

# 1*H*-Pyrrolo[3,2-*b*]pyridine GluN2B-Selective Negative Allosteric Modulators

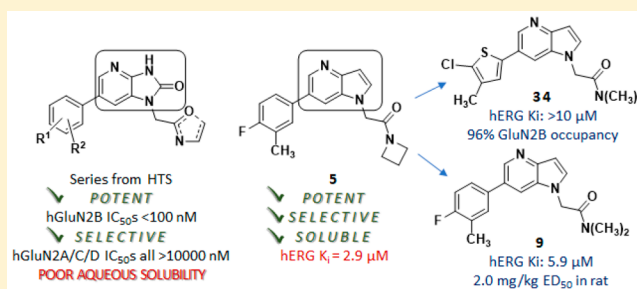
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## Supporting Information

**ABSTRACT:** Herein, we disclose a series of selective GluN2B negative allosteric modulators containing a 1*H*-pyrrolo[3,2-*b*]pyridine core. Lead optimization efforts included increasing brain penetration as well as decreasing cytochrome P450 inhibition and hERG channel binding. The series was also optimized to reduce metabolic turnover in human and rat. Compounds **9**, **25**, **30**, and **34** have good in vitro GluN2B potency and good predicted absorption, but moderate to high projected clearance. They were assessed in vivo to determine their target engagement. All four compounds achieved >75% receptor occupancy after an oral dose of 10 mg/kg in rat. Compound **9** receptor occupancy was measured in a dose–response experiment, and its ED<sub>50</sub> was found to be 2.0 mg/kg.

**KEYWORDS:** Azaindole, NR2B, Lead Optimization, Autoradiography



N-Methyl-D-aspartate (NMDA) receptors are a critical component of the functional machinery of the glutamatergic network in the mammalian central nervous system (CNS), operating as the mediators of fast excitatory neurotransmission. Channel activation occurs in a voltage-dependent manner following glutamate release and relief of a physiological Mg<sup>2+</sup> block of the channel, which initiates the ion flux. Once the channel is open, it is highly permeable to sodium, potassium, and calcium ions. Overactivation of the receptor results in excessive Ca<sup>2+</sup> influx and may lead to excitotoxicity.<sup>1,2</sup>

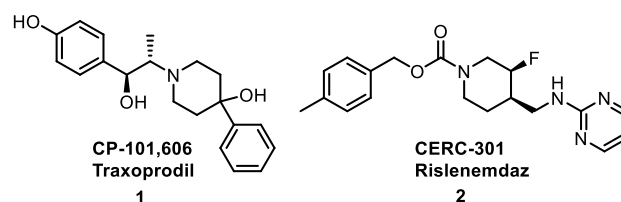
Structurally, an NMDA receptor functions as a tetramer comprising two GluN1 and two GluN2 subunits. GluN1 is expressed ubiquitously throughout the CNS, while GluN2 has four subunits A–D that have distinct localizations predicated upon expression patterns of the four genes that encode for them. Each of the GluN2 subunits exhibit unique biochemical properties and have distinct physiological roles.<sup>3</sup>

Clinical and preclinical research on NMDAR subunit selective modulators has overwhelmingly been focused on GluN2B functional antagonists.<sup>4</sup> Selective GluN2B modulation by a small molecule inhibitor is distinct from a channel blocking inhibitor in that selective GluN2B inhibitors have been found to utilize a GluN2B extracellular allosteric site positioned at the interface of GluN1 and GluN2B. The site is not present in subunits GluN2A, GluN2C, and GluN2D,<sup>5</sup> and inhibitors that bind at this site can have at least 1000-fold selectivity over GluN2A, GluN2C, and GluN2D subunits.<sup>6</sup>

Therapeutically, the NMDA receptors have received an increasing amount of attention since the end of the 20th

century, in part due to positive outcomes reported in patients diagnosed with major depressive disorder following the administration of NMDAR antagonists.<sup>7,8</sup> Proof-of-concept clinical trials have shown GluN2B-selective negative allosteric modulators (NAMs) to be therapeutic in treatment-resistant depression (TRD). Traxoprodil (CP-101,606; Chart 1) was shown to be efficacious in Phase II clinical trials for TRD, achieving a response in 60% of subjects diagnosed with TRD and in a current depressive episode.<sup>9</sup> QT interval prolongation was also observed, and trials with Traxoprodil were ceased.<sup>10,11</sup>

