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EDITED BY

Abel Santamaria,
Manuel Velasco Suárez Instituto
Nacional de Neurología y
Neurocirugía, Mexico

REVIEWED BY

Faheem Ullah,
University of Science Malaysia,
Malaysia
Maria L. De Ceballos,
Spanish National Research Council
(CSIC), Spain

*CORRESPONDENCE

Simon M. Ametamey
simon.ametamey@pharma.ethz.ch
Ruiqing Ni
ruiqing.ni@uzh.ch

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Evaluation of cannabinoid type 2 receptor expression and pyridine-based radiotracers in brains from a mouse model of Alzheimer's disease

Vasil Kecheliev¹, Francesco Spinelli², Adrienne Herde²,
Ahmed Haider², Linjing Mu^{2,3}, Jan Klohs⁴,
Simon M. Ametamey^{2*} and Ruiqing Ni^{1,4*}

¹Institute for Regenerative Medicine, University of Zurich, Zurich, Switzerland, ²Department of Chemistry and Applied Biosciences, ETH Zurich, Zurich, Switzerland, ³Department of Nuclear Medicine, University Hospital Zurich, Zurich, Switzerland, ⁴Institute for Biomedical Engineering, University of Zurich and ETH Zurich, Zurich, Switzerland

Neuroinflammation plays an important role in the pathophysiology of Alzheimer's disease. The cannabinoid type 2 receptor (CB₂R) is an emerging target for neuroinflammation and therapeutics of Alzheimer's disease. Here, we aim to assess the alterations in brain CB₂R levels and evaluate novel CB₂R imaging tracers in the arcA β mouse model of Alzheimer's disease amyloidosis. Immunohistochemical staining for amyloid- β deposits (6E10), microgliosis (anti-Iba1 and anti-CD68 antibodies), astrocytes (GFAP) and the anti-CB₂R antibody was performed on brain slices from 17-month-old arcA β mice. Autoradiography using the CB₂R imaging probes [¹⁸F]RoSMA-18-d6, [¹¹C]RSR-056, and [¹¹C]RS-028 and mRNA analysis were performed in brain tissue from arcA β and non-transgenic littermate (NTL) mice at 6, 17, and 24 months of age. Specific increased CB₂R immunofluorescence intensities on the increased number of GFAP-positive astrocytes and Iba1-positive microglia were detected in the hippocampus and cortex of 17-month-old arcA β mice compared to NTL mice. CB₂R immunofluorescence was higher in glial cells inside 6E10-positive amyloid- β deposits than peri-plaque glial cells, which showed low background immunofluorescence in the hippocampus and cortex of 17-month-old arcA β mice. *Ex vivo* autoradiography showed that the specific binding of [¹⁸F]RoSMA-18-d6 and [¹¹C]RSR-056 was comparable in arcA β and NTL mice at 6, 17, and 24 months of age. The level of *Cnr2* mRNA expression in the brain was not significantly different between arcA β and NTL mice at 6, 17, or 24 months of age. In conclusion, we demonstrated

pronounced specific increases in microglial and astroglial CB₂R expression levels in a mouse model of AD-related cerebral amyloidosis, emphasizing CB₂R as a suitable target for imaging neuroinflammation.

KEYWORDS

Alzheimer's disease, positron emission tomography, astrocyte, microglia, cannabinoid type 2 receptor (CB₂R), neuroinflammation, autoradiography

Introduction

Abnormal accumulation of amyloid-beta (A β) aggregates in Alzheimer's disease (AD) leads to a cascade of pathophysiological changes, including neuroinflammation, microvascular alterations, synaptic dysfunction, and neuronal loss. Neuroinflammation, including gliosis and increased levels of complements, cytokines, and chemokines, plays an important role in the development of AD (Rapic et al., 2013; Heneka et al., 2015; López et al., 2018; Xin et al., 2020). Microglia are resident macrophages in the central nervous system (CNS) that are important for maintaining brain homeostasis (Heneka et al., 2015) but have also been implicated in the pathophysiology of AD (Deczkowska et al., 2018; Kiani Shabestari et al., 2022). Increased numbers of astrocytes and microglia were observed in the vicinity of A β plaques in post-mortem AD mouse model brains and patients with AD (Heneka et al., 2015). Recent single-cell sequencing transcriptomics for disease-associated microglia (DAM) indicates the presence of transcriptionally distinct and neurodegeneration-specific microglial profiles with potential significance in AD signatures, including TREM2, CD33, and ApoE (Keren-Shaul et al., 2017; Deczkowska et al., 2018; Song and Colonna, 2018; Grubman et al., 2021).

Positron emission tomography (PET) ligands for detecting neuroinflammation microgliosis and astrocytosis in AD for understanding the disease mechanism. Early neuroinflammation has been reported in AD and in amyloidosis animal models (Rodriguez-Vieitez et al., 2015; Kreisl et al., 2020; Biechele et al., 2021; Ni et al., 2021b). Among these, PET ligands for translocator protein (TSPO) are the most widely used for detecting neuroinflammation. Previous TSPO PET imaging studies have shown microglial activation preceding A β deposition in several animal models, such as APP23, J20, APPSL70, App^{NL-G-F}, and PS2APP mice (Sacher et al., 2019; Biechele et al., 2021). However, limitations in the complex cellular locations, polymorphisms, and non-specific binding of TSPO and whether TSPO measures microglial proliferation or activation remain to be addressed (Leng and Edison, 2021; Zhou et al., 2021). Novel specific PET tracers for visualizing microgliosis, especially the DAM subtype, are highly desired.

Cannabinoid type 2 receptors (CB₂Rs) are mainly expressed by immune cells, including monocytes and macrophages. In the

CNS, CB₂Rs are mainly expressed on microglia at low levels under physiological conditions and are upregulated in acute inflammatory conditions, such as ischemic stroke (Cristino et al., 2020). CB₂Rs are essential to induce Toll-like receptor-mediated microglial activation (Reusch et al., 2022). Activation of CB₂R offers neuroprotective effects, such as reducing A β -induced neuronal toxicity (Köfalvi et al., 2016; Navarro et al., 2018; Wang et al., 2018b; Scheiner et al., 2019; Zhao et al., 2020), suppressing microglial activation (Ehrhart et al., 2005; Ramírez et al., 2005), restoring cognitive capacity (Wu et al., 2017), and ameliorating novel object recognition in animal models of amyloidosis (Li et al., 2019). Thus, CB₂R has been of therapeutic interest in AD (Köfalvi et al., 2016). However, the expression levels of CB₂R in animal models of AD amyloidosis have not been extensively characterized. CB₂R has been shown to be increased and involved in A β pathology in 5 \times FAD (López et al., 2018; Zhang and Chen, 2018) and J20 mouse models of AD amyloidosis (Koppel et al., 2014) but reduced in the brains of 3 \times Tg mice (with both A β and tau pathology) and in aging C57B6 mice (Wang et al., 2018a).

