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MOLECULAR AND DEVELOPMENTAL NEUROSCIENCE

The type 1 cannabinoid receptor is highly expressed in embryonic cortical projection neurons and negatively regulates neurite growth *in vitro*

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Abstract

In the rodent and human embryonic brains, the cerebral cortex and hippocampus transiently express high levels of type 1 cannabinoid receptors (CB₁Rs), at a developmental stage when these areas are composed mainly of glutamatergic neurons. However, the precise cellular and subcellular localization of CB₁R expression as well as effects of CB₁R modulation in this cell population remain largely unknown. We report that, starting from embryonic day 12.5, CB₁Rs are strongly expressed in both reelin-expressing Cajal-Retzius cells and newly differentiated postmitotic glutamatergic neurons of the mouse telencephalon. CB₁R protein is localized first to somato-dendritic endosomes and at later developmental stages it localizes mostly to developing axons. In young axons, CB₁Rs are localized both to the axolemma and to large, often multivesicular endosomes. Acute maternal injection of agonist CP-55940 results in the relocation of receptors from axons to somato-dendritic endosomes, indicating the functional competence of embryonic CB₁Rs. The adult phenotype of CB₁R expression is established around postnatal day 5. By using pharmacological and mutational modulation of CB₁R activity in isolated cultured rat hippocampal neurons, we also show that basal activation of CB₁R acts as a negative regulatory signal for dendritogenesis, dendritic and axonal outgrowth, and branching. Together, the overall negative regulatory role in neurite development suggests that embryonic CB₁R signaling may participate in the correct establishment of neuronal connectivity and suggests a possible mechanism for the development of reported glutamatergic dysfunction in the offspring following maternal cannabis consumption.

Introduction

Marijuana is the most widely used illicit drug among women of reproductive age (Fried & Smith, 2001; Ebrahim & Gfroerer, 2003). Δ^9 -tetrahydrocannabinol, the major psychoactive substance of marijuana and hashish, readily crosses the placental barrier (Hutchings *et al.*, 1989) and prenatal marijuana exposure has been shown to result in negative neurobehavioral consequences in the offspring (reviewed by Fried & Smith, 2001). Accordingly, it has been shown that prenatal cannabinoid exposure transiently or permanently alters the function of GABAergic (Garcia-Gil *et al.*, 1999; Berghuis *et al.*, 2007), dopaminergic (Walters & Carr, 1986, 1988; Rodriguez de Fonseca *et al.*, 1991; Bonnin *et al.*, 1995, 1996), serotonergic (Molina-Holgado *et al.*, 1996) and opioidergic (Vela *et al.*, 1998) neuronal systems. Specifically, several aspects of executive function appear to be

negatively associated with *in-utero* cannabis exposure. Clinical and empirical evidence indicates that executive functions are primarily subserved by the prefrontal region of the brain but may also involve other structures such as the hippocampus (Barch, 2005). The predominant neuronal type of cortical structures are glutamatergic neurons and it has been demonstrated that prenatal cannabinoid exposure is also associated with changes in glutamate transmission in the offspring (Mereu *et al.*, 2003; Antonelli *et al.*, 2004, 2005; Castaldo *et al.*, 2007). Although functionally important type 1 cannabinoid receptors (CB₁Rs) are also expressed, albeit at relatively low levels, in adult glutamatergic axon terminals (Katona *et al.*, 2006; Kawamura *et al.*, 2006), a strikingly strong and transient elevation of CB₁R binding and mRNA expression was described in the cerebral cortex at developmental stages when this structure is almost entirely composed of developing glutamatergic neurons (reviewed in Fernandez-Ruiz *et al.*, 1999). However, the resolution of these pioneering studies was not sufficient to identify the CB₁R-expressing cells. Importantly, white matter areas containing the developing axons of glutamatergic neurons also showed strong and transient expression of functional CB₁R, first detected by binding of labeled cannabinoid

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ligands and by agonist-induced GTP γ S binding (Fernandez-Ruiz *et al.*, 1999). Axonal expression was recently confirmed by immunohistochemistry (Berghuis *et al.*, 2007) but precise spatio-temporal patterns of CB₁R expression in young telencephalic glutamatergic neurons remain largely unknown. Subcellular distribution, which is strongly related to CB₁R activation levels (Leterrier *et al.*, 2004, 2006), is also unknown for CB₁Rs as well as for the majority of embryonic G-protein-coupled receptors.

In the present study, using optical and electron microscopy, we first describe, at high resolution, CB₁R localization in the cerebral cortex and hippocampus from embryonic day (E)12.5 to postnatal day (P)5. Next, by using pharmacological and mutational modulation of CB₁R activity in isolated cultured hippocampal neurons, we show that the modification of CB₁R activation significantly modulates outgrowth of dendrites and axons. Our results suggest that embryonic CB₁R signaling participates in the correct establishment of neuronal connectivity and propose a possible mechanism for the development of reported glutamatergic dysfunction in the offspring following maternal cannabis consumption. A part of these results was previously reported as a conference poster (Vitalis *et al.*, 2004).

Materials and methods

Animals

Animal procedures were conducted in strict compliance with approved institutional protocols and in accordance with the provisions for animal care and use described in the European Communities council directive of November 24, 1986 (86/609/EEC). Sprague–Dawley rats (Janvier) were used for dissociated cell culture experiments. Mice (OF1 or C57/Bl6 strains) were purchased from Charles River. Animals expressing green fluorescent protein (GFP) under the control of the 5-HT_{3A} promoter (5-HT_{3A}:GFP) were purchased from The Gene Expression Nervous System Atlas Project (Rockefeller University, New York, NY, USA) and were obtained by homologous recombination of a bacterial artificial chromosome containing the 5-HT_{3A} gene that will be fully described elsewhere (Ksenija Vucurovic and Tania Vitalis, unpublished observation). Genitors were phenotyped using UV illumination. The day of vaginal plug detection was counted as E0.5 for the mice and E0 for the rats.

Reagents

Mouse monoclonal antibodies raised against reelin (G10, 1 : 500, a kind gift of Dr Goffinet, Université Catholique de Louvain), neuron-specific beta III tubulin (Tuj-1) (1 : 2000, Berkeley Antibody Company, Richmond, CA, USA), the 67 kDa isoform of glutamic acid decarboxylase (1 : 5000, MAB5406, Chemicon International), microtubule-associated protein 2 (clone HM-2, Sigma, St Louis, MO, USA) and the FLAG epitope (clone M1, Sigma) were used. Rabbit polyclonal antibody raised against the C-terminal portion of the CB₁ receptor (C-Ter; 1 : 200 for embryonic tissue and 1 : 1000 for culture experiments) was produced by Eurogentec (Seraing, Belgium) and has been characterized previously (Leterrier *et al.*, 2004). Alexa-Fluor-conjugated (1 : 400–1 : 600) secondary antibodies were from Invitrogen. DsRed2 encoding plasmid was from Clontech (Mountain View, CA, USA). Our validated CB₁-enhanced green fluorescent protein (EGFP) and FLAG-CB₁-EGFP constructs were used as previously described (Leterrier *et al.*, 2004, 2006). Mutants displaying either enhanced agonist affinity (T210I) or enhanced inverse agonist affinity (T210A) relative to the wild-type constructs (D'Antona *et al.*, 2006) were obtained using Quikchange mutagenesis (Stratagene) of

the FLAG-CB₁-EGFP plasmid. The following chemicals were used for pharmacological treatments: CB₁R agonists CP-55940 (Tocris, Bristol, UK) and (*R*)-(-)-[2,3-dihydro-5-methyl-3-(4-morpholinylmethyl)pyrrolo[1,2,3-de]-1,4-benzoxazin-6-yl]-1-naphthalenyl-methanone mesylate [WIN55,212-2 (WIN); Sigma], and CB₁R inverse agonist 1-(2,4-dichlorophenyl)-5-(4-iodophenyl)-4-methyl-*N*-4-morpholinyl-1*H*-pyrazole-3-carboxamide (AM281, Tocris). All culture media and additives were from Invitrogen.

Acute treatments of pregnant dams

Pregnant dams bearing E16.5 embryos received a single injection of the CB₁R agonist CP-55940 (0.7 mg/kg, *i.p.*; Sigma) or vehicle 2 and/or 12 h before killing. Embryos were dissected out in cold phosphate buffer (0.1 M) and were immersed overnight in 4% paraformaldehyde in 0.1 M phosphate buffer, pH 7.4.

Radioactive in-situ hybridization

The CB₁R cRNA probe (a kind gift of Dr Beat Lutz, Johannes Gutenberg University, Mainz) corresponded to the full-length protein. The plasmid was linearized with BamHI for antisense RNA synthesis by T7 polymerase and with EcoRI for sense RNA synthesis by T3 polymerase. The transcription was carried out using the Promega kit and probes were labeled with ³⁵S-UTP (> 1000 Ci/mmol; Amersham). Hybridization was performed on fresh frozen brain sections (15 μ m thick) as described in Fontaine & Changeux (1989). Slides were dipped in photographic emulsion (NTB2, Kodak). After 5–10 days the emulsion was developed and sections were Nissl-stained (0.25% thionin solution).

