

$\alpha 4\beta\delta$ GABA_A receptors are high-affinity targets for γ -hydroxybutyric acid (GHB)

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γ -Hydroxybutyric acid (GHB) binding to brain-specific high-affinity sites is well-established and proposed to explain both physiological and pharmacological actions. However, the mechanistic links between these lines of data are unknown. To identify molecular targets for specific GHB high-affinity binding, we undertook photolinking studies combined with proteomic analyses and identified several GABA_A receptor subunits as possible candidates. A subsequent functional screening of various recombinant GABA_A receptors in *Xenopus laevis* oocytes using the two-electrode voltage clamp technique showed GHB to be a partial agonist at $\alpha\beta\delta$ - but not $\alpha\beta\gamma$ -receptors, proving that the δ -subunit is essential for potency and efficacy. GHB showed preference for $\alpha 4$ over $\alpha(1,2,6)$ -subunits and preferably activated $\alpha 4\beta 1\delta$ (EC₅₀ = 140 nM) over $\alpha 4\beta(2/3)\delta$ (EC₅₀ = 8.41/1.03 mM). Introduction of a mutation, $\alpha 4F71L$, in $\alpha 4\beta 1(\delta)$ -receptors completely abolished GHB but not GABA function, indicating nonidentical binding sites. Radioligand binding studies using the specific GHB radioligand [³H](*E,RS*)-(6,7,8,9-tetrahydro-5-hydroxy-5H-benzocyclohept-6-ylidene)acetic acid showed a 39% reduction (*P* = 0.0056) in the number of binding sites in $\alpha 4$ KO brain tissue compared with WT controls, corroborating the direct involvement of the $\alpha 4$ -subunit in high-affinity GHB binding. Our data link specific GHB forebrain binding sites with $\alpha 4$ -containing GABA_A receptors and postulate a role for extrasynaptic $\alpha 4\delta$ -containing GABA_A receptors in GHB pharmacology and physiology. This finding will aid in elucidating the molecular mechanisms behind the proposed function of GHB as a neurotransmitter and its unique therapeutic effects in narcolepsy and alcoholism.

γ -hydroxybutyric acid receptor | γ -hydroxybutyric acid high-affinity binding sites | $\alpha 4$ -subunit knockout | photoaffinity ligand

The GABA metabolite γ -Hydroxybutyric acid (GHB) is present in micromolar concentrations in the mammalian brain, where it has been proposed to act as a neurotransmitter (1). Additionally, GHB is a drug of abuse (Fantasy) and a registered drug for treating narcolepsy (2) and alcoholism (3). GHB binds to at least two distinct populations of low- and high-affinity binding sites in the brain (4). When GHB is ingested in high doses and reaches millimolar concentrations in the brain, it induces behavioral effects such as sedation, motor incoordination and hypothermia (3). These actions are largely mediated by metabotropic GABA_B receptors, because effects are prevented by GABA_B receptor antagonist pretreatment (5) and completely abolished in GABA_{B(1)} KO mice (6). In addition to the validated GABA_B receptor effects and other suggested receptors (7), GHB binds with nanomolar to micromolar affinity to a remarkably abundant protein of distinct spatial distribution and ontogenesis (4), representing an additional functional target. Interestingly, this high-affinity binding protein is preserved in GABA_{B(1)} KO mice (6) and can be specifically probed with [³H](*E,RS*)-(6,7,8,9-tetrahydro-5-hydroxy-5H-benzocyclohept-6-ylidene)acetic acid ([³H]NCS-382) (6) and [¹²⁵I]4-hydroxy-4-[4-(2-iodobenzoyloxy)phenyl]butanoate ([¹²⁵I]

BnOPh-GHB) (8). Furthermore, several reports point to GHB-induced effects that cannot be consequences of GABA_B receptor activation alone: Fos expression studies with GHB indicate a unique pattern of neuronal activation, which in several ways, is different from the pattern produced by the GABA_B receptor agonist baclofen (9). Numerous effects induced by GHB, including sedation, catalepsy (10), increased striatal dopamine release, changes in EEG pattern (11), discriminative stimulus properties (12), and reinforcing effects (13), are dose-dependently decreased by pretreatment with the GHB receptor-specific ligand NCS-382. Additionally, ataxia seems to be mediated through the high-affinity GHB sites (14). The reported euphoric effect of GHB and its therapeutic effect in narcolepsy cannot be mimicked by baclofen (3) and thus, might involve other targets. Drug discrimination studies also show that rats are able to distinguish between GHB and baclofen, further supporting that the effects and mechanisms of the two drugs are different (15). Taken together, these findings strongly suggest that GHB acts at targets in addition to the GABA_B receptor.

