



Re-exploration of the mGlu₁ PAM Ro 07-11401 scaffold: Discovery of analogs with improved CNS penetration despite steep SAR



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ABSTRACT

This letter describes the re-exploration of the mGlu₁ PAM Ro 07-11401 scaffold through a multi-dimensional, iterative parallel synthesis approach. Unlike recent series of mGlu₁ PAMs with robust SAR, the SAR around the Ro 07-11401 structure was incredibly steep (only ~6 of 200 analogs displayed mGlu₁ PAM activity), and reminiscent of the CPPHA mGlu₅ PAM scaffold. Despite the steep SAR, two new thiazole derivatives were discovered with improved in vitro DMPK profiles and ~3- to 4-fold improvement in CNS exposure (K_p s 1.01–1.19); albeit, with a ~3-fold diminution in mGlu₁ PAM potency, yet comparable efficacy (~5-fold leftward shift of the glutamate concentration–response curve at 10 μ M). Thus, this effort has provided additional CNS penetrant mGlu₁ PAM tools in a different chemotype than the VU0486321 scaffold. These compounds will permit a better understanding of the pharmacology and therapeutic potential of selective mGlu₁ activation, while highlighting the steep SAR challenges that can often be encountered in GPCR allosteric modulator discovery.

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Efforts towards the development of positive allosteric modulators (PAMs) of the metabotropic glutamate receptor subtype 1 (mGlu₁) were pioneered by Knoflach and co-workers at Roche, resulting in **1–4** (Fig. 1).^{1–3} These small molecule PAMs, coupled with data generated with negative allosteric modulators (NAMs) of mGlu₁,^{4,5} highlighted issues with species differences due to a single amino acid in rat versus human mGlu₁,^{6,7} and thus **4**, a PAM active on both human and rat mGlu₁, emerged as a valuable tool compound, despite modest CNS penetration (K_p = 0.29 and high protein binding (f_u < 0.01). For over a decade, **4** was the only in vivo tool compound to study selective mGlu₁ activation.^{8–11} Based on recent genetic data implicating *GRM1* in schizophrenia,^{12–14} coupled with data showing that the adverse effect liabilities of group I metabotropic glutamate receptors (mGluRs) are mediated by mGlu₅ and not mGlu₁,¹⁵ our lab has launched a program to develop the next generation of mGlu₁ PAMs.^{14–18} In the past year, we have reported on the discovery and optimization of novel mGlu₁ PAMs **5–7** with improved potency (EC_{50} s < 20 nM), DMPK profiles (f_u s > 2.0% unbound) and CNS penetration (K_p s > 1)

to afford new avenues for target validation and to assess the therapeutic potential of selective mGlu₁ activation.^{14–18}

Previously, revisiting the mGlu₄ PAM (–)-PHCCC scaffold led to the discovery of improved tool compounds.¹⁹ Therefore, over a decade after its discovery, we felt it was prudent to revisit the Ro 07-11401 scaffold in an effort to develop an in vivo tool compound within this series with improved disposition to account for any chemotype or ligand-biased pharmacology and expand the repertoire of validation tools for mGlu₁.

In our functional assays, Ro 07-11401 (**4**) was an equipotent mGlu₁ PAM on both rat (EC_{50} = 276.5 nM, pEC_{50} = 6.56 ± 0.08, 109 ± 3% Glu Max) and human (EC_{50} = 246.0 nM, pEC_{50} = 6.61 ± 0.08, 96 ± 3% Glu Max), but with a modest disposition profile (vide infra).^{14–18} Thus, we pursued a multi-dimensional optimization plan (Fig. 2) to explore SAR around **4** in an attempt to improve disposition by identifying replacements for the weakly basic oxadiazole (effectively neutral with the pendant CF₃ moiety) with analogs **9** and the lipophilic 9H-xanthene moiety in analogs **8**. Once identified, optimal moieties would then be combined.

The synthesis was straightforward (Scheme 1). A variety of commercially available 2-amino-4-substituted oxazoles **10** were coupled under HATU conditions with a diverse array of carboxylic acids to provide analogs **8** in yields ranging from 18% to 54%. Sim-

