

Design and synthesis of a new series of 4-alkylated 3-isoxazolol GABA_A antagonists

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Abstract

A number of analogues of the low-efficacy partial GABA_A agonist 5-(4-piperidyl)-3-isoxazolol (4-PIOL), in which the 4-position of the 3-isoxazolol ring is substituted by different groups, were synthesized and tested as GABA_A receptor ligands. While alkyl and benzyl substitution provided affinities and antagonist potencies comparable to those of 4-PIOL, diphenylalkyl and naphthylalkyl substitution resulted in marked increase in both affinity and potency. The 2-naphthylmethyl and the 3,3-diphenylpropyl analogues showed antagonist potencies comparable or markedly higher than that of the standard antagonist SR 95531. Molecular modeling studies exposed a large cavity in the vicinity of the 4-position of 4-PIOL, in which there seems to be additional sites for specific receptor interactions.

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4-Aminobutyric acid (GABA), the major inhibitory neurotransmitter in the central nervous system (CNS), inhibits neuronal activity through two classes of GABA receptors, ionotropic GABA_A and GABA_C receptors and metabotropic GABA_B receptors [1,2]. Besides playing an important role in central transmission processes, these receptors, especially GABA_A receptors, are believed to be involved in certain neurological and psychiatric disorders and are therapeutic targets in certain diseases [3,4]. In order to pharmacologically characterize this receptor class, a number of GABA_A ligands (Fig. 1), such as the highly selective GABA_A agonists isoguvacine, muscimol [5] and 4,5,6,7-tetrahydroisoxazolo[5,4-*c*]pyridin-3-ol (THIP) [5,6] and the low-efficacy partial agonist 5-(4-piperidyl)-3-isoxazolol (4-PIOL) [7] have been developed. 2,3'-(Carboxypropyl)-3-amino-6-(*p*-methoxyphenyl)pyridazinium bromide (SR 95531) [8], now used as a standard

antagonist for GABA_A receptors, represents another structural class of ligands.

Introduction of alkyl groups into the 4-position of the 3-isoxazolol ring of muscimol severely inhibits interaction with the GABA_A receptor recognition site. This is illustrated by the very low affinity for GABA_A receptor of 4-Me-muscimol (IC₅₀ > 100 nM) [9] compared with that of muscimol (IC₅₀, 0.006 μM) [10]. In contrast, GABA_A recognition site tolerates introduction of alkyl groups into the 4-position of the 3-isoxazolol ring of 4-PIOL [11,12]. These structure–activity relationships indicate that the binding modes of the GABA_A agonists, muscimol and THIP, and in particular 4-PIOL, are different.

There is strong evidence that an arginine residue at GABA_A receptor recognition site is directly involved in the binding of the anionic part of the receptor ligand [13–15]. Based on this observation, a hypothesis has been proposed concerning the binding modes of the bioactive conformations of muscimol and 4-PIOL as illustrated in Fig. 2 [11]. In these binding modes, the two 3-isoxazolol rings do not overlap. This means that the 4-position of the 3-isoxazolol ring in muscimol does not

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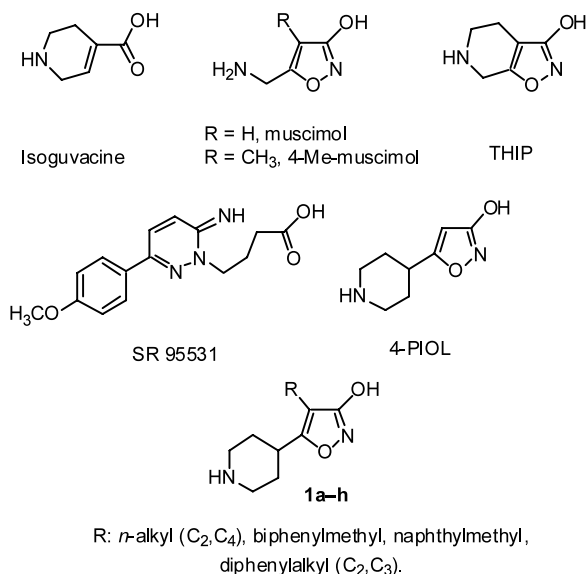


Fig. 1. Structures of GABA_A agonists isoguvacine, muscimol, 4-methylmuscimol and THIP, the GABA_A antagonist SR 95531, the low-efficacy partial GABA_A agonist 4-PIOL and a general structure of the new isoxazolols **1a–1h**.

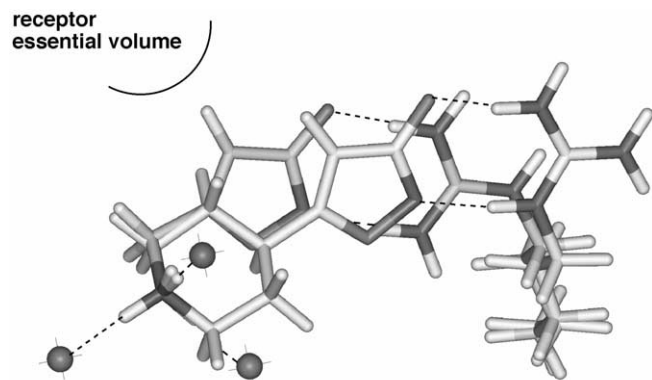


Fig. 2. A pharmacophore model for GABA_A receptor agonists showing the proposed binding modes of muscimol (dark grey bonds) and 4-PIOL (light grey bonds) and their interactions with two different conformations of an arginine residue. The dark grey spheres indicate sites to which the ammonium group in muscimol interacts via hydrogen bonds.

correspond to the 4-position in the 3-isoxazolol ring of 4-PIOL during interaction of muscimol and 4-PIOL with the receptor recognition site.

