

Discovery of "Molecular Switches" within a Series of mGlu₅ Allosteric Ligands Driven by a "Magic Methyl" Effect Affording Both PAMs and NAMs with *In Vivo* Activity, Derived from an M₁ PAM Chemotype

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0.94), MPEP-site binding NAM **28d** (VU6044766) that displayed anxiolytic activity in a mouse marble burying assay. These data further support the growing body of literature concerning the existence of G protein-coupled receptor (GPCR) allosteric privileged structures, and the value and impact of subtle methyl group walks, as well as the highly productive fluorine walk, around allosteric ligand cores to stabilize unique GPCR conformations.

KEYWORDS: positive allosteric modulator (PAM), negative allosteric modulator (NAM), metabotropic glutamate receptor 5 (mGlu₅), CNS penetration, structure–activity-relationship (SAR), privileged structure

INTRODUCTION

G Protein-Coupled Receptors (GPCRs) are highly flexible membrane-bound proteins that are continuously sampling new active and inactive state ensembles with unique signaling profiles.¹⁻⁴ Allosteric ligands, both positive and negative allosteric modulators (PAMs and NAMs), are able to stabilize these unique receptor ensembles for therapeutic benefit, targeting less evolutionarily conserved binding (often inducedfit) sites, deep within the transmembrane binding domain.⁵⁻⁹ We were the first to report on the phenomenon of "molecular switches" among allosteric ligands, wherein subtle structural changes to the ligand can induce significant changes in receptor-ligand interaction and/or receptor conformation leading to changes in the mode of pharmacology, i.e., PAM to NAM, or changes in family subtype selectivity, i.e., mGlu₅ PAM to mGlu₃ NAM.¹⁰⁻¹⁴ More recently, and building on the classical concept of "privileged structures" of GPCR orthosteric ligands,¹⁵ a similar concept is emerging for GPCR allosteric modulator ligands.^{16,17} Here, we note two different concepts: (1) cooperativity-driven subtype selectivity, wherein an allosteric ligand binds to all members of a family but appears

to be a PAM of a single subtype due to high cooperativity with the orthosteric agonist (while other subtypes show neutral cooperativity) and (2) chemistry-driven pharmacology, where subtle changes to an mGlu₅ PAM ligand, for example, can lead to a potent mGlu₃ NAM ligand. Once recognized, it has become relatively straightforward to convert multiple mGlu₅ PAM chemotypes into mGlu₃ NAMs.^{13,14}

Of the eight currently identified metabotropic glutamate receptors (mGlu₁₋₈), mGlu₅ has received recent attention as a promising target for multiple CNS disorders and is one of the most studied, both preclinically and clinically.^{18,19} Several mGlu₅ NAMs have advanced into human testing, demonstrating proof-of-concept in numerous CNS disorders; however, historical NAMs with little chemotype diversity have exhibited

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Figure 1. Structures of representative mGlu₅ PAMs (1-4) and NAMs (5-8) that have displayed in vivo efficacy.

narrow therapeutic windows and dose-limiting psychotomimetic-like effects.^{14,21} Additionally, the prototypical heterobiaryl acetylene chemotype found in many of these compounds has long been associated with short half-life compounds and hepatotoxicity.^{20,21} Thus, novel, nonacetylene chemotypes are in high demand. In contrast, mGlu₅PAMs have proven more challenging and laden with pharmacological caveats (Figure 1).^{8,12} Early on, it became clear that mGlu₅ ago-PAMs and PAMs with large fold-shifts displayed neurotoxicity in preclinical rodent models, despite robust antipsychotic and pro-cognitive efficacy; thus, "pure" PAMs, devoid of agonist activity, became the target of choice. Ultimately, signal bias afforded an approach to achieve the benefits of mGlu₅ PAMs without the adverse effect liability.

In this case, VU0409551, a novel mGlu₅ PAM, exhibited distinct stimulus bias and selectively potentiated mGlu₅ coupling to $G_{\alpha q}$ -mediated signaling (but not mGlu₅ modulation of NMDAR currents or NMDAR-dependent synaptic plasticity in the rat hippocampus) and yielded robust antipsychotic-like and cognition-enhancing activity without neurotoxicity.^{22,23} However, the VU0409551 scaffold is an exception, and our Center continues to search for new mGlu₅ PAM chemotypes devoid of agonism and with the potential for signal bias. Here, we detail efforts toward new mGlu₅PAM and NAM chemotypes,

derived unexpectedly from an M_1 PAM scaffold,²⁴ and the impact of a "magic methyl"^{25–27} as a "molecular switch" to provide centrally active mGlu₅ PAMs and NAMs within a highly conserved series of 7-alkoxy-thieno[3,2-*b*]pyridine-5-carboxa-mides.

RESULTS AND DISCUSSION

A back-up lead optimization campaign for our M1 PAM VU-319/ACP-319 program led to the in-depth profiling of numerous scaffolds in an effort to identify "pure" M1 PAMs devoid of agonist activity and associated undesired cholinergic adverse events.^{24,28} One such campaign identified a true "needle in the haystack" with VU6007477 (9) as a rare member of a series of M_1 PAMs based on a pyrrolo[2,3-b]pyridine carboxamide core that displayed no $M_{1}\xspace$ agonism and a lack of cholinergic toxicity; however, K_p was modest and overall DMPK was modest to poor.²⁴ This led to a broader survey of alternate 5,6-heterobicyclic cores (Figure 2), and the identification of potent M_1 PAMs derived from a thieno[3,2-b]pyridine core, exemplified by VU6005549 (10). Amide scans of 10 led to numerous active moieties, yet the DMPK profiles were all unacceptable for further progression down the lead optimization flowchart. The benzylic CH₂ was shown to be a metabolic hotspot, but attempts to block this proved ineffective; therefore,

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Figure 2. Optimization of an M_1 PAM 9 into a novel mGlu₅ PAM series (e.g., 12) devoid of agonism, yet structurally similar to an mGlu₅ NAM (8), VU0424238 (auglurant).

we replaced the carbon linker with an oxygen atom to provide heterobiaryl ethers such as 11 and 12. To our surprise, both 11 and 12 were inactive as M_1 PAMs; however, we recognized the chemotype as one having high homology with our prototypical mGlu₅ NAM (13), VU0424238 (8, auglurant).²¹ Thus, we screened 11 and 12 in our mGlu₅ triple-add screening protocol (to simultaneously capture agonist, PAM and NAM activity), and both compounds proved to be weak mGlu₅ PAMs (EC₅₀s of 9.7 and 2.8 μ M, for 11 and 12, respectively), with no agonist (EC₅₀ > 30 μ M) activity. While this was not expected (an M₁ PAM switching to an mGlu₅ PAM), this result identified a new mGlu₅ PAM scaffold devoid of agonist activity.

