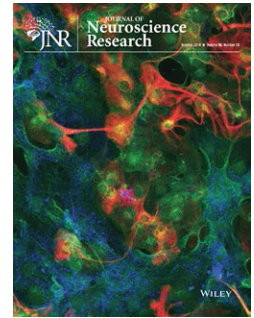


## RESEARCH ARTICLE



# Neuronal and synaptic morphological alterations in the hippocampus of cannabinoid receptor type 1 knockout mice

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## Abstract

Cannabinoid receptor type 1 (CB1R) modulates synaptic activity and is widely distributed in brain areas such as the hippocampus, cerebellum, cerebral cortex, and striatum, among others. CB1R is involved in processes such as memory, learning, motor coordination, and mood. Genetic deletion of CB1R causes behavioral alterations. In this work, we evaluated neuronal morphology and synaptic structure in the hippocampus of adult male CB1R knockout mice (CB1R<sup>-/-</sup>). Morphological changes in the CB1R<sup>-/-</sup> hippocampus evidenced a decrease in the expression of cytoskeletal proteins neurofilaments 160 KDa, neurofilaments 200 KDa, and microtubule-associated protein 2. CA1 neurons showed decreased arborization and changes in synaptic structure such as lower thickness of postsynaptic density and a reduction in synaptophysin levels. Results obtained in the present work provide evidence of the participation of CB1R in the establishment of neuronal structure and networks that could have an important role in neuronal plasticity. In addition, these changes observed in CB1R<sup>-/-</sup> could be correlated with behavioral alterations reported.

## KEYWORDS

cannabinoid receptor type 1, cannabinoid system, RRID:AB\_477523, RRID:AB\_477193, RRID:AB\_477261, RRID:AB\_477257, RRID:AB\_143162, RRID:AB\_476738, RRID:AB\_10015282, RRID:AB\_2340770, neuronal plasticity, synapse

## 1 | INTRODUCTION

The endocannabinoid system, which has been conserved evolutionarily from plants to vertebrates (Elphick & Egertová, 2005; Solymosi & Köfalvi, 2017), modulates synaptic transmission and consists of cannabinoid receptors type 1 (CB1R) and type 2 (CB2R), its

endogenous ligands (anandamide and 2-arachidonoylglycerol) and enzymes involved in its metabolism (Lambert & Fowler, 2005).

CB1R is widely expressed in the central nervous system in vertebrates, both in neurons and glial cells (Facchinetti, Del Giudice, Furegato, Passarotto, & Leon, 2003; Metna-Laurent & Marsicano, 2015; Molina-Holgado et al., 2002), mainly in the cerebral cortex, cerebellum, hippocampus, basal ganglia, striatum, and amygdala (Hu

**Abbreviations:** CB1R, cannabinoid receptor type 1; CB1R<sup>-/-</sup>, CB1R knockout mice; CB1R<sup>+/+</sup>, CB1R wild-type mice; CB2R, cannabinoid receptor type 2; DG, dentate gyrus; MAP2, microtubule-associated protein 2; NF160, neurofilaments 160 KDa; NF200, neurofilaments 200 KDa; PB, phosphate buffer; SD, standard deviation; Syn, synaptophysin; WB, Western blot.

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& Mackie, 2015). CB1R is one of the most abundant G protein-coupled receptors in the central nervous system and is mainly presynaptically located in axonal terminals.

In mammals, CB1R participates in the regulation of motor coordination, anxiety, memory, and appetite (Marsicano & Lafenetre 2009). Activation of CB1R modulates cell excitability, neurotransmitter release, transient or permanent synaptic transmission, thus participating in short- and long-term plasticity by different signaling pathways (Castillo, Younts, Chávez, & Hashimoto, 2012; Ligresti, De Petrocellis, & Di Marzo, 2016; Xu & Chen, 2015). CB1R also modulates voltage-dependent ion channels and inhibits voltage-gated  $Ca^{2+}$  channels (N- and P/Q-type) and activates rectifying K channels (Ligresti et al., 2016).

The hippocampus has demonstrated an essential role in cognitive and emotional functions, involving plasticity process such as structural and functional changes and generation of new neurons (Fanselow & Dong, 2010; Gould, 2009). The hippocampus has behavioral, anatomical, and functional segmentation into separate structural; dorsal hippocampus is involved in learning, memory, and exploration, while the ventral hippocampus modulates emotional and affective processes (Fanselow & Dong, 2010). CB1R is highly expressed in hippocampus and mediates learning and memory functions (Marsicano & Lafenetre, 2009), regulates mood states (Marco & Viveros, 2009) and synaptic changes associated (Alger, 2009).

Alteration in CB1R signaling produced changes in emotional behavior showing anhedonic state (Burokas et al., 2014; Sanchis-Segura, Cline, Marsicano, Lutz, & Spanagel, 2004), impaired extinction of aversive memory (Marsicano et al., 2002; Martin, Ledent, Parmentier, Maldonado, & Valverde, 2002), and an increase in the conditioned fear responses (Jacob, Marsch, Marsicano, Lutz, & Wotjak, 2012). Several studies have demonstrated an anxiogenic behavior in CB1R knockout mice (CB1R<sup>-/-</sup>) or upon pharmacological blockade of CB1R under normal conditions (Bowers & Ressler, 2016; Martin et al., 2002). Hypothalamic-pituitary-adrenal axis dysregulation has been observed in CB1R<sup>-/-</sup> (Barna, Zelena, Arszovszki, & Ledent, 2004), anxiety-like behavior similar to that detected in naive mice exposed to stress conditions (Hill, Hillard, & McEwen, 2011). In addition, CB1R<sup>-/-</sup> presents alterations in long-term synaptic plasticity in corticolimbic structures such as amygdala and hippocampus (Bohme, Laville, Ledent, Parmentier, & Imperato, 2000; Jacob et al., 2012; Marsicano et al., 2002; Monory, Polack, Remus, Lutz, & Korte, 2015) and an increase in serotonin level in prefrontal cortex in normal conditions, but these decrease after fluoxetine is administered (Aso et al., 2009; Ferreira et al., 2012). As functional alterations found in CB1R<sup>-/-</sup> correlate with a decrease in CB1R receptor expression in animal models of depression (Hill et al., 2008), CB1R<sup>-/-</sup> has been postulated as a model of depression (Valverde & Torrens, 2012). Hay and collaborators indicate that disruption of conserved regulatory sequence (ECR1) in *cnr1*, that contained a polymorphism, reduced CB1R expression in the hippocampus and decreased anxiety-related behaviors (Hay et al., 2019).

CB1R is implicated in the process of long-term plasticity associated with spatial learning and memory. Deletion of CB1R can

### Significance

Cannabinoid receptor type 1 (CB1R) is expressed in the central nervous system, mainly in cerebral cortex, hippocampus, cerebellum, and striatum at presynaptic level. CB1R plays a modulatory role in synaptic activity regulating the neurotransmitter release. The blockade of CB1R, by genetic or pharmacological approach, induces neuronal plasticity alterations. Here, we demonstrated that genetic ablation of CB1R induces morphological alterations in neuronal cytoskeleton and synapse, which correlates with morphological alterations observed in chronic exposure to CB1R agonist. These data support that CB1R participates also in the neuronal feature development.

result in increased long-term potentiation in hippocampus (Bohme et al., 2000; Jacob et al., 2012), but contraries' alteration has been observed in spatial memory in these mice (Albayram et al., 2011; Jacob et al., 2012; Lin et al., 2011; Mikics et al., 2006). In addition, treatment with CB1R antagonist induces increased long-term potentiation (Hoffman et al., 2007; Slanina, Roberto, & Schweitzer, 2005) and enhanced spatial memory (de Oliveira, Pasqualini, Diehl, Molina, & Quillfeldt, 2008; Jacob et al., 2012; Lichtman, 2000; Lin et al., 2011; Reich, Mohammadi, & Alger, 2008; Sink et al., 2010). Also, cannabis users present memory deficit (Ilan, Smith, & Gevins, 2004; Schweinsburg, Brown, & Tapert, 2008; Solowij et al. 2002) and this could correlate with reduction in the expression of CB1R (Hirvonen et al., 2012; Rotter et al., 2013; Villares, 2007) and smaller volume of the hippocampus (Lorenzetti, Chye, Silva, Solowij, & Roberts, 2019; Schacht, Hutchison, & Filbey, 2012).

The expression levels of the CB1R in humans are associated with different polymorphisms in the *cnr1* gene (Zhang et al., 2004), many of them have been correlated with psychiatric disorders (Hillard, Weinlander, & Stuhr, 2012) such as anxiety, depression, and psychosis (Juhász et al., 2009; Lazary et al., 2009; Suárez-Pinilla et al., 2015), and with alteration in learning and memory (Fairfield et al., 2018; Ruiz-Contreras et al., 2011, 2014).

Endocannabinoid signaling participates in the development of thalamocortical projections, and CB1R ligands disrupt its normal formation (Itami et al., 2016). CB1R<sup>-/-</sup> mice and pyramidal cell-specific conditional mutant mice develop deficits in neuronal progenitor proliferation and axon fasciculation. Moreover, pharmacological blockade of CB1R during development induces axonal disruption (Mulder et al., 2008; Tortoriello et al., 2014; Watson, Chambers, Hobbs, Doherty, & Graham, 2008) and impairs dendritic elongation (Tapia et al., 2017). In utero blockade of CB1R by CB1R siRNA electroporation alters neuronal migration during cortical development, and increases seizures induced by convulsant pentylenetetrazol (Díaz-Alonso et al., 2017). *In vitro* studies of cortical neurons have shown that the activation of the CB1R produces a decrease in the axonal diameter and in the growth of neurites, product of changes

in dynamics of the neuronal cytoskeleton (Njoo, Agarwal, Lutz, & Kuner, 2015; Tortoriello et al., 2014). Chronic treatment with CB1R agonist during adolescence causes a deterioration in spatial memory in adulthood, which is correlated with the decrease in the expression of pre- and postsynaptic components, less arborization, and a lower number of spines observed in granular neurons of the hippocampus (Rubino et al., 2009). Moreover, chronic exposure to the nonselective cannabinoid agonist, WIN55,212-2, in the adult induces a decrease in MAP2 expression in the CA1 area of the hippocampus (Lawston, Borella, Robinson, & Whitaker-Azmitia, 2000) and a decrease in the number of spines in neurons (Candelaria-Cook & Hamilton, 2014).