Immunohistochemistry on brain sections

In order to analyse expression of the CB₁R protein, brains were obtained at embryonic (E12.5, E14.5 and E16.5) and postnatal (P0 and P5) stages. Embryos were placed overnight in 4% paraformaldehyde in 0.1 M phosphate buffer, pH 7.4, then cryoprotected, embedded into gelatine (7%)/sucrose (10%), frozen into isopentane (–40°C) and stored at –80°C until processing. Postnatal animals were deeply anesthetized with intraperitoneal pentobarbital injection (150 mg/kg body weight) and perfused transcardially with 4% paraformaldehyde. Brains were processed as described above. Blocks were sectioned coronally with a cryostat (20 μ m thick). Sections were incubated overnight at 4°C with the primary antibodies, washed and incubated with Alexa 568-conjugated goat anti-rabbit and/or Alexa 488-conjugated anti-mouse antibodies (1 : 200). Sections were finally mounted in Vectashield (Vector, CA, USA) containing 4'6-diamidino-2-phenylindole 2 HCl (DAPI) (1 : 1000).

After fixation, for the analysis of the effects of chronic treatments, brains were embedded in 0.1 M phosphate buffer/agarose (3%), pH 7.4, and sectioned with a vibratome (45 μ m). Sections were permeabilized and incubated overnight at 4°C with the primary antibodies, then washed and incubated with secondary antibodies. Sections were washed, nuclei were labeled with bisbenzimidazole (1 μ g/mL) and sections were finally mounted in Vectashield (Vector).

Ultrastructural pre-embedding immunocytochemistry

Embryonic mouse brains obtained from untreated or treated dams (see above) were quickly dissected as described above and immediately

fixed by immersion in either 4% paraformaldehyde with 0.1% glutaraldehyde in 0.1 M phosphate buffer or 2% paraformaldehyde plus 1% glutaraldehyde, first for 1 h at room temperature (23°C) and then overnight at 4°C. Brains were then embedded in 3% agarose, cut with a vibratome and 150- μ m-thick coronal sections were collected. Two aldehyde quenching steps in 0.1% sodium borohydrate and 0.1 M glycine preceded the blocking step of non-specific sites in 5% normal goat serum + 5% bovine serum albumin. An overnight incubation at room temperature was performed in anti-CB₁R (1 : 500 in phosphate-buffered saline). For immunoperoxidase procedures, a biotinylated anti-rabbit IgG (Vector) was applied as secondary antibody (1 : 200 in phosphate-buffered saline, 2 h) and the ABC peroxidase complex (Vectastain Elite, Vector) was used for amplification. Revelation was performed with 0.05% diaminobenzidin as the chromogen. For immunogold labelings, a 5 h incubation in ultra-small gold conjugate F(ab')₂ fragments of goat anti-rabbit IgG (1 : 200; Aurion, Netherlands) was followed by extensive washing, 10 min postfixation in 2% glutaraldehyde and a silver enhancement reaction (R-Gent SE-ME kit, Aurion). Some sections were submitted to a gold-toning procedure to protect silver deposits from osmium displacement. After OsO₄ postfixation (2% for immunoperoxidase labeling and 1% for gold revelation) and 2% uranyl acetate *en-bloc* staining, immunostained sections were dehydrated in graded acetone and finally embedded in Durcupan (Fluka, Switzerland) resin. Observations were made with a Phillips CM120 electron microscope and digitalized with a Morada camera.

Hippocampal neuronal cultures

Hippocampal neuronal cultures were performed essentially as described by Goslin & Banker (1989) with some modifications (Jolimay *et al.*, 2000). Briefly, hippocampi of rat embryos were dissected at E17. After trypsinization, dissociation was achieved with a fire-polished Pasteur pipette. Cells were counted and plated on poly-D-lysine-coated (Sigma) 15-mm-diameter glass coverslips at a density of 400–500 cells/mm². The plating medium was Neurobasal supplemented with 2% B27 and containing Glutamax I (0.5 mM) and penicillin G (10 U/mL)/streptomycin (10 g/mL). At 4 h after plating, the coverslips were transferred into Petri dishes containing supplemented Neurobasal medium that was conditioned for 24 h on an 80% confluent glial layer. Neurons were transfected after 3 days *in vitro* (DIV) using Lipofectamine 2000 (Invitrogen), following the manufacturer's instructions. They were treated and processed at 1 DIV after transfection. For pharmacological treatments, ligands dissolved in dimethylsulfoxide were added directly to the culture medium. The highest final concentration reached was 0.2% dimethylsulfoxide; control experiments with up to 0.5% dimethylsulfoxide have shown the absence of effects on neuronal morphology and on the cellular distribution of CB₁Rs (data not shown).

Immunocytochemistry on cultured neurons

For immunocytochemistry, neurons were briefly rinsed with Dulbecco's phosphate-buffered saline (DPBS) (Invitrogen) and fixed in DPBS containing 4% paraformaldehyde and 4% sucrose. After permeabilization with 5 min incubation in DPBS containing 0.1% Triton X-100 and blocking for 30 min in antibody buffer (DPBS supplemented with 2% bovine serum albumin and 3% normal goat serum), neurons were incubated with primary antibodies diluted 1 : 500–1 : 5000 in antibody buffer for 1–2 h at room temperature. After DPBS rinses, neurons were labeled with secondary antibodies

diluted to 1 : 400 in antibody buffer for 1 h at room temperature. Coverslips were finally mounted in Mowiol (Calbiochem, La Jolla, CA, USA) eventually containing Hoechst 33342 diluted to 1 : 1000.

Microscopy and morphometry

Wide-field images were taken on a Leica DM-R microscope using a dry 20 \times , NA 0.7 objective and QImaging (Burnaby, British Columbia, Canada) QICAM. In all cases, emission and excitation filters proper to each fluorophore were used sequentially and the absence of cross-talk between different channels was checked with selectively labeled preparations. Neurites were outlined and measured using an assisted semiautomatic method (NeuronJ) (Meijering *et al.*, 2004). For neurons at DIV3, axon, primary and secondary dendrites were outlined and measured for their number and length. For neurons at DIV10, IMAGEJ (<http://rsb.info.nih.gov/ij/index.html>) was used to delete the somatodendritic area and the function 'skeletonize' was used to reduce all axonal arborescence to a single pixel network. Pixels were then counted to estimate the extent of the axonal network. All measurements were analysed and plotted using the PRISM software (GraphPad, San Diego, CA, USA). Significance of differences between various conditions was calculated using one-way ANOVA with Newman-Keuls post-tests for computing p estimates.

Results

CB₁R expression in the developing cortico-hippocampal formation

First, by using *in-situ* hybridization, confocal microscopy and electron microscopy, we determined the precise cellular distribution of CB₁Rs in the developing cerebral cortex and hippocampus from E12.5 until birth. Between E12.5 and E13.5 CB₁R mRNA and protein were expressed by a large proportion of newly differentiated neurons of the cortical preplate as indicated by the co-expression of CB₁R with the neuron-specific and early-differentiation marker class III beta-tubulin (Tuj-1) (Menezes & Luskin, 1994) (Fig. 1A1–A3 and C). We have not observed significant CB₁R protein or mRNA expression in the ventricular zone. In the preplate, CB₁R was expressed by a subpopulation of reelin-expressing Cajal-Retzius pioneer neurons (G10-positive cells in Fig. 1D1–D3). Both soma and growing neuritic processes were decorated by a punctuated intracellular CB₁R immunolabeling (Fig. 1C, see also Supporting information, Fig. S1 and Video S1). Electron microscopic analysis showed that this punctuate labeling corresponded to CB₁Rs that were exclusively associated with the external membrane of large electrolucent endosomes (200–400 nm) (Fig. 1E–H).

By E14.5, CB₁R mRNA was still strongly expressed in the entire cortical plate (data not shown). Immunohistochemistry showed that the major proportion of CB₁R protein was gradually reallocated into growing axons of maturing projection neurons (Fig. 2A). Thus, in the more mature lateral parts of cerebral cortex, CB₁R immunolabeling was mainly localized in the intermediate zone, where Tuj-1-positive (Fig. 2B1–B3) corticofugal axons are located (Del Río *et al.*, 2000). At the same time, in the less mature medial parts of the cortical region, which will later give rise to the hippocampus, punctuate CB₁R immunolabeling was localized to somatic endosomes of newly differentiated neurons that did not yet show axonal processes. Thus, electron microscopy indicated that, in the medial part of the cortex, CB₁R immunolabeling was associated with large and complex endosomes in the soma (Fig. 2C and D), similar to the distribution observed at E12.5, as shown in Fig. 1. In the axonal compartment,