To optimize mGlu₅ PAM 12, we initially centered our efforts on evaluating the structure-activity relationship (SAR) in the northern amide region through incorporating a diverse array of methylene-linked heteroaryl rings, with particular focus in walking a variety of substituents (i.e., halogen, alkyl, etc.) around the ring. As depicted in Scheme 1, the synthesis of these amide analogues 17 was conducted in a straightforward fashion in 5steps.²⁹ Starting from commercially available 7-chlorothieno-[3,2-b]pyridine 13, oxidation with m-CPBA provided the requisite N-oxide 14 to facilitate cyanation using trimethylsilyl cyanide and dimethylcarbamyl chloride. In the following step, the resulting intermediate 15 underwent microwave-assisted nucleophilic aromatic substitution with 5-hydroxy-2-methylpyridine, providing pyridyl-ether 16 in 82% yield. Desired analogs 17 were obtained following nitrile hydrolysis and standard amide formation (i.e., HATU) with a variety of commercially available aminomethylpyridines.



"Reagents and conditions: (a) *m*-CPBA, CH_2Cl_2 , rt, 19 h, 92–98%; (b) Me₃SiCN, Me₂NCOCl, CH_2Cl_2 , 44 h, 82%; (c) 5-hydroxy-2methylpyridine, K₂CO₃, DMF, microwave 150 °C, 15 min, 78–82%; (d) conc. HCl, reflux, 1.5 h, 88%; (e) NH₂R, HATU, DIPEA, DMF, 60 °C, 23–74%.

In short order, 24 synthesized amide derivatives 17 were screened using a calcium fluorescence "triple add" protocol which is able to simultaneously identify agonists, NAMs, and PAMs. The resulting SAR for this series was "shallow", a common observation during optimization of allosteric modulators, with almost all analogs 17 devoid of mGlu₅ PAM activity^{22,23} (see Supporting Information, Table S1). The only active compounds from this series resulted from the classic



Figure 3. Summary of structural changes conducted in the methylpyridine region of hit compound 12, with mGlu₅ activities of selected analogs 17.

Scheme 2. Synthesis of Pyridine Core Analogs 21^a



"Reagents and conditions: (a) $Zn(CN)_2$, $Pd(PPh_3)_4$, DMF, 120 °C, 2–4 h, 30–63% (b) 5-hydroxy-2-methylpyridine, K_2CO_3 , DMF, microwave 150 °C, 15–30 min, 73–92% (c) conc. HCl, reflux; (d) (5-fluoropyridin-2-yl)methanamine, HATU, DIPEA, DMF, 50 °C, 24–56% (2-steps).





^aReagents and conditions: (a) 5-hydroxy-2-methylpyridine, K₂CO₃, DMF, 100 °C, 2 h, 47%; (b) NaOH (2M), MeOH, reflux, 75 °C, 4 h (c) (5-fluoropyridin-2-yl)methanamine, HATU, DIPEA, DMF, 50 °C, 12% (2-steps).

"fluorine walk" (Figure 3), with substitution at the pyridine 5position (17a) affording a 5-fold improvement in potency in comparison to hit compound 12 (EC₅₀ = 0.532 μ M vs EC₅₀ = 2.83 μ M, respectively). Keeping the 5-flourine in place, we also investigated the effect of incorporating an additional nitrogen atom into the ring (17b); however, this was unfavorable, as was changing the position of the pyridine nitrogen (17c). Given these results, compound 17a was chosen as our new lead for further development.

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Figure 4. Summary of mGlu₅ activities of all synthesized core analogs 21 and 24.

Scheme 4. Synthesis Analogs 26, 27, and 28^a



^aReagents and conditions: (a) conc. HCl, reflux, 1 h; (b) (5-fluoropyridin-2-yl)methanamine, HATU, DIPEA, DMF, 60 °C, 1–3 h, 29–55% (2-steps); (c) 5-amino-2-methylpyridine, Xantphos, $Pd_2(dba)_3$, Cs_2CO_3 , 1,4-Dioxane, microwave 150 °C, 1.5 h, 53%; (d) B_2pin_2 , $PdCl_2(PPh_3)_2$, KOAc, 1,4-Dioxane, 100 °C, 1 h; (e) 5-(bromomethyl)-2-methyl-pyridine, $Pd(dppf)Cl_2$ ·DCM, Cs_2CO_3 , THF/H₂O, 100 °C, 10 h (f) Pyridinol, KOtBu, DMSO, 70 °C, 10 h, 41% (2-steps).

In the next stage of our optimization protocol, we sought to explore changes to the central core of 17a, with a particular focus on core truncation in an effort to identify the minimum required pharmacophore for mGlu₅ modulation. To prepare picolinamide derivatives 21a-d, a nitrile was first installed to the core via palladium-catalyzed Negishi cross-coupling using the requisite dichloropyridine 18 and zinc cyanide (Scheme 2). From the resulting picolinonitriles 19, including commercially available 4chloropicolinonitrile, the remaining steps were analogous to those described for the synthesis of analogs 17 (*vide supra*). Alternatively, derivative 24 was prepared starting from the appropriate methyl ester 22, followed by nucleophilic aromatic substitution, ester hydrolysis and amide coupling (Scheme 3).²⁹

As shown in Figure 4, truncation of the core to the pyridine **21d** was met with a complete loss of activity, highlighting the importance of the bicyclic core for mGlu₅ modulation. Interestingly, activity was partially restored by the addition of a methyl group at the 6-position (**21a**); however, this was not observed with methyl substitution at the 5-position (**21b**) or with 5,6-disubstitution (**21c**). Additionally, the quinoline

central core 24, a common bioisostere of the thienopyridine, afforded good potency with similar efficacy to 17a. While the central core SAR did provide alternative options, the original thienopyridine central core remained optimal.

