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Review

Cannabis and Endometriosis: The Roles of the Gut Microbiota and the Endocannabinoid System

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Abstract: There is a growing interest in the endocannabinoid system and the gut microbiota. Both areas have been suggested to play a role in endometriosis, however this is yet to be studied comprehensively. The purpose of this narrative review is to identify the gaps in endometriosis research in the context of these two areas. It underscores the limited knowledge regarding the involvement of the endocannabinoid system and gut microbiota in endometriosis. Further research in these areas is essential to unlock the therapeutic potential of cannabis and enhance the understanding of endometriosis pathogenesis.

Keywords: endometriosis; the endocannabinoid system; CB₁; CB₂; gut microbiota

Introduction

Endometriosis is a disease defined by the presence of endometrial-like tissue outside of the uterus[1]. Endometriosis is estimated to occur in up to 10% of women and those assigned female at birth during their reproductive years [2,3], with 11% of women being diagnosed by age 44 in Australia [4]. [4]. [2,3], with 11% of women being diagnosed by age 44 in Australia [4]. [4]. Endometriosis-related chronic pelvic pain refers to a variety of pain symptoms including dysmenorrhea (period pain), dyspareunia (pain during sexual intercourse), fatigue, dyschezia (pain on bowel motions, and dysuria (pain on urination)[5–7]. [5–7]. People with endometriosis are often diagnosed with various comorbidities including irritable bowel syndrome, rheumatoid arthritis, psoriasis, anxiety, depression[8] and chronic fatigue syndrome[9]. The debilitating symptoms of endometriosis impact social activities, work and career progression, finances, academic studies, mental health, emotional health and sexual/romantic relationships, ultimately impacting quality of life [5,10–13]. Furthermore, the economic burden of endometriosis impacts the Australian economy, costing AUD\$30,000 per woman per year, with the most predominant factor being lost productivity, directly correlated with pain severity. Internationally this varies depending on different national economies, from US\$1459 to US\$20,239 per year.

One of the known mechanisms through which endometriosis symptoms occur is a result of repetitive deposition of endometrial-like tissue and active breakdown of endometriotic lesions, which in turn leads to an inflammatory cascade alongside the development of adhesions and scar tissue, as well as other factors, causing pelvic pain.[1,14]. The inflammation-mediated pain has been correlated with high levels of cyclooxygenase-2 (Cox-2) and tumour necrosis factor alpha (TNF- α) in spinal cords and brains, alongside peripheral macrophages in a murine model[15].

Endometriosis is often characterized by the altered efficiency of progesterone and estrogen hormones, leading to progesterone resistance, as observed in people with endometriosis not responding to the use of progestins, and excess estrogen levels. This depicts an imbalance which initiates local infiltration of immune cells and inflammation[16,17]. Newly established cells resulting

from inflammation activate various pathways of cell proliferation, angiogenesis, metastasis and invasion. As estrogen and progesterone receptors are responsible for proliferation and differentiation in the endometrium, imbalances result in changes in the expression of estrogen-metabolising enzymes, promoting progesterone resistance in endometriotic lesions[18].

Current treatments for endometriosis are generally considered suboptimal by those with the disease [19,20] with concerns about lack of effectiveness and problematic side effects of many medications for pelvic pain [21]. Opioid analgesics are not recommended for chronic pelvic pain due to both a lack of efficacy and safety concerns with respect to ongoing use[22]. However, despite this they continue to be prescribed; people with endometriosis have a four times greater risk of chronic opioid use compared to those without[23], and opioids are often prescribed alongside benzodiazepines[24]. Both opioids[25] and benzodiazepines[26] present significant risk of cognitive impairment, addiction and severe withdrawal symptoms, with a combination of these drugs significantly increasing the risk of overdose[27]. Surgery is considered a viable and common effective treatment [28] but often has significant costs, long waiting times [29], and substantial recurrence rates, even with expert endometriosis surgeons[30]. Access to surgery is even more reduced in developing countries and remote and rural locations[31].

Due to these issues, novel, pain management options are considered an urgent research and clinical priority in endometriosis[32–34]. Limited access to surgery often leads to people with endometriosis employing self-management strategies. The use of cannabis, either illicitly or legally, is becoming a relatively popular self-management strategy in those with endometriosis, with substantial self-report data on reduction of symptoms [35]. This review explores the potential mechanisms of action by which cannabis may modulate endometriosis symptoms.

The growing interest in the gut microbiota and its influence on various metabolic and inflammatory diseases poses the question of gut microbiota involvement in modulation and/or pathogenesis of endometriosis. Studies have also reported interactions between gut microbes and the endocannabinoid system (ECS). Both of these aspects will be explored in this review.

