

Original Article

Radiosynthesis and preclinical evaluation of a carbon-11 labeled PET ligand for imaging metabotropic glutamate receptor 7

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Abstract: Metabotropic glutamate receptor 7 (mGlu₇) is a G protein-coupled receptor that is preferentially found in the active zone of neurotransmitter release in the central nervous system (CNS). mGlu₇ plays a vital role in memory, learning, and neuronal development, rendering it a potential target for treating epilepsy, depression, and anxiety. The development of noninvasive imaging ligands targeting mGlu₇ could help elucidate the functional significance of mGlu₇ and accelerate drug discovery for neurological and psychiatric disorders. In this report, a novel carbon-11 labeled positron emission tomography (PET) tracer designated [¹¹C]18 (codenamed MG7-2109) was synthesized via ¹¹C-methylation in 23% decay-corrected radiochemical yield (RCY). *In vitro* serum stability, serum protein binding, *in vitro* autoradiography and *ex vivo* biodistribution studies of [¹¹C]18 were conducted. Preliminary PET imaging results revealed a homogeneous distribution of [¹¹C]18 and rapid clearance in rodent brains. This study provides valuable insights into the development of mGlu₇-targeted PET tracer based on an isoxazolo(5,4-c)pyridine scaffold.

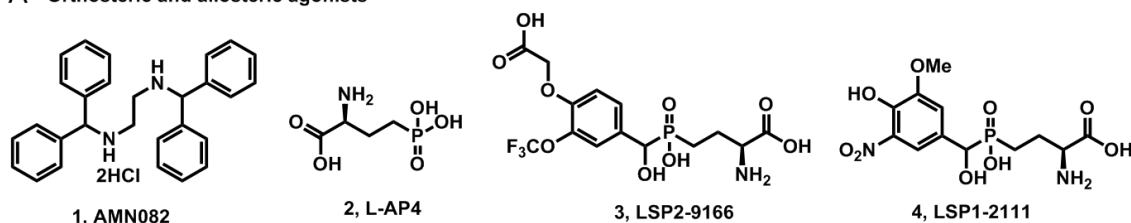
Keywords: mGlu₇, negative allosteric modulator, positron emission tomography, radioligand

Introduction

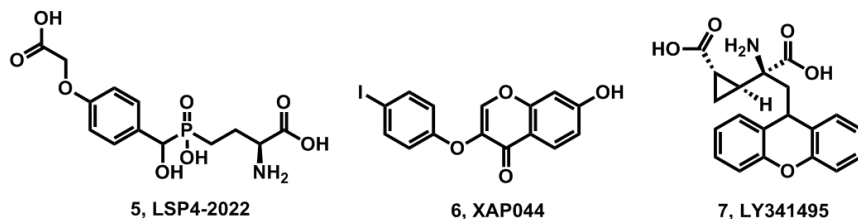
Glutamate is the primary excitatory neurotransmitter in the mammalian central nervous system (CNS), orchestrating physiological processes involved in memory formation, synaptic plasticity, and neurodevelopment. Depending on their structures and physiological functions, two classes of glutamate receptors are identified as metabotropic glutamate receptors (mGlu) and ionotropic glutamate receptors (iGlu) [1]. iGlu are ligand-gated ion channels that regulate the excitatory neurotransmission rapidly, whereas mGlu are G protein-coupled receptors modulating signal transduction cascades, including the second messengers, ion channels and other independent pathways [2, 3]. To date, eight mGlu subtypes are reported and further divided into three subgroups according to sequence homology, cell signaling transduction and pharmacology [4, 5]. Group I (mGlu₁ and mGlu₅) are mainly expressed postsynaptic to activate phospholipase C or adenylyl cyclase (ACs). In general, both group II (mGlu₂ and mGlu₃) and III (mGlu₄ and mGlu_{6,8}) are located pre- and postsynaptic and have similar mechanism of action, which inhibit ACs and regulate ion channels [1, 6].

Among group III, mGlu₇ is widely distributed in the CNS. The striatum, hippocampus, thalamus and neocortex are the most abundant regions with mGlu₇ expression. As auto- or hetero-receptors, mGlu₇ activation results in an attenuated release of the endogenous neurotransmitters, glutamate or gamma-aminobutyric acid (GABA) in the pre-synaptic regions of glutamatergic or GABAergic terminals, respectively [3, 5, 7, 8]. Several reports uncovered that mGlu₇ plays an important role in the processes of learning and memory [9-11]. Of note, dysregulation of mGlu₇ signaling can be observed under some neuropathological conditions, such as Alzheimer's disease (AD) [12], Parkinson's disease (PD) [13], Huntington's disease (HD) [14] and Rett syndrome [15]. Thus, mGlu₇ has been studied as a potential therapeutic target in several animal models of neurodegenerative diseases [16-20]. In addition to endogenous agonists like glutamate, several exogenous mGlu₇ agonists and antagonists have been found and used in mGlu₇ studies (Figure 1A, 1B). In recent years, several allosteric mGlu₇ modulators, both positive and negative, have attracted more attention and some progress has been made owing to their high selectivity and reduced side effects (shown in Figure 1C, 1D) [6, 7, 21,

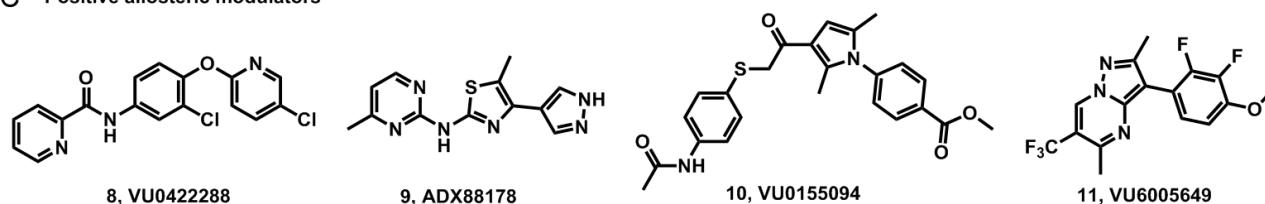
A Orthosteric and allosteric agonists



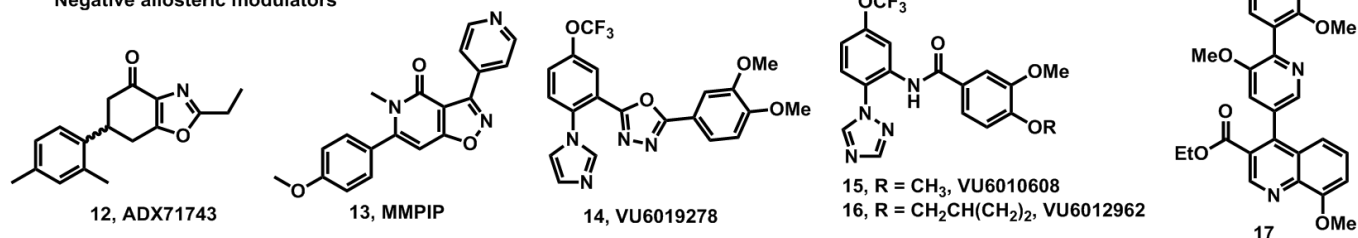
B Antagonists



C Positive allosteric modulators



D Negative allosteric modulators

Figure 1. Representative mGlu₇ modulators.

