



Delta FosB and AP-1-mediated transcription modulate cannabinoid CB₁ receptor signaling and desensitization in striatal and limbic brain regions

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ABSTRACT

Repeated Δ⁹-tetrahydrocannabinol (THC) administration produces cannabinoid type 1 receptor (CB₁R) desensitization and downregulation, as well as tolerance to its *in vivo* pharmacological effects. However, the magnitude of CB₁R desensitization varies by brain region, with CB₁R in the striatum and its output nuclei undergoing less desensitization than other regions. A growing body of data indicates that regional differences in CB₁R desensitization are produced, in part, by THC-mediated induction of the stable transcription factor, ΔFosB, and subsequent regulation of CB₁R. The purpose of the present study was to determine whether THC-mediated induction of ΔFosB in the striatum inhibits CB₁R desensitization in the striatum and output nuclei. This hypothesis was tested using bitransgenic mice with inducible expression of ΔFosB or ΔcJun, a dominant negative inhibitor of AP-1-mediated transcription, in specific forebrain regions. Mice were treated repeatedly with escalating doses of THC or vehicle for 6.5 days, and CB₁R-mediated G-protein activation was assessed using CP55,940-stimulated [³⁵S]GTPγS autoradiography. Overexpression of ΔFosB in striatal dopamine type 1 receptor-containing (D₁R) medium spiny neurons (MSNs) attenuated CB₁R desensitization in the substantia nigra, ventral tegmental area (VTA) and amygdala. Expression of ΔcJun in striatal D₁R- and dopamine type 2 receptor (D₂R)-containing MSNs enhanced CB₁R desensitization in the caudate-putamen and attenuated desensitization in the hippocampus and VTA. THC-mediated *in vivo* pharmacological effects were then assessed in ΔcJun-expressing mice. Tolerance to THC-mediated hypomotility was enhanced in ΔcJun-expressing mice. These data reveal that ΔFosB and possibly other AP-1 binding proteins regulate CB₁R signaling and adaptation in the striatum and limbic system.

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Abbreviations: AMYG, amygdala; BSA, bovine serum albumin; CaMKIIα, calcium/calmodulin-dependent protein kinase II; CB₁R, cannabinoid type 1 receptor; CBLM, cerebellum; CP55,940, (–)-cis-3-[2-hydroxy-4-(1,1-dimethylheptyl)phenyl]-trans-4-(3-hydroxypropyl)cyclohexanol; CPU, caudate-putamen; D₁R, dopamine type 1 receptor; D₂R, dopamine type 2 receptor; ERK, extracellular signal-regulated kinase; GP, globus pallidus; GDP, guanosine diphosphate; GTPγS, guanosine 5'-(gamma-thio)triphosphate; HIP, hippocampus; MSN, medium spiny neuron; NAC, nucleus accumbens; PFC, prefrontal cortex; SN, substantia nigra; THC, Δ⁹-tetrahydrocannabinol; VTA, ventral tegmental area.

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1. Introduction

Δ⁹-Tetrahydrocannabinol (THC), the main psychoactive constituent of marijuana, elicits behavioral effects by activating cannabinoid CB₁ receptors (CB₁R) in the central nervous system [1,2]. Repeated THC administration produces CB₁R desensitization, measured as a reduction in receptor-mediated G-protein or effector activity [3], which occurs concomitantly with tolerance to THC-mediated *in vivo* effects [4]. CB₁R desensitization varies in magnitude by brain region depending on the dose and duration of repeated cannabinoid administration. Moreover, the brain regional profile of desensitization corresponds with the development of tolerance to specific cannabinoid-mediated responses [3]. For example, CB₁R desensitization in the dorsal striatum and its output nuclei (i.e., globus pallidus, substantia nigra) is generally lower in

magnitude and requires higher drug doses and/or longer treatment duration than other brain regions [3,5,6]. Studies using post-mortem autoradiography or in vivo imaging in brains from marijuana users compared to non-users have also revealed a smaller decrease in CB₁R levels in the caudate-putamen compared to other regions, including hippocampus [7,8]. These findings are consistent with reports in human marijuana users that showed greater tolerance to the memory impairing effects of THC, which are hippocampal-dependent, than to motor impairment and subjective “high”, which involve striatal circuits [9]. These data suggest the potential functional and translational relevance of regional differences in CB₁R adaptation, but the regulatory mechanisms that underlie these region-specific effects are not known.

We have proposed that regional differences in the interaction of CB₁Rs with specific signaling and regulatory proteins contribute to region-specific differences in CB₁R adaptation [3,10,11] and recently suggested that the induction of transcription factors following repeated THC administration plays a modulatory role in CB₁R desensitization [12]. This idea was based, in part, on the demonstration that an inverse regional relationship exists between THC-mediated CB₁R desensitization and induction of Δ FosB [13]. For example, repeated THC treatment induces Δ FosB expression in the striatum, a region that generally shows lower CB₁R desensitization. Δ FosB belongs to the Fos family of transcription factors, which dimerize with Jun proteins to produce an AP-1 complex that regulates the transcription of target genes [14,15]. Δ FosB, a truncated splice variant of FosB, is stable and accumulates with repeated drug administration [16]. Inducible transgenic overexpression of Δ FosB in dopamine type 1 receptor (D₁R) positive striatal medium spiny neurons (MSNs) showed enhanced rewarding effects of drugs of abuse and natural rewards [17,18]. In contrast, expression of Δ cJun, a dominant negative inhibitor of Δ FosB-mediated transcription, in both D₁R and dopamine type 2 receptor (D₂R) positive MSNs reduced cocaine- and morphine- induced condition place preference [19,20]. Δ FosB produces functional effects by regulating the expression of target genes that include certain receptors and signaling proteins [21], which can affect receptor-mediated signaling. For example, inducible transgenic overexpression of Δ FosB in the nucleus accumbens enhanced mu opioid, but not CB₁, receptor-mediated G-protein activity and inhibition of adenylyl cyclase [22]. Although Δ FosB might not play a pivotal role in CB₁R signaling acutely, it might modulate CB₁R desensitization. The current study tested the hypothesis that repeated THC administration to transgenic mice that inducibly overexpress Δ FosB or its transcriptional inhibitor Δ cJun in the forebrain will display a difference in CB₁R desensitization, as assessed by CP55,940-stimulated [³⁵S]GTP γ S binding. Additionally, tolerance to common pharmacological effects (hypomotility, antinociception, catalepsy, and hypothermia) of THC was quantified in transgenic mice treated repeatedly with vehicle or THC.