1.1. Endometriosis and the Endocannabinoid System

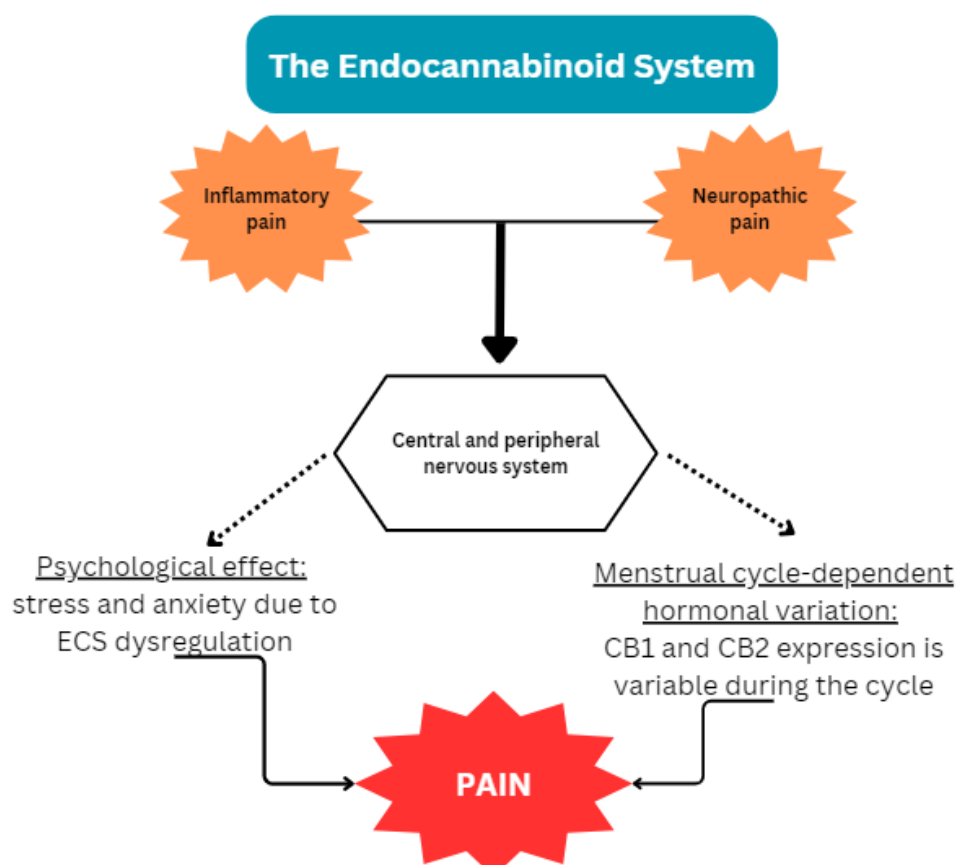


Figure 1. The endocannabinoid system (ECS) in pain. As the ECS is spread throughout the central and peripheral nervous system, a psychological effect is reported as a result of endocannabinoid dysregulation. This is coupled with the menstrual-cycle dependent hormonal variation occurring due to varying levels of cannabinoid receptor 1 (CB₁) and cannabinoid receptor 2 (CB₂). This figure is attributed to [36].

Research into the *Cannabis* genus during the 1990s contributed to the scientific discovery of ECS [37]. The ECS is a complex signaling system that is comprised of three major components: G-protein-coupled cannabinoid receptors (CB₁, CB₂, endocannabinoids (endogenously produced cannabinoids, including anandamide (AEA) and 2-arachidonoyl glycerol (2-AG)), ion channel transient receptor potential vanilloid 1 (TRPV1)[38] and the enzymes involved in the synthesis and catabolism of endocannabinoids [39–41].

Table 1. Key endocannabinoids and related information.

Endocannabinoid	Abbreviation	Roles
Cannabinoid receptor 1	CB ₁	G-protein coupled receptor found on cells in the central and peripheral nervous system – involved in processes of mood, appetite, learning, memory and pain.
Cannabinoid receptor 2	CB ₂	G-protein coupled receptor found on cells in the central and peripheral nervous system – expressed during active inflammation
Anandamide	AEA	Agonist towards CB ₁ and CB ₂ receptors
Arachidonoyl glycerol	2-AG	Agonist towards CB ₁ and CB ₂ receptors
Transient receptor potential vanilloid 1	TRPV1	Ion channel involved in endocannabinoid signalling
N-oleoylethanolamine	OEA	Fatty acid ethanolamine – involved in food intake, inflammation and pain
N-palmitoylethanolamine	PEA	Low affinity for CB ₁ , CB ₂ and TRPV1 – induces peripheral antinociception upon activating CB ₁ and CB ₂
Fatty acid amide hydrolase	FAAH	Metabolises endogenous ligands
Monoglyceride lipase	MGL	Metabolises endogenous ligands

Research to date has demonstrated that the ECS is involved in homeostasis and regulation *via* neuromodulatory activity, physiological processes such as digestion [42], immune function [43], nociception [44], appetite regulation[45], cardiovascular and respiratory function[46] and sleep wake cycles[47].

The ECS influences pain modulation, making it a potential target for the treatment of chronic pain conditions such as endometriosis (Figure 1). Activation of CB₁ and CB₂ receptors by endocannabinoids or exogenous cannabinoids can suppress nociceptive processing and induce analgesia[48–51]. Similar observations were made in preclinical studies showing the analgesic effects of cannabinoid agonists on neuropathic pain[52–54]. However, clinical trials have presented mixed results in terms of the efficacy of cannabinoids in pain modulation. In a prospective randomized placebo-controlled trial, smoked cannabis (3.56% THC – participants smoked three cigarettes daily over a 4-day period) reduced daily pain experienced by adults with HIV-associated sensory neuropathy[55]. In contrast, the administration of Sativex, a low-dose THC (2.7mg/100µL) and CBD (2.5mg/100µL) combination oro-mucosal spray (used over an 8-week study period – drugs were titrated per patient) in adults with multiple sclerosis showed non-significant differences between the treatment and placebo group[56]. As studies often employ different modes of administration, which affects bioavailability, the resulting effect of cannabis varies, therefore further studies will be required to fully understand the mechanisms and therapeutic potential of the ECS.

Endocannabinoid receptors have been found throughout female reproductive tissue[41] and the use of exogenous cannabinoids may alleviate endometriosis-associated pain [57,58]. The ECS also regulates endometriosis mechanisms such as inflammation, angiogenesis, apoptosis, endometrial hyperproliferation and fibrosis [41,57–63], and this may be a potential pathway by which endometriosis lesions re-occur post surgery.

