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Synthesis and biological evaluation of carbon-11- and fluorine-18-labeled 2-oxoquinoline derivatives for type 2 cannabinoid receptor positron emission tomography imaging $\stackrel{\text{type}}{\approx}$

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Abstract

Introduction: The type 2 cannabinoid (CB_2) receptor is part of the endocannabinoid system and has been suggested as a mediator of several central and peripheral inflammatory processes. Imaging of the CB_2 receptor has been unsuccessful so far. We synthesized and evaluated a carbon-11- and a fluorine-18-labeled 2-oxoquinoline derivative as new PET tracers with high specificity and affinity for the CB_2 receptor. **Methods:** Two 2-oxoquinoline derivatives were synthesized and radiolabeled with either carbon-11 or fluorine-18. Their affinity and selectivity for the human CB_2 receptor were determined. Biological evaluation was done by biodistribution, radiometabolite and autoradiography studies in mice.

Results: In vitro studies showed that both compounds are high affinity CB_2 -specific inverse agonists. Biodistribution study of the tracers in mice showed a high in vivo initial brain uptake and fast brain washout, in accordance with the low CB_2 receptor expression levels in normal brain. A persistently high in vivo binding to the spleen was observed, which was inhibited by pretreatment with two structurally unrelated CB_2 selective inverse agonists. In vitro autoradiography studies with the radioligands confirmed CB_2 -specific binding to the mouse spleen. **Conclusion:** We synthesized two novel CB_2 receptor PET tracers that show high affinity/selectivity for CB_2 receptors. Both tracers show favourable characteristics as radioligands for central and peripheral in vivo visualization of the CB_2 receptor and are promising candidates for primate and human CB_2 PET imaging.

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1. Introduction

The type 2 cannabinoid receptor (CB₂ receptor) is part of the endocannabinoid system, together with the type 1 cannabinoid receptor (CB₁ receptor), endogenous ligands, transporters and enzymes. Most research has focused on the CB₁ receptor, which is responsible for the psychoactive

* Corresponding author. Tel.: +32 16330447; fax: +32 16330449. *E-mail address:* guy.bormans@pharm.kuleuven.be (G.M. Bormans). effects of cannabis and a multitude of other CNS functions (for review, see Ref. [1]) and for which PET radioligands have been developed recently [2,3].

The CB₂ receptor is related to the immune system and expressed in the spleen, lymph nodes and Peyer's patches [4]. In normal brain, only low CB₂ receptor expression levels were observed [5]. However, in some pathological conditions, the CB₂ receptor is up-regulated. Peripheral CB₂ receptor up-regulation has been demonstrated in tumors [6], endometrial inflammation [7] and human and mouse atherosclerotic plaques [8]. In the brain, CB₂ receptor upregulation is linked to neuroinflammation [9] and was observed in the spinal cord of a mouse model for

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amyotrophic lateral sclerosis [10], plaques of demyelination in multiple sclerosis patients [11], but also in a rat stroke model [12] and on senile amyloid plaques in Alzheimer patients [13]. The CB₂ receptor seems to play an important role in the regulation of several biological processes, such as bone modeling [14], skin cancer development [15] and immunity [16].

To date, a number of nonselective tritiated radioligands such as $[{}^{3}H]CP55,940$ or $[{}^{3}H]WIN55,212$ with affinity for the human CB₂ receptor are available for in vitro use. Also, synthesis of a near-infrared dye-labeled CB₂ receptor ligand for noninvasive in vivo imaging was reported [17]. We recently described the development of a carbon-11-labeled CB₂ selective PET tracer, which did not penetrate the blood– brain barrier (BBB) [18]. Therefore, the use of this tracer will be limited to peripheral visualization of the CB₂ receptor. The goal of this recent study was to develop a tracer which crosses the BBB in order to enable the future visualization of CB₂ receptor up-regulation in the above-described neuroinflammation pathologies.

In the present study, we have synthesized and studied the biological behaviour of carbon-11- or fluorine-18-labeled 2-oxoquinolines as PET tracers for in vivo visualization of the CB_2 receptor.

2. Materials and methods

2.1. General conditions

All reagents and solvents were purchased from Aldrich (Steinheim, Germany), Acros (Geel, Belgium) or ABCR (Karlsruhe, Germany), and were used without further purification.

¹H NMR spectra were recorded on a Bruker AVANCE 300-MHz spectrometer (Bruker AG, Faellanden, Switzerland) using $CDCl_3$ or $DMSO-d_6$ as solvent. Chemical shifts are reported in parts per million relative to tetramethylsilane (δ =0). Coupling constants are reported in Hertz. Splitting patterns are defined by s (singlet), d (doublet), dd (double doublet), t (triplet) or m (multiplet). High-performance liquid chromatography (HPLC) purification and analysis were performed on a Merck Hitachi L6200 intelligent pump (Hitachi, Tokyo, Japan) connected to a UV spectrometer (Waters 2487 dual γ absorbance detector) set at 254 nm. Radioactivity in the HPLC eluate was detected using a 2-in. NaI(Tl) scintillation detector connected to a single-channel analyzer. Data were acquired and analyzed using the RaChel (Lablogic, Sheffield, UK) or GINA Star (Raytest, Straubenhardt, Germany) data acquisition system. Quantification of radioactivity for biodistribution and radiometabolite studies was done using a gamma counter [3-in. NaI(Tl) well crystal] coupled to a multichannel analyzer and mounted in a sample changer (Wallac 1480 Wizard 3 in., Wallac, Turku, Finland). The values are corrected for background radiation and physical decay during counting. Mass measurement

was performed on a time-of-flight mass spectrometer (LCT, Micromass, Manchester, UK) equipped with an orthogonal electrospray ionization interface. Accurate mass determination was done by co-infusion with a 10 μ g/ml solution of Kryptofix as an internal lock mass. Acquisition and processing of data were done using Masslynx software (version 3.5, Micromass).

2.2. Synthesis

Synthesis of Compounds 1, 2, 3, 4, 5 and 6 was based on methods described by Raitio et al. [19] with introduction of small modifications.

