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S1P₃ Receptor Mediates the Proinflammatory Effect of the Endocannabinoid 2-Arachidonoylglycerol in Endometriotic Epithelial Cells

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ABSTRACT

Endometriosis is a chronic inflammatory disease characterized by the ectopic implantation of endometrium outside the uterus associated with pelvic pain and infertility. The molecular mechanisms involved in the pathogenesis of endometriosis are complex and far from being fully elucidated. We recently showed that the signaling of the bioactive sphingolipid sphingosine 1-phosphate (S1P) is deeply dysregulated in endometriosis. The endocannabinoids anandamide (AEA) and 2-arachidonoylglycerol (2-AG), via ligation to G-protein coupled receptors, CB1, CB2, and GPR18 as well as the cation channel TRPV1, play a crucial role in the modulation of pain and inflammation. Here, the role of endocannabinoid signaling in endometriosis and its possible cross talk with the S1P signaling axis has been investigated. It has been found that CB1, CB2, GPR18, TRPV1 as well as the enzymes involved in endocannabinoid metabolism are expressed in endometriotic lesions. Furthermore, the effect of 2-AG and AEA in the modulation of inflammation has been established in human endometriotic epithelial cells. 2-AG, but not methanandamide (MAEA), the nonhydrolyzable AEA analogue, induced a marked increase in the expression of cyclooxygenase 2 and various pro-inflammatory interleukins (IL-1 β , IL-6 and IL-8). Interestingly, S1P₃, whose expression is augmented by 2-AG, is crucial for transducing the biological action of the endocannabinoid. Indeed, S1P₃ pharmacological blockade or its specific silencing impaired the pro-inflammatory action of 2-AG. In conclusion, these findings demonstrate, for the first time, the occurrence of a functional interplay between endocannabinoids and S1P signaling in endometriosis, paving the way for novel pharmacological strategies to treat the disease.

1 | Introduction

The endocannabinoid system (ECS) has emerged as an important regulator of pain, neuromodulation, and inflammation [1,2].

The ECS is composed of the endocannabinoids anandamide (AEA) and 2-arachidonoylglycerol (2-AG), the cannabinoid G-protein-coupled receptors (CB1, CB2, GPR18, and GPR55) and the transient receptor potential cation channel subfamily

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V member 1 (TRPV1) [3]. In addition, the enzymes responsible for endocannabinoid metabolism such as the biosynthetic enzymes diacylglycerol-lipase (DAGL) and N-acylphosphatidyl ethanolamines-phospholipase D (NAPE-PLD), as well as the catabolic enzymes fatty acid amide hydrolase (FAAH) and monoacylglycerol lipase (MAGL) are components of ECS. Cannabinoid receptors are expressed in several tissues including the reproductive organs [4, 5]. Interestingly, AEA and 2-AG are elevated in the plasma of endometriosis-affected women and increased peritoneal fluid levels of 2-AG positively correlated with prostaglandin E2 (PGE2), suggesting a possible link between ECS and inflammatory pain [6]. Endometriosis is a chronic benign gynecological endocrine disease characterized by fatigue, heavy menstrual flow, and infertility. The disease, affecting ~10% of women of reproductive age, is defined by the presence of endometrium-like tissue outside the uterus [7]. Recently, a key role of the bioactive sphingolipid sphingosine 1-phosphate (S1P) in endometriosis pathogenesis has been solidly demonstrated [8–11]. Notably, the concentration of S1P is significantly increased in the peritoneal fluid of women with endometriosis [12], and a deep dysregulation of S1P metabolism and signaling has been reported in endometriotic lesions [13, 14].

S1P is involved in the regulation of fundamental cellular and tissue responses such as inflammation, cell survival, migration, and tumorigenesis [15]. S1P synthesis is catalyzed by sphingosine kinases (SK1 and SK2) that phosphorylate sphingosine. The multifaceted effects of S1P mainly rely on the so-called “inside-out” mechanism of action, that implies its binding to five specific G-protein-coupled receptors, S1P_{1–5}, after the export into the extracellular microenvironment mediated by transporters such as Spinster homolog 2 (Spns2) [16, 17].

Current endometriosis treatments comprise surgical and hormonal therapies [18], each linked respectively to high rates of recurrence and side effects in long-term administration [19, 20]. Thus, the comprehension of the molecular mechanisms implicated in endometriosis pathogenesis might lead to the identification of innovative nonhormonal pharmacological targets for the disease. Although not extensively investigated, literature data support the existence of a functional cross talk between ECS and S1P signaling [21]. For instance, S1P treatment has been shown to modulate CB2 and TRPV1 levels in C2C12 myoblasts [22] and the activation of SK mediates the hypotensive response to AEA in anesthetized mice [23].

The data presented here show for the first time that 2-AG—but not methanandamide (MAEA), a non-hydrolyzable AEA analogue—induces a pro-inflammatory response in endometriotic cells. Our findings also reveal that the S1P-specific receptor isoform S1P₃, whose expression is augmented by 2-AG, is critically involved in the pro-inflammatory effect of the endocannabinoid. This work provides the first experimental evidence of a functional interplay between endocannabinoids and S1P signaling in endometriosis, pointing at new pharmacological targets for disease treatment.

2 | Materials and Methods

2.1 | Materials

All biochemicals, TRI reagent, cell culture reagents, Nutrient Mixture F-12 Ham (F12), Dulbecco's Modified Eagle Medium (DMEM), fetal bovine serum (FBS), L-glutamine, Penicillin/Streptomycin, phosphatase inhibitor cocktail, protease inhibitor cocktail, 2-AG, bovine serum albumin (BSA), VPC23019, TY-52156, PF-543, the specific siRNA for SK1, SK2, S1P₁, S1P₂, S1P₃ and the scramble siRNA were purchased from Merck Life Sciences (Burlington, MA, USA). C17 sphingosine and C17-S1P were obtained from Avanti Polar Lipids (Alabaster, AL, USA).

