

Changes in GABA_A Receptor Gene Expression Associated with Selective Alterations in Receptor Function and Pharmacology after Ethanol Withdrawal

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Changes in the expression of subunits of the GABA type A (GABA_A) receptor are implicated in the development of ethanol tolerance and dependence as well as in the central hyperexcitability associated with ethanol withdrawal. The impact of such changes on GABA_A receptor function and pharmacological sensitivity was investigated with cultured rat hippocampal neurons exposed to ethanol for 5 d and then subjected to ethanol withdrawal. Both ethanol treatment and withdrawal were associated with a marked decrease in the maximal density of GABA-evoked Cl[−] currents, whereas the potency of GABA was unaffected. Ethanol exposure also reduced the modulatory efficacy of the benzodiazepine receptor agonists lorazepam, zolpidem, and zaleplon as well as that of the inverse agonists Ro 15-4513 and FG 7142, effects that were associated with a reduced abundance of mRNAs encoding the receptor subunits $\alpha 1$, $\alpha 3$, $\gamma 2L$, and $\gamma 2S$. Ethanol withdrawal restored the efficacy of lorazepam, but not that of low concentrations of zolpidem or zaleplon, to control values. Flumazenil, which was ineffective in control neurons, and Ro 15-4513 each potentiated the GABA response after ethanol withdrawal. These effects of withdrawal were accompanied by upregulation of the $\alpha 2$, $\alpha 3$, and $\alpha 4$ subunit mRNAs as well as of the $\alpha 4$ protein. Diazepam or γ -hydroxybutyrate, but not baclofen, prevented the changes in both GABA_A receptor pharmacology and subunit mRNA levels induced by ethanol withdrawal. Changes in GABA_A receptor gene expression induced by prolonged exposure to and withdrawal of ethanol are thus associated with altered GABA_A receptor function and pharmacological sensitivity.

Key words: GABA_A receptor; ethanol; tolerance; dependence; gene expression; hippocampal neurons; patch clamp; γ -hydroxybutyrate; diazepam

Introduction

Certain acute pharmacological actions of ethanol, including its anxiolytic, sedative, ataxic, anticonvulsant, and general anesthetic effects, may be exerted through facilitation of the function of GABA type A (GABA_A) receptors in specific brain regions (Faingold et al., 1998; Grobin et al., 1998; Ueno et al., 2001; Aguayo et al., 2002; Chester and Cunningham, 2002). GABA_A receptors are ligand-gated Cl[−] channels and mediate fast inhibitory transmission in the mammalian CNS (Mehta and Ticku, 1999; Vicini, 1999). They are heteromeric complexes of five subunits that belong to various classes: $\alpha 1$ –6, $\beta 1$ –4, $\gamma 1$ –3, δ , ϵ , π , θ , and $\rho 1$ –3 (Barnard et al., 1998; Whiting et al., 1999; Sieghart and Sperk, 2002). These subunits are expressed in a region- and ontogeny-dependent manner in the brain and generate a large number of GABA_A receptor subtypes that differ not only in sub-

unit composition but in their physiological and pharmacological properties (Sieghart, 1995; McKernan and Whiting, 1996; Barnard et al., 1998; Hevers and Luddens, 1998).

Prolonged exposure to ethanol results in the development of dependence and of tolerance to its behavioral actions (Suwaki et al., 2001). Altered GABA_A receptor function, characterized by a reduced responsiveness to GABA, tolerance to ethanol, cross-tolerance to benzodiazepines and barbiturates, as well as sensitization to neurosteroids and inverse agonists, is thought to underlie these chronic effects of ethanol (Ticku and Burch, 1980; Allan and Harris, 1987; Morrow et al., 1988; Sanna et al., 1993; Devaud et al., 1996). The molecular mechanisms responsible for adaptation of GABA_A receptors to long-term ethanol exposure remain unclear but may involve changes in cell surface density (Ticku and Burch, 1980), in post-translational protein modification (Kumar et al., 2002), or in subunit expression (Mhatre et al., 1993; Devaud et al., 1995, 1997; Follesa et al., 2003).

Long-term ethanol treatment results in a decrease in the abundance of the $\alpha 1$ subunit mRNA (Montpied et al., 1991; Devaud et al., 1995; Follesa et al., 2003) and an increase in that of $\alpha 4$, $\alpha 6$, $\gamma 1$, and $\gamma 2S$ mRNAs (Devaud et al., 1995) in the cerebral cortex and cerebellum. Changes in GABA_A receptor subunit expression after chronic ethanol exposure were also demonstrated

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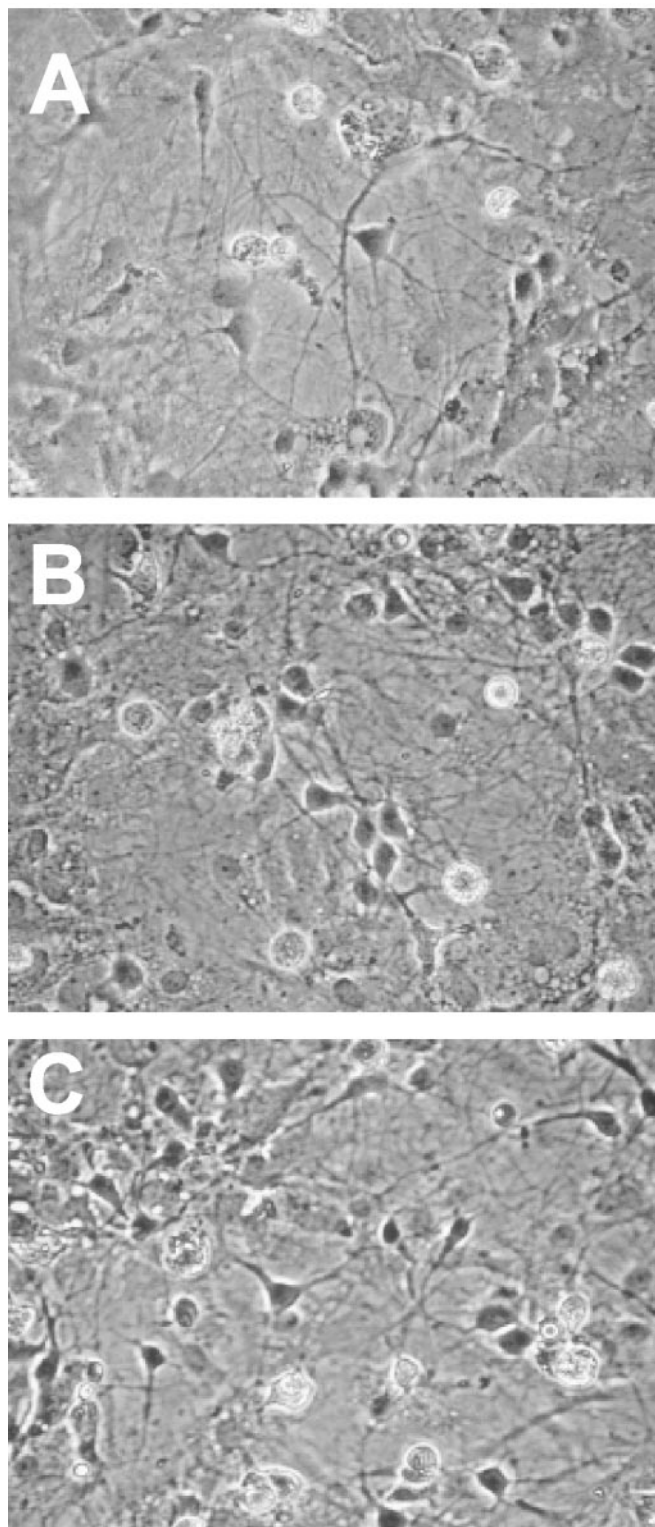


Figure 1. Light micrographs of live rat hippocampal neurons in culture. Cells were cultured for 5 d in the absence (A) or presence (B) of 100 mM ethanol; some ethanol-treated cells were then incubated for an additional 6 hr in the absence of ethanol (C). Magnification, 200 \times .

in the hippocampus, a region considered to be less sensitive with regard to the acute ethanol effect on GABA_A receptor function. Chronic ethanol exposure decreased α 1 subunit expression, increased that of the α 4 subunit (Mahmoudi et al., 1997; Matthews et al., 1998; Cagetti et al., 2003), and did not alter that of the γ 2

Table 1. Effects of chronic ethanol treatment and ethanol withdrawal on the total RNA content of hippocampal neurons

Experiment	Total RNA (micrograms per dish)		
	Control	Chronic EtOH	EtOH withdrawal
1	14.41 \pm 2.03	12.88 \pm 2.32	15.40 \pm 3.51
2	13.49 \pm 3.02	15.93 \pm 5.95	12.97 \pm 5.65
3	16.85 \pm 1.41	15.05 \pm 2.66	16.25 \pm 3.53
4	14.37 \pm 2.15	16.25 \pm 2.12	17.23 \pm 2.79
5	18.69 \pm 4.28	16.56 \pm 1.15	15.36 \pm 3.57

Cells were cultured in 35 mm dishes for 5 d in the absence (control) or presence (chronic EtOH) of 100 mM ethanol; some ethanol-treated cells were also incubated for an additional 6 hr in the absence of ethanol (EtOH withdrawal). Total RNA was isolated from the cells and quantitated. Data are expressed as micrograms of RNA per culture dish and are means \pm SE of three dishes from five randomly selected experiments.

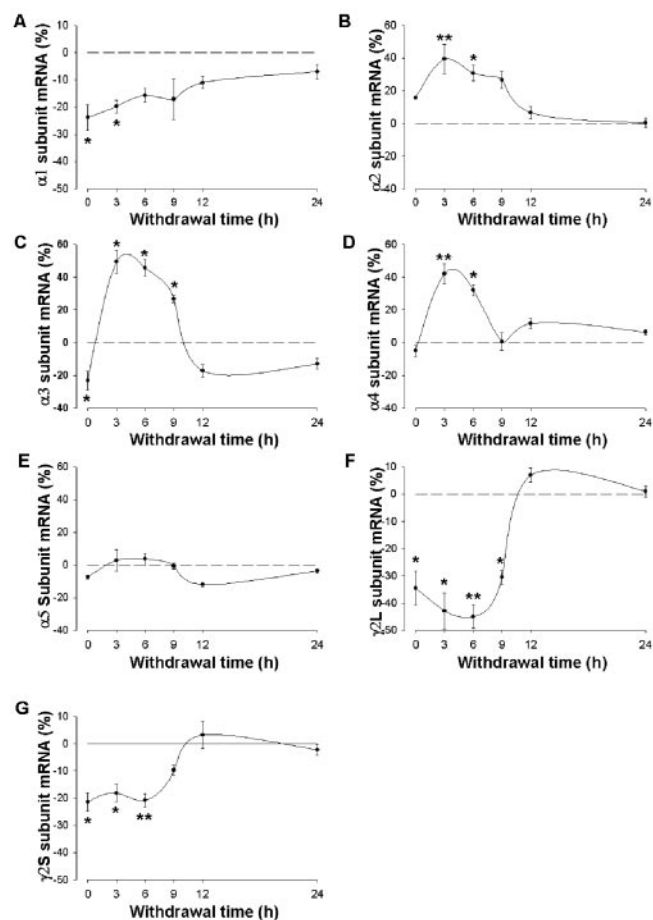


Figure 2. Time course of the effects of ethanol withdrawal on the abundance of GABA_A receptor subunit mRNAs in hippocampal cells. Cells were incubated first for 5 d with 100 mM ethanol and then for the indicated times in ethanol-free medium. The amounts of GABA_A receptor α 1 (A), α 2 (B), α 3 (C), α 4 (D), α 5 (E), γ 2L (F), and γ 2S (G) subunit mRNAs were then determined by RNase protection assay. Data are means \pm SE of 6–13 values from three independent experiments and are expressed as a percentage of the corresponding value for control cultures incubated in the absence of ethanol for 5 d. * p < 0.05, ** p < 0.001 versus control.

subunit (Matthews et al., 1998). This pattern of changes was different from that of the cerebral cortex (Matthews et al., 1998). Chronic intermittent ethanol treatment also downregulated δ subunit expression and upregulated that of the γ 1 and γ 2 subunits in the hippocampus (Cagetti et al., 2003).

