

Expression and distribution of metabotropic GABA receptor subtypes GABA_BR1 and GABA_BR2 during rat neocortical development

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Abstract

To understand the possible contribution of metabotropic γ -aminobutyric acid receptors (GABA_BR) in cortical development, we investigated the expression pattern and the cellular and subcellular localization of the GABA_BR1 and GABA_BR2 subtypes in the rat neocortex from embryonic day 14 (E14) to adulthood. At the light microscopic level, both GABA_BR1 and GABA_BR2 were detected as early as E14. During prenatal development, both subtypes were expressed highly in the cortical plate. Using double immunofluorescence, GABA_BR1 colocalized with GABA_BR2 in neurons of the marginal zone and subplate, indicating that these proteins are coexpressed and could be forming functional GABA_BRs during prenatal development *in vivo*. In contrast, only GABA_BR1 but not GABA_BR2 was detected in the tangentially migratory cells in the lower intermediate zone. During postnatal development, immunoreactivity for GABA_BR1 and GABA_BR2 was distributed mainly in pyramidal cells. Discrete GABA_BR1-immunopositive cell bodies of interneurons were present throughout the neocortex. In addition, GABA_BR1 but not GABA_BR2 was found in identified Cajal-Retzius cells in layer I. At the electron microscopic level, immunoreactivity for GABA_BR1 and GABA_BR2 was found in dendritic spines and dendritic shafts at extrasynaptic and perisynaptic sites throughout postnatal development. We further demonstrated the presynaptic localization of GABA_BR1 and GABA_BR2, as well as the association of the receptors with asymmetrical synaptic junctions. These results indicate potentially important roles for the GABA_BRs in the regulation of migratory processes during corticogenesis and in the modulation of synaptic transmission during early development of cortical circuitry.

Introduction

γ -Aminobutyric acid (GABA) is the main inhibitory neurotransmitter in the cerebral cortex. The response to GABA is mediated via activation of both ionotropic and metabotropic receptors (Macdonald & Olsen, 1994; Misgeld *et al.*, 1995). Metabotropic GABA receptors (GABA_BRs) are coupled to intracellular signal transduction mechanisms via G-proteins and mediate slow inhibitory postsynaptic potentials (Misgeld *et al.*, 1995). Activation of GABA_BRs can modulate multiple signal transduction pathways, mainly activating K⁺ channels, inhibiting voltage-dependent Ca²⁺ channels, stimulating phospholipase A2, inhibiting adenylate cyclase and regulating inositol phospholipid hydrolysis (Misgeld *et al.*, 1995).

Two different cDNAs encoding GABA_BRs have been cloned to date: GABA_BR1, which exist in alternatively spliced forms designated R1a–e, and GABA_BR2 (Bowery & Brown, 1997; Kaupmann *et al.*, 1997, 1998a; Isomoto *et al.*, 1998; Pfaff *et al.*, 1999; Schwarz

et al., 2000). It has been suggested that heterodimeric assembly of GABA_BR1a/b and GABA_BR2 subtypes is required to provide fully functional receptors (Jones *et al.*, 1998; Kaupmann *et al.*, 1998a; White *et al.*, 1998; Kuner *et al.*, 1999).

Light microscopic immunocytochemistry, *in situ* hybridization and receptor autoradiographic studies have shown that GABA_BRs are abundant in the cerebral cortex of adult animals (Bowery *et al.*, 1987; Chu *et al.*, 1990; Turgeon & Albin, 1994; Kaupmann *et al.*, 1997; Bischoff *et al.*, 1999; Margeta-Mitrovic *et al.*, 1999; Muñoz *et al.*, 2001). Electron microscopic studies revealed that GABA_BRs are present in both excitatory and inhibitory synapses and also located outside the synaptic membrane specialization, at both postsynaptic and presynaptic locations (Gonchar *et al.*, 2001; Kulik *et al.*, 2002). Electrophysiological and pharmacological studies have also characterized pre- and postsynaptic inhibitory functions of GABA_BRs in the neocortex (Howe *et al.*, 1987; Deisz *et al.*, 1993, 1997; Varela *et al.*, 1997). Presynaptic GABA_BRs are thought to suppress neurotransmitter release by depressing Ca²⁺ influx via voltage-activated Ca²⁺ channels, whereas postsynaptic GABA_BRs decrease neuronal excitability by activating an inwardly rectifying K⁺ (Kir3) conductance (Kaupmann *et al.*, 1998b).

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Little information is available about the expression of GABA_BRs in the developing neocortex. Nevertheless, several lines of evidence suggest that GABA_BR play important roles during cortical development. First, GABA_BRs modulate the excitatory actions of glutamate in developing cortical neurons through the regulation of glutamate-activated calcium transients (Obrietan & van den Pol, 1999) and second, specific GABA_BR agonists stimulated migration of neurons (Behar *et al.*, 1998), whereas specific antagonists inhibited migration (Behar *et al.*, 2000).

Information regarding the distribution of GABA_BRs in the cerebral cortex is a prerequisite for studies designed to elucidate the contribution of GABA_BRs to developmental processes and to cortical function. However, the distribution and localization of GABA_BR subtypes during corticogenesis are incompletely characterized. To better understand the mechanisms by which GABA_BRs mediate the physiological effects of GABA on developing cortical neurons, the goal of the present study was to characterize the expression pattern, and the cellular and the precise subcellular localization of GABA_BR1 and GABA_BR2 in the rat neocortex during pre- and postnatal development.

Materials and methods

Thirty-nine Wistar rats from embryonic day 14 (E14) to adulthood, obtained from the Animal House Facilities of the University Miguel Hernández in Alicante and the Department of Human Anatomy and Genetics in Oxford were used in the present study. The care and handling of the animals prior to and during the experimental procedures followed European Union and UK Home Office regulations, and were approved by the Animal Care and Use Committees of the authors' institutions.

For each developmental stage, the animals used ($n = 3$ for each stage) were from different litters and were grouped as follows: E14, E16, E18, postnatal day 0 (P0, day of birth), P3, P5, P7, P10, P12, P15, P21, P30 and P60.

Antibodies and controls

Affinity-purified polyclonal antibodies against GABA_BR1a/b and GABA_BR2 were raised in rabbits and guinea pig, respectively. The characteristics and specificity of the antibody against GABA_BR1/b have been described elsewhere (Kulik *et al.*, 2002). The antibody against GABA_BR2 was obtained from Chemicon (Temecula, CA, USA). To identify neuronal populations during cortical development, we used a monoclonal antibody, G10, against Reelin (De Bergeyck *et al.*, 1998) gift of Dr A.M. Goffinet (Neurobiology Unit, University of Namur School of Medicine, Belgium), a monoclonal antibody, 12E3, against PSA-NCAM protein (the polysialylated form of the neuronal cell adhesion molecule NCAM; Seki and Arai, 1991), gift of Dr Tatsunori Seki (Department of Anatomy, Juntendo University School of Medicine, Tokyo, Japan), and a monoclonal antibody against calbindin (Chemicon, Temecula, CA, USA).

To test method specificity in the procedures for light and electron microscopy, the primary antibody was omitted or replaced with 5% (v/v) normal serum of the species of the primary antibody. Under these conditions, no selective labelling was observed. Staining patterns were also compared to those obtained by calretinin and calbindin (Chemicon, Temecula, CA, USA); only the antibodies against GABA_BR1 and GABA_BR2 labelled consistently the plasma membrane of cells. When double labelling was used, some sections were always incubated with only one primary antibody and the full complement of secondary antibodies to test for any cross-reactivity.

Other sections were incubated with two primary antibodies and one secondary antibody, followed by the full sequence of signal detection. No cross-labelling was detected that would influence the results.

Immunocytochemistry for metabotropic GABA receptors

Light microscopic procedures

Fetuses were collected by caesarean section after anaesthesia of the dam with an intraperitoneal (i.p.) injection of a Rompun–Imalgene mixture (0.1 mL/kg body weight). Animals were deeply anaesthetized by hypothermia (E14 to P5) or by i.p. injection of Rompun–Imalgene (1 : 1, 0.1 mL/kg body weight) and then the hearts were exposed surgically for perfusion fixation (López-Bendito *et al.*, 2001, 2002). After perfusion, tissue blocks containing the neocortex were dissected and washed thoroughly in 0.1 M phosphate buffer (PB, pH 7.4). Coronal, 60 µm sections were cut with a Vibratome and collected in 0.1 M PB. For the immunoperoxidase method, sections were incubated in 10% normal goat serum (NGS) in 50 mM Tris buffer (pH 7.4) containing 0.9% NaCl (TBS), with 0.2% Triton X-100, for 1 hour. Sections were then incubated for 48 hours with affinity-purified polyclonal antibodies against GABA_BR1a/b or GABA_BR2 at a final protein concentration of 1–2 µg/mL in TBS containing 1% NGS. After washes in TBS, the sections were incubated for 2 hours in biotinylated goat anti-rabbit IgG (Vector Laboratories, Burlingame, CA, USA) diluted 1 : 50 in TBS containing 1% NGS. They were then transferred into avidin–biotin–peroxidase complex (ABC kit, Vector Laboratories, Burlingame, CA, USA) diluted 1 : 100 and left for 2 hours at room temperature. Peroxidase enzyme activity was revealed using 3,3'-diaminobenzidine tetrahydrochloride (DAB; 0.05% in TB, pH 7.4) as the chromogen and 0.01% H₂O₂ as substrate. Finally, the sections were air-dried and coverslipped prior to observation with a Leica DMRS photomicroscope equipped with differential interference contrast optics.