In the final stage of our SAR exploration, we turned our attention to the southern pyridyl-ether region as another site of diversification. In addition to investigating alternate linker atoms in place of the ether, i.e., NH and CH₂, we also investigated changes to the pyridine ring in this region by altering both the position of the methyl substituent as well as the position of the nitrogen. To prepare these derivatives, we modified our original synthetic strategy toward 17 (vide supra) to allow for late-stage exploration of the southern region (Scheme 4). Starting from nitrile 15, hydrolysis followed by amide coupling provided intermediate 25 in good overall yield, which served as a common intermediate from which all analogs in this region were derived. The linker variants 26 and 27 were produced from 25 using either Buchwald-Hartwig cross-coupling with 5-amino-2-methylpyridine, or via Suzuki-Miyaura cross-coupling between the boronic ester of 25 and 5-(bromomethyl)-2-methyl-pyridine.

Alternatively, the pyridyl-ether analogs **28** were prepared using nucleophilic aromatic substitution with the requisite pyridinols.²⁹

As shown in Table 1, changing the linker was not tolerated, with both linker variants 26 and 27 resulting in a significant loss

Table 1. Structures and mGlu₅ Activities of Analogs 17a, 26, 27 and 28^a

R				
Compound No.	R-group	mGlu ₅ IC ₅₀ (μ M) ^a	%Glu min	
26	HN CH3	>30	N/A	
27	CH3	>30	N/A	
28a ^b	O CH ₃	0.260	3	
28b ^b	H ₃ C N	0.203	2	
28c	CH3 CH3	>30	N/A	
28d ^c	N N	0.112	2	
28e	CH3	1.9	80	

^{*a*}Calcium mobilization assays with HEK293 cells expressing human mGlu₅ performed in the presence of an EC₈₀ fixed concentration of glutamate; values represent the mean from one independent experiment performed in triplicate. ^{*b*}Values represent the mean from two (n = 2) independent experiments performed in triplicate. ^{*c*}Values represent the mean from three (n = 3) independent experiments performed in triplicate.

in activity. In contrast, very interesting results were obtained from the subtle modifications that were employed to the southern pyridine ring. Unexpectedly, almost all changes in this region engendered a "molecular switch"^{10–14} to afford mGlu₅ NAMs by a regiosiomeric "magic methyl" effect.^{25–27} Most notably, removal of the 6-methyl substituent (**28d**) resulted in a potent NAM (IC₅₀ = 0.112 μ M, 2% Glu_{max}), as did moving the methyl substituent to the 3- and 4-positions (compounds **28a** and **28b**, respectively). In contrast, incorporating the methyl in the 2-position had deleterious effect on potency (**28c**). As has been noted in previous reports describing mGlu₅ allosteric ligands, such modulation resulting from slight perturbations in structure raises concerns over putative metabolites and their potential to engender opposing modes of mGlu₅ modulation. Additionally, shifting the pyridine nitrogen *ortho* relative to the linker (28e) also provided an mGlu₅ NAM, however potency was significantly reduced in comparison to 28a, 28b, and 28d.

Evaluation of the DMPK profiles of compounds **28a**, **28b**, and **28d** supported the selection of **28d** to progress into mouse behavioral studies given its slightly more favorable clearance and plasma protein binding (mCL_{hep} = 82.4 mL/min/kg; mf_{u,p} = 0.062) in comparison to those of **28a** (mCL_{hep} = 85.1 mL/min/kg; mf_{u,p} = 0.042) and **28b** (mCL_{hep} = 86.5 mL/min/kg; mf_{u,p} = 0.046).²⁹

Primary *in vitro* pharmacology and DMPK of PAM 7a and NAM 28d

On the basis of these exciting results, both PAM 17a (VU6036486) and NAM 28d (VU6044766) were progressed for further profiling in order to assess their potential as *in vivo* tool compounds (Table 2). In addition to potent activity at mGlu₅, both compounds were found to demonstrate functional selectivity over the other mGlu receptors (mGlu_{1-4,7,8} > 30 μ M). To determine whether these compounds interacted at the common 2-methyl-6-(phenylethynyl)-pyridine (MPEP) allosteric binding site, displacement studies were conducted using

Table 2. Summary of In Vitro Pharmacology	and DMPK Data
for Compounds 17a and 28d	

Property	PAM 17a	NAM 28d
Physicochemical		
MW	394	380
cLogP	2.41	2.22
tPSA	77	77
In vitro Pharmacology		
hmGlu5 EC ₅₀ (μ M)	0.52	N/A
hmGlu5 IC ₅₀ (μ M)	N/A	0.11
$mGlu_5 K_i (\mu M)$	>10	0.16
In vitro PK		
CL _{int} (mL/min/kg)		
Rat	152	314
Mouse	ND	970
Human	173	47.0
CL_{hep} (mL/min/kg)		
Rat	47.8	57.2
Mouse	ND	82.4
Human	18.7	14.5
$f_{\mathrm{u,p}}$		
Rat	0.048	0.115
Mouse	ND	0.062
Human	0.056	0.068
$\operatorname{Rat} f_{\mathrm{u},\mathrm{b}}$	0.030	0.035
In vivo Brain Distribution		
Rat IV PBL cassette		
K _p	0.83 ^a	0.94 ^b
K _{p,uu}	0.52 ^a	0.29 ^b
In vivo PK		
IV dose (mg/kg)	0.2 (rat)	1.0 (mouse)
CL_{p} (mL/min/kg)	36.9	141
V _{ss} (L/kg)	2.20	8.2
Elim. $t_{1/2}$ (h)	1.37	0.69
PO dose (mg/kg)	10 (rat)	10 (mouse)
C_{\max} (ng/mL)	152	ND
t_{\max} (h)	0.50	ND
AUC _{inf} (h*ng/mL)	371	ND
F (%)	8.22	ND

^a0.20 mg/kg dose. ^b0.25 mg/kg dose.

the radioligand $[{}^{3}H]$ 3-methoxy-5-pyridin-2-ylethynylpyridine ($[{}^{3}H]$ methoxyPEPy) which is a close structural analog of MPEP. As illustrated in Figure 5, the mGlu₅ NAM **28d**



Figure 5. Radioligand competition binding experiments with 17a (VU6036486), **28d** (VU6044766), and $[^{3}H]$ -3-methoxy-5-(pyridin-2-ylethynyl)pyridine, showing an interaction with the classical MPEP allosteric site on mGlu₅ (n = 1).

