Role of vitamin D on gut microbiota in cystic fibrosis

Mansi Kanhere, Emory University
Benoit Chassaing, Georgia State University
Andrew Gewirtz, Emory University
Vin Tangpricha, Emory University

Journal Title: Journal of Steroid Biochemistry and Molecular Biology
Volume: Volume 175
Publisher: Elsevier: 12 months | 2018-01-01, Pages 82-87
Type of Work: Article | Post-print: After Peer Review
Publisher DOI: 10.1016/j.jsbmb.2016.11.001
Permanent URL: https://pid.emory.edu/ark:/25593/tmsqb

Final published version: http://dx.doi.org/10.1016/j.jsbmb.2016.11.001

Copyright information:
© 2016 Elsevier Ltd
This is an Open Access work distributed under the terms of the Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International License (http://creativecommons.org/licenses/by-nc-nd/4.0/).

Accessed January 21, 2020 10:04 PM EST
Role of vitamin D on gut microbiota in cystic fibrosis

Mansi Kanhere¹, Benoit Chassaing², Andrew T. Gewirtz², and Vin Tangpricha³,⁴

¹Division of Endocrinology, Department of Pediatrics, Emory University School of Medicine, Atlanta, GA, USA
²Center for Inflammation, Immunity, & Infection, Institute for Biomedical Sciences, Georgia State University
³Division of Endocrinology, Metabolism, and Lipids, Department of Medicine, Emory University School of Medicine, Atlanta, GA, USA
⁴Atlanta VA Medical Center, Decatur, GA, USA

Abstract

This review explores the potential for vitamin D to favorably alter the gut microbiota, given emerging evidence of the role of vitamin D in controlling mucosal inflammation in the gut. It will focus on cystic fibrosis (CF) patients, a population with both vitamin D deficiency due to gut malabsorption and an altered gut microbiota composition. Recent evidence shows that vitamin D acts to maintain the integrity of the gut mucosal barrier by enhancement of intercellular junctions that control mucosal permeability and reduction of pro-inflammatory cytokines such as IL-8. In addition, vitamin D receptor-mediated signaling has been shown to inhibit inflammation-induced apoptosis of intestinal epithelial cells. As a result of these effects on the intestinal mucosa, maintenance of sufficient vitamin D status may be essential for the development of a healthy gut microbiota, particularly in conditions defined by chronic mucosal inflammation such as CF. We hypothesize here that using high dose of vitamin D may be used to favorably manipulate the aberrant mucosa seen in patients with CF. This may result in improved clinical outcomes in association with a low inflammatory environment that allows beneficial bacteria to outcompete opportunistic pathogens. Current evidence is sparse but encouraging, and additional evidence is needed to establish vitamin D as a therapeutic approach for gut microbiota modification.

Keywords

vitamin D; intestinal microbiota; cystic fibrosis; vitamin D receptor; inflammation

Corresponding Author: Vin Tangpricha, MD, PhD, 101 Woodruff Circle NE- WMRB1301, Atlanta GA 30322, Ph (404) 727-7254, Fax (404) 592-6257, vin.tangpricha@emory.edu.

Conflicts of interest: None

Publisher's Disclaimer: This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final citable form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.
Introduction

Cystic fibrosis (CF), an autosomal recessive condition, is caused by mutations in the cystic fibrosis transmembrane conductance regulator (CFTR) gene. CFTR is an ion channel important for the maintenance of chloride transport, essential for normal functioning of epithelial surfaces of lung and pancreas[1]. CF is characterized by recurrent lung infections, malabsorption and failure to thrive related to abnormal mucosa and chronic inflammation in the respiratory and digestive systems.

A common co-morbidity of CF is vitamin D deficiency, resulting from fat malabsorption secondary to exocrine pancreatic insufficiency, inadequate sun exposure, and inadequate dietary vitamin D intake [2]. An important extra-skeletal function of vitamin D is its role in immune regulation, such as upregulation of antimicrobial peptides expression (cathelicidins and B-defensins) and turning off effector T cells Th1 and Th17 [3-5]. More recently, vitamin D was identified as a factor influencing the gut microbiota, through the vitamin D receptor (VDR) [6, 7]. Mechanisms underlying this association involve the role of VDR in maintaining the integrity of the intestinal mucosal barrier [8].

