

In Vivo Induction of P-Glycoprotein Function can be Measured with [¹⁸F]MC225 and PET

Lara García-Varela, Manuel Rodríguez-Pérez, Antía Custodia, Rodrigo Moraga-Amaro, Nicola A. Colabufo, Pablo Aguiar, Tomás Sobrino, Rudi A.J.O. Dierckx, Aren van Waarde, Philip H. Elsinga, and Gert Luurtsema*

Cite This: *Mol. Pharmaceutics* 2021, 18, 3073–3085

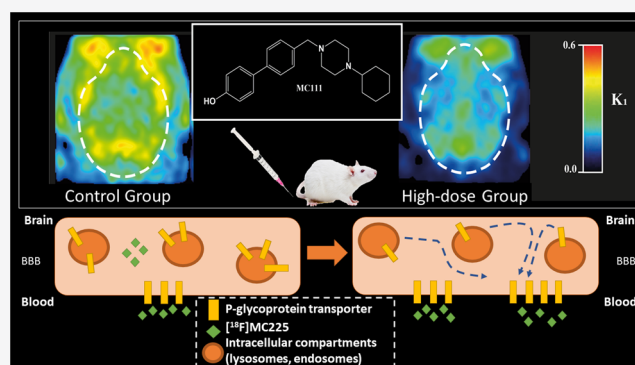
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ABSTRACT: P-Glycoprotein (P-gp) is an efflux pump located at the blood–brain barrier (BBB) that contributes to the protection of the central nervous system by transporting neurotoxic compounds out of the brain. A decline in P-gp function has been related to the pathogenesis of neurodegenerative diseases. P-gp inducers can increase the P-gp function and are considered as potential candidates for the treatment of such disorders. The P-gp inducer MC111 increased P-gp expression and function in SW480 human colon adenocarcinoma and colo-320 cells, respectively. Our study aims to evaluate the P-gp inducing effect of MC111 in the whole brain *in vivo*, using the P-gp tracer [¹⁸F]MC225 and positron emission tomography (PET). Eighteen Wistar rats were treated with either vehicle solution, 4.5 mg/kg of MC111 (low-dose group), or 6 mg/kg of MC111 (high-dose group). Animals underwent a 60 min dynamic PET scan with arterial-blood sampling, 24 h after treatment with the inducer. Data were analyzed using the 1-tissue-compartment model and metabolite-corrected plasma as the input function. Model parameters such as the influx constant (K_1) and volume of distribution (V_T) were calculated, which reflect the *in vivo* P-gp function. P-gp and pregnane xenobiotic receptor (PXR) expression levels of the whole brain were assessed using western blot. The administration of MC111 decreased K_1 and V_T of [¹⁸F]MC225 in the whole brain and all of the selected brain regions. In the high-dose group, whole-brain K_1 was decreased by 34% (K_1 -high-dose = 0.20 ± 0.02 vs K_1 -control = 0.30 ± 0.02 ; $p < 0.001$) and in the low-dose group by 7% (K_1 -low-dose = 0.28 ± 0.02 vs K_1 -control = 0.30 ± 0.02 ; $p = 0.42$) compared to controls. Whole-brain V_T was decreased by 25% in the high-dose group (V_T -high-dose = 5.92 ± 0.41 vs V_T -control = 7.82 ± 0.38 ; $p < 0.001$) and by 6% in the low-dose group (V_T -low-dose = 7.35 ± 0.38 vs V_T -control = 7.82 ± 0.37 ; $p = 0.38$) compared to controls. k_2 values did not vary after treatment. The treatment did not affect the metabolism of [¹⁸F]MC225. Western blot studies using the whole-brain tissue did not detect changes in the P-gp expression, however, preliminary results using isolated brain capillaries found an increasing trend up to 37% in treated rats. The decrease in K_1 and V_T values after treatment with the inducer indicates an increase in the P-gp functionality at the BBB of treated rats. Moreover, preliminary results using brain endothelial cells also sustained the increase in the P-gp expression. In conclusion, the results verify that MC111 induces P-gp expression and function at the BBB in rats. An increasing trend regarding the P-gp expression levels is found using western blot and an increased P-gp function is confirmed with [¹⁸F]MC225 and PET.



KEYWORDS: ABC transporters, brain Imaging, efflux transporters, P-glycoprotein, P-gp inducers

INTRODUCTION

P-glycoprotein (P-gp) is an efflux transporter that belongs to the ATP-binding cassette (ABC) transporter family and is located at the luminal side of the endothelial cells at the blood–brain barrier (BBB) and other blood-tissue barriers.¹ P-gp pumps a wide variety of compounds, including endogenous and exogenous substances from the brain to the circulation, by adenosine triphosphate (ATP)-driven transport.² Thereby, P-gp located at the BBB contributes to the protection of the brain from harmful substances and maintains the homeostasis needed

Received: April 14, 2021
Revised: June 24, 2021
Accepted: June 24, 2021
Published: July 6, 2021



for the appropriate neural function.³ However, several factors can affect P-gp function or expression such as inflammatory responses, stress, therapeutic drugs, and diet.⁴

A decline in the P-gp function at the BBB can increase the concentration of neurotoxic compounds inside the brain, which can be related to the onset and progression of several neurodegenerative diseases such as Alzheimer's (AD). It has been reported that P-gp is involved in the clearance of A β peptides since the inhibition of P-gp decreased extracellular levels of A β secretions *in vitro*.⁵ Moreover, an inverse correlation between A β deposition and expression of P-gp was found in the BBB of elderly nondemented subjects.⁶ Also, positron emission tomography (PET) imaging studies performed in AD patients found an increase of [¹¹C]verapamil uptake inside the brain compared to healthy volunteers, which indicates a decreased P-gp function.⁷

Several studies have already reported the use of P-gp inducers to restore the function of P-gp in neurodegenerative diseases.^{8,9} P-gp induction does not occur via direct binding to the P-gp transporter, and P-gp expression and function can be regulated by different mechanisms including the activation of several transcription factors and post-transcriptional and post-translational mechanisms.^{10–12} Recently, it has been proposed that an increase in P-gp expression at the cell surface can be caused by relocating intracellular P-gp protein to the plasma membrane.^{10,11} Among the transcription factors, the nuclear receptors pregnane xenobiotic receptor (PXR) and constitutive androstane receptor (CAR) have been identified as key factors in the P-gp upregulation.¹³ The PXR and CAR can be activated by endogenous or exogenous compounds.¹⁴ PXR is activated not only by steroids, chemotherapeutics, HIV protease inhibitors, glucocorticoids, and anticonvulsants but also by dietary compounds and natural steroids such as pregnenolone and progesterone.¹⁵ CAR is activated by endogenous ligands such as bile acids and also by drugs, dietary compounds, and pollutants.¹⁶

Since P-gp inducers can play an important role in the treatment of neurodegenerative diseases,^{8,9} several preclinical studies have already reported the efficacy of such drugs. A study performed in a transgenic mouse model of human AD found a reduced P-gp function in these animals compared to wild-type mice. After treatment with pregnenolone-16 α -carbonitrile for 7 days, PXR was activated, and consequently, the P-gp function was restored resulting in an increase of A β clearance from the brain.¹⁷ 1,25-Dihydroxyvitamin D3-enriched diet also induced the P-gp function and reduced the A β deposition in mice models of AD. In contrast, AD model mice fed with a vitamin D-deficient diet showed a reduction in P-gp expression and increased A β plaques in the brain as well as increased cognitive deficits.¹⁸ The administration of St. John's Wort, which is a herbal drug used in the treatment of depression, to transgenic AD mice enhanced the cerebral P-gp function and reduced A β accumulation.¹⁹

