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Structure–activity relationship study of pyridazine derivatives as glutamate transporter EAAT2 activators

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ABSTRACT

Excitatory amino acid transporter 2 (EAAT2) is the major glutamate transporter and functions to remove glutamate from synapses. A thiopyridazine derivative has been found to increase EAAT2 protein levels in astrocytes. A structure–activity relationship study revealed that several components of the molecule were required for activity, such as the thioether and pyridazine. Modification of the benzylthioether resulted in several derivatives (**7-13**, **7-15** and **7-17**) that enhanced EAAT2 levels by >6-fold at concentrations <5 μ M after 24 h. In addition, one of the derivatives (**7-22**) enhanced EAAT2 levels 3.5–3.9-fold after 24 h with an EC₅₀ of 0.5 μ M.

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Glutamate is a major neurotransmitter in the mammalian central nervous system (CNS) and essential for normal brain function including cognition, memory, and learning. However, the extracellular concentration of glutamate must remain below excitotoxic levels ($\sim 1 \mu M$) to avoid overstimulation of glutamate receptors, leading to neuronal damage or death.¹ Excitotoxicity has been associated with multiple acute neurological conditions such as ischemic stroke, epilepsy, and trauma, as well as chronic adultonset neurodegenerative disorders such as Alzheimer's disease and amyotrophic lateral sclerosis (ALS).^{2–6} One potential approach to preventing excitotoxicity is to enhance glutamate reuptake. Excitatory amino acid transporter 2 (EAAT2) is the major glutamate transporter and functions to remove glutamate from synapses.⁷ An increase in EAAT2 protein expression and function may provide a means to prevent insufficient glutamate reuptake and consequently reduce neuronal damage.



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In an effort to identify small molecules that can increase EAAT2 protein expression, a high-throughput screen of approximately 140,000 compounds was previously conducted in our laboratories using a cell-based enzyme-linked immunosorbent assay.⁸ The hits identified from this screening provide starting points for further optimization in order to arrive at pharmacologically useful molecules for studying the role of EAAT2 in excitotoxicity induced neuronal injury and potentially as therapeutic agents.

The thiopyridazine **1** was confirmed to show a dose-dependent increase in EAAT2 protein levels after 24 h exposure. Herein, we report the structure–activity relationship (SAR) study of **1** for elevating EAAT2 protein levels.

Many of the pyridazine analogues utilized in the SAR study were prepared using the method depicted in Scheme 1. Treatment of ketone **2** with glyoxylic acid (**3**) and K₂CO₃ gave **4**, which was used directly without purification. It was allowed to react with hydrazine in acetic acid at 100 °C to yield the desired pyridazinone **5** as an off-white solid after recrystallization from ethyl acetate.⁹ Direct alkylation of **5** gave **8** in good yields. Intermediate **5** was also converted into pyridazinethione **6** in the presence of P₂S₅ in pyridine at 120 °C.¹⁰ Alkylation of **6** provided **7**, which could be subsequently oxidized to sulfone **9** with 3-chloroperoxybenzoic acid (m-CPBA) in CH₂Cl₂.¹¹

A series of additional analogues **14–17** was prepared using the methodology outlined in Scheme 2. Recently, 2-pyridyl *N*-methyliminodiacetic acid (MIDA) boronate (**10**) has been reported as an air-stable slow-release reagent with high cross-coupling efficiency even with heteroaryl chlorides.¹² Therefore, this material was used

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Scheme 1. Reagents and conditions: (a) K_2CO_3 , H_2O ; (b) NH_2NH_2 , AcOH, 100 °C (20%); (c) P_2S_5 , pyridine, 120 °C (80%); (d) R^2Br , K_2CO_3 , DMF (90%); (e) m-CPBA, CH₂Cl₂ (80%).

in palladium-mediated cross-couple reactions with heteroaryl chlorides **11–13** to obtain **14–16**, respectively. In the case of **13** transesterification occurred during the cross-coupling reaction to give the isopropyl ester **16**, which was then allowed to react with alkylamines in ethanol at 85 °C to generate amides **17**.

