

GABA_A agonists and partial agonists: THIP (Gaboxadol) as a non-opioid analgesic and a novel type of hypnotic

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Received 14 June 2004; accepted 30 June 2004

Abstract

The GABA_A receptor system is implicated in a number of central nervous system (CNS) disorders, making GABA_A receptor ligands interesting as potential therapeutic agents. Only a few different classes of structures are currently known as ligands for the GABA recognition site on the hetero-pentameric GABA_A receptor complex, reflecting the very strict structural requirements for GABA_A receptor recognition and activation. A large number of the compounds showing agonist activity at the GABA_A receptor site are structurally derived from the GABA_A agonists muscimol, THIP (Gaboxadol), or isoguvacine, which we developed at the initial stage of the project. Using recombinant GABA_A receptors, functional selectivity has been shown for a number of compounds, including THIP, showing subunit-dependent potency and maximal response. The pharmacological and clinical activities of THIP probably reflect its potent effects at extrasynaptic GABA_A receptors insensitive to benzodiazepines and containing $\alpha_4\beta_3\delta$ subunits. The results of ongoing clinical studies on the effect of the partial GABA_A agonist THIP on human sleep pattern show that the functional consequences of a directly acting agonist are distinctly different from those seen after administration of GABA_A receptor modulators, such as benzodiazepines. In the light of the interest in partial GABA_A receptor agonists as potential therapeutics, structure-activity studies of a number of analogues of 4-PIOL, a low-efficacy partial GABA_A agonist derived from THIP, have been performed. In this connection, a series of GABA_A ligands has been developed showing pharmacological profiles ranging from low-efficacy partial GABA_A agonist activity to selective antagonist effect. © 2004 Elsevier Inc. All rights reserved.

Keywords: GABA_A receptor; Agonist; Partial agonist; Functional selectivity; THIP (Gaboxadol); 4-PIOL; Muscimol; Clinical studies; Novel hypnotic effects; Non-opioid analgesia

1. GABA receptors: multiplicity, structure and function

The discovery of GABA in the early fifties and the identification of the alkaloid bicuculline [1] and its quaternized analogue bicuculline methochloride (BMC) [2] as competitive GABA antagonists in central nervous system (CNS) tissues initiated the pharmacological characterization of GABA receptors. The subsequent design of isoguvacine, 4,5,6,7-tetrahydroisoxazolo[5,4-c]pyridin-3-ol (THIP, Gaboxadol) [3] (Fig. 1) and piperidine-4-sulphonic

acid (P4S) [4] as a novel class of specific GABA agonists further stimulated studies of the pharmacology of the GABA receptors.

The GABA analogue baclofen did, however, disturb the picture of a uniform class of GABA receptors. Baclofen, which was designed as a lipophilic analogue of GABA capable of penetrating the blood-brain barrier (BBB), is an antispastic agent, but its GABA agonistic effect could not be antagonized by BMC. In the early eighties, Bowers and co-workers demonstrated that baclofen, or rather (*R*)-(–)-baclofen, was selectively recognized as an agonist by a distinct sub-population of GABA receptors, which was named GABA_B receptors [5]. Cloning of the GABA_BR1 and GABA_BR2 receptors and the identification of functional heterodimeric GABA_B receptors have greatly stimulated GABA_B receptor research [6,7]. The “classical”

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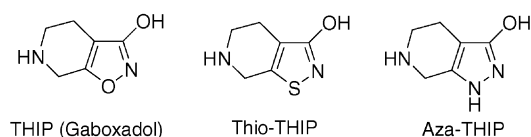


Fig. 1. Structures of the GABA_A agonist/partial agonist THIP, the GABA_A partial agonist/antagonist Thio-THIP, and the GABA_C antagonist Aza-THIP.

BMC-sensitive GABA receptors were designated GABA_A receptors. This receptor classification represents an important step in the development of the pharmacology of GABA.

During this period, the exploration of the GABA receptors was dramatically intensified by the observation that the binding site for the benzodiazepines (BZDs) [8,9] was associated with the GABA_A receptors. After the cloning of a large number of GABA_A receptor subunits, this area of the pharmacology of GABA continues to be in a state of very rapid development [10].