There are significant associations between *cnr1* polymorphisms and changes in specific brain structure such as reduced caudate nucleus and thalamus volumes and white matter (Suárez-Pinilla et al., 2015). Also, *cnr1* polymorphisms may predispose smaller hippocampal volume after heavy cannabis use (Lorenzetti et al., 2019; Schacht et al., 2012). These morphological brain abnormalities are consistent with the link to some psychiatric disorders. A recent stereological postmortem study in depression and schizophrenia subjects indicates that the volume and number of cells in different areas of hippocampus are reduced (Chen et al., 2019).

As changes in behavior are usually accompanied by morphological changes in neurons and synapses in related areas such as the hippocampus and the amygdala, this work evaluated neuronal morphology, neuronal cytoskeleton, and synaptic structure in the hippocampus using CB1R<sup>-/-</sup> mice. The purpose of this work is to describe the neuronal histological changes that are produced by the deficiency of CB1R that could explain the alterations in long-term synaptic plasticity, the behavioral changes described in CB1R<sup>-/-</sup> mice.

## 2 | METHODS

### 2.1 | Animals

Fifteen male CD1 CB1R wild-type (CB1R<sup>+/+</sup>) and 15 male CB1R knockout mice (CB1R<sup>-/-</sup>) were used in this study. CB1R<sup>-/-</sup> mice were generated by disruption of the *cnr1* gene as previously described by Ledent et al. (1999) and kindly donated by Prof. Dr. György M. Nagy (Hungarian Academy of Science and Semmelweis University, Budapest, Hungary). Heterozygote-heterozygote (CB1R<sup>+/-</sup>) mating produced CB1R<sup>+/+</sup> and CB1R<sup>-/-</sup> littermates. At postnatal day 21, male offspring were weaned and obtain tail samples for genotyping. Animals were genotyped by polymerase chain reaction (PCR) technique using primers "CB1" (forward: 5' CATCATCACAGATTCTATGTAC; WT reverse: 5' GAGGTGCCAGGAGGGAACC) and "Neo" (mutant forward: 5' GCAGGATCTCTGTATCATCTACC; mutant reverse: 5' GATGCTCTTCGTCCAGATCATCC). CB1R<sup>+/+</sup> and CB1R<sup>-/-</sup> amplicons are ~360 base pairs and ~250 base pairs, respectively (See Figure S1). After genotyping, 60-day-old male mice were chosen by simple random selection from litters. Animals were housed five or six male offspring of each litter per cage, in a humidity- and temperature-controlled environment (12 hr/12 hr light/dark cycle), with free access

to standard laboratory rodent food and water. Animal treatments were in accordance with CICAL protocols (Institutional Committee for the Care and Use of Laboratory Animals, School of Medicine, University of Buenos Aires, RES (CD) 2198/2015).

### 2.2 | Immunohistochemistry

Five CB1R<sup>+/+</sup> mice and five CB1R<sup>-/-</sup> mice were fixed by intracardiac perfusion with 4% (w/v) paraformaldehyde in 0.1 M phosphate buffer (PB) pH 7.4 and postfixed for 4 hr in the same solution. Coronal sections of the brains were cut with a vibratome (40- $\mu$ m thick) and stored in 50% (v/v) glycerol solution at -20°C until use. Coronal serial sections containing hippocampus were selected from five different mice of each group.

Immunohistochemistry was performed with following primary antibodies: mouse anti-Synaptophysin (Syn, 1:1,000, Sigma-Aldrich, St Louis, MO, USA, Cat# S5768, RRID:AB\_477523); mouse anti-microtubule-associated protein 2 (MAP2, 1:1,000, Sigma-Aldrich, Cat# M4403, RRID:AB\_477193); mouse anti-neurofilament 160 kDa (NF160, 1:1,000, Sigma-Aldrich, Cat# N2787, RRID:AB\_477261); mouse anti-neurofilament 200 kDa (NF200, 1:1,000, Sigma-Aldrich, Cat# N0142, RRID:AB\_477257). The same antibodies against Syn, NF160, NF200, and MAP2 were used in previous studies from our laboratory (Caltana, Saez, Aronne, & Brusco, 2015; Evrard et al., 2006; Tagliaferro et al., 2006).

For immunofluorescent labeling, after incubating with primary antibodies, brain serial slices (four to six slices per animal) were washed in phosphate saline buffer and incubated for 4 hr with fluorescent secondary antibodies anti-mouse IgG Alexa 568 (1:500, Invitrogen, Carlsbad, CA, USA, RRID:AB\_143162). Sections were later counterstained with Hoechst 33342 (2  $\mu$ g/ml, Sigma-Aldrich) to label nuclei and coverslipped with 70% (v/v) glycerol mounting medium.

Images were acquired on an Axiolab epifluorescence microscope (Carl Zeiss, Göttingen, Germany) equipped with a Q-Color3 CCD camera (Olympus, Tokyo, Japan). Counting and morphometry were performed using ImageJ (NIH, <https://imagej.net>, RRID:SCR\_003070) software. Brain areas analyzed were the stratum radiatum of CA1 and CA2, CA3 and the molecular layer of the dentate gyrus (DG), those areas were identified and defined according to the Mouse Brain in Stereotaxic Coordinates Atlas (Franklin & Paxinos, 2007). In accordance with Kohara et al. (2014), Chevaleyre and Siegelbaum (2010) and Botcher, Falck, Thomson, and Mercer (2014), we visually identified the CA2 area as the region with the thickest somatic layer located between CA3, characterized by its distinct mossy fiber terminal layer, and CA1, characterized by its compact cell body layer.

In order to ensure objectivity, for each set of experiments, all measurements were performed on coded slides, in blind conditions, by two observers (DS and LRC). Measurements of both groups of CB1R<sup>+/+</sup> and CB1R<sup>-/-</sup> sections were carried out in standardized conditions (at the same session, in the same day, by the same observer). In each tissue section, each microscopic field was selected within

the limits of each anatomical area of interest to be morphometrically analyzed.

In order to evaluate MAP2, NF160+, and NF200+ fibers, the total area of the immunolabeled fibers was related to the total area of the corresponding microscopic field (20× primary magnification), rendering a relative area parameter.

Optical intensity of Syn expression was analyzed in the stratum radiatum of CA1 and CA2, CA3 and the molecular layer of the DG. Optical intensity was analyzed quantitatively using an image analysis system (ImageJ). Color images were acquired for each hippocampal area by a digital camera connected to the fluorescent microscope. The images were transferred to black-white images, and mean light transmission was measured on a scale of 0 to 256 relative units. This measurement was repeated 15 times (50  $\mu\text{m}^2$ ) on each image and averaged. To normalize the variations in immunostaining strength between the slides, the intensity of each individual layer was divided by the intensity of the corpus callosum in each section, which was selected for background levels of immunoreactivity because of the absence of axon terminals (Jung et al., 2009).

### 2.3 | Electron microscopy

Five CB1R<sup>+/+</sup> mice and five CB1R<sup>-/-</sup> mice were fixed by intracardiac perfusion with 4% (w/v) paraformaldehyde and 0.25% (v/v) glutaraldehyde in PB (Tagliaferro et al., 2006). Dissected sections of 1 mm<sup>3</sup> corresponding to CA1 area of hippocampus were post-fixed in 1% (w/v) osmium tetroxide in the same PB for 30 min. After dehydration in ethanol gradient, tissues were contrasted with 5% (w/v) uranyl acetate and then embedded in Durcupan (Fluka AG, Chemische Fabrik, Buchs SG, Switzerland). After sections were embedded in Durcupan, 1- $\mu\text{m}$  slices were obtained and toluidine blue stained, to select similar areas of CA1 stratum radiatum. Then ultrathin sections (50 nm) were obtained and stained with lead citrate (Reynolds, 1963). Images were acquired on a Zeiss 109 transmission electron microscope (TEM) (Carl Zeiss, Oberkochen, Germany) and photographed with a GATAN CCD camera (Pleasanton, CA, USA).

For electron microscopy images, synapse number per 100  $\mu\text{m}^2$  was measured in CA1 stratum radiatum (CB1R<sup>+/+</sup>: 49 areas, CB1R<sup>-/-</sup>: 45 areas). For synapse morphology, the number of synaptic vesicles, synaptic curvature (ratio of the synaptic length in the synaptic gap and the distance between the synaptic limits), and the thickness of the postsynaptic density (CB1R<sup>+/+</sup>: 121 synaptic terminals and CB1R<sup>-/-</sup>: 122 synaptic terminals) were measured using ImageJ.

### 2.4 | Golgi stain

Five CB1R<sup>+/+</sup> mice and five CB1R<sup>-/-</sup> mice brains were dissected and fixed with 4% (w/v) paraformaldehyde during 24 hr. They were later immersed in 3% (w/v) potassium dichromate solution (Sigma-Aldrich)

for 3 days in the dark, and then transferred into 2% (w/v) silver nitrate solution (Sigma-Aldrich) for 1 day in the dark at room temperature (modified from Bayram-Weston, Olsen, Harrison, Dunnett, & Brooks, 2016).

At the end of this time, 100- $\mu\text{m}$  thick sections were cut with a vibratome in distilled water, and sections were collected in 0.3% (w/v) gelatin solution. Finally, sections were mounted on slides, dehydrated by air-drying and coverslipped using Canada Balsam-type mounting medium.

Golgi-stained brain slices were analyzed using an inverted microscope DSU OlympusIX83. Proximal neuronal complexity and primary dendrites' number of representative pyramidal neurons (10–12 slices per animal, four to five neurons per slice, from three animals per phenotype) from hippocampal CA1 region were analyzed as previously described (Sholl, 1953; Spilker et al., 2016). Sholl analysis was performed by analyzing dendritic branching, number of apical and basal dendrites, and dendritic diameter. To determine spine density, the number of spines on segments of at least 100  $\mu\text{m}$  of dendritic length/neuron was counted. The number of spines was measured in the first 100  $\mu\text{m}$  from the first branching point (secondary dendrites) of the principal dendrite. Dendritic spine were classified according their morphology in mature shaped spines, which included Type I (stubby) and Type II (mushroom) and Type III (immature-shaped thin) (Ferrerias et al., 2017). For quantitative analysis of their branching patterns and spine density NeuroJ plugin ImageJ software and Cell target software were used (Garcia-Segura & Perez-Marquez, 2014).