Microbiota refers to the complex community of microorganism that colonize the human body, including the skin, respiratory, gastrointestinal and urogenital systems [9]. The intestinal microbiota contains $10^{14}$ microorganisms with more than 3 million genes, collectively referred to as the microbiome [10]. This complex and dynamic microbial community plays a central role in human health, including macronutrient metabolism, nutrient capture and immune function. In addition to non-modifiable factors that affect the microbiota, such as maternal microbial composition and host genetics, it is profoundly influenced by modifiable factors such as diet and medications, especially antibiotics [11]. Moreover, disruption of the gut microbiota (or dysbiosis) has been linked to both intestinal inflammation, such as inflammatory bowel disease, and extra-intestinal conditions associated with chronic inflammation and metabolic dysfunction, such as obesity and metabolic syndrome[12-16]. There is also emerging evidence that aberrant intestinal mucosa in CF patients leads to a state of intestinal dysbiosis [17, 18], likely due to a combination of abnormal mucus secretion and repetitive antibiotic exposures for recurrent sino-pulmonary infections treatment.

Given the emerging studies on the influence of vitamin D on gut microbiota, this review explores the current evidence for modifying the gut microbiota to influence clinical outcomes in CF. Vitamin D supplementation may provide an easy and inexpensive potential option for positive modulation of the gut microbiota that would lead to improvement of CF-related symptoms.

Methods

A PubMed search was performed to select relevant publications, using the following search terms: dysbiosis, gut microbiota, cystic fibrosis, vitamin D, probiotics, mucosal inflammation and vitamin D receptor. We examined all studies in humans and in rodent models.
Altered gut microbiota in CF

Emerging research suggests that intestinal microbiota may be disrupted in individuals with CF [17], with multiple independent groups showing a significant decrease in species richness and enrichment of potentially pathogenic species in individuals with CF compared to healthy controls [18-22]. Key findings of these studies have been reviewed in TABLE 1.

1. Review of murine studies—Brodie et al. studied the gut microbiota composition in CFTR knock out mice compared to wild type mice (WT) and found that CFTR KO mice displayed significantly less bacterial richness and diversity compared to WT animals. Moreover, they found that the loss of functional CFTR is associated with an increase in pathogenic bacteria, such as Mycobacteria and Bacteroides fragilis, and a decrease in bacteria that provide protection at mucosal surfaces, such as Acinetobacter lwofii and members of the order Lactobacilliales [23]. Similarly, Bazett et al. looked at the intestinal microbiome in CFTR knockout mice with three different phenotypes and found alterations in intestinal structure, bacterial overgrowth, and dysbiosis as compared to wild type mice[24].

Review of human studies—Cross-sectional and longitudinal observation studies looking specifically at the gut microbiota in CF have indicated a trend of suppression in the potentially beneficial Bifidobacterium species accompanied by a trend of increase in potentially pathogenic organisms -Fusobacterium and Enterobacteriaceae species [18, 21]. Members of the Bifidobacterium species are considered a marker of intestinal health and a decrease in abundance has been associated with intestinal diseases such as inflammatory bowel disease and colorectal cancer and extra-intestinal diseases, such as obesity and asthma[25]. In a recent study examining gut microbial communities in children with CF compared to age matched healthy controls, Nielsen et al. found that gut dysbiosis in CF patients was present in early childhood with further digression with advancing age. They observed an increase in the relative abundance of Ruminococcaceae family and Alistipes genus (potentially important beneficial strains for mucosal health) with increasing age in healthy children, but not in CF. In contrast, this analysis revealed that potential pathogenic organisms, such as Escherichia and Shigella (Proteobacteria), Enterococcus, Veillonella, Megasphaera, Clostridium group XI and Blautia were more abundant in CF compared to healthy controls[19].