Human endometriotic epithelial 12Z cell line (Cat. No. T0764) and the Applied Cell Extracellular Matrix were purchased from Applied Biological Materials Inc. (Richmond, BC, Canada). Bradford protein assay, Tris/Glycine/SDS, the EveryBlot Blocking Buffer, the Clarity western ECL substrate and the trans-blot turbo PVDF membrane were obtained from Bio-Rad (Hercules, CA, USA). Anti-S1P₃ antibody was obtained from Abcam Ltd. (Cambridge, UK). Anti-SK1, anti-SK2, anti-phospho-SK2 (Thr578), and anti-phospho-SK1 (Ser225) antibodies were purchased from ECM Biosciences LLC (Versailles, KY, USA). Anti-Spns2 antibody was purchased from FabGennix International Inc. Anti-CB1, anti-CB2, and Methanandamide (MAEA) were purchased from Cayman Chemical (Ann Arbor, Michigan, USA) and anti-GPR18 from Invitrogen, Thermo Fisher Scientific (Waltham, MA, USA). Anti-TRPV1 was obtained from OriGENE Technologies Inc. (Rockville, MD, USA). RayBio C-Series Human Inflammation Array C1 was from RayBiotech (Peachtree Corners, Georgia, USA). Anti-GAPDH antibodies and secondary antibodies conjugated to horseradish peroxidase, were obtained from Santa Cruz Biotechnology (Santa Cruz, CA). AllPrep DNA/RNA FFPE kit was purchased from Qiagen (Hilden, Germany). TaqMan Universal Master Mix II, TaqMan gene expression assays, Lipofectamine RNAiMAX, High-capacity cDNA reverse transcription kit and the UltraVision LP Detection System HRP Polymer and DAB Plus Chromogen were obtained from Thermo Fisher Scientific Inc. (Waltham, MA, USA).

2.2 | Sample Collection

The gene expression study ($n=20$) and the immunohistochemistry (IHC) analysis ($n=15$) were conducted on endometriotic lesions obtained from an independent cohort of patients. Control endometrial samples ($n=15$) used for gene expression analysis were acquired using diagnostic hysteroscopy procedures. Clinical and imaging investigations were performed to exclude a diagnosis of endometriosis or other uterine disorders in controls. All the samples were histologically characterized. The diagnostic hysteroscopy or surgical procedure for lesion removal was performed during the proliferative phase and all hormonal treatments had been stopped at least 2–3 months before surgery. The endometrial cycle phase was confirmed by histologic analysis of endometrial biopsies. There were no differences in age, pregnancy and parity between the study and control groups. The study protocol was approved by the Institutional Review Board (number 13742) and all patients provided written informed consent.

2.3 | Cell Culture and Treatment

Human endometriotic epithelial 12Z cells were cultured in dishes coated with the extracellular matrix using a 1:1 mixture of DMEM:F12, supplemented with 10% FBS, 2 mM L-glutamine, 100 U/mL penicillin/streptomycin at 37°C in 5% CO₂. For the experiments, cells were serum-starved overnight in a medium supplemented with 1 mg/mL fatty acid-free BSA. When requested, cells were pretreated with pharmacological antagonists or an equivalent volume of vehicle (DMSO) for 45 min prior to treatment with 2-AG or the corresponding vehicle (ethanol).

2.4 | Cell Transfection

Endometriotic epithelial 12Z cells were transfected with siRNA duplexes using Lipofectamine RNAiMAX, according to the manufacturer's instructions [24, 25]. siRNAs diluted in the mixture of DMEM:F12 were incubated with Lipofectamine RNAiMAX at room temperature (RT) for 20 min and then added to cells to a final concentration of 50 nM, in DMEM:F12 supplemented with 10% FBS. After 30 h cells were serum-starved overnight and stimulated with 10 μM 2-AG.

2.5 | Quantitative Real Time PCR (qPCR)

Total RNA was extracted from Formalin-Fixed, Paraffin-Embedded (FFPE) tissue samples using the AllPrep DNA/RNA FFPE kit and from endometriotic epithelial 12Z cells employing TRI-reagent following the manufacturer instructions, and reverse-transcribed using the high-capacity cDNA reverse transcription kit, as directed by the manufacturer. TaqMan gene expression assays were used to quantify target gene mRNAs in triplicate on a CFX96 Touch qPCR Detection System (Bio-Rad, Hercules, CA, USA). The target sequences were amplified alongside the reference gene β-actin [14]. The 2^{-ΔCt} and 2^{-ΔΔCt} methods were used to calculate the relative expression of mRNA [26, 27].

2.6 | Immunohistochemistry

Tissue samples were fixed with formalin, paraffin embedded and 3 μm sections were suitably sliced from each block and then stained for IHC analysis with specific antibodies against CB1 (1:50), CB2 (1:50), GPR18 (1:50), and TRPV1 (1:50) as previously described [28]. The procedure started with deparaffinization and rehydration. The slides were heated in a microwave at 95°C–97°C in a citrate buffer (pH 6.0) for a total of 20 min and cooled to RT for antigen retrieval. Then the slides were incubated with the primary antibodies at 4°C overnight. To detect the antigen–antibody complexes the “UltraVision LP Detection System HRP Polymer and DAB Plus Chromogen” kit was used according to the manufacturer's instructions: tissue slices were covered with the “Primary Antibody Enhancer” for 10 min at RT followed by 15 min with the secondary antibody conjugated to horseradish peroxidase. The bound antibody complexes were stained for 3 min 40 s with diaminobenzidine and then mounted. The slides were analyzed at the Histopathology and

Molecular Diagnostics of Careggi University Hospital, Florence. Images were obtained by Optikam PRO6 Digital Camera (C-P6), OPTIKA microscope Italy, B-383PLi.

2.7 | Western Blot (WB) Analysis

Cells were collected in 50 mM Tris, pH 7.5, 120 mM NaCl, 6 mM EGTA, 1 mM EDTA, 20 mM NaF, 15 mM Na₄P₂O₇, 1% Nonidet, with the addition of protease and phosphatase inhibitor cocktail and then incubated at 4°C for 30 min. Cells were centrifuged for 10000 g, 15 min at 4°C, and the supernatant containing protein was collected. Proteins were used to perform SDS/PAGE and WB. PDVF membranes were incubated at 4°C overnight with the primary antibodies and then with specific secondary antibodies at RT for 1 h. The binding of the antibodies with the specific proteins was detected by chemiluminescence employing Amersham Imager 600 (GE Healthcare, Buckinghamshire, UK).

2.8 | Human Inflammation Array

Pro-inflammatory cytokine secretion into the media was quantified by RayBio C-Series Human Inflammation Array C1. Endometriotic epithelial cells were cultured and treated with 2-AG or MAEA for 24 h before conditioned media were collected according to the manufacturer's instructions. After the incubation of the media with the antibody arrays' membranes, chemiluminescence signals were obtained employing Amersham Imager 600. Signal intensities were used to quantify inflammatory factors' secretion using ImageJ software.

2.9 | Sphingosine Kinase in Cell Assay

Endometriotic epithelial cells were plated and upon reaching 70% confluence, were serum-starved overnight. Cells were incubated with the exogenous substrate C17 sphingosine (5 μM) and contemporaneously stimulated or not with 10 μM 2-AG for 30 min. Cells were then harvested in methanol and subsequently added to internal standard for C17-S1P (10 pmol d17 S1P in methanol). Samples were vortexed, precipitated overnight at –80°C, followed by a 5 min centrifuge at 21 300 g, 4°C and then the supernatant was analyzed by liquid chromatography–tandem mass spectrometry (LC–MS/MS).

