

Ulcerative Colitis Impairs the Acylethanolamide-Based Anti-Inflammatory System Reversal by 5-Aminosalicylic Acid and Glucocorticoids

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Abstract

Studies in animal models and humans suggest anti-inflammatory roles on the N-acylethanolamide (NAE)-peroxisome proliferators activated receptor alpha (PPAR α) system in inflammatory bowel diseases. However, the presence and function of NAE-PPAR α signaling system in the ulcerative colitis (UC) of humans remain unknown as well as its response to active anti-inflammatory therapies such as 5-aminosalicylic acid (5-ASA) and glucocorticoids. Expression of PPAR α receptor and PPAR α ligands-biosynthetic (NAPE-PLD) and -degrading (FAAH and NAAA) enzymes were analyzed in untreated active and 5-ASA/glucocorticoids/immunomodulators-treated quiescent UC patients compared to healthy human colonic tissue by RT-PCR and immunohistochemical analyses. PPAR α , NAAA, NAPE-PLD and FAAH showed differential distributions in the colonic epithelium, lamina propria, smooth muscle and enteric plexus. Gene expression analysis indicated a decrease of PPAR α , PPAR γ and NAAA, and an increase of FAAH and iNOS in the active colitis mucosa. Immunohistochemical expression in active colitis epithelium confirmed a PPAR α decrease, but showed a sharp NAAA increase and a NAPE-PLD decrease, which were partially restored to control levels after treatment. We also characterized the immune cells of the UC mucosa infiltrate. We detected a decreased number of NAAA-positive and an increased number of FAAH-positive immune cells in active UC, which were partially restored to control levels after treatment. NAE-PPAR α signaling system is impaired during active UC and 5-ASA/glucocorticoids treatment restored its normal expression. Since 5-ASA actions may work through PPAR α and glucocorticoids through NAE-producing/degrading enzymes, the use of PPAR α agonists or FAAH/NAAA blockers that increases endogenous PPAR α ligands may yield similar therapeutics advantages.

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Introduction

Ulcerative colitis (UC) is a chronic relapsing inflammation of the colonic tissue caused by the influence of complex genetic and environmental interactions [1]. Different pro-inflammatory factors, including reactive oxygen and nitrogen metabolites, eicosanoids, platelet-activating factors and cytokines, are upregulated and actively contribute to an exacerbated intestinal immune response to otherwise innocuous stimuli [2,3]. For the management of human UC, several drugs are currently used as 5-aminosalicylic acid (5-ASA), glucocorticoids, anti-TNF α and immunomodulators as thiopurines, which interfere with pro-inflammatory cascades and effectively down-regulate the overstated inflammatory response [4–6]. Evidences suggested that the anti-inflammatory effect of 5-ASA can be mediated by the

activation of peroxisome proliferator-activated receptors (PPARs), such as PPAR α and PPAR γ [7–11], which are highly expressed in the intestinal and colonic mucosa by both epithelial cells and macrophages [12–14]. UC patients seem to have reduced levels of PPAR γ in their colonic epithelium and similar deficiencies were observed in colitis mouse models, but only in macrophages of the lamina propria [15,16]; confirming the beneficial effects of PPAR γ agonists on the attenuation of colon inflammation [17,18]. Less information is available on PPAR α and β/δ receptors.

Recent studies in experimental colitis suggested that PPAR α ligands also have anti-inflammatory properties, which was enhanced after glucocorticoid treatment, but was weakened in PPAR α -null mice [19–23]. Moreover, 5-ASA is able to induce PPAR α expression and promote its translocation to the nucleus in

Table 1. Characteristics of UC patients.

Case	Gender	Age	Smoking habits	Alcohol intake gr/day	BMI kg/m ²	Year at diagnosis	CRP [§]	UC Extension*	Mayo score at diagnosis	MTWSI [¶]	Extraintestinal manifestations	Treatment: Induction of remission	Treatment: Maintenance
1	Female	35	No	No	24.97	2006	1.4	E3	2	Moderate	No	5-ASA+ glucocorticoids	5-ASA
2	Female	29	No	No	26.10	2006	4.4	E3	2	Moderate	Apthous stomatitis	5-ASA+ glucocorticoids+ azathioprine	5-ASA+ azathioprine
3	Male	29	Yes	No	21.88	2006	7.4	E3	3	Severe	No	5-ASA+ cyclosporine	5-ASA+ azathioprine
4	Female	28	Yes	No	30.86	2006	1.2	E3	2	Moderate	No	5-ASA+ glucocorticoids	5-ASA
5	Female	46	Yes	No	28.00	2006	1.2	E3	3	Moderate	No	5-ASA+ glucocorticoids	5-ASA
6	Female	38	No	No	23.87	2006	0.5	E3	1	Mild	No	5-ASA	5-ASA
7	Male	69	No	40	22.00	2006	0.8	E3	1	Mild	No	5-ASA+ glucocorticoids	5-ASA
8	Male	20	No	No	22.98	2006	0.2	E3	3	Severe	No	5-ASA+ cyclosporine	5-ASA+ azathioprine
9	Male	23	No	No	25.01	2007	8.8	E3	2	Moderate	Arthritis	5-ASA+ glucocorticoids	5-ASA+ azathioprine
10	Female	26	Yes	No	23.42	2006	2	E3	3	Severe	Erythema nodosum	5-ASA+ glucocorticoids	5-ASA
11	Male	37	No	No	22.00	2006	0.2	E3	1	Mild	No	5-ASA	5-ASA
12	Male	48	No	100	21.24	2006	0.4	E2	2	Moderate	No	5-ASA+ glucocorticoids	5-ASA
13	Male	34	No	No	22.86	2006	33.9	E3	3	Severe	No	5-ASA+ glucocorticoids	Azathioprine+ Infliximab
14	Male	61	No	No	23.26	2006	8.9	E3	2	Severe	No	5-ASA+ glucocorticoids	5-ASA
15	Female	28	No	No	23.05	2007	1.4	E3	2	Mild	Arthritis	5-ASA	infliximab
16	Male	26	No	No	24.30	2008	0.4	E3	2	Moderate	No	5-ASA+ glucocorticoids	5-ASA
17	Male	39	No	No	22.52	2007	0.6	E2	2	Moderate	No	5-ASA+ glucocorticoids	5-ASA
18	Male	17	Yes	No	22.53	2006	3	E3	2	Moderate	No	5-ASA+ glucocorticoids	5-ASA
19	Male	62	Yes	No	25.27	2006	1.2	E3	3	Moderate	No	5-ASA+ glucocorticoids	5-ASA
20	Male	30	No	No	22.86	2006	14.9	E3	2	Moderate	Arthritis	glucocorticoids+ azathioprine	azathioprine

Table 1. Cont.

Case	Gender	Age	Smoking habits	Alcohol intake gr/day	BMI kg/m ²	Year at diagnosis	CRP ^s	UC Extension*	Mayo score at diagnosis	MTWSI [§]	Extraintestinal manifestations	Treatment: Induction of remission	Treatment: Maintenance
21	Female	42	No	No	27.34	2007	0.9	E3	2	Mild	No	5-ASA+ glucocorticoids	5-ASA+ azathioprine
22	Male	73	Yes	No	26.95	2007	8.5	E2	2	Moderate	No	5-ASA+ glucocorticoids	5-ASA
23	Male	44	No	No	23.98	2006	2.9	E2	2	Moderate	No	5-ASA+ glucocorticoids+ cyclosporine	5-ASA+ azathioprine
24	Female	62	No	No	24.00	2006	0.6	E3	1	Moderate	No	5-ASA	5-ASA

^sC-reactive protein at diagnosis (mg/L);

*Ulcerative colitis extension by Montreal criteria;

[§]Modified Truelove–Witts Severity index (MTWSI): Mild 11–15 points, Moderate 16–20 points, Severe 21–27 points. doi:10.1371/journal.pone.0037729.t001

an animal model of irradiation-induced intestinal inflammation [11]. PPAR α is specifically expressed in the more differentiated colonic epithelial cells facing the intestinal lumen of the small intestine and colon [12–14]. Thus, PPAR α has been proposed to participate in the intestinal epithelial barrier system; absence of PPAR α expression resulted in an increase of tight junction permeability associated with apoptosis in an animal model of experimental colitis [20]. PPAR α signaling system is an anti-inflammatory system composed of the PPAR α receptor and its endogenous ligands, the N-acyl ethanolamides oleoylethanolamide (OEA) and palmitoylethanolamide (PEA). It also includes the enzymes involved in their biosynthesis and release, such as N-acyl phosphatidylethanolamide-specific phospholipase D (NAPE-PLD), as well as mechanisms for cellular uptake and hydrolysis, such as fatty acid amide hydrolase (FAAH) and N-acyl ethanolamide-hydrolyzing acid amidase (NAAA) [24–26]. Increased PPAR α expression or enhanced PPAR α ligand production can attenuate inflammatory process observed in current animal models of experimental colitis. For instance, treatment with FAAH antagonists or genetic ablation of FAAH protected against colitis inflammation [27–29]. Thus, PPAR α system is positioned to exert a putative role in many of the points where homeostasis breaks in UC, although the anti-inflammatory role of PPAR α remains to be determined in humans.

The aim of the present study is to analyze the expression and distribution of components of the acylethanolamide-PPAR α anti-inflammatory system such as PPAR α receptor and the enzymes involved in endogenous ligand degradation (FAAH and NAAA) and biosynthesis (NAPE-PLD) in the normal human colonic tissue compared to untreated active UC at disease onset and after achieving remission, according to clinical and endoscopic criteria, and depending on treatment received (5-ASA, glucocorticoids and/or immunomodulators).