Several CB₂R ligands have been developed and evaluated (Ni et al., 2019b), including [¹¹C]NE40 (Vandeputte et al., 2012), [¹¹C]A-836339 (MDTC) (Pottier et al., 2017; Du et al., 2022), [¹⁸F]MA3 (Attili et al., 2019), [¹⁸F]FC0324 (Caillé et al., 2017), [¹⁸F]JHU94620 (Moldovan et al., 2016), [¹⁸F]LU13 (Gündel et al., 2022), [¹⁸F]DM102 (Modemann et al., 2022), [¹⁸F]CRA13 (Hassan et al., 2020), [¹¹C]RS-016 (Meletta et al., 2017), [¹¹C]RS-028 (Haider et al., 2018), [¹¹C]RSR-056 (Slavik et al., 2015), and [¹⁸F]RoSMA-18-d6 (Haider et al., 2020). Thus far, only one in-human *in vivo* CB₂R PET using [¹¹C]NE40 (Ahmad et al., 2016) in patients with AD and healthy controls has been reported, showing no group difference. Only the tracer [¹¹C]A-836339 has been evaluated in an AD animal model: Increased [¹¹C]A-836339 uptake was observed in the cortex, cerebellum and whole brain of J20 mice compared to wild-type mice (Savonenko et al., 2015); another [¹¹C]A-836339 microPET study showed that the uptake was blockable in the cortex of APP/PS1 mice (Horti et al., 2010).

The aim of the current study was to assess the alterations in CB₂R and distribution in the brain of the arcA β mouse model of AD amyloidosis and to evaluate the recently developed pyridine-derived CB₂R tracers [¹¹C]RS-028, [¹⁸F]RoSMA-18-d6, and [¹¹C]RSR-056, which exhibit

subnanomolar affinity and high selectivity toward CB₂R (Ni et al., 2021a).

Materials and methods

Animals

Twenty transgenic arcA β mice overexpressing the human APP695 transgene containing the Swedish (K670N/M671L) and Arctic (E693G) mutations under control of the prion protein promoter at 6, 17, and 24 months of age and 20 age-matched non-transgenic littermates (NTLs) of both sexes were used in this study (Knobloch et al., 2007; Merlini et al., 2011; Ni et al., 2022). The arcA β mouse model exhibits parenchymal plaque as well as cerebral amyloid angiopathy and shows impaired cerebrovascular functions (Ni et al., 2018, 2019a). Paper tissue and red mouse house (Tecniplast[®], Buguggiate VA, Italy) shelters were placed in cages for environmental enrichment. All experiments were performed in accordance with the Swiss Federal Act on Animal Protection and were approved by the Cantonal Veterinary Office Zurich ZH082/18.

For mRNA and autoradiography, arcA β and age-matched NTL mice at 6, 17, and 24 months of age were anesthetized under 5% isoflurane and decapitated. One brain hemisphere from arcA β mice and NTLs was immediately frozen in liquid nitrogen and stored at -80°C as described earlier (Ni et al., 2021a). The other half of the brain hemisphere was embedded in Tissue Tekv (Sakura[®] Finetek, Torrance, CA, USA), frozen, and stored at -80°C for autoradiography. For immunofluorescence staining, mice were perfused under ketamine/xylazine/acepromazine maleate anesthesia (75/10/2 mg/kg body weight, i.p. bolus injection) with ice-cold 0.1 M phosphate-buffered saline (PBS, pH 7.4, Sigma Aldrich, Burlington, VT, USA) and 4% paraformaldehyde in 0.1 M PBS (pH 7.4), fixed for 2 h in 4% paraformaldehyde (pH 7.4), and then stored in 0.1 M PBS (pH 7.4) at 4°C .

mRNA isolation and real-time polymerase chain reaction

Total mRNA isolation of the brain tissue from arcA β and age-matched NTL mice at 6, 17, and 24 months of age was performed according to the protocols of the Isol-RNA Lysis Reagent (5 Prime Sciences, Montreal, Canada) and the bead-milling TissueLyser system (Qiagen, Hilden, Germany) (Ni et al., 2021a). A QuantiTect[®] Reverse Transcription Kit (Qiagen) was used to generate cDNA. The primers (Microsynth, Balgach, Switzerland) used for quantitative polymerase chain reaction (qPCR) are summarized in **Supplementary Table 1**. Quantitation of *Cnr2* mRNA expression was performed with the DyNAmo Flash SYBR[®] Green qPCR Kit (Thermo Fisher Scientific, Waltham, MA, USA) using a 7900 HT Fast Real-Time

PCR System (Applied Biosystems, Waltham, MA, USA). The amplification signals were detected in real time, which permitted accurate quantification of the amounts of the initial RNA template over 40 cycles according to the manufacturer's protocol. All reactions were performed in duplicate within three independent runs, and each reaction was normalized against the expression of beta-actin. Quantitative analysis was performed using SDS Software (v2.4) and a previously described $2^{-\Delta\Delta\text{Ct}}$ quantification method (Livak and Schmittgen, 2001). The specificity of the PCR products of each run was determined and verified with SDS dissociation curve analysis.

Immunofluorescence

For immunohistochemical analysis, brain tissue from arcA β and age-matched NTL mice at 17 months of age was used. Coronal brain sections (40 μm) were cut around Bregma 0 to -2 mm and stained with anti-A β antibody 6E10, anti-ionized calcium-binding adapter 1 (Iba1) and anti-CD68 for microgliosis, GFAP for astrocytes and anti-CB₂R antibody as previously described (Kecheliev et al., 2022; **Supplementary Table 2**). The CB₂R antibody used in this study targets amino acid region 300–350 on murine CB₂R protein expressed by cells of hematopoietic origin. Sections were mounted with Prolong Diamond mounting media. Imaging occurred at $\times 20$ magnification using an Axio Observer Z1 slide scanner (Zeiss, Oberkochen, Germany) using the same acquisition settings for all brain slices and at $\times 63$ magnification using a Leica SP8 confocal microscope (Leica, Wetzlar, Germany). The images were analyzed by a person blinded to the genotype using QuPath and ImageJ (NIH, USA). The colocalization of CB₂R with plaque (6E10 channel), GFAP+ astrocytes or Iba1+ microglia in the cortex and hippocampus was determined on $\times 63$ -magnification images. The amount of CB₂R immunofluorescence within these masks was determined by measuring the mean CB₂R intensity as well as its integrated density (factor of area and average intensity).

