GABA receptors: pharmacological potential and pitfalls

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Abstract

Gamma-amino butyric acid (GABA), the major inhibitory neurotransmitter in the mammalian central nervous system, plays a key role in the regulation of neuronal transmission throughout the brain, affecting numerous physiological and psychological processes. Changes in GABA levels provoke disbalance between excitatory and inhibitory signals, and are involved in the development of numerous neuropsychiatric disorders. GABA exerts its effects via ionotropic (GABA_A) and metabotropic (GABA_B) receptors. Both types of receptors are targeted by many clinically important drugs that affect GABAergic function and are widely used in the treatment of anxiety disorder, epilepsy, insomnia, spasticity, aggressive behaviour, and other pathophysiological conditions and diseases. Of particular importance are drugs that modulate GABA_A receptor complex, such as benzodiazepines, barbiturates, neuroactive steroids, intravenous and inhalational anesthetics, and ethanol. Molecular interactions and subsequent pharmacological effects induced by drugs acting at GABA_A receptors are extremely complex due to structural heterogeneity of GABAA receptors and existence of numerous allosterically interconnected binding sites and various chemically distinct ligands that are able to bound to them. There is a growing interest in the development and application of subtype-selective drugs that will achieve specific therapeutic benefits without undesirable side effects. The aim of this review is to briefly summarize the key pharmacological properties of GABA receptors, and to present selected novel findings with the potential to open new perspectives in the development of more effective therapeutic strategies.

Keywords: GABA, GABA receptor pharmacology, benzodiazepines, neurosteroids

Introduction to GABA receptor diversity and pharmacological complexity

As the principal mediator of inhibitory synaptic transmission in the mammalian brain, GABA exerts its effects and shapes behaviour via two different types of GABA receptors. While fast ionotropic GABA_A receptors belong to the family of ligand gated ion channels, metabotropic GABA_B receptors are receptors with seven transmembrane domains coupled to G-proteins. Activation of GABA receptors by GABA induces membrane hyperpolarization, reduces the frequency of the generation of action potentials, and results in inhibition. Mainly local-circuit interneurons that constitute 15-20% of all cortical neurons predominantly use GABA as the neurotransmitter [1-3].

Most of its effects GABA exerts via GABA_A receptors. GABA_A receptors may be embedded in postsynaptic membrane where they mediate transient and fast synaptic inhibition that occurs in milliseconds, or may be located at the extrasynaptic places where respond to ambient concentrations of GABA and mediate long-term inhibition. In fast synaptic inhibition, GABA released from the presynaptic membrane terminals rapidly diffuses across the synaptic cleft and binds to binding sites at postsynaptic GABA_A receptors. Following binding, these receptors undergo a rapid conformational change that opens integral chloride channel and permit the flow of chloride ions down their chemical gradient, across the postsynaptic membrane. This ensures propagation of neurotransmission and represents basis of neural communication [4,5].

In contrast to GABA that directly activates and opens ion channels, some drugs, such as competitive antagonist bicuculline, selectively block GABA_A receptors and prevent inhibitory action of GABA. Muscimol, the principal psychoactive constituent of the mushroom *Amanita muscaria*, also acts as a potent, selective agonist of GABA_A receptors. A variety of clinically important drugs, including anticonvulsants, anxiolytics, general anesthetics, barbiturates, ethanol and neuroactive steroids, as well as drugs important as research tool such as convulsive picrotoxin, allosterically modulate GABA-induced activation of GABA_A receptors by acting on distinct binding sites on the receptor and modulate GABA action [5,6]. These drug interactions, together with structural diversity of GABA_A receptors, shape the complex nature of GABA_A receptor pharmacology, although simultaneously offer a great opportunity for the design of novel and more selective compounds for improved therapeutic applications [7]. Unfortunately, severe problems are related to the long-term use of drugs acting at GABA_A receptors, including loss of efficacy, and development of tolerance

and dependence that largely limit the time window of their desirable therapeutic benefits [4,8-11].

GABA also achieves inhibitory effects by acting at G-protein-coupled metabotropic GABA_B receptors. The slow, longer-lasting inhibition of GABA_B receptors is mediated by indirect gating of either potassium or calcium channels via second messengers. Characteristic agonists of GABA_B are the GABA analog baclofen and 3-aminopropylphosponous acid (3-APPA; CGP27492), while saclofen, phaclofen and 2-hydroxysaclofen act as antagonists of GABA_B receptors [12].

Old nomenclature also included GABA_C receptors that are expressed predominantly in retina. However, International Union of Basic and Clinical Pharmacology has recommended that GABA_C receptors should be assumed as one of the many isoforms of GABA_A receptors [5].

GABAA receptors and their subunits

Ubiquitously expressed GABA_A receptors are the major receptors of inhibitory neurotransmission across the central nervous system (CNS) that play a major role in virtually all brain physiological functions. GABA_A receptors are heteropentameric ligand-gated proteins that form a channel specific for chloride ions as an integral part. GABA_A receptors are members of the Cys-loop ligand-gated ion channel superfamily due to the presence of a disulphide-bridged loop in the extracellular domain. They share structural and functional homology with other members of the family. In addition to GABA_A receptors, the main pentameric receptors within the Cys-loop superfamily are nicotinic acetylcholine (nACh), 5-hydroxytryptamine type 3 (5HT₃) and glycine receptors [13,14].

GABA_A receptors are assembled from an impressive repertoire of different homologous subunits that are encoded from the pool of 19 different genes. Six α (α_{1-6}), three β (β_{1-3}), three γ (γ_{1-3}), δ , ε , θ , π and three ρ (ρ_{1-3}) subunits have been identified by molecular cloning techniques [6,14-16]. In some cells expression of more than eight different isoforms has been demonstrated [5]. Throughout the CNS some subunits are broadly expressed, while others exhibit more restricted expression profile [17,18]. For example, the most abundant α_1 subunit is expressed almost ubiquitously in the brain, α_5 is highly expressed in the hippocampus, α_6 subunit is expressed only in granule cells of the cerebellum, while ρ subunits, the major molecular components of the formerly called GABA_C receptors, are expressed mainly in the retina [19-22]. Expression of β_2 subunit strongly correlates with that of $\alpha 1$, while expression pattern of $\beta 3$ subunit resembles to that of $\alpha 2$. Isoform $\gamma 2$ is the most widely distributed isoform of γ subunit [17].

GABA_A receptor subtypes composed of different subunits represent distinct receptor populations that exhibit unique functional and pharmacological properties, and are differentially regulated at the transcriptional, post-transcriptional and translational levels [23]. GABA_A receptor subtypes are expressed in a specific spatio-temporal pattern in the developing and adult CNS. Some combinations of subunits, such as $\alpha 1\beta 2\gamma 2$, the most prevalent adult isoform, are broadly distributed in different brain areas, while the expression of some other subtypes of GABA_A receptors is restricted to certain regions [24].

Mature subunits share a common topological organization, and have approximately 450 amino acid residues in length. Subunits consist of a large extracellular N-terminal domain that contains already mentioned Cys loop, followed by four transmembrane domains (TM1-TM4), and a short extracellular C-terminal domain. TM2 domains of all five subunits line the lumen of ion channel. There is a large phosphorytable intracellular loop between TM3 and TM4 domains of each subunit that participates in protein-protein interactions and plays an important role in the regulation of GABA_A receptor function. Some of the intracellular proteins that interact with GABA_A receptors are GABA_A receptor associated protein (GABARAP), Golgi-specific DHHC zinc finger protein (GODZ) phospholipase C-related, catalytically inactive proteins PRIP-1 and PRIP-2, Plic-1, radixin, GABA_A receptors interacting factor 1 (GRIF-1), and brefeldin A-inhibited GDP/GTP exchange factor 2 (BIG2). These proteins interact with particular subunits at GABA_A receptor complex and modulate broad spectrum of GABA_A receptors activities, such as trafficking enhancement, surface stabilization, receptor internalization, anchoring of receptors in the cytoskeleton, and regulation of modification of GABA_A receptors [5,25].

Chromosomal mapping techniques revealed that the majority of genes coding for the subunits of GABA_A receptors are localized into four clusters on chromosomes 4, 5, 15 and X in the human genome, and within these clusters genes have conserved transcriptional orientation [26]. Additional diversity of receptor structure is accomplished by alternative splicing of some of the subunit mRNAs and by RNA editing that might affect kinetics, subunit assembly and cell-surface expression of GABA_A receptors [27]. For example, the γ_2 subunit appears in three alternatively spliced variants termed γ_2 L, γ_2 S and γ_2 XL [28], although physiological and pharmacological relevance of this diversity is still not completely resolved. Nova, a neuron-specific RNA binding protein, regulates alternative splicing of γ_2 subunit by acting on an intronic splicing enhancer which lies far downstream of the regulated

exon, and general dysregulation of Nova's splicing enhancer function may underlie the neurologic defects seen in patients with Nova's absence [29].

Only a limited number of mutations have been found in GABA_A receptor subunit genes that are specific for different diseases. For example, point mutations in the $\alpha 1$ and $\gamma 2$ subunits are found in patients with genetic epilepsies, while single nucleotide polymorphisms (SNPs) in the gene encoding the $\alpha 2$ subunit are involved in alcohol dependence and illicit drug dependence. The gene encoding the $\beta 1$ subunit also has been linked to alcohol dependence and to bipolar disorder. The genes encoding the $\alpha 1$, $\alpha 6$, $\beta 2$ and π subunits have been linked to schizophrenia [4].