Potential interest in gut microbiota manipulation for CF management—The gut microbiota is involved in both regulatory and pro-inflammatory immune pathways, via interaction with intestinal mucosal receptors [26, 27]. Although direct evidence is limited, recent studies imply a potential role of the gut microbiota in CF management, via affecting inflammatory and immune response. It is well established that repeated exacerbations of pulmonary symptoms in persons with CF are associated with a progressive decline in lung function. Carmody et al. described that the onset of symptoms of exacerbation may be preceded by marked shifts in the baseline microbial communities [28]. The suppressed species richness found by Duytschaever et al. [18] suggest that biodiversity of gut microbiota remains lower in children with CF compared to healthy siblings. A growing body of evidence suggests that microbial diversity is an important marker of gut health, with a low
diversity associated with intestinal inflammation and several diseases, from inflammatory bowel disease to obesity [16, 29-31]. Further, in patients with CF, Madan et al. have shown that gut microbiota diversity decrease and loss of beneficial microbial species were associated with the first CF pulmonary exacerbation, as well as with initial Pseudomonas colonization. Thus, it may be reasonable to hypothesize that interventions that alter the gut microbiota in a beneficial way may have a potential role in clinical management in CF. Efrati et al. found that probiotic supplementation in patients with CF who had mild-moderate lung disease and were chronically infected with Pseudomonas aeruginosa resulted in a significant decline in the rate of pulmonary exacerbations for 6 months post-treatment compared to the prior two years, with none of the patients experiencing pulmonary exacerbation during the probiotic treatment [32]. In a similar study, Guarino et al. assessed the effect of the probiotic Lactobacillus rhamnosus strain GG (LGG) on pulmonary outcomes in nineteen children with CF through a prospective, randomized, placebo-controlled, crossover study. They found a reduction in pulmonary exacerbations and hospital admissions and an improvement in the forced respiratory volume in 1 second (FEV1) in patients treated with LGG [33]. While the exact mechanism is unclear, these studies suggest that probiotics may affect the gut microbiota by locally improving intestinal barrier function [34] and preventing membrane barrier disruption by pathogens [35]. These data suggest that manipulation of gut microbiota have the potential to improve respiratory outcomes in CF.

Vitamin D and the gut microbiota

A growing body of research suggests that gut epithelial VDR signaling plays a central role in maintaining the integrity of the intestinal mucosa. The proposed mechanism is through the inhibition of inflammation-induced epithelial cell apoptosis and enhancement of intercellular junctions [8, 36].

Review of murine studies—Studies in murine models report that vitamin D deficiency increased the severity of colitis and bacterial numbers in the colons of mice, with these deleterious effects being reversed by vitamin D supplementation [37] [38]. DSS(Dextran sulfate sodium) increases mucosal permeability and causes disruption of the mucosal barrier in intestinal epithelial cells, thus the DSS induced experimental colitis model has been widely used to study the effects of vitamin D on mucosal injury[39]. TABLE 2 reviews studies that looked at the effects of VDR and 1,25(OH)2D3 on protecting mice from DSS induced colitis. These studies elucidate that vitamin D deficiency predisposes mice to severe colitis as compared to their wild type counterparts [38, 40, 41]. Ooi et al. also demonstrated that treatment of 1,25(OH)2D3 deficient mice with 125(OH)2D3 decreased the severity of colitis[40]. Further, Zhang et al. showed that 1,25(OH)2D3 prevented DSS induced disruption of tight junctions in colonic mucosa[42]. This data provides compelling evidence that vitamin D has an important role in maintaining epithelial integrity and regulating mucosal injury. It can thus be postulated that vitamin D deficiency leads to a disruption of the intestinal epithelial barrier and thus predisposes to dysbiosis. This was demonstrated in a recent study by Sun et al. who found that the lack of VDR in a VDR KO mice model is associated with intestinal dysbiosis and that VDR regulates the composition and functions of the gut microbiota. Precisely, they found a dramatic reduction of lactic acid bacteria in VDR knockout mice, associated with an increased abundance of Bacteroides and Clostridium [6].
Clostridium has been associated with intestinal virulence and enteritis and Bacteroides, may predispose the mice colon to cancer [43, 44].

Further support of the role of vitamin D in intestinal dysbiosis is provided by the work of Sun et al. who showed that decreased expression of intestinal epithelial VDR disturbs microbial homeostasis, leading to fewer butyrate producing bacteria and predisposing VDR knock out mice to chemically-induced colitis [7]. Butyrate, a byproduct of carbohydrate breakdown by microbiota, has a well established role in preventing mucosal inflammation and inhibiting colorectal cancer [45, 46]. Moreover, Manor et al. found an increased capacity for catabolism of butyrate in the gut microbiota of children with CF, which may lead to increased mucosal inflammation [47]. In addition, Sun et al. found that administration of the bacterial product butyrate increased intestinal VDR expression and suppressed inflammation in a colitis model. They propose that lack of VDR causes dysbiosis by impairing ATG16L1, an important regulator for autophagy [7].

Review of human studies—While human studies are limited, similar observations have been made, whereby high dose vitamin D₃ supplementation showed a beneficial effect on the human gut microbiota with a marked reduction in typical opportunistic pathogens and an increase in phylotype richness[48]. Healthy volunteers given vitamin D₃ supplementation experienced changes in the gut microbiota in the upper GI tract; decreased relative abundance of pathogenic species including Pseudomonas, Escherichia and Shigella and increased bacterial richness [48]. The possible mechanisms by which vitamin D may serve as a therapeutic intervention to decrease intestinal mucosal inflammation and thus influence gut microbiota are illustrated in Figure 1.