22]. However, these molecules have not been used as clinical drugs for reasons including low subtype-selectivity, poor bioavailability and inability to cross the blood-brain barrier (BBB).

Positron emission tomography (PET) is a highly sensitive imaging technology for pre-clinical and clinical functional molecular imaging. After the appropriate radiolabeled ligand is injected in a non-pharmacological dose, three-dimensional images, including the concentration and location information of the radioligand, could be obtained to reveal the physiological state of the target through non-invasive data collection and reconstruction [23-25]. Although the development of mGlu₇-selective PET tracers will contribute to the mechanism study of mGlu₇-associated disease and drug discovery, only one PET tracer derived from mGlu₇ negative allosteric modulator (NAM), [¹¹C]MMPIP was developed and evaluated in rodents. Due to its moderate affinity to mGlu₇ (26 nM) [26] and the existing radioactive metabolites in the brain, [¹¹C]MMPIP cannot be used for the quantitative assessment of mGlu₇,

[27]. Therefore, compound 18 (MG7-2109) (Figure 2) [26], with improved affinity to mGlu₇ (reported IC₅₀ = 12 nM) was selected as a candidate compound in this work. Radiosynthesis of [¹¹C]18 followed by *in vitro* autoradiography, *ex vivo* biodistribution and PET studies on rodent brains were conducted to assess its performance for imaging mGlu₇ in the brain.

Materials and methods

General

Unless otherwise stated, reagents, solvents, and chemicals were purchased from commercially available vendors and used without further purification. The chemical synthesis of compound 18 (MG7-2109) and corresponding precursor 19 were synthesized according to the previous report [26]. Nuclear magnetic resonance (NMR) spectra were recorded in an AVANCE NEO 400MHZ spectrometer. ¹H NMR chemical shifts (δ) are reported in parts per million (ppm) relative to TMS with the residual solvent peak

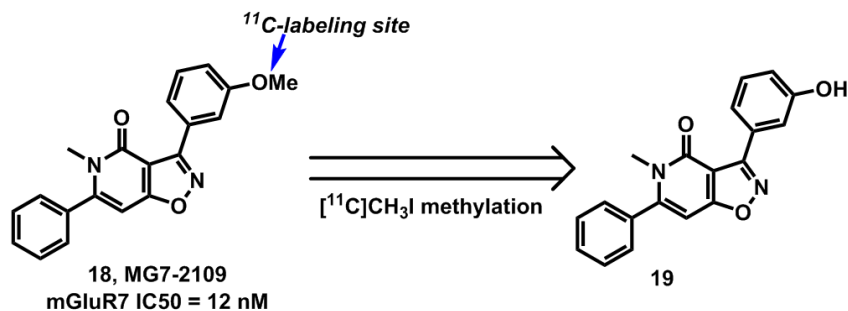


Figure 2. Proposed mGlu₇-selective PET radioligand and the responding precursor.

used as an internal reference. Abbreviations used are singlet (s), doublet (d), triplet (t), quartet (q), and multiplet (m). High-performance liquid chromatography-mass spectrometry (HPLC-MS) was performed on Shimadzu LCMS-2010EV single four-stage rod mass spectrometer ESI electric spray ion source or Agilent single four-stage rod 6120B ESI electric spray ion source.

Chemistry

(E)-3-methoxybenzaldehyde oxime **21**: A solution of 3-methoxybenzaldehyde **20** (5.0 g, 36.8 mmol) and TEA (5.6 g, 55.1 mmol) and hydroxylamine hydrochloride (3.8 g, 55.1 mmol) in EtOH/H₂O (50 mL/50 mL) was stirred at room temperature overnight. Then the reaction was extracted with water and EtOAc. The organic phase was washed brine, then dried on Na₂SO₄ and concentrated in vacuum. The residue was purified by silica gel column (petroleum ether/EtOAc = 8/1) to give *(E)*-3-methoxybenzaldehyde oxime **21** (6.0 g, yield 90%) as a colorless oil.

(Z)-*N*-hydroxy-3-methoxybenzimidoyl chloride **22**: To a solution of *(E)*-3-methoxybenzaldehyde oxime **21** (3.7 g, 24.5 mmol) in DMF (50 mL) was added NCS (2.9 g, 22 mmol) in batches at 0°C. Then the solution was stirred at 0°C to room temperature overnight. The solution was extracted with water and EtOAc. The EtOAc phase was washed with brine, dried over Na₂SO₄ and concentrated in vacuum to give *(Z)*-*N*-hydroxy-3-methoxybenzimidoyl chloride **22** (3.7 g, yield 82%) as a colorless oil without further purification.

(Z)-3-(pyrrolidin-1-yl)but-2-enoate **24**: A solution of pyrrolidine (2.5 g, 35.0 mmol) and ethyl 3-oxobutanoate **23** (4.7 g, 35.5 mmol) in toluene (25 mL) was stirred at 115°C for 2 h with Dean-stark. Then the solution was concentrated in vacuum to give ethyl *(Z)*-3-(pyrrolidin-1-yl)but-2-enoate **24** (6.0 g, yield 71%) as a yellow oil for next step without further purification.

ethyl 3-(3-methoxyphenyl)-5-methylisoxazole-4-carboxylate **25**: To a solution of ethyl *(Z)*-3-(pyrrolidin-1-yl)but-2-enoate **24** (6.0 g, 33.0 mmol) and TEA (6.0 g, 60.0 mmol) in DCM (40 mL) was added dropwise *(Z)*-*N*-hydroxy-3-methoxybenzimidoyl chloride **22** (3.7 g, 20.0 mmol) in

DCM (40 mL) at 0°C. Then the solution was stirred at room temperature overnight. The solution was diluted with DCM, washed with 1*N* HCl and brine successively, dried over Na₂SO₄ and concentrated in vacuum. The residue was purified by silica gel column (petroleum ether/EtOAc = 2/1) to give 1.8 g crude and then purified by reversed-phase column (MeOH = 0-95%) to give ethyl 3-(3-methoxyphenyl)-5-methylisoxazole-4-carboxylate **25** (350.0 mg, yield 6%) as a yellow oil. ¹H NMR (400 MHz, DMSO-*d*₆)

δ 7.42 - 7.38 (m, 1H), 7.17 - 7.15 (m, 2H), 7.11 - 7.08 (m, 1H), 4.18 (q, *J* = 8.0 Hz, 2H), 3.80 (s, 1H), 2.71 (s, 3H), 1.16 (t, *J* = 8.0 Hz, 3H).