Finally, several additional analogues designed to evaluate replacement of the pyridazine in **1** with phenyl and pyridyl moieties were prepared using the methodology outlined in Scheme 3. Suzuki coupling of 2-bromopyridine (**18**) with various boronic acids afforded **19** and **20**.¹³ Treatment of **20** with P_2S_5 in pyridine at 120 °C provided **21** in 71% yield. Alkylation of **21** with 2-methylbenzyl bromide in the presence of K_2CO_3 in DMF generated **22**.

All of the derivatives of **1** were initially evaluated in PA-EAAT2 cells³ (a primary astrocyte line stably expressing EAAT2 mRNAs) following compound (10 μ M) incubation for 4 and 24 h before harvesting and measuring EAAT2 levels by Western blot analysis. The fold increases in EAAT2 protein levels relative to DMSO controls are reported (Tables 1–4).

Replacement of the 2-pyridyl with 3-pyridyl (**7-1**, **7-5** and **7-8**), 4-pyridyl (**7-2**, **7-9** and **7-11**) or phenyl (**7-6**) resulted in equal or reduced activity (Table 1). Removing either one or both of the nitrogen atoms in the pyridazine resulted in loss of activity (Table 2). Collectively, these results suggested that both the 2-pyridyl and the pyridazine were required for enhancing EAAT2 protein expression.



Scheme 2. Reagents and conditions: (a) Pd₂(dba)₃, XPhos, Cu(OAc)₂, K₂CO₃, DMF/*i*-PrOH (4/1), 100 °C; (b) H₂NR, ethanol, 85 °C (90%).



Scheme 3. Reagents and conditions: (a) Pd(PPh₃)₄, Na₂CO₃, CH₃CN/H₂O (1/1), 75 °C, 4-benzylthiophenylboronic acid (54%); (b) Pd(PPh₃)₄, Na₂CO₃, CH₃CN/H₂O (1/1), 75 °C, 6-hydroxypyridine-3-boronic acid pinacol ester (70%); (c) P₂S₅, pyridine, 120 °C (71%); (d) 2-methylbenzyl bromide, K₂CO₃, DMF (76%).

Table 1

Effects of the modified pyridyl and benzyl substituent on EAAT2 protein levels

Compound	\mathbb{R}^1	R ²	Fold increase for EAAT2 ^a	
			4 h	24 h
1	2-Py	2-Cl-6-F-Bn	na ^b	2.0 ± 0.8
7-1	3-Py	2-Cl-6-F-Bn	na	2.3 ± 0.8
7-2	4-Py	2-Cl-6-F-Bn	na	1.9 ± 0.9
7-3	4-Me-Ph	2-Cl-6-F-Bn	1.2 ± 0.2	2.3 ± 0.7
7-4	2-Py	2-Me-Bn	1.7 ± 0.1	3.5 ± 0.3
7-5	3-Py	2-Me-Bn	1.5 ± 0.4	na
7-6	Ph	2-Me-Bn	1.1 ± 0.1	2.0 ± 0.4
7-7	2-Py	2-Cl-Bn	2.1 ± 1.4	4.0 ± 0.3
7-8	3-Py	2-Cl-Bn	1.2 ± 0.4	1.2 ± 0.3
7-9	4-Py	2-Cl-Bn	1.5 ± 0.2	1.7 ± 0.3
7-10	2-Py	4-Me-Bn	2.1 ± 1.1	2.6 ± 0.4
7-11	4-Py	4-Me-Bn	1.3 ± 0.1	2.4 ± 0.6

^a Compound concentration of 10 μM.

^b na = not active.

Table 2

Effects of modifications of the pyridazine on EAAT2 protein levels

$X = N \qquad Y = Z \qquad X$							
Compound	Х	Y	Ζ	R	Fold increase for EAAT ^a		
					4 h	24 h	
7-4	S	Ν	Ν	2-Me-Bn	1.7 ± 0.1	3.5 ± 0.3	
22	S	С	Ν	2-Me-Bn	1.3 ± 0.6	na ^b	
8-1	0	Ν	Ν	2-Me-Bn	2.0 ± 0.9	2.5 ± 0.5	
15	0	Ν	С	2-Me-Bn	1.5 ± 0.3	na	
7-12	S	Ν	Ν	Bn	1.3 ± 0.2	3.0 ± 0.4	
19	S	С	С	Bn	1.2 ± 0.5	1.1 ± 0.4	

R

^a Compound concentration of 10 μM.