In connection with the design of conformationally restricted analogues of GABA, another “disturber of the peace” appeared on the GABA scene, namely *cis*-4-aminobut-2-enoic acid (CACA). This compound and the structurally related GABA analogue, *cis*-2-aminomethylcyclopropanecarboxylic acid (CAMP) are GABA-like neuronal depressants, which are not sensitive to BMC, and they bind to a class of GABA receptor sites, which neither recognize isoguvacine nor (*R*)-(-)-baclofen. The phosphinic acid analogue of isoguvacine, TPMPA [11] and, more recently, 4,5,6,7-tetrahydropyrazolo[5,4-*c*]pyridin-3-ol (Aza-THIP) [12] (Fig. 1) have been shown to be selective antagonists at GABA_C receptors [13]. These receptors have been named GABA_C receptors or non-GABA_A, non-GABA_B (NANB) receptors for GABA [11,13].

The GABA_A receptor complex is a pentameric structure formed by co-assembly of subunits from seven different classes (α_{1-6} , β_{1-3} , γ_{1-3} , δ , ϵ , θ , ρ_{1-3}). The five subunits, which in native receptors usually are constituted by two α , two β and a γ , δ or ϵ subunit, are situated in a circular array surrounding a central chloride-permeable pore [14].

Several studies have demonstrated the involvement of GABA and GABA_A receptors in diseases like seizures, depression, anxiety and sleep disorders [14,15]. GABA and other directly acting GABA_A receptor agonists (GABA-mimetics) bind specifically to a recognition site located at the interface between an α and a β subunit [16–18] whereas the classical BZDs, such as diazepam, flunitrazepam, as well as the novel BZD ligands with a non-benzodiazepine structure as for example zaleplon, zolpidem, zopiclone and indiplon, bind to an allosteric site located at the interface between an α and a γ subunit [16,18,19].

The quantitatively most predominant GABA_A receptor subunit combination throughout the brain is comprised of $\alpha_1\beta_{2/3}\gamma_2$ subunits, primarily synaptically located [20–23].

However, recently, much attention has been focused on the less abundant but ostensibly functionally important receptor subtypes with a more restricted location and specialized function. Thus, the primarily extrasynaptically located $\alpha_4\beta_3\delta$ receptors, which are found in high concentrations in hippocampus and thalamus but also in neocortex [22,24], seem to play a key role in sleep. Results from our ongoing research projects have suggested that these receptors are the main target for THIP (Gaboxadol), which at present is subjected to clinical studies as a hypnotic.

Whereas Gaboxadol behaves as a partial agonist ($E_{\max} = 70\%$) with a fairly low potency ($EC_{50} = 238 \mu\text{M}$) at $\alpha_1\beta_3\gamma_{2S}$ receptors expressed in *Xenopus* oocytes [17,25], it acts as a highly potent ($EC_{50} = 1.3 \mu\text{M}$) full agonist in the rat cortical wedge preparation [26]. Based on a series of in vitro and in vivo pharmacological studies [27,28] it is hypothesized that Gaboxadol and the BZDs interact with two distinct receptor populations: the benzodiazepines with synaptically located $\alpha_1\beta_3\gamma_{2S}$ receptors and Gaboxadol with extrasynaptically located $\alpha_4\beta_3\delta$ receptors. Hence, the effects exerted at these separate populations will sum in an additive rather than a supra-additive manner. A functional selectivity for $\alpha_4\beta_3\delta$ receptors at which Gaboxadol acts as a highly potent agonist with an E_{\max} of 165%, also explains that apparent potent and efficacious behaviour of Gaboxadol in the rat cortical wedge preparation.

$\alpha_4\beta_3\delta$ receptors have proven difficult to express in recombinant system and therefore only a few studies investigating the pharmacology of this receptor construct have been published. However, recently a novel murine L(tk) cell line stably expressing $\alpha_4\beta_3\delta$ receptors has been extensively used to study the basic pharmacology of this receptor subtype [29,30]. Other groups have published results from more specific studies in *Xenopus* oocytes expressing $\alpha_4\beta_{2/3}\delta$ receptors [31,32]. More recently, a broad spectrum of agonists have been found to behave as full and partial agonists in $\alpha_{1-6}\beta_x\gamma_2$ receptors expressed in *Xenopus* oocytes [17,25,33].

2. GABA_A agonists

The basically inhibitory nature of the central GABA neurotransmission prompted the design and development of different structural types of GABA agonists. The conformational restriction of various parts of the molecule of GABA and bioisosteric replacements of the functional groups of this amino acid have led to a broad spectrum of specific GABA_A agonists. Some of these molecules have played a key role in the development of the pharmacology of the GABA_A receptor, or rather receptor family [34,35].

