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The gastrointestinal tract – a central organ of cannabinoid signaling in health and disease

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Background and Purpose

In ancient medicine, extracts of the marijuana plant Cannabis sativa were used against diseases of the gastrointestinal (GI) tract. Today, our knowledge of the ingredients of the Cannabis plant has remarkably advanced enabling us to use a variety of herbal and synthetic cannabinoid compounds to study the endocannabinoid system (ECS), a physiologic entity that controls tissue homeostasis with the help of endogenously produced cannabinoids and their receptors. After many anecdotal reports suggested beneficial effects of Cannabis in GI disorders, it was not surprising to discover that the GI tract accommodates and expresses all the components of the ECS. Cannabinoid receptors and their endogenous ligands, the endocannabinoids, participate in the regulation of GI motility, secretion, and the maintenance of the epithelial barrier integrity. In addition, other receptors, such as the transient receptor potential cation channel subfamily V member 1 (TRPV1), the peroxisome proliferator-activated receptor alpha (PPARa) and the G-protein coupled receptor 55 (GPR55), are important participants in the actions of cannabinoids in the gut and critically determine the course of bowel inflammation and colon cancer. The following review summarizes important and recent findings on the role of cannabinoid receptors and their ligands in the GI tract with emphasis on GI disorders, such as irritable bowel syndrome, inflammatory bowel disease and colon cancer.

Keywords

Cannabis; cannabinoid receptors; colon cancer; GPR55; IBD; IBS

The endocannabinoid system in the GI tract

Cannabis has a long history as a traditional therapeutic agent for the treatment of abdominal pain and gut dysfunction. This beneficial effect is based on the fact that the gastrointestinal (GI) tract is endowed with cannabinoid (CB) receptors and their endogenous ligands.

Together they make up the endocannabinoid system (ECS), a physiologic entity that controls

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Disclosures

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homeostasis in the gut. There is also a wide range of cannabinoid compounds of exogenous origin. Next to herbal cannabinoids, such as ⁹-tetrahydrocannabinol (⁹-THC), cannabidiol, tetrahydrocannabivarin, cannabichromene, cannabigerol and others, there is a large array of synthetic cannabinoids. In general, cannabinoid compounds can be divided into five distinct classes, i.e. classical cannabinoids (e.g., ⁹-THC); non-classical cannabinoids (e.g., CP-55,940); indoles (e.g., WIN55,212), eicosanoids, and antagonist/inverse agonists (e.g., rimonabant) (1). For a detailed description of the ECS in the gut, the reader is referred to more comprehensive reviews (2,3).

In short, the ECS consists of the CB receptors 1 and 2 (CB₁, CB₂), their endogenous ligands ("endocannabinoids") as well as their degrading and synthesizing enzymes. CB₁ receptors can be found throughout the GI tract. There, they are predominantly located in the enteric nervous system (ENS) (4) and the epithelial lining (5). Additionally, CB₁ is found in extrinsic fibers of the ENS, plasma cells, and in smooth muscle cells of blood vessels within the colonic wall (6,7). Within the ENS, the CB₁ receptor is expressed prejunctionally in cholinergic, but not nitrergic neurons, explaining why CB₁ activation can depress excitatory transmitter release (8). CB₂ receptors are mainly present in immunocytes, myenteric plexus neurons, and in epithelial cells during ulcerative colitis (7,9). In addition to CB receptors, the orphan G-protein coupled receptor 55 (GPR55) and the transient receptor potential cation channel subfamily V member 1 (TRPV1) are endocannabinoid-responsive receptors and may be responsible for non-CB₁/CB₂ receptor effects of cannabinoids in the GI tract and are therefore regarded as part of an expanded ECS (10,11). PPAR receptors, in particular PPARα and PPARγ, are also responsive to herbal, synthetic and endogenous cannabinoids and may mediate many of the analgesic and anti-inflammatory effects observed in cannabinoid treatment [rev. in (12)]. The abovementioned receptors are present in the GI tract, e. g. on nerve terminals of extrinsic primary afferents (TRPV1) (2), and the ENS and enterocytes (PPARa, GPR55) (2,13).

Endocannabinoids are short-lived bioactive lipids and produced "on demand". Arachidonoyl ethanolamine (anandamide) and 2-arachidonoylglycerol (2-AG) are among the best characterized endocannabinoids and are synthesized by N-acyl phosphatidylethanolamine phospholipase D (NAPE-PLD) and diacylglycerol lipases (DAGL), respectively. They are degraded by specific enzymes: anandamide primarily by fatty acid amide hydrolase (FAAH) and 2-AG by monoglyceride lipase (MGL; or monoacylglycerol lipase, MAGL) (rev. in (3)). In the GI tract, FAAH and MGL were shown to be expressed in epithelial cells, the ENS, and in immune cells during ulcerative colitis (6,7,14). Endocannabinoids may be also degraded by cyclooxygenase-2 (COX-2) and lipoxygenase to give rise to prostaglandin ethanolamides, glyceryl prostaglandins, hydroxyeicosatetraenoic acid and hydroperoxyeicosatetraenoic acid derivatives (15,16). In contrast to the degrading enzymes, the synthesizing enzyme of anandamide, NAPE-PLD, and of 2-AG, DAGL α and β , have been observed in epithelial, myenteric plexus and lamina propria cells, and also in the smooth muscle layer (7).

