



# Research progress on the carcinogenesis mechanism of inflammation in ulcerative colitis: a narrative review

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**Objective:** In this review, we have summarized the influence of inflammation-related pathological mechanisms on the development of ulcerative colitis (UC) to colorectal cancer (CRC).

**Background:** UC is a chronic inflammatory bowel disease (IBD) of unknown etiology that affects the colon and rectum. Long-term inflammation of UC may greatly increase the risk of CRC, and the secretion of inflammatory factors and sustained inflammation may be key drivers of UC-associated CRC progression. Compared with the general population, the risk of CRC in patients with UC is 2.4 times higher, and the mortality rate of patients with UC is higher than that of those with sporadic CRC. The use of non-steroidal anti-inflammatory drugs can reduce the probability of UC transforming into CRC.

**Methods:** Literatures about inflammation and UC were extensively reviewed to analyze and discuss.

**Conclusions:** We believe that the mechanism of continuous inflammation that promotes cancer in UC may be the result of the mutual influence of intestinal microbes, inflammatory signals, and tissue remodeling. The invasion of intestinal microorganisms activates inflammatory signals and promotes the secretion of inflammatory factors which intensifies the remodeling of the extracellular matrix (ECM) and recruits immune cells. Eventually, a mutually engendering circuit of microbial invasion, release of inflammatory mediators, and remodeling of ECM is formed, which triggers continuous inflammation and promotes development of CRC.

**Keywords:** Ulcerative colitis (UC); Crohn's disease (CD); gut microbes; inflammatory signals

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## Introduction

Ulcerative colitis (UC) is a common chronic inflammatory bowel disease (IBD) which happens in part or all of the mucosa and submucosa of the colon. It is characterized

by urgent bowel movements, tenesmus, bloody diarrhea, abdominal pain, and fatigue (1). In the past 20 years, with the rapid industrialization in the south and east of China, the incidence of UC in these areas has risen swiftly. The

incidence of UC in these industrialized areas is close to that of developed countries (2). The exact pathogenesis of UC is still unknown, although currently the main influencing factors are understood to include immune response disorders, changes in intestinal microbiota, genetic susceptibility, and environmental factors (3). In spite of the growing efforts of researchers in response to the increasing incidence of UC, its management is still limited. According to the guidelines of the American Gastroenterology Association (AGA), in addition to surgical treatment when necessary, corticosteroids should be used to prevent inflammation in the acute phase. Similarly, long-term treatments are mainly steroidal and non-steroidal anti-inflammatory drugs and immunomodulators (4,5). These drugs often have a wide range of side effects, and some remain controversial.

Along with the suffering caused by UC, the long-lasting inflammation in UC could possibly greatly increase the risk of colorectal cancer (CRC). Overall, the incidence of CRC in patients with UC is about 2.4-fold higher than people without UC (6). Perhaps UC patients themselves are prone to CRC-inducing genetic mutations. Some researchers have estimated that the incidence of CRC in UC patients after 30 years of UC diagnosis could be as high as 20% or more (7). Considering this bias, the incidence of CRC in UC patients has modestly decreased, especially in developed countries (6). However, as the incidence of UC has been increasing significantly with the process of social industrialization (2), the number of CRC cases potentially induced by UC is increasing rapidly, especially in developing countries like China.

The secretion of inflammatory factors and the persistence of inflammation may be key promoting forces in the progression of UC-related CRC, including in its occurrence, development, and progression. Intestinal microbial infection, inflammatory signaling activation, inflammatory cell infiltration, and inflammatory cytokine release in intestinal epithelial cells could all support the CRC promoting inflammation. The CRC promoting inflammation could further lead to chronic inflammation, inflammatory hyperplasia, abnormal hyperplasia, and tumor development (8). For example, in patients diagnosed with UC for the first time, interleukin (IL)-17A, a typical inflammatory cytokine, can predict the severity of the disease very well, and also the course of the disease in the future 3 years (9). It may also be involved in the occurrence and development of CRC, as it was shown to be more highly expressed in CRC than in UC, and rarely detected in normal

