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Two pools of Triton X-100-insoluble GABA_A receptors are present in the brain, one associated to lipid rafts and another one to the postsynaptic GABAergic complex

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Abstract

Rat forebrain synaptosomes were extracted with Triton X-100 at 4°C and the insoluble material, which is enriched in postsynaptic densities (PSDs), was subjected to sedimentation on a continuous sucrose gradient. Two pools of Triton X-100-insoluble GABA_A receptors (GABA_ARs) were identified: I) a higher-density pool ($\rho = 1.10\text{--}1.15$ g/ml) of GABA_ARs that contains the $\gamma 2$ subunit (plus α and β subunits) and that is associated to gephyrin and the GABAergic postsynaptic complex; and II) a lower-density pool ($\rho = 1.06\text{--}1.09$ g/ml) of GABA_ARs associated to detergent-resistant membranes (DRMs) that contain α and β subunits but not the $\gamma 2$ subunit. Some of these GABA_ARs contain the δ subunit. Two pools of GABA_ARs insoluble in Triton X-100 at 4°C were also identified in cultured hippocampal neurons: I) a GABA_AR pool that forms clusters that colocalize with gephyrin, and remains Triton X-100-insoluble after cholesterol depletion, and II) a GABA_AR pool that is diffusely distributed at the neuronal surface, that can be induced to form GABA_AR clusters by capping with an anti- $\alpha 1$ GABA_AR subunit antibody, and that becomes solubilized in Triton X-100 at 4°C after cholesterol depletion. Thus, there is a pool of GABA_ARs associated to lipid rafts that is non-synaptic and that has a subunit composition different from that of the synaptic GABA_ARs. Some of the lipid raft-associated GABA_ARs might be involved in tonic inhibition.

Keywords

lipid rafts; detergent resistant membrane (DRM); γ -aminobutyric acid type-A receptor; hippocampal cultures; gephyrin; postsynaptic density (PSD)

INTRODUCTION

The γ -Aminobutyric acid type A receptors (GABA_ARs) concentrate at the postsynaptic membrane of GABAergic synapses both in the brain and in culture (Craig et al. 1994; Somogyi et al. 1996; Nusser et al. 1995, 1998; Fritschy et al. 1998; Sassoe-Pognetto et al. 2000; Christie et al. 2002a,b). There are also GABA_ARs at the neuronal cell surface that are not associated with GABAergic synapses (Nusser et al. 1998; Brickley et al. 1999; Christie et al. 2002a,b, 2006; Fritschy and Brunig 2003; Wei et al. 2003). Several detergents, including Triton X-100, have been employed for the solubilization of GABA_ARs from brain membranes, as a first step

for the purification and biochemical characterization of the solubilized GABA_ARs. There is also a population of brain GABA_ARs that resists detergent extraction, which has remained largely non-characterized and is the focus of the present study.

Several methods have been described for the preparation of postsynaptic densities (PSDs) from the excitatory glutamatergic synapses with Gray's type-I morphology (type-I PSDs). These methods are based on the insolubility of these structures in various detergents (Cotman et al. 1974; Carlin et al. 1980; Walikonis et al. 2000). The preparation of type-I PSDs has allowed the identification and characterization of many of the protein components of these structures (Kennedy 1997; Walikonis et al. 2000; Yoshimura et al. 2004; Peng et al. 2004; Jordan et al. 2004; Collin et al. 2006). In contrast, there are no methods in the literature describing the preparation of PSDs from the inhibitory GABAergic synapses with Gray's type-II morphology (or type-II PSDs). In this communication we have devised a method for the preparation of a brain fraction enriched in type-II GABAergic postsynaptic densities, based on their insolubility in Triton X-100 at 4°C. This fraction is enriched in GABA_ARs and gephyrin, proteins that concentrate at the GABAergic postsynaptic complex. Gephyrin is a cytoplasmic scaffold protein that concentrates at the PSD of the glycinergic and GABAergic synapses. Gephyrin binds to microtubules and is essential for the postsynaptic clustering of the glycine receptors and is involved in the clustering of some GABA_ARs (Feng et al. 1998; Essrich et al. 1998; Kneussel et al. 1999, 2001; Christie et al. 2002a; Levi et al. 2004).