Alternate sections were used for immunofluorescence experiments using a laser scanning confocal microscope (Leica TSLCM, Austria). Primary antibodies used in combination with a polyclonal antibody against GABA_BR1 were: guinea pig polyclonal anti-GABA_BR2; monoclonal anti-PSA-NCAM, diluted 1 : 1000; monoclonal anti-Reelin antibody, G10, diluted 1 : 1000 and monoclonal anti-calbindin antibody diluted 1 : 1000. Sections were incubated overnight at 4 °C in a mixture of primary antibodies in TBS containing 1% NGS and 0.1% Triton X-100. Sections were then washed in TBS and incubated for two hours in a mixture of secondary antibodies coupled to the cyanine-derived fluorochromes Cy2 or Cy3 (Amersham, UK). These secondary antibodies were used at a dilution of 1 : 200 in TBS with 0.1% Triton X-100. After extensive TBS washes, the sections were mounted on gelatinized slides and coverslipped with Vectashield (Vector Laboratories, Burlingame, CA, USA).

Each labelling combination was analysed in sections from two to four prenatal or postnatal animals. Co-localization of GABA_BR1 with GABA_BR2 or with the neurochemical markers Reelin and calbindin was assessed by confocal laser microscopy (Leica TLSCM, Austria). For each marker, a semiquantitative estimation of the proportion of double labelled neurons was performed by analysing samples of 120–160 cells, collected in 8–10 coronal sections through the neocortex. Images were stored digitally and analysed using appropriate software supplied by the microscope manufacturer. Brightness and contrast were adjusted for the whole frame; no part of a frame was enhanced or modified in any way.

Electron microscopic procedures

A similar procedure to that described earlier (López-Bendito *et al.*, 2001, 2002) was used.

Pre-embedding immunoperoxidase and immunogold methods. Immunocytochemical reactions for the pre-embedding method were carried out as described above, but with the omission of Triton X-100 in all steps. Sections for the pre-embedding immunogold method were incubated together with those used for peroxidase reaction up to the secondary antibody stage. They were then incubated in goat anti-rabbit or goat anti-guinea pig IgG coupled to 1.4 nm gold particles (Nanoprobes Inc., Stony Brook, NY) diluted 1 : 100 in TBS containing 1% NGS. After several washes in phosphate-buffered saline (PBS), the sections were postfixed in 1% glutaraldehyde dissolved in the same buffer. They were washed in double distilled water, followed by silver enhancement of the gold particles with an HQ Silver kit (Nanoprobes Inc., Stony Brook, NY).

The peroxidase-reacted sections and the gold–silver-labelled sections were treated with OsO₄ (1% in 0.1 M PB), block-stained with uranyl acetate, dehydrated in graded series of ethanol and flat-embedded on glass slides in Durcupan (Fluka, Reidel-deHaën, UK) resin. Regions of interest were cut at 70–90-nm-thick sections using an ultramicrotome (Reichert Ultracut E, Leica, Austria). Ultrathin sections were contrasted and analysed in a Jeol-100 electron microscope.

Results*Expression of GABA_BRs during prenatal development*

To study the tissue expression of GABA_BRs during prenatal cortical development we carried out pre-embedding immunohistochemical studies at different ages.

GABA_BR1

At E14, GABA_BR1 immunoreactivity was detected in cells bodies located in the mantle layer of the septal and ganglionic eminences (Fig. 1A–D). The intensity of immunoreactivity was strongest at the palaeocortical primordium, at the surface of the lateral and medial ganglionic eminences. Immunoreactivity at the neocortical primordium was less intense (Fig. 1C). The receptor expression level increased with development. Thus, at E16 we found strong immunoreactivity for GABA_BR1 in the habenula, dorsal thalamus, thalamic eminence, hypothalamus, in a circumscribed area surrounding the forming internal capsule and in the marginal zone (MZ) of the neocortical primordia (Fig. 1E). In the MZ, we could distinguish at least two populations of cells immunoreactive for GABA_BR1: large rounded cells located deeply in that layer, and superficial cells running horizontally (Fig. 1H). In the neocortical primordium, the cortical plate (CP) and the subplate (SP) were also immunoreactive (Fig. 1E–G). In addition, GABA_BR1 immunoreactivity was also present in the tangential migratory cells in the intermediate zone (Fig. 1G and I).

At E18 we found increased levels of GABA_BR1 immunoreactivity in the same regions as in previous ages (Fig. 1J–L). Immunoreactivity in the CP was observed in apical dendrites of SP neurons or in ascendent or descendent fibres running in this zone (Fig. 1K and L). GABA_BR1 immunoreactivity was also present in axons running in the intermediate zone, just below the SP (see Fig. 1K and L).

GABA_BR2

At E14, GABA_BR2 immunoreactivity was found at very low levels in most telencephalic areas (Fig. 2A), though it was intense in cells localized in the preplate (PP, Fig. 2A and B). The receptor expression level increased with development. Thus, at E16 we found strong GABA_BR2 immunoreactivity in the neocortical MZ, CP and SP (Fig. 2C–E). In addition, we observed immunolabelled axons in the intermediate zone, as well as numerous immunopositive processes in the ventricular zone (VZ, Fig. 2D and C). However, we did not detect GABA_BR2 immunoreactivity in cell bodies of tangentially migratory neurons in the intermediate zone.

At E18, GABA_BR2 immunoreactivity was strong in the neocortex (Fig. 2F–I), and found in axons running in the lower intermediate zone, as well as in dendrites crossing the CP vertically (Fig. 2G–I). The pyramidal cell layer in and above the SP was devoid of GABA_BR2 immunoreactivity. In the VZ, GABA_BR2 immunoreactivity was stronger than at earlier ages, especially in vertical processes located in the lower part, possibly related with radial glial cell processes (Fig. 2G and I).

Co-localization of GABA_BR1 and GABA_BR2

In order to investigate whether GABA_BR1 and GABA_BR2 were coexpressed in individual neurons during development, we carried out double-immunofluorescence experiments (Fig. 3). At E16, we observed, in the cortical MZ, that around 95% of the cells immunoreactive for GABA_BR2 were also labelled for GABA_BR1 (Fig. 3), resembling pioneer neurons of the MZ (Meyer *et al.*, 1998) and SP neurons. The tangentially migratory cells located in the intermediate zone immunoreactive for GABA_BR1, were not labelled for GABA_BR2 (Fig. 3B, also see Fig. 1G).

Characterization of neurons expressing GABA_BRs

In order to establish which neuronal populations express GABA_BR1 at early stages of development, we carried out double-immunofluorescence experiments for the receptor and for specific neuronal markers using confocal microscopy. We used PSA-NCAM (Fig. 4B), a specific protein of tangentially migratory neurons in the cortical primordium (Seki & Arai, 1991); calbindin (CB, Fig. 4E), a specific protein expressed in interneurons at early stages of development (Anderson *et al.*, 1997); and Reelin (Fig. 4H), an extracellular matrix glycoprotein that is synthesized and secreted by Cajal-Retzius cells (D'Arcangelo *et al.*, 1995).

We observed an intense labelling for PSA-NCAM in cells located in the intermediate zone, SP and MZ (Fig. 4B), displaying the morphology of tangentially migrating neurons. A quantitative survey showed that around 97% of the cells immunopositive for PSA-NCAM in the MZ and intermediate zone also showed GABA_BR1 immunoreactivity (Fig. 4C), suggesting that the receptor was expressed in the majority of the tangentially migratory cells. However, not all cells with GABA_BR1 immunoreactivity also expressed PSA-NCAM, indicating that the receptor might be also present in a population of nonmigratory neurons.

Using a monoclonal antibody against calbindin, we found intense labelling in cells located in the cortical MZ, corresponding to large rounded cells (Fig. 4E), resembling pioneer neurons. Quantitative studies showed that at least 83% of those cells also showed GABA_BR1 immunoreactivity.