tissues (10). Anti-inflammatory therapeutics such as anti-IL-17A can significantly alleviate inflammation in UC and reduce the risk of CRC in UC mice model (11). Exploring the mechanism of inflammation in the carcinogenesis of UC may help prevent and treat UC-related CRC. In this review, we have summarized the current research progress in the pathogenesis of CRC carcinogenesis with respect to infection, tissue remodeling, inflammatory cytokine secretion, and inflammation activation in UC. We propose that mutual influence of intestinal microbes, inflammatory signals, and tissue remodeling all contribute to the continuous inflammation that promotes UC-related CRC. We present the following article in accordance with the Narrative Review reporting checklist (available at <https://dx.doi.org/10.21037/apm-21-3138>).

### CRC and gut microbiota

Including both UC and Crohn's disease (CD), IBD is considered an abnormal mucosal immune response of the body to the gut microbiota under the influence of environmental factors, pathogens, and genetic susceptibility (12). Under physiological conditions, the intestinal barrier is composed of epithelial cells, extracellular matrix (ECM) components, and a mucus layer with a complex physical and chemical structure. The barrier could protect the intestine from infection and unnecessary attack of the innate and the adaptive immune system. When UC occurs, the intestinal mucosal barrier is attenuated, which allows intestinal microorganisms to invade into the mucosa (13). In more than 80% of the colon specimens of UC patients, intestinal bacteria were found in the lesions where the mucosa was destroyed (14). In mouse models of colitis induced by azomethane (AOM) or dextran sodium sulfate (DSS), intestinal microorganisms were detected in the mucosal layer much earlier than leukocyte infiltration or changes in epithelial structure (15). These findings indicate that the invasion of intestinal microorganisms may be an important driving force for the occurrence of UC.

Gut microbes are made up of a large number of different bacteria that produce a variety of metabolites. A well-established feature of CRC development is that most cancers originate as precursors to benign polyps (16). In the majority of cases, CRC develops from adenomatous polyps through dysplastic alterations that are strongly influenced by the microbiota (17). It is estimated that at least 0.2% to 8.3% of colorectal polyps are malignant (18). A variety of molecular changes may occur during the

transformation of these polyps from benign to malignant tumors, so the ability to study metabolites produced by the microbiome could lead to new discoveries about the development and progression of this cancer. Patients with UC are prone to colorectal cancer and dysplastic polyps, as well as sporadic adenomas (7). Previous studies have assessed the incidence of dysplastic lesions in UC patients and found that the incidence of adenomatous polyps in IBD is extremely low, only about 2% (19,20). This may be due to the beneficial protection of chemical drugs in UC and the inhibition of the inflammatory process of UC in adenoma development (21). Although a low incidence of dysplastic polyps is observed in patients with UC, this should not preclude close monitoring of this high-risk population.

Gut microbes may also play an important role in the development of UC-related CRC. There is an interaction between the intestinal flora and monocyte-like macrophages. The typical interactive effect is the activation of the Toll-like receptor (TLR) of immune cells with bacteria-derived lipopolysaccharide (LPS). Activation of TLR mediates the secretion of inflammatory cytokines such as IL-1 $\beta$  and IL-17, which in turn activates T cell helper cells and the corresponding cancer-promoting inflammatory response (22). Intestinal microbes can also affect the inflammation and tumor transformation of cells with their metabolites. There is a reduced diversity of gut microbes in each single patient with IBD. The influence of disease processes, alterations of disease on the microbiome, and differences in the diet and drug use of patients all possibly bring about big differences in the microbial metabolome between different UC patients. Metabolites of the microbes can be used as a fine indicator of the disease course. Conversely, specific metabolites from UC patients, such as fecal calprotectin (FC), can also be used as a biomarker in diagnosing UC (23). In patients with UC who received microbiota transplantation, changes in bacterial species and metabolites were also observed as the condition improved (24). Among the many metabolites, linoleic acid and 12-hydroxy-8,10-octadecadienoic acid were implicated to have strong correlations with the development of UC-related CRC, suggesting that these metabolites can be used as key biomarkers for the development of UC-related CRC (25). The composition and metabolism of gut microbes are the decisive factors in the occurrence of UC-related CRC. However, whether either the gut microbes or their metabolites should be responsible for the inflammatory and carcinogenesis effects of UC is still needs further research (23).