Methods

Ethics statement

Biopsies and colonic resection samples used for the present study were obtained after a written inform consent from all the patients, as requested by the clinical guides of Hospital del Mar. Research procedures were approved by the Hospital del Mar and Hospital Carlos Haya Clinical Research and Ethics Committee and were conducted according to the principles expressed in the Declaration of Helsinki.

Subjects

We selected retrospectively 24 consecutive patients diagnosed from January 2006 to December 2007 of a first flare of UC, with extensive or left-side extension according to the Montreal classification (E2 and E3) [30]. UC was defined by the criteria of Lennard-Jones [31]. All patients had to achieve clinical and endoscopic remission after medical treatment according to Truelove–Witts index (<6 points) [32] and endoscopic Mayo score (score = 0) [33] during 12 months of diagnosis. We excluded patients with distal UC according to Montreal classification (E1) and patients without clinical and endoscopic remission criteria after treatment. Thus, we obtained several endoscopic samples of UC mucosa from each patient collected before any treatment (active group), and after medical treatment and endoscopic remission (quiescent group).

Colonic samples were retrieved from tissue bank of Pathology Service at the Hospital del Mar. Data from each patients were collected retrospectively from medical records including age, sex, smoking habits and alcohol history, body mass index and

Table 2. Clinical characteristics of control group.

Case	Age at diagnosis	Gender	Smoking habit	Alcohol	*BMI kg/m ²	Co-morbidity	Medical treatment	Colon cancer location	Year at diagnosis
1	74	Male	No	No	23.20	Diabetes mellitus; dislipemia; hypertension	Tonvastatine Metformine	Sigmoid colon	2006
2	76	Female	No	No	20.80	Diabetes mellitus; hypertension	Gilbenclamide Enalapril	Rectal	2006
3	68	Male	No	No	28.10	Diabetes mellitus; hypertension	Enalapril Atenolo	Rectal	2006
4	78	Male	No	No	31.98	Diabetes mellitus; hypertension	Gilbenclamide Enalapril	Right colon	2006
5	80	Male	No	No	27.30	Diabetes mellitus; dislipemia; hypertension	Metformine Insulin Bisoprolol	Cecum	2006
6	74	Female	No	No	28.23	Hypertension	Enalapril	Sigmoid colon	2006
7	56	Male	No	No	19.81	No	No	Sigmoid colon	2006
8	61	Female	No	No	24.25	No	No	Rectal	2006
9	68	Female	No	No	28.98	No	No	Sigmoid colon	2006
10	74	Male	No	No	29.09	Diabetes mellitus; hypertension	Metformine Enalapril	Sigmoid colon	2006
11	79	Female	No	No	26.40	Hypertension	Enalapril	Right colon	2006
12	56	Male	Yes	No	30.91	Dislipemia	Simvastatine	Sigmoid colon	2006
13	62	Female	No	No	26.75	Hypertension Atrial fibrillation	Digoxine; Warfarine	Rectal	2006
14	80	Male	No	No	21.64	No	No	Right colon	2006
15	69	Female	No	No	21.50	No	No	Rectal	2006
16	79	Male	No	No	34.41	No	No	Cecal	2006
17	68	Female	No	No	26.74	No	No	Sigmoid colon	2006
18	76	Male	No	No	26.12	Diabetes mellitus;hypertension	Metformine Enalapril;	Right colon	2006
19	67	Female	No	No	22.00	No	No	Descending colon	2006
20	67	Male	No	No	22.00	No	No	Sigmoid colon	2006
21	78	Male	No	No	23.80	Diabetes mellitus	No	Right colon	2006
22	75	Female	No	No	20.60	Hypertension	Enalapril	Sigmoid colon	2006

*BMI: body mass index Kg/m²; Medical treatment at the time to take samples of colon mucosa.
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Table 3. Primers sequences used for RT-PCR.

Gene symbol (name)	Oligosense (5'→3')	Oligoantisense (5'→3')	GenBank® accession no.	Product size (bp)	Annealing temperature (C°)
SP1	AGCAGGATGGTTCTGGTCAA	AGGTGATGTTCCCATTCAGG	NM_138473.2	210	54.0
NAPE-PLD	CACGGTAATGGTGGAAATGG	GTCAGATGGTCATAGTGGTTG	NM_001122838.1	178	57.0
FAAH	CCCAGATGGAACATTACAGG	CAGGATGACTGGTTTTCAGG	NM_001441.2	187	57.6
NAAA	ATGGCGCAAGTCATCGGGGA	TGAAGTCACACATGCCGCGGA	NM_014435.3	127	59.0
PPAR α	ATCACGGACACGCTTTCAC	GGTCGCACTGTGCATACACC	NM_001001928.2	220	58,9
PPAR γ	TGCCATCAGGTTTGGCGGA	AATGTTTTGCCAGGCCCGGA	NM_138712.3	118	61.4
iNOS	TCAGCAAGCAGCAGAATGAG	ATAATGGACCCAGGCAAGA	NM_000625.4	210	63.3

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extraintestinal manifestations, date of diagnosis, disease location (Montreal classification), endoscopic lesions (Mayo clinic score) and clinical score according Truelove and Witts index at onset, and medical treatment received to induce remission after diagnosis: 5-aminosalicylates (3 cases), glucocorticoids (15 cases), and/or the immunomodulators cyclosporine A and azathioprine (6 cases) (Table 1).

For the control group, we selected retrospectively 22 patients who were pathohistologically given a definite diagnosis of colorectal cancer and who had not received preoperative radiotherapy or chemotherapy treatment and underwent colonic resections for colorectal cancer. Several colonic resections were obtained from each patient at least 10 cm from the tumour (control group). We confirmed histopathologically the absence of microscopic alterations (Table 2).

Colonic samples were frozen at -80°C for molecular analysis ($N=7-8$) or fixated with 4% paraformaldehyde in 0.1 M phosphate buffered saline (PBS) by immersion and included in paraffin until immunohistochemical analysis ($N=22-24$). The analysis of the immunostaining patterns was carried out at transmural planes of the normal and pathological colonic tissue by comparing it with hematoxylin-eosin staining.

mRNA isolation and quantitative RT-PCR analysis

In order to evaluate the mRNA expression we collected prospectively 7 colonic endoscopic biopsies from patients with a first flare of active UC and 8 colonic resections, at least 10 cm from the tumour, of patients with colorectal cancer (control group). Colonic resections were divided into mucosa, containing both epithelium and lamina propria, and submucosa layers, containing smooth muscle and enteric plexi. Reverse transcript reaction was carried out from 4 μg of mRNA using the Transcriptor Reverse Transcriptase kit and random hexamer primers (Transcriptor RT, Roche Diagnostic GmbH, Mannheim, Germany). Quantitative real-time reverse transcription polymerase chain reaction (quantitative RT-PCR) was performed using a CFX96TM Real-Time PCR Detection System (Bio-Rad Laboratories, Hercules, CA, USA), and the SYBR Green detection format (FastStart Universal Master Kit, Roche, Mannheim, Germany). Each reaction was run in duplicate and contained 5 μl of cDNA. Quantification was carried out with the classic standard curve method run at the same time. We analyzed the housekeeping genes SP1 transcription factor and βACTIN , selecting the most suitable according to their homogeneity (Figure S1). Absolute values from each sample were normalized with regard to the housekeeping gene SP1. Primers for PCR reaction were designed based on NCBI database sequences of human

reference mRNA (Table 3), checked for specificity with BLAST software from NCBI website (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>) and synthesized by Invitrogen.

Western blotting

In order to evaluate the presence of PPAR α , NAAA, NAPE-PLD and FAAH in the colon mucosa we collected prospectively 8 colonic resection of control patients processed as previously described [34,35]. Each blotted membrane lane was incubated separately with the specific rabbit anti-PPAR α (1:100; Fitzgerald, cat. no. RDI-PPARAabrx), rabbit anti-NAAA (1:1000; R&D Systems, cat. no. AF4494), rabbit anti-NAPE-PLD (1:100) and rabbit anti-FAAH (1:100) antibodies [35], overnight at 4°C . Western blots showed that each primary antibody detects a protein of the expected molecular weight (see Methods S1).

Immunohistochemistry

We analyzed the distribution of PPAR α , NAAA, NAPE-PLD and FAAH in the normal colonic tissue and in the active and quiescent UC mucosa by immunohistochemistry, following methods previously described in Marquez et al [35]. Sections were incubated overnight at room temperature with rabbit anti-PPAR α antibody (diluted 1:75; Fitzgerald), rabbit anti-NAAA (diluted 1:200; R&D Systems), rabbit anti-NAPE-PLD antibody (diluted 1:100) and rabbit anti-FAAH (diluted 1:100). Then, sections were incubated in a biotin-conjugated donkey anti-rabbit immunoglobulin (Amersham) diluted 1:500 for 1 hour, and incubated in ExtrAvidin peroxidase (Sigma) diluted 1:2000 for 1 hour. We revealed immunolabeling with 0.05% diaminobenzidine (DAB; Sigma), 0.05% nickel ammonium sulphate, and 0.03% H_2O_2 in 0.1 M phosphate-buffered saline (pH 7.4). Sections were dehydrated in ethanol, cleared in xylene, and coverslipped with Eukitt mounting medium (Kindler GmbH and Co., Freiburg, Germany).