Subunit composition determines pharmacological effects

It is still not resolved which subunits assemble together to form different isoforms of GABA_A receptors that really exist in nature [14]. Numerous expression studies using recombinant receptors indicated that receptors composed of α , β , and γ subunits form GABA_A receptors with a pharmacological profile that mimic properties of the majority of native receptors. Despite the fact that the immense number of different pentameric combinations can be assembled from the pool of native protein subunits, it seems that the prevalent form of native GABA_A receptors is the combination of two α , two β and a single γ , δ or ϵ subunit [30,31]. As mentioned before, isoform $\alpha 1\beta 2\gamma 2$ is considered as the most prevalent subtype of adult GABA_A receptors, with $\gamma 2\beta 2\alpha 1\beta 2\alpha 1$ counter-clockwise arrangement surrounding central ion pore when viewed from the synaptic cleft [32]. It is estimated that approximately 60% of all GABA_A receptors have the subunit composition $\alpha 1\beta 2\gamma 2$, approximately 15–20% have the $\alpha 2\beta 3\gamma 2$ combination, approximately 10–15% have the $\alpha 3\beta n\gamma 2$ combination, approximately 5% have the $\alpha 4\beta n\gamma$ or $\alpha 4\beta n\delta$ combination, less than 5% have the $\alpha 5\beta 2\gamma 2$ combination and, likewise, less than 5% have the $\alpha 6\beta 2/3\gamma 2$ combination [4]. Furthermore, it is possible for GABA_A receptors to contain two different α subunits. Thus, $\alpha 1\alpha 6\beta x\gamma 2$ and $\alpha 1 \alpha 6 \beta x \delta$ receptors have been found in the cerebellum [33].

Composition and arrangement of GABA_A receptors subunits shapes the synaptic response to GABA, as well as electrophysiological, functional and pharmacological properties of GABA_A receptors, including potency and efficacy of different drugs to exert their effects [6,34,35]. For example, Zezula et al. [36] systematically investigated properties of recombinant GABA_A receptors formed after transfection of human embryonic kidney (HEK) 293 cells with α 1-, β 3-, or γ 2-subunits, or with various combinations of these subunits. From

all possible subunit combinations, high affinity binding sites for GABA were formed in cells transfected with $\alpha 1\beta 3$ - or $\alpha 1\beta 3\gamma 2$ -subunits only. Binding of benzodiazepines were induced in cells after transfection with $\alpha 1\gamma 2$ - or $\alpha 1\beta 3\gamma 2$ subunits, whereas binding sites for convulsant [³⁵S]t-butylbicyclophosphorothionate (TBPS) were found in cells transfected with $\beta 3$ -, $\beta 3\gamma 2$ -, $\alpha 1\beta 3$ -, or $\alpha 1\beta 3\gamma 2$ -subunits. Binding of [³⁵S]TBPS was inhibited by pentobarbital (barbiturate), alphaxalone (anesthetic neuroactive steroid) and propofol (anesthetic) with a potency which differed in cells transfected with $\beta 3$ -, $\beta 3\gamma 2$ -, $\alpha 1\beta 3\gamma 2$ -subunits, demonstrating that receptors with different subunit composition exhibit distinct pharmacological properties.

Tonic and phasic inhibition by GABAA receptors

Some subunits, such as δ subunits, are placed only at extrasynaptic places where GABA_A receptors are activated by low ambient GABA concentrations present in the extracellular fluid, and mediate a so-called tonic inhibition that is relatively non-desensitizing. Isoforms containing γ subunit are expressed predominantly at synapses and mediate rapidly desensitizing phasic GABAergic neurotransmission. Synaptic GABA_A receptors are activated transiently after the release of GABA from presynaptic vesicles, while extrasynaptic GABA_A receptors are in general activated continuously, controlling neuronal excitability and the strength of synaptic transmission [37-41].

Tonic and phasic conductances underlie different physiological and behavioural processes [4]. Receptors containing δ subunit regulate numerous behavioural functions, including memory, nociception and anxiety, and may modulate neurogenesis. Taking into account that they are highly responsive to sedative-hypnotic and sleep-promoting drugs, general anesthetics, alcohol and neuroactive steroids, they are considered as potential therapeutic targets for the treatment of memory impairment, insomnia, mood disorders, pain, and post-stroke recovery [39,41-44].

Furthermore, extrasynaptic GABA_A receptors with δ subunit exhibit a pronounced variability in expression levels under different physiological and pathophysiological conditions that might reshape behaviour and change neuronal susceptibility to different drugs [41]. Modulation of tonic inhibition might be of clinical relevance as extrasynaptic GABA_A receptors may diminish seizure susceptibility and states of anxiety during the estrous cycle. In particular, in late diestrus in mice (high-progesteron phase), increased expression of δ -containing receptors increases tonic inhibition, suggesting that consequent reduction of

neuronal excitability might be beneficial in women with catamenial epilepsy and premenstrual dysphoric disorder [45]. The large increase in progesterone-derived neurosteroids during pregnancy, and their sharp decline at parturition, may also have considerable effects on GABA_A receptors during pregnancy and postpartum. Both tonic and phasic inhibition decrease in pregnant mice due to down-regulation of δ and γ 2 subunits, and are rebound immediately after delivery. Mice which do not exhibit δ -subunit regulation throughout pregnancy show depression-like and abnormal maternal behaviours [46]. On the other hand, expression of GABA_A receptors containing δ subunit is diminished in depression and schizophrenia [47,48].

Endogenous ligands of GABAA receptors

In addition to GABA, neurosteroids are the most important endogenous ligands of GABA_A receptors. Certain neurosteroids potently and specifically enhance function of GABA_A receptors, and consequently exert anxiolytic, analgesic, anticonvulsant, sedative, hypnotic and anesthetic properties, and contribute to behavioural effects of psychoactive substances [49]. In the CNS, steroids can be synthesized *de novo* from cholesterol, during stress, pregnancy and after ethanol consumption, and these steroids are called neurosteroids, or can be converted from steroid precursors from periphery [50-52]. *De novo* synthesized neuroactive steroids locally modify neuronal activity in a paracrine manner, and can influence mood and behaviour in various physiological and pathophysiological conditions. GABA-modulatory actions of the pregnane steroids are highly selective, with their actions being brain region- and neuron-dependent. Furthermore, the sensitivity of GABA_A receptors to neurosteroids is not static but can dynamically change. The molecular mechanisms determining this neuronal specificity include different GABA_A receptor isoforms, protein phosphorylation and local steroid metabolism and synthesis [53,54].

Relatively recently, it has been discovered that endocannabinoid 2-arachidonoyl glycerol (2-AG) potentiates GABA_A receptors at low concentrations of GABA, specifically in receptors containing β 2 subunit [55]. 2-AG allosterically potentiates GABA_A receptors via a binding site located in transmembrane segment TM4 of the β 2 subunit GABA_A receptors [56]. The discovery of a direct action of the 2-AG on the β 2-containing GABA_A receptors offers the potential for selective pharmacological intervention that would reduce motility and possibly affect sedation and state of anxiety [56]. Furthermore, it is shown that 2-AG acts synergistically with the neurosteroid tetrahydrodeoxycorticosterone (THDOC; 3 α ,21-

dihydroxy- 5α -pregnan-20-one) and modulates extrasynaptic receptors that respond to neurosteroids [55].

Modulation of GABAA receptors by pharmacological agents

GABA_A receptors are targeted by many clinically important drugs. Namely, besides binding site(s) for GABA, GABA_A receptors possess binding sites for a variety of clinically important substances, such as benzodiazepines and benzodiazepine-site ligands, barbiturates, intravenous and volatile anesthetics, ethanol and neuroactive steroids, all of which achieve at least some of their pharmacological effects after binding to GABA_A receptor complex [6,17,57]. These drugs modulate effects of GABA and affect GABAergic transmission, exerting a wide spectrum of pharmacological effects and inducing numerous conformational states of GABA_A receptors.

Diverse approaches have been applied in last years to more precisely identify various binding sites on GABA_A receptor complex, as this knowledge is prerequisite for the design of subtype specific, selective drugs of clinical relevance. Study performed on the comparative models of the extracellular and transmembrane domains of GABA_A receptors in the agonist-free state has revealed that a loose packing of these domains results in a large amount of solvent-accessible space, probably offering a potential explanation for such complex pharmacology and great flexibility of GABA_A receptors [58].

Drugs acting at benzodiazepine binding sites can only modulate GABA_A receptors, while barbiturates, neuroactive steroids and anesthetics at higher concentrations are able to directly activate GABA_A receptors in the absence of GABA [31,59-63]. Binding sites for all of these drugs are allosterically coupled. Therefore, drug that fits or binds to any of these multiple binding pockets induces conformational changes of the receptor complex and modulates GABA-elicited response [64,65]. For some of these multiple binding sites ligands have been identified, whereas for others sites, ligands are unknown. Similarly, the sites of action of large number of compounds that allosterically modulate GABA_A receptors are still not identified [66].

Increased activity of GABA_A receptor activity leads to anxiolysis, ataxia, myorelaxation, sedation, hypnosis, anesthesia and anterograde amnesia, whereas decrease in GABA_A receptor activity leads to increased vigilance, memory enhancement, anxiety and seizures [6,67]. GABA_A receptors are therefore widely used in the treatment of anxiety

disorders, insomnia, epilepsy, restlessness, and aggressive behaviours [17,68]. Recently, Rudolph and Möhler [69] gave comprehensive overview of the role of GABA_A receptor subtypes on cognitive and emotional behaviour, particularly to cognitive dysfunction and Down syndrome, anxiety disorders, depression, schizophrenia, and autism. They highlighted that a partial inverse agonist acting at the α 5-subunit-containing GABA_A receptors is in a clinical trial in individuals with Down syndrome. Substances that selectively reduce function of α 5-containing GABA_A receptors are considered as potential cognition enhancers for Alzheimer's syndrome and other dementia, consistent with genetic studies implicating GABA_A receptors in learning performances [70]. With regard to anxiety disorders, nonsedative anxiolytics based on the modulation of α 2- and α 3-subunit-containing GABA_A receptors have been established in clinical proof-of-concept trials. Furthermore, Rudolph and Möhler [69] emphasized the GABA hypothesis of depression and new options for antidepressant drug development. They also pointed out that cognitive symptoms in schizophrenia are attributed to a cortical GABAergic deficit, and that dysfunctional GABAergic inhibition contributes to the pathophysiology of autism spectrum disorders.