We are also showing that there is another pool of GABA_ARs that resists solubilization with Triton X-100 at 4°C and is associated with detergent-resistant membranes or DRMs (Brown and Rose 1992). DRMs have been operationally defined equivalent of lipid rafts (Brown and London 2000). The DRMs, like lipid rafts, are enriched in cholesterol and sphingomyeline, and contain lipid-anchored and some integral proteins (Schroeder et al. 1994; Brown and Rose 1992; Brown and London 1997; London and Brown 2000). Lipid rafts play a coordinating role in several signal transduction pathways (Simon and Toomre 2000). Several neurotransmitter receptors have been shown to be associated to lipid rafts (Bruses et al. 2001; Zhu et al. 2006; Becher et al. 2001; Suzuki et al. 2001; Hering et al. 2003; Dalskov et al. 2005).

In this communication we show that in both brain and hippocampal cultured neurons there are two pools of Triton X-100-insoluble GABA_ARs: one that concentrates in the postsynaptic GABAergic complex, and does not seem to be associated with DRMs or lipid rafts and another one that is non-synaptic and is associated with DRMs and lipid rafts.

MATERIALS AND METHODS

Antibodies

Several anti-GABA_AR subunit antibodies were raised and affinity-purified in our laboratory: rabbit anti-rat $\alpha 1$ (to amino acids a.a. 1–15), rabbit anti-rat $\alpha 2$ (to a.a. 417–423), rabbit anti-rat $\alpha 3$ (to a.a. 1–13), and guinea pig or rabbit anti-rat $\gamma 2$ (to a.a. 1–15). We used these anti- $\gamma 2$ antibodies for immunofluorescence or immunogold. The mouse monoclonal antibody (mAb) anti- $\beta 2/3$ (clone 62-3G1) was also generated in our laboratory to the affinity-purified bovine GABA_ARs. The specificities of the GABA_AR subunit antibodies have been determined previously by ELISA, immunoblotting, displacement by antigenic peptide, rat brain immunohistochemistry, immunofluorescence of hippocampal cultures and of HEK 293 cells transfected with various subunits, and in some knock out and knock down mutant mice (De Blas et al. 1988; Vitorica et al. 1988; Moreno et al. 1994; Miralles et al. 1999; Homanics et al. 1999; Christie et al. 2002a, b, 2006; Riquelme et al. 2002; Charych et al. 2004a, b; Li et al. 2005a, b; Chandra et al. 2005; Serwanski et al. 2006). Rabbit anti- $\beta 2$ and anti- $\beta 3$ antibodies were from Chemicon (Temecula, CA). The rabbit anti- $\gamma 2$ used in the immunoblots was from Alpha Diagnostic (San Antonio, TX). Three rabbit anti- δ antibodies were used. One was from

Phosphosolutions (Aurora, CO), another one from Novus Biologicals (Littleton, CO), and a third one was an affinity-purified anti- δ antibody raised in our laboratory to a.a. 390–402 of the rat δ subunit. The rabbit anti-gephyrin antiserum or affinity purified-antibodies used in the immunoblots were prepared as previously described (Kawasaki et al 1997), and the mAb to gephyrin (clone 7a) used in immunofluorescence and EM immunogold was from Synaptic systems (Gottingen, Germany) or Cedarlane Laboratories (Ontario, Canada). Other antibodies used were mAb to Thy-1 (clone OX7, Sera Lab, Loughborough, Leicestershire, England), mAb to Flotillin-1 (clone 18, BD Transduction Laboratories, San Diego, CA), mAb to Transferrin Receptor (clone H68.4, Zymed, San Francisco, CA), mAb to PSD-95 (Clone K28/43, Upstate Biotechnology Inc, Lake Placid, NY), and rabbit anti-Caveolin-1 (Santa Cruz Biotechnology, Santa Cruz, CA). For immunoblots, goat anti-rabbit or anti-mouse IgG antibodies and rabbit or mouse peroxidase-anti-peroxidase complexes were from MP Biomedicals (Aurora, OH). Fluorophore-labeled or colloidal gold-labeled species-specific anti-IgG secondary antibodies made in donkey were from Jackson ImmunoResearch Laboratories (West Grove, PA).