Acylethanolamides other than anandamide, like palmitoylethanolamide (PEA) and oleoylethanolamide (OEA), can be classified as endocannabinoid-like compounds. They do not directly activate CB receptors but they can activate GPR55 (predominantly PEA) and

GPR119 (only OEA) and are able to influence the signaling of anandamide via an entourage effect (17). PEA and OEA also activate PPARα and are present in high levels within the gut. Both of them are degraded by FAAH, however, PEA is preferentially degraded by another amidase, N-acylethanolamine-hydrolyzing acid amidase (NAAA), which is strongly expressed in immune cells and active particularly in the intestine, suggesting a potentially pathophysiological role in the GI tract (rev. in (17)). In summary, the GI tract is able to locally produce its own endocannabinoid ligands according to its physiological needs and may rapidly react to disturbances in the gut to maintain homeostasis.

Cannabinoids in GI motility and secretion

Cannabinoids affect gut motility mainly by activating CB₁ receptors present on enteric neurons (18). Activation of CB₁ receptors results in the inhibition of acetylcholine release which consequently causes a decrease of intestinal smooth muscle contractility and peristalsis (19). Early studies demonstrated that the plant-derived CB receptor agonist THC, the main component of *Cannabis*, decreases intestinal transit and inhibits electrically evoked contractions in guinea pig explants (20,21). Synthetic CB receptor agonists likewise reduce gastric emptying, upper GI transit, and colonic propulsion (reviewed in (2)). In contrast, rimonabant (SR141716), an inverse agonist of CB₁ receptors, increased electrically-evoked contractions and peristalsis in isolated intestinal segments (22,23), as well as intestinal motility in vivo (24). Although CB2 receptors are expressed in the ENS, they are suggested to play a minor role in the regulation of gut motility under basal conditions but might become important under pathophysiological settings (9). Indeed, JWH-133, a CB₂ receptor agonist, but not arachidonyl-2'-chloroethylamide (ACEA), a CB₁ receptor agonist, attenuated gut transit dose-dependently in the inflamed gut of rats, an effect that was prevented by a CB₂ receptor antagonist (25). There is also increasing evidence that GPR55 is involved in the regulation of gut motility since its agonist O-1602 was able to slow down whole gut transit in mice (13). Both PEA and OEA inhibit intestinal transit in mice but the mode of action is unclear because neither CB receptors nor PPARa seem to be involved in that process (26,27); however, in a mouse model of postinflammatory IBS (mustard oilinduced), inhibition of transit by PEA could be blocked with a CB₁ receptor antagonist, but was not significantly modified with a PPARa antagonist (28).

Acute inhibition of endocannabinoid-synthesizing or - degrading enzymes also modulates intestinal motility. Thus, inhibition of DAGL α was able to normalize gut motility in a mouse model of genetically-induced constipation (29). Pharmacological inhibition of FAAH or MGL led to a decrease in gut motility through mechanisms that involved a rise in anandamide or 2-AG levels, respectively, and the activation of CB₁ receptors (14,30,31). Interestingly, FAAH-deficient mice did not show alterations in basal gut motility; however, pharmacological inhibition or genetic deletion of FAAH normalized endotoxin-induced hypermotility (31). Taschler et al. demonstrated that MGL-deficient mice did not show alterations in basal gut motility but that they were insensitive to CB receptor agonist treatment due to desensitization of intestinal CB₁ receptors (30).

It has to be noted that also the gut brain-axis may account for the regulation of gut motility by cannabinoids. For instance, intracerebroventricular injection of the CB receptor agonist

WIN55,212-2 attenuated whole gut transit in mice (32). Additional evidence that gut motility might be regulated by central CB receptors was provided by Vianna *et al.* who showed that deletion of CB₁ receptors specifically in the vagal nerves of mice caused an increase in GI motility (33). Similar to rodents, CB₁ receptors are functionally present in the human small and large intestines (34–36). Thus, WIN55,212-2 and ACEA inhibited electrically-evoked contractions in a healthy human colon and this effect was completely blocked by rimonabant (37). Also 2-AG and anandamide were shown to inhibit acetylcholine-induced contractions in explants of human colonic longitudinal and circular muscle, however, this effect could not be blocked with CB₁ or CB₂ antagonists (38). The authors suggested a non-cannabinoid or alternative cannabinoid pathway mediating this effect (38). It is possible that the non-CB effects by anandamide may have been brought about by GPR55 which causes relaxation in the murine colon (13).

There is evidence that cannabinoids play an important role in the regulation of gastric and intestinal secretion in rodents and humans. Studies revealed that cannabinoids reduce the production of gastric acid secretion by activating CB₁ receptors (19). In mice, intestinal hypersecretion induced by cholera toxin was reduced by CB₁ receptor activation (39). In another study, pharmacological inhibition or genetic deletion of FAAH provided beneficial effects against diclofenac-induced gastric irritation (40). In contrast, enhanced secretion was observed in humans treated with the CB₁ antagonist rimonabant (41). In summary, a large body of evidence demonstrates that (endo-) cannabinoids affect physiologic functions of the gut, a property that could be therapeutically exploited. Activation of CB₁ receptors by increased levels of endocannabinoids and, as a consequence, a slowed gut motility might have beneficial effects for patients with symptoms of hypermotility. On the other hand, inhibition of endocannabinoid synthesis or blockade of CB₁ receptors might enhance gut motility in GI disorders associated with constipation. If central side effects of cannabinoids could be overcome, modulation of cannabinoid levels would certainly represent a valuable pharmacological approach for the treatment of GI disorders. Another possibility could be the use of non-psychotropic cannabinoids like cannabidiol, which has been described as a ligand of many receptors including GPR55, TRPV2, PPARy and 5-HT1A but not of CB receptors (but might modulate their actions) (42). Cannabidiol has shown relaxant effects on croton oil- and sepsis-induced hypermotility in mice (43,44).