Cannabinoid receptors, specifically CB₁ receptors, are expressed on both the somata and fibres of sensory and sympathetic neurons that innervate endometriotic growths[64]. Activation of CB₁ receptors has been linked to a reduction in endometriosis-induced hyperalgesia, while blocking CB₁ receptors increases pain[65]. Moreover, elevated levels of endocannabinoids, such as AEA and 2-AG, have been observed in people with endometriosis[41]. However, the expression of CB₁ receptors in endometrial stromal cells is lower in endometriosis compared to healthy controls, suggesting a negative feedback loop that may impair the ability of endocannabinoids to control pain[66]. The levels of endocannabinoids in people with endometriosis vary across the menstrual cycle, and this is thought to be influenced by the on-demand synthesis and degradation of enzymes[67]. In the secretory phase, however, CB₁ regulation is absent, reflecting an impaired response to progesterone levels[68]. Table 1 outlines the findings of these key studies.

1.1.1. Endocannabinoid Deficiency in Endometriosis

Endometriosis has been described by some researchers as an “endocannabinoid deficiency”[68]. Plasma levels of endocannabinoid ligands AEA, 2-AG, *N*-oleoylethanolamine (OEA) and *N*-palmitoylethanolamine (PEA) fluctuate in people with endometriosis[68]. When associating these endocannabinoids with endometriosis-associated pain severity, a correlation was noted between both increased endocannabinoid ligands in plasma and decreased local CB₁ receptor expression in people with endometriosis.[68]. These levels were studied in association with common endometriosis symptoms, where it was identified that elevated levels of AEA was linked with moderate-to-severe dysmenorrhea while elevated levels of PEA were found in people with moderate-to-severe dyspareunia[68]. This indicates a potential dysregulation in the ECS, suggesting an altered signalling process in response to endometriosis-related pain.

1.1.2. Potential Interplay of the ECS and Endometriosis

The peritoneal microenvironment is often studied in light of the establishment and progression of endometriosis. Inflammatory cytokines (interleukin-1 β (IL-1 β), interleukin-6 (IL-6), insulin-like growth factor (IGF-1) and tumour necrosis factor- α (TNF- α) are observed to be at a higher concentration in the peritoneal fluid of people with endometriosis [40,69–71]. The production of proinflammatory cytokines may occur as a result of disease-modified macrophages producing IGF-1, as high concentrations of IGF-1 were found in the peritoneal fluid of people with endometriosis[15]. This in turn leads to people experiencing hyperalgesia, which has been correlated with altered expression of TRPV1 in the peritoneum of people with endometriosis [72]. Levels of 2-AG and AEA have also been observed to be significantly higher in the peritoneal fluid of people with endometriosis, and this was associated with relatively higher abdominal pain[73]. The protective role of the ECS was observed in people with endometriosis, where immunohistochemistry revealed that cannabinoid agonists inhibit endometrial cell proliferation[61]. In contrast, previous studies have also shown the involvement of the ECS in endometriosis modulation and progression. ECS modulation in innervation of ectopic uterine growths was demonstrated through the abundance of CB₁ receptors on sensory and sympathetic fibres innervating ectopic growths, dorsal root ganglia and coeliac ganglia[41]. This was also observed in the epithelial cells in ovarian endometriotic lesions[74]. Such studies demonstrate the complex and contradictory role of the ECS in pain modulation, warranting more research in this area to understand the mechanisms behind the interaction of the ECS and endometriosis.

1.1.3. Exogenous cannabinoids

The intricate and multifaceted role of the ECS in pain modulation has spurred investigations into the potential impact of exogenous cannabinoids for pain management. While clinical trials are yet to be conducted, preclinical studies have explored the effects of the exogenous cannabinoid delta-9-tetrahydrocannabinol (THC) on pain management. In murine models of endometriosis, repetitive administration of botanically-derived THC demonstrated protective properties[75]. It hindered the growth of ectopic endometrial tissue while alleviating mechanical hypersensitivity in the caudal abdominal region. In an experimental mouse model varying concentrations of cannabidiol (CBD) were found to significantly reduce endometriotic implant surface area, alongside proinflammatory cytokine levels, including IL-6 and TNF- α [76]. A similar study showed that CBD administration

reduced endometriotic lesion diameter, volume and area *in vivo*, demonstrating antioxidant effects by reducing lipid peroxidation[77]. These promising findings highlight the potential therapeutic benefits of CBD and THC for endometriosis-associated pain, warranting the need for human studies.

Table 2. A summary of recent studies on endometriosis and the endocannabinoid system.

Author	Year	Type of study	Endocannabinoid/cannabinoid	Methodology	Findings
Andrieu, et al.	2022	Human – endometriosis	2-AG and AEA	LCMS analysis of endocannabinoids ELISA analysis of cytokines	An increase in abdominal pain was associated with a high level of 2-AG and a low level of AEA in the peritoneal fluid of people with endometriosis
Sanchez, et al.	2016	Human - endometriosis	CB ₁	PCR analysis of menstrual cups of people with endometriosis	The availability of endocannabinoids in the human endometrium is regulated by the availability of enzymes across the menstrual cycle. The absence of CB ₁ regulation in endometriosis in the secretory phase might reflect an impaired response to progesterone.
Rocha, et al.	2011	Human - endometriosis	TRPV1	Immunohistochemistry analysis of rectouterine peritoneum	There is a correlation between greater expression of TRPV1 in the peritoneum of people with endometriosis who experience chronic pelvic pain compared to those with endometriosis who do not.
Bilgic, et al.	2017	Human endometrial archive samples	Cannabinoid agonists	Immunohistochemistry analysis	Cannabinoid agonists inhibit endometriotic cell proliferation, modulating apoptosis of endometriotic cells.
Okten, et al.	2023	Female Wistar albino female rats	CBD	Immunohistochemical staining	CBD reduced endometriotic implant surface area and proinflammatory cytokine levels.
Escudero-Lara, et al.	2020	Female C57Bl/6J mice	THC	Immunostaining	Repetitive administration of THC hindered ectopic endometriotic growths, alleviating hypersensitivity.
Genovese, et al.	2022	Sprague-Dawley rats	CBD	Histological analysis ELISA	CBD reduced endometriotic lesion diameter and demonstrated antioxidant effects, as viewed by downregulated expression of MMP-9, iNOS and TGFβ).