### 2.5 | Western blot (WB)

Five CB1R<sup>+/+</sup> mice and five CB1R<sup>-/-</sup> mice were anesthetized with a solution containing ketamine/xylazine (100  $\mu\text{g}/\text{g}$ ; 10  $\mu\text{g}/\text{g}$  dissolved in saline solution) and decapitated, then the brain and hippocampi were dissected. The samples were homogenized in RIPA homogenization buffer (PMSF 1 mM, EDTA 1 mM, Na<sub>3</sub>VO<sub>4</sub> 1 mM, NaF 10 mM) with protease inhibitors and centrifuged at 14,000 rpm for 10 min at 4°C, after which the supernatant was collected. Samples (50  $\mu\text{g}$  protein) were filled into an 8% SDS-PAGE gel; the proteins were divided by electrophoresis and then transferred to nitrocellulose membranes, which were then incubated with primary antibodies (Guadagnoli, Caltana, Vacotto, Gironacci, & Brusco, 2016). The antibodies used for immunoblot were as follows: mouse anti-Syn (1:1,000, Sigma); mouse anti-MAP2 (1:2,000, Sigma); mouse anti-NF160 (1:1,000, Sigma); mouse anti-NF200 (1:1,000, Sigma); rabbit anti-actin (1:1,000, Sigma-Aldrich Cat# A5060, RRID:AB\_476738) (Figure S2). After being washed, the blot was incubated with a 1:2,000 dilution of a peroxidase-coupled donkey anti-rabbit polyclonal antiserum (Jackson ImmunoResearch Laboratories Inc. Cat#711-035-152, RRID:AB\_10015282) or peroxidase-coupled donkey anti-mouse polyclonal antiserum (Jackson ImmunoResearch Laboratories Inc. Cat# 715-035-150, RRID:AB\_2340770) for 1 hr at room temperature. Finally, bound enzymatic activity was detected by the enhanced chemiluminescence system (ECL+) from Amersham.

## 2.6 | Statistical analysis

Sixty-day-old CB1R<sup>+/+</sup> and CB1R<sup>-/-</sup> male mice were randomly selected from litters after genotyping. The number of animals (n) was calculated according to Arifin and Zahiruddin (2017) following the 3Rs rule (Russell & Burch, 1959). For immunohistochemistry, three separate experiments were run for each immunostaining study. Individual experiments were composed of 6 to 10 coronal serial sections of each animal from each group. Three to five fields were measured from each hippocampus in each coronal section of each animal by stereological analysis method using ImageJ (<https://imagej.nih.gov/ij/plugins/grid.html>).

For WB, three separate experiments were run for each immunostaining study. Individual experiments were composed of five animals from each group.

Statistical analysis was performed using GraphPad Prism 4 (GraphPad Software Inc., La Jolla, CA, USA). In immunostaining, WB, and electron microscopy images quantification, statistical significance was assessed by unpaired student's *t* test to compare the means of CB1R<sup>+/+</sup> and CB1R<sup>-/-</sup> mice. The same analysis was performed in dendritic diameter, dendritic branching, and number of spines in Golgi stain to compare the means of CB1R<sup>+/+</sup> and CB1R<sup>-/-</sup> mice. In the case of dendritic intersections in Golgi study, statistical significance was assessed by two-way analysis of variance (ANOVA) followed by Bonferroni posttest to evaluate the effect of genotype in number of prolongations and distance from neuron soma. *p* values <0.05 were considered significant. Reported values represent the mean ± SD of experiments performed for each marker and each hippocampal area.

In order to simplify graphic presentation, values of WB were expressed as percentages of the CB1R<sup>+/+</sup> group (value = 1 for WB).

## 3 | RESULTS

### 3.1 | Neurofilament expression analysis

Neurofilaments are the most important components of the axonal cytoskeleton and their elastic properties help maintain the markedly asymmetric shape of neurons. As neurofilament dynamic remodeling is essential for axonal growth and maintenance, the study of neurofilament expression reveals whether alterations occur in neuronal functioning (Evrard & Brusco, 2011; Yuan, Rao, & Nixon, 2012).

The expression of NF160 and NF200 was analyzed in the stratum radiatum of the hippocampal areas CA1, CA2, CA3, and in the molecular layer of the DG (Figures 1a,b and 2a,b).

A reduction was observed in the area covered by NF160+ and NF200+ fibers in the CA1 area in CB1R<sup>-/-</sup> as compared to CB1R<sup>+/+</sup> (NF160 = CB1R<sup>+/+</sup>: 16.11 ± 3.54 vs. CB1R<sup>-/-</sup>: 12.09 ± 1.42, unpaired *t* test, *t*(8) = 2.356, *p* = 0.047, *n* = 5 and NF200 = CB1R<sup>+/+</sup>: 31.58 ± 2.02 vs. CB1R<sup>-/-</sup>: 24.18 ± 3.44, unpaired *t* test, *t*(8) = 4.14,

*p* = 0.029, *n* = 5) (Figures 1c,d,l and 2c,d,l). However, no differences were found between groups in the expression of NF160 and NF200 in the stratum radiatum of CA2 area (NF160 = CB1R<sup>+/+</sup>: 20.32 ± 5.37 vs. CB1R<sup>-/-</sup>: 16.17 ± 5.65, unpaired *t* test, *t*(8) = 1.19, *p* = 0.686, *n* = 5 and NF200 = CB1R<sup>+/+</sup>: 14.80 ± 4.75 vs. CB1R<sup>-/-</sup>: 13.16 ± 2.31, unpaired *t* test, *t*(8) = 0.69, *p* = 0.319, *n* = 5) (Figures 1e,f,m and 2e,f,m).

A reduction in the area covered by NF160+ fibers in CB1R<sup>-/-</sup> was detected in the CA3 area in the stratum radiatum (NF160 = CB1R<sup>+/+</sup>: 10.67 ± 1.91 vs. CB1R<sup>-/-</sup>: 7.05 ± 1.24, unpaired *t* test, *t*(8) = 3.54, *p* = 0.045, *n* = 5) (Figure 1g,h,n), with no changes for NF200 (NF200 = CB1R<sup>+/+</sup>: 16.32 ± 6.53 vs. CB1R<sup>-/-</sup>: 11.11 ± 2.48, unpaired *t* test, *t*(8) = 1.66, *p* = 0.096, *n* = 5) (Figure 2g,h,n).

Finally, no significant changes were observed in the area of NF160+ and NF200+ fibers in the molecular layer of the DG (NF160 = CB1R<sup>+/+</sup>: 14.54 ± 3.49 vs. CB1R<sup>-/-</sup>: 11.93 ± 4.89, unpaired *t* test, *t*(8) = 0.96, *p* = 0.473, *n* = 5 and NF200 = CB1R<sup>+/+</sup>: 20.97 ± 3.67 vs. CB1R<sup>-/-</sup>: 19.77 ± 3.28, unpaired *t* test, *t*(8) = 0.54, *p* = 0.289, *n* = 5) (Figures 1i,j,o and 2i,j,o).

WB analyses in CB1R<sup>-/-</sup> showed a reduction in the expression of NF160 in whole hippocampus with respect to CB1R<sup>+/+</sup> (NF160 = CB1R<sup>+/+</sup>: 1.00 ± 0.28 vs. CB1R<sup>-/-</sup>: 0.46 ± 0.34, unpaired *t* test, *t*(8) = 2.69, *p* = 0.027, *n* = 5) (Figure 1k), which may correspond to the changes observed in each of the hippocampal areas studied. No differences were observed in total NF200 expression in the hippocampus (NF200 = CB1R<sup>+/+</sup>: 1.00 ± 0.24 vs. CB1R<sup>-/-</sup>: 0.92 ± 0.23, unpaired *t* test, *t*(8) = 0.52, *p* = 0.616, *n* = 5) (Figure 2k).

### 3.2 | MAP2 expression analysis

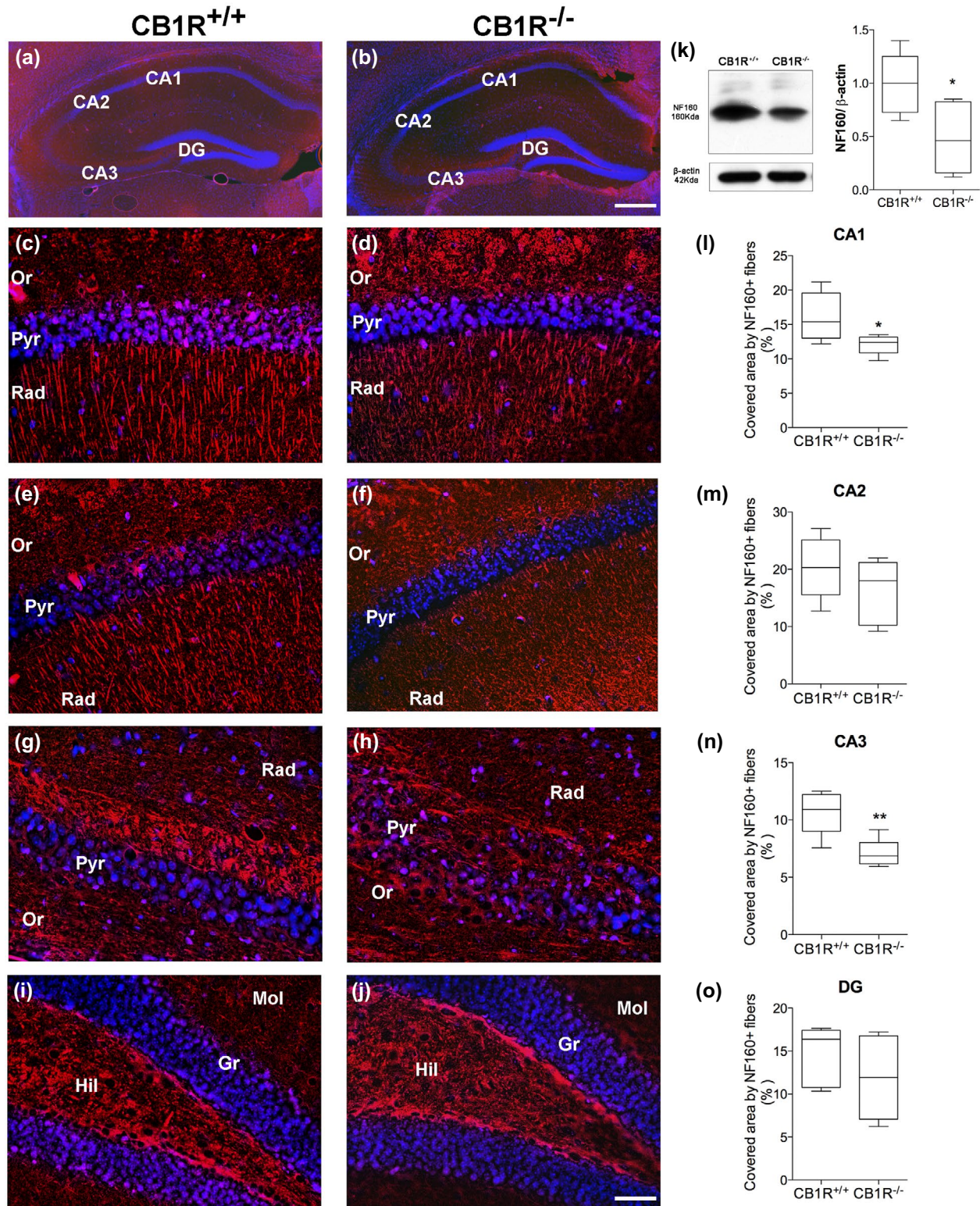
MAP2 is a microtubule-associated protein whose expression is restricted to the neuronal cell body and dendrites (Dehmelt & Halpain, 2005). The MAP2 protein participates in processes such as arborization and dendritic extension, being commonly used as a specific marker of dendrites.