Subcellular fractionation of the rat brain and preparation and fractionation of the “One Triton PSD” fraction

The preparation of the crude mitochondrial/synaptosomal P2, synaptosomal SYN and “One Triton PSD” fractions was done according to Carlin et al. (1980). Three-month-old Sprague-Dawley male rats were used in these and other fractionation experiments. All steps were carried out at 4°C and all the fractions were suspended in 50 mM Tris-HCl pH 7.4, aliquoted and stored at –70°C. In this procedure, the synaptosomal fraction (SYN) was collected between the 1.0 M and 1.2 M sucrose interphase after sedimentation through a discontinuous sucrose gradient. This fraction was suspended in 0.32 M sucrose by dilution with 6 mM Tris-HCl pH 8.1, solubilized with 0.5% Triton X-100 for 15 minutes and centrifuged at 32,800× g for 1 hour. The resulting pellet is the “One Triton PSD” fraction, named by others (Walikonis et al. 2000). The “One Triton PSD” pellet was suspended in 0.32 M sucrose and loaded onto a continuous sucrose gradient 0.32–2.0 M in 1 mM NaHCO₃, centrifuged at 201,800× g for 16 hours, and fractionated in 1 ml fractions. Sucrose density was measured by refractometry. Cholesterol was measured by using the Amplex Red cholesterol assay kit (Invitrogen, Carlsbad, CA) following the manufacturer’s instructions. Immunoblots were done according to De Blas and Cherwinski (1983).

The synaptosomal plasma membrane (SPM) fraction was prepared after lysis of the crude synaptosomal fraction followed by flotation by centrifugation in a discontinuous sucrose gradient as described by Jones and Matus (1974).

Isolation of the DRM fraction from SPM and cholesterol depletion

The SPM fraction (4 mg protein) in 50 mM Tris-HCl pH 7.4 was centrifuged at 16,000× g for 10 min at 4°C and the pellet was suspended in 2 ml of TNE buffer (1% Triton X-100 in 50 mM Tris-HCl, pH 7.4, 150 mM NaCl, 5 mM EDTA, 1 mg/L Leupeptin, 0.1 mM PSMF) at a protein to detergent ratio of 1:5 (w/w). The mixture was incubated on ice for 20 min, followed by homogenization with 3 strokes in a hand-operated glass/glass Dounce homogenizer, incubation on ice for 10 min, and centrifugation at 125,000× g for 30 mins. The Triton X-100 soluble supernatant (TS) was collected, and the Triton X-100 insoluble pellet (TI) was suspended in TNE buffer (50 mM Tris-HCl pH 7.4, 150 mM NaCl, 5 mM EDTA, 1 mg/L Leupeptin, 0.1 mM PSMF) at 4°C. Alternatively, after extraction of SPM with TNE buffer, the sample was mixed with the same volume of 80% (w/v) sucrose and 4 ml were overlaid with 6ml of 35% (w/v) and 2ml of 5% (w/v) sucrose, respectively. The upper two layers contained 0.05% Triton X-100. After flotation by centrifugation at 18,800× g for 16 hours at 4°C, the DRM fraction was collected at the 5%/35% sucrose interphase. For cholesterol depletion, the SPM fraction was incubated with or without 0.5% saponin in TNE buffer on ice

for 20 minutes. The cholesterol-depleted SPM fraction was subjected to the aforementioned extraction with TNEX, followed by flotation by centrifugation in the 5%/35%/40% discontinuous sucrose gradient. After centrifugation, the gradient was fractionated in 12 fractions, 1 ml each.