Muscimol, a constituent of the mushroom *Amanita muscaria*, has been extensively used as a lead for the design of different classes of GABA analogues (Fig. 2). The 3-isoxazolol carboxyl group bioisostere of muscimol

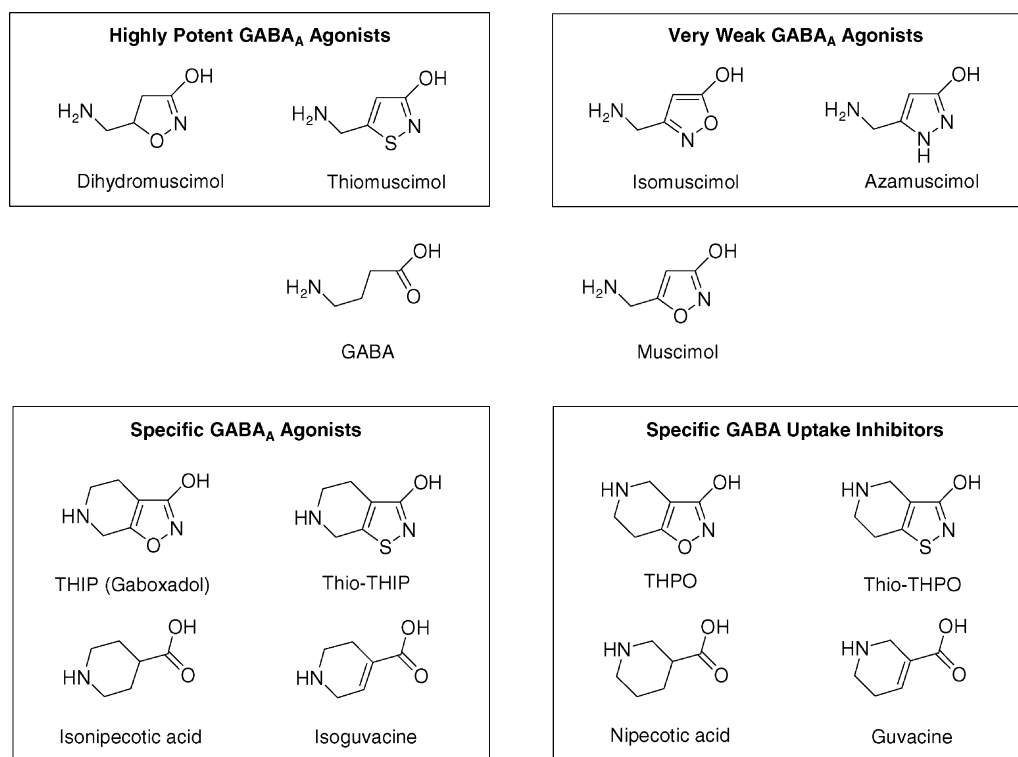


Fig. 2. Comparison of the structures of some GABA_A agonists and GABA uptake inhibitors.

can be replaced by a 3-isothiazolol or 3-hydroxyisoxazoline group to give thiomuscimol and dihydromuscimol, respectively, without significant loss of GABA_A receptor agonism [36]. (*S*)-Dihydromuscimol is the most potent GABA_A agonist so far described [37]. The structurally related muscimol analogues, isomuscimol and azamuscimol, on the other hand are virtually inactive, emphasizing the very strict structural constraints imposed on agonist molecules by the GABA_A receptors [36].

The conversion of muscimol into THIP (Gaboxadol) (Figs. 2 and 3) [3] and the isomeric compound 4,5,6,7-tetrahydroisoxazole[4,5-*c*]pyridin-3-ol (THPO) effectively separated GABA_A receptor and GABA uptake affinity, THIP being a specific GABA_A agonist and THPO a GABA uptake inhibitor (Fig. 2) [38].

Using THIP as a lead, a series of specific monoheterocyclic GABA_A agonists, including isoguvacine and isonipectic acid, was developed (Fig. 2) [3,36]. Whereas Thio-THIP (Fig. 1) shows GABA_A agonist effects on cat spinal neurons [39], recent studies using human brain

recombinant GABA_A receptors have disclosed very low-efficacy partial agonism/antagonism of Thio-THIP [40].