(VU6044766) induced full displacement of radioligand binding, suggesting an interaction with the allosteric MPEP site ($K_i = 0.16 \ \mu$ M). Interestingly, PAM **17a** (VU6036486) exhibited weak binding at concentrations up to 100 μ M, indicating an interaction at either the classical MPEP binding site and/or an alternative overlapping site.^{22,23}

As shown in Table 2, compounds 17a and 28d were found to have good free fraction in plasma ($f_{u,s} > 0.05$) and in brain ($f_{u,s} > 0.03$); however, they were predicted to have high hepatic clearance across all species tested. The CNS penetration of both compounds was assessed using a rat plasma/brain level (PBL) cassette assay which revealed good exposure in brain for both PAM 17a ($K_p = 0.83$, $K_{p,uu} = 0.52$) and NAM 28d ($K_p = 0.94$, $K_{p,uu} = 0.29$). These data prompted further evaluation of discrete IV and PO pharmacokinetics for both compounds. PAM 17a displayed moderate clearance (36.9 mL/min/kg), a half-life of 1.37 h with a moderate volume of distribution (2.2 L/kg). Despite the promising *in vivo* data, the compound had low oral bioavailability (8%). For NAM 28d, the *in vivo* mouse study revealed superhepatic clearance (141 mL/min/kg), a half-life of 0.69 h and a high volume of distribution (8.2 L/kg). Of note, there was no measurable drug in blood following mouse PO administration. $^{29}\,$

In Vivo Pharmacology

With a relatively potent mGlu₅ PAM (17a) and NAM (28d) with acceptable PK profiles and brain exposure, we were now poised to determine if the mGlu₅ modulation observed in our *in vitro* cellular assays would be mirrored by *in vivo* behavioral studies. To evaluate PAM 17a, we assessed its ability to reverse amphetamine-induced hyperlocomotion (AHL) which is used as a predictor of antipsychotic activity.^{22,23} In this study, PAM 17a (VU6036486) was administered as a 30 or 60 mg/kg i.p. dose 30 min prior to a 0.75 mg/kg s.c. dose of amphetamine. As shown in Figure 6, PAM 17a reversed AHL by ~30% at a dose of 60 mg/kg.²⁹

To assess NAM **28d**, we employed a marble burying assay to evaluate anxiolytic activity. Naive mice are well-known to bury foreign objects, e.g., glass marbles, in deep bedding.²¹ This behavior can be inhibited with pretreatment of low doses of certain benzodiazepines or selective serotonin reuptake inhibitors and has also been proven to be sensitive to a diverse array of mGlu₅ NAM chemotypes.²¹ As shown in Figure 7, a 30 mg/kg i.p. dose produced a significant reduction in marble burying relative to the vehicle-treated group, suggesting anxiolytic activity.²⁷

CONCLUSION

In summary, we have described the multidimensional optimization of a novel mGlu₅ PAM and mGlu₅ NAM chemotype based on a thieno [3,2-b] pyridine core scaffold, derived from an M1 PAM ligand. Despite "shallow" SAR for the majority of our campaign, modification to the northern amide region effected a 5-fold improvement in potency of the original initial hit 2, resulting in compound 17a (VU6036486, $EC_{50} =$ 0.52 μ M). Most interestingly, "molecular switches" were discovered through slight structural modifications to the southern pyridyl-ether region, resulting in the discovery of multiple mGlu₅ NAMs, the most potent being compound 28d (VU6044766, IC₅₀ = 0.11 μ M). This report highlights that "molecular switches" can result from small structural changes, such as a "magic methyl", and thus highlights the need to thoroughly investigate and develop chemical SAR as well as characterize metabolites of mGlu₅ allosteric ligands as these may



Figure 6. Dose-dependent reversal of amphetamine-induced hyperlocomotion in rats by PAM 17a; n = 7-8 male Sprague–Dawley rats per treatment group; vehicle = 10% Tween 80 in H₂O; *p < 0.05, **p < 0.01 vs vehicle/amphetamine (AMPH) control group.



Figure 7. Inhibition of marble burying with NAM **28d** (VU6044766) administered at a dose of 30 mg/kg; n = 14 male CD-1 mice per treatment group; 15 min pretreatment, 30 min burying time; vehicle = polysorbate 80 (IP); *p < 0.05 versus vehicle control group. VU0424238 was used as a positive control and administered at a dose of 30 mg/kg.

result in opposing modes of pharmacology. Finally, these data further strengthen the position that there are GPCR allosteric privileged structures and that these scaffolds can be employed to access allosteric ligands for multiple GPCR targets.

EXPERIMENTAL SECTION

Chemistry

All compounds were purified to \geq 95% as determined by analytical LCMS (214 nm, 254 nm and ELSD) as well as ¹H and ¹³C NMR and HRMS. The general chemistry, experimental information, and syntheses of all other compounds, along with detailed pharmacology, and DMPK methods are supplied in the Supporting Information.

