

ORIGINAL RESEARCH

The Impact of CBI Receptor on Inflammation in Skeletal Muscle Cells

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Background: Various factors trigger the inflammatory response and cytokine activation in skeletal muscle. Inflamed muscle will exhibit significant levels of inflammation and cytokine activity. Interleukin-6 (IL-6), a pro-inflammatory cytokine, exerts pleiotropic effects on skeletal muscle. Endocannabinoid produced by all cell types binds to a class of G protein-coupled receptors, in particular cannabinoid CB1 receptors, to induce skeletal muscle actions.

Objective: The purpose of this research was to discover whether activation of cannabinoid CB1 receptors in L6 skeletal muscle cells may promote IL-6 gene expression.

Materials and Methods: L6 skeletal muscle cells were cultured in 25 cm² flasks and quantitative reverse transcription-polymerase chain reaction (probe-based) utilised to quantify IL-6 gene expression levels among different treatment settings.

Results: Arachidonyl-2'-chloroethylamide (ACEA) 10 nM, a persistent selective CB1 receptor agonist, promotes IL-6 gene expression in a time-dependent manner. Rimonabant 100 nM, a selective cannabinoid CB1 receptor antagonist, blocks the impact of ACEA. However, insulin does not change IL-6 gene expression.

Conclusion: For the first time, a unique link between ACEA and IL-6 up-regulation has been established; IL-6 up-regulation generated by ACEA is mediated in skeletal muscle through cannabinoid CB1 receptor activation. As a result, cannabinoid CB1 receptors may be useful pharmaceutical targets in the treatment of inflammation and related disorders in skeletal muscle tissues.

Keywords: cannabinoid CB1 receptors, skeletal muscle, interleukin 6, ACEA, inflammation

Introduction

An endogenous signalling molecule is recognised as a crucial local regulator of tissue function. Cannabinoid modulates cellular processes via attaching to the two cell surface G protein-coupled receptor subtypes: CB1 and CB2. The CB1 receptor is the major receptor responsible for the effects of the endocannabinoid system in metabolic functions. CB1 receptors are the most common GPCRs expressed in the brain, but are also expressed in peripheral tissues such as adipose, liver, pancreas and skeletal muscle. On the other hand, CB2 receptors were found to be predominantly abundant in immune system. Indeed, CB2 receptors were expressed within the spleen, tonsils, and thymus. Cannabinoid possibly plays many essential functions in physiological processes such as glucose homeostasis and insulin sensitivity. As of now, nothing is known about the pharmacology and biological function(s) of the CB1 cannabinoid receptor. Cannabinoid has recently been shown to have a considerable effect on inflammation in response to diverse stressors.

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cannabinoid systems emerge as major factors in inflammatory regulation processes. ^{14,15} Targeting components, specifically the CB1 receptor, may aid in developing possible treatments for people with inflammatory diseases and related disorders, including insulin resistance in type II diabetes. ¹²

Inflammation plays a key role in the pathogenesis of disorders involving skeletal muscle failure. 16 Interleukin-6 (IL-6) is an inflammatory cytokine and mediator that affects immune and nonimmuno-regulation in a variety of cell types and tissues outside of the immune system, including skeletal muscle tissue. 12,17-19 IL-6 is a physiologically active cytokine that regulates inflammatory responses, skeletal muscle cell proliferation, differentiation, growth, and metabolism.²⁰ Recent data has shown that IL-6 is generated and released from skeletal muscle cells in the absence of inflammation and by complex signalling cascades. ^{21–23} IL-6 is then believed to be a myokine because of its role in skeletal muscle growth, myogenesis, and energy metabolism regulation. ^{23–25} Based on several investigational studies, prolonged IL-6 administration of mice results in insulin resistance in these animals.^{21–23} Furthermore, investigations in both animals and humans have found that cannabinoids have also been connected with type II diabetes and obesity.²⁶ Cannabinoids and CB1 receptor modulation do, however, influence downstream inflammatory pathways, in particular IL-6 in several tissues.27,28