Radiosynthesis and autoradiography

[¹⁸F]RoSMA-18-d6 (affinity $K_i = 0.8$ nM, CB₂R/CB₁R $> 12,000$), [¹¹C]RSR-056 and [¹¹C]RS-028 were synthesized and purified as described previously (Slavik et al., 2015; Haider et al., 2018, 2020; Ni et al., 2021a) and formulated with 5% ethanol in water. The molar activities were 156–194, 52.3, and 86.7–178 GBq/ μmol for [¹⁸F]RoSMA-18-d6, [¹¹C]RSR-056, and [¹¹C]RS-028, respectively. The radiochemical purity for all three radioligands was $>99\%$. Autoradiography was performed as described previously (Haider et al., 2020). Dissected mouse brains were embedded in Tissue Tek, cut into 10 μm thick sagittal sections on a cryostat (Cryo-Star HM-560MV; Microm, Thermo Scientific, Waltham,

MA, USA) and stored at -80°C . We first compared the percentage of specific binding of [^{18}F]RoSMA-18-d6, [^{11}C]RSR-056, and [^{11}C]RS-028 using brain tissue slices from the same arcA β mice at 17 months of age ($n = 3$). Next, brain tissue slices from the arcA β and age-matched NTL mice at 6 months of age ($n = 5, 6$), 17 months of age ($n = 4, 5$), and 24 months of age ($n = 5, 5$) were used in [^{18}F]RoSMA-18-d6 autoradiography. Brain tissue slices from the arcA β and age-matched NTL mice at 6 months of age ($n = 3, 3$), 17 months of age ($n = 3, 3$), and 24 months of age ($n = 3, 3$) were used in [^{11}C]RSR-056 autoradiography. Autoradiography binding data from the same mice was analyzed using correlation analysis.

For [^{18}F]RoSMA-18-d6 and [^{11}C]RSR-056 autoradiography, slices were thawed on ice and preconditioned in ice-cold buffer (pH 7.4) containing 50 mM TRIS Sigma Aldrich, Burlington, VT, USA, 5 mM MgCl_2 Sigma Aldrich, Burlington, VT, USA, and 0.1% fatty acid-free bovine serum albumin (BSA, Sigma Aldrich, Burlington, VT, USA). The tissue slices were dried and then incubated with 1 ml of the corresponding radioligand (0.5 nM [^{18}F]RoSMA-18-d6 and 2 nM [^{11}C]RSR-056) for 15 min at room temperature in a humidified chamber. For blockade conditions, the selective CB $_2$ R antagonist GW405833 (10 μM , Sigma Aldrich, Burlington, VT, USA) was added to the solution containing the radioligand. A standard was placed to calibrate the radioligand concentration for calculation of protein level. The slices were washed with ice-cold washing buffer (pH 7.4) containing 50 mM TRIS, 5 mM MgCl_2 , 0.1% fatty acid-free BSA, and ice-cold distilled water. For [^{11}C]RS-028 autoradiography, slices were thawed on ice and preconditioned in ice-cold buffer (pH 7.4) containing 50 mM TRIS, 5 mM MgCl_2 , 2.5 mM ethylenediaminetetraacetic acid (EDTA, Sigma Aldrich, Burlington, VT, USA) and 0.1% fatty acid-free BSA. The tissue slices were dried and then incubated with 1 ml of the [^{11}C]RS-028 (3 nM) for 15 min at room temperature in a humidified chamber. For blockade conditions, the selective CB $_2$ R antagonist GW405833 (10 μM) was added to the solution containing the radioligand. A standard was placed to calibrate the radioligand concentration for calculation of protein level. Specific binding was calculated as the difference between total and blockade condition. After drying, the slices were exposed to a phosphorimager plate (FUJIFILM, Tokyo, Japan) for 30 min, and the film was scanned in a BAS5000 reader (FUJIFILM).

Statistics

Group comparisons in multiple brain regions were performed by using two-way ANOVA with Sidak's *post-hoc* analysis (GraphPad Prism 9, GraphPad, San Diego, CA, USA). Comparisons for CB $_2$ R inside plaque, peri-plaque and parenchymal were performed by using one-way ANOVA with Tukey's *post-hoc* analysis. All data are presented as the mean \pm standard deviation. Significance was set at $*p < 0.05$.

Results

Increased cannabinoid type 2 receptor expression with proliferation of microglia and astrocytes in the brains of arcA β compared to non-transgenic littermate

First, the regional CB $_2$ R level, the cellular source and the expression of CB $_2$ R (mean immunofluorescence on the area occupied by the selected marker) were assessed in the brains of arcA β mice and NTL mice at 17 months of age. CB $_2$ R immunofluorescence intensity was increased approximately 4–10-fold in the cortex (4.45 ± 0.25 vs. 0.46 ± 0.25 , $p < 0.0001$), hippocampus (4.82 ± 0.10 vs. 0.95 ± 0.27 , $p < 0.0001$), and thalamus (1.87 ± 0.31 vs. 0.41 ± 0.05 , $p < 0.0001$) of arcA β mice compared to NTL mice at 17 months of age ($n = 3$ per group) (Figures 1, 2A,B). The background signal of CB $_2$ R is low in the parenchyma (outside astrocytes/microglia) (Figures 1, 2H). CB $_2$ R signal density was calculated as the factor of average intensity and area of fluorescent image pixels.

Colocalization analysis indicated that CB $_2$ R signal density was upregulated on both Iba1 $^{+}$ microglia (379855.49 ± 35254.48 vs. 6486.02 ± 2773.22 , $p < 0.0001$) and GFAP $^{+}$ astrocytes (250994.60 ± 31974.33 vs. 19568.63 ± 12282.96 , $p < 0.0001$) in the brains of arcA β mice compared to NTL mice at 17 months of age ($n = 3$ per group) (Figure 2E). Furthermore, the average CB $_2$ R signal intensity is increased in image pixels indicating both Iba1 $^{+}$ microglia (55.25 ± 3.76 vs. 182.44 ± 11.10 , $p < 0.0001$) and GFAP $^{+}$ astrocytes (121.37 ± 11.80 vs. 7.02 ± 0.79 , $p < 0.0001$) of arcA β mice compared to NTL mice at 17 months of age ($n = 3$ per group) (Figure 2F).