Finally, in order to investigate whether the horizontal neurons expressing GABA_BR1 in the cortical layer I were Cajal-Retzius cells, we used Reelin (Fig. 4H). In prenatal animals, we did not detect colocalization of GABA_BR1 and Reelin. However, in postnatal

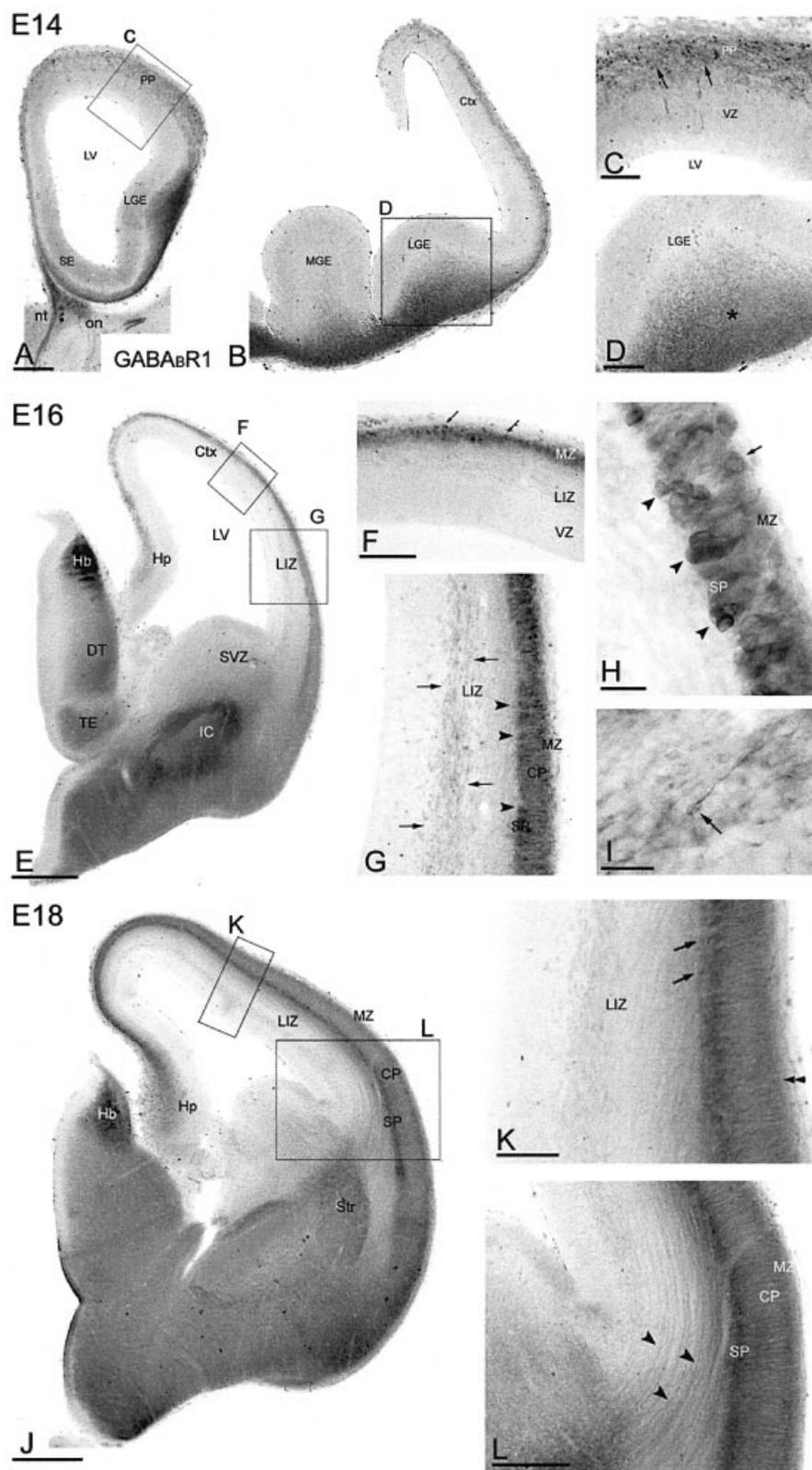


FIG. 1. Immunoreactivity for GABA_BR1 during prenatal development using a pre-embedding immunoperoxidase method. Rostral (A) and mid-caudal (B) sections of the telencephalon at E14 showing GABA_BR1 immunoreactivity in the mantle zones of the septal (SE) and of both medial (MGE) and lateral (LGE) ganglionic eminences, and in the preplate (PP) and marginal zone (MZ) of the cortical anlage. Ctx, cortex; nt, *nervus terminalis* on, olfactory nerve. (C) Detail of the telencephalic area boxed in panel A, showing strong GABA_BR1 immunolabelling in PP cells (arrows) and weaker labelling in the ventricular zone (VZ). (D) Detail of the LGE area boxed in panel B. The mantle layer of the LGE contained a large number of cells expressing GABA_BR1 (asterisk). (E) Mid-caudal coronal section of the telencephalon at E16 showing a strong GABA_BR1 immunolabelling in the habenula (Hb), dorsal thalamus (DT), internal capsule (IC) and cortex (Ctx). Hp, hippocampus; SVZ, subventricular zone; LIZ, lower intermediate zone. (F) Detail of the cortex boxed in panel E. GABA_BR1 immunolabelling was observed in large rounded cells of the cortical MZ (arrows). (G and I) GABA_BR1 immunoreactivity was also strong in the tangential migratory cells located in the lower part of the intermediate zone (LIZ) (e.g. arrows). (G and H) GABA_BR1 immunoreactivity was also detected in cells located in the subplate (SP) (arrowheads), forming groups in this cortical layer (arrowheads). (J) Mid-caudal coronal section of the telencephalon at E18 showing a strong GABA_BR1 immunoreactivity in the SP, CP and MZ of the neocortex, and in the tangentially migratory neurons in the LIZ. Str, striatum. (K and L) Detail of the areas boxed in panel J, showing GABA_BR1 immunopositive cells in the SP (arrows) and MZ (double arrowhead), as well as in axons located immediately beneath the SP (arrowhead). on, olfactory nerve; LV, lateral ventricle; DT, dorsal thalamus; IC, internal capsule; TE, thalamic eminence. Scale bars, 200 μ m (A–B and L); 100 μ m (C, D, F, G and K); 500 μ m (E and J); 30 μ m, (H and I).

animals (P2), the estimate of colocalization showed a 97% overlap for GABA_BR1- and Reelin-immunoreactivity in cortical layer I cells (Fig. 4I), suggesting that the neurons expressing GABA_BR1 are indeed Cajal-Retzius cells.

Expression of GABA_BR_s during postnatal development

To study the tissue expression of GABA_BR_s during postnatal development we carried out pre-embedding immunohistochemical studies at different ages up to adulthood.