3-(3-methoxyphenyl)-*N*,5-dimethylisoxazole-4-carboxamide **26**: A solution of ethyl 3-(3-methoxyphenyl)-5-methylisoxazole-4-carboxylate **25** (550.0 mg, 2.1 mmol) and methylamine/MeOH (10 mL) in sealed tube was stirred at 85°C overnight. The solution was concentrated in vacuum to give 3-(3-methoxyphenyl)-*N*,5-dimethylisoxazole-4-carboxamide **26** (500.0 mg, yield 96%) as a light-yellow solid. ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.31 (brs, 1H), 7.43 - 7.39 (m, 1H), 7.25 - 7.23 (m, 2H), 7.09 - 7.07 (m, 1H), 3.80 (s, 3H), 2.74 (s, 3H), 2.53 (s, 3H).

3-(3-methoxyphenyl)-*N*-methyl-5-(2-oxo-2-phenylethyl)isoxazole-4-carboxamide **27**: To a solution of 3-(3-methoxyphenyl)-*N*,5-dimethylisoxazole-4-carboxamide **26** (250.0 mg, 1.0 mmol) in dry THF (40 mL) *n*-BuLi (1.0 mL, 2.5 M/L, 2.2 mmol) was added dropwise at -70°C under N₂. Then the solution was stirred at -70°C for 1.5 h. Then methyl benzoate **7** (305.0 mg, 2.2 mmol) in dry THF (1 mL) was added dropwise at -65°C and the solution was stirred at this temperature for 1.5 h. Then the solution was stirred at room temperature. The solution was poured into ice water and extracted with water and EtOAc. The EtOAc phase was washed with brine, dried over Na₂SO₄ and concentrated in vacuum. The residue was purified by flash (petroleum ether/EtOAc = 2/1) to give 3-(3-methoxyphenyl)-*N*-methyl-5-(2-oxo-2-phenylethyl)isoxazole-4-carboxamide **27** (160 mg, yield 45%) as a light yellow oil.

3-(3-methoxyphenyl)-5-methyl-6-phenylisoxazolo[4,5-*c*]pyridin-4(5*H*)-one (**18**): To a solution of 3-(3-methoxyphenyl)-*N*-methyl-5-(2-oxo-2-phenylethyl)isoxazole-4-carboxamide **27** (160.0 mg, 0.4 mmol) in toluene (8 mL), *p*-Toluenesulfonic acid monohydrate (86.0 mg, 0.4 mmol) was added, then the solution was stirred at 110°C overnight. The solution was diluted with EtOAc. The EtOAc phase was washed with saturated Na₂CO₃ and brine, dried over Na₂SO₄ and concentrated in vacuum to give 3-(3-methoxyphenyl)-5-methyl-6-phenylisoxazolo[4,5-*c*]pyridin-4(5*H*)-one **18** (120.0 mg, yield 79%) as a brown solid. ¹H NMR (400 MHz, Chloroform-*d*₃) δ 7.99 - 7.91 (m,

2H), 7.55 - 7.50 (m, 3H), 7.45 - 7.38 (m, 3H), 7.06 (ddd, $J = 8.3, 2.7, 1.0$ Hz, 1H), 6.54 (s, 1H), 3.91 (s, 3H), 3.42 (s, 3H). Purity: 97%. HPLC-MS, $m/z = 333.0$ ($M + 1$).

3-(3-hydroxyphenyl)-5-methyl-6-phenylisoxazolo[4,5-c]pyridin-4(5H)-one (19): To a solution of 3-(3-methoxyphenyl)-5-methyl-6-phenylisoxazolo[4,5-c]pyridin-4(5H)-one 18 (100.0 mg, 0.3 mmol) in dry DCM (20 mL) was added BBr_3 (0.3 mL) at 0°C, then the solution was stirred at 0°C for 20 min and then warmed to room temperature stirred for 7 h. The solution was diluted with DCM and poured into ice water. The DCM phase was washed with brine, dried over Na_2SO_4 and concentrated in vacuum. The residue was purified by reversed-phase column chromatography (MeOH = 0-95%) to give 3-(3-hydroxyphenyl)-5-methyl-6-phenylisoxazolo[4,5-c]pyridin-4(5H)-one 19 (40.0 mg, yield 42%) as a white solid. ^1H NMR (400 MHz, Chloroform- d_3) δ 7.95 - 7.89 (m, 1H), 7.79 (dt, $J = 7.9, 1.2$ Hz, 1H), 7.52 (dt, $J = 4.4, 2.9$ Hz, 3H), 7.44 - 7.32 (m, 3H), 7.00 (ddd, $J = 8.2, 2.7, 1.0$ Hz, 1H), 6.57 (s, 1H), 3.40 (s, 3H). Purity: 97%. HPLC-MS, $m/z = 319.1$ ($M + 1$).

Pharmacology

Cell line generation: In order to generate human mGlu₇ stable cell lines to be used for thallium flux assays, human mGlu_{7A} was prepared by PCR amplification (from plasmid GRM70000001 from the cDNA Resource Center) of the entire coding sequence of each receptor and cloning into pIRES puro 3 (Invitrogen). The cloning sites were NheI/NotI. HEK GIRK cells, generously provided by Lily Jan (University of California San Francisco, San Francisco, CA), were transfected with 24 μg of DNA using Fugene6 (Promega), stable transfectants were selected with 1000 ng/mL puromycin dihydrochloride (Sigma-Aldrich, St. Louis, MO), and polyclonal human mGlu₇ GIRK cell lines were established. Cells were maintained following selection in 45% DMEM, 45% Ham's F12, 10% FBS, 100 units/mL penicillin/streptomycin, 20 mM HEPES, pH 7.3, 1 mM sodium pyruvate, 2 mM glutamine, 700 $\mu\text{g}/\text{mL}$ G418 (Mediatech, Inc., Herndon, VA), and 600 $\mu\text{g}/\text{mL}$ puromycin (growth media) at 37°C in the presence of 5% CO_2 . All cell culture reagents were purchased from Invitrogen Corp. (Carlsbad, CA) unless otherwise noted.

Human mGlu₇ thallium flux in vitro assays: Compound activity at mGlu₇ was assessed using thallium flux through GIRK channels, a method that has been previously described in detail [28]. For Schild analysis experiments a full concentration-response of L-AP4 was utilized in the absence and presence of fixed concentrations of MG7-2109. For potency evaluation experiments an EC_{80} concentration of L-AP4 was utilized. Data were analyzed using Excel (Microsoft Corp, Redmond, WA). The slope of the fluorescence increase beginning 5 s after thallium/agonist addition and ending 15 s after thallium/agonist addition was calculated, corrected to vehicle and maximal L-AP4 control slope values, and plotted using either XLfit (ID Business Solutions Ltd.) or Prism software (GraphPad

Software, San Diego, CA) to generate concentration-response curves. Potencies were calculated from fits using a four-point parameter logistic equation.