Abbreviations: 2-AG = 2-arachidonoyl glycerol; AEA = anandamide; CB₁ = cannabinoid receptor 1; TRPV1 = transient receptor vanilloid 1; CBD = cannabidiol; THC = delta-9-tetrahydrocannabinol; LCMS = liquid chromatography-mass chromatography; ELISA = enzyme-linked immunosorbent assay; PCR = polymerase chain reaction; MMP-9 = matrix metalloproteinase-9; iNOS = inducible nitric oxide synthase; TGFβ = transforming growth factor beta.

1.2. The Endocannabinoid System and the Gut Microbiota

The gut microbiota refers to the collection of trillions of microorganisms, including bacteria, archaea, viruses and fungi. The gut microbiota plays a crucial role in human health and disease, impacting immunomodulation and inflammatory processes[78].

In humans, the gut microbiota is dominated by four bacterial phyla: *Firmicutes*, *Bacteroidetes*, *Actinobacteria* and *Proteobacteria*[79]. As the phylotypic composition of these can vary amongst individuals[80], it can be implied that each host has a unique biological relationship with its gut microbiota, thereby influencing the risk of disease [81,82]. The composition of the gut microbiota in physiological processes also changes with age, implicating long-term health outcomes.

Given the nascent nature of this field of research, there is currently a scarcity of studies examining the relationship between the ECS, the gut microbiota and endometriosis. However, studies have shown the involvement of the ECS in the gut microbiota in metabolic and inflammatory disorders.

The gut is part of the ECS. CB₁ receptors have been localised in the gut epithelium, smooth muscle, submucosal myenteric plexus and myenteric ganglia[83,84]. CB₂ receptors have been detected in the plasma cells and macrophages of the gastrointestinal (GI) mucosa and submucosa, as well as intestinal epithelial cells in the GI mucosa[83,85]. While CB₁ receptor activation is associated with appetite regulation and relief from nausea and vomiting, CB₂ receptor activation mediates inflammation[86]. Endocannabinoids are known to be synthesised in various parts of the gut and their levels fluctuate, based on metabolic and inflammatory status. Endocannabinoids and exogenous cannabinoids exert opposite effects on gut permeability. For instance, when examining decreased permeability as a result of inflammation, it was demonstrated that 2-AG and AEA increased permeability, while THC and CBD decreased permeability[87,88]. Recent literature on the link between the ECS and gut microbiota is summarised in Table 2.

1.2.1. Interactions between gut microbial communities and endocannabinoids

While the interactions between gut microbiota and endocannabinoids has mainly been investigated in preclinical models, studies have linked these interactions to beneficial effects in disease states such as inflammatory bowel disorder (IBD). The administration of *Lactobacillus acidophilus* in mice led to an increase of CB₂ expression in intestinal epithelial cells, resulting in analgesic effects, thus decreasing visceral pain[85]. Furthermore, the administration of a probiotic mixture containing *Bifidobacteria*, *Lactobacilli* spp. and *Streptococcus thermophilus* in zebrafish resulted in an upregulation of CB₁ and CB₂ expression, which then led to anti-inflammatory effects[89,90].

Akkermansia muciniphila is a Gram-negative anaerobic mucus-degrading bacterium, abundantly found in healthy intestinal mucosa. *A. muciniphila* has been found to modulate gut barrier integrity. Its protective effects have been cited in relation to the ECS. A study administering *Akkermansia muciniphila* in mice fed a high-fat diet led to an increase in 2-AG, 2-OG and 2-PG levels[91]. *A. muciniphila* has been shown to regulate CB₁ mRNA in Caco-2 cells through increased production in outer membrane vesicles, which prevent the development of metabolic disorders such as obesity[92]. Moreover, a decrease in CB₁ activity resulted in reduced circulating lipopolysaccharide (LPS) levels, thus improving the inflammatory cytokine profile and intestinal permeability. This occurred alongside enhanced *A. muciniphila* and *Lachnospiraceae* levels in the gut. The gut microbiota profile of obesity is characterised by a *Firmicutes*:*Bacteroidetes* ratio, whereby there is an increased abundance of *Firmicutes* and a reduced abundance of *Bacteroidetes*. When chronically treating obesogenic mice with THC, this ratio was shifted and an increase in the abundance of *A. muciniphila* was observed, demonstrating the protective effects of exogenous cannabinoids [93]. Further investigation is required to understand how *A. muciniphila* interacts with endocannabinoids and the outcomes of this on inflammation-associated disorders.

Studies have also examined the relationship between the ECS and gut microbial metabolites. For instance, endocannabinoids have been found to mediate the anti-inflammatory effects of SCFAs. This association was observed in an exercise intervention[94] where increase in SCFAs (including butyrate) and SCFA-producing bacteria (such as *Bifidobacterium*) were correlated with decrease in proinflammatory cytokines TNF- α and IL-6. Furthermore, an increased abundance of SCFA-producing bacteria, such as *Bifidobacterium*, was positively associated with increases in

endocannabinoids AEA, PEA and OEA, which were correlated with CB₁ receptor levels. These studies provided an indication of how the association between ECS and the gut microbiota can be potentially utilised to improve gut inflammation.

1.2.2. The ECS-gut-brain axis

As CB₁ receptors are found in the CNS, the ECS is involved in signalling, modulating various physiologic and homeostatic processes. While the ECS-gut-brain axis has not been studied extensively, recent studies have considered the involvement of the ECS-gut-brain axis in exercise interventions, processes of food intake and disease states such as Alzheimer's disease.

