

TEMPORAL AND REGIONAL REGULATION OF α 1, β 2 AND β 3, BUT NOT α 2, α 4, α 5, α 6, β 1 OR γ 2 GABA_A RECEPTOR SUBUNIT MESSENGER RNAS FOLLOWING ONE-WEEK ORAL FLURAZEPAM ADMINISTRATION

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Abstract—The effect of prolonged benzodiazepine administration on GABA_A receptor subunit (α1-6, β 1-3, γ 2) messenger RNAs was investigated in the rat hippocampus and cortex, among other brain areas. Rats were orally administered flurazepam for one week, a protocol which results in benzodiazepine anticonvulsant tolerance in vivo, and in the hippocampus in vitro, in the absence of behavioral signs of withdrawal. Autoradiographs of brain sections, hybridized with [35S]oligoprobes in situ, were examined immediately (day 0) or two days after drug treatment, when rats were tolerant, or seven days after treatment, when tolerance had reversed, and were compared to sections from pair-handled, vehicle-treated controls. α1 subunit messenger RNA level was significantly decreased in CA1 pyramidal cells and dentate granule cells at day 0, an effect which persisted only in CA1 neurons. Decreased "α1-specific" silver grain density over a subclass of interneurons at the pyramidal cell border suggested concomitant regulation of interneuron GABA_A receptors. A reduction in β3 subunit messenger RNA levels was more widespread among hippocampal cell groups (CA1, CA2, CA3 and dentate gyrus), immediately and two days after treatment, and was also detected in the frontal and parieto-occipital cortices. Changes in β2 subunit messenger RNA levels in CA1, CA3 and dentate gyrus cells two days after ending flurazepam treatment suggested a concomitant up-regulation of β2 messenger RNA. There was a trend toward an increased level of α 5, β 3 and γ 2 subunit messenger RNAs in CA1, CA3 and dentate gyrus cells, which was significant for the β 3 and γ 2 subunit messenger RNAs in the frontal cortex seven days after ending flurazepam treatment. There were no flurazepam treatment-induced changes in any other GABAA receptor subunit messenger

The messenger RNA levels of three ($\alpha 1$, $\beta 2$ and $\beta 3$) of nine GABA_A receptor subunits were discretely regulated as a function of time after ending one-week flurazepam treatment related to the presence of anticonvulsant tolerance, but not dependence. The findings suggested that a localized switch in the subunit composition of GABA_A receptor subtypes involving these specific subunits may represent a minimal requirement for the changes in GABA_A receptor-mediated function recorded previously at hippocampal CA1 GABAergic synapses, associated with benzodiazepine anticonvulsant tolerance. © 1999 IBRO. Published by Elsevier Science Ltd.

Key words: benzodiazepines, GABA, hippocampus, dependence, tolerance, in situ hybridization.

Several classes of clinically effective anticonvulsants exert their actions at different allosteric modulatory sites on the GABA_A receptor (cf. Ref. 62). Continual use of the benzodiazepine class leads to a reduction in clinical effectiveness, i.e. tolerance. It is well established that benzodiazepine tolerance is associated with a dysfunction of GABAergic neurotransmission (for reviews see Refs 3, 21 and 30). GABA_A receptors gate fast inhibitory synaptic transmission via an integral chloride ion channel. The native receptor is a pentamer with unknown stoichiometry, though it is probably composed of 2α , 2β and a γ (or δ) subunit. 29,39 Multiple variants of the five GABA_A receptor subunit families have

GABA_A receptors have a dense, heterogeneous distribution throughout the CNS, particularly in the cortex and hippocampus. ⁴⁸ At least five patterns of GABA_A receptor subunit association have been identified *in vivo*^{5,17,18,41,44,52,75} and certain subunit combinations have been associated with specific neuronal populations. ^{23,41,44} The heterogeneity and subcellular compartmentalization of GABA_A receptor subunits suggest that multiple, functionally distinct GABA_A receptor subtypes may be assembled on individual neurons. ⁴⁴

A change in expression of the genes encoding GABA_A receptor subunits, and thus a switch in the subunit composition of GABA_A receptor subtypes, has been proposed to contribute to the dysfunction at

been identified ($\alpha 1-6$, $\beta 1-4$, $\gamma 1-4$, $\delta 1$, $\epsilon 1$), which may confer different Cl⁻ channel properties and distinct GABA and benzodiazepine pharmacological properties. ^{13,14,39,57,63,70,71}

^{*}To whom correspondence should be addressed. Abbreviations: DG, dentate gyrus; SO, stratum oriens; SP, stratum pyramidale; SR, stratum radiatum; SSC, standard saline

α₁ Subunit

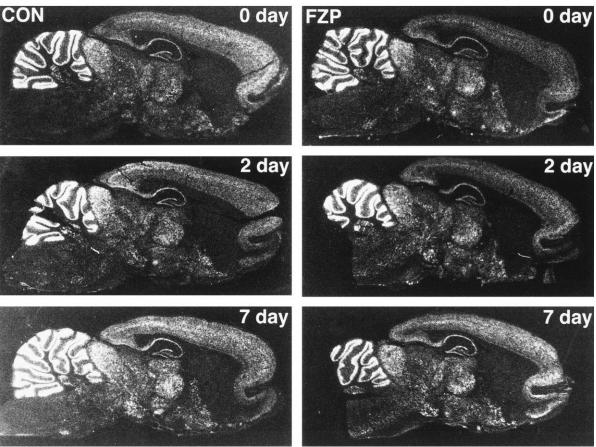


Fig. 1. Representative dark-field photomicrographs of autoradiographs of sagittal brain sections hybridized with an antisense [35 S]oligoprobe for the α 1 subtype mRNA of the GABA_A receptor. Sections (10 μ m) were cut from one-week flurazepam-treated (FZP) rats killed immediately (day 0), two or seven days after ending oral drug administration and compared to sections from matched control rats (CON). Immediately after the end of treatment, there was a significant decrease in the relative gray level on autoradiographs localized over CA1 pyramidal cells and dentate granule cells. This difference only persisted in CA1 pyramidal cells two days after treatment and was no longer present seven days after ending treatment, when rats are no longer tolerant *in vivo* (Table 1).

GABAergic synapses associated with benzodiazepine tolerance. Variable expression of receptor subunits could alter GABA receptor function and GABA or benzodiazepine pharmacology, and could be manifested behaviorally as a decreased CNS sensitivity to the drug following repeated or prolonged administration. Studies aimed establishing such a role for changes in GABAA receptor subunit composition associated with benzodiazepine tolerance have yielded conflicting findings. 20,31,33,42,73 One potential difficulty may be that some benzodiazepine treatments produce not only tolerance, but also physical dependence. It could then become difficult to differentiate molecular mechanisms underlying reduced neuronal responsiveness to benzodiazepines, i.e. functional tolerance, from similar or additional changes in GABA_A receptors occurring in response to repeated or continued receptor activation, but associated with physical dependence. Dependence may be manifested as behavioral hyperexcitability following drug withdrawal or precipitated, i.e. by antagonist administration. Another difficulty in interpreting the results of a variety of previous studies is the identification of a common anatomical locus or loci mediating functional tolerance. Moreover, whereas studies of tolerance mechanisms in heterologous expression systems following prolonged agonist exposure may yield important insights into the possible relationship between biochemical and molecular mechanisms underlying pharmacological or functional changes at GABA_A receptors, these mechanisms cannot be directly ascribed to tolerance *in vivo*.