Increased cannabinoid type 2 receptor associated with 6E10-positive A β plaque in the brains of arcA β compared to non-transgenic littermate

Increased 6E10 immunofluorescence intensity was observed in the cortex (3.48 ± 0.22 vs. 0.39 ± 0.15 , $p < 0.0001$) and hippocampus (6.80 ± 0.77 vs. 0.42 ± 0.20 , $p < 0.0001$) of arcA β mice compared to NTL mice at 17 months of age ($n = 3$ per group) and was comparable in the thalamus (0.95 ± 0.73 vs. 0.30 ± 0.20 , $p = 0.2936$) (Figures 1, 2G). In the brains of arcA β mice at 17 months of age, CB $_2$ R immunofluorescence was located on microglia and astrocytes both inside/within plaques (Figure 1). The intra-plaque (98.27 ± 4.31 , $p < 0.0001$) and peri-plaque (42.38 ± 9.84 , $p = 0.0007$) glial CB $_2$ R levels were both 20-fold and 10-fold that in the parenchyma (4.5 ± 0.24) of the arcA β mice ($n = 3$ per group), respectively. The glial-CB $_2$ R mean fluorescence intensity inside plaque was higher than that located

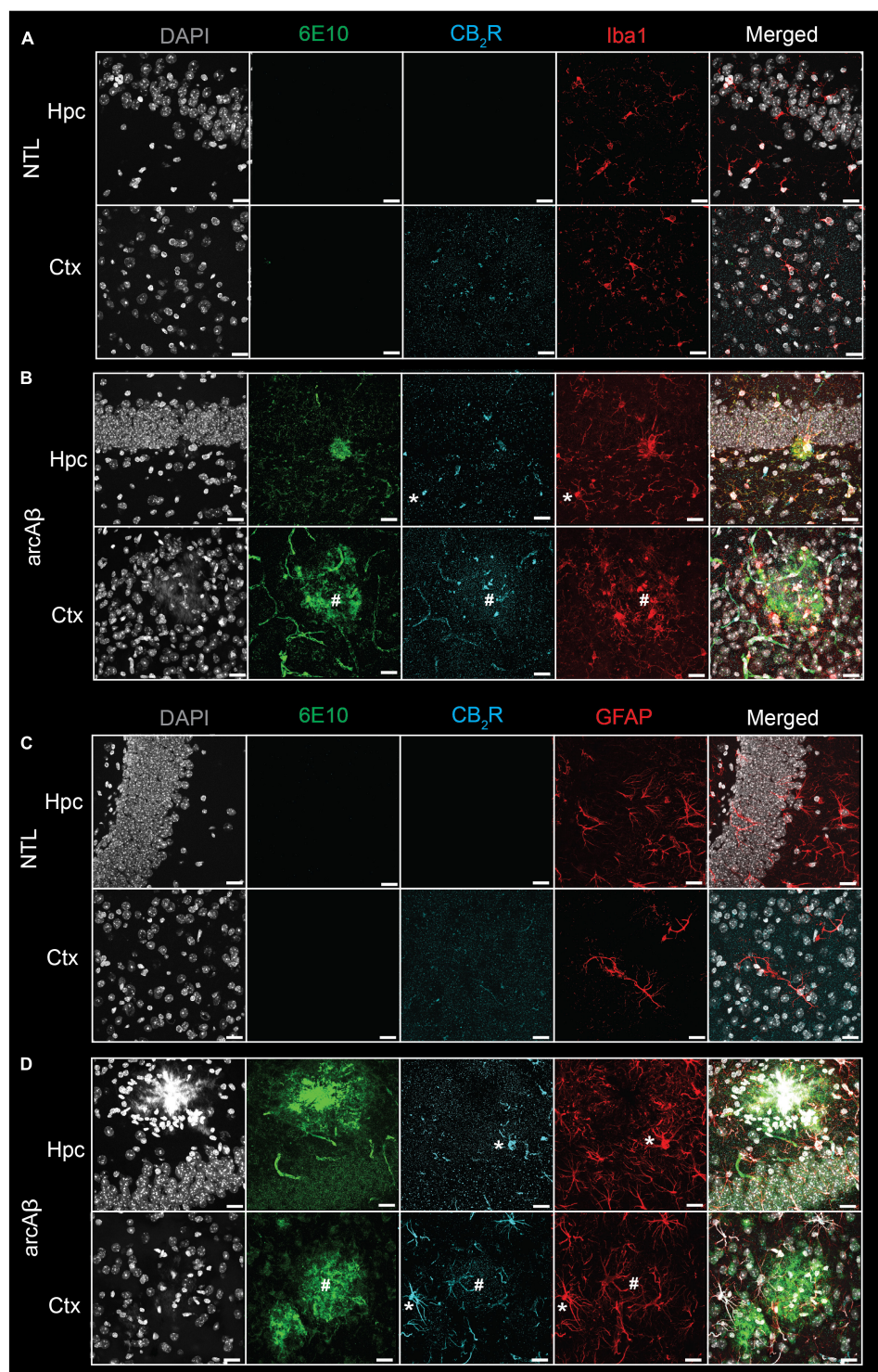
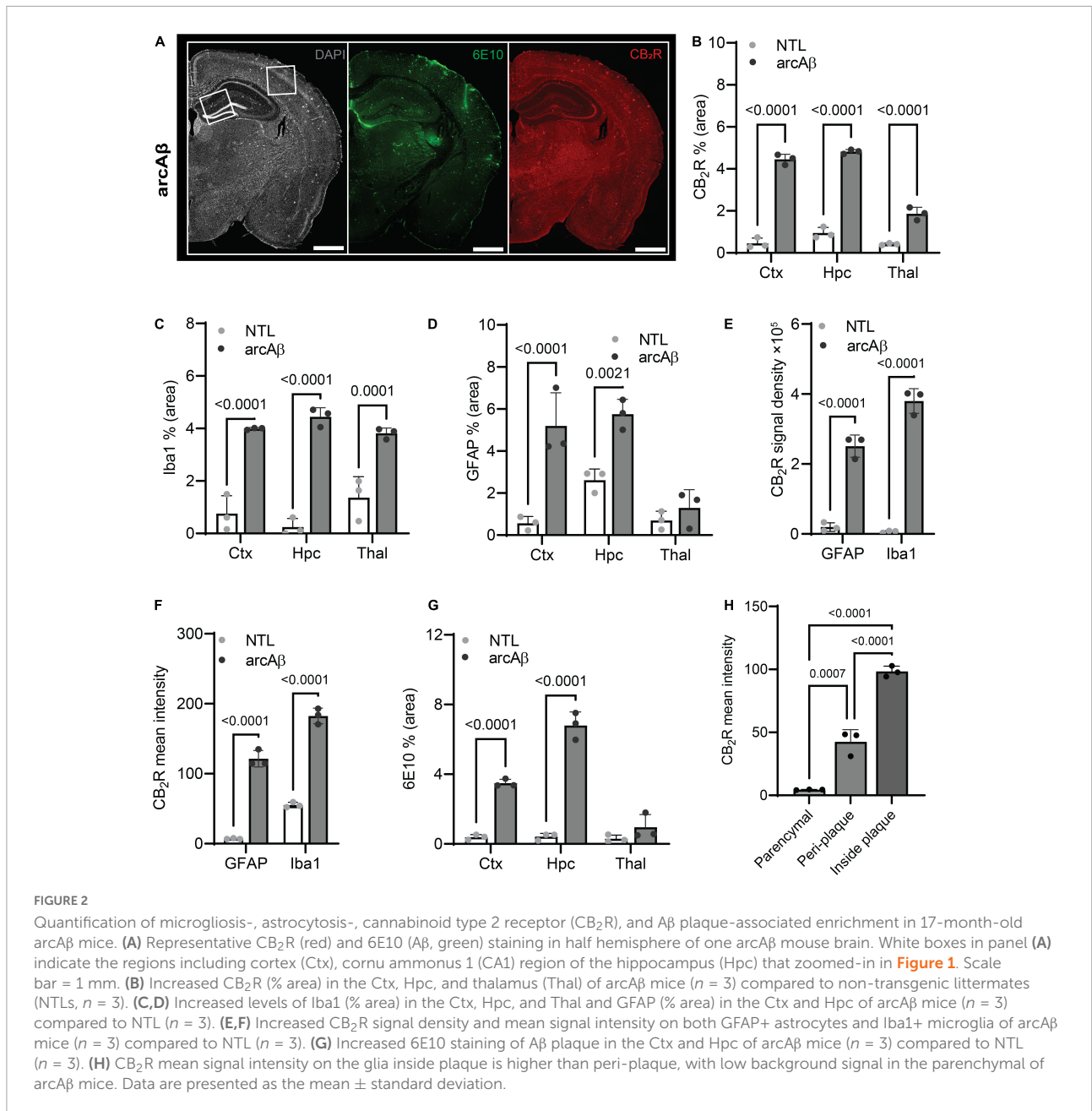


