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Toll-like receptor-4 (TLR4) promotes the development of colitisassociated colorectal tumors

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Abstract

Background and Aim— Chronic inflammation is a risk factor for colon cancer in patients with ulcerative colitis (UC). The molecular mechanisms linking inflammation and colon carcinogenesis are incompletely understood. We tested the hypothesis that TLR4 is involved in tumorigenesis in the setting of chronic inflammation.

Methods—Tissues from UC patients with cancer were examined for TLR4 expression. Colitisassociated neoplasia was induced using azoxymethane (AOM) injection followed by dextran sodium sulfate (DSS) treatment in TLR4-deficient (TLR4–/–) or wild-type (WT) mice. Inflammation, polyps, and microscopic dysplasia were scored. Cox-2 and PGE₂ production were analyzed by realtime PCR, immunohistochemistry, or enzyme immunoassay. EGFR phosphorylation and amphiregulin production were examined by Western blot analysis and ELISA, respectively.

Results— We show that TLR4 is overexpressed in human and murine inflammation-associated colorectal neoplasia. TLR4–/– mice were markedly protected from colon carcinogenesis. Mechanistically, we show that TLR4 is responsible for induction of Cox-2, increased PGE₂ production and activation of EGFR signaling in chronic colitis. Amphiregulin, an EGFR ligand, was induced in a TLR4, COX-2-dependent fashion and contributes to activation of EGFR phosphorylation in colonic epithelial cells.

Conclusion—TLR4 signaling is critical for colon carcinogenesis in chronic colitis. TLR4 activation appears to promote the development of colitis-associated cancer by mechanisms including enhanced Cox-2 expression and increased EGFR signaling. Inhibiting TLR4 signaling may be useful in prevention or treatment of colitis-associated cancer.

The authors have no conflicts of interest to disclose.

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Introduction

Inflammation is considered a risk factor for many common malignancies including cancers of the lung ¹, breast ², and colon ³. Colon cancer is the third most common cancer and the third leading cause of cancer-related mortality in the United States ⁴. The link between inflammation and colon cancer offers the possibility of identifying novel ways to prevent cancer. However, the molecular mechanisms whereby chronic inflammation predisposes to cancer remain elusive.

The clearest link between inflammation and colon cancer is seen in patients with inflammatory bowel disease (IBD)⁵. Colorectal cancer is one of the most serious complications of IBD, accounting for increased mortality in these disorders ⁶. The severity of inflammation correlates with the risk of colorectal cancer in patients with IBD ⁷, ⁸. Consistent with a role of inflammation in colorectal neoplasia, animal models have demonstrated that the multifunctional transcription factor NF- κ B is required for colorectal neoplasia ⁹. Therefore, focusing on the relationship between chronic inflammation and carcinogenesis may provide insights into the pathogenesis of colitis-associated cancer (CAC) and possibly sporadic colorectal cancer.

The colon contains 100 trillion bacteria, and commensal bacteria have been implicated in the development of sporadic colorectal cancer ¹⁰. Commensal bacteria may promote colorectal cancer by a variety of mechanisms including generation of reactive oxygen intermediates resulting in chromosomal instability ¹¹. Bacteria are required for eliciting chronic inflammation and colon cancer in animal models of CAC ^{12, 13}. Therefore, we have focused on TLR4 and its role in CAC. TLR4 is normally expressed at low levels in the intestinal mucosa ^{14–16} while it is up-regulated in patients with IBD ¹⁷. In acute colitis, we previously demonstrated that TLR4 is a potent inducer of cyclooxygenase-2 (Cox-2) expression ¹⁸. TLR4 may also be important for evasion of tumor surveillance ¹⁹. Altogether, these data raise the intriguing possibility that TLR4 promotes colon cancer in the setting of chronic inflammation, and this serves as the focus of the present study.

In the current study, we demonstrate first that TLR4 is over-expressed in human colon cancers arising in chronic ulcerative colitis. Over-expression of TLR4 was then confirmed in an animal model of inflammation-induced colon tumorigenesis wherein healthy, wild-type (WT) mice were given azoxymethane (AOM) to induce colonic neoplasia in the setting of chronic inflammation. Importantly, mice genetically lacking TLR4 were protected against colon tumorigenesis in this animal model of inflammation-induced carcinogenesis. Mechanistically, we show that TLR4 is required for expression of Cox-2 and enhanced PGE₂ production in chronic colitis. TLR4-dependent tumorigenesis was also associated with activation of EGFR signaling. These findings are significant because both Cox-2 and EGFR have been linked to the development of colon tumors ²⁰, ²¹. The current results provide important insights into the previously unrecognized role of TLR4 signaling in CAC and strengthen the rationale for developing chemopreventive therapies that target TLR4.

Materials and Methods

Animal studies

TLR4^{-/-} mice were purchased from Oriental Bio Service, Inc. (Kyoto, Japan). All knockout mice were backcrossed to C57BI/6J mice at least 8 generations. C57BL/6J mice were obtained from Jackson Laboratory as controls (Jackson Laboratory, Bar Harbor, Maine). Mice were kept in specific-pathogen free (SPF) conditions and fed by free access to a standard diet and water. All experiments were done according to Mount Sinai School of Medicine animal experimental ethics committee guidelines.