Furthermore, the properties of GABA_A receptors specific to interneurons may differ significantly from those found on projecting neurons, offering the possibility of developing interneuron-specific drugs that is of great therapeutic interest as more and more neurological and psychiatric disorders are linked to malfunction or deficits of interneurons [3]. Cells of glial origin also express GABA_A receptor subunit isoforms and form functional ion channels that are modulated by classical GABA_A receptor drugs, including diazepam and anesthetics etomidate and propofol [71].

Effects of GABA and GABA agonists at GABAA receptor

Besides GABA, other compounds also act at GABA recognition sites. Muscimol, 4,5,6,7-tetrahydroisoxazolo(5,4-*c*)pyridine-3-ol (THIP, a bicyclic muscimol analog) and isoguvacine are structural analogs of GABA, and have direct GABA mimetic effects, while alkaloid bicuculline competitively antagonises effects of GABA at GABA binding sites. Muscimol has been widely used as a valuable tool to study pharmacology of GABA_A receptors, and was the lead compound in the development of a range of GABAergic agents [72-74]. Behavioural effects of muscimol and THIP are preferentially mediated via high-affinity agonist binding sites at extrasynaptic receptors containing α 4 and δ subunits. For

THIP at high concentrations, the superagonistic behaviour on $\alpha 4\beta 3\delta$ receptors is achieved due to its ability to increase the duration of longer channel openings and their frequency, resulting in longer burst durations, while muscimol-related moderate superagonist behaviour is caused by reduced receptor desensitisation [75]. The ability to specifically increase the efficacy of extrasynaptic receptor activation by selected exogenous agonists might have important therapeutic implications, particularly under circumstances when synaptic inhibition is compromised, enabling manipulation of different components of inhibitory circuits [75-77]. Furthermore, it was found that chronic activation of GABA_A receptors by muscimol may exert neuroprotective effects against A β -induced apoptotic neuronal death [78].

Radioligand binding studies indicated that GABA and GABA agonists stimulate binding of benzodiazepine agonists, and inhibit binding of inverse agonists at benzodiazepine binding site. Vice versa, GABA binding is stimulated by benzodiazepines, barbiturates, neuroactive steroids and anesthetics. Picrotoxin and other convulsants allosterically inhibit binding of GABA at GABA_A receptors [79-83]. Studies with [³H]muscimol and [³H]GABA have revealed two kinds of agonist binding sites on the GABA_A receptor that differ in affinity. Biochemical and pharmacological experiments have suggested that high- and low-affinity sites might be interchangeable conformational variants of the same receptor complex. In most studies, functional responses to GABA have been observed only at micromolar concentrations, while nanomolar concentrations are needed to occupy the high-affinity binding sites. Hence, it is possible that the high-affinity sites represent a desensitized form of GABA_A receptors or otherwise non-functional binding sites [76,84].

The binding site for GABA at GABA_A receptor is located at the extracellular interface between the α and the β subunit, and a number of amino acids residues in the binding pocket have been identified that are important for pharmacological activity, subtype selectivity, assembly, trafficking and surface expression [84-87]. At the major, and the best characterized subtype of GABA_A receptor that is assembled from α , β and γ subunits, there are two binding sites for GABA. When two molecules of GABA bind to receptor complex, the receptor channel opens, and chloride ions rush into the postsynaptic neuron. However, as one pair of these otherwise identical subunits is flanked by γ and β (site 1) and the other by α and γ (site 2), different environments affect the binding sites. In $\alpha 1\beta 2\gamma 2$ receptors, site 2 has an approximately threefold higher affinity for GABA than site 1, whereas muscimol and bicuculline show some preference for site 1. Opening of the chloride channel also occurs for receptors occupied with a single agonist molecule but is promoted approximately 60-fold in those occupied by two agonists [88]. After occupancy of GABA binding sites, the large portion of the receptor undergo conformational changes that trap agonist inside the binding pocket, a deep narrowing cleft that probably may constrict during channel gating [64].

The potency of GABA in opening chloride channels is determined by subunit composition. For example, in one study aimed to characterize the potency of GABA at human recombinant δ -containing extrasynaptic GABA_A receptors expressed in Xenopus oocytes, it is found that $\alpha 4/\delta$ -containing receptors display high sensitivity to GABA, with mid-nanomolar concentrations activating $\alpha 4\beta 1\delta$ and $\alpha 4\beta 3\delta$ receptors. In the majority of oocytes expressing $\alpha 4\beta 3\delta$ subtypes, GABA produced a biphasic concentration-response curve, and activated the receptor with low and high concentrations, while at $\alpha 4\beta 2\delta$ receptors, GABA exerted low micromolar activity. An analysis of 10 point mutations in the large N-terminal extracellular domains in $\alpha 4\beta 3\delta$ receptors shows that GABA interacts with amino acids different to those reported for $\alpha 1\beta 2\gamma 2$ GABA_A receptors suggesting that GABA may have different binding modes for extrasynaptic δ -containing GABA_A receptors compared to their synaptic counterparts [89].

Effects of drugs acting at benzodiazepine binding sites at GABAA receptor

Due to their pharmacological and clinical relevance, benzodiazepines have attracted much attention as allosteric modulators of GABA_A receptors. Various ligands acting at benzodiazepine binding site can enhance (positive modulators or agonists) or reduce (negative modulators or inverse agonists) the actions of GABA on GABAergic transmission with a different range of efficacy (from full to partial). Antagonists are devoid of intrinsic activity, but can inhibit effects of both positive and negative modulators [6].

Benzodiazepines and other substances acting at the benzodiazepine binding site are the most commonly prescribed drugs in therapeutic use due to their desirable anxiolytic, myorelaxant, sedative/hypnotic and anticonvulsant effects. Classical benzodiazepines, such as diazepam and flunitrazepam, do not open channel directly, but allosterically potentiate activation of GABA_A receptor containing either $\alpha 1$, -2, -3 or -5 subunits. This means that after their binding, conformational changes at the receptor confer to increased affinity for channel gating by GABA at both agonist sites. Accordingly, in electrophysiological recordings classical benzodiazepines do not affect the maximal current amplitude, but GABA

concentrations required for channel opening are shifted to lower values. In contrast to them, negative allosteric modulators shift GABA concentration curve to the right [5].

The subunit composition of the GABA_A receptor complex largely determines pharmacological properties at benzodiazepine binding site. In fact, pharmacological profile is predominantly determined by α and γ subunits isoforms. It has been shown that different receptor conformations, in particular $\gamma\beta\alpha1\beta\alpha6$, $\gamma\beta\alpha6\beta\alpha1$, $\gamma\beta\alpha1\beta\alpha1$ and $\gamma\beta\alpha6\beta\alpha6$, have their own pharmacological signature and that diazepam efficacy is determined exclusively by the α - subunit neighbouring the γ 2-subunit [35]. The most applied concept is that the α 1 β x γ 2receptors, called type I BZ receptors, preferentially bind full benzodiazepine agonists with high affinity, while $\alpha 2/3/5\beta x\gamma 2$ -receptors, classified as type II BZ receptors, have approximately 10-fold lower affinity for benzodiazepine agonists [24,81]. Sedative, anterograde amnesic and partly anticonvulsant properties are attributed to al-containing receptors, while the presence of $\alpha 2$ and/or $\alpha 3$ subunits determines anxiolytic and myorelaxant activity of benzodiazepines [90-92]. This means that compound that selectively modulates receptors with $\alpha 2$ and/or $\alpha 3$ subunits, but not $\alpha 1$ -containing receptors, would be a nonsedating anxiolytic, suitable for the treatment of generalised anxiety disorder [93]. Receptors containing $\alpha 5$ subunits confer to learning and temporal and spatial memory. The $\alpha 5$ subunit of the GABA_A receptor is localized mainly to the hippocampus of the mammalian brain. It is shown that the largely extrasynaptic $\alpha 5$ GABA_A receptors in hippocampal pyramidal cells are implicated as control elements of the temporal association of threat cues in trace fear conditioning [94]. Furthermore, the α 5 -/- mice showed a significantly improved performance in a water maze model of spatial learning, indicating that α 5-containing receptors play a key role in cognitive processes [95]. However, as pointed by Sigel and Steinmann [5], behaviour is a complex phenomenon, and caution is needed in an attempt to define the role of individual GABA_A receptor subunits, as most probably several types of GABA_A receptors determines even the simple behavioural traits. Receptors containing $\alpha 4$ and $\alpha 6$ subunits are generally thought as insensitive to classical benzodiazepine agonists [96,97], although they retain a high affinity for the partial inverse benzodiazepine agonist Ro 15-4513 [98]. Unusually, specific high-affinity binding of the benzodiazepine agonist flunitrazepam to $\alpha 6\beta 2\gamma 2$ -containing GABA_A receptors was shown [99], but flunitrazepam action on $\alpha 6\beta 2\gamma 2$ -receptors was opposite to its action on $\alpha 1\beta 2\gamma 2$ -receptors where it exhibits an inverse agonistic effect in a GABA-dependent manner.

Benzodiazepines bind with high-affinity to a binding site located at the α/γ subunit interface in the extracellular domain of GABA_A receptor [65,100]. The binding site for benzodiazepines is formed by one of α subunits (α 1, α 2, α 3 or α 5) and a γ subunit (typically the γ 2 subunit, which is present in approximately 90% of all GABA_A receptors) [4]. Studies on recombinant GABA_A receptors indicated that co-expression of α , β and γ subunits is required for arrangement of high-affinity binding sites at GABA_A receptor complex [101]. Hanson and Czajkowsky [102] have identified that Gln182-Arg197 region of γ 2 subunit is part of the allosteric pathway that allows propagation of structural changes induced by positive benzodiazepine modulators through the protein to the channel domain. Furthermore, several amino acid residues on α and γ subunits have been identified that are crucial for benzodiazepine activity and form benzodiazepine binding pocket [86,103-105]. It is considered that antagonists bind in a pocket partly overlapping with the agonist site, although they can extend further into the solvent accessible cavity [106].