Structurally and behaviorally, GHB, in many ways, resembles its endogenous precursor GABA. Thus, in addition to effects at GABA_B receptors, ionotropic GABA_A receptors have also been studied as possible GHB targets. The role of GABA_A receptors in mediating effects of GHB has been controversial, in part because of the heterogeneity of this receptor class and the lack of recombinant functional studies performed; thus, the large number of subtype combinations that can be formed from the numerous known subunits [$\alpha(1-6)$, $\beta(1-3)$, $\gamma(1-3)$, δ , ϵ , θ , π , and $\rho(1-3)$] (16) have not been investigated.

Depending on composition, GABA_A receptors can be found at both synaptic and extrasynaptic locations and mediate phasic and tonic inhibition, respectively (17). The majority of GABA_A receptors contain a γ -subunit, and these receptors can be found at both synaptic and extrasynaptic locations, whereas the δ -subunit predominates on peri- and extrasynaptic locations (18, 19), most commonly accompanied by $\alpha(4/6)$ -subunits.

In 1987, the work by Snead and Nichols (20) reported evidence for coupling of the GHB binding site to a GABA-gated chloride channel; whereas effects of GHB on the major GABA_A (synaptic) receptors have been refuted (21, 22), effects at

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extrasynaptic receptors have, until now, not been systematically investigated. In fact, several studies infer a role for GHB at extrasynaptic GABA_A receptors, such as a correlation between elevated GHB levels and increased tonic extrasynaptic inhibition through GABA_A receptors (23). More specifically, effects involved receptor subtypes containing α 4- and δ -subunits (24–26).

In this study, we have exploited an in-house-developed, high-affinity, and selective GHB photoligand (8, 27, 28) to cross-link and partially purify the high-affinity GHB binding protein from rat brain cortex, with several GABA_A subunits emerging as candidate proteins. Functional studies in *Xenopus laevis* oocytes and radioligand binding studies in KO mouse brain tissue verified α 4 β δ -receptors as high-affinity targets for GHB. Thus, we present direct molecular evidence for a GHB–GABA_A receptor interaction in both recombinant and native systems.

Results

Proteomics Identify GABA_A Receptor Subunits as Candidates for High-Affinity GHB Binding Sites. Using an engineered GHB photoaffinity ligand (28), we previously determined the high-affinity GHB binding protein to be ~50 kDa (8). In attempt to provide better validity in the identifications made by the proteomics analysis, we also subjected samples to limited proteolysis by time-dependent treatment with proteinase K, which resulted in minor bands of ~28, ~21, and ~18 kDa (Fig. 1). High-resolution orbitrap mass spectrometric analysis by nanoscale liquid chromatography–tandem MS spectra of each of these bands identified several GABA_A receptor subunits: α 1, α 2, α 3, α 5, β 1, β 2, β 3, and γ 1 (Table S1 and Dataset S1).

GHB Is a Partial Agonist at Particular Subtypes of Recombinant GABA_A Receptors. Prompted by the proteomics results, we systematically investigated the effects of GHB at different human recombinant GABA_A receptor subtypes expressed in *X. laevis* oocytes. At various synaptic and extrasynaptic receptor combinations (α 1 β 2 γ 2L, α 5 β 3 γ 2L, α 2 β 1 δ , α 1 β 3 δ , and ρ 1), GHB at concentrations of 1 mM

and higher was without effect. However, when oocytes were injected with the combination of α 4-, β (1–3)-, and δ -subunits, GHB induced inward currents, exhibiting both an intriguing dependence on α 4/ δ for efficacy and β 1 for potency (Table 1). GHB activated α 4 β 1 δ receptors with high nanomolar potency [EC_{50} = 140 nM (30–660)], inducing a maximum current of $74 \pm 10\%$ relative to GABA (Fig. 2A and C). At α 4 β 1 and α 4 β 1 γ 2L, 3 mM GHB elicited a small response ($3 \pm 0.2\%$, $P = 0.01$ and $2 \pm 2\%$, $P = 0.31$, Z test compared with control, respectively) (Fig. 2B and Table 1). In oocytes injected with RNA for neither α 4 and β 1 nor α 4, β 1, and δ did we find any indication that GHB could antagonize the GABA response (Fig. S1). Substitution of the β 1-subunit with β 2 or β 3 led to a slight reduction in the relative agonist efficacy (53–76%) but a >7,000-fold reduction in potency [EC_{50} = 8.4 mM (4.0–17) and 1.0 mM (0.6–2.8), respectively] (Fig. 2C). Construction of a current–voltage (I–V) curve in *X. laevis* oocytes expressing α 4 β 1 δ -receptors confirmed that GHB activated a chloride channel (Fig. 2D). Effects of GHB were undetectable in α 4 β 3-, α 4 β 3 γ 2L-, and α 4 β 2 γ 2L-receptors (Table 1). By contrast, when coexpressing α 1-, α 2-, or α 6-subunits with β 1 δ , currents induced by GHB were dramatically reduced (α 1 and α 6) compared with α 4 (Table 1) or completely absent (α 2).