These changes are associated in most cases with alterations in the amounts of subunit proteins (Devaud et al., 1997), but it has remained unclear whether they are directly correlated with the

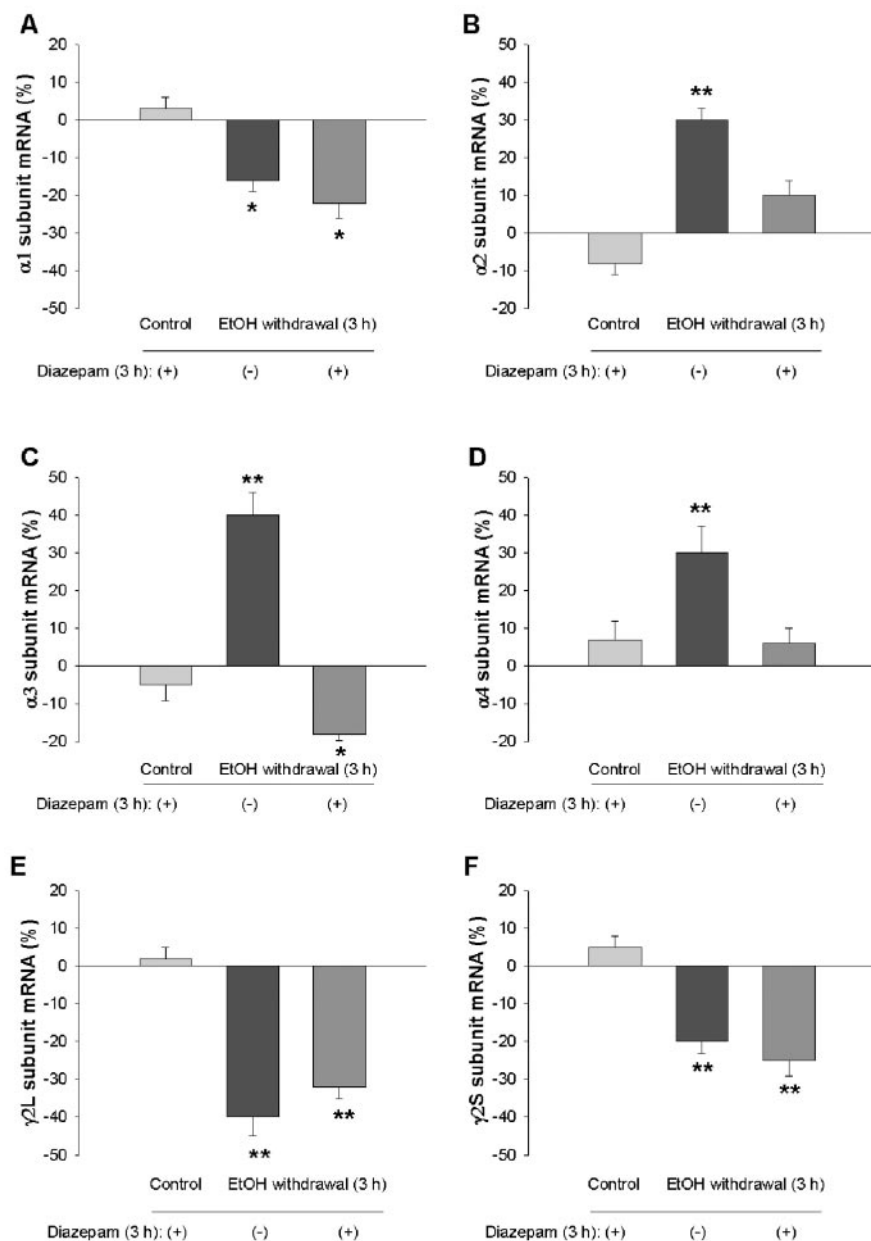


Figure 3. Prevention by diazepam of changes in the abundance of GABA_A receptor subunit mRNAs induced by ethanol withdrawal. Hippocampal neurons were incubated first for 5 d with 100 mM ethanol and then for 3 hr in ethanol-free medium in the absence or presence of 10 μ M diazepam. Cells incubated for 5 d in the absence of ethanol were also exposed to diazepam for 3 hr. The abundance of GABA_A receptor α 1 (A), α 2 (B), α 3 (C), α 4 (D), γ 2L (E), and γ 2S (F) subunit mRNAs was then determined by RNase protection assay. Data are means \pm SE of 9–12 values from three independent experiments and are expressed as a percentage of the corresponding value for control cultures incubated for 5 d in the absence of ethanol. * p < 0.05, ** p < 0.001 versus control. EtOH, Ethanol.

changes in GABA_A receptor function or pharmacological sensitivity that result from chronic exposure to and subsequent withdrawal of ethanol.

With the use of rat hippocampal neurons in primary culture, we have further evaluated the effects of prolonged exposure to and abrupt withdrawal of ethanol on GABA_A receptor function, expression, and responsiveness to ligands selective for different receptor subtypes. The use of cultured hippocampal neurons allowed us, at variance with laboratory animals, to overcome the difficulty of establishing the precise onset of ethanol withdrawal caused by the possibility of the rapid washout of ethanol.

Materials and Methods

Primary culture of hippocampal neurons. Animal care and handling throughout all experimental procedures were in accordance with the European Communities Council Directive of 24 November 1986 (86/609/EEC). The experimental protocols were also approved by the Animal Ethics Committee of the University of Cagliari. Primary cultures of hippocampal neurons were prepared from Sprague Dawley rats on postnatal days 1–3 as described previously (Costa et al., 2000), with minor modifications. Pups were killed by decapitation, and the hippocampus was removed and transferred to a culture dish containing Neurobasal A medium (Invitrogen, San Diego, CA) supplemented with 10% heat-inactivated fetal bovine serum (Sigma, St. Louis, MO), 25 μ M glutamate, 0.5 mM glutamine, penicillin (100 U/ml), streptomycin (0.1 mg/ml), and amphotericin B (0.25 μ g/ml). The tissue was chopped with scissors, and the resulting fragments were transferred to a sterile tube and gently dissociated by repeated passage through a Pasteur pipette with an opening of 0.5 mm. The dissociated cells were plated either in 35 mm culture dishes (4×10^6 cells) that had been coated with poly-L-lysine hydrobromide (100 μ g/ml; 30–70 kDa) (Sigma) for measurement of GABA_A receptor subunit mRNAs or in multi-well dishes containing 12-mm-round glass coverslips coated with poly-L-lysine (6×10^5 cells) for electrophysiological recording or immunocytofluorescence analysis. Cells were cultured in a humidified incubator containing 5% CO₂ at 37°C. Twenty-four hours after plating, fetal bovine serum was replaced by B-27 supplement (Invitrogen), and glutamate was removed from the medium after 3 d in culture.

Ethanol treatment. After 5 d in culture, cells were exposed continuously for the next 5 d to 100 mM ethanol according to the procedure of Smothers et al. (1997), with minor modifications. Ethanol was added directly to the culture medium, and to prevent daily fluctuations in its concentration caused by evaporation, culture dishes and multi-well dishes with coverslips were kept inside a sealed sterile plastic container (pre-equilibrated with 5% CO₂/95% air) along with an isomolar concentration of ethanol in an open beaker. Control cultures were kept likewise in a sealed sterile plastic container not containing ethanol. To further ensure the consistency of the ethanol concentration, the culture medium was replaced daily for both ethanol-treated and control cells. Because daily replacement of medium could have potential effects on the network properties of cells, we ran preliminary experiments with untreated cells in which a culture condition with daily changes of medium, starting from day 5 to days 10–12, was compared with a condition of no medium replacement for the same period of days. The subsequent measurement of several parameters such as total amount of mRNA extracted, functional and pharmacological responses of GABA_A receptors, morphology of neurons, and abundance of glial cells, revealed no apparent alteration produced by this procedure (results not shown). Because the daily culture medium replacement actually began on day 5 of culture, it is likely that at this stage hippocampal neurons may have already reached a sufficient degree of maturation and differentiation, becoming much less sensitive to this procedure.

The concentration of 100 mM ethanol used in our study to treat cultured hippocampal cells was chosen on the basis of its efficacy in both acutely potentiating GABA_A receptor function and producing changes in GABA_A receptor subunit gene expression, as determined in pilot experiments (results not shown). Such a concentration of ethanol has been used in previous studies in our laboratory (Follesa et al., 2003) as well as in other laboratories (Smother's et al., 1997; van Zundert et al., 2000).

To assess the effects of chronic ethanol treatment, cultured cells exposed to ethanol, as described above, were used immediately after the removal of ethanol from the culture medium (0 hr of withdrawal). In withdrawal experiments, the ethanol-containing medium was replaced after 5 d by ethanol-free medium containing (or not) diazepam (10 μM), baclofen (100 μM), or γ-hydroxybutyrate (GHB) (1, 10, or 100 mM) with or without SCH 50911 (100 μM), after which the cells were incubated for an additional 3–24 hr. Ethanol, GHB, baclofen, and SCH 50911 were dissolved in medium, whereas diazepam was dissolved in dimethyl sulfoxide and subsequently diluted to the desired concentration in culture medium. Control neurons were treated with the corresponding vehicle. All experimental groups were compared with control cells maintained in culture for the same number of days or hours during ethanol withdrawal. Thus, each experimental group had its respective control processed at the same time.

Riboprobe preparation. GABA_A receptor subunit cDNAs were prepared as described previously (Follesa et al., 1998) by reverse transcription and PCR. In brief, cDNA prepared from rat brain (1–10 ng) was subjected to amplification with TaqDNA polymerase (2.5 U) (Perkin-Elmer/Cetus, Norwalk, CT) in 100 μl of standard buffer (100 mM Tris-HCl, pH 8.3, 500 mM KCl, 15 mM MgCl₂, 0.01% gelatin) containing 1 μM each of specific sense and antisense primers and 200 μM of each deoxynucleoside triphosphate. The primer pairs for the various receptor subunits were designed to include cDNA sequences with the lowest degree of homology among the different subunits (Follesa et al., 1998). The reaction was performed in a thermal cycler (Eppendorf) for 30 cycles of 94°C for 45 sec and 60°C for 1 min, with a final extension at 72°C for 15 min. The reaction products were separated by electrophoresis, visualized by staining with ethidium bromide, excised from the gel, purified, and cloned into the pAMP 1 vector (Invitrogen). The resulting plasmids were introduced into *Escherichia coli* DH5α and subsequently purified from the bacterial cells, and the cDNA inserts were sequenced with a Sequenase DNA sequencing kit (USB, Cleveland, OH). The determined nucleotide sequences were 100% identical to the respective previously published sequences. Plasmids containing the cDNA fragments corresponding to the various GABA_A receptor subunits were linearized with restriction enzymes (Follesa et al., 1998) and used as templates for the appropriate RNA polymerase (SP6 or T7) to generate [α -³²P]UTP-labeled cRNA probes for RNase protection assays.

