**Fusobacterium nucleatum**

Contributes to the Carcinogenesis of Colorectal Cancer by Inducing Inflammation and Suppressing Host Immunity

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Abstract

The presence of *Fusobacterium nucleatum* (*F. nucleatum*) in the gut is associated with the development of colorectal cancer (CRC). *F. nucleatum* promotes tumor development by inducing inflammation and host immune response in the CRC microenvironment. Adhesion to the intestinal epithelium by the cell surface proteins FadA, Fap2 and RadD expressed by *F. nucleatum* can cause the host to produce inflammatory factors and recruit inflammatory cells, creating an environment which favors tumor growth. Furthermore, *F. nucleatum* can induce immune suppression of gut mucosa by suppressing the function of immune cells such as macrophages, T cells and natural killer cells, contributing the progression of CRC.

Introduction

The human intestine is home to more than 100 trillion microbes, forming a unique genome that changes with human nutrition status, geographic location, and even age [1,2]. Gut microbiota is in harmony with the human body, affecting human health and shaping the immune system of the host [3]. Imbalance in gut microbiota can lead to a variety of diseases, such as colitis, colorectal cancer (CRC), infection, food allergies, obesity, diabetes, cardiovascular atherosclerosis, bone metabolic diseases, Parkinson’s and neurodegenerative diseases [4].

The colon has been colonized by the largest density of microorganisms [5]. Recent studies have reported that certain pathogenic bacteria in the colon are associated with CRC [5,6]. It’s possible to screen CRC by detecting tumor-associated microorganisms in the feces [7]. CRC is the third most common tumor in the world, causing significant morbidity and mortality [3]. Unfortunately, the mechanism of this malignancy has not been fully explained, but inflammation is a recognized risk factor [8]. CRC is a chronic disease which can arise from other intestinal inflammatory conditions [6].

In recent years, many studies have been conducted on the correlation between inflammatory microorganisms and CRC. The occurrence of intestinal inflammatory diseases such as colitis is related to the metastasis of intestinal microbes [2]. Colitis occurs when microbes turn from a ‘eubiotic’ to a ‘dysbiotic’ state [2]. Chronic bacterial infection of the colon is a driving factor in tissue inflammation, increasing the risk of developing CRC [9].

Gut Microbiota and Host Immunity

Human bodies are constantly exposed to a diverse array of microbes, as well as their metabolite byproducts [10]. The gut microbiome influences the development of the immune system of the host, and conversely the immune system regulates the microbe composition in the gut [5,11]. Microorganisms in the gut play a key role in the activation, training and regulation of the host immune system [12]. The communication between the intestinal microflora and the host’s immune system begins at a very early stage of development [13]. Intestinal microbes can recruit immune cells and initiate inflammatory reactions, which play a direct role in promoting the maturation of the immune system [10,14]. The intestinal...
mucosal immune system is a protective barrier system composed of cellular (i.e. epithelial- and mesoderm-derived immune cells) and non-cellular components (e.g. antimicrobial peptides, cytokines and antibodies), which can resist microbial attack [15].

On the one hand, intestinal microbes affect the host immune system through secretion of metabolites (e.g. butyrate, L-tryptophan, indole, bile acids and retinoic acid), signaling pathways (e.g. Toll-like receptors and Nod-like receptors), and small noncoding RNAs (e.g. miRNA) [9,16]. On the other hand, the intestinal immune system exposes bacteria to the host, reducing pathological outcomes, and modulating the stratification of bacteria in the epithelial barrier [17]. The intestinal immune system affects the composition of bacteria, whereas the bacteria promote the development of the intestinal immune system. Disruption of the relationship between intestinal bacteria and the host immune system will affect the overall health of the host [15,17]. Intestinal epithelial cells recognize pathogen-associated molecular structures (such as lipopolysaccharides and flagella) through their surface Toll-like receptors. This triggers the maturation of antigen-presenting cells (such as dendritic cells), the initiation of immune responses, and the release of inflammatory factors, which are associated with the development of CRC [18]. This dysregulation of the host immune system represents a potential mechanism for the effect of intestinal microbes on the development and progression of CRC [19].