Double immunofluorescence

Paraffin-embedded sections of colonic tissue were analyzed for the presence of NAAA, NAPE-PLD and FAAH in plasma cells (CD38+), B lymphocytes (CD19+), T lymphocytes (CD3+) and macrophages (CD14+) of the lamina propria of control and UC colitis groups. Sections were incubated overnight at room temperature in a cocktail containing rabbit anti-NAAA, NAPE-PLD or FAAH antibody (see above) and mouse monoclonal anti-human CD14-IgG1 conjugated to R-phycoerythrin-Cy7 (eBioscience, San Diego, CA, USA, cat. no. 25-0149), anti-human CD3-IgG1 conjugated to R-phycoerythrin-Cy7 (eBioscience, cat. no. 25-0038) or conjugated to eFluor[®] 450 (eBioscience, cat.

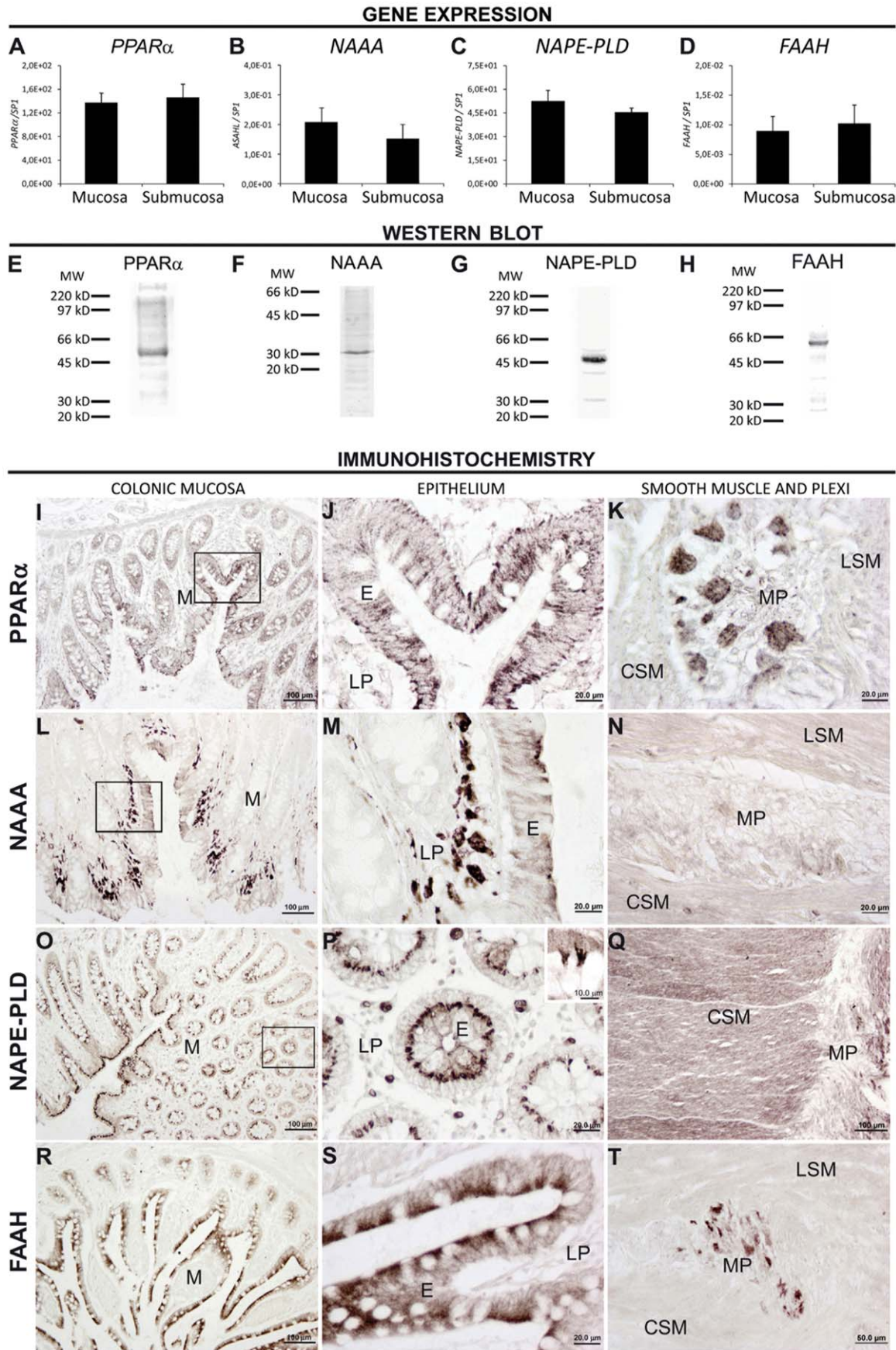


Figure 1. RT-PCR (A–D), Western blot (E–H) and immunohistochemical analyses (I–T) showing the presence and distribution of PPAR α , NAAA, NAPE-PLD and FAAH in the normal human colonic tissue. Gene expressions of PPAR α , NAAA, NAPE-PLD and FAAH were detected in both mucosa (epithelium and lamina propria) and submucosa (smooth muscle and plexi) layers (A–D), and confirmed by high-magnification photomicrographs of their protein expression by immunohistochemistry (I–T). Western blots of protein extracts from human colonic

tissue showed prominent immunoreactive bands of the expected size for PPAR α (52 kDa), NAAA (31 kDa), NAPE-PLD (46 kDa) and FAAH (62 kDa). Positions of molecular markers (MW) are indicated at the left (E–H). Abbreviations: CSM, circular smooth muscle; E, epithelium; LSM, longitudinal smooth muscle; LP, lamina propria; M, mucosa; MP, myenteric plexus.
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Table 4. Rating scale that summarizes the immunohistochemical distribution in the normal human colonic tissue (n = 24)¹.

	Epithelium	Lamina propria	Smooth muscle	Myenteric plexus
PPAR α	+++	–	–	++
NAAA	+	+++	+	–
NAPE-PLD	++	+	+++	–
FAAH	++	+	–	++

¹Gray-scale values measured in single epithelium, lamina propria, muscular layers and plexi are represented on an arbitrary rating scale of the immunoreactivity of each structure. Symbols are as follows: high (+++), moderate (++) , low (+) and without immunoreactivity (–).
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no. 48-0038), anti-human CD19-IgG2a conjugated to R-Phycoerythrin (Immunostep, Salamanca, Spain, cat. no. 19PE1-100T) or anti-human 38-IgG1 conjugated to fluorescein isothiocyanate (Immunostep, cat. no. 38F-100T). Then, the sections were incubated for 2 hours at room temperature in secondary donkey anti-rabbit IgG-Cy3 antibody (dilution 1:300; Jackson Immuno-research Laboratories, West Grove, PA, USA, cat. no. 711-165-152) or goat anti-rabbit IgG-FITC antibody (dilution 1:300; Jackson Immuno-research Laboratories, cat. no. 111-095-003).

Quantification of mucosa immunostaining

For epithelium, we carried out a densitometrical quantification for each protein. For lamina propria, we evaluated the type and the number of immunostained immune cells per area (μm^2) analyzed. In addition, quantification was segregated depending on UC severity and treatment received: 5-ASA, glucocorticoids, and/