A number of analogues of 4-PIOL have been synthesized with substituents in the 4-position of the 3-isoxazolol ring in order to further investigate the steric tolerance of this position [12]. Substituents of different size and structural flexibility such as alkyl, phenylalkyl, diphenylalkyl and naphthylalkyl were explored.

Pharmacological characterization of the synthesized compounds was carried out using receptor binding assays and by electrophysiological experiments using whole-cell patch-clamp techniques. The results from these studies are exemplified in Table 1 [12]. Substitution

Table 1
Receptor binding and in vitro electrophysiological data

R		[³ H]muscimol binding ^a K _i (μM)	Electrophysiology ^b IC ₅₀ (μM)
4-PIOL	H	9.1	110
1a	CH ₃ CH ₂ -	6.3	26
1b	CH ₃ (CH ₂) ₃ -	7.7	3.0
1c		3.8	4.0
1d		0.10	0.48
1e		0.049	0.37
1f		0.4	0.71
1g		0.36	0.81
1h		0.068	0.02
SR 95531		0.074	0.24

^a Standard receptor binding on rat brain synaptic membranes, *n* = 3. ^b Two electrode voltage-clamp recordings on human α₁β₂γ_{2S} GABA_A receptors expressed in oocytes.

of the 4-position with alkyl or benzyl groups, as for compounds **1a–1c**, resulted in affinities and potencies comparable with those of 4-PIOL. Interestingly, introduction of more bulky groups such as biphenylalkyl, diphenylalkyl and naphthylalkyl groups, as exemplified by the 4-biphenylmethyl, 1-naphthylmethyl, 2-naphthylmethyl, 2,2-diphenylethyl and the 3,3-diphenylpropyl analogues (**1d–1h**), is not only tolerated but resulted in a marked increase both in affinity and potency, as shown in Table 1.

Using cultured cerebral cortical neurones in the electrophysiological testing, the pharmacology of the 4-PIOL analogues in the absence or presence of the specific GABA_A receptor agonist isoguvacine was studied [12]. The results demonstrated that the structural modifications led to a change in the pharmacological profile of the compounds from moderately potent low-efficacy partial GABA_A receptor agonist activity to a potent and selective antagonist effect. The 2-naphthylmethyl and the 3,3-diphenylpropyl analogues, **1e** and **1h**, respectively, showed an antagonist potency comparable to or markedly higher than that of the standard GABA_A antagonist SR 95531. Only the methyl and ethyl analogues retained detectable ability to induce an agonist effect shown on cultured cerebral neurones [12].

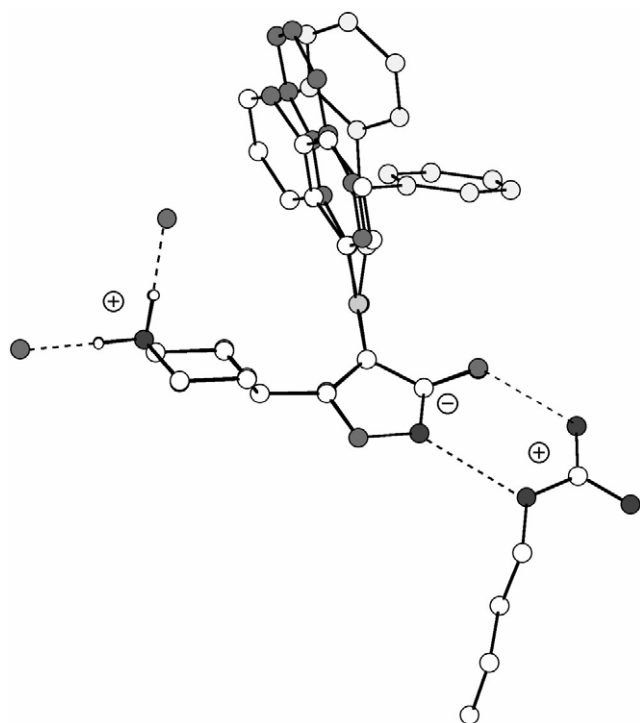


Fig. 3. Superimposition of compounds **1d** (unfilled atoms), **1e** (dark grey atoms) and **1h** (light grey atoms) in their proposed bioactive conformations.

These structure–activity studies seem to support the proposed hypothesis concerning the distinct binding mode of 4-PIOL, implying that the 4-position in 4-PIOL does not correspond to the 4-position in muscimol (Fig. 2). Thus, a cavity of considerable binding capacity seems to exist at the 4-PIOL recognition site of GABA_A receptor. Molecular modeling studies of the two high-affinity compounds containing a 2-naphthylmethyl and a 3,3-diphenylpropyl substituent, **1e** and **1h**, respectively, and the less active 1-naphthylmethyl analogue, **1d**, indicate that this proposed binding cavity may be exploited in two directions (Fig. 3). In both these positions, an aromatic ring seems to be highly favourable for the receptor affinity.

This series of alkylated and arylalkylated 4-PIOL analogues has provided new information on the agonist

and competitive antagonist binding site in GABA_A receptor and is going to be exploited in further development of the pharmacophore model. In addition, these compounds could serve as useful tools for studies of GABA_A receptor structure and mechanisms.

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