Radiochemistry

$[^{11}\text{C}]\text{CH}_3\text{I}$ was prepared from $[^{11}\text{C}]\text{CO}_2$ in cyclotron and then transferred into a precooled reaction vial (-15 to -20°C) containing the precursor 19 (0.95 mg), NaOH (0.5 M, 6.3 μL), and DMF (300 μL). After the transfer, this reaction mixture was heated at 80°C for 5 min. Then HPLC buffer ($\text{CH}_3\text{CN}/\text{H}_2\text{O} = 70/30$, 0.1% Et_3N , 0.5 mL) was added and then injected into HPLC for purification (column: CAPCELL PAK UG80 C18 column, 10 mm ID \times 250 mm; buffer: $\text{CH}_3\text{CN}/\text{H}_2\text{O}$ (70/30, 0.1% Et_3N); flow rate: 5.0 mL/min). $[^{11}\text{C}]\text{18}$ was obtained in $23 \pm 4\%$ radiochemical yield (decay-corrected, $n = 5$) at the end of synthesis (EOS) based on $[^{11}\text{C}]\text{CO}_2$. $[^{11}\text{C}]\text{18}$ was generated with excellent radiochemical purity (> 99%) and molar activity (201 ± 111 GBq/ μmol , $n = 5$).

In vitro stability and protein binding

In vitro serum stability: 400 μL of mouse, rat, non-human primate, and human serum were preincubated at 37°C for 5 min. Subsequently, $[^{11}\text{C}]\text{18}$ (approximately 1.5 mCi) was added to the samples and incubated at 37°C for 30 min, followed by the addition of 400 μL ice-cold MeCN to terminate the enzymatic reactions, as well as vortexing and centrifuging at 10,000 \times g for 5 min. The percentage of intact parent and radiometabolites was assessed by radio-HPLC using buffer $\text{CH}_3\text{CN}/\text{H}_2\text{O}$ (70/30, 0.1% Et_3N). As a negative control, the same experiment is carried out with PBS instead of serum.

Plasma protein binding: The plasma protein binding was conducted according to a previous report [29]. Briefly, 1.5 mCi of $[^{11}\text{C}]\text{18}$ was added per 150 μL of plasma that has been preincubated under 37°C for 5 min. The samples were incubated at room temperature for 10 min. To each 150 μL of radiotracer-plasma solution was added 300 μL of ice-cold PBS and all samples were briefly vortexed. 300 μL of each sample was added into the filters with a size cutoff of 10 kDa, and the samples were centrifuged at 21000 g for 15 min at 4°C. The filters were then washed with 300 μL PBS at 21000 g for 20 min at 4°C; 300 μL cold PBS was used to wash the tube and collect all the filtrates to obtain the protein fraction. The radioactivity of the protein fraction and filtrate was measured in a gamma counter (Wizard, PerkinElmer), and the free fraction f_u was calculated according to the following equation:

$$f_u = 1 - (A_{\text{protein}}/A_{\text{total}})$$

In vitro autoradiography

In brief, the rat brain sections (20 μm) were preincubated with Tris-HCl buffer (50 mM) at room temperature for 20 min, followed by incubation with $[^{11}\text{C}]\text{18}$ (1 $\mu\text{Ci}/\text{mL}$) for 30 min. For blocking studies, compound 18 (1 μM) or MMPIP

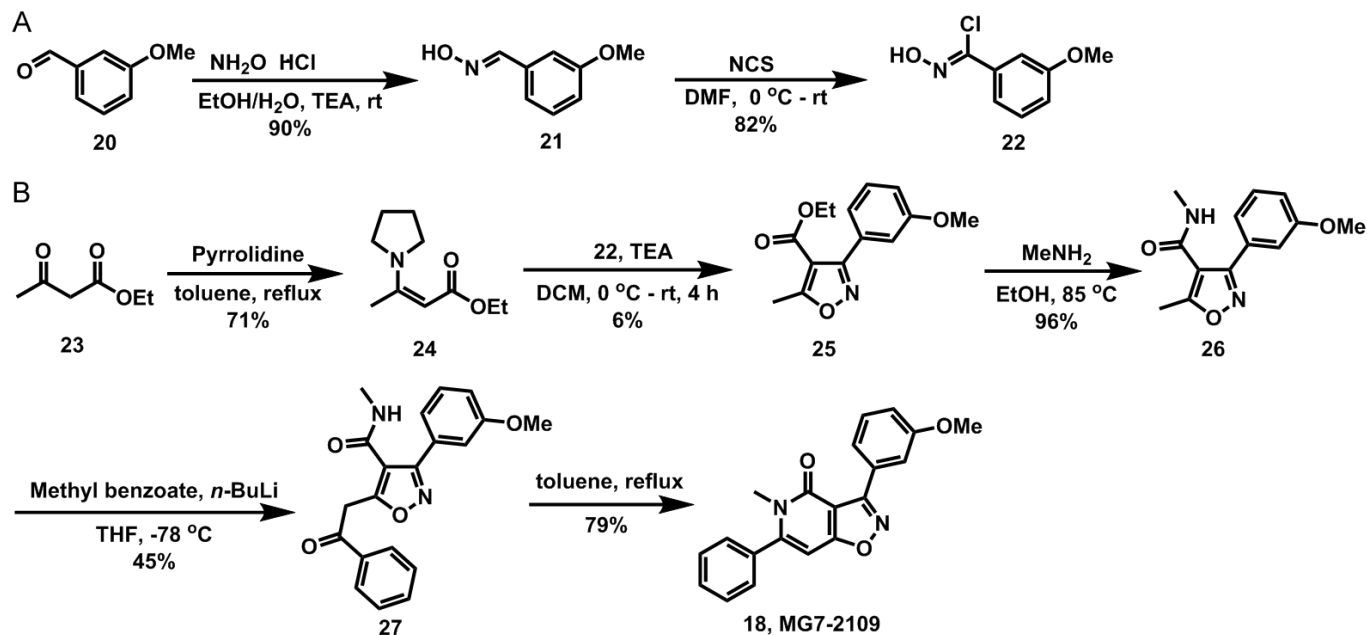


Figure 3. Chemical synthesis of 18 (MG7-2109).

(1 μM) was added to the incubation solution. Then the brain sections were washed with an ice-cold thrice for 2 min and dipped in cold distilled water for 10 s. Subsequently, the brain slices were dried and positioned on imaging screens (BAS-MS2025, GE Healthcare, USA). Then the screen was scanned by an Amersham Typhoon 5 analyzer system. The radioactivity values were normalized as a percentage of the radioactivity relative to the control.

PET imaging

An Inveon PET scanner (Siemens) was used for dynamic PET imaging studies and [^{14}C]18 (ca. 1.1 mCi/0.2 mL) was intravenously injected via the tail vein via a pre-installed catheter, and dynamic PET images were acquired in a 3D mode for a period of 60 min. The dynamic PET images were reconstructed using ASIPro VW software, and volumes of interest were calculated.

Whole-body distribution

In brief, [^{14}C]18 (50 $\mu\text{Ci}/0.1 \text{ mL}$) was injected intravenously into each mouse via the tail vein. At the time point of 5, 15, 30 and 60 min, mice were sacrificed by cervical dislocation and the tissues of interest were collected and weighted. An automatic gamma counter (PerkinElmer, USA) was used for determining each organ's decay-corrected radioactivity.