Invasion of Fusobacterium nucleatum Contributes to the Carcogenesis of CRC

Fusobacterium nucleatum is a Gram-negative, anaerobic oral commensal bacterium that is associated with a variety of human diseases, including periodontal disease [20], Alzheimer’s disease [21], brain abscess [22], cardiovascular disease [23], miscarriage [24] and inflammatory bowel disease [25]. Recently, F. nucleatum has been proposed to be associated with CRC [26]. F. nucleatum promotes the occurrence of CRC through several virulence mechanisms: colonization, invasion, and modulation of host immune response [27].

F. nucleatum bacteria interact with each other by expressing a variety of different virulence factors, and can adhere to many different mammalian cell types, including epithelial and endothelial cells, polymorph nuclear neutrophils, monocytes, erythrocytes, fibroblasts, and natural killer (NK) cells [28,29]. The cell surface protein FadA is a key virulence factor in F. nucleatum which regulates adhesion and invasion of the bacterium. The expression of FadA gene in human CRC specimens was significantly higher than that in adjacent normal tissues [30]. This protein enables F. nucleatum to bind E-cadherin in CRC and epithelial cells, activate the β-catenin pathway, and induce the expression of transcription factors lymphoid enhancer factor (LEF)/T cell factor (TCF) which promote tumor cell growth [30,31]. It’s recently reported that FadA can up-regulate Wnt/β-catenin modulator Annexin A1 expression through E-cadherin [32]. FadA can also bind to endothelial cells VE-cadherin, which is a linker molecule on endothelial cells [33]. This combination alters the integrity of the endothelium, increases the permeability of the endothelium, and allows the bacteria to overcome the blood brain barriers, placental barriers, and colonize different parts of the body [34]. Outer membrane vesicles (OMVs) from F. nucleatum can degrade E-cadherin, thus promoting bacterial invasion and tumor metastasis [35].

In addition, F. nucleatum also has two other outer membrane proteins, Fap2 and RadD [36]. The lectin Fap2 can bind Gal-GalNAc, a polysaccharide overexpressed in CRC. This binding of Fap2 facilitates colonization of F. nucleatum and explicates fusobacteria abundance in CRC [37]. RadD can mediate communication between F. nucleatum and other bacterial species, contributing to the formation of multispecies biofilms [36,38], which has been shown to be associated with proximal colon cancer [39].

F. nucleatum-Induced Inflammation Contributes to CRC Development

There has been a growing body of literature suggesting a link between chronic inflammation and CRC, in which gastrointestinal inflammation may promote CRC development [30]. Increased evidence suggests that F. nucleatum can shape the inflammatory microenvironment in the CRC, promoting tumor growth and metastasis [40]. For example, F. nucleatum can stimulate reactive oxygen species (ROS) production, inducing inflammatory responses in CRC cells [41]. Infection of CRC cells with F. nucleatum increased the expression of miR21, a pathogenic role in chronic inflammation and colitis-associated colon cancer, therefore promoting tumor cell proliferation and invasive activity [8,42].

Adherence of F. nucleatum to CRC cells via FadA stimulates the release of inflammatory factors, such as NF-κB, IL-6, IL-8, IL-10 and IL-18, which promote cell proliferation in CRC [30]. In patients with F. nucleatum infection, strong humoral immunity is induced, and antibodies against F. nucleatum, IgA and IgG are present [43]. F. nucleatum infection increases the infiltration of inflammatory cells, such as macrophages, dendritic cells, and granulocytes, which create a pro-inflammatory microenvironment that is conducive to the occurrence of CRC [44]. Macrofages infected with F. nucleatum can induce the release of inflammatory cytokines [45,46]. Natural cytotoxic receptor NKP46 of NK cell can directly recognize F. nucleatum through its surface ligand, secreting TNF-α to aggravate inflammation [20]. Some inflammatory response signatures were specific to F. nucleatum but not to other bacteria found in CRC tissues, such as IL1β, IL24, PTGS2 (COX-2), IL8, IL6 and TNF, which were enriched in F. nucleatum-infected CRCs [44].

F. nucleatum is prevalent in gastrointestinal inflammation diseases especially inflammatory bowel disease (IBD) [25]. F. nucleatum strains originating from IBD patients were significantly more invasive than strains isolated from healthy tissues [25]. Highly invasive F. nucleatum isolates derived from the inflamed area of human Crohn’s disease triggered high expression of MUC2 and TNF-α in colon cancer cells [47]. In IBD patients, the release of IL-1β and TNF-α can damage colon cells and impair epithelial integrity, which increases the chance of contact between F. nucleatum and the colon epithelium [48]. This may partially explain why patients with IBD are susceptible to CRC. Mechanisms of adhesion, invasion and inflammation mediated by F. nucleatum in CRCs were summarized in Table 1.