Potential role for vitamin D to modify gut microbiota and pulmonary outcomes in CF—Crites et al. showed that loss of functional CFTR in intestinal epithelial cells leads to a pro-inflammatory status specifically increased IL-6 and IL-8 secretion [49]. Studies reviewed above establish that an underlying state of chronic inflammation in CF is associated with an altered intestinal microbiota. It can thus be postulated that vitamin D may attenuate intestinal inflammation and thus favorably modify the altered gut microbiota seen in the CF gut. This theory is supported by the work of Morin et al. who looked at the effects of vitamin D₃ and its metabolites in CFTR knockdown intestinal epithelial cells. Interestingly, they observed that 1,25(OH)²D₃ leads to an inhibition of IL-8 and reduces cytokine-induced NF-κB nuclear translocation thus resulting in a suppression of inflammatory mediators[50].

The role for vitamin D in affecting pulmonary function was proposed by the Third National Health and Nutrition Survey (NHANES III) data, which showed an association between serum 25-hydroxyvitamin D levels and pulmonary function in healthy adults[51]. Further, Grossman et al. found that in adults hospitalized for a pulmonary exacerbation, a single, oral bolus of cholecalciferol was associated with better antibiotic therapy-free days, hospital-free days and 1 year survival compared to subjects in the placebo group [52]. In a longitudinal study by McCauley et al. in the pediatric population, higher 25(OH)D levels were associated with lower rates of pulmonary exacerbations[53]. Since these studies show that vitamin D has a role in pulmonary outcomes in the non-CF population it may be reasonable to
hypothesize that these effects of vitamin D may be inferred to affect lung function in CF. In fact, recent studies suggest that vitamin D may have a role on pulmonary function and even pulmonary infections in CF. Sexauer et al. found that serum 25(OH)D levels showed a significant correlation with forced expiratory volume in 1 second (FEV 1) and forced vital capacity (FVC) in a group of 597 children and adults with CF [54]. In addition, Vanstone et al. found a significant association between serum 25(OH)D levels and annual frequency of pulmonary exacerbations in pediatric CF patients [55].

**Conclusion**

Review of studies above suggests that intestinal dysbiosis is a hallmark of CF and vitamin D therapy in CF patients with vitamin D deficiency may result in restoration of gut microbial homoeostasis, with reduction in opportunistic pathogens and increase in microbiota diversity. Given the potential role of vitamin D on pulmonary function, this in turn may have implications for the improvement of clinical outcomes in patients with CF, with reduction in respiratory infections and pulmonary exacerbations, and ultimately survival improvement. Above studies support the hypothesis that intestinal microbial dysbiosis is predominant in CF. However, the clinical significance of dysbiosis in CF are not entirely understood. There are also no studies to date to evaluate if vitamin D deficiency predisposes CF patients to greater dysbiosis and no clinical trials to show an improvement in gut microbial homeostasis with replacement with vitamin D. Further research is required to look at the role of vitamin D deficiency on modifying the gut microbiota in patients with CF and if supplementation with vitamin D can manipulate this altered microbiota.

**Acknowledgments**

Financial support and sponsorship: Supported by a grant from the National Center for Advancing Translational Sciences of the National Institutes of Health under Award Number UL1TR000454 (VT). The content is solely the views of the authors and does not necessarily represent the official views of the National Institutes of Health.

**References**

2. Chesdachai S, Tangpricha V. Treatment of vitamin D deficiency in cystic fibrosis. The Journal of steroid biochemistry and molecular biology. 2015


41. Froicu M, Cantorna MT. Vitamin D and the vitamin D receptor are critical for control of the innate immune response to colonic injury. BMC immunology. 2007; 8:5. [PubMed: 17397543]


Highlights

- Vitamin D plays a vital role in maintaining the integrity of the gut epithelium.
- The mechanism underlying this association involves the reduction of proinflammatory cytokines such as IL-8 and inhibition of inflammation-induced apoptosis of intestinal epithelial cells by vitamin D.
- Intestinal dysbiosis is prevalent in CF owing to multiple factors including malabsorption of nutrients and recurrent antibiotics.
- VDR regulates the composition and functions of the gut microbiota and decreased expression of intestinal epithelial VDR disrupts microbial homeostasis.
- Vitamin D could be a potential treatment option for altering dysbiosis in CF, but additional trials are needed to evaluate this.
Figure 1. A schematic representation of mechanisms by which vitamin D may decrease intestinal mucosal inflammation and thus alter dysbiosis in CF
Description: Microbial dysbiosis is a hallmark of the CF gut underlying to chronic mucosal inflammation. Vitamin D has been proposed to decrease intestinal inflammation through the inhibition of inflammation-induced epithelial cell apoptosis and enhancement of intercellular junctions that are important in maintaining mucosal permeability. In addition it reduces cytokine-induced NF-κB nuclear translocation thus resulting in a suppression of inflammatory mediators. Together, these effects may restore gastrointestinal homeostasis and prevent intestinal dysbiosis in CF.
### Table 1