## Chart 1. GluN2B-Selective NAMs Assessed in Clinical Proof of Concept Trials for Depression



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Another GluN2B-selective NAM, CERC-301 (Chart 1) has been assessed in a pilot trial<sup>12</sup> and two Phase II trials in patients with major depressive disorder (MDD).<sup>13,14</sup> Both Phase II trials failed to meet their primary end points; although in the latter of the two Phase II studies it was noted that on Day 2 subjects averaged a 3.47-point improvement over placebo on the Hamilton Depression Rating Scale (HDMS). Based on the data reported for CP-101,606 and prior to the results reported with CERC-301, medicinal chemistry efforts toward the discovery of novel GluN2B-selective NAMs were initiated at Janssen with the goal of identifying an orally available small molecule with putative therapeutic use in patients with TRD.

A high throughput screen (HTS) of the Janssen global screening deck yielded a potent and selective series of GluN2B NAMs containing a 1,3-dihydro-2*H*-imidazo[4,5-*b*]pyridin-2-one core. Compound 3, for example, was found to have a human GluN2B IC<sub>50</sub> of 22 nM and was selective for the GluN2B-subunit (2A, 2C, 2D IC<sub>50</sub> all >10 μM). It also displaced [<sup>3</sup>H]-Ro 25-6981 with a K<sub>i</sub> of 13 nM, suggestive of allosteric binding. Compound 3 has adequate drug-like physicochemical properties with cLogP = 1.9, LE = 0.39, and MPO = 4.6. It was soon determined, however, that poor solubility and poor epithelial cellular uptake of dihydro-2*H*-imidazo[4,5-*b*]pyridin-2-one core compounds would be challenging to overcome. We attributed these undesirable properties to the cyclic urea that is contained within the central core. We were pleased to find that adjusting the ring system to a pyrrole-fused pyridine (1*H*-pyrrolo[3,2-*b*]pyridine) resulted in a more soluble, potentially more brain penetrant, GluN2B-active compound (4).

Initial optimization efforts focused on replacing the cyclopropyl amide. The hydrogen bond donor on the secondary amide was thought to be contributing to efflux mechanisms reported in a MDCK-MDR1 assay. After a slight tuning of the phenyl substituents and replacement of the secondary cyclopropyl amide with a tertiary amide (azetidine), compound 5 emerged (cLogP = 3.2, LE = 0.45) (Figure 1). It

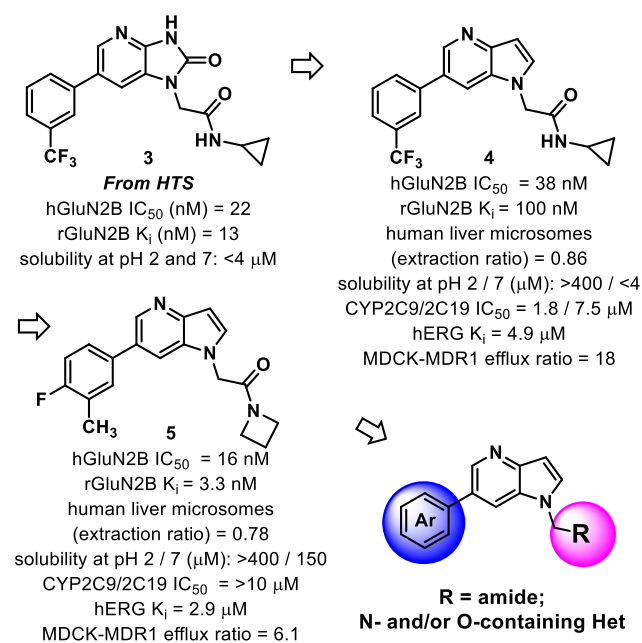


Figure 1. HTS hits and initial core optimization of fused pyridines.

had significantly improved rat GluN2B binding affinity compared to 4 and was selective over the 2A, 2C, and 2D isoforms (IC<sub>50</sub>s all >10 μM). Compound 5 also had good solubility and good cellular uptake but exhibited hERG channel binding, as measured by dofetilide binding. Even so, its overall favorable profile merited further structure–activity relationship (SAR) investigations. 4-Fluoro-3-methylphenyl was set as the aryl group left-hand side when conducting modifications to the amide piece of the molecule. This choice was made due to excellent GluN2B binding values obtained for preliminary compounds containing this moiety (compound 5 is one such example [K<sub>i</sub> = 3.3 nM]).