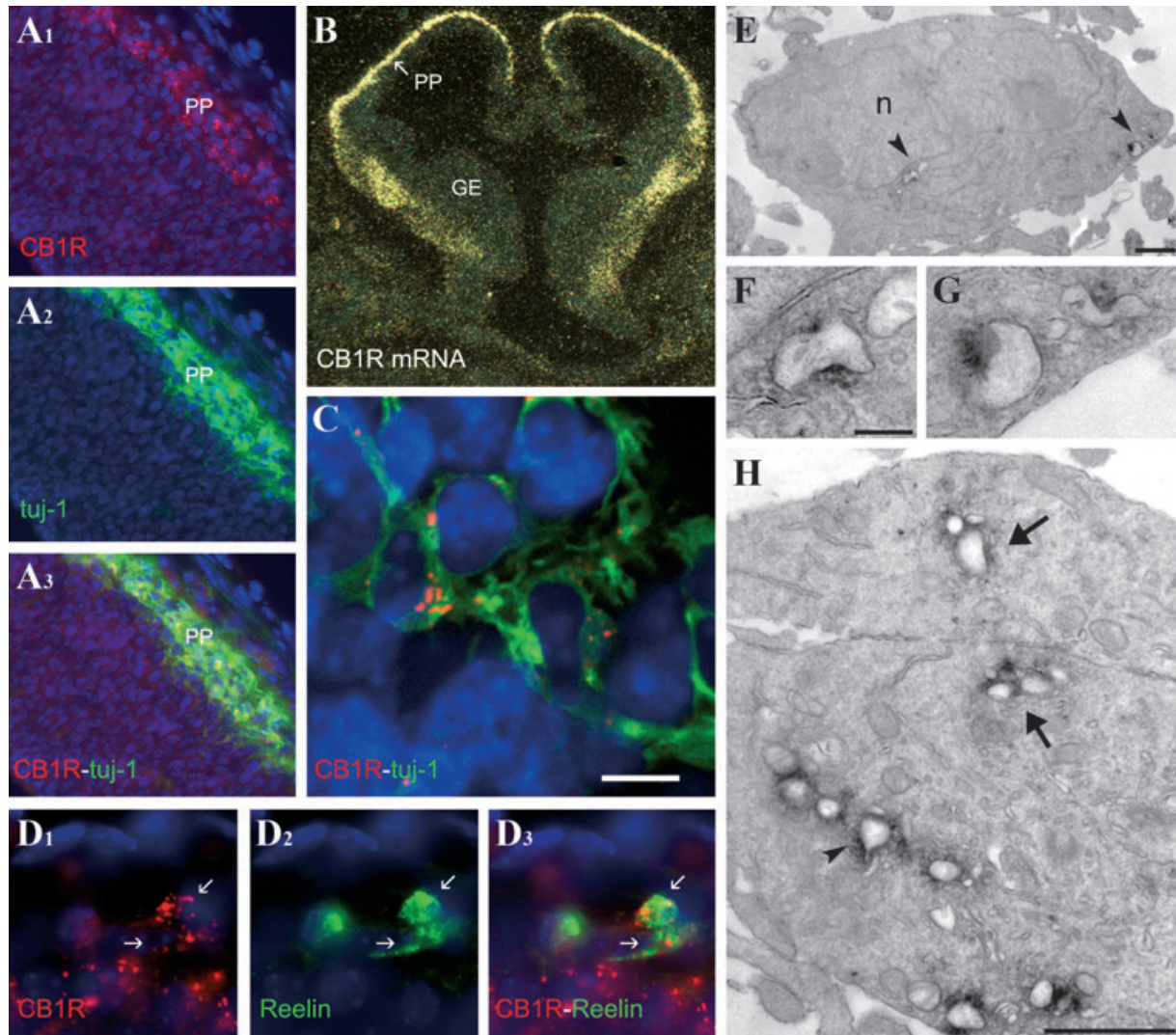


FIG. 1. Localization of CB₁R in cortical neurons at E12.5. (A1–A3) Immunolabeling of CB₁R (A1; red) in Tuj-1-positive neurons (A2; green) of the preplate (PP). Sections are counterstained with DAPI (blue). (B) *In-situ* hybridization showing CB₁R mRNA in the telencephalon. Note the absence of CB₁R mRNA in the ganglionic eminence (GE). (C) Confocal view showing punctiform CB₁R labeling in the somata and neurites of preplate cells. See also supporting Video S1. (D1–D3) Co-localization of CB₁R (red) with reelin (green) in the cortical marginal zone (arrows). (E–H) Ultrastructural CB₁R immunoperoxidase labeling of preplate cortical neurons. Please note that the dissociated aspect of neurons is an artifact of the electron microscopic procedure, due to the fragility of embryonic brain tissue at this early stage. (E) Neuronal soma with an indented nucleus (n). Arrowheads point to three CB₁R-labeled endosomes that are enlarged in F and G. (F and G) Note that CB₁R immunoprecipitate is localized on the cytoplasmic side of the limiting membrane of large endosomes. Their lumen is electrolucent and contains membranous or more frequently ‘dusty’ debris. (H) Soma profiles of two contiguous preplate neurons containing numerous CB₁R-positive large endosomes. These are rather circular in shape and some appear to correspond to fusing endosomes (arrows). The arrowhead indicates an endosome with a tubular extension. Bar in C = 150 μ m for A1–A3, 1.25 mm for B, 20 μ m for D1–D3 and 8 μ m for C. Bars in E and H = 1 μ m. Bar in F (same for G) = 0.2 μ m.

CB₁R immunolabeling was associated with the plasma membrane (Fig. 2E and G), accompanied by labeling of the outer membrane of small clear lumen vesicles (Fig. 2F, H and I). However, as the diaminobenzidine precipitate used to reveal immunolabeling in this experiment may also result in artifactual labeling of membranes, we have also performed immunogold detection of CB₁R. The resulting localization of gold beads confirmed that CB₁Rs were localized to small clear lumen vesicles and were also more or less regularly disposed along the axonal plasma membrane (Fig. 2E). Finally, between E12.5 and E14.5, CB₁R immunolabeling was absent from the ganglionic eminences where most cortical GABAergic neurons are generated (Fig. 2A).

By E16.5, CB₁R mRNA was strongly expressed in the entire cortical plate and hippocampus (Fig. 3A), which are mainly

composed of the perikarya of glutamatergic projection neurons at this stage (Rakic, 2006). Interestingly, the localization of CB₁R protein was almost completely segregated from the localization of CB₁R mRNA, with the protein being translocated to developing fiber tracts such as the fornix, anterior commissure, hippocampal commissure, fimbria hippocampi, corpus callosum and individual cortico-fugal axons outside the above fiber tracts (Fig. 3B and C). At this stage, in the somato-dendritic compartment, the remaining CB₁R immunolabeling was associated with the external surface of large endosomes similar to those described at earlier stages. In the axonal compartment (Fig. 3E), CB₁R labeling was associated with intra-axonal endosomal organelles including multivesicular bodies (Fig. 3G and H) and with the surface of the axons (Fig. 3F). At this stage, CB₁R immunolabeling was not detected in the