Molecular Pharmacology of the GHB α 4 β 1 δ -Receptor Interaction. The pharmacology of the α 4 β 1 δ subtype in relation to GHB was further investigated. The response induced by 30 μ M GHB was completely blocked by coapplication of the GABA_A receptor antagonist gabazine (1 μ M) (Fig. 3A). In attempts to delineate the GHB binding site, we investigated the role of a conserved α -subunit Phe residue (F64 in α 1 and F71 in α 4). This residue lines the GABA binding site of α β γ -receptors, and when mutated to Leu, it gives a marked (>200-fold) increase in GABA EC_{50} at α 1 β 2 γ 2 GABA_A receptors (29). Interestingly, introduction of a mutated α 4F71L-subunit completely abolished GHB activity at α 4 β 1 and α 4 β 1 δ compared with WT receptors (Fig. 3B). By contrast, GABA EC_{50} values were only increased about fivefold for α 4 (F71L) β 1 ($P = 0.058$) and α 4(F71L) β 1 δ ($P = 0.001$) (Table 1).

Radioligand Binding Studies to Link GHB Binding and Function. To link GHB function at α 4 β δ -receptors with the GHB high-affinity binding site, we probed numerous GABA_A ligands for their ability to inhibit high-affinity GHB binding ($[^3H]NCS-382$ binding) (Table S2). Only one of these ligands, gabazine, inhibited binding with an IC_{50} in the mid micromolar range, whereas the IC_{50} for GABA was in the low millimolar range (Fig. 4A). Additionally, by autoradiography, we found that gabazine inhibited $[^{125}I]BnOph-GHB$ binding in a regionally specific manner similar to GHB (Fig. 4B) (8). Next, we investigated $[^3H]NCS-382$ binding in brain tissue from α 4 and δ KO mice and WT littermates. Whereas $[^3H]NCS-382$ binding did not differ statistically between δ KO and WT mice (Fig. 4C and D), a significant reduction in B_{max} in α 4 KO membranes compared with WT was observed ($P < 0.01$) (Fig. 4E and F and Table 2, mean \pm SEM).

Discussion

The present study identifies α 4 β δ -receptors, particularly α 4 β 1 δ -receptors, as high-affinity targets for GHB, which are likely to represent the elusive GHB receptor. The dependence of α 4- and δ -subunits for eliciting a GHB-induced response and the remarkable selectivity for the β 1-subunit serve as a base for hypotheses regarding the unique properties of GHB. Our finding is supported by several studies, indicating a possible role for GHB at extrasynaptic α 4-containing GABA_A receptors. A study using metabolic fingerprinting recently reported that the actions of GHB are similar to several ligands acting at extrasynaptic GABA_A receptors (25). A link between GHB and α 4 has also been indicated by the ability of GHB to antagonize increases in α 4 mRNA levels induced by ethanol withdrawal (24) and the association of

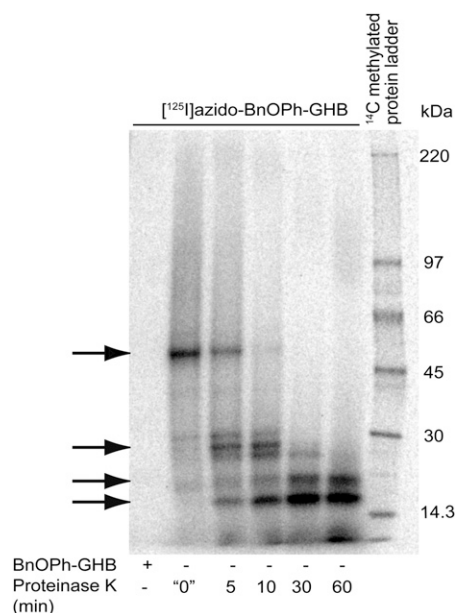


Fig. 1. Photoaffinity labeling of high-affinity GHB binding sites from rat brain and isolation of target proteins. SDS/PAGE separations of $[^{125}I]$ azido-BnOph-GHB radiophotoaffinity-labeled partially degraded binding proteins using time-dependent Proteinase K limited proteolysis. Arrows indicate the four protein bands isolated for proteomics analysis (top to bottom: ~50, ~28, ~21, and ~18 kDa).