In light of the structural similarity of THIP and Thio-THIP (Fig. 1) the markedly different pharmacology of these compounds is noteworthy and emphasizes the strict structural requirements of GABA_A receptors. The p*K_a* values of THIP (4.4; 8.5) and Thio-THIP (6.1; 8.5) [39] are different, and a significant fraction of the molecules of the latter compound must contain a non-ionized 3-isothiazolol group at physiological pH. Furthermore, the different degree of charge delocalization of the zwitterionic forms of THIP and Thio-THIP and other structural parameters of these two compounds may have to be considered in order to explain their different potency and efficacy at GABA_A receptors.

A series of cyclic amino acids derived from THPO, including nipectic acid [38] and guvacine [41], was developed as GABA uptake inhibitors. Whereas nipectic acid and guvacine potently inhibit neuronal as well as glial GABA uptake [42], THPO interacts selectively with the latter uptake system. Thio-THPO is slightly weaker than THPO as an inhibitor of GABA uptake [39].

3. Partial GABA_A agonists

Under clinical conditions where stimulation of the GABA_A receptor system may be relevant, partial agonists displaying a relatively high efficacy may be particularly useful. The level of efficacy needed may be dependent on

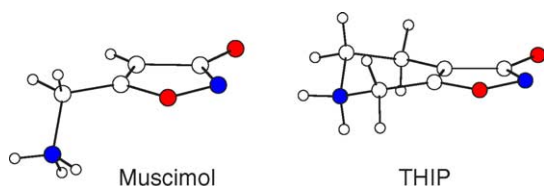


Fig. 3. The structures of muscimol and THIP as determined by X-ray crystallographic analyses (from The Chemical Record 2002;2: 419–30).

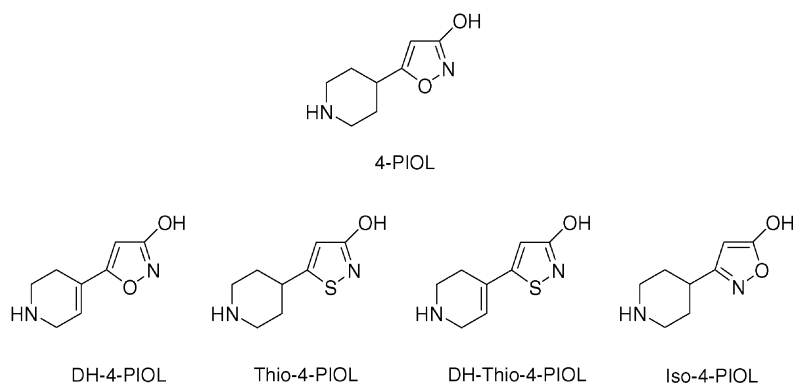


Fig. 4. Structures of the low-efficacy partial GABA_A agonist, 4-PIOL, and a number of analogues.

the disease in question. The potent analgesic effects of THIP (Gaboxadol) (see later section) seem to indicate that this relatively high level of efficacy [43,44] is close to optimal with respect to treatment of pain.

Analogously, very low-efficacy GABA_A agonists showing predominant antagonist profiles may have clinical interest in conditions where a reduction in GABA_A receptor activity may be needed. Such compounds may produce sufficient GABA_A receptor activity to avoid side effects.

The non-fused THIP analogue, 5-(4-piperidinyl)isoxazol-3-ol (4-PIOL) (Fig. 4), is a low affinity GABA_A agonist [45], which has been characterised as a partial GABA_A agonist using functional patch-clamp techniques on cultured cerebral cortical and hippocampal neurones [46,47]. In cortical neurones, the action of 4-PIOL was compared with those of the GABA_A agonist isoguvacine and the GABA_A antagonist BMC [46]. Based on these studies, it is concluded that 4-PIOL is a low-efficacy partial agonist showing a predominant GABA_A antagonist profile, being about 30-fold weaker than BMC as an antagonist at the GABA_A receptors. Recently, 4-PIOL has been proposed to possess some subtype-specific characteristics [48]. On recombinant GABA_A receptors of the $\alpha_1\beta_2\gamma_2$ subtype expressed in HEK-293 cells, 4-PIOL acted as a weak agonist, whereas it was devoid of activity in the $\alpha_6\beta_2\gamma_2$ receptor subtype.

