



Research Article

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The Effects of Cannabinolic Acid Amides on TNF- α and IL-6 Expression in Lipopolysaccharide Activated Human Peripheral Blood Mononuclear Cells

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Abstract

Introduction: The effects of tetrahydrocannabinolic acid (THCA) and cannabidiolic acid (CBDA) amide derivatives 1-9 on TNF- α and IL-6 expression and cell proliferation were investigated on lipopolysaccharide (LPS) activated human peripheral blood mononuclear cells (PBMCs).

Materials and Methods: Compounds 1-9 were tested on LPS activated human PBMCs and several inflammatory endpoints monitored. Negative controls included PBMC only, LPS and dexamethasone with LPS. After a 72-hour incubation, PBMC supernatants were collected to measure IL-6 and TNF- α levels by ELISA. Cell pellets were harvested, and cell proliferation analyzed with the CTG assay.

Results: Compounds 1 and 2 increased TNF- α expression and inhibited cell proliferation, while unsubstituted amides 5, 6 decreased the production of TNF- α compared to LPS-treated cells. None of the compounds tested had a noticeable effect on LPS induced IL-6 expression.

Conclusion: The change of substitution in the amide group yields THCA and CBDA derivatives with different and sometimes opposing activities with respect to TNF- α expression with a general lack of effect on IL-6 output. The function of several of these compounds as potential activators of the immune response to viral infections and antitumor activity, as well as their immunosuppressive potential warrants further investigation and may highlight the contributions of such novel cannabinolic acid-based drugs to pharmacology.

Keywords: Cannabinoids, THCA, TNF- α , IL-6, LPS, PBMCs

Introduction

The spread of the Covid-19 virus in the world has significantly intensified research into effective treatments for this disease. Since progression of the disease includes two stages - infection with the virus and activation of host immunity, with further dissemination of the infection, a "cytokine storm" arises which is characterized by

an uncontrolled release of inflammatory cytokines, predominantly involving TNF- α and IL-6. Unfortunately, the main drugs used against COVID-19 do not provide effective treatment. The search for new drugs that can effectively and quickly block the expression of TNF α , IL-6 and other inflammatory cytokines has therefore become the focus of the fight against this severe disease [1].



The use of cannabinoids as new potential anti-inflammatory drugs for the treatment of COVID-19 has drawn the attention of researchers from the beginning of the pandemic [2-4]. Cannabinoids bind to cannabinoid receptors CB1 and CB2 thereby promoting the expression of pro- and anti-inflammatory cytokines in immune cells. To date cannabinoid studies have predominantly focused on tetrahydrocannabinol, and particularly on cannabidiol, as these are the most affordable and stable products isolated from marijuana [5]. The potential medical uses of their precursors tetrahydrocannabinolic (THCA) and cannabidiolic (CBDA) acids have been disregarded because their pure forms are difficult to isolate, and they tend to decarboxylate. It is only recently that several studies have been devoted to their biological activities their ability to inhibit the release of cyclooxygenases during inflammation [6,7].

The great opportunities, that may be captured by the synthetic transformation of the carboxyl group, also presage the significance of these acids as potential novel drug precursors [8,9]. This is particularly relevant in the context of the new technology which allows to extract these acids from plant materials using ion-exchange resins and yields a purity suitable for further synthesis [10]. The current work is an activity study of several of these previously described THCA and CBDA amides, specifically focusing on their effects on LPS-induced TNF- α and IL-6 secretion as well as on proliferation of human PBMCs. The choice of TNF- α and IL-6 was based on widespread reports of their significant contribution to cytokine pro-inflammatory activities during the "cytokine storm" induced by COVID-19 viral infection [11].

Materials and Methods

Compounds 1-9 were obtained from THCA and CBDA [10] in accordance with methods of their synthesis [8].