Table 1. Two-electrode voltage clamp recordings of GHB-elicited currents from *X. laevis* oocytes expressing different $\alpha(1,4,6)$ - and $\alpha 4F71L$ -containing GABA_A receptors

	$I/I_{GABA, max} \pm SEM$	EC_{50} (95% CI; M)	n_H (95% CI)	n
$\alpha 4\beta 1\delta$	0.74 ± 0.1	1.4×10^{-7} ($0.3-6.6 \times 10^{-7}$)	0.52 (0.09-0.95)	5
$\alpha 4\beta 1\gamma 2L$	$0.02 \pm 0.02^*$	†	†	4
$\alpha 4\beta 1$	$0.03 \pm 0.002^*$	†	†	4
$\alpha 4\beta 2\delta$	0.53 ± 0.06	8.4×10^{-3} ($4.0-17 \times 10^{-3}$)	1.6 (0.52-1.8)	4
$\alpha 4\beta 2\gamma 2L$	†	†	†	5
$\alpha 4\beta 2$	‡	‡	‡	22
$\alpha 4\beta 3\delta$	0.76 ± 0.08	1.0×10^{-3} ($0.6-2.8 \times 10^{-3}$)	1.3 (0.6-1.2)	6
$\alpha 4\beta 3\gamma 2L$	†	†	†	5
$\alpha 4\beta 3$	†	†	†	5
$\alpha 1\beta 1\delta$	$0.02 \pm 0.01^{\S}$	¶	¶	8
$\alpha 6\beta 1\delta$	$0.06 \pm 0.02^{\S}$	¶	¶	10
Mutant data				
$\alpha 4F71L\beta 1$ GHB	†	†	†	6
$\alpha 4F71L\beta 1\delta$ GHB	†	†	†	10
$\alpha 4F71L\beta 1$ GABA	§	7.2×10^{-7} ($5.5-9.3 \times 10^{-7}$)	0.89 (0.7-1.1)	4
$\alpha 4F71L\beta 1\delta$ GABA	§	1.8×10^{-7} ($1.3-2.6 \times 10^{-7}$)	1.03 (0.7-1.4)	6

*Normalized current at 3 mM GHB.

†Because of high estimated EC_{50} value, no curve could be generated.

‡GABA-induced currents were not high enough to estimate GHB responses.

§GABA concentrations used to determine maximum current for each subunit were, respectively, 10 μ M, 1 mM, 10 μ M, 1 μ M, 1 mM, 10 μ M, 10 μ M, 1 mM, 10 μ M, 10 mM, and 1 mM.

¶ I/I_{GABA} not applicable.

chronically elevated GHB levels seen in succinic semialdehyde dehydrogenase KO mice with increased tonic inhibition at extrasynaptic GABA_A receptors (23).

Similar to other δ -subunit-preferring GABA_A agonists, such as 4,5,6,7-tetrahydroisoxazolo[5,4-c]pyridin-3-ol (THIP or gaboxadol) (30), we find that the δ -subunit plays a clear potentiating role for GHB function at recombinant $\alpha 4\beta \delta$ -receptors. It is tempting to speculate that this finding could be physiologically relevant. Indeed, in δ KO mice, systemic administration of GHB fails to induce spike-and-wave discharges and epileptic absence seizures (26), supporting that the δ -subunit is also required for GHB-induced responses in vivo. In contrast to its essential role for

function, the δ -subunit seems not to be crucial for binding, because no significant difference in binding levels was found between WT and δ KO mice. By contrast, the $\alpha 4$ -subunit seems to be an important determinant for [³H]NCS-382 binding, which was indicated by a 39% reduction of B_{max} in $\alpha 4$ KO compared with WT tissue. The residual [³H]NCS-382 binding may be explained by compensatory mechanisms such as up-regulation of other GABA_A receptor subunits (31) or the ability of [³H]NCS-382 to bind to additional GABA_A interfaces. This latter point is supported by the ability of GHB to activate not only $\alpha 4\beta 1(\delta)$ but also $\alpha 1\beta 1\delta$ - and $\alpha 6\beta 1\delta$ -receptors. Although a compensatory mechanism in δ KO mice could possibly mask an involvement of the δ -subunit in ligand binding, our proteomics data do not support this hypothesis.