Cannabinoid modulates several inflammatory cells, such as gastrointestinal inflammation. ^{29,30} Moreover, it appears that cytokine network processes are regulated by cannabinoid signalling.¹⁴ Cannabinoids enhance the secretion of IL-6 from different cells. 14,15 Previous studies have demonstrated that skeletal muscle tissue produces IL-6. 12,19 Increasing evidence indicates that skeletal muscle is implicated in the inflammatory process via secreting cytokines.²³ The effects of cannabinoids on muscle cell cytokine expression have not been evaluated. At the same time, cannabinoids are currently being explored for several disorders where inflammatory modulation is important. 11,13 Generally, research indicates that CB1 receptors, in particular, play a key role in inflammation.^{29,30} Indeed, stimulation of the CB1 receptor increases human macrophage, epithelial cell, and GI tract inflammation.^{29,30} Extensive in-vitro and in-vivo studies have demonstrated significant pro-inflammatory and/or antiinflammatory actions for all cannabis activities. 16,31,32 This receptor's function in skeletal muscle cells remains unknown.

Despite an emerging interest in the endocrine skeletal muscle, the role of the endocrine skeletal muscle's functional cannabinoid system in downstream inflammatory signalling is unknown. In an attempt to address this issue, a selective agonist of the cannabinoid CB1 receptors, ACEA and a selective antagonist of the cannabinoid CB1 receptors, rimonabant were investigated for their effects on IL-6 mRNA expression.

Materials and Methods

Materials

Tocris Bioscience (UK) supplied the insulin, ACEA, and rimonabant, while Santa Cruz (USA) provided the dimethylsulphoxide reagent. Thermo Scientific Company (USA) supplied the Maxima Probe qPCR Master Mix (2X) and Thermo Scientific RevertAid First Strand cDNA Synthesis, while Qiagen (Germany) was the provider of the RNeasy Mini Total RNA Purification kits and RNase-Free DNase Set. Applied Biosystem (USA) provided Trizol and charcoal-stripped serum, while FBS (fetal bovine serum) was obtained from Capricorn Scientific (USA). Sigma Company (Germany) supplied horse serum, and Ham-F 10 was obtained from PAA Company (USA). Dulbecco's modified essential medium (DMEM) was supplied by Caisson (USA).

Cell Culture

The American Type Culture Collection (USA) provided the L6 skeletal muscle cell line from rats along with the myoblast cell line, originating from cells, maintained in the form of an attached monolayer culture in DMEM, with a high glucose level (4500 mg/L) with L-glutamate which had been supplemented by 10% (v/v) heat-inactivated FBS and 100 µg/mL of penicillin-streptomycin. Incubation of the cells took place at a temperature of 37°C using a 5% carbon dioxide atmosphere at 90% humidity. Passaging of the cells occurred at around 60%-70% confluency while changing the medium was carried out thrice weekly, as shown in Figure 1. The confluent cells underwent 14 days of further culturing in 25 cm² flasks in order to allow myotubes to form, in alignment with the protocols outlined in my previous publications, 48,49 albeit with minor modifications (Figure 1). After around two weeks of culturing, the 70%– 90% confluent myotubes were then exposed to 2% (v/v) delipidated serum for a period of 5 hours before undergoing starvation for a further 19 hours. Figure 1 then shows that the cells were treated for varying periods of time (1, 3, 5, and 24 hours) with vehicle (0.1% DMSO), rimonabant 100nM, ACEA 10nM, and insulin 100nM. The ACEA and rimonabant cells had undergone pre-treatment using rimonabant for

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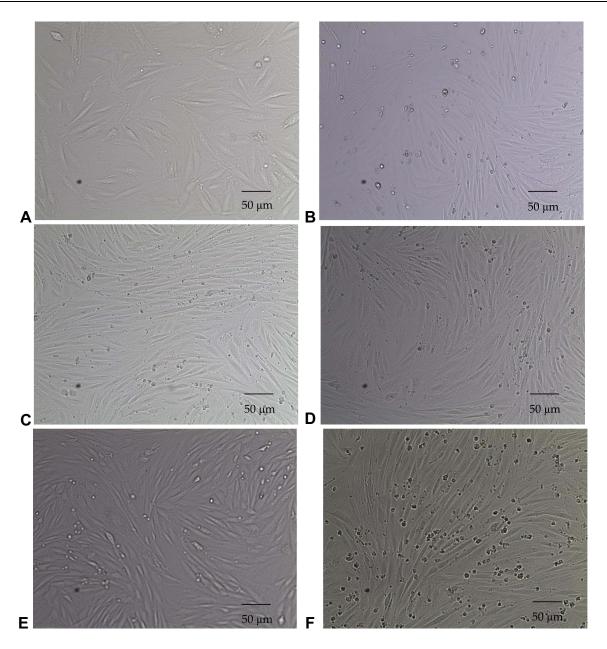