Radioligand binding assays

For [³H]Flunitrazepam ([³H]FNZ) binding, 100 µg of protein of each fractions were incubated in triplicate with 10 nM [³H]FNZ in 50 mM Tris-HCl, pH 7.4, in a final volume of 500 µl for 30 min at 4°C. Nonspecific binding was determined by displacement with 10 µM clonazepam. For [³H]Muscimol ([³H]MUS) binding assay, all fractions were washed thoroughly with 50 mM Tris-HCl, pH 7.4 buffer by centrifugation at 34,500× g five times to eliminate endogenous GABA. Each fraction (150 µg protein) was incubated in triplicate with 40 nM [³H]MUS in 50 mM Tris-HCl, pH 7.4, for 30 minutes at 4°C in the dark. Non-specific binding was determined in the presence of 40 µM GABA. At the end of the incubation periods, 1.6 mg bovine γ-globulin and 0.32 ml of 30% polyethylene glycol (PEG) were added, incubated for 5 min at 4°C, and filtered through 24 mm Whatman glass-microfiber GF/B filters that had been previously washed with 8% PEG. The filters were washed twice with 5 ml of ice-cold 8% PEG, dried and radioactivity was determined with a liquidscintillation counter.

Immunofluorescence of hippocampal cultures and Triton X-100 extraction

Low-density hippocampal cultures were prepared according to Goslin et al. (1998) as described elsewhere (Christie et al. 2002a, b; Christie and De Blas 2003). Briefly, dissociated neurons from embryonic day 18 Sprague-Dawley rat hippocampi were plated at low density (3,000–8,000 cells per 18 mm diameter circular coverslip) and maintained in glial cell conditioned medium for 19–22 days. Immunofluorescence on fixed or live cells has been described elsewhere (Christie and De Blas 2002, Christie et al. 2006). Briefly, for fixed-cell labeling (no-capping), neurons on coverslips were fixed in 4% paraformaldehyde, 4% sucrose in phosphate buffered saline (PBS) for 15 minutes at room temperature (RT) followed by cell permeabilization with 0.25% Triton-X-100 in PBS for 5 minutes and incubation with 5% donkey serum in PBS for 30 minutes. Fixed and permeabilized neurons were incubated with a mixture of the rabbit anti-GABA_AR antibody and mAb to gephyrin for 2 hours. After washing, cells were incubated with a mixture of species-specific secondary anti-rabbit and anti-mouse IgG antibodies, made in donkey and conjugated to Texas Red or FITC fluorophores, in 0.25% Triton X-100 in PBS for 1 hour at RT. The coverslips were washed with PBS and mounted using Prolong anti-fade mounting solution (Invitrogen, Carlsbad, CA).

For “one-step capping”, the procedure has been described elsewhere (Christie et al. 2006). Briefly, live cells were surface-labeled by incubation with the primary rabbit or guinea pig anti-GABA_AR antibody (0.5–2 µg/ml) in culture medium at 37°C for 45 minutes in a 5% CO₂ atmosphere. After washing the cells were fixed and permeabilized, as described above, followed by incubation with mouse mAb to gephyrin or to Thy-1 and by incubation with a mixture of the fluorophore-labeled secondary antibodies. For Triton X-100 extraction, after “one-step capping” with the primary antibody to GABA_ARs, the cultures were incubated with 0.5% Triton X-100 in 20 mM phosphate buffer, pH 7.4, (PB) at 4°C or 37°C for 10 min. Alternatively, neurons were cholesterol-depleted, by incubation with 5 mM methyl-β-cyclodextrin (MβCD) at 37°C for 20 min, prior to incubation with 0.5% Triton X-100 in 20 mM PB at 4°C. The cells were fixed and processed as described above.