The finding that $\alpha 4F71$ is a key residue in mediating GHB activity at $\alpha 4\beta 1\delta$ -receptors holds interesting implications for additional dissection of the GHB binding site. The only fivefold reduction in GABA EC_{50} inflicted by the mutation indicates that GABA interacts differently in δ -containing compared with γ -containing receptors, which was recently reported (32), and it suggests that GHB and GABA could have distinct but overlapping binding sites. This finding would also explain why GHB displays partial agonism at $\alpha 4\beta 1\delta$ -receptors compared with GABA but does not antagonize the GABA response. Alternatively, the maximal response for GABA in oocytes injected with RNA for $\alpha 4$, $\beta 1$ and δ could well be the result of a mixture of currents evoked from individual $\alpha 4\beta 1$ - and $\alpha 4\beta 1\delta$ -receptors (32), which would underestimate GHB efficacy. It is also possible that posttranslational modifications occur or variant stoichiometric forms of the $\alpha 4\beta 1\delta$ -receptors are expressed that have different pharmacological properties. This finding would result in relatively larger errors in the efficacy of compounds for $\alpha 4\beta 1\delta$ -receptors compared with $\alpha 4\beta 1$ -receptors, which seems to be the case for GHB.

From the current data, we propose that NCS-382/GHB binds to the β - α interface in a site also recognized by GABA and gabazine but notably, in a way alternative to their high-affinity binding modes (33). Similar to effects of THIP (30), the δ -subunit plausibly increases GHB sensitivity by inducing allosteric effects in relation to either receptor binding or gating. The remarkable abundance of [³H]NCS-382 binding sites in fore- and midbrain regions (20-30

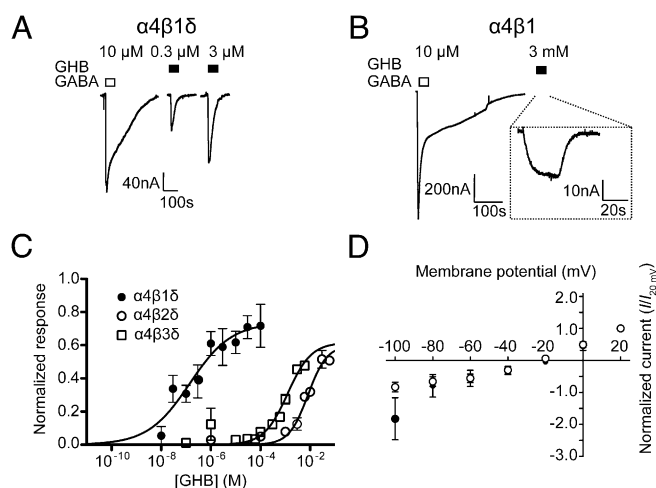


Fig. 2. Pharmacological characterization of GHB at recombinant $\alpha\beta\delta$ -receptors in *X. laevis* oocytes. Representative GHB current traces at (A) $\alpha 4\beta 1\delta$ and (B) $\alpha 4\beta 1$ GABA_A receptors. (C) Concentration response curve at $\alpha 4\beta 1\delta$ -receptors normalized to GABA_{max} (means \pm SEM; $n = 4-6$). (D) GHB (closed circles) and GABA (open circles) I-V relationships at $\alpha 4\beta 1\delta$ -receptors ($n = 7$). Reversal potential was not significantly different for GABA and GHB currents ($V_{rev} = -25.2 \pm 3.9$ and -27.8 ± 4.8 , respectively; $P = 0.68$, Student t test).