Figure 1 Images depicting L6 skeletal muscle myoblasts from rats, differentiated into L6 skeletal muscle myotubes (passage 7). (A) Myoblasts obtained on day 3 of the tissue culture in 10% FBS Ham F-10 media (10X). (B) Cells obtained on day 4 of the tissue culture in 6% horse serum Ham F-10 media (10X). (C) Myotubes obtained on day 5 of the tissue culture in 2% horse serum Ham F-10 media (10X). (D) Myotubes obtained after 4 hours of tissue culture in 2% delipidated serum Ham F-10 media (10X). (E) Myotubes obtained after 19 hours of cell starvation in only Ham F-10 media (10X), (scale bar 50μm).

10 minutes before adding ACEA. After the treatment, the cells were washed using ice-cold PBS, before lysing using Trizol (2mL per flask).

Extraction of RNA and Synthesis of cDNA

The L6 skeletal muscle cells of rats were placed in 25 cm² flasks and scraped in 2 mL of ice-cold Trizol, whereupon the RNA was separated and isolated in line with the guidelines of

the manufacturer. RNeasy purification columns (Qiagen, Germany) were then used to carry out the RNA clean-up and on-column DNAse digestion. A spectrophotometer (JENWAY Genova Nano) was then used to assess the concentration and purity of the RNA. In order to carry out the cDNA synthesis, a quantity of 500 ng of total RNA underwent reverse transcription via RevertAid First Strand cDNA Synthesis in a process taking 5 minutes using a total volume of 20 μ L at a temperature of 25°C, before the temperature was increased to 42°C for the

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Table I List of Gene Primer and Probe Sequences

| Gene | Sequences (5'3') |
|--------------|---|
| IL-6 | Probe: 5'-CTCTCCGCAAGAGACTTCCAGCCAGTT-3' Forward primer: 5'-GCCCTTCAGGAACAGCTATGA-3' Reverse primer: 5'-TGTCAACAACATCAGTCCCAAG A-3' |
| TATA- BOX | Probe 5'- TCCCAAGCGGTTTGCTGCAGTCA -3' Forward Primer 5'- TTCGTGCCAGAAATGCTGAA-3' Reverse Primer 5'- GTTCGTGGCTCTCTTATTCTCA TG -3' |

subsequent one hour period. Finally, the termination of the reaction took place for 5 minutes at 70°C. Gene expression was then quantified using the relative standard curve approach on the basis of the TagMan quantitative real-time PCR (qRT-PCR). For this process, the preparation of the samples was carried out using a total reaction volume of 25 µL (comprising 13 μL Maxima Probe qPCR Master Mix 2X reagent, 1.5 μL of forward primer (10 µM), 2.5 µL Probe (2 µM), 1.5 µL of reverse primer (10 mM), 5 µL of water, and 5 µL of cDNA). A 7500 fast real-time PCR system (Applied Biosystems, USA) was employed to complete the qRT-PCR analysis, while the determination of the gene expression was made by considering the relationship to the reference gene, TATA. Primer Express software (Applied Biosystems, USA) was used in the case of probes and primers for all genes, as shown in Table 1, with the design and synthesis performed by Integrated DNA Technologies, Inc. (USA). The standard curve approach was employed, using a slope ranging from -3.2 to -3.6 with R^2 values exceeding 99%, reflecting efficiency of amplification approaching 100%.

Data Analysis

Data are presented in the form of mean \pm SEM following the generation of triplicate or quadruplicate wells from no fewer than three experimental groups. Data analysis of mRNA data employed one-way ANOVA and a Tukey's test. The GraphPad Prism, version 5.03 (GraphPad Software Inc) was used for all analyses, and the statistical significance level was determined to be P < 0.05.