### TLR4/nuclear factor kappa B pathway

Microorganisms activate inflammatory lesions in UC. The glycolipid LPS is widely found in the cell walls of Gram-negative bacteria. It targets TLR of epithelial cells under infection, which further activates downstream inflammatory pathways such as nuclear factor kappa B (NF- $\kappa$ B) (26). As the most important inflammation-controlling hub that regulates the transcription of cytokines, cytokine receptors, and adhesion molecules, NF- $\kappa$ B participates in the regulation of multiple pathological processes of UC and CRC, including proliferation, apoptosis, metastasis, and drug resistance (27). Inhibitor of NF- $\kappa$ B kinase subunit beta (IKK $\beta$ ) is a key regulator of NF- $\kappa$ B activation. In the AOM/DSS-induced UC mouse model, the IKK $\beta$  knockout mice were less likely to develop CRC. Similarly, compared with healthy controls, in the bone marrow-specific IKK $\beta$  knockout mice, tumorigenesis was inhibited, and the pro-inflammatory cytokines in the intestinal mucosa such as IL-1 $\beta$ , ICAM, IL-6, MIP-2, KC, COX-2, and matrix metalloproteinases (MMP)-9 were significantly reduced (28). To summarize, both the UC lesions and immune cells could be activated by classical and non-classical NF- $\kappa$ B signals to promote the inflammation and progression of UC.

In multiple cancers, including CRC, NF- $\kappa$ B has been characterized as a tumor promoting regulator. Recent studies have shown that a variety of potential CRC suppressive chemicals, such as curcumin, dexamethasone, resveratrol, evodia, and so on, can inhibit NF- $\kappa$ B directly or indirectly. The inhibition of NF- $\kappa$ B with these chemicals could suppress tumor proliferation, inflammation, metastasis, and drug resistance, and could promote its apoptosis (27). As NF- $\kappa$ B interacts extensively with other cancer-promoting processes and molecules, it can regulate tumor progression and metastasis by promoting the induction of angiogenesis-related genes (29) (e.g., S100B, a common inflammatory protein in tumors). The levels of S100B in healthy control, peritumoral, UC, and CRC tissues increased with aggravation of the lesion. By directly binding to wtp53 and phosphorylating NF- $\kappa$ B, S100B up-regulates pro-inflammatory inducible nitric oxide synthase (iNOS), vascular endothelial growth factor (VEGF), and IL-6, inhibits apoptosis-inducing wtp53 and Bax, and forms a pro-inflammatory/angiogenic and anti-apoptotic environment. Inhibition of S100B has been shown to inactivate NF- $\kappa$ B and reduce CRC incidence (30). Other NF- $\kappa$ B interacting molecules mainly include IL-6/STAT3 (31), Notch, MMP, Cyclin, survivin, Fas, interleukin, TNF, and so on (27).

There have been many studies on NF- $\kappa$ B and tumors, but currently there is no effective and specific anti-tumor therapy that targets NF- $\kappa$ B in clinical practice. As a disease characterized by persistent inflammation, UC may be more likely to benefit from therapies such as anti-NF- $\kappa$ B in reducing the incidence of CRC.

### Tumor necrosis factor- $\alpha$

The tumor necrosis factor (TNF)- $\alpha$  protein is a main inflammatory mediator and is mainly expressed in infiltrating macrophages in UC (8,32). Immunoreactive TNF- $\alpha$  protein can be detected in the colon of patients with active UC and advanced CRC, but not in normal mucosa; similar effects are also seen in AOM/DSS-induced UC mouse models (33). Both the TNF- $\alpha$  chimeric monoclonal antibody (mAb) and inhibitor can significantly limit the level of inflammation in the colon and reduce the number and size of tumors in AOM/DSS challenged mice (33,34). Clinically, TNF- $\alpha$  mAb has a good effect on IBD patients, including those with UC and CD (35), which further illustrates the importance of TNF- $\alpha$  to CD.