(1'R,2'R)-2,6-dihydroxy-5'-methyl-4-pentyl-2'-(prop-1-en-2-yl)-1',2',3',4'-tetrahydro-[1,1'-biphenyl]-3-carboxamide 6

Yield 48%. ¹H NMR (CDCl₃), δ : 0.88-0.91(3H, t, CH); 1.33-1.35(4H, m, CH); 1.62-1.65(3H, m, CH); 1.70(3H, s, CH); 1.78-1.99(5H, m, CH); 2.10-2.17(2H, bd, CH); 2.2.19-2.23(1H, m, CH); 2.36-2.45(1H, m, CH); 2.71-2.74(2H, m, CH); 4.08(1H, bs, CH); 4.39(1H, s, CH); 4.52(1H, s, CH); 5.58(1H, s, CH); 5.94(2H, s, NH); 6.41(1H, s, CH); 11.94(1H, s, OH). ¹³C NMR (CDCl₃), δ : 13.98, 18.87, 22.45, 23.71, 27.86, 30.25, 30.85, 31.78, 35.21, 46.63, 107.57, 110.39, 111.15, 115.096, 124.08, 140.13, 140.83, 147.36, 158.83, 161.17, 175.31. Molecular ions [M-H]⁺=358 were observed, consistent with the molecular formula C₂₂H₃₁N₃O₃. Effects of these compounds were tested on human PBMCs challenged with Lipopolysaccharide (LPS) by monitoring several inflammatory

endpoints.

Materials and Equipment

VICTOR Nivo Multimode Microplate Reader (PerkinElmer, VICTOR Nivo); Lymphoprep™ (Stemcell, Cat. No. SV30010, Lot. No. J170012); SepMate™-50 (STEMCELL Technologies, Cat. No. A10491-01); Lipopolysaccharides from Escherichia coli O111:B4 (Sigma, Cat. No L2630-10MG); Dexamethasone (Alfa Aesar, Cat. No A17590); Human TNF- α ELISA (Biolegend, Cat. No 430204); Corning® 96-well Flat Clear Bottom White (Corning, Cat. No 3903); Cell Titer-Glo® Luminescent Cell Viability Assay (Promega, Cat. No G7572). Blood samples were collected from healthy volunteers who had signed the Informed Consent Form (ICF) according to protocols reviewed and approved by Chempartner Institutional Ethic Committee (IEC).

PBMC Isolation

Human blood samples from individual donors were diluted with an equal volume of sterile Phosphate Buffered Saline (PBS) and mixed by gentle agitation. Fifteen milliliters of Lymphoprep medium were transferred into a new 50 mL centrifuge tube, whilst ensuring that the Ficoll and blood (30 mL) volume ratio was kept at 1:2 and the diluted blood sample was then carefully added onto the surface of the Ficoll medium. Centrifugation resulted in the appearance of four layers. From top to bottom of the tube these layers corresponded to plasma, mononuclear cells, Ficoll medium and RBCs. After removing the plasma, mononuclear cells were transferred into a new sterile centrifuge tube. Sterile PBS buffer was added to the collected PBMCs for washing at a volume ratio of 3:1. PBMCs were washed with 5-10 mL PBS twice more before counting cells with a cytometer.

Cells were centrifuged at 350x g, 10 min, 20°C, with an acceleration and deceleration setting of 5 and resuspended in RPMI 1640 complete medium (containing 10% Fetal Bovine Serum FBS) for the assay. Cell density was adjusted to a final concentration of 2E6 cells/mL with RPMI 1640 complete medium. Fresh PBMC cell suspensions (100 μ L/well) were seeded into 96-well plates (#3903) and a dilution series of compounds and 0.1 μ g/mL LPS added to each well (in triplicates). Controls included 3 wells with PBMCs only, 3 wells with 0.1 μ g/mL LPS and 2 wells with 1 μ g/mL dexamethasone (Dex) and 0.1 μ g/mL LPS. After 72 hours of incubation, the supernatant was collected for Tumor Necrosis Factor- α (TNF- α) measurement by Enzyme-linked Immunosorbent Assay (ELISA), and cell pellets were harvested to measure cell proliferation using the CellTiter-Glo® Luminescent Cell Viability Assay (CTG).