2.10 | Statistical Analysis

Statistical analysis was performed with the use of Student's *t*-test, Mann–Whitney test, one-way or two-way ANOVA followed by Bonferroni post hoc test. Graphical representations were generated using GraphPad Prism 10.0 (GraphPad Software) (San Diego, CA, USA).

3 | Results

We initially determined the expression of CB1, CB2, GPR18, and TRPV1 in endometriotic lesions of different localizations. The real-time PCR analysis reported in Figure 1A shows that all the

investigated receptors are expressed at mRNA level in the analyzed lesions. Moreover, the expression levels of each receptor were comparable to those observed in endometrial samples from healthy women (Figure 1A). The expression of CB1, CB2, GPR18, and TRPV1 was confirmed at the protein level, as demonstrated by the representative IHC images shown in Figure 1B: the staining of the analyzed receptors is clearly detectable both in the stroma and in the glands. For all the analyzed receptors, strong immunoreactivity was observed in the glandular epithelium (black arrows), while the stromal cells showed lower signal (red arrows). In addition, the expression of the enzymes involved in 2-AG and AEA synthesis, NAPE-PLD and DAGL respectively, as well as in endocannabinoid catabolism, FAAH and MAGL that degrade 2-AG and AEA, respectively, was demonstrated in endometriotic lesions (Figure 1C). In particular, mRNA levels of NAPE-PLD and MAGL were found to be significantly increased in endometriotic lesions compared to healthy endometrial tissue (Figure 1C).

In order to characterize the biological action of endocannabinoids in endometriosis, epithelial endometriotic 12Z cells, were employed. We first investigated whether the cells express the endocannabinoid metabolic enzymes and the receptors. qPCR

analysis showed that both the biosynthetic (DAGL and NAPE-PLD) and the catabolic (FAAH and MAGL) enzymes were expressed in endometriotic epithelial cells (Figure S1A). Moreover, the expression of the receptors CB1, CB2, GPR18 and TRPV1 was observed at the mRNA level (Figure S1A) and further confirmed at the protein level by WB analysis (Figure S1B).

Then, we evaluated whether endocannabinoids are able to regulate the inflammatory response in endometriotic epithelial cells. To this aim, the mRNA levels of cyclooxygenase 2 (COX2), interleukin-1 β (IL-1 β), IL-6 and IL-8 were examined in the cells treated for 24 h with increasing concentrations (2.5 μ M, 5 μ M, 10 μ M) of 2-AG and MAEA, the nonhydrolyzable anandamide analogue. By real-time PCR analysis, it was found that 2-AG potently increased the expression of COX2, IL-1 β , IL-6 and IL-8 with a maximal effect at 10 μ M (Figure 2A). On the contrary, the treatment with MAEA did not significantly affect the expression of the analyzed inflammatory factors at any of the tested concentrations (Figure 2A).

In order to confirm these data, we then examined whether the treatment with the endocannabinoids modulates the