Recently, a new promising compound, 4'-((4-cyclohexylpiperazin-1-yl)methyl)-[1,1'-biphenyl]-4-ol (MC111) has been identified as a P-gp inducer by *in vitro* studies. MC111 activity was assessed in colo-320 cells. After treatment with MC111, an increase in [³H]verapamil efflux was observed, which indicates an increase in the P-gp function. Western blot studies also indicated that MC111 increases P-gp expression in SW480 human colon adenocarcinoma cells.²⁰ Furthermore, this new compound also activates the breast cancer resistance protein (BCRP) function and expression in colo-320 and MCF7 cells.²⁰

The dual activity of MC111 may be beneficial for the treatment of several neurodegenerative diseases where not only an impaired P-gp function and expression has been observed but also alterations in the BCRP and other ABC transporters.^{21,22}

This study aims to demonstrate the P-gp induction effect of MC111 *in vivo* at the BBB of healthy rats. For this purpose, we used positron emission tomography (PET) imaging, which is a noninvasive imaging technique that quantifies the biochemical process *in vivo*, and the tracer [¹⁸F]MC225 for measuring the P-gp function. [¹⁸F]MC225 is considered as a weak P-gp substrate and therefore, it shows relatively high uptake in the brain at baseline conditions when the P-gp transporter is completely functional. This characteristic of [¹⁸F]MC225 could be beneficial for imaging increased P-gp function upon treatment with an inducer when the tracer uptake in the brain is expected to decrease.^{23,24}

In the present study, we will evaluate the effect of an inducer for the first time *in vivo*, after its administration to living subjects. To this aim, a dynamic PET scan will be made in eighteen healthy male rats. The kinetics of [¹⁸F]MC225 will be assessed by the 1-tissue compartment model (1-TCM) and changes in the P-gp function will be reflected by the influx constant (K_1) and volume of distribution (V_T). Moreover, the P-gp expression level in brain tissue will be evaluated postmortem using western blots. The results may yield insight into the potential therapeutic application of MC111 in diseases where the P-gp function/expression is decreased.

■ EXPERIMENTAL SECTION

Chemicals. 6,7-Dimethoxy-2-(3-(5-(2-((methylsulfonyl)oxy)ethoxy)-1,2,3,4-tetrahydronaphthalen-1-yl)propyl)-1,2,3,4-tetrahydroisoquinolin-2-ium methanesulfonate salt (mesylate [¹⁸F]MC225 precursor) was purchased from Syncom (Syncom, Groningen, the Netherlands). The inducer MC111 (4'-((4-cyclohexylpiperazin-1-yl)methyl)-[1,1'-biphenyl]-4-ol) was supplied by the University of Bari, Italy. Chemicals were purchased from Sigma-Aldrich (Sigma-Aldrich, St. Louis, MO) and isoflurane from Pharmachemie (Pharmachemie, Haarlem, the Netherlands).

Tracer Production. [¹⁸F]MC225 was synthesized by a one-step reaction.²⁵ Briefly, the radiosynthesis was automated and performed in Synthera modules (IBA RadioPharma Solutions, Ottignies-Louvain-la-Neuve, Belgium). After drying the [¹⁸F]-Fluoride, the mesylate [¹⁸F]MC225 precursor dissolved in dimethylformamide (DMF) was added to the vial and the mixture was heated at 140 °C for 30 min. After purification on a semipreparative high-performance liquid chromatography (HPLC) system, the radiotracer was reformulated to 15% ethanol in sterile 0.9% NaCl solution. Quality control procedures were performed in an ultraperformance liquid chromatography (UPLC) system, as previously described.²⁴

Animals. Eighteen healthy male Wistar rats (RjHan: WI) were obtained from Janvier Labs (Janvier Labs, Le Genest-Saint-Isle, France) (281 \pm 9 g). Before starting the experiments, rats were acclimatized to the new environment for at least 7 days. Rats were housed in groups of 2 in a temperature-controlled room (21 \pm 2 °C) with 12 h dark/12 h light cycles. Water and standard laboratory chow were available *ad libitum*. All animal experiments were performed in compliance with Dutch and EU regulations. The experimental protocol was approved by the National Committee on Animal Experiments of the Netherlands (CCD, the Hague) and the Institutional Animal Care and Use Committee of the University of Groningen (CCD license

The amygdala, cerebellum, corpus callosum, medial geniculate, mesencephalic, septum, superior colliculus, striatum, cortex, hippocampus, hypothalamus, midbrain, brainstem, thalamus, basal ganglia, and a whole-brain region were selected as volumes of interest (VOI) for this study. The radioactivity concentration in each VOI (KBq/mL) was calculated and corrected for the injected dose (KBq) and the bodyweight of the animal (g) and thus, it was expressed as the standardized uptake value (SUV).

Input Function Analysis. Time-activity curves (TAC) of whole blood, plasma, and plasma-corrected for metabolites were calculated using the blood measurement and expressed in SUV. The metabolite-corrected plasma TAC–SUV was obtained by multiplying the SUV values of the plasma samples by the percentage of the parent fraction in each time-point.

The first-order rate constant of elimination (K_e) was calculated by fitting a single exponential curve to the metabolite-corrected plasma TAC–SUV of each subject, by an iterative nonlinear least-squares approach using GraphPad software: $Y = Y_0 \times \exp(-K_e \times X)$, where Y is the metabolite-corrected SUV value in plasma; Y_0 is the Y -intercept; K_e is the first-order rate constant of elimination, and X is the time.³¹

Pharmacokinetic Modeling. Measured radioactivity in whole blood and plasma as well as the percentage of the [¹⁸F]MC225 parent tracer were used as input functions for the pharmacokinetic analysis (PMOD Technologies).

The 1-tissue compartmental model (1-TCM) has been selected as the model of choice to analyze the [¹⁸F]MC225 data in rats and the cerebral volume of blood was fixed to 5%.^{24,29} The TACs were corrected for blood delay. First, the blood delay of the whole-brain region TAC was estimated and then this value was fixed for the rest of the brain regions.

The influx rate constant K_1 is considered as the best parameter to measure the P-gp at the BBB,^{24,32,33} thus the study evaluated the changes in K_1 caused by the administration of the treatment. Since the volume of distribution (V_T) can also be used to assess changes in the P-gp function, this parameter was also calculated.²⁴ Some studies reported changes in the efflux constant k_2 due to the P-gp inhibition,^{34,35} thus the effect of the treatment on k_2 values was also evaluated.

Parametric Images. Parametric images representing K_1 , V_T , and k_2 values of three rats from different groups (control, low-dose, and high-dose group) were used for illustrative purposes. To this aim, the metabolite-corrected plasma TAC of each subject was used as the input function for 1-TCM analysis (vB = 5%), using PMOD.

Western Blot Analysis. P-glycoprotein (P-gp) and pregnane X receptor (PXR) expression levels in whole-brain tissue were quantified by means of western blot.

Protein extraction was carried out by adding cold RIPA buffer (Sigma-Aldrich) with the protease inhibitor cocktail (Sigma-Aldrich) to brain slices in an approximated 10:1 proportion (volume/weight), and tissue disruption was performed in a TissueLyser II (Qiagen, Hombrechtikon, Switzerland). Then, tissue lysates were centrifuged for 30 min at 20 000g, and supernatants were collected to perform western blot. Protein concentration in the lysates was quantified using a micro BCA Protein assay kit (Thermo Fisher, Waltham, MA).