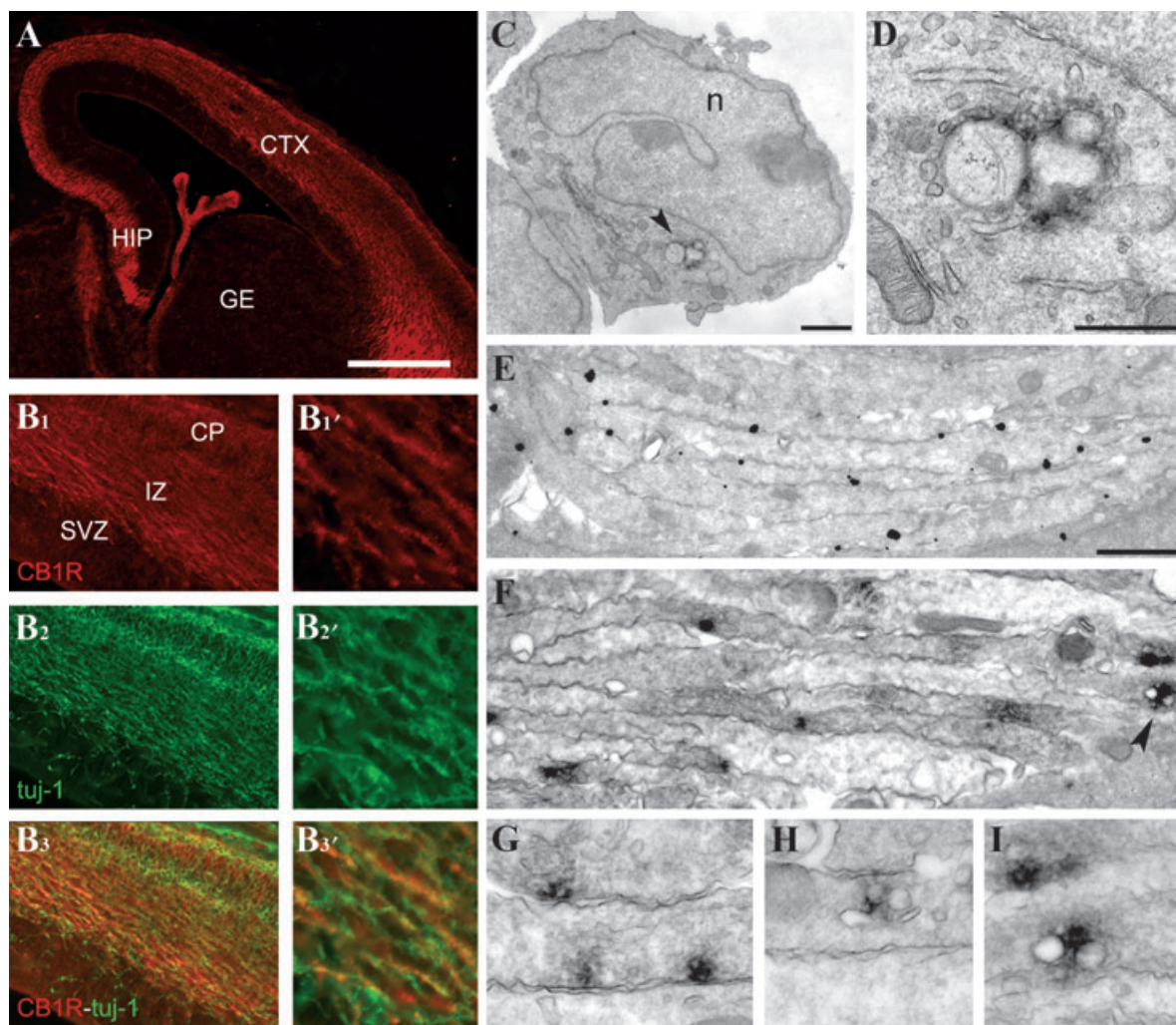


FIG. 2. Localization of CB₁Rs on projection neurons at E14.5. (A) Coronal section showing immunolabeling of CB₁R in the cortex (CTX) and hippocampus (HIP). Note the lack of immunolabeling in the ganglionic eminence (GE). (B1–B3) Co-localization of CB₁R (red) with Tuj-1 (green) in corticofugal axons in the intermediate zone (IZ). (B1'–B3') Higher magnification of B1–B3. Note that all CB₁R-positive fibers express Tuj-1. (C and D) Subcellular distribution of CB₁R immunolabeling in neuronal somata. (C) Representative example of a CB₁R-positive cortical plate neuron with a deeply indented nucleus (n). CB₁R labels a group of three connected round endosomes (arrowhead), further enlarged in D. Note that CB₁R precipitate remains restricted to the endosomal limiting membrane. (E–I) Subcellular distribution of CB₁R immunoreactivity in corticofugal axons. (E) The majority of CB₁R-labeling gold beads are attached to the axolemma. (F) Scattered distribution of CB₁R immunoprecipitate, either localized at the axolemma or, less frequently, within the axoplasm where it is usually found close to the outer membrane of small clear lumen vesicles (arrowhead, enlarged view in I). (G) High magnification of axolemma labeling. (H and I) CB₁R-positive round vesicles. Bar in A = 600 μ m for A, 250 μ m for B1–B3 and 80 μ m for B1'–B3'. Bars in C and E (same for F) = 1 μ m. Bar in D = 0.5 μ m and is also for G–I. CP, cortical plate; SVZ, subventricular zone.

subventricular zone where most GABAergic neurons are tangentially migrating (data not shown).

In order to investigate if CB₁R proteins abundantly expressed in embryonic fiber tracts are functionally competent, we took advantage of the fact that, in neurons, agonist-mediated activation of functional G-protein-coupled receptors and more specifically of CB₁R receptors ultimately results in internalization and retrograde somato-dendritic translocation (Coutts *et al.*, 2001; Letierrier *et al.*, 2004, 2006). Agonist-induced internalization is an event far downstream of G-protein-coupled receptor activation, which requires pharmacologically accessible receptors in a functionally relevant context allowing an appropriate ligand-mediated conformation change. Subsequent phosphorylation of specific threonine and serine residues leads to uncoupling of G-proteins and is followed by interaction with a cascade of scaffolding proteins (such as arrestins and chlatrin) that ultimately results in endocytosis and intracellular translocation (Hanyaloglu &

von Zastrow, 2008). Thus, agonist-induced internalization is a strong indication that the investigated G-protein-coupled receptor is indeed functionally competent. Cannabinoid analogs such as Δ^9 -tetrahydrocannabinol readily cross the placental barrier (Hutchings *et al.*, 1989) and thus we expected that systemic treatment of pregnant dams with cannabinoid agonist would result in internalization of functional CB₁Rs in the embryonic brain. Indeed, subacute intraperitoneal treatment of pregnant dams (two injections at 12 and 2 h before killing) at E16.5 with the CB₁R agonist CP-55940 resulted in a striking redistribution of CB₁Rs toward the somato-dendritic compartments of projection neurons. Thus, in CP-55940-treated animals, only residual CB₁R immunostaining could be observed in the hippocampal fimbria, and hippocampal projection neuron perikarya were decorated by a punctuated CB₁R immunolabeling (Fig. 3D). We have observed a similar redistribution in the neocortex (data not shown). A single injection of CP-55940 (2 h before killing) led to a

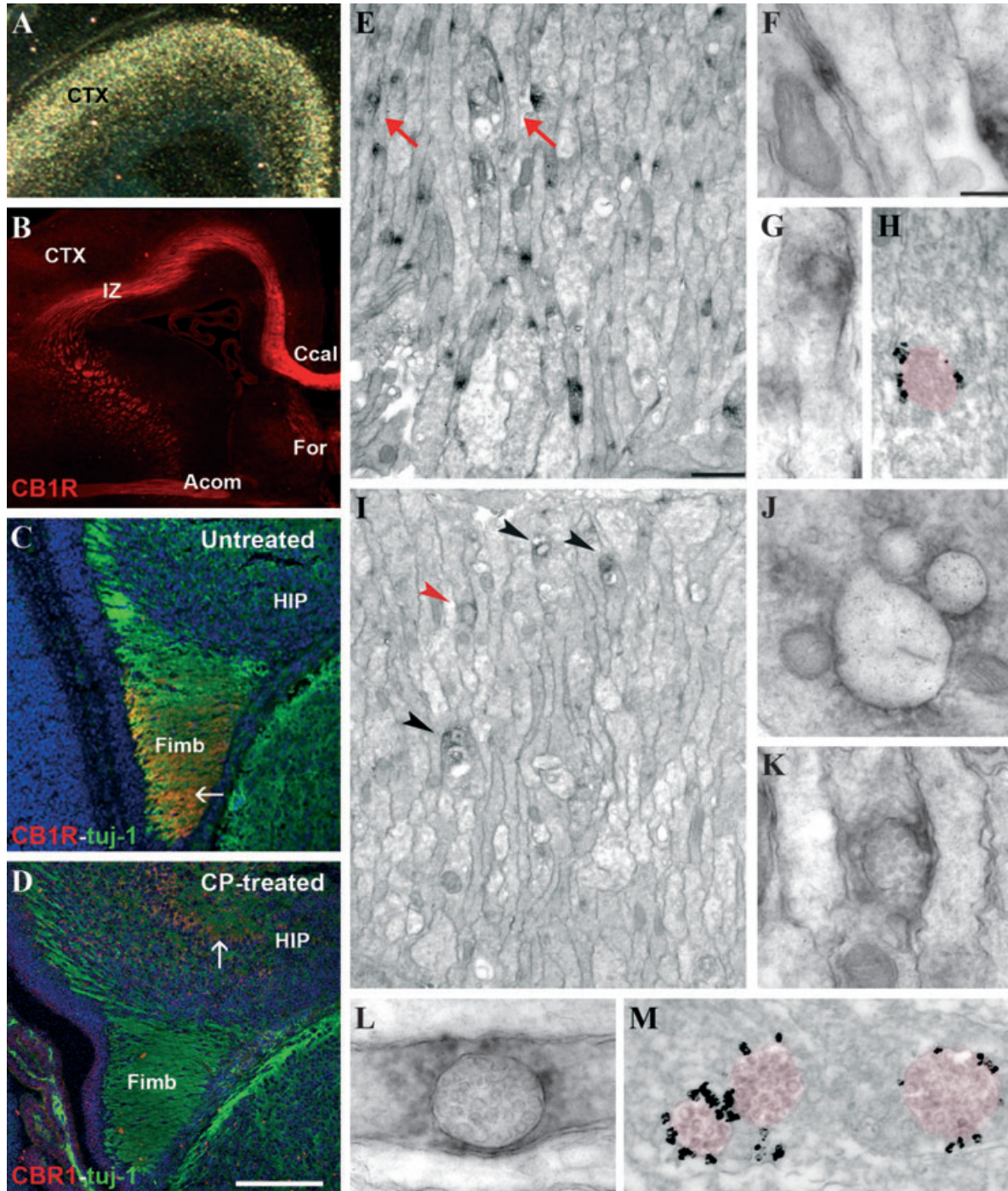


FIG. 3. Localization of CB₁R_s at E16.5 is modified by treatment with CB₁R agonist. (A) *In-situ* hybridization showing a high level of CB₁R mRNA expression in the cortex (CTX). (B) Coronal section showing CB₁R immunolabeling mainly restricted to axonal tracts, such as the intermediate zone (IZ), corpus callosum (Ccal), anterior commissure (Acom) and fornix (For). Maternal treatment with the CB₁R agonist CP-55940 (CP) (D) induces modification in the location of CB₁R immunolabeling (red) in comparison with control condition (C). Axonal tracts are labeled with Tuj-1 (green) and sections are counterstained with DAPI (blue). (C) In the control condition, CB₁R (red) is exclusively located in the fimbria (Fimb). (D) Following CP-55940 treatment CB₁R is translocated to the somatic compartment of hippocampal neurons. Ultrastructural CB₁R labeling of corticofugal axonal tracts in control embryonic brain (E-H) vs. CB₁R agonist-treated embryos (I-M). (E) Corticofugal axon in a control embryo. Scattered CB₁R precipitates label either the axonal membrane or intra-axonal endosomal organelles. Red arrows point to two areas enlarged in F and G. (F) Axolemma labeling. (G) CB₁R labels the limiting membrane of a round endosome with an electrolucent lumen. (H) Immunogold labeling of the limiting membrane of an intra-axonal multivesicular endosome (light red mask for clarity), the other type of CB₁R-positive endosome found along axons. (I-M) Corticofugal axon labeling following CP-55940 administration. (I) Compared with control, labeling of the axolemma is reduced and immunopositivity is mostly found on organelles within the axonal lumen (arrowheads), most of them being multivesicular endosomes. These are more numerous and usually larger than in control axons. (J) Connected endosomes suggesting fusion events from the soma of a neocortical neuron. (K) Enlarged view of the multivesicular endosome indicated with the red arrowhead in I. (L) Another immunoperoxidase-labeled multivesicular body whose diameter nearly equals that of the axon. (M) Three multivesicular bodies aligned in a single axonal shaft and surrounded by gold beads. These results are representative for two independent experiments. Bar in D = 625 μ m for A, 700 μ m for B, 300 μ m for C and D. Bar in E (also for I) = 1 μ m. Bar in F (also for G, H and J-M) = 0.2 μ m. HIP, hippocampus.

