of TFPI-2 mRNA in bladder cancer cells. Reverse transcription-quantitative polymerase chain reaction analysis of TFPI-2 revealed increased TFPI-2 mRNA expression in T24 bladder cancer cells treated with different doses of EGCG compared with the PBS-treated control. **P<0.01 vs. the PBS control group. TFPI-2, tissue factor pathway inhibitor 2; EGCG, epigallocatechin gallate; NS, no significance; NL, healthy adjacent bladder tissue.

San Jose, CA, USA) according to the manufacturer's instructions. T24 bladder cancer cells were harvested and HEPES binding buffer containing Annexin V-FITC and PI was applied for 15 min at room temperature. Samples were analyzed using a FACSCanto[™] flow cytometer (BD Biosciences) to determine the percentage of early apoptotic cells (Annexin V⁺/PI⁻).

Cell invasion assay. A Transwell assay was used to investigate the invasive abilities of the T24 bladder cancer cells. Briefly, Transwell inserts were coated with Matrigel (1:2 dilution; BD Biosciences). Prior to placing the inserts into the chambers, the chambers were filled with complete media. Approximately $1x10^6$ cells/ml were seeded in 300 µl of serum-free RPMI-1640 medium with the addition of EGCG in the inserts, which were subsequently placed into the chambers. After 72 h of incubation, the inserts were stained with crystal violet at room temperature for 30 min and the number of invasive cells was counted using a light microscope at a magnification of x100 (Nikon 80i Eclipse; Sendai, Japan).

Statistical analysis. Data are presented as the mean \pm standard deviation. All experiments were performed in triplicate and were two-tailed. The unpaired t-test was used to determine the statistical significance of differences between two groups. For comparisons between more than two groups, one-way analysis

of variance with a post hoc Geisser-Greenhouse correction was used. P<0.05 was considered to indicate a statistically significant difference. Statistical analyses were performed using SPSS v17 software (SPSS, Inc., Chicago, IL, USA).

Results

Methylation of the TFPI-2 gene in bladder cancer is decreased by EGCG. Our group has previously demonstrated decreased expression of TFPI-2 in renal cell carcinoma and bladder cancer (17,21). The present study investigated whether this decrease in expression was a result of DNA methylation. A total of 24 bladder cancer samples were analyzed, which revealed a significantly higher methylation level in the TFPI-2 gene compared with healthy adjacent bladder tissue (P<0.01; Fig. 1A). DNA methylation levels of the TFPI-2 gene in the bladder cancer cells treated EGCG were significantly decreased compared with the control cells (P<0.0001; Fig. 1B).

EGCG restores the expression of TFPI-2 in bladder cancer cells. Our group has previously reported that EGCG restores TFPI-2 expression levels in renal cell carcinoma (17). The present study investigated the effect of EGCG on the expression of TFPI-2 in bladder cancer (Fig. 2). EGCG significantly increased TFPI-2 mRNA expression compared with the control



Figure 3. EGCG restores the expression of TFPI-2 protein in bladder cancer cells. Western blotting for TFPI-2 revealed an increased TFPI-2 protein level in T24 bladder cancer cells treated with different doses of EGCG. ****P<0.0001 vs. the PBS control group. TFPI-2, tissue factor pathway inhibitor 2; EGCG, epigallocatechin gallate; NL, healthy adjacent bladder tissue.



Figure 4. EGCG increases the apoptosis and decreases the viability of bladder cancer cells. The results of the apoptosis assay of the (A) PBS-treated control, (B) 10 μ m EGCG and (C) 20 μ m EGCG groups demonstrated increased apoptosis in the EGCG treated cells. (D) The cell viability assay revealed that 20 μ m EGCG significantly inhibited bladder cancer cell growth. **P<0.01 vs. the PBS control group. EGCG, epigallocatechin gallate; NS, no significance; FITC, fluorescein isothiocyanate; PI, propidium iodide.

group (P<0.01) in a dose-dependent manner. Notably, normal bladder tissue exhibited a similar level of TFPI-2 expression compared with T24 bladder cancer cells. The protein level of TFPI-2 in bladder cancer cells treated with different doses of EGCG was then investigated (Fig. 3), which revealed that the

protein level of TFPI-2 was significantly increased in response to 20 μ m EGCG compared with the control group (P<0.0001). By contrast, the protein level of TFPI-2 in T24 bladder cancer cells was marked lower compared with that in adjacent healthy bladder tissue.