One-week oral flurazepam treatment produces tolerance to benzodiazepine actions to suppress pentylenetetrazole-induced seizures *in vivo*, ^{59,60} but not dependence. ⁶⁷ Tolerance was also demonstrated *in vitro* in a model system established in the hippocampus of flurazepam-treated rats, by a reduced

sensitivity of GABA-mediated responses in CA1 cells to benzodiazepine receptor agonists. 78,81,83 GABA_A receptor agonist potency was also reduced in CA1 pyramidal cells. 78 Moreover, among the principal hippocampal cell types studied following chronic benzodiazepine treatments with a range of electrophysiological approaches, GABAergic function was shown to be selectively impaired in CA1 pyramidal cells. 53,54,79-82 Furthermore, the temporal pattern of the changes in GABA function detected in CA1 pyramidal cells following one-week flurazepam treatment 53,81,82,84 paralleled the appearance and disappearance of anticonvulsant tolerance in vivo. 59,60 Using in situ hybridization methods, we had previously demonstrated localized decreases in $\alpha 1$, but not $\alpha 5$ and $\gamma 2$, mRNA levels in the cortex and CA1 region of the hippocampus two days after ending flurazepam administration. In order to establish whether the expression of specific GABA_A receptor subunits may be discretely regulated with a similar timecourse as the temporal pattern of tolerance detected in vivo and in vitro, the relative expression of nine GABA_A receptor subunit (α 1, α 2, α 4, α 5, α 6, β 1, β 2, β 3 and γ 2) mRNAs was investigated in the hippocampus, among other brain areas.

EXPERIMENTAL PROCEDURES

Chronic benzodiazepine treatment

After a two-day adaptation period, male Sprague-Dawley rats (180-200 g initial weight; Harlan, Indianapolis, IN) were offered flurazepam hydrochloride in 0.02% saccharin water for one week (100 mg/kg for three days, 150 mg/kg for four days). Rats treated using this protocol are tolerant, 59,60 but show no signs of spontaneous or precipitated withdrawal.67 Though rats are offered what are apparently relatively "high" doses of flurazepam, benzodiazepine levels in the brain achieved during one-week oral flurazepam administration, expressed in diazepam equivalents (161.1 \pm 35.9 ng diazepam/g brain; 0.57 μ M), ^{53,66,77,78} are similar to or less than those achieved with other chronic benzodiazepine treatments (200–275 ng diazepam/g brain; $0.7{-}1.0~\mu \dot{M}).^{20,42,58}$ At the end of the drug administration period, rats were killed for brain section preparation. Rats were killed immediately (day 0) after ending flurazepam administration or offered saccharin water for two or seven days until they were killed. Control rats received the saccharin solution for the same lengths of time. Changes in mRNA levels were anticipated to be at or near maximal immediately after drug removal, or still decreased two days after drug removal, when impaired GABAA receptor inhibition was detected in the hippocampus in the absence of residual benzodiazepine metabolites.⁷⁷ Seven days after the end of treatment, when tolerance has reversed, the electrophysiological correlates of tolerance have also disappeared. 53,77,81,82,

In accordance with the Public Health Service Policy on Human Care and Use of Laboratory Animals and the Animal Welfare Act, all efforts were made to minimize animal suffering, to reduce the numbers of animals used and to utilize alternatives to *in vivo* techniques.

In situ hybridization histochemistry

Oligonucleotide probes. Oligoprobes (45-mers) complimentary to rat cDNA amino acid residues (α1, 342–356;³⁴

α2, 341–357;³⁵ α3, 361–375;⁴⁰ α4, 15–30 of the signal peptide;⁷⁴ α5, 355–369,⁴⁰ α4 according to Khrestchatisky *et al.*;³⁴ α6, 342–356;³⁸ β1, 382–396;⁷⁹ β2, 382–396;⁷⁹ β3, 380–394;⁷⁹ γ2, 338–352⁶¹) were synthesized by Oligos Etc. (Willsonville, OR) according to Wisden *et al.*⁷⁵ Probes were 3' end-labeled with [³⁵S]dATP (12.5 mCi/ml; E. I. Dupont, Boston, MA) with terminal deoxytransferase (Boehringer Mannheim, Indianapolis, IN). ³⁵S-Labeled oligoprobe was separated from unincorporated [³⁵S]dATP by centrifugation on a Bio-Spin chromatography column (BioRad, Richmond, CA). Labeling efficiency ranged from 67% to 90% for all oligoprobes, except the α3 probe (\sim 50%). Coupled with the low levels of the α3 subunit mRNA, particularly in the hippocampus, ^{72,75} preliminary experiments indicated that the very weak α3 antisense [³⁵S]oligoprobe hybridization signal on autoradiographs could not be reliably quantified. Therefore, between-group comparisons with the α3 probe were not carried out.

Tissue preparation. Brains were dissected and rapidly frozen in isopentane in an acetone-dry ice bath. Sagittal serial sections (10 µm) were cut through the hippocampal formation, 2.4-2.9 mm lateral to the midline, according to Paxinos and Watson.⁵⁰ Sections were thaw-mounted on to poly-L-lysine-coated slides and stored at -70°C until processed for in situ hybridization histochemistry. Prior to hybridization, sections were rapidly brought to room temperature under vacuum and fixed for 5 min in 4% paraformaldehyde in 0.1 M sodium phosphate buffer (pH 7.4). For each oligoprobe, parallel serial sections were selected from flurazepam-treated and control groups based on their lateral level of section. 50 Sections were rinsed 3×5 min in sodium phosphate buffer. Sections were immersed for 10 min in 0.1 M triethanolamine containing 0.25% acetic anhydride then rinsed for 10 min in 2 × standard saline citrate (SSC; $1 \times SSC = 0.15 \text{ M}$ NaCl, 0.15 M sodium citrate, pH 7.0), dehydrated through an ethanol series (70%, 80%, 95%), defatted in chloroform and stored under 95% ethanol at -20° C until used. For each [35S]oligoprobe, one section per rat from both experimental and control groups were batch processed in parallel and handled identically across all time-points.

Hybridization. Sections were air-dried and prehybridized with 80 µl/slide of hybridization buffer [50% (v/v) formamide, 4 × SSC, 1 × Denhardt's reagent, 500 mg/μl heat denatured Herring sperm DNA, 250 ml yeast tRNA and 10% dextran sulfate containing 10 mM dithiothreitol] under parafilm in a humidified $(2 \times SSC)$ chamber. Slides treated with RNase for 30 min at 37°C prior to prehybridization were used as negative controls.⁶⁶ Sections were hybridized at 42°C in a humidified chamber with 50 ml/slide hybridization buffer containing 10 mM dithiothreitol and 1 × 10⁶ d.p.m. ³⁵S-labeled antisense oligoprobe/section under glass coverslips with parafilm bridges. Additional slides were incubated with sense $\alpha 1$ and $\beta 3$ [35S]oligoprobes. Coverslips were removed in 2 × SSC. Sections were washed to a final stringency of $0.5 \times SSC$ at $52-60^{\circ}C$ according to the calculated oligoprobe $T_{\rm m}$. Sections were dehydrated for 5 min in 70% and 95% ethanol containing 300 mM ammonium acetate (pH 5.5), followed by 100% ethanol. Slides were dried in a vacuum desiccator and apposed to Biomax MR film (Eastman Kodak Co., Rochester, NY). The development of the image on the film was tracked with sections hybridized with increasing ratios of oligoprobe labeled with [35S]dATP to "cold" dATP (40%, 80% and 100%) to help insure that the signal was adequate, but not saturated. Coronal sections, hybridized with the 35 S-labeled $\alpha1$ antisense probe, were dipped in NTB-2 emulsion (Eastman Kodak Co., Rochester, NY) as described previously 66 for grain counting over presumptive interneurons in the hippocampal CA1 region. Autoradiographic film and emulsion were