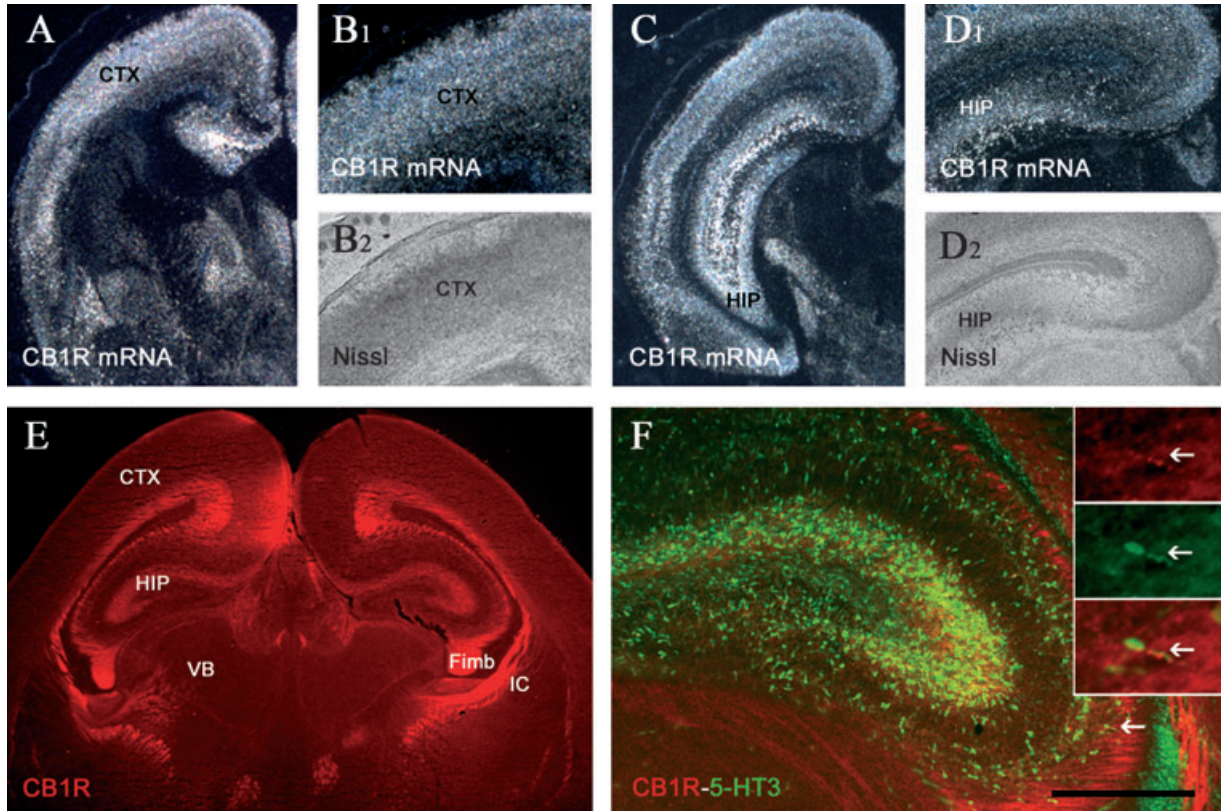


FIG. 4. Localization of CB₁R at birth (P0). (A) *In-situ* hybridization showing a high level of CB₁R expression in coronal sections of the rostral telencephalon. (B1 and B2) High-resolution images of the somatosensory cortex (CTX) shown in A. (B1) Expression of the CB₁R mRNA. (B2) Bright-field view of the same region. (C) *In-situ* hybridization showing high level of CB₁R expression in the caudal telencephalon. (D1 and D2) High resolution of the hippocampal formation shown in C. (D1) Expression of the CB₁R mRNA. (D2) Bright-field view of the same region. Note that in the hippocampal formation both diffuse and spotty cellular expression could be observed (D1). (E and F) CB₁R immunolabeling of an intermediate coronal section. Note that CB₁R is strongly expressed in the hippocampus (HIP) and in numerous axonal tracts, such as the fimbria (Fimb) and cortical axons passing through the internal capsule (IC). (F) Coronal section of the hippocampus of a mouse expressing green fluorescent protein (GFP) under the control of the 5-HT_{3A} promoter stained for CB₁R (red). Boxes show neurons expressing both GFP and CB₁R located in the CA3 region (F, arrow). Bar: A–C, 3 mm; B1, B2, D1 and D2, 1.5 mm; E, 1.8 mm; F, 450 μ m; insets in F, 70 μ m; VB; ventrobasal thalamic nucleus.

partial redistribution of the receptor (data not shown). At the electron microscopic level the distribution of CB₁R was also strikingly different in CP-55940-treated and control (vehicle-injected) embryos. In contrast to control animals (Fig. 3E and H), CP-55940-treated embryos showed a lack of CB₁R immunolabeling on the axonal surface and a remarkable increase in the number of CB₁R-positive multivesicular bodies in the axonal tracts that occupied the full diameter of the relatively large axons (Fig. 3I and K–M). In addition, numerous large CB₁R-positive endosomes with a polycyclic profile were seen in the somato-dendritic compartments, suggesting frequent fusion events between these labeled organelles (Fig. 3J).

At birth (P0), both glutamatergic and GABAergic cell populations expressed CB₁R (Fig. 4), as shown by using a bacterial artificial chromosome mouse line in which the GFP is expressed under the control of the 5-HT_{3A} promoter (Fig. 4F). In these mice, a large proportion of GFP-positive interneurons expressed CB₁R. This cell population corresponds to young GABAergic neurons that will later express cholecystokinin and vasoactive intestinal peptide (Vucurovic and Vitalis, unpublished results). In adult rat neocortex and hippocampus, CB₁R is preferentially expressed in this cell population (Katona *et al.*, 1999, 2000; Morales *et al.*, 2004; Bodor *et al.*, 2005; Hill *et al.*, 2007). In addition, similarly to what was observed at E16.5, CB₁R immunolabeling was also observed in numerous axonal tracts at P0. The mature location of CB₁R became predominant around P5 as previously reported in the rat (in supporting Fig. S1) (Morozov & Freund, 2003). From this stage, strong CB₁R mRNA expression was

mostly restricted to interneurons located in superficial and deep neocortical layers and in the hippocampus (in supporting Fig. S1), with CB₁R immunolabeling localized to a dense local axonal network but no CB₁R immunolabeling detected in axonal tracts.

In conclusion, CB₁Rs are expressed very early by the majority of postmitotic projection neurons in the hippocampus and cerebral cortex. CB₁Rs are gradually translocated in the growing axons of these projection neurons. In these axons, CB₁Rs are localized on the axonal plasma membrane but an important proportion of CB₁Rs is localized in endosomal structures. Axonal CB₁Rs in the embryonic brain display an adequate physiological response to agonist stimulation (internalization followed by somato-dendritic translocation, as previously reported for CB₁R in cultured hippocampal neurons by Letierrier *et al.*, 2006), suggesting that embryonic CB₁Rs are functionally competent. Importantly, these results also show that maternal cannabis consumption may lead to pharmacological activation of CB₁Rs in the embryonic brain.

Pharmacological or mutational activation or blockade of CB₁R activity exerts opposite effects on the morphology of cultured hippocampal neurons

Cerebral CB₁R function has been mostly interpreted in the context of the regulation of synaptic transmission (Freund *et al.*, 2003). However, the presence of functionally competent CB₁Rs on newly differentiated

neurons and in growing axons indicates that CB₁Rs may participate in the regulation of neuronal development, as was also suggested by recent reports (Harkany *et al.*, 2007, 2008; Watson *et al.*, 2008). In order to precisely quantify the effects of CB₁R activation, we first used low-density cultures of neurons isolated from the embryonic hippocampus, a brain structure expressing high levels of CB₁Rs. Rat hippocampal cultures represent a well-characterized model system that was successfully used in numerous studies to investigate neuronal development (Dotti *et al.*, 1988; Bradke & Dotti, 2000). In this system, at DIV4, neurites are already differentiated into axons and dendrites, which have acquired their morphological and structural features (Stage 3; Craig & Banker, 1994).

In order to prepare primary neuronal cultures with a relatively high proportion of glutamatergic neurons, we isolated neurons from the hippocampus of rats at E17 (Carnegie stage 22, corresponding to E15.5 in mice). At this age, tangentially migrating GABAergic interneurons have not yet reached the primordium of hippocampus in high numbers (Jimenez *et al.*, 2002). After 4 days in culture (DIV4), simultaneous immunodetection of CB₁R and the 67 kDa isoform of glutamic acid decarboxylase, a marker of GABAergic interneurons, was performed to investigate the expression of endogenous CB₁Rs. We found that the majority (74.4 ± 0.3%) of the neurons expressed CB₁R and, among this population, only a minority (4.6 ± 0.7%) were GABAergic, showing that CB₁Rs are mostly expressed by glutamatergic neurons at this early stage (Fig. 5). Endogenous CB₁Rs were located in vesicles in the somato-dendritic compartment as well as in axons and in axonal growth cones. This subcellular distribution was strongly reminiscent of the distribution previously shown in older neuronal cultures, where CB₁Rs are mostly expressed by GABAergic neurons (Irving *et al.*, 2000; Coutts *et al.*, 2001; Letierrier *et al.*, 2004, 2006; McDonald *et al.*, 2007).