For “two-step capping”, we followed the procedure described elsewhere (Levi et al. 2004, Christie et al. 2006). Briefly, live cells were sequentially incubated with the primary rabbit anti-α1 antibody for 45 min and the fluorophore-conjugated secondary donkey anti-rabbit IgG antibody (10 µg/ml) for 30 min at 37°C, followed by fixation, permeabilization, and incubation

with the mAb to gephyrin and subsequent fluorophore-conjugated secondary donkey anti-mouse IgG antibody. Detergent extraction after “two-step capping” was done by incubating the cultures with 0.5% Triton X-100 at 4°C before the fixation step, as for “one-step capping”. Some cultures were treated with M β CD prior to Triton X-100 extraction as for “one-step capping”.

Image acquisition and analysis

Fluorescence images were collected using a 60 \times pan-fluor objective on a Nikon Eclipse T300 microscope with a Sensys KAF 1401E CCD camera, driven by IPLab 3.0 (Scanalytics, Fairfax, VA) acquisition software. For qualitative analysis, images were processed with Photoshop 7.0 (Adobe, San Jose, CA) as described elsewhere (Li et al. 2005a, b, Christie et al. 2006).

Quantification of GABA_AR and gephyrin clusters

For each determination, 30 to 40 dendrites were randomly selected from 15–20 cells (2 dendrites/cell) from 3 to 4 separate experiments. A 100 μ m-long segment from each dendrite was used for quantification. We did not quantify the cluster density in terms of dendritic area because detergent treatment of non-fixed cells led to somewhat thinner dendrites than when the detergent extraction was done after fixation (see also Allison et al. 2000). To determine the number of clusters per 100 μ m of dendritic length, the maximum intensities of the fluorophore channel were normalized and the low intensity and diffuse non-clustered background fluorescence signal seen in the dendrites was subtracted. Cluster co-localization in two different fluorescence channels was determined by overlaying the images. A cluster in a fluorescence channel was considered to co-localize with a cluster in the other channel when >66% of surface of one of the clusters overlapped with the other cluster. Measurements of cluster size were performed using IPLabs 3.5 software. The 8-bit images were segmented, on the basis of fluorescence intensity levels, to create a binary mask that maximized the number of clusters and minimized the coalescence of individual clusters. Values were averaged per 100 μ m and reported as mean \pm standard error of the mean (SEM). Statistical comparison was made by Student's t test.

Electron microscopy immunogold

It was done according to the procedure of Hunt et al. (1996). Briefly, subcellular fractions were fixed in 4% paraformaldehyde, 0.1% glutaraldehyde and suspended in 1.5% low melting point agarose. The solidified agarose was cut into \sim 1mm³ blocks, and the blocks were incubated with a single antibody (or a mixture of two primary antibodies), followed by incubation with a colloidal gold-conjugated secondary antibody (6 nm or 12 nm diameter for Fig. 1) or a mixture of two colloidal gold-conjugated secondary antibodies (10 nm and 18 nm diameter for Fig. 2). The blocks were then incubated in 1% glutaraldehyde in 0.12 M sodium phosphate buffer, pH 7.4 for 15 min, followed by incubation in 1% osmium tetroxide for 30 min at RT, and 2% uranyl acetate for 30 min at RT, dehydration in a series of ethanol, infiltration, and embedding in Spurr's resin at RT. Thin sections were cut on a Reichardt ultramicrotome and collected on Formvar-coated grids. The sections were contrasted by incubation with 2% uranyl acetate and 0.3% lead citrate, 3 min each, and photographed in a Philips 300 transmission electron microscope.