While investigating the microbiota-dependent gut-brain pathway on exercise, it was shown that the CB₁-expressing TRPV1 sensory neurons are triggered by fatty acid amide metabolites from gut bacteria such as *Lachnospiraceae* and *Eubacterium* [95]. The protective effects of *Lachnospiraceae* have been well-studied in various diseases, including ulcerative colitis[96,97]. This led to analgesic effects observed after exercise, demonstrating the interaction of intestinal microbial colonization and peripheral CB₁ signaling.

Endocannabinoids such as AEA and CB₂ receptors, and enzymes such as fatty acid amide hydrolase (FAAH) influence retrograde signalling (observed in Alzheimer's Disease) in the brain. Here, inhibitory feedback regulates neurotransmitter release[98]. Moreover, as AEA induces gut permeability, its overexpression is thought to cause a 'leaky' gut, resulting in metabolic endotoxemia[99]. As this condition leads to the release of toxins such as LPS which may cross the epithelial barrier, an inflammatory signalling pathway may develop, impacting the CNS and resulting in neuroinflammation[100].

CB₁ binding leads to increased food intake. CB₁ receptors are present in cells of the lining of the intestinal epithelium. When CB₁ receptor activity is heightened, in the small-intestinal epithelium, the release of cholecystokinin-8 (CCK-8) is inhibited, resulting in delayed satiation and overeating (in diet-induced obesity)[101], highlighting the indirect control mechanism of endocannabinoids in gut-brain neurotransmission.

1.2.3. The ECS in IBD

The use of cannabis for abdominal pain relief and other IBD symptoms is prevalent, with studies reporting use amongst 70-95% of people with IBD [86,102–104]. However, several studies have reported similar results when using endocannabinoids and/or exogenous cannabinoids in therapies for IBD. In analysis of biopsies of paediatric patients, AEA levels were found to be significantly decreased in inflamed IBD mucosa[105]. A significant increase in the expression of CB₂ receptors has also been observed in the biopsies of Crohn's disease ileum and in rectum biopsies of ulcerative colitis, colocalised with T-lymphocyte infiltration [106] (Figure 2). Similarly, a murine IBD studies demonstrated that increasing the availability of the endogenous CB₁ and CB₂ receptor agonists, diminishes visceral pain[107–109]. Nonetheless, the issue remains that such implications have not translated to clinical trials, as noted in one study where people with IBD using cannabis reported higher incidences of abdominal pain and arthralgias[110]. Further research is required to develop insights into what may be causing these negative side effects and utilisation of the ECS as a therapy for IBD.

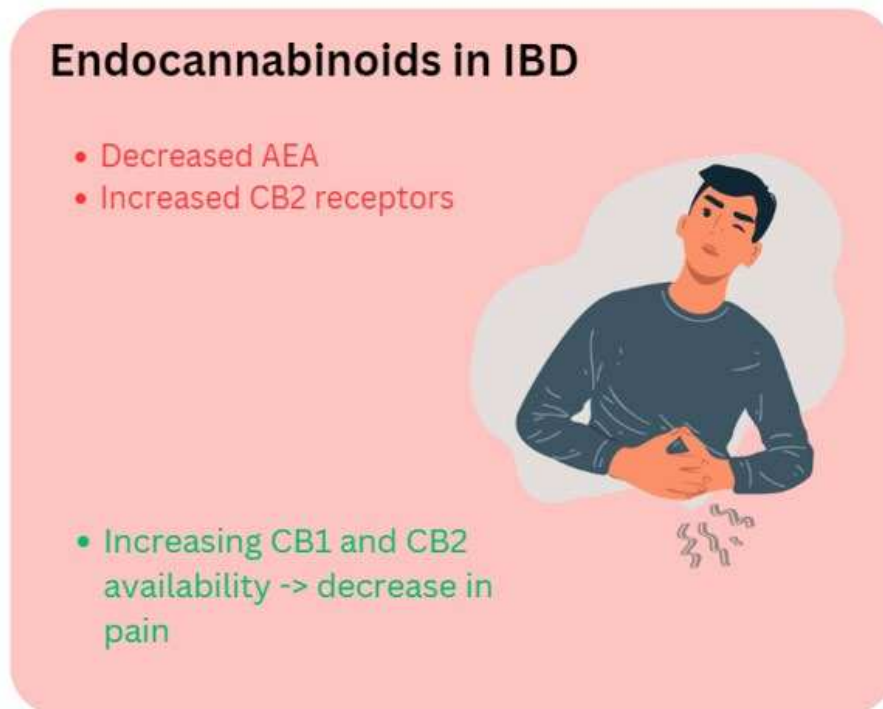


Figure 2. A decrease in AEA and increase in CB₂ receptors is reported in IBD. An increase in CB₁ and CB₂ availability leads to pain alleviation.

There is a high prevalence of inflammatory bowel syndrome (IBS) and its associated symptoms in people with endometriosis, in one study 52% (194/373) of those with endometriosis had diagnosed IBS[111], where those experiencing minimal-mild endometriosis often report more severe IBS symptoms, as compared to those with moderate-severe endometriosis. Similarly, of 160 people with diagnosed IBS, 59 had a history or recent diagnosis of endometriosis[112]. People with endometriosis are reported to have a threefold increase in likelihood of developing IBS[113]. Such studies demonstrate the importance of understanding the mechanisms through which inflammatory bowel diseases and endometriosis interact and thus the need for treatments targeting both diseases.

Table 3. A summary of recent literature on endocannabinoids and gut microbes.