Next, cultured hippocampal neurons were transfected with CB₁R-EGFP. Subneuronal localization of the transfected CB₁R mirrored that of the native CB₁R and the transfected neurons were also predominantly of the glutamatergic phenotype (Benson *et al.*, 1994) (data not shown). We then analysed the modifications of the axonal and dendritic morphology of individual neurons co-transfected with the structural marker DsRed2 and with CB₁R-EGFP. Neuronal CB₁Rs have been shown to display significant spontaneous activation, which is partly mediated by endocannabinoids synthesized in the same neuron (Turu *et al.*, 2007). This offers the opportunity to both enhance and inhibit steady-state CB₁R activation. Neurons were

treated with the vehicle, the agonist WIN (200 nM) or the antagonist/inverse agonist AM281 (200 nM) for 24 h. We found that treatment with AM281 (200 nM) dramatically increased the total number of primary dendrites (to 160.8 ± 10.5%) and the total dendritic length (to 205.3 ± 23%; Fig. 6A and B). These effects were concentration-dependent with half-maximal effective concentration (EC₅₀) at 0.8 ± 0.3 nM of AM281 (Fig. 6C). Applications of WIN induced the opposite effect, without reaching significance threshold (Fig. 6B). Co-application of WIN abolished AM281 effects, indicating that the effects of AM281 were mediated by CB₁Rs (Fig. 6B). The axonal compartment did not show significant modification upon overnight pharmacological treatments. Interestingly, however, when WIN treatment (200 nM) was maintained between DIV3 and DIV10, the total arbor length of the axons was greatly decreased (to 55.4 ± 5.0%) (Fig. 6D). At this time point, AM281-treated neurons still showed a significantly larger number of dendrites (Fig. 6E). We obtained similar effects when the overnight incubations were realized using incubation media that had not been conditioned with glial cells, showing that endocannabinoids potentially present in the conditioned medium have no significant effect on neuronal development (data not shown). Finally, neurons transfected only with the structural marker DsRed2 only responded slightly to pharmacological treatments, demonstrating not only that the morphological effects were specifically mediated by the transfected CB₁Rs but also that endogenous CB₁Rs, present in the majority (75%) of the neurons, had a limited influence on neuronal morphology in our experimental conditions (Fig. 6B).

The above results indicate that pharmacological activation and inhibition of transfected CB₁Rs significantly modify neuronal structure. However, pharmacological treatments act indiscriminately on every cell in the culture and thus the measured alterations may result from an indirect effect, mediated for example by non-neuronal cells, such as astrocytes, which are present in the culture in low numbers. Thus, we have confirmed the above results by co-transfecting sister cultures of DIV3 hippocampal neurons with DsRed2 and either CB₁R-EGFP or one of the two point mutants of CB₁R that were recently reported by the group of D. Kendall to display either enhanced (T210I) or reduced (T210A) constitutive activity (D'Antona *et al.*, 2006) as compared with non-mutated or wild-type CB₁Rs. We found that expression of T210I for 24 h (Fig. 7B) significantly reduced the number and length of primary and secondary dendrites (down to 67.8 ± 8.8% and 67.0 ± 10.0%, respectively, Fig. 7D), as compared

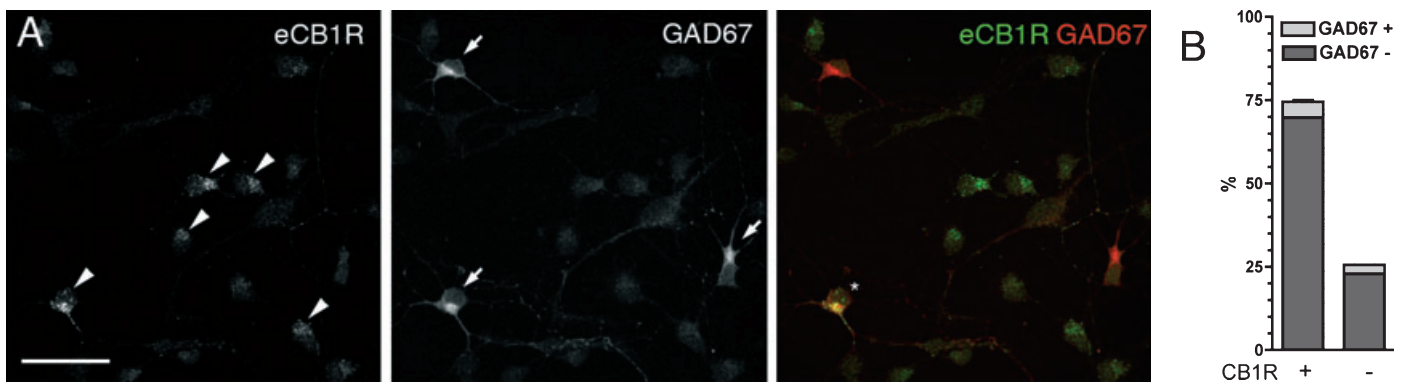


Fig. 5. Endogenous CB₁Rs are highly expressed in cultured glutamatergic hippocampal neurons at DIV4. (A) Immunohistochemical detection shows that at DIV4 the majority of cultured hippocampal neurons express endogenous CB₁ receptors (eCB₁Rs) (arrowheads). Simultaneous detection of the 67 kDa isoform of glutamic acid decarboxylase (GAD67) shows that only about 4–5% of the neurons are GABAergic (arrows) and that around half of these neurons express eCB₁R (asterisk). Bar 50 μm. (B) Quantification shows that around 75% of cultured hippocampal neurons are glutamatergic and express endogenous CB₁R at DIV4. Results are mean ± SEM.

