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Ancient nuclear receptor VDR with new functions: microbiome and inflammation

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Abstract

Nuclear receptor vitamin D receptor (VDR) regulates most of the biological actions of the active vitamin D metabolite, 1α ,25-dihydroxyvitamin D₃ (1,25(OH)₂D₃). VDR is highly expressed in intestine and is a critical regulator of proliferation and differentiation, intestinal barrier function, innate immunity, and host defense in the gut. Evidence strongly supports a protective effect of vitamin D and VDR in inflammatory bowel diseases (IBD). Emerging evidence demonstrates low VDR expression and dysfunction of vitamin D/VDR signaling in patients with IBD. In the current review, we summarize recent progress made in vitamin D/VDR signaling in genetic regulation, immunity, and microbiome in both ulcerative colitis and Crohn's disease. We cover the mechanisms of intestinal VDR in regulating inflammation through inhibiting the NF- κ B pathway and activating autophagy. Recent studies suggest that the association of VDR SNPs with immune and intestinal pathology may be gender-dependent. We emphasize the tissue specificity of VDR and its gender and time-dependent effects. Furthermore, we discuss potential clinical application and future direction of vitamin D/VDR in prevention and treatment of IBD.

Keywords

Inflammatory Bowel Diseases; immunity; infection; bacteria; microbiome; myeloid; nuclear receptor; *Salmonella*

Introduction

The vitamin D receptor is an ancient nuclear receptor that has been highly conserved in birds, fish, and mammals.¹ Epidemiological studies have demonstrated that vitamin D deficiency is associated with various human diseases.² Vitamin D absorbed from the diet or produced in the skin after exposure to sunlight is then converted to 25(OH) VD by the liver and to the active vitamin D metabolite 1α ,25-dihydroxyvitamin D3 (1,25(OH)₂D₃) by the kidney.³ The vitamin D receptor (VDR) regulates most of the biological actions of the active vitamin D metabolite, 1α ,25-dihydroxyvitamin D₃ (1,25(OH)₂D₃). VDR is highly expressed in the small intestine and colon (The Human Protein Atlas https://www.proteinatlas.org/

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ENSG00000111424-VDR/tissue).4,5 VDR plays critical roles in proliferation and differentiation, intestinal barrier function, innate immunity, and host defense in the gut.⁶

It is acknowledged that a complex interplay of genetic risk factors, environmental triggers, immune responses, and intestinal bacteria contributes to the pathogenesis of IBD,^{7–9} which includes Crohn's disease (CD) and ulcerative colitis (UC). Low vitamin D status has been observed in IBD patients.^{10–12} A north-south gradient in rates of CD suggests that vitamin D deficiency due to insufficient sunlight in regions farther from the equator may be an environmental trigger contributing IBD pathogenesis. Vitamin D also influences the course and severity of IBD.¹¹ Evidence strongly supports a protective effect of vitamin D and VDR in IBD. Low VDR expression and dysfunction of vitamin D/VDR signaling in reported in both CD and UC patients.^{13,14} We show that a low intestinal epithelial VDR protein level is associated with impaired autophagy function, accompanied by a reduction in the mRNA and protein levels of *Atg1611*, an IBD risk gene and regulator of autophagy and the microbiome. ¹³

In the current review, we will summarize recent progress made in human *Vdr* gene variation in IBD, vitamin D/VDR signaling in transcriptional regulation, immunity, and microbiome in IBD. We will cover the mechanisms of VDR in regulating inflammation and discuss potential clinical applications of vitamin D/VDR in prevention and treatment of IBD. The vitamin D/VDR pathway is essential in maintaining homeostasis and signaling between the microbiota and host in intestinal inflammation.

We performed an electronic literature search of papers written in English in the MEDLINE database via PubMed. Searches included combinations of the following terms: vitamin D, vitamin D receptor, microbiome/microbiota, inflammatory bowel disease, inflammation, autophagy, nuclear factor kappa B. Papers without clear relevance to the role of VDR signaling in inflammation or microbiota regulation were excluded.

