

Excitatory Amino Acid Receptor Ligands: Resolution, Absolute Stereochemistry, and Enantiopharmacology of 2-Amino-3-(4-butyl-3-hydroxyisoxazol-5-yl)propionic Acid

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(*RS*)-2-Amino-3-(4-butyl-3-hydroxyisoxazol-5-yl)propionic acid (Bu-HIBO, **6**) has previously been shown to be an agonist at (*RS*)-2-amino-3-(3-hydroxy-5-methylisoxazol-4-yl)propionic acid (AMPA) receptors and an inhibitor of CaCl₂-dependent [³H]-(*S*)-glutamic acid binding (*J. Med. Chem.* **1992**, *35*, 3512–3519). To elucidate the pharmacological significance of this latter binding affinity, which is also shown by quisqualic acid (**3**) but not by AMPA, we have now resolved Bu-HIBO via diastereomeric salt formation using the diprotected Bu-HIBO derivative **11** and the enantiomers of 1-phenylethylamine (PEA). The absolute stereochemistry of (*S*)-Bu-HIBO (**7**) (ee = 99.0%) and (*R*)-Bu-HIBO (**8**) (ee > 99.6%) were established by an X-ray crystallographic analysis of compound **15**, a salt of (*R*)-PEA, and diprotected **8**. Circular dichroism spectra of **7** and **8** were recorded. Whereas **7** (IC₅₀ = 0.64 μM) and **8** (IC₅₀ = 0.57 μM) were equipotent as inhibitors of CaCl₂-dependent [³H]-(*S*)-glutamic acid binding, neither enantiomer showed significant affinity for the synaptosomal (*S*)-glutamic acid uptake system(s). AMPA receptor affinity (IC₅₀ = 0.48 μM) and agonism (EC₅₀ = 17 μM) were shown to reside exclusively in the *S*-enantiomer, **7**. Compounds **7** and **8** did not interact detectably with kainic acid or *N*-methyl-D-aspartic acid (NMDA) receptor sites. Neither **7** nor **8** affected the function of the metabotropic (*S*)-glutamic acid receptors mGlu₂ and mGlu_{4a}, expressed in CHO cells. Compound **8** was shown also to be inactive at mGlu_{1α}, whereas **7** was determined to be a moderately potent antagonist at mGlu_{1α} (K_i = 110 μM) and mGlu_{5a} (K_i = 97 μM). Using the rat cortical wedge preparation, the AMPA receptor agonist effect of **7** was markedly potentiated by coadministration of **8** at 21 °C, but not at 2–4 °C. These observations together indicate that the potentiation of the AMPA receptor agonism of **7** by **8** is not mediated by metabotropic (*S*)-glutamate receptors but rather by the CaCl₂-dependent (*S*)-glutamic acid binding system, which shows the characteristics of a transport mechanism. After intravenous administration in mice, **7** (ED₅₀ = 44 μmol/kg) was slightly more potent than AMPA (**1**) (ED₅₀ = 55 μmol/kg) and twice as potent as Bu-HIBO (**6**) (ED₅₀ = 94 μmol/kg) as a convulsant, whereas **8** was inactive. After subcutaneous administration in mice, Bu-HIBO (ED₅₀ = 110 μmol/kg) was twice as potent as AMPA (ED₅₀ = 220 μmol/kg) as a convulsant. Since **7** and Bu-HIBO (EC₅₀ = 37 μM) are much weaker than AMPA (EC₅₀ = 3.5 μM) as AMPA receptor agonists *in vitro*, the presence of a butyl group in the molecules of Bu-HIBO and **7** seems to facilitate the penetration of these compounds through the blood–brain barrier.

Introduction

(*S*)-Glutamic acid, which is the main excitatory neurotransmitter in the central nervous system (CNS), and other excitatory amino acids (EAAs) operate through four different classes of receptors. In addition to the three heterogeneous classes of ionotropic EAA (iGlu) receptors, named *N*-methyl-D-aspartic acid (NMDA), (*RS*)-2-amino-3-(3-hydroxy-5-methylisoxazol-4-yl)propionic acid (AMPA), and kainic acid receptors,^{1–3} a heterogeneous class of metabotropic EAA (mGlu) receptors has been shown to have important functions in the central excitatory neurotransmission processes.⁴ It is now generally agreed that iGlu as well as mGlu recep-

tors play important roles in the healthy as well as the diseased CNS and that all subtypes of these receptors are potential targets for therapeutic intervention in a number of diseases.^{5,6}

EAA receptors are involved in the mechanisms of long-term potentiation, which is believed to play an important role in learning and memory functions, and the deficits of these functions in Alzheimer patients may, to some extent, be caused by hypoactivity at iGlu and/or mGlu receptors in the brain.^{7–10} There also is growing evidence of an implication of EAA receptors in schizophrenia.^{11,12} As in Alzheimer's disease (AD), the role of these receptors in the etiology and the clinical manifestations of schizophrenia is still very incompletely understood, but there is evidence to suggest that hypoactivity at EAA receptors also is a factor of impor-

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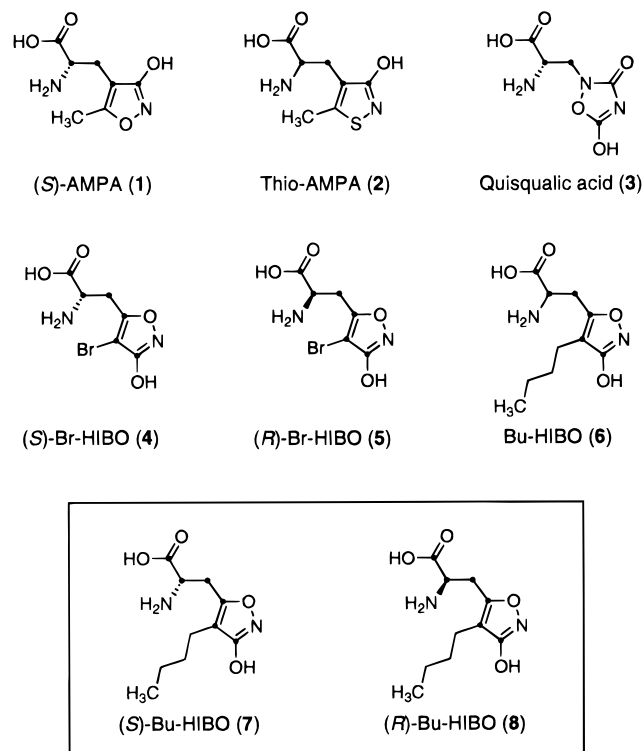


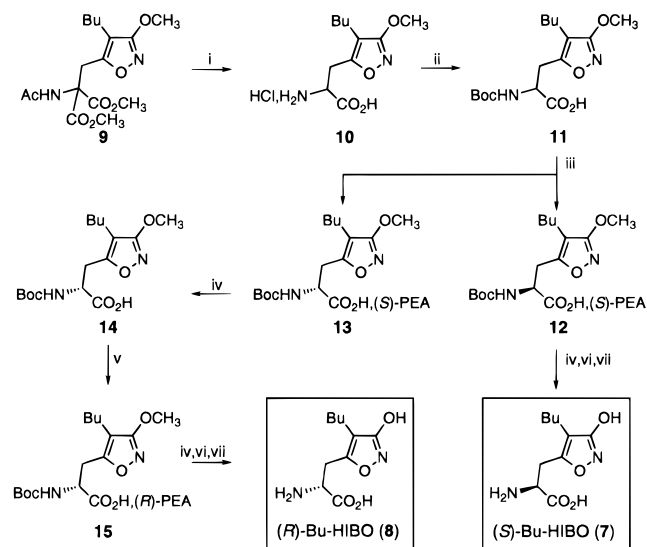
Figure 1. Structures of a number of heterocyclic analogues and homologues of glutamic acid.