The stratum radiatum of the hippocampal areas CA1 and CA2 revealed a reduction in the area covered by MAP2+ fibers in CB1R<sup>-/-</sup> mice as compared to CB1R<sup>+/+</sup> mice (CA1 = CB1R<sup>+/+</sup>: 30.85 ± 6.33 vs. CB1R<sup>-/-</sup>: 23.70 ± 2.42, unpaired *t* test, *t*(8) = 2.35, *p* = 0.046, *n* = 5 and CA2 = CB1R<sup>+/+</sup>: 14.44 ± 3.92 vs. CB1R<sup>-/-</sup>: 9.47 ± 1.49, unpaired *t* test, *t*(8) = 2.64, *p* = 0.029, *n* = 5) (Figure 3a-m), while the stratum radiatum of CA3 and the molecular layer of the DG rendered no differences between groups (CA3 = CB1R<sup>+/+</sup>: 14.77 ± 4.31 vs. CB1R<sup>-/-</sup>: 15.29 ± 3.37, unpaired *t* test, *t*(8) = 0.21, *p* = 0.83, *n* = 5 and GD = CB1R<sup>+/+</sup>: 25.88 ± 4.32 vs. CB1R<sup>-/-</sup>: 23.93 ± 3.15, unpaired *t* test, *t*(8) = 0.81, *p* = 0.4395, *n* = 5) (Figure 3g,h,n,i,j,o).

WB analyses in whole hippocampus showed a reduction in MAP2 expression in CB1R<sup>-/-</sup> mice compared to CB1R<sup>+/+</sup> (MAP2 = CB1R<sup>+/+</sup>: 1.00 ± 0.33 vs. CB1R<sup>-/-</sup>: 0.49 ± 0.26, unpaired *t* test, *t*(8) = 2.71, *p* = 0.026, *n* = 5) (Figure 3k), which could correspond with the decrease observed in morphological analyses in the area covered by MAP2+ fibers in the areas CA1 and CA2.

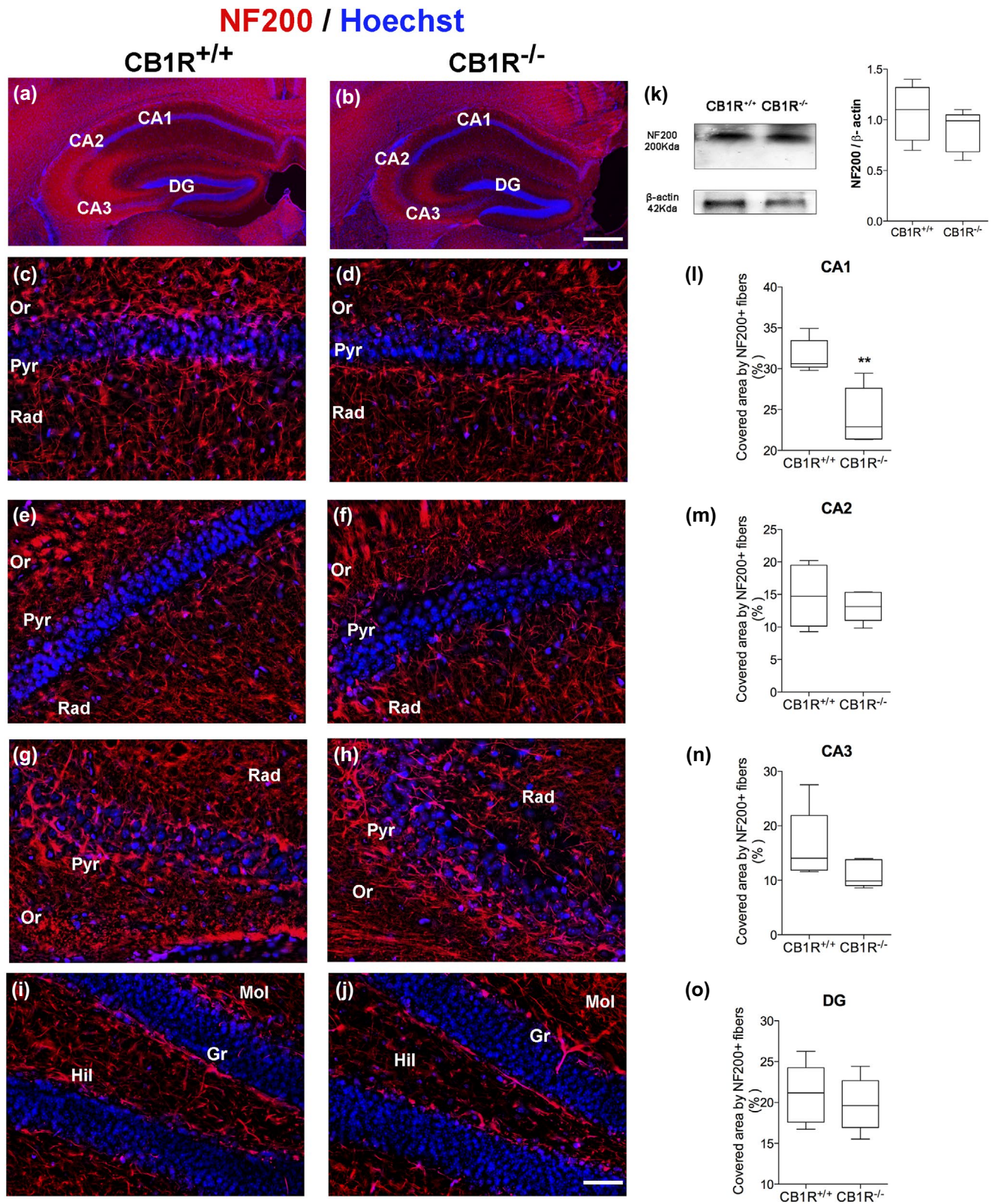


## NF160 / Hoechst



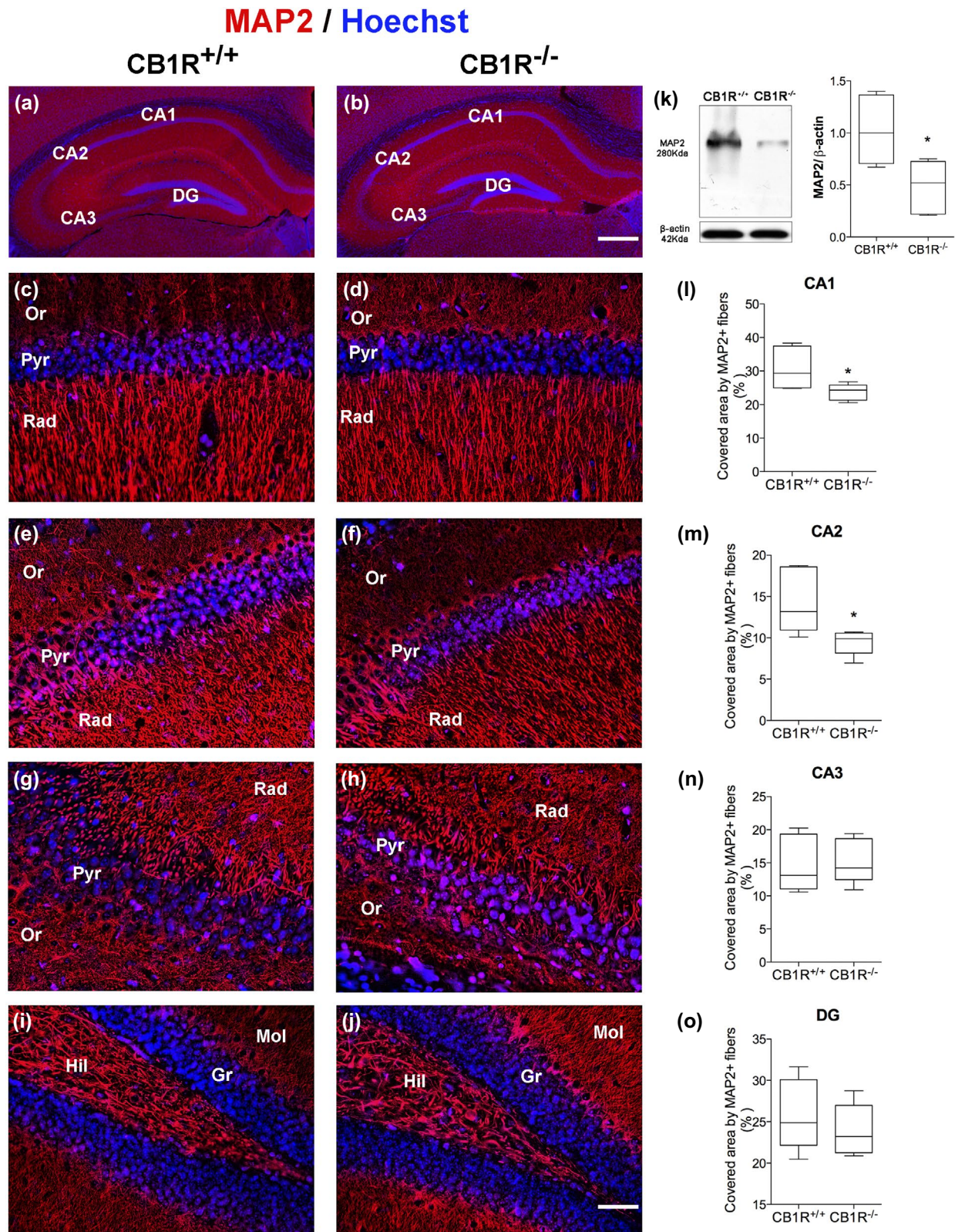
**FIGURE 1** NF160 expression in the hippocampus. Immunofluorescence and semiquantification of NF160+ fiber area in CA1 (c, d, and l), CA2 (e, f and m), CA3 (g, h, and n) and the DG (i, j and o) (bars = 30 μm) from CB1R<sup>+/+</sup> (a) and CB1R<sup>-/-</sup> (b) (bars = 300 μm). Unpaired t-test, columns represent mean ± SD (n = 5/group), \*p < 0.05. CA, cornus ammonis; DG, dentate gyrus; Gr, granular layer; Hil, hilus; Mol, molecular layer; Or, stratum oriens; Pyr, pyramidal layer; Rad, stratum radiatum. (k) NF160 protein WB in the hippocampus. Relative optical density of NF160/β-actin. Unpaired t-test, bars represent mean ± SD (n = 5/group), \*p < 0.05