FIGURE 1

Increased cannabinoid type 2 receptor (CB₂R) in microglia and astrocytes associated with amyloid-beta deposits in 17-month-old arcAβ mice. **(A,B)** Brain tissue sections of non-transgenic littermate (NTL, *n* = 3) and arcAβ mice (*n* = 3) were stained for Aβ (6E10 antibody, green), CB₂R (cyan), and Iba1 (red) in the hippocampus (Hpc) and cortex (Ctx). Increased CB₂R and Iba1 immunoreactivity inside and surrounding the plaque. **(C,D)** Staining for Aβ (6E10, green), CB₂R (cyan), and GFAP (red) in the Hpc and Ctx. Nuclei were counterstained with DAPI (white). Increased CB₂R and GFAP immunoreactivity inside and surrounding Aβ plaques. *Localization of CB₂R on microglia or astrocytes outside plaque, #Colocalization of CB₂R on microglia or astrocytes within plaque. Scale bar = 20 μm. CB₂R immunoreactivity was detected on both microglia and astrocytes.



peri-plaque of the arcAβ mice ($p < 0.0001$, $n = 3$ per group) (Figures 1, 2H).

Increased Iba1+ and CD68+ microglia and GFAP+ astrocytes in the brains of arcAβ mice compared to non-transgenic littermate mice

Next, the levels of activated microglia using Iba1 and CD68 and astrocytes using GFAP were assessed in the brains of arcAβ mice and NTL mice at 17 months of age ($n = 3$ per group).

Increased numbers of microglia (Iba1% area) were observed in the vicinity of Aβ plaques and were upregulated in the cortex (3.99 ± 0.04 vs. 0.76 ± 0.68 , $p < 0.0001$), hippocampus (4.44 ± 0.35 vs. 0.24 ± 0.33 , $p < 0.0001$), and thalamus (3.81 ± 0.21 vs. 1.37 ± 0.80 , $p = 0.0001$) of arcAβ mice at 17 months of age compared to NTL mice ($n = 3$ per group). Similarly, an increased GFAP% area was associated with plaque in the cortex (5.29 ± 1.57 vs. 0.56 ± 0.33 , $p < 0.0001$) and hippocampus (5.75 ± 0.72 vs. 2.61 ± 0.53 , $p = 0.0021$) of arcAβ mice at 17 months of age compared to NTL mice ($n = 3$ per group), but not in the thalamus (1.29 ± 0.87 vs. 0.70 ± 0.44 , $p = 0.7933$) (Figures 1, 2C,D). CD68 is a lysosomal protein

expressed at high levels by activated microglia and at low levels by resting microglia in the CNS. Reactive microglia indicated by increased CD68 surrounding amyloid plaques (6E10) were observed and increased in the cortex (2.74 ± 0.50 vs. 1.03 ± 0.29 , $p = 0.0003$) of arcA β mice at 17 months of age compared to NTL mice ($n = 3$ per group) and were comparable in the hippocampus (1.57 ± 0.40 vs. 0.82 ± 0.35 , $p = 0.0734$) and thalamus (1.61 ± 0.19 vs. 1.10 ± 0.37 , $p = 0.2874$) (Figure 3).

No difference in whole-brain levels of [¹⁸F]RoSMA-18-d6 and [¹¹C]RSR-056 specific binding or *Cnr2* expression between arcA β and non-transgenic littermate mice of different ages

Autoradiography using [¹⁸F]RoSMA-18-d6 (0.5 nM), [¹¹C]RSR-056 (2 nM), and [¹¹C]RS-028 (3 nM) was performed on sagittal arcA β brain tissue at 17 months of age slides to assess the radioligand specificity. The concentrations of the radioligands used in autoradiography were determined based on previous publication (at half binding affinity of the ligand) [¹¹C]RS-028 (Haider et al., 2018), [¹¹C]RSR-056 (Slavik et al., 2015), and [¹⁸F]RoSMA-18-d6 (Haider et al., 2020). For the brain tissues, less than 50% of binding sites were blocked in the presence of CB₂R antagonist GW405833 (10 μ M). The percentage of specific binding is significantly lower than percentages previously reported for the spleen for these radioligands, likely due to the limited number of CB₂R binding sites in the brain (Slavik et al., 2015; Haider et al., 2018, 2020). [¹⁸F]RoSMA-18-d6 ($40.3 \pm 9.2\%$) showed a higher percentage of specific binding than [¹¹C]RSR-056 ($32.0 \pm 7.8\%$) and [¹¹C]RS-028 ($32.0 \pm 12.8\%$, Supplementary Figure 1) on arcA β ($n = 3$) and NTL ($n = 3$) mouse brain tissue at 17 months of age 3.