^b na = not active.

Next, the sulfur linker was examined (Table 3). Replacing the sulfur with oxygen generally yielded less active derivatives. An

Table 4

Table 3

Effects of modifications of the sulfur linker on EAAT2 protein levels

N=N X^R

Compound	Х	R	Fold increase for EAAT2 ^a	
			4 h	24 h
1	S	2-Cl-6-F-Bn	na ^b	2.0 ± 0.8
8-2	0	2-Cl-6-F-Bn	1.1 ± 0.3	2.9 ± 0.4
7-4	S	2-Me-Bn	1.7 ± 0.1	3.5 ± 0.3
8-1	0	2-Me-Bn	2.0 ± 0.9	2.5 ± 0.5
7-13	S	2,6-Di-Me-Bn	na	6.5 ± 1.0
8-3	0	2,6-Di-Me-Bn	2.1 ± 1.2	3.2 ± 1.4
7-14	S	1-(2-Cl-6-F-phenyl)ethyl	1.7 ± 0.9	5.5 ± 1.0
8-4	0	1-(2-Cl-6-F-phenyl)ethyl	1.6 ± 0.2	3.0 ± 0.7
7-15	S	2-(2-Cl-6-F-phenylethyl)	1.9 ± 0.7	6.7 ± 1.5
8-5	0	2-(2-Cl-6-F-phenylethyl)	2.2 ± 0.4	3.1 ± 1.2
7-16	S	2,4-Di-F-Bn	2.4 ± 1.5	3.8 ± 0.8
8-6	0	2,4-Di-F-Bn	na	2.3 ± 0.6
7-17	S	2,4-Di-Me-Bn	1.9 ± 0.8	6.4 ± 1.0
8-7	0	2,4-Di-Me-Bn	na	2.0 ± 0.9
6-1	S	Н	na	2.0 ± 0.8
5-1	0	Н	1.5 ± 0.5	1.6 ± 1.0
14	NH	2-Me-Bn	1.7 ± 1.1	2.6 ± 0.5
9-1	SO ₂	2-Cl-6-F-Bn	1.4 ± 0.3	2.8 ± 0.4
9-2	SO_2	2-Me-Bn	2.3 ± 0.4	1.1 ± 0.2
9-3	SO_2	2-Cl-Bn	1.6 ± 0.3	na
17-1	CONH	<i>n</i> -Pr	na	1.2 ± 0.2
17-2	CONH	2-Me-Bn	na	1.4 ± 0.2

^a Compound concentration of 10 µM.

^b na = not active.

analogue containing a NH linker (14) showed weaker activity. Likewise, oxidation of the sulfur to a sulfone (9-1, 9-2, and 9-3) was also detrimental. Finally, replacing the sulfur with an amide moiety (17-1 and 17-2) was not tolerated. Collectively, these results indicated that the sulfur linker was optimal.

Finally, the benzyl group was examined (Table 4). Compared to 1, 2-chloro, 3-chloro and 2,6-dichloro substitutions improved the potency by 2-fold (7-7, 7-18 and 7-22), but 2,3-dichloro, 2,4-di-