The necessary volume of cell lysate containing a total amount of 25 μ g of protein was subjected to sodium dodecyl sulfate–polyacrylamide gel electrophoresis (SDS-PAGE) in 4–15% Criterion TGX precast midi protein gel (BioRad, Hercules, CA) using a constant voltage of 140 V. Then, proteins were

transferred onto a poly(vinylidene difluoride) (PVDF) membrane (Millipore, Cork, Ireland) using a Trans-Blot semidry system (BioRad) with a limited voltage of 25 V and 180 mA for 2 h. After the blotting step, membranes were blocked for 1 h with a 3% bovine serum albumin (BSA) solution (in Tris-Chloride Buffer with a 0.1% of Tween 20 (TBST)). Once blocked, target proteins were detected through incubation with primary antibodies against P-gp (Rabbit monoclonal to P-glycoprotein ab170904, Abcam, Cambridge, U.K.), PXR (Rabbit polyclonal to PXR ab192579, Abcam), β -Actin (Mouse monoclonal to β -Actin ab8226, Abcam), and histone H3 (Rabbit polyclonal to H3, ab1791, Abcam) diluted 1:1000, 1:1000, 1:5000, and 1:1000, respectively, in TBS with 3% of BSA. Primary antibodies were incubated overnight at room temperature under agitation, and then membranes were washed three times in TBST to remove the excess of primary antibodies and avoid unspecific signaling. As secondary antibodies, an HRP-conjugated Goat anti-Rabbit IgG (P044801-2, Dako, Denmark) and HRP-conjugated Rabbit anti-Mouse IgG (P026002-2) were used. They were diluted to 1:5000 in TBS with 3% BSA and incubated for 1 h at room temperature under agitation. Eventually, after washing, the HRP activity was revealed with a Pierce ECL western blotting substrate (Thermo Fisher) and detected in a Chemi Doc MP imaging system (BioRad). WB results were analyzed by measuring the mean gray value of protein bands delimited in ROIs using ImageJ software (Rasband, WS). The relative expression of P-gp to β -Actin and PXR to histone H3 was calculated for each sample, and the average of samples for each group was normalized to the control.

As previously mentioned, additional experiments were performed to evaluate the changes in the P-gp expression level in isolated brain capillaries. Animals were sacrificed, and brains were immediately collected in ice-cold Dulbecco's phosphate-buffered saline (DPBS) (Cytiva, Global Life Sciences Solutions, Marlborough, MA). The isolation of the capillaries was performed, as previously described by Hartz et al.^{26,27} Briefly, meninges were carefully removed, and the olfactory bulb, cerebellum, and white matter structures were separated from the cortex. The cortex was homogenized in DPBS with Ficoll PM400 (Cytiva, Global Life Sciences Solutions) and centrifuged at 5800g for 20 min at 4 °C. The supernatant was discarded, and the pellet was resuspended in 1% BSA–DPBS buffer. The suspension was filtered through a 200 μ m mesh and after several washing steps, the filtrate was centrifuged at 1500g for 3 min at 4 °C. Afterward, the pellet was flash-frozen and stored at –80 °C. The amount of protein in each sample was also quantified using a micro BCA protein assay kit, and western blot studies were performed loading 40 μ g of the protein in each well. Western blots using the brain isolated endothelial cells were carried out following the previously described protocol.

Statistical Analysis. Descriptive data are presented as mean \pm standard deviation (SD), whereas the results of the statistical analysis are shown as mean \pm standard error (SE) unless otherwise indicated. IBM SPSS 23 (Chicago, IL) was used for the statistical analysis. Generalized estimated equation (GEE) analysis with the independent matrix³⁶ was used to assess the differences in the SUV–TAC of the blood and brain. Generalized linear models (GLM) were used to evaluate the differences in SUV_{40–60 min}, pharmacokinetic parameters (V_T , K_1 , k_2), and blood kinetics (K_e) among the groups, independently for each region. P -values < 0.05 were considered statistically significant. Relative changes among the groups were calculated, using the control as the reference.

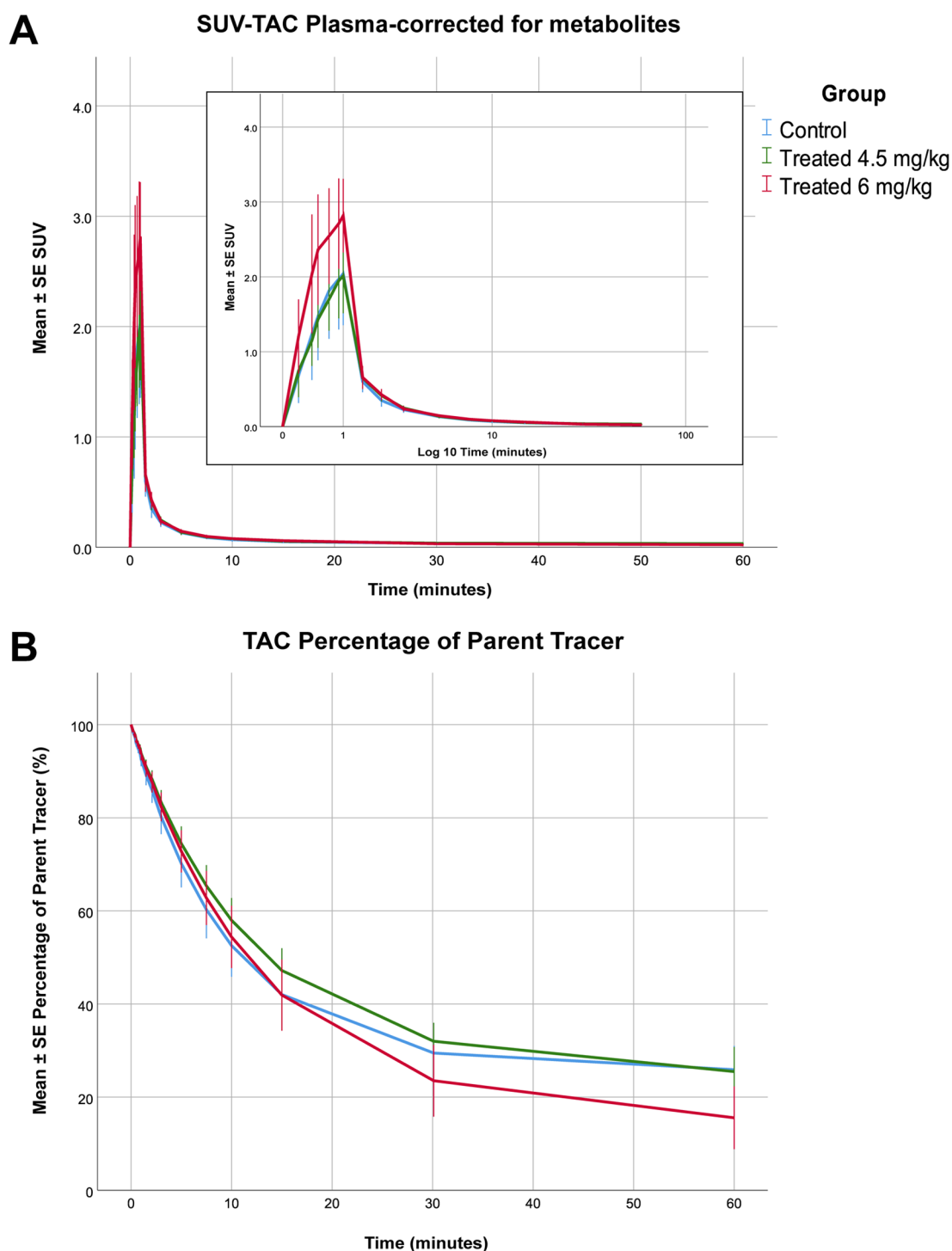


Figure 2. Metabolite-corrected SUV–TACs for plasma of the control (blue), low-dose (green), and high-dose groups (red) (A) and the percentage of plasma radioactivity representing [^{18}F]MC225 as a function of time (B).