tance in the latter CNS disorder.^{12–14} In AD as well as schizophrenia, agonists or partial agonists at EAA receptors, including AMPA receptors, may have therapeutic interest, and furthermore, compounds capable of enhancing EAA receptor activation are potential drugs in these disorders.¹⁵

A large number of specific and full AMPA receptor agonists have been developed,^{16,17} including (*S*)-AMPA (**1**)^{18,19} and (*RS*)-2-amino-3-(3-hydroxy-5-methylisothiazol-4-yl)propionic acid (Thio-AMPA, **2**)²⁰ (Figure 1). The naturally occurring amino acid, quisqualic acid (**3**) having the *S*-configuration, is an equally potent AMPA agonist, but this amino acid also interacts with other iGlu receptors,²¹ a number of mGlu receptor subtypes,²² and transport mechanisms associated with EAA-operated synapses.²³ An equally complex pharmacological profile is shown by (*S*)-2-amino-3-(4-bromo-3-hydroxyisoxazol-5-yl)propionic acid [(*S*)-Br-HIBO, **4**], which is an agonist at AMPA receptors²⁴ and an agonist/antagonist at receptors coupled to phosphoinositide (PI) hydrolysis.^{25,26} The AMPA receptor agonism of **4** is enhanced *in vitro*²⁷ and *in vivo*²⁴ by (*R*)-Br-HIBO (**5**), which is inactive at iGlu and mGlu receptors^{24,26} but is equipotent with **4** as an inhibitor of CaCl₂-dependent glutamic acid binding.²⁴ The mechanisms underlying this enhancement effect, of potential therapeutic interest, are very incompletely understood.^{24,27}

Since Bu-HIBO (**6**) shows an *in vitro* pharmacological profile²⁸ qualitatively and quantitatively very similar to that of Br-HIBO,^{21,29} and since the lipophilic butyl group of **6** may facilitate the penetration of this compound through the blood–brain barrier (BBB), the enantiomers of **6** were chosen as model compounds for studies of the mechanisms involved in the enhancement effect of **5** described above. We here report the resolution, determination of absolute stereochemistry, and

Scheme 1^a



^a Reagents: (i) 2 M HCl; (ii) (Boc)₂O, Et₃N; (iii) (*S*)-PEA, EtOH–Et₂O; (iv) extraction; (v) (*R*)-PEA, EtOH–Et₂O; (vi) 48% HBr; (vii) Et₃N.

molecular and *in vivo* pharmacology of (*S*)-Bu-HIBO (**7**) and (*R*)-Bu-HIBO (**8**).

Results

Resolution. The resolution of Bu-HIBO (**6**) was performed through a diastereomeric salt formation using the diprotected derivative **11**. This key intermediate was prepared from methyl 2-(acetamidomethyl)-3-(4-butyl-3-methoxyisoxazol-5-yl)-2-(methoxycarbonyl)propanoate (**9**)²⁸ through a selective deprotection of the dimethyl acetamidomalonate moiety followed by Boc protection of the primary amino group using di-*tert*-butyl dicarbonate (Scheme 1). A salt primarily consisting of compound **12** was obtained by adding (*S*)-1-phenylethylamine [(*S*)-PEA] to a solution of **11**. After three recrystallizations of the crude product, no change in the optical rotation of **12** was observed, and no sign of the diastereomeric impurity, compound **13**, could be detected in the ¹H NMR spectrum of compound **12** (δ_{OMe} of **12**, 3.90 ppm; δ_{OMe} of **13**, 3.86 ppm). To partly resolved **14**, obtained from the mother liquor of the first precipitation of compound **12**, was then added (*R*)-PEA following a procedure similar to that used for the preparation of **12**. Crude **15** precipitated, and after two recrystallizations no change in optical rotation was seen.

To establish the absolute stereochemistry of **7** and **8**, an X-ray crystallographic analysis of compound **15** was performed. Furthermore, circular dichroism (CD) spectra of **7** and **8** were recorded and compared to those of the respective Br-HIBO enantiomers, **4** and **5**,²⁴ and to those of the *S*- and *R*-forms of 2-amino-3-(3-hydroxyisoxazol-5-yl)propionic acid (HIBO).³⁰ The CD spectra of **7** and **8** (Figure 2) are mirror images, demonstrating that the two compounds are enantiomers. In agreement with earlier observations in acidic solution,^{31–33} the *S*-form **7** shows a relatively strong positive Cotton effect at 220 nm ($\Delta\epsilon = +0.2 \text{ m}^2/\text{mol}$). The structurally related 3-isoxazolol amino acids, (*S*)-Br-HIBO²⁴ and (*S*)-HIBO,³⁰ also show a positive Cotton effect at 220 nm of approximately the same magnitude (not illustrated).

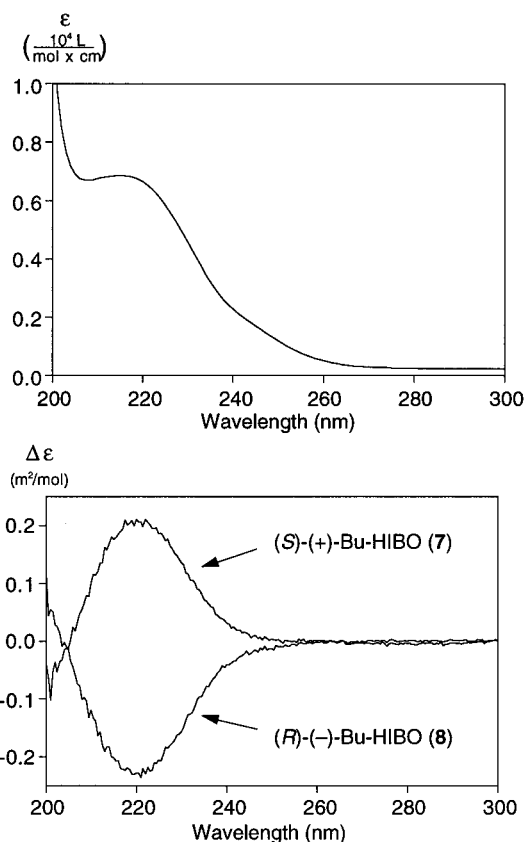


Figure 2. UV spectrum of (*S*)-Bu-HIBO (**7**) (top) and CD spectra of **7** and (*R*)-Bu-HIBO (**8**) (bottom) ($c = 0.16$ mM, 0.1 M HCl).

Chiral HPLC Analysis. The enantiomeric excess (ee) of (*S*)-Bu-HIBO (**7**) was determined to 99.0% using an (*S*)-proline ligand-exchange column. In agreement with earlier observations using this type of column,³⁴ (*R*)-Bu-HIBO (**8**) elutes prior to the *S*-enantiomer ensuring accurate quantification of the minor peak. For determination of the stereochemical purity of **8**, the Crownpak CR(-) column was used. As expected,³⁵ the elution order of the Bu-HIBO enantiomers on this chiral column was reversed as compared to that determined using the (*S*)-proline column. When analyzing a sample of **8** on the Crownpak CR(-) column, no trace of **7** could be seen. However, after spiking the sample with a small amount of Bu-HIBO, the *S*-enantiomer was detectable. The ee of the spiked sample of **8** was 99.6%, indicating that the isolated **8** had a stereochemical purity > 99.6% ee.

X-ray Crystallographic Analysis of Compound 15. The X-ray analysis established the absolute configuration of the *N*-protected α -amino carboxylate moiety of **15** to be *R*, determined relative to (*R*)-PEA used in the diastereomeric salt formation. A perspective drawing of the molecular structure of **15** with atomic labeling is shown in Figure 3. The molecular packing of **15** is depicted in Figure 4.