4. 4-PIOL analogues as GABA_A antagonists

Introduction of alkyl groups into the 4-position of the 3-isoxazolol ring of muscimol and THIP severely inhibits interaction with the GABA_A receptor recognition site as illustrated in Fig. 5. Thus, 4-Me-muscimol is three to four orders of magnitude weaker than muscimol as an inhibitor of GABA_A receptor binding, whereas 4-Et-muscimol [49], and also 4-Me-THIP [50] are inactive. In contrast, the GABA_A recognition site tolerates introduction of alkyl groups into the 4-position of the 3-isoxazolol ring of 4-PIOL [51]. These structure-activity relationships indicate that the binding modes of the GABA_A agonists, muscimol and THIP and in particular of the low-efficacy partial agonist 4-PIOL are different.

There is strong evidence that an arginine residue at the GABA_A receptor recognition site is directly involved in the binding of the anionic part of the receptor ligand [52]. Based on this observation, a hypothesis has been proposed concerning the binding modes of the bioactive conformations of muscimol and 4-PIOL [51] as illustrated in Fig. 6. 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 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.

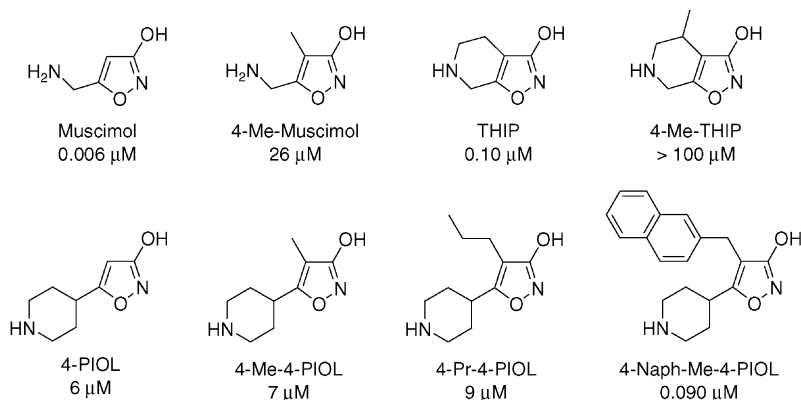


Fig. 5. GABA_A agonist binding data (IC₅₀, μM) for muscimol, THIP, 4-PIOL and some analogues of these GABA_A agonist ligands.

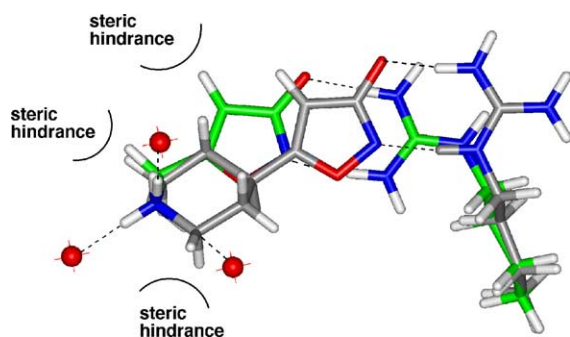


Fig. 6. A pharmacophore model for GABA_A receptor agonists showing the proposed binding modes of muscimol (green bonds) and 4-PIOL (light grey bonds) and their interactions with two different conformations of an arginine residue. The red spheres indicate sites to which the ammonium group in muscimol interacts via hydrogen bonds (from The Chemical Record 2002;2: 419–30).

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 [53]. The results from these studies are exemplified in Fig. 5. Substitution of the 4-position with alkyl or benzyl groups resulted in affinity and potencies comparable with those of 4-PIOL. Interestingly, however, introduction of more bulky groups such as diphenylalkyl and naphthylalkyl groups, as exemplified by the 2-naphthylmethyl analogue, 4-Naph-Me-4-PIOL are not only tolerated but resulted in a marked increase both in affinity and potency.

Using whole-cell patch-clamp techniques on 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 [53]. 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 potent and selective antagonist effect. The 2-naphthylmethyl and the 4-biphenylmethyl analogues (Fig. 7) showed antagonist potency comparable with or markedly higher than that of the standard GABA_A antagonist gabazine.

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 (Figs. 5 and 6). Thus, a cavity of considerable binding capacity seems to exist at the 4-PIOL recognition site of the GABA_A receptor.

Molecular modeling studies of the two high affinity compounds containing a 2-naphthylmethyl and a 3,3-diphenylpropyl substituent, and the less active 1-naphthylmethyl and 4-biphenylmethyl analogues, indicate that this proposed binding cavity may be exploited in two directions (Fig. 7). In both of these positions an aromatic ring seems to be highly favourable for the receptor affinity [53].