Results

Chemistry

The synthesis of 18 (MG7-2109) was started from the formation of oxime **21** from 3-methoxybenzaldehyde **20**, fol-

lowed by chlorination of obtained **21** to afford intermediate **22** in 82% yield (Figure 3A). On the other hand, condensation of ethyl acetoacetate **23** with pyrrolidine afforded enamine **24** in 71% yield. Cycloaddition of oxime **22** and enamine **24** delivered isoxazole **25** in 6% yield. Ammonolysis of **25** with methylamine provided amide **26** in 96% yield, which reacted with methyl benzoate to give intermediate **27** in the presence of excess *n*-BuLi at -78 °C. Cyclization of ketone **27** in boiling toluene afforded the "cold" isoxazolopyridone derivative **18** (MG7-2109) in 79% yield (Figure 3B).

Pharmacology

As shown in Figure 4A, MG7-2109 (30 μM , 10 μM , 3333 nM, 1111 nM, 370 nM, 123 nM, or 41 nM), dose-dependently decreased the maximal response of L-AP4 (a group III-selective mGlu agonist) toward mGlu₇ in the G protein-coupled inwardly rectifying potassium (GIRK) thallium assay, consistent with a noncompetitive mechanism of action. The potencies (IC_{50} s) for inhibiting an EC_{80} L-AP4 response in the mGlu₇ GIRK assay for MG7-2109 and MMPIP were evaluated using the mGlu₇ NAM XAPO44 as a positive control. The results revealed MG7-2109 and MMPIP exhibit NAM activity with IC_{50} values of 81 nM and 223 nM, respectively, while the positive control XAPO44 displayed a potency of 162 nM in the human mGlu₇ GIRK assay (Figure 4B). Pharmacokinetic parameters of MG7-2109 in Sprague Dawley (SD) rats appeared rapid metabolic clearance in blood and brain after a single intravenous administration (Figure 4C). The off-target screening of MG7-2109 to numerous classical targets in CNS including receptors, transporters and ion channels was conducted. No significant off-target binding of MG7-2109 was observed towards 59 CNS targets (< 50% inhibition) at

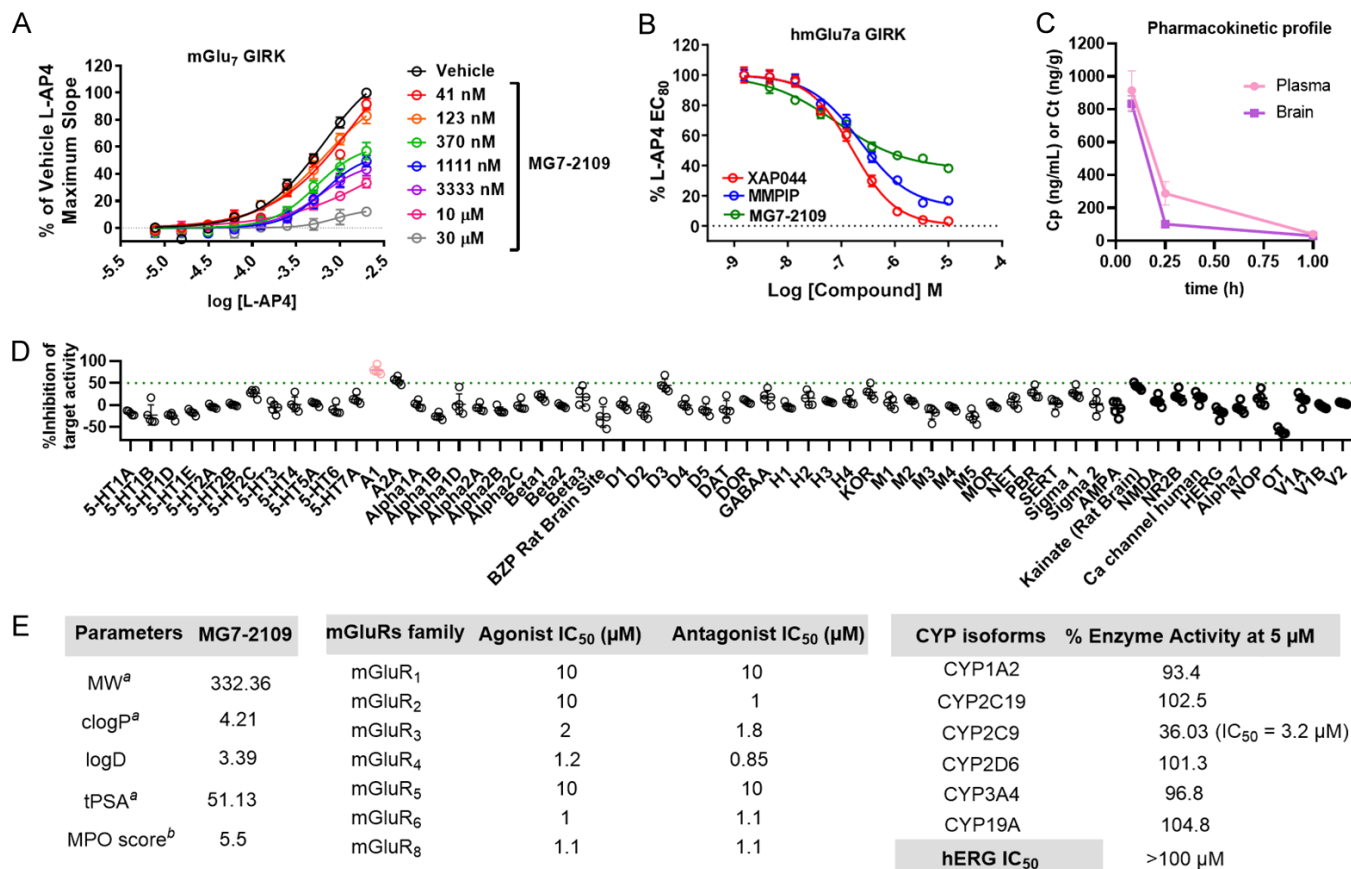


Figure 4. A. MG7-2109 noncompetitively decreases the maximal L-AP4 response in mGlu₇ GIRK thallium flux assays. B. *In vitro* evaluation of the potencies of MG7-2109, MMPIP, and the control mGlu₇ antagonist XAP044 in mGlu₇ GIRK thallium flux assays. C. Plasma and brain concentration-time profiles of MG7-2109 following a single intravenous administration to SD rats (male, dose: 1 mg/kg). D. Off-target pharmacological evaluation of MG7-2109 against major CNS targets at a concentration of 10 μM. E. Physicochemical properties and inhibitory activity of MG7-2109. ^aValues were calculated with ChemDraw 21.0 software. ^bValues were predicted with ACD/labs.

the concentration of 10 μM, except A1 (Ki = 644 nM) (Figure 4D). As summarized in Figure 4E, the cLogP value of MG7-2109 was predicted as 4.21, and the LogD value was determined to be 3.39 through the ‘shake-flask’ method. The topological polar surface area (tPSA) and multiparameter optimization (MPO) score of MG7-2109 were assessed as 51.13 and 5.5, respectively. Additionally, the agonists or antagonists IC₅₀ values of MG7-2109 against other members of the mGlu family were measured in the range of 0.85 μM to 10 μM. The inhibition of cytochromes P450 (CYP) enzymes was assessed by using 5 μM of MG7-2109. Weak inhibition (IC₅₀ > 5 μM) was observed for 1A2, 2C19, 2D6, 3A4 and 19A isoforms, while the IC₅₀ of 2C9 isoform was measured as 3.2 μM. Meanwhile, the hERG IC₅₀ of MG7-2109 was determined to be > 100 μM.