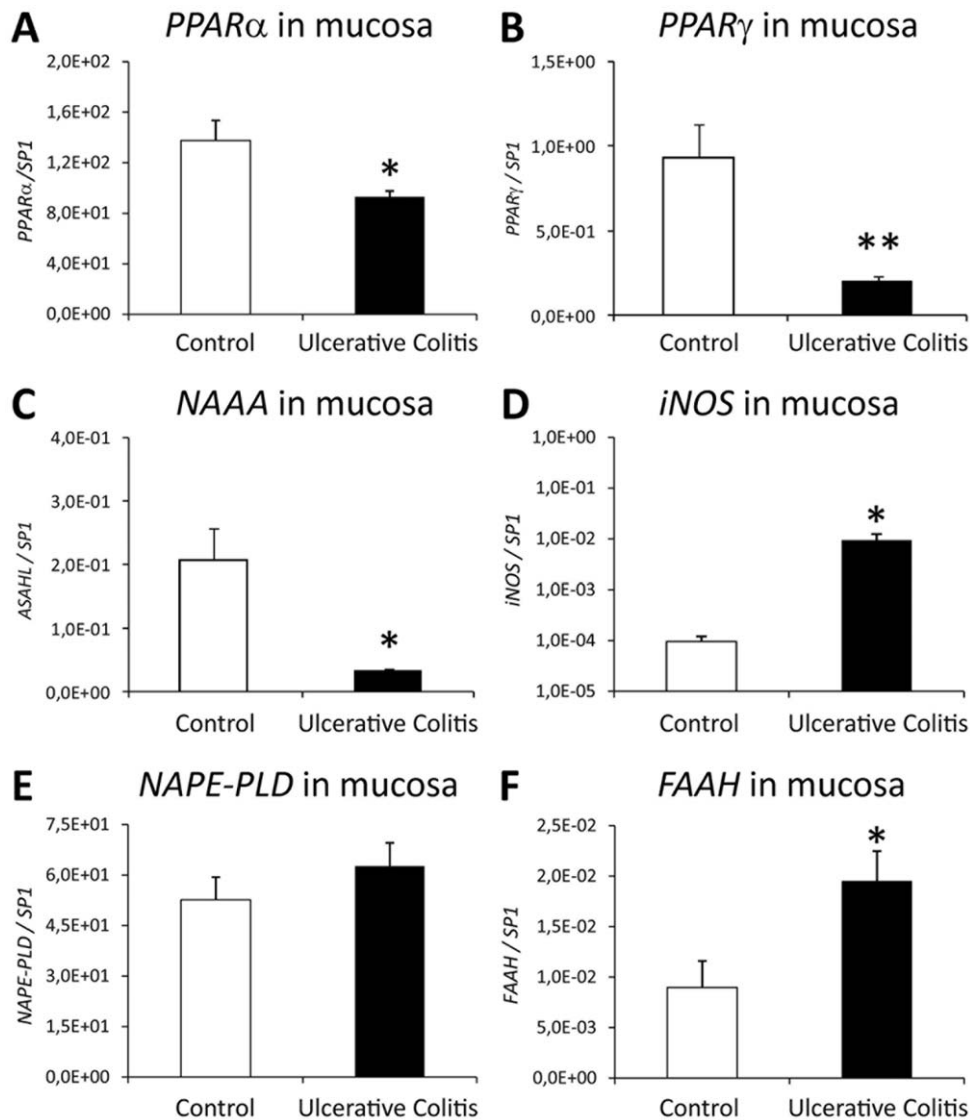


Figure 2. Relative quantification of PPAR α (A), PPAR γ (B), NAAA (C), iNOS (D), NAPE-PLD (E) and FAAH (F) gene expression in the colonic mucosa of active UC patients compared to human healthy colonic tissue (control). Absolute values were normalized with regard to the housekeeping gene SP1. Active UC at disease onset showed lower PPAR α , PPAR γ and NAAA gene expression, but higher iNOS and FAAH gene expression compared to control. No change was detected for NAPE-PLD gene expression. Student *t* test (N = 8): **P* < 0.05, ***P* < 0.01 versus control group.
doi:10.1371/journal.pone.0037729.g002

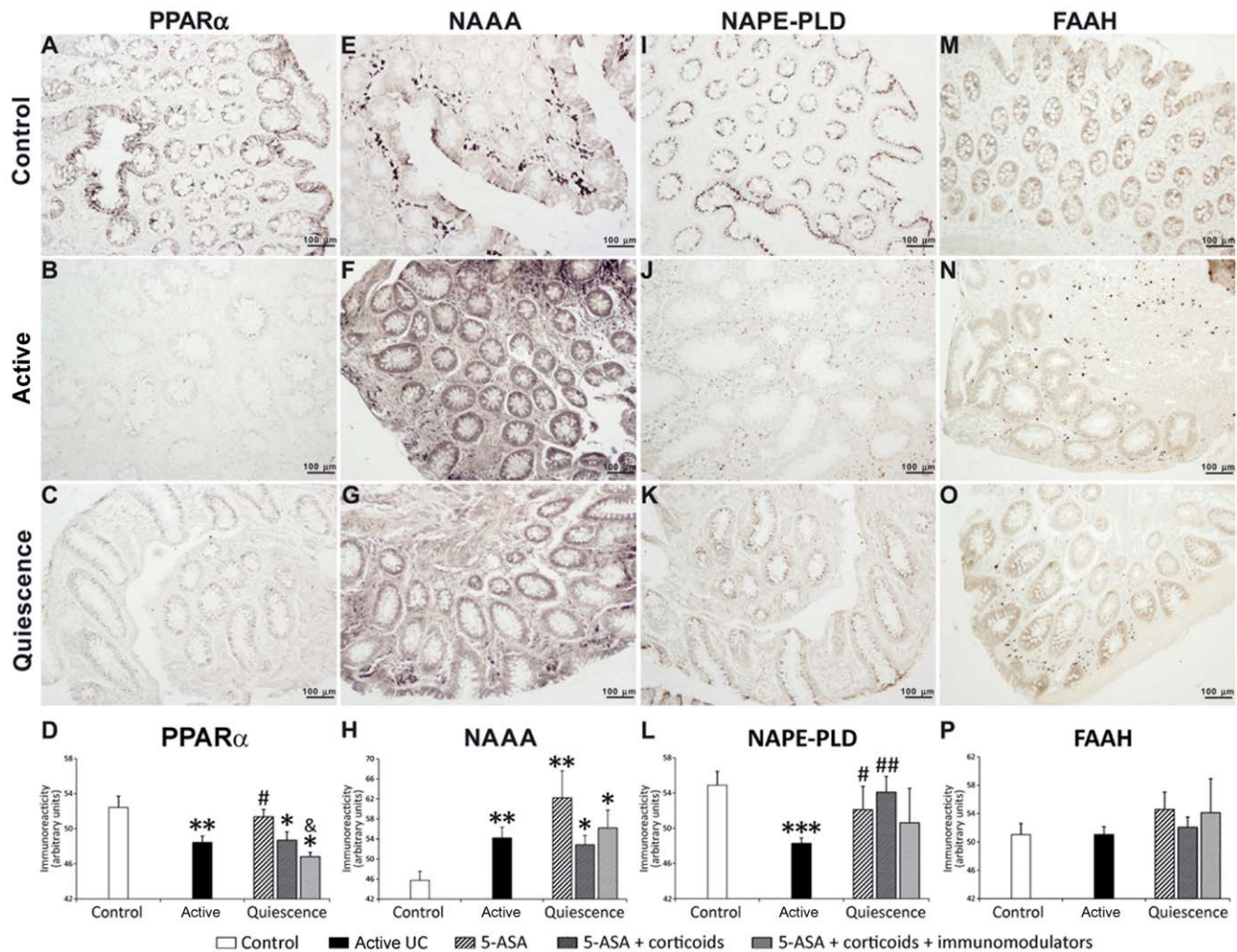


Figure 3. Representative photomicrographs and densitometrical quantification of PPAR α (A–D), NAAA (E–H), NAPE-PLD (I–L) and FAAH (M–P) immunoreactivity in human healthy (control; A, E, I, M), active UC (B, F, J, N) and quiescent UC (C, G, K, O) colonic epithelium depending on treatment. Active UC at disease onset showed a decrease in PPAR α and NAPE-PLD immunoreactivity, but an increase in NAAA immunoreactivity. PPAR α immunoreactivity showed significant differences depending on treatment. Treatment with 5-ASA in UC patients restored completely PPAR α and NAPE-PLD protein levels to control ones. However, PPAR α expression dropped again in UC patients treated with 5-ASA and glucocorticoids, and 5-ASA, glucocorticoids and immunomodulators compared to control group. In contrast, treatment with 5-ASA and glucocorticoids increased NAPE-PLD immunoreactivity. No changes were observed in FAAH immunoreactivity in the epithelium of control, active (untreated) UC and quiescent (treated) UC patients. Mann-Whitney U and Wilcoxon tests (N = 22–24): * P < 0.05, ** P < 0.01, *** P < 0.001 versus control group; # P < 0.05, ## P < 0.01 versus UC group; &#P < 0.05 versus 5-ASA-treated quiescent UC group. doi:10.1371/journal.pone.0037729.g003

or immunomodulators. Digital high-resolution microphotographs were taken under the same conditions of light and brightness/contrast by an Olympus BX41 microscope equipped with an Olympus DP70 digital camera and a Metal Halide epifluorescence system (Olympus Europa GmbH, Hamburg, Germany).

Statistical analysis

Data were analyzed using SPSS 15.0 software (Statistical Package for the Social Sciences Inc., Chicago, Illinois, USA). Results are expressed as mean \pm S.E.M. Differences between groups were evaluated using Student t test for parametric observation and Mann-Whitney U and Wilcoxon tests for non parametric observations. A P value of P < 0.05 was considered statistically significant.

Results

Presence and distribution of PPAR α , NAAA, NAPE-PLD and FAAH in the normal human colonic tissue

The normal colonic tissue showed gene expression of PPAR α , NAAA, NAPE-PLD and FAAH in the mucosa, including epithelium and lamina propria, and the submucosa layers, containing smooth muscle and enteric plexi (Figs. 1A–D). Protein extracts from normal colonic tissue confirmed the presence of protein levels of PPAR α , NAAA, NAPE-PLD and FAAH. They appeared as prominent immunoreactive bands of expected molecular masses at \sim 52 kDa for PPAR α , \sim 31 kDa for NAAA, \sim 46 kDa for NAPE-PLD and \sim 62 kDa for FAAH (Figs. 1E–H).

Results of the immunohistochemical distribution were summarized in a rating scale (Table 4). PPAR α immunoreactivity was observed in the colonic epithelium of both absorptive and goblet

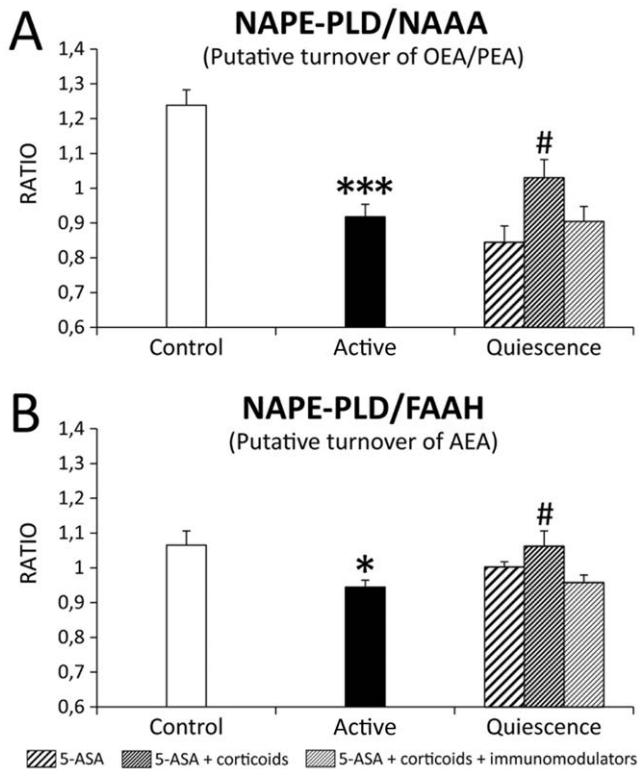


Figure 4. Estimation of acylethanolamide turnover by analyzing the NAPE-PLD/NAAA (A) and NAPE-PLD/FAAH (B) ratios in the epithelium of the untreated active and treated quiescent UC patients in comparison with control patients. Active UC produced a significant decrease in NAPE-PLD/NAAA and NAPE-PLD/FAAH ratios, which were restored to control levels only after the treatment with 5-ASA and glucocorticoids. Mann-Whitney U and Wilcoxon tests (N = 22–24): * $P < 0.05$, *** $P < 0.001$ versus control group; # $P < 0.05$ versus active UC group.
doi:10.1371/journal.pone.0037729.g004