RESULTS

GABA_ARs and gephyrin are enriched in the “One Triton PSD” fraction

Two types of synapses, with very distinct morphology, are present in the brain. These are excitatory Gray's type-I and inhibitory Gray's type-II synapses. Synaptosomes are pinched-off presynaptic terminals that are formed after homogenization of the brain. Some

synaptosomes have an attached postsynaptic membrane with a prominent postsynaptic density derived from type-I synapses (Cotman et al. 1974; Matus et al. 1980; Hunt et al. 1996). Since we are interested in studying GABA_A receptors and GABAergic synapses, we have investigated whether type-II GABAergic synapses are also present in the classical synaptosomal preparation. We have found that this is the case (Fig. 1A–C). This fraction had synaptosomes with type-II synaptic morphology whose postsynaptic elements became decorated with gold particles corresponding to antibodies to the GABAergic postsynaptic markers $\alpha 1$ subunit of the GABA_AR (Fig. 1A) or gephyrin (Fig. 1B–C) as shown by electron microscopy (EM) immunogold. The type-II synaptic morphology (Fig. 1A–C) included light postsynaptic density and flattened synaptic vesicles in the presynaptic axon terminal. The gold particles corresponding to anti- $\alpha 1$ GABA_AR subunit were located at the extracellular side of the vesicularized postsynaptic membrane (Fig. 1A arrowheads), as expected for an antibody that recognizes an epitope localized at the extracellular N-terminus of $\alpha 1$. In contrast, the gold particles corresponding to an antibody that recognizes the postsynaptic cytoplasmic protein gephyrin were found at the expected cytoplasmic side of the postsynaptic membrane (Fig. 1B–C arrowheads). Note in Fig. 1A that the colloidal gold-conjugated secondary antibody could not penetrate the intact part of the synaptic cleft, thus preventing the immunolabeling of the GABA_ARs present in this part of the cleft. No permeabilizing detergent was used in the experiments aiming to preserve the morphological integrity of the structures. However, gold immunolabeling with anti- $\alpha 1$ occurred in the adjacent region of the postsynaptic membrane that became separated from the presynaptic terminal after homogenization. These results indicate that the classical synaptosomal preparation also contains synaptosomes, derived from GABAergic synapses, with type-II morphology.

Other groups have developed methods for the purification of type-I postsynaptic densities (Cotman et al. 1974; Carlin et al. 1980; Walikonis et al. 2000) by taking advantage of the insolubility of type-I PSDs in Triton X-100. One of these fractions, enriched in type-I PSDs, has been called “One Triton PSD” fraction (Walikonis et al. 2000). We have now explored whether the “One Triton PSD” fraction is also enriched in GABA_A receptors and gephyrin, two proteins that concentrate at the GABAergic postsynaptic complex. If so, it would indicate that type-II PSDs are also present in the “One Triton PSD” fraction.

Immunoblots (Fig. 1G) showed that there was a progressive enrichment in various GABA_AR subunits ($\alpha 1$, Mr = 51000; $\alpha 2$, Mr = 52000; $\beta 2/3$, Mr = 54000–57000 and $\gamma 2$, Mr = 46000) and the two main isoforms of gephyrin (Mr = 93000–98000) from the crude mitochondrial/synaptosomal fraction (P2) to synaptosomal (SYN) to synaptosomal plasma membrane (SPM) to “One Triton PSD” (PSD) fraction. These results indicate that the “One Triton PSD” fraction is enriched in GABAergic receptors and gephyrin, two markers of GABAergic synapses. As others have shown (Hunt et al. 1996; Kennedy 1997), the “One Triton PSD” fraction, as expected, was also enriched in PSD-95 (Mr = 95000), a marker for type-I PSDs. We do not know if the faint protein band (Mr = 47000), which migrates slightly faster than the main $\alpha 2$ protein band (Mr = 52000) and that is absent from the PSD fraction, is an immature form of the $\alpha 2$ subunit (ie. a non-glycosylated peptide) that is not present at synapses or a proteolytic fragment that is extracted by Triton X-100.