RNA extraction and measurement of GABA_A receptor subunit mRNAs. Total RNA was isolated from cultured hippocampal cells with an RTN kit (Sigma) and quantified by measurement of absorbance at 260 nm. An RNase protection assay for the semiquantitative measurement of the GABA_A receptors α1, α2, α3, α4, α5, γ2L, and γ2S subunit mRNAs was

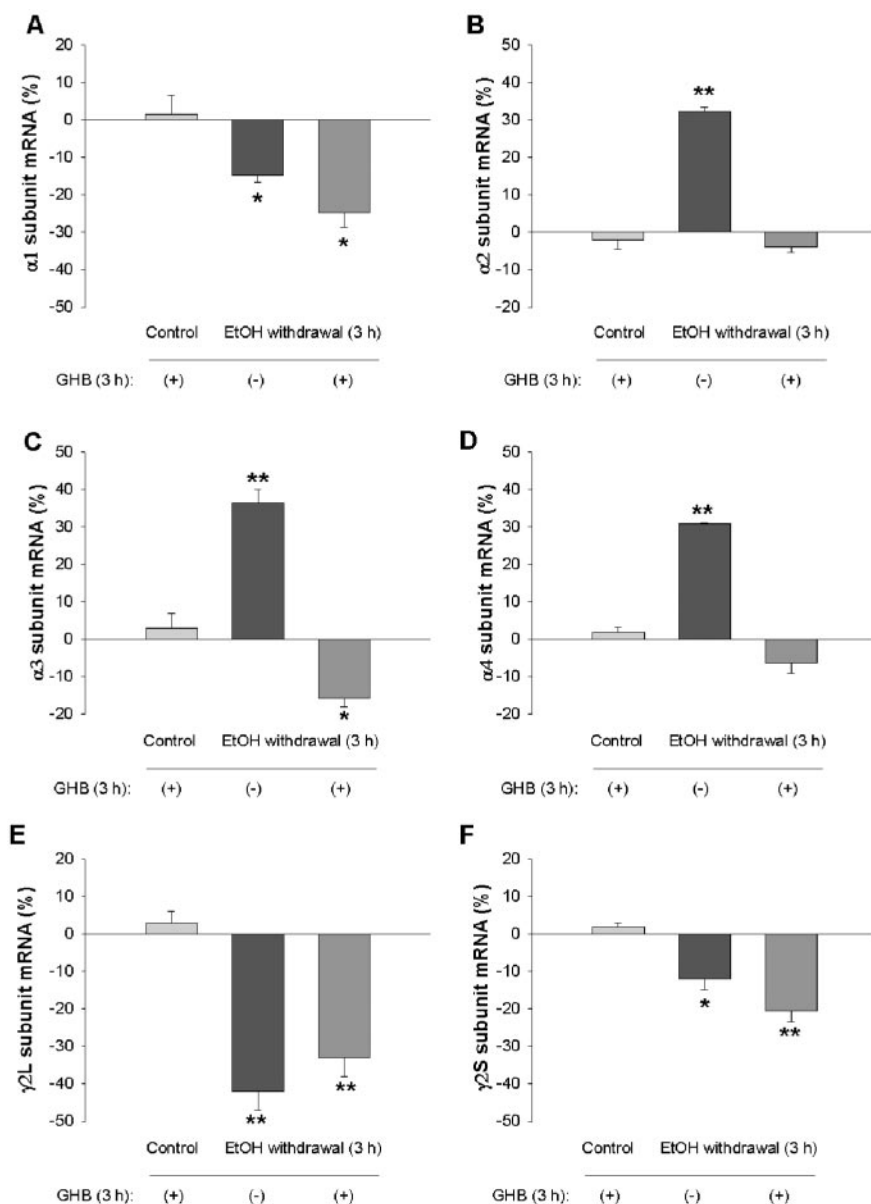


Figure 4. Prevention by GHB of changes in the abundance of GABA_A receptor subunit mRNAs induced by ethanol withdrawal. Hippocampal neurons were incubated first for 5 d with 100 mM ethanol and then for 3 hr in ethanol-free medium in the absence or presence of 100 mM GHB. Cells incubated for 5 d in the absence of ethanol were also exposed to GHB for 3 hr. The abundance of GABA_A receptor α1 (A), α2 (B), α3 (C), α4 (D), γ2L (E), and γ2S (F) subunit mRNAs was then determined by RNase protection assay. Data are means ± SE of 9–12 values from three independent experiments and are expressed as a percentage of the corresponding value for control cultures incubated for 5 d in the absence of ethanol. **p* < 0.05, ***p* < 0.001 versus control.

performed as described (Follesa et al., 1998). In brief, 15 μg of total RNA was dissolved in 20 μl of hybridization solution containing 150,000 cpm of ³²P-labeled cRNA probe for a specific GABA_A receptor subunit (6 × 10⁷ to 7 × 10⁷ cpm/μg) and 15,000 cpm of ³²P-labeled cyclophilin cRNA (1 × 10⁶ cpm/μg). Cyclophilin is expressed widely among tissues, including the brain, and its gene is most likely regulated in an “on or off” manner; cyclophilin mRNA was thus used as an internal standard for our measurements (Follesa et al., 1998). The hybridization reaction mixtures were incubated at 50°C overnight and then subjected to digestion with RNase, after which RNA–RNA hybrids were detected by electrophoresis (on a sequencing gel containing 5% polyacrylamide and urea) and autoradiography. The amounts of GABA_A receptor subunit mRNAs and cyclophilin mRNA were determined by measurement of the optical density of the corresponding bands on the autoradiogram with a densitometer (model GS-700; Bio-Rad, Hercules, CA), which was calibrated to detect

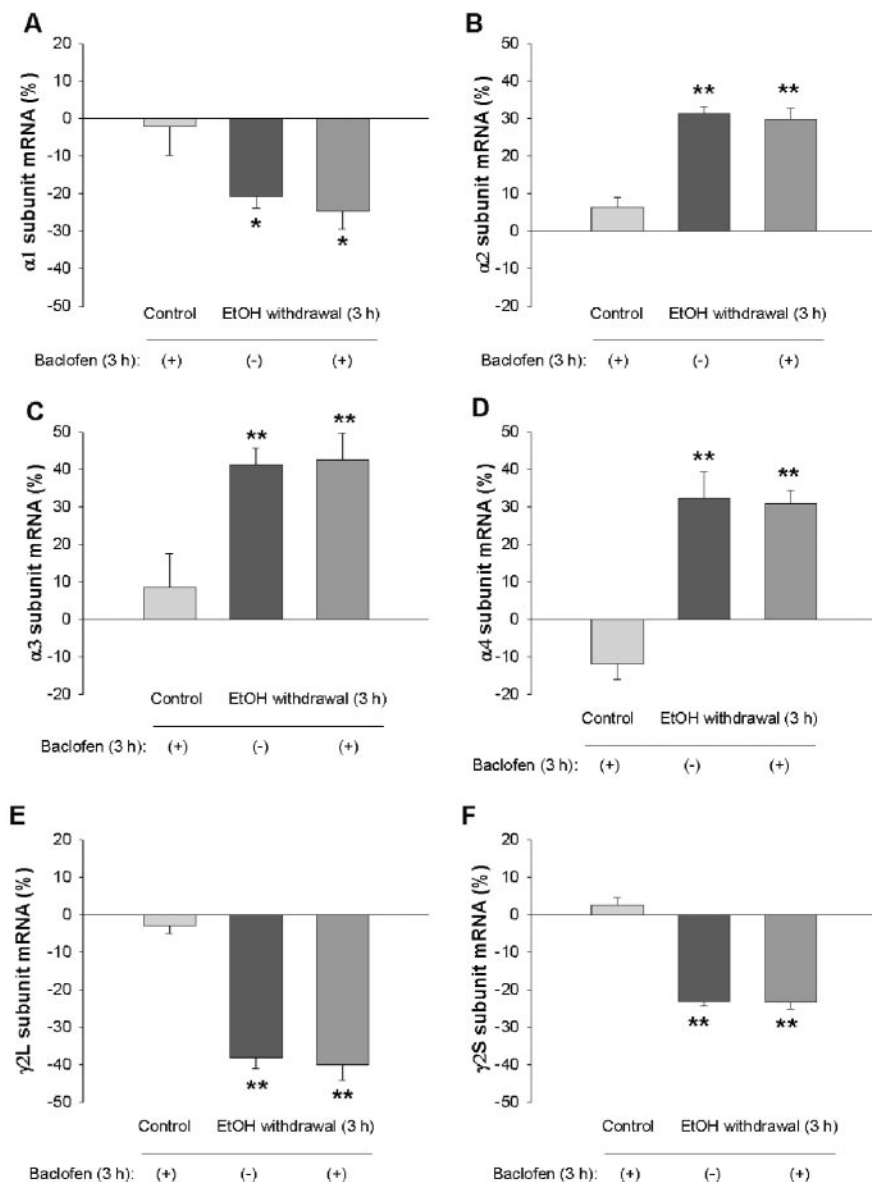


Figure 5. Lack of effect of baclofen on changes in the abundance of GABA_A receptor subunit mRNAs induced by ethanol withdrawal. Hippocampal neurons were incubated first for 5 d with 100 mM ethanol and then for 3 hr in ethanol-free medium in the absence or presence of 100 μ M baclofen. Cells incubated for 5 d in the absence of ethanol were also exposed to baclofen for 3 hr. The abundance of GABA_A receptor α 1 (A), α 2 (B), α 3 (C), α 4 (D), γ 2L (E), and γ 2S (F) subunit mRNAs was then determined by RNase protection assay. Data are means \pm SE of 9–12 values from three independent experiments, each performed in triplicate or quadruplicate, and are expressed as a percentage of the corresponding value for control cultures incubated for 5 d in the absence of ethanol. * p < 0.05, ** p < 0.001 versus control.

saturated values so that all measurements were in the linear range. The data were normalized by dividing the optical density of the protected fragment for each receptor subunit mRNA by that of the respective protected fragment for cyclophilin mRNA. The amount of mRNA was therefore expressed in arbitrary units.

Immunoblot analysis. Hippocampal neurons were homogenized in a solution containing 10 mM Tris-HCl, pH 7.4, 0.32 M sucrose, 5 mM EDTA, and 0.1 mM phenylmethylsulfonyl fluoride, and the homogenate was centrifuged at $1000 \times g$ for 10 min. The resulting supernatant was then centrifuged for 20 min at $12,000 \times g$, and the crude membrane pellet so obtained was washed three times with homogenization buffer and stored at -20°C until use. Portions of the crude membrane fraction (40 μ g of protein) were incubated for 5 min at 95°C in 20 μ l of SDS sample buffer and then subjected to SDS-PAGE on 10% minigels (Mini Protean II; Bio-Rad). The separated proteins were transferred electrophoretically

to a nitrocellulose membrane and subjected to immunoblot analysis with goat polyclonal antibodies (1 μ g/ml) generated in response to an extracellular epitope (peptide N1–19) of the rat GABA_A receptor α 4 subunit (Santa Cruz Biotechnology, Santa Cruz, CA). The membrane was incubated with the antibodies in the absence or presence of the N1–19 peptide (10 μ g/ μ g of antibody) (Santa Cruz Biotechnology). Immune complexes were detected with an ECL detection kit (Amersham Biosciences, Little Chalfont, UK).