cells (Figs. 1I, J). The immunoreactivity filled the epithelial cells, showing prominent staining in the apical and basal portions. We did not detect PPAR α immunoreactivity in immune cells of the lamina propria, the muscularis mucosae, the muscularis externa (circular and longitudinal smooth muscle) and the serosa (Figs. 1I, J). Numerous PPAR α -immunopositive ganglion cells were evident only in the myenteric plexi (Figure 1K). Moderate to low intensity of NAAA immunostaining was observed in the colonic epithelium (Figs. 1L, M). Interestingly, we detected numerous NAAA immune cells in the lamina propria, which showed a variety of shapes and sizes (Fig. 1M). Muscularis mucosae, muscularis externa, plexi and serosa showed very weak staining for NAAA (Fig. 1N). Intense NAPE-PLD immunoreactivity was widely distributed in the colonic epithelium, being prominent in the perinuclear portion of the absorptive cells (Figs. 1O, P, inset). Some positive plasma cells were also observed in the lamina propria (Fig. 1P). Strong NAPE-PLD immunostaining defined both layers of muscularis externa, but low immunostaining was detected in fibers of the myenteric plexi (Fig. 1Q). FAAH immunoreactivity was mainly detected in the colonic epithelium, which shows higher expression in the apical portion of the epithelial cells (Figs. 1R, S). A low number of immunoreactive immune plasma cells were observed in the lamina propria and no staining was detected in the muscularis mucosae, the muscularis externa and the serosa. However, we can

observe a specific FAAH immunoreactivity in nervous cells of the myenteric plexi (Fig. 1T).

Quantification of PPAR α , PPAR γ , NAAA, iNOS, NAPE-PLD and FAAH gene expression in the mucosa of UC patients

In order to evaluate any changes on the expression of PPAR α signaling system in the colonic mucosa (epithelium and lamina propria) of UC patients, we analyzed the relative differences in the mRNA levels of selected genes such as PPAR α , PPAR γ , NAAA, iNOS, NAPE-PLD and FAAH in the UC mucosa, containing epithelium and lamina propria, by quantitative RT-PCR. We detected significantly lower levels of PPAR α ($P < 0.05$), PPAR γ ($P < 0.01$) and NAAA ($P < 0.05$) mRNA in the mucosa of UC patients compared to that of control ones (Figs. 2A–C). In contrast, iNOS and FAAH gene expression was significantly higher in the mucosa of UC patients ($P < 0.05$) (Figs. 2D, F). We observed no change in the levels of NAPE-PLD mRNA between both groups (Fig. 2E).

Densitometrical quantification of PPAR α , NAAA, NAPE-PLD and FAAH immunoreactivity in the epithelium of UC patients depending on treatment

Figure 3 shows representative microphotographs showing qualitative differences of the immunohistochemical expression of PPAR α , NAAA, NAPE-PLD and FAAH in the colonic epithelium of control, active and quiescent groups. Results corresponding to the quantification of immunoreactivity are shown in Figures 3D, H, L, P respectively. We detected a decrease of PPAR α and NAPE-PLD immunoreactivity in the epithelium of UC patients compared to that of control ones ($P < 0.01$ and $P < 0.001$ respectively) (Figs. 3D, L). In contrast, NAAA immunoreactivity was more prominent in the epithelium of active UC patients ($P < 0.01$) (Fig. 3H). No change was detected in FAAH immunoreactivity in the epithelium between active UC and control groups (Fig. 3P). In order to address the disease severity, we analyzed the NAPE-PPAR α signaling system depending on the clinical score (mild, moderate and severe) in active UC patients (Figure S2). UC patients with moderate clinical score showed a significant reduction ($P < 0.05$) of PPAR α immunohistochemical expression (Figure S2A). When NAPE-PLD immunoreactivity was analyzed, we detected significant decreases in UC patients with mild ($P < 0.01$), moderate ($P < 0.01$) and severe ($P < 0.001$) clinical score (Figure S2B). However, FAAH immunoreactivity was not affected (Figure S2C). Finally, UC patients with moderate ($P < 0.05$) and severe ($P < 0.01$) clinical score showed significant increases in NAAA immunoreactivity (Figure S2D). We also analyzed the possible effect of gender and smoking habits on the NAPE-PPAR α signaling system in active UC patients. We did not detect differences between females and males or between smokers and non-smokers in the immunohistochemical expression in the epithelium of active UC patients (Figures S3 and S4).

We also quantified the immunoreactivity in the colonic epithelium of quiescent UC patients depending on the treatment received: 5-ASA (3 cases), 5-ASA and glucocorticoids (15 cases), or 5-ASA, glucocorticoids and immunomodulators (6 cases). 5-ASA treatment produced an increase of PPAR α immunoreactivity in the colonic epithelium of active UC patients ($P < 0.05$), but not when UC patients were treated with 5-ASA in combination with other drugs (Fig. 3D). Thus, no difference in PPAR α immunoreactivity was observed in UC patients treated with 5-ASA plus glucocorticoids only or 5-ASA, glucocorticoids and immunomodulators with respect to untreated active UC ones. Interestingly, the decrease in PPAR α immunoreactivity observed in the epithelium

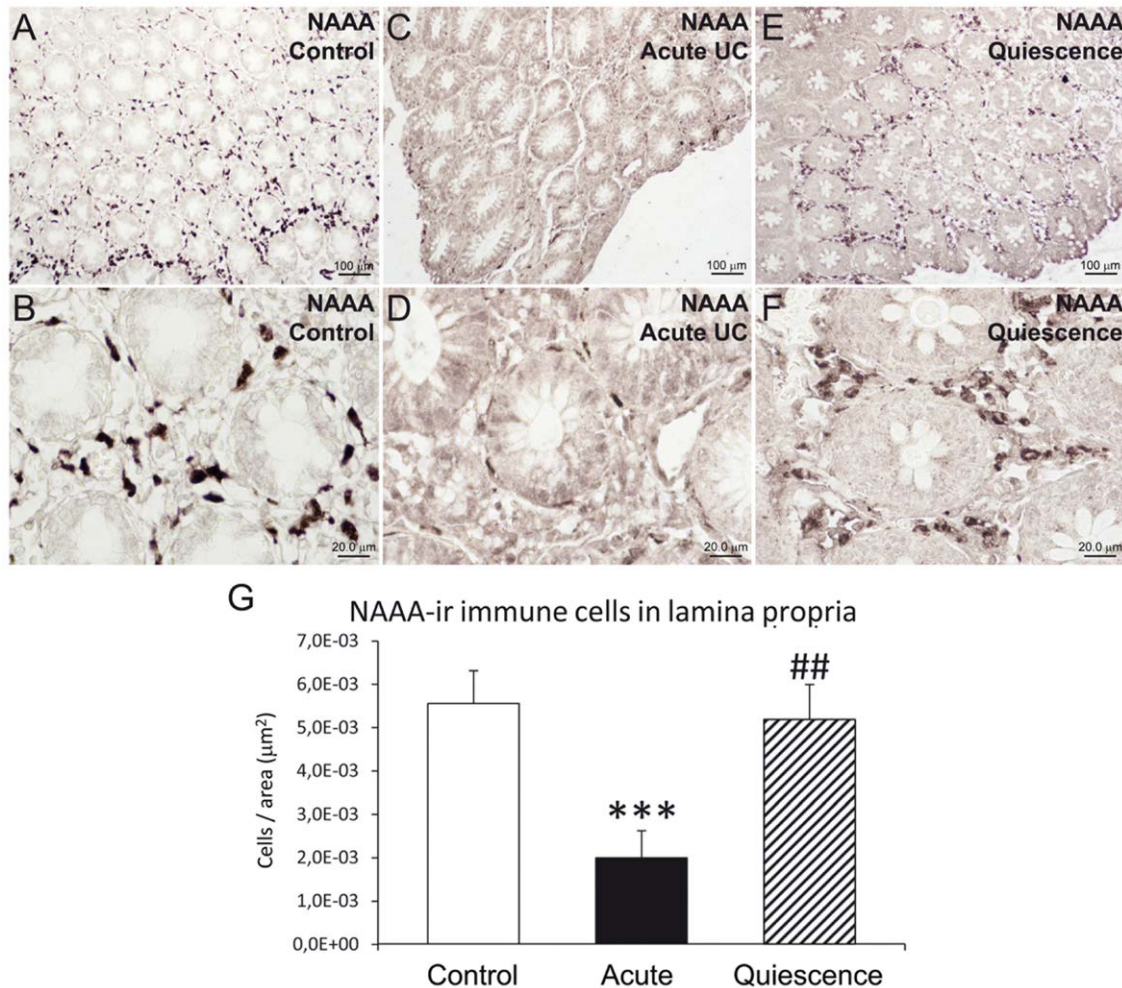


Figure 5. Analysis of the number of NAAA-ir cells per area (μm^2) in the lamina propria of control group compared to acute UC patients and after 5-ASA and corticosteroid treatment (quiescent group). A-F: Representative high-magnification photomicrographs showing NAAA immunostaining in the lamina propria. G: Acute UC at disease onset was associated with a significant decrease in the number of NAAA-ir cells in the infiltrate of the lamina propria. The number of NAAA-ir cells was increased to control levels after treatment (5-ASA and corticoids). Mann-Whitney U and Wilcoxon tests (N = 15–22): *** $P < 0.001$ versus control group; ## $P < 0.01$ versus acute UC group. doi:10.1371/journal.pone.0037729.g005

of UC patients treated with 5-ASA, glucocorticoids and immunomodulators was significant compared to that of UC patients treated with 5-ASA only ($P < 0.05$). However, we cannot detect significant changes in NAAA immunoreactivity in the epithelium of quiescent UC patients treated with any of the drugs, being similar to that of active UC patients (Fig. 3H). Regarding NAPE-PLD immunoreactive levels, there was a significant increase to control levels in UC patients treated with 5-ASA ($P < 0.05$) or 5-ASA and glucocorticoids ($P < 0.01$) compared to the active UC patients (Fig. 3L). However, a wide variability in the intensity of NAPE-PLD immunoreactivity was detected in UC patients treated with 5-ASA, glucocorticoids and immunomodulators. Finally, we cannot observe any difference in FAAH immunoreactivity in the UC epithelium after treatment (Fig. 3P).