The specific activities of [³H]Muscimol ([³H]MUS) and [³H]Flunitrazepam ([³H]FNZ) binding to the GABA- and benzodiazepine-binding sites of the GABA_ARs (located in the boundary between the α/β and α/γ subunits respectively) also showed enrichment in the PSD fraction over the other fractions (Fig. 1H). The specific activity of [³H]MUS binding to PSD (1.75 ± 0.09 pmol/mg protein, mean \pm SEM) was not significantly different from that of SPM (1.36 ± 0.16 pmol/mg protein, $p = 0.10$), but was significantly higher than SYN (0.52 ± 0.08 pmol/mg protein, $p = 0.0005$) and P2 (0.51 ± 0.04 pmol/mg protein, $p = 0.0002$). For [³H]FNZ binding, the specific activity of PSD (1.70 ± 0.13 pmol/mg protein) was not significantly

different from that of SPM (1.39 ± 0.14 pmol/mg, $p = 0.15$), but was significantly higher than P2 (0.50 ± 0.04 pmol/mg protein, $p = 0.0001$) and SYN (0.66 ± 0.19 pmol/mg protein, $p = 0.0054$) (Fig. 1H). The radioligand-binding activity indicates that the GABA_ARs in the PSD and other fractions are fully assembled, since only assembled receptors have the binding sites for the two radioligands. This could not be ascertained by the immunoblotting assay, which does not differentiate between assembled and non-assembled GABA_AR subunits.

The EM immunogold of the PSD fraction showed that gold particles representing antibodies to $\alpha 1$ (arrowheads, Fig. 1D) or gephyrin (arrowheads, Fig. 1E) decorated amorphous structures of medium electron-density (empty arrows, Fig. 1D and E) but not the high electron-dense structures with type-I PSD morphology (filled arrows, Fig. 1D and E). The structures with the type-I PSD morphology (filled arrows, Fig. 1F) were frequently decorated with gold particles corresponding to anti-PSD-95 antibody (arrowhead, Fig. 1F). The type-II PSDs could only be identified by immunogold. Contrary to type-I PSDs, the type-II PSDs could not be identified just by morphological criteria.

The results are consistent with the notion that the “One Triton PSD” fraction is enriched in fully assembled GABA_ARs and gephyrin and that this fraction contains not only the prominent and electron-dense type-I glutamatergic PSDs, but also the more amorphous and less electron-dense type-II GABAergic PSDs (empty arrows Fig. 1D, E), structures that are decorated with the anti-GABA_ARs and anti-gephyrin antibodies (arrowheads Fig. 1D, E).

Two pools of GABA_ARs are present in the “One Triton PSD” fraction: one associated with DRMs and another one associated with GABAergic postsynaptic densities