Richter et al. [107] described a binding mode for diazepam and analogs. By using this binding hypothesis and virtual screening approaches, they identified numerous known ligands of benzodiazepine-binding site from different structural classes and predicted potential new chemotypes for this site, thus demonstrating its suitability for drug discovery and structure-based drug design which is of major therapeutic importance. According to the model, although hundreds of ligands from different structural classes can bind to the benzodiazepine-binding sites, only a minority is bound to the same bioactive conformation as diazepam.

Evidence exists that some benzodiazepines can also bind to additional binding sites on GABA_A receptor complex. Walters et al. [108] demonstrated that in the presence of low concentrations of GABA, diazepam produces a biphasic potentiation of $\alpha 1\beta 2\gamma 2$ -receptor channels, with distinct components in the nanomolar and micromolar concentration ranges. Mutations at equivalent residues within the second transmembrane domains (TM2) of α , β and γ subunits abolished only the micromolar component, while nanomolar component, which depends on the presence of $\gamma 2$ subunit, was not affected. Similarly, converse mutation of the corresponding TM2 residue and a TM3 residue within $\rho 1$ subunits confers diazepam sensitivity on homo-oligomeric $\rho 1$ -receptor channels that are otherwise insensitive to benzodiazepines. Furthermore, effect of micromolar component on the GABA-induced current was not inhibited by flumazenil, an antagonist of benzodiazepine binding sites. As these transmembrane domains are important for effects of anesthetics, and taking into account anesthetic properties of diazepam, it might be that low potency modulation is elicited via anesthetic binding site [66].

In contrast to diazepam that act as positive allosteric modulator in both nanomolar and micromolar range, β -carbolines, such as methyl-6,7-dimethoxy-4-ethyl- β -carboline (DMCM), act as negative allosteric modulators at high affinity sites, but potentiate effects of GABA via the low-affinity binding sites [109]. The apparent reversal of inhibition and potentiation by high concentrations of DMCM is insensitive to flumazenil, $\beta 2/3$ subunit dependent, and probably elicited via the binding site for anticonvulsant loreclezole in the transmembrane domain. Moreover, Baur et al. [110] found evidence for the presence of a third site for benzodiazepines that prevents modulation of GABA_A receptors via the classical benzodiazepine site. The novel site potentially contributes to the high degree of safety to some of these drugs. Their results indicate that this site may be located at the α/β subunit interface.

In addition, it is found that the specific effects of imidazobenzodiazepine Ro15-4513 in antagonizing ethanol-stimulated ³⁶Cl⁻ uptake and behaviour can be completely blocked by benzodiazepine receptor antagonists. As other benzodiazepine receptor inverse agonists fail to antagonize the actions of ethanol *in vitro* or *in vivo*, a novel binding site for Ro15-4513 is also hypothesized [111]. Hanchar et al. [112] demonstrated that extrasynaptic δ subunit-containing receptors bind the imidazo-benzodiazepines flumazenil and Ro15-4513 with high affinity, contrary to the widely held belief that these receptors are benzodiazepine insensitive. Recently, it was identified that in addition to benzodiazepine binding site, Ro15-4513 interacts with high affinity to sites at the $\alpha 4/6+\beta 3$ - interface (each subunit has a plus and minus side determined by the absolute arrangement of subunits around central pore) and thus inhibits ethanol effects [98].

As for some potential binding sites no interacting drugs have been identified, Ramerstorfer et al. [113] established a steric hindrance procedure for the identification of drugs acting at the extracellular $\alpha 1+\beta 3$ – interface, which is homologous to the benzodiazepine binding site at the $\alpha 1 + \gamma 2$ - interface, while GABA-binding sites are located at the $\beta + \alpha$ interface [66]. Among screened compounds, anxiolytic pyrazoloquinoline 2-pmethoxyphenylpyrazolo[4,3-c]quinolin-3(5H)-one (CGS 9895) was able to enhance GABAinduced currents at a1B3 receptors from rat. Thus, at nanomolar concentrations CGS 9895 behaves as an antagonist of high-affinity, benzodiazepine binding site, but exerts a lowpotency positive modulatory action at the extracellular $\alpha+\beta$ - interface of $\alpha1\beta3\gamma2$ and $\alpha1\beta3$ receptors. Other experiments indicated that the CGS 9895 effect was dependent on α and β subunit isoforms forming the interface. Since CGS 9895, even at high concentrations, was only able to enhance GABA-induced currents but not directly activate these receptors, it is likely that drugs acting via the $\alpha+\beta-$ interface only have GABA-modulatory properties, like the benzodiazepines do. In contrast to benzodiazepines, these drugs will interact with receptors composed of $\alpha\beta$, $\alpha\beta\gamma$, and $\alpha\beta\delta$ subunits. Hence, they would exhibit a much broader anticonvulsive action than benzodiazepines, which might be of clinical importance for the treatment of epilepsy and development of subtype-selective drugs.

Based on results obtained with crystal structure of ELIC, prokaryotic homolog of GABA_A receptor, it is possible that at least some benzodiazepines, such as flurazepam, may directly interact with the GABA-binding site at $\beta+\alpha$ - interface at high concentration and displace GABA from its binding site, although these effects are observed only at high concentrations and further experiments are needed to confirm if this site has a functional role in GABA_A receptors [66,114].

Unfortunately, despite their proven clinical efficacy, positive modulators of benzodiazepine binding sites at GABA_A receptor possess a relatively narrow window between doses that produce anxiolytic effects and those that cause sedation, and are associated with the development of tolerance and physical and psychological dependence and a potential for abuse [115-117]. Hence, strategy to resolve the disadvantages of classical full agonists was directed toward development of partial agonists with lower intrinsic efficacy. A number of non-selective partial agonists with reduced intrinsic efficacy, including bretazenil, pazinaclone and abecarnil, were described. In general, although they have a large window between anxiolytic and sedative doses and their dependence and abuse liabilities were much lower, they failed to translate into clinical benefit [93]. More recently, the imidazolone derivatives imepitoin was shown to act as low-affinity partial agonist at the benzodiazepine site of the GABA_A receptor, and was approved for the treatment of epilepsy in epileptic dogs due to its more favorable pharmacokinetic profile in dogs versus humans, reactivating again the interest for partial benzodiazepine agonists for the treatments of human epilepsy [118].

Regarding development of tolerance and dependence to benzodiazepine-site ligands after prolonged administration, diverse effects at the molecular and cellular level have been described, including receptor desensitization and changes in the number and subunit composition of surface GABA_A receptors, often leading to decreased ability of benzodiazepines to potentiate effects of GABA [17,119-124].

Effects of convulsants at GABAA receptor

Very early picrotoxin was identified as a noncompetitive inhibitor of GABAA receptor. Binding studies further revealed that numerous compounds from heterogeneous chemical affinity for picrotoxin-binding sites. including classes posses tbutylbicyclophosphorothionate (TBPS), *t*-butylbicycloorthobenzoate (TBOB), pentylenetetrazole, and some insecticides (such as dieldrin and lindane). They inhibit GABAinduced influx of chloride ions acting as allosteric antagonists at GABA_A receptor complex [125-128].

TBPS partially and allosterically inhibits binding of GABA and positive modulators of GABAA receptors, but facilitates binding of negative modulators. Similarly, modulators of GABA_A receptor function affect [³⁵S]TBPS binding. GABA exhibits biphasic effect on the allosteric modulation of [35S]TBPS binding. Thus, [35S]TBPS binding was allosterically inhibited by high concentrations of GABA, GABA agonists and other positive modulators (various benzodiazepine receptor agonists, anesthetics and pentobarbital) and promoted in the presence of negative modulators such as ethyl-β-carboline-3carboxylate (β-CCE). On the contrary, GABA applied at low concentrations enhanced [³⁵S]TBPS binding [6,36,129-132]. In addition, the effects of the anesthetic steroid and pentobarbital on [³⁵S]TBPS binding were markedly altered by GABA, which at 2 µM increased their maximal effects and half-maximal concentrations. On the contrary, GABA did not affect changes in [³⁵S]TBPS binding produced by various benzodiazepine receptor agonists, indicating more pronounced functional coupling of the GABA sites with those for the steroid and the barbiturate, as compared with the benzodiazepine site. Hence, it appears that the degree of [³⁵S]TBPS binding in the presence of GABA reflects the functional state of GABAA receptors and may be useful for characterization of allosteric interactions between various sites on the receptor [6,133].

Inhibitory effects of GABA on [35 S]TBPS binding are determined by α subunit isoform. It has been shown that expression of recombinant $\alpha 6\beta 2\gamma 2$ GABA_A receptors produces [35 S]TBPS binding sites that are about 10-fold more sensitive to inhibition by GABA than were those inherent to $\alpha 1\beta 2\gamma 2$ GABA_A receptors. Similarly, the neurosteroid 5 α pregnan-3 α -ol-20-one (allopregnanolone) affected the binding in both $\alpha 1\beta 2\gamma 2$ and $\alpha 6\beta 2\gamma 2$ receptors, but inhibition was greater in $\alpha 6$ -containing than in $\alpha 1$ -containing receptors, also suggesting differential coupling of both GABA and neurosteroid sites with the convulsant site [134].

At first, it was suggested that picrotoxin's site of action is probably within the channel pore as mutations in TM2 domains (that form chloride channels) produced currents that were insensitive to picrotoxin [135]. By convention, Cys-loop receptor TM2 residues are numbered

from intracellular 0 to extracellular 20 positions. Although the exact location of picrotoxin binding to ionophore is still unknown, its sensitivity to mutations in residues 2/3 and 6 of TM2 indicates that the site contains residues 2–6. In particular, it is suggested that picrotoxin enters deep inside the ionophore pore, binds with its hydrophobic moiety to residue 2 of TM2 (close to the pore) and forms hydrogen bounds with residue 6 in the middle of TM2. Residue 15 is probably important for interplay between GABA and picrotoxin binding sites. Other studies implicated residues 9 and 15 in the regulation of channel properties, including desensitization, stabilization of open states and gating. This indirectly supports the possibility of a second "modulatory" (allosteric) binding site of picrotoxin including residues 15–19 [128].