Table 1. Time-dependent $\alpha 1$, $\beta 2$ and $\beta 3$ subunit messenger RNA levels in flurazepam-treated rat brain

| | | | α1 subunit | | | β2 subunit | | | β3 subunit | |
|---|-----------------------------------|---|--|--|--|---|--|---|--|---|
| | | 0 days | 2 days | 7 days | 0 days | 2 days | 7 days | 0 days | 2 days | 7 days |
| Groups | Control FZP treated | n = 8 $n = 8$ | n = 7 $n = 8$ | n = 8 $n = 8$ | n = 8 $n = 8$ | n = 8 $n = 7$ | n = 7 $n = 8$ | $n=8\\n=8$ | n = 7 $T = n$ | n = 8 $n = 8$ |
| Hippocampus proper CA1 pyramidal cells | Control FZP treated P value | $79.1 \pm 2.7 \\ 67.7 \pm 4.3 \\ 0.03*$ | $67.0 \pm 3.3 \\ 59.6 \pm 2.4 \\ 0.04*$ | $71.6 \pm 5.1 \\ 67.4 \pm 4.3 \\ 0.53$ | 40.1 ± 5.0 39.4 ± 7.8 0.93 | 33.4 ± 3.4 43.8 ± 3.7 $0.05*$ | 42.9 ± 6.3 45.6 ± 6.9 0.76 | 81.9 ± 6.1 62.3 ± 7.1 0.04* | 98.7 ± 4.3 82.3 ± 6.4 0.04* | 80.4 ± 3.7 89.0 ± 5.0 0.16 |
| CA2 pyramidal cells | Control FZP treated P value | 70.9 ± 6.7 69.7 ± 5.6 0.89 | $61.3 \pm 4.3 \\ 61.1 \pm 1.4 \\ 0.87$ | $64.1 \pm 4.4 \\ 61.6 \pm 3.5 \\ 0.65$ | 30.3 ± 4.6 30.3 ± 6.9 1.0 | $27.2 \pm 3.2 \\ 26.9 \pm 3.1 \\ 0.93$ | 31.6 ± 3.8 39.0 ± 5.2 0.25 | 72.3 ± 6.8 61.6 ± 4.6 0.04* | 92.0 ± 5.5 75.8 ± 3.9 0.02** | $77.8 \pm 2.5 \\ 80.5 \pm 3.4 \\ 0.51$ |
| CA3 pyramidal cells | Control FZP treated P value | $53.7 \pm 3.0 \\ 46.2 \pm 3.3 \\ 0.10$ | 46.4 ± 2.1 45.6 ± 2.3 0.66 | $46.7 \pm 3.5 \\ 48.1 \pm 2.3 \\ 0.73$ | $27.9 \pm 2.7 \\ 25.4 \pm 6.4 \\ 0.69$ | 28.4 ± 3.8 39.0 ± 3.2 0.04* | $24.6 \pm 4.0 \\ 30.8 \pm 4.3 \\ 0.29$ | 86.5 ± 5.9 70.6 ± 4.3 0.04* | 97.7 ± 4.0 84.3 ± 2.9 0.04* | 82.7 ± 4.2 97.4 ± 3.6 0.01* |
| Dentate gyrus Polymorph cells (CA4 | Control | 45.5 ± 3.8 | 30.9 ± 0.9 | 38.9 ± 3.3 | 24.3 ± 3.0 | 22.5 ± 3.0 | 24.3 ± 4.0 | 93.6 ± 6.7 | 98.7 ± 4.2 | 88.2 ± 5.2 |
| pyramidal cells) | FZP treated P value | 37.5 ± 3.4 0.11 | 32.0 ± 2.1 0.37 | 39.2 ± 2.2 0.94 | 25.2 ± 6.4 0.89 | 25.9 ± 2.8 0.40 | 28.9 ± 4.0 0.40 | 74.0 ± 6.3 0.11 | $81.7 \pm 3.9 \\ 0.001*$ | 92.6 ± 4.7 0.51 |
| Granule cells | Control FZP treated P value | 72.2 ± 3.2 58.1 ± 4.1 0.01* | $55.3 \pm 3.1 \\ 51.0 \pm 3.3 \\ 0.26$ | $65.1 \pm 4.1 \\ 63.7 \pm 4.5 \\ 0.82$ | 32.5 ± 3.4 32.7 ± 7.0 0.97 | 29.4 ± 2.9 37.9 ± 2.6 0.04* | 34.0 ± 4.5 34.9 ± 4.7 0.88 | $113.9 \pm 7.2 \\ 89.7 \pm 8.1 \\ 0.03*$ | $127.5 \pm 3.1 \\ 108.9 \pm 3.7 \\ 0.001*$ | 105.7 ± 5.6 113.6 ± 3.8 0.23 |
| Cortex Frontal | Control FZP treated P value | $53.7 \pm 5.9 \\ 44.6 \pm 7.2 \\ 0.31$ | 47.5 ± 5.0 46.6 ± 3.0 0.48 | 54.8 ± 5.0 45.6 ± 5.5 0.20 | 30.9 ± 3.6 22.3 ± 6.1 0.20 | $23.0 \pm 3.3 \\ 23.5 \pm 2.8 \\ 0.91$ | $24.8 \pm 9.0 \\ 22.4 \pm 2.6 \\ 0.46$ | 35.1 ± 4.3 22.0 ± 2.6 0.01* | 50.5 ± 3.6 49.5 ± 2.4 0.80 | 24.2 ± 9.0 34.0 ± 3.1 0.02* |
| Parieto-occipital | Control FZP treated P value | 47.7 ± 3.1 41.9 ± 1.6 0.09 | 42.8 ± 4.0 33.6 ± 3.0 0.24 | $47.1 \pm 2.3 \\ 41.1 \pm 3.3 \\ 0.13$ | $25.1 \pm 2.2 \\ 23.0 \pm 3.6 \\ 0.60$ | $22.9 \pm 2.6 \\ 16.4 \pm 2.2 \\ 0.06$ | $27.4 \pm 4.3 \\ 20.3 \pm 3.5 \\ 0.38$ | 31.1 ± 3.1 24.0 ± 3.6 0.14 | 48.6 ± 4.0 41.9 ± 2.0 0.14 | $25.7 \pm 3.0 25.0 \pm 2.7 0.86$ |
| Caudate-putamen | Control FZP treated P value | 111 | 111 | 111 | 111 | 111 | 111 | 29.7 ± 3.8 21.0 ± 1.7 0.03* | 49.1 ± 2.7 45.1 ± 0.6 0.15 | $26.4 \pm 3.2 \\ 30.8 \pm 2.4 \\ 0.27$ |
| Thalamus | Control FZP treated P value | $53.8 \pm 2.6 \\ 51.7 \pm 2.9 \\ 0.57$ | 44.2 ± 3.6 45.3 ± 3.1 0.63 | $51.8 \pm 3.5 \\ 57.1 \pm 3.3 \\ 0.26$ | 45.4 ± 6.6 44.0 ± 6.3 0.87 | 50.3 ± 4.2 44.2 ± 2.6 0.21 | 59.2 ± 6.4 58.3 ± 6.5 0.78 | 111 | 111 | 111 |
| Cerebellum Granule cells | Control FZP treated P value | 140.8 ± 4.5 132.5 ± 3.0 0.12 | 121.1 ± 4.7 115.5 ± 7.7 0.52 | 133.4 ± 6.4 124.1 ± 3.9 0.21 | 90.4 ± 119 96.8 ± 14.4 0.72 | 69.4 ± 12.8 61.1 ± 5.0 0.52 | 86.7 ± 11.0 105.6 ± 12.2 0.38 | $74.2 \pm 5.9 \\ 79.7 \pm 8.7 \\ 0.60$ | 92.6 ± 7.7 74.2 ± 4.9 0.05* | 75.2 ± 3.6 72.0 ± 8.1 0.70 |
| | | | | | | | | | | |