2.2.1. 2-Nitro-3-hydroxy-4-methoxybenzaldehyde (1)

To a mixture of isovanillin (4.45 g, 29.25 mmol) in anhydrous dichloromethane (50 ml) at -70°C, 58.4 ml of a 0.5 M nitronium tetrafluoroborate solution in sulfolane was added and stirred for 1 h. The reaction mixture was allowed to heat up to -20° C and further reacted for 24 h at -20° C. Water (30 ml) was added dropwise and the mixture was allowed to reach room temperature (RT). The reaction mixture was extracted with diethylether (3×100 ml) and the combined diethylether fractions were washed with water (50 ml) and brine (50 ml) and dried over MgSO₄. After evaporation of the solvent, the crude reaction product was purified by silica gel column chromatography using hexane/ EtOAc 1:1 as the eluent to yield a yellow oil (3.28 g, 57%). MS (ES)⁻: m/z=196 (M-H)⁻. ¹H NMR (DMSO) δ 10.807 (1H, br s), 9.756 (1H, s), 7.577 (1H, d, ${}^{3}J_{H-H}$ =8.4 Hz), 7.344 $(1H, d, {}^{3}J_{H-H}=8.5 \text{ Hz}), 3.988 (3H, s).$

2.2.2. 2-Nitro-3-butyloxy-4-methoxybenzaldehyde (2)

Bromobutane (6.12 ml, 57.00 mmol) was added dropwise to a mixture of **1** (3.20 g, 16.23 mmol) and potassium carbonate (7.88 g, 56.99 mmol) in anhydrous dimethylformamide (100 ml). The reaction mixture was stirred at 100°C for 2 h. After filtration and dilution with water (50 ml), the filtrate was extracted with hexane/EtOAc 1:1 (3×80 ml). The combined extracts were washed with water (60 ml) and brine (60 ml), dried over MgSO₄ and the solvent was evaporated. Further purification was done by silica gel column chromatography using hexane/EtOAc 19:1 as the eluent to yield **2** as a yellow oil (3.21 g, 78%). MS (ES)⁺: m/z=254(M+H)⁺. ¹H NMR (CDCl₃) δ 9.789 (1H, s), 7.635 (1H, d, ³*J*_{H-H}=8.6 Hz), 7.108 (1H, d, ³*J*_{H-H}=8.6 Hz), 4.113 (2H, t, ³*J*_{H-H}=6.5 Hz), 3.999 (3H, s), 1.695 (2H, m), 1.437 (2H, m), 0.949 (3H, t, ³*J*_{H-H}=7.3 Hz).

2.2.3. 2-Amino-3-butyloxy-4-methoxybenzaldehyde (3)

To a solution of **2** (3.00 g, 11.85 mmol) in ethanol/acetic acid/water (2:2:1, 80 ml), iron powder (1.98 g, 35.45 mmol) and concentrated HCl (0.6 ml) were added. The reaction mixture was refluxed (110°C) for 15 min, stirred at RT for 1 h, filtrated and water (60 ml) was added to the resulting filtrate, and the solution was extracted with ethylacetate (3×60 ml). The combined ethylacetate extracts were washed with saturated NaHCO₃ (3×60 ml) and brine (60 ml), dried

over MgSO₄ and the solvent was evaporated. Further purification was done by silica gel column chromatography using hexane/EtOAc 9:1 to yield **3** as a light yellow oil (1.25 g, 47%). MS (ES)⁺: m/z=224 (M+H)⁺. ¹H NMR (CDCl₃) δ 9.741 (1H, s), 7.216 (1H, d, ³J_{H-H}=8.8 Hz), 6.376 (1H, d, ³J_{H-H}=8.8 Hz), 6.269 (2H, s), 3.949 (2H, t, ³J_{H-H}=6.7 Hz), 3.902 (3H, s), 1.760 (2H, m), 1.512 (2H, m), 0.978 (3H, t, ³J_{H-H}=7.3 Hz).

2.2.4. 2-Oxo-7-methoxy-8-butyloxy-1,2-dihydroquinoline-3carboxylic acid methyl ester (4)

To a solution of **3** (1.22 g, 5.46 mmol) in methanol (25 ml), dimethyl malonate (1.89 ml, 16.54 mmol), piperidine (1.35 ml, 13.67 mmol) and 30 µl acetic acid were added. The reaction mixture was refluxed overnight and extracted with ethylacetate (3×15 ml). The combined EtOAc extracts were washed with water (20 ml) and brine (20 ml), dried over MgSO₄ and the solvent was evaporated. Purification of **4** was done by silica gel column chromatography using dichloromethane: CH₃OH gradient mixtures (0% to 1% CH₃OH) as the eluent yielding **4** as a white solid (0.55 g, 33%). MS (ES)⁺: m/z=306 (M+H). ¹H NMR (DMSO) δ 10.984 (1H, s), 8.463 (1H, s), 7.569 (1H, d, ³J_{H-H}=8.9 Hz), 7.055 (1H, d, ³J_{H-H}=8.9 Hz), 3.964 (2H, t, ³J_{H-H}=6.8 Hz), 3.908 (3H, s), 3.783 (3H, s), 1.742 (2H, m), 1.401 (2H, m), 0.918 (3H, t, ³J_{H-H}=7.4 Hz).

2.2.5. 2-Oxo-7-methoxy-8-butyloxy-1,2-dihydroquinoline-3carboxylic acid (5)

5 (0.50 g, 1.64 mmol) was dissolved in ethanol (19 ml) and HCl 2 M was added (12 ml). The mixture was stirred at 60°C for 7 h, cooled and filtrated. The filter residue was dried under vacuum to give **5** as a yellow-white powder (0.26 g, 55%). MS (ES)⁺: m/z=292 (M+H).

2.2.6. 2-Oxo-7-methoxy-8-butyloxy-1,2-dihydroquinoline-3carboxylic acid cyclohexylamide (**6**)

To a solution of 5 (0.13 g, 0.45 mmol) in anhydrous toluene (10 ml), thionylchloride (0.11 ml, 1.51 mmol) was added. The reaction mixture was refluxed for 3 h. The toluene was evaporated, anhydrous toluene was added again and the evaporation was repeated. The resulting acyl chloride was without further purification dissolved in dichloromethane (10 ml) and added dropwise to a solution of cyclohexylamine (0.07 g, 0.67 mmol) and triethylamine (0.113 g, 1.12 mmol) in dichloromethane (10 ml). The reaction mixture was stirred for 2 h at RT. Water (30 ml) was added; the dichloromethane layer was separated from the water layer which was extracted with dichloromethane (3×20 ml). The resulting dichloromethane extracts were washed with brine (30 ml) and dried using MgSO₄. The solvent was evaporated and the residue was further purified by silica gel column chromatography using heptane/EtOAc 2:1 as the eluent yielding **6** as a white solid (0.15 g, 90%). 1 H NMR (CDCl₃) δ 9.595 (1H, d, ${}^{3}J_{H-H}$ =7.8 Hz), 9.126 (1H, s), 8.859 (1H, s), 7.454 (1H, d, ³J_{H-H}=8.8 Hz), 6.932 (1H, d, ³J_{H-H}=8.9 Hz), 4.145 (2H, t, ³J_{H-H}=6.8 Hz), 3.974 (3H, s),

1.256–2.002 (15H, m), 0.994 (3H, t, ${}^{3}J_{H-H}$ =7.4 Hz). MS (ES)⁺ Accurate mass: $[C_{21}H_{27}N_2O_4+Na]^+$ theoretical mass 394.1869 Da and found 394.1829 Da.