Human Vdr gene variation in IBD

The human *Vdr* gene is located on chromosome 12q. It is composed of promoter and regulatory regions (1a–1f) and exons 2–9, which encode 6 domains of the full length VDR protein. When 1,25(OH)₂D₃, the active form of vitamin D also known as calcitriol, binds to VDR protein, VDR is stabilized and translocates to the nucleus. Ligand-bound VDR associates with the retinoic acid receptor (RXR) via the dimerization domains and functions as a transcription factor¹⁵. The 1,25(OH)₂D₃–VDR–RXR complex binds to the vitamin D response elements (VDRE) in the promoters of target genes via the VDR DNA-binding domain and activates the expression of target genes, including the autophagy regulator ATG16L1,¹³ barrier protein claudin 2,¹⁶ scaffold protein Axin1,¹⁷ and anti-microbial peptides cathelicidin,¹⁸ and defensin- $\beta 2$.¹⁹

Susceptibility to IBD is associated with polymorphisms in the Vdr gene.^{20–22} Numerous single nucleotide polymorphisms (SNPs) are found in the Vdr gene. Five are of particular interest in IBD research: the non-synonymous *FokI* and synonymous *BsmI*, *ApaI*, *TaqI* and *Tru9I*. These SNPs, each named for the restriction endonuclease site it disrupts, have been

reported to be associated with increased susceptibility to colorectal cancer.^{15,23} Studies on the role of these SNPs in IBD susceptibility have been contradictory.²⁴ Recent studies suggest that the association of *Vdr* SNPs with immune and intestinal pathology may be gender-dependent. The *Taq1* SNP is more frequent in male UC patients.²⁵ Another study found that the haplotype "baT", which contains the ancestral alleles for *BsmI, ApaI*, and *TaqI*, protects against adenomatous polyps only in females, while the *FokI* SNP increases risk particularly in males.²⁶ There is also gender disparity in the association of *FokI* with lumbar pathology. The Ff allele of *Fok1* is protective in females while ff is required for protection in males.²⁷ Gender-specific effects of *Vdr* SNPs is also observed in other immune-related diseases, including multiple sclerosis²⁸ and type I diabetes.²⁹

Relevance of intestinal VDR in IBD

Although VDR is expressed in numerous tissues, intestinal epithelial expression is of particular importance in protecting against IBD. The expression level of VDR is critical in IBD pathology and treatment. Intestinal VDR expression decreases in both UC and CD patients,¹⁴ which reduces the anti-inflammatory effect of the vitamin D analogue. TNF-a, a pro-inflammatory cytokine associated with IBD, is known to reduce the expression level of intestinal VDR.³⁰ Constitutive RAS signaling, which is commonly observed in colon cancer, downregulates VDR expression and thus inhibits vitamin D signaling.³¹ We need also consider the heterogeneity of VDR and its variable expression in physiological and pathological conditions in the intestine when assessing the efficacy of vitamin D / vitamin D analogue treatments in IBD.

Although it is possible that decreased VDR is a consequence of chronic inflammation in IBD, in experimental models of colitis *Vdr* whole body knockout mice are known to develop a severe colitis.³² This can be ameliorated by introducing transgenic expression of VDR in intestinal epithelial cells. Additionally, wild-type mice with transgenic overexpression VDR in the epithelium are less susceptible to chemical-induced models of colitis.¹⁴ This protection is associated with decreased epithelial production of pro-inflammatory cytokines. We have shown that bacterial product butyrate or probiotics protect against colitis by increasing the expression of VDR at the mRNA and protein level, thus activating VDR signaling in epithelial cells.^{13,33}

Conditional knockout of intestinal epithelial *Vdr* alters the microbiota that promotes more severe inflammation in chemical-induced models of colitis. These conditional knockouts have impaired Paneth cell function marked by deficient autophagy and decreased production of antimicrobial peptides (AMPs).¹³ Dietary vitamin D deficiency also impairs AMP production, resulting in dysbiosis and worsened high fat diet-induced metabolic disease.³⁴ VDR signaling in Paneth cells may be critical in preventing dysbiosis and maintaining mucosal health.