Thus, [¹⁸F]RoSMA-18-d6 (0.5 nM) and [¹¹C]RSR-056 (2 nM) were selected for further experiments to examine the CB₂R levels in arcA β and NTL at 6, 17, and 24 months of age by autoradiography of mouse brain slices. As no specific regional pattern of [¹⁸F]RoSMA-18-d6 (at 0.5 nM) and [¹¹C]RSR-056 (at 2 nM) specific binding was observed in arcA β and NTL mice at 6, 17, and 24 months of age, the specific binding level was analyzed using the whole hemisphere region-of-interest. No difference was observed in brain [¹⁸F]RoSMA-18-d6 (at 0.5 nM) levels between NTL and arcA β mice at 6 months of age (0.19 ± 0.03 vs. 0.18 ± 0.06 pmol/g tissue, $n = 5, 6$), 17 months of age (0.22 ± 0.01 vs. 0.19 ± 0.01 pmol/g tissue, $n = 3, 5$), and 24 months of age (0.20 ± 0.04 vs. 0.22 ± 0.06 pmol/g tissue, $n = 5, 5$) (Figures 4A,B). Similarly, for [¹¹C]RSR-056 (2 nM), no difference in radioactivity accumulation was observed in the brains of NTL and arcA β mice at 6 months of age (0.11 ± 0.03 vs. 0.12 ± 0.02 pmol/g tissue, $n = 5, 6$), 17 months of age (0.18 ± 0.05 vs. 0.13 ± 0.02 pmol/g, $n = 3,$

5), and 24 months of age (0.14 ± 0.09 vs. 0.15 ± 0.03 pmol/g, $n = 5, 5$) (Figures 4C,D). There was a robust correlation between [¹¹C]RSR-056 (2 nM) specific binding and [¹⁸F]RoSMA-18-d6 (at 0.5 nM) specific binding in arcA β and NTL mouse brains at 6, 17, and 24 months of age (Spearman rank, $r = 0.8042$, $p = 0.0025$) (Figure 4E).

Next, the mRNA expression levels of *Cnr2* were evaluated in the left hemisphere from the same cohort of arcA β and NTL mice in autoradiography at 6, 17, and 24 months of age that were assessed by [¹⁸F]RoSMA-18-d6 and [¹¹C]RSR-056 autoradiography ($n = 5$ –6/age group). No significant difference was observed in *Cnr2* mRNA expression between the NTL and arcA β mice at 6 months of age (1.92 ± 1.72 vs. 1.83 ± 0.79), 17 months of age (4.65 ± 6.30 vs. 4.20 ± 4.43), and 24 months of age (1.33 ± 0.92 vs. 3.36 ± 3.07) (Figure 4F).

Discussion

Here, we demonstrated an increase in local CB₂R expression levels in arcA β mice at 17 months of age compared to NTL mice and evaluated novel PET tracers [¹¹C]RSR-056 and [¹⁸F]RoSMA-18-d6 for detecting brain CB₂R changes in arcA β mice at 6, 17, and 24 months of age. Increased CB₂R fluorescence intensities and numbers of microglia and astrocytes inside/surrounding A β plaques were observed in arcA β mice compared to NTL mice at 17 months of age. However, no significant difference in CB₂R levels was observed at the whole-brain level measured either by using autoradiography or by mRNA analysis in arcA β compared to NTL mice at 6, 17, and 24 months of age.

Cannabinoid type 2 receptor has been an emerging target for imaging neuroinflammation partly due to its low expression levels under physiological conditions and upregulation under acute inflammatory conditions (Stella, 2010). The CB₂R fluorescence intensity was greatly increased in arcA β mice compared to NTL mice and was higher inside plaque than peri-plaque and in the parenchyma of arcA β mice. This observation is different from a previous publication of a significant increase in CB₂R intensities compared to the core of plaques (radius ≤ 7 μ m) (Savonenko et al., 2015). In addition, recent studies have reported the astroglial and neuronal expression of CB₂R in mouse and rat brains in addition to its expression on microglia by using immunostaining and RNAscope techniques (Van Sickle et al., 2005; Gong et al., 2006; Zarruk et al., 2012; Li and Kim, 2015; Savonenko et al., 2015; Yamagishi et al., 2019; Galán-Ganga et al., 2021). Savonenko et al. (2015) reported expression of CB₂R in neurons; and astrocytes in the brain from J20 amyloidosis mouse model at 12 months of age, in addition to its microglial expression. In our results from the immunofluorescence staining, CB₂R expression on both astrocytes and microglia was increased significantly in arcA β mice compared to the negligible level