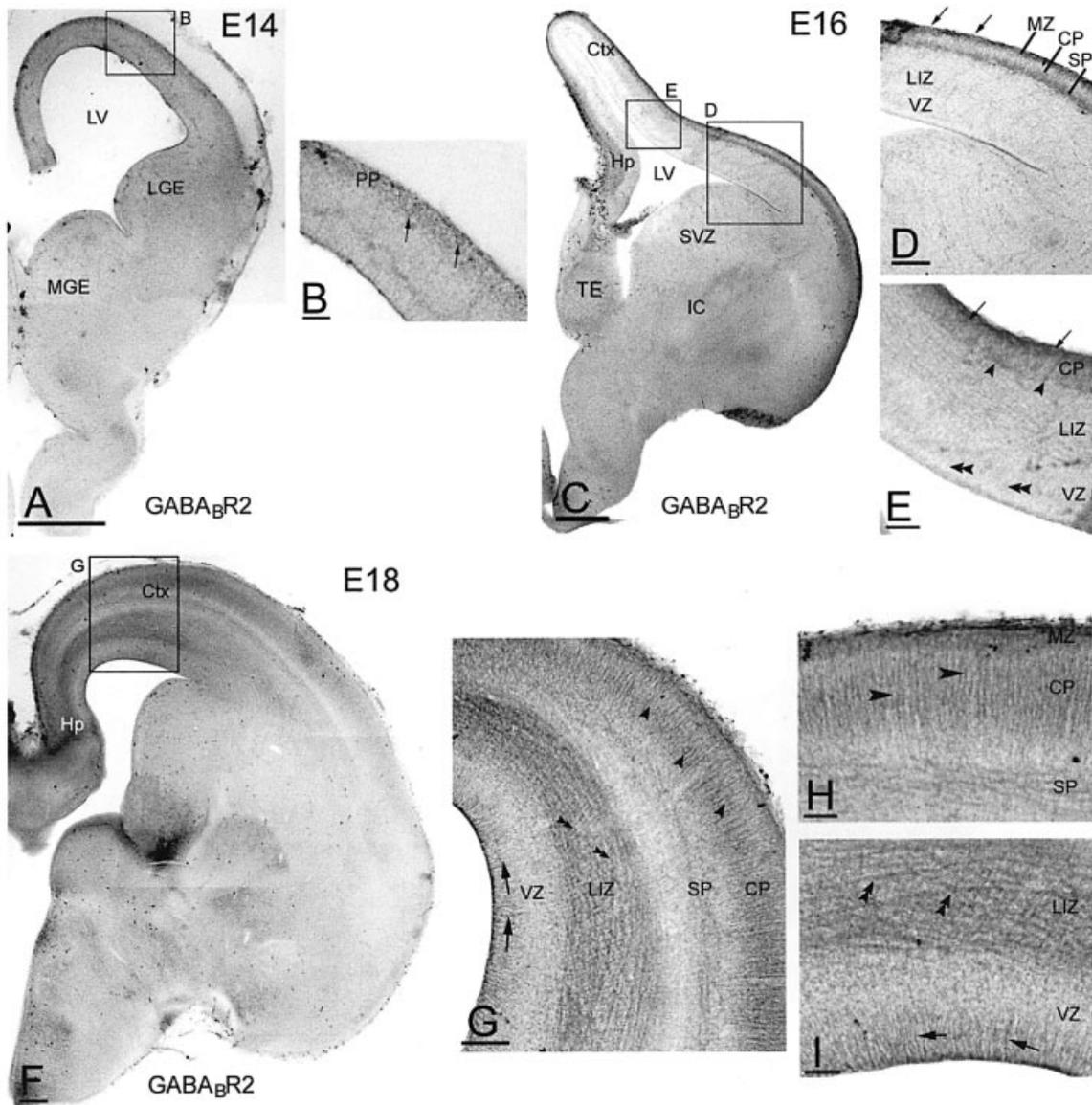


FIG. 2. Immunoreactivity for GABA_BR2 during prenatal development using a pre-embedding immunoperoxidase method. (A) Coronal section at E14 showing GABA_BR2 immunoreactivity in the preplate (PP) of the cortex, with no detectable immunoreactivity observed in other areas. (B) Detail of the PP boxed in panel A. GABA_BR2 immunoreactivity was observed in cells located throughout this cortical stratum. (C–E) Mid-caudal coronal section at E16 showing GABA_BR2 immunoreactivity in different telencephalic areas. (D and E) Detail of two cortical areas boxed in panel C, showing GABA_BR2 immunoreactivity in cells located in the subplate (SP, arrowheads), cortical plate (CP) and marginal zone (MZ, arrows), as well as in processes and cells distributed in the ventricular zone (VZ, double arrowheads). (F–I) Mid-caudal coronal section at E18 showing GABA_BR2 immunoreactivity in different telencephalic areas. (G) Detail of boxed area in panel F. Immunoreactivity was localized in CP cells (arrowheads). Axons immunoreactive for GABA_BR2 were detected immediately above the VZ (double arrowhead). Immunoreactivity for GABA_BR2 in the VZ was more evident at E18 than at earlier ages (arrows). Ctx, cortex; MGE, medial ganglionic eminence; LGE, lateral ganglionic eminence; Hp, hippocampus; DT, dorsal thalamus; LIZ, lower intermediate zone; IC, internal capsule; SVZ, subventricular zone; LV, lateral ventricle. Scale bars, 200 μm (A, C and F); 50 μm (B, E, H and I); 100 μm (D and G).

GABA_BR1

At birth, we found high expression levels in all layers of the neocortex (Fig. 5A). Thus, GABA_BR1 immunoreactivity was detected mainly in cortical layer I neurons (Fig. 6B) having the typical fusiform morphology and horizontal orientation of Cajal-Retzius cells. In supragranular layers, GABA_BR1 was observed in vertical processes most likely corresponding to the apical dendrites of layers

IV–V pyramidal cells (Fig. 5A). This characteristic pattern of distribution changed during development. Thus, at around P7, we observed strong labelling in somata (Fig. 5C and F) and apical dendrites (Fig. 5E) of layer V pyramidal cells, and in layer VIb cells (Fig. 5A, F, I and N).

At P7, GABA_BR1 immunolabelling was found in interneurons of supragranular layers, most of them located between layers I and II/III (Fig. 5D and G). GABA_BR1 immunolabelling was still found in

FIG. 3. Co-localization of GABA_BR1 and GABA_BR2 during prenatal development as demonstrated using double immunofluorescence. Low magnification of a neocortical region at E16 double-labelled for GABA_BR1 (A, red) and GABA_BR2 (B, green). (C) Overlay of panels A and B. Note the high degree of colocalization of these GABA_BRs at the MZ. However, cells in the LIZ appear immunoreactive for GABA_BR1 but not for GABA_BR2 (arrowheads). (D–F) High magnification of MZ region labelled for both GABA_BRs. Virtually all cells in that region showed colocalization of the two receptor subtypes. Scale bars, 10 μ m (A–F).

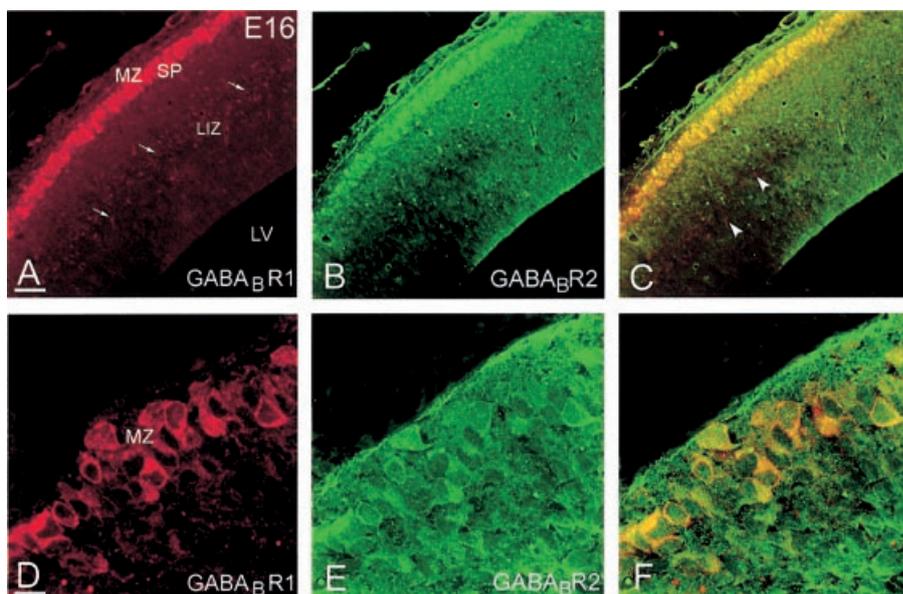
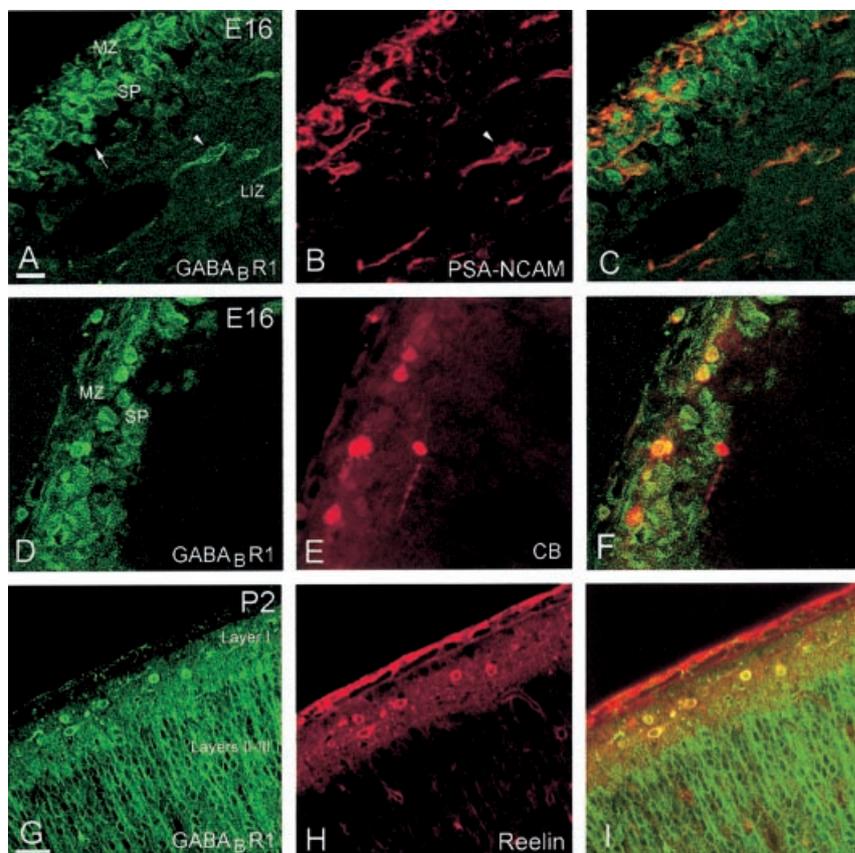


FIG. 4. Characterization of cell populations expressing GABA_BR1 during pre- and postnatal development. Double-immunofluorescence for GABA_BR1 and PSA-NCAM, calbindin and Reelin. (A–C) Immunofluorescence for GABA_BR1 (A, green) and PSA-NCAM (B, red) in the neocortex at E16. (C) Overlay of panel A and B. Almost all the PSA-NCAM positive cells expressed GABA_BR1 (e.g., arrowheads). (D–F) Immunofluorescence for GABA_BR1 (D, green) and calbindin (E, red), showing that almost all calbindin positive cells in the MZ were positive for GABA_BR1 (F, overlay). (G–I), Immunofluorescence for GABA_BR1 (G, green) and Reelin (H, red) in layer I of the neocortex at P2. I, Overlay of panel G and H. GABA_BR1 immunoreactivity was observed throughout all layers of the cortex. The specific protein of Cajal-Retzius cells, Reelin, was restricted to layer I neurons (see panel N). Virtually all cells expressing Reelin also showed GABA_BR1. Scale bars, 20 μ m (A–I); 40 μ m (G–I).



putative Cajal-Retzius cells (Fig. 5D), though at P10 the number of those neurons decreased (Fig. 5G) and disappeared shortly after that age. GABA_BR1 immunoreactivity was also observed in layer V pyramidal cells and their apical dendrites (Fig. 5C, E and F).