Immunocytofluorescence analysis. Cells cultured on coverslips were washed three times with PBS, fixed for 1 hr at room temperature with 4% paraformaldehyde in PBS, washed three times with TN buffer (50 mM Tris-HCl, pH 7.5, 150 mM NaCl), and permeabilized for 1 hr at room temperature with TN-T buffer (0.1% Triton X-100 in TN buffer) containing 0.5% dried skim milk. Nonspecific binding sites for avidin and biotin were blocked by incubation of the cells for 15 min at room temperature, first with avidin D blocking solution and then with biotin blocking solution (Vector, Burlingame, CA). The cells were then incubated overnight at 4°C with the goat polyclonal antibodies (1:500 dilution in TN-T) to the GABA_A receptor α 4 subunit. After several washes with TN-T buffer, the cells were incubated for 1 hr at room temperature, first with biotin-conjugated donkey antibodies (1:200 in TN-T) to goat IgG (Jackson ImmunoResearch Laboratories, West Grove, PA) and then with tetramethyl rhodamine isothiocyanate-conjugated streptavidin (Jackson ImmunoResearch Laboratories) diluted to a concentration of 2 μ g/ml in TN-T. The cells were washed extensively with TN buffer, and a coverslip was then applied with permanent aqueous mounting medium (Sigma). Quantitative analysis was performed using a Leica four-dimensional confocal laser scanning microscope with an argon–krypton laser as described previously (Spiga et al., 2003). Confocal images were generated using PL Floustar 100 \times oil (numerical aperture = 1.3). Each frame was acquired eight times and then averaged to obtain noise-free images. Three-dimensional reconstructions of cells were obtained with the “maximum intensity” algorithm that was used on optical sections, usually at consecutive intervals of 0.5–1 μ m, and were imaged through the depth of the labeled neurons and saved as image data set and processed with Scanware 4.2a Leica. All confocal images were white-labeled on a black background, in a gray scale ranging from 0 (black) to 255 (white). For morphometric and statistical analysis on three-dimensional reconstructed images, Bioscan Optimas version 6.5 software was used. The area of the body of the cell is obtained by marking its profile, excluding all dendritic trunks with a spline of 64 intervals. This yields the bounded area in calibrated square units (square micrometers). Perimeter values are evaluated similarly, and the total boundary length is expressed in micrometers. Major axis length is a real value, which is obtained from area objects giving the length of the major axis in calibrated units. Breadth is a real value, which is extracted from area objects giving the sum of the maximum distance of the boundary from either side of the major axis in calibrated units. Circularity is a real value of the ratio of the squared perimeter divided by the area (i.e., perimeter squared/area). This is a dimensionless number with a minimum value of 4π (12.57) achieved only for circular boundaries.

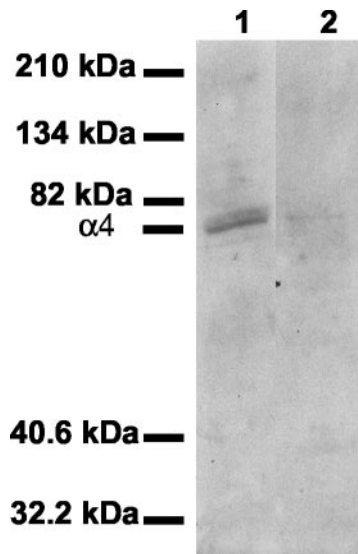


Figure 6. Immunoblot analysis of the GABA_A receptor α_4 subunit in hippocampal neurons. A crude membrane fraction (40 μ g of protein) of cultured hippocampal neurons was subjected to immunoblot analysis with antibodies specific for the GABA_A receptor α_4 subunit in either the absence (lane 1) or presence (lane 2) of the α_4 peptide N1–19. The positions of molecular size standards and of the α_4 protein are indicated.

Whole-cell electrophysiological recording. Immediately before recording, coverslips were transferred to a perfusion chamber (Warner Instruments, Hampden, CT), and neurons were visualized with a Nikon upright microscope equipped with Nomarski optics. Large neurons with a pyramidal shape and well defined dendritic processes were selected for electrophysiological recording (see Fig. 1). The membrane potential was clamped at -60 mV with an Axopatch 200-B amplifier (Axon Instruments, Foster City, CA). The resting membrane potential for the studied neurons was approximately -60 mV. Recording pipettes (borosilicate capillaries with a filament; outer diameter, 1.5 mm) (Sutter Instruments, Novato, CA) were prepared with a two-step vertical puller (Sutter Instruments) and had resistances between 4 and 6 M Ω . Pipette capacitance and series resistance were compensated, the latter at 60%. Currents through the patch-clamp amplifier were filtered at 2 kHz and digitized at 5.5 kHz with commercial software (pClamp 8.1; Axon Instruments).

The external solution contained (in mM): 130 NaCl, 5 KCl, 2 CaCl₂, 1 MgCl₂, 10 HEPES-NaOH, pH 7.3, and 11 glucose. The internal solution contained (in mM): 140 CsCl, 2 MgCl₂, 1 CaCl₂, 10 EGTA, 10 HEPES-CsOH, pH 7.3, and 2 ATP (disodium salt). Drugs were applied with a fast-exchange flow-tube perfusion system driven by a motor (Warner Instruments). Agonists were applied at intervals of 30 sec. All experiments were performed at room temperature (23–25°C). Data were analyzed by pClampfit 8.01 software (Axon Instruments). Modulation of GABA-evoked Cl⁻ currents by drugs is expressed as percentage change, $[(I'/I) - 1] \times 100\%$, where I is the average of control responses obtained before drug application and after drug washout, and I' is the average of the agonist-induced responses obtained from the same cell in the presence of drug. Nonlinear regression analysis of GABA dose–response relations determined from the average peak current amplitude was performed with Prism software (version 4, Graphpad) according to the equation $I = I_{\min} + (I_{\max} - I_{\min}) / (1 + 10^{[\log(\text{EC}_{50} - X)]/n_H})$, where I_{\min} and I_{\max} are the minimal and maximal responses to GABA, respectively, EC_{50} is the concentration of GABA that produces 50% of the maximal response, X is the test concentration of GABA, and n_H is the Hill coefficient.

Statistical analysis. Data are presented as means \pm SE. The statistical significance of differences was assessed by one-way ANOVA followed by Scheffé's test. A p value of <0.05 was considered statistically significant.

Results

Neither continuous exposure to 100 mM ethanol for 5 d nor subsequent withdrawal of ethanol for 6 hr appeared to affect the

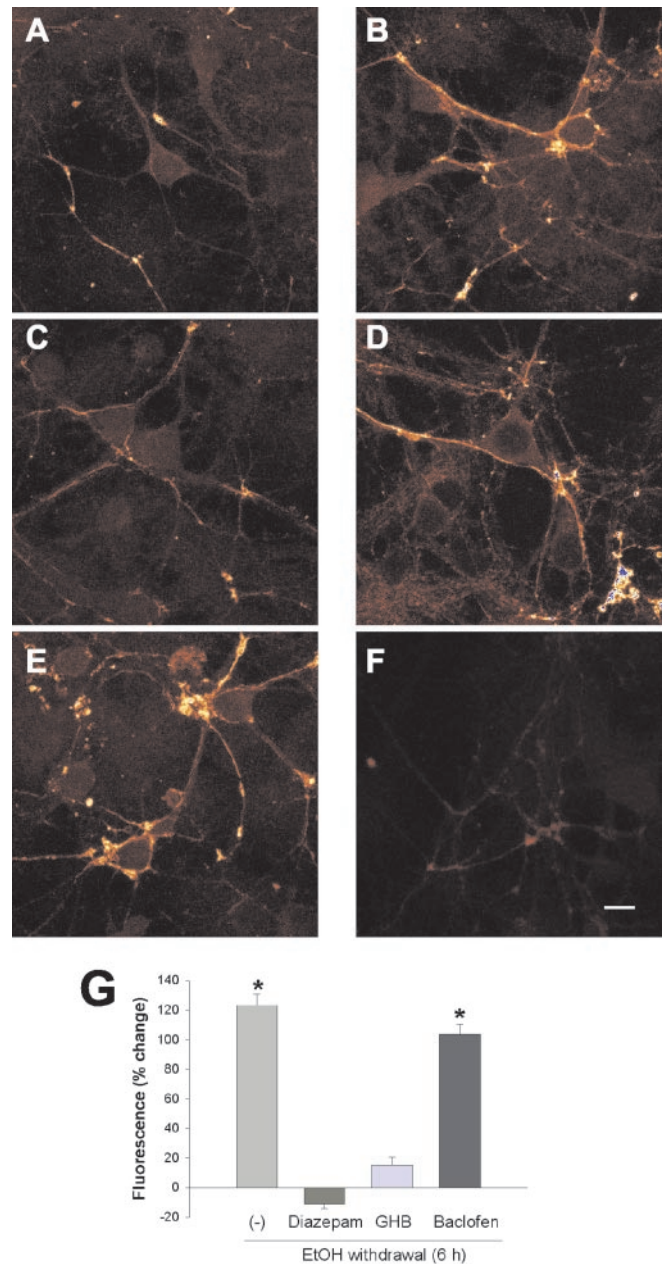


Figure 7. Ethanol withdrawal–induced increase in the abundance of the GABA_A receptor α_4 subunit in hippocampal neurons and its inhibition by diazepam or GHB. Cells were cultured for 5 d in the absence (A) or presence of 100 mM ethanol; the ethanol-treated cells were subsequently incubated for an additional 6 hr in ethanol-free medium in the absence (B) or presence of 10 μ M diazepam (C), 100 mM GHB (D), or 100 μ M baclofen (E). All cells were then subjected to immunocytofluorescence analysis with antibodies to an extracellular epitope (peptide N1–19) of the α_4 subunit. F, Control cells were also subjected to analysis with the specific antibodies in the presence of the N1–19 peptide (10 μ g/ml). Representative images are shown. Scale bar, 10 μ m. G, Semicquantitative determination of the abundance of the α_4 subunit as determined by image analysis of the immunocytofluorescence data. Results are expressed as percentage change in fluorescence intensity relative to control cells and are means \pm SE of values for at least 40 randomly selected cells for each experimental group and in three independent experiments. * $p < 0.001$ versus control neurons.

gross morphology of cultured rat hippocampal neurons (Fig. 1). This conclusion was supported by analysis of morphometric parameters by confocal laser-scanning microscopy (see Table 2). In addition, ethanol treatment and withdrawal did not affect the amount of total RNA in these cells (Table 1).