Bond lengths and angles of both the anion and cation of **15** are all within expected ranges,³⁷ the geometry of the 3-methoxyisoxazole moiety being consistent with the general pattern of 3-alkoxy- and 3-hydroxyisoxazoles (ref 38 and references therein). The isoxazole ring is planar within the limits of experimental error. The exocyclic oxygen and carbon atoms O2, C6, and C13 are

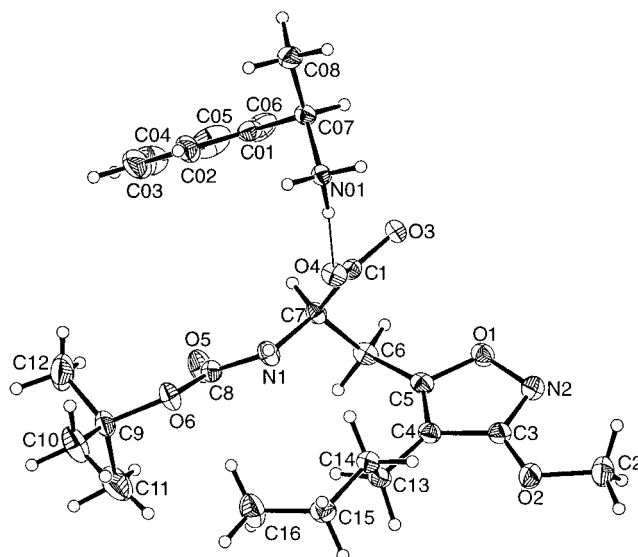


Figure 3. Perspective drawing³⁶ of compound **15** with atomic labeling of non-hydrogen atoms. Displacement ellipsoids of non-hydrogen atoms are shown at the 50% probability level. Hydrogen atoms are represented by spheres of arbitrary size. A hydrogen bond between the cation and anion is indicated with a thin line.

Table 1. Selected Torsion Angles (deg) for (*R*)-1-Phenylethylammonium (*R*)-3-(4-Butyl-3-methoxyisoxazol-5-yl)-2-[(*tert*-butyloxycarbonyl)amino]propionate (**15**)^a

Anion			
N2-C3-O2-C2	-4.3(2)	C7-N1-C8-O5	14.0(2)
C3-O2-C2-H2 _A	-175.0(1)	C7-N1-C8-O6	167.3(1)
C4-C5-C6-C7	72.0(2)	N1-C8-O6-C9	-176.2(1)
C1-C7-C6-C5	42.2(2)	O5-C8-O6-C9	2.5(2)
N1-C7-C6-C5	-80.3(2)	C8-O6-C9-C10	-178.0(9)
C5-C6-C7-H7	158.0(1)	C8-O6-C9-C11	64.7(2)
O4-C1-C7-C6	-125.6(1)	C8-O6-C9-C12	-60.4(2)
N1-C7-C1-O4	-0.8(2)	C3-C4-C13-C14	71.0(2)
N1-C7-C1-O3	178.7(1)	C4-C13-C14-C15	-174.1(1)
C6-C7-N1-C8	-88.2(2)	C13-C14-C15-C16	-65.5(2)
C1-C7-N1-C8	147.2(1)		
Cation			
C06-C01-C07-N01	162.2(2)	C02-C01-C07-H07	-2.0(1)
C06-C01-C07-C08	-60.0(2)		

^a Estimated standard deviations are given in parentheses.

at distances of $-0.022(2)$, $-0.046(3)$, and $0.023(2)$ Å, respectively, from this plane. The conformation of compound **15** is described by selected torsion angles listed in Table 1. An intramolecular hydrogen bond is found in the anion between the amide and carboxylate groups (N1-H1 \cdots O4) (Table 2).

The crystal packing (Figure 4 and Table 2) can be described as a collection of infinite anion/cation/anion "sandwiches" along the *a* axis. Within each "sandwich" the ammonium group of a cation is hydrogen-bonded to the carboxylate groups of three symmetry-related anions. Furthermore a few close contacts [C-H \cdots O(N)], which may be classified as hydrogen bonds,³⁹ are observed within each "sandwich". Between the "sandwiches" only one close contact (C-H \cdots O=C) is observed.

In Vitro Binding Affinities. The affinities of (*S*)-Bu-HIBO (**7**) and (*R*)-Bu-HIBO (**8**) at different recognition sites have been studied using [³H][3-(2-carboxypiperazin-4-yl)propyl]phosphonic acid ([³H]CPP),⁴⁰ [³H]-glycine,⁴¹ [³H]AMPA,⁴² [³H]kainic acid,⁴³ and [³H]-(*S*)-glutamic acid⁴⁴ (Table 3). Neither **7** nor **8** showed any detectable affinity ($IC_{50} > 100$ μ M) in the [³H]CPP, [³H]-

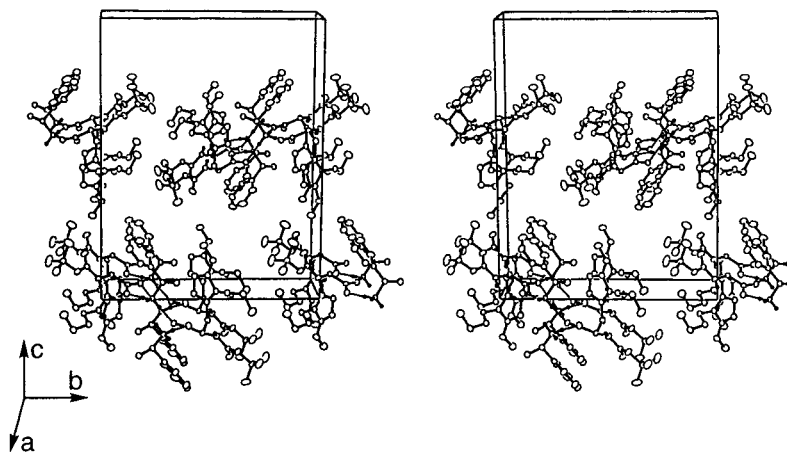


Figure 4. Stereoscopic view³⁶ of the packing of compound **15**. Displacement ellipsoids of non-hydrogen atoms are shown at the 50% probability level. For clarity only hydrogen atoms (represented by spheres of arbitrary size) bonded to nitrogen atoms are shown. Intermolecular hydrogen bonds are indicated by thin lines.

Table 2. Hydrogen Bond Geometries and Close Contacts (Å, deg) for (*R*)-1-Phenylethylammonium (*R*)-3-(4-Butyl-3-methoxyisoxazol-5-yl)-2-[(*tert*-butyloxycarbonyl)amino]propionate (**15**)^{a,b}

D-H...A	D-H	H...A	D...A	DHA
Intramolecular				
N1-H1...O4 ⁱ	0.91(2)	2.13(2)	2.578(2)	109(2)
Intermolecular				
N01-H01A...O4 ⁱ	0.98(2)	1.74(2)	2.718(2)	173(2)
N01-H01B...O3 ⁱⁱ	0.92(2)	1.89(2)	2.802(2)	171(2)
N01-H01C...O3 ⁱⁱⁱ	0.91(2)	1.91(2)	2.808(2)	167(2)
C2-H2A...O5 ^{iv}	1.02(2)	2.51(2)	3.054(2)	113(2)
C10-H10B...O5 ⁱⁱ	0.98(2)	2.52(2)	3.463(3)	162(2)
C07-H07...N2 ⁱⁱⁱ	0.97(2)	2.54(2)	3.392(2)	147(1)
C08-H08B...O4 ⁱⁱⁱ	0.98(2)	2.54(2)	3.444(2)	153(2)
C06-H06...O3 ⁱⁱⁱ	1.08(2)	2.55(2)	3.471(2)	143(2)
O1...O4 ^v			2.996(2)	

^a Symmetry codes: (i) x, y, z ; (ii) $1 + x, y, z$; (iii) $1/2 + x, 1/2 - y, -z$; (iv) $3/2 - x, -y, -1/2 + z$; (v) $-1 + x, y, z$. ^b Estimated standard deviations are given in parentheses.