The expression of TNF receptor (R2) is also upregulated in IBD patients and AOM/DSS challenged mice (34,36). However, the up-regulated subtype TNFR2 cannot not activate the FADD/caspase 8/caspase 3 pathway to promote apoptosis. Simultaneously, the activation of inflammation and proliferation with TNFR2 is enhanced (37). Furthermore, TNFR2 could activate NF- $\kappa$ B and other inflammatory pathways to form a pro-inflammatory positive feedback (34). In recent years, it has been discovered that the expression of metastasis-associated in colon cancer 1 (*MACC1*) gene may be initiated by the activation of TNF- $\alpha$  dependent NF- $\kappa$ B p65, which promotes the activation of the transcription factor c-Jun. These interactions may jointly promote the occurrence and development of UC-related CRC (8).

### IL-6

Although the role of TNF- $\alpha$  in the development of UC-related CRC has been widely accepted, TNF- $\alpha^{-/-}$  mice have been found to still develop UC and related CRC with a similar severity to normal mice under the stimulation of AOM/DSS (38). There are multiple ways of inflammation activation, which could all play important roles in the progression of UC and related CRC. Similar to TNF- $\alpha$ , IL-6 is also a typical multifunctional NF- $\kappa$ B regulatory cytokine. The circulating and intestinal IL-6 levels of

patients with IBD are elevated, and the level of IL-6 in the lesion is higher than that in normal tissues (39,40). In the AOM/DSS mice model, IL-6 expression also increased with the aggravation of the lesion. Specifically, the number of tumors in IL-6 $^{-/-}$  mice decreased, and the size of tumors were smaller (41,42). These studies indicated that IL-6 may be a key promoter in the development of UC and related CRC.

After IL-6 binds to sIL-6R, its receptor of soluble form, it mediates chronic inflammation both in UC and CRC through trans-signal transduction (43). In epithelial cells of mucosa associated with UC and CRC, IL-6R messenger RNA (mRNA) is down-regulated, while the expression of secreted sIL-6R is increased, which would promote tumorigenesis of CRC (44). Other targets of IL-6 include JAK/STAT3, Shp-2-Ras, PI3K/AKT, and so on. Interactions between IL-6 and these targets can further activate other inflammatory factors such as TNF- $\alpha$ . In short, IL-6 may mainly bind to its soluble form receptor sIL-6R to promote the occurrence and development of UC and CRC.

### Other major CRC promoting inflammatory signals

#### *The PGE2/COX-2 pathway*

Cyclooxygenase (COX) mainly includes COX-1 and COX-2, and its signal transduction is related to tumorigenesis (45). The expression of COX-2 only happens in response to certain pro-inflammatory stimuli. Pro-inflammatory cytokines induce the expression of COX2 and PGE2 and participate in vasodilation, leading to inflammation. It is also involved in inducing the expression of BCL-2, inhibiting apoptosis, and promoting tumor progression (46). In human CRC endothelial cells, PGE2 directly enhances the expression of CXCL1, a chemokine that can activate and recruit neutrophils and induce angiogenesis (47).

#### *The IL-23/Th17 pathway*

Dendritic cells and other antigen-presenting cells produce IL-23, a member of the IL-12 cytokine family. Since IL-23 receptors (IL23R) could regulate the differentiation of helper T cells (Th17), they are widely associated with chronic inflammatory diseases (26). Ustekinumab, an IL-12 and IL-23 monoclonal antibody, can be used for psoriasis, psoriatic arthritis, and CD, and also shows a certain clinical effect on UC (48,49).



## ECM

In UC, series of aggressive metabolites and inflammatory mediators accumulate in the inflamed intestinal mucosa, leading to tissue damage (50). Tissue damage repair, especially immune response-enabled ECM degradation and remodeling, may be one of the most important pathologies of UC (51). On the one hand, ulcers in the pathological manifestations of UC itself tell a story of tissue damage repair, especially ECM damage and remodeling. On the other hand, the secretion of aggressive metabolites and inflammatory mediators is activated in UC, including nitric oxide, prostaglandins, oxygen free radicals, histamine, MMPs, and other proteases. These aggressive metabolites and inflammatory mediators may be the natural responsive products of the immune system to infection and trauma. However, the metabolites and inflammatory mediators could also aggravate tissue damage, perturb ECM remodeling, and perpetuate the disease in UC (52). The disorder of ECM remodeling in UC may be an important factor leading to progression of UC and UC-related CRC. For example, MMPs, under physiological conditions, are mainly involved in embryogenesis and angiogenesis, allowing inflammatory cells to migrate to the damaged sites of the tissue. Under pathological conditions, these MMP-mediated migration processes may also induce ulcer healing disorders, sustained inflammation, and even cancer (53). Under continuous inflammatory signals such as the invasion of intestinal bacteria on the lesion and the secretion of inflammatory factors in tissues, the transcription, translation, secretion, and activation of MMPs are enhanced, basement membrane and ECM are hydrolyzed, and chemokines are secreted (52,54). Such a process may form a continuous positive feedback loop that drives UC.