Author	Year	Type of study	Endocannabinoid/cannabinoid + microbiota	Methodology	Findings
Strisciuglio, et al.	2023	Human- Crohn's disease	CB ₂ receptors	Western blot immunofluorescence	Increased expression of CB ₂ receptors in ileum of people with Crohn's disease.
Vijay, et al.	2021	People with knee osteoarthritis	2-AG, OEA, AEA, PEA <i>Bifidobacterium</i> , <i>Coprococcus</i> , <i>Faecalibacterium</i> , <i>Colinsella</i>	Metabolomic analysis Gut microbiome analysis Gene expression assay	An association between increased levels of SCFAs with circulating levels of endocannabinoids, higher microbiome diversity and low levels of proinflammatory <i>Colinsella</i> .
Pagano, et al.	2019	Pediatric patients with ulcerative colitis and male adult CD1 mice	Cannabidiol TRPA1	RT-PCR	In a TRPA1 antagonist manner, cannabidiol regulates systemic inflammation and intestinal permeability.
Di Sabatino, et al.	2011	Human- Crohn's disease Mucosal samples	AEA	HPLC-MS Wound healing scratch assay Immunohistochemistry	Significantly low levels of AEA in inflamed gut mucosa.
Grill, et al.	2019	C57BL/6 mice	CB ₁ receptors	<i>In situ</i> hybridisation Immunohistochemistry	Changes in gene expression of CB ₁ and CB ₂ receptors, GPR-55 and monoglycerol lipase in the gut after LPS treatment, demonstrating involvement in intestinal and systemic inflammation. These were observed in comparison to CB ₁ and MGL knockout mice. High expression of CB ₁ receptors in the submucosal and myenteric plexus. Reduced MGL expression in the ileum following LPS treatment. GPR-55 mRNA present alongside T-cell and macrophage markers in the ileum of healthy and treated mice.
Argueta & DiPatrizio	2017	Male C57BL/6 mice	CB ₁ receptor, 2-AG and AEA	LCMS Gene expression analysis	In diet-induced obesity, the increase of CB ₁ receptor activity inhibits CCK-8, resulting in delayed satiation and overeating. Blocking CB ₁ receptor activity resulted in decreased LPS activity, enhancing anti-inflammatory effects by increasing the abundance of <i>A. muciniphila</i> , <i>Lachnospiraceae</i> and <i>Erysipelotrichaceae</i> .
Mehrpouya-Bahrami, et al.	2017	Male C57BL/6 mice	<i>Akkermansia muciniphila</i> , <i>Lachnospiraceae</i> , <i>Erysipelotrichaceae</i> ,	16s RNA metagenomics	

Author(s)	Year	Sex	Strain	Intervention	Assays	Findings
Cluny, et al.	2015	Male mice	C57BL/6N	<i>Firmicutes</i> , <i>Bacteroidetes</i> <i>A. muciniphila</i> THC	qPCR	Chronic administration of THC in obesogenic mice increased the <i>Firmicutes</i> : <i>Bacteroidetes</i> ratio and the abundance of <i>A. muciniphila</i> .
Sakin, et al.	2015	Adult male Balb-C mice and Sprague-Dawley rats	CB ₁ and CB ₂ receptors		Colorectal distension test Nociceptive testing	Availability of CB ₁ and CB ₂ receptors diminishes visceral pain.
Kiran, Rakib, Moore & Singh	2022	Female mice	C57BL/6	CB ₂ inverse agonist SMM-189	Flow cytometry analysis Western blot analysis Histology Fibre photometry analysis DRG extraction, culture and calcium imaging	CB ₂ inverse agonist SMM-189 suppressed colitis, while ameliorating the loss of body weight, reducing the inflammatory disease score and disease severity.
Dohnalova, et al.	2022	C57BL/6J mice	CB ₁		Amplex flurometry analysis PCR, qPCR, RNA-seq Transcriptional profiling	Fatty acid amide metabolites trigger CB ₁ -expressing TRPV1 sensory neurons, thus elevating dopamine levels during exercise.
Jamontt, Molleman, Pertwee & Parsons	2010	Male Charles River Wister rats	Distal colon tissue	CBD + THC	<i>In vitro</i> evaluation MPO assay BCA protein assay	CBD and THC reduced inflammation and functional disturbances by reducing the release of TNF α , IFN γ and nitric oxide <i>in vitro</i> and <i>in vivo</i> .
Borelli, et al.	2009	Male ICR mice	CBD		Western blot ELISA LCMS	CBD reduced colon injury, decreased expression of inflammatory markers, including nitric oxide synthase and reactive oxygen species.
Alhamoruni, et al.	2012	Caco-2 cells	CB ₁ and CB ₂ receptors, TRPV1, PPAR γ and PPAR α THC and CBD		Measurements of transepithelial resistance electrical	THC and CBD accelerated recovery of cytokine-induced intestinal permeability.
Distrutti, et al.	2014	Zebrafish	<i>Bifidobacteria</i> , <i>Lactobacilli</i> spp. and <i>Streptococcus thermophilus</i>		TUNEL assay	Administration of a probiotic mixture containing <i>Bifidobacteria</i> , <i>Lactobacilli</i> spp. and <i>Streptococcus thermophilus</i> in zebrafish led to an increase in CB ₁ and CB ₂ expression.
Gioacchini, et al.	2017	Adult male zebrafish	<i>Bifidobacteria</i> , <i>Lactobacilli</i> spp. and <i>S. thermophilus</i> <i>Bacteroidetes</i> and <i>Actinobacteria</i>		RT-PCR Immunohistochemistry	Administration of a probiotic mixture containing <i>Bifidobacteria</i> , <i>Lactobacilli</i> spp. and <i>S. thermophilus</i> in aged zebrafish resulted in an increased abundance of <i>Bacteroidetes</i> and <i>Actinobacteria</i> .

alongside increases in CB₁, demonstrating anti-inflammatory effects.