Table 3. Receptor Binding and Electrophysiological Data [pIC₅₀ and pEC₅₀ values ± SEM ($n = 3$) in parentheses]

compound	IC ₅₀ (μM)		EC ₅₀ (μM)
	[³ H]AMPA	[³ H]-(<i>S</i>)-glutamic acid ^a	cortical wedge
AMPA	0.04 ^b (7.41 ± 0.02)	>100	3.5 ^b (5.48 ± 0.03)
quisqualic acid (3)	0.08 (7.10 ± 0.04)	0.025 (7.60 ± 0.05)	7.3 (5.15 ± 0.09)
Br-HIBO	0.60 ^c	0.20 ^c	23 (4.65 ± 0.07)
(<i>S</i>)-Br-HIBO (4)	0.34 ^c	0.22 ^c	21 (4.67 ± 0.024)
(<i>R</i>)-Br-HIBO (5)	32 ^c	0.15 ^c	>1000
Bu-HIBO (6)	1.8 (5.76 ± 0.05)	0.63 (6.20 ± 0.05)	37 (4.43 ± 0.02)
(<i>S</i>)-Bu-HIBO (7)	0.48 (6.33 ± 0.04)	0.64 (6.19 ± 0.05)	17 (4.77 ± 0.024)
(<i>R</i>)-Bu-HIBO (8)	>1000	0.57 (6.25 ± 0.01)	>1000

^a In the presence of 2.5 mM CaCl₂. ^b Reference 20. ^c Reference 24.

glycine, or [³H]kainic acid binding experiments. **8** did not show any affinity in [³H]AMPA binding (IC₅₀ > 100 μM), whereas **7** (IC₅₀ = 0.48 μM) was slightly more effective than Bu-HIBO (**6**) (IC₅₀ = 1.8 μM) as a displacer of [³H]AMPA binding. In the CaCl₂-dependent [³H]-(*S*)-glutamic acid binding assay,⁴⁴ **7** and **8** were approximately equipotent (Table 3), thus displaying no

stereoselectivity for this site. The effects of AMPA and **6–8** on synaptosomal (*S*)-glutamic acid uptake system(s) using [³H]-(*R*)-aspartic acid as a nonmetabolizable substrate were studied following an earlier described procedure.⁴⁵ In agreement with published data, the following IC₅₀ values for a series of reference compounds were determined: (*S*)-aspartic acid (10 μM), (*R*)-aspartic acid (13 μM), (*S*)-glutamic acid (18 μM), and (*R*)-glutamic acid (>300 μM). Under these test conditions, none of the compounds under study, AMPA and **6–8**, showed detectable inhibitory effects (IC₅₀ > 300 μM).

In Vitro Electrophysiology. Electrophysiological experiments were performed using the rat cortical wedge preparation.^{46,47} In accordance with the affinities in [³H]AMPA receptor binding, **7** was a selective and potent AMPA receptor agonist (EC₅₀ = 17 μM) (Figure 5A), whereas **8** was inactive (EC₅₀ > 1000 μM) when administered alone (Table 3). The depolarization induced by **7** was completely blocked by the AMPA receptor antagonist 6-nitro-7-sulfamoylbenzo[*l*]quinoxaline-2,3-dione (NBQX) (5 μM) and was insensitive to the NMDA antagonist CPP (5 μM). Coapplication of **8** (100 μM) and **7** at 21 °C resulted in a leftward parallel shift of the dose–response curve to **7** (Figure 5B) yielding an EC₅₀ value of 5 μM (pEC₅₀ = 5.27 ± 0.077) as compared to 17 μM for **7** alone [pEC₅₀ = 4.77 ± 0.024; paired *t*-test, $p < 1\%$ compared with **8** (100 μM) and **7**]. When these experiments were carried out at 2–4 °C, rather than at 21 °C, this enhancement of the depolarizing effect of **7** by coadministration of **8** was markedly reduced (Figure 5C). Thus, at 2–4 °C an EC₅₀ value of 15 μM (pEC₅₀ = 4.82 ± 0.084) for the depolarization of **7**, administered alone, was determined, whereas this value was only reduced to 12 μM (pEC₅₀ = 4.92 ± 0.032; paired *t*-test, nonsignificant) when **7** was coadministered with **8** (100 μM).

Application of **8** (100 μM), 10–15 min after termination of an application of **7** (50 μM), resulted in an NBQX-sensitive and CPP-insensitive response approximately 35% of the maximum response to AMPA. This priming of **8** (100 μM) was stronger when quisqualic acid (100 μM) was used as an agonist instead of **7**. Thus, pretreatment of the slices with 100 μM quisqualic acid resulted in an NBQX-sensitive and CPP-insensitive

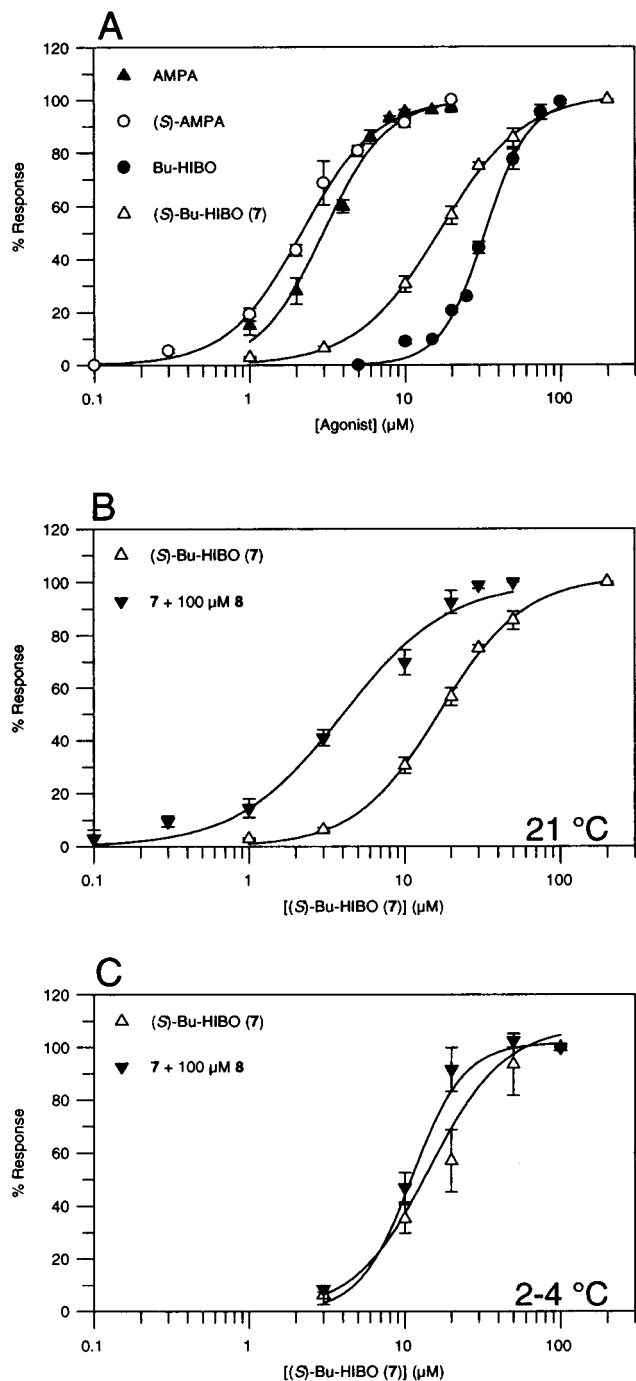


Figure 5. Dose–response curves obtained from the rat cortical wedge preparation for AMPA, (S)-AMPA (**1**), Bu-HIBO (**6**), and (S)-Bu-HIBO (**7**) at 21 °C (A) and for **7** alone and in the presence of a fixed concentration of (R)-Bu-HIBO (**8**) (100 μM) at 21 °C (B) and at 2–4 °C (C).

response close to the maximum response to AMPA (not illustrated).

