The expression and function of Toll-like receptors 3 and 9 in human colon carcinoma

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Abstract. Toll-like receptors (TLRs) are pattern-recognition receptors that are important in immune signaling. TLR recognition of various viral components including double-stranded RNA (TLR3) and unmethylated CpG-DNA (TLR9) plays a crucial role in cell survival. However, TLR expression and function in colon carcinoma cells are not well clarified. We investigated the expression of TLR3 and TLR9 in colon carcinoma cells using immunohistochemical methods. The function of TLR3 and TLR9 signaling in carcinoma cell lines was studied by direct cell stimulation with, or by cell transfection of, polyinosinic-polyctydylidic acid (Poly I:C), a synthetic form of dsRNA, and by cell stimulation with CpG-oligodeoxynucleotides (ODNs), respectively. Positive TLR3 and TLR9 immunohistochemical staining was observed in 91 and 86% of human hepatocellular carcinoma (HCC) tissues, respectively. Cell surface stimulation of TLR3 with Poly I:C did not affect cell viability but it did activate NF-κB activity. By contrast, stimulation of intracellular TLRs with transfected Poly I/C significantly induced apoptosis. Cell surface stimulation of TLR9 with CpG-ODNs promoted cell proliferation, and, furthermore, these CpG-ODN TLR9 agonists reduced the cytotoxicity of the anticancer drug adriamycin. Cell surface expression of TLR3 and TLR9 in colon carcinoma cells plays an important role in cell survival. In addition, the proapoptotic activity of intracellularly expressed TLR3 may provide the possibility of using TLR agonists as novel clinical cytotoxic agents against colon carcinoma cells.

Introduction

Toll-like receptors (TLRs) play a critical role in innate immunity against microbial pathogens, as well as in the subsequent induction of adaptive immune responses (1-3). These receptors are the major pattern-recognition transducers in response to microbial intruders such as bacteria, protozoa, fungi or viruses (4). Stimulation of TLRs induces a range of innate and adaptive immune responses through cytokines, interferons, chemokines and cell surface molecules, in addition to increasing the effector functions of TLR-expressing cells.

TLR2 and TLR4 play a key role in the recognition of various bacteria. Viral glycoproteins are recognized by TLRs 2 and 4, virus-related double-stranded RNA (dsRNA) by TLR3 and single-stranded RNA (ssRNA) by TLRs 7 and 8. TLR9 recognizes specific oligodeoxyxynucleotide (ODN) sequences that consist of unmethylated CpG-ODNs, which are frequently present in bacterial and viral DNA (5).

TLRs are broadly distributed in various cells of the immune system, including polymorphonuclear phagocytes, monocytes, dendritic cells and natural killer cells, as well as in some epithelial and endothelial cells (6,7). The specific subcellular localization of TLRs remains unclear. TLRs 1, 2, 4, 5 and 6 are present in the plasma membrane. TLRs 3, 7, 8 and 9 are mainly present in endosomes and are likely to signal from acidic endosomes (8).

Signaling pathways activated by specific TLRs are largely dictated by the adaptor proteins that are recruited to the intracellular domain of the TLR. MyD88 is involved in the majority of TLR pathways, except for TLRs 3 and 8. TLR3 signaling depends solely on its binding to the adaptor protein termed the Toll/Interleukin-1 receptor domain-containing adaptor-inducing IFN-β (TRIF) adaptor protein. Binding of TRIF to TLR3 leads to the activation of NF-κB and IRF3 transcription factors, thereby inducing the antiviral interferon response (9,10).

In humans, TLR9 is expressed in B-lymphocytes, monocytes and plasmacytoid dendritic cells. TLR9 recognizes specific oligodeoxyxynucleotide (ODN) sequences consisting of unmethylated CpG-ODNs, which are frequently present in bacterial and viral DNA (5).

TLRs have recently been found to be expressed in various normal epithelial and cancer cells (11-18). Of these cells, gastrointestinal epithelial cells in particular are constantly exposed to external microbial pathogens or viruses. Therefore, TLRs play a critical role not only in immune defense but also in other biological functions.

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In addition, we have previously reported that human hepatocellular carcinoma (HCC) cells express TLR3 and TLR9 and that these receptors function mainly in cell survival and in anti-apoptotic pathways (15,16). Previous reports indicated that colorectal cancer cells express TLR2, TLR3, TLR4, TLR5 and TLR9. These findings indicated that TLRs might play an important role also in colorectal cancer cells.