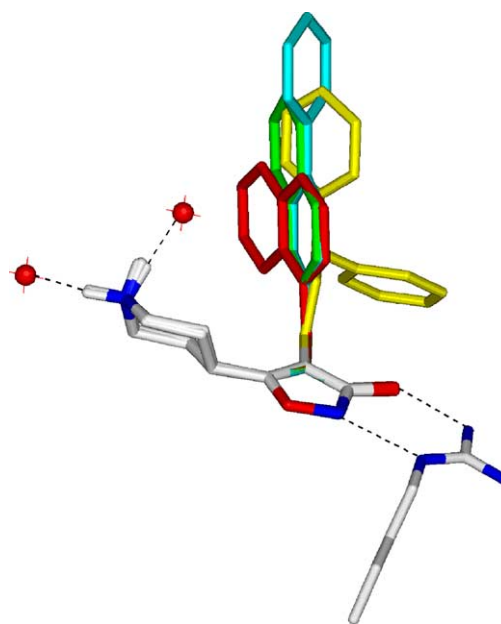


Fig. 7. Proposed bioactive conformations for the high affinity 1-naphthylmethyl, 2-naphthylmethyl, 3,3-diphenylpropyl, and 4-biphenylmethyl analogues of 4-PIOL (from The Chemical Record 2002;2: 419–30).

As mentioned earlier, the GABA binding site in the GABA_A receptor is assumed to be located at the interface between α and β subunits [54]. It has been speculated that the GABA_A antagonists bind to and stabilize a distinct inactive receptor conformation. In case of the 4-arylalkyl substituted 4-PIOL analogues it may be speculated that the large cavity accommodating the 4-substituent is located in the space between these subunits. GABA_A receptors belong to the same superfamily as the nicotinic acetylcholine receptors. It has been proposed that the mechanism for ligand-induced channel opening in nicotinic acetylcholine receptors involves rotations of the subunits in the ligand binding domain [55]. Assuming that the GABA_A receptors utilize a similar mechanism for channel opening, large substituents may interfere with the channel opening resulting in antagonistic effects of the compounds [53].

5. Behavioural and clinical effects of the partial GABA_A agonist THIP (Gaboxadol)

5.1. Analgesia and anxiety

The involvement of central GABA_A receptors in pain mechanisms and analgesia has been thoroughly studied, and the results have been discussed and reviewed [34,56]. The demonstration of potent antinociceptive effects of the specific and metabolically stable partial GABA_A agonist THIP (Gaboxadol) in different animal models and the potent analgesic effects of Gaboxadol in man greatly stimulated studies in this area of pain research. Gaboxadol-induced analgesic effects were shown to be insensi-

tive to the opioid antagonist naloxone indicating that these effects are not mediated by the opioid receptors [57].

Gaboxadol and morphine are approximately equipotent as analgesics, although their relative potencies are dependent on the animal species and experimental models used. Acute injection of Gaboxadol potentiates morphine-induced analgesia, and chronic administration of Gaboxadol produces a certain degree of functional tolerance to its analgesic effects. In contrast to morphine, Gaboxadol does not cause respiratory depression. Clinical studies on post-operation patients, and patients with chronic pain of malignant origin have disclosed potent analgesic effects of Gaboxadol, in the latter group of patients at total doses of 5–30 mg (i.m.) of Gaboxadol.

In cancer patients and also in patients with chronic anxiety [58] the desired effects of Gaboxadol were accompanied by side effects, notably sedation, nausea, and in a few cases euphoria. The side effects of Gaboxadol have, however, been described as mild and similar in quality to those of other GABA-mimetics [58]. This combination of analgesic and anxiolytic effects of Gaboxadol obviously has therapeutic prospects.

The neuronal and synaptic mechanisms underlying Gaboxadol- and, in general, GABA-induced analgesia are still only incompletely understood. The insensitivity of Gaboxadol-induced analgesia to naloxone has been consistently demonstrated. GABA-induced analgesia does not seem to be mediated primarily by spinal GABA_A receptors but rather by GABA mechanisms in the fore-brain, and it appears also to involve neurones in the midbrain. The naloxone-insensitivity and apparent lack of dependence liability of GABA_A agonist-mediated analgesia suggest that GABAergic drugs may play a role in future treatment of pain. Furthermore, it has been suggested that pharmacological manipulation of GABA mechanisms may have some relevance for future treatment of opioid drug addicts [56].