The intestinal barrier is critical in maintaining intestinal homeostasis. Vitamin D treatment has been found to promote barrier function in *in vitro* and *in vivo* studies.^{35–37} Loss of VDR expression in intestine leads to decreased production of the barrier protein claudin 2¹⁶ in cells without any stimulation. VDR also promotes barrier integrity by preventing intestinal

epithelial apoptosis.³⁸ These studies suggest that VDR signaling may protect against IBD by promoting intestinal barrier function. Further study is warranted to determine if intestinal VDR influences the expression of additional barrier proteins.

IL-10 knockout mice have lower intestinal VDR expression and spontaneously develop a microbiota-dependent colitis, which is more severe in Vdr/IL-10 double knockouts.³⁹ Transgenic expression of human VDR in the mouse epithelium ameliorates *IL-10* knockout colitis,³⁸ whereas 1,25(OH)₂D₃ treatment does not.⁴⁰ VDR expression reduces epithelial pro-inflammatory cytokine production and immune cell infiltration into the mucosa,³⁸ indicating that epithelial VDR signaling is critical in modulating the immune response in IBD. Thus, VDR could be both a cause (e.g. gene variation and conditional knockout) and a consequence (reduced intestinal VDR protein) of chronic inflammation.

Vitamin D/VDR regulates immunity

VDR regulates both innate and adaptive immune cells

As described above, epithelial VDR expression alone is able to ameliorate genetic and chemical models of colitis. However, VDR expression in both innate and adaptive immune cells also affects the inflammatory response. Myeloid-specific *Vdr* knockout mice are more susceptible to DSS-induced colitis.⁴¹ Deletion of VDR in macrophages and granulocytes mildly affected colitis-associated symptoms but greatly increased pro-inflammatory cytokine expression in the inflamed colon, suggesting a prominent role for innate immune cell VDR signaling in regulating intestinal inflammation.⁴¹ Stimulation of macrophages with 1,25(OH₂)D₃ abrogates their pro-inflammatory and T cell-recruiting cytokine production.⁴²

VDR signaling promotes anti-inflammatory cytokine secretion and promotes tolerogenic dendritic cell and regulatory T cell differentiation to reduce inflammation.^{43,44} The severe inflammation observed in *Vdr/IL-10* double knockout mice is attributed to an augmented pro-inflammatory T cell response.³⁸ VDR signaling in T cells influences differentiation. Treatment with 1,25(OH)₂D₃ prevents pathogenic Th17 differentiation, along with decreased expression of the pro-inflammatory cytokines IFN γ and IL-17.⁴⁵

Mechanisms: VDR signaling, autophagy, and NF-_KB activation

The VDR target gene *Atg1611*¹³ is an IBD risk gene that regulates intestinal homeostasis. ATG16L1 is a key autophagy protein that mediates autophagosome formation. The *Atg1611-T300A* mutation increases degradation of the ATG16L1 protein and is associated with increased risk of CD. In intestinal inflammation, the *T300A* risk allele results in impaired bacterial clearance and dysbiosis.⁴⁶ We have shown that absence of intestinal epithelial VDR leads to dysbiosis and susceptibility to colitis by reducing ATG16L1 signaling and dampening inflammatory responses. ^{13, 33} Fecal microbiota analysis showed that loss of intestinal VDR shifts the microbiota to a pro-inflammatory profile. These data are consisted with our findings in human IBD samples: decreased VDR in intestine is associated with reduced ATG16L1 expression.¹³

Impaired autophagy in myeloid cells caused by myeloid-specific knockout of *Atg16l1* contributes to intestinal inflammation.^{47,48} These mice exhibit increased susceptibility to

DSS-induced colitis due to impaired autophagy. Both macrophages and dendritic cells deficient in ATG16L1 have impaired processing of MHC class II antigens and thus impaired CD4+ T cell activation.^{47,48} ATG16L1 is also critical for T cell function. Targeted deletion of ATG16L1 in CD4+ T cells results in increased Th2 expansion and impaired Treg development leading to spontaneous intestinal inflammation.⁴⁹ Overall, induction of *Atg1611* expression by VDR¹³ in intestinal epithelial cells may play a critical role in maintaining intestinal homeostasis.