Preparation of Compound Solutions

Compounds were diluted with RPMI 1640 complete medium according to the layout. Compounds were diluted to a working concentration of 10 μM with RPMI 1640 complete medium and further diluted in two 10-fold serial dilution steps. Compounds were prepared at a 5-fold concentration (50 μM). Diluted compounds (50 $\mu\text{L}/\text{well}$) were added to the indicated wells, and plates incubated at 37 $^{\circ}\text{C}$, 5% CO_2 incubator for 72 hrs.

LPS induced cytokine release assay: LPS (0.1 $\mu\text{g}/\text{mL}$, 100

μL) was added to each well. Dex (1 $\mu\text{g}/\text{mL}$) was included as a positive control. Plates were incubated at 37 $^{\circ}\text{C}$, 5% CO_2 for 72 hrs. Supernatant was collected to measure IL-6 and TNF- α by ELISA. Cell proliferation was measured by CTG.

Statistical Analysis

Data analysis was performed using the Graphpad Prism 6.0 software. Data is represented as mean value with standard error of the mean (SEM). Data of all experiments presented in (Figures 1-3) are one-way ANOVA analyses compared to the LPS treated group.

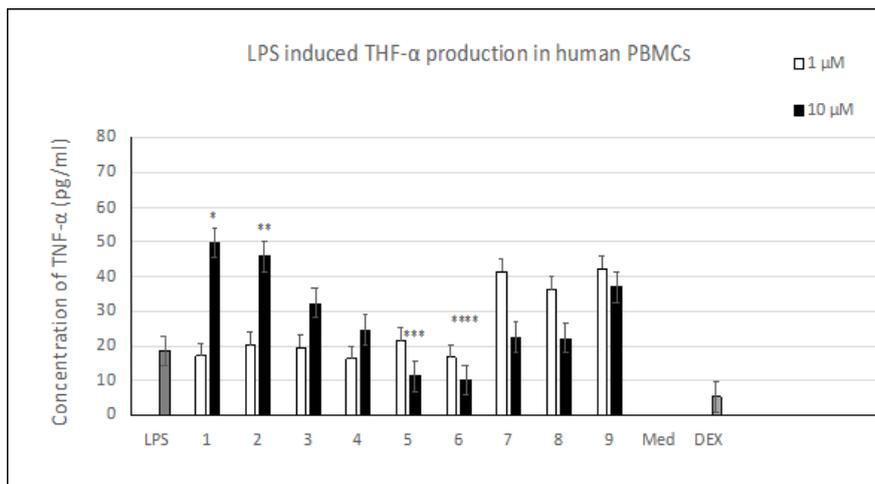


Figure 1: Influence of the amino group structures of compounds 1-9 on LPS induced TNF- α production (pg/ml) in human PBMCs (LPS – 0.1 $\mu\text{g}/\text{ml}$; DEX – 1.0 $\mu\text{g}/\text{ml}$ + 0.1 $\mu\text{g}/\text{ml}$ LPS; One-way ANOVA analysis: *p, **p, ***p, ****p < 0.01 when compared to the LPS treated group; n=3.

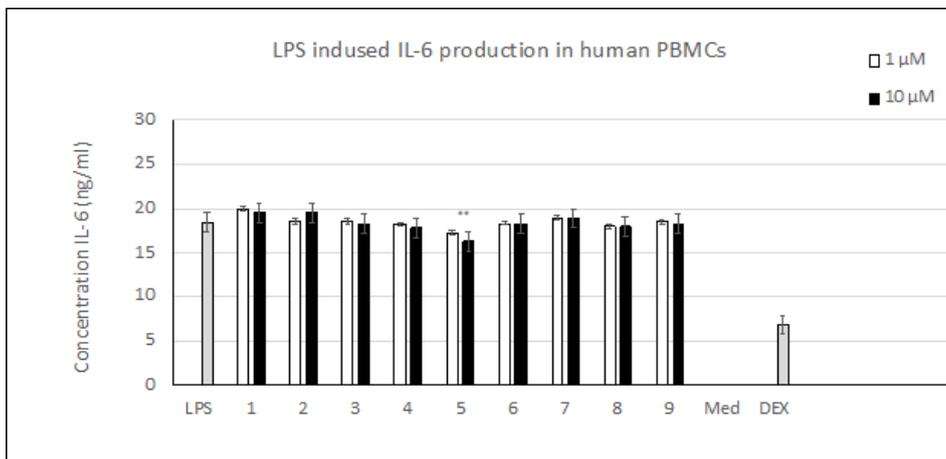


Figure 2: LPS induced IL-6 production (ng/ml) in human PBMCs (LPS – 0.1 $\mu\text{g}/\text{ml}$; DEX – 1.0 $\mu\text{g}/\text{ml}$ + 0.1 $\mu\text{g}/\text{ml}$ LPS; One-way ANOVA analysis: **p < 0.005 when compared to the LPS treated group; n=3.