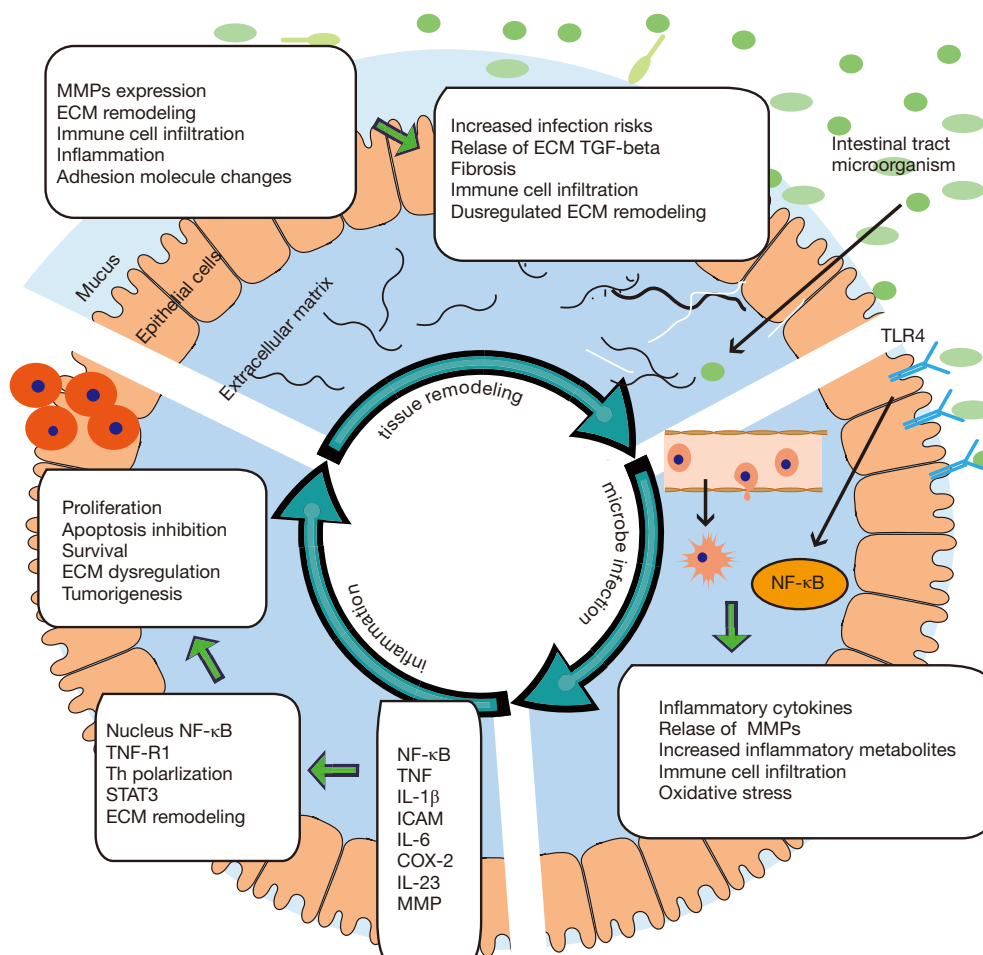
The content and way of combination of ECM components also have a great influence on local inflammation in UC. For example, the ECM component hyaluronic acid hardly adheres to white blood cells under normal conditions. When inflammation occurs, hyaluronic acid forms a complex with trypsin inter- $\alpha$  inhibitor, which attracts a large amount of white blood cells to activate inflammation (55). A similar mechanism is also found in the early stage of IBD tissue, where change in content and way of combination of ECM components happens to enable leukocyte infiltration, adhesion, and activation (56). Other ECM components including laminin, heparan sulfate proteoglycan, fibronectin, keratin, and dermatan sulfate have widely been reported as closely related to the occurrence and development of