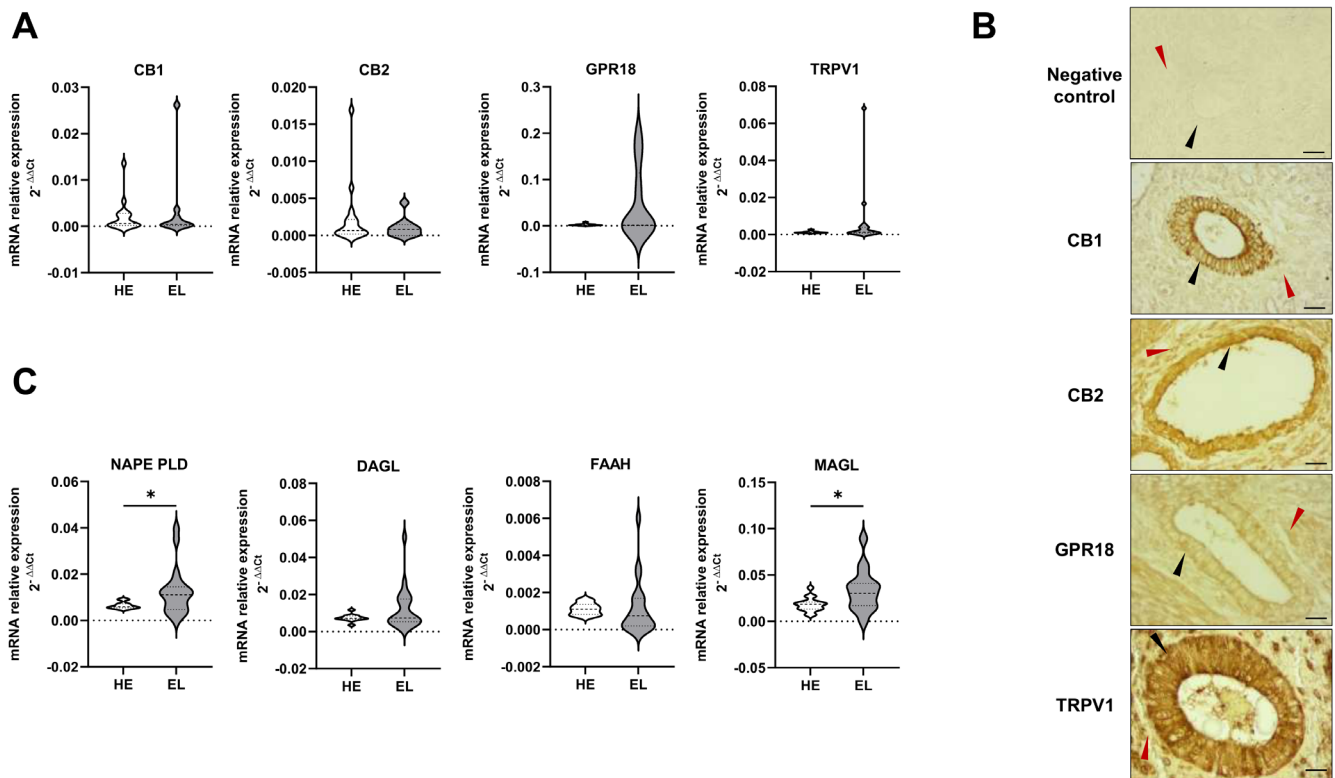


FIGURE 1 | Cannabinoid receptors and enzymes are expressed in human endometriotic lesions. (A) qPCR analysis was performed using TaqMan Gene Expression Assay probes specific for cannabinoid receptors CB1, CB2, GPR18, and TRPV1 in the endometrium of healthy women (HE) ($n = 15$) or endometriotic lesions (EL) of different localization ($n = 20$: 7 ovarian endometriosis and 13 deep infiltrating endometriosis). Results were analyzed employing the $2^{-\Delta\Delta C_t}$ method. (B) Representative immunohistochemical images of CB1, CB2, GPR18 and TRPV1 expression in endometriotic lesions ($n = 15$: 11 ovarian endometriosis and 4 deep infiltrating endometriosis). The staining with DAB produced a brown precipitate at the site of antibody binding, localized both in epithelial (black arrows) and stromal (red arrows) cells of the endometriotic lesions. The negative control was obtained by processing tissue sections in parallel with the same IHC protocol, omitting the primary antibody. Magnification: $\times 40$, scale bar: 30 μ m. (C) qPCR analysis was performed using TaqMan Gene Expression Assay probes specific for cannabinoid enzymes NAPE-PLD, DAGL, FAAH, and MAGL in the endometrium of healthy women (HE) ($n = 15$) or endometriotic lesions (EL) of different localization ($n = 20$: 7 ovarian endometriosis and 13 deep infiltrating endometriosis). Results were analyzed employing the $2^{-\Delta\Delta C_t}$ method. NAPE-PLD and MAGL mRNA levels are significantly increased in EL compared to HE (Student's t -test * $p < 0.05$).

extracellular release of functional pro-inflammatory factors. To this aim, the conditioned media obtained from endometriotic cells treated for 24h with 2-AG or MAEA were analyzed employing an antibody array that allows the simultaneous detection of multiple cytokines. In agreement with the results obtained by real-time PCR, 2-AG significantly increased the extracellular release of various pro-inflammatory cytokines, namely IL-1 β , IL-2, IL-6, IL-7 and IFN γ (Figure 2B). In contrast, MAEA treatment did not significantly affect the extracellular release of these inflammatory factors in 12Z cells (Figure 2B).

Given the key role of S1P in modulating the inflammatory process in endometrial cells [29], we next examined whether cell treatment with 10 μ M 2-AG for 24h regulated the expression of molecules involved in S1P metabolism and signaling. Data shown in Figure 3A show that 2-AG significantly augmented the mRNA levels of SK1, of the receptor isoform S1P $_3$ and of the specific transporter Spns2, suggesting the occurrence of an interplay between 2-AG and S1P signaling in endometriotic cells. In accordance, WB analysis performed in cells treated with 10 μ M 2-AG for 24h showed that S1P $_3$ protein levels were significantly increased by the endocannabinoid (Figure 3B). On the contrary, the 2-AG-induced increase in mRNA levels was not mirrored by a parallel increase in the protein content of SK1 and Spns2, at least at the examined time point (Figure 3B). Of note, our group previously reported a significant upregulation of S1P $_3$ both at the mRNA and protein levels in endometriotic

lesions compared to endometrium from healthy controls [10, 14]. Interestingly, 2-AG was responsible for a significant increase in CB1 mRNA levels while the expression levels of the other endocannabinoid receptors were not affected (Figure S2).

Next, the potential role of 2-AG-dependent upregulation of S1P $_3$ in the pro-inflammatory action of the endocannabinoid was examined. For this purpose, the expression of the pro-inflammatory factors was evaluated in endometriotic cells challenged with 2-AG in the presence or absence of TY-52156 (10 μ M), S1P $_3$ specific antagonist, or VPC23019 (10 μ M), pharmacological antagonist of S1P $_1$ /S1P $_3$. The obtained results show that the enhanced expression of COX2, IL-1 β , IL-6 and IL-8 elicited by 2-AG was significantly reduced by preincubation with TY-52156 (Figure 4A) or VPC23019 (Figure S3), suggesting a crucial role for S1P $_3$ in transmitting the pro-inflammatory action of 2-AG.

In order to confirm these findings, the RNA interference approach was employed to efficaciously reduce the expression levels of S1P $_1$, S1P $_2$ or S1P $_3$ (Figure S4A). Notably, the selective knockdown of S1P $_3$ impaired the 2-AG-induced increase of COX2, IL-1 β and IL-6 transcription while the downregulation of the other two receptor isoforms did not alter 2-AG action (Figure 4B). Interestingly, the selective knockdown of S1P $_1$ and S1P $_3$ significantly reduced the increase of IL-8 levels elicited by 2-AG, whereas the silencing of S1P $_2$ enhanced the