Abbreviations: CB₂ = cannabinoid receptor 2; 2-AG = 2-arachidonoyl glycerol; OEA = N-oleoylethanolamine; AEA = anandamide; PEA = N-palmitoylethanolamine; TRPA-1 = transient receptor potential ankyrin type 1; CB₁ = cannabinoid receptor 1; THC = delta-9-tetrahydrocannabinol; PPAR γ = peroxisome proliferator-activated receptor gamma; PPAR α = peroxisome proliferator-activated receptor alpha; CBD = cannabidiol; RNA = ribonucleic acid; HPLC-MS = high protein liquid chromatography – mass spectrometry; RT-PCR = reverse transcriptase polymerase chain reaction; PCR = polymerase chain reaction; qPCR = quantitative polymerase chain reaction; LCMS = liquid chromatography mass spectrometry; TUNEL = terminal deoxynucleotidyl transferase dUTP nick end labelling; DRG = dorsal root ganglia; RNA-seq = RNA sequencing; SCFA = short-chain fatty acid; GPR55 = G-couple protein receptor 55; MGL = monoacyl glycerol lipase (MGL); CCK-8 = cholecystokinin-8; LPS = lipopolysaccharide; IFN γ = interferon gamma; TRPV-1 = transient receptor vanilloid 1; TNF- α = tumour necrosis factor alph.1.3. Endometriosis and the Gut Microbiota.

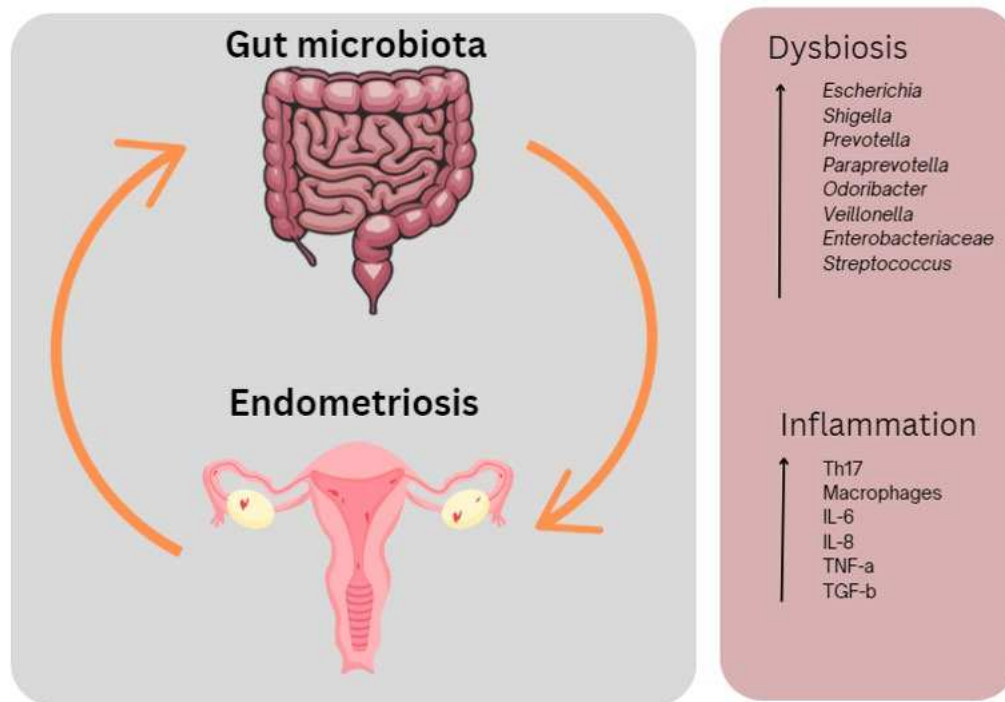


Figure 3. The interplay between the gut microbiota and endometriosis. Dysbiosis in the gut and inflammation occurring both in the gut and in the peritoneal cavity results in endometriosis-associated symptoms including pain and fatigue.

A bidirectional relationship between endometriosis and gut microbiota has been proposed (Figure 3). Current literature on the potential role of gut microbiota in endometriosis is summarised in Table 3. As gut microbes and their metabolites are involved in various immune, metabolic and epithelial functions, imbalances in the gut microbiota can trigger an inflammatory response through specific inflammatory immune cell recruitment, proinflammatory cytokine production and compromised immune surveillance. This has been observed in different disease states such as IBD [114], psoriasis[115] and cancer[116].

1.3.1. Microbial dysbiosis in endometriosis

In the endometriotic state, larger percentages of bacterial phyla *Proteobacteria*, *Verrucomicrobia*, *Streptococcus* or *Fusobacteria* have been reported. Increases in *Enterobacteriaceae*, *Streptococcus* and *E. coli* have been identified as dominating phyla in endometriosis cohorts[117]. Recently, *Fusobacterium* is suggested to contribute to the pathogenesis of endometriosis[118]. The dominance of *Shigella* has also been noted[119]. Such microbial communities are known to be involved in the degradation of estrogen by producing β -glucuronidase and β -glucosidase[120], resulting in the development of a high estrogen environment, promoting the progression of endometriosis[121,122].

An increase in *E. coli* is attributed as a biomarker of endometriosis and this has led to the “bacterial contamination” theory. High levels of *E. coli* have been found in faecal[123] and menstrual blood samples[124] of participants with endometriosis. *E. coli* is known to trigger TLR4 mediated growth and progression of endometriosis, resulting in pelvic inflammation[125]. This occurs through the production of LPS, triggering secretion of secondary inflammatory mediators such as NF- κ B, in the peritoneal cavity, therefore, resulting in the development and progression of endometriosis.