Recent reports have suggested that the neoplastic process may sabotage TLR signaling pathways to favor cancer progression. TLRs on tumor cells facilitate their evasion from immune surveillance via suppression of T-cell proliferation and natural killer cell activity. These studies suggest that TLR signaling in tumor cells is associated with the progression of cancer (12,14). However, the functions of TLRs in colon carcinoma cells are not well understood. We therefore investigated the expression of TLR3 and TLR9 in colon carcinoma cell lines and in primary human normal and carcinogenic colon tissues, and attempted to clarify the function of TLR3 and TLR9 in colon carcinoma cells.

Materials and methods

Cells and HCC tissues. The colon adenocarcinoma cells Colo320 and SW480 were purchased from the American Type Culture Collection (Rockville, MD, USA). All cells were cultured in DMEM at 37°C, supplemented with 1% penicillin/streptomycin (Gibco BRL, Grand Island, NY, USA) and 10% heat-inactivated fetal calf serum (Gibco BRL).

A total of 42 colon carcinoma tissues (9 non-tumor tissues, and 8 tissues with metastasis from colon carcinoma tissues) were obtained from tissue array slides (SuperBioChips Laboratories, Seoul, Korea). We obtained informed consent from all patients prior to the subsequent use of their resected tissues. Resected tissues were frozen immediately at -80°C or were fixed in 10% formalin.

Reagents. Polyinosinic-polycytidylic acid (Poly I:C), which was used as a human TLR3 ligand, was obtained from Sigma (St. Louis, MO, USA), and Lipofectamine LTX (Lipo) was obtained from Invitrogen (Carlsbad, CA, USA). The type C CpG oligonucleotide (ODN M362: 5'-tgc tgc tgc ttg caa gca gct tga t-3'), which was used as a human TLR9 ligand, was obtained from Invitrogen (Carlsbad, CA, USA) and anti-α-tubulin monoclonal antibody (Imgenex, San Diego, CA, USA) was used as a control. The non-stimulatory oligonucleotide ODN M362 (ODN M362 control: 5'-tgc ttc tgc tgg taa gca gct tga t-3'), which was used as a human TLR9 ligand, was obtained from Invitrogen (Carlsbad, CA, USA). The anti-α-tubulin monoclonal antibody (Imgenex, San Diego, CA, USA) was used as a control. The non-stimulatory oligonucleotide ODN M362 (ODN M362 control: 5'-tgc ttc tgc tgg taa gca gct tga t-3'), which was used as a human TLR9 ligand, was obtained from Invitrogen (Carlsbad, CA, USA).

Immunohistochemical staining. Immunohistochemical staining of TLR3 and TLR9 expression in colon carcinoma and non-colon carcinoma tissues was performed using the labeled streptavidin-biotin method. Deparaffinized sections were heated for 5 min at 120°C in a pressure cooker to reactivate the antigen. Sections were blocked and incubated with anti-TLR3 or 9 antibodies (Santa Cruz Biotechnology, Santa Cruz, CA, USA), overnight at 4°C. The sections were then incubated with a second biotinylated antibody, followed by an avidin-biotin-peroxidase complex. The sections were then developed in a substrate solution of 0.01% 3,3'-diaminobenzidine-hydrogen peroxidase and counterstained with 10% hematoxylin.

Table I. Expression of TLR3 in colon carcinoma and non-tumor tissues.

<table>
<thead>
<tr>
<th>Histology</th>
<th>Absent or weak (%)</th>
<th>Moderate (%)</th>
<th>Strong (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Non-tumor tissue</td>
<td>1 (11.1%)</td>
<td>8 (88.9%)</td>
<td>0 (0%)</td>
</tr>
<tr>
<td>Colon carcinoma</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Poorly differentiated</td>
<td>1 (33.3%)</td>
<td>2 (66.7%)</td>
<td>0 (0%)</td>
</tr>
<tr>
<td>Moderately differentiated</td>
<td>4 (6.4%)</td>
<td>36 (57.1%)</td>
<td>23 (36.5%)</td>
</tr>
<tr>
<td>Well differentiated</td>
<td>3 (12.0%)</td>
<td>15 (60.0%)</td>
<td>7 (28.0%)</td>
</tr>
</tbody>
</table>