Activity at mGlu Receptors in Vitro. Based on pharmacology, signal transduction pathway, and sequence homology, the eight mGlu receptor subtypes have been divided into three groups: mGlu₁/mGlu₅, mGlu₂/mGlu₃, and mGlu₄/mGlu₆/mGlu₇/mGlu₈.^{4,22} In this study, mGlu₁, mGlu₂, and mGlu₄ were chosen as representative subtypes for these three groups. Initially, we tested **7** and **8** alone and in the presence of (S)-glutamic acid (1 mM). As seen in Figure 6, none of the compounds displayed any activity as agonists or

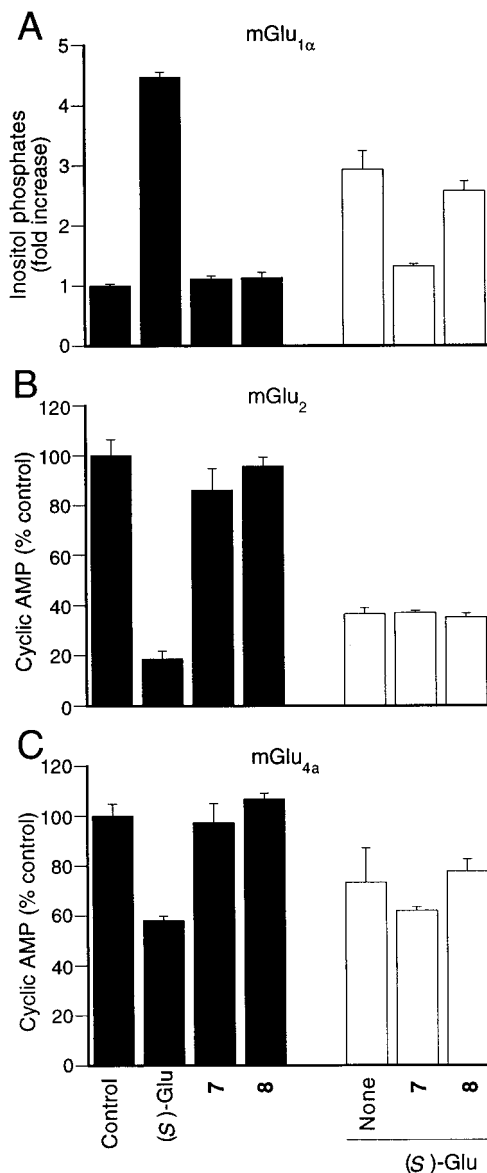


Figure 6. Agonist and antagonist activities at CHO cells expressing mGlu_{1α}, mGlu₂, and mGlu_{4a}. A: In agonist assays, mGlu_{1α}-expressing cells were incubated with ligands at a concentration of 1 mM for 20 min. For antagonist assays, cells were preincubated with ligand (1 mM) for 20 min and then incubated with ligand (1 mM) for 20 min in the presence of 20 μM (S)-Glu. Total IP formation was determined by ion-exchange assay and the fold increase in IP level calculated, compared to control cells (incubated in buffer only). B and C: In agonist assays, mGlu₂- and mGlu_{4a}-expressing cells were incubated with ligands (1 mM) for 10 min in the presence of 10 μM forskolin. In antagonist assays, cells were preincubated with ligand (1 mM) for 20 min and then incubated with ligand (1 mM) for 10 min in the presence of 20 μM (mGlu₂) or 50 μM (mGlu_{4a}) (S)-Glu. Cyclic AMP levels were measured by an RIA assay (Amersham) and set to 100% in control cells (incubated in buffer only). Data represent means (±SD) of representative experiments performed in triplicate.

antagonists at cell lines expressing mGlu₂ or mGlu_{4a}. Furthermore, **7** as well as **8** were inactive as agonists at the mGlu₁ receptor subtype. When tested as antagonists at the mGlu_{1α}-expressing cell line, **7** inhibited the response induced by (S)-glutamic acid, whereas **8** had no effect (Figure 6A). As seen in Figure 7, **7** was able to antagonize the response of (S)-glutamic acid in a dose-dependent manner with an IC₅₀ value of 245 μM

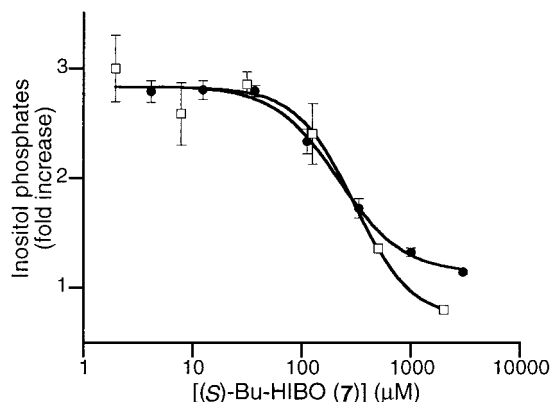


Figure 7. Concentration–response curves for the antagonist (*S*)-Bu-HIBO (**7**) in the presence of 20 or 10 μM (*S*)-Glu at CHO cells expressing mGlu_{1 α} (●) or mGlu_{5a} (□), respectively. For further details, see Figure 6.

Table 4. Convulsive Potencies in Mice

compound	ED ₅₀ ($\mu\text{mol/kg}$)	
	subcutaneous	intravenous
AMPA	220 ^a	55
Bu-HIBO (6)	110	94
(<i>S</i>)-Bu-HIBO (7)	nt	44
(<i>R</i>)-Bu-HIBO (8)	nt	>350

^a Reference 20. nt, not tested.

corresponding to a K_i value of 110 μM ($pK_i = 3.96 \pm 0.01$). **7** was also tested for antagonistic effect at mGlu_{5a}, which along with mGlu_{1 α} forms group I of mGlu receptors. As shown in Figure 7, **7** displayed very similar antagonistic effects at mGlu_{5a} as seen for mGlu_{1 α} with an IC₅₀ value of 305 μM corresponding to a K_i value of 97 μM ($pK_i = 4.02 \pm 0.04$).

In Vivo Pharmacology. The convulsant activities of AMPA, **6**, and enantiomerically pure **7** and **8** were studied in mice after subcutaneous and/or intravenous administration (Table 4). As compared to AMPA, **6** was the more potent convulsant after subcutaneous administration and the less potent convulsant after intravenous administration. The convulsive potency of **6** was unaffected by the route of administration, whereas AMPA was 4 times more potent when given intravenously than when given subcutaneously. Due to limited amounts of compounds, **7** and **8** were only tested intravenously. After this administration, **7** was approximately twice as potent as **6**, whereas **8** did not affect the animals even at the highest dose tested (80 mg/kg).

Discussion

The symptoms of Alzheimer and schizophrenic patients may, to some extent, reflect hypoactivity at iGlu and, perhaps, mGlu receptors.^{7–14} Rapid agonist-induced desensitization of iGlu receptors, notably AMPA receptors,^{48–50} and possible excitotoxic effects^{51,52} in vivo of full iGlu receptor agonists may make it difficult to administer such compounds safely to patients.^{10,15} On the other hand, compounds capable of sensitizing or of blocking desensitization of iGlu receptors may have therapeutic interest.⁵³

Whereas AMPA²⁹ and a number of AMPA analogues^{16,17} specifically activate AMPA receptors as full agonists, other structurally related AMPA agonists

derived from the ibotenic acid homologue, HIBO,¹⁶ show more complex pharmacological profiles. Thus, like (*S*)-AMPA (**1**),¹⁸ (*S*)-Br-HIBO (**4**)²⁴ (Figure 1) is a potent AMPA receptor agonist, but **4** also interacts with mGlu receptors coupled to PI hydrolysis.²⁶ Whereas (*R*)-AMPA is inactive,^{18,19} **4** and (*R*)-Br-HIBO (**5**) are equipotent inhibitors of CaCl₂-dependent [³H]-(*S*)-glutamic acid binding (Table 3), and **5** has been shown to enhance excitation produced by **4** and quisqualic acid (**3**), but not that by AMPA.^{24,27}