7-Chlorothieno[3,2-6]pyridine 4-oxide (14)

To a solution of 7-chlorothieno[3,2-b]pyridine (2.00 g, 11.8 mmol) in dichloromethane (50 mL) was added 3-chloroperoxybenzoic acid (3.17 g, 14.2 mmol) in equal portions over a 5 min period. After 19 h at room temperature, a solution of $Na_2S_2O_3$ (10% aqueous) was added to the reaction, and the organic layer was separated and washed with a solution of K₂CO₃(10% aqueous). The aqueous layer was extracted with dichloromethane (3x), and the combined organic layers washed with brine, dried over Na2SO4 (anhydrous), filtered, and concentrated under reduced pressure to afford the title compound as a pale-yellow solid, which was used in the following step without further purification (2.12 g, 11.0 mmol, 93% yield). ¹H NMR (400 MHz, CDCl₃) $\delta = 8.26$ (d, J = 6.7 Hz, 1H), 7.87 (d, J = 5.7 Hz, 1H), 7.75 (d, J = 5.7 Hz, 1H),7.23 (d, J = 6.7 Hz, 1H). ¹³C NMR (101 MHz, CDCl₃) $\delta = 147.1, 136.4,$ 136.0, 131.7, 126.8, 120.1, 120.0. LC-MS (ESI+): $R_{\rm T} = 0.432 \text{ min}, 97\%$ at 215 nm and >98% at 254 nm, $m/z = 186.3 [M + H]^+$. HR-MS (TOF, ESI+): for C_7H_4CINOS [M], calc'd 185.9775 [M + H]⁺, found 185.9770.

7-Chlorothieno[3,2-b]pyridine-5-carbonitrile (15)

To a solution of 14 (2.11 g, 11.3 mmol) in dichloromethane (50 mL) was added trimethylsilyl cyanide (2.80 mL, 22.7 mmol). After 15 min at room temperature, dimethylcarbamoyl chloride (2.00 mL, 22.7 mmol) was added dropwise. After 19 h at room temperature, an additional amount of trimethylsilyl cyanide (2.80 mL, 22.7 mmol) and dimethylcarbamoyl chloride (2.00 mL, 22.7 mmol) and dimethylcarbamoyl chloride (2.00 mL, 22.7 mmol) were added. After a total of 44 h, a solution of K₂CO₃(10% aqueous) was added to the reaction mixture, and the organic layer separated. The aqueous layer was extracted with dichloromethane (3x), and the combined organic layers were washed with brine, dried over Na₂SO₄(anhydrous), filtered, and concentrated under reduced pressure. Purification by silica-gel flash chromatography (0–10% EtOAc/hexanes) afforded the title compound as a pale-yellow solid (1.86 g, 9.26 mmol, 82% yield). ¹H NMR (400 MHz, CDCl₃) δ = 7.99 (d, *J* = 5.5 Hz, 1H), 7.69–7.62 (m, 2H). ¹³C NMR (101 MHz, CDCl₃) δ = 157.6, 139.3, 136.3, 134.4, 131.6,

126.0, 122.0, 116.9. LC-MS (ESI+): $R_T = 0.762 \text{ min}$, 97% at 215 nm and 98% at 254 nm, $m/z = 195.3 \text{ [M + H]}^+$. HR-MS (TOF, ESI+): for $C_8H_3ClN_2S$ [M], calc'd 194.9778 [M + H]⁺, found 194.9776.

7-((6-Methylpyridin-3-yl)oxy)thieno[3,2-b]pyridine-5-carbonitrile (16)

To a microwave vial were added 15 (1.00 g, 5.14 mmol), 3-hydroxy-6methylpyridine (0.673 g, 6.17 mmol), potassium carbonate (1.49 g, 10.8 mmol), and DMF (15 mL). The vial was capped and irradiated in a microwave reactor at 150 °C for 15 min. Upon completion, the reaction was diluted with water and dichloromethane, passed through a phase separator washing with dichloromethane (3x), and the collected organic phase was concentrated to dryness. Purification by silica-gel flash chromatography (0-50% EtOAc/hexanes) afforded the title compound as a white powder (1.15 g, 4.21 mmol, 82% yield). ¹H NMR (400 MHz, MeOD) δ = 8.48 (d, J = 2.7 Hz, 1H), 8.21 (d, J = 5.5 Hz, (10) 111, 7.75 (dd, J = 8.5, 2.9 Hz, 1H), 7.62 (d, J = 5.5 Hz, 1H), 7.51 (d,8.5 Hz, 1H), 7.13 (s, 1H), 2.63 (s, 3H). 13 C NMR (101 MHz, MeOD) δ = 161.6, 160.7, 157.9, 149.9, 142.3, 135.9, 133.6, 131.7, 127.0, 126.7, 125.8, 118.2, 109.0, 23.2. LC-MS (ESI+): $R_T = 0.604 \text{ min}$, > 98% at 215 and 254 nm, $m/z = 268.4 [M + H]^+$. HR-MS (TOF, ESI+): for $C_{14}H_9N_3OS$ [M], calc'd 268.0539 [M + H]⁺, found 268.0538.

N-((5-Fluoropyridin-2-yl)methyl)-7-((6-methylpyridin-3-yl)oxy)thieno[3,2-*b*]pyridine-5-carboxamide (17a, VU6036486)

Step 1. A mixture of compound 16 (2.05 g, 7.67 mmol) and concentrated HCl (37% aqueous, 75.6 mmol) was heated at 100 °C for 1.5 h. After the heat was removed, the reaction was neutralized with NaOH (6 M, aqueous) at room temperature, and the water and solvent were removed under reduced pressure. The crude residue was dissolved in 10% MeOH/DCM, the undissolved salt was filtered, and the collected filtrate was concentrated under reduced pressure to afford the title compound as an off-white solid, which was used in following step without further purification (2.01 g, 6.74 mmol, 88% yield). ¹H NMR (400 MHz, DMSO) δ = 8.53 (d, J = 2.9 Hz, 1H), 8.31 (d, J = 5.5 Hz, 1H), 7.77 (dd, J = 8.5, 2.9 Hz, 1H), 7.74 (d, J = 5.5 Hz, 1H), 7.45 (d, J = 8.5 Hz, 1H), 7.15 (s, 1H), 2.55 (s, 3H). ¹³C NMR (101 MHz, DMSO) δ = 166.1, 160.1, 158.4, 156.2, 149.0, 147.8, 142.1, 133.9, 129.4, 125.3, 124.5, 124.1, 103.8, 23.5. LC-MS (ESI+): $R_T = 0.417 \text{ min}$, 96% at 215 and 254 nm, $m/z = 287.4 [M + H]^+$. HR-MS (TOF, ESI+): for $C_{14}H_{10}N_2O_3S[M]$, calc'd 287.0485 $[M + H]^+$, found 287.0485.