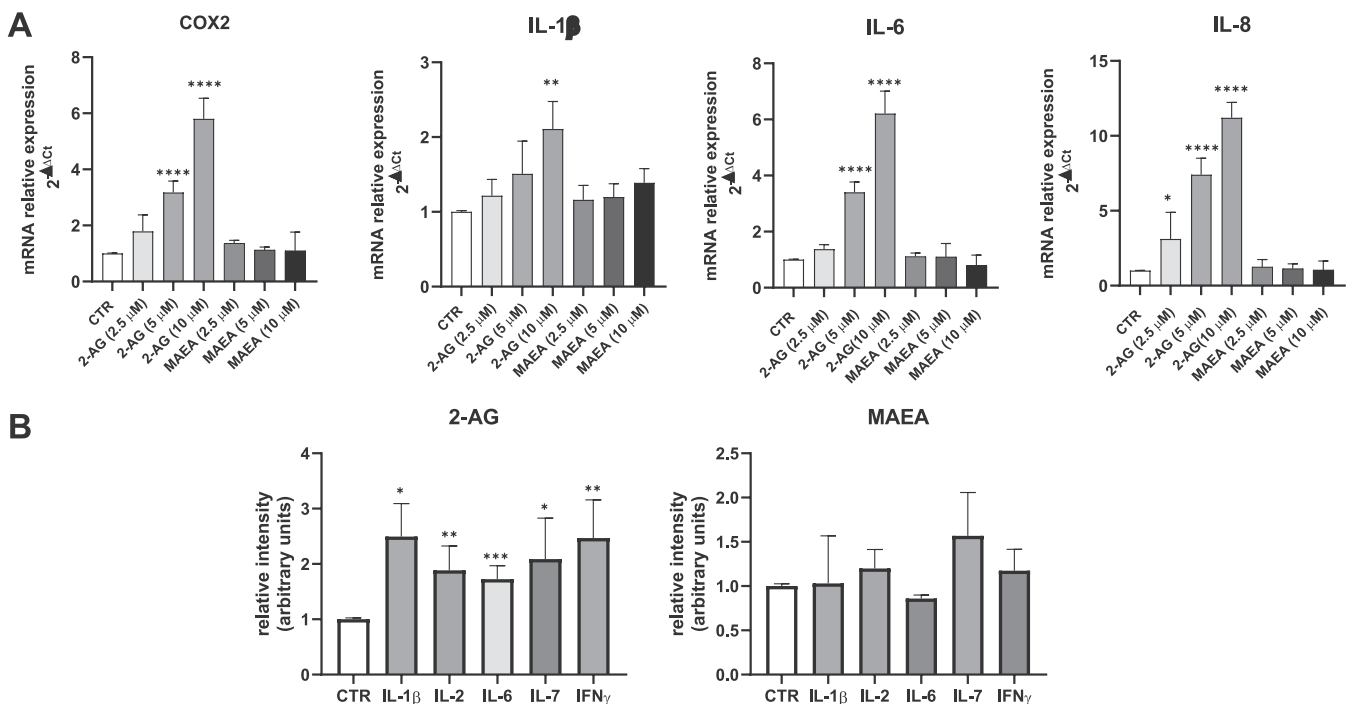


FIGURE 2 | 2-AG, but not MAEA, increases the expression and release of different pro-inflammatory factors in endometriotic epithelial cells. (A) Endometriotic epithelial cells were serum-starved for 18h and treated with increasing concentrations of 2-AG and MAEA for 24h. mRNA quantitative analysis of COX2, IL-1 β , IL-6 and IL-8 was performed by qPCR. Results, analyzed with the $2^{-\Delta\Delta C_t}$ method, were obtained using β -Actin as a housekeeping gene and individual inflammatory factors of the unchallenged specimen as a reference gene. 2-AG increases COX2, IL-1 β , IL-6 and IL-8 expression in a statistically significant manner (One-way ANOVA, * $p < 0.05$; ** $p < 0.01$; **** $p < 0.0001$). (B) Endometriotic epithelial cells were serum-starved for 18h and treated with 10 μ M 2-AG or 2.5 μ M MAEA for 24h. The obtained conditioned media were screened for the content of proinflammatory markers using the Human Inflammation Array as described in Section 2. Results were expressed as fold increase in respect to control. 2-AG induces the extracellular release of IL-1 β , IL-2, IL-6, IL-7 and IFN γ in a statistically significant manner (Student's t -test * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$).

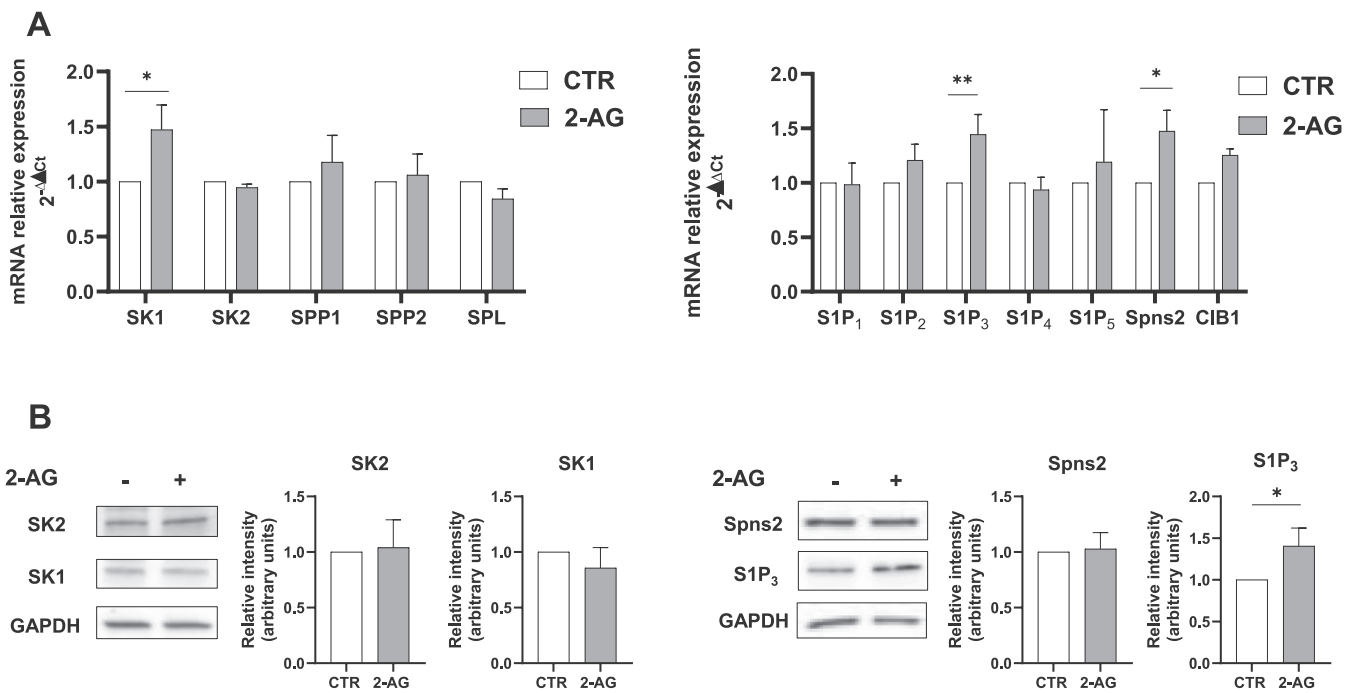


FIGURE 3 | 2-AG modulates S1P signaling axis in endometriotic epithelial cells. Endometriotic epithelial cells were serum-starved for 18 h and treated with 10 μ M 2-AG for 24 h. (A) mRNA quantitative analysis of S1P metabolic enzymes (SK1, SK2, SPP1, SPP2 and SPL) as well as molecules implicated in S1P signaling (S1P₁₋₅, the specific transporter Spns2 and the SK1-activating protein CIB1) was performed by qPCR. Results, analyzed with the 2^{-ΔΔCt} method, were obtained using β -Actin as a housekeeping gene and individual targets of the unchallenged specimen as a reference gene. 2-AG increases SK1, S1P₃, and Spns2 mRNA levels in a statistically significant manner (Student's *t*-test **p* < 0.05; ***p* < 0.01). (B) Protein lysates were analyzed using SDS-PAGE electrophoresis and WB, using specific anti-SK1, anti-SK2, anti-S1P₃, anti-Spns2 and anti-GAPDH antibodies. The histograms represent the densitometric analysis of 4 independent experiments. Data are the mean \pm SD and are reported as band intensity normalized to the expression of GAPDH, fold change over control (set as 1). 2-AG increases S1P₃ protein content in a statistically significant manner (Student's *t*-test **p* < 0.05).

endocannabinoid effect, suggesting positive and negative roles of these receptor isoforms in the specific modulation of IL-8 expression (Figure 4B).