2. Materials and methods

2.1. Materials

THC, [(–)-cis-3-[2-hydroxy-4-(1,1-dimethylheptyl)phenyl]-trans-4-(3-hydroxypropyl) cyclohexanol] (CP55,940) and SR141716A were provided by the Drug Supply Program of the National Institute on Drug Abuse (Rockville, MD). [³⁵S]Guanosine 5'-(gamma-thio)triphosphate (GTP γ S; 1250 Ci/mmol), [³H]SR141716A (54 Ci/mmol) and Kodak Biomax MR film were purchased from PerkinElmer Life Sciences (Boston, MA). Bovine serum albumin (BSA), guanosine diphosphate (GDP) and doxycycline were purchased from Sigma-Aldrich (St. Louis, MO). ScintiSafe Econo

1 scintillation fluid was obtained from Research Products International (Mount Prospect, IL). Whatman glass fiber filters were purchased from Brandel (Gaithersburg, MD). All other reagent grade chemicals were obtained from Sigma Chemical Co. or Fisher Scientific.

2.2. Subjects and drug treatments

Subjects were male, bitransgenic *NSE-tTA* \times *TetOp- Δ FosB* mice and *NSE-tTA* \times *TetOp-FLAG- Δ c-Jun* mice with brain-region specific, tetracycline-regulated inducible expression of either Δ FosB or Δ cJun, respectively [19,23]. Δ FosB or Δ cJun expression is controlled by adding doxycycline to the drinking water, which prevents Δ FosB/ Δ cJun expression. Therefore, omission of doxycycline from the drinking water allows Δ FosB/ Δ cJun to be expressed. In mice that overexpress Δ FosB, expression is found in D₁R-positive MSNs in the caudate-putamen and nucleus accumbens, deep layers of cerebral cortex and hippocampus [23]. In mice that overexpress Δ cJun, expression occurs in both D₁R and D₂R positive MSNs in the caudate-putamen and nucleus accumbens, parietal cortex and hippocampus [19]. Δ cJun is a dominant negative functional inhibitor of AP-1-mediated transcription, thus this model provides a strategy to block the effects of THC-induced Δ FosB expression. Mice were housed four to six per cage and maintained on a 12-h light:12-h dark cycle in a temperature-controlled environment (20–22 °C) with food and water available ad libitum. Mice were maintained on drinking water that contained doxycycline (100 μ g/ml) throughout gestation and were either taken off doxycycline for 8 weeks prior to experiments to induce expression of Δ FosB or Δ cJun (Δ FosB-ON, Δ cJun-ON) or maintained on doxycycline (control; Δ FosB-OFF, Δ cJun-OFF). After 8 weeks with/without doxycycline, mice were injected subcutaneously twice daily (08:00 and 16:00) for 6 days with vehicle (1:1:18 solution of ethanol, emulphor and saline) or THC doses that increased every 2 days (10–30–60 mg/kg, injection) [5]. On day 7, mice for autoradiographic studies received only the morning injection of THC or vehicle, and 24 h later brains were collected for CP55,940-stimulated [³⁵S]GTP γ S binding. This THC treatment regimen was employed because it produces CB₁R desensitization throughout the brain, including the striatum, and therefore should reveal whether Δ FosB expression alters CB₁R desensitization. All experiments were performed with the approval of the Institutional Animal Care and Use Committee at Virginia Commonwealth University in accordance with the National Institutes of Health guide for the care and use of Laboratory animals 8th edition.

2.3. Agonist-stimulated [³⁵S]GTP γ S autoradiography

Assays were conducted as previously published [10,24]. Coronal brain sections (20 μ m) were cut on a cryostat maintained at –20 °C, thaw-mounted onto gelatin-coated slides and stored desiccated at 4 °C overnight. Sections were collected at levels that included (1) prefrontal cortex, (2) nucleus accumbens, (3) caudate-putamen, (4) globus pallidus, (5) hippocampus and amygdala (including central, basolateral and basomedial nuclei), (6) VTA, (7) substantia nigra, and (8) cerebellum. Slides were stored desiccated at –80 °C until use. On the day of the assay, slides were brought to room temperature, rinsed in 50 mM Tris–HCl buffer (pH 7.4) with 3 mM MgCl₂, 0.2 mM EGTA and 100 mM NaCl (TME Buffer) for 10 min at 25 °C. Slides were transferred to TME Buffer + 0.5% BSA with 2 mM GDP and 10 mU/ml adenosine deaminase for 15 min at 25 °C. Slides were then incubated in TME Buffer + 0.5% BSA containing 0.04 nM [³⁵S]GTP γ S, 2 mM GDP, 10 mU/ml adenosine deaminase and 3 μ M CP55,940 or vehicle (ethanol) for 2 h at 25 °C. CP55,940 was selected because it does not stimulate [³⁵S]GTP γ S binding in brain sections from CB₁R knockout mice [25], thereby

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