2.2.7. 2-Oxo-7-hydroxy-8-butyloxy-1,2-dihydroquinoline-3carboxylic acid cyclohexylamide (7)

The reaction proceeded under nitrogen atmosphere. To a solution of diphenylphosphine (0.12 ml, 0.7 mmol) in anhydrous THF (0.7 ml) at 0°C, butyllithium (0.60 ml of a 1.6 M solution in hexane) was slowly added. The reaction mixture was stirred for 30 min at RT. A solution of 6(0.20 g)0.54 mmol) in anhydrous THF (1 ml) was slowly added to the reaction mixture, which was further stirred for 3 h. The reaction was quenched by slowly adding water (10 ml) and NaOH (pH > 11). The reaction mixture was extracted with diethylether $(2 \times 10 \text{ ml})$ and the collected diethylether fractions were extracted with 0.1 M NaOH (20 ml). The combined water fractions were acidified by HCl 0.1 M until the solution turned milky white. The solution was extracted with diethylether $(3 \times 20 \text{ ml})$ and the combined diethylether fractions washed with water (20 ml) and brine (20 ml), and dried using MgSO₄. The excess solvent was evaporated and purification was done by silica gel column chromatography using CH₂Cl₂/CH₃OH gradient mixtures (0% to 1% CH₃OH) as eluent yielding 7 as a white solid (0.055 g, 29%). ¹H NMR (CDCl₃) δ 9.612 (1H, d, ³J_{H-H}=7.9 Hz), 9.004 (1H, s), 8.900 (1H, s), 8.848 (1H, s), 7.383 (1H, d, ³J_H- $_{H}$ =8.7 Hz), 6.947 (1H, d, $^{3}J_{H-H}$ =8.7 Hz), 4.073 (2H, t, $^{3}J_{H-H}$ = $_{H}$ =6.8 Hz), 1.210–2.005 (15H, m), 1.007 (3H, t, $^{3}J_{H-H}$ =7.4 Hz). MS $(ES)^+$ Accurate mass: $[C_{20}H_{25}N_2O_4+Na]^+$ theoretical mass 380.1712 Da and found 380.1724 Da.

2.2.8. 2-Oxo-7-fluoroethoxy-8-butyloxy-1,2-dihydroquinoline-3-carboxylic acid cyclohexylamide (8)

To a solution of 7 (0.080 g, 0.22 mmol) in butanone (6 ml), NaI (0.0033 g, 0.022 mmol), potassium carbonate (0.046 g, 0.33 mmol) and fluoroethylbromide (FEtBr; 0.043 g, 0.34 mmol) were added and the reaction mixture was refluxed for 3 h. The solvent was evaporated and purification was done by silica gel column chromatography using heptane/EtOAc gradient mixtures (0% to 5% EtOAc) yielding **8** as a white solid (0.037 g, 41%). ¹H NMR (CDCl₃) δ 9.580 (1H, d, ³*J*_{*H*-*H*}=8.1 Hz), 9.125 (1H, s), 8.862 (1H, s), 7.439 (1H, d, ³*J*_{*H*-*H*}=47.3 Hz, ³*J*_{*H*-*H*}=4.0 Hz), 4.370 (2H, dt, ³*J*_{*H*-*H*}=6.7 Hz), 1.256-2.004 (15H, m), 0.996 (3H, t, ³*J*_{*H*-*H*}=7.4 Hz). MS (ES)⁺ Accurate mass: [C₂₂H₂₉FN₂O₄+Na]⁺ theoretical mass 427.2003 Da and found 427.2013 Da.

2.2.9. 1-(2,4-Dichlorophenyl)-6-fluoro-N-piperidin-

1-yl-1,4-dihydroindeno[1,2-c]pyrazole-3-carboxamide (9)

Synthesis was performed according to the methods described by Mussinu et al. [20] yielding **9** as a white solid. ¹H NMR (CDCl₃) δ 7.663 (1H, d, ⁴*J*_{*H*-*H*}=2.2 Hz), 7.630 (1H, br s), 7.549 (1H, d, ³*J*_{*H*-*H*}=8.5), 7.470 (1H, dd, ³*J*_{*H*-*H*}=8.8 Hz and ⁴*J*_{*H*-*H*}=2.2 Hz), 7.267 (1H, d, ³*J*_{*H*-}

 $_{H}$ =9.7 Hz), 6.915–6.949 (2H, m), 3.891 (2H, s), 2.887 (4H, t, $^{3}J_{H-H}$ =5.3 Hz), 1.761 (4H, m), 1.441 (2H, m). Accurate mass: $[C_{22}H_{19}N_{4}OCl_{2}F + H]^{+}$ theoretical mass 445.0993 Da and found 445.0990 Da.

2.3. Radiosynthesis

2.3.1. Synthesis of [¹¹C]-6

2.3.1.1. Synthesis of $[{}^{11}C]CH_3I$, $[{}^{11}C]methyl$ triflate and $\int^{11} C$ -6. Carbon-11 was produced in a Cyclone 18/9 cyclotron (IBA, Louvain-la-Neuve, Belgium) in the form of 11 CH₄ by irradiation of a mixture of N₂ with 5% H₂ with 18-MeV protons. ¹¹CH₄ was transferred from the cyclotron target to a home-made synthesis module housed in a lead shielded cabinet where ¹¹CH₄ was trapped on a column filled with Porapak N (divinylbenzene/vinyl pyrolidone polymer), cooled in liquid nitrogen. The loop was allowed to warm to RT. ¹¹CH₄ was swept off the column with a stream of helium and was mixed with I₂ vapour at 100°C, after which the mixture passed through a quartz column heated to 650°C to yield ¹¹CH₃I [21]. ¹¹CH₃I was used as such or was passed through a column (150×3 mm) filled with silver triflate (50-mm silver triflate between two times 50-mm quartz wool) heated to 180°C, yielding the more reactive [¹¹C]methyl triflate. A stream of helium containing the alkylating agent was bubbled through a solution of 200 μ g 7 and 2–4 mg Cs₂CO₃ in 200 μ l DMF. The reaction mixture was heated during 2 min at 90°C, diluted with 1.8 ml 0.05 M ammonium acetate buffer pH 6.9 containing 40% ethanol and applied onto a semipreparative XTerra RP18 column (5 μm, 7.8×150 mm; Waters) which was eluted with 0.05 M ammonium acetate buffer (pH=6.9)/ EtOH (40:60 v/v, 2 ml/min). With this system, retention time is 14.8±0.5 min for $[^{11}C]$ -6 (*n*=9).