At P15, the strong GABA_BR1 labelling of pyramidal cell somata and apical dendrites decreased dramatically, while an intense

neuropilar staining appeared in all cortical layers, especially in layer V (Fig. 5I). In addition, we also observed many GABA_BR1-immunoreactive interneurons with multipolar and bipolar (Fig. 5I and J) morphology distributed throughout all cortical layers (Fig. 5J).

At P21–P30, the adult levels of GABA_BR1 immunoreactivity in the neocortex were reached. Immunolabelling was intense in the

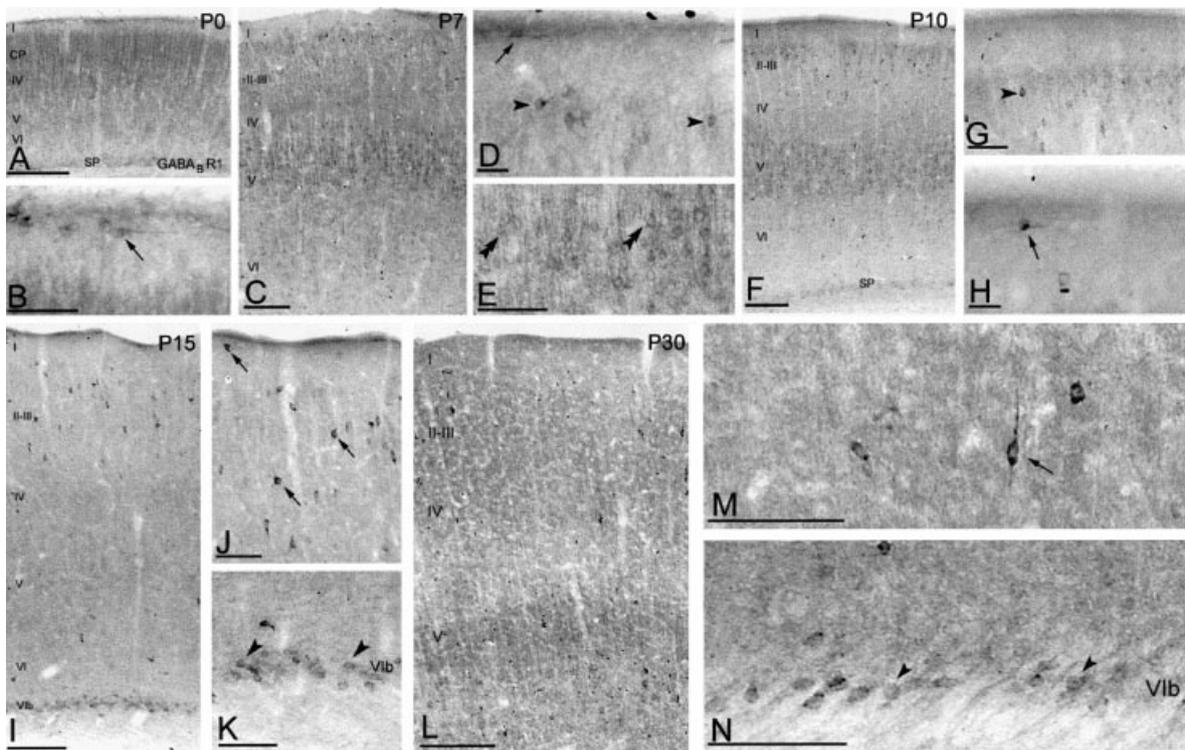


FIG. 5. Immunoreactivity for GABA_BR1 in the neocortex during postnatal development using a pre-embedding immunoperoxidase method. (A) At P0, GABA_BR1 immunolabelling was detected in somata and apical dendrites of neocortical pyramidal cells. Immunolabelling was intense in layers II/III and in cells located in layer IV. (B) Detail of the GABA_BR1 immunoreactivity in cells with a horizontal morphology located in layer I (arrow). (C–E) At P7, strong GABA_BR1 immunoreactivity was found in somata and apical dendrites of layer V pyramidal neurons (E, double arrowheads). GABA_BR1 immunoreactive cells were detected in layer I (D, arrow), as well as in interneurons in the upper part of layer II/III (D, arrowheads). (F–H) At P10, intense GABA_BR1 immunolabelling was observed in layer V cells, layer II/III interneurons (arrowheads) and in cells located in layer VIb. The number of immunopositive cells in layer I considerably decreased with age, and only a few cells could still be detected at P10 (arrow). (I–K) At P15, GABA_BR1 immunoreactivity was found in interneurons throughout the neocortex, though particularly strong in supragranular layers (J, arrows), as well as in cells in layer VIb (K, arrowheads). A neuropilar staining was also detected throughout the neocortex. (L–N) At P30, the number of interneurons immunoreactive for GABA_BR1 decreased (M, arrow), although the neuropilar immunoreactivity remained similar to that at earlier ages. We also detected immunopositive cells located in layer VIb (arrowheads). Scale bars, 200 μ m (A, C, F, I and L), 20 μ m (B and D); 100 μ m (E, G, J, K, M and N).

neuropil of layers I, II/III and V (Fig. 5L), as well as a large proportion of immunoreactive interneuron somata and dendritic processes throughout all cortical layers (Fig. 5M).

GABA_BR2

During postnatal ages, we also observed a high expression of GABA_BR2 in the neocortex. At P2, we found an intense GABA_BR2 immunolabelling in most cortical layers (Fig. 6A and B), and particularly strong in the neuropil of layer I (Fig. 6B). In supragranular layers, GABA_BR2 was detected in vertical processes, probably apical dendrites of infragranular pyramidal cells (Fig. 6B), though this characteristic pattern of distribution changed during development (Fig. 6). Thus, at P7 and P10, the strongest neuropilar staining was detected in layer I (Fig. 6C and D), but from P15 onwards, all layers reached similar levels of GABA_BR2 immunoreactivity (Fig. 6E and F). We did not detect immunoreactive somata of pyramidal cells or interneurons in any layer at any developmental age.

Subcellular localization of GABA_BRs during pre- and postnatal development

To investigate the precise subcellular localization of GABA_BR1 and GABA_BR2 during pre- and postnatal development we carried out electron microscopic investigations using the pre-embedding immunoperoxidase and immunogold techniques.

GABA_BR1

During prenatal development we concentrated on the SP and CP, where the intensity of immunostaining at E18 was the strongest. GABA_BR1 immunoreactivity was found postsynaptically in SP cells with an elongated morphology and only one dendritic process (Fig. 7) running parallel to the pial surface. In all the cases, the peroxidase reaction end-product was observed forming patches and associated with the plasma membrane (Fig. 7A and B) or intracellular organelles (Fig. 7C and D).

During postnatal development, GABA_BR1 immunoreactivity was associated to the plasma membrane and the endoplasmic reticulum (Fig. 8A–C) of layer I cells with an ultrastructure resembling that of Cajal-Retzius cells (Fig. 8A). Strong GABA_BR1 immunoreactivity was also found in the somata (Fig. 8D) and dendritic trees (Fig. 8D–F) of a subset of interneurons in all layers of the neocortex, invariably associated with the plasma membrane and intracellular organelles. Peroxidase reaction end-product for GABA_BR1 filled dendritic spines of pyramidal cells throughout all cortical layers (Fig. 8G and H). The dendritic shafts of pyramidal cells were also immunolabelled, but the density of the reaction product was consistently lower than that observed within the spines.