Cannabinoids in emesis and nausea

The dorsal vagal complex (DVC) in the brainstem is the site responsible for the vomiting reflex while the neural circuitry responsible for nausea is less known. CB receptors and particularly FAAH and MGL are present in the DVC and area postrema suggesting an important role of endocannabinoids in the control of emesis (45–47). Cannabis has been traditionally used as an antiemetic agent, and exogenous cannabinoids are presently prescribed for people with chemotherapy-induced nausea and vomiting (48). However, due to central side effects, cannabinoids are not used as first line drugs.

The endocannabinoids anandamide and 2-AG have been shown to reduce emesis in experimental models (46). Drugs that can raise endocannabinoid levels without causing the typical cannabinoid agonist-induced central side effects are therefore potential options to

treat emesis. The FAAH inhibitor URB597 reduced LiCl-induced emesis via ${\rm CB_1}$ and ${\rm CB_2}$ receptors (46). Reduction of emesis by the MGL inhibitor JZL184 was shown to be sensitive to ${\rm CB_1}$ antagonism (49). Also cannabidiol showed anti-emetic and anti-nausea effects in animal models, the effects were brought about by indirect agonism of 5-HT $_{\rm 1A}$ somatodendritic autoreceptors in the dorsal raphe nucleus (50).

The role of endocannabinoids has been investigated more recently in detail in the conditioned gaping model in rats and results indicate that 2-AG and the visceral insular cortex (VIC) could play an important role in nausea (51). Exogenous 2-AG, but not exogenous anandamide, applied by bilateral intra VIC infusion, dose-dependently suppressed conditioned gaping (51). The effect could not be blocked with the CB₁ antagonist AM251, but instead with the COX inhibitor indomethacin (51). Interestingly, bilateral VIC infusion with the MGL inhibitor MJN110 also suppressed conditioned gaping but here, the effect could be blocked with AM251 (52).

Endocannabinoids have been clearly established as important messengers in the neuronal network that controls vomiting and nausea. Interference with endocannabinoid degradation may represent a valuable therapeutic approach not only against emesis but also against anticipatory nausea in chemotherapy patients.

Cannabinoids and functional bowel disorders

Irritable bowel syndrome (IBS) and functional dyspepsia are the most frequent functional bowel disorders encountered globally. The previous view that functional GI disorders lack histopathological and biochemical alterations has been challenged by studies demonstrating low grade inflammation, increased presence of immune and mast cell, changes in the epithelial barrier, and bacterial overgrowth in IBS patients. These alterations together with a derangement of the gut-brain axis may be involved in the development of visceral hyperalgesia and motility disturbances. The predominant presence of CB₁ receptors along the gut-brain axis may allow cannabinoids to positively influence derangements along this axis (3,53). The role of the ECS in IBS has been already described in a previous review by Storr&Sharkey (53). Here, more recent results will be summarized and discussed.

IBS: visceral hypersensitivity and the ECS

Symptoms of IBS, such as abdominal pain, discomfort, and altered bowel habits, have been previously linked with visceral hypersensitivity and aberrant 5-hydroxytryptamine (5-HT) signaling (53). Feng et al. explored the link between 5-HT and the ECS and observed increased levels of 5-HT, but a decrease in anandamide, in the duodenal mucosa of patients with postinfectious IBS (PI-IBS) (54). Using a rat model, they showed that acute luminal administration of 5-HT into the duodenum induced anandamide release via vagal 5-HT₃ receptors, whereas chronic 5-HT treatment decreased anandamide levels via 5-HT₃, indicating that 5-HT may be involved in the regulation of intestinal anandamide content. In addition, luminally-applied CB₁ receptor agonists attenuated 5-HT-induced hyperalgesia (54). In IBS-D (diarrhea-predominant) patients, no changes in anandamide levels but a decrease in PEA was observed in comparison to healthy subjects. The decrease was associated with abdominal pain (55). The IBS-D patients also had an increase in 2-AG while

IBS-C patients had higher levels of OEA (55). It is interesting that levels of PEA were also found decreased in a mouse model of inflammation-induced hypermotility (croton oilinduced) (56). The decrease was reduced by a non-psychoactive Cannabis extract, cannabichromene, in a CB receptor-independent manner (56). In contrast, in a mouse model of postinflammatory IBS (mustard oil-induced), PEA slowed gut transit, an effect that was dependent on CB₁ receptors (28). By use of a trinitrobenzenesulfonic acid (TNBS)-induced model of visceral hypersensitivity, Iwata et al. showed that a CB₂ receptor agonist was effective in improving pain thresholds in a dose-dependent manner without signs of central CB₁ receptor activation (40,57). Considering these data it is possible that low levels of endocannabinoids in IBS patients may contribute to hyperalgesia and abdominal pain and cause perturbations in the bowel motility which could be improved by endo- or exocannabinoids via CB- and possibly non-CB receptor pathways. This leads to the idea that FAAH inhibitors could be valuable therapeutics against PI-IBS and possibly other forms of IBS. In accordance with this concept, several studies reported that pharmacological inhibition of FAAH and also MGL significantly reduced visceral nociception in rodent models of colorectal distension and acetic acid-induced abdominal stretching (40,58,59). In this context it is worth to mention the role of mast cells in IBS. Activated mast cells have been shown to correlate with abdominal pain in IBS (60). Since mast cells express CB receptors and are also targets of PEA (61), which is thought to modulate mast cells activation, endocannabinoids may regulate activity of mast cells and hence interfere with IBS symptoms like abdominal pain; however, this remains to be shown.