Following previously established methods for inducing colonic neoplasia ²², six to ten week old gender-matched mice were injected with 7.4mg/kg of AOM (Sigma, St. Louis, MO) intraperitoneally (i.p.) at the beginning of the experiment (day 0). After 14 days, mice were treated with 3% DSS (MW 36–50 kDa: ICN, Aurora, Ohio) in their drinking water for 7 days. This was followed by 14 days of normal water, another 7 days of 3% DSS treatment, and then normal water for an additional 14 days. During the DSS treatment and recovery phase, body weights, stool consistency, and stool occult blood were monitored, as described previously 23.

Mice were sacrificed on day 56. Colons were removed, opened longitudinally and stained with 1% Alcian Blue (Sigma, St. Louis, MO). High-resolution mucosal pictures were obtained using a Leica MZ APO dissecting microscope (Leica, Bannockburn, IL) with SONY Power HAD DXC-970MD 3CCD Color digital video camera, and Scion Image 1.60 image software (Scion Co., Frederick, MD). After macroscopic assessment, cecum, proximal, and distal parts of the colon were fixed in 10% buffered formalin, embedded in paraffin, sectioned and stained with hematoxylin and eosin. Histological assessment was performed by two independent gastrointestinal pathologists (R.X., H.C.) blinded to the mouse genotype and treatment. Severity of mucosal inflammation was graded using a standard scoring system (Table 1). Dysplastic lesions were determined by previously established criteria ²⁴. To quantify the microscopic extent of dysplasia, paraffin-embedded colons were cut in 5 µm thick serial sections and every 20th section was analyzed for dysplasia. Number, size, and the percentage of the mucosal surface area containing dysplasia were determined under the microscope. The size of the lesions was calculated using a scale micrometer on the microscope.

Cell lines and reagents

The human colon cell line SW480 (1×10^6 cells/well) was maintained in Dulbecco's modified Eagle's medium supplemented with 2% heat-inactivated FCS, 2mM L-glutamine, penicillin/ streptomycin and was incubated in 6-well plates overnight at 37 °C in a 5% CO₂ humidified incubator. The next day, cells were incubated with Ultrapure lipopolysaccharide (LPS, 2µg/ml), *Eschericha coli* 0111: B4 (Invivogen, San Diego, CA) for varying time periods.

Real Time PCR

Total RNA was isolated using RNA Bee (Tel-Test, Inc., Friendwood, TX) according to the manufacturer's instructions. A total of 1 μ g RNA was used as the template for single strand cDNA synthesis utilizing the Transcriptor First Strand cDNA Synthesis Kit (Roche, Indianapolis, IN) according to the manufacturer's instructions. Quantitative real-time PCR was performed for Cox-2, and β -actin using TaqMan probes. The primers and probes used in this study can be found on-line in Supplemental material. The cDNA was amplified using TaqMan universal PCR Master Mix (Roche, Indianapolis, IN) on an ABI Prism 7900HT sequence detection system (Applied Biosystems, Foster City, CA), programmed for 95°C for 10 minutes, then 40 cycles of: 95 °C for 15 seconds, 60 °C for 1 minute. The amplification results were analyzed using SDS 2.2.1 software (Applied Biosystems, Foster City, CA) and the gene of interest was normalized to the corresponding β -actin results. Data were expressed as fold induction relative to the lowest gene product amplified.

RNA interference (RNAi)

SW480 cells were plated at a density of 1.5×10^5 cells/well in 12-well plates 24hours before the first transfection. TLR4 small interfering RNA (siRNA) oligonucleotide corresponding to the sequence (GGUAAGGAAUGAGCUAGUAUU) was purchased from Dharmacon (Chicago, IL). Fifty nM of siRNA were transfected twice every 24 hours with X-trim gene siRNA transfection reagent (Roche, Indianapolis, IN) as per the manufacturer's instructions. Forty-eight hours after the first transfection, cells were stimulated with LPS (2µg/ml) for the

indicated period of time. Negative control siRNA (50 nM), which has no significant homology to any known gene sequences from mouse, rat, or human, was used as a control (Ambion, Austin, TX).

Western Blot Analysis

De-identified human colon resections were obtained under the auspices of the Mount Sinai Medical Center Institutional Review Board. Snap frozen tumor and surrounding colonic tissues from six ulcerative colitis patients, and actively inflamed colonic tissues from four ulcerative colitis patients undergoing colectomy were used. Mouse colon samples were taken at the time of sacrifice and frozen at -80° C. Details of tissue preparation and antibodies used for Western blot are included in Supplemental material. The mouse TLR4 band intensity was calculated using NIH image 1.62 by normalizing with the intensity of the corresponding β -actin band.