2.3.2. Synthesis of [¹⁸F]-8

2.3.2.1. Production of [¹⁸F]fluoride. [¹⁸F]Fluoride was produced from a $[{}^{18}O(p,n){}^{18}F]$ nuclear reaction by irradiation on 97% enriched H₂¹⁸O (Rotem HYOX18, Rotem Industries, Beer Sheva, Israel) using 18-MeV protons. After irradiation, $[^{18}F]F^-$ was separated from $[^{18}O]H_2O$ by trapping on a SepPak Light Accell plus QMA anion exchange cartridge (Waters, Milford, USA), which was previously preconditioned by successive washing with 0.5 M K_2CO_3 solution (10 ml) and water (2×10 ml). The [¹⁸F]F⁻ was eluted from the cartridge with a solution containing 2.47 mg potassium carbonate and 27.9 mg Kryptofix 222 in 0.75 ml of H₂O-CH₃CN (5:95 v/v) into a reaction vessel. The solution was heated to 96°C and the solvent was evaporated with a stream of nitrogen. [¹⁸F]F⁻ was further dried by adding 1 ml of anhydrous acetonitrile and subsequent evaporation at 96°C with a stream of nitrogen.

2.3.2.2. Synthesis of $[^{18}F]$ fluoroethylbromide and $[^{18}F]$ -8. 2-Bromoethyl triflate (BrCH₂CH₂OTf) (5 µl) in

o-dichlorobenzene (0.7 ml) was added to the reaction vessel containing dried [¹⁸F]F⁻ and potassium–kryptofix complex [22]. The reaction vial was then heated to 110°C and the resulting [¹⁸F]fluoroethylbromide ([¹⁸F]FEtBr) was distilled with a stream of helium and passed through an ascarite column (6×150 mm) in a reaction vial containing 200 µg 7 and 2–4 mg Cs₂CO₃ in 200 µl DMF. The reaction mixture was heated at 90°C for 10 min, diluted with 1.8 ml ammonium acetate buffer 0.05 M pH 6.9 containing 30% ethanol and injected on a semipreparative XTerra RP18 column (5 µm, 7.8×150 mm; Waters) which was eluted with 0.05 M ammonium acetate buffer (pH=6.9)/EtOH (40:60 v/v, 1.8 ml/min). With this system, retention time is 15.4±1.4 min for [¹⁸F]-**8** (*n*=4).

2.4. Quality control

Quality control for [¹¹C]-6 and [¹⁸F]-8 using authentic 6 or 8 as a reference was done by HPLC on an XTerra RP18 column (5 μ m, 4.6×250 mm, Waters) eluted with 0.05 M ammonium acetate buffer pH 6.9/acetonitrile (40:60 v/v, 1 ml/min). Using this system, retention time is 12.8±0.1 min for [¹¹C]-6 (*n*=5) and 11.9±0.1 min for [¹⁸F]-8 (*n*=4).

2.5. Partition coefficient

Twenty-five microliters of a solution of the HPLCisolated [¹¹C]-6 or [¹⁸F]-8 was added to a test tube containing 2 ml of 1-octanol and 2 ml of 0.025 M phosphate buffer pH 7.4. The test tube was vortexed at RT for 2 min and then centrifuged at 2700×g for 10 min. A 100-µl aliquot was taken from the 1-octanol phase and a 900-µl aliquot from the aqueous phase, taking care to avoid cross contamination between the phases. The separate aliquots were transferred into tared vials and the volume added was calculated from the mass of the aliquots and the specific density (ρ) of the phase, assuming that $\rho_{\text{buffer}}=1.000$ g/ml and $\rho_{1-\text{octanol}}=0.827$ g/ml. The radioactivity of the aliquots was counted using an automatic γ counter. The partition coefficient (P) was calculated as [radioactivity (cpm/ml) in 1-octanol]/[radioactivity (cpm/ ml) in phosphate buffer pH 7.4].

2.6. Biodistribution studies

The experiments in mice were carried out in compliance with the national laws relating to the conduct of animal experimentation and approved by the local animal ethics committee. All biodistribution studies were conducted in male National Murine Research Institute (NMRI) mice (37–50 g). Mice were anesthetized with isoflurane (2% in oxygen). The solution of the HPLC-purified product was diluted with saline to a concentration of approximately 90 MBq/ml or 7.4 MBq/ml (for [¹¹C]-6 or [¹⁸F]-8, respectively). An aliquot of 100 μ l was injected via a tail vein. The animals were sacrificed by decapitation, and the organs and body parts were dissected and weighed. The activity in the dissected organs and blood was measured

using a gamma counter. For calculation of total blood, muscle and bone radioactivity, blood, muscle and bone mass were assumed to be 7%, 40% and 12%, respectively, of the body mass [23,24].

2.7. Plasma and brain metabolites

The in vivo metabolic stability of the tracers in plasma and brain was studied in male NMRI mice. 9.25 MBq or 0.74 MBq (for $[^{11}C]$ -6 or $[^{18}F]$ -8, respectively) was injected through a tail vein, and the mice were sacrificed by decapitation at 2, 10 or 30 min pi (two to three mice per time point). For determination of plasma metabolites, blood was collected into a BD vacutainer (containing 7.2 mg K2EDTA; Becton Dickinson, Franklin Lakes, NJ, USA). The samples were then centrifuged at $1837 \times g$ for 5 min, to separate plasma. Plasma was mixed with 10 µl of reference compound 6 or 8, respectively (1 mg/ml) and injected onto an Oasis HLB column (Hydrophilic-Lipophilic Balanced; 4.6×20 mm, Waters) [25]. The proteins of the plasma matrix were washed from the Oasis column with 10 ml of water, which was collected as two 5-ml fractions. The outlet of the Oasis column was then connected to an analytical XTerra RP18 column (5 µm, 4.6×250 mm; Waters), and both columns in series were then eluted using 0.05 M ammonium acetate pH 6.9/acetonitrile (30/70 v/v, 1 ml/min). Brain samples were collected and homogenized in 4 ml of CH₃CN/ H_2O (50/50 v/v) followed by centrifugation at 1840×g for 10 min. The supernatant was treated with 1 ml of CH₃CN followed by centrifugation as described above. The supernatant is filtered through a 0.2-µm filter (Pall Corporation, Ann Arbor, MI, USA) and 1 ml of this solution was mixed with 10 μ g of reference compound 6 or 8 (1 mg/ml solution in CH₃CN) followed by injection onto an HPLC system which consisted of an XTerra RP18 column (5 µm, 4.6×250 mm; Waters) and ammonium acetate buffer 0.05 M pH 6.9/acetonitrile 40/60 (v/v) as the mobile phase (1 ml/min). After passage through an in-line UV detector, all HPLC eluates were collected as 1-ml fractions and their radioactivity was measured using a gamma counter.