Dillon et al. [136] provided evidence that picrotoxin and TBPS interact with GABAbound and -resting receptors, but their affinity for GABA-bound receptors is about 10 times greater, largely due to a markedly increased association rate to the multi-liganded receptors. Accordingly, TBPS and picrotoxin are considered as open channel blockers. However, [³⁵S]TBPS binds to recombinant receptors in the absence of GABA [130]. Hence, it does not gain access to its binding site via the open pore but through alternative routes regulated from the agonist binding site [137]. It is shown that picrotoxin displaceable [³⁵S]TBPS binding to $\alpha 1\beta 2\gamma 2$ GABA_A receptors might occur in the absence of GABA, suggesting that access to the binding site is independent of activation. In fact, bicuculline-sensitive spontaneous gating contributes to [³⁵S]TBPS binding in the absence of GABA by providing access to the channel. Accordingly, a decrease in a spontaneous gating reduces accessibility of TBPS to its binding site [132]. GABA application during picrotoxin or TBPS administration enhanced $\alpha 1\beta 2\gamma 2$ receptor blockade, and this GABA-dependent component of TBPS blockade accounts for the stimulation of [³⁵S]TBPS binding to $\alpha 1\beta 2\gamma 2$ receptors seen with low concentrations of GABA, while application of GABA at concentrations that cause significant steady-state desensitization reduces [³⁵S]TBPS binding [132].

Effects of anesthetics at GABAA receptor

General anesthetics are used to induce a reversible loss of consciousness in surgical patients and to relieve pain. GABA_A receptors are the major site of action of clinically used intravenous anesthetics such as etomidate, propofol, barbiturates and neuroactive steroids. On the other hand, clinically used volatile anesthetics such as isoflurane and enflurane, and long-

chain alcohols, presumably act via a multitude of targets, GABA_A receptors being just one of them [4,63,138,139].

In general, GABA_A receptors are positively modulated by clinical doses of anesthetics, while at higher concentrations anesthetics can directly activate GABA_A receptors [63,140]. Accordingly, many studies were aimed to identify binding site(s) for anesthetics on GABAA receptors, which are generally considered as the main target for propofol and etomidate action as already mentioned. Belelli and co-authors [141] have shown that the ability of etomidate to modulate and activate GABA_A receptors *in vitro* is uniquely dependent upon the β subunit subtype present within the receptor, i.e. receptors containing β^2 - or β^3 -, but not β^1 subunit, are highly sensitive to etomidate. They identified a region distal to the extracellular Nterminal domain as a determinant of the selectivity of etomidate: mutation of Asn present within the channel domain of the β 3 subunit to Ser (the homologous residue in β 1) strongly suppressed the GABA-modulatory and GABA-mimetic effects of etomidate. Later, it was confirmed that the same point mutation (Asn265Met) completely abolished the suppression of noxious-evoked movements in response to propofol and etomidate in a knock-in mouse, while only slightly decreased in response to volatile anesthetics enflurane and halothane. Mice also displayed a profound reduction in the loss of righting reflex duration (a surrogate for loss of consciousness) in response to intravenous but not volatile anesthetics. Accordingly, electrophysiological recordings revealed that anesthetic agents were significantly less effective in enhancing GABA-induced currents in cortical brain slices derived from mutant mice, thus identifying the key molecular determinant of behavioural responses evoked by intravenous anesthetics [142].

Studies on recombinant receptors have revealed that isoforms of α , γ and δ subunits affect propofol modulation of the GABA_A receptor, although it seems likely that the propofol binding site involves the β subunit (see below). In order to investigate the role of α subunit in the modulatory effects of propofol, Krasowski et al. [143] used whole-cell patch clamp recordings from cells expressing only $\alpha 1\beta 3\gamma 2$ or $\alpha 6\beta 3\gamma 2$ subtypes, and found that at clinically relevant concentrations, propofol potentiated submaximal GABA currents in $\alpha 1\beta 3\gamma 2$ receptors to a far greater degree than in $\alpha 6\beta 3\gamma 2$ receptors. Isoform of α subunit influenced the efficacy of propofol for modulation, but not its potency. On the contrary, direct gating of the chloride channel was significantly larger in the $\alpha 6\beta 3\gamma 2$ receptors. A same trend for potentiation and direct gating was observed for other anesthetics. Similarly, pharmacological properties of human recombinant GABA_A receptors were studied in Xenopus oocytes expressing $\alpha 1\beta 2$, $\alpha 1\beta 2\gamma 2L$, or $\alpha 2\beta 2\gamma 2L$ receptor isoforms. In all receptor isoforms tested, propofol potentiated the GABA-activated currents in a concentration-dependent manner, and was able to directly activate all three receptor isoforms. Addition of the $\gamma 2L$ subunit to the $\alpha 1\beta 2$ isoform reduced receptor sensitivity to direct activation, while replacement of the α 1-subunit with the α 2subunit increased receptor sensitivity to propofol's direct effects, indicating that isoforms of α and γ subunits have the ability to influence both the direct and modulatory actions of propofol on GABA_A receptor function [144]. Jones et al. [140] reported that $\gamma 2$ subunit is not a prerequisite for activation of GABAA receptors by propofol or for its potentiation of GABAactivated currents, but showed that it may contribute to the efficacy of propofol as a GABAA receptor activator. Furthermore, the findings that propofol reduced desensitization and prolonged deactivation of receptors containing y2L subunit and enhanced peak currents or prolonged deactivation of receptors containing δ subunit suggest that propofol-induced enhancement of both phasic and tonic inhibition may contribute to its anesthetic effect in the brain [145]. In accordance with mentioned studies, mutations in α and γ subunits change modulatory effects of propofol at the GABA_A receptors [146,147], while particular residues in β subunits, distributed throughout the membrane-spanning region of the receptor, as well as in the extracellular domain, determine the propofol's actions. Thus, a point mutation in the β 1 subunit (Met286Trp) abolished potentiation of GABA by propofol, but did not alter direct activation of the receptor by high concentrations of propofol [148]. The same mutation had the same effect in the β^2 subunit, probably by altering the dimensions of a binding pocket for propofol and related alkylphenol general anesthetics [149]. One mutation in B2 subunit (Tyr444Trp) was particularly interesting, as it selectively suppressed the ability of propofol to enhance receptor function, while retaining normal sensitivity to etomidate [150].

Although various anesthetics do not bind to the same binding site and exhibit distinct structural requirements for modulatory and direct actions, it seems that main sites of action are located close to each other, within the transmembrane domain [151]. The influence of the large intracellular loop in propofol sensitivity has also been recognized. Moraga-Cid et al. [147] have shown that mutation of a conserved Phe385 residue within the α 1 large intracellular loop significantly reduced propofol enhancement, although propofol-hyposensitive mutant receptors retained their sensitivity to other allosteric modulators such as alcohols, etomidate, trichloroethanol, and isoflurane. At the single-channel level, the ability of propofol to increase open probability was significantly reduced, altogether suggesting a new role of the large intracellular loop in allosteric modulation, and providing new insight into the propofol-mediated modulation of GABA_A receptor complex. By using a novel propofol

analogue photolabeling reagent, Yip et al. [152] have identified a novel binding site for propofol in β 3 homopentamers and α 1 β 3 heteropentamers. The binding site is located within the β subunit, at the interface between the transmembrane domains and the extracellular domain, and lies close to known determinants of anesthetic sensitivity in transmembrane segments TM1 and TM2.

The similar approach was used to determine binding site for etomidate. By using radiolabeled, photoreactive etomidate analogs, residues α Met-236 and β Met-286 in the α TM1 and β TM3 transmembrane helices, were identified as a part of single binding pocket in the transmembrane domain at the β + α - interface, close to the extracellular domain, bellow the GABA-binding site. Such localization of the etomidate binding site to an inter-subunit, not an intra-subunit binding pocket was an interesting finding, suggesting that binding sites at subunit interfaces may be a feature not only for GABA and benzodiazepines, but also for etomidate and other anesthetics [153,154]. As photolabeled etomidate analog was also displaced by isoflurane, Li et al. [139] suggested that inhalation anesthetics also interact with this binding site. More precisely, in their study photolabeling was inhibited by low millimolar anesthetic concentrations of propofol, barbiturates, and isoflurane, but not by octanol or ethanol. Inhibition by barbiturates and propofol was only partial, consistent with allosteric interactions, whereas isoflurane inhibition was nearly complete, suggesting competitive interactions.

Effects of barbiturates at GABAA receptor

Response of $GABA_A$ receptors may also be modulated by barbiturates. They have been used in the past due to their anticonvulsant, anxiolytic, sedative, and hypnotic actions, but they exert serious side effects, including profound depression of CNS activity with the induction of pronounced sedation, cognitive slowing or confusion, hyperactivity in children and a lethal risk in a case of overdose [155-157].

As they bind to GABA_A receptor with low affinity, most of their actions have been analysed indirectly by studying interactions with other binding sites. Barbiturates enhance binding of GABA and benzodiazepines, whereas inhibit binding at convulsant binding site [158]. Depending on the concentration, barbiturates act in three different ways, and their effects are proportional to anesthetic potency. At low μ M (sub-anesthetic) concentrations they potentiate GABA-induced effects (modulatory effect) and prolong duration of open conformation of chloride channel, possibly by stabilizing open state(s). In higher μ M (anesthetic) concentrations (app. >50 μ M) they directly open chloride channel (agonistic effect), whereas at very high mM concentrations block GABA-induced current (inhibitory effect) [159,160]. More precisely, when co-applied with low concentrations of GABA, barbiturates increased the mean open time. The increase in channel open time results in greater chloride current flux and increased likelihood that channel openings will summate, producing larger inhibitory currents [157]. In the absence of GABA, high concentrations of barbiturates may directly activate GABA_A receptor chloride currents, although with lower efficacy than GABA [60]. All these effects are probably mediated by separate mechanisms. In support of this conclusion, it was demonstrated that mutation of Thr262 in the transmembrane domain of β 1 subunits at α 1 β 1 recombinant GABA_A receptors abolished modulatory effect, without affecting direct agonistic or inhibitory effects of pentobarbital [161].