Step 2. To a solution of carboxylic acid (6.00 mg, 21.0 μ mol) in DMF (0.30 mL) was added N,N-diisopropylethylamine (9 µL, 52.4 μ mol), HATU (15.9 mg, 41.8 μ mol), and 5-fluoropyridin-2-yl)methanamine (2.91 mg, 23.1 μ mol), and the reaction was heated at 60 °C. After 3 h, the reaction was removed from the heat, and once it was at room temperature, it was filtered and directly purified by reverse phase HPLC (10-40% over 4 min, Method 1) to afford the title compound as an off-white residue (6.20 mg, 15.4 μ mol, 74% yield). ¹H NMR (400 MHz, CDCl₃) δ = 8.86 (t, J = 6.1 Hz, 1H), 8.46–8.42 (m, 2H), 7.84 (d, *J* = 5.5 Hz, 1H), 7.58 (d, *J* = 5.5 Hz, 1H), 7.47 (s, 1H), 7.44 (dd, *J* = 8.5, 2.9 Hz, 1H), 7.39–7.34 (m, 2H), 7.26 (d, J = 8.4 Hz, 1H), 4.75 (d, J = 6.0 Hz, 2H), 2.63 (s, 3H). ¹³C NMR (101 MHz, CDCl₃) δ = 164.6, 161.0, 158.9 (d, J = 255.2 Hz), 157.9, 156.4, 153.3 (d, J = 3.9 Hz), 150.5, 148.3, 142.2, 137.6 (d, J = 23.9 Hz), 132.3, 129.3, 125.3, 125.1, 124.5, 123.7 (d, J = 18.5 Hz), 123.1 (d, J = 4.3 Hz), 102.2, 44.5 (d, J = 1.6 Hz), 23.9. LC-MS (ESI+): $R_T = 0.752 \text{ min}$, > 98% at 215 and 254 nm, m/z =395.3 [M + H]⁺. HR-MS (TOF, ESI+): for C₂₀H₁₅FN₄O₂S [M], calc'd 395.0973 [M + H]⁺, found 395.0976.

7-Chloro-*N*-((5-fluoropyridin-2-yl)methyl)thieno[3,2-*b*]pyridine-5-carboxamide (25)

Step 1. A mixture of compound **15** (100 mg, 0.514 mmol) and concentrated HCl (37% aqueous, 0.850 mL, 10.2 mmol) was heated to 100 °C. After 1 h, the reaction was removed from the heat and concentrated under reduced pressure to provide the carboxylic acid intermediate, which was used in the following step assuming quantitative yield and without further purification.

Step 2. To a solution of carboxylic acid (0.514 mmol) in DMF (2.0 mL) were added *N*,*N*-diisopropylethylamine (0.224 mL), HATU (391

mg, 1.03 mmol), and (5-fluoropyridin-2-yl)methanamine (71.2 mg, 0.565 mL), and the reaction was heated to 60 °C. After 3 h, the reaction was removed from the heat, diluted with water and dichloromethane once it was at room temperature, and passed through a phase separator with washing with dichloromethane $(3\times)$, and the combined organic phases were concentrated by drying. The crude product was purified by silica-gel column chromatography (0-50% EtOAc/hexanes) to afford the title compound as a white solid (99.2 mg, 0.277 mmol, 55% yield). ¹H NMR (400 MHz, CDCl₃) δ = 8.87 (t, J = 5.8 Hz, 1H), 8.46 (t, J = 1.8 Hz, 1H), 8.23 (s, 1H), 7.88 (d, J = 5.5 Hz, 1H), 7.60 (d, J = 5.5 Hz, 1H), 7.40 (d, I = 2.3 Hz, 1H), 7.39 (d, I = 1.8 Hz, 1H), 4.81 (dd, I = 5.9, 1.0 Hz, 2H). ¹³C NMR (101 MHz, CDCl₃) δ = 164.1, 158.9 (d, J = 255.2 Hz), 156.0, 153.1 (d, J = 3.9 Hz), 149.1, 139.6, 137.6 (d, J = 24.0 Hz), 135.5, 132.9, 125.7, 123.9 (d, J = 18.5 Hz), 123.2 (d, J = 4.3 Hz), 117.4, 44.4 (d, J = 1.6 Hz). LC-MS (ESI+,): $R_T = 0.844$ min, 90% at 215 nm and 92% at 254 nm, $m/z = 322.2 [M + H]^+$. HR-MS (TOF, ESI+): for $C_{14}H_9ClFN_3OS [M]$, calc'd 322.0212 $[M + H]^+$, found 322.0212.

N-((5-fluoropyridin-2-yl)methyl)-7-(pyridin-3-yloxy)thieno-[3,2-*b*]pyridine-5-carboxamide (28d, VU6044766)