Mean (\pm S.E.M.) relative gray level. FZP, flurazepam; n, number of rats. *Significant difference between control and flurazepam-treated groups, $P \le 0.05$.
Note: image acquisition conditions were identical within, and not between, temporal groups; therefore, comparisons of relative gray level should be made between experimental groups, not between temporal groups.

β₃ Subunit

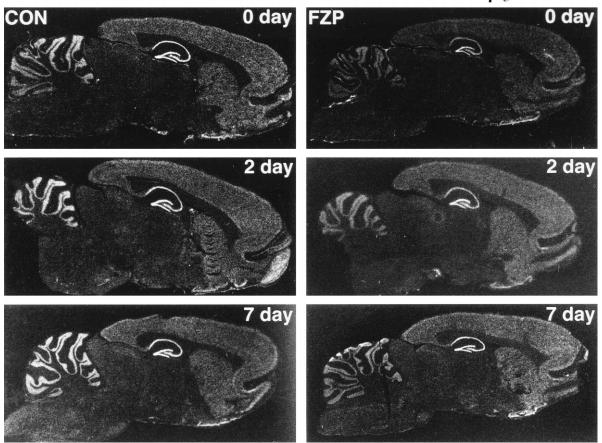


Fig. 2. Representative dark-field photomicrographs of autoradiographs of sagittal brain sections hybridized with an antisense [35 S]oligoprobe for the β 3 subunit mRNA of the GABAA receptor. Sections (10 μ m) were cut from one-week flurazepam-treated (FZP) rats killed immediately (day 0), two or seven days after ending oral drug administration and compared to sections from matched control rats (CON). Immediately after the end of treatment, there was a significant decrease in the relative gray level on autoradiographs localized over all principal cells in the hippocampus [CA1–CA3 pyramidal cells, polymorph (CA4) cells and granule cells], as well as the frontal cortex and caudate—putamen. The decreases in relative β 3 subunit mRNA levels persisted in all regions two days after treatment. Seven days after stopping drug administration, there was an increase in mRNA levels in the CA3 region and in the frontal cortex (see Table 1).

developed as described previously. 66 Sections were stained with Cresyl Echt Violet (Fluka, Switzerland) for comparison with the hybridization signal.

Image analysis. Images on film were acquired using NIH Image Software (version 1.59) with a high-resolution CCD camera (Sierra Scientific, Sunnyvale, CA) mounted over a light box (Northern Light, Imaging Research, St Catherines, Ontario, Canada), which provided constant illumination. The overall illumination was adjusted so that the distribution of relative grey values, i.e. the number of pixels in the image as a function of grey value (0–256), fell within the limits of the system, typically within 30–220 gray value units. Once established, the settings remained constant for all images acquired within a particular experimental protocol. Images, acquired as described, were digitized and stored on an external Bernoulli disk (Iomega, Roy, UT) for later analysis using a Macintosh Quadra 950 computer.

Gray level measurements (per mm²), which reflected mRNA levels, were made on digitized images by delimiting an area of interest, freehand, using predetermined criteria to

define the region or subregion. For example, to measure the density on the image over the CA1 pyramidal cell layer, a perpendicular line was drawn from the outermost edge of the dentate gyrus (DG) at the hilus and from the lateral edge of the dorsal blade. Within this defined area, the area delineated was limited to the innermost 90% of the total area to reduce the possible contribution of a signal from neighboring areas. Similar criteria were derived for each area measured. If necessary, e.g., to define cortical layers, the image was compared to an image of the Cresyl Echt Violet-stained section from the same rat. Background gray level was determined over the corpus collosum, a white matter area. All data are expressed as mean ± S.E.M. of the raw grey level values. Comparison of regional mRNA levels was made by ANOVA. Individual comparisons were made with orthogonal contrasts.

Dark-field images of silver grains detected over interneuron subclasses, defined by their location within the CA1 region [stratum oriens (SO), SO/stratum pyramidale (SP) border, SP or stratum radiatum (SR)], were used to detect the $\alpha1$ mRNA level in coronal sections exposed to NTB-2 emulsion. Images were captured with a DAGE

α_2 - γ_2 Subunits

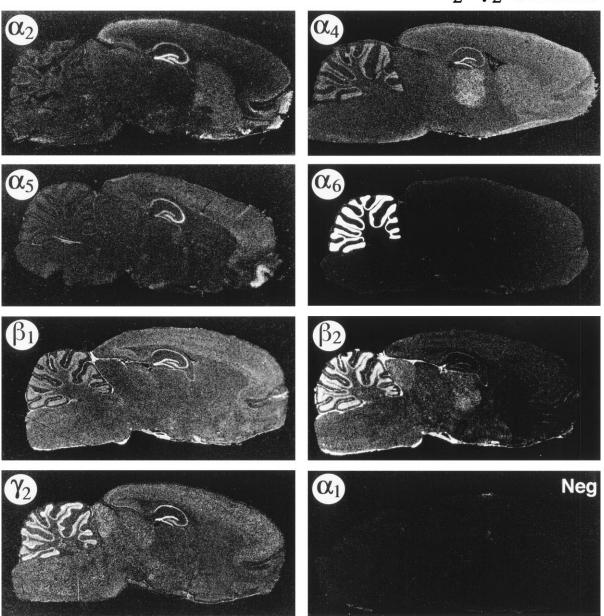


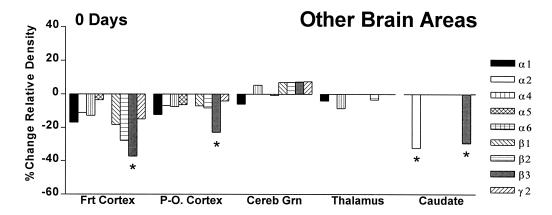
Fig. 3. Representative dark-field photomicrographs of autoradiographs of control rat sagittal brain sections hybridized with antisense [35 S]oligoprobes for the $\alpha 2$, $\alpha 4$, $\alpha 5$, $\alpha 6$, $\beta 1$, $\beta 2$ and $\gamma 2$ subunit mRNAs of the GABAA receptor. All temporal groups for each [35 S]oligoprobe were processed identically as described in the Experimental Procedures. With the exception of $\beta 2$ subunit mRNA levels, none of these GABAA receptor subunit mRNAs were regulated by one-week flurazepam treatment as indicated by a comparison of relative gray level values; therefore, additional representative autoradiographs of flurazepam-treated rat brains are not shown. There was a significant increase in the relative levels of the $\beta 2$ mRNA two, but not seven, days after the end of drug administration localized to CA1 and CA3 pyramidal cells and dentate granule cells (see Table 1). A control section hybridized with the sense [35 S]oligoprobe for the $\alpha 1$ subunit mRNA, used as the negative (NEG) for $\alpha 1$ subunit mRNA levels, is also shown.