IBS: stress, pain and the ECS

Chronic stress can induce visceral hyperalgesia via the hypothalamic–pituitary–adrenal axis and probably adds to the pain that IBS patients perceive. Recent work in rat models has shown that chronic stress causes reciprocal changes in 2-AG and COX-2/FAAH levels in L6–S2, but not L4–L5 dorsal root ganglia (DRGs) (62). Moreover, CB₁ receptors were downregulated while TRPV1 receptors were upregulated in L6–S2 but not in L4–L5 DRGs, indicating region-specific changes in primary sensory fibers innervating the distal colon (62). A report suggests that epigenetic regulation in the DRG neurons could be responsible for these changes: while chronic stress was associated with methylation in the promoter region of the Cnr1 gene (encodes the CB₁ receptor), histone acetylation at the Trpv1 promoter and expression of the TRPV1 receptor were increased (63). These findings point out that reciprocal changes in the endovanilloid and endocannabinoid system occur in visceral sensory fibers and that these changes could contribute to hyperalgesia and abdominal pain.

Stress and visceral pain may be also regulated by the ECS within the CNS. It is known that chronic stress reduces levels of anandamide (but increases 2-AG) in the brain and downregulates CB₁ receptors, and that these changes may contribute to the stress response (64). In line with this, both the FAAH inhibitor PF 3845 and the dual FAAH/MAGL inhibitor JZL 195 were effective in inflammatory and mechanically evoked visceral pain models suggesting that an increase in endocannabinoid levels alleviates visceral pain (59). A more thorough description of this topic is given in (65).

IBS: genetic variations and the ECS

Genetic variations of ECS components (CB receptors, synthesizing/degrading enzymes) may be associated with the pathogenesis of functional bowel disorders. Polymorphism in the FAAH gene (C385A) leads to a mutant FAAH enzyme and reduces breakdown of anandamide (66). A study in patients with constipation predominant (C-) IBS, D- and M-(mixed) IBS, with chronic abdominal pain and functional dyspepsia, showed a clear association of the non-wild type FAAH genotype with functional bowel disease phenotypes and with accelerated colonic transit in IBS-D patients (67). However, no statistically significant association between the FAAH genotype and sensation measurements was observed (67). A polymorphism in the CNR1 gene, rs806378, was found to be significantly associated with IBS symptom phenotype, colonic transit in IBS-D, and sensation rating of gas, but not with pain (68). In line with a possible role of CNR1 variants in the development of IBS symptoms, allele frequencies of AAT triplet repeats in CNR1 were observed to be associated with IBS in a study of a Korean population (69). Similar results, namely the detection of eight CNR1 alleles with AAT triplet repeats, were reported in a Chinese IBS cohort, whereas no association could be detected between C385A FAAH polymorphism and IBS pathogenesis (70). Interestingly, FAAH activity was recently determined in whole colon samples from patients who underwent colectomy for slow transit constipation (71). The results revealed a strong decrease in activity in these patients as compared to individuals free of transit dysfunction (71). The FAAH enzyme, therefore, seems to be a key molecule for the regulation of endocannabinoid levels and colon motility, but not for GI pain sensation.

Effect of CB receptor agonists in IBS patients

Thus, it seems that CB receptor activation in IBS has potential therapeutic value, but probably only in IBS-D patients with genetic variations of ECS components.

Functional dyspepsia

There is good indication that the ECS may be involved in functional dyspepsia. Tack et al. have previously shown that early satiety and symptoms of functional dyspepsia are caused by a disturbed gastric accommodation (76). In addition, hypersensitivity to gastric balloon

distension was observed to be present in a subset of patients with functional dyspepsia (77). A cross-over, randomized, controlled clinical trial in healthy individuals now demonstrated that CB₁ receptor antagonist rimonabant was able to inhibit meal-induced gastric accommodation, but did not affect fasting gastric compliance or sensitivity to gastric balloon distension, indicating that gastric accommodation is controlled by endocannabinoids (78). However, it was not clear from the study whether the ECS controls accommodation via centrally-mediated pathways or via the ENS. A new study has recently addressed the question as to whether CB₁ receptors in the brain are involved in functional dyspepsia and could demonstrate that increased availability to a CB₁ receptor radioligand was predominantly found in brain regions involved in the regulation of visceral pain and satiety (79). These findings would argue for a role of central CB receptors in the regulation of gastric accommodation in humans. It is, therefore, possible that both, central and peripheral CB receptors are involved in the development of functional GI disorders, and that pharmacological manipulation of exclusively peripheral CB receptors may not provide full benefit for patients with these disorders.