RESULTS

Tracer Production. The final product was obtained in $5.6 \pm 2.9\%$ yield (mean \pm SD) ($n = 9$) (decay-corrected) in an average synthesis time of 121 ± 24 min (mean \pm SD). The radiochemical purity of [^{18}F]MC225 was $95.3 \pm 2.7\%$ (mean \pm SD) and the molar activity higher than 80 TBq/mmol (mean \pm SD).

Plasma Kinetics and Metabolism of [^{18}F]MC225. No significant differences between the control and the treated animals were found concerning the parent fraction in plasma.

However, statistical analysis revealed that the (metabolite-corrected) SUV–TAC of [^{18}F]MC225 in plasma was higher in the high-dose group than in the other two groups. The SUV–TAC of the tracer was 44% ($p = 0.042$) higher in the high-dose group than in the control group (high-dose group = 0.97 ± 0.10 , low-dose group = 0.67 ± 0.06 , control group = 0.70 ± 0.10) (Figure 2).

The statistical analysis found a significant increase in the K_e in the 6 mg/kg treated group. The K_e from the high-dose group increased by 25% compared to the control group (K_e high-dose

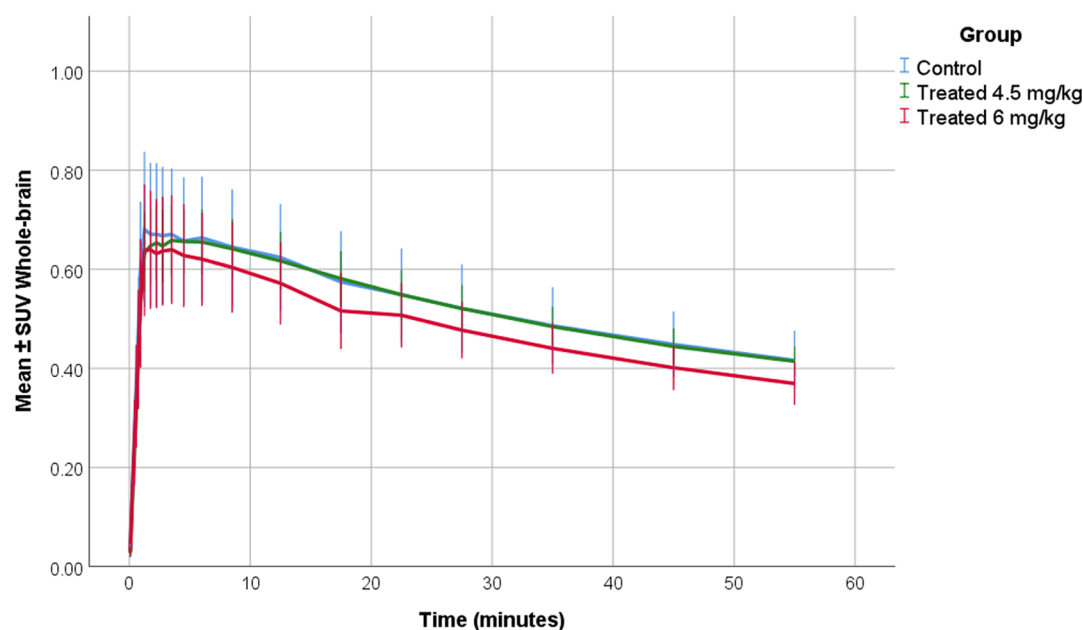


Figure 3. SUV–TACs of the whole brain in the control (blue color), low-dose (4.5 mg/kg) (green color), and high-dose groups (6 mg/kg) (red color).

group = $0.048 \pm 0.003 \text{ s}^{-1}$ vs K_e control group = $0.039 \pm 0.003 \text{ s}^{-1}$, $p = 0.014$). However, no significant differences were found between the control group and the low-dose group (K_e low-dose group = $0.035 \pm 0.003 \text{ s}^{-1}$ vs K_e control group = $0.039 \pm 0.002 \text{ s}^{-1}$, $p = 0.35$).

Brain Kinetics of [^{18}F]MC225. SUV–TAC of the Whole Brain. SUV–TACs of the whole brain were calculated. In all groups, [^{18}F]MC225 showed an initially high uptake followed by a subsequent decrease (Figure 3). In all groups, the peak was observed at $3.3 \pm 2.4 \text{ min}$ (mean \pm SD) after injection of the tracer. Statistical analysis did not find any significant differences in the SUV–TAC of the whole brain among the groups.

SUV_{40–60 min} of the Whole-Brain Region. Since the last part of the PET scans shows more stable SUV values, a mean SUV value from 40 to 60 min was calculated for all of the animals (SUV_{40–60 min}). The SUV_{40–60 min} calculated using the last 20 min of the scan decreased in the high-dose group with regard to the control group, but this difference was not statistically significant (SUV_{40–60 min} control = 0.45 ± 0.02 ; SUV_{40–60 min} low-dose = 0.45 ± 0.02 ; SUV_{40–60 min} high-dose = 0.40 ± 0.02).

Pharmacokinetic Modeling. A low dose of 4.5 mg/kg MC111 decreased the V_T and K_1 values compared to the control group, however, the decrease was not statistically significant. The higher dose (6 mg/kg) caused a significant reduction in the K_1 and V_T values in all of the brain regions compared to the control group. The analysis also found significant differences in K_1 and V_T between the high-dose group and the low-dose group. Figure 4 shows the V_T , K_1 , and k_2 values of all of the brain regions in all of the experimental groups.

The whole-brain region K_1 significantly decreased by 34% ($p < 0.001$) after the high-dose treatment compared to the control group, and the whole-brain region V_T also decreased by 24% ($p < 0.001$). A similar trend was observed in all brain regions. The largest change in the K_1 was observed in the septum, the superior colliculus, and the cortex region with a decrease of 39, 38, and 36% compared to the control group, respectively. In the case of V_T , the largest changes were found in the brainstem, the thalamus, and the superior colliculus with reductions of 30, 27,

and 26%, respectively (more information in Table 1). The analysis did not find significant differences in k_2 values between the low-dose group and the control in any of the brain regions. However, k_2 values decreased in the high-dose group compared to the control.

Parametric Images. Figure 5 shows parametric maps of K_1 , V_T , and k_2 in three representative subjects of each group using 1-TCM.

Western Blot. Western blot analysis using the whole brain showed no statistical differences between the control and treated groups. There was a noticeable intragroup variability regarding P-gp analysis, but no significant differences were found (Figure 6). In PXR analysis, there was a tendency to lower levels of PXR in the high-dose group (Figure 7 and 8).

Preliminary results using western blot analysis in isolated brain endothelial cells showed an increasing trend up to 37% ($p = 0.087$) in treated rats compared to control rats.