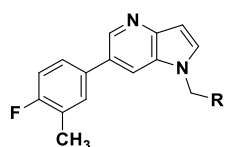
A representative set of 6-(4-fluoro-3-methylphenyl)-1*H*-pyrrolo[3,2-*b*]pyridines containing varied substituents tethered from the 1-position of the core is shown (i.e., *N*-pyrrolo-substituted, Table 1). The examples containing an amide were all found to have good or very good binding affinities, with rat K<sub>i</sub> values <100 nM (5–9, Table 1). The azetidines (5 and 8) and the pyrrolidine (7) had better human IC<sub>50</sub> and rat K<sub>i</sub> values than the piperidine amide (6). All of the examples have high or moderately high turnover in human and rat liver microsomes (LM). In general, as the number of saturated carbons on the amide increased, a decrease in the compounds' stability in LM was noted (5–9). As it follows, compound 9, which contains a dimethyl amide, was found to be the most stable LM. However, ER is still relatively high (0.7, human) and suggestive of a moderately high clearance of 9 in human.

Many examples were found to have cytochrome P450 (CYP) inhibition, particularly of 2C9 (6–8, 10, 12–15), but also significantly of 2C19 (7, 10, 12–15) and 1A2 (10, 12–15). In general, the examples containing a tethered amide were the weakest inhibitors; however, the examples containing larger amides did inhibit CYPs 2C9 and 2C19. Interestingly, the extent of 2C9 inhibition was found to be proportional to the size of the amide group.

Compounds 5, 8–10, and 13–15 displayed hERG binding. Further cardiovascular safety derisking of the compounds was not conducted since the compounds demonstrated additional disadvantages that we later found could be overcome (vide infra). For example, either a high efflux ratio (8) and/or CYP inhibition (8, 10, 13–15) were measured for the majority of the hERG binders.

Aromatic and aliphatic heterocycles were evaluated as amide replacements (10–16). Methyl 2-pyridine 10 had moderate GluN2B potency, but was unstable in LM, inhibited CYPs, and had hERG binding. Oxadiazoles 11–12 and an oxazole (13) had moderate GluN2B potency and low stability in LM. Compound 11 was unique to the set of methyl-2-heterocycles because it did not inhibit CYPs (up to 10 μM). Compound 11 also had no hERG binding, while 8–10 and 13–15 had moderate to high hERG binding with K<sub>i</sub> values < 6 μM. Compounds 14–16 are small cyclic ethers or aliphatic rings with moderate hGluN2B potency. The two examples analyzed in LM were found to be unstable (14–15).

The dimethylamide was the R-group used in the next round of SAR optimization (Table 2). It was chosen as a follow-up to 9, which had good GluN2B potency, low potential for drug–drug interaction (DDI), and among the lowest extraction ratios measured (Table 1). Several substituted 1*H*-pyrrolo[3,2-*b*]pyridines were made using the azetidine amide as well, but generally they were less stable and/or less potent than their corresponding dimethylamides (19 and 28 are the two azetidine examples included here).

**Table 1. SAR of 1-Position Nitrogen Substituents of 1*H*-Pyrrolo[3,2-*b*]pyridines**

cp d	R	IC <sub>50</sub> <sup>a</sup>	ratK <sub>i</sub> <sup>b</sup>	CYP450 IC <sub>50</sub> <sup>c</sup> (μM)	LM ER hu/r <sup>d</sup>	hERG K <sub>i</sub> <sup>e</sup>	Ef-flu x <sup>f</sup>
5		16	3.3	All >10	0.78 / 0.78	2.9	6.1
6		155	35	2C9 1.2	-- / >0.9	--	--
7		41 <sup>g</sup>	7.7	2C9 2.7 2C19 8.2	0.88 / 0.88	>10	6.7
8		30 <sup>h</sup>	10	2C9 8.7	0.74 / 0.66	2.9	9.6
9		25 <sup>i</sup>	11 <sup>j</sup>	All >10	0.7 / 0.65	5.9	5.4
10		409	66	1A2 5.7 2C19 4.7 2C9 3.2	0.94 / >0.90	1.4	--
11		47	88	All >10	0.87 / 0.65	>10	--
12		180	78	1A2 1.2 2C19 4.9 2C9 2.9	0.91 / 0.81	>10	--
13		91 <sup>g</sup>	117	1A2 1.5 2C19 0.6 2C9 7.1	0.93 / 0.85	0.9	1.1
14		67	75	1A2 0.9 2C19 2.5 2C9 3.4	>0.95 / >0.9	2.4	--
15		402	ND	1A2 0.4 2C19 0.3 2C9 2.6	>0.95 / >0.9	1.1	--
16		182	109	--	--	--	--