2.8. Spleen autoradiography

For ex vivo studies, the spleen of vehicle-pretreated (control) or ligand-pretreated mice was removed 30 min pi of the tracer (9.25 or 0.74 MBq for $[^{11}C]$ -6 or $[^{18}F]$ -8, respectively) and washed with saline to remove blood pool activity. The tissues were then quickly frozen in isopentane cooled with liquid nitrogen to -50°C and cut with a cryotome (Shandon Cryotome FSE, Thermo Fisher, Waltham, MA, USA) into 50-µm-thick serial sections mounted on slides.

For in vitro studies, the spleen of NMRI mice was removed and sections of 20 μ m were cut. The tissue was preincubated for 10 min at RT with TRIS 50 mM pH 7.4 containing 5% bovine serum albumin (BSA) to remove endogenous ligands. Each slide was incubated with radio-

active solution (260 kBq in 200 μ l TRIS 50 mM pH 7.4, 5% BSA) and in the presence/absence of 'cold' ligand (6 or 9, 10 μ M) for 10 min at RT and rinsed twice (2×5 min) with TRIS 50 mM pH 7.4 containing 1% BSA and 5% ethanol at 4°C. After a quick dip in water at RT, the slides were dried.

Autoradiograms were obtained by exposing the slides overnight to a high-performance phosphor screen (Canberra-Packard, Meriden, CT, USA). The images were analyzed with Optiquant software (Canberra-Packard) and the radioactivity concentration in the autoradiograms was expressed in digital light units (DLU) per square millimeters.

2.9. Pretreatment and statistics

All compounds were dissolved (0.2 mg/ml) in aqua ad injectabilia containing 10% DMSO and 5% Tween 80. Injection of 1 mg/kg 'cold' compound was performed intraperitoneally 30 min before tracer injection. All control animals were pretreated with the same volume of vehicle.

All statistical analyses were performed with the unpaired two-sided Student's t test. A P value of less than .005 was considered statistically significant.

2.10. Competition binding assay and $[^{35}S]$ -GTP γS assay

2.10.1. Cell culture

All cell culture media and supplements were obtained from Invitrogen (Merelbeke, Belgium). Transfected CHO cells stably expressing the human CB₁ or CB₂ receptors (CHO-CB₁ and CHO-CB₂), kindly provided by Euroscreen (Gosselies, Belgium), were maintained using Ham's F12 medium supplemented with 10% foetal calf serum, 500 µg/ ml geneticin G418, 100 IU/ml penicillin, 100 µg/ml streptomycin, 2.5 µg/ml fungizone/amphotericin B and 2 mM L-glutamine. At confluence, cells were trypsinized for dilutions. Cells were cultured at 37°C in an atmosphere of humidified air and 5% CO₂.

2.10.2. Membrane preparation

Cells were lysed in ice-cold homogenization buffer containing 50 mM Tris-HCl, pH 7.4. The homogenate was centrifuged at $15,000 \times g$ for 10 min. The resulting membrane pellet was washed twice under the same conditions, resuspended in the same buffer and frozen (-80°C) in aliquots until use. Protein content was determined using the Bradford method with bovine serum albumin as standard [26].

2.10.3. Competition binding assay

The assay was performed as previously described [27]. Briefly, competition experiments were conducted on 40 μ g CHO-CB₁ or CHO-CB₂ membrane preparation incubated with 1 nM [³H]-SR141716A (Amersham, Roosendaal, The Netherlands) or [³H]-CP55,940 (NEN Life Sciences, Zaventem, Belgium) for determination of the affinity for hCB₁R and hCB₂R, respectively, and with decreasing concentrations of various competition ligands, in plastic tubes containing 0.5 ml final volume of binding buffer (50 mM Tris-HCl, 3 mM MgCl₂, 1 mM EDTA, 0.5% bovine serum albumin, pH 7.4). Nonspecific binding was determined in the presence of 10 μ M HU 210, a nonselective CB₁R and CB₂R agonist (Tocris Bioscience, Bristol, United Kingdom). After incubation for 1 h at 30°C, the solutions were filtered through 0.5% polyethyleneimine-pretreated glass fiber filters (Whatman, Maidstone, UK).

2.10.4. $\int_{-35}^{35} S - GTP \gamma S Assay$

Experiments (n=3, duplicates) were carried out in a 0.5ml total volume of buffer (50 mM Tris-HCl, 3 mM MgCl₂, 1 mM EDTA, 100 mM NaCl, 0.1% BSA, pH 7.4) containing 20 μ M GDP, 40 μ g protein samples from membrane preparation, 0.05 nM [³⁵S]-GTP γ S (Amersham, Roosendaal, the Netherlands) and test compounds [27]. Tubes were incubated 1 h at 30°C before filtration through nonpretreated glass fiber filters. Nonspecific binding was determined using 100 μ M Gpp(NH)p. The radioactivity was measured by liquid scintillation.

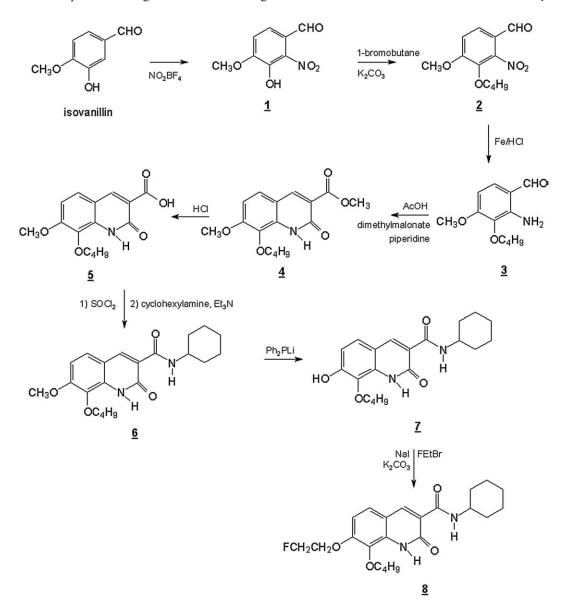
2.11. Data analysis

IC₅₀, EC₅₀ and efficacy were determined by nonlinear regression analysis performed using GraphPad Prism software (San Diego, CA, USA). K_i values were calculated following the Cheng–Prusoff equation: K_i =IC₅₀/(1+L/K_d), where L is the radioligand concentration [28].