To gain further insight into the mechanism by which 2-AG exploits S1P signaling to exert its pro-inflammatory effect, we extensively examined the possible involvement of SKs, the enzymes responsible for S1P synthesis. Indeed, on the basis of the S1P “inside-out” mechanism of action, the sphingolipid generated following SK activation acts as a ligand of its receptors after being exported outside the cells [16, 17]. Since 2-AG treatment was unable to enhance SK protein content (Figure 3B), we examined whether the short-term regulation of SK1 and SK2 via their phosphorylation [30, 31] was implicated in the endocannabinoid action. Data obtained by WB analysis using specific anti-phospho-SK1 or anti-phospho-SK2 antibodies showed that cell challenge with 10 μ M 2-AG for different time intervals (5 to 30 min) did not affect SK1 or SK2 phosphorylation, ruling out SK1/SK2 activation by the endocannabinoid (Figure 5A). In agreement, the cell assay of sphingosine kinase activity further excluded the activation of SK after 2-AG treatment (Figure 5B). Indeed, the intracellular levels of S1P quantified by LC-MS/MS were not affected by the treatment with 2-AG (Figure 5B). Moreover, in order to further verify the involvement of SK in mediating the 2-AG effect, SK1 and SK2 were efficaciously knocked down by employing siRNA technology (Figure S4B). As depicted in Figure 5C, the 2-AG-induced expression of COX2, IL-1 β , IL-6

and IL-8 was unaffected by the downregulation of SK1 or SK2, thus demonstrating that the pro-inflammatory action of 2-AG did not depend on either SK1 or SK2 in endometriotic epithelial cells.

In conclusion, our findings support the view that the specific S1P₃ isoform transduces the pro-inflammatory action of 2-AG in endometriotic epithelial cells and establishes the rationale for the exploitation of S1P₃ targeting as an innovative nonhormonal approach to counteract endometriosis.

4 | Discussion

Endometriosis is a chronic disease with life-impacting symptoms and high prevalence in women of reproductive age [32]. Inflammation is a cardinal feature of endometriosis, strongly linked with its pathogenesis and the development of symptoms [33].

The cannabinoid receptors have long been known to play a role in inflammatory regulation through both pro-inflammatory and anti-inflammatory mechanisms, depending on the cellular environment and pathological state [34]. Here, CB1, CB2, GPR18 and TRPV1 were found to be expressed in healthy endometrium and endometriotic lesions. The expression of CB1, CB2 and TRPV1 has been previously reported in the

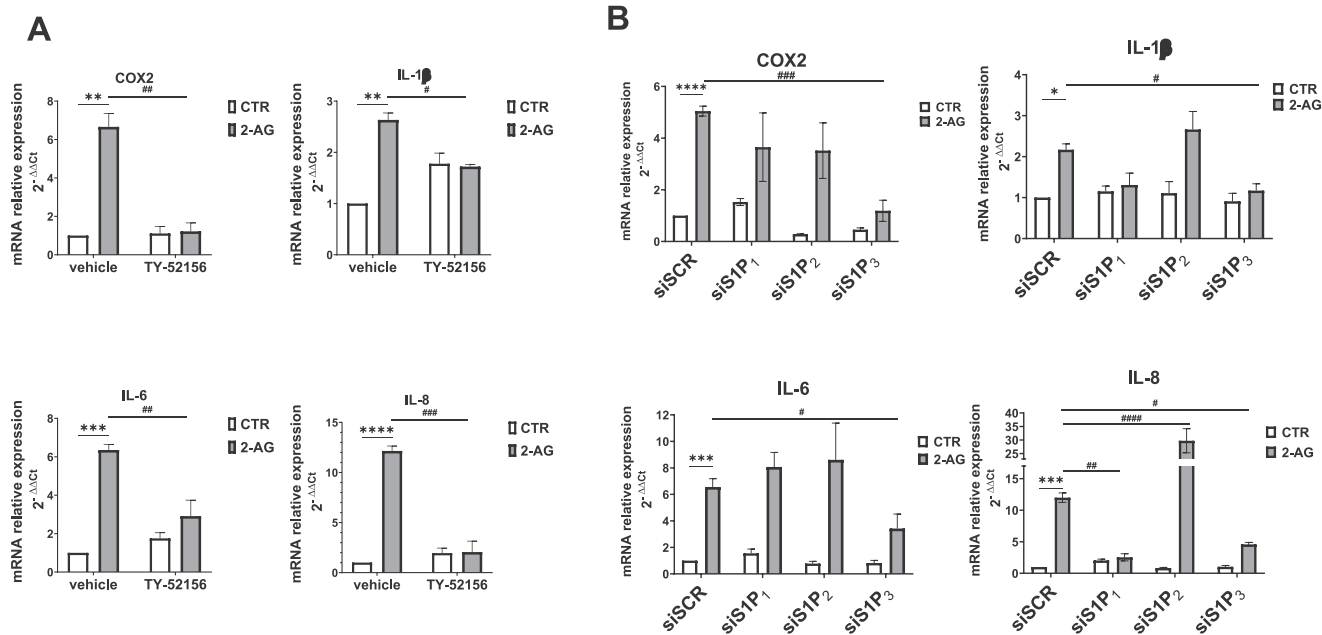


FIGURE 4 | 2-AG pro-inflammatory action relies on S1P₃. (A) Serum-starved endometriotic epithelial cells were pretreated or not with the S1P₃ antagonist TY-52156 (10 μM) for 45 min before being challenged with 10 μM 2-AG for 24 h. mRNA quantitative analysis of COX2, IL-1β, IL-6, and IL-8 was performed by qPCR. Results, analyzed with the 2^{-ΔΔCt} method, were obtained using β-Actin as a housekeeping gene and individual inflammatory factors of the unchallenged specimen as a reference gene. The blockade of S1P₃ on 2-AG-induced inflammatory effect (***p* < 0.01; ****p* < 0.001, *****p* < 0.0001) was statistically significant by two-way ANOVA followed by Bonferroni's post hoc test (#*p* < 0.05; ##*p* < 0.01; ###*p* < 0.001). (B) Endometriotic epithelial cells transfected with SCR-, S1P₁-, S1P₂- and S1P₃-siRNA were serum-starved prior to being challenged with 10 μM 2-AG for 24 h. mRNA quantitative analysis of COX2, IL-1β, IL-6 and IL-8 was performed by qPCR. Results, analyzed with the 2^{-ΔΔCt} method, were obtained using β-Actin as a housekeeping gene and individual inflammatory factors of the unchallenged specimen as a reference gene. The effect of S1P₃ downregulation in the reduction of 2-AG-induced inflammatory effect (**p* < 0.05; ****p* < 0.001, *****p* < 0.0001) was statistically significant by two-way ANOVA followed by Bonferroni's post hoc test (#*p* < 0.05; ##*p* < 0.01; ###*p* < 0.001; *****p* < 0.0001).