Detection of TLR 3 and 9 protein expression by immunoblotting. The expression of TLRs 3 and 9, and that of the loading control α-tubulin, in colon adenocarcinoma cell lines (Colo320, SW480) was analyzed by immunoblotting. Briefly, after incubation of the cells on 6-well plates (Nunc™ Brand Products, Denmark) for 48 h, the cells were washed twice with phosphate-buffered saline (PBS) and were then lyed by the addition of SDS sample buffer (50 mmol/l Tris-HCl, pH 6.8, 2.5% SDS, 5% glycerol, 5% 2-mercaptoethanol and 0.01% bromphenol blue). Equal amounts of extracted proteins were separated by SDS-PAGE and were then transferred to PVDF membranes (Millipore, Billerica, MA, USA). Blots were blocked by incubation in Tris-HCl (pH 7.5) containing 5% milk and 0.1% Tween-20 for 30 min at room temperature, and were probed overnight at 4°C with the primary antibodies. The following primary antibodies were used: anti-TLR3 or TLR9 monoclonal antibody (Imgenex, San Diego, CA, USA) and anti-α-tubulin monoclonal antibody (Oncogene Research Products, San Diego, CA, USA). The antibodies were diluted in 5% milk or in BSA in Tris-HCl (pH 7.5) containing 0.1% Tween-20. The immunoblots were then probed with horseradish peroxidase-conjugated anti-mouse immunoglobulin G (IgG) (diluted 1:1000 in 1% milk or in BSA in Tris-HCl, pH 7.5) (Amersham Biosciences, Buckinghamshire, UK). After the final wash, signals were detected using an ECL kit (Amersham Pharmacia Biotech).

Detection of cell viability and apoptosis assay. The colon carcinoma cell lines were seeded at a density of 1.0×10⁴ cells/well in 96-well flat-bottom microtiter plates (Corning Glass Works, Corning, NY, USA) and were incubated at 37°C in 5% CO₂. After incubation for 24 h, reagents were added, and the plates were incubated for a further 48 h. To assess the viability of the colon carcinoma cells, a 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) assay was performed.
using the Cell Titer 96-assay kit (Promega, Madison, WI, USA) according to the manufacturer's instructions. For analysis of apoptosis, a total of 2x10^4 SW 480 cells were cultured in a chamber slide for 24 h followed by no addition, or by the addition of 1 µg/ml of either Poly I:C or Lipo, or of 1 µg/ml Poly I:C plus Lipo. After incubation for 24 h, cell nuclei were stained with 4',6-diamidino-2-phenylindole (DAPI; Sigma) and the cells were analyzed under a fluorescent microscope.

**NF-κB activity assays.** The NF-κB activity assay was performed using the Dual-Glo™ Luciferase Assay System (Promega) and the pGL4.32 (luc2P/NF-κB-RE/Hygro) Vector (Promega) according to the manufacturer's instructions.

**Results**

**TLR3 is expressed in colon carcinoma cell lines and colon tissues.** As shown in Fig. 1, TLR3 expression was detected by western blotting in both of the colon carcinoma cell lines tested. TLR3 expression in primary human colon tissue was then investigated by immunohistochemical staining of both non-colon carcinoma and colon carcinoma lesions (Fig. 2, representative staining). While there was a clear positive TLR3 signal in 91/100 colon carcinoma cases (91%), no major differences in TLR3 expression were observed between different histological tumor grades (Table I). TLR3 staining was detected not only in the cytoplasm, but also in the cell membrane. Few differences were noted in the TLR3 staining patterns between non-tumor and tumor tissues.

![Figure 2. Immunohistochemical analysis of TLR3 expression in human colon tissues. A representative human colon carcinoma (A) and a non-colon carcinoma (B) tissue specimen were immunohistochemically stained using an anti-TLR3 antibody. A control, human normal sigmoid colon specimen (C) was immunohistochemically stained using an isotype matched mouse IgG. Staining was carried out using a streptavidin-biotin labeling method. TLR3 staining was observed in the cytoplasm as well as in the cell membrane (original magnification, x400).](image)