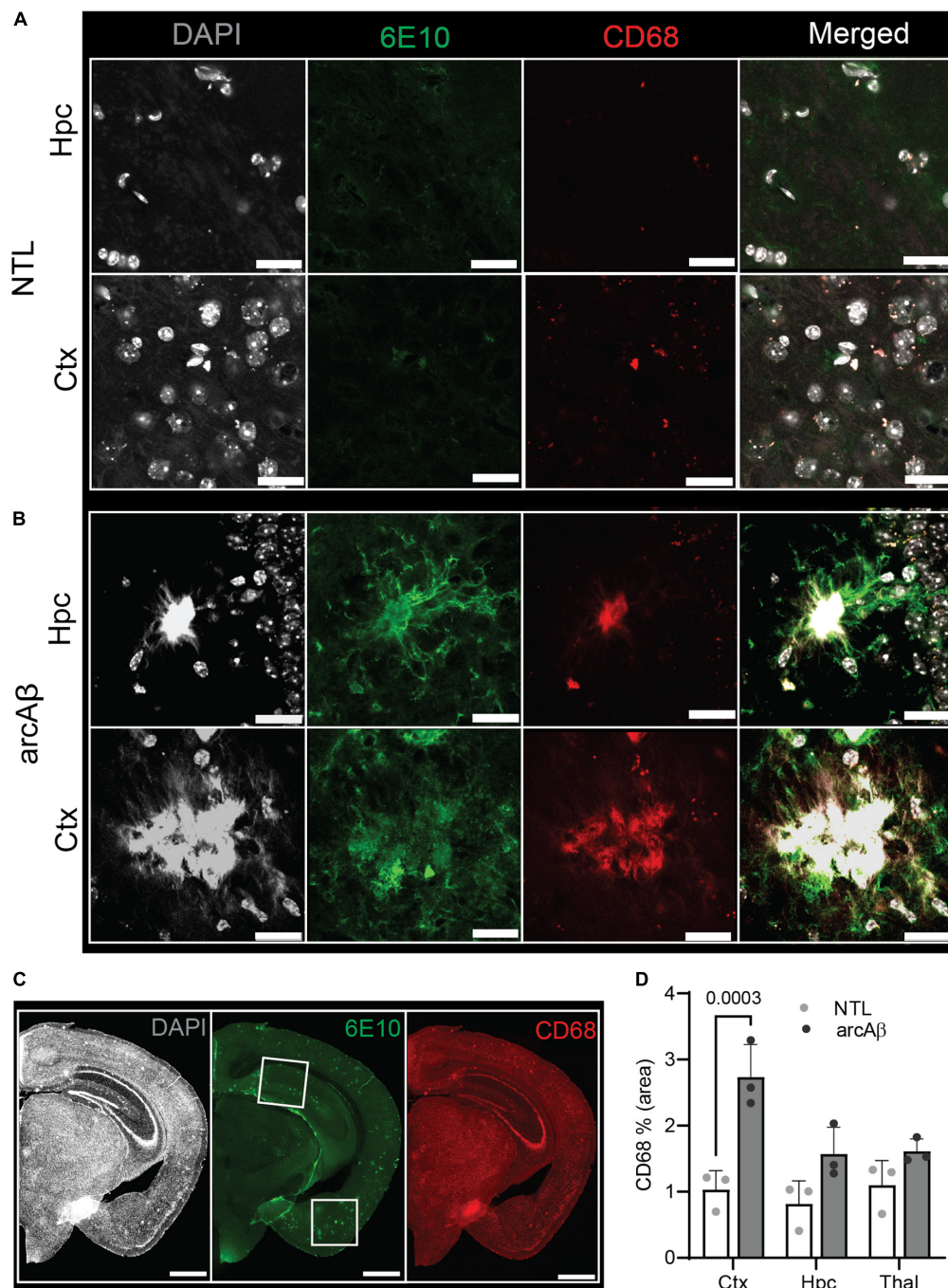


FIGURE 3

Microglial activation in the 17-month-old arcAβ mouse brain. (A–C) Brain tissue sections of non-transgenic littermate (NTL, $n = 3$) and arcAβ mice ($n = 3$) were stained for 6E10 (green)/CD68 (red) in the hippocampus (Hpc) and cortex (Ctx). White boxes in panel (C) indicate the regions including cortex (Ctx), cornu ammonus 1 (CA1) region of the hippocampus (Hpc) that zoomed-in in panels (A,B). Nuclei were counterstained with DAPI (white). Scale bar = 20 μm (A,B), =1 mm (C). (D) Quantification of CD68 signals in the Hpc, Ctx, and thalamus (Thal) of arcAβ mice compared to NTL mice confirmed microglial activation in arcAβ mice.

in NTL mice at 17 months of age (Figures 1, 2). Concerns have been raised regarding the specificity of CB₂R antibodies used in immunohistochemical staining. Specific neuronal subpopulations of CB₂R have been shown by using fluorescence

in situ hybridization and proximity ligand assays in non-human primates (Sierra et al., 2015). However, previous study has reported that CB₂-GFP expression is colocalized with Iba1 staining of microglia but not with NeuN staining of neuron