**FIGURE 2** NF200 expression in the hippocampus. Immunofluorescence and semiquantification of NF200+ fiber area in CA1 (c, d, and l), CA2 (e, f, and m), CA3 (g, h, and n), and the DG (i, j, and o). (bars = 30  $\mu$ m) from CB1R<sup>+/+</sup> (a) and CB1R<sup>-/-</sup> (b) (bars = 300  $\mu$ m). Unpaired *t*-test, columns represent mean  $\pm$  SD (*n* = 5/group), \**p* < 0.05. CA, cornus ammonis; DG, dentate gyrus; Gr, granular layer; Hil, hilus; Mol, molecular layer; Or, stratum oriens; Pyr, pyramidal layer; Rad, stratum radiatum. (k) NF200 protein WB in the hippocampus. Relative optical density of NF200/ $\beta$ -actin. Unpaired *t*-test bars represent mean  $\pm$  SD (*n* = 5/group), \**p* < 0.05





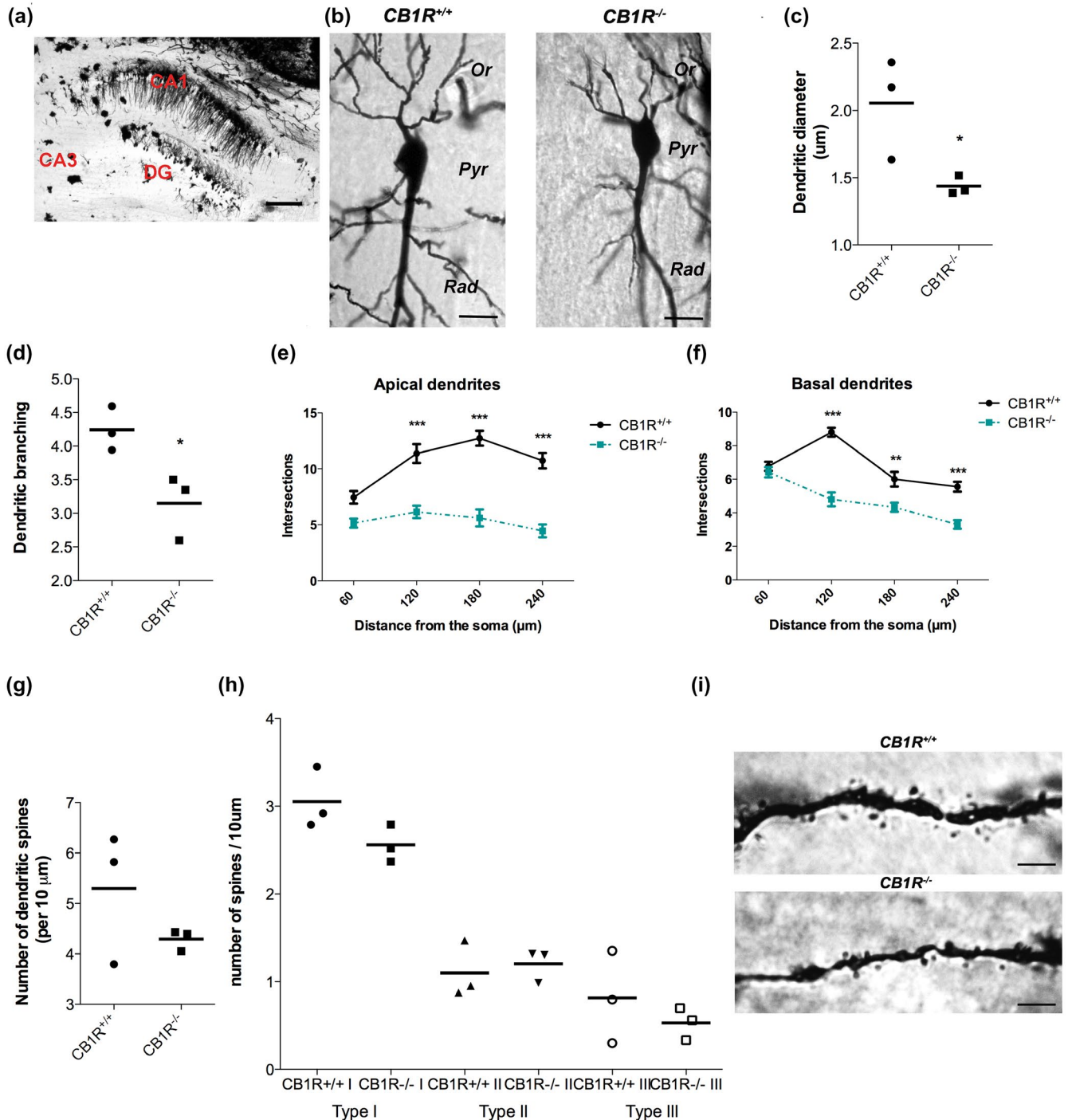
**FIGURE 3** MAP2 expression in the hippocampus. Immunofluorescence and semiquantification of MAP2+ fiber area in CA1 (c, d, and l), CA2 (e, f, and m), CA3 (g, h, and n), and the DG (i, j, and o) (bars = 30  $\mu$ m) from CB1R<sup>+/+</sup> (a) and CB1R<sup>-/-</sup> (b) (bars = 300  $\mu$ m). Unpaired t-test, columns represent mean  $\pm$  SD ( $n = 5$ /group), \* $p < 0.05$ , \*\* $p < 0.01$ . CA, cornus ammonis; DG, dentate gyrus; Gr, granular layer; Hil, hilus; Mol, molecular layer; Or, stratum oriens; Pyr, pyramidal layer; Rad, stratum radiatum. (k) MAP2 protein WB in the hippocampus. Relative optical density MAP2/ $\beta$ -actin. Unpaired t-test, bars represent mean  $\pm$  SD ( $n = 5$ /group), \* $p < 0.05$



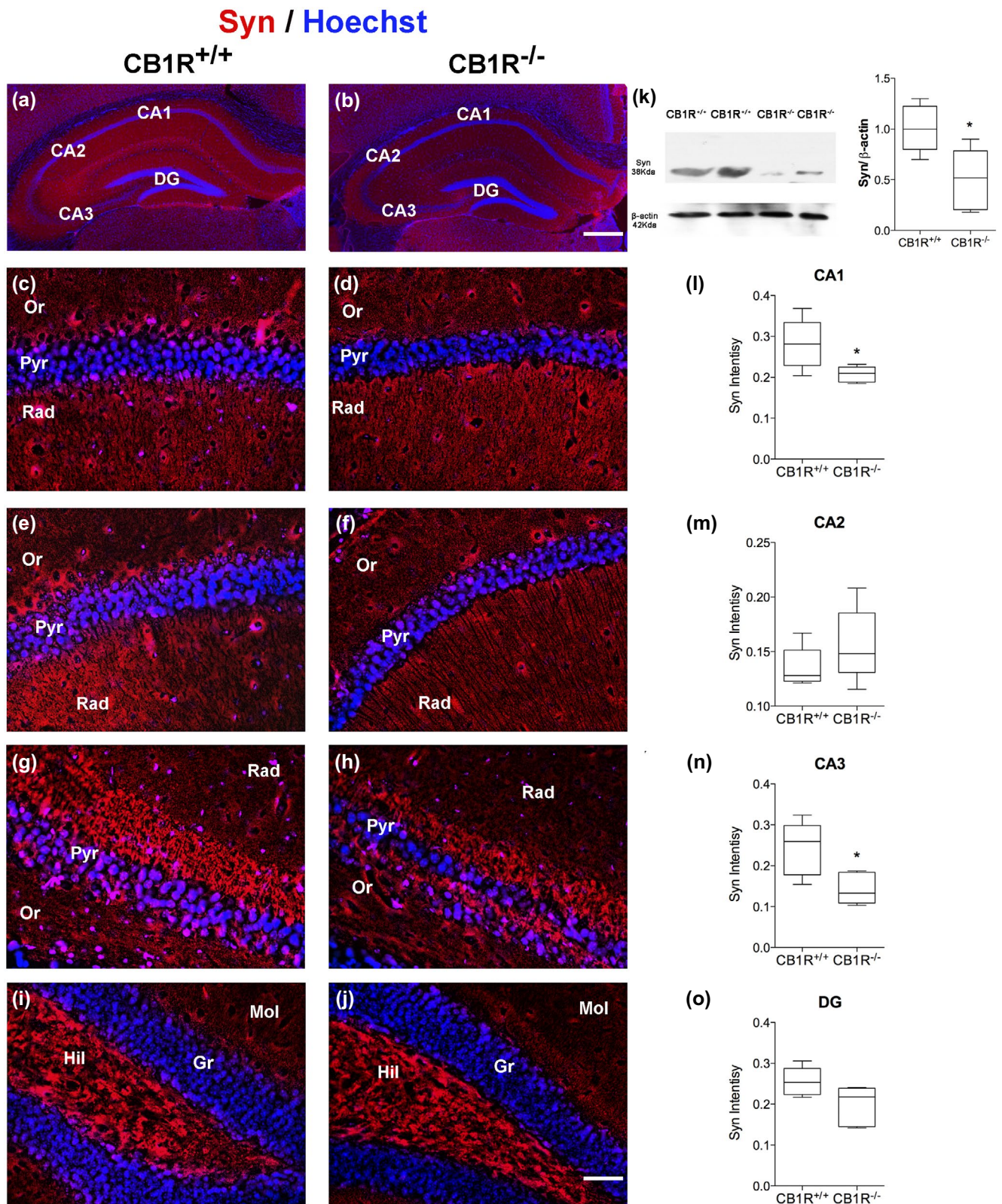
### 3.3 | Neuronal morphological analysis