Table 5

Fold increase and EC₅₀ values of EAAT2 protein levels of selected analogues^a

Compound	R	Fold increase for EAAT2				EC ₅₀ (μM)	
		1 μM, 24 h		10 µM, 24 h			
		β-actin ^b	GAPDH ^c	β -actin ^b	GAPDH ^c	β -actin ^b	GAPDH ^c
7-4	2-Me-Bn	1.0 ± 0.8	1.0 ± 0.3	3.5 ± 0.3	3.1 ± 0.2	2.6 ± 0.8	1.8 ± 0.3
7– 7	2-Cl-Bn	1.1 ± 0.5	1.6 ± 0.4	4.0 ± 0.3	4.5 ± 0.6	3.3 ± 0.5	3.2 ± 0.5
7-13	2,6-Di-Me-Bn	2.0 ± 0.2	2.0 ± 0.4	6.5 ± 1.0	6.4 ± 0.9	2.6 ± 0.2	2.4 ± 0.4
7-14	1-(2-Cl-6-F-phenyl)ethyl	1.5 ± 0.5	1.8 ± 0.4	5.5 ± 1.0	5.2 ± 1.2	3.9 ± 0.5	2.4 ± 0.4
7-17	2,4-Di-Me-Bn	2.0 ± 0.3	2.2 ± 0.6	6.4 ± 1.0	6.0 ± 0.9	2.7 ± 0.3	1.8 ± 0.6
7-22	2,6-Di-Cl-Bn	3.0 ± 0.1	3.5 ± 0.0	3.9 ± 0.4	3.5 ± 0.5	0.5 ± 0.1	0.5 ± 0.0
7-16	2,4-Di-F-Bn	1.0 ± 0.2	1.0 ± 0.5	3.8 ± 0.8	3.7 ± 0.6	2.6 ± 0.2	2.6 ± 0.5
7-30	2,4,6-Tri-Me-Bn	1.5 ± 0.4	1.6 ± 0.4	4.7 ± 0.5	4.8 ± 0.6	2.4 ± 0.4	2.1 ± 0.4
7-15	2-(2-Cl-6-F-phenylethyl)	1.2 ± 0.3	1.2 ± 0.2	6.7 ± 1.5	5.3 ± 1.2	3.3 ± 0.3	3.3 ± 0.2
7-31	2-(2-Cl-phenylethyl)	1.0 ± 0.7	1.0 ± 0.1	4.8 ± 0.6	4.5 ± 0.6	4.7 ± 0.8	5.1 ± 0.1
7-33	1-(1,2,3,4-Tetrahydronaphthalenyl)	2.4 ± 0.3	2.8 ± 0.4	5.3 ± 1.1	4.5 ± 0.6	1.9 ± 0.3	1.7 ± 0.4
7-18	3-Cl-Bn	4.0 ± 0.4	4.2 ± 0.1	4.1 ± 0.5	3.8 ± 0.5	1.7 ± 0.4	0.5 ± 0.1

^a Data represent the mean values of four separate experiments.

 $^{\rm b}$ Data standardized by $\beta\text{-actin.}$

^c Data standardized by GAPDH.

Effects of the benzyl group on EAAT2 protein levels



Compound	R	Fold increase for EAAT2 ^a		
		4 h	24 h	
1	2-Cl-6-F-Bn	na ^b	2.0 ± 0.8	
7-7	2-Cl-Bn	2.1 ± 1.4	4.0 ± 0.3	
7-18	3-Cl-Bn	2.1 ± 0.3	4.1 ± 0.5	
7-19	2,3-Di-Cl-Bn	na	2.4 ± 0.4	
7-20	2,5-Di-Cl-Bn	1.4 ± 0.1	2.5 ± 0.4	
7-21	2,4-Di-Cl-Bn	1.8 ± 0.5	na	
7-22	2,6-Di-Cl-Bn	1.5 ± 0.8	3.9 ± 0.4	
7-23	2-F-Bn	2.2 ± 0.1	3.1 ± 0.2	
7-24	4-F-Bn	2.5 ± 0.6	2.6 ± 0.3	
7-16	2,4-Di-F-Bn	2.4 ± 1.5	3.8 ± 0.8	
7-25	2,6-Di-F-Bn	2.1 ± 1.4	2.2 ± 0.7	
7-26	2,4,6-Rri-F-Bn	1.6 ± 1.0	2.4 ± 0.6	
7-4	2-Me-Bn	1.7 ± 0.1	3.5 ± 0.3	
7-27	3-Me-Bn	2.0 ± 0.2	2.8 ± 0.5	
7-28	4-Me-Bn	2.1 ± 1.1	2.6 ± 0.4	
7-29	2,4-Di-Me-Ph	1.2 ± 0.3	2.2 ± 0.3	
7-17	2,4-Di-Me-Bn	1.9 ± 0.8	6.4 ± 1.0	
7-13	2,6-Di-Me-Bn	na	6.5 ± 1.0	
7-30	2,4,6-Rri-Me-Bn	1.7 ± 0.3	4.7 ± 0.5	
7-15	2-(2-Cl-6-F-phenylethyl)	1.9 ± 0.7	6.7 ± 1.5	
7-31	2-(2-Cl-phenylethyl)	1.7 ± 0.2	4.8 ± 0.6	
7-32	2-(2-Me-phenylethyl)	1.2 ± 0.3	2.8 ± 0.4	
7-33	1-(1,2,3,4-Tetrahydronaphthalenyl)	na	5.3 ± 1.1	
7-14	1-(2-Cl-6-F-phenyl)ethyl	1.7 ± 0.9	5.5 ± 1.0	
7-34	2-Cl-4-F-Ph	1.3 ± 0.1	2.5 ± 0.7	
7-35	2-Cl-4-F-Bn	1.6 ± 0.6	1.7 ± 0.4	
7-12	Bn	1.3 ± 0.2	3.0 ± 0.5	
7-36	Et	1.6 ± 0.7	1.9 ± 0.9	
6-1	Н	na	2.0 ± 0.8	