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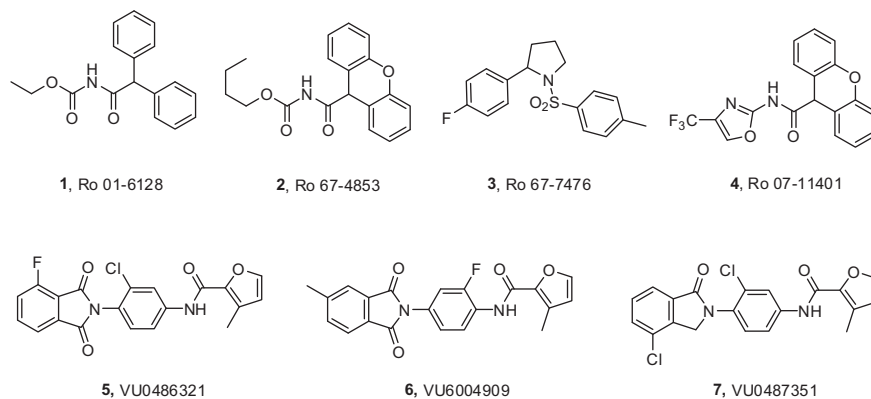


Figure 1. Structures of representative mGlu₁ PAMs **1–7**.

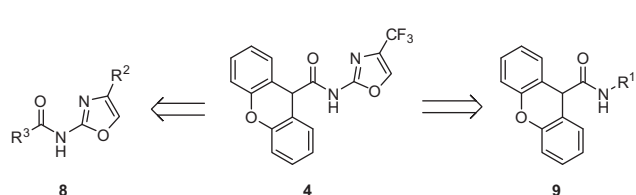


Figure 2. Chemical optimization plan to identify replacements for the lipophilic oxadiazole with analogs **9**, and the lipophilic 9H-xanthene with analogs **8**.

ilarly, the 9H-xanthene-9-carboxylic acid **11** was coupled under HATU conditions to a variety of 5- and 6-membered heterocyclic amines to deliver analogs **9** in 11–94% yields. For both series, many of the amines and acids coupled poorly due to a combination of steric and stereoelectronic effects, as well as a range of poor solubility.

In all, over 200 analogs of **8** and **9** were synthesized and triaged via a 10 μ M single point screen on human mGlu₁, using an EC₂₀ concentration of glutamate, prior to full concentration–response curves (CRCs) on both human and rat mGlu₁. Surprisingly, all analogs **8** (Fig. 3) were uniformly inactive mGlu₁ PAMs (no potentiation of an EC₂₀ of glutamate at a concentration of 10 μ M), indicating that the 9H-xanthene was a critical pharmacophore. The SAR was remarkably steep, and reminiscent of the steep SAR encountered with the non-MPEP, mGlu₅ PAM CPPHA,²⁰ wherein virtually any modification led to a complete loss of PAM activity. Obviously, these data cast doubt on the success of the campaign with analogs **9**, wherein the 9H-xanthene amide was held constant. While SAR once again was steep, active analogs did result (Table 1); however, functionalized pyrazoles, oxazoles, oxadiazoles, thiophenes, piperidines, azetidines, cycloalkyl, a structurally

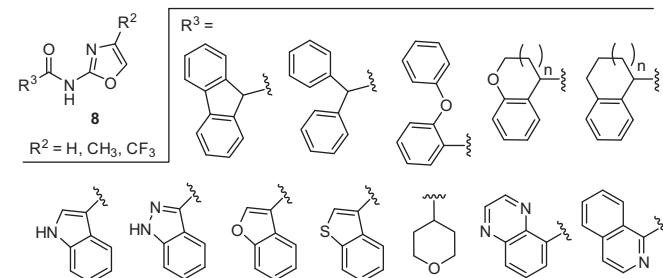


Figure 3. Representative 9H-xanthene amide replacement analogs **8** that are inactive mGlu₁ PAMs.

Table 1
Structures and activities for analogs **9**

9

Compd	R ¹	hmGlu ₁ EC ₅₀ (μ M) ^a [% Glu Max \pm SEM]	mGlu ₁ pEC ₅₀ (\pm SEM)
9a		0.71 [110 \pm 6]	6.14 \pm 0.12
9b		0.90 [89 \pm 3]	6.04 \pm 0.07
9c		1.4 [75 \pm 3]	5.85 \pm 0.07
9d		2.7 [101 \pm 7]	5.57 \pm 0.11
9e		>10 [54 \pm 7]	<5
9f		4.8 [96 \pm 11]	5.32 \pm 0.20
9g		>10 [68 \pm 8]	<5
9h		1.7 [58 \pm 3]	5.77 \pm 0.11

Scheme 1. Reagents and conditions: (a) R²CO₂H, DIEA, DMF, 60 $^{\circ}$ C, 18–54%; (b) H₂NR¹, HATU, DIEA, DCE, rt, 11–94%.

^a Calcium mobilization mGlu₁ assays, values are average of three ($n = 3$) independent experiments performed in triplicate.

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