Although barbiturate sensitivity does not require a specific subunit composition, their effects are also determined by subunit composition [57,157]. Isoforms of α subunit affect efficacy, but not affinity of barbiturates in potentiation of GABA response. In contrast to benzodiazepines that are ineffective at GABA_A receptors containing α 4 or α 6 subunit, barbiturates elicit modulatory effects at receptors containing these isoforms [97,159], and the isoform of α subunit influences the level of potentiation [162]. Related to agonistic activity, the type of α subunit present determines both the degree of affinity and efficacy observed. Receptors containing α 6 isoform produce maximum direct response to barbiturate pentobarbitone, larger than that obtainable with maximum GABA, and larger than that obtainable with other α subunits. GABA_A receptors containing α 6 subunits also have higher affinity for direct activation by pentobarbitone. Furthermore, the direct effect of pentobarbitone can be blocked by picrotoxin, but not by competitive antagonists, such as bicuculline [97,159]. Surprisingly, unlike receptors containing α 6 isoform, α 4 β 1 γ 2 receptors did not elicit any direct activation of GABA_A receptors by pentobarbital [97].

Some amino acid residues have been identified with major impact on barbiturate activity. An invariant proline residue in transmembrane domain TM1 is included in a consensus motif of all GABA_A receptor subunits. Proline in TM1 segment of β 1 subunit affects the linkage between GABA binding and channel gating and is critical for barbiturate enhancement at recombinant GABA_A receptors, without affecting enhancement by diazepam or neurosteroid alfaxalone [163]. Gly219 near the TM1 at β 2 subunit is also identified as important for allosteric (modulatory) effects of pentobarbital since pentobarbital-induced enhancement of [³H]muscimol and [³H]flunitrazepam binding in receptors containing the β_2 (Gly219Phe) point mutation displayed a significantly reduced efficacy in modulation.

Furthermore, functional analysis of pentobarbital-enhanced GABA currents recorded with whole-cell patch clamp demonstrated that this mutation eliminates the potentiating effect of the anesthetic. Interestingly, the $\alpha 1\beta 2$ (Gly219Phe) $\gamma 2$ receptors also were more sensitive to direct channel activation by pentobarbital, suggesting that Gly219 may be important for conformational or allosteric interactions of channel gating by GABA and barbiturates [164].

Chiara et al. [165] used a photoreactive barbiturate analog to determine residues important for barbiturate binding site. It photolabeled sites at the $\alpha+\beta$ - and $\gamma+\beta$ - subunit interfaces in the transmembrane domain, near the synaptic side. These binding sites are distinct but homologous to the etomidate sites at the two $\beta+\alpha$ - subunit interfaces, as all are located at the same depth in the transmembrane domain. They demonstrated that photoreactive barbiturate and etomidate derivatives are highly selective for their own sites, indicating that there are two structurally related, but pharmacologically distinct, classes of inter-subunit general anesthetic-binding sites in the transmembrane domain of human $\alpha 1\beta 3\gamma 2$ GABA_A receptors. They hypothesized that binding at any of these homologous inter-subunit sites is sufficient for anesthetic action. Namely, a wide range of general anesthetic structures target these four sites but with variable selectivity, which offers an explanation of the puzzling lack of well defined structure-activity relationships among general anesthetics. More importantly, these findings indicate that it may be possible to synthesize general anesthetics with specificity for sites between specific subunits in the transmembrane domain of pentameric GABA_A receptors, with a hope to target specific nerve pathways and behaviours in a subunit-dependent manner in the future [165].

Effects of neurosteroids at GABAA receptors

Inhibitory neurotransmission mediated by GABA_A receptors can be modulated by endogenous neurosteroids, mostly metabolites of progesterone and deoxycorticosterine, including allopregnanolone and tetrahydrodeoxycorticosterone (THDOC). In general, they potently enhance function of synaptic and extrasynaptic GABA_A receptors by an allosteric mechanism [31,40,166,167]. Neurosteroids possess distinct, characteristic effects on the membrane potential and current conductance mainly via potentiation of GABA_A receptors at low doses, and direct activation of receptor chloride channel at higher concentrations. As a results, upon administration neurosteroids exert anxiolytic, analgesic, anticonvulsive, sedative and hypnotic effects, while applied at higher doses may induce a state of general anesthesia

[40,168-171]. Although the most important effects of neurosteroids are mediated via GABA_A receptors, they also exert various effects on an array of ligand-gated ion channels and distinct G-protein-coupled receptors via nongenomic mechanisms, including the *N*-methyl-D-aspartate (NMDA), alpha-amino-3-hydroxy-5-methyl-4-isoxazole-propionic acid (AMPA), kainate, glycine, serotonin, sigma type-1, and nicotinic acetylcholine receptors [156,172]. Besides modulation of diverse receptors, the other functions attributed to specific neurosteroids include neuroprotection, for example against glutamate-induced excitotoxicity, and induction of neurite outgrowth, dendritic spines and synaptogenesis, and contribution to the behavioural effects of psychoactive drugs [49,51,173-175].

As emphasized before, neurosteroids exhibit two modes of action at GABAA receptors. They allosterically modulate channel opening, and at high concentrations act as GABA-mimetics and directly open the chloride channel [31,53,176]. Two discrete binding sites in the receptor's transmembrane domains have been identified that mediate the potentiating and direct activation effects of neurosteroids. Neurosteroids potentiate GABA responses from a cavity formed by the α -subunit transmembrane domains, while direct receptor activation is achieved by interfacial residues between α and β subunits and is additionally enhanced by steroid binding to the potentiation site, indicating that activation of GABA_A receptors by neurosteroids relies on the occupancy of both sites [177]. Further work provided a more detailed kinetic and pharmacological characterization of the effects of mutations on channel activation and modulation by neurosteroids, leading to a model where residues in the TM1 membrane-spanning domain shape the binding surface to accommodate a variety of structurally distinct neurosteroids [178]. The potentiating effect of neurosteroids can be mediated by steroid interactions with its site within the same β - α pair that mediates receptor activation as well as the opposite β - α pair [179]. Chisary et al. [180] have shown that neurosteroids require a membranous route of access to transmembrane-domain binding sites that might have implications for the design of novel neuroactive steroids because their lipid solubility and related accessibility are probably the key determinants of receptor modulation [180]. This also suggests that by virtue of their high lipid solubility, µM concentrations of neurosteroids may be achieved locally [31].

Neuroactive steroids may act both as positive and negative modulators of GABA_A receptor function. Applied at concentrations below 300 nM, they are positive allosteric neuromodulators as they facilitate conformational transition of the GABA-gated chloride channel to an open state, increase mean channel open time and consequently increase GABA-elicited chloride currents [49,131,181-183]. Furthermore, they modulate binding of GABA

agonists, increase binding of positive modulators at benzodiazepine binding sites, and enhance TBPS binding in the absence of GABA [184-187]. On the contrary, sulphated endogenous steroids like pregnenolone sulphate and dehydroepiandrosterone sulphate (DHEAS) act as negative modulators at steroid binding site when applied at μ M concentrations [59]. For various neuroactive steroids sulfation at C-3 reverses the direction of modulation from positive to negative, suggesting that sulfation could be an important control point for the activity of endogenous neurosteroids. As interactions of positive and negative steroid modulators are not competitive, they probably act via distinct binding sites at GABA_A receptor. This implies that negative and positive steroid modulators can act independently or coordinately to regulate GABA-mediated inhibition in the CNS [188]. Antagonistic neurosteroids may shorten the channel open time and enhance GABA-induced desensitization [6].

Similarly to other binding sites, subunit assembly determines sensitivity and pharmacological effects of neuroactive steroids. Studies on recombinant GABA_A receptors have revealed that presence of δ subunit confers increases sensitivity to neurosteroid modulation by affecting intrinsic gating and desensitization kinetics [181,189]. Thus, at concentrations known to occur *in vivo*, neuroactive steroids specifically enhance a tonic inhibitory conductance in central neurons that is mediated by extrasynaptic δ subunit-containing GABA_A receptors [190]. In addition, expression of extrasynaptic GABA_A receptors is dynamically regulated by neuroactive steroids. For example, it is suggested that neuroactive steroids mediate a stress-induced enhancement in the expression of $\alpha 4\beta \delta$ receptors, probably contributing to observed stress-related memory impairment [41,44,191]. Potency and efficacy of neuroactive steroids is also determined by other subunits. Although these differences are not very pronounced, even small differences in receptor sensitivity might be physiologically relevant. Namely, allopregnanolone concentrations in plasma are 3-10 nM in physiological conditions, and increase only to 30-60 nM during stress [168].

Interestingly, it is confirmed that endogenous neurosteroids profoundly affect neuronal activity at particular receptor isoform. For example, it is shown that $\alpha 1\beta 2\delta$ subtype of GABA_A receptors presumably expressed in the forebrain, exhibits an extremely small GABA-mediated current in the absence of the modulator, but in the presence of THDOC exerts a profound inhibitory influence on neuronal activity by increasing maximum current amplitude and decreasing half-maximal concentration (EC₅₀) of GABA [192]. Hence, depending on the subunit composition, GABA-modulatory effects of physiological levels of the neurosteroid are not uniformly experienced throughout the CNS, or even within the same brain region. In

addition, these modulatory effects are determined by the phosphorylation status of the GABA_A receptor, or associated proteins, and by local steroid metabolism [31,49,54,193].

As endogenous ligands, neuroactive steroids may modulate diverse physiological and pathophysiological conditions, regulating GABA_A receptor-associated functions and behaviour. Altered neurosteroid levels are associated with psychiatric and neurological disorders, including premenstrual dysphoric disorder, premenstrual syndrome, menstrual migraine, postpartum depression, pain disorder, catamenial epilepsy, major depression, schizophrenia, alcohol dependence and anxiety [51,177,194]. The pathogenesis of these diseases remains unclear, partly due to the lack of useful animal models to study such complex disorders [46]. As mentioned before, pregnancy is associated with a pronounced enhancement in progesterone-derived neurosteroid levels, with sharp decline after delivery. Because neuroactive steroids exert anxiolytic effects, neurosteroid withdrawal could play role in the pathophysiology of postpartum depression [195]. Synthetic neurosteroids that exerts better bioavailability and efficacy, as well as drugs that enhance neurosteroid synthesis and lack unwarranted side effects of benzodiazepines, are therefore considered as potential therapeutics in the treatment of anxiety, epilepsy and other brain disorders [156,195-197].