Finally, we also observed GABA_BR1 immunoreactivity in presynaptic elements, distributed throughout all cortical layers and

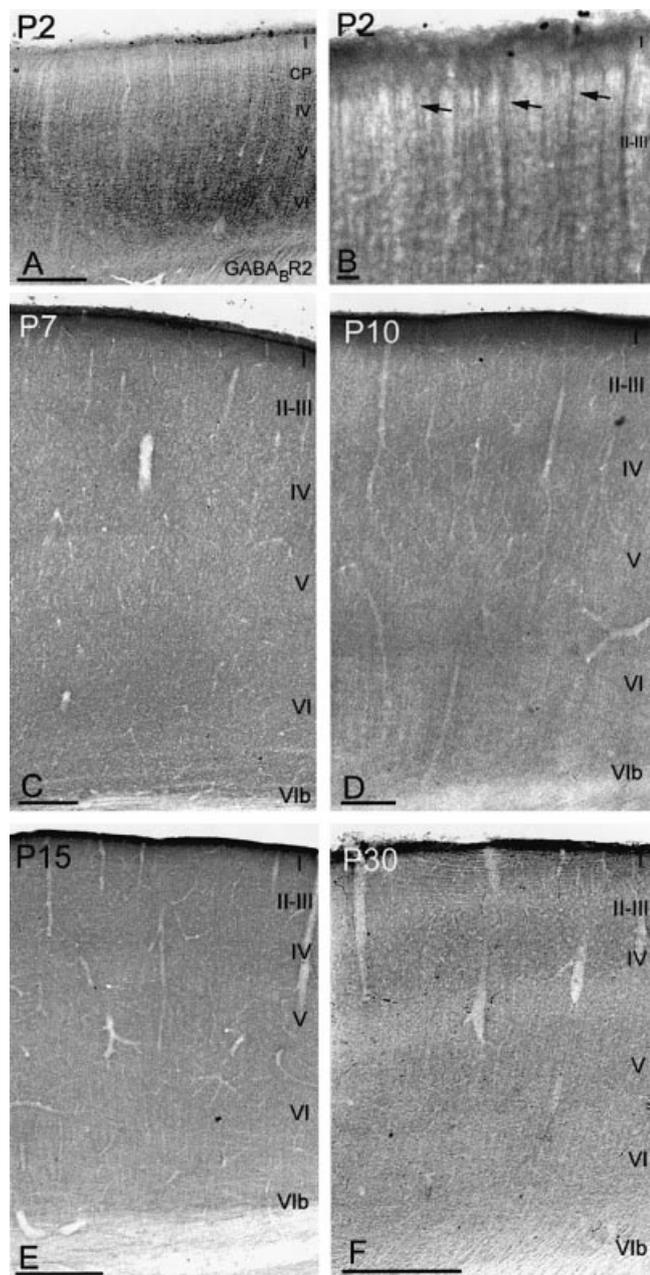


FIG. 6. Immunoreactivity for GABA_BR2 in the neocortex during postnatal development using a pre-embedding peroxidase method. (A) At P2, GABA_BR2 immunolabelling was intense in layers I and II/III and in cells located in layer IV. (B) Detail of the GABA_BR2 immunoreactivity in upper layers of the neocortex showing labelled apical dendrites of pyramidal cells (arrows). (C–D) At P7 and P10, GABA_BR2 immunoreactivity was intense in layer I. In the remaining layers, a uniform neuropilar GABA_BR2 immunostaining was observed. (E–F) From P15 to adulthood, GABA_BR2 immunoreactivity was moderate to intense throughout the neocortex. Scale bars, 100 μ m (A, C and D); 20 μ m (B); 300 μ m (E and F,).

postnatal ages (Fig. 8I and J). Peroxidase reaction end-product filled axon terminals establishing asymmetrical synapses (e.g., see Fig. 8I). A few immunolabelled symmetrical synaptic contacts were also observed. Immunoparticles were also found on axonal membranes and occasionally at the presynaptic active zone of putative glutamatergic synapses (Fig. 8K).

GABA_BR2

The distribution of GABA_BR2 immunoreactivity in the neocortex during postnatal development was almost identical to the distribution of GABA_BR1. Thus, peroxidase reaction end-product for GABA_BR2 filled dendritic spines and dendritic shafts of pyramidal cells (Fig. 9A–G). Immunoreactive dendrites established asymmetrical synaptic contacts with putative glutamatergic axon terminals (Fig. 9A, D, E and G). Immunoparticles were mostly detected on the extrasynaptic plasma membrane of pyramidal cells spines (Fig. 9H and G) and also at the edge of asymmetrical synapses (Fig. 9I). GABA_BR2 immunoreactivity was also detected in pre-synaptic elements throughout development (Fig. 9B and F). Immunoparticles were located on axonal membranes, as well as at the presynaptic active zone of putative glutamatergic synapses (Fig. 9J).

Discussion

The present study provides the first detailed description of the expression pattern, and precise cellular and subcellular localization of GABA_BR in the rat neocortex during prenatal and postnatal development. Using conventional light and confocal microscopy we have shown that GABA_BR1 and GABA_BR2 are expressed from very early stages of cortical development and are widely distributed in the neocortex during development in pyramidal and nonpyramidal cells. The colocalization of the two receptor subtypes in the same embryonic cortical cells suggests that GABA_BR form heteromeric complexes and that these cells express functional GABA_BR from very early stages of development. Furthermore, immunoelectron microscopy revealed an association of GABA_BR1 and GABA_BR2 with putative glutamatergic synapses and demonstrated their post-synaptic and pre-synaptic localization in the neocortex during development. The similar subcellular localization of the two receptor subtypes suggest that during cortical morphogenesis, functional GABA_BR expressed in different neuronal compartments may play an important role in maturation and cortical organization events.

Expression of GABA_BRs in embryonic cortical cells

GABA_BR1 and GABA_BR2 were expressed from very early stages of neocortical development, though they differed mainly in the intensity of immunolabelling within the same brain areas. This early presence of GABA_BR in the neocortex is in line with a recent study showing that embryonic cortical cells contain mRNA and protein for GABA_BR1, and that this receptor subtype is functional at E17 in both cortical plate and ventricular zone cells (Behar *et al.*, 2001). Our data further demonstrated that GABA_BR1 is expressed in cortical plate cells even earlier, at E14, and showed that it is present in the tangentially migratory interneurons coming from the medial ganglionic eminence.

Previous *in vitro* studies, using neurons dissociated from immature cortical regions, demonstrated that GABA could act as a chemo-attractant, stimulating embryonic nerve cell movement (Behar *et al.*, 1996, 1998). The same authors reported that GABA released by cortical cells stimulates postmitotic layer II/III cells to migrate radially into the cortical plate via activation of saclofen- and picrotoxin-sensitive receptors (Behar *et al.*, 2000). Taken together, these and our results suggest that both routes of neuronal migration, radial and tangential, might be modulated via activation of GABA_BR.

Our immunohistochemical studies have indicated that a majority of MZ and CP cells coexpress GABA_BR1 and GABA_BR2 during prenatal development. It has been suggested that the coassembly of

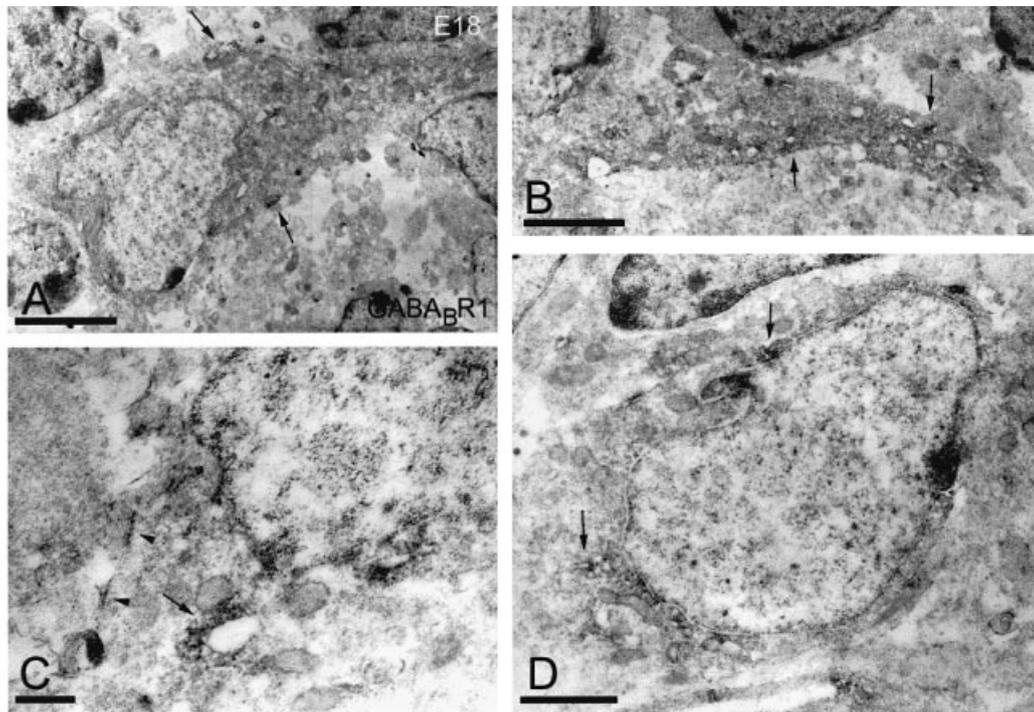


FIG. 7. Electron micrographs of the neocortex showing immunoreactivity for GABA_BR1 in subplate neurons at E18, using a pre-embedding immunoperoxidase method. (A–D) Peroxidase reaction end-product (arrows) was concentrated near the plasma membrane of cell bodies (A and D) and processes (B) of cells located in that stratum. GABA_BR1 immunoreactive synaptic contacts onto these cells were also observed (C, arrowheads). Scale bars, 2 μm (A–B); 0.3 μm (C); 10 μm (D).