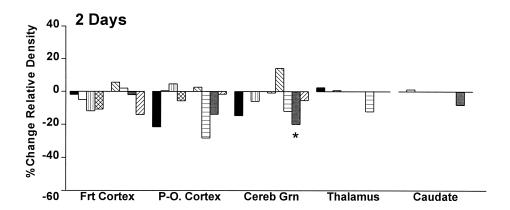
camera mounted on a Nikon Optiphot microscope (×125). Results obtained using the density threshold option of the image software were empirically determined to be equivalent to silver grain counting over bright-field images. ⁶⁶ Relative grain density was defined as the "pixel" area covered by grains as a fraction of total pixel area of the interneuron soma of interest. Background density over the corpus collo-

sum was subtracted from the total grain density to determine " $\alpha 1$ subunit-specific" grain density.

RESULTS

As illustrated in Figs 1-3, the relative distributions





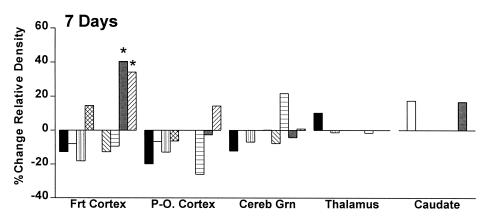


Fig. 4. Percentage change in relative density (gray level) on autoradiographs over principal hippocampal cell layers [CA1-CA3 pyramidal cells, polymorph cells (PC; formerly CA4 pyramidal cells) and DG cells]. Sections from rats killed immediately (day 0), two or seven days after ending oral flurazepam administration or their matched controls were handled identically and were batch processed according to the in situ hybridization procedures decribed in the Experimental Procedures. Flurazepam-treated and their matched control sections from each temporal group were exposed to the same film. Digitized images over autoradiographs were acquired with NIH image software, maintaining the same illumination intensity across all sections on each film. Statistical analyses were performed on raw gray level measurements (see Table 1). A significant decrease in α1 subunit mRNA levels, reflected in the percentage change in relative density immediately (day 0) after treatment, was localized to CA1 pyramidal cells and DG cells. The decrease only persisted in CA1 neurons two days after ending flurazepam treatment. A reduction in \(\beta \) subunit mRNA level, reflected in the change in relative density, was present in all principal hippocampal cells, immediately and two days after treatment. In contrast to the decreased β3 mRNA level, a significant increase in β2 subunit mRNA level was found in CA1, CA3 and granule cells two days after treatment. This trend persisted seven days after treatment. Whereas the γ 2 subunit mRNA level was not significantly down-regulated immediately after stopping treatment, a similar trend toward an increase was seen two and seven days after flurazepam treatment. This biphasic trend was noted across all subunits, with the \(\beta \) subunit being the most notable exception.

| Table 2 of Su | bunit messenger | RNA level i | n interneurons | in the CA | 1 region |
|---------------|-----------------|-------------|----------------|-----------|----------|
| | | | | | |

| | | CA1 layer | | | | |
|------------------------|--|---|--|----------------------------------|--|--|
| | SO | SO/SP border | SP | SR | | |
| Control FZP treated | $100.6 \pm 3.1 (17)$ $100.9 \pm 3.2 (17)$ | $115.5 \pm 2.9 (15)$ $101.0 \pm 3.4 (7)$ | $109.2 \pm 2.2 (44)$ $108.7 \pm 2.0 (49)$ | 94.4 ± 7.8 (5) 95.7 ± 3.7 (7) | | |
| P value | 0.94 | 0.01* | 0.87 | 0.87 | | |

Mean (± S.E.M.) relative silver grain density. Number of cells/group is given in parentheses. FZP, flurazepam.

of the nine GABA_A receptor subunit mRNAs evaluated were similar to those described by others. 49,52,75 The antisense [35 S]oligoprobe signal was specific to neurons and was very low to absent over white matter regions. Background labeling was uniform over RNase-treated sections and low over sections treated with sense strands for the $\alpha 1$ (Fig. 3) or $\beta 3$ oligoprobes, as reported previously for each of the same oligoprobes. 75 All statistical analyses were made using raw gray level values (see Table 1). For comparison of the changes in the mRNA levels among the nine GABA_A receptor subunits, the data in the text are expressed as percentages of the control mRNA level.

α1 Subunit messenger RNA level

Dark-field photomicrographs of $\alpha 1$ subunit mRNA distribution in parasagittal brain sections are shown in Fig. 1. As described previously, the relative distribution of the $\alpha 1$ subunit mRNA signal in rat hippocampal neurons was CA1 > DG \geq CA3 > polymorph cells (formerly CA4). The $\alpha 1$ mRNA level was similar in the frontal, striatal and parietal cortices, and the thalamus. The $\alpha 1$ mRNA signal over the cerebellar granule cell layer was two-fold greater than that over hippocampal granule cells.

The $\alpha 1$ subunit mRNA level was significantly decreased in the CA1 pyramidal cells (- 14.4%) and DG cells (- 19.5%) immediately (day 0) after the end of treatment (Table 1, Fig. 4). The significant decrease in $\alpha 1$ mRNA level persisted only in CA1 pyramidal cells two days after ending flurazepam treatment (- 11%) and was reversed seven days after ending treatment (- 5.8%). The decreased mRNA level in cortical areas did not reach significance at any time-point. There were no significant changes detected in $\alpha 1$ mRNA levels in other brain areas, i.e. the cerebellum and thalamus (Table 1).

In emulsion-coated sections, very densely labeled $\alpha 1$ -positive interneurons were scattered throughout the neuropil and within and bordering the pyramidal cell layer. 22,34,66 Interneurons were classified according to their location within the hippocampal formation, i.e. SO, SO/SP border, SP and SR. Dark-field image analysis of silver grains over individual interneurons indicated a selective decrease in $\alpha 1$ mRNA

level only in interneurons located at the SO/SP border (Table 2).

α2 Subunit messenger RNA level

The $\alpha 2$ subunit mRNA signal was moderate over CA1 pyramidal cells in comparison to CA3 pyramidal cells, polymorph cells and DG cells, the latter being of a relatively high density (Fig. 3). The level of $\alpha 2$ mRNA was relatively low in cortical areas and in the caudate–putamen, and negligible or absent in other brain areas in the parasagittal plane, i.e. the cerebellum, thalamus and collicular regions. The relatively large significant decrease in the $\alpha 2$ mRNA level in the caudate–putamen immediately after ending flurazepam treatment (-32%) was reversed both two (+1.3%) and seven days (+17.4%) after ending treatment. No significant changes in $\alpha 2$ mRNA levels were found in any other brain areas at any time-point (Fig. 5).