3. Results

3.1. Synthesis and radiosynthesis

The synthesis of reference compounds and precursor is illustrated in Scheme 1. Synthesis of Compounds 1, 2, 3, 4, 5 and 6 was based on methods described by Raitio et al.



Scheme 1. Chemical synthesis of 6, 7 and 8.

[19]. Compound **6** was demethylated using lithium diphenylphosphide as a dealkylating agent. This method provides selective dealkylation of the methoxy group, as such avoiding cleavage of the butyloxy group [29]. Compound **8** was synthesized by reacting **7** with FEtBr in basic conditions.

Radiosynthesis of $[^{11}C]$ -6 was performed by bubbling $[^{11}C]CH_3I$ or $[^{11}C]$ methyl triflate through a solution of 7 in DMF in basic conditions (Scheme 2). When the reaction was done at RT (using $[^{11}C]CH_3I$), a low radiochemical yield of 4.2% was obtained. However, the radiochemical yield after heating the reaction mixture for 2 min at 90°C was 28±13% (average of 17 labeling reactions, % relative to starting $^{11}CH_3I$ activity) or $32\pm11\%$ (average of seven labeling reactions, % relative to starting $^{11}CH_3I$ activity) or $32\pm11\%$ (average of seven labeling reactions, % relative to starting $^{11}CH_3$ triflate activity) with a specific activity around 37,000 GBq/mmol and a radiochemical purity >99%. Using $[^{11}C]$ methyl triflate as alkylating agent does not seem to provide any advantage over using $[^{11}C]CH_3I$.

Radiolabeling of **8** was done by reacting [¹⁸F]FEtBr with **7** in DMF in the presence of Cs_2CO_3 (Scheme 2). At RT, no formation of [¹⁸F]-**8** was observed. After 10 min at 90°C, an average yield of 8.7% was obtained (*n*=4, EOS). Further heating did not improve the yield. The radio-chemical purity was >99% with a specific activity around 10,000 GBq/mmol.

3.2. Partition coefficient, polar surface area

The log*P*, calculated log*P* and tPSA of $[^{11}C]$ -6 and $[^{18}F]$ -8 are stated in Table 1 [30,31]. For comparison, the calculated log*P* and tPSA are given for 2-oxo-7-methoxy-8-pentyloxy-1,2-dihydroquinoline-3-carboxylic acid cyclohexylamide

Table 1 Log*P*, calculated log*P* and tPSA of $[^{11}C]$ -6, $[^{18}F]$ -8 and reference

compound 10	6		
Compound	LogP	Calculated logP	tPSA (Å ²)

Compound	LogP	Calculated logP	$tPSA(Å^2)$
[¹¹ C]-6	3.9±0.1	3.5±0.4	80.42
[¹⁸ F]- 8	3.2±0.2	3.8±0.5	80.42
10	_	4.1±0.4	80.42

(10), which is a potent CB_2 inverse agonist [32] and on which the synthesis of our tracers was based.

3.3. Affinity studies

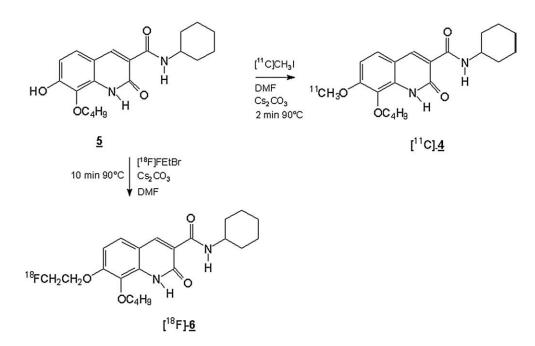
The affinity (K_i) of **6** and **8** for the human CB₂ receptor was 9.6 and 35.8 nM, respectively. They were highly selective towards the CB₁ receptor ($K_ihCB_1 > 1000$ nM). Both compounds behaved as inverse agonists in a [^{35}S]-GTP γS assay displaying a comparable potency (EC₅₀=11.8±2.2 and 12.5±3.2 nM for **6** and **8**, respectively) and negative efficacy (-69% and -78% relative to basal specific [^{35}S]-GTP γS binding for **6** and **8**, respectively).

3.4. Biodistribution studies

To determine specific CB_2 receptor binding, **9** was used as competing agent. This compound was described by Mussinu et al. [20] to have high affinity for the mouse CB_2 receptor (K_imCB₂=0.23 nM) and good selectivity over the mouse CB_1 receptor (K_imCB₁=1268 nM).

3.4.1. Biodistribution studies of $[^{11}C]$ -6 and $[^{18}F]$ -8 in mice

The distribution of $[^{11}C]$ -6 and $[^{18}F]$ -8 in mice was highly similar (Tables 2 and 3). Both compounds were cleared from



Scheme 2. Radiosynthesis of [¹¹C]-6 and [¹⁸F]-8.

Table 4

Table 2 Biodistribution of [¹¹C]-6 in control mice

Organ	Organ %ID±S.D.ª		SUV±S.D. ^b	,
	2 min	60 min	2 min	60 min
Kidneys	12.0±1.0	4.0±0.4	6.7±0.2	2.2±0.3
Urine	$0.1{\pm}0.1$	$0.9{\pm}0.5$		
Liver	38.2±2.7	21.2±1.0	$8.0{\pm}0.4$	4.2±0.2
Spleen	0.9±0.1	0.7±0.1	3.8±0.2	3.1±0.5
Pancreas	1.0±0.3	0.6±0.2	2.1±0.5	1.2 ± 0.2
Lungs	1.8±0.5	1.0 ± 0.1	2.5±0.5	1.3±0.2
Heart	1.7±0.2	0.5±0.1	4.0±0.3	1.0 ± 0.1
Intestines	10.5±0.6	29.8±3.7		
Stomach	1.6±0.3	$1.8{\pm}1.0$		
Blood	7.7±0.4	4.6±0.5	1.1±0.1	0.7 ± 0.1
Brain	1.7±0.3	$0.4{\pm}0.0$	1.9±0.3	$0.4{\pm}0.0$

Data are expressed as mean±S.D.; n=4 per time point.

^a Percentage of injected dose calculated as cpm in organ/total cpm recovered.