Enzyme linked immunosorbent assay (ELISA)

For the amphiregulin ELISA, SW480 (1×10^6 cells/well) were plated in 6 well plates. Cells were treated with LPS (2µg/ml) for the indicated periods. For *ex vivo* colonic tissue cultures, 100 mg of tissue from each part of the colon (not including the polypoid lesions) were cultured for 24 hours in 12 well flat bottom plates in serum free RPMI 1640. Supernatants were harvested for measurement of amphiregulin or TNF- α . ELISA (R&D Systems, Minneapolis, MN) was performed per the manufacturer's instructions.

Immunofluorescent and immunohistochemical studies

Paraffin-embedded human colectomy specimens of normal colon (n=11) (obtained from resection edge of sporadic colorectal cancers) or colitis-associated dysplasia or cancer (n=15; all patients with ulcerative colitis) were stained with biotinylated mouse monoclonal antihuman TLR4 (1:500, eBioscience, San Diego, CA) overnight at 4°C, followed by streptavidin-FITC (10 μ g/ml, eBioscience, San Diego, CA) for 1 hour at room temperature. Details of staining methodology and antibodies used are found in Supplemental material.

Assessment of proliferation

Colonic tissue sections were examined for cell proliferation (bromodeoxyuridine: BrdU labeling) in the polypoid lesions as well as surrounding colonic mucosa. Mice were injected with 120 mg/kg of BrdU (Sigma, St. Louis, MO) i.p., 90 minutes prior to sacrificing, and colonic tissues was stained for BrdU using a BrdU staining kit (Zymed Laboratories Inc, South San Francisco, CA) according to the manufacturer's instructions. The number of BrdU-positive cells per well-oriented crypt were calculated in every 3 crypts for each colon segment at high magnification under light microscopy.

NF-kB Assay

For measurement of NF- κ B activation, nuclear extracts were isolated from snap frozen mouse colonic tissues using a nuclear extraction kit (Active Motif, Carlsbad, CA) according to the manufacturer's instructions. After measurement of protein concentration by the Bradford method, 1µg of nuclear extract was used to measure the DNA-binding activity of NF- κ B p65 using TransAM NF- κ B p65 Chemiluminescent kit (Active Motif, Carlsbad, CA) according to the manufacturer's instructions. Chemiluminescent intensities were calculated as relative light units (RLU) and normalized with the mean RLU from untreated animals.

Measurement of PGE₂

Production of PGE₂ in the tissue culture supernatant was determined using a monoclonal EIA kit (Cayman, Ann Arbor, MI) according to the manufacturer's instructions and as described

previously ^{18, 25}. Briefly, colonic samples from TLR4–/– and WT mice were washed in cold PBS containing penicillin, streptomycin, and fungizone (100U/ml each). 100 mg tissue fragments from the distal part of the colon closest to the anus were cultured for 24 hours in 12 well flat bottom plates in serum free RPMI 1640. Culture supernatants were harvested for PGE₂ measurement.

Statistical analysis

Data were presented as mean (\pm SD). The significance was analyzed by Student t-test and Fisher's exact test with Microsoft Excel and StatView software, respectively. *P* values were considered significant when < 0.05.

Results

TLR4 expression is up-regulated in colitis-associated tumors

TLR4 expression is known to be low in the normal colon but increased in IBD ¹⁶, ^{26–28}. We hypothesized that TLR4 is up-regulated in colitis-associated tumors. Colon tumor specimens from patients with UC with dysplasia or cancer were examined for TLR4 expression by Western blot analysis and immunofluorescent staining (Figure 1A, B). Non-dysplastic colon tissue from the same UC patients was used as a control. In addition, we examined samples from patients with active UC undergoing colectomy to assess the effect of moderate to severe inflammation on expression of TLR4. Non-dysplastic colon had low expression of TLR4 protein whereas samples of protein from the matched tumor specimens demonstrated higher expression of TLR4 (Figure 1A). Colon samples from active UC demonstrated expression of TLR4 comparable to the affected, non-dysplastic colon in UC cancer patients (Figure 1A). Immunostaining showed that, in normal colon, only cells at the base of the crypt and scattered cells in the lamina propria express TLR4 (Figure 1B). Colitis-associated tumors showed predominantly epithelial staining of TLR4 with varying degrees of TLR4 expression in cells of the lamina propria.