α4 Subunit messenger RNA level

The pattern of the $\alpha 4$ mRNA signal in the hippocampus was relatively lower and more uniform in all pyramidal cell layers (Fig. 3) in comparison to that in DG cells, where it was twofold greater. $\alpha 4$ mRNA levels were relatively low in the cortex, cerebellar granule cells and thalamus. Chronic flurazepam treatment did not alter levels of $\alpha 4$ mRNA in any brain region at any of the time-points measured (Figs 4, 5).

α5 Subunit messenger RNA level

At the brain level examined, the $\alpha 5$ mRNA signal was relatively exclusive to the hippocampus and cortex (Fig. 3). The patterns of the $\alpha 5$ mRNA signal generated over hippocampal cells were similar to those described previously in emulsion-coated brain sections: CA3 > polymorph (CA4) cells > CA2 = CA1 \gg DG.⁶⁶ The relative density in the cortex was similar to that in the dentate. As with the $\alpha 4$ subunit mRNA, there were no significant difference in $\alpha 5$ subunit mRNA levels post-flurazepam treatment (Figs 4, 5). However, there was a non-significant trend toward increased levels in DG cells (Fig. 4) and the frontal cortex (Fig. 5).

^{*}Significant difference between control and flurazepam-treated groups, $P \le 0.05$.

α6 Subunit messenger RNA level

As anticipated, the $\alpha6$ subunit mRNA signal was intense and present exclusively in cerebellar granule cells (Fig. 3). Chronic flurazepam treatment had no effect on the level of $\alpha6$ mRNA (Figs 4, 5).

β1 Subunit messenger RNA level

The level of $\beta 1$ subunit mRNA was high in the hippocampus, with lower levels measurable in the cortex and in cerebellar granule cells (Fig. 3). The CA2 and the dentate region had very high levels of $\beta 1$ subunit mRNA, CA2 pyramidal cells slightly higher than granule cells. Relatively high levels of $\beta 1$ mRNA were also detected in CA1 and CA3 neurons, as well as polymorph cells. Chronic flurazepam treatment had no effect in regulating $\beta 1$ subunit mRNA levels (Figs 4, 5).

β2 Subunit messenger RNA level

β2 Subunit mRNA levels were relatively low in the hippocampus (CA1 > CA2 = DG = CA3 =polymorph cells) and in the cortex, being most prominent in cerebellar granule cells (Fig. 3). The levels in the thalamus were somewhat higher than those in the CA1 region (Table 1). Though there was no effect of chronic flurazepam treatment immediately after the end of flurazepam treatment, \(\beta 2\) mRNA level was significantly up-regulated in CA1 (+31.2%) and CA3 (+37.2%) pyramidal cells and in DG cells (+ 29.2%) two days after ending treatment (Fig. 4). In the cortex, on the other hand, there was a non-significant trend toward a decrease in mRNA level immediately and two days after ending flurazepam treatment in the frontal and parieto-occipital cortices, respectively (Fig. 5, Table 1). There was no evidence of β2 subunit mRNA regulation in the other brain areas.

β3 Subunit messenger RNA level

The relative levels of $\beta 3$ subunit mRNA in the hippocampus were as follows: DG \gg polymorph cells > CA1 = CA3 > CA2 (Table 1, Fig. 2). The β3 mRNA level was relatively low in cortical areas and slightly lower in the caudate-putamen. Chronic flurazepam treatment induced time-dependent decreases in β 3 subunit mRNA expression in several brain areas. Immediately after oral flurazepam administration was stopped, there were significant reductions in all pyramidal cell types (CA1, -24.0%; CA2, -14.8%; CA3, -18.4%) and in DG cells (-21.2%) (Fig. 4, Table 1). There was a large, significant reduction in β3 mRNA level in the frontal cortex (-37.3%), which did not reach significance in the parieto-occipital cortex (-22.8%). A significant reduction in β 3 mRNA level (-29.2%) was also detected in the caudate-putamen, but not in the cerebellum (Fig. 5, Table 1). Two days after

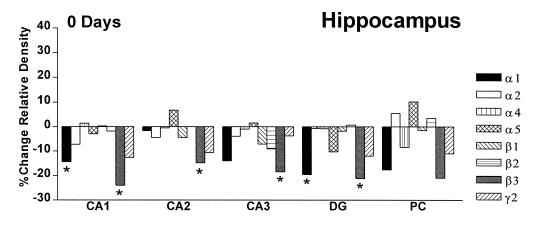
flurazepam was withdrawn, the reductions in $\beta3$ mRNA level persisted in the hippocampus. Seven days after ending flurazepam administration, several brain areas showed significant increases in $\beta3$ subunit mRNA levels, notably CA3 pyramidal cells (+17.8%) and frontal cortical cells (+34.2%). A similar trend was seen in other hippocampal cell groups and in the caudate–putamen, which did not reach significance (Fig. 5, Table 1).

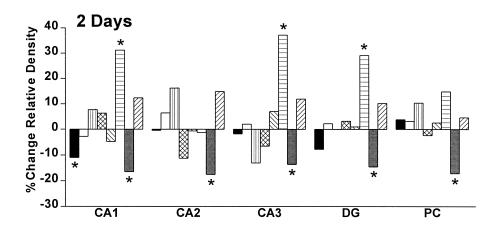
γ2 Subunit messenger RNA level

The level of $\gamma 2$ subunit mRNA was relatively uniform across the hippocampus, being somewhat higher in DG cells (Fig. 3). The level of $\gamma 2$ subunit mRNA in cerebellar granule cells was comparable to that in hippocampal granule cells. The levels in the cortex were approximately 50% of those in the hippocampus. One-week flurazepam treatment had no measurable effect on $\gamma 2$ mRNA expression in any brain area evaluated. None the less, as with the $\beta 3$ mRNA expression, there was a trend toward increased expression seven days after flurazepam was withdrawn, which was significant in the frontal cortex (+34.2%; Fig. 5).

DISCUSSION

The hypothesis that a switch in subunit composition of the GABAA receptor may underlie benzodiazepine tolerance has received support from numerous studies. Nevertheless, reports have been contradictory, and with the exception of the α 2 and γ1 mRNAs, changes in the levels of expression of each of the known α , β and γ subunit mRNAs have been detected following various in vivo benzodiazepine treatment regimens which result in anticonvulsant tolerance in rodents. For example, α1 subunit mRNA expression was decreased in the cerebral cortex^{26,27,29,31,33,66,76} and hippocampus^{29,31,66,76} following protracted benzodiazepine administration studies, and was unchanged some others. 26,33,46,85,86 Levels of $\gamma2$ or $\beta3$ subunit mRNA were down-regulated 33,55,86 expression unchanged.46 In contrast, other investigators reported up-regulation of subunit mRNAs. 27,31,47 One factor complicating the interpretation of findings is the inability to separate the molecular changes related to benzodiazepine tolerance from those associated with dependence. This divergence of findings may also be due to the benzodiazepine selected for chronic treatment, the dose, frequency and method of drug administration, and the time of analysis post-treatment. Moreover, changes in mRNA and protein expression following long-term benzodiazepine administration have been discretely localized^{31,66} and may have gone undetected in previous northern blot or solution hybridization studies. ^{26–28,31,33,46,47,55,56,76,85} By focusing studies in the hippocampus, some of these problems can be