Microbiota and the ECS

A change in the microbiotic population of the gut may alter the permeability and lead to metabolic endotoxemia and hence to metabolic disorders associated with obesity. Endocannabinoids are involved in the regulation of energy metabolism and food intake and communicate in this respect with the microorganisms of the gut (80). The epithelial lining expresses CB receptors and they are most likely involved in these mechanisms. 2-AG and PEA cause an increase in epithelial barrier function ("gate keeper") while anandamide is thought to be a "gate opener" (81). Thus, the intestinal ECS may have an important role in the control of microbial products entering the bloodstream and in the development of metabolic diseases. A detailed review on this topic is given in (81).

Dysbiosis (alteration in the composition of gut microbiota) has been also suggested as one of the potential causes of IBS, especially in the case of PI-IBS (82). It is known that antibiotic therapy provides certain benefits for IBS patients (83), however, it is not quite clear how eradication of bacteria could contribute to symptom relief. In this context it is interesting that Lactobacillus acidophilus NCFM could induce CB2 receptor expression in the rodent gut mucosa (84). When applying NCFM in a model of chronic colonic hypersensitivity, it caused analgesia which was abrogated by i.p. blockade with AM630, suggesting that CB₂ receptors may provide a link between gut microbiota and visceral hypersensitivity (84). However, in a human trial, CB₂ receptors were not found to be upregulated in colonic mucosal biopsies from persons that were given Lactobacillus acidophilus NCFM over a period of 21 days (85). On the other hand, treatment of mice with antibiotics reduced painrelated responses to i.p. application of acetic acid or intracolonic capsaicin (86). The effect was accompanied by a small rise in CB₂ receptor transcripts in colon tissue, as well as a decrease of CB₁ and mu-opioid receptors. Additionally, total luminal bacterial counts correlated with CB receptor expression (86) suggesting a possible interaction between microbial products and CB receptors.

Cannabinoids and intestinal inflammation

Chronic inflammatory conditions of the GI tract are known as inflammatory bowel disease (IBD) and occur in two major forms, ulcerative colitis (UC) and Crohn's disease (CD). IBD is thought to originate from a complex interaction of the gut microbiota (or their products) with the epithelial barrier, based on the genetic background and the immune system of the host (87). To investigate the role of cannabinoids in IBD, mostly animal models that rely on chemically-induced mucosal inflammation have been used.

The endocannabinoid system as a therapeutic target in IBD

Evidence gathered from several studies in rodents points to a therapeutic relevance of the ECS in IBD. As reviewed by Izzo & Sharkey (2) and Alhouayek & Muccioli (88), endocannabinoid signaling is largely enhanced in the inflamed intestine. Expression of CB₁ (89) and CB₂ receptors (90), and of anandamide (91) were increased, whereas FAAH levels were reduced in the initial phase of colitis (92). Pharmacological strategies to enhance endocannabinoid levels, either by inhibition of endocannabinoid degradation (92-94) or of the transport across the plasma membrane (91,92) ameliorated inflammation. In particular, inhibition of FAAH by PF-3845 (94) and FAAH/COX blockade by ARN2508 (95) dramatically reduced damage in experimental colitis models. In the latter study, raised levels of anandamide, PEA and OEA were measured that most likely contributed to the beneficial effect (95). A recent work by Alhouayek et al showed that inhibition of NAAA, which preferentially degrades PEA, caused significant improvement of experimental colitis suggesting that PEA is an important acylethanolamide in the regulation of intestinal inflammation (96). In accordance, oral administration of PEA (which is interestingly sold as an over-the-counter drug and advertised to mitigate symptoms of GI disorders) exerted antiinflammatory effects in the gut (97). Experiments on cultured human colonic biopsies derived from UC patients showed that PEA caused a decrease in expression and release of inflammatory mediators which was dependent on PPARa (98).

Activation of the CB₁ (89) or CB₂ receptor (90,99) with specific agonists also protected from colitis. Accordingly, genetic ablation or pharmacological antagonism of CB₁ (89,100) or CB₂ receptors (90,100) left mice more susceptible to intestinal inflammation. Moreover, treatment with ⁹-THC was reported to reduce colitis in rats (101). The limitations of using Cannabis for treatment of gut inflammation, however, are the psychoactive effects that arise from activation of CB₁ receptors in the brain. Investigation of pharmacologically active cannabinoids with low or no affinity for CB₁ receptors and of atypical cannabinoids would be therefore of high interest. Indeed, it has been shown that cannabidiol and cannabigerol, two non-psychotropic ingredients of Cannabis, have proven beneficial in various models of intestinal inflammation (101-105). Also, the atypical cannabinoid O-1602 was reported to reduce disease severity in a CB₁-/ CB₂ receptor-independent way by inhibiting neutrophil recruitment (106). Recently, GPR55, which is part of the "expanded" ECS, has been investigated in experimental colitis. A pro-inflammatory role of GPR55 could be established because genetic deletion of GPR55 and treatment with the GPR55 antagonists CID16020046 or ML-191 alleviated intestinal inflammation (97,106,107). In this context, cannabidiol, which is known to act as a GPR55 antagonist (108), showed inhibition of GI inflammation

in an LPS-induced model by targeting enteric reactive gliosis (103). Interestingly, only parts of the beneficial effects of cannabidiol in this model were mediated by PPAR γ (103) raising the possibility that GPR55 could have been involved in this effect. Cannabidiol may also exert a protective effect on the intestinal barrier. In a Caco-2 cell monolayer stimulated by EDTA, cannabidiol concentration-dependently caused rapid recovery of the barrier and this effect was inhibited by a CB $_1$ antagonist (109). Since cannabidiol has no affinity to CB $_1$ receptors, the authors argued that cannabidiol could have antagonized CB $_1$ -mediated increases in permeability mediated by locally produced endocannabinoids (109). Activation of CB $_2$ receptors also attenuated cytokine-evoked mucosal damage in human colonic explants (110).