DISCUSSION

The present study was designed to determine the effect of MC111 on the P-gp function *in vivo* using the P-gp tracer [^{18}F]MC225 and PET. Our analysis highlights the induction effect of MC111 on the P-gp function at the BBB of healthy rats. This finding was in line with the results that were previously observed *in vitro*.²⁰ Moreover, the results demonstrate that [^{18}F]MC225 can be used to measure increases in the P-gp function reflected in decreases in K_1 and V_T values of [^{18}F]MC225 in all brain regions. Western blot analyses performed using the isolated capillaries cells of the rats found an increasing trend regarding the P-gp expression levels in rats treated with the inducer.

The blood analysis revealed that the treatment with MC111 did not alter the metabolism of the tracer since the analysis did not find significant differences in the percentages of radiometabolites among the groups. However, the treatment with 6 mg/kg dose of MC111 increased the (metabolite-corrected) concentration of the tracer in plasma compared to the control group and low-dose group. However, the elimination of the

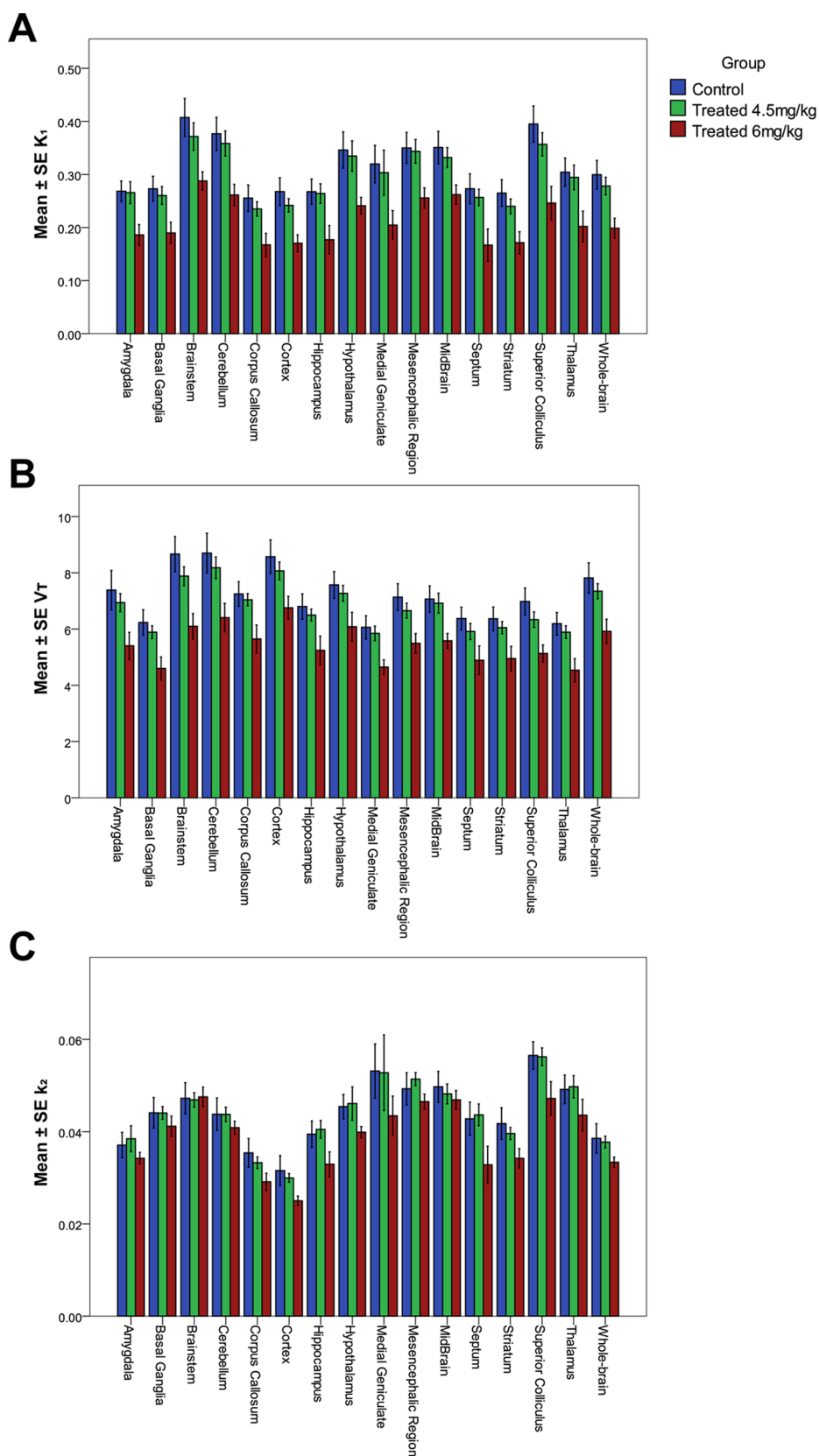


Figure 4. Mean \pm SE values of K_1 (A), V_T (B), and k_2 (C) in the control (blue), low-dose (4.5 mg/kg) (green), and high-dose (6 mg/kg) (red) groups.

tracer from plasma (K_e) was also higher in the high-dose group than in the other two groups, which means that the washout of the tracer from the blood was faster in the high-dose group. Therefore, it seems that although the concentration of the tracer

in plasma was higher in the high-dose group, the washout of the tracer was also faster. One rat (from the high-dose group) showed very high concentration levels of [^{18}F]MC225 in blood

Table 1. Mean \pm SE K_1 and V_T Values of All Selected Brain Regions in All Experimental Groups, Percentages of Change Relative to the Control Group, and p -Values