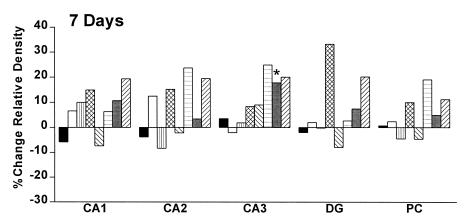


Fig. 5. Percentage change in relative density (gray level) on autoradiographs over brain areas with significant GABA_A receptor subunit mRNA level (frontal cortex, parieto-occipital cortex, cerebellar granule cells, thalamus and caudate-putamen). Sections from rats killed immediately (day 0), two or seven days after ending oral flurazepam administration or their matched controls were handled identically and were batch processed according to the in situ hybridization procedures described in Experimental Procedures. Flurazepam-treated and their matching control sections from each temporal group were exposed to the same film. Digitized images over autoradiographs were acquired with NIH Image software, maintaining the same illumination intensity across all sections on each film. Statistical analyses were performed on raw gray level measurements (see Table 1). A significant decrease in β3 subunit mRNA level, reflected in the percentage change in relative gray level immediately after treatment, was localized to the frontal cortex. This decrease did not persist two days after ending treatment. There was a rebound increase in $\beta 3$ subunit mRNA level seven days after ending flurazepam treatment. β3 Subunit mRNA level was also regulated biphasically in the caudate-putamen, but the rebound seven days after ending drug administration was not significant. Though $\gamma 2$ subunit mRNA level was not significantly decreased immediately or two days after ending treatment, a significant increase in $\gamma 2$ subunit mRNA level paralleled that of the \$\beta\$3 subunit mRNA in the frontal cortex. No other subunit mRNAs showed significant changes in their levels in any brain area measured.

circumvented, and there is an opportunity to relate changes in subunit mRNA expression to changes in GABA_A receptor function in a well-defined neuronal population.

Relationship between GABA_A receptor subunit messenger RNA expression and in vivo tolerance

The present in situ hybridization study was carried out on rats made tolerant to benzodiazepines in vivo, but not dependent. In our hippocampal model system, this treatment was also found to result in tolerance *in vitro*^{78,83} and to significantly impair CA1 pyramidal cell GABAergic function. 77,78,80–84 Changes in the levels of $\alpha 1$, $\beta 2$ and $\beta 3$ subunit mRNAs were evident when tolerance was present (day 0 and two days), but had reversed when tolerance was absent (seven days). The discrete reduction in the level of α1 subunit mRNA in CA1 pyramidal cells two days after ending flurazepam treatment replicated our previous findings using silver grain analysis. 66 However, the time-course of the change in α1 subunit mRNA expression does not parallel that of benzodiazepine receptor down-regulation. In autoradiographic binding studies, significant, localized reductions in [3H]flunitrazepam10,67,68 and [3H]zolpidem binding (unpublished observation) in the CA1 region of the hippocampus were transient, i.e. had reversed by two days after drug removal, though rats were still tolerant in vivo and in vitro. Conversely, GABA/benzodiazepine binding sites in the CA1 region remained "uncoupled" until tolerance was reversed. 10 Regulation of the α1 subunit has been implicated in mediating both receptor down-regulation and decreased allosteric coupling between GABA and benzodiazepine binding sites, yet its role is unresolved. 7,36,37,56,65 The benzodiazepine binding pocket is probably located at the interface of the α and γ subunits, 64 and recent studies using α/γ chimeras have also implicated two different domains on the γ subunit in regulating high-affinity benzodiazepine binding and allosteric coupling, respectively. 6 Other subunit mRNAs (α3, $\alpha 4$, $\alpha 5$, $\beta 1$), including the $\gamma 2$ subunit, have been shown to be regulated with acute, more protracted or more intense flurazepam treatments, 46,47,85,86 or following diazepam^{27,31} or lorazepam³³ administration. Changes in $\gamma 2$ subunit mRNA expression have been more closely linked with changes in benzodiazepine binding density. 85 None the less, changes in the mRNA expression of these subunits was not required to demonstrate anticonvulsant tolerance in in vivo 59,60 or in in vitro hippocampus 32,78,83 following flurazepam administration. In addition, unlike one-week oral flurazepam administration, other benzodiazepine treatment protocols which result in benzodiazepine tolerance and in a broader range of changes in GABA_A receptor subunit mRNA expression may also be associated with benzodiazepine dependence. 31,73

Studies of benzodiazepine site ligands which do not induce tolerance^{28,31} may help clarify the conflicting findings of studies of mRNA expression following various chronic benzodiazepine treatments, since drugs which have limited toleranceinducing potential, such as abercarnil²⁷ and imidazenil,31 did not induce similar changes in mRNA expression. Chronic treatment with the α 1-selective imidazopyridine, zolpidem, which also has a low tolerance liability, predictably down-regulated α1 subunit mRNA expression. However, since zolpidem has a negligible affinity for the $\alpha 4$ subunit, the up-regulation of $\alpha 4$ and $\beta 1$ mRNA expression was unanticipated.²⁸ Holt et al.^{27,28} proposed that some GABAA receptor subunit changes found following allosteric modulator treatments are the result of direct drug-receptor interactions and that changes in GABA tone may also affect steady-state levels of mRNA as the result of coordinated translation of GABAA subunit genes, found to be organized in clusters on human chromosomes (4p13– 4q11: α 2, α 4, β 1 and γ 1; 5q32–5q33: α 1, α 6, β 2 and γ 2; 15q11–15q13: α 5, β 3, γ 3).

Rebound overexpression of GABA_A receptor subunit messenger RNAs

The lack of a significant change in the expression of the α5 subunit mRNA in the hippocampus and cortex, 66 at a time when rats are tolerant, replicates our previous findings. Expression of $\alpha 5$ mRNA, among other GABAA receptor subunit mRNAs, e.g., α 3 and α 4, was increased in whole brain or cortex following various chronic benzodiazepine treatments. 27,31,47 Conversely, a decrease in γ 2 mRNA expression was reported following the same or other protocols.^{31,85} Nevertheless, such subunit mRNA changes were not a prerequisite for in vivo or in vitro tolerance development during oneweek flurazepam administration. Alternatively, the increase in \(\beta \) and \(\gamma \) mRNA in the frontal cortex and the trend toward the increased $\alpha 5$, $\beta 3$ and $\gamma 2$ mRNA expression in CA1, CA3 and DG cells, at a time when tolerance is no longer present in vivo (seven days), suggests that an $\alpha 5\beta 3\gamma 2$ receptor, a preferred hippocampal GABAA receptor subtype, (cf. Refs 41 and 44) might subsequently be overexpressed. Moreover, since dependence does not occur following one-week flurazepam treatment, the up-regulation of the β 3 and γ 2 subunit mRNAs in the frontal cortex likely represents a rebound, rather than a withdrawal phenomenon. A similar trend, increased expression seven days after ending flurazepam treatment, was shown for the majority of GABA_A receptor subunit mRNAs examined. A role for biphasic switches in mRNA expression has not been established. Switches in the expression of certain subunit combinations occur during development, 19 perhaps reflecting both GABA's early neurotrophic role⁶⁹ and its subsequent role as an

inhibitory neurotransmitter in the hippocampus.⁸ For example, in specific brain areas, $\alpha 3$, $\alpha 5$, $\beta 3$ and $\gamma 2$ subunit mRNAs peak during embryonic and early postnatal development, then rapidly decline. Thus, some of the switches in subunit subtypes detected following chronic benzodiazepine treatment may represent a coordinated control of the translation of specific GABA_A receptor subtypes, analogous to the proposal of Holt *et al.*,^{27,28} and may not be directly related to tolerance development.