To further examine the role of TLR4 in colitis-associated tumorigenesis, we used a well-established model of CAC ^{24, 29}. In this model, animals receive AOM, a genotoxic agent, followed by two cycles of DSS to induce inflammation. Using the AOM-DSS model, all WT mice developed polypoid or flat tumors. The microscopic features recapitulate the features of human dysplasia and polyps in colitis patients ³⁰. We hypothesized that TLR4 expression would be increased in colorectal neoplasia in this CAC model. To investigate this possibility, we first examined mRNA expression of various TLRs in tumor tissues and surrounding nondysplastic mucosa by real-time PCR (Table 2). Compared to the untreated colon, all TLRs examined demonstrated increased expression with inflammation. Of the TLRs tested, however, only TLR4 was increased in the tumor tissue compared with the surrounding inflamed mucosa. We then examined TLR4 mRNA expression under various conditions, e.g. untreated, acute DSS, chronic DSS (at day 56), AOM alone, AOM+DSS at day 56 (Figure 1C). Although TLR4 mRNA expression is increased following acute or chronic DSS compared with untreated mice, the highest level of expression was seen in tumors caused by AOM+DSS. Expression of TLR4 protein by Western blot was significantly elevated in tumors that developed in WT mice compared to the surrounding non-dysplastic mucosa (Figure 1D). Immunofluorescent staining for TLR4 also showed an increase in TLR4 expression in polypoid tumors (Figure 1E). Little TLR4 staining was seen in the normal epithelium or in the epithelium in chronic colitis. Muscularis propria stains for TLR4 consistent with a previous report 31 . We conclude that colorectal tumors over-express TLR4 and that TLR4 may play a role in tumorigenesis.

TLR4 promotes development of colorectal tumors in the setting of chronic colitis

To examine the role of TLR4 in colitis-associated tumorigenesis, we first addressed whether absence of TLR4 altered the susceptibility to developing CAC. TLR4–/– and WT mice were treated with AOM-DSS as described in Materials and Methods and developed weight loss and bleeding during DSS treatment (Figure 2A, B). We found a striking difference between WT and TLR4–/– mice with respect to development of polyps and dysplasia. All WT mice grossly showed multiple polypoid lesions but no such visible lesions were seen in TLR4–/– mice (Figure 2C). When examined microscopically, all WT mice (n=18) had at least one dysplastic lesion and up to 17 lesions per mouse (Figure 2C, D). In contrast, only five of eighteen TLR4–/– mice (27.8%) had one or two dysplastic lesions (Figure 2C, D). A significant increase in the number of dysplastic lesions was observed between WT and TLR4–/– mice (P < 0.001) (Figure 2D, Table 3).

The size and severity of dysplasia were also different between WT mice and TLR4–/– mice (Table 3). The few dysplastic lesions found in TLR4–/– mice were small, flat, and low grade. By contrast, 48% of dysplastic lesions (47/97) in WT mice were polypoid. BrdU staining of colon showed that proliferation was greatly increased in dysplasia and the surrounding epithelium of WT mice compared to non-dysplastic mucosa in TLR4–/– mice (Figure 2E). The number of BrdU positive cells was significantly higher in WT dysplastic (33.4 positive cells/100 epithelial cells \pm 8.3SD) and non-dysplastic mucosa (20.4 cells \pm 6.2SD) compared with the non-dysplastic mucosa in TLR4–/– mice (14.3 cells \pm 3.8SD). These results demonstrate that TLR4 is required for initiation and growth of colorectal tumors in a mouse model of CAC.

NF-KB activation is decreased in acute but not chronic DSS colitis in TLR4-/- mice

Having demonstrated the link between TLR4 and colorectal tumors, we turned our attention to the mechanism by which this may occur. The severity of inflammation correlates with the risk of colorectal cancer in patients with IBD ⁷, ⁸. Using a scoring system for chronic colitis that examines features seen in human colitis, there was no significant difference between TLR4 -/- and WT mice with respect to overall severity of colitis (Figure 3). A common connection between inflammation and carcinogenesis is the activation of signaling pathways that promote cell proliferation and increase cell survival such as the transcription factor NF- κ B ⁹. TNF- α is a pro-inflammation and carcinogenesis ³². Therefore, we examined transcriptional activity of NF- κ B and TNF- α production in the colonic tissues from the CAC model. There were no significant differences in either NF- κ B activation or TNF- α production between WT and TLR4 -/- mice in the CAC model (Figure 3).

We reasoned, however, that inflammation at earlier time points might be different between WT and TLR4–/– mice. We have previously shown that following acute DSS treatment, TLR4 –/– mice have significantly decreased recruitment of neutrophils ²³. Although the total inflammatory scores following acute DSS treatment were similar between WT and TLR4–/– mice, there was a small but significant decrease in the sub-score for acute inflammation in TLR4–/– mice compared with WT mice (Figure 3B). Thus, we examined NF- κ B activation and TNF- α production after seven days of DSS treatment. WT mice had significantly higher activity of NF- κ B and increased production of TNF- α compared with TLR4–/– mice in the acute phase of colitis (Figure 3B). These results indicate that the decrease in inflammatory mediators such as NF- κ B activation and TNF- α production in TLR4–/– mice may contribute to the subsequent protection from colorectal dysplasia. These data also highlight that in spite of expression of many other TLRs during inflammation (Table 2), the absence of TLR4 results in a decreased ability to mount a robust pro-inflammatory response.