UC (57). The TGF- $\beta$  protein is often considered the main regulatory protein of ECM remodeling. Recent studies have asserted that simply inhibiting the expression of TGF- $\beta$  in epithelial cells in the inflammatory microenvironment may lead to the occurrence of aggressive CRC (58). In colonic epithelial component-specific MMP-9 transgenic homozygous mice, reactive oxygen species accumulated and DNA damage reduced, which inhibited the occurrence and development of CRC (59). Tripartite motif-containing 21 (TRIM21) is a protein that regulates the immune response of the intestinal mucosa. The expression of TRIM21 is reduced in CRC patients. In the TRIM21 highly expressed mouse model, the tumor cell proliferation-related gene Ki67; the tissue remodeling and angiogenesis-related genes MMP10, HIF1- $\alpha$ , COX2, Ang4; and the pro-inflammatory cytokines IL-6, TNF- $\alpha$ , and IL-1 $\beta$  are all significantly up-regulated. While TRIM21 expression leads to down-regulation of the tumor cell adhesion-related gene E-cadherin and inflammatory cytokines IL-10, TGF- $\beta$ , and Foxp3, which inhibits tumor occurrence and development (60). All these findings indicate that disordered ECM and its interaction with inflammatory signals may be effective targets in the treatment of CRC. In short, ECM changes or remodeling participate in many aspects of UC. However, the importance of ECM in the development of UC and related CRC has not attracted enough attention, much about its role in the development of UC-related CRC requires further investigation.

## Summary

For the prevention of the UC-related CRC, timely control of the disease, regular screening, and healing of the damaged mucosa as soon as possible are the best strategies (6). Besides treatment when in the acute phase, often requires sustained use of non-steroidal anti-inflammatory drugs to control chronic inflammation, otherwise uncontrolled chronic mucositis may become a potent driving force for the development of UC and related CRC.

A variety of factors including intestinal microbes, inflammatory mediators such as NF- $\kappa$ B, TNF- $\alpha$ , IL-6, PEG2, and so on, and the remodeling of ECM all have direct or indirect effects on the pathology and inflammation of UC. Generally, the remodeling of ECM makes it easier for microorganisms to penetrate the mucus layer and invade epithelial cells. The invasion of intestinal microbes activates TLR4 and other inflammatory signal transduction in the body to promote the secretion of inflammatory factors, and



**Figure 1** Risk factors for the pathology and inflammation of UC. NF- $\kappa$ B, nuclear factor kappa B; TNF, tumor necrosis factor; IL, interleukin; MMPs, matrix metalloproteinases; ECM, extracellular matrix; COX-2, cyclooxygenase-2; TNF-R1, TNF receptor 1; ICAM, intercellular cell adhesion molecule.

activation of sustained inflammation. Then, inflammation and cytokines aggravate ECM remodeling and recruit immunity cells. Eventually, a mutually positive feedback of microbial invasion, inflammation, and ECM remodeling is established, which triggers continuous inflammation which may be responsible for the continuous inflammation in UC and related CRC (*Figure 1*).

Different subtypes of IBD, including UC and CD have many similarities or even overlaps in the location, pathological features, and treatment methods (1). The lesions of UC often appear in the epithelial layer of the colon only, progress in a spreading way, and are more likely to induce CRC. In contrast, CD spreads throughout the intestinal wall, the lesions are diffuse, and are more likely to spread to the small intestine side. Despite these distinctions,

especially in inflammation activation, tissue remodeling, and gene mutation, the 2 diseases are very similar (52,61). Therefore, the mechanism of inflammation in UC and related CRC summarized in this review possibly also aligns with CD. Similar to how controlling inflammation can reduce the risk of CRC, controlling inflammation may also help reduce CD-related cancers such as small bowel cancer.

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