| region | group | K_1 (mL/(mL min)) | | | V_T (mL/mL) | | |
|----------------------|-----------|---------------------|---------------------------------|------------|-----------------|---------------------------------|------------|
| | | mean \pm SE | changes relative to control (%) | p -value | mean \pm SE | changes relative to control (%) | p -value |
| amygdala | control | 0.27 \pm 0.02 | | | 7.38 \pm 0.47 | | |
| | 4.5 mg/kg | 0.27 \pm 0.02 | −0.94 | 0.92 | 6.94 \pm 0.47 | −6.03 | 0.50 |
| | 6 mg/kg | 0.19 \pm 0.02 | −30.62 | 0.00 | 5.41 \pm 0.51 | −26.81 | 0.00 |
| basal ganglia | control | 0.27 \pm 0.02 | | | 6.23 \pm 0.33 | | |
| | 4.5 mg/kg | 0.26 \pm 0.02 | −4.63 | 0.62 | 5.89 \pm 0.33 | −5.53 | 0.46 |
| | 6 mg/kg | 0.19 \pm 0.02 | −30.43 | 0.00 | 4.6 \pm 0.36 | −26.19 | 0.00 |
| brainstem | control | 0.41 \pm 0.03 | | | 8.66 \pm 0.43 | | |
| | 4.5 mg/kg | 0.37 \pm 0.03 | −8.80 | 0.31 | 7.88 \pm 0.43 | −9.02 | 0.20 |
| | 6 mg/kg | 0.29 \pm 0.03 | −29.33 | 0.00 | 6.1 \pm 0.48 | −29.63 | 0.00 |
| cerebellum | control | 0.38 \pm 0.02 | | | 8.7 \pm 0.49 | | |
| | 4.5 mg/kg | 0.36 \pm 0.02 | −4.77 | 0.58 | 8.18 \pm 0.49 | −6.01 | 0.45 |
| | 6 mg/kg | 0.26 \pm 0.03 | −30.60 | 0.00 | 6.41 \pm 0.54 | −26.37 | 0.00 |
| corpus callosum | control | 0.26 \pm 0.02 | | | 7.25 \pm 0.34 | | |
| | 4.5 mg/kg | 0.23 \pm 0.02 | −8.00 | 0.43 | 7.04 \pm 0.34 | −2.80 | 0.68 |
| | 6 mg/kg | 0.17 \pm 0.02 | −34.47 | 0.00 | 5.65 \pm 0.37 | −22.08 | 0.00 |
| cortex | control | 0.27 \pm 0.02 | | | 8.57 \pm 0.41 | | |
| | 4.5 mg/kg | 0.24 \pm 0.02 | −9.65 | 0.29 | 8.07 \pm 0.41 | −5.89 | 0.38 |
| | 6 mg/kg | 0.17 \pm 0.02 | −36.39 | 0.00 | 6.76 \pm 0.45 | −21.19 | 0.00 |
| hippocampus | control | 0.27 \pm 0.02 | | | 6.8 \pm 0.35 | | |
| | 4.5 mg/kg | 0.26 \pm 0.02 | −1.33 | 0.90 | 6.5 \pm 0.35 | −4.48 | 0.54 |
| | 6 mg/kg | 0.18 \pm 0.02 | −33.77 | 0.00 | 5.24 \pm 0.38 | −22.94 | 0.00 |
| hypothalamus | control | 0.35 \pm 0.03 | | | 7.57 \pm 0.37 | | |
| | 4.5 mg/kg | 0.33 \pm 0.03 | −3.30 | 0.75 | 7.27 \pm 0.37 | −3.96 | 0.57 |
| | 6 mg/kg | 0.24 \pm 0.03 | −30.33 | 0.00 | 6.08 \pm 0.41 | −19.61 | 0.01 |
| medial geniculate | control | 0.32 \pm 0.03 | | | 6.06 \pm 0.29 | | |
| | 4.5 mg/kg | 0.3 \pm 0.03 | −5.13 | 0.72 | 5.85 \pm 0.29 | −3.59 | 0.59 |
| | 6 mg/kg | 0.2 \pm 0.04 | −35.96 | 0.02 | 4.65 \pm 0.32 | −23.34 | 0.00 |
| mesencephalic region | control | 0.35 \pm 0.02 | | | 7.14 \pm 0.33 | | |
| | 4.5 mg/kg | 0.34 \pm 0.02 | −1.86 | 0.83 | 6.65 \pm 0.33 | −6.75 | 0.31 |
| | 6 mg/kg | 0.26 \pm 0.02 | −26.91 | 0.00 | 5.49 \pm 0.37 | −23.00 | 0.00 |
| midbrain | control | 0.35 \pm 0.02 | | | 7.07 \pm 0.34 | | |
| | 4.5 mg/kg | 0.33 \pm 0.02 | −5.43 | 0.52 | 6.92 \pm 0.34 | −2.13 | 0.75 |
| | 6 mg/kg | 0.26 \pm 0.02 | −25.27 | 0.00 | 5.58 \pm 0.37 | −21.06 | 0.00 |
| septum | control | 0.27 \pm 0.02 | | | 6.37 \pm 0.35 | | |
| | 4.5 mg/kg | 0.26 \pm 0.02 | −5.95 | 0.60 | 5.91 \pm 0.35 | −7.23 | 0.35 |
| | 6 mg/kg | 0.17 \pm 0.02 | −38.90 | 0.00 | 4.89 \pm 0.38 | −23.28 | 0.00 |
| striatum | control | 0.26 \pm 0.02 | | | 6.36 \pm 0.32 | | |
| | 4.5 mg/kg | 0.24 \pm 0.02 | −9.46 | 0.33 | 6.05 \pm 0.32 | −5.00 | 0.48 |
| | 6 mg/kg | 0.17 \pm 0.02 | −35.30 | 0.00 | 4.95 \pm 0.35 | −22.18 | 0.00 |
| superior colliculus | control | 0.39 \pm 0.03 | | | 6.98 \pm 0.33 | | |
| | 4.5 mg/kg | 0.36 \pm 0.03 | −9.66 | 0.30 | 6.34 \pm 0.33 | −9.21 | 0.17 |
| | 6 mg/kg | 0.25 \pm 0.03 | −37.67 | 0.00 | 5.13 \pm 0.36 | −26.42 | 0.00 |
| thalamus | control | 0.3 \pm 0.02 | | | 6.19 \pm 0.31 | | |
| | 4.5 mg/kg | 0.29 \pm 0.02 | −3.23 | 0.76 | 5.89 \pm 0.31 | −4.82 | 0.49 |
| | 6 mg/kg | 0.2 \pm 0.03 | −33.61 | 0.00 | 4.54 \pm 0.34 | −26.70 | 0.00 |
| whole-brain | control | 0.3 \pm 0.02 | | | 7.82 \pm 0.38 | | |
| | 4.5 mg/kg | 0.28 \pm 0.02 | −7.18 | 0.42 | 7.35 \pm 0.38 | −5.99 | 0.38 |
| | 6 mg/kg | 0.2 \pm 0.02 | −33.68 | 0.00 | 5.92 \pm 0.41 | −24.29 | 0.00 |

reaching SUV values of 6 in blood and plasma. This value had been considered as an outlier and removed from the analysis.

Analysis of the whole-brain SUV–TACs did not find significant differences among the experimental groups. The SUV calculated using the last 20 min of the scan showed a reduction of 10% in the high-dose group compared to the control, however, this reduction in the $SUV_{40-60 \text{ min}}$ was not statistically significant. These results refute the use of SUV for

quantification of P-gp induction and support the use of pharmacokinetic parameters.

Following previous recommendations, the kinetics of [^{18}F]MC225 was analyzed using 1-TCM.²⁴ The statistical analysis found a significant decrease in the K_1 and V_T values in animals treated with a high-dose of MC111 compared to the control and low-dose groups, which indicates an increase in the P-gp function. The whole-brain region K_1 in the high-dose group decreased by 34% ($p < 0.001$) compared to the control group

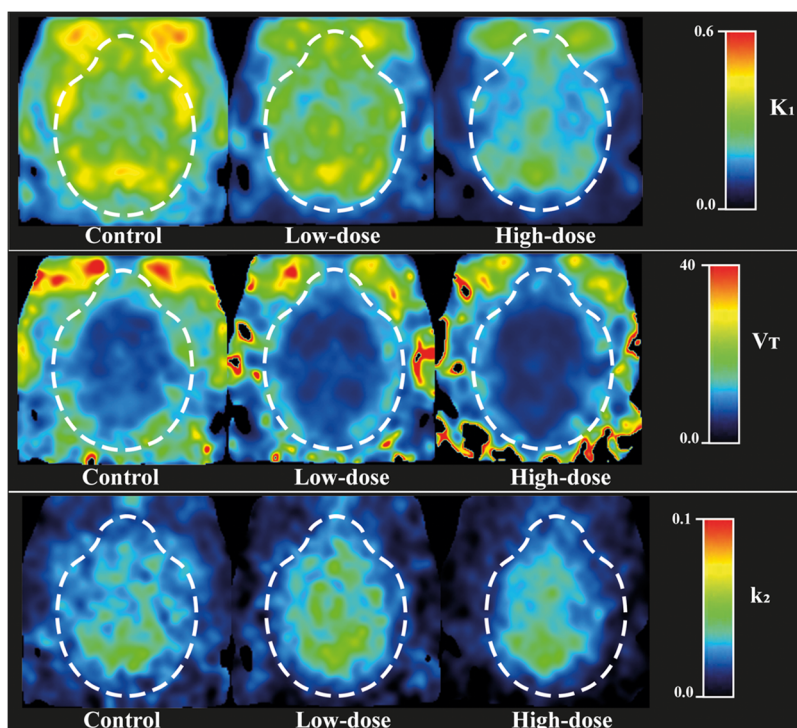


Figure 5. Parametric images calculated using 1-TCM and 60 min scan duration of three representative subjects of each group: control, low-dose (4.5 mg/kg), and high-dose (6 mg/kg).