Mucosal Cox-2 expression and PGE₂ production are decreased in TLR4-/- mice in the CAC model

Cox-2 is critical for development of colorectal neoplasia ²⁰. We hypothesized that decreased expression of Cox-2 in TLR4–/– mice was associated with protection against tumorigenesis. Using real-time PCR, we found that mucosal Cox-2 expression was significantly decreased in TLR4–/– mucosa compared to WT mucosa in the CAC model (Figure 4A). Mucosal Cox-2 expression was found in lamina propria cells and to a lesser extent in IEC by immunofluorescent staining (Figure 4B). We confirmed by double staining that the majority of Cox-2 positive lamina propria cells were also positive for the macrophage marker CD68 (Figure 4C). We also performed double-staining of Cox-2 and a marker of myofibroblasts and found Cox-2-expressing myofibroblasts at the base of the crypts (Figure 4C).

Given that TLR4–/– mice have decreased expression of mucosal Cox-2 compared to WT mice, we investigated whether PGE₂ production was also reduced. PGE₂ production is important for both the formation and growth of colorectal tumors ³³. We found that PGE₂ production by colonic tissue from TLR4–/– mice was significantly less than in WT mice (Figure 4D). Given the well-established link between Cox-2 and colon carcinogenesis ²⁰, these data suggest that TLR4 signaling may promote the development of CAC, in part, by inducing Cox-2 expression and PGE₂ production.

Phosphorylation of EGFR is decreased in TLR4-/- mucosa

Several studies have suggested that PGE_2 stimulates colorectal carcinogenesis in part by activation of EGFR signaling ³⁴, ³⁵. We reasoned that TLR4–/– mice might be protected against colorectal carcinogenesis in chronic colitis because of decreased EGFR activation. Consistent with this hypothesis, levels of phosphorylated EGFR were significantly lower in colonic mucosa from TLR4–/– mice compared with WT mice (Figure 5A).

Additional studies were carried out to elucidate the mechanism by which activation of TLR4 stimulates EGFR tyrosine kinase activity. Amphiregulin, an EGFR ligand, is believed to play a role in colon carcinogenesis ³⁶ and can induce COX-2 ³⁷. Hence, experiments were carried out to determine whether amphiregulin plays a role in TLR4-mediated activation of EGFR. Treatment with LPS caused a several-fold increase in production of amphiregulin in SW480 cells (Figure 5B). In contrast to nonspecific siRNA, LPS-mediated induction of amphiregulin was abrogated by siRNA to TLR4 (Figure 5C). We next addressed whether ligand binding was involved in LPS-mediated stimulation of EGFR tyrosine kinase activity. We found that antibody blockade of the EGFR ligand-binding site as well as a neutralizing antibody to amphiregulin abrogated LPS-mediated activation of EGFR (Figure 5D).

Given the compelling evidence that amphiregulin links the TLR4 pathway with activation of EGFR in human colonocytes, we hypothesized that TLR4–/– mice have reduced levels of amphiregulin in the colon. Mucosal amphiregulin release measured by ELISA was also significantly reduced in TLR4–/– mice compared with WT mice in the CAC model (Figure 5E). Collectively, these results strongly suggest that the reduction in EGFR tyrosine kinase activity in colonic mucosa from TLR4–/– mice (Figure 5A) is likely to be explained at least, in part, by reduced production of amphiregulin.

Discussion

The present study establishes a molecular link between TLR4 and colon cancer in the setting of chronic inflammation (Figure 6). We demonstrate for the first time that TLR4 is highly expressed in colon cancers from patients with long-standing ulcerative colitis and in colonic tumors in a murine model of CAC. Our studies show a critical role for TLR4 in development

of colitis-associated dysplasia since, in the absence of TLR4, colonic polyps do not occur. The ligand for TLR4, LPS, is abundant in the colonic lumen raising the intriguing possibility that colonic bacteria induce growth of colonic tumors through activation of TLR4. Chronic infection or chronic inflammation resulting in TLR4 activation may occur in other common malignancies, such as Helicobacter pylori and gastric cancer ³⁸. In addition, hyaluronan, a non-bacterial TLR4 ligand present in chronically inflamed tissues, may activate TLR4 and contribute to carcinogenesis ³⁹. Our results highlight the possibility of interfering with the host-microbial innate immune response to prevent or treat cancer.

NF- κ B activation in both colonocytes and macrophages has been shown to be important in development of CAC and supports the connection between inflammation and colon cancer ⁹. In Greten *et al.*'s study examining the contribution of epithelial versus macrophage-dependent activation of NF- κ B on colonic tumorigenesis, epithelial NF- κ B activation was necessary for high tumor numbers whereas macrophage-dependent NF- κ B activation contributed to tumor size. NF- κ B can be activated by many upstream stimuli including TLRs. Our studies suggest that TLR4 may be a principal receptor, upstream of NF- κ B, promoting the development of colonic tumors especially during the acute inflammatory stage. In our model, TLR4 expression is found in both epithelial cells as well as lamina propria mononuclear cells in neoplastic areas. *In vitro*, TLR4 can directly induce proliferation of IEC through EGFR activation ¹⁸. *In vivo*, there are likely to be multiple pathways leading to dysregulated epithelial proliferation in response to TLR4.