Effects of ethanol at GABAA receptors

Ethanol is often classified as modulator of GABA_A receptor activity. It is without doubt that very high concentrations of ethanol modulate activity of diverse membrane proteins, including GABA_A receptors [198]. However, it is not quite clear whether social ethanol intake (<20 mM) really exerts effects on human behaviour via GABA_A receptors [5]. Wallner and co-authors [199] have demonstrated that GABA_A receptors might be enhanced by ethanol concentrations that are reachable by moderate, social consumption. They found that GABA_A receptors responsive to these low concentrations require presence of δ -subunit, which is thought to be associated exclusively with α 4- and α 6-subunits, and the β 3-subunit. Their finding also suggests that extrasynaptic GABA_A receptors are primary targets for ethanol. Later, Borghese and co-workers [198] failed to replicate the sensitivity of α 4 β 3 δ GABA_A receptors to low concentrations of ethanol at rat and human recombinant GABA_A receptors expressed in Xenopus oocytes. In addition, in their study ethanol at 30 mM concentration did not affect tonic GABA-mediated currents in dentate gyrus reported to be mediated by α 4 β 3 δ subtype of GABA_A receptors. Similarly, in another study, α 6 β 3 δ GABA_A receptors were expressed in Xenopus oocytes and were not modulated by physiological concentrations (up to 30 mM) of ethanol [200].

On the other hand, prolonged ethanol consumption that leads to the development of alcohol dependence induces changes of GABAA receptor subunits at transcriptional and translational levels in brain area-specific manner, including reduction in δ subunit expression in the orbitofrontal cortex, cerebellum, hippocampus and amygdala. Changes in receptor composition might have important consequences for GABAergic neurotransmission, plasticity and behaviour including impairment in memory, anxiety, and executive and motor functions [201-203]. Furthermore, it was found that ethanol at concentrations of 20 to 100 mM stimulates GABA-mediated uptake of chloride ions in isolated brain vesicles, and this effect can be blocked by the imidazobenzodiazepine Ro15-4513. Pre-treatment of rats with Ro15-4513 blocks the anticonflict activity of lower doses of ethanol, and behavioural intoxication observed with higher doses of ethanol. These effects of Ro15-4513 in antagonizing ethanolstimulated ³⁶Cl⁻ uptake and behaviour can be completely blocked by benzodiazepine binding site antagonists, indicating that neuropharmacological actions of ethanol (20-100 mM) may be mediated via central GABA receptors [111]. Similarly, it was shown that [³H]Ro15-4513 binding is inhibited only by those benzodiazepine-site ligands that reverse the behavioural alcohol antagonism of Ro15-4513 such as flumazenil and β-carboline-3-carboxylate ethyl ester (β -CCE), but not by any classical benzodiazepine agonists. Additional experiments indicated that ethanol and Ro15-4513 occupy mutually exclusive binding sites. Since only Ro15-4513, but not flumazenil, can inhibit ethanol effects, and taking into account that Ro15-4513 differs from flumazenil by only a single group at the C7 position of the benzodiazepine ring, it is assumed that this group in Ro15-4513 might be the area that overlaps with the alcohol-binding site. Hence, it turns out that many of the behavioural effects of ethanol at relevant physiological concentrations are mediated via ethanol/Ro15-4513-sensitive GABAA receptors [112].

Wallner and co-workers [98] showed that differences in alcohol sensitivity toward β subunits are determined by the extracellular N-terminal part of the protein. By using point mutations, they discovered that the β 3 alcohol selectivity is determined by a single amino acid residue in the N-terminus that differs between β subunits (β 3Tyr66, β 2Ala66, β 1Ser66). The β 3Tyr66 residue is located in a region which in γ subunits contributes to the imidazobenzodiazepine binding site at the classical α + γ 2- subunit interface. Furthermore, they proposed a model in which β 3 and δ containing GABA_A receptors form a unique ethanol site at the α 4/6+ β 3- subunit interface. As this site is homologous to the classical benzodiazepine

binding site, it also has high affinity for a few selected benzodiazepine site ligands including alcohol antagonistic Ro15-4513 that bears large moiety at the C7 position of the benzodiazepine ring. They also suggested that large moieties at the C7-benzodiazepine ring compete with alcohol for its binding pocket at a $\alpha 4/6+\beta 3$ - ethanol/Ro15-4513 site, thus finally providing explanation for the competitive relationship between ethanol and imidazobenzodiazepine antagonists.

Interactions of flavonoids with GABAA receptor

Flavonoids represent a heterogeneous class of plant secondary metabolites. They exert a wide-range of health-promoting effects including antioxidative, anti-inflammatory, cardioprotective, anticarcinogenic, antidiabetic and neuroprotective activities [204-206]. Like many neuroactive drugs, they achieve their effects, at least in part, by modulating GABAA receptors [207,208]. Radioligand binding studies indicated that flavonoids exert a selective affinity for the benzodiazepine binding site at GABA_A receptor complex, mostly acting as partial agonists [209]. Although it is initially thought that they act on classical, "high-affinity" benzodiazepine binding sites, many flavonoid actions are not inhibited by classical benzodiazepine antagonist flumazenil. For example, Hall et al. [210] have shown that 6methoxyflavanone and 6-methoxyflavone act as flumazenil-insensitive positive allosteric modulators of GABA responses at human recombinant $\alpha 1\beta 2\gamma 2L$ and $\alpha 2\beta 2\gamma 2L$ GABAA receptors. Functional electrophysiological studies suggest that flavonoids act on GABAA receptors via two separate mechanisms: by acting on flumazenil-sensitive high-affinity site and an alternative site that may be the flumazenil-insensitive low-affinity benzodiazepine site [208]. However, as in aforementioned study 6-methoxyflavanone inhibited [³H]flunitrazepam binding whilst remaining unaffected by flumazenil, this might suggest a novel allosteric binding site that is independent of both the high-affinity and low affinity benzodiazepine binding sites [210].

More importantly, some flavonoids may directly open certain subtypes of GABA_A receptors in the absence of GABA [208]. Behavioural studies demonstrated anxiolytic effects of flavonoids in animal models. Some of the tested flavonoids (e.g. chrysin) have anxiolytic effects similar to diazepam, but not associated with myorelaxant, sedative or amnesic actions, suggesting that they could lead to improved therapeutic drugs in the treatment of anxiety [211,212]. However, in addition to positive modulation, flavonoids may also achieve negative

effects on GABA_A receptors [213,214]. Some flavonoids are positive neuromodulators when applied in the presence of low concentrations of GABA, but in the presence of high concentrations of GABA act as negative modulators [215]. A range of natural flavonoids act as negative modulators of receptors containing ρ -subunits.

Recently, it was found that quercetin and its glycosides inhibit GABA-induced inward current at recombinant GABA_C receptors. These inhibitory effects of quercetin and quercetinglycosides on GABA-induced inward current were noncompetitive and membrane voltageinsensitive, indicating that quercetin and its glycosides regulate GABA_C receptor channel activity through interaction with a different site from that of GABA [216]. This study also provides evidence that the number of carbohydrate attached to quercetin might play an important role in the regulation of GABA_C receptor channel activity. On the contrary, 6methoxyflavanone and 6-methoxyflavone were inactive as modulators at human recombinant receptors containing ρ1 subunits [210].

Interestingly, menthol, a naturally occurring compound in the essential oil of mint leaves, also act as a positive allosteric modulator of recombinant GABA_A receptors. In particular, menthol (150-750 μ M) produced a concentration-dependent prolongation of spontaneous GABA_A receptor-mediated inhibitory postsynaptic currents in the periaqueductal grey (PAG) neurons, although menthol actions were unaffected by the benzodiazepine antagonist flumazenil. Menthol also enhanced a tonic current, which was sensitive to the GABA_A receptor antagonists picrotoxin and bicuculline, but unaffected by gabazine and GABA_C receptor antagonist (1,2,5,6-tetrahydropyridine-4-yl)-methyl-phosphonic acid (TPMPA). In addition, menthol potentiated currents induced by the extrasynaptic GABA_A receptor agonist THIP, altogether indicating that menthol positively modulates both synaptic and extrasynaptic populations of GABA_A receptors in native PAG neurons [217].

GABA_C receptors

GABA_C receptors are considered as a specific subtype of GABA_A receptors due to their structural homology, although they differ from GABA_A receptors by biochemical, pharmacological and physiological properties [218-220]. They are predominantly expressed in the retina where play a unique functional role in retinal signal processing, but are also expressed throughout brain and in periphery [221,222]. GABA_C receptors are involved in numerous processes across the CNS, including vision, olfactory senses, sleep, memory and cognitive functions, hormone secretion and pain perception, with promising potential for the treatment of myopia, sleep disorders, memory and learning enhancement, and fear and anxiety-related disorders [222,223]. Although understanding of the role of GABA_C receptors and processes triggered by ligand–receptor interactions in neurons are still limited because of the lack of adequate pharmacological tools, some progress has been made following synthesis and pharmacological evaluation of the selective fluorescent and biotinylated probes for ρ_1 GABA_C receptors that hopefully will bring new knowledge regarding the binding site and GABA_C receptors flexibility, and be useful tools for localizing, visualizing, and studying the physiopathological processes of GABA_C receptors [224].

In mammals, there are three isoforms of ρ subunits (ρ 1- ρ 3) that form homooligomeric (formed by ρ 1, ρ 2 or ρ 3 isoform) or pseudoheteromeric (made up of a combination of ρ 1 and ρ 2 isoforms or ρ_2 and ρ_3 isoforms) GABA_C receptor pentamers [220,224]. In humans, only two isoforms are expressed: ρ 1 and ρ 2.