Results

Cannabinoid CBI Receptors Induced IL-6 mRNA Expression in Rat L6 Skeletal Muscle Myotubes

Treatment with ACEA (10 nM) for 3 hours significantly upregulated the IL-6 gene expression (p<0.05) but these responses were blocked by Rimonabant (100 nM) while treatment with Rimonabant (100 nM) for 24 hours dramatically decreased the IL-6 gene expression (p<0.01). ACEA (10 nM) completely reversed this reaction (p<0.01) and insulin had no effect on IL-6 mRNA expression (Figure 2).

Discussion

This study's novel findings are that the IL-6 expression by skeletal muscle cells is escalated by stable cannabinoid analogue ACEA. The cannabinoid CB1 receptor subtype acts as the mediator for this ACEA effect. An analysis of available literature shows that this is the inaugural study exploring the cannabinoid analogue and its receptor subtype effect on inflammatory cytokine expression by skeletal muscle cells. Therefore, this study has the potential to add to available knowledge by presenting a novel contrivance for cannabinoid analogue's role in skeletal muscle cytokine network. Numerous papers provide proof to show that skeletal muscle cells collected from various species contain cannabinoid receptors.³³ Apart from CB1 receptor gene expression, with nuclear receptors (NR4A) gene expression used as a functional downstream signalling readout of CB1 receptors (Data are not published yet), there is proof that functional cannabinoid CB1 receptors exist in the cells of the skeletal muscle. Cell surface cannabinoid receptors have also been noted to mediate numerous cannabinoid physiologic roles.³⁴ In the present study. I present proof that the effect of cannabinoid analogue ACEA on IL-6 expression is mediated by the cannabinoid CB1 receptors subtype. From my results, it can be concluded that the IL-6 expression is increased by the selective agonist ACEA. This very strong agonist, at a concentration of 10 nM, has the potential to completely stimulate their cognate receptors in the absence of any substantial stimulation of the cannabinoid CB1 receptors subtype, rimonabant. When looked at together, these conclusions present robust evidence regarding cannabinoid CB1 receptors' role in the IL-6 upregulation initiated by ACEA.

To differentiate which receptor mediates this effect, ACEA was also used since ACEA is more selective for CB1 receptor than CB2 receptor by around 2000 times.³⁵ ACEA was found to increase IL-6 mRNA expression in rat L6 myotubes. Interestingly, this effect induced by ACEA was blocked by rimonabant, a selective CB1 receptor antagonist/inverse agonist.36 This finding suggests that IL-6 activation was mediated by activation of the CB1 receptor since ACEA is a selective CB1 receptor agonist

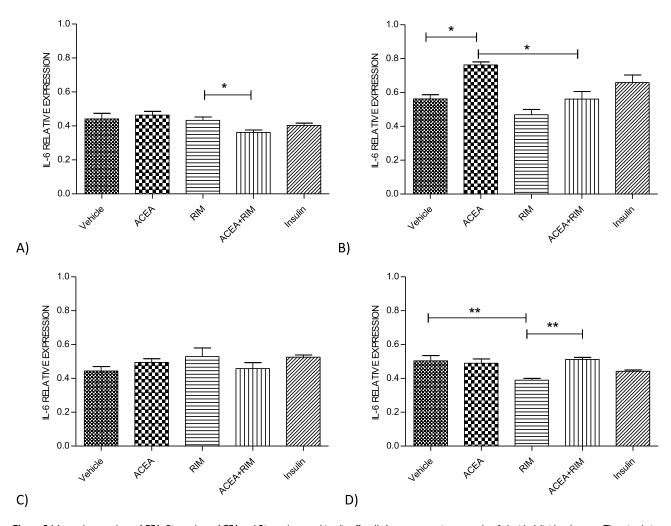


Figure 2 L6 muscle myotubes—ACEA, Rimonabant, ACEA and Rimonabant, and insulin affect IL-6 gene expression; myotubes fed with delipidated serum. The stimulation time covered I to 24 hours, and IL-6 mRNA levels, relative to TATA-Box, was evaluated by quantitative real-time PCR (qRT-PCR) (I00 nM). The following scenarios explain the stimulation process conducted: (A) stimulation was done for I hour. (B) Stimulation was done for 3 hours. (C) Stimulation was applied for up to 5 hours. (D) Stimulation was applied for up to 24 hours. The data were reported as the mean ± SEM of three separate groups. (n=3; *denotes p=0.05 and **denotes p=0.01). Data were investigated by conducting one-way ANOVA test and Tukey's test.