![Figure 3. Effect of exogenous and endogenous Poly I:C on colon carcinoma cell viability. The indicated colon carcinoma cell lines were incubated without or with different concentrations of Poly I:C in the absence [Lipo (-)] or presence ([Lipo (+)]) of 2.5 µl/ml Lipofectamine LTX for 48 h following which cell viability was assessed using the MTT assay. Cell viability in the absence of treatment was set at 1. Data shown are the means ± SD of six independent experiments.](image)
Transfected Poly I:C-induced apoptosis in colon carcinoma cells. Cellular TLR3 is stimulated by Poly I:C. We therefore next investigated the cytotoxicity of non-transfected or of transfected Poly I:C for these colon carcinoma cells. Stimulation of both cell lines with various concentrations of non-transfected Poly I:C for 48 h resulted in no change in cell viability, whereas stimulation of the cells with transfected Poly I:C for 48 h resulted in a decrease in cell viability in a dose-dependent manner in the SW480 cell line (Fig. 3). We then determined whether transfection of Poly I:C induced apoptosis in these SW480 cells by staining the nuclei with DAPI. Whereas apoptosis was not induced in the cells following treatment with non-transfected Poly I:C, typical apoptotic features of the nuclei were observed in the colon carcinoma cells that were treated with transfected Poly I:C (Fig. 4).

Poly I:C-induced NF-κB activation. Since TLR3 signaling is known to modulate NF-κB signaling pathways we further examined if activation of TLR3 by Poly I:C stimulation mediated induction of NF-κB in the two cell lines. NF-κB activity following cell surface stimulation with Poly I:C was assayed using a luciferase assay and was compared to that in the absence of stimulation. Stimulation of the cells with Poly I:C resulted in increased NF-κB activity in both the SW480 and the Colo320 cell lines compared to the non-stimulated cells (Fig. 5). These results indicated that cell surface stimulation with Poly I:C activates the NF-κB signaling pathway.

TLR9 is expressed in colon carcinoma cell lines and colon tissues. As shown in Fig. 6, TLR9 protein expression was detected in both colon carcinoma cell lines by western blotting. TLR9 expression in colon tissue was further investigated by immunohistochemical staining of both non-colon carcinoma tissue and colon carcinoma lesions (Fig. 7, representative staining). While 86/100 colon carcinoma cases (86%) displayed a clear positive TLR9 signal, no major differences in TLR9 signals were observed between different histological grades of the tumors (Table II). TLR9 staining was detected not only in the cytoplasm, but also in the cell membrane.

CpG-ODNs affect cell proliferation. In order to determine the biological significance of the signaling that occurs via the cell surface TLR9 in colon carcinoma cells, we investigated the cytotoxicity of treatment of the two colon carcinoma cell lines with the TLR9 ligand CpG-ODNs. As shown in Fig. 8,
stimulation of the two cell lines with CpG-ODNs for 48 h increased the viability of SW480 cells. This result suggested that cell surface stimulation with CpG-ODNs might affect cell proliferation and survival in colon carcinoma cells.

CpG-ODNs reduce the cytotoxicity of ADM. We next examined possible interactions between the effect of CpG-ODNs and that of the anticancer reagent, adriamycin (ADM), on the viability of the colon carcinoma cell line Colo320. Cell surface stimulation of Colo320 cells with a combination of 1 µM CpG-ODNs and 2 µg/ml ADM for 48 h resulted in an increase in cell viability of ~23% compared to ADM treatment alone. Furthermore, cell surface stimulation with a combination of 1 µM CpG-ODNs and 5 µg/ml ADM for 48 h resulted in an increase in cell viability of ~12% compared to stimulation with ADM alone (Fig. 9). These results suggest that CpG-ODNs might contribute to a reduction in the cytotoxicity of ADM.

CpG-ODNs activate the NF-κB signaling pathway. Since TLR9 signaling is known to modulate NF-κB signaling pathways we further examined if CpG-ODN stimulation of TLR9 mediated induction of NF-κB activity. Induction of NF-κB activity following cell surface stimulation with 1 µM CpG-ODNs was assayed using a luciferase reporter assay and was compared to the NF-κB activity of non-stimulated cells. NF-κB activity of both colon carcinoma cell lines was increased by treatment with 1 µM CpG-ODNs (Fig. 10).

Discussion

TLRs are essential for immune defense against microbes and viruses. TLR agonists enhance tumor immunotherapy by stimulating TLR signaling in immune cells thereby activating both innate and adaptive immune responses. By contrast, recent studies have indicated that some tumor cells express TLRs and that TLR expression is associated with cancer risk, suggesting that TLRs may also play important roles in tumor biology.