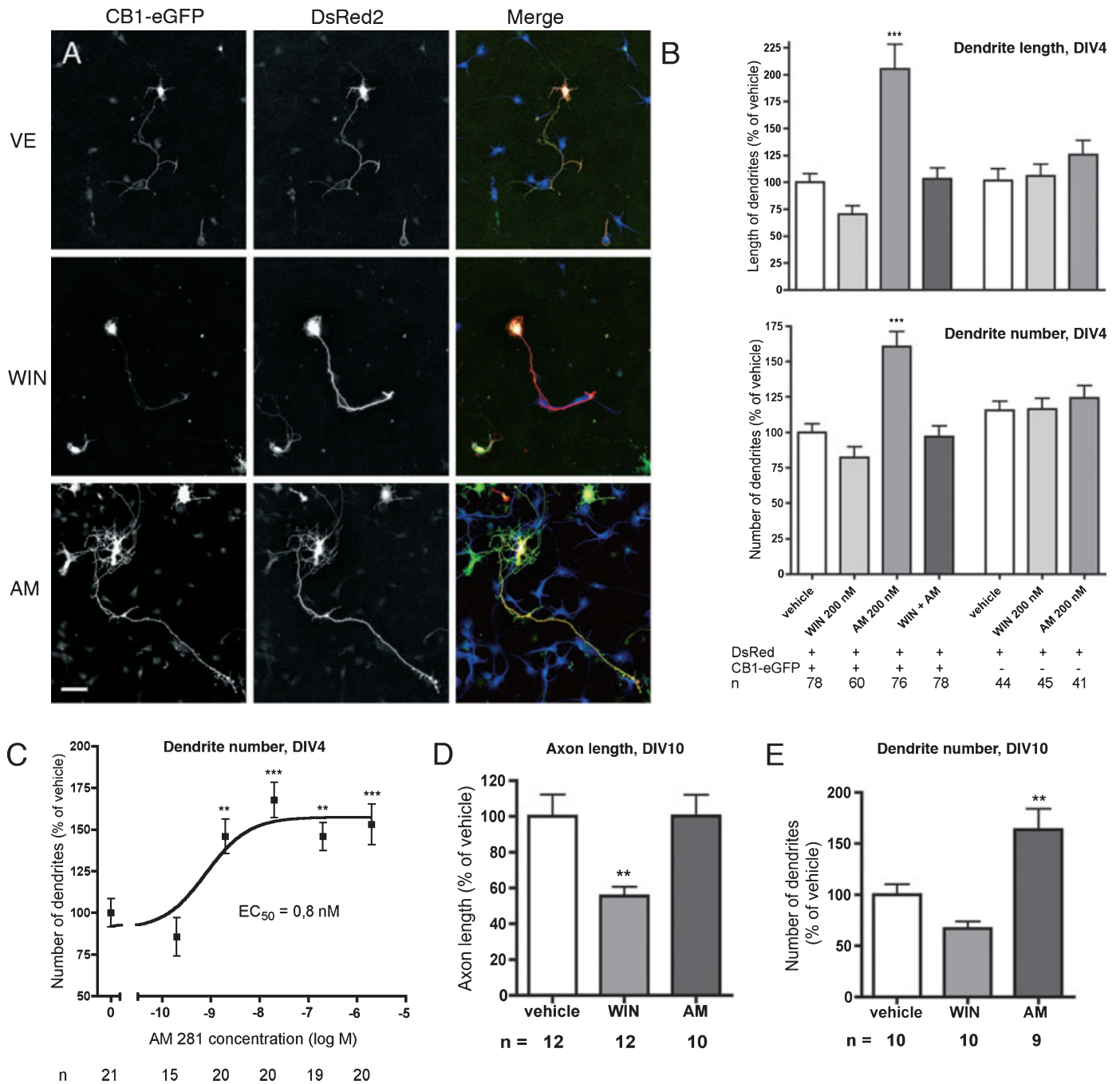


FIG. 6. Pharmacological activation or blockade of transfected CB₁R_s inversely modifies neuronal structure. Cultured hippocampal neurons were transfected with CB₁R-EGFP and the structural marker DsRed2 at DIV3. After treatment with 200 nM of the CB₁R agonist WIN or 200 nM of the inverse agonist AM281 (AM), the neurite structure of CB₁R-EGFP-expressing cultured hippocampal neurons is profoundly modified in opposite directions. (A and B) Overnight agonist treatment (WIN) reduces both the length and number of primary dendrites, whereas inverse agonist (AM) treatment results in a highly significant augmentation of these parameters as compared with vehicle (VE). Simultaneous application of WIN counteracts AM effects. In control neurons transfected only with the structural marker DsRed2, there is no significant morphological change with this treatment protocol. On the merged image, CB₁R is labeled in green, DsRed in red and the neuronal marker microtubule-associated protein 2 in blue. Results are pooled from four independent experiments and are expressed as mean ± SEM. (C) The effect of overnight AM281 treatment is dose-dependent with an EC₅₀ of 0.8 nM. (D and E) Prolonged (7 days) treatment with 200 nM WIN resulted in shorter axons and reduced dendrite number, whereas treatment with 200 nM AM281 had opposite effects on the number of dendrites. Results are representative of at least two independent experiments and are expressed as mean ± SEM. ***P* < 0.01 and ****P* < 0.001 vs. vehicle.

with the expression of wild-type CB₁R (Fig. 7A). On the contrary, expression of the hypoactive T210A mutant of CB₁R (Fig. 7C) significantly increased the length of axons (up to 160.1 ± 13.2%) and also slightly increased the number and length of primary dendrites (to 110.9 ± 6.3 and 116.4 ± 9.3%, respectively, Fig. 7D).

In conclusion, we have shown that pharmacological or mutational manipulation of CB₁R activity significantly modifies neuronal morphology of cultured hippocampal neurons. CB₁R activation exerts an overall negative effect on the development of both axons and dendrites. Finally, the spontaneous activation of transfected wild-type

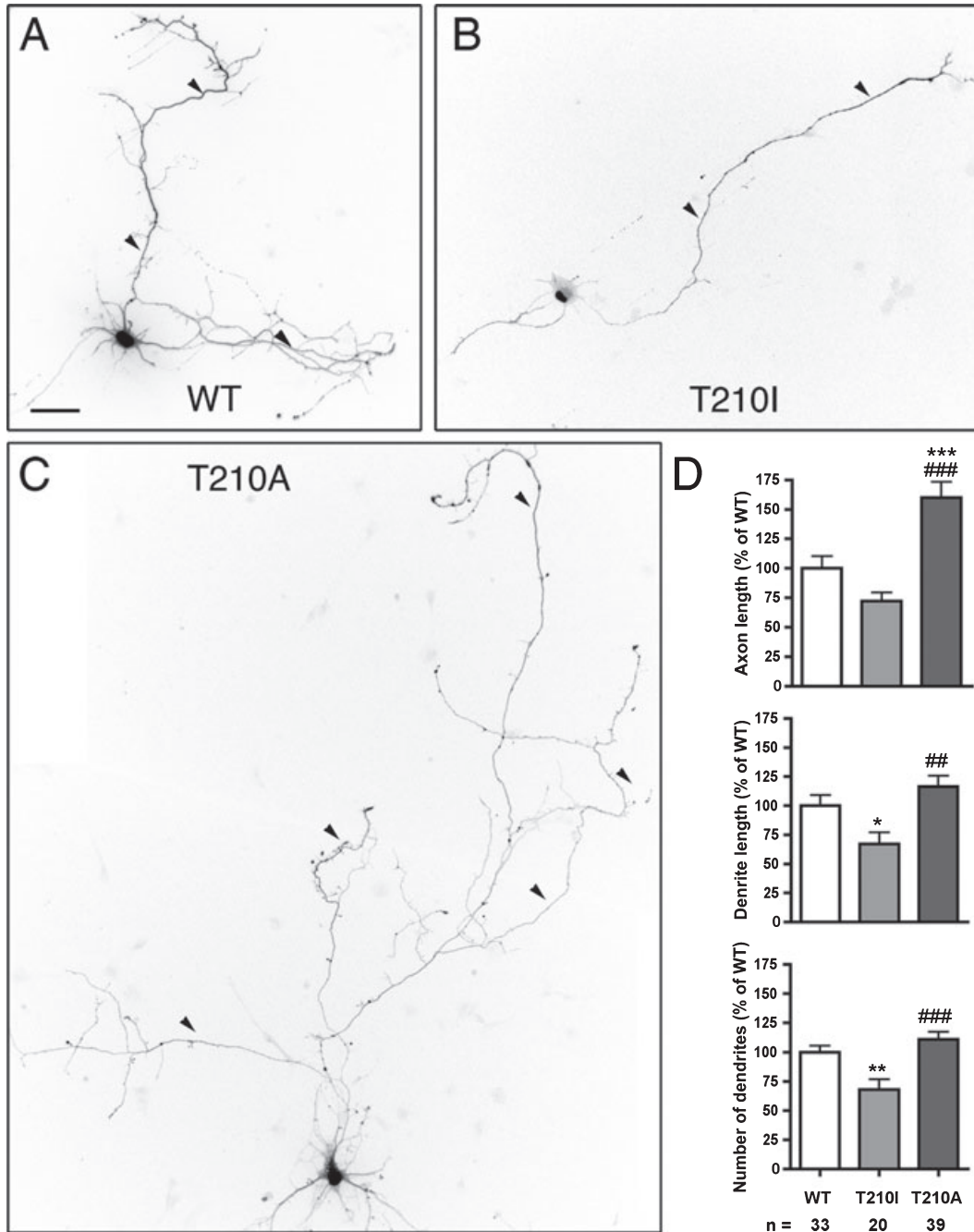


FIG. 7. Transfection of point mutants of CB₁Rs that display different levels of constitutive activity results in modified dendritic arborization and axon outgrowth. (A–C) Cultured hippocampal neurons were co-transfected at DIV3 with the structural marker DsRed2 and either wild-type CB₁R-EGFP (wild-type) or the T210I-EGFP over-active point mutant or the T210A-EGFP low-activity mutant. At DIV4, neurons were fixed and neuron morphology was quantified by NeuronJ. Bar 50 μm. (D) Mutational activation or inactivation exerts significant opposite effects on dendritic and axonal outgrowth. Axons are indicated by arrowheads. Results are pooled from two independent experiments and are expressed as mean ± SEM. **P* < 0.05, ***P* < 0.01 and ****P* < 0.001 vs. vehicle. ##*P* < 0.01 and ###*P* < 0.001 vs. T210I-CB₁R.

CB₁Rs has a significant negative effect on the neuritic development of isolated hippocampal neurons, as demonstrated by the effect of AM281 treatment or the morphological difference between neurons transfected with wild-type CB₁R vs. T210I-CB₁R.

Discussion

We report that, starting from E12.5, CB₁Rs are strongly expressed in both reelin-expressing Cajal-Retzius cells and newly differentiated

postmitotic glutamatergic neurons of the mouse telencephalon. CB₁R protein is localized first to somato-dendritic endosomes and at later developmental stages it localizes mostly to developing axons (see supporting Fig. S2). In young axons, CB₁Rs localize to both the axolemma and large, often multivesicular endosomes. We also show that maternal treatment with cannabinoid agonist results in internalization of embryonic CB₁Rs, suggesting functional competence. Finally, by using *in-vitro* assays we demonstrate that activation of CB₁R signaling acts as a negative regulatory signal for dendritic and axonal outgrowth.

CB₁R expression in the developing telencephalon

Strong and transient CB₁R binding and mRNA expression were described in the rodent cerebral cortex at early developmental stages (reviewed in Fernandez-Ruiz *et al.*, 1999). Later at E14-P4, during the period of intense axonal and dendritic outgrowth (Lopez-Bendito *et al.*, 2003), CB₁R mRNA is still highly expressed by cortical and hippocampal areas, and, importantly, white matter areas containing the developing axons of these neurons also show transiently strong expression of functional CB₁Rs (Fernandez-Ruiz *et al.*, 1999). This transiently strong expression in telencephalic glutamatergic neurons was also suggested by studies in human fetuses showing CB₁R binding in axon-rich regions such as the capsula interna and pyramidal tracts (Mato *et al.*, 2003; Fride, 2004). Axonal expression was recently confirmed by immunohistochemistry (Berghuis *et al.*, 2007) and in the present study we provide the first comprehensive high-resolution description of CB₁R localization during the early stages of cortical and hippocampal development. Thus, we show that CB₁Rs are highly expressed from the earliest stages of neuronal differentiation by postmitotic preplate neurons at E12.5. Indeed, starting from the earliest stage investigated, CB₁R mRNA and protein expression was restricted to Tuj-1-immunopositive committed neurons and was never observed in the ventricular zone, indicating that, in contrast to adult progenitor cells (Jin *et al.*, 2004; Aguado *et al.*, 2005, 2006; Jiang *et al.*, 2005), in the rodent embryonic forebrain CB₁R expression is restricted to postmitotic neurons. This is in line with data reported in the chick and zebrafish embryonic brainstem and spinal cord (Watson *et al.*, 2008) during the preparation of the present article. We also identify CB₁R expression in a subpopulation of reelin-secreting Cajal-Retzius cells. Cajal-Retzius cells participate in the regulation of the phenotype of radial glial cells and the inside-out formation of the cortical plate (reviewed by Super *et al.*, 1998), as well as in the development of cortical dendrites (MacLaurin *et al.*, 2007). These results suggest that CB₁R signaling could participate in the initial steps of cortical plate and hippocampal formation and in the development of the glutamatergic projections. We also show that, at later developmental stages, CB₁Rs are gradually translocated to axon tracts composed of fascicles of corticofugal axons. In contrast with the work of the group of Harkany (Berghuis *et al.*, 2007), reporting functional significance of CB₁R expression in migrating GABAergic neurons before birth and CB₁R expression in 'GABAergic interneurons during late gestation as they were undergoing intracortical tangential or radial migration', we did not find notable expression of CB₁R mRNA or CB₁R protein in the ganglionic eminences or in tangentially migrating interneurons in the subventricular zones. Finally, our data indicate that, at birth, CB₁R shows an expression pattern intermediate between embryonic and adult phenotype, the latter being firmly acquired only after P5.