Although the “One Triton PSD” fraction is enriched in PSDs, it could also contain DRMs and lipid rafts, which are also insoluble in Triton X-100 at 4°C (Suzuki 2002). To ascertain whether the GABA_ARs present in this fraction are associated with DRMs, or with GABAergic PSDs, or with both, we subjected the PSD fraction to sedimentation in a 0.32–2.0 M continuous sucrose gradient. After centrifugation, the main protein peak concentrated in fractions 9–11 at $\rho = 1.18$ – 1.23 g/ml (Fig. 2A). A smaller protein peak was found in fraction 4 ($\rho = 1.08$ g/ml). Immunoblots containing the same amount of protein per lane (Fig. 2B) showed that the low-density fractions 3–5 ($\rho = 1.06$ – 1.09 g/ml) were enriched in the GABA_AR α and β subunits and in the lipid raft markers Thy-1 ($M_r = 21000$), Caveolin-1 ($M_r = 24000$) and Flotillin-1 ($M_r = 48000$), as well as in cholesterol (Fig. 2A), indicating that fractions 3–5 are enriched in DRMs and lipid rafts. The extrasynaptic GABA_AR δ subunit ($M_r = 54000$) also concentrated in the low-density fractions, although the intensity of the δ subunit protein band was considerably weaker than that of the various α and β GABA_AR subunits. This result, corroborated with three different anti- δ subunit antibodies, was expected since it is known that the δ subunit is expressed at low levels in the rat forebrain (see discussion). For this reason we did not further study the δ subunit in our forebrain preparations. Transferrin receptor (TrsfR $M_r = 95000$), which is absent from DRMs and lipid rafts (Harder et al. 1998; Jing et al. 1990), concentrated in fractions 8–12 (Fig. 2B). The various α and β GABA_AR subunits were also present in fractions 8–12, although to a lesser extent than in the fraction enriched in lipid raft markers. The results in Fig. 2B were normalized to the same amount of protein per lane, to show the relative enrichment in each fraction. Even when considering the distribution of the total amount of each GABA_AR subunit in the gradient, the DRM fractions 3–5 contained 38% of all $\alpha 1$, 28% of $\alpha 2$, 21% of $\alpha 3$, 49% of $\beta 2$, 46% of $\beta 3$ and 33% of all the δ present in the gradient. These results showed that in spite of the relatively small amount of protein present in these fractions, they concentrated a significant proportion of all GABA_ARs. In contrast to the α , β and δ subunits, only 8% of all $\gamma 2$ or 12% of all gephyrin were found in the DRM fractions 3–5. The EM immunogold labeling of combined DRM fractions 3 and 4, enriched in lipid raft markers, showed vesicular structures (Fig. 2C, crossed arrows) similar to the low-density DRMs

observed by others at the EM level in DRM preparations from various tissue sources (Suzuki et al. 2002; Besshoh et al. 2005). In addition, we have found that these vesicles were frequently decorated with anti- β 2/3 (empty arrowheads) and anti- α 1 (filled arrowheads) immunogold particles (Fig. 2C), indicating that the vesicular DRMs contain α 1- and β 2/3-GABA_AR, subunits that are enriched in the DRMs as immunoblots show (Fig. 2B).

The γ 2 GABA_AR subunit and gephyrin were enriched in fractions 6–8 (Fig. 2B), which contained 30% of all the γ 2 and 38% of all the gephyrin in the gradient. The PSD-95, marker for type-I PSDs, mainly distributed in fractions 8–12. These fractions contained 83% of all the PSD-95 in the gradient. Since the γ 2 GABA_AR subunit and gephyrin are postsynaptic GABAergic markers (Craig et al. 1996; Giustetto et al. 1998; Kneussel et al. 1999; Christie et al. 2002b; van Rijnsoever et al. 2005), and since the γ 2 subunit is essential for the postsynaptic clustering of GABA_ARs (Essrich et al. 1998; Schweizer et al. 2003; Li et al. 2005b), we concluded that fractions 6–8 ($\rho = 1.10$ – 1.15 g/ml) are enriched in type-II PSDs and fractions 8–12 ($\rho = 1.15$ – 1.24 g/ml) are enriched in type-I PSDs, with some overlap in the enrichment of the two types, particularly in fraction 8. Thus, type-II PSDs can be partially separated from type-I PSDs in fractions 6 and 7 due to the lower density of the former, which might be related to the more amorphous and less electron-dense quality of type-II PSDs. It is worth noting that fractions 9–12 still carry a considerable amount of the total γ 2 subunit (58%) and gephyrin (51%). Nevertheless, fractions 9–12 show no enrichment (per mg of protein) in these GABAergic markers due to the high concentration in these fractions of proteins insoluble in Triton X-100. In contrast, fractions 9–12 are highly enriched (per mg of protein) in PSD-95.

The EM of fraction 7 (Fig. 2D–F), which is enriched in GABAergic postsynaptic markers γ 2-GABA_AR and gephyrin, showed