Radiochemistry

As shown in Figure 5, demethylation of 18 using boron tribromide gave the desired phenol precursor 19 in 42% yield. The deprotonation of the hydroxyl group of precursor 19 with NaOH promoted the carbon-11 methylation reaction at 80 °C for 5 min. The desired radiotracer [¹¹C]18

was obtained in 23% decay-corrected radiochemical yield (RCY) with more than 99% radiochemical purity at the end of synthesis (EOS).

In vitro formulation stability and protein binding

In vitro stability studies indicated that [¹¹C]18 has excellent stability after incubation with different species of serum (mice, rat, NHP and human) at 37 °C for 30 min (Figure 6A). In addition, [¹¹C] MG7-2109 demonstrated lower protein binding in mice, rat and NHP plasma (> 40% free fraction), while higher protein binding of [¹¹C]MG7-2109 in human serum was observed (Figure 6B).

In vitro autoradiography studies

In vitro autoradiography studies of [¹¹C]18 were performed on the rat brain sections under baseline and blocking conditions. Unfortunately, the ARG results revealed nearly homogeneous radiotracer distribution under baseline conditions, and no obvious blocking effects were detected by using cold compound 18 (1 μM, self-blocking) and MMPIP (1 μM) as blocking reagents (Figure 7).

mGlu₇ radiotracer

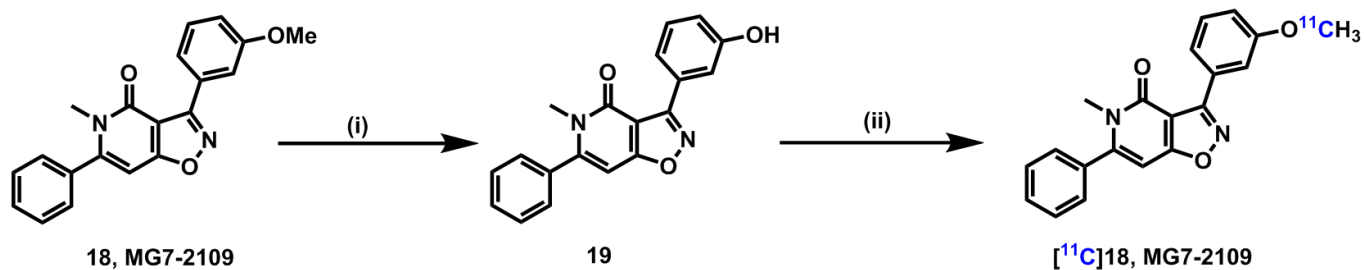


Figure 5. i. BBr₃, DCM, 0 °C, 2 h, 42%; ii. [¹¹C]CH₃I, 0.5 M NaOH, DMF, 80 °C, 5 min, 23% decay-corrected RCY for [¹¹C]18.

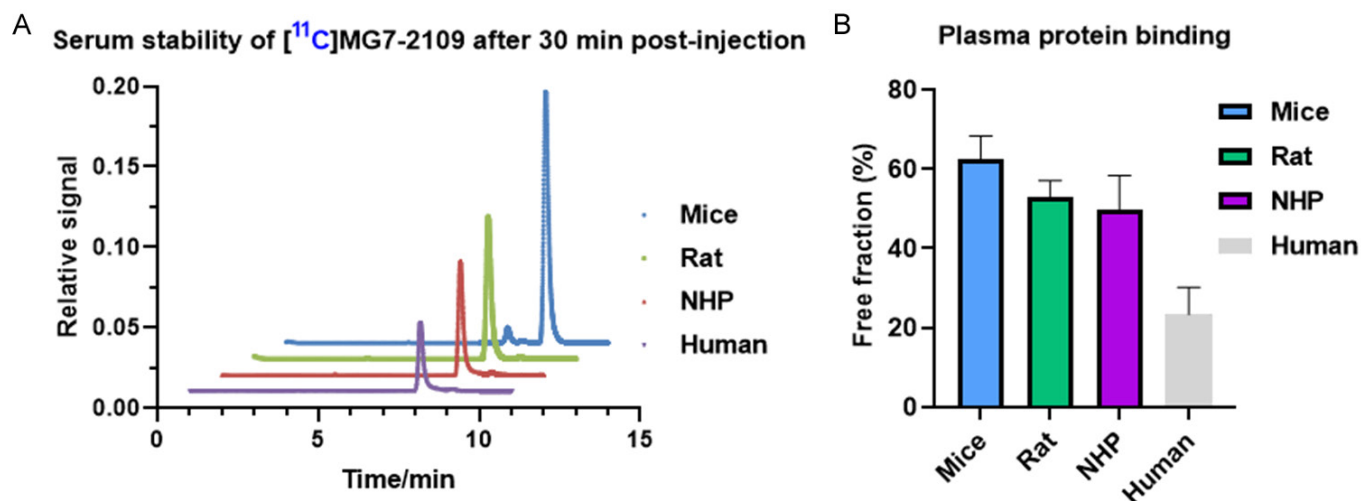


Figure 6. A. Formulation stability of [¹¹C]MG7-2109 in serum of mice, rats, NHPs, and humans at 30 min; B. Free fraction of [¹¹C]MG7-2109 in plasma, n = 3. NHP = non-human primates.

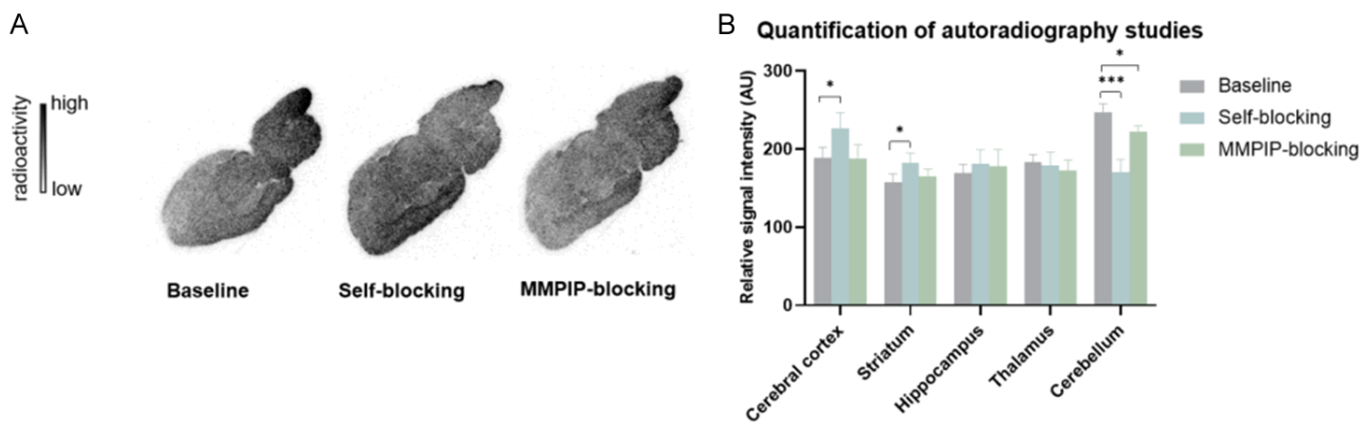


Figure 7. *In vitro* autoradiography of [¹¹C]18. A. Representative images for baseline and blocking (1 μM) autoradiography studies with [¹¹C]18. B. Quantification of autoradiography studies with [¹¹C]18. All data were referred to as mean ± SD, n ≥ 3. Asterisks were used to indicate statistical significance: *P ≤ 0.05 and ***P ≤ 0.001.