endometrium and endometriotic lesions [35–39], while this study represents the first experimental evidence as regards the expression of the receptor GPR18 in endometriosis. Our findings are in accordance with a previous work by Sanchez and coworkers [40] showing that CB1, CB2 and TRPV1 transcript levels were not different between endometrial stromal cells from endometriosis-affected women and healthy controls in the proliferative phase. Conversely, a significant increase in TRPV1 mRNA and protein levels was observed in rectosigmoid deep infiltrating endometriosis nodules [41]. Based on immunohistochemical analyses, endocannabinoid receptors CB1 and CB2 were found to be downregulated in endometriotic tissues [42]. Analogously, in accordance with the literature [39, 40, 42] we here demonstrate the expression of the endocannabinoid metabolic enzymes NAPE-PLD, DAGL, FAAH and MAGL in healthy endometrium and endometriotic lesions. Limited literature data are available as regards alterations in the expression of endocannabinoid-metabolizing enzymes in endometriotic lesions in respect to controls. In particular, it was shown that FAAH and NAPE-PLD transcripts did not significantly differ between endometrial stromal cells from endometriosis-affected women and healthy controls [40] whereas NAPE-PLD, DAGL, FAAH and MAGL protein levels were reduced in endometriotic lesions [42]. It will be interesting to analyze, in future studies, whether the expression of ECS components varies across endometriotic lesions of different anatomical localizations, with the aim of correlating the dysregulation of the ECS system to disease phenotype and severity.

In this study 2-AG, but not the non-hydrolyzable anandamide analogue, MAEA, was found to strongly upregulate mRNA levels of COX2, enzyme responsible for the conversion of arachidonic acid into prostanoids including prostaglandins, prostacyclin and thromboxane, all lipid mediators critically involved in the regulation of inflammation and pain perception [43]. Moreover, 2-AG augmented the release of pro-inflammatory cytokines IL-1β, IL-6 and IL-8 in endometriotic epithelial cells, confirming the role of ECS in the modulation of inflammation. Differential biological roles of endocannabinoid species have been previously extensively reported. For instance, 2-AG, but not AEA, exerted neuroprotective effects on granule cells in the dentate gyrus of the hippocampus [44], while only AEA inhibited amyloid β aggregation [45].

In accordance with the here reported proinflammatory role of 2-AG in endometriotic cells, increased levels of the endocannabinoid in the peritoneal fluid of women affected by endometriosis positively correlated with PGE2 concentration, a key mediator of inflammation [6]. Moreover, in endometrial inflammation the selective activation of CB2 was demonstrated to be associated with the release of nitric oxide, a key player in immune regulation and inflammation [46]. 2-AG has also been reported to increase the production of chemokines in HL-60 cells [47], while its seminal plasma levels are higher in the presence of inflammation [48]. However, in animal models of endometriosis, treatment with the two main constituents of the plant *Cannabis sativa* able to bind the cannabinoid receptors,

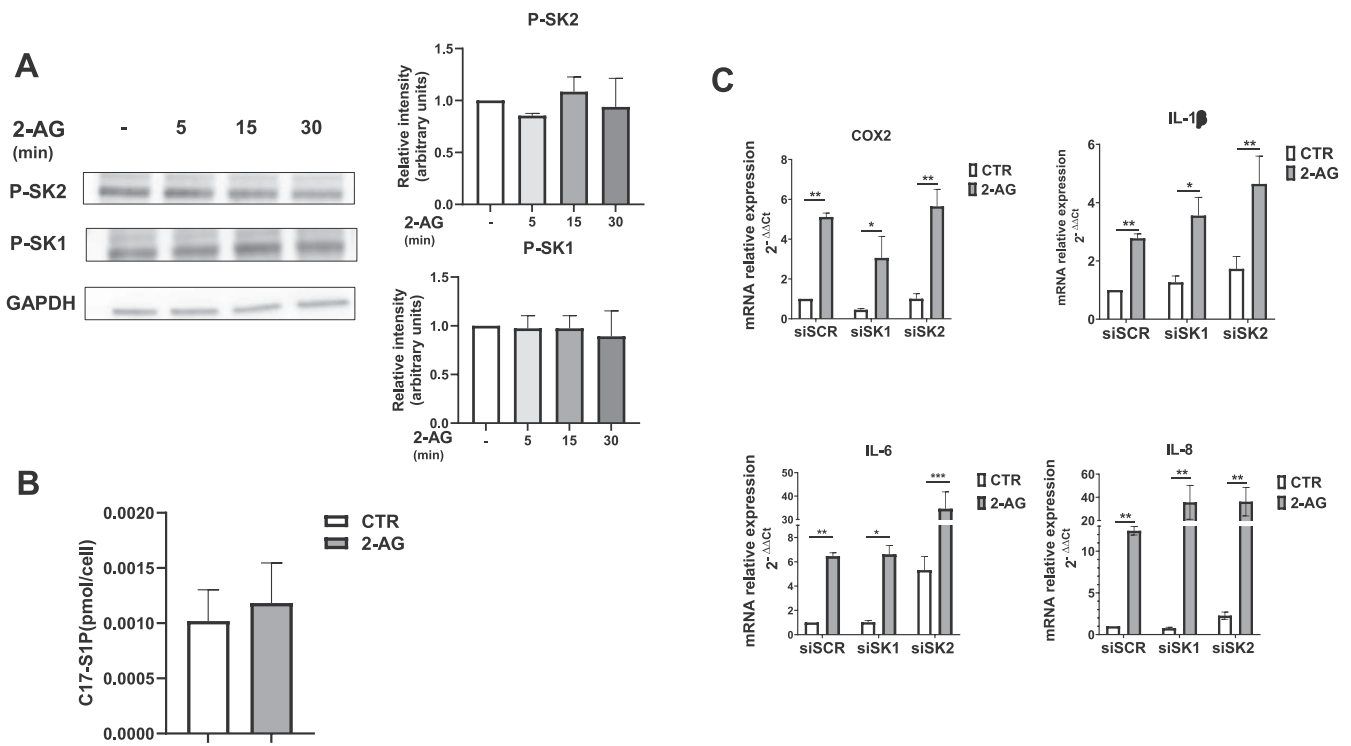


FIGURE 5 | 2-AG pro-inflammatory action is SK-independent in endometriotic epithelial cells. (A) Serum-starved endometriotic epithelial cells were treated with 10 μ M 2-AG for different time intervals (5, 15 and 30 min). Protein lysates were analyzed using SDS-PAGE electrophoresis and WB, using specific anti-phospho-SK1 (P-SK1), anti-phospho-SK2 (P-SK2) and anti-GAPDH antibodies. The histograms represent the densitometric analysis of three independent experiments. Data are the mean \pm SD and are reported as band intensity normalized to the expression of GAPDH, fold change over control (set as 1). (B) Serum-starved endometriotic epithelial cells were treated for 30 min with 10 μ M 2-AG before being harvested and then subjected to C17-S1P quantification by LC-MS/MS as described in the Section 2. Results are the mean \pm SEM of three independent experiments and are reported as pmol of S1P normalized on cell number. (C) Endometriotic epithelial cells transfected with SCR-, SK1- and SK2-siRNA were serum-starved prior to being challenged with 10 μ M 2-AG for 24 h. mRNA quantitative analysis of COX2, IL-1 β , IL-6 and IL-8 was performed by qPCR. Results, analyzed with the 2^{- $\Delta\Delta$ Ct} method, were obtained using β -Actin as a housekeeping gene and individual inflammatory factors of the unchallenged specimen as a reference gene. The effect of SK1 or SK2 downregulation did not significantly affect the 2-AG inflammatory effect (* p < 0.05; ** p < 0.01; *** p < 0.001).