^a Compound concentration of 10 μM.

^b na = not active.

chloro and 2,5-dichloro analogues (**7-19**, **7-21** and **7-20**) were equivalent to **1**. In the case of fluorine substitutes, only the 2-fluoro

and 2,4-difluoro derivatives (7-23 and 7-16) demonstrated improved activity. 4-Fluoro, 2,6-difluoro, and 2,4,6-trifluoro analogues (7-24, 7-25, and 2-26) did not result in significant improvement. Replacement of the halogens with methyls (7-4, 7-13, 7-17, and 7-30) gave increased potency. Compounds containing 2-methyl substituted benzyl groups significantly increased EAAT2 protein level, where as the 3- or 4-methylbenzyl derivatives (7-27 and 7-28) were less potent. Increasing the tether length between the phenyl and the sulfur to an ethylene generally yielded more potent analogues (7-15 and 7-31 vs 1 and 7-7, respectively) with one noted exception (compare 7-32 to 7-4). However, truncating the linker resulting in a diarylthioether decreased potency (7-29 vs 7-17). Interestingly, adding substitutes on the carbon linker (7-14 and 7-33) increased activity about 3-fold. Finally, other changes gave compounds with essentially the same potency as 1 (7-12, 7-36, and 6-1).

In order to further characterize twelve of the most active analogues from the initial assessment, they were evaluated in a 6-point dose–response assay (0.1, 0.3, 1.0, 3.3, 10.0, and 30.0 μ M). The fold increases at 1 and 10 μ M, as well as the EC₅₀ values are summarized in Table 5. In addition, these values were determined relative to two unrelated proteins (e.g., β -actin and GAPDH). Several compounds (**7-13**, **7-17**, **7-31** and **7-18**) were able to increase EAAT2 levels >6-fold at 10 μ M compared to β -actin or GAPDH, but with EC₅₀ values in the 2.6–3.3 μ M range. Analog **7-22** demonstrated the lowest EC₅₀ of 0.5 μ M with a maximum increase of EAAT2 levels of 3- to 4-fold.

In summary, a series of pyridazine derivatives were synthesized and evaluated for increasing EAAT2 protein levels. Several compounds were found to significantly increase EAAT2 protein levels (>6-fold), such as **7-13**, **7-15** and **7-17**. Derivative **7-22** increased EAAT2 protein levels to a lower extent (3.5–3.9-fold), but with a lower EC₅₀ value (0.5 μ M). These compounds will provide useful tools for further assessing the role of glutamate excitotoxicity in cellular systems and potentially in animal models of acute and chronic neurodegeneration. In addition, these probe molecules may also be beneficial in determining the biological mechanisms for regulating EAAT2 levels.

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Supplementary data

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.bmcl.2011.08.009.

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