We have now resolved Bu-HIBO (**6**) and characterized the pharmacology of (*S*)-Bu-HIBO (**7**) and (*R*)-Bu-HIBO (**8**) (Figure 1) in vitro and in vivo. No stereoselectivity for CaCl₂-dependent [³H]-(*S*)-glutamic acid binding is observed for **7** and **8**. This is in analogy to the results previously obtained for **4** and **5**, both of which show slightly higher affinity for this binding site.²⁷ Enhancement of **7**-induced depolarizations of rat cortical neurons by the enantiomer **8**, showing no depolarizing effect per se, was observed, and this effect was subjected to further characterization. The pronounced enhancement effect was demonstrated using a fixed concentration of **8** (100 μM), which increased the potency of **7** more than 3-fold (Figure 5B). Comparison of the dose–response curves for **7** and the racemic compound **6** reveals that the enhancement effect is concentration-dependent, since at low concentrations **7** was more potent than the racemate **6**, whereas **6** and **7** became approximately equipotent at higher concentrations (Figure 5A). Furthermore, this enhancement effect of **8** turned out to be temperature-dependent, as lowering of the test temperature from 21 °C (Figure 5B) to 2–4 °C (Figure 5C) virtually eliminated the enhancement. This latter observation is in agreement with the general behavior of uptake/transport mechanisms.

Whereas the degree of selectivity of the interaction of (*S*)-Br-HIBO (**4**) with mGlu receptors coupled to PI hydrolysis is unclear,^{25,26} **7** was shown to antagonize exclusively the group I mGlu receptor, mGlu_{1 α} , without affecting detectably the group II and III mGlu receptors, mGlu₂ and mGlu_{4a}, respectively (Figure 6A). Like Br-HIBO,²¹ neither **7** nor **8** showed significant effects on the synaptosomal (*S*)-glutamic acid uptake system, which is labeled by [³H]-(*R*)-aspartic acid.

Since **8** does not interact detectably with the synaptosomal uptake system(s) generally accepted to be responsible for the removal of neurotransmitter (*S*)-glutamic acid from the synaptic cleft^{45,54} or with any of the iGlu or mGlu receptors studied, it seems unlikely that these synaptic mechanisms are involved in the enhancement phenomenon observed for **8** (Figure 5B). As tentatively proposed for the enantiomers of Br-HIBO,²⁷ it may be concluded that the CaCl₂-dependent (*S*)-glutamic acid binding system mediates this enhancement and that this system represents an energy-requiring presynaptic transport mechanism of as yet unknown physiological relevance. We envisage that quisqualic acid (**3**) and both enantiomers of Br-HIBO and Bu-HIBO are substrates for this transport mechanism. The AMPA receptor-active compounds **3**,¹⁶ (*S*)-Br-HIBO (**4**),²⁴ and (*S*)-Bu-HIBO (**7**) (Table 3) may be taken up in this transport system and/or associated vesicular structures and released by subsequent treatment with the receptor-inactive enantiomers (*R*)-Br-

HIBO (**5**) and (*R*)-Bu-HIBO (**8**). Such mechanisms may explain the priming effects of **3** and **7**, whereas **8** showed excitatory effects after pretreatment with **3** or **7**. This effect of **8** was more pronounced after pretreatment with **3**, which shows higher affinity for the CaCl₂-dependent [³H]-(*S*)-glutamic acid binding site as compared to **7** (Table 3).

We have previously shown that Thio-AMPA (**2**) is more active than AMPA after systemic administration in mice.²⁰ This observation probably reflects that **2** penetrates the BBB more easily due to the lipophilic character of the sulfur atom and the higher p*K*_a value of the isothiazol-3-ol nucleus of **2** (7.00) than that of the 3-isoxazolol nucleus of AMPA (5.12).²⁰ We anticipated that the presence of a butyl group in the molecules of **6–8** would facilitate the penetration of these compounds through the BBB. We have shown that although **6** and **7** are markedly weaker than AMPA as AMPA agonists (Table 3), **7** is slightly more potent and **6** slightly weaker than AMPA as convulsants after intravenous administration in mice. Compound **8**, which is assumed to penetrate the BBB to approximately the same extent as **6** and **7**, was inactive in this animal model (Table 4). In light of the proposed mechanism of action of **8**, this compound may, however, be able to produce gentle facilitation of the central (*S*)-glutamic acid neurotransmission system, but more detailed animal behavioral studies of compounds **7** and **8** must await the synthesis of further quantities of compound.

Experimental Section

Chemistry. General Procedures. Thin-layer chromatography (TLC) was performed on silica gel F₂₅₄ plates (Merck). Evaporations were performed under vacuum on a rotary evaporator at approximately 15 mmHg. Light petroleum was the fraction boiling at 40–60 °C. Melting points were determined in capillary tubes and are not corrected. Elemental analyses were performed by Mr. G. Cornali, Microanalytical Laboratory, LEO Pharmaceutical Products, Copenhagen, or by Mr. P. Hansen, Department of General and Organic Chemistry, University of Copenhagen, and are within ±0.4% of the theoretical values. ¹H NMR spectra were recorded on a Bruker AC-200 F spectrometer. TMS or 1,4-dioxane (δ 3.70) was used as internal standard for spectra recorded in CDCl₃ or D₂O, respectively. *J* values are given in Hz. Optical rotations were measured in thermostated cuvettes on a Perkin-Elmer 241 polarimeter. UV and CD spectra were recorded at room temperature in 1.0-cm cuvettes on a Perkin-Elmer Lambda 2 spectrophotometer and on a Jasco J-720 spectropolarimeter, respectively.

(*RS*)-2-Amino-3-(4-butyl-3-methoxyisoxazol-5-yl)propionic Acid Hydrochloride (10**).** A suspension of methyl 2-acetamido-3-(4-butyl-3-methoxyisoxazol-5-yl)-2-(methoxycarbonyl)propionate (**9**)²⁸ (2.3 g, 6.46 mmol) in aqueous hydrochloric acid (100 mL, 2 M) was refluxed for 24 h. The resulting solution was evaporated and reevaporated from toluene (tol). The residue was crystallized (AcOH–Et₂O) to give **10** (1.55 g, 86%, hygroscopic): mp 146–148 °C dec; ¹H NMR (D₂O) δ 4.28 (t, *J* = 6.1, 1H), 3.89 (s, 3H), 3.31 (d, *J* = 6.1, 2H), 2.25 (t, *J* = 7.4, 2H), 1.3–1.5 (m, 2H), 1.1–1.3 (m, 2H), 0.82 (t, *J* = 7.2, 3H). Anal. (C₁₁H₁₉ClN₂O₄·0.75H₂O) C, H, Cl, N.

(*RS*)-3-(4-Butyl-3-methoxyisoxazol-5-yl)-2-[(*tert*-butyloxycarbonyl)amino]propionic Acid (11**).** To a solution of **10** (1.0 g, 3.59 mmol) and Et₃N (1.75 mL, 12.6 mmol) in H₂O (30 mL) was added a solution of di-*tert*-butyl dicarbonate (1.05 mL, 4.56 mmol) in THF (30 mL). After the mixture was stirred for 72 h at room temperature, THF was evaporated, and the aqueous mixture was cooled, acidified with hydrochloric acid (pH ~ 3), and immediately extracted with AcOEt

(3 × 60 mL). The combined organic phases were dried, filtered, and evaporated. The residue was recrystallized (AcOEt–light petroleum) to give **11** (697 mg). The mother liquor furnished an additional 336 mg of recrystallized (cyclohexane) **11**. Total amount of **11** 1.03 g (84%): mp 88–91 °C; ¹H NMR (CDCl₃) δ 5.38 (d, *J* = 8.0, 1H), 4.59 (m, 1H), 3.96 (s, 3H), 3.24 (m, 2H), 2.22 (t, *J* = 7.4, 2H), 1.55–1.15 and 1.43 (m, 4H and s, 9H), 0.90 (t, *J* = 7.0, 3H). Anal. (C₁₆H₂₆N₂O₆) C, H, N.