^b Standardized uptake values calculated as (radioactivity in cpm in organ/ weight of the organ in grams)/(total counts recovered/body weight in grams).

plasma via the hepatobiliary system where $[^{18}F]$ -8 was cleared more rapidly [14% vs. 21% of injected dose (ID) in the liver and 53% vs. 30% in the intestines 60 min pi for $[^{18}F]$ -8 and $[^{11}C]$ -6, respectively]. Both compounds showed a high initial kidney uptake, but no urinary excretion was observed. Except for the spleen, no major organ retained one of the studied tracers. The uptake of $[^{11}C]$ -6 in spleen was higher than that of $[^{18}F]$ -8 [standardized uptake values (SUVs) of 3.8 and 2.2 at 2 min pi, respectively]. For both tracers, more than 80% of the initial spleen uptake was still present at 60 min pi. Bone uptake was high for $[^{18}F]$ -8.

3.4.2. Biodistribution studies of $[^{11}C]$ -6 in mice pretreated with 6 or 9

Spleen mass did not vary significantly between the study groups. Pretreatment of the mice with 6 or 9 thirty minutes

Table 3 Biodistribution of $[^{18}F]$ -8 in control mice

Organ	% ID±S.D. ^a			SUV±S.D. ^b		
	2 min	60 min	120 min	2 min	60 min	120 min
Kidneys	11.3±1.2	2.8±0.3	2.7±0.2	6.2±0.9	1.4±0.2	1.4±0.1
Urine	0.1 ± 0.0	1.6 ± 0.4	1.2±1.3			
Liver	37.5±3.6	13.8±1.0	13.6±1.6	6.7 ± 0.9	2.3±0.3	2.3±0.2
Spleen	0.7 ± 0.1	$0.6{\pm}0.1$	0.5 ± 0.1	2.2 ± 0.6	1.9 ± 0.2	$1.9{\pm}0.5$
Pancreas	0.8 ± 0.1	0.3 ± 0.0	$0.4{\pm}0.1$	2.0 ± 0.2	0.6 ± 0.1	0.6 ± 0.1
Lungs	1.7±0.3	0.5 ± 0.1	0.5 ± 0.1	2.2 ± 0.4	0.6 ± 0.0	$0.7{\pm}0.1$
Heart	1.5 ± 0.1	$0.2{\pm}0.0$	0.2 ± 0.0	3.4 ± 0.4	0.5 ± 0.0	$0.4{\pm}0.1$
Intestines	10.8 ± 1.1	53.1±2.7	55.2±2.6			
Stomach	1.4±0.3	$1.0{\pm}0.6$	1.0±0.5			
Blood	5.8±0.4	2.7±0.1	2.2±0.9	0.8 ± 0.1	$0.4{\pm}0.0$	0.3±0.1
Brain	1.8±0.5	0.1 ± 0.0	0.1 ± 0.0	1.4±0.3	0.1±0.0	0.1 ± 0.0
Muscle	15.0±3.6	11.7±1.6	9.2±1.0	$0.4{\pm}0.1$	0.3±0.0	$0.2{\pm}0.0$
Bone	6.3±0.4	11.3±1.3	12.0±3.2	0.5 ± 0.0	$0.9{\pm}0.1$	1.0±0.3

Data are expressed as mean±S.D.; n=4 per time point.

^a Percentage of ID calculated as cpm in organ/total cpm recovered.

^b Standard uptake values calculated as (radioactivity in cpm in organ/ weight of the organ in grams)/(total counts recovered/body weight in grams).

60 min ^e 3.8±0.7 20.3±2.1	60 min ^c 1.7±0.2	60 min ^d	60 min ^e
	1.7±0.2	1.8+1.0	
20.3±2.1		1.0±1.0	1.7±0.3
	3.7 ± 0.2	3.8 ± 0.9	3.5 ± 0.5
$0.5 \pm 0.0^{*}$	2.8±0.3	$0.8 \pm 0.1*$	1.4±0.1*
1.0 ± 0.3	1.1 ± 0.1	1.4±0.3	1.2±0.3
0.7 ± 0.0	1.0 ± 0.1	0.8 ± 0.2	0.9 ± 0.1
0.5±0.3	0.8 ± 0.1	$0.9{\pm}0.4$	0.9 ± 0.5
25.1±1.8			
1.1±0.2			
3.9 ± 0.6	0.5 ± 0.0	0.6 ± 0.1	0.5 ± 0.1
2.740.0	0.3 ± 0.0	0.3 ± 0.0	0.3±0.2
	3.9±0.6 0.4±0.2	3.9±0.6 0.5±0.0	3.9±0.6 0.5±0.0 0.6±0.1

Data are expressed as mean±S.D.; n=4 per time point.

^a Percentage of injected dose calculated as cpm in organ/total cpm recovered.

^b Standardized uptake values calculated as (radioactivity in cpm in organ/weight in grams)/(total counts recovered/body weight in grams).

^c Pretreatment vehicle.

- ^d Pretreatment **6**.
- ^e Pretreatment 9.

* P<.005 with respect to the vehicle pretreated at the same time point.

before tracer injection resulted in significantly lower spleen values, whereas for the other major organs, no difference was observed (Table 4; *P*<.005).

The vehicle used for injection of the 'cold' product had no influence on the biodistribution data compared to untreated animals (Tables 2 and 4). When SUVs were compared, both groups were not significantly different (P<.05, Bonferroni corrected).

3.5. Plasma and brain metabolite analysis

Both tracers were rapidly metabolized in plasma with less than 20% intact product left at 30 min pi. Only radiometabolites more polar relative to intact tracer were found. In brain, a relatively small amount (<20%) of radiometabolites was observed (Table 5). In view of the large fraction of radiometabolites in plasma, this may be due to the presence of radiometabolites in the vascular compartment of the brain. Further studies using brain perfusion prior to brain radiometabolite quantification will be required to ascertain whether radiometabolites pass the BBB.

Table 5

Fraction of intact tracer of $[^{11}C]$ -6 and $[^{18}F]$ -8 determined in the plasma and brain of normal mice at 2, 10 or 30 min pi

	[¹¹ C]- 6	[¹⁸ F]-8	
Plasma			
2 min	68±3	86±5	
10 min	24±1	41±6.	
30 min	17±0	12±1	
Brain			
2 min	95±0	92±1	
10 min	ND	82±3	

Data are expressed as mean \pm S.D.; n=2 or 3 per time point. ND=not determined.