We previously found that human tumor cell lines and tumor tissues, particularly human HCCs, express multiple TLRs (13-16). In the present study, immunohistochemical analysis...
revealed that 91.2% of colon carcinoma tissues as well as two different colon carcinoma cell lines express TLR3 in addition to its expression in non-tumor tissues such as cirrhotic or normal tissues. We previously reported the high prevalence of TLR3 in human HCC. In these previous reports, flow cytometric analysis and immune fluorescence staining indicated that TLR3 was clearly expressed both on the cell surface and in the cytoplasm of HCC cells (15). TLRs 3, 7, 8 and 9, which recognize nucleic acid ligands, are known to be expressed in the endosomes and in the endoplasmic reticulum. In the present study, a similar staining pattern of TLRs, i.e., on the cell surface and in the cytoplasm, was observed in colon carcinoma tissues as was observed in the HCCs of the previous study.

The present study indicated that stimulation of SW480 colon carcinoma cells with non-transfected Poly I:C resulted in no change in cell viability, whereas stimulation of the cells with transfected Poly I:C resulted in decreased cell viability in a dose-dependent manner. These results indicate that activation of intracellular TLR3 may predominate over that of cell surface TLR3 in this cell line.

However, the expression and function of TLR3 in cancer cells are not well understood. Although the findings of the present study indicate that cell surface stimulation of TLR3 did not affect cell viability, cell surface stimulation with Poly I:C did activate the NF-κB signaling pathway. These results suggest that cell surface TLR3 may be functional. In cancer cells, upon activation of NF-κB, the NF-κB dimers typically enter the nucleus and induce the production of cytokines, growth factors and antiapoptotic proteins. It appears that, in tumors, NF-κB can convert inflammatory stimuli into tumor cell survival and growth signals (13-16).

Stimulation of intracellular TLR3 by transfection of the cells with Poly I:C showed that transfected Poly I:C caused apoptotic cell death in colon carcinoma cells in a dose-dependent manner. These findings indicate that TLR3 signals may be linked to apoptotic signals. The combined results indicate that TLR3 signals are different depending on the localization of TLR3 i.e., whether it is on the cell surface or intracellular, and that these signals play an important role in cell survival and cell death. Similar observations have been reported for several other types of cancer cells (13-16).

Regarding TLR9, we found that TLR9 expression, similar to TLR3 expression, was prevalent in human colon carcinoma cells. There was a clear positive TLR9 signal in 86 of 100 colon carcinoma cases (86%). TLR9 staining was detected not only in the cytoplasm, but also in the cell membrane. We previously reported that there was high expression of TLR9 in human HCC. In that previous study, western blot analysis of subcellular fractions, and flow cytometric analysis of intact cells, clearly demonstrated the expression of TLR9 on both the cell surface and in the cytoplasm of human HCC cells (16). In the present study, the staining pattern of TLR3 in colon carcinoma cells was similar to that of TLR9 in the previous study of HCCs.

We then investigated the function of TLR9 in colon carcinoma cells. We demonstrated that stimulation with CpG-ODNs increased the cell viability of SW480 cells. This result suggested that cell surface stimulation with CpG-ODNs might affect cell proliferation and survival in these colon carcinoma cells. This result is consistent with the fact that the enhanced cell proliferation and survival of HCC cells (16). We further found that activation of TLR9 with CpG-ODNs in both colon carcinoma cell lines upregulated NF-κB activity. In general, engagement of the TLR9 signaling pathway leads to the activation of two major transcription factors that have central roles in innate immunity, i.e., NF-κB and IRF-7. TLR9 requires the adaptor molecule MyD88 for initiation of these signals, and MyD88 can directly associate with and activate IRF-7, leading to type I-IFN production (19,20). NF-κB usually plays an important role in regulating immune and inflammatory responses, apoptosis and oncogenes (21,22). Thus the CpG-ODN-activation of NF-κB activity is consistent with other studies of TLR9 signaling.

Finally, we found that cell surface stimulation with CpG-ODNs reduced the cytotoxicity of ADM. We previously showed that cell surface stimulation with CpG-ODNs might contribute to a reduction in the cytotoxicity of ADM towards HepG2 cells via the upregulation of apoptosis inhibitors (16). We therefore speculated that the reduction in ADM cytotoxicity in colon carcinoma cells by CpG-ODNs may be mediated by the same mechanism.

In conclusion, functional TLR3 and TLR9 are expressed in colon carcinoma cells. TLR3 and TLR9 activate NF-κB and the activation of these TLRs is closely related to cell death and survival. Further evaluation of the possible roles and regulation of TLR3 and TLR9 are critical for controlling cell death, proliferation and immune escape of malignant cells.

References