Cannabis for the treatment of IBD?

Questionnaires among IBD patients revealed that *Cannabis* is commonly used as a self-medication to relieve IBD-related symptoms like abdominal pain, diarrhea, and loss of appetite (111,112). A retrospective study reported significant improvements in 21 out of 30 CD patients after *Cannabis* use (113). In a small prospective placebo-controlled study of CD patients, a beneficial clinical response was achieved in 10 out of 11 subjects in the treatment group (114). A more recent questionnaire confirmed that the use of *Cannabis* subjectively improved pain and other symptoms in IBD patients, but also pointed out that *Cannabis* use for more than six months was a strong predictor in CD patients for requiring surgery (115).

Despite these interesting findings, the exact mechanisms how the ECS operates in IBD have not yet been unraveled but evidence gathered so far points to an overall protective role (Fig. 1). The up-regulation of ECS components possibly constitutes an attempt to restore homeostatic balance (3). Cannabinoids have been shown to influence the recruitment of immune cells to the site of intestinal inflammation (93,106,107) and to reduce the release of pro-inflammatory cytokines, i.e. TNF- α , IFN- γ , IL-1 β and IL-6 (93,102,103,105). Activation of the CB₁ receptor might also lead to enhanced wound closure during colitis (5). Of particular interest are recent findings that gut microorganisms may influence the expression of intestinal ECS components (81). 2-AG and PEA were mostly associated with beneficial effects on the gut-barrier function (81). The crosstalk between gut microbiota and the ECS is therefore worthy to be further examined in future studies.

Collectively, cannabinoids show great potential in the treatment of IBD and further research is warranted to gain a better insight into the mechanistic actions of (endo-) cannabinoids.

Cannabinoids and colon cancer

Differential expression of components of the ECS in colorectal cancer (CRC) was first reported by Ligresti et al. (116). In this study, anandamide and 2-AG contents were found to be higher (3-fold and 2-fold, respectively) in CRC lesions as compared to normal colonic mucosa and, interestingly, their levels were higher in adenomatous polyps than in carcinomas (116). Increased endocannabinoid synthesis in CRC was also reported in a more recent study (117). Here, anandamide, as well as its synthesizing enzyme NAPE-PLD, were up-regulated approximately 2-fold in cancer tissues. Intriguingly, mRNA expression and activity levels of FAAH were also increased. Most likely, as a consequence of increased

FAAH activity, elevated levels of arachidonic acid, the main product of anandamide and 2-AG degradation, were also detected (117). In another study, the main degrading enzyme of 2-AG, MGL, was also found increased in CRC specimens (118).

Examination of CB₁ expression revealed a down-regulation of mRNA levels in 18 out of 19 colon cancer samples as compared to adjacent non-neoplastic colon mucosa (119). The reason for this silencing was found to be DNA hypermethylation at CpG islands around the transcription start site of CNR1. In parallel to the epigenetic regulation, also protein levels of CB₁ receptors were reduced in colon cancer specimens as shown by Western blotting (119). These findings were corroborated by Cianchi et al. who reported CB₁ receptor expression to be higher in normal colonic epithelium than in colonic tumor tissue (120). However, a comprehensive study describing the correlation between CB₁ receptor immunoreactivity and patient outcome conducted in 534 Korean patients found no differences in overall survival between patients with carcinomas of either high or low CB₁ receptor immunoreactivity (121). Distant metastasis was found to be lower in patients with high CB₁ receptor expression, but there were no differences in lymph node metastasis, tumor invasion, or tumor size. Surprisingly, in stage IV patients, high CB₁ immunoreactivity even correlated with a poorer survival rate (121). Similar observations were made in a cohort of 487 Swedish patients (122). There, high CB₁ expression was reported to correlate with poorer disease-specific survival in stage II microsatellite stable CRC patients (122). Reduced overall survival has also been reported for patients who were either homo- or heterozygous for the 1359 G/A single nucleotide exchange in the CNR1 gene although it is not yet known how this polymorphism affects cannabinoid signaling (123). CB₂ receptor mRNA expression was found in 28.6% of CRC samples and significantly correlated with lymph node involvement (124), however, no consistent data on protein expression were available. So far, the human studies indicate increased endocannabinoid activity in colon cancer while the role of CB receptors remains less clear.

Cannabinoids reduce carcinogenesis in animal models of colon cancer

In mice, colon cancer can be induced either chemically or, for instance, by germline mutation of the adenomatous polyposis coli (Apc) gene. $Apc^{Min/+}$ mice spontaneously develop multiple polyps in the intestine. Additional knock out of Cnr1 or inhibition of the CB_1 receptor with AM251 in these mice caused a strong increase in intestinal polyp burden, whereas activation of CB_1 receptors with methanandamide significantly reduced the number of polyps (119). Genetic deletion of Cnr2 (the gene encoding CB_2 receptor), had no effect on polyp growth in this model (119). Chemically, colon cancer develops after multiple intraperitoneal injections of the carcinogen azoxymethane (AOM). In this model, anandamide and 2-AG were found increased in the colon of AOM-treated mice (125). In addition, inhibition of FAAH with N-arachidonoyl-serotonin (AA-5-HT) reduced the development of precancerous lesions, and furthermore, the non-selective, synthetic CB_1/CB_2 receptor agonist, HU210, was able to mimic this effect (125).