Further research is required to unravel the intricate dynamics between the microbiota and its influence on endometriosis progression. Such investigations would not only grow the understanding of the way in which the microbiota promote endometriosis pathogenesis, but also the development of targeted therapies to improve the quality of life of those with endometriosis.

1.3.2. Microbial dysbiosis and endometriosis symptoms

Research has demonstrated correlations with specific microbes and bacterial phyla impacting symptoms in those with endometriosis. Previous studies have shown the estrogen-gut-brain axis, which is thought to influence the development of chronic stress in people with endometriosis *via* the activation of β -adrenergic signalling[126]. This has been associated with dysbiosis on a genus level, particularly decrease in *Paraprevotella*, *Odoribacter*, *Veillonella* and *Ruminococcus* [126] which are often viewed as biomarkers of endometriosis[127]. A murine study showed that a decrease in *Ruminococcus* is negatively correlated with apoptosis of endometriotic epithelial cells and increased IL-6 levels, resulting in peritoneal inflammation[128]. *Prevotella* has been found in high abundance in people with endometriosis, especially in those experiencing gastrointestinal symptoms, and is associated with constipation, bloating, flatulence, vomiting and nausea[129]. As many studies report a correlation between gut microbiota imbalances and endometriosis, continued and extensive research is essential in understanding the full potential of harnessing the gut microbiota for the treatment of endometriosis and its associated symptoms.

The vaginal microbiota is dominated by healthy *Lactobacillus*, which maintains an acidic and protective environment, preventing the growth of pathogenic bacteria. However, in endometriosis, the protective environment maintained by vaginal microbiota is impacted by an abundance of *Gardnerella*, *Escherichia*, *Shigella*, *Ureoplasma*[119], *Streptococcus*, *Moraxellae*, *Staphylococcus* and *Enterobacteria*, coupled with a lowered abundance of *Lactobacillus*[130]. These imbalances have been associated with endometriosis-associated pelvic pain[131,132]. An understanding of the vaginal microbiota in endometriosis, in correlation with the ECS, is required for the characterisation of treatments that may be prescribed for people with endometriosis.

Table 4. A summary of recent literature on the gut microbiota and people with endometriosis.

Author	Year	Microbiota	Methodology	Findings
Svensson, et al.	2021	<i>Prevotella</i> , <i>Bacilli</i> , <i>Bacteriodia</i> , <i>Clostridia</i> , <i>Coriobacteria</i> and <i>Gammaproteobacter</i>	16s rRNA sequencing	<i>Prevotella</i> has been associated with common endometriosis gastrointestinal symptoms such as constipation, flatulence, bloating, vomiting and nausea. People with endometriosis were found to have a high abundance of <i>Lactococcus</i> (<i>Bacilli</i>), lower abundance of <i>Odoribacter</i> and higher abundance of <i>Prevotella</i> .
Sandstrom, et al.	2020	<i>Ruminococcus</i>	16s rRNA sequencing	Decrease in <i>Ruminococcus</i> correlated with an increase in IL-6 in a murine model of endometriosis, resulting in peritoneal inflammation. Absence of <i>Atopobium</i> in vaginal and cervical microbiota Increased <i>Gardnerella</i> in cervical microbiota
Ata, et al.	2019	Complete absence of <i>Atopobium</i> , <i>Gardnerella</i> , <i>Streptococcus</i> , <i>Escherichia</i> , <i>Shigella</i> , <i>Ureoplasma</i>	PCR amplification 16s rRNA sequencing	Dominant gut microbiota in endometriosis group – <i>Escherichia</i> and <i>Shigella</i> Predominant population of lower genital tract – <i>Lactobacillus</i> <i>Alloprevotella</i> significantly decreased in the cervix
Xu, et al.	2017	<i>Paraprevotella</i> , <i>Odoribacter</i> , <i>Veillonella</i> and <i>Ruminococcus</i>	16s rRNA sequencing Immunohistochemistry	The development of chronic stress in people with endometriosis occurs through the activation of β -adrenergic signalling, which occurs as a result of dysbiosis – decrease in specific unknown genus.
Khan, et al.	2010	<i>E. coli</i>	ELISA of macrophages from peritoneal fluid and epithelial/stromal cells from biopsy specimens of eutopic/ectopic endometria of women with and without endometriosis RT-PCR	Menstrual blood of people with endometriosis has a higher concentration of <i>E. coli</i> compared to healthy controls. An infiltration of macrophages in eutopic/ectopic endometria of people with endometriosis was noted.

Abbreviations: rRNA = ribosomal ribonucleic acid; ELISA = enzyme-linked immunosorbent assay; RT-PCR = reverse transcriptase polymerase chain reaction; IL-6 = interleukin-6.

6. Conclusions and Future Directions

This review summarises the relationship between endometriosis, the ECS and the gut microbiota. While studies have not completely deciphered the molecular basis of these relationships, current literature demonstrates the vital roles that the ECS and the gut microbiota play in potential implications in preventing the development of endometriosis as well as effective therapeutic strategies. The mechanisms of the ECS in both endometriosis and the gut microbiota were explored. Further research understanding the mechanisms, influence on inflammation and analysis of endocannabinoids and exogenous cannabinoids is required for the development of treatments of various diseases including endometriosis. While the impact of cannabinoids in endometriosis has been investigated, future studies are needed to comprehend how cannabinoids influence endometriosis-related pain and symptoms and how this can be implemented in the clinic, following efficacy and safety studies. This may be done through implementation of exogenous cannabinoids, which have been shown to have protective effects. An understanding of various inflammatory diseases was used to show the potential therapeutic role of the gut microbiota, while paving new avenues for further research. Finally, the relationship between the gut microbiota and endometriosis was explored, highlighting the need for further investigations of how cannabinoids may influence gut microbiota in endometriosis.

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