Intestinal inflammation is largely mediated by activation of NF- κ B. VDR is known to downregulate NF- κ B signaling and ameliorate intestinal inflammation through its interaction with NF- κ B and its inhibitor IKK β .⁵⁰ Interaction of VDR with NF- κ B p65 blocks nuclear translocation to lessen the inflammatory response. Loss of VDR in intestinal epithelial cells and MEF cells leads to increased NF- κ B signaling, which promotes the proinflammatory cytokine production.⁵ In macrophage cells, vitamin D activation of VDR promotes interaction of VDR with the NF- κ B p50 subunit prevents LPS-induced proliferation by preventing colocalization of NF- κ B p50 with KLF5.⁵¹ Similarly, vitamin D treatment can prevent LPS-induced placental inflammation by promoting dimerization of VDR with NF- κ B p65.⁵² Thus, suppression of NF- κ B signaling by VDR ameliorates inflammation, which supports the use of vitamin D supplementation in inflammatory diseases like IBD.

Singh *et al* ⁵³ has broadened our understanding of the interaction of VDR and NF- κ B and further supports a critical role of VDR in regulating inflammatory disease. By integrating disease-associated SNPs identified by GWAS with VDR genomic binding sites identified by Chip-Seq, the authors identified 42 SNPs associated with immune phenotypes and VDR binding. Interestingly, most SNPs were not within canonical DR3-type VDREs that bind VDR-RXR. Rather, the transcription factor most frequently found to bind these 42 SNPs was NF- κ B. Activated VDR and NF- κ B share many genomic binding sites, including 22 of the 42 disease-associated SNPs. The data suggests that VDR likely influences transcription indirectly by interacting with NF- κ B and other non-RXR transcription factors. The identification of non-VDRE SNPs bound by VDR opens new avenues of study in assessing the mechanism of VDR signaling in disease.

VDR signaling regulates the gut microbiota

The human microbiome modulates numerous aspects of host physiology and is a critical factor in the development of IBD.^{54,55} Vitamin D may benefit the intestinal microbiome⁵⁶ and improve glucose homeostasis in diabetes.⁵⁷ We are pioneers in studying how intestinal VDR expression influences the microbiome.^{5,13} From the results of our studies utilizing mouse models of *Salmonella*-colitis, and mono-association of the commensal *E. coli* F18 in originally germ-free mice models, we have begun to understand that enteric bacteria activate VDR signaling.⁵ We have demonstrated that VDR expression protects the host from invasive pathogens and maintains intestinal homeostasis.^{5,58} Our recent study⁵⁹ demonstrates that human *Vdr* gene variation is a key host factor influencing the gut microbiome. Meta-analysis in humans shows *Parabacteroides* to be the most significant taxon correlated with *Vdr gene*. Knockout of *Vdr* in mice also leads to changes in *Parabacteroides* abundance.⁵⁹ In

the whole body *Vdr* knockout mice, *Lactobacillus* was depleted in the fecal stool, whereas *Clostridium* and *Bacteroides* were enriched. We identify several important pathways (e.g. metabolism, detoxification, infections, signal transduction, and cancer) affected by *Vdr* status.⁶⁰

Intriguingly, we also demonstrates that intestinal epithelial *Vdr* conditional knockout (*Vdr* ^{IEC}) leads to dysbiosis.¹³ We have reported profound alterations in the gut microbiome profile with increased abundance of *Bacteroidaceae* in *Vdr* ^{IEC} mice.¹³ Likewise, patients with IBD have altered mucosal colonization by the *Bacteroidaceae*.⁶¹ The *Vdr* ^{IEC} mice had dysbiosis and abnormal Paneth cells without any induction of colitis, suggesting the genetic role of intestinal epithelial *Vdr* in the development of chronic inflammation. The microbiota may play a role in the vulnerability of the *Vdr* ^{IEC} mice to colitis. The co-housing data suggest that the absence of intestinal epithelial VDR confers a transmissible risk for DSS-induced colitis. Thus, intestinal VDR contributes to host protection against inflammation.