Acylethanolamide producing/degrading enzyme ratio in the colonic epithelium

In order to analyze whether the differential immunohistochemical expression of either acylethanolamide producing or degrading enzymes may have resulted in an altered PPAR α endogenous ligand tone in the untreated active and treated quiescent UC

epithelium, we calculated the ratios between NAPE-PLD and NAAA expressions, and between NAPE-PLD and FAAH expressions. These ratios can suggest possible changes of OEA/PEA levels (Fig. 4). The main result of these analysis was that there was a significant decrease of both NAPE-PLD/NAAA ($P < 0.001$) and NAPE-PLD/FAAH ($P < 0.05$) ratios in the epithelium of untreated active UC patients. Interestingly, we detected an increase of NAPE-PLD/NAAA and NAPE-PLD/FAAH ratios (both at $P < 0.05$) only in the epithelium of quiescent UC patients treated with 5-ASA and glucocorticoids, but not with 5-ASA or 5-ASA, glucocorticoids and immunomodulators (Fig. 4).

NAAA, NAPE-PLD and FAAH immunoreactive cells in the lamina propria of UC patients and after treatment

The number of NAAA and FAAH immunoreactive cells in the lamina propria showed significant changes in active UC patients and after treatment (quiescent group). We found a significant low number of NAAA-ir cells (2.75-fold; $P < 0.001$) in the infiltrate of active UC patients, but was completely restored, similar to control level, after treatment (Fig. 5). Performing double immunofluorescence, NAAA expression was found in CD19-positive (+) B

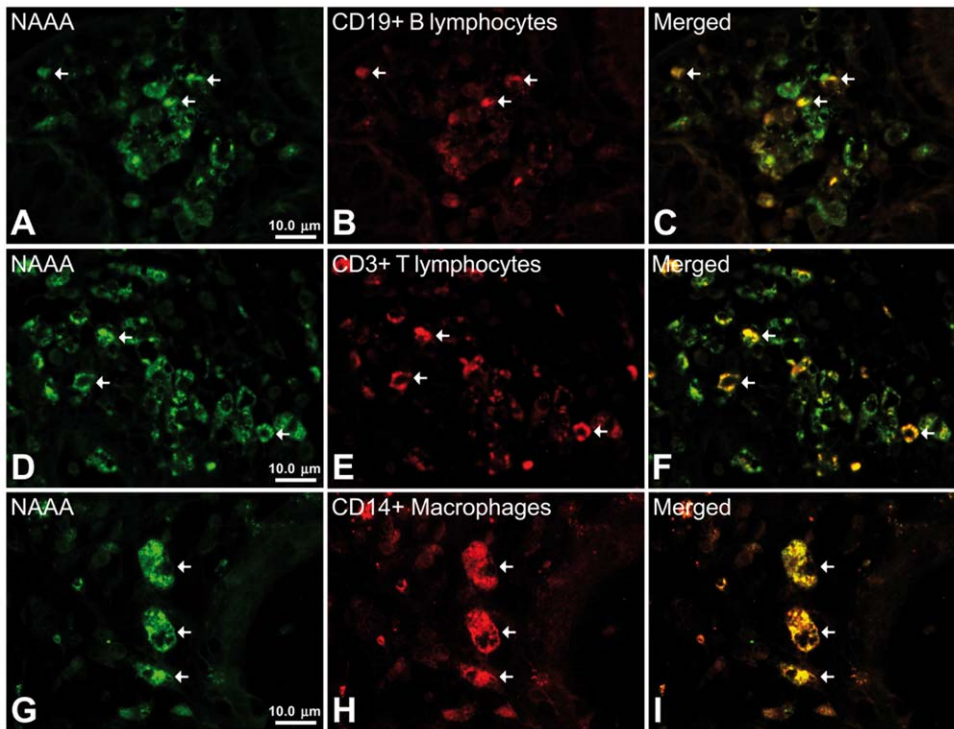


Figure 6. Representative high-magnification photomicrographs showing double immunofluorescence for NAAA, CD19, CD3 and CD14 in order to characterize the immune cells in the mucosa infiltrate of UC patients. NAAA immunofluorescence was observed in CD19+ B lymphocytes (A–C), CD3+ T lymphocytes (D–F) and CD14+ macrophages (G–I). doi:10.1371/journal.pone.0037729.g006

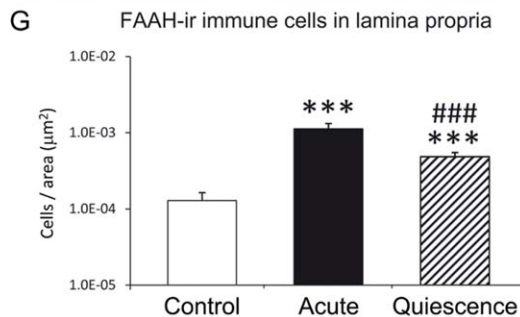
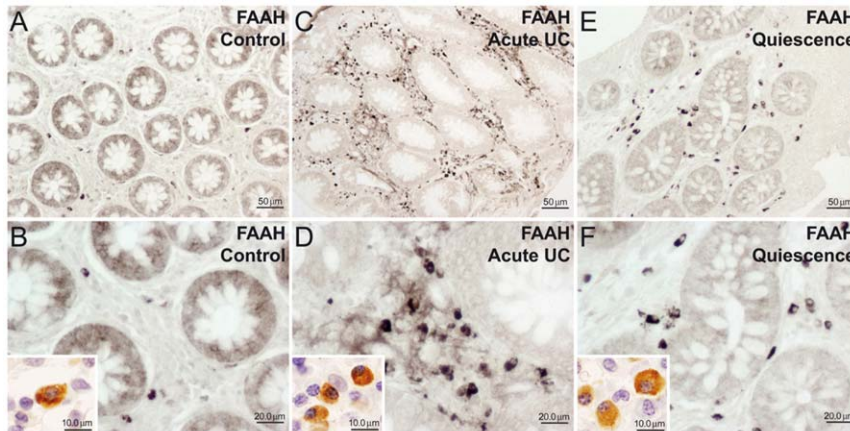


Figure 7. Analysis of the number of FAAH-ir cells per area (μm^2) in the lamina propria of acute and quiescent (5-ASA and corticoid-treated) UC patients compared to control ones. A–F: Representative high-magnification photomicrographs showing FAAH immunostaining in the lamina propria. G: Acute UC at disease onset was associated with a dramatic increase in the number of FAAH-ir cells in the infiltrate of the lamina propria. The number of FAAH-ir cells was significantly dropped after treatment, but do not reach control levels. Mann-Whitney U and Wilcoxon tests (N = 15–22); *** $P < 0.001$ versus control group; ### $P < 0.001$ versus acute UC group. doi:10.1371/journal.pone.0037729.g007

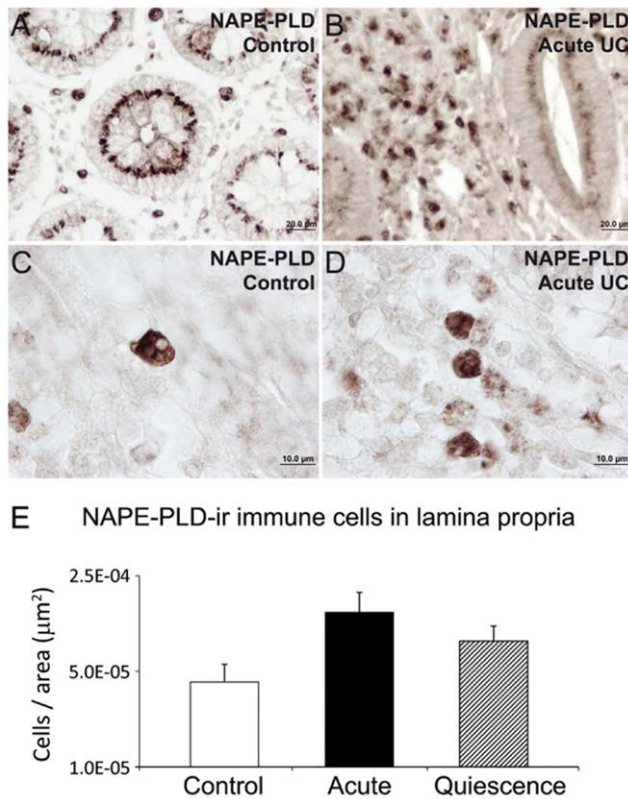


Figure 8. Analysis of the number of NAPE-PLD-ir cells per area (μm^2) in the lamina propria of acute and quiescent (5-ASA and corticoid-treated) UC patients compared to control ones. A–D: Representative high-magnification photomicrographs showing NAPE-PLD immunostaining in the lamina propria. E: No significant change in the number of NAPE-PLD-ir cells was observed in the infiltrate of the lamina propria of acute UC patients and after treatment (5-ASA and corticoids). Mann-Whitney U and Wilcoxon tests (N = 15–22). doi:10.1371/journal.pone.0037729.g008