Antitumorigenic effects in the AOM model were also observed with non-psychotropic cannabinoids. For instance, cannabidiol was shown to reduce the formation of aberrant crypt foci (ACF), polyps, and tumors in the colon and the AOM-induced up-regulation of p-Akt

(126). It also counteracted caspase-3 inactivation. In colorectal carcinoma cell lines, it protected DNA from oxidative damage and it reduced cell proliferation in a CB₁-, TRPV1- and PPARγ-antagonists sensitive manner (126). A "cannabidiol botanical drug substance" (a *Cannabis sativa* extract with high content of cannabidiol) had similar effects in the same model, reducing ACF, polyp and tumor formation via CB₁ and CB₂ receptor activation (127), whereas treatment with cannabigerol reduced the number of ACFs only (128). In yet another murine model, in which colitis-associated colon cancer was induced through the application of AOM and dextran sulfate sodium (DSS), the atypical cannabinoid O-1602 showed antitumorigenic properties (129). The drug reduced the number and area of tumors by 30% and 50%, respectively. In addition, activation of the oncogenic transcription factor STAT3 was decreased while pro-apoptotic factors p53 and Bax were increased in O-1602 treated mice (129). Perhaps surprisingly, one study showed that antagonism of CB₁ receptors with rimonabant reduced the formation of ACFs with 4 or more crypts in mice with AOM-induced colon cancer (130).

Potential applications of cannabinoids and related substances have also been studied in xenograft models. The semi-synthetic cannabinoid quinone HU-331 (131) and the hexahydrocannabinol analogue LYR-8 (132) reduced tumor growth of xenografts derived from HT-29 cells. Likewise, the CB₂ receptor agonist CB13 inhibited the growth of DLD-1 derived tumors (120). A "cannabidiol botanical drug substance" (127) and cannabigerol (128) decelerated or even halted the growth of HCT116 xenografts, respectively.

Anticarcinogenic mechanisms of cannabinoids: reduction of cancer cell proliferation and inhibition of angiogenesis and metastasis

Cannabinoids have been shown to exert anti-proliferative effects on colon cancer cells through apoptosis via activation of CB₁/CB₂ receptors, or through receptor-independent mechanisms (rev. in (133)). The molecular mechanisms underlying the induction of apoptosis upon CB₁/CB₂ receptor activation have been discussed in detail by Velasco et al. (134). Briefly, de novo synthesis of the pro-apoptotic sphingolipid ceramide (120), downregulation of the protein survivin (inhibitor of apoptosis) (119), inhibition of PI3K/Akt signaling (135,136), and induction of endoplasmic reticulum stress that leads to autophagymediated cell death (136), have all been reported. Notably, cannabinoids with low or no affinity for CB receptors (like cannabidiol and O-1602) are also known to exert antiproliferative effects, although the underlying mechanisms have not yet been fully clarified (126,127,129). A cannabinoid-like compound LYR-8, for instance, was demonstrated to decrease angiogenesis in a xenograft model using chick chorioallantoic membranes (132). Concomitantly, the expression of factors that modulate the tumor microenvironment, like vascular endothelial growth factor, COX-2, and hypoxia-inducible factor 1a was reduced in this model (132). Inhibition of MGL, either pharmacologically or through silencing with siRNA, attenuated the invasion of colon cancer cells (118), suggesting a role of endocannabinoid degrading enzymes in CRC progression. Importantly, adhesion and migration of highly metastatic colon cancer cells was shown to be diminished after treatment with cannabidiol or a GPR55 inhibitor (108).

In conclusion, data obtained so far point to a deregulation of the ECS in colon cancer that could be interpreted as an attempt to restore the original healthy state. Despite controversial data on the role of the ECS in human colon cancer, promising preclinical data on the reduction in tumor growth by typical and atypical cannabinoid compounds warrant further exploration on the cause of ECS deregulation in colon carcinogenesis. It should be of prime interest to investigate known and hitherto unknown components of the ECS to better understand the complexity of CB receptor signaling by endocannabinoids and the regulation of their synthesizing and degrading enzymes.

Concluding remarks

Cannabinoids have a long history of being used to treat diseases or to alleviate symptoms. In modern medicine, this is not fully translated, and cannabinoids or cannabinoid-derived drugs are rarely used mainly due to the lack of clinical trials supporting such use. Over the last decades, cannabinoid research was driven by basic scientists who characterized pharmacological actions of cannabinoids, who discovered the ECS with all its constituents, and who taught us how activation or blockade at different sites may be helpful for the treatment of GI diseases. The GI tract is one of the regions where cannabinoid signaling is involved in many physiological and pathophysiological regulatory mechanisms, this is now clearly understood. The last decade has added more translational studies, and we have learned where cannabinoids are involved in pathophysiological states and human disease and where and how cannabinoids alter physiological or pathophysiological conditions. Through a recent meta-analysis we are also better informed on side effects associated with cannabinoid treatment. The analysis revealed that there was an increased risk of short-term adverse events with cannabinoids, mostly dizziness, dry mouth, nausea, fatigue, somnolence, euphoria, drowsiness, but also cardiac (1.42; 0.58-3.48; odds ratio; 95% CI) and hepatobiliary (3.07; 0.12-76.29; odds ratio; 95% CI) disorders were among them (137). Nevertheless, the opportunities are multifold with targeting the numerous involved receptors with agonists and antagonists, and with targeting synthesizing and degrading mechanisms. To harvest the potential therapeutic effects is now challenging, but based on the broad cannabinoid platform built by basic researchers, clinical trials are urgently wanted. From a scientist's perspective and all the caveats in mind, it seems to be a matter of time when cannabinoid compounds will be used in the treatment of GI disease