Vitamin D deficiency in the diet induces intestinal dysbiosis; notable changes include increased *Helicobacter hepaticus* and decreased *Akkermansia municiphila* populations in the gut.³⁴ *H. hepaticus* is known to induce colon cancer and colitis in mice with impaired immunity,⁶² while low *A. municiphila* is decreased in mouse models of colitis and low population level is associated with ulcerative colitis.⁶³ *A. municiphila* is also known to improve barrier function and metabolic health.^{64,65}

The *Vdr* gene, as part of the innate immune response, transcribes for antimicrobial peptides and autophagy-related proteins (e.g. ATG1611),¹³ which are responsible for the deterrence and elimination of infection, and determination of gut microbiome. Vitamin D deficiency reduces production of alpha defensin Defa1 and MMP7 in Paneth cells.³⁴ The Intestinal epithelial *Vdr* conditional knockout (*Vdr* ^{IEC}) leads to dysbiosis.¹³ A low intestinal epithelial VDR protein level led to dysbiosis and impaired autophagy function, accompanied by a reduction in the mRNA and protein levels of ATG1611, an IBD risk gene and regulator of autophagy and the microbiome. Further, Wang *et al.* reported that human *Vdr* gene variation is a key host factor influencing the gut microbiome.⁵⁹ Therefore, vitamin D/VDR is essential in homeostasis and signaling between the microbiota and host in intestinal inflammation.

Gut bacteria can regulate the immune response in a VDR-dependent fashion. We have shown that the protective effect of probiotics against colitis is mediated by epithelial VDR signaling.³³ The secondary bile acid lithocholic acid, which is produced by *Clostridium* bacteria in the gut lumen, inhibits the Th1 immune response and suppresses IFN γ and IL-2 production by activating VDR in T cells.⁶⁶ *Vdr* knockout mice have lower *Clostridium* in the gut,⁶⁰ illustrating the influence of crosstalk between the microbiome and VDR signaling in immunity. Butyrate, a short chain fatty acid produced by gut microbes, can increase epithelial VDR expression and thus decrease production of pro-inflammatory cytokines.¹³

Taken together, these studies provide new insights into the mechanism of VDR regulating in bacterial-host interactions and inflammation.

Vitamin D and human intestinal inflammation

A recent study has demonstrated that serum levels of 25-hydroxy vitamin D of 35 ng/mL or less in UC patients UC in clinical remission are associated with disease relapse. Patients in remission with serum level <35ng/mL had more basal inflammation, were more likely to relapse, and relapsed sooner than patients with levels >35ng/mL.⁶⁷ Increasing levels of vitamin D might reduce their risk for UC relapse.

A small study of patients in remission from CD found that daily oral supplementation with 10,000 IU of 1a,25-dihidroxy vitamin D_3 conferred greater protection against relapse than a smaller dose of 1000 IU/day over 12 months.⁶⁸ A similar study in UC patients with serum vitamin D levels under 30ng/mL found that daily 4000 IU supplementation over 90 days improved patient quality of life but did not significantly improve clinical UC disease activity.

To our knowledge, there is no report on the direct role of vitamin D supplementation on microbiome in IBD patients. A recent study shows that high-dose vitamin D supplementation in humans markedly reduces opportunistic pathogens (e.g. *Gammaproteobacteria*) and increases phylotype richness.⁷⁰ It is believed that vitamin D₃ supplementation reduces inflammation in the intestinal environment, thus diminishing the competitive advantage of opportunistic pathogens (e.g. *Escherichia/Shigella* spp. or *Pseudomonas* spp). In return, the low inflammatory environment allows beneficial bacteria to outcompete pathogens, resulting in increased phylotype richness. These studies show that vitamin D may promote a 'healthy' microbiota profile that potentially confer protection against intestinal inflammation. It will be interesting to learn the change of human VDR protein after using vitamin D supplementation and its role in regulating gut microbiome in health and inflammation.