<sup>a</sup>IC<sub>50</sub> values are reported in nM and were determined by a calcium mobilization assay in inducible CHO T-Rex cells heterologously expressing the hGluN1a/GluN2B receptor. Values reported are the mean of three experiments unless otherwise stated; the SEM for their pIC<sub>50</sub> values are all ≤ ±0.2. <sup>b</sup>K<sub>i</sub> values are reported in nM and were determined using a radioligand competitive binding assay in rat cortex membranes using [<sup>3</sup>H]-5. Values reported are the mean of three experiments, unless otherwise stated. <sup>c</sup>CYP IC<sub>50</sub> values were obtained from human liver microsomes for six isoforms: 1A2, 2C19, 2C8, 2C9, 2D6, 3A4. <sup>d</sup>Stability in human and rat liver microsomes. Data reported as hepatic extraction ratio. <sup>e</sup>K<sub>i</sub> values are reported in μM and were determined using a radioligand ([<sup>3</sup>H]dofetilide) competitive binding assay to the hERG protein expressed in HEK-293 cells. <sup>f</sup>Efflux ratio was measured using MDCK cells transfected with the P-glycoprotein (MDR-1). The efflux ratio is reported as the ratio of velocities (B–A/A–B). <sup>g</sup>The mean of four experiments. <sup>h</sup>The mean of nine experiments. <sup>i</sup>The mean of five experiments. <sup>j</sup>The mean of ten experiments

One potential weak point identified in the preliminary in vitro characterization of dimethylamide **9** was its moderate level of hERG binding (IC<sub>50</sub> = 5.9 μM). It was therefore just serendipity that most examples in Table 2 had hERG IC<sub>50</sub> > 10 μM, despite them all containing the dimethylamide R-group (data not shown). The single exception is **19** (hERG IC<sub>50</sub> = 5.3), which is an azetidine amide.

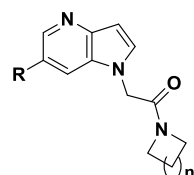
Compounds **17–23** demonstrate the basic SAR of a single trifluoromethyl substituent on the phenyl (**17–20**) or pyridine (**21–23**) R-group. Substitution *meta* to the ring attachment was preferred for good potency (**18–19**), although the presence of a pyridine ring reduced potency dramatically (**18** vs **21–23**). Three-position substitution on phenyl tended to generate compounds with good GluN2B potencies (**18–19**, **24–27**, **29**). The thiophenes (**30–34**) had moderate to excellent human IC<sub>50</sub> and rat K<sub>i</sub> values.

Compounds **18**, **21**, **24–27** and **29–35** were moderately stable in LM and generally had low DDI potential. Compound **27** additionally demonstrated excellent rat K<sub>i</sub> (5.1 nM), but because it also had a high efflux ratio (B–A/A–B = 16) it was deprioritized. Compounds **9**, **18**, **25**, **30**, and **34** were selected for in vivo studies, based on their robust rat K<sub>i</sub> values and good all-around in vitro ADME.

Rat pharmacokinetic (PK) experiments were conducted with **9**, **18**, **25**, **30** and **34** dosed at 1.0 mg/kg i.v. and 5.0 mg/kg p.o. Calculated bioavailability values ranged from 23 to >100%. The five compounds had moderate to high clearance (CL). It is of note that CL values for **18** and **34** were 167 and 90 mL/min/kg, respectively, higher than rat hepatic clearance (~70 mL/min/kg), suggesting the potential for extra-hepatic clearance mechanisms. Despite moderate to high CL values, compounds **9**, **25**, **30**, and **34** were selected for target engagement studies and brain level assessment in rats after oral dosing. They had excellent rat K<sub>i</sub> values, making them good candidates for rat receptor occupancy studies.