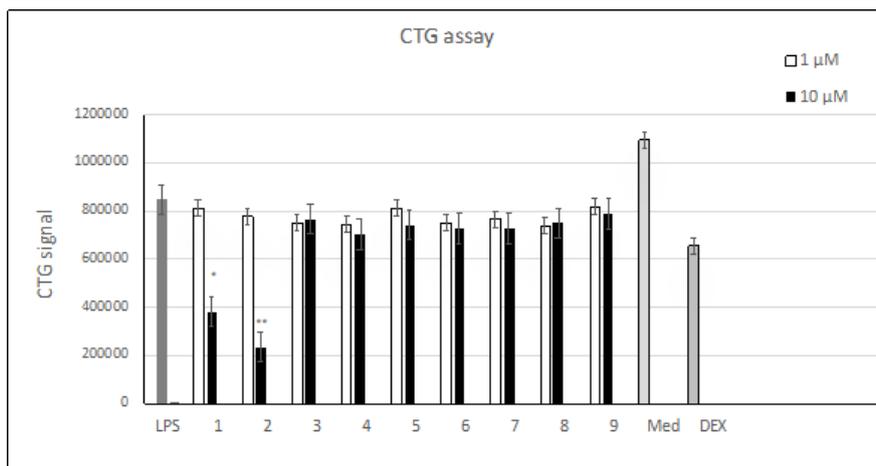


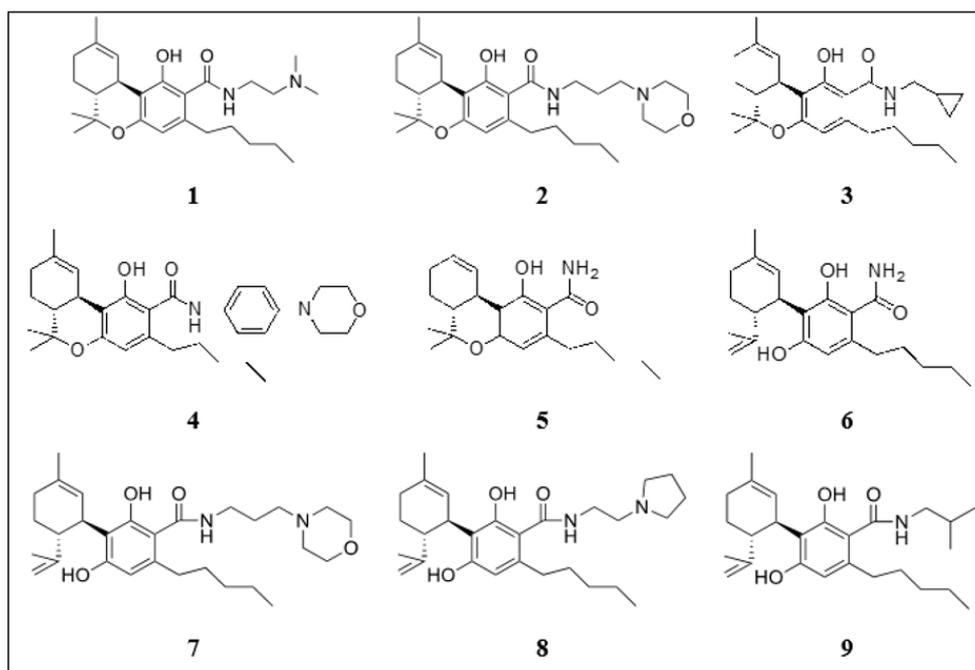
Figure 3: Cell proliferation in the LPS induced PBMC system (Med- Medium Ctrl; LPS – 0.1 μg/ml; DEX – 1.0 μg/ml + 0.1 μg/ml LPS; One-way ANOVA analysis: * $p < 0.005$, ** $p < 0.001$ when compared to the LPS treated group; $n = 3$).