Abbreviation: IL-6, Interleukin 6.

at the concentration used in the present study.³⁵ This study also provided strong evidence that CB1 receptor is a functionally active receptor in skeletal muscle. This is in line with a previous study that also found that CB1 receptor was found to be expressed in skeletal muscle.^{11,12,33}

Through the present study, it has been shown that when cannabinoid CB1 receptors are activated, the expression of interleukin-6 (IL-6) mRNA is escalated. This possibly crucial mechanism/process that has a vital influence on skeletal muscle function regulation is still unknown. Sometimes scholars perceive skeletal muscle as an endocrine organ. This is because skeletal muscle discharges an inflammation mediator known as Interleukin-6 and is responsible for increasing metabolism when human

skeletal muscle is at rest.³⁷ It has been concluded that obese individuals tend to have high levels of Interleukin-6, which has also been linked to Type II diabetes.³⁸ It has also been concluded that IL-6 increases the oxidation of fatty acids and transportation of glucose in L6 myotubes.³⁹ Cannabinoid agonist WIN55, 212 inhibited the release of IL-6 in the pancreatic acini of rats.⁴⁰

In the present study, IL-6 mRNA expression in rat L6 skeletal muscle myotubes was triggered by ACEA stimulation of cannabinoid CB1 receptors. Such results correspond with those of another study that used Theiler's murine encephalomyelitis virus-infected astrocytes to show that anandamide CB1 receptor agonist potentiated the production of IL-6.⁴¹ When these results are considered, it seems possible that the secretion of IL-6

inflammatory cytokine in skeletal muscle myotubes and the consequent modulation of fatty acid and glucose metabolism is increased by the activation of the CB1 receptor. This implies that inflammatory cytokines modulation, especially IL-6, via the inhibition and/or activation of skeletal muscle CB1 receptors could potentially change the metabolism of fatty acids or glucose, insulin resistance, proliferation, differentiation, muscle growth, myogenesis and apoptosis, which could have a positive impact on obesity, diabetes, and sarcopenia. Moreover, the present study supported an evidence about the role of cannabinoid CB1 receptors in IL-6 regulation in L6 skeletal muscle cells, which is consistent with previous study that showed an increased IL-6 level in skeletal muscle-specific CB1Rknockout (Skm-CB1-/-) mice and demonstrated important benefits to insulin resistance, metabolism, and muscle mass in absence of skeletal muscle cannabinoid CB1 receptors in diet-induced obese and aged mice. 42 And also, in this case, it has demonstrated that the pharmacological antagonism of CB1 by Rimonabant, but not its agonism by ACEA, exerts anti-inflammatory effects in dystrophic mice, 43 which is line with this current research study.

The present study reports that the effects of ACEA (in Figure 2) can be noted in skeletal muscle cells. It could also be brought to the fore that the ACEA stimulations lasting 5 and 24 hours did not work, while it did on the 3-hour ones are on cells (Figure 2). This can be credited to the fact that cannabinoid CB1 receptors, being G-protein coupled receptors, require a specific period for coupling the G-protein before it can target the expression of the IL-6 mRNA gene through inducing that transit downstream signalling pathway. Notwithstanding the reality that ACEA cell concentration does not have the capability to elucidate the difference in the effect of the 1, 5 and 24 hours treatment, a crucial factor may still be duration exposure. The indication that expression density/level of cannabinoid CB1 receptors in the skeletal muscle could meet a necessary condition for precise cannabinoid CB1 receptors activator action to change the cannabinoid CB1 receptor-responsive genes expression.