To a reaction vial under an atmosphere of nitrogen were added 3pyridinol (8.7 mg, 91.4 µmol) and DMSO (0.300 mL), followed by potassium tert-butoxide (10.2 mg, 91.4 μ mol). After 15 min at room temperature, 25 (15.0 mg, 45.7 µmol) in DMSO (0.100 mL) was added, and the reaction was heated to 70 °C. After 16 h, the reaction was removed from the heat, filtered, and directly purified by reversephase HPLC (25-55%, Method 2) to provide the title compound as a white residue (8.2 mg, 21.1 μ mol, 46% yield). ¹H NMR (400 MHz, $CDCl_3$) $\delta = 8.88$ (t, J = 5.4 Hz, 1H), 8.62 - 8.55 (m, 2H), 8.48 - 8.42 (m, 1H), 7.86 (d, J = 5.5 Hz, 1H), 7.60 (d, J = 5.5 Hz, 1H), 7.56 (ddd, J = 8.3, 2.8, 1.4 Hz, 1H), 7.52 (s, 1H), 7.44 (ddd, J = 8.4, 4.7, 0.7 Hz, 1H), 7.41–7.34 (m, 2H), 4.77 (d, J = 6.0 Hz, 2H). ¹³C NMR (101 MHz, $CDCl_3$) $\delta = 164.5$, 160.5, 158.9 (d, J = 255.2 Hz), 158.0, 153.2 (d, J =3.8 Hz), 150.6, 150.6, 146.9, 142.9, 137.5 (d, J = 24.1 Hz), 132.5, 128.8, 125.4, 125.3, 124.9, 123.9 (d, J = 18.4 Hz), 123.2 (d, J = 4.3 Hz), 102.6, 44.4. LC-MS (ESI+): $R_{\rm T}$ = 0.750 min, > 98% at 215 and 254 nm, m/z = 381.0 $[M + H]^+$. HR-MS (TOF, ESI+): for $C_{19}H_{13}FN_4O_2S$ [M], calc'd 381.0816 [M + H]⁺, found 381.0819.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acsbiomedche-mau.1c00024.

Experimental procedures and spectroscopic data for selected compounds, detailed pharmacology and DMPK methods (PDF)

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Notes

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ABBREVIATIONS

mGlu₅, metabotropic glutamate receptor subtype 5; CRC, concentration—response-curve; NAM, negative allosteric modulator; PAM, positive allosteric modulator; PBL, plasma:brain level; SAR, structure—activity-relationships

REFERENCES

(1) Wotten, D.; Christopoulos, C.; Marti-Solano, M.; Babu, M. M.; Sexton, P. M. Mechanisms of signalling and biased agonism in G protein-coupled receptors. *Nat. Rev. Mol. Cell Biol.* **2018**, *19*, 638–653. (2) Hilger, D.; Masureel, M.; Kobilka, B. K. Structure and dynamics of

GPCR signaling complexes. *Nat. Struct. Mol. Biol.* **2018**, *25*, 1–34. (3) Hauser, A. S.; Attwood, M. M.; Rask-Andersen, M.; Schioth, H. B.; Gloriam, D. E. Trends in GPCR drug discovery: new agents, targets and indications. *Nat. Rev. Drug Discovery* **2017**, *16*, 829–842.

(4) Niswender, C. M.; Conn, P. J. Metabotropic glutamate receptors: physiology, pharmacology, and disease. *Annu. Rev. Pharmacol. Toxicol.* **2010**, *50*, 295–322.

(5) Conn, P. J.; Christopolous, A.; Lindsley, C. W. Allosteric Modulators of GPCRs as a Novel Approach to Treatment of CNS Disorders. *Nat. Rev. Drug Discovery* **2009**, *8*, 41–54.

(6) Melancon, B. J.; Hopkins, C. R.; Wood, M. R.; Emmitte, K. A.; Niswender, C. M.; Christopoulos, A.; Conn, P. J.; Lindsley, C. W. Allosteric Modulation of 7 Transmembrane Spanning Receptors: Theory, Practice and Opportunities for CNS Drug Discovery. J. Med. Chem. 2012, 55, 1445–1464.

(7) Wenthur, C. J.; Gentry, P. R.; Mathews, T. P.; Lindsley, C. W. Drugs for allosteric sites on receptors. *Annu. Rev. Pharmacol. Toxicol.* **2014**, *54*, 165–184.

(8) Conn, P. J.; Lindsley, C. W.; Meiler, J.; Niswender, C. M. Opportunities and challenges in the discovery of allosteric modulators of GPCRs for the treatment of CNS disorders. *Nat. Rev. Drug Discovery* **2014**, *13*, 692–708.

(9) Lindsley, C. W.; Emmitte, K. A.; Hopkins, C. R.; Bridges, T. M.; Gregory, K. A.; Niswender, C. M.; Conn, P. J. Practical strategies and concepts in GPCR allosteric modulator discovery: Recent advances with metabotropic glutamate receptors. *Chem. Rev.* **2016**, *116*, 6707–6741.

(10) Sharma, S.; Kedrowski, J.; Rook, J. M.; Smith, J. M.; Jones, C. K.; Rodriguez, A. L.; Conn, P. J.; Lindsley, C. W. Discovery of Molecular Switches that Modulate Modes of mGluR5 Pharmacology *In Vitro* and *In Vivo* Within a Series of Functionalized 5-(phenylethynyl)pyrimidines'. J. Med. Chem. **2009**, *52*, 4103–4106.

(11) Lindsley, C. W.; Philip, S. Portoghese Medicinal Chemistry Lectureship: Drug Discovery Targeting Allosteric Sites. J. Med. Chem. 2014, 57, 7485–7498 2013.

(12) Wood, M. R.; Hopkins, C. R.; Brogan, J. T.; Conn, P. J.; Lindsley, C. W. 'Molecular switches' on allosteric ligands that modulate modes of pharmacology. *Biochemistry* **2011**, *50*, 2403–2410.

(13) Wenthur, C. J.; Niswender, C. M.; Morrison, R.; Daniels, J. S.; Conn, P. J.; Lindsley, C. W. Discovery of (*R*)-(2-fluoro-4-((4methoxyphenyl)ethynyl)phenyl)(3-hydroxypiperidin-1-yl)methanone (ML337), an mGlu₃ selective and CNS penetrant negative allosteric modulator (NAM). *J. Med. Chem.* **2013**, *56*, 3713–3718.

(14) Engers, J. L.; Rodriguez, A. L.; Konkol, L. C.; Morrison, R. D.; Thompson, A. D.; Byers, F. W.; Blobaum, A. L.; Chang, S.; Loch, M. T.; Niswender, C. M.; Daniels, J. S.; Jones, C. K.; Conn, P. J.; Lindsley, C. W.; Emmitte, K. A. Discovery of VU0650786: A selective and CNS penetrant negative allosteric modulator of metabotropic glutamate receptor subtype 3 with antidepressant and anxiolytic activity in rodents. J. Med. Chem. 2015, 58, 7485–7500.