Results

Compounds 1-9 were obtained by amination of the carbonyl group of the THCA and CBDA aromatic rings [8] (Table 1). Evaluation of TNF- α levels induced by LPS. As shown on Figure 1, the addition of 0.1 μg/mL LPS induced an obvious increase in the TNF- α level in human PBMCs. THCA derivatives 1-4 promoted the production of TNF- α compared to LPS-treated cells. Compound 5 and its CBDA analog, compound 6, resulted in a dose-dependent decrease in TNF- α . Compounds 7-9 increased the production of TNF- α at a concentration of 1 μM and decreased TNF- α levels at

a concentration of 10 μM. Evaluating IL-6 levels induced by LPS. As shown on Figure 2 the addition of 0.1 μg/mL of LPS induced an obvious increase in the level of IL-6 in human PBMCs, as was the case for TNF- α . Compound 5 dose-dependently decreased the production of IL-6 compared to LPS-treated cells, but this value did not exceed 10% at a compound concentration of 10 μM. The remaining substances did not have any noticeable effects. Cell proliferation in the LPS induced PBMC system. As shown in (Figure 3), at 10 μM, compounds 1 and 2 inhibited cell proliferation 2.5 to 3-fold compared to LPS-control cells. The remaining compounds show a slight decrease in toxicity compared to dexamethasone.

Table 1: Structures of THCA and CBDA amide compounds 1-9.



Discussion

TNF- α is a pro-inflammatory cytokine and appears during an injury response or bacterial and viral infections. Other important features are its ability to cause hemorrhagic necrosis of tumors *in vivo* in combination with its ability to kill tumor cells *in vitro* [12]. IL-6 is another major proinflammatory cytokine with a variety of effects on the immune system, particularly in the development of long-term immune activation and contributes to the progression of the COVID-19 induced “cytokine storm” [11]. IL-6 also plays an important role in the onset and development of many types of cancers such as pancreatic and colorectal cancers [13].

The structure of the amide substituent in compounds 1-9 had a significant effect on the expression of TNF- α (Figure 1). According to the data obtained, all compounds examined can be subdivided into two groups depending on the change in TNF- α expression induced by compound concentrations of 1 μ M and 10 μ M. The first group includes THCA derivatives 1-4, the second – unsubstituted THCA amide 5 and CBDA amides 6-9.

The ability of compounds 1 and 2 to induce the expression of TNF- α together with their inhibition of cell proliferation (Figure 3) is consistent with the previously reported antitumor activity of these compounds [8,14]. The role of TNF- α in tumor suppression was discussed right from its discovery in 1975, but to date its function as an anticancer agent remains problematic [15,16]. TNF- α is also released by CBDA amides 7-9 at a concentration of 1 μ M with further suppression at 10 μ M. These dual results obtained for compounds 7-9 suggest that these compounds may act via several mechanisms, as was the case for CBD and its analogues with a 2,2-dimethylheptyl substituent [17,18]. Compounds 5, 6 exhibited quite a pronounced suppression of TNF- α expression, which together with their low cytotoxicity may mark these compounds as potentially useful anti-inflammatory drugs. The data presented in (Figure 2), shows that except for compound 5 none of the cannabinoids tested altered the expression of proinflammatory cytokine IL-6. A slight decrease (of up to 10%) depending on the concentration tested was observed in the case of unsubstituted THCA amide 5.

Conclusion

Changing the substitution of the amide group of THCA and CBDA derivatives resulted in different and sometimes opposing activities with respect to the expression of TNF- α , with a general lack of effect on IL-6 output. The function of several of these compounds as potential activators of the immune response to viral infections and antitumor activity, as well as their immunosuppressive potential warrants further investigation and may highlight the contributions of such novel cannabinolic acid-based drugs to pharmacology.

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Author Disclosure Statements

The author has no conflicts of interests, and no competing financial interests exist.

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