Embryonic CB₁Rs are localized to endosomes before acquiring functional expression on the axolemma

At the ultrastructural level CB₁R was almost exclusively associated with the outer membrane of large and complex endosomes in preplate neurons. At later stages, CB₁Rs were still localized to complex endosomes in the somato-dendritic region of cortical plate neurons but in developing axonal tracts a significant number of CB₁Rs were localized to round and multivesicular endosomes, accompanied by expression on the axonal plasma membrane. Endosomal staining was associated with the cytoplasmic side of the organelles in accordance with the fact that the anti-CB₁R antibody used was raised against the C-terminus of the receptor. Notably, a similar, mostly endosomal

localization was also reported in the somato-dendritic region of adult cortical and hippocampal neurons (Katona *et al.*, 2000; Bodor *et al.*, 2005), as well as in the somato-dendritic region of mature cultured hippocampal neurons (Leterrier *et al.*, 2006; McDonald *et al.*, 2007) and in cell lines (Leterrier *et al.*, 2004; D'Antona *et al.*, 2006; Ellis *et al.*, 2006; Wu *et al.*, 2008), where this endosomal localization of CB₁Rs is the result of intense constitutive endocytosis. This cycling is tightly associated with basal receptor activation, partly resulting from autocrine activation by endogenous 2-arachidonoylglycerol (2-AG) produced by diacylglycerol lipases (Turu *et al.*, 2007). Thus, the finding that CB₁Rs are mostly endosomal in the embryonic brain raises the possibility that these receptors are significantly activated in the embryonic brain. Interestingly, in embryonic axons, where CB₁Rs are strongly internalized, both diacylglycerol lipases α and β are highly expressed (Bisogno *et al.*, 2003; Watson *et al.*, 2008), suggesting that CB₁Rs may be significantly activated by a putative autocrine or paracrine production of 2-AG, leading to permanent activation and internalization of CB₁Rs in both the somato-dendritic region and axons. This endocytic stimulation is not maximal, as shown by the effect of acute agonist treatment, which leads to a more enhanced endosomal phenotype, characterized by complete depletion of CB₁R from the axonal surface and enhanced formation of multivesicular endosomes, ultimately leading to significant translocation from axons to the somato-dendritic compartment of glutamatergic neurons. Interestingly, in mature neurons CB₁Rs appear to be efficiently recycled after moderate activation but more sustained activation directs CB₁Rs to the lysosomal degradation pathway through interaction with G-protein-coupled receptor-associated sorting protein (GASP-1) protein (Martini *et al.*, 2007; Tappe-Theodor *et al.*, 2007). Thus, extrapolation of these results to our data would suggest that CB₁Rs, moderately activated by autocrine 2-AG in embryonic axons, are possibly efficiently recycled locally, whereas more pronounced activation by exogenous cannabinoid ligands possibly leads to somato-dendritic translocation followed by lysosomal degradation.

It should also be noted that considering axonal multivesicular endosomes solely as predegradative and degradative compartments is probably an oversimplification, as we have often found CB₁Rs localized to multivesicular endosomes in untreated animals in the embryonic brain (present study) as well as in untreated DIV9 cultured hippocampal neurons (Leterrier *et al.*, 2006). Interestingly, recent data suggest that these endosomes may be specialized intermediates characteristic to the somatodendro-axonal transcytotic pathway, which is responsible for the correct axonal targeting of axolemmal proteins such as NgCam (Yap *et al.*, 2008) or CB₁R (Leterrier *et al.*, 2006). The definition of the exact role of these organelles clearly necessitates further investigation.

In conclusion, our results indicate that: (i) CB₁Rs in the embryonic brain are functional, as was also previously indicated by efficient GTP γ S binding in the rat (Berrendero *et al.*, 1998) and human brain (Mato *et al.*, 2003); (ii) CB₁Rs are moderately but significantly activated by endocannabinoids, presumably through autocrine production of 2-AG; and (iii) maternal cannabinoid consumption may lead to over-activation and subneuronal translocation of embryonic CB₁Rs.

Pharmacological activation or blockade of CB₁Rs inversely modifies the morphology of pyramidal neurons

What is the physiological role of activated CB₁Rs in developing projection axons, well before the establishment of synaptic functions?

Our results show that stimulation or inhibition of the CB₁R activation state by pharmacological ligands or point mutation of T210 inversely regulates neuron morphology. Interestingly, the significant effect of antagonist/inverse agonist treatment as well as the difference between wild-type and T210A-CB₁R indicate that transfected wild-type CB₁Rs are sufficiently activated to exert a notable inhibiting effect, without addition of an exogenous agonist. Because the effects were identical in neurons incubated with glia-conditioned or with fresh unconditioned medium, the source of this basal activation is probably either the constitutive activity of CB₁Rs, the cell autonomous autocrine production of 2-AG (Turu *et al.*, 2007) or a combination of both. The dominant effect of CB₁R activation in young DIV3 neurons is the inhibition of dendritogenesis, dendritic outgrowth and, to a lesser extent, axonal outgrowth, the latter effect being more pronounced at later developmental stages. Previously, CB₁Rs were shown *in vitro* both to positively regulate fibroblast growth factor (FGF)-stimulated axonal growth of cerebellar neurons (Williams *et al.*, 2003) and negatively regulate neurite development of GABAergic neurons (Berghuis *et al.*, 2005). In addition, recent *in-vivo* data show that, in chick and zebrafish embryos, CB₁R gene knockdown results in defective axonal growth and fasciculation (Watson *et al.*, 2008). Finally, and rather at variance with our direct *in-vitro* data, in CB₁R knockout mice the spine density and field area of basal dendrites of layer III pyramidal neurons are significantly reduced as compared with wild-type mice (Ballesteros-Yanez *et al.*, 2007). Our results, obtained in isolated hippocampal neurons, suggest that the dominant cell-autonomous effect of CB₁R stimulation (pharmacological or mutational) is the inhibition of both axon and dendrite development. It should, however, be kept in mind that our results were obtained by using transfected neurons and that effects obtained by pharmacological manipulation of endogenous CB₁Rs, although pointing to the same direction, did not reach the significance threshold in our experimental system. Previously, the existence of powerful compensatory mechanisms and high functional plasticity were proposed to explain the lack of marked developmental central nervous system deficits in the available null mutant mice, deficient for different components of the endocannabinoid system (Harkany *et al.*, 2008).

Taken together, the overall negative regulatory role in the neurite development of cortical projection neurons suggests that embryonic CB₁R signaling is one of the numerous players that help the establishment of correct neuronal connectivity. Data reported in the present study point to an interesting putative mechanism to explore in order to better understand the development of reported glutamatergic dysfunction in offspring following maternal cannabis consumption.

Note added in revision

During the revision process of this manuscript, Mulder *et al.* (2008) have published a study that includes interesting data on CB₁R expression and function in glutamatergic neurons of the embryonic rodent brain. The expression pattern of CB₁Rs is well correlated with data reported in the present study, with the exception of the proliferative subventricular zone where, at variance with our data, they report significant levels of CB₁R expression. *In vitro*, Mulder *et al.* (2008) find that endogenous CB₁Rs inhibit dendritic development of cultured cortical neurons but, at variance with the present report where we used hippocampal neurons, they report a stimulating role for CB₁Rs in axonal growth. Finally, Mulder *et al.* (2008) present qualitative data on the *in-vivo* significance of embryonic CB₁R expression by showing that pharmacological or genetic ablation of CB₁Rs results in deficits of neuronal progenitor proliferation and axon fasciculation.

Supporting information

Additional supporting information may be found in the online version of this article:

Fig. S1. Location of CB₁R on cortical neurons at P5 and in adult mice.
Fig. S2. Summary of CB₁R expression profile during early cortical development in mice.

Video S1. Three D reconstruction of a confocal stack from the cortical preplate of an E12.5 embryo.

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Abbreviations

2-AG, 2-arachidonoylglycerol; CB₁R, type 1 cannabinoid receptor; DAPI, 4',6-diamidino-2-phenylindole 2 HCl; DIV, days *in vitro*; DPBS, Dulbecco's phosphate-buffered saline; E, embryonic day; EGFP, enhanced green fluorescent protein; GFP, green fluorescent protein; P, postnatal day; Tuj-1, neuron-specific beta III tubulin; WIN, WIN55,212-2.

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