(*S*)-1-Phenylethylammonium (*S*)-3-(4-Butyl-3-methoxyisoxazol-5-yl)-2-[(*tert*-butyloxycarbonyl)amino]propionate (12**).** To a solution of **11** (1.51 g, 4.41 mmol) in EtOH (5 mL) were added (*S*)-(-)-PEA (589 μL, 4.62 mmol) and subsequently Et₂O (5 mL). On standing at room temperature, **12** precipitated. The crystals formed were collected and dried in vacuo. The salt was recrystallized three times from a mixture of EtOH and Et₂O (1:1) to give **12** (690 mg, 34%): mp 174–176 °C; [α]_D²⁵ +22.4° (*c* = 1.04, MeOH); ¹H NMR (CDCl₃) δ 7.32 (m, 5H), 5.69 (d, *J* = 6.1, 1H), 4.28 (q, *J* = 6.8, 1H), 4.02 (q, *J* = 6, 1H), 3.90 (s, 3H), 3.07 (dd, *J* = 6, 14, 1H), 2.61 (dd, *J* = 6, 14, 1H), 2.19 (t, *J* = 7.1, 2H), 1.48 (d, *J* = 6.8, 3H), 1.45–1.15 and 1.40 (m, 4H and s, 9H), 0.90 (t, *J* = 6.9, 3H). Anal. (C₂₄H₃₇N₃O₆) C, H, N.

(*R*)-1-Phenylethylammonium (*R*)-3-(4-Butyl-3-methoxyisoxazol-5-yl)-2-[(*tert*-butyloxycarbonyl)amino]propionate (15**).** The mother liquor from a first crystallization of a batch of **12** was evaporated. The residue consisted of a 4:1 mixture of the diastereomeric salts **13** and **12**, respectively: ¹H NMR (CHCl₃) δ_{OMe} 3.86 (compound **13**), 3.90 (compound **12**). The mixture of the diastereomeric salts (437 mg, 0.94 mmol) was dissolved in CHCl₃ (40 mL), and H₂O (25 mL) and AcOH (3 mL) were added. The phases were separated, and the aqueous phase was extracted with CHCl₃ (40 mL). The combined organic phases were dried, filtered, evaporated, and reevaporated from tol. The partly resolved **14** crystallized by addition of light petroleum. The resulting crystals of **14** (238 mg, 0.69 mmol) were collected by filtration, dried in vacuo, and processed as described above for **12** using (*R*)-(+)-PEA (100 μL, 0.78 mmol). Crude **15** was recrystallized twice from EtOH and Et₂O (1:1) to give 180 mg of **15** (0.39 mmol, 57%): mp 175–176 °C; [α]_D²⁵ –22.0° (*c* = 1.00, MeOH). The ¹H NMR spectrum of **15** was virtually identical with that of **12**. Anal. (C₂₄H₃₇N₃O₆) C, H, N.

(*S*)-(+)-2-Amino-3-(4-butyl-3-hydroxyisoxazol-5-yl)propionic Acid Hemihydrate [(*S*)-Bu-HIBO, **7].** The salt **12** (650 mg, 1.40 mmol) was dissolved in CHCl₃ (50 mL) and treated twice with a mixture of H₂O (25 mL) and AcOH (4 mL). The organic phase was dried, filtered, evaporated, and reevaporated from tol. The residue was dissolved in 48% aqueous hydrobromic acid (10 mL) and refluxed for 30 min. After evaporation and reevaporation (twice) from H₂O, the residue was dissolved in H₂O (3 mL), and Et₃N was added until pH ~ 3–4. Compound **7** (131 mg, 39%) was precipitated by addition of EtOH (10 mL). Recrystallization from H₂O gave the hemihydrate of **7** (85 mg, 25%): mp 227 °C dec; ee = 99.0%; [α]_D²⁵ +19.0° (*c* = 0.58, 1 M HCl); Δε (220 nm) = +0.2 m²/mol; ¹H NMR [D₂O–CF₃COOH (19:1)] δ 4.41 (t, *J* = 6.3, 1H), 3.34 (d, *J* = 6.3, 2H), 2.26 (t, *J* = 7.7, 2H), 1.41 (m, 2H), 1.26 (m, 2H), 0.83 (t, *J* = 7.2, 3H). Anal. (C₁₀H₁₆N₂O₄·0.5H₂O) C, H, N.

(*R*)-(-)-2-Amino-3-(4-butyl-3-hydroxyisoxazol-5-yl)propionic Acid Hemihydrate [(*R*)-Bu-HIBO, **8].** Compound **8** was obtained from **15** (480 mg, 1.04 mmol) by a method analogous to that used for the preparation of **7**. Precipitated **8** (121 mg, 51%) was recrystallized from H₂O to give the hemihydrate of **8** (90 mg, 37%): mp 236 °C dec; ee > 99.6%; [α]_D²⁵ –21.1° (*c* = 0.52, 1 M HCl); Δε (220 nm) = –0.2 m²/mol. The ¹H NMR spectrum of **8** was virtually identical with that of **7**. Anal. (C₁₀H₁₆N₂O₄·0.5H₂O) C, H, N.

Determination of Enantiomeric Purity. Chiral HPLC using a Crownpak CR(-) column (4.0 × 150 mm; Daicel) was performed at 37 °C. The column was eluted with 0.4 mL/min aqueous HClO₄ (pH 2.0). The HPLC instrumentation used consisted of a Jasco 880 pump, a Rheodyne 7125 injector equipped with a 20 μL loop, and a Waters M480 UV detector,

set at 210 nm, attached to a Merck-Hitachi D-2000 chromatographic integrator. Chiral ligand-exchange HPLC was performed on a column (4.6 × 120 mm) containing Cu²⁺-complexed (*S*)-proline as chiral selector. The silica-based packing material was synthesized following a published procedure.³⁴ The column was connected to an HPLC instrumentation consisting of a Waters M510 pump, a Waters UK6 injector, and a Waters M480 UV detector, set at 210 nm, attached to a Merck-Hitachi D-2000 chromatographic integrator. The column was thermostated at 50 °C and eluted at 1.0 mL/min with aqueous KH₂PO₄ (50 mM, pH 4.6)/2-propanol (4:1). All compounds injected were dissolved in H₂O (approximately 0.2 mg/mL).

X-ray Analysis of (*R*)-1-Phenylethylammonium (*R*)-3-(4-Butyl-3-methoxyisoxazol-5-yl)-2-[(*tert*-butyloxycarbonyl)amino]propionate (15). Crystal data: C₂₄H₃₇N₃O₆, *M_r* = 463.57; colorless needles (2-propanol); orthorhombic, space group *P*2₁2₁2₁ (No. 19); *a* = 6.446(2) Å, *b* = 17.506(3) Å, *c* = 22.465(6) Å; *V* = 2535(1) Å³, *Z* = 4, *D_c* = 1.215 Mg m⁻³; *F*(000) = 1000, $\mu(\text{Cu } K\alpha) = 0.714 \text{ mm}^{-1}$, *T* = 122 K; crystal size 0.11 × 0.13 × 0.30 mm.

Data Collection and Processing. Diffraction data were collected at low temperature (122 ± 0.5 K) on an Enraf-Nonius CAD-4 diffractometer with graphite monochromated Cu *K* α radiation ($\lambda = 1.54184 \text{ \AA}$).⁵⁵ Unit cell dimensions were determined by least-squares refinement of the setting angles of 20 reflections (θ : 37–42°). Intensity data were collected in the $\omega/2\theta$ scan mode ($\theta \leq 75^\circ$, *hkl*, *h-k-l*, and *-h-k-l*). The intensities of five reflections monitored every 10⁴ s showed an average decrease in the intensities of 8%. The orientation of the crystal was checked every 600 reflections. Data were reduced using the data reduction package DREADD.^{56,57} The data were corrected for Lorentz and polarization effects and decay, but no absorption correction was made. A total of 8331 reflections were averaged (*R*_{int} = 0.016) according to the point-group symmetry 222, resulting in 5196 unique reflections. The number of reflections with $F^2 > 2\sigma(F^2)$ is 4801.