GABA_BR1 and GABA_BR2 subtypes into heteromeric complexes is required for the translocation of the receptor from the synthetic site to the cell surface (Margeta-Mitrovic *et al.*, 2000) and also for agonist-induced signalling (Couve *et al.*, 1998; Jones *et al.*, 1998; Kaupmann *et al.*, 1998a; White *et al.*, 1998; Kuner *et al.*, 1999). Furthermore, colocalization of GABA_BR1 and GABA_BR2 has been observed in the CNS (Kaupmann *et al.*, 1998a; Billinton *et al.*, 2000; Gonchar *et al.*, 2001; Kulik *et al.*, 2002) in adult animals. However, no colocalization has been demonstrated in embryonic cortical cells yet. Therefore, our finding, that both GABA_BR subtypes are expressed in the MZ and CP cells, suggests that membrane-bound GABA_BR are functionally active at an early stage of neuronal development.

In the neocortex, the precise functional significance of GABA_BR during development remains unclear. Obrietan & van den Pol (1998, 1999) postulated that in embryonic cells, GABA_BR play a crucial inhibitory role regulating Ca²⁺ rises elicited by GABA_A activation and/or by inhibiting glutamate receptor-mediated Ca²⁺ transients. Alterations in intracellular Ca²⁺ are involved in many processes essential for neuronal development, such as growth cone motility, neurite outgrowth, neuronal migration and synapse formation (Komuro & Rakic, 1996; Mattson & Kater, 1987; Nelson *et al.*, 1990). Our finding of abundant GABA_BR1 and GABA_BR2 immunoreactivity in the embryonic cortex supports a potentially important role for GABA_BR in a wide array of physiological processes during corticogenesis.

Developmental distribution of GABA_BRs

The immunohistochemical distribution pattern of GABA_BR1 and GABA_BR2 in the neocortex is consistent with earlier *in situ* hybridization (Kaupmann *et al.*, 1998a; Bischoff *et al.*, 1999;

Durkin *et al.*, 1999; Clark *et al.*, 2000) and autoradiographic studies (Bowery *et al.*, 1987; Chu *et al.*, 1990; Turgeon & Albin, 1994).

GABA_BR1, but not GABA_BR2, was observed in Cajal-Retzius cells and the tangential migratory cells during prenatal development. Physiological responses following activation of GABA_BR require the coassembly of GABA_BR1 and GABA_BR2 (Jones *et al.*, 1998; Kaupmann *et al.*, 1998b; White *et al.*, 1998; Kuner *et al.*, 1999). The possible expression of homomeric GABA_BR1 by Cajal-Retzius cells and the tangential migratory cells in the lower intermediate zone (LIZ) suggests either that they are nonfunctional GABA_BR or that additional subtypes or isoforms of GABA_BR and/or other proteins (White *et al.*, 2000) exist that have not yet been identified. The absence of functional coupling to postsynaptic ion channels (White *et al.*, 1998; Martin *et al.*, 1999) is supported by our preliminary electrophysiological data on Cajal-Retzius cells that revealed no activation of outward current after application of baclofen (G. López-Bendito, R. Luján and O. Paulsen, unpublished observations). Overall, the spatiotemporal expression of GABA_BR1 reported here in the MZ/layer I may be an important mechanism to regulate Ca²⁺ dynamics, which is thought to play a role in the regulation of gene expression and control of developmental events.

Interestingly, GABA_BR1 showed a developmental reorganization in cortical pyramidal cells peaking around P10. Thus, during the first postnatal week, GABA_BR1 was located preferentially in pyramidal cell somata and dendrites projecting to layer I. At P10–P15, the receptor became more uniformly distributed on the surface of the pyramidal cells, though mainly localized in dendritic spines and associated with synapses. A similar redistribution has also been described for group I mGluRs in the same neocortical cell types during development (López-Bendito *et al.*, 2002).

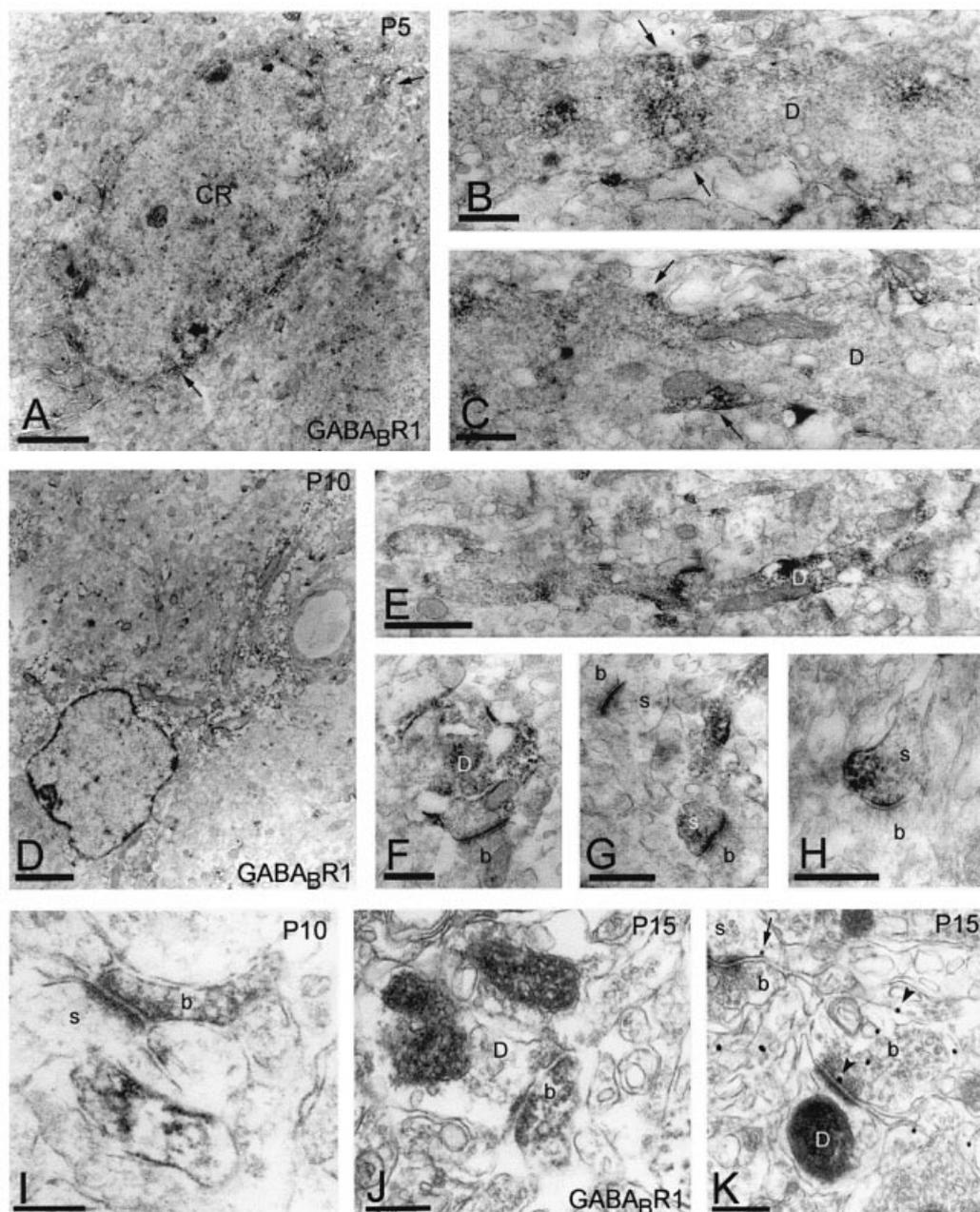


FIG. 8. Electron micrographs showing GABA_BR1 immunoreactivity in the neocortex during postnatal development, using pre-embedding immunoperoxidase and immunogold methods. (A–C) Presence of GABA_BR1 in Cajal-Retzius cells (CR) at P5. Note the large cell body and the abundant rough endoplasmic reticulum. Immunolabelling was concentrated near the plasma membrane of cell bodies and processes (arrows). (D–F) At P10, GABA_BR1 immunoreactivity was observed mainly in somata and dendrites (white 'D') of interneurons in layer II/III. (G and H) Dendritic spines (s) of pyramidal cells immunolabelled for GABA_BR1. (I and J) GABA_BR1 immunoreactivity in axon terminals establishing asymmetrical synapses with spines and dendritic shafts ('D') in layer I (I) and II/III (J) at P10 and P15, respectively. (K) Immunogold particles were localized on the extrasynaptic plasma membrane of dendritic spines (s, arrow) and axon terminals (b, arrowheads). White 'D' and small black 'D' show dendrite. Scale bars, 2 μ m (A and D); 0.5 μ m (B, C and F); E, 1 μ m.

Interneurons expressed GABA_BR during development in the neocortex, with the staining for GABA_BR1 being present in somata and extending uniformly throughout the dendritic arbour. However, although at light microscopic level the staining for GABA_BR2 was not detected in cell bodies of interneurons, electron microscopy showed that they expressed this subtype indeed. The GABA_BR1-containing interneurons showed different morphology and location throughout the neocortex, suggesting the existence of different subset of interneurons expressing GABA_BR. This is in agreement with

findings of Gonchar *et al.* (2001) who reported that parvalbumin-, calretinin- and somatostatin-containing interneurons of the visual cortex colocalized with GABA_BR1a/b.