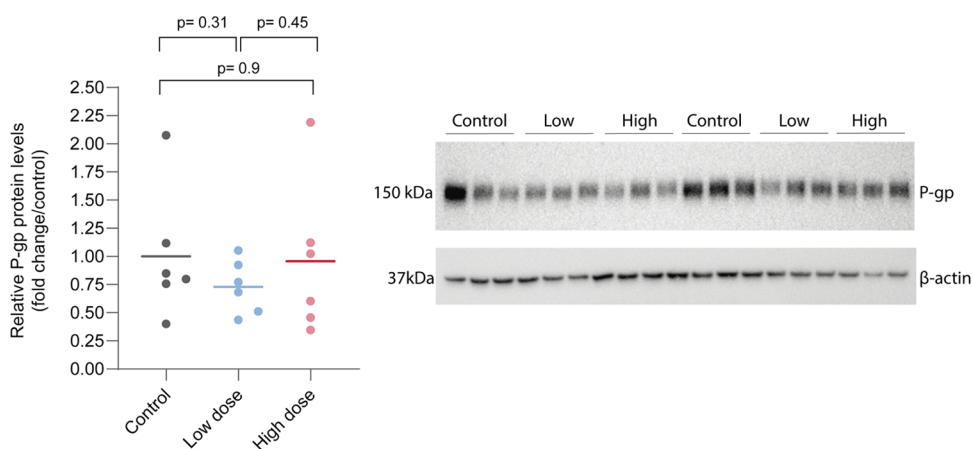


Figure 6. P-gp expression levels and western blot bands corresponding to P-gp (band close to 160 kDa predicted molecular weight) and β -actin as the load control (band close to 42 kDa predicted molecular weight).

and by 29% ($p = 0.005$) compared to the low-dose group. The V_T of the whole-brain region in the high-dose group decreased by 24% ($p = 0.001$) compared to the control and by 19% ($p = 0.011$) compared to the low-dose group. Similar changes were observed in all of the selected brain regions. As was also observed in previous studies, changes in K_1 are larger than the change in V_T . For this reason, K_1 has been identified as a more sensitive parameter to measure the P-gp function at the BBB in rats and in nonhuman primates,^{24,37} however, V_T can also be used as a surrogate parameter to estimate the P-gp function.²⁴

Many therapeutic drugs have shown an induction effect on the P-gp transporter.³⁸ Subchronic morphine administrations induced P-gp expression in brain microvessels of rats.³⁹ In the mentioned study, rats were treated with increasing doses of morphine by intraperitoneal injections (ip) twice a day for 5 consecutive days. Six hours after the last dose of morphine, the

levels of P-gp were not higher but after 24 h of abstinence, the P-gp expression increased 1.5 times compared to control animals.³⁹ Long-term exposure (60 days) to phenobarbital, phenytoin, carbamazepine, and valproic acid (antiepileptic drugs) increased the P-gp expression levels and P-gp function in rat brain microvascular endothelial cells *in vitro*.⁴⁰ P-gp expression in brain capillaries of mice was also increased 24 h after the administration of rifampin (50 mg/kg). That study also observed an increase of P-gp expression and function *in vitro* using isolated brain capillaries after the exposure to rifampicin and hyperforin, two PXR-ligands.⁴¹ After the treatment with the corticosteroid dexamethasone for 4 days at dose 25 mg/kg ip, brain capillaries of rats showed an increase in the P-gp expression.⁴² The previously described studies evaluated the P-gp function *in vitro* using endothelial cells acquired from brain postmortem. However, our study confirms the P-gp induction

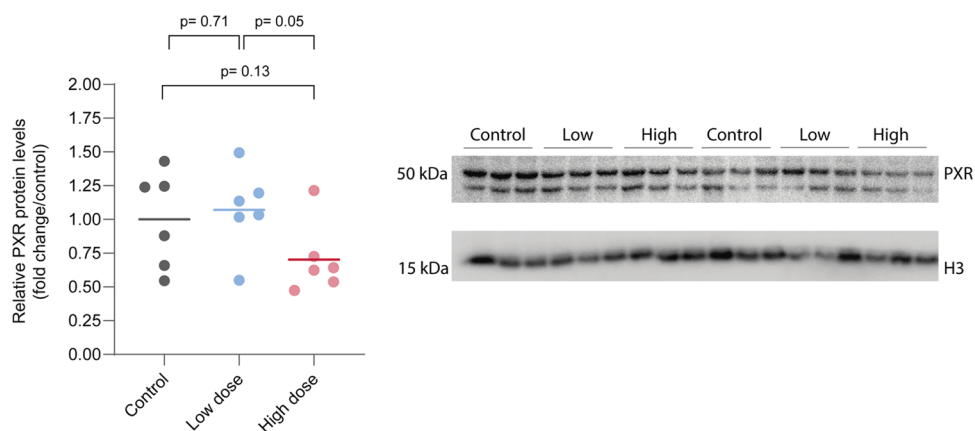


Figure 7. PXR expression levels and western blot bands corresponding to PXR (two bands close to 50 kDa predicted molecular weight) and histone H3 as the load control (band close to 17 kDa predicted molecular weight). Notice that in our analysis only the 50 kDa band was quantified.

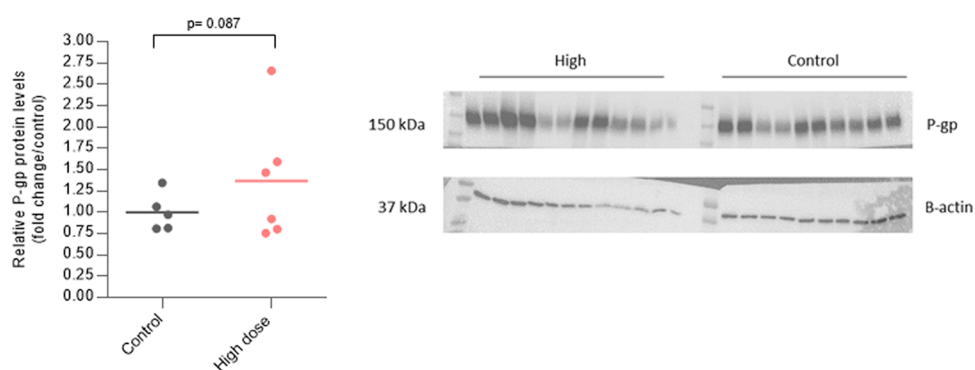


Figure 8. P-gp expression levels in isolated brain endothelial cells and western blot bands corresponding to P-gp (band close to 160 kDa predicted molecular weight) and β -actin as the load control (band close to 42 kDa predicted molecular weight).

effect of MC111 *in vivo* using PET imaging. This compound has been selected from an extensive P-gp/BCRP library and *in vitro* studies have already shown that MC111 increases the P-gp and BCRP function and expression.²⁰ Several studies have suggested a synergetic role between the two efflux transporters.¹⁰ Moreover, the expression of these two transporters is usually increased in various neurodegenerative diseases.²² Therefore, it is expected that the dual action of MC111 may be beneficial for the treatment of AD.