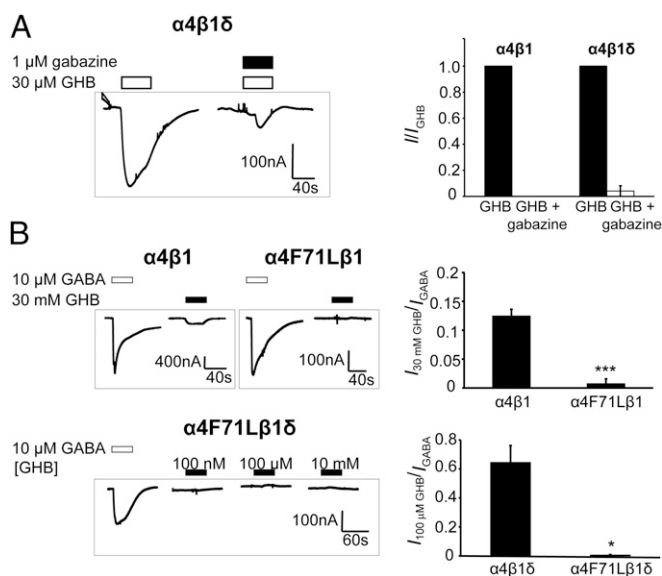


Fig. 3. Abolishment of GHB response by gabazine coapplication and $\alpha 4F71L$ point mutation. (**A, Left**) Representative gabazine inactivation trace of GHB currents at $\alpha 4\beta 1\delta$. (**A, Right**) Summarized data displaying fraction of GHB current at $\alpha 4\beta 1\delta$ (means \pm SEM; $n = 3$). (**B, Left**) Representative traces of GABA and GHB-elicited currents from $\alpha 4\beta 1$, $\alpha 4(F71L)\beta 1$ (Upper), and $\alpha 4(F71L)\beta 1\delta$ (Lower). (**B Right**) Summary (means \pm SEM) of 30 mM GHB effects at $\alpha 4\beta 1$ vs. $\alpha 4(F71L)\beta 1$ ($***P = 6.3 \times 10^{-5}$) and 100 μ M GHB at $\alpha 4\beta 1\delta$ vs. $\alpha 4(F71L)\beta 1\delta$ ($*P = 0.011$).

pmol/mg; present study) (6) may, thus, be explained by binding to a number of β - α interfaces that seem functionally silent in the absence of the δ -subunit, an observation also supported by the variety of α - and β -subunits in our proteomics analyses.

The markedly higher GHB sensitivity at $\beta 1$ -containing vs. $\beta 2/3$ -containing $\alpha 4\delta$ GABA_A receptors makes analogs designed specifically for the high-affinity GHB binding site (27, 34) interesting as pharmacological tools for studying $\beta 1$ -containing receptors. To date, only few such compounds have been reported (35, 36). The preference for $\beta 1$ may, in part, explain the sleep-mediating effects of GHB, because the endogenous pathway of sleep has been shown to depend mainly on $\beta 1$ -containing GABA_A receptors, whereas anesthesia and hypnosis are mediated through $\beta 3$ -containing receptors (37, 38). $\beta 1$ is associated with extrasynaptic receptors (39) and abundantly found in the hippocampus and cortex (40), where high-affinity GHB binding sites are predominantly located (6). Although less abundant, $\alpha 4$ and δ are also present in hippocampus and cortex (40). $\beta 1$ (and to a lesser degree, $\alpha 4$ and δ) is present in several brain regions important for modulation of sleep and facilitation of EEG synchronization such as the reticular thalamic nucleus (37), which may underlie the unique therapeutic effect of GHB in narcolepsy. Furthermore, a correlation has been shown between activity at $\beta 1$ -subunit-containing GABA_A receptors and ataxia, which is a proposed GHB receptor-mediated effect, because it is induced by GHB receptor-specific ligands and not antagonized by GABA_B antagonists (14). The $\alpha 4\beta 1\delta$ -subtype has been identified as an important target for endogenous neurosteroids, and receptor expression is modulated by fluctuating levels of these subtypes throughout the estrus cycle, potentially causing great sex differences in response to $\alpha 4\beta 1\delta$ -receptor ligands (41, 42). Furthermore, $\alpha 4\beta 1\delta$ up-regulation has been associated with increased anxiety and hyperalgesia because of disinhibition of GABAergic output neurons (43). Whether GHB effects are affected by neurosteroids, age, and sex remains to be investigated.

Overall, we have provided correlation between GHB high-affinity binding sites and $\alpha 4\beta 1\delta$ -receptors in vitro. Extensive elaborate studies are now needed to clarify the contribution of extrasynaptic $\alpha 4\beta 1\delta$ -receptors to the in vivo pharmacological effects of GHB.

Materials and Methods

Chemical Compounds. GHB and GABA were purchased from Sigma. NCS-382 and gabazine (SR 95531) hydrobromide were from Tocris Bioscience. Lithium BnOPh-GHB was prepared in-house as described previously (27). [³H]NCS-382 (20 Ci/mmol) was purchased from Biotrend. The radioligands [¹²⁵I]BnOPh-GHB and [¹²⁵I]azido-BnOPh-GHB were prepared as previously described (28, 44).

Proteomics. The photoaffinity radiolabeling with [¹²⁵I]azido-BnOPh-GHB of rat membrane was carried out exactly as previously described (8, 28) and treated with proteinase K (Sigma-Aldrich) and denaturing gel electrophoresis. The isolated gel bands were prepared for liquid chromatography-MS. Additional details in *SI Materials and Methods*.