Figure 5. EGCG inhibits the invasion of bladder cancer cells. The results of the Transwell invasion assay at a magnification of x100 of the (A) PBS-treated control, (B) 10 μ m EGCG and (C) 20 μ m EGCG groups wells. (D) Quantification of the Transwell invasion assay revealed that 10 and 20 μ m EGCG treatment significantly inhibited bladder cancer cell invasion. ***P<0.001 vs. the PBS control group. EGCG, epigallocatechin gallate.

EGCG increases the apoptosis, and inhibits the viability and invasion of bladder cancer cells. Given the restoration of TFPI-2 expression in the T24 bladder cancer cells following EGCG treatment, the effect of EGCG was further investigated in vitro. This revealed that the apoptosis of bladder cancer cells was notably increased by 20 μ m EGCG (Fig. 4A-C). In addition, 20 μ m EGCG inhibited bladder cancer cell viability (P<0.01 vs. the control group; Fig. 4D). However, 10 μ m EGCG did not exhibit any significant inhibitory effect on T24 cell viability (Fig. 4D). The effect of EGCG on the invasive ability of the bladder cancer cells was then investigated, which demonstrated that EGCG significantly inhibited invasion compared with the control group (P<0.01; Fig. 5).

Discussion

Although substantial progress has been made in the treatment of bladder cancer, the outcome of patients with invasive bladder cancer has not substantially improved in the last 20 years. Novel treatments that target specific tumor-associated targets, including TFPI-2, could minimize adverse advents and enhance treatment efficacy. However, the role of TFPI-2 in bladder cancer remains unclear. Our group has previous identified that TFPI-2 is downregulated in bladder cancer (21). In addition, the present study identified that the TFPI-2 gene was hypermethlyated in bladder cancer. It was thus hypothesized that inhibition of TFPI-2 methylation via the administration of EGCG could restore TFPI-2 expression and inhibit tumor growth. Several previous studies have investigated the effect of tea extracts, including ECGC, in cancer therapy (22,23). Hypermethylated tumor suppressors are a common characteristic of cancer, and bioactive flavonoids that are natural extracts from fruits, vegetables and other plants have been proposed to decrease hypermethylation (24). EGCG, an antioxidant polyphenol flavonoid isolated from green tea, has been identified to serve a chemopreventive role by certain studies (25). The effect of EGCG in atherosclerosis and neurodegenerative diseases, in addition to cancer, is being investigated. Specifically, EGCG has been suggested to function via the following mechanisms in cancer: i) Inducing apoptosis (26-28); ii) inhibiting epidermal growth factor receptor-mediated signaling pathways (29,30); iii) inhibiting signaling between vascular endothelial growth factor and MMPs, thus compromising angiogenesis (31,32); iv) inhibiting telomerase and DNMT (33-36); and v) reducing oxidative stress levels (35).

To the best of our knowledge, our group is the first to demonstrate that TFPI-2 could be one of the genes for which DNA methylation is inhibited by EGCG. A previous study by our group investigating renal cell carcinoma demonstrated a similar result concerning TFPI-2 hypermethylation (17). The present study demonstrated that methylation of the TFPI-2 gene is prevalent in bladder cancer. Previous investigations of prostate cancer and TFPI-2 have revealed that the methylation of TFPI-2 is a common feature of urological malignancies (4). In addition, it has previously been shown that TFPI-2 expression is inversely associated with progression of malignancies in general (1-11). Notably, coagulation abnormalities are a common characteristic of cancer. Tissue factor (TF) promotes clotting and thus promotes the progression of malignancies (37). TFPI-2 inhibits TF and thus cancer growth. Aberrant methylation of TFPI-2 promoter CpG islands in human cancer and cancer cell lines is widely accepted to be responsible for the downregulation of TFPI-2 expression during cancer progression (4). Additionally, an aberrantly spliced variant of TFPI-2 mRNA that leads to an untranslated form of TFPI-2 is also responsible for loss of TFPI-2 function (38).

In conclusion, the results of the present study indicate that ECGC restores TFPI-2 expression in bladder cancer, thus 3518

inhibiting invasion and viability, and increasing apoptosis. Therefore, ECGC is novel therapeutic candidate for the treatment of patients with bladder cancer and requires further study.

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