Table 2. Morphometric parameters of rat hippocampal neurons subjected to ethanol withdrawal in the presence of diazepam, GHB, or baclofen

Experimental group	Area (μm^2)	Perimeter (μm)	Circularity	Maximum axis length (μm)	Breadth (μm)
Control ($n = 33$)	173 \pm 9	51.2 \pm 1.3	15.6 \pm 0.3	19.0 \pm 0.5	13.7 \pm 0.5
Withdrawal ($n = 41$)	171 \pm 12	51.8 \pm 2.0	16.3 \pm 0.3	19.0 \pm 0.8	13.6 \pm 0.5
Withdrawal + diazepam ($n = 23$)	158 \pm 9	47.7 \pm 1.3	14.6 \pm 0.2	17.4 \pm 0.5	13.1 \pm 0.4
Withdrawal + GHB ($n = 25$)	169 \pm 10	51.1 \pm 1.5	16.4 \pm 0.3	18.5 \pm 0.7	13.2 \pm 0.4
Withdrawal + baclofen ($n = 22$)	171 \pm 11	50.5 \pm 1.7	15.2 \pm 0.2	18.0 \pm 0.8	13.5 \pm 0.5

Cells were cultured for 5 d in the absence (control) or presence of 100 mM ethanol, after which the ethanol-treated cells were incubated for an additional 6 hr in ethanol-free medium in the absence or presence of diazepam (10 μM), GHB (100 mM), or baclofen (100 μM). The cells were then examined by confocal laser-scanning microscopy for determination of the indicated morphometric parameters. Data are means \pm SE of values determined from the indicated number (n) of cells.

Effects of chronic ethanol treatment on GABA_A receptor gene expression

The abundance of GABA_A receptors $\alpha 1$, $\alpha 2$, $\alpha 3$, $\alpha 4$, and $\alpha 5$ mRNAs as well as that of the mRNAs for the two splice variants of the $\gamma 2$ subunits, $\gamma 2L$ and $\gamma 2S$, were determined by RNase protection assay after continuous exposure of cultured hippocampal neurons to 100 mM ethanol for 5 d. Ethanol induced a significant decrease in the amount of $\alpha 1$, $\alpha 3$, $\gamma 2L$, and $\gamma 2S$ subunit mRNAs relative to control values (Fig. 2*A, C, F, G*). ANOVA revealed a significant main effect of chronic ethanol treatment on mRNA levels for $\alpha 1$ ($F_{(1,16)} = 21.83$; $p < 0.001$), $\alpha 3$ ($F_{(1,20)} = 9.48$; $p < 0.006$), $\gamma 2L$ ($F_{(1,11)} = 17.31$; $p < 0.002$), and $\gamma 2S$ ($F_{(1,21)} = 34.46$; $p < 0.001$).

In contrast, chronic ethanol exposure did not significantly affect the abundance of $\alpha 2$ ($F_{(1,24)} = 3.21$; $p < 0.067$), $\alpha 4$ ($F_{(1,16)} = 0.31$; $p < 0.588$), and $\alpha 5$ ($F_{(1,24)} = 4.26$; $p < 0.075$) mRNAs (Fig. 2*B, D, E*).

Effects of ethanol withdrawal on GABA_A receptor gene expression

To determine the effect of ethanol withdrawal on GABA_A receptor gene expression, we incubated hippocampal neurons first with 100 mM ethanol for 5 d and then in the absence of ethanol for 3–24 hr. The abundance of the $\alpha 1$, $\gamma 2L$, and $\gamma 2S$ subunit mRNAs, which was reduced after ethanol treatment, remained significantly decreased, relative to control values, 3 hr after removal of ethanol (Fig. 2*A, F, G*) ($\alpha 1$, $F_{(5,36)} = 5.29$, $p < 0.001$; $\gamma 2L$, $F_{(5,18)} = 18.11$, $p < 0.001$; $\gamma 2S$, $F_{(5,42)} = 12.13$, $p < 0.001$). The amount of the $\alpha 1$ subunit mRNA had returned to control values 6 hr after ethanol removal (Fig. 2*A*). The amounts of the $\gamma 2L$ and $\gamma 2S$ subunit mRNAs remained significantly decreased 6 hr after ethanol withdrawal but had returned to control values by 9–12 hr (Fig. 2*F, G*). In contrast, the abundance of $\alpha 2$, $\alpha 3$, and $\alpha 4$ mRNAs was markedly increased, relative to control values, in response to ethanol withdrawal, with the maximal effects being apparent 3 hr after ethanol removal; the amounts of these mRNAs had returned to control values by 9–12 hr after ethanol withdrawal (Fig. 2*B–D*). ANOVA revealed a significant main effect of ethanol withdrawal on mRNA levels for $\alpha 2$ ($F_{(5,52)} = 7.97$; $p < 0.001$), $\alpha 3$ ($F_{(5,34)} = 17.18$; $p < 0.001$), and $\alpha 4$ ($F_{(5,37)} = 23.46$; $p < 0.001$). Discontinuation of ethanol treatment had no effect ($F_{(5,47)} = 2.04$; $p < 0.088$) on the abundance of the $\alpha 5$ subunit mRNA (Fig. 2*E*).

Effects of diazepam, GHB, and baclofen on ethanol withdrawal-induced changes in GABA_A receptor gene expression

Benzodiazepines, GHB, and the GABA_B receptor agonist baclofen reduce withdrawal symptoms and the craving for ethanol both in human alcoholics and in ethanol-dependent laboratory animals (Fadda et al., 1989; Gallimberti et al., 1989; Addolorato et

al., 1996, 2002; Agabio et al., 1998; Lejoyeux et al., 1998; Colombo et al., 2000). We therefore examined the effects of diazepam, GHB, and baclofen on the changes in GABA_A receptor gene expression observed during ethanol withdrawal. Hippocampal neurons were incubated first for 5 d with 100 mM ethanol and then for 3 hr in ethanol-free medium containing 10 μM diazepam, 1–100 mM GHB, or 100 μM baclofen. Diazepam prevented the changes in the abundance of the $\alpha 2$, $\alpha 3$, and $\alpha 4$ subunit mRNAs induced by ethanol withdrawal (Fig. 3). Similarly, GHB at 100 mM (Fig. 4), but not at 1 or 10 mM (data not shown), prevented the ethanol withdrawal-induced changes in subunit mRNA levels. In contrast, the amounts of $\alpha 1$, $\alpha 2$, $\alpha 3$, $\alpha 4$, $\gamma 2L$, and $\gamma 2S$ mRNAs apparent 3 hr after ethanol withdrawal were not affected by the presence of baclofen (Fig. 5). Diazepam, GHB, or baclofen had no significant effect on the amounts of the various subunit mRNAs in neurons not exposed to ethanol (Figs. 3–5).

Effect of ethanol withdrawal on $\alpha 4$ subunit abundance

Recent studies strongly suggested a pivotal role for the increased expression of the $\alpha 4$ subunit associated with withdrawal from ethanol (Devaud et al., 1995, 1997; Mahmoudi et al., 1997; Matthews et al., 1998; Cagetti et al., 2003; Follsea et al., 2003) as well as other positive allosteric modulators of the GABA_A receptors (Smith et al., 1998b; Follsea et al., 2000). Therefore, to determine whether the increase in the abundance of the $\alpha 4$ subunit mRNA induced by discontinuation of ethanol exposure was associated with a similar increase in the amount of the encoded protein at the cell surface, we subjected hippocampal neurons to immunocytofluorescence analysis with a confocal laser-scanning microscope. The specific antibodies were generated in response to an extracellular epitope (peptide N1–19) of the rat $\alpha 4$ subunit and were characterized by immunoblot analysis of a crude membrane fraction of hippocampal neurons, in which they recognized a single protein of ~ 70 kDa (Fig. 6).

A low level of $\alpha 4$ subunit immunoreactivity was detected in control hippocampal neurons, as expected (Pirker et al., 2000); it was localized mostly in the perinuclear region, at the cell membrane, and in association with dendrites and synapses (Fig. 7*A*). In neurons subjected to chronic exposure to ethanol and subsequent ethanol withdrawal for 6 hr, the abundance of the $\alpha 4$ subunit was markedly increased ($F_{(1,78)} = 37.57$; $p < 0.001$) compared with that apparent in control cells (Fig. 7*B, G*). This effect was most pronounced for the $\alpha 4$ protein localized at the cell membrane and in association with dendrites and synapses. Replacement of ethanol with 10 μM diazepam (Fig. 7*C, G*) or 100 mM GHB (Fig. 7*D, G*), but not with 100 μM baclofen (Fig. 7*E, G*), prevented the increase in $\alpha 4$ subunit expression induced by ethanol withdrawal. No immunoreactivity was detected on incubation of neurons with the antibodies to $\alpha 4$ together with the N1–19 peptide antigen (Fig. 7*F*).

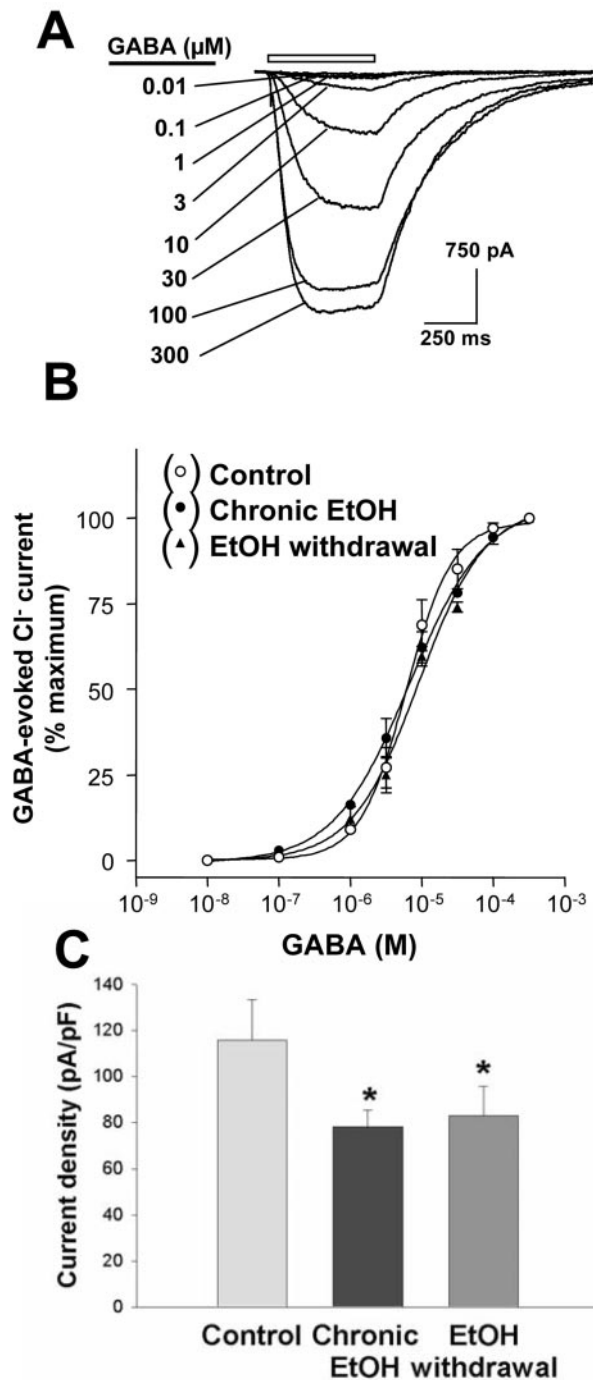


Figure 8. Effects of chronic exposure to and subsequent withdrawal of ethanol on GABA-induced Cl⁻ currents in cultured hippocampal neurons in the whole-cell patch-clamp configuration. *A*, Representative tracings recorded from a control cell showing Cl⁻ currents evoked by GABA at concentrations of 0.01–300 μM. The period of GABA application is indicated by the horizontal bar. *B*, Dose–response curves for GABA-evoked Cl⁻ currents in control cells, cells exposed to 100 mM ethanol for 5 d, and cells subjected to ethanol withdrawal for 6 hr. Data were normalized with respect to the maximal current amplitude apparent at 300 μM GABA and are means ± SE of values from 6–10 neurons. *C*, Maximal current density for GABA_A receptor-mediated currents in neurons of the three experimental groups described in *B*. Data are means ± SE of values from 18–23 neurons. **p* < 0.05 versus control neurons.