lymphocytes, CD3+ T lymphocytes and CD14+ macrophages (Fig. 6). In contrast, the number of FAAH-ir cells of the lamina propria increased dramatically (10-fold; $P < 0.001$) in active UC patient (Fig. 7). After treatment, the number of FAAH-ir cells decreased significantly in quiescent UC patients ($P < 0.001$), but did not reach control level. We did not find change in the number of NAPE-PLD-ir cells in the lamina propria of active UC patients and after treatment (Fig. 8). Nearly all the FAAH-ir cells in the infiltrate expressed the plasma cell-specific CD38, which include B lymphocytes and natural killer cells (Figs. 9A–C). NAPE-PLD expression was mainly found in CD38+ plasma cells and CD3+ T lymphocytes (Figs. 9 D–I).

Discussion

The key findings of this study are to demonstrate that profound changes in the acylethanolamide-PPAR α anti-inflammatory system are produced in human UC. Overall the findings suggest that active UC deactivate this anti-inflammatory system while 5-ASA/glucocorticoids treatment restores its normal expression (Table 5). The process involves both receptors and enzymes for acylethanolamides.

Considering PPAR α receptor we found that it is mainly expressed in the human colonic epithelium, but not in immune cells of the lamina propria [14]. Interestingly, colonic mucosa

(epithelium and lamina propria) in active UC patients at disease onset showed a significant down-regulation of both PPAR α and PPAR γ mRNA expression in colonic mucosa of active UC patients. These data indicate that, not only PPAR γ , but also PPAR α , are implicated in the pathophysiology of the human colonic inflammation. We also detected an over-expression of iNOS mRNA, a pro-inflammatory mediator that produces nitric oxide species and leads oxidative stress and cell death [11,36,37]. This enzyme is under the active control of PPAR α receptor since PPAR α agonists enhance its degradation [38]. Immunohistochemical results demonstrated that PPAR α mRNA down-expression in the UC mucosa correlated with PPAR α protein down-expression in the UC epithelium. Moreover, only 5-ASA treatment increased immunohistochemical expression of PPAR α to control expression level, but not when UC patients were treated with 5-ASA in combination with glucocorticoids and/or immunomodulators.

5-ASA is structurally related to nonsteroidal anti-inflammatory drugs that shares molecular targets including inflammation, proliferation and/or apoptosis [39–41]. 5-ASA inhibits inflammation by scavenging free radicals and thus interfering with the arachidonic acid metabolism [42]. Recent studies indicated that the anti-inflammatory effect of 5-ASA is mediated by the activation of PPAR γ [9–11], a nuclear receptor whose agonists can suppress or delay inflammation effectively by inhibiting multiple steps in NF- κ B and AP-1 signaling pathways [7,8] and attenuating the production of nitric oxide (iNOS) and macrophage-derived cytokines such as TNF α , IL-1 and IL-6 in mouse models of colitis [16,43,44]. Moreover, Linard et al. [11] showed that 5-ASA is able to induce PPAR α , PPAR γ and RXR α co-expression and promote their translocation to the nucleus in an animal model of irradiation-induced intestinal inflammation. In the present study, we demonstrated that 5-ASA specifically increased the expression of PPAR α in the human UC epithelium suggesting that, not only PPAR γ , but also PPAR α can be a key receptor for the potent anti-inflammatory effect of 5-ASA in the human UC [9,10]. At this time, nothing at all is known about the regulation of PPAR α expression and much more studies are needed to elucidate the anti-inflammatory mechanisms of 5-ASA.

Others components of the PPAR α signaling system, such as NAPE-PLD, FAAH and NAAA, are expressed in the healthy colonic epithelium and immune cells of the colonic lamina propria in humans [35]. NAPE-PLD is one of several N-acylethanolamide-biosynthesis enzymes that catalyze the release of N-acylethanolamide (NAE) from N-acyl-phosphatidylethanolamine (NAPE), converting endogenous lipids into chemical signals like oleoylethanolamine (OEA), palmitoylethanolamine (PEA) and anandamide (AEA) [41,42,45,46]. Some studies showed that the biological activity of PEA, such as anti-inflammatory and analgesic activities [47], and OEA, such as food intake [48–50], are mediated by non-cannabinoid receptors among which PPAR α is probably the most important [51–53]. In mammalian tissues, three enzymes responsible for hydrolyses of NAEs to fatty acids and ethanolamine have been identified: FAAH-1, FAAH-2 (human isozyme) and NAAA [24,54–57]. Thus, it has been shown that selective FAAH or NAAA inhibitors produced an anti-inflammatory effect [26–29]. Interestingly, FAAH and NAAA have different catalytic properties and substrate specificity [24]. FAAH is catalytically active at neutral and alkaline pH and shows the highest reactivity with anandamide, followed by OEA and PEA [58]. In contrast, NAAA activity is optimum at pH 4.5–5, being inactive at alkaline pH, and hydrolyzes PEA much faster than others NAEs [24,55]. Therefore, alterations of FAAH and NAAA activity can be as a result of variations of luminal pH in colonic inflammation, and it is

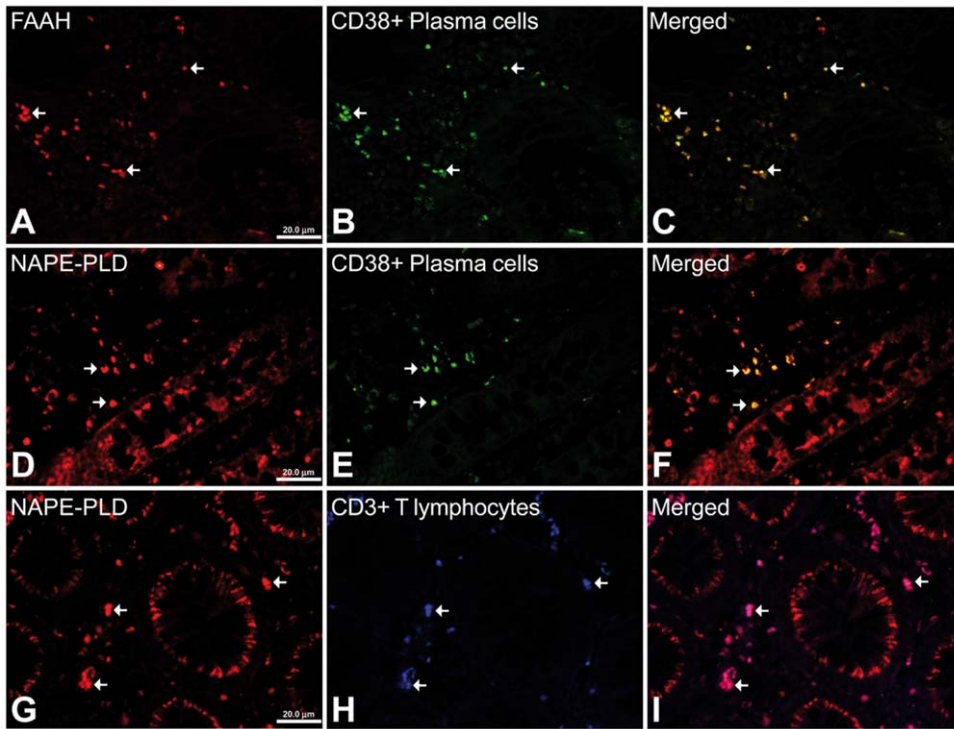


Figure 9. Representative high-magnification photomicrographs showing double immunofluorescence for FAAH, NAPE-PLD, CD38 and CD3 in order to characterize the immune cells in the mucosa infiltrate of UC patients. Nearly all FAAH immunofluorescent cells are plasma cell-specific CD38 (A–C). NAPE-PLD immunofluorescence was observed in both CD38+ plasma cells (D–F) and CD3+ T lymphocytes (G–I). doi:10.1371/journal.pone.0037729.g009

conceivable that reduced intracolonic pH in active UC impairs the anti-inflammatory effects of PPAR endogenous agonists [59].

In the present study, we demonstrated that mRNA and protein expression of NAPE-PLD, FAAH and NAAA was partially altered in active colitis, and immunohistochemical expression of these enzymes was partially restored after treatment (quiescent colitis) in a tissue-dependent manner (epithelium and immune cells of the lamina propria). Overall, the present data suggested that both increase of NAAA expression and lack of change in FAAH expression in the UC epithelium agree with a substantial reduction of luminal pH in the colon of UC patients [59]. Therefore, NAPE-PLD down-expression and NAAA over-expression in UC epithelium might let to a net reduction in NAEs turnover (specifically

PEA) in the epithelium, leading the attenuation of the anti-inflammatory response via the activation of PPAR receptors (Fig. 10). Interestingly, inflammation associated with osteoarthritis and rheumatoid arthritis showed a lower concentration of PEA in the synovial fluid compared to non-inflamed normal volunteers [60]. Changes observed in NAPE-PLD, FAAH and NAAA mRNA expression in the mucosa (epithelium and lamina propria) correlated completely with changes observed in the number of immunoreactive cells in the lamina propria of UC patients, but not with their immunohistochemical expression in the UC epithelium (see Table 5). These discrepancies can be explained by a higher expression of these enzymes in the immune cells during UC

Table 5. Summary of the changes detected in PPAR α signaling system (PPAR α , NAPE-PLD, FAAH and NAAA) in the colonic epithelium and lamina propria of active UC patients and after treatment (quiescent UC patients)¹.