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Abbreviations

ACEA arachidonyl-2'-chloroethylamide

ACF aberrant crypt foci

2-AG 2-arachidonoylglycerol

AOM azoxymethane

AA-5-HAT *N*-arachidonoyl-serotonin

CB cannabinoid

CI confidence interval

CRC colorectal cancer

CD Crohn's disease

COX-2 cyclooxygenase-2

DAGL diacylglycerol lipase

DRG dorsal root ganglion

DVC dosal vagal complex

ECS endocannabinoid system

ENS enteric nervous system

FAAH fatty acid amide hydrolase

GI gastrointestinal

GPR55 G-protein coupled receptor 55

GPR119 G-protein coupled receptor 119

5-HT 5-hydroxytryptamine

IBD inflammatory bowel disease

IBS irritable bowel syndrome

MGL monoglyceride lipase

NAPE-PLD N-acyl phosphatidylethanolamine phospholipase D

NAAA N-acylethanolamine-hydrolyzing acid amidase

OEA oleoylethanolamide

PEA palmitoylethanolamide

PPARa peroxisome proliferator-activated receptor alpha

PPARγ peroxisome proliferator-activated receptor gamma

9-THC 9-tetrahydrocannabinol

TNBS trinitrobenzenesulfonic acid

TRPV1 transient receptor potential cation channel subfamily V member 1

UC ulcerative colitis

VIC visceral insular cortex.

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Key points

• The endocannabinoid system (ECS) represents an important homeostatic entity of the gut that consists of cannabinoid receptors, their endogenous ligands (the "endocannabinoids"), and their synthesizing/degrading enzymes.

- A large number of studies have confirmed that the ECS is crucially involved in the control of motility, secretion and mucosal integrity of the gut and may even determine the course of intestinal inflammation and cancer. The ECS provides many drug targets for human gastrointestinal disorders, such as irritable bowel syndrome, inflammatory bowel disease and colon cancer.
- Conduction of clinical trials and translation into clinical application of cannabinoids are important future goals in this field.

Expression of the endocannabinoid system in the human GI tract

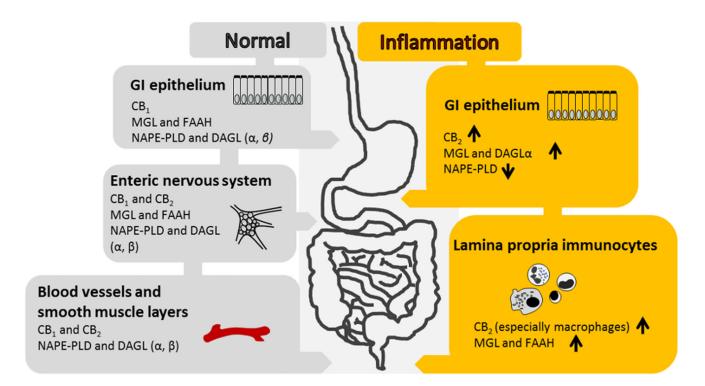


Fig. 1.Expression of receptors and synthesizing/degrading enzymes of the endocannabinoid system (ECS) in the normal and acutely inflamed human gastrointestinal (GI) tract. Data were taken from Wright et al. (5) and Marquéz et al. (7). CB₁, CB₂, cannabinoid receptors 1 and 2; FAAH, fatty acid amide hydrolase; MGL, monoacylglycerol lipase; NAPE-PLD, N-acyl phosphatidylethanolamine phospholipase D; DAGL, diacylglycerol lipase.

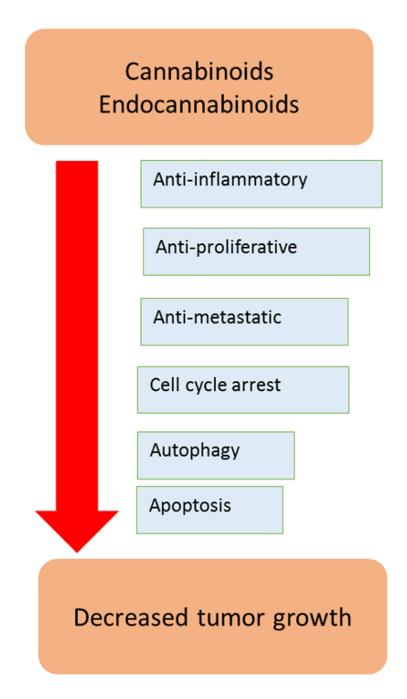


Fig. 2. (Endo-) cannabinoids exert various anti-tumorigenic effects in colon cancer. For a more detailed description of molecular mechanisms in which cannabinoids and endocannabinoids could play a role, the reader is referred to Velasco et al. (134).