Electrophysiology. Expression of recombinant GABA_A receptors in *X. laevis* oocytes. Human cDNAs containing the $\alpha 1$ -, $\alpha 2$ -, $\alpha 4$ -, $\alpha 5$ -, $\alpha 6$ -, $\beta 1$ -, $\beta 2$ -, $\beta 3$ -, δ -, and $\gamma 2L$ -subunits were subcloned into vectors suitable for mRNA transcription and injected into oocytes to express the desired GABA_A subtypes as previously described (45). Stages V–VI oocytes were microinjected with 0.5–5 ng mRNA using the following ratios of mRNA: $\alpha 4\beta 1\delta$ (1:1:5); $\alpha 4\beta(2/3)\delta$ (5:1:5); $\alpha 4\beta(1-3)\gamma 2L$ (1:1:1:10); and $\alpha(1, 2)\beta 1\delta$ (5:1:5). After injection, oocytes were maintained at 18 °C in the ND96 wash solution [96 mM NaCl, 2 mM KCl, 1 mM MgCl₂, 1.8 mM CaCl₂, 5 mM Hepes (hemisodium salt) augmented with 50 mg/mL gentamycin and tetracycline]. The $\alpha 4F71L$ mutant was generated by Quik-Change mutagenesis (Stratagene) according to the manufacturer's instructions. The correct identity of the mutant construct was confirmed by DNA sequencing. **Two-electrode voltage clamp recordings.** Whole-cell currents were measured 3–5 d after injection of cRNA by two-electrode voltage clamp (Digidata 1200, Geneclamp 500B amplifier) together with a Powerlab/200 (AD Instruments) and Chart version 3.5. Oocytes were voltage-clamped at -60 mV.

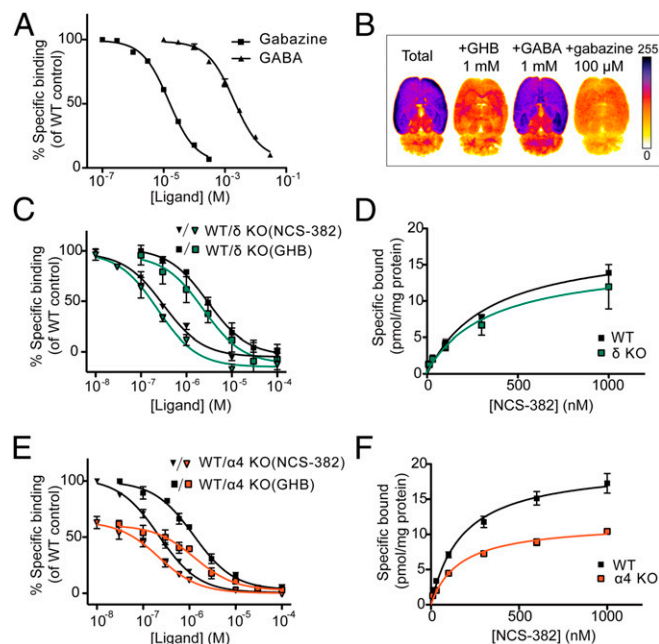


Fig. 4. GHB high-affinity radioligand binding to rat and KO mouse brain preparations. (**A**) Gabazine and GABA inhibition of [³H]NCS-382 (16 nM) binding to rat brain homogenate (pK_i 4.7 ± 0.11 and 2.7 ± 0.021 , respectively; $n = 3$). (**B**) Autoradiograms of [¹²⁵I]BnOPh-GHB (100 pM) binding to horizontal brain sections ($n = 2$). (**C** and **E**) Inhibition by GHB and NCS-382 and (**D** and **F**) saturation of [³H]NCS-382 binding (16 nM) to membrane preparations from $\alpha 4$ - and δ -subunit KO mouse brains, respectively (means \pm SEM), showing significantly lower binding in $\alpha 4$ KO vs. WT ($P = 0.0056$).

Table 2. [³H]NCS-382 binding data from $\alpha 4$ and δ KO brain tissue

	GHB IC ₅₀ (pIC ₅₀ ± SEM; M)*	NCS-382 IC ₅₀ (pIC ₅₀ ± SEM; M) [†]	B _{max} (pmol/mg protein)
$\delta^{-/-}$	2.12 · 10 ⁻⁶ (5.68 ± 0.06)	2.24 · 10 ⁻⁷ (6.70 ± 0.11)	10.1 ± 3.01
$\delta^{+/+}$	3.12 · 10 ⁻⁶ (5.51 ± 0.05)	2.65 · 10 ⁻⁷ (6.62 ± 0.10)	14.3 ± 2.93
$\alpha 4^{-/-}$	1.22 · 10 ⁻⁶ (5.97 ± 0.16)	2.16 · 10 ⁻⁷ (6.67 ± 0.03)	12.2 ± 0.89 [‡]
$\alpha 4^{+/+}$	1.19 · 10 ⁻⁶ (5.94 ± 0.09)	2.20 · 10 ⁻⁷ (6.67 ± 0.06)	20.1 ± 1.90

*n = 3.