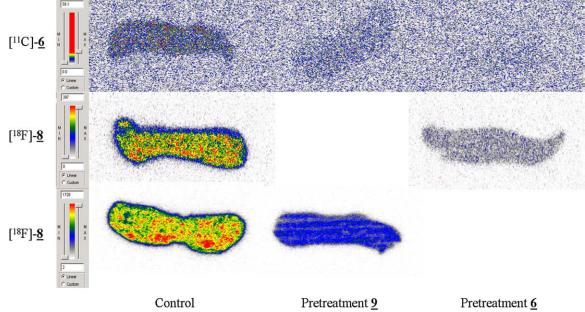


Fig. 1. Ex vivo autoradiography of mouse spleen. Spleen from mice pretreated with vehicle (control) or cold competitor either 9 or 6 (pretreated groups) and injected with either $[^{11}C]$ -6 or $[^{18}F]$ -8.

3.6. Ex vivo and in vitro spleen autoradiography

Quantification of the autoradiography images (comparison DLU/mm² for whole spleen slice) obtained from spleen sections of mice injected with the radiotracers showed that after pretreatment with **9**, $[^{11}C]$ -**6** spleen uptake was about one third of the spleen uptake in vehicle-injected mice. This result is in accordance with the biodistribution data where pretreatment with **9** reduced the spleen SUV more than two times at 60 min pi. Pretreatment with **6** resulted in a 10–15 times lower spleen uptake. However, spleen slices obtained from the mice after pretreatment with **6** were barely distinguishable from the background so that quantification may not be accurate. Even so, pretreatment with **6** gave also in the biodistribution study a more pronounced fall in spleen uptake compared to pretreatment with **9**. When using $[^{18}F]$ -**8**, pretreatment with **6** inhibited the spleen activity to about one fifth of the spleen uptake in control mice. $[^{18}F]$ -**8** spleen uptake values obtained after pretreatment with **9** were about threefold lower than control values (Fig. 1).

The ex vivo autoradiography data were confirmed by in vitro autoradiography in which mouse spleen sections were incubated with [¹¹C]-6 in the presence or absence of CB₂-specific compound 6 or 9 (10 μ M) (Fig. 2). The presence of 10 μ M of 6 or 9 resulted in a binding inhibition of about 35%.

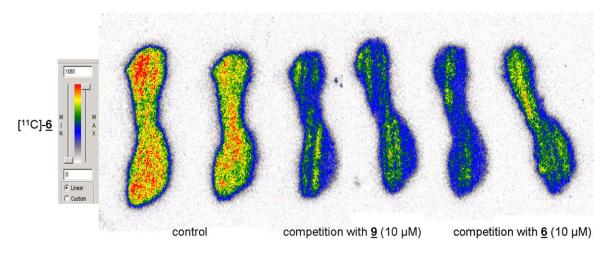


Fig. 2. In vitro autoradiography on mouse spleen.

4. Discussion

We have, for the first time, demonstrated radiosynthesis of a high-affinity CB_2 selective tracer which passes the BBB. Selection of the tracer was based on literature data describing several 2-oxoquinoline derivatives as high and selective CB_2 receptor ligands (e.g., **10** [32]). We shortened the lipophilic carbon tail from a pentoxy to a butoxy group, which resulted in a decrease of 0.6 in calculated log*P* value with the aim to decrease non specific binding related to the lipophilicity of the tracer (Table 1).

Synthesis of the precursor 7 was quite challenging. Indeed, we tried several dealkylation agents without success; however, using lithium diphenylphosphide as dealkylation agent gave a selective cleavage of the methoxy group with a reasonable yield of 29%. The yield using [¹¹C]CH₃I or [¹¹C] methyl triflate was much higher than using [¹⁸F]FEtBr in the same labeling conditions. Note that the labeling procedure with F-18 was not yet optimised.

The affinity for the human CB₂ receptor was in the low nanomolar range, thereby comparable to the affinity of the nonselective CB₂/CB₁ agonist HU-210 (7.3 \pm 0.9) and the selective CB₂ inverse agonist SR144528 (51.7 \pm 4.8) when tested in the same conditions [33]. This indicates that **6** and **8** bind to the human CB₂ receptor with high affinity. However, the fluoroethoxy group on position 7 seems to have a slightly negative effect on the affinity data when compared to a methoxy group in this position. Note that species differences in CB₂ receptor binding data have been reported in the literature [34]. The studied molecules are structurally related to JTE-907 of which a higher mouse CB₂ receptor affinity compared to the human has been reported (1.55 vs. 35.9 nM, respectively) [35]. One could therefore speculate the affinity for the mouse CB₂ receptor to be higher for both tracers.

The ideal PSA value for molecules to penetrate the BBB is between 0 and 90 [36], whereas the $\log P$ is ideally between 1 and 3 [37]. Although the $\log P$ value of both tracers is higher than 3, they both show good uptake in mouse brain (1.7–1.8% of ID at 2 min pi) and efficient washout from brain.

Surprisingly, **8** seems to be less lipophilic than **6**, as it is characterized by both a lower log*P* value and a shorter retention time on the same analytical HPLC system. Therefore, background noise due to nonspecific binding to brain structures could be lower for $[^{18}F]$ -**8**. This is confirmed by the biodistribution data where brain washout (2 min/ 60 min ratio ID) is higher for $[^{18}F]$ -**8** (ratio=18) compared to $[^{11}C]$ -**6** (ratio=4.3).

Both tracers are cleared from plasma by the hepatobiliary system as can be expected from their lipophilicity. They show persistent spleen uptake, an organ which is known for the high levels of CB₂ expression [4]. The spleen uptake was found to be inhibited in vitro and in vivo by both **6** and the structurally unrelated CB₂-specific compound **9**, strongly indicating that the observed spleen uptake is CB₂ specific. The rather slow uptake of $[^{18}F]$ -**8** in bone suggests in vivo defluorination of the tracer rather than CB₂-specific bone uptake.

Although both tracers are swiftly and extensively metabolised, this metabolism pattern does not seem to preclude CB₂ receptor binding as indicated by the specific spleen uptake and retention.

As discussed in the Introduction, neuroinflammation results in marked CB_2 receptor up-regulation, most likely due to up-regulation in activated microglial cells [9]. Therefore, these tracers could be useful for imaging the CB_2 receptor in the described pathological conditions in the brain. Future work will include imaging of neuroinflammation in several animal models, e.g., such as for multiple sclerosis and amyotrophic lateral sclerosis followed by clinical studies.

5. Conclusion

We successfully synthesized two novel CB_2 receptor PET tracers that show high affinity for the human CB_2 receptor and in vivo specific CB_2 receptor binding in the spleen of mice. Therefore, these tracers are promising candidates for the evaluation of the CB_2 receptor in vivo and will be further investigated in preclinical and human studies.

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