Relationship between changes in subunit messenger RNA and protein levels

The localized decreases in the steady-state levels of $\alpha 1$ and $\beta 3$ subunit mRNAs in the cortex and hippocampus of flurazepam-treated rats immediately after the drug was withdrawn was reflected in discrete, parallel changes in $\alpha 1$ and $\beta 3$, but not $\alpha 2$, $\beta 1$ or $\gamma 2$, subunit protein in the same brain **Ouantitative** immunohistochemical studies indicated a significant reduction in $\alpha 1$ subunit antibody staining density in the dendritic areas of CA1, CA3 and DG cells at the same time-point. β3 subunit antibody staining density was similarly decreased, but was more widespread in dentate areas. A comparison of the time-course of GABAA receptor subunit protein expression relative to mRNA expression will be needed to establish whether changes in protein expression persist at a time when tolerance is still detected in the hippocampus in vitro. 78,83 Preliminary experiments have suggested that the significant decrease in $\alpha 1$ subunit protein persists in the SO of the CA1 region, e.g., two days after ending flurazepam treatment (unpublished observations). Establishing a correlation between mRNA and protein levels is particularly important, since it is unknown whether changes in mRNA expression signal subsequent changes in protein translation or were regulated in a feedback manner. For example, exposure of chick cortical neurons to GABA resulted in a delayed decline in α1 subunit mRNA⁴ and α1 subunit translation.⁴³ Whether a similar repression of mRNA expression occurs following benzodiazepine exposure is not yet known.65 The strong correlation between changes in the levels of specific subunit mRNAs and subunit proteins in CA1 pyramidal cells suggests that the decreased levels of mRNA expression may represent a switch in GABA_A receptor subunit composition and may be an important mechanism contributing to functional changes at the GABAergic postsynapse on CA1 pyramidal cells associated with benzodiazepine tolerance. Nevertheless, though the findings suggest that the β 2 subunit could switch with the β 3 subunit, no concomitant increases in α subunit mRNAs or proteins have been identified to suggest which α subunit might substitute for a decreased $\alpha 1$ subunit expression. Whether post-translational mechanisms may also relate to the decreased protein expression^{36,37,56,65} or additionally contribute to changes in postsynaptic GABA function also requires further investigation.

Relationship between changes in GABA_A receptor subunit messenger RNA expression and CA1 pyramidal cell function

The role of a switch in GABA_A receptor subunits in mediating changes in GABAergic function is still uncertain. Decreased al subunit mRNA and protein expression in CA1 pyramidal cells⁶⁶ was associated with functional changes in these neurons following long-term benzodiazepine treatment.^{77,80–82} The decreased potency of zolpidem to potentiate GABA currents in dissociated CA1 pyramidal cells of chronic diazepam-treated rats³² and the reduced ability of zolpidem to potentiate the decay of CA1 pyramidal cell miniature inhibitory postsynaptic currents⁸³ suggests that the function of α1-containing GABA_A receptor subtypes may be selectively impaired in the hippocampus of flurazepam-tolerant rats. The increased levels of $\beta2$ subunit mRNA in CA1 pyramidal cells two days after flurazepam administration was stopped, together with the decreased levels of \$3 subunit mRNA and protein expression, 11 could account for the 2.5-fold decrease in GABA_A agonist potency in the CA1 region of the hippocampus of flurazepam-tolerant rats.⁷⁸ One study indicated a similar 2.5-fold increase in GABA affinity in β3 vs β1 or β2 subunit-containing recombinant receptors, 15 though another report indicated no effect of B subunit substitution.²⁴ Conversely, a switch in the level of β2 and β3 mRNA subunits was also found in DG cells after chronic flurazepam treatment, yet in comparison to CA1 pyramidal neurons, this hippocampal cell group does not show a similar impairment in GABAergic function, i.e. a profound decrease in miniature inhibitory postsynaptic current amplitude or a reduction in unitary C1⁻ channel conductance. 53,81 Reports have suggested that the β subunit of the GABA_A receptor may have an important function to target receptors to the subsynaptic membrane and may play a role in regulating receptor localization or clustering during development or associated with plastic changes at GABAergic synapses. 12,51 The possible functional role of a switch in β subunits in benzodiazepine-tolerant rats remains uncertain.

Regulation of $GABA_A$ receptor subunit messenger RNA in hippocampal interneurons

Whole-cell electrophysiological studies in CA1 pyramidal cells in flurazepam-treated hippocampus also showed that spontaneous, but not miniature inhibitory postsynaptic current, frequency was reduced two days after the drug was withdrawn. These findings suggest that a presynaptic mechanism, a change in interneuron excitability, rather than a

change in GABA release mediated at GABAergic terminals, may be involved in the alteration in GABA function in tolerant rats.⁸¹ The prominent localization of the $\alpha 1$ subunit protein on certain GABAergic interneuron populations^{22,45} and its regulation in other principal hippocampal cell populations (Fig. 1, Table 1)66 prompted the limited evaluation of α1 subunit mRNA levels over anatomically-defined interneuron subgroups. The reduction in α1 subunit mRNA levels in cells located at the SO/SP border (Table 2) is the first demonstration of the regulation of GABA_A receptor mRNAs on presumptive interneurons, and warrants the further study of electrophysiologically, immunocytochemically and morphologically identified interneuron subpopulations 16,22,25,45 in the hippocampal CA1 region of chronic benzodiazepinetreated rats. Study of the regulation of other subunit mRNAs, e.g., the β2 subunit, on hippocampal interneurons and of the functional consequences of the regulation of mRNA expression on interneurons will also be important.

CONCLUSIONS

The levels of three (α 1, β 2 and β 3) of nine GABA_A receptor subunits were regulated as a function of time after ending one-week flurazepam treatment, a chronic benzodiazepine treatment which induces anticonvulsant tolerance but does not result

in physical dependence. The same chronic benzodiazepine treatment was shown in previous electrophysiological studies to have profound consequences on hippocampal function and to induce benzodiazepine tolerance in the hippocampus in vitro. The findings of in situ hybridization studies suggested that changes in the levels of GABAA receptor mRNAs are discretely regulated in specific cortical regions, and in particular in hippocampal CA1 pyramidal neurons, with a pattern consistent with the appearance and disappearance of benzodiazepine anticonvulsant tolerance in vivo and in vitro. Exploratory studies suggested that postsynaptic GABAA receptors on selected interneuron populations were also regulated. Additional evidence has suggested that changes in the expression of specific GABA_A receptor subunit mRNAs may mirror changes in the translation of their respective subunit proteins. Taken together, these findings suggest that regulation of α1, β2 and β3 subunits on GABA_A receptors, in particular in the CA1 region of the hippocampus, may be a minimal requirement for initiating benzodiazepine anticonvulsant tolerance.

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