The intestinal epithelium is in intimate contact with the underlying lamina propria containing macrophages and subepithelial myofibroblasts; these cells have been shown to express TLR4 ^{15, 16}. Tumor-associated macrophages play a clear role in supporting tumor growth in a variety of systems ⁴⁰. Some of the effects on the intestinal epithelium may be mediated by TLR4dependent signals derived from the stroma. We show that Cox-2 expression and PGE₂ levels are markedly different between WT versus TLR4-/- colons. Immunostaining demonstrates expression of Cox-2 predominantly in tumor-associated macrophages and myofibroblasts, but we know from our previous work that IEC also express Cox-2 in a TLR4-dependent way following acute injury 18 . The finding of increased PGE₂ production in our model is consistent with previous evidence of increased levels of PGE_2 and deregulated expression of enzymes involved in the synthesis and catabolism of PGs in inflamed human colonic mucosa and colon cancers ^{41, 42}. Recent work by Brown et al. demonstrates that mesenchymal stromal cells express Cox-2 and migrate to colonic epithelial progenitors during injury in a MyD88dependent fashion ⁴³. Without the migration of PGE₂-producing stromal cells, epithelial proliferation is impaired. In addition, recent work demonstrates that bacteria can produce reactive oxygen intermediates that induce Cox-2 expression in macrophages leading, in turn, to chromosomal instability ¹¹. These data highlight the interplay of bacteria and host cells in promoting colorectal cancer.

The reciprocal interaction between the epithelium and the stroma is complex. PGE_2 can stimulate the release of EGFR ligands, such as amphiregulin ³⁶. Thus, bioactive lipids produced by the stroma can amplify IEC proliferation through induction of trophic growth factors. At least *in vitro*, we show that amphiregulin is the principal contributor to EGFR activation. We have not shown, however, that the decrease in amphiregulin in TLR4–/– mice is the reason that EGFR activation is decreased. Likewise, EGFR signaling has been shown previously to increase Cox-2 expression and PGE₂ secretion resulting in a positive feedback loop that contributes to deregulated cell proliferation⁴⁴. More work is necessary to understand whether a positive feedback loop between EGFR and Cox-2 is at play in our model. In summary, several well-established mediators of human colon carcinogenesis, namely Cox-2, PGE₂, and the EGFR, are modulated by TLR4 signaling.

For our studies, we have chosen the AOM-DSS model of CAC 24 . AOM is a colonic genotoxic carcinogen that is extensively used for the investigation of colorectal carcinogenesis in rodents. Although by itself, AOM does not cause dysplasia or cancer in C57/BL6 mice, in combination with repeated cycles of DSS, it increases the incidence of dysplastic lesions 29 . In particular, this model has been used to interrogate the inflammation-cancer link in the colon 9 . Although no single animal model suitably reproduces all the features of human CAC, increased Cox-2 expression and enhanced PGE₂ levels characterize this model and are reminiscent of human CAC 45 .

Another question raised by our work is whether the effect of TLR4 is unique or whether other TLRs can contribute to development of malignancy. The colonic microbiota is diverse and should activate many TLRs. It is possible that other TLRs have the potential to induce dysplasia but that their contribution is weaker than TLR4. The prediction would be that the absence of *all* TLR signaling would also prevent colon cancer. We have evaluated whether CAC could be induced in MyD88–/– mice and all 15 mice succumbed from colitis following the first cycle of DSS (data not shown). MyD88–/– mice are fragile and may die as a result of sepsis, hemorrhage, or both, at least in part because MyD88 and TLR signaling are necessary for repair of the intestinal epithelium. It also suggests that targeting MyD88 as a therapeutic or chemopreventive intervention is likely to be impractical.

In summary, our results point to a critical role of TLR4 in development of colon cancer. Based on this study, we postulate that targeted inhibition of TLR4 may be effective in preventing development of colon cancer in IBD. TLR4 inhibitors are currently being evaluated in clinical trials of sepsis ⁴⁶. One of the issues with the use of TLR4 antagonists is that our previous work suggests that during acute injury, TLR4 signaling is beneficial. Perhaps the value of TLR4 antagonists would be in patients with quiescent colitis with dysplasia or at high risk for dysplasia. Ideally, TLR4 signaling would be dampened but not eliminated. Cox-2 inhibitors are effective in reducing the risk of adenomatous polyps but unfortunately long-term use is associated with increased cardiovascular complications ^{47, 48}. Therefore, new approaches to preventing colon carcinogenesis are required. Our studies will permit rational therapies to be developed that combine blockade of TLR4 and other signaling pathways, such as EGFR signaling, in the prevention or treatment of CAC.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

TLR	toll-like receptor
WT	wild-type
DSS	dextran sodium sulfate
PAMP	pathogen associated molecular pattern
LPS	lipopolysaccharide
Cox	cyclooxygenase
EGFR	epidermal growth factor receptor
PG	

prostaglandin

IEC

intestinal epithelial cells





A. Expression of TLR4 in normal mucosa and tumors from patients with UC. Samples of tumor tissue (n=4) and corresponding surrounding non-dysplastic colon (n=4) were examined for TLR4 protein by Western blot analysis. Colonic tissues from patients with active ulcerative colitis were also examined (Inflamed) (n=4). 25 μ g/lane of protein was loaded per lane. β -actin staining is used as a loading control.