Since P-gp function can be altered after several treatments, it is important to monitor such alterations to prevent decreases in drug efficacy or drug toxicities. Moreover, since P-gp inducers have the ability to restore the P-gp function and thus can be a potential treatment for neurodegenerative diseases where the P-gp function is decreased, PET imaging with P-gp tracers may allow the evaluation of the effect of promising inducer drug candidates.

Tracers for imaging P-gp function such as [¹¹C]verapamil and [¹¹C]-*N*-desmethyl-loperamide show very low uptake inside the brain at baseline conditions due to their strong affinity toward the transporter protein, thus hampering the evaluation of P-gp induction.^{43–46} However, weak P-gp substrates such as [¹⁸F]MC225 show higher baseline brain uptake levels than [¹¹C]verapamil²⁴ and thus have been proposed as adequate tracers to measure P-gp induction. The results of our studies confirm the ability of [¹⁸F]MC225 to measure increases in the P-gp function because the treatment with the P-gp inducer MC111 significantly decreased the K_1 and V_T values of [¹⁸F]MC225 in

the high-dose group compared to the control group, indicating an increase in the P-gp function in these rats.

Increases in the P-gp function do not always mean increases in the P-gp expression, for this reason, it is important to assess simultaneously the P-gp expression level and function to identify the mechanism of action of a drug. For instance, western blot and flow cytometry analysis indicated that colchicine increased the P-gp expression in caco-2 cells without increasing the P-gp function.⁴⁷ In the present study, at first, the P-gp expression level was assessed postmortem using the whole-brain tissue of the rats, however, these results did not find any change in the P-gp expression among the groups. To adequately evaluate the P-gp expression level at the BBB of rats and to not overestimate the P-gp levels, the brain endothelial cells must be isolated because the P-gp transporter is also expressed at low levels by nonendothelial cell types and because the volume of the brain capillaries represents approximately 1% of the total brain volume.^{27,48,49} Thus, additional western blot experiments were performed to determine the P-gp expression levels in isolated brain endothelial cells of the rats. The preliminary results have shown a tendency to higher P-gp expression levels in rats treated with MC111 compared to control rats, however, this increase was not statistically significant. Additional data is needed to confirm these results and to examine the mechanism of action of this novel compound. Moreover, P-gp transporters are also expressed in intracellular compartments, whereas only the transporter molecules on the cell surface are functional.^{50,51} Therefore, it is important to use antibodies that specifically recognize an external epitope of the P-gp transporter to avoid

the detection of P-gp stored in intracellular vesicles.^{27,52,53} For this reason, more research is required to distinguish the expression level of the P-gp transporter at the membrane of the cell, which will be more related to the functionality of the transporter, from the intracellular P-gp. Our study also evaluated the PXR expression level in the whole-brain tissue since we have hypothesized that MC111 may increase the P-gp expression by the activation of the PXR. However, the analysis did not detect any increase in the expression levels of this nuclear receptor. Interestingly, in the PXR western blot, a band between 37 and 50 kDa was also found in addition to the expected PXR 50 kDa band. This could be due to the presence of alternative splicing variants. Isoforms 1b and 2b of 43 and 39 kDa, respectively, were reported in uniprot database but further analysis is needed to elucidate this issue.⁵⁴ Notice that in our analysis only the 50 kDa band quantification is shown. In future studies, the PXR expression levels will be also determined in isolated brain cells to find out if MC111 is also a PXR ligand.

CONCLUSIONS

In conclusion, this study shows the ability of MC111 to induce the P-gp expression and function at the BBB of rats. An increasing trend regarding the P-gp expression levels in isolated brain capillaries is found using western blot. Moreover, an increased P-gp function is confirmed with [¹⁸F]MC225 and PET. This study demonstrates that the administration of a 6 mg/kg dose of MC111 can be safely administered to rats to induce the P-gp expression and function at the BBB. Moreover, the results also verify the ability of [¹⁸F]MC225 to measure increases in the P-gp function of rats. [¹⁸F]MC225 may be the first radiofluorinated tracer able to monitor increases and decreases of the P-gp function at the mammalian BBB. Moreover, this technique may be used to screen potential P-gp inducers that can be applied as experimental treatments for several neurodegenerative diseases.

AUTHOR INFORMATION

Corresponding Author

Gert Luurtsema – Department of Nuclear Medicine and Molecular Imaging, University of Groningen, University Medical Center Groningen, 9713 GZ Groningen, The Netherlands; Email: g.luurtsema@umcg.nl

Authors

Lara García-Varela – Department of Nuclear Medicine and Molecular Imaging, University of Groningen, University Medical Center Groningen, 9713 GZ Groningen, The Netherlands; orcid.org/0000-0001-9803-4708

Manuel Rodríguez-Pérez – Clinical Neurosciences Research Laboratory, Health Research Institute of Santiago de Compostela (IDIS), 15706 Santiago de Compostela, Spain

Antía Custodia – Clinical Neurosciences Research Laboratory, Health Research Institute of Santiago de Compostela (IDIS), 15706 Santiago de Compostela, Spain

Rodrigo Moraga-Amaro – Department of Nuclear Medicine and Molecular Imaging, University of Groningen, University Medical Center Groningen, 9713 GZ Groningen, The Netherlands

Nicola A. Colabufo – Dipartimento di Farmacia-Scienze del Farmaco, Università degli Studi di Bari, I-70125 Bari, Italy; orcid.org/0000-0001-5639-7746

Pablo Aguiar – Department of Nuclear Medicine and Molecular Imaging Group, Clinical University Hospital, IDIS Health Research Institute, 15706 Santiago de Compostela, Spain

Tomás Sobrino – Clinical Neurosciences Research Laboratory, Health Research Institute of Santiago de Compostela (IDIS), 15706 Santiago de Compostela, Spain; orcid.org/0000-0002-9760-8690

Rudi A.J.O. Dierckx – Department of Nuclear Medicine and Molecular Imaging, University of Groningen, University Medical Center Groningen, 9713 GZ Groningen, The Netherlands

Aren van Waarde – Department of Nuclear Medicine and Molecular Imaging, University of Groningen, University Medical Center Groningen, 9713 GZ Groningen, The Netherlands

Philip H. Elsinga – Department of Nuclear Medicine and Molecular Imaging, University of Groningen, University Medical Center Groningen, 9713 GZ Groningen, The Netherlands

Complete contact information is available at:

<https://pubs.acs.org/10.1021/acs.molpharmaceut.1c00302>

Author Contributions

L.G.-V. performed the synthesis of the tracer and the preclinical studies, analyzed the data, conducted the pharmacokinetic analysis, and wrote the manuscript with help from all authors. M.R.-P. performed the western blot studies and commented on the final manuscript. A.C. helped in the isolation of the brain capillaries and the western blot studies. R.M.-A. provided assistance during the *in vivo* studies. N.A.C. contributed to the design of the experiment, provided the P-gp inducer (MC111), and commented on the final manuscript. P.A. and T.S. helped in the design of the western blot studies and commented on the final manuscript. A.v.W. contributed to the design of the experiment and commented on the final manuscript. RAJOD and PEH commented on the final manuscript. G.L. contributed to the design of the experiment, helped in the interpretation of the data, and in the preparation of the manuscript.

Notes

The authors declare no competing financial interest.

The data sets used and analyzed during the current study are available from the corresponding author on reasonable request.

ACKNOWLEDGMENTS

The authors would like to thank Jürgen W.A. Sijbesma and Anna Schildt for their assistance during the preclinical studies.

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