D9-tetrahydrocannabinol and cannabidiol, was shown to ameliorate the endometriosis-associated pain [49] and inflammation [50], respectively, pointing at differential effects exerted by phytocannabinoids and endocannabinoids. It is worth noting that both AEA and 2-AG exert anti-inflammatory effects reducing the production of pro-inflammatory cytokines in immune cells [51–53]. In addition, 2-AG decreased the expression of COX2 elicited by different inflammatory stimuli in hippocampal neurons [54, 55] and leptin-induced reactive oxygen species formation in hypothalamic neurons [56].

Here, an interplay between ECS and S1P signaling in the production of inflammatory factors has been revealed in endometriosis. Indeed, 2-AG was found to modulate the S1P signaling axis since the treatment with the endocannabinoid upregulated the mRNA and protein content of S1P₃, while it augmented SK1 and Spns2 only at the transcriptional level. The reported findings showing S1P signaling modulation by ECS are in line with a previous study by Standoli et al. [57] in which it was demonstrated that pharmacological stimulation of CB2 counteracted the LPS-induced increase of SK1 and SK2 transcription in microglial cells. Reciprocally, ECS signaling regulation by S1P has been shown since the sphingolipid augmented the expression of

TRPV1, while it reduced CB2 levels in skeletal muscle cells [22]. Interestingly, here S1P₃ was demonstrated to mediate the pro-inflammatory action of 2-AG since its pharmacological inhibition or downregulation by gene silencing significantly reduced the elevated expression of COX-2, IL-1 β , IL-6 and IL-8 elicited by the endocannabinoid. These results provide compelling evidence that the upregulation of the analyzed pro-inflammatory molecules by 2-AG relies on S1P₃ engagement. However, this does not exclude that, in a reciprocal manner, the 2-AG-induced upregulation of S1P₃ is at least in part mediated by pro-inflammatory factors, thereby supporting a positive feedback mechanism that sustains inflammation. Notably, our research group has already provided multiple pieces of evidence that S1P₃ is crucially involved in endometriosis pathogenesis. Indeed, this receptor subtype has been found to be upregulated in endometriotic lesions [10, 14] and to positively correlate with the fibrosis extent of the disease [10]. Moreover, S1P₃ mediated the pro-fibrotic action of S1P [10] as well as the pro-invasive phenotype elicited by neuropeptide S [9] in endometriotic epithelial cells. Notably, in relation to the regulation of IL-8 levels, the data presented in this study demonstrated not only the involvement of S1P₃, similarly to what was observed for the other cytokines, but also a further positive role for S1P₁ and a negative regulatory role for S1P₂. Indeed,

when this latter receptor isoform was silenced in endometriotic epithelial cells, IL-8 levels were significantly increased following treatment with 2-AG, compared to control cells. This role of S1P₂ contrasts with previous findings in extravillous trophoblast-derived HTR-8/SVneo cells, where IL-8 release was dependent on the S1P₂/Rho signaling axis, highlighting that such regulatory mechanisms are highly context-specific [58].

Compelling evidence from the literature indicates that various extracellular cues elicit their biological effects through the regulation of SK1 and SK2, causing enhanced production of S1P, that after its export outside the cell can engage its cognate receptors S1PR [59]. Crucially here, 2-AG was demonstrated to be unable to enhance the catalytic activity of SK and consistently, SK-specific downregulation did not interfere with its pro-inflammatory action, thus ruling out the involvement of SK/S1P₃ inside-out signaling in endometriotic epithelial cells. On the contrary, SK1 activation induced by the endocannabinoid AEA was required to recruit S1P₃ and mediate vasorelaxation in the rat coronary artery [60] and the SK1/S1P regulatory axis was necessary for the rapid hypotension induced by AEA in anesthetized mice [23].

The action of 2-AG likely depends on its interaction with one or more specific receptors, found to be expressed in the endometriotic epithelial cells. However, our preliminary experimental approaches aimed at identifying the cannabinoid receptor involved in mediating the pro-inflammatory effect of 2-AG have so far not yielded robust results. Since the data presented here show that SK1/SK2 is not required to mediate the effects of 2-AG, one possible explanation for the activation of S1P₃ by 2-AG is a direct interaction between cannabinoid and S1P₃ receptors. Further studies are required to clarify the molecular mechanism by which 2-AG leads to S1P₃ transactivation in endometriotic cells. Of note, S1PR and CB1/CB2 belong to class A of GPCR and share 20% sequence identity. Notably, S1P₅ has been reported to exert a negative regulation of the tumorigenic effect induced by CB2 in glioblastoma cells: bioluminescence resonance energy transfer analysis highlighted that this S1P receptor subtype strongly and specifically interacts with CB2 [61].

Collectively, these findings enhance our understanding of the molecular mechanisms involved in the development of endometriosis pathogenesis, highlighting the critical role of S1P₃ in mediating the pro-inflammatory action of 2-AG in endometriotic epithelial cells. S1PR modulators have emerged as promising therapeutics for various immune-mediated diseases, including multiple sclerosis, inflammatory bowel disease and psoriasis, by modulating lymphocyte trafficking and reducing tissue inflammation [62]. Since we recently demonstrated that S1P₃ is involved in the fibrotic and invasive traits of endometriotic cells [9, 10], the here presented data further support the potential of targeting S1P₃ as a novel nonhormonal therapeutic strategy for endometriosis treatment.

Author Contributions

M.R., M.P., I.S., L.R. and E.N. performed the experiments and contributed to data analysis. P.B., F.P., C.B. and C.D. conceived or designed the experiments. C.B., C.D. and F.C. participated in data analysis. P.B., C.B.

and C.D. wrote the manuscript. F.P. revised the paper. All authors read and approved the final manuscript.

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Conflicts of Interest

The authors declare no conflicts of interest.

Data Availability Statement

All data generated or analyzed during this study are included in this published article. The analyzed datasets and materials used in the current study are available from the corresponding authors on reasonable request.

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Supporting Information

Additional supporting information can be found online in the Supporting Information section. **Figures S1-S4:** fsb271255-sup-0001-FigureS1-S4.pdf.