GABA_C receptors are sensitive neither to bicuculline (characteristic ligand of GABA_A receptors) nor to baclofen (characteristic ligand of GABA_B receptors) [222,225]. They are more sensitive to GABA than GABA_A receptors, and at least three GABA molecules are required to activate the GABA_C receptor [222]. When activated, they have a smaller chloride conductance, longer channel opening time and desensitize less readily in the presence of GABA then GABA_A receptors [218]. Benzodiazepines and barbiturates do not modulate GABA_C receptors [218]. Neuroactive steroids may modulate GABA-induced current at GABA_C receptors, in particular at p1 receptor channels, in both positive and negative manner, although the modulation occurs with relatively high concentrations of neuroactive steroids and is more prominent in the presence of low concentrations of GABA [226]. Characteristic agonists of GABA_C receptors are cis enantiomer of 4-aminocrotonic acid (CACA) and (+)cis-2-aminomethylcyclopropane carboxylic acid ((+)-CAMP) [218]. Furthermore, GABAC receptors are much less sensitive to GABA_A receptor antagonist gabazine, but can be selectively antagonized by (1,2,5,6-tetrahydropyridine-4-yl)-methyl-phosphinic acid and 3-(TPMPA), 3-aminopropyl(methyl) phosphinic acid (3-APMPA), aminopropylphosphonic acid (3-APA), indicating that agonist/antagonist binding pockets of GABA_A and GABA_C receptors are not the same [220,227,228]. Compound imidazole-4-acetic acid exhibits a ρ subunit-dependent pharmacological profile: at $\rho 1$ and $\rho 3$ receptors acts as a potent antagonist, while at ρ^2 receptors as a potent partial agonist [222].

N-terminal half of the ρ subunits has been shown to mediate formation of homo- and heterooligomeric GABA_C receptors, and specific sequence within the N-terminus of the ρ 1 subunit involved in the assembly process has been determined [229]. Furthermore, several

key structural elements that determine specific pharmacological response of GABA_C receptors have been found [228]. Namely, mutational studies, including those directed toward N-terminal domain and transmembrane domain TM4, have revealed residues that change sensitivity to agonists or make GABA_C complex inactive, and contribute to the binding pocket determining properties of GABA binding [221,230,231]. Thus, Tyr102 at p1 subunit was identified as part of GABA binding domain, and probably the important residue for coupling agonist binding to channel opening [232]. Barbiturate sensitivity was imparted by mutation of Trp328 at p1 subunit, located within the transmembrane domain TM3. It also seems that this residue plays an important role in agonist-dependent activation, suggesting a functional interconnection between the GABA and pentobarbital activation domains [233]. GABAC receptors that contain ρ^2 subunit exhibit decreased sensitivity to picrotoxin [234]. On ρ^1 receptor it is shown that the mechanism of picrotoxin effects is compatible with an allosteric inhibition and receptor activation was a prerequisite for antagonism [235]. Difference in picrotoxin sensitivity of p1 and p2 homo-oligomers is determined by a single residue. Interestingly, it was found that this amino acid in the putative channel domain (TM2) of GABA p1 receptors influences picrotoxin sensitivity, and also meditates agonist binding by an allosteric mechanism [236].

GABA_B receptors

Metabotropic GABA_B receptors are G-protein coupled receptors that mediate slow and prolonged inhibitory neurotransmission in the brain [237]. They are widely expressed and distributed in the CNS, although GABA_A sites generally outnumber GABA_B sites [238]. GABA_B receptors differ from GABA_A receptors by their structural and functional properties, but also exhibit numerous pharmacological effects, including central muscle relaxation, epileptogenesis, suppression of drug craving, antinociception, cognitive impairment and inhibition of hormone release [12,239].

GABA_B receptors can be located presynaptically and postsynaptically. Stimulation of presynaptic GABA_B receptors decreases conductance of calcium ions via voltage-gated calcium channels. Consequently, activation of presynaptic GABA_B autoreceptors induces inhibition of GABA release, while presynaptic GABA_B heteroreceptors suppress release of other neurotransmitters and bioactive peptides. Depending on whether synaptic terminal releases an inhibitory or excitatory neurotransmitter, presynaptic GABA_B receptors will facilitate or suppress neuronal excitability thus playing important role in tuning various

synapses [12,240,241]. On the other hand, postsynaptic GABA_B receptors are coupled via Gproteins to inwardly rectifying potassium channels and underlie slow inhibitory postsynaptic currents. Postsynaptic GABA_B receptors stimulate efflux of potassium ions that hyperpolarizes neuronal membrane and shunts excitatory currents. As presynaptic and postsynaptic GABA_B receptors are dominantly located at extrasynaptic sites, their activation requires patterns of presynaptic activity that ends in simultaneous GABA spillover from neighbouring synapses and elevations of ambient GABA, as it occurs during epileptic seizures. Furthermore, the association of GABA_B receptors with glutamatergic synapses suggests their important role in the modulation of glutamatergic neurotransmission [240,242,243].

The GABA_B receptor was the first heteromeric G-protein coupled receptor identified. It functions as an obligatory heterodimer assembly: both GABA_{B1} and GABA_{B2} subunits are necessary to form a functional GABA_B receptor, and neither of these subunits is functional on its own [244] (Kaupmann et al. 1998). Soon after the cloning of both subunits, it was demonstrated that GABA_{B2} is required for GABA_{B1} to reach the cell surface by masking an endoplasmic reticulum retention signal of GABA_{B1} [245]. However, GABA_{B2} is not only required for the correct trafficking of GABA_{B1}, but also for the proper functioning of the receptor. In particular, GABA_{B1} is involved in ligand recognition via its N-terminal extracellular domain and it binds GABA [246]. Although GABA_{B2} subunit does not constitute a binding site for any natural GABA_B ligand [247], it enhances agonist affinity [248], and is required for receptor activation. In addition, GABA_{B2} subunit is responsible for G-protein coupling [249].

There are two physiologically significant isoforms of $GABA_{B1}$ subunit, $GABA_{B1a}$ and $GABA_{B1b}$ that differ in N-terminal region. Expression of these two subunits is developmentally regulated at the transcription level. They are transcribed from the same gene after activation of alternative promoters [250]. Accordingly, there are two major subtypes of $GABA_B$ receptor, one formed from $GABA_{B2}$ and $GABA_{B1a}$ subunits, and the other formed from $GABA_{B2}$ and $GABA_{B1a}$ subunits [240].

 $GABA_{B1}$ and $GABA_{B2}$ subunits are structurally homologous and both possess two main domains: a heptahelical membrane domain that is responsible for recognition and activation of G-proteins (HD domain), and a large extracellular "Venus flytrap" domain (VFT domain) involved in ligand binding. Both domains oscillate between different conformational states, and these allosteric transitions are essential for receptor function and offer numerous possibilities for the allosteric regulation of receptor activity. GABA and other agonists bind exclusively at the VFT domain of the GABA_{B1} subunit, but VFT domain of the GABA_{B2} subunit is necessary for the activation of whole receptor. On the other hand, GABA_{B2} HD contains molecular determinants required for G-protein coupling, and by trans-activation mechanism, binding of GABA at VFT domain of GABA_{B1} subunit lead to activation of HD domain of GABA_{B2} subunit [251-253]. However, the HD of GABA_{B1} improves coupling efficacy. Conversely, although GABA_{B1} extracellular domain is sufficient to bind GABA_B ligands, the extracellular domain of GABA_{B2} increases the agonist affinity on GABA_{B1}, and is necessary for agonist activation of the receptor. Altogether, this indicates that multiple allosteric interactions between the two subunits are required for wild-type functioning of the GABA_B receptor [251].

Geng et al. [254] presented the crystal structures of heterodimeric GABA_B complex consisting of extracellular VFT domains of GABA_{B1} and GABA_{B2} subunits in the apo, agonist-bound and antagonist-bound forms. The apo and antagonist-bound structures represent the resting state of the receptor; while the agonist-bound complex corresponds to the active state. They found that both subunits adopt an open conformation at rest, but only GABA_{B1} VFT closes on agonist-induced receptor activation. Furthermore, they revealed a unique activation mechanism for GABA_B receptor that involves the formation of a novel heterodimer interface between subunits.

GABA_B receptors are not modulated by benzodiazepines, barbiturates, or steroids, and are not sensitive to bicuculline [218,225]. Characteristic agonists of GABA_B receptors are baclofen, a lipophilic derivative of GABA, and 3-aminopropylphosponous acid (3-APPA; CGP27492), while saclofen, phaclofen and 2-hydroxysaclofen act as antagonists of GABA_B receptors [12]. Baclofen was introduced to the market in 1972 and is used to treat spasticity and skeletal muscle rigidity in patients with spinal cord injury, multiple sclerosis, amyotrophic lateral sclerosis, and cerebral palsy. However, although GABA_B agonists showed promising therapeutic effects in a whole range of other indications, they exhibit numerous side effects, including sedation, tolerance, and muscle relaxation [237,256].

Conclusions

GABA_A receptors mediate most of the fast synaptic inhibition in the mammalian brain and are targeted by many clinically important drugs. They are subject to modulation at a variety of allosteric sites, with pharmacology dependent on receptor subunit combination. In addition to GABA, naturally occurring steroids can potently and specifically enhance GABA_A receptor function in a direct manner, and consequently exert anxiolytic, analgesic, anticonvulsant, sedative, hypnotic and anesthetic properties. GABA_A receptors are thus important for function and plasticity of the CNS. Analysis of the specific roles of GABA_AR subtypes reveals their involvement in the pathophysiology of major CNS disorders, and opens novel perspectives for therapeutic intervention. Further pharmacological studies will contribute to more complete understanding of numerous interactions between various ligands and their binding sites on GABA receptors that might improve current pharmacological approach in treating various diseases.

Acknowledgments

This work was supported by Croatian Ministry of Science, Education and Sports to MJJ and JV.

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