In vivo PET imaging study

The *in vivo* PET imaging study in rat brains was conducted and the results are illustrated in **Figure 8**. Consistent with the ARG results, a homogeneous uptake of [¹¹C]18 was observed followed by rapid clearance in rat brains.

Whole-body biodistribution study

The biodistribution study was performed in mice and the radioactivity levels of [¹¹C]18 in major organs were measured at 5-, 15-, 30- and 60-min post-tracer administration (**Figure 9**). The results showed high uptake (%ID/g >

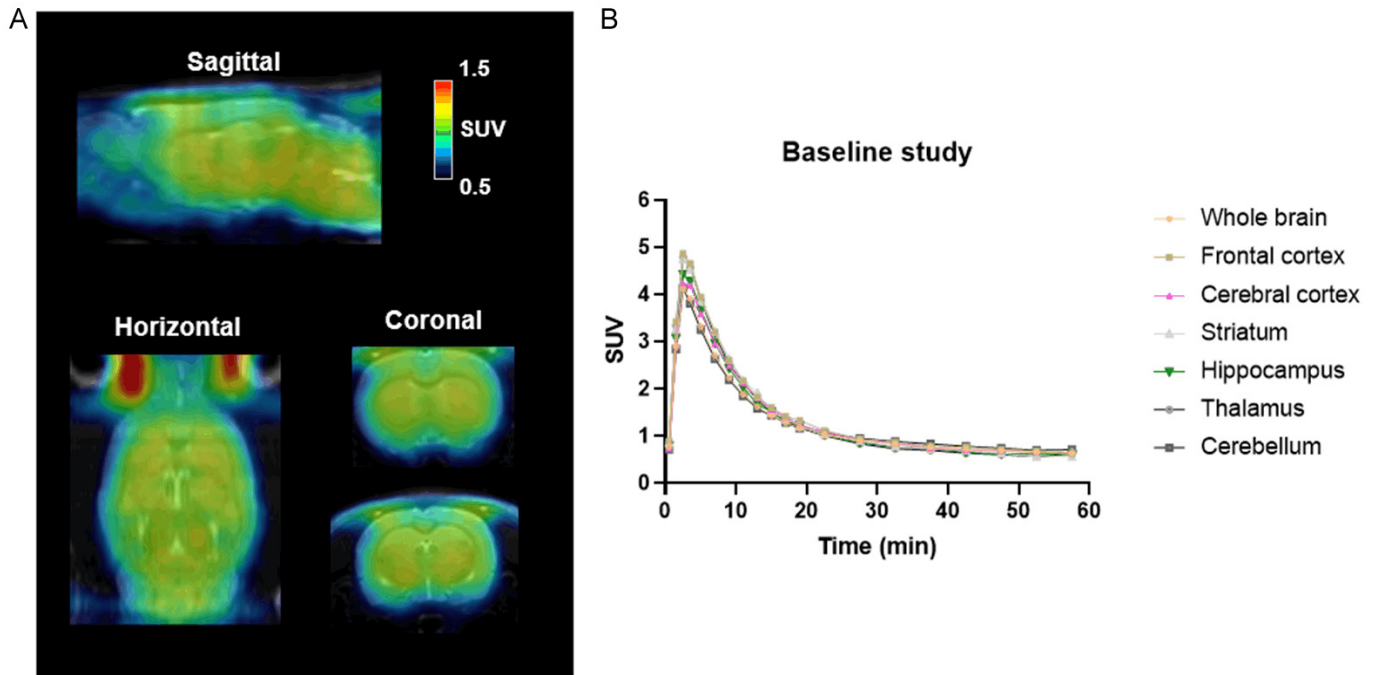


Figure 8. A. Summed PET images (5-20 min) [¹⁴C]18 in rat brains under baseline conditions. B. TACs of [¹⁴C]18 in various brain regions of interest.

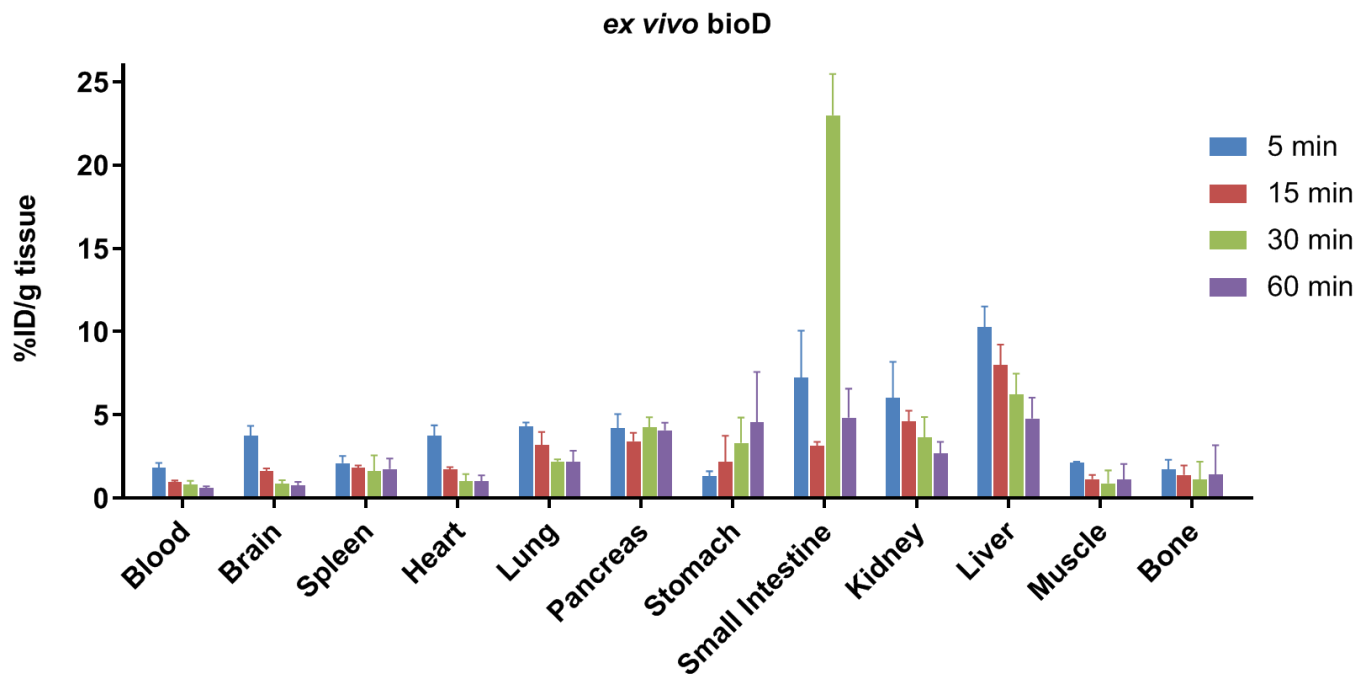


Figure 9. Whole-body ex vivo biodistribution studies of [¹⁴C]18 in CD-1 mice. All data are referred to as mean ± SD, n ≥ 3.