F. nucleatum-Induced Immune Suppression Promotes CRC Development

Macrophages

F. nucleatum modulates the tumor immune environment by amplifying bone marrow-derived cells [49] such as tumor-associated macrophages, which play an important role in tumor invasion and metastasis [50]. Meanwhile, the tumor microenvironment can affect the heterogeneity of macrophages, which can differentiate from pro-inflammatory M1-phenotype to a tumor-promoting M2-phenotype [51,52]. Our recent study revealed that F. nucleatum displayed an immunosuppressive effect by promoting M2 polarization of macrophages in F. nucleatum-related CRCs, possibly through the TLR4/IL-6/p-STAT3/c-MYC signaling pathway [53]. F. nucleatum
induces infiltration of M2 macrophages in the colorectal environment, thus forming a tumor-promoting microenvironment \[54,55\]. The metabolite of \( F. \) nucleatum, butyric acid, can induce apoptosis in monocytes/macrophages and lymphocytes by activating free fatty acid receptors \[56\]. In addition, \( F. \) nucleatum can invade macrophages and induce the expression of indoleamine2,3-dioxygenase on the cell surface, creating a toxic microenvironment which impairs the function of peripheral blood lymphocytes, thereby allowing macrophages to escape cytotoxic T lymphocyte attack \[55\].

**T cells**

\( T \) cell activity can be inhibited by the virulence factors of \( F. \) nucleatum \[36,57\]. \( F. \) nucleatum abundance is negatively correlated with CD3\(^+\)T cell density in CRC \[1\]. Additionally, we have shown that a high abundance of \( F. \) nucleatum in CRC is associated with lower numbers of CD4\(^+\)T cells \[58\]. Decreased T cell density in CRC could be explained by apoptotic cell death and arrested proliferation of T cells induced by \( F. \) nucleatum \[59–61\]. For example, \( F. \)

\[Table 1. \textit{Fusobacterium nucleatum} induced invasion and inflammation contributes to colorectal cancer\]

| Virulence factor | Function | Mechanisms | References |
|------------------|----------|------------|------------|
| Fn infection     | Pro-inflammatory microenvironment infiltration | Inducing inflammatory cells | [49]       |
|                  | Inflammatory cytokines production | Accumulation of reactive oxygen species | [41]       |
|                  | Cell proliferation and invasion | Increasing the expression of miR21 | [8,42]     |
| FadA             | Cell proliferation | Activating the β-catenin pathway | [30,31]    |
|                  | | Up-regulating Wnt/β-catenin modulator Annexin A1 | [32]       |
| Fap2             | Bacterial colonization | Binding endothelial cell VE-cadherin | [33,34]    |
|                  | Bacterial colonization | Binding Gal-GalNAc overexpressed in CRC | [37]       |
| RadD             | Biofilms formation | Mediating communication between Fn and other bacteria | [36,38]    |
| LPS              | Inflammatory cytokines production | Activating immune cells | [55]       |
| OMVs             | Bacterial invasion and tumor metastasis | Degrading E-cadherin | [35]       |

\( Fn, Fusobacterium nucleatum; CRC, colorectal cancer; OMV, outer membrane vesicles. \)

Figure 1. Infiltrating immune cell populations in human \( F. \) nucleatum related colorectal cancer. High abundance of \( F. \) nucleatum within colon cancer tissue (A) and matched metastatic lymph nodes (B) detected by immunofluorescence. High density of immune cells (CD3\(^+\), CD68\(^+\), CD83\(^+\), and NE cells) within the environment of \( F. \) nucleatum-positive colon cancers (immunofluorescence). NE, neutrophils.
**Fusobacterium nucleatum** inhibitory protein (FIP) can inhibit human T cell activation by arresting cells in the G1 phase of the cell cycle [60]. A recent study revealed that the association of *F. nucleatum* with tumor-infiltrating lymphocytes (TIL) differed by MSI status of CRC. The presence of *F. nucleatum* was negatively associated with TIL in MSI-high tumors, but positively in non-MSI-high tumors [62].