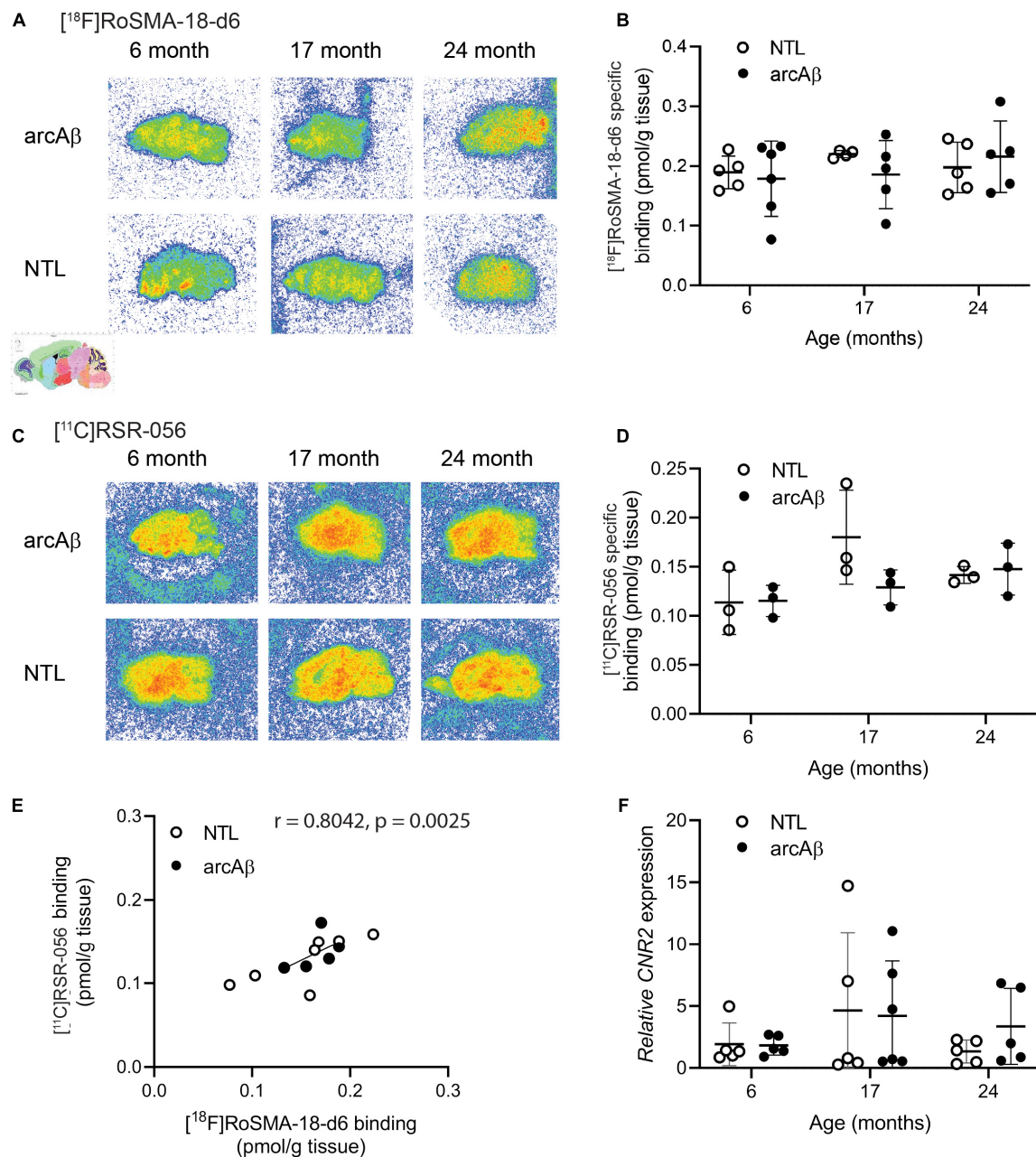


FIGURE 4

Comparable regional $[^{18}\text{F}]\text{RoSMA-18-d6}$ and $[^{11}\text{C}]\text{RSR-056}$ specific binding in the brains of arcA β and non-transgenic mice at 6, 17, and 24 months of age. **(A)** Representative $[^{18}\text{F}]\text{RoSMA-18-d6}$ (0.5 nM) autoradiographic images (total binding) of sagittal brain sections of arcA β and non-transgenic littermate (NTL) mice at 6 months of age ($n = 5, 6$), 17 months of age ($n = 4, 5$), and 24 months of age ($n = 5, 5$). **(C)** Representative $[^{11}\text{C}]\text{RSR-056}$ (2 nM) autoradiographic images (total binding) of sagittal brain sections of arcA β and non-transgenic littermate (NTL) mice at 6 months of age ($n = 3, 3$), 17 months of age ($n = 3, 3$), and 24 months of age ($n = 3, 3$). **(B,D)** Quantification of specific binding of $[^{18}\text{F}]\text{RoSMA-18-d6}$ (0.5 nM), and $[^{11}\text{C}]\text{RSR-056}$ (2 nM) in the whole sagittal brain slice in panels **(A,C)**. Two-way ANOVA, arcA β vs. NTL. **(E)** Robust correlation between specific binding of $[^{18}\text{F}]\text{RoSMA-18-d6}$ (0.5 nM) and $[^{11}\text{C}]\text{RSR-056}$ (2 nM) autoradiograms on arcA β and NTL mouse brain hemispheres (Spearman rank, $r = 0.8042, p = 0.0025$). **(F)** *Cnr2* expression in arcA β and NTL mouse brain hemisphere homogenates at 6 months of age ($n = 5, 5$), 17 months of age ($n = 5, 6$), and 24 months of age ($n = 5, 5$).

or GFAP staining of astrocyte in CB2-GFP BAC transgenic mice (Lundt et al., 2015). López et al. (2018) reported that CB₂R-dependent- enhanced green fluorescent protein (EGFP)

expression is limited to plaque-associated microglial cells but is absent in neurons and astrocytes in CB2^{EGFP/f/f}/5 × FAD mice.

Although *Cnr2* expression in AD APP/PS1 has been reported to be upregulated, great variation between animals and a low fold increase lead to insignificance in comparison (Aso et al., 2013; Vidal-Palencia et al., 2022). Recent gene expression analysis showed that regional *Cnr2* expression differs between male/female APP/PS1 mice (Vidal-Palencia et al., 2022). Here, *Cnr2* expression was analyzed using homogenates of half hemispheres of arcA β and NTL mice with further dissection. No difference in *Cnr2* expression between arcA β and NTL mice of different ages was observed.

For preclinical imaging, high variabilities in imaging of brain CB₂R levels among animal models of neuroinflammation were reported from previous studies. Upregulated levels of brain CB₂R have been demonstrated in transient middle cerebral artery occlusion ischemic stroke mice using [¹⁸F]RoSMA-18-d6 (Ni et al., 2021a) and in senescence-accelerated SAMP10 mice using [¹¹C]NE40 (Yamagishi et al., 2019). Another study by PET using [¹¹C]A-836339 in a lipopolysaccharide-injected rat model did not report changes in tracer uptake following neuroinflammation (Pottier et al., 2017). MicroPET using [¹¹C]A-836339 showed increased uptake in the brain areas with A β depositions in a J20 mouse model of AD (Ni et al., 2019b). In the only reported study in patients with AD using PET, Ahmad et al. (2016) reported lower CB₂R availability in A β -positive AD patients than in healthy controls assessed by PET using [¹¹C]NE40 and [¹¹C]PIB, respectively. No relationship between [¹¹C]NE40 and cerebral A β load was observed in this study.

[¹¹C]RSR-056 and [¹⁸F]RoSMA-18-d6 showed 32 and 40% specific binding in the AD mouse brain, respectively, and there was no difference between arcA β and NTL mice. One of the difficulties is the low CB₂R expression level in the brain and the low number of binding sites. Using the same tracers, [¹¹C]RS-028 (Haider et al., 2018), [¹¹C]RSR-056 (Slavik et al., 2015), and [¹⁸F]RoSMA-18-d6 (Haider et al., 2020), lower non-specific binding has been shown in post-mortem spleen and spinal cord tissues from patients with amyotrophic lateral sclerosis than in those from healthy controls. Further development of CB₂R tracers of even higher affinity to overcome the low number of binding sites (B_{max}) is desired. In addition, as species differences exist regarding CB₂R brain expression, further studies on post-mortem brain tissues from patients with AD will provide information on CB₂R disease relevance.

There are several limitations in this study: (1) Negative control: CB₂R has been considered difficult to validate by using immunohistochemical staining. Further studies using CB₂R knockout mice showing absence of staining and binding of radioligand will be confirmative. (2) Sample size: Number of animals included in the mRNA and autoradiography experiment of this brief report is low. Further study using larger sample size will improve the accuracy of the results. (3) *In vivo* imaging: Autoradiography provides information on the probe

binding specificity and identifies potential regions of interest with validation from immunohistochemical characterization. Due to a lack of difference from autoradiography, *in vivo* measurement has not been performed in the AD mouse models using the radioligands investigated.

Conclusion

In conclusion, increases in CB₂R immunofluorescence intensity on the glia were detected in the brains of arcA β mice compared to NTL mice and were associated with A β deposits. Further improvement of the binding properties of CB₂R PET tracers will be needed to detect subtle changes in CB₂R in an AD animal model.

Data availability statement

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found in the article/**Supplementary material**.

Ethics statement

The animal study was reviewed and approved by the Cantonal Veterinary Office Zurich.

Author contributions

RN, JK, SA, and AHa designed the study. VK performed the staining and microscopy. LM synthesized the radioligands. FS performed the mRNA analysis. RN performed the autoradiography. VK, FS, LM, and RN performed data analysis. VK and RN wrote the initial manuscript. All authors read and approved the final manuscript.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fnagi.2022.1018610/full#supplementary-material>

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