5.2. THIP (Gaboxadol) and sleep disorders

GABAergic compounds acting at the barbiturate site, the neurosteroid site, or the BZD site of the GABA_A receptor complex have been used as hypnotics for years. Most of these hypnotics interact with all subtypes of receptors. However, as α_4 and δ containing receptors do not bind BZDs, effect of BZDs at these receptor subtypes is absent [59]. It is generally agreed that the sedative effect of BZDs is mediated primarily via α_1 containing receptors, whereas side effects related to amnesia may be mediated by α_5 , primarily located in the hippocampal region [60]. However, short-acting α_1 selective BZD ligands like zaleplon, zopiclone and zolpidem do produce memory impairment and hangover effects [61], suggesting that even an α_1 selective compound with a very short half life may produce side effects. The reason for this side effect profile may well be a consequence of the high degree of GABA receptor

activation caused by the positive GABA_A receptor modulator. The massive activation of GABA receptors will influence several other systems, ultimately resulting in a general acute modification of the overall function of the CNS. Electroencephalographic (EEG) measurements during sleep support this hypothesis. A normal sleep pattern involves a complex variation of different degrees of sleep, ranging from light sleep via deeper sleep stages to the dream-associated rapid eye movement (REM) stage of sleep. Present understanding of sleep quality and the relation to EEG patterns is still limited, however, not only the duration of REM sleep but also the transitions between the different sleep stages are important [61]. The effect of BZDs at the sleep micro architecture is not limited to the onset of REM sleep, which is delayed.

In contrast to the observations with BZDs, barbiturates and neurosteroids, a series of studies have shown that this perturbation of the sleep micro architecture may not arise with compounds acting directly at the GABA_A recognition site [62,63]. In this respect, Gaboxadol is an interesting compound. Gaboxadol appears to improve the quality of sleep, as measured using behavioural parameters. The compound shows no effect on the onset of REM sleep, as measured using EEG [62,63]. Furthermore, studies in elderly patients, suffering from significant reduction in non-REM sleep, showed that Gaboxadol was able to normalize the sleep pattern [64]. Following dosing with Gaboxadol, patients did not experience the hangovers or impaired attention as reported for BZDs. Similar results have been obtained with muscimol, with the GABA uptake inhibitor Tiagabine [65], and with the glia-selective GABA uptake inhibitor, THPO (Fig. 2) [66], strongly suggesting that the functional consequences of a direct acting agonist or enhanced synaptic GABA concentration are different from those seen with GABA_A receptor modulators.

6. Conclusion

Molecular biology studies have revealed a high degree of structural heterogeneity of the GABA_A receptors. Development of subtype selective or specific compounds is of key importance for the understanding of the physiological and pathological roles of different GABA receptor subtypes and may lead to valuable therapeutic agents. Studies along these lines have, so far, been complicated by the lack of information about the topography of the recognition site(s) of the GABA_A receptor complex. In the absence of a crystal structure of the GABA_A receptor complex, 3D-pharmacophore models, based on the analysis of known receptor ligands, have been useful tools in the design of new ligands. Recombinant receptors play a key role in the pharmacological characterization of ligands for the GABA_A receptor. However, problems regarding subunit composition of native receptors, are far from being fully elucidated. Furthermore, the subtypes of receptors

involved in different disorders are still largely unknown. Although the functional consequences of modifications of subunit compositions, so far, is unpredictable, it has been shown that functional selectivity is obtainable for a number of GABA_A agonists. An important aspect is the selection of subunit combinations relevant for the prediction of in vivo activity.

Neither full GABA_A agonists nor antagonists may be useful therapeutic agents. Whereas the high-efficacy partial GABA_A agonist Gaboxadol shows very potent non-opioid analgesic effects and a novel type of hypnotic effects in the human clinic, it seems likely that partial GABA_A agonists showing lower levels of efficacy such as 4-PIOL analogues, may have therapeutic interest in certain CNS disorders such as schizophrenia. Information about the basic mechanism of receptor–ligand interaction resulting in partial agonism is, however, still not available, making design of new partial GABA_A agonists on a rational basis difficult.

Acknowledgement

This work was supported by grants from the Lundbeck Foundation, the Danish Medical Research Council, the Centre for Drug Design and Transport, the NeuroScience PharmaBiotec Centre and the Danish State Biotechnology Programme. The secretarial assistance of Mrs. Anne Nordly is gratefully acknowledged.

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