Target engagement was measured using ex vivo receptor autoradiography. Time dependency was evaluated after oral administration of a 10 mg/kg solution dose. The animals were sacrificed at different time points (0.25, 0.5, and 2 h) after drug administration. Brain sections were prepared and briefly incubated with the radiolabeled compound 3-[<sup>3</sup>H] 1-(azetidin-1-yl)-2-[6-(4-fluoro-3-methyl-phenyl)pyrrolo[3,2-*b*]pyridin-1-yl]ethanone (i.e., a tritiated version of **5**).<sup>15</sup> Thiophene **34** had the highest level of GluN2B occupancy (96%) despite its very high CL value. Brain concentrations of **34** at 30 min were high (802 ng/mL). Interestingly, free brain fraction measured in vitro for **24** in rat was only 0.75%, or 6 ng/mL (20 nM) at the 0.5 h time point. For comparison, rat brain protein binding for **9** and **30** were 96.06% and 98.27% bound, respectively, but they did not reach quite as robust occupancies as **34** (Table 3). Compound **9** GluN2B occupancy was steady over the 2 h time course, which is consistent with its moderately low rat CL (Table 3).

Compound **9** was selected for assessment in a dose–response ex vivo GluN2B occupancy study. It was prioritized in part due to the absence of a thiophene structural alert that is present in **30**.<sup>16</sup> We set 70% GluN2B occupancy as the level of target engagement desired for compound advancement. Dose–response analysis ex vivo GluN2B occupancy was measured at 30 min after oral dosing (from 0.01 to 30 mg/kg, (Figure 2)). Level of GluN2B occupancy at each dose was measured and ED<sub>50/70</sub> values were calculated. The measured ED<sub>50</sub> and EC<sub>70</sub> values for compound **9** were 2.0 and 3.4 mg/kg, respectively, in

Table 2. SAR of the 6-Aryl Position 1*H*-Pyrrolo[3,2-*b*]pyridines

For n=0, amide is dimethylamide

cp d	n	R	IC <sub>50</sub> <sup>a</sup>	rat K <sub>i</sub> <sup>b</sup>	CYP450 IC <sub>50</sub> (μM) <sup>c</sup>	LM ER hu/r <sup>d</sup>	B-A / A-B <sup>f</sup>
17	0		464	--	--	--	--
18	0		81	15	All >10	0.77 / 0.65	4.0
19	1		37	15	2C9 5.9	0.87 / 0.80	5.3
20	0		4460	--	--	--	--
21	0		145	--	All >10	0.70 / 0.50	--
22	0		520	--	--	--	--
23	0		362	--	--	--	--
24	0		44	41	All >10	0.75 / 0.73	5.0
25	0		33	12	All >10	0.66 / 0.68	4.1
26	0		17	25	All >10	0.73 / 0.72	5.9

cp d	n	R	IC <sub>50</sub> <sup>a</sup>	rat K <sub>i</sub> <sup>b</sup>	CYP450 IC <sub>50</sub> (μM) <sup>c</sup>	LM ER hu/r <sup>d</sup>	B-A / A-B <sup>f</sup>
27	0		18	5.1	All >10	0.60 / 0.55	16
28	1		375	79	--	--	--
29	0		15	54	All >10	0.54 / 0.56	5.9
30	0		47	16	All >10	0.75 / 0.48	3.5
31	0		54	36	All >10	0.75 / 0.67	--
32	0		29	52	All >10	0.68 / 0.67	3.5
33	0		32	27	All >10	0.71 / 0.63	3.4
34	0		21	4.3	2C19 8.4	0.82 / 0.74	3.5
35	0		23	46	2C9 9.3	0.77 / 0.68	--

<sup>a-j</sup> See Table 1.

Table 3. Rat PK and GluN2B Occupancy Data for 9, 18, 25, 30, and 34

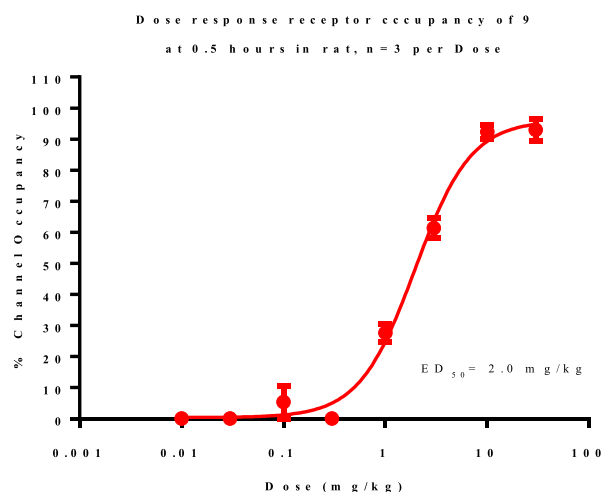
	i.v. PK (1 mg/kg)		p.o. PK (5 mg/kg)				GluN2B occupancy (10 mg/kg p.o.)				
	Cl (mL/min/kg)	V <sub>ss</sub> (L/kg)	C <sub>max</sub> (ng/mL)	AUC <sub>inf</sub> (h·ng/mL)	t <sub>max</sub> (h)	F (%)	% occupancy timecourse (h) <sup>a</sup>			C <sub>max-brain</sub> (ng/mg)	T <sub>max</sub> (h)
							0.25	0.5	2.0		
9	32	0.7	1380	1587	0.42	60	79	83	82	1426	0.5
18	167	1.6	227	372	0.50	64					
25	48	0.5	546	571	0.25	32	77	69	32	818	0.25
30	25	0.5	2330	4728	0.67	140	77	84	76	1972	0.5
34	90	1.4	184	226	0.30	23	80	96	41	802	0.5