Studies on dendritic intersections and spine density in CA1 hippocampal area, using Golgi technique, showed a significant effect of genotype in the number of dendrites (A two-way ANOVA test number of dendrites  $\times$  genotype; apical dendrites =  $F(1, 16)$ :

33.38,  $n = 3$ ,  $p = 0.01$ ; basal dendrites =  $F(1, 16)$ : 7.04,  $n = 3$ ,  $p < 0.01$ ) (Figure 4a,b,e,f) these data are correlated with the reduction in dendritic branching in  $CB1R^{-/-}$  ( $CB1R^{+/+}$ :  $4.24 \pm 0.327$  vs.  $CB1R^{-/-}$ :  $3.51 \pm 0.48$ , unpaired  $t$  test,  $t(4) = 3.23$ ,  $p = 0.003$ ,  $n = 3$ ) (Figure 4b,d). However,  $CB1R$  deficiency induce a reduction in dendritic diameter ( $CB1R^{+/+}$ :  $2.05 \pm 0.37$  vs.  $CB1R^{-/-}$ :  $1.43 \pm 0.06$ ,



**FIGURE 4** Neuronal arborization and dendritic spines in CA1 pyramidal neurons. Golgi staining representative images (a, b, and i; bars = 200, 30, and 5  $\mu\text{m}$ , respectively), dendritic diameter (c), dendritic branching (d). Unpaired  $t$ -test, columns represent mean  $\pm$  SD ( $n = 3$ /group), \* $p < 0.05$ . Number of apical and basal dendrites of pyramidal neuron (e and f) Two-way ANOVA and Bonferroni posttest, columns represent mean  $\pm$  SD ( $n = 3$ /group), \*\* $p < 0.01$ , \*\*\* $p < 0.001$ . Number of dendritic spines per  $100 \mu\text{m}$  (g), types of spines (h). Unpaired  $t$ -test, columns represent mean  $\pm$  SD ( $n = 3$ /group), \* $p < 0.05$



**FIGURE 5** Syn expression in the hippocampus. Immunofluorescence and semiquantification of Syn+ intensity in CA1 (c- d and l), CA2 (e- f and m), CA3 (g- h and n), and the DG (i- j and o) (bars = 30  $\mu$ m) from CB1R<sup>+/+</sup> (a) and CB1R<sup>-/-</sup> (b) (bars = 300  $\mu$ m). Unpaired t-test, columns represent mean  $\pm$  SD ( $n = 5$ /group), \* $p < 0.05$ . CA, cornus ammonis; DG, dentate gyrus; Gr, granular layer; Hil, hilus; Mol, molecular layer; Or, stratum oriens; Pyr, pyramidal layer; Rad, stratum radiatum. (k) Syn protein WB in the hippocampus. Relative optical density Syn/ $\beta$ -actin. Unpaired t-test, bars represent mean  $\pm$  SD ( $n = 5$ /group), \* $p < 0.05$



unpaired *t* test,  $t(4) = 2.28$ ,  $p = 0.048$ ,  $n = 3$  (Figure 4c,i) but did not alter the number of dendritic spines (CB1R<sup>+/+</sup>:  $5.29 \pm 1.31$  vs. CB1R<sup>-/-</sup>:  $4.29 \pm 0.20$ ,  $t(4) = 1.29$ ,  $p = 0.26$ ,  $n = 3$ ) (Figure 4g,i) also no difference in spine type were observed (Type I: CB1R<sup>+/+</sup>  $3.05 \pm 0.21$  vs. CB1R<sup>-/-</sup>  $2.56 \pm 0.21$ , unpaired *t* test,  $t(4) = 2.08$ ,  $p = 0.10$ ,  $n = 3$ ; Type II: CB1R<sup>+/+</sup>:  $1.1 \pm 0.32$  vs. CB1R<sup>-/-</sup>  $1.2 \pm 0.18$ , unpaired *t* test,  $t(4) = 0.48$ ,  $p = 0.65$ ,  $n = 3$ ; Type III: CB1R<sup>+/+</sup>:  $0.81 \pm 0.52$  vs. CB1R<sup>-/-</sup>:  $0.53 \pm 0.18$ , unpaired *t* test,  $t(4) = 0.88$ ,  $p = 0.42$ ,  $n = 3$ ) (Figure 4h,i).

### 3.4 | Synaptophysin expression analysis

Synaptophysin is a protein involved in vesicular formation and recycling, which is widely used as a marker for synaptic vesicles.

A reduction in Syn intensity was observed in the stratum radiatum of CA1 and CA3 areas of the hippocampus in CB1R<sup>-/-</sup> (Figure 5a,b; CA1 = CB1R<sup>+/+</sup>:  $0.28 \pm 0.06$  vs. CB1R<sup>-/-</sup>:  $0.20 \pm 0.01$ , unpaired *t* test,  $t(8) = 2.62$ ,  $p = 0.036$ ,  $n = 5$  and CA3 = CB1R<sup>+/+</sup>:  $0.24 \pm 0.06$ , vs. CB1R<sup>-/-</sup>:  $0.14 \pm 0.03$ , unpaired *t* test,  $t(8) = 2.90$ ,  $p = 0.019$ ,  $n = 5$ ) (Figure 5c,d,l,g,h,n), while no differences were detected in the stratum radiatum of CA2 area or DG (CA2 = CB1R<sup>+/+</sup>:  $0.13 \pm 0.018$  vs. CB1R<sup>-/-</sup>:  $0.15 \pm 0.03$ , unpaired *t* test,  $t(8) = 1.21$ ,  $p = 0.258$ ,  $n = 5$  and DG = CB1R<sup>+/+</sup>:  $0.25 \pm 0.034$ ; CB1R<sup>-/-</sup>:  $0.819 \pm 0.049$ , unpaired *t* test,  $t(8) = 2.17$ ,  $p = 0.061$ ,  $n = 5$ ) (Figure 5e,f,m,i,j,o).

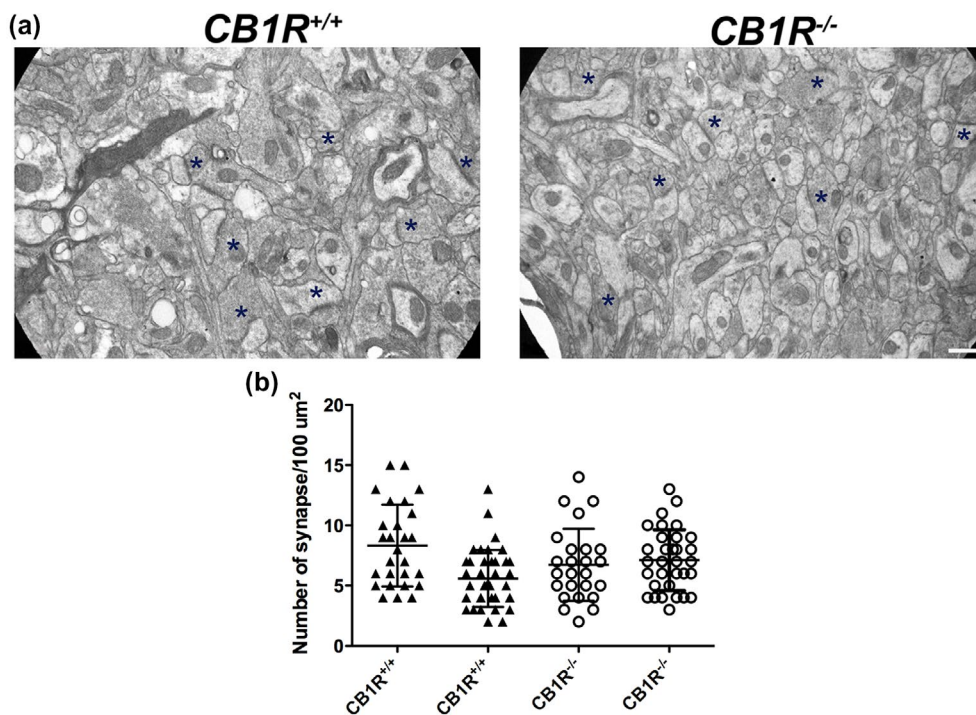
Syn protein content in the whole hippocampus was lower in CB1R<sup>-/-</sup> mice than that in CB1R<sup>+/+</sup> (Syn = CB1R<sup>+/+</sup>:  $1.00 \pm 0.22$  vs. CB1R<sup>-/-</sup>:  $0.49 \pm 0.29$ , unpaired *t* test,  $t(8) = 3.13$ ,  $p = 0.014$ ,  $n = 5$ ) (Figure 5k).