Studies examining alteration of gut microbiota in patients with CF

<table>
<thead>
<tr>
<th>TYPE OF STUDY</th>
<th>STUDY DESIGN (n)</th>
<th>EFFECT ON MICROBIOTA</th>
<th>CLINICAL SIGNIFICANCE</th>
<th>REFERENCE</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cross-sectional and Longitudinal</td>
<td>Compared fecal microbiota between patients with CF (21) and healthy siblings (24)</td>
<td>Increased <em>Enterobacteriaceae Gammaproteobacteria</em> Decreased <em>Bacteroides Prevotella</em></td>
<td>Not assessed</td>
<td>[18]</td>
</tr>
<tr>
<td>Prospective longitudinal analysis</td>
<td>Gut microbiota samples collected from birth to 34 months (13)</td>
<td>Increased <em>Not applicable</em> Decreased <em>Parabacteroides Bacteroides</em> Bifidobacterium</td>
<td>Decline in the abundance of Parabacteroides s prior to initial Pseudomonas aeruginosa colonization ($p &lt; 0.001$)</td>
<td>[22]</td>
</tr>
<tr>
<td>Prospective longitudinal analysis</td>
<td>Comparison of gut microbiota between children with CF (23) aged from 0–18 years old and healthy controls (35)</td>
<td>Increased <em>Proteobacteria Enterococcus, Veillonella Megasphaera Clostridium group XI Blautia</em> Decreased <em>Ruminococcaceae Alistipes</em></td>
<td>Not assessed</td>
<td>[19]</td>
</tr>
<tr>
<td>Prospective longitudinal analysis</td>
<td>Analysis of gut microbiota from patients with CF and healthy siblings (21)</td>
<td>Decreased <em>Bifidobacterium Clostridium cluster XIVa</em></td>
<td>Not assessed</td>
<td>[21]</td>
</tr>
</tbody>
</table>
Table 2
Role of VDR/1,25(OH)₂D₃ in protection from experimental colitis

<table>
<thead>
<tr>
<th>MURINE MODEL AND GENOTYPE</th>
<th>STUDY DESIGN</th>
<th>STUDY FINDINGS</th>
<th>REFERENCE</th>
</tr>
</thead>
<tbody>
<tr>
<td>C57BL/6 VDR KO vs. WT mice</td>
<td>• Exposed to DSS *</td>
<td>• Increased mortality of the VDR KO mice at doses of DSS that only caused a mild form of colitis in WT mice.</td>
<td>[41]</td>
</tr>
<tr>
<td></td>
<td></td>
<td>• Administration of 1,25(OH)₂D₃ resulted in decreased weight loss and improved colonic injury scores in the DSS-induced inflammation in WT mice compared to controls.</td>
<td></td>
</tr>
<tr>
<td>C57BL/6 Cyp27b1 (CYP) KO and WT mice</td>
<td>• Mice were exposed to DSS to induce colitis • 1,25(OH)₂D₃ was supplemented to some of the mice in the CYP KO group.</td>
<td>• CYP KO mice were more susceptible to DSS colitis and had greater intestinal permeability after exposure to DSS. • CYP KO mice treated with 1,25(OH)₂D₃ showed decreased severity of colitis as compared to their untreated counterparts.</td>
<td>[40]</td>
</tr>
<tr>
<td>C57BL/6 mice</td>
<td>Mice were fed on vitamin D deficient or vitamin D sufficient diets and then treated with DSS to induce colitis</td>
<td>• DSS induced weight loss and colitis was more prominent in vitamin D deficient mice • Vitamin D deficient mice showed an inability to contain colonic enteric bacteria even in the absence of experimental colitis</td>
<td>[38]</td>
</tr>
</tbody>
</table>

*DSS colitis model refers to mice with experimental colitis produced by administration of 2.5% dextran sodium sulfate (DSS) in drinking water.

ABBREVIATIONS: VDR KO- Vitamin D receptor knockout, WT – wild type, Cyp KO-Cyp27B1 knock out (unable to produce 1,25(OH)₂D₃ due to defective 1α-hydroxylase enzyme)