[†]n = 5.[‡]P = 0.0056, unpaired Student t test.

The recording microelectrodes were filled with 3 M KCl and had resistance between 0.2 and 1 M Ω . Increasing concentrations of GABA were applied until a plateau in the current was reached. GABA_AR subtypes were evaluated for GHB activation by the application of GHB up to 3–10 mM. Gabazine was dissolved in 10 mM stock concentration in H₂O. I-V curves were generated by holding the oocyte at varying membrane potentials and applying either 10 mM GABA or 30 μ M GHB. The peak currents on agonist application were measured, and only oocytes that had full I-V curves measured for both GABA and GHB were included in the analysis.

Radioligand Binding Studies. Tissue and membrane preparation. Membranes were prepared as described previously (46). δ KO and WT brain tissue (cerebral cortex) was from adult male and female mice (2–4 mo; C57BL/6J \times 129Sv/SvJ) (47). Tissue from four KO mice (one female and three males) and four WT males was pooled for membrane preparations. $\alpha 4$ KO and WT brain tissue (midbrain + cerebral cortex) was from adult male mice (aged 4–6 mo) of a mixed strain 129 \times 1/S1 \times C57BL/6J genetic background (48). Two individual pools of $\alpha 4$ KO membranes prepared from three and two male KO mice were used.

[³H]NCS-382 homogenate binding assay and [¹²⁵I]BnOPh-GHB autoradiography. The [³H]NCS-382 binding assay was performed in 96-well plate format modified from the original report (49, 50) using 35–50 μ g protein (well-washed membranes), 16 nM [³H]NCS-382, and test compound in 200 μ L total incubation buffer per well (triplicates). Nonspecific binding was determined with 1 mM GHB. The reaction was terminated by rapid filtration through GF/C unifilters, and radioactivity was measured on a Packard TopCount NXT Microplate Scintillation Counter (PerkinElmer). Autoradiography was performed as previously reported (8), except that the incubation buffer was 50 mM Tris-HCl buffer (pH 7.4). In brief, horizontal brain slices (20 μ m; bregma = 4.5–4.8 mm) (51) from male Sprague-Dawley rats (~250 g; Charles River) were preincubated for 30 min in buffer and then incubated at room temperature for 30 min with 100 pM [¹²⁵I]BnOPh-GHB in the absence (total binding) or presence of competing compound. Sections were briefly washed, dried for 1 h at room temperature, and fixed in paraformaldehyde. Dried sections were exposed to a BAS-2040 phosphor imaging plate for 1.5 h at room temperature and scanned on a BAS-2500 bioimaging analyzer.

Data Analysis. Pharmacological data and statistical analyses were carried out using GraphPad Prism 5.0 (GraphPad Software Inc). Autoradiograms were analyzed using ImageJ V.1.43s (<http://rsbweb.nih.gov/ij>). The amplitude of

each current response to GABA (*I*) was normalized to the amplitude of the maximum current response to GABA (*I*_{max}). Normalized dose–response curves were analyzed by nonlinear regression. Mean parameters of each curve were derived from at least three oocytes.

I-V curves were analyzed according to Eq. 1,

$$I = I_0 \left(\frac{1 - V}{V_{rev}} \right), \quad [1]$$

where *I*₀ is the current at 0 mV and *V*_{rev} is the reversal potential. To control for rectification, *V*_{rev} was determined using data from –40 to +20 mV. Curves were normalized to each agonist rather than both to GABA to correct for changes in GHB efficacy resulting from the differing proportion of receptors without δ -subunit incorporation expressed in individual oocytes [i.e., for GHB, *I*(30 μ M GHB)/*I*(30 μ M GHB, +20 mV); for GABA, *I*(10 μ M GABA)/*I*(10 μ M GABA, +20 mV)]. The reversal potential between GHB and GABA was compared with a Student t test.

Inhibition curves and homologous displacement binding curves were analyzed by nonlinear regression. *B*_{max} values were estimated from homologous (NCS-382) competition curves exactly as described (52). For visual interpretation, saturation curves were constructed and fitted using the equation for one-site saturation. Calculations were based on the equation, where added ligand is the sum of [³H]NCS-382 and [NCS-382], and specific disintegrations per minute (DPM) was calculated from specific counts per minute (CPM) values and a known counting efficiency of 0.375 (TopCount NXT). KO and WT data are presented as mean values \pm SEM (*n*) and compared by unpaired, two-tailed Student t test.

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