B. Immunofluorescent staining for TLR4 in human colitis-associated cancer. Normal human colon tissue (2 representative examples of 11 patients) (Top panels) and colon cancer in patients with ulcerative colitis (2 representative examples of 15 patients) (middle panels) were stained for TLR4.

C. TLR4 mRNA expression in the AOM-DSS model. Normal colon (n=10), acute phase after 7 days of DSS (n=10), day 56 after injection of AOM (n=5), day 56 of chronic DSS colitis after two cycles of DSS (n=5), non-dysplastic mucosa in AOM+DSS treated mice at day 56 (n=5), and tumor tissues taken from AOM+DSS treated mice at day 56 were analyzed for TLR4 mRNA expression by real-time PCR. P < 0.05 for tumor tissue versus non-dysplastic mucosa, non-dysplastic mucosa versus normal colon, and chronic DSS versus normal mucosa.

D. Western blot analysis for TLR4 in murine colon samples. Immunoblots of tissue lysates (25 μ g/lane) prepared from colonic samples of tumor versus corresponding surrounding nondysplastic mucosa (n=4). Graph shows ratios of TLR4 band intensity normalized with β -actin band (ND= Non-dysplastic mucosa, T= Tumor). Mean ratios of non-dysplastic mucosa and tumor are shown at the end (**P*<0.0001).

E. Immunofluorescent staining for TLR4 in the mouse CAC model. Serial sections of tumors or surrounding mucosa are stained with H&E for histology and immunofluorescent staining for TLR4. Mesenteric lymph node (MLN) is used as a positive control; mesenteric lymph node from a TLR4–/– mouse is used as a negative control.



Figure 2. TLR4–/– mice are protected from colonic tumors in the setting of inflammation A, B. AOM and DSS were administered as shown. Weight change (A) and stool blood (B) were examined during the study period. The data represent the average (\pm SD) of four independent experiments with a total of 36 mice (TLR4–/– (n=18), and WT controls (n=18)) (*p<0.05).

C. Tumors in the CAC model. Colons were cut open longitudinally and the mucosal surface was stained with 1% Alcian blue. All WT mice showed multiple polypoid lesions (**a**, **b**) but not the TLR4–/– mice (**f**, **g**). Microscopically, two examples of WT dysplasias are shown at lower (**c**) and higher (**d**) magnifications. Non-dysplastic mucosa from a WT mouse is shown in **e**. Most TLR4–/– mucosa following AOM-DSS appeared as in panel **h**. An example of a dysplastic lesion in a TLR4–/– mouse is shown at lower (**i**) and higher (**j**) magnifications. **D**. Incidence of dysplasia. The number of dysplastic lesions was counted per mouse under the microscope. Data are expressed as mean \pm SEM (*P* < 0.0001).

E. Proliferation in the CAC model. BrDU incorporation was assessed using anti-BrDU Ab staining. Representative pictures are taken from a WT dysplastic lesion (a), and WT non-dysplastic mucosa (b). Mucosa of AOM-DSS treated TLR4–/– mice (c) is also shown.



Figure 3. Comparison of inflammatory activity in TLR4-/- versus WT mice

A. Histology scores in WT versus TLR4–/– mice at day 56 following AOM+DSS (left panel). Table 1 demonstrates the criteria used for scoring. The total histology score was similar in WT and TLR4–/– mice. NF- κ B activation state (middle panel) and TNF- α secretion (right panel) in WT versus TLR4–/– mice at day 56 following AOM+DSS. There are no significant differences between WT and TLR4–/– mice in NF- κ B activation or TNF- α production in the intestine.

B. Histology scores in WT versus TLR4–/– mice at day 7 following DSS. The acute and chronic sub-scores for the indices are shown. NF- κ B activation state and TNF- α secretion in WT versus TLR4–/– mice after seven days of DSS treatment. TLR4–/– mice have significantly decreased NF- κ B activation and TNF- α secretion.

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Figure 4. Mucosal Cox-2 expression is decreased in TLR4-/- mice in the CAC model

A. TaqMan real-time PCR was used to compare mucosal expression of Cox-2 mRNA in WT and TLR4–/– mice in the CAC model (n=6 each). Data are represented as mean \pm SD of relative values of expression (*P < 0.05).

B. Immunofluorescent staining for Cox-2 in the colon. Representative pictures show Cox-2 positive cells (green - FITC) in lamina propria cells (arrowhead) and IEC (arrow) in WT mice and TLR4–/– mice.

C. Double immunostaining for Cox-2 (green - FITC) and the macrophage marker CD68 (red - TRITC) in WT mucosa (top panel) or anti-fibroblast antibodies (red-TRITC) (bottom panel). Arrows indicate double positive cells (right most panels).

D. PGE₂ production by colonic tissue from TLR4–/– or WT mice in the CAC model. Colonic tissues from TLR4–/– (n=9) and WT mice (n=10). Data are shown as mean \pm SD. (*P < 0.05).