Effect of chronic ethanol withdrawal on morphometric parameters

The use of confocal microscopy allowed us to also carefully evaluate the effect of ethanol withdrawal on cell morphology. The

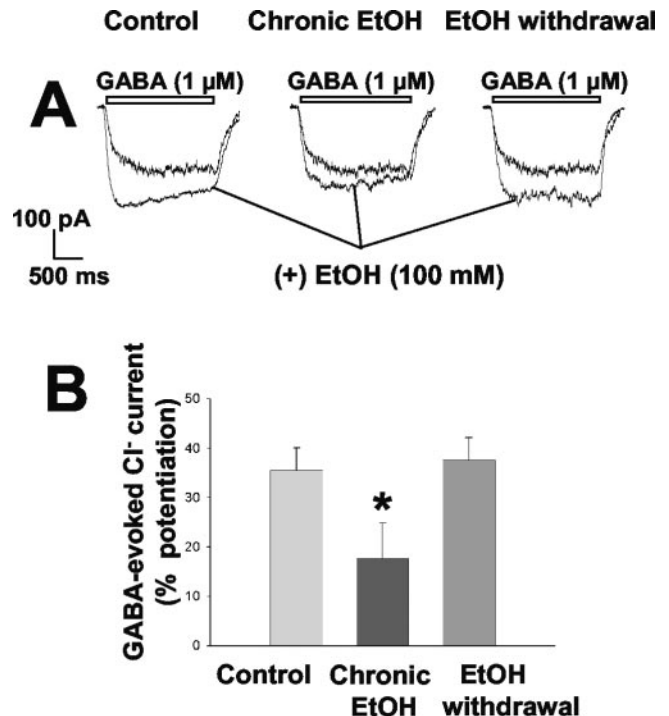


Figure 9. Acute modulatory action of ethanol on GABA_A receptor function in hippocampal neurons subjected to chronic exposure to or withdrawal of ethanol. *A*, Representative tracings of Cl⁻ currents induced by 1 μM GABA in a control cell, a cell exposed to 100 mM ethanol for 5 d, and a cell subjected to ethanol withdrawal for 6 hr. GABA was administered in the absence or presence of 100 mM ethanol. *B*, Quantitation of the acute effects of 100 mM ethanol on GABA-evoked Cl⁻ currents in the three experimental groups. Data are expressed as percentage potentiation of the GABA response and are means ± SE of values from 8–13 neurons. **p* < 0.05 versus value for control cells.

morphometric parameters (area, perimeter, circularity, length, and breadth) measured in cultured hippocampal neurons from control and ethanol withdrawal groups, the latter either in the absence or presence of diazepam (10 μM), GHB (100 mM), or baclofen (100 μM), did not show any statistically significant variation (Table 2).

Effects of chronic exposure to and withdrawal of ethanol on GABA-evoked Cl⁻ currents

We next examined whether the changes in GABA_A receptor gene expression induced in hippocampal neurons by chronic exposure to and subsequent withdrawal of ethanol result in alterations in GABA_A receptor function or sensitivity to various modulatory drugs. For these electrophysiological experiments, the effects of withdrawal were evaluated 6 hr after discontinuation of ethanol exposure (5 d, 100 mM) because preliminary data revealed that the studied changes were maximal at this time. We first determined the dose–response relation for GABA-evoked Cl⁻ currents recorded from hippocampal neurons in the whole-cell patch-clamp configuration (Fig. 8*A,B*). Data were normalized with respect to the maximal current induced by 300 μM GABA. Calculation of EC₅₀ values revealed that chronic exposure to (6.8 ± 1.1 μM) or subsequent withdrawal of (9.0 ± 1.2 μM) ethanol had no significant effect ($F_{(2,21)} = 1.88$; *p* < 0.177) on the apparent potency of GABA compared with the value for control cells (6.1 ± 1.1 μM). The Hill coefficients calculated for control neurons and cells subjected to chronic ethanol treatment or to ethanol withdrawal were 1.3, 1.0, and 1.1, respectively. The maximal density of GABA_A receptor-mediated current (current/ca-

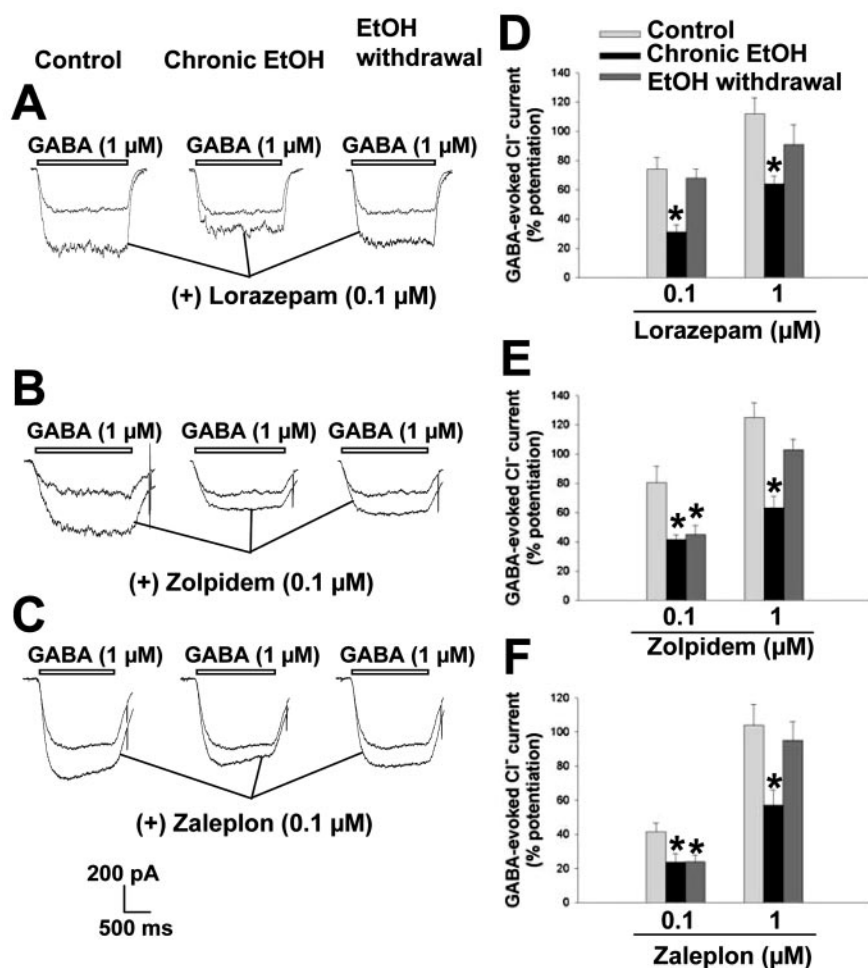


Figure 10. Modulatory action of benzodiazepine receptor agonists on GABA_A receptor function in hippocampal neurons subjected to chronic exposure to or withdrawal of ethanol. *A–C*, Representative tracings of Cl[−] currents induced by 1 μM GABA in control cells, cells exposed to 100 mM ethanol for 5 d, and cells subjected to ethanol withdrawal for 6 hr. GABA was administered in the absence or presence of lorazepam (*A*), zolpidem (*B*), or zaleplon (*C*), each at a concentration of 0.1 μM. *D–F*, Quantitation of the respective modulatory effects of the three test drugs (0.1 or 1 μM) on GABA-evoked Cl[−] currents in the three experimental groups. Data are expressed as percentage potentiation of the GABA response and are means ± SE of values from 9–23 neurons. **p* < 0.05 versus value for control cells.

pacitance) was reduced significantly ($F_{(2,58)} = 6.08$; $p < 0.004$) by 33 or 28% after chronic ethanol treatment or ethanol withdrawal, respectively (Fig. 8C).

Effects of chronic exposure to and withdrawal of ethanol on acute ethanol modulation of GABA_A receptor function

Previous electrophysiological studies have shown that ethanol enhances GABA_A receptor function in hippocampal neurons (Reynolds et al., 1992; Aguayo and Pancetti, 1994; Poelchen et al., 2000), whereas others reported no significant ethanol potentiating effect (Proctor et al., 1992a,b; Soldo et al., 1994). Thus, we next measured the acute effect of ethanol on GABA responses in cultured hippocampal neurons from the control, chronic ethanol exposure, and ethanol withdrawal groups. Acute application of 100 mM ethanol potentiated Cl[−] currents induced by 1 μM GABA (~EC₁₀) in control neurons by 36 ± 5% (Fig. 9). This acute effect of ethanol was inhibited significantly in cells chronically exposed to ethanol (18 ± 8% potentiation) but not in those subjected to ethanol withdrawal (38 ± 5% potentiation) ($F_{(2,28)} = 6.46$; $p < 0.005$).

Effects of chronic exposure to and withdrawal of ethanol on GABA_A receptor sensitivity to benzodiazepine receptor ligands

Given that the pharmacology of benzodiazepine receptor ligands is dependent on the subunit composition of GABA_A receptors, especially with regard to the α and γ subunits (Pritchett et al., 1989; Barnard et al., 1998), we next examined the impact of the changes in subunit mRNA abundance induced by chronic exposure to and withdrawal of ethanol on benzodiazepine receptor pharmacology in hippocampal neurons. We evaluated the effects of positive (lorazepam, zolpidem, zaleplon) and negative (Ro 15-4513, FG 4172) modulators of GABA_A receptors as well as of flumazenil, a competitive antagonist of the benzodiazepine receptor, on control neurons as well as on those subjected to long-term treatment with or withdrawal of ethanol.

The benzodiazepine lorazepam markedly potentiated (74 ± 8 and 112 ± 11% at 0.1 and 1 μM, respectively) the Cl[−] current induced by 1 μM GABA in control neurons (Fig. 10*A,D*). The efficacy of this benzodiazepine was reduced significantly (~50%) in neurons subjected to chronic ethanol treatment compared with that in control neurons. In neurons subjected to ethanol withdrawal, however, the modulatory efficacy of lorazepam was restored to a level similar to that apparent in control neurons (0.1 μM lorazepam, $F_{(2,41)} = 14.72$, $p < 0.001$; 1 μM lorazepam, $F_{(2,52)} = 9.52$, $p < 0.001$). The imidazopyridine zolpidem (Fig. 10*B,E*) and the pyrazolopyrimidine zaleplon (Fig. 10*C,F*), both of which are selective for GABA_A receptors containing the α1 subunit (Sanna et al., 2002), also potentiated GABA-evoked Cl[−] currents in

a concentration-dependent manner in control neurons. Consistent with the lower receptor affinity and modulatory potency of zaleplon compared with zolpidem (Damgen, 1999; Sanna et al., 2002), the potentiating effect of zaleplon in control neurons was smaller than that of zolpidem at a concentration of 0.1 μM but not at 1 μM. The potentiating effects of both compounds at both concentrations tested were significantly reduced (~50%) in neurons subjected to chronic ethanol treatment. In neurons subjected to ethanol withdrawal, whereas the effects of zolpidem and zaleplon at 0.1 μM remained reduced at levels similar to those apparent after chronic ethanol exposure, at the higher concentration (1 μM) both drugs potentiated the GABA response to an extent similar to that observed in control neurons (0.1 μM zolpidem, $F_{(2,30)} = 11.90$, $p < 0.001$; 1 μM zolpidem, $F_{(2,31)} = 14.75$, $p < 0.001$; 0.1 μM zaleplon, $F_{(2,27)} = 4.99$, $p < 0.014$; 1 μM zaleplon, $F_{(2,27)} = 5.58$, $p < 0.009$).