In the colorectal tumor microenvironment, *F. nucleatum* can release short-peptides (formylmethionyl-leucyl-phenylalanine) and short-chain fatty acids (butyrate, propionate, and acetate) which lead to recruitment of myeloid-derived suppressor cells (MDSCs) [48]. MDSCs can regulate immune response by suppressing CD4+ T helper cell function, inhibiting T cell proliferation, and inducing T cell apoptosis [48,63]. *F. nucleatum* can also induce human lymphocyte death through Fap2 and RadD [36]. Fap2 of *F. nucleatum* can inhibit human T cell activation by directly interacting with TIGIT, an inhibitory receptor present on various T cells [57]. Moreover, *F. nucleatum* can interact directly with monocytes, which may recruit T helper 17 cells and T regulatory cells through CCL20/CCR6 pathway, promoting CRC formation [64].

**NK Cells**

All human NK cells express the Fap2 receptor TIGIT, which recognizes poliovirus receptor (PVR) and nectin-2 as ligands [65]. This binding of NK cells to *F. nucleatum* inhibits the killing activity of NK cells, thereby promoting the formation of colorectal tumors [35,57,66].

**Dendritic Cells**

The infiltration of CD103+ dendritic cells (DCs) was increased in tumors from *F. nucleatum*-fed mice compared with control group [44]. This population of DCs can promote the expansion of Foxp3+ regulatory T cells, a CD4+ T cell subset that inhibits cytotoxic and effector T cells, therefore diminishing anti-tumor immunity [67].

**Tumor-Associated Neutrophils**

Amount of tumor-associated neutrophils (TANs) in intestinal tumors of *F. nucleatum*-fed mice was significantly increased compared with controls [44]. It’s recently reported that TANs play a role in tumor progression and in the regulation of anti-tumor immunity [68]. Increased TANs in CRC associate with malignant phenotype and predict poor prognosis of patients with CRC [69]. These findings suggest that *F. nucleatum* may suppress antitumor immunity through inducing the infiltration of TANs in CRC.

Representative images of infiltrating immune cells in *F. nucleatum*-related CRC were shown in Figure 1. Adhesion, invasion, inflammation and immune suppression mediated by *F. nucleatum* in CRC was sketched in Figure 2. Mechanisms of immune suppression induced by *F. nucleatum* in CRC were summarized in Table 2.

### Table 2: *Fusobacterium nucleatum* induces immune suppression in colorectal cancer

| Immune cells | Function | Mechanism | References |
|--------------|----------|-----------|------------|
| Macrophages  | M2 polarization  | Activating TLR4 signaling pathway | [53] |
|              | Apoptosis      | Butyric acid activating free fatty acid receptors | [56] |
|              | Escape T lymphocyte attack | Impairing the function of peripheral blood lymphocytes | [55] |
| Lymphocytes  | Reducing CD3+ T cells | Correlated with TOX expression | [1] |
|              | Reducing CD4+ T cells | Arresting cells in the G1 phase | [60] |
|              | Inhibiting proliferation | Interaction with TIGIT | [57] |
|              | Inhibiting activation | Recruitment of MDSCs | [48,63] |
|              | Apoptosis      | Butyric acid | [56] |
| NK cells     | Inhibition of NK cell cytotoxicity | Fap2 binding TIGIT molecule | [35,57] |
| DCs          | Dampening anti-tumor immunity | Promoting the expansion of regulatory T cells | [67] |
| TANs         | Dampening anti-tumor immunity | Increasing the number of TANs | [44] |

TOX, thymocyte selection-associated high-mobility group box; MDSCs, myeloid-derived suppressor cells; NK, natural killer; DCs, dendritic cells; TANs, tumor-associated neutrophils.
Conclusion
The presence of *F. nucleatum* as symbiotic bacteria in the human intestinal tract has been confirmed to be related to the development of CRC. *F. nucleatum* promotes CRC through different virulence mechanisms, such as adhesion to the intestinal epithelium and inducing inflammatory and immune responses in the host. The resistance reactions induced in the host by *F. nucleatum* induce an inflammatory environment in the host, and promote the recruitment of inflammatory cells as well as the secretion of inflammatory factors. This response to *F. nucleatum* creates a microenvironment which favors tumor growth. Furthermore, *F. nucleatum* can induce immune suppression of gut mucosa by suppressing the function of immune cells such as macrophages, T cells and NK cells, contributing to the progression of CRC.

Conflicts of Interest
Authors declare no Conflict of Interests for this article.

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