	Gene expression	Immunohistochemical expression in epithelium		Number of immunoreactive cells in lamina propria	
	Active untreated-UC	Active untreated-UC	Quiescent treated-UC	Active untreated-UC	Quiescent treated-UC
PPAR α	↓ (*)	↓ (**)	↑ (*) (5-ASA)	nc	nc
NAPE-PLD	nc	↓ (***)	↑ (***) (5-ASA+ glucocortic.)	nc	nc
FAAH	↑ (*)	nc	nc	↑ (***)	↓ (***)
NAAA	↓ (*)	↑ (**)	nc	↓ (***)	↑ (**)

¹Symbols are as follows: increased expression (↑), decreased expression (↓), no change (nc). Statistical significance was represented by.

*P<0.05,

**P<0.01 and

***P<0.001.

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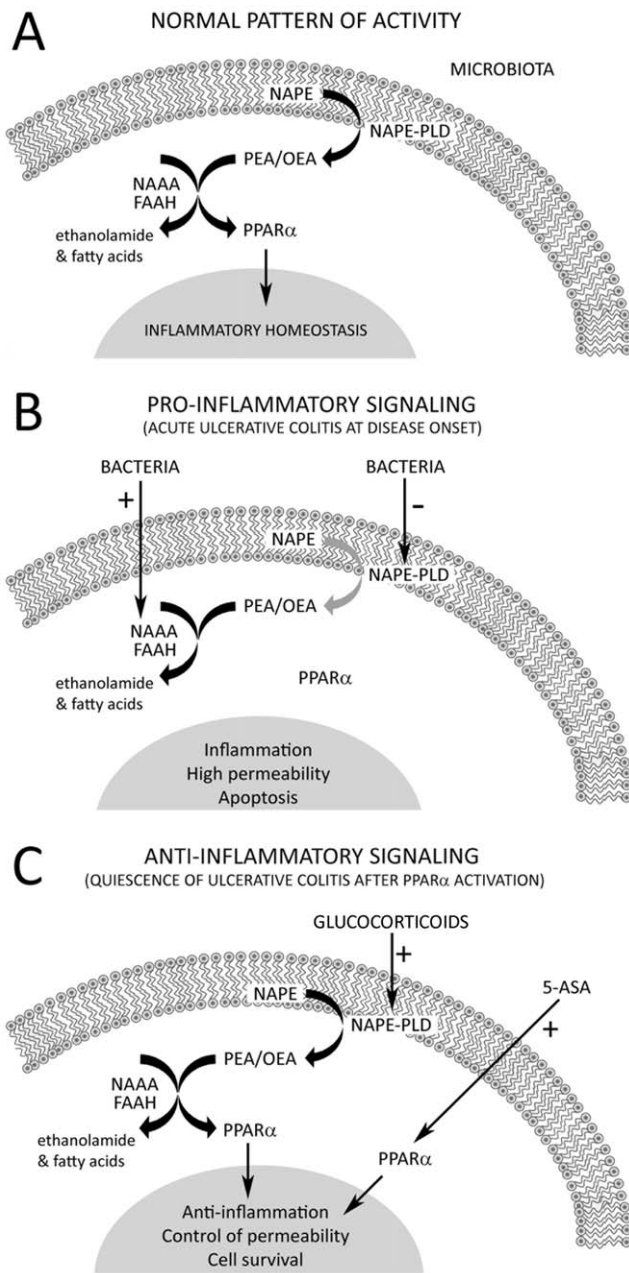


Figure 10. Schematic drawings that hypothesize the normal pattern of NAE-PPAR α activity (A), and the pro- and anti-inflammatory NAE-PPAR α signaling that may occur in the colonic epithelial cells of active UC at disease onset (B) and after a putative treatment with 5-ASA and/or glucocorticoids (C).
doi:10.1371/journal.pone.0037729.g010

infiltration, but also the different roles of the NAE-PPAR α signaling system in colonic epithelium and lamina propria.

In the lamina propria of healthy human colon, we found that the number of NAAA-ir immune cells was 50-fold higher than the number of FAAH-ir immune cells. These data can suggest a higher rate of PEA hydrolysis in comparison with AEA hydrolysis. In the lamina propria of active UC, we found that the number of FAAH-ir immune cells increased up to 10-fold, whereas the number of NAAA-ir immune cells decreased up to 2.75-fold, suggesting a concomitant increase of AEA hydrolysis as well as

decrease of PEA hydrolysis. These results can be related with the fact that AEA activates cannabinoid (CB1 and CB2) receptors, whereas PEA is inactive on these receptors, but activates PPAR α [51], playing different roles in inflammatory activation. Previous biochemical and immunochemical analysis demonstrated NAAA expression in macrophage cells of the rat lung and brain [25]. Here, we showed that NAAA is predominantly expressed in macrophages and B and T lymphocytes in the lamina propria of UC patients. Most FAAH-ir cells in the lamina propria of UC patients expressed CD38, a surface glycoprotein found in plasma B and natural killer cells, and this result agrees with previous studies showing FAAH activity in lymphocytes [61]. NAPE-PLD-ir cells in the lamina propria of UC patients were CD38+ plasma cells and CD3+ T lymphocytes, but not CD14+ macrophages, contrary to expectation after pro-inflammatory stimuli [62].

UC-specific treatments produced tissue-dependent impairments in the expression of PPAR α signaling system. NAPE-PLD, but not NAAA or FAAH, responded to treatment in the epithelium, while NAAA and FAAH, but not NAPE-PLD, responded to treatment in the immune cells of the lamina propria of UC patients. 5-ASA produced an increase of NAPE-PLD immunohistochemical expression (similar to control levels) in the quiescent UC epithelium, which was enhanced after corticosteroid treatment. Interestingly, the analysis of the NAPE-PLD/NAAA and NAPE-PLD/FAAH ratios suggested an increase of NAEs production in the UC epithelium after 5-ASA/corticosteroid treatment, but not when patients were treated exclusively with 5-ASA. It is clear that 5-ASA treatment leads to an increase of NAPE-PLD and PPAR α expression, so probably both 5-ASA and the concomitant over-production of NAEs via glucocorticoids can enhance an anti-inflammatory response in the epithelium of UC patients by the activation of PPAR α (Fig. 10). This hypothesis agrees with previous data indicating that glucocorticoids generate anti-inflammatory regulatory responses by promoting arachidonic acid-containing lipid biosynthesis [63]. Treatment also increases the number of NAAA-ir immune cells, reaching control levels and, probably, normalizing PEA hydrolysis. However, the significant decrease of FAAH-ir immune cells after treatment did not reach control levels, so there may be still an over-degradation of AEA in the lamina propria of UC patients.

We must pay attention on two limitations related with the cohort of patients used in the present study. As a result of prioritizing clinical, endoscopic and histopathological considerations to obtain a homogeneous cohort, control and UC groups were not-age matched. Additionally, smoker patients and patients and controls from both genders were included in the study. However, these factors cannot be included in additional analysis because of the size of the cohort, designed to be a within-subject design (patients were they own control for quiescence status).

In conclusion, our results indicated that PPAR α , NAPE-PLD, FAAH and NAAA form part of a key lipid signaling system that regulates UC-activated inflammatory response in human. 5-ASA, through PPAR α receptor, and glucocorticoids, through acylethanolamide producing/degrading enzymes, reduces colitis-associated inflammation suggesting PPAR α agonists or FAAH/NAAA inhibitors as potential drugs for the treatment of inflammatory bowel diseases in human.

Supporting Information

Figure S1 Housekeeping gene expressions of SP1 transcription factor (A) and β ACTIN (B) represented by the threshold cycles (C(t)). We cannot detect differences in gene expression between control and active UC patients. Student *t*-test

(N = 7–8): SP1, $F = 0.291$, $P = 0.734$; β ACTIN, $F = 0.388$, $P = 0.597$.)
(TIF)

Figure S2 Densitometrical quantification of PPAR α (A), NAPE-PLD (B), FAAH (C) and NAAA (D) immunoreactivity in human healthy (control) and active UC colonic epithelium depending on severity (mild, moderate and severe). Mann-Whitney U and Wilcoxon tests (N = 22–24): * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$ versus control group.
(TIF)

Figure S3 Densitometrical quantification of PPAR α (A), NAPE-PLD (B), FAAH (C) and NAAA (D) immunoreactivity in active UC colonic epithelium depending on gender. No statistical difference was observed. Student t -test (N = 22–24).
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Figure S4 Densitometrical quantification of PPAR α (A), NAPE-PLD (B), FAAH (C) and NAAA (D) immunoreactivity in active UC colonic epithelium depending on smoking habits. No statistical difference was observed. Student t -test (N = 22–24).
(TIF)

Methods S1 mRNA isolation and quantitative RT-PCR analysis.
(DOC)

Author Contributions

Conceived and designed the experiments: JS MA FRdF. Performed the experiments: JS YRZ LM. Analyzed the data: JS YRZ MI PR FJBS. Contributed reagents/materials/analysis tools: FJBS MA FRdF. Wrote the paper: JS FRdF.

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