Consistent with its pharmacological profile of a pure antagonist devoid of intrinsic activity, flumazenil (3 μM) did not significantly affect GABA-evoked Cl[−] currents in either control neurons or neurons subjected to long-term treatment with ethanol

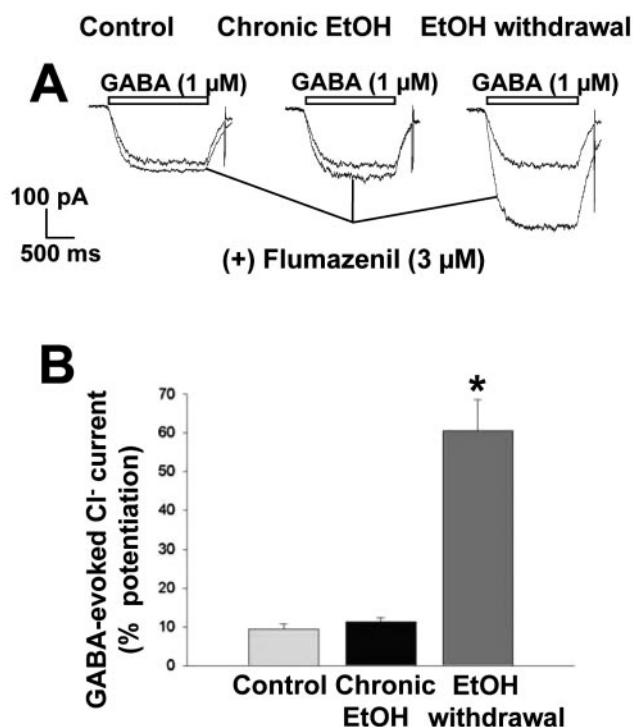


Figure 11. Agonist-like action of the benzodiazepine receptor antagonist flumazenil in hippocampal neurons subjected to ethanol withdrawal. *A*, Representative tracings of Cl⁻ currents induced by 1 μM GABA in a control cell, a cell exposed to 100 mM ethanol for 5 d, and a cell subjected to ethanol withdrawal for 6 hr. GABA was administered in the absence or presence of flumazenil (3 μM). *B*, Quantitation of the modulatory effect of flumazenil on GABA-evoked Cl⁻ currents in the three experimental groups. Data are expressed as percentage potentiation of the GABA response and are means ± SE of values from 8–25 neurons. **p* < 0.01 versus value for control cells.

(Fig. 11). Flumazenil, however, markedly potentiated (60 ± 8%) the GABA response in neurons subjected to ethanol withdrawal ($F_{(2,40)} = 33.68$; $p < 0.001$).

We next investigated the effects of two inverse agonists of the benzodiazepine receptor, the benzodiazepine derivative Ro 15-4513 and the β-carboline FG 7142. Ro 15-4513 (3 μM) markedly inhibited (80 ± 5%) GABA-evoked Cl⁻ currents in control neurons (Fig. 12*A,C*). This inhibitory effect was significantly reduced (25 ± 13%) in neurons subjected to chronic ethanol exposure, and Ro 15-4513 potentiated the GABA response (55 ± 13%) after ethanol withdrawal ($F_{(2,24)} = 81.25$; $p < 0.001$), again consistent with the notion that discontinuation of long-term ethanol treatment induces an increase in the density of α₄-containing GABA_A receptors. FG 7142 (3 μM) inhibited (56 ± 5%) GABA-evoked Cl⁻ currents in control neurons (Fig. 12*B,C*). This effect was significantly reduced in neurons subjected to chronic ethanol treatment or to subsequent withdrawal ($F_{(2,20)} = 11.12$; $p < 0.001$).

Effects of diazepam, GHB, and baclofen on GABA_A receptor pharmacology during ethanol withdrawal

Finally, we examined whether diazepam, GHB, and baclofen were able to block the changes in pharmacological sensitivity of GABA_A receptors induced by ethanol withdrawal. Incubation of neurons with either 10 μM diazepam (Fig. 13) or 100 mM GHB (Table 3) during the 6 hr period of ethanol withdrawal resulted in a significant inhibition of the modulatory efficacy of 1 μM lorazepam and reversed the agonist-like actions of 3 μM flumazenil

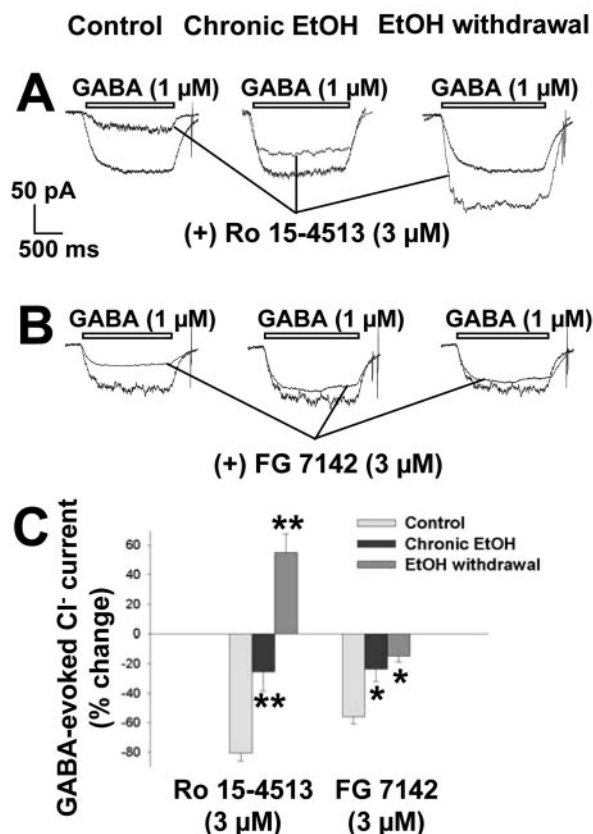


Figure 12. Modulatory effects of the benzodiazepine receptor inverse agonists Ro 15-4513 and FG 7142 on GABA_A receptor function in hippocampal neurons subjected to chronic exposure to or withdrawal of ethanol. *A, B*, Representative tracings of Cl⁻ currents induced by 1 μM GABA in control cells, cells exposed to 100 mM ethanol for 5 d, and cells subjected to ethanol withdrawal for 6 hr. GABA was administered in the absence or presence of Ro 15-4513 (*A*) or FG 7142 (*B*), each at a concentration of 3 μM. *C*, Quantitation of the modulatory effects of Ro 15-4513 and FG 7142 on GABA-evoked Cl⁻ currents in the three experimental groups. Data are expressed as percentage change in the GABA response and are means ± SE of values from 5–12 neurons. **p* < 0.05, ***p* < 0.01 versus value for control cells.

and 3 μM Ro 15-4513 (1 μM lorazepam, $F_{(1,23)} = 50.73$, $p < 0.001$; 3 μM flumazenil, $F_{(1,20)} = 39.14$, $p < 0.001$; 3 μM Ro 15-4513, $F_{(1,20)} = 118.57$, $p < 0.001$).

To determine whether the action of GHB might involve an interaction with GABA_B receptors, we exposed neurons to both 100 mM GHB and 100 μM SCH 50911, a competitive antagonist of the GABA_B receptor, during ethanol withdrawal. SCH 50911 failed to antagonize the inhibitory effect of GHB on the changes in GABA_A receptor sensitivity to lorazepam, flumazenil, or Ro 15-4513 induced by ethanol withdrawal (Table 3). Furthermore, baclofen (100 μM) did not mimic this action of GHB (Table 3).

Discussion

A correlation of alterations in GABA_A receptor function and pharmacological sensitivity with specific changes in receptor gene expression after prolonged ethanol treatment has been explored previously (Devaud et al., 1996; Kang et al., 1998; Cagetti et al., 2003; Follsea et al., 2003).

We have shown further that chronic ethanol exposure and its subsequent discontinuation induce marked changes in GABA_A receptor function and pharmacological responsiveness to both nonselective and selective benzodiazepine receptor ligands. Furthermore, these changes may result from alterations in the abundance of mRNAs encoding α1–4 and γ2 subunits as well as in

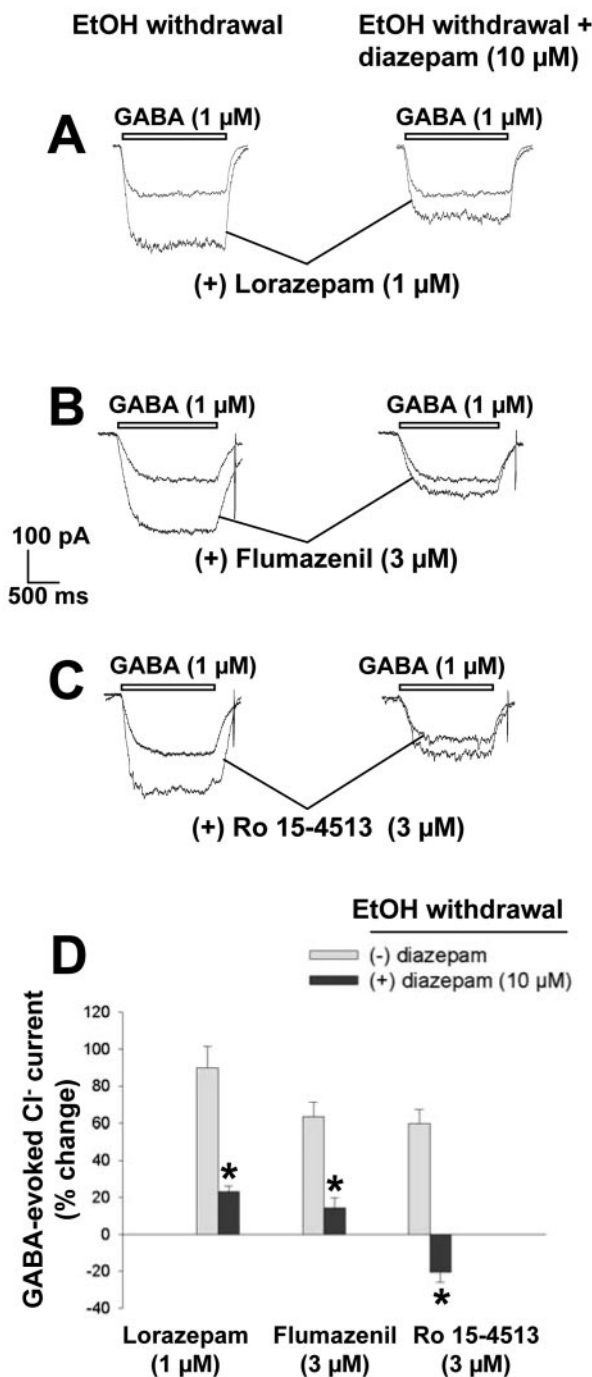


Figure 13. Effects of diazepam on the changes in GABA_A receptor sensitivity to lorazepam, flumazenil, or Ro 15-4513 induced by ethanol withdrawal. *A–C*, Representative traces of Cl[−] currents induced by 1 μM GABA in hippocampal neurons that had been incubated first for 5 d with 100 mM ethanol and then for 6 hr in ethanol-free medium in the absence or presence of 10 μM diazepam. GABA was applied in the absence or presence of 1 μM lorazepam (*A*), 3 μM flumazenil (*B*), or 3 μM Ro 15-4513 (*C*). *D*, Quantitation of the modulatory effects of lorazepam, flumazenil, and Ro 15-4513 on GABA-evoked Cl[−] currents in the two experimental groups. Data are expressed as percentage change in the GABA response and are means ± SE of values from 9–16 neurons. **p* < 0.01 versus the corresponding value for cells subjected to ethanol withdrawal in the absence of diazepam.

that of the α4 protein. The pattern of changes in GABA_A receptor subunit mRNA levels observed in our study differs to some extent from that of other previous reports, particularly regarding the γ2 subunit, the expression of which decreased in our study after

prolonged ethanol exposure and withdrawal, was unchanged after 14 or 40 d of ethanol treatment (Matthews et al., 1998), or increased in chronic intermittent ethanol-treated rats (Cagetti et al., 2003).