Subcellular localization of GABA_B Rs

The current view of the subcellular distribution of GABA_BR in the CNS is mainly based on data obtained in the adult cerebellum (Kaupmann *et al.*, 1998a; Fritschy *et al.*, 1999; Ige *et al.*, 2000; Kulik *et al.*, 2002), ventrobasal thalamus (Kulik *et al.*, 2002) and visual

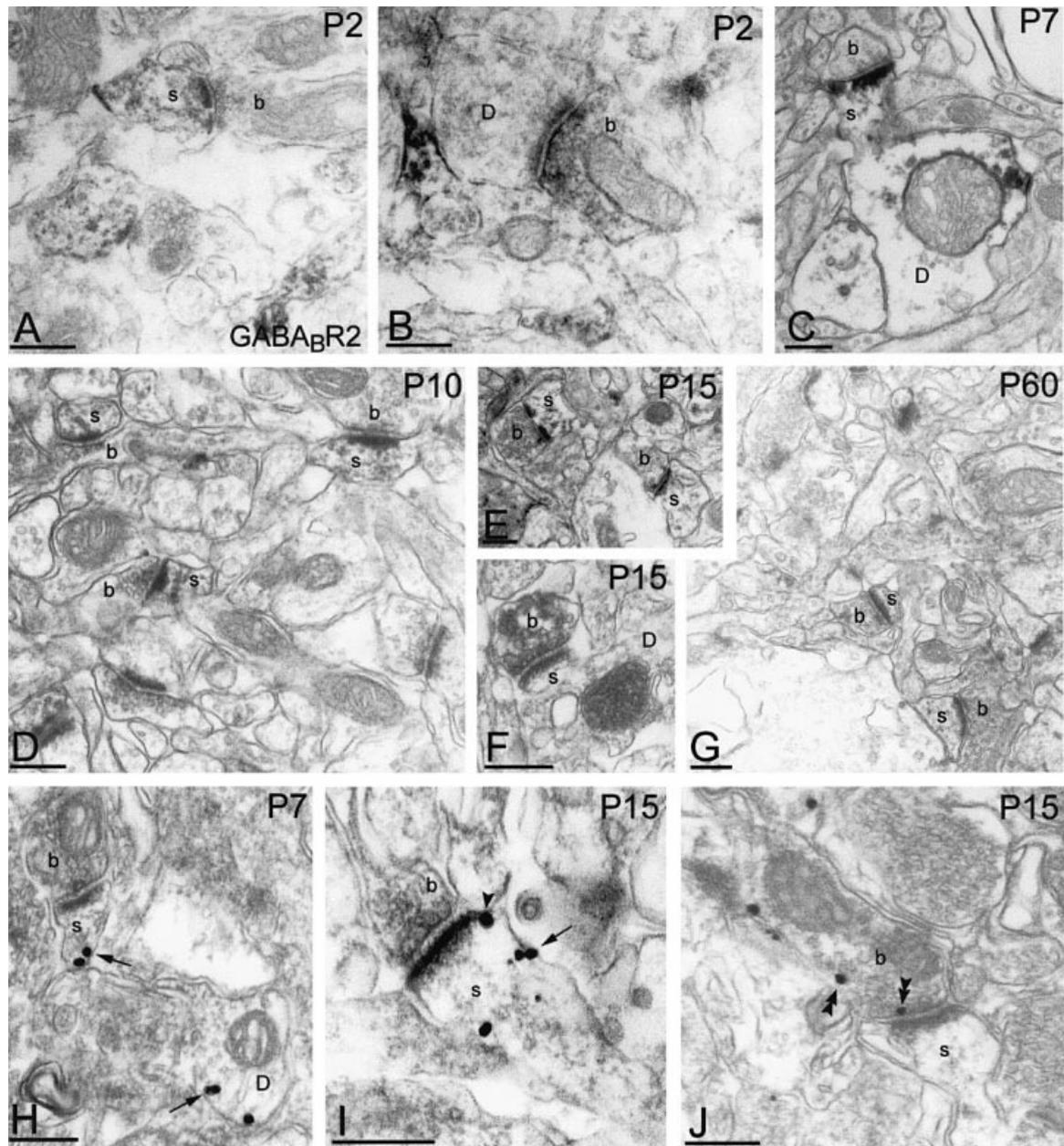


FIG. 9. Electron micrographs showing GABA_BR2 immunoreactivity in the neocortex during postnatal development, using pre-embedding immunoperoxidase and immunogold methods. (A and B) At P2, GABA_BR2 immunolabelling concentrated in dendritic spines (s) of pyramidal cells, as well as in some axon terminals (b) establishing synapses with dendritic spines or dendritic shafts ('D') in layer I. (C) Dendrite in layer II/III immunolabelled for GABA_BR2 at P7. Peroxidase reaction end-product concentrated near the plasma membrane and in the spine (s) attached to the shaft ('D'). (D–G) GABA_BR2 immunoreactivity was consistently present in dendritic spines (s) of pyramidal cells throughout the neocortex in postnatal development until adulthood, as well as in axon terminals establishing asymmetrical synapses with dendritic spines (s) or shafts ('D'). (H–I) Immunogold particles were localized in the extrasynaptic (arrows) plasma membrane of dendritic spines (s) and shafts ('D') of pyramidal cell, as well as in perisynaptic sites (I, arrowhead) associated with asymmetrical synapses. (J) Presynaptically, gold particles were also present along the plasma membrane of axon terminals (b, double arrowheads) establishing asymmetrical synapses with dendritic spines (s) of pyramidal cells. Black small 'D' indicates dendrite. Scale bars, 0.25 μ m (A–J).

cortex (Gonchar *et al.*, 2001). Those studies showed that GABA_BR1 and GABA_BR2 were found at extrasynaptic sites, at the edges of asymmetrical synapses and occasionally over the synaptic specialization. Our finding that a high proportion of dendritic spines of cortical pyramidal cells were immunoreactive for both GABA_BR1 and GABA_BR2 is consistent with previous studies (Benke *et al.*, 1999; Bischoff *et al.*, 1999; Fritschy *et al.*, 1999; Gonchar *et al.*,

2001; Muñoz *et al.*, 2001). The majority of GABA_BR were localized on cortical dendritic spines, mainly at extrasynaptic sites or at the edge of the synaptic specialization. The similar subcellular localization of GABA_BR1 and GABA_BR2 on neuronal elements of the neocortex presented in our study further suggests colocalization of these receptors in different subcellular compartments of pyramidal and nonpyramidal cells during development. In addition, our finding

directly demonstrated that pyramidal cells and interneurons contained both postsynaptic and presynaptic GABA_BR.

Several studies have shown that baclofen suppressed evoked excitatory postsynaptic potentials (EPSPs) and IPSPs, suggesting that GABA_BR mediate presynaptic inhibition of excitatory and inhibitory neurotransmission in different brain regions (Pittaluga *et al.*, 1987; Brugger *et al.*, 1993). In the neocortex, we found GABA_BR1 and GABA_BR2 in axon terminals establishing asymmetrical synapses during cortical development. This finding is consistent with electrophysiological studies that showed depression of excitatory responses by GABA_BR agonist in neocortex (Howe *et al.*, 1987; Deisz *et al.*, 1997; Varela *et al.*, 1997).

The association of GABA_BR with asymmetrical synapses, presumably glutamatergic, in adult and during development raises the question of what is the source of GABA to activate these receptors. One possible source of GABA to activate GABA_BR is the spillover from neighbouring interneurons (Isaacson *et al.*, 1993). Another possibility is the activation through taurine, a weak agonist at GABA_BR. Taurine is present in the cortex and has been suggested to act as a chemoattractant for neurons during cortical development via mechanisms involving GABA_BR (Behar *et al.*, 2001). Finally, GABA_BR may be activated by nonsynaptically released GABA. In the neocortex, this mechanism of activation has recently been suggested for presynaptic GABA_BR on pyramidal cell axon terminals (Zilberter *et al.*, 1999). Regardless of the mechanism of activation of GABA_BR, the colocalization of GABA_BR1 and GABA_BR2 at excitatory synapses suggests that they are functional and it is likely that they are involved in developmental physiological events.

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Abbreviations

CP, cortical plate; Ctx, cortex; DAB, 3,3'-diaminobenzidine; DT, dorsal thalamus; E, embryonic day; GABA_BR, metabotropic γ -aminobutyric acid receptor; Hb, habenula; Hp, hippocampus; IC, internal capsule; LGE, lateral ganglionic eminence; LIZ, lower intermediate zone; MGE, medial ganglionic eminence; MZ, marginal zone; NGS, normal goat serum; nt, *nervus terminalis*; on, olfactory nerve; P, postnatal day; PB, phosphate buffer; PBS, phosphate buffer saline; PP, preplate; PSA-NCAM, polysialylated form of the neuronal cell adhesion molecule NCAM; SE, septal eminence; SP, subplate; SVZ, subventricular zone; TB, Tris buffer; TBS, Tris-buffered saline; TE, thalamic eminence; VZ, ventricular zone.

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