Structure Determination and Refinement. The structure was solved by direct methods using SHELXS-86.⁵⁸ Full-matrix least-squares refinements (SHELXL-93)⁵⁹ were performed on F^2 minimizing $\sum w(F_o^2 - F_c^2)^2$ with anisotropic displacement parameters for the non-H atoms. The positions of all the H atoms were located in subsequent difference electron density maps. Positional and isotropic displacement parameters were refined for all the H atoms, except for the methyl H atoms of the *tert*-butyl group, since a couple of these H atoms tended to give unreasonable geometries. The H atoms of the *tert*-butyl group were included in idealized positions (riding model) with fixed isotropic displacement parameters equal to 1.5 *U*_{eq} for the parent C atom.

The refinement (5196 all unique data, 413 parameters) converged at *R_F* = 0.0381, *wR_F²* = 0.0837; $w^{-1} = \sigma^2(F_o^2) + (0.0421P)^2 + 0.3141P$, where $P = (F_o^2 + 2F_c^2)/3$; *S* = 1.033. The final shift/esd was 0.000. Min and max values in the final difference electron density map were -0.20 and 0.19 e Å⁻³. The *R*-configuration of the protected α -amino acid moiety was determined relative to the known (*R*)-PEA. The absolute configuration was supported by refinement of the Flack absolute structure parameter ϵ in the final structure factor calculation $\epsilon = 0.00(14)$.⁶⁰ The large standard deviation of ϵ is due to the atoms present in the compound (C, H, N, O), which produce minor anomalous scattering. Complex neutral-atom scattering factors were taken from ref 61 and used as contained in SHELXL-93.⁵⁹

Receptor Binding. Affinity for NMDA, AMPA, and kainic acid receptor sites was determined using the ligands [³H]-CPP,⁴⁰ [³H]glycine,⁴¹ [³H]AMPA,⁴² and [³H]kainic acid,⁴³ respectively. The membrane preparations used in all four receptor binding experiments were prepared according to the method of Ransom and Stec.⁶² [³H]-(*S*)-Glutamic acid receptor binding was performed as described previously.⁴⁴

In Vitro Electrophysiology. A rat cortical wedge preparation for determination of EAA-evoked depolarizations described by Harrison and Simmonds⁴⁶ was used in a modified version, as described previously.⁴⁷

Cell Culture. The Chinese hamster ovary (CHO) cell lines expressing mGlu_{1a}, mGlu₂, mGlu_{4a}, and mGlu_{5a} were maintained as described previously.^{63–66} The cell lines were grown in a humidified 5% CO₂/95% air atmosphere at 37 °C in DMEM containing a reduced concentration of (*S*)-glutamine (2 mM), 1% (*S*)-proline, penicillin (100 U/mL), streptomycin (100 mg/mL), and 10% dialyzed fetal calf serum (all GIBCO, Paisley, Scotland). Two days before assay 1.8 × 10⁶ cells were divided into the wells of 24-well plates.

Measurement of PI Hydrolysis and Cyclic AMP Formation. PI hydrolysis was measured essentially as described previously.^{67,68} Briefly, the cells were labeled with [³H]inositol (2 $\mu\text{Ci/mL}$) 24 h prior to the assay. For agonist assay, the cells were incubated with ligand dissolved in PBS–LiCl for 20 min, and agonist activity was determined by measurement of ³H-labeled mono-, bis-, and trisinositol phosphates by ion-exchange chromatography. For antagonist assay, the cells were preincubated with the ligand dissolved in PBS–LiCl for 20 min prior to incubation with ligand and 20 μM (mGlu_{1a}) or 10 μM (mGlu_{5a}) (*S*)-glutamic acid. The antagonist effect was then determined as the inhibitory effect of the (*S*)-glutamic acid-mediated response. *K_i* values were calculated from IC₅₀ values by the use of the Cheng–Prusoff equation.⁶⁹ The assay of cyclic AMP formation was performed as described previously.^{67,68} Briefly, the cells were incubated for 10 min in PBS containing the ligand, 10 μM forskolin, and 1 mM 3-isobutyl-1-methylxanthine (IBMX) (both Sigma Chemicals, St. Louis, MO). The agonist activity was then determined as the inhibitory effect of the forskolin-induced cyclic AMP formation. For antagonist assay, the cells were preincubated with ligand dissolved in PBS containing 1 mM IBMX for 20 min prior to a 10-min incubation in PBS containing the ligand (1 mM), (*S*)-glutamic acid [20 μM (mGlu₂) or 50 μM (mGlu_{4a})], forskolin (10 μM), and IBMX (1 mM). All experiments were performed at least twice in triplicate.

Synaptosomal (*S*)-Glutamic Acid Uptake. The experiments were carried out essentially as described.⁴⁵ In short: Rat cortical synaptosomes were prepared from male Sprague–Dawley rats (200–300 g) using a previously described method.⁷⁰ The final synaptosomal pellet was suspended in assay buffer at a final concentration of 2 mg of original tissue/mL. Uptake of [³H]-(*R*)-aspartic acid was measured in a buffer containing 128 mM NaCl, 10 mM glucose, 5 mM KCl, 1.5 mM NaH₂PO₄, 1.77 mM CaCl₂, 1 mM MgSO₄, and 10 mM Tris (pH 7.4). Following a 5-min preincubation of the synaptosomes, the assay was initiated by adding 200 μL of membrane suspension to microtiter plates containing 25 μL of test substance and 25 μL of 3 μM [³H]-(*R*)-aspartic acid [3 μM nonradioactive (*R*)-aspartic acid plus 30 nM [³H]-(*R*)-aspartic acid]. Total binding was determined in the presence of buffer, and nonspecific binding was determined in the presence of 1 mM (*R*)-aspartic acid. Following 2 min of incubation, the reaction mixture was filtered through a 96-well Whatman GF/B filter using the Packard filtermate system and rinsed with 3 × 250 μL of ice-cold buffer. Radioactivity was determined using the Packard topcount system.

In Vivo Convulsant Activity. Convulsant activity was determined in male mice (NMRI/BOM, SPF, Bomholtgaard, Denmark) weighing 24–26 g. The mice were kept in groups of 15 in plastic cages (40 × 25 × 15 cm). The compounds were dissolved in saline and administered subcutaneously or intravenously. AMPA was given intravenously as a hydrobromide salt in doses of 1.25, 5, and 20 mg/kg. Bu-HIBO (**6**), also as a hydrobromide salt,²⁸ was given in doses of 10, 20, and 40 mg/kg (subcutaneous) or 1.25, 5, 20, and 40 mg/kg (intravenous). (*R*)-Bu-HIBO (**8**) was given in doses of 40 and 80 mg/kg (intravenous) and (*S*)-Bu-HIBO (**7**) in doses of 5, 20, and 40 mg/kg (intravenous). Each group consisted of 5–10 mice. The animals were observed for 30 min for the presence or absence of clonic/tonic convulsions. ED₅₀ values were calculated by log–probit analysis.

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Supporting Information Available: For (*R*)-1-phenylethylammonium (*R*)-3-(4-butyl-3-methoxyisoxazol-5-yl)-2-[(*tert*-butyloxycarbonyl)amino]propionate (**15**), tables listing final atomic coordinates, equivalent isotropic (non-H atoms) or isotropic (H atoms) displacement parameters, anisotropic displacement parameters for the non-H atoms, and selected bond lengths and angles (6 pages); a list of structure factors (10 pages). Ordering information is given on any current masthead page.

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