### 3.5 | Synaptic ultrastructural analysis

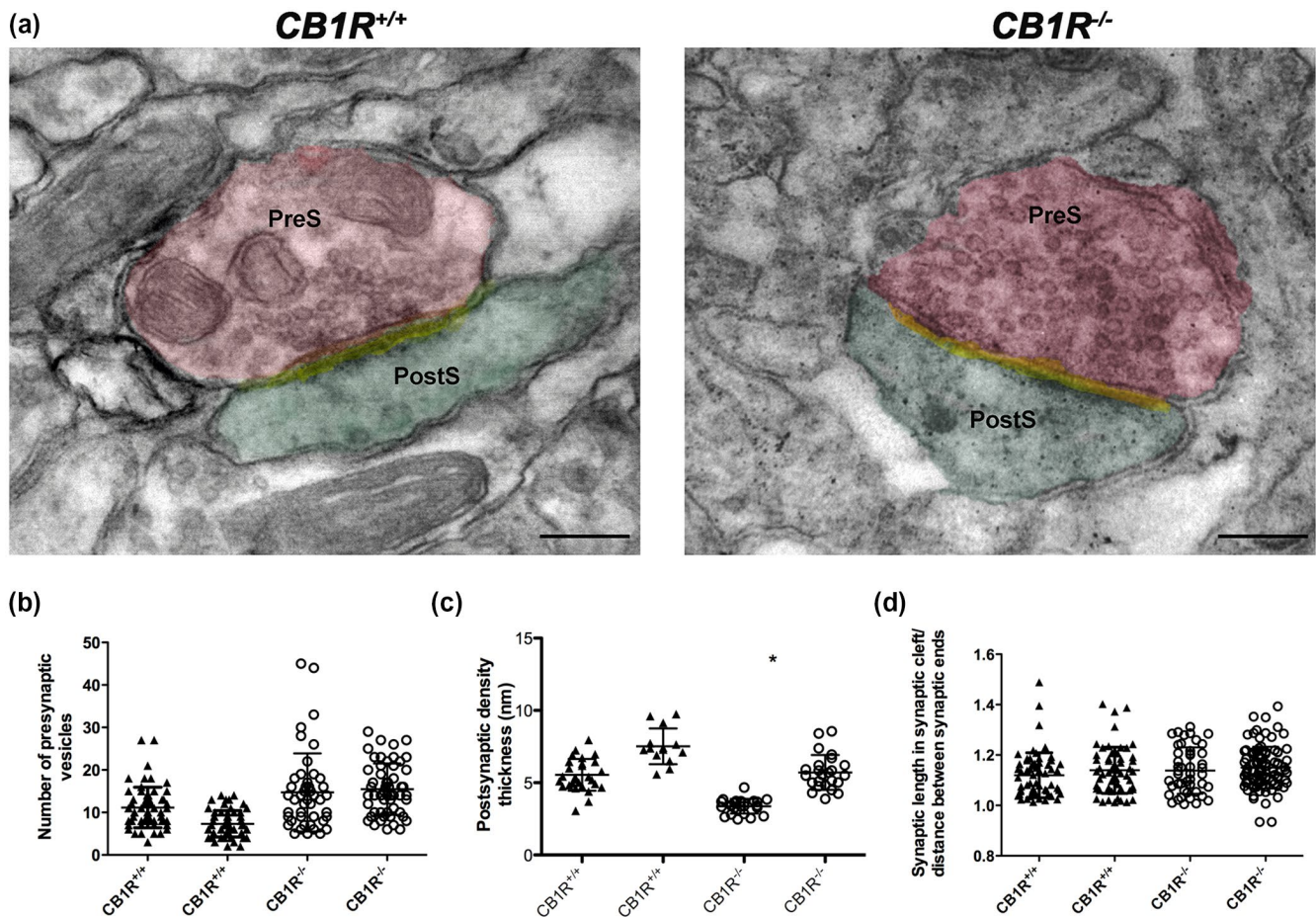
As observed in electron microscopy studies, CB1R deficiency did not alter the number of synapses in the stratum radiatum of CA1 area of the hippocampus (CB1R<sup>+/+</sup>:  $6.02 \pm 2.20$  vs. CB1R<sup>-/-</sup>:  $6.90 \pm 2.09$ , unpaired *t* test,  $t(2) = 0.40$ ,  $p = 0.724$ ,  $n = 2$ ) (Figure 6a,b). Regarding synaptic morphology, no differences were found in the number of synaptic vesicles (CB1R<sup>+/+</sup>:  $7.68 \pm 2.91$  vs. CB1R<sup>-/-</sup>:  $13.76 \pm 4.70$ , unpaired *t* test,  $t(2) = 1.15$ ,  $p = 0.260$ ,  $n = 2$ ) or synaptic curvature (CB1R<sup>+/+</sup>:  $1.13 \pm 0.09$  vs. CB1R<sup>-/-</sup>:  $1.15 \pm 0.08$ , unpaired *t* test,  $t(2) = 0.12$ ,  $p = 0.885$ ,  $n = 2$ ) (Figure 7a,b,d). However, CB1R<sup>-/-</sup> evidenced a decrease in the thickness of the postsynaptic density (CB1R<sup>+/+</sup>:  $6.55 \pm 1.46$  vs. CB1R<sup>-/-</sup>:  $3.19 \pm 1.25$ , unpaired *t* test,  $t(2) = 3.02$ ,  $p = 0.038$ ,  $n = 2$ ) (Figure 7a,c).

## 4 | DISCUSSION

The physiological importance of the endocannabinoid system lies in its role as a neuromodulator of the synaptic activity, participating in central nervous system development, synaptic plasticity, and neurogenesis (Galve-Roperh, Palazuelos, Aguado, & Guzman, 2009; Oudin, Hobbs, & Doherty, 2011). Several reports showed that a disruption on endocannabinoid signaling induce alteration in plasticity (Heifets & Castillo, 2009). Adult CB1R<sup>-/-</sup> mice showed a decreased anandamide (AEA) in hippocampus produced by an increase in AEA membrane transporter (AMT) and its hydrolase, fatty acid amide hydrolase (FAAH) (Maccarrone et al., 2002). Also, repeated



**FIGURE 6** Synapses in stratum radiatum of CA1 area. Transmission electron microscopy (a) (bars = 0.5 μm). Asterisks represent synapses. Number of synapses per 100 μm<sup>2</sup> (b). Unpaired *t*-test, columns represent mean ± SD ( $n = 2$ /group), \* $p < 0.05$



**FIGURE 7** Synaptic ultrastructure in stratum radiatum of CA1 area. Transmission electron microscopy of synapses (a). Number of presynaptic vesicles (b). Postsynaptic density thickness (c). Relation between synaptic length in the synaptic cleft and distance between the synaptic ends (d) (bars = 200 nm). Unpaired *t*-test, columns represent mean  $\pm$  SD ( $n = 2$ /group), \* $p < 0.05$ . PreS, presynaptic neuron; PostS, postsynaptic neuron

FAAH inhibitor treatment reduced neuronal proliferation with decreased expression of CB1R in the hippocampus (Rivera et al., 2015). Moreover, inactivation of MAGL induce increased spine density in pyramidal neuron and improves basal synaptic transmission and long-term synaptic plasticity in the hippocampus (Chen et al., 2012).

All these evidences show that the dysregulation of the endocannabinoid system induces changes in neuronal plasticity. Here we focus on the effect of CB1 deficiency on neuronal and synaptic architecture. Particularly, the changes in neuronal plasticity via CB1R have been associated with its participation in the modulation of the remodeling of the cytoskeleton both in neuronal development and in adult life (Njoo et al., 2015; Tortoriello et al., 2014; Zimmermann et al., 2018). These changes involve alterations in the stability of microtubules and in the dynamics of the actin filaments and are correlated with the determination of the axonal diameter and the growth of neurites.

CB1R is expressed in the pyramidal cell layer of the CA1-CA3 region and highly expressed in the stratum radiatum of these areas and in the molecular layer of the DG. Furthermore, inputs from glutamatergic pyramidal cells of the entorhinal cortex terminate in all areas of the hippocampus and hypothalamic glutamatergic

projections terminate mostly in the molecular layer (Steindel et al., 2013). Specifically, CB1R is present in sites where basket cell terminals surround pyramidal and granular cells. Lower CB1R expression is observed in the interconnecting network of inhibitory interneurons innervating the dendritic tree of pyramidal or granular cell bodies in the molecular layer, radiatum, and oriens (Katona et al., 1999).

Steindel and colleagues showed that CB1R deficiency in cortical glutamatergic neurons (Glu-CB1-KO mice) resulted in no apparent change in the CB1R expression pattern in the hippocampal formation but specific deletion of CB1R from GABAergic neurons induced a loss of CB1R protein in the hippocampus, especially in the stratum radiatum of the CA1-CA3 regions, and in the outer two-thirds of the molecular layer of the DG (Steindel et al., 2013).

Taking in account that CA1-CA3 hippocampal areas receive principally forebrain GABAergic inputs, changes observed in this study could due to CB1R deficiency in local hippocampal neurons in addition to CB1R deficiency in neuron inputs from other brain areas.

$CB1R^{-/-}$  mice exhibit behavioral alterations similar to those described in animal models of stress or depression, as endocannabinoid signaling in corticolimbic structures such as the prefrontal cortex,



amygdala, and hippocampus plays a key role in depression and stress response (Lee, Hill, & Lee, 2016; Smaga, Bystrowska, Gawliński, Przegaliński, & Filip, 2014). CB1R deficiency produce alterations in learning and memory process and deficit in long-term plasticity in the hippocampus, the principal area involved in cognitive function (Bannerman et al., 2014; Marsicano & Lafenetre, 2009). Some of the behavioral changes in CB1R<sup>-/-</sup> are associated with changes in the function and morphology of structures of the limbic system and the serotonergic system, among others.

Low dose of delta-9-tetrahydrocannabinol (THC) improves altered cognitive functions and increased hippocampal spine density and enhances the expression of synaptic proteins of 12-18 months aged mice (Bilkei-Gorzo et al., 2017). Young CB1R<sup>-/-</sup> mice do not show memory or cognitive impairment when compared with CB1R<sup>+/+</sup> mice. In contrast, decline in cognitive functions and learning is accelerated in mature and older CB1R<sup>-/-</sup> mice. These modifications are also accompanied by neuronal loss in hippocampus (Albayram et al., 2011; Bilkei-Gorzo et al., 2005). Chronic treatment with CB1R cannabinoid agonist during adolescence produces a decrease in arborization and spine density in pyramidal neurons of the prefrontal cortex (Miller et al., 2018; Renard et al., 2016) and in granular neurons of the hippocampus in the adult (Rubino et al., 2009). Controversially, chronic treatment with THC during adulthood increases neuronal arborization in prefrontal cortex (Kolb, Gorny, Limebeer, & Parker, 2006; Kolb, Li, Robinson, & Parker, 2018).

In the present work, morphological changes in the hippocampus of CB1R<sup>-/-</sup> were evidenced as a decrease in the expression of cytoskeletal proteins, as well as lower dendritic arborization in the stratum radiatum of CA1 area, without changes in length. In addition, in the absence of CB1R, changes in synaptic structure were observed as lower thickness of postsynaptic density and a reduction in Syn levels. However, no changes were detected in the number of dendritic spines, synapses, or synaptic vesicles in neurons in the CA1 area of CB1R<sup>-/-</sup> mice.