Conclusion and Future Direction

In conclusion, the vitamin D/VDR signaling is involved in the genetic, environmental, immune, and microbial aspects of IBD, making it a significant protein of interest in understanding IBD pathology and in developing treatments (Fig. 1). As a transcription factor, VDR regulates the expression and signaling of target genes contributing to intestinal inflammation and dysbiosis, including *Atg1611*. Genetic variation of the *Vdr* gene and its targets are associated with inflammation. VDR also directly influences both the host immune system and microbiome through its expression in intestinal epithelial and immune cells. Using vitamin D supplement is conferred protection against relapse of IBD. Thus, intestinal VDR expression and vitamin D/VDR signaling protect again intestinal inflammation and dysbiosis.

The recent progress in microbiome and genetics analyses brings new insights into an old topic on vitamin D/VDR in inflammation. We believe that the following areas should be considered for the future studies and clinical application of vitamin D/VDR in prevention and treatment of IBD.

- Development of novel strategies for the prevention and treatment of human IBD by restoring the healthy host-microbiome interactions through vitamin D/VDR actions.
- Exploration of interactions between the microbiota and vitamin D/VDR that may affect the response to therapies
- Consideration of VDR expression on the response to microbiome-targeted therapies, including fecal microbiota transplant (FMT)
- Exploration of novel roles of vitamin D/VDR in other GI diseases

VDR signaling in the intestine and immune system promotes mucosal homeostasis and protects against inflammatory disease. The current research progress indicates that vitamin D/VDR will likely be a potential therapeutic target for IBD. Understanding the tissue-specific roles of vitamin D/VDR in intestine may offer a diagnostic/prognostic indicator in IBD.

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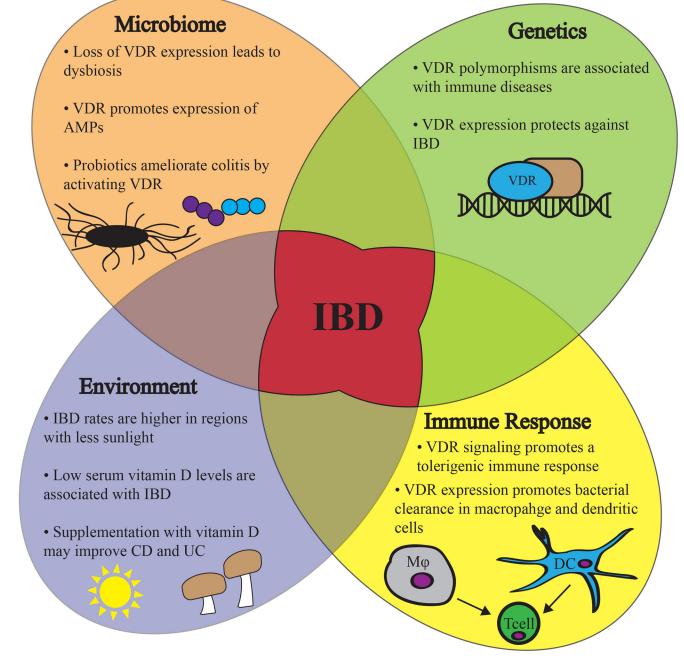


Figure 1.

A working model of VDR signaling involved in the pathogenesis of IBD.

Vitamin D/VDR regulate the genetic, environmental, immune, and microbial aspects in IBD, thus making VDR a significant host factor in IBD pathology and in developing treatments. Susceptibility to IBD is associated with polymorphisms in the *Vdr* gene. Moreover, VDR expression is significantly decreased in IBD patients. Vitamin D deficiency may be an environment trigger contributing to the pathogenesis of IBD. Vitamin D supplement may reduce the relapse of IBD. Vitamin D/VDR appears to be an important immunological

regulator of IBD. The vitamin D/VDR signaling promotes microbial and mucosal homeostasis and protects against inflammatory diseases.