Determining the specific IL-6 transcripts at a precise time point that results in the production of IL-6 is a challenge. Nonetheless, it is vital to propose that the connection between protein synthesis and the content of RNA in skeletal muscle cells is not always a fixed linear correlation.44 It can be argued that the IL-6 expression changes in inflammation as seen in the current study could be fleeting and therefore not be represented by the protein expression changes, and this is insignificant with regards to the development of inflammation as a reaction to stable cannabinoid analogue ACEA. The leading reason this study did not include the measurement of total protein is that the treatment duration was short, which in keeping with my main objective, was used for the identification of transcriptional events coming before inflation begins as a result of being activated by cannabinoid CB1 receptors. Previously, it has been indicated⁴⁵ that the majority of the inflammatory transcripts studied (like IL-6) in skeletal muscle cells do not translate into protein during the timeframe specified. This is why future studies involving longer durations (ie, more than 24 hours) are suggested to allow for exploring the translated changes in reaction to treatment. This could deliver a better chance to explore the changes taking place at different intervals as durations are made longer (up to four days). On this basis, it becomes apparent that cannabinoid receptors, especially cannabinoid CB1 receptors, constitute crucial molecular targets for treating inflammation using cannabinoid-based therapies. Methods that use therapies based on cannabinoid receptors will rely on more studies exploring cannabinoid CB1 receptors in skeletal muscle signalling mechanisms, treatment duration and timing, and adverse and beneficial effects monitoring.

Surprisingly, in the current study, insulin 10 nM did not change IL-6 expression in skeletal muscle cells using these time frame. This is not in line with other studies which showed that insulin induced IL-6 expression in adipocytes. 46 Therefore, it is suggested in the future studies to try higher concentration and shorter time scale, in particular insulin is considered as an important hormone affected by inflammation and/or crosstalk with cannabinoid receptors in terms of biological and metabolic functions. 11,12 Of note, the mechanism which is proposed on how cannabinoid CB1 receptor is regulated IL-6 might be through direct regulation (Gi-cAMP-PKA),²⁷ or through indirect regulation of Akt via a Gi/PI3K-Akt/NF-kB pathway, 28 or via PI3K/Akt/mTOR signaling, 47 although further studies are required to explore these suggested mechanisms. It can be noted that the levels of mRNA could be a reflection of turnover and signalling protein transcription, although this might be happened at different time frame. Nevertheless, it can be suggested that there is a need for more studies assessing protein expression level (intracellular), release and activation

(phosphorylation). Added to this, more work is required in exploring some functional measurements like interleukin release using L6 skeletal muscle cells or primary skeletal muscle cells. More work is also suggested to assess the direct effect of cannabinoid CB1 receptor on IL-6 using knock out of CB1 receptor in L6 skeletal muscle cells or to mimic the muscle physiology using knock out of CB1 receptor in primary myoblast isolated from mice or rats. This could provide more evidence with regards to this signalling. Other areas that could benefit from more studies include assessing more important genes connected to this signalling in skeletal muscle, including TNFalpha.

Conclusions

In the present study, I have demonstrated that when cannabinoid CB1 receptors are activated, the expression of IL-6 increases in a way that is influenced by time. Such findings deliver a novel mechanism characterised by cannabinoid analogue playing the role of a pro-inflammatory mediator in the skeletal muscle tissue. The findings from the present study also imply that there may be a possible therapeutic use of cannabinoid CB1 receptor antagonist at acute early states for skeletal muscle dysfunction related to inflammation. My findings point to skeletal muscle cell cannabinoid CB1 receptor as a therapeutic target, and expand its potential to include anti-inflammatory effects in diabetes, obesity, and sarcopenia.

Significance

This study describes a previously undiscovered signalling system in skeletal muscle cells that involves cannabinoid CB1 receptors. Furthermore, it explains a previously unknown process of muscle inflammation to reveal another possible treatment target for metabolic diseases associated with inflammation.

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Author Contributions

The author declares that this work was done by the author named in this article and all liabilities pertaining to claims relating to the content of this article will be borne by the author. All experimental work, study design, statistical analysis, interpretation of data, writing and revising the manuscript was performed by Mansour Haddad. The author has read and approved the final manuscript and has agreed on the journal to which the article has been submitted.

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Disclosure

The author declares that he has no conflicts of interest.

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