The organization of the neuronal cytoskeleton plays an important role both in neuronal structure and functionality and synaptic plasticity. Neurofilaments are the major components of the neuronal cytoskeleton, and are present in dendrites and particularly abundant in the axon (Yuan et al., 2012). In turn, MAP2 is a protein that maintains the stability of neuronal microtubules and is present in dendrites, where it participates in processes such as dendritic arborization and dendritic extension (Dehmelt & Halpain, 2005).

The present report shows a decrease in NF160+ and NF200+ fibers in the stratum radiatum of the CA1 area, in NF160+ fibers in stratum radiatum of the CA3 area, and in MAP2+ fibers in the stratum radiatum of the CA1 and CA2 areas of the hippocampus of CB1R<sup>-/-</sup>. In contrast, no alterations were detected in the expression of neurofilaments or MAP2 in the molecular layer of the DG in the absence of CB1R. Changes in the expression of neurofilaments and MAP2 in the stratum radiatum of the hippocampus evidence the participation of CB1R in the establishment and maintenance of the neuronal cytoskeleton and reinforce the notion of CB1R participation in the

dynamics of the neuronal cytoskeleton. Changes in neuronal cytoarchitecture observed in the present study is in agreement with previous studies in different animal models with CB1R agonist, consider that long-term administration of cannabinoid agonists produces a downregulation of receptor (Nogueras-Ortiz & Yudowski, 2016; Sim-Selley, 2003; Sim-Selley et al., 2006; Wu et al., 2008). Treatment with THC reduces the expression of microtubules and microfilament in cell cultures (Tahir, Trogadis, Stevens, & Zimmerman, 1992; Wilson Jr., Tahir, Mechoulam, Zimmerman, & Zimmerman, 1996). Tagliaferro et al. (2006) have reported an increase in NF160, NF200, and MAP2 in the CA1 area induced by chronic treatment with WIN 55,212-2, a cannabinoid agonist. A proteomic study in hippocampi of animal exposed to THC shows altered expression of cytoskeleton and structural proteins (Quinn et al., 2008) and this may reflect modification in the microtubule cytoskeleton reported by cannabinoid treatment. In addition, a decrease in NF200 expression has been reported in both normal and pathological conditions in CB1R<sup>-/-</sup> (Jackson, Pryce, Diemel, Cuzner, & Baker, 2005). Taking into account previous studies and the results reported here, CB1R could play a fundamental role in the organization of the cytoskeleton, participating in the formation and stability of synaptic connections.

Dendrites are the most numerous postsynaptic elements of the central nervous system, and their length and branching are associated with neuronal plasticity. The neuronal branching was reduced in apical and basal dendrites in pyramidal neurons of CA1 area in CB1R<sup>-/-</sup>. Additionally, CA1 pyramidal neurons present lower dendritic branching and dendrite diameter in absence of CB1R. Moreover, a decrease was observed in the expression of MAP2 in the stratum radiatum of CA1 and CA2 areas, which showed that CB1R is participating in the stabilization of the dendritic cytoskeleton and participates in the process of arborization. Similar changes in morphology including a decrease in dendritic arborization and number of spines have been reported in a recent study where CB1R was specifically deleted in GABAergic neurons in the CA1 area of the hippocampus (Monory et al., 2015). The absence of CB1R specifically in new hippocampal neurons during adult life produces alterations in neuritogenesis as a reduction in dendritic length and a lower density of dendritic spines (Zimmermann et al., 2018).

However, in the present study no differences were observed in the number and type of dendritic spines, which depends mainly of actin microfilaments (Gordon-Weeks & Fournier, 2014). In addition, the decrease in the number of primary prolongations and the alterations in dendrite structure and the possible changes in the stability of microtubules determined by a decrease in MAP2 protein indicate morphological alterations due to CB1R deficiency.

Reduction of neuronal and dendritic arborization with lower dendritic diameter could alter dendrites, spine, and synapse stabilization in the absence of CB1R. The morphology of dendritic spine is related with different in synaptic function (Arellano, Benavides-Piccione, DeFelipe, & Yuste, 2007), changes in proportion of spine type correlate with different synaptic efficiency and also represent forms of synaptic plasticity in the hippocampus (Bourne & Harris, 2008; Popov et al., 2004). In the present work, we report a reduction

in dendritic branching and lower dendrite diameter but changes in proportion or number of dendritic spine were not observed, this is differential effect in structure of dendrite by deletion of CB1R and could be related with different participation in the process of dendritogenesis and spinogenesis.

The alterations observed in the neuronal cytoskeleton and the morphology of pyramidal neurons in the CA1 area of CB1R<sup>-/-</sup> could produce deficiencies in synaptic communication. In the present study, the analysis of neuronal connections in the stratum radiatum of CA1 area revealed no differences in the number of synapses between CB1R<sup>+/+</sup> and CB1R<sup>-/-</sup>. However, CB1R deficiency was seen to produce structural changes such as lower Syn expression and lower thickness of postsynaptic density.

The implication of CB1R in the synaptic structure has also been reported in the work of Tagliaferro et al. (2006), which shows an increase in the number of vesicles as a result of treatment with a cannabinoid agonist and hence supports the hypothesis of a direct effect of cannabinoids on synaptic stability (Karanian, Brown, Makriyannis, Kosten, & Bahr, 2005). More recently, Buceta et al. show that cerebellar lobules of CB1R<sup>-/-</sup> mice contain greater length of synapsis and lower vesicle density, representing ultrastructural adaptations in the absence of CB1R (Buceta et al., 2019). However, in the present work, the number of synaptic vesicles detected by electron microscopy did not show differences between CB1R<sup>-/-</sup> and CB1R<sup>+/+</sup>.

Since, Syn is not essential for neurotransmission but modulate the efficiency of the synaptic vesicle cycle (Gordon & Cousin, 2016). In Syn-deficient mice, hippocampal neurons exhibited defective synaptic vesicle endocytosis both during and after neuronal activity (Kwon & Chapman, 2011). Complete loss of Syn impairs learning and memory (Schmitt, Tanimoto, Seeliger, chaeffel, & Leube, 2009) but does not affect viability (Eshkind & Leube, 1995). Our data indicate that a decrease in Syn may not imply a lower number of synaptic vesicles, but a decrease in Syn expression in each synaptic vesicle. Stratum radiatum of CA3 area of CB1R<sup>-/-</sup> also showed a reduction in Syn expression as compared to CB1R<sup>+/+</sup>, which, together with the smaller area covered by NF160 fibers, may respond to the lower density of neurons reported in that area of adult CB1R<sup>-/-</sup> (Albayram et al., 2011; Bilkei-Gorzo et al., 2005). On the other hand, the decrease observed in the thickness of the postsynaptic density in the stratum radiatum of CA1 area of CB1R<sup>-/-</sup> could be related to the changes observed in cytoskeletal components. In other words, alterations in cytoskeletal dynamics may lead to changes in the interaction of actin filaments with postsynaptic density proteins (Gordon-Weeks & Fournier, 2014; Okabe, 2007). This type of morphological change in the CA1 area of the hippocampus has also been studied in rats under conditions of hypoxia, which also showed a decrease in the thickness of postsynaptic density (Von Lubitz & Diemer, 1983). Changes in postsynaptic density may reflect alterations in the density of postsynaptic receptors and in proteins associated with receptor-mediated signaling, which lead to defective neuronal communication. Therefore, CB1R deficiency can be thought to cause both presynaptic and postsynaptic changes

affecting synaptic transmission. Alterations in neuronal structural and synaptic vesicle protein in CA1-CA3 area (Tables S1 and S2) may be related to changes in synaptic communication of intrahippocampal connectivity observed in CB1R<sup>-/-</sup> (Bohme et al., 2000; Jacob et al., 2012) and deficit memory reported in mice evaluated in the present work (Mikics et al., 2006).

In sum, the present work demonstrates that CB1R deficiency produces changes at the morphological level in the hippocampus, one of the areas involved in learning, memory, and mood regulation. The reduction in the expression of cytoskeletal components of the different hippocampal areas of CB1R<sup>-/-</sup> evidences the participation of CB1R in neuronal cytoskeleton consolidation. Morphological and synaptic structure alterations in the CA1 area demonstrate the requirement of CB1R signaling for the correct establishment of synaptic connections. On the whole, the results in the present work provide evidence of the participation of CB1R in neuronal plasticity from a morphological analysis and provide further support for a large number of studies showing CB1R regulation of short- and long-term plasticity.

## Declaration of Transparency

The authors, reviewers and editors affirm that in accordance to the policies set by the *Journal of Neuroscience Research*, this manuscript presents an accurate and transparent account of the study being reported and that all critical details describing the methods and results are present.

## CONFLICT OF INTEREST

Authors declare no conflict of interest.

## AUTHOR CONTRIBUTIONS

Delia Soriano: Investigation, Methodology, Conceptualization, Formal Analysis, Writing – Original Draft Preparation. Marina Vacotto: Investigation. Alicia Brusco: Conceptualization, Funding Acquisition, Writing – Review & Editing. Writing – Review & Editing. Laura Caltana: Conceptualization, Funding Acquisition, Writing – Review & Editing.

## PEER REVIEW

The peer review history for this article is available at <https://publons.com/publon/10.1002/jnr.24694>.

## DATA AVAILABILITY STATEMENT

The data that support the findings of this study are available from the corresponding author upon reasonable request.

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## SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section.

**FIGURE S1** (a) PCR for CB1R and Neo showing different genotypes. (b) Western Blot of CB1R, showing no expression in CB1R<sup>-/-</sup>. (c) Immunohistochemistry of CB1R and NeuN showing no expression of CB1R in neurons in CB1R<sup>-/-</sup>. (d) Immunohistochemistry of CB1R and S100 showing no expression of CB1R in astrocytes in CB1R<sup>-/-</sup>

**FIGURE S2** Full Blot of NF160 (a), NF200 (b), MAP2 (c) and Syn (d)

**TABLE S1** Summary of neuronal changes in hippocampus produced by CB1R deletion

**TABLE S2** Summary of synaptic changes in hippocampal CA1 area produced by CB1R deletion

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