<sup>a</sup>Ex vivo GluN2B labeling was expressed as the percentage of GluN2B labeling in corresponding brain areas of vehicle-treated animals.

rat. The plasma concentration associated with 70% GluN2B occupancy was 798 ng/mL total, or 31 ng/mL free for 9.

Finally, compounds 9, 18, 25, 30, and 34 were assessed qualitatively for their propensity to produce reactive metabolites in human liver microsomes supplemented with

glutathione (GSH) and NADPH. All compounds were identified by LC-MS/MS to form at least three GSH adducts with net additions to parent mass of +GSH - 2H + O, +GSH - 2H + O - CH<sub>2</sub>, and +GSH - 2H + 2O. Further studies



**Figure 2.** Dose versus receptor occupancy at 0.5 h in rat in a dose–response experiment of **9**.

would be necessary to better understand the nature and potential of these metabolites.

To summarize, compounds **9** and **30** have the highest plasma and brain exposures of the 1*H*-pyrrolo[3,2-*b*]pyridines assessed in vivo (Table 3). Compound **34**, although a very high clearing compound with relatively low plasma exposure in PK, has robust brain exposure. It is the only compound that displays full receptor occupancy, even though it is very highly protein bound (99.25% in the brain). Its advantage may have resulted from a superior rat  $K_i$  value. Unfortunately, GSH adducts have been identified for all of the promising 1*H*-pyrrolo[3,2-*b*]pyridines, warranting additional SAR investigations and in-depth metabolite characterization.

In conclusion, 1*H*-pyrrolo[3,2-*b*]pyridine GluN2B NAMs, which are chemically distinct from literature compounds, were identified from an HTS. Optimization efforts included improving solubility and permeability of the compounds, first by changing the core and then modifying the secondary amide, which eliminated two hydrogen bond donors. The series was also optimized for a lower metabolic turnover and CYP inhibition potential. Compounds **9**, **25**, **30**, and **34** had favorable rat metabolic stability and were taken on to rat PK and receptor occupancy studies. The four achieved >75% receptor occupancy after oral dosing of 10 mg/kg. In addition, a dose–response target engagement study was conducted to determine the  $ED_{70}$  and  $EC_{70}$  of **9** (3.4 mg/kg and 798 ng/mL, respectively). GSH adducts were identified for the compounds of the series, which would be prudent to further characterize prior to compound advancement.

## ■ ASSOCIATED CONTENT

### Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: [10.1021/acsmchemlett.8b00542](https://doi.org/10.1021/acsmchemlett.8b00542).

General experimental protocols and analytical data for compounds **5**–**35**, protocols for pharmacological assays, PPB data, time vs concentration curves for GluN2B occupancy of **9**, **30**, and **34**, GSH adduct metabolic scheme, and characterization data for [ $^3H$ ]-**5** (PDF)

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### Author Contributions

The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript.

### Notes

The authors declare no competing financial interest.

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## ■ ABBREVIATIONS

NMDA, *N*-methyl-*D*-aspartate; NAM, negative allosteric modulator; TRD, treatment-resistant depression; MDD, major depressive disorder; HDMS, Hamilton Depression Rating Scale; LE, ligand efficiency; MPO, multiparameter optimization; LM, liver microsomes; CYP, cytochrome P450; MDCK, Madin–Darby canine kidney epithelial cells; DDI, drug–drug interaction; PK, pharmacokinetic; CL, clearance; RO, receptor occupancy; GSH, glutathione

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