3) in the brain, heart and lung at 5 min, followed by gradual clearance in 60 min post tracer administration. The high tracer uptake in small intestine (> 20 %ID/g) and liver (> 10 %ID/g) were also detected.

Discussion

To date, [¹⁴C]MMPiP is the only characterized mGlu₇-targeted radiotracer. Although *in vitro* ARG study demon-

strated specific binding of [¹⁴C]MMPiP with mGlu₇, PET imaging of mGlu₇ in the living brain was not successful due to its low affinity to mGlu₇. In this study, we aim to evaluate an analog of MMPiP compound 18 with improved binding activity to mGlu₇ for PET imaging of mGlu₇. The synthesis of reference compound 18 was accomplished starting from the commercially available methoxybenzaldehyde 20 through seven steps and afforded the desired product 18 (MG7-2109) in 1.5% overall yield (Figure 3).

Schild analysis of the activity of **18** towards a concentration response of the group III mGlu agonist L-AP4 in the human mGlu₇ GIRK thallium flux assay is consistent with **18** acting as an mGlu₇ NAM. The inhibition potency (IC₅₀) of XAPO44, MMPIP and **18** against an mGlu₇ L-AP4 EC₈₀ response was also evaluated in the mGlu₇ GIRK thallium flux assay. Although the IC₅₀ values (MMPIP IC₅₀: 223 nM and **18** IC₅₀: 81 nM) diverged compared to the reported data (MMPIP IC₅₀: 26 nM and **18** IC₅₀: 12 nM) [30], a consistent inhibition trend was observed (**Figure 4B**). Of note, the reported IC₅₀ values for MMPIP and MG7-2109 were performed in a different assay utilizing the rat mGlu₇ receptor (calcium mobilization assays utilizing the promiscuous G protein G_{α15}) than the data reported herein. The pharmacokinetics of **18** was assessed via a single intravenous administration to SD rats (1 mg/kg) at 0.08, 0.25- and 1-hour time point (**Table S1**). The brain/plasma ratio of MG7-2109 ranged between 0.36 to 0.92, indicating that **18** has sufficient CNS penetration [31]. The off-target screening against 60 enzymes, GPCRs and ion channels indicated no significant off-target activity of **18** at the concentration of 10 μM. CNS MPO score is widely used to predict the probability of a drug candidate crossing the BBB, and MPO score ≥ 4 is highly desirable. The MPO score of **18** was calculated as 5.5, suggesting it has favorable BBB penetration properties. The selectivity assay for **18** towards mGlu₇ group I and other group II subtypes revealed that **18** is a selective compound for mGlu₇. The assay of CYP450s and hERG channel inhibitory properties is essential to evaluate drug toxicity and potential drug-drug interactions. The low inhibitory effect of CYP450s (IC₅₀ ≥ 3.2 μM) and hERG (IC₅₀ > 100 μM) demonstrated a favorable safety profile of MG7-2109.

Based on the structure of **18**, the methyl group on the phenol ring is proposed as the labeling position [32, 33]. Indeed, the precursor **19** could be easily synthesized from **18** by treatment with BBr₃. The carbon-11 methylation reaction was carried out by deprotonation of the hydroxyl group of precursor **19** with NaOH, yielding the desired radiotracer [¹¹C]**18** in 23% RCY. [¹¹C]**18** identity was confirmed by co-injection on HPLC with the standard **18**, and radiochemical purity was > 99%. *In vitro* stability tests of [¹¹C]**18** were performed in mice, rat, NHP and human serum, and then analyzed by radio-HPLC to determine the fraction of parent tracer. To investigate the efficacy and toxicity of [¹¹C]**18**, free drug concentrations in different types of plasma were measured by serum protein binding and fraction unbound (fu). While fewer differences in protein binding affinity were observed for mice, rat, and NHP, lower fu value of [¹¹C]**18** in human plasma were detected. *In vitro* ARG studies using rat brain sections were conducted to assess the specific binding of [¹¹C]**18**. Under baseline conditions, the highest radioactivity was found in the cerebellum. Other regions like cortex and thalamus had relatively lower levels of tracer uptake. However, blocking studies with excess **18** and MMPIP did not significantly reduce the radioactive signals, indicating limited specific binding of [¹¹C]**18** with mGlu₇. *In vivo* PET imaging

study in the rat brain revealed a homogeneous distribution of [¹¹C]**18** and rapid clearance, which was in accordance with limited specific binding observed in ARG results. The biodistribution study in mice was performed to assess the radioactivity distribution in different tissues/organs. High radioactivity was detected in the small intestine, liver, and kidney, which indicated that [¹¹C]**18** may be excreted through the hepatobiliary and urinary system.

Conclusion

In this work, a novel carbon-11 labeled radiotracer [¹¹C]**18** (MG7-2109) was synthesized via ¹¹C-methylation in 23% decay-corrected RCY. Comprehensive *in vitro* stability, protein binding, autoradiography, and *ex vivo* biodistribution studies were conducted to evaluate the performance characteristics of [¹¹C]**18** for specific binding to mGlu₇. Although this tracer was not successful for PET imaging of mGlu₇, this study offers valuable insights for developing mGlu₇-targeted radiotracer based on the isoxazolo(5,4-c)pyridine scaffold.

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Disclosure of conflict of interest

None.

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mGlu₇ radiotracer

Table S1. Individual plasma and brain concentrations and brain-Kp of 18 (MG7-2109) in male Sprague Dawley rats following a single intravenous administration (Dose: 1 mg/kg)

Time (hr)	Animal ID	Plasma concentration (ng/mL)	Brain concentration (ng/g)	Brain-Kp
0.08	1	1022.78	796.77	0.78
	2	935.73	887.16	0.95
	3	786.46	820.23	1.04
	Mean	914.99	834.72	0.92
	SD	119.52	46.90	0.13
	%CV	13	6	14
0.25	4	206.01	89.10	0.43
	5	333.68	88.98	0.27
	6	324.91	125.07	0.38
	Mean	288.20	101.05	0.36
	SD	71.31	20.80	0.09
	%CV	25	21	24
1	7	39.85	28.59	0.72
	8	57.77	40.98	0.71
	9	22.59	22.50	1.00
	Mean	40.07	30.69	0.81
	SD	17.59	9.42	0.16
	%CV	44	31	20

LLOQ: 2.02 ng/mL for plasma and 1.01 ng/mL for brain.