Figure 5. TLR4 signal regulates EGFR tyrosine phosphorylation and amphiregulin expression in colonocytes

A. Western blot analysis of phosphorylated EGFR and EGFR in the colon. Results from three representative samples obtained from WT or TLR4–/– mice are shown. 25µg/lane of protein was loaded per lane. The membrane was sequentially probed for phospho-EGFR and EGFR. Beta-actin was used to demonstrate equal protein loading.

B. LPS induces the release of amphiregulin protein. SW480 cells were treated with vehicle (control) or LPS (2 μ g/ml) for the indicated periods. Amphiregulin concentration in

supernatants was measured by ELISA. Data are expressed as mean \pm SD of relative values of expression in three individual experiments with triplicate samples (*P < 0.05).

C. LPS-mediated induction of amphiregulin mRNA is TLR4-dependent. SW480 cells were transiently transfected with control siRNA or siRNA against TLR4 and then stimulated with LPS ($2\mu g/ml$) for the indicated periods of time. Untransfected control samples were not LPS

treated. Levels of amphiregulin mRNA were determined by real time PCR. The data are represented as mean \pm SD of relative values of expression in three individual experiments with triplicate samples (*P < 0.05). Inset: Western blot shows TLR4 expression with medium alone, control siRNA, and TLR4 siRNA in SW480 cells. The last bar is THP-1 cells as a positive control.

D. LPS-mediated activation of EGFR is ligand-dependent. SW480 cells were pre-treated with antibodies to the ligand-binding site of EGFR, anti-amphiregulin antibody or control IgG for two hours. Subsequently, cells were stimulated with LPS ($2\mu g/ml$) for 30 minutes. Blots of whole cell lysates ($25 \mu g/lane$) were sequentially probed for phospho-EGFR and EGFR. The data are one representative experiment of three with similar results. β -actin was used as an internal control for protein loading.

E. TLR4 deficiency is associated with reduced colonic mucosal production of amphiregulin. The production of amphiregulin was measured in colonic mucosa from WT (n=10) and TLR4 -/- (n=8) mice. Data are expressed as mean \pm SD (*P < 0.05).



Figure 6. Model of TLR4-mediated colon carcinogenesis

TLR4 expression is increased in chronic intestinal inflammation. TLR4 signaling in response to LPS induces Cox-2 expression and PGE_2 production. PGE_2 through its receptors (EP) can act in a paracrine or autocrine fashion on colonocytes to stimulate the expression and release of amphiregulin, an EGFR ligand. EGFR signaling is associated with increased proliferation of colonocytes. Likewise, TLR4 expression in tumor-associated macrophages may also respond to LPS by inducing Cox-2 and PGE₂, which may then act on the epithelium to stimulate proliferation of colonocytes.

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Scoring system for of	chronic mucosal inf	flammation				
Histological scoring*						
1	2	3	4	5	6	7
Crypt damage	Acute	Edema	Erosions/	Chronic inflammatory index	Epithelial regeneration	Distortion/
	inflammatory		ulceration	(mononuclear cell infiltration)		branching
	index (neutrophils/					
	eosinophils)					
intact = 0	0-4	0-4	0-4	normal distribution $= 0$	no tissue repair $= 3$	normal crypt= 0
basal $1/3 = 1$				intermediate $=1$	Surface epithelium not intact =2	mild = 1
basal $2/3 = 2$				full thickness of	Slight epithelial injury $= 1$	moderate $= 2$
entire $loss = 3$				lymphoplasmacytosis =2	Complete regeneration or normal tissue =	severe $= 3$
Loss of crypt + epi= 4					0	

Expression of TLRs in the AOM-DSS model.

TLRs	Normal mucosa (untreated)	Non-dysplastic surrounding mucosa	Tumor	P value [*]
TLR2	2.4 ± 2.0	4975.0 ± 6835.7	206.0 ± 582.9	0.207
TLR3	359.2 ± 238.1	8555.8 ± 14520.5	6700.0 ± 17380.6	0.197
TLR4	2.7 ± 3.0	48.1 ± 36.3	1538.7 ± 1546.0	0.027
TLR5	19.7 ± 5.8	375.9 ± 706.2	318.9 ± 836.2	0.453
TLR9	100.0 ± 76.9	446.8 ± 533.1	300.1 ± 410.7	0.298

Table 2

*Comparing tumor with inflamed surrounding mucosa.

Table 3

Incidence and size of polyps

	WT (n=18)	TLR4-/- (n=18)	P value
Incidence (%)	100	27.8	0.027
Tumors/animal (range)	$5.4 \pm 1.2 \ (1 - 17)$	$0.4 \pm 0.2 (0 - 2)$	0.00016
Tumor size (mm) (range)	$3.6 \pm 0.4 \ (1 - 10)$	$1.3 \pm 0.3 \ (0.5 - 1)$	0.02
Percentage of mucosal surface with tumor (range)	22.6% ± 3.1 (1 – 65)	$4.4\% \pm 0.6 (1 - 5)$	0.0025