GABA_A receptor function after prolonged ethanol exposure and its discontinuation

Exposure of hippocampal neurons to ethanol induced a marked reduction in the maximal Cl[−] current density attributable to GABA_A receptors. This effect was still apparent 6 hr after discontinuation of ethanol exposure. Given that the potency of GABA was not significantly affected by either treatment, this reduction in current density appears to be caused by a decrease in the cell surface density of GABA_A receptors. This conclusion is consistent with our observation that the abundance of α1, γ2L, and γ2S subunit mRNAs was reduced after chronic ethanol exposure and its subsequent discontinuation and is in line with recently published data showing that ethanol exposure reduces cell surface expression of α1 subunit-containing GABA_A receptor subtypes (Kumar et al., 2003). Indeed, a marked decrease in the efficiency of α1βγ2 receptor assembly has been demonstrated in the α1 subunit knock-out mice (Sur et al., 2001; Kralic et al., 2002). Furthermore, given that ethanol withdrawal was also associated with an increased abundance of α2, α3, and α4 subunit mRNAs, such treatment, in addition to maintaining a reduction in GABA_A receptor density, likely results in a shift in the pattern of receptor subunit assembly.

Differential effects of nonselective and selective benzodiazepine receptor ligands after prolonged ethanol exposure and its abrupt discontinuation

The reduction in the abundance of the α1, α3, γ2L, and γ2S subunit mRNAs induced by prolonged ethanol exposure may be responsible for the associated functional uncoupling between the neurotransmitter binding site and the modulatory benzodiazepine recognition site of the GABA_A receptor, as is apparent from the loss of pharmacological efficacy of benzodiazepine receptor ligands. Consistent with previous evidence (Morrow et al., 1988; Buck and Harris, 1990; Sanna et al., 1993), the efficacy of lorazepam, zolpidem, and zaleplon was significantly reduced after prolonged exposure of hippocampal neurons to ethanol. The negative modulatory efficacy of the benzodiazepine receptor inverse agonists Ro 15-4513 and FG 7142 was also reduced in the ethanol-treated cells.

Discontinuation of chronic ethanol exposure induced distinct changes in the sensitivity of GABA_A receptors to the various benzodiazepine recognition site ligands. These changes appear to be dependent on the alterations in GABA_A receptor subunit expression triggered by such treatment. Potentiation of GABA-evoked

Table 3. Effects of GHB and GABA_B receptor-specific drugs on ethanol withdrawal-induced changes in GABA_A receptor sensitivity to benzodiazepine receptor ligands

Drug	GABA-evoked Cl [−] current (% change)			
	Vehicle	GHB	GHB + SCH 50911	Baclofen
Lorazepam	91 ± 14	23 ± 3*	37 ± 5*	109 ± 14
Flumazenil	63 ± 8	14 ± 5*	10 ± 3*	46 ± 5
Ro 15-4513	59 ± 8	−20 ± 5*	−27 ± 6*	54 ± 8

Currents induced by 1 μM GABA were measured in hippocampal neurons that had been incubated first for 5 d with 100 mM ethanol and then for 6 hr in ethanol-free medium in the absence or presence of 100 mM GHB (with or without 100 μM SCH 50911) or 100 μM baclofen. GABA was applied in the absence or presence of 1 μM lorazepam, 3 μM flumazenil, or 3 μM Ro 15-4513. Data are expressed as percentage change in the GABA response and are means ± SE of values from 12–22 neurons. **p* < 0.05 versus the corresponding value for ethanol withdrawal in the absence or drug (vehicle).

Cl⁻ currents by lorazepam was restored to control levels 6 hr after ethanol withdrawal. Lorazepam is a classic benzodiazepine derivative that binds with similar high affinities and modulates GABA-evoked Cl⁻ currents with similar potencies at GABA_A receptors containing α 1–3 or α 5 subunits in combination with β and γ 2 subunits; it does not exhibit substantial affinity for receptors containing α 4 or α 6 subunits (Sieghart, 1995; Barnard et al., 1998). Restoration of the modulatory efficacy of lorazepam to control values after discontinuation of ethanol exposure may thus result from increased expression of the α 2 and α 3 subunits.

Consistent with this line of reasoning, the efficacies of zolpidem and zaleplon at 0.1 μ M, a concentration at which both drugs *in vitro* modulate selectively GABA responses mediated by α 1-containing receptors (Sanna et al., 2002), remained reduced after ethanol withdrawal at a level similar to that observed during chronic ethanol exposure. At a higher concentration (1 μ M), at which these drugs, like lorazepam, act nonselectively at α 1-, α 2-, or α 3-containing receptors (Sanna et al., 2002), the efficacies of zolpidem and zaleplon were restored to control levels after ethanol withdrawal. The inhibitory action of FG 7142, a β -carboline with preferential affinity for α 1-containing receptors (Sieghart, 1995; Barnard et al., 1998), was also reduced by chronic ethanol exposure and remained so after ethanol withdrawal.

Given that GABA_A receptor subtypes containing α 1 or α 2 subunits mediate the sedative and anxiolytic effects of benzodiazepines, respectively (Mohler et al., 2002), our results are consistent with the reduced sedative efficacy of benzodiazepines in human alcoholics as well as with the ability of these drugs to reduce the anxiogenic effect of ethanol withdrawal (Sellers et al., 1983; Lejoyeux et al., 1998).

Upregulation of the α 4 subunit during ethanol withdrawal

Consistent with previous observations in the cerebral cortex, hippocampus, and cerebellar granule cells (Devaud et al., 1995; Devaud et al., 1997; Mahmoudi et al., 1997; Matthews et al., 1998; Cagetti et al., 2003; Follesa et al., 2003), we have shown that ethanol withdrawal resulted in a marked increase in the abundance of the α 4 subunit mRNA in cultured hippocampal neurons. This effect was accompanied by a pronounced increase in the amount of the α 4 protein that was apparent predominantly at the cell membrane and in association with dendrites and synapses.

Upregulation of the α 4 subunit is also induced by withdrawal of benzodiazepine receptor ligands (Follesa et al., 2001, 2002) or neurosteroids (Smith et al., 1998a,b; Follesa et al., 2000), suggesting that it might play an important role in the cellular hyperexcitability and anxiety-like behavior apparent in both animals and humans during withdrawal from these positive allosteric modulators of GABA_A receptors.

GABA exhibits a high affinity but relatively low efficacy at GABA_A receptors containing the α 4 subunit; these receptors are also insensitive to classic benzodiazepines and possess high affinity for flumazenil and Ro 15-4513 (Wafford et al., 1996; Whittemore et al., 1996). Consistent with recent data showing an increased sensitivity to Ro 15-4513 associated with an enhanced expression of the α 4 subunit after chronic ethanol exposure in chronic intermittent ethanol-treated rats (Cagetti et al., 2003), we showed that this compound and flumazenil, a partial inverse agonist and antagonist, respectively, in control cells, both acted as positive modulators of GABA responses after discontinuation of prolonged ethanol treatment, consistent with the notion that ethanol withdrawal increases the abundance of GABA_A receptors containing the α 4 subunit. Flumazenil ameliorates ethanol with-

drawal symptoms such as anxiety and hyperexcitability in animals and human alcoholics (File et al., 1989; Buck et al., 1991; Gerra et al., 1991; Nutt et al., 1993; Moy et al., 1997), an effect that has been proposed to result from blockade of the action of a putative endogenous benzodiazepine receptor ligand endowed with inverse agonist activity. Our data now suggest that this effect of flumazenil might also be attributable to an increase in the density of α 4-containing receptors associated with ethanol withdrawal.

Prevention of ethanol withdrawal-induced molecular and functional effects by diazepam or GHB

We have now shown that the presence of diazepam or GHB, drugs used to treat ethanol dependence, craving, and withdrawal syndrome (Addolorato et al., 1996; Lejoyeux et al., 1998) during the period of ethanol withdrawal, prevented the increases in the abundance of α 2, α 3, and α 4 subunit mRNAs, the upregulation of the α 4 protein, the restoration of lorazepam efficacy, and the conferment of positive modulatory action on both flumazenil and Ro 15-4513 induced by such withdrawal. Neither drug, however, altered the molecular and functional changes associated with prolonged exposure to ethanol, including the reduced abundance of α 1, α 3, γ 2L, and γ 2S subunit mRNAs and the reduced efficacy of lorazepam and other GABAergic modulators.

The inhibitory effects of diazepam during ethanol withdrawal are consistent with its mechanism of action at the GABA_A receptor, whereas the mechanism by which GHB elicits its effects is not clear. Despite its similarities to GABA and GABAergic drugs in terms of chemical structure and pharmacological profile, GHB does not possess activity at GABA_A receptors (Serra et al., 1991; Feigenbaum and Howard, 1996; Follesa et al., 2003). GHB has been suggested to exert its central depressant effects by increasing the synthesis and extracellular concentration of GABA in specific brain regions (Gobaille et al., 1999). Furthermore, administration of GHB, like that of ethanol (Morrow et al., 2001), has been shown to increase the formation of neuroactive steroids in rats (Barbaccia et al., 2002), an effect mediated by GABA_B receptors. The accumulation of neuroactive steroids in the brain would be expected to result in an increased GABAergic tone; however, the GABA_B receptor antagonist SCH 50911 failed to inhibit the action of GHB during ethanol withdrawal, and the GABA_B receptor agonist baclofen did not mimic GHB action, suggesting that GABA_B receptors do not contribute to the effects of GHB in our experimental model.

Conclusions

We have shown that prolonged exposure to and subsequent withdrawal of ethanol are associated with marked, specific, and opposite changes in GABA_A receptor subunit gene expression as well as in receptor function and pharmacological sensitivity in cultured rat hippocampal neurons. Downregulation of GABA_A receptors and a reduction in the efficacy of various benzodiazepine receptor ligands induced by prolonged ethanol treatment are associated with a reduced expression of α 1, α 3, γ 2L, and γ 2S subunits. In contrast, an increase in the abundance of α 4-containing receptors induced by ethanol withdrawal may be an important determinant of withdrawal syndrome and is blocked by drugs that are effective in the treatment of ethanol dependence.

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