(15) Evans, B. E.; Rittle, K. E.; Bock, M. G.; DiPardo, R. M.; Freidinger, R. M.; Whitter, W. L.; Lundell, G. F.; Veber, D. F.; Anderson, P. S.; Chang, R. S. L.; Lotti, V. J.; Cerino, D. J.; Chen, T. B.; Kling, P. J.; Kunkel, K. A.; Springer, J. P.; Hirshfieldt, J. Methods for drug discovery: development of potent, selective, orally effective cholecystokinin antagonists. J. Med. Chem. **1988**, *31*, 2235–2246.

(16) Davis, D. C.; Bungard, J. D.; Chang, S.; Rodriguez, A. L.; Blobaum, A. L.; Boutaud, O.; Melancin, B. J.; Niswender, C. M.; Conn, P. J.; Lindsley, C. W. Lead optimization of the VU0486321 series of mGlu₁ PAMs. Part 4: SAR reveals positive cooperativity across multiple mGlu receptor subtypes leading to subtype *un*selective PAMs. *Bioorg. Med. Chem. Lett.* **2021**, *37*, 127724.

(17) Berizzi, A. E.; Bender, A. M.; Conn, P. J.; Lindsley, C. W.; Sexton, P. M.; Langmead, C. J.; Christopoulos, A. Structure-activity relationships of *pan*-G-_{q/11} coupled muscarinic acetylcholine receptor positive allosteric modulators'. *ACS Chem. Neurosci.* **2018**, *9*, 1818–1828.

(18) Bennett, K. A.; Christopher, J. A.; Tehan, B. G. Structure-based discovery and development of metabotropic glutamate receptor 5 negative allosteric modulators. *Adv. Pharmacol.* **2020**, *88*, 35–58.

(19) Arsova, A.; Mollr, T. C.; Hellyer, S. D.; Vedel, L.; Foster, S. R.; Hansen, J. L.; Brauner-Osborn, H.; Gregory, K. J. Positive Allosteric Modulators of Metabotropic Glutamate Receptor 5 as Tool Compounds to Study Signaling Bias. *Mol. Pharmacol.* **2021**, *99*, 328– 341.

(20) Emmitte, K. A. mGlu5 negative allosteric modulators: a patent review (2013–1016). *Expert Opin. Ther. Pat.* **2017**, *27*, 691–706.

(21) Felts, A. S.; Rodriguez, A. L.; Blobaum, A. L.; Morrison, R. D.; Bates, B. S.; Thompson, A. D.; Rook, J. M.; Byers, F. W.; Chang, S.; Venable, D. F.; Luscombe, V. B.; Tamagnan, G. D.; Bridges, T. M.; Niswender, C. M.; Daniels, J. S.; Jones, C. K.; Conn, P. J.; Lindsley, C. W.; Emmitte, K. A. Discovery of N-(5-fluoropyridin-2-yl)-6-methyl-4(pyrimidin-5-yloxy)picolinamide (VU0424238): A novel negative allosteric modulator of metabotropic glutamate receptor subtype 5 selected for clinical development. J. Med. Chem. 2017, 60, 5072-5085.

(22) Rook, J. M.; Xiang, Z.; Lv, X.; Ghosal, A.; Dickerson, J.; Bridges, T. M.; Johnson, K. A.; Bubser, M.; Gregory, K. J.; Vinson, P. N.; Byun, N.; Stauffer, S. R.; Daniels, J. S.; Niswender, C. M.; Lavreysen, H.; Mackie, C.; Conde; Ceide, S.; Alcazar, J.; Bartolome, J. M.; Macdondald, G. J.; Steckler, T.; Jones, C. K.; Lindsley, C. W.; Conn, P. J. Biased mGlu₅ positive allosteric modulators provide in vivo efficacy without potentiating mGlu₅ modulation of NMDAR currents. *Neuron* **2015**, *86*, 1029–1040.

(23) Conde-Ceide, S.; Martin-Martin, M. L.; Alcazar, J.; Manka, T.; Tong, H. M.; Garcia-Barrantes, P. M.; Lavreysen, H.; Mackie, C.; Vinson, P. N.; Daniels, S. J.; Menges, A.; Niswender, C. M.; Jones, C. K.; Macdonald, G. J.; Steckler, T.; Conn, P. J.; Stauffer, S. R.; Bartolome-Nebreda, J. M.; Lindsley, C. W. Discovery of VU0409551/JNJ-46778212: An mGlu₅ Positive Allosteric Modulator Clinical Candidate Targeting Schizophrenia. *ACS Med. Chem. Lett.* **2015**, *6*, 716–720.

(24) Engers, J. L.; Childress, E. S.; Long, M. F.; Capstick, R. A.; Luscombe, V. B.; Cho, H. P.; Dickerson, J. W.; Rook, J. M.; Blobaum, A. L.; Niswender, C. M.; Conn, P. J.; Lindsley, C. W. VU6007477, a novel M₁ PAM based on a pyrrolo[2,3-*b*]pyridine carboxamide core devoid of cholinergic side effects'. *ACS Med. Chem. Lett.* **2018**, *9*, 917–922.

(25) Zhao, J.; Zhang, D.; Stashko, M. A.; DeRyckere, D.; Vasileiadi, E.; Parker, R. E.; Hunter, D.; Lu, Q.; Zhang, Y.; Norris-Drouin, J.; Li, B.; Drewry, D. H.; Kireev, D.; Graham, D. K.; Earp, H. S.; Frye, S. V.; Wang, X. Highly selective MERTK inhibitors achieved by a single methyl group. *J. Med. Chem.* **2018**, *61*, 10242–10254.

(26) Leung, C. S.; Leung, S. S.; Tirado-Rives, J.; Jorgensen, W. Methyl Effects on Protein-Ligand Binding. *J. Med. Chem.* **2012**, *55*, 4489-4500.

(27) Schonherr, H.; Cernak, T. Profound Methyl Effects in Drug Discovery and a Call for New C-H Methylation Reactions. *Angew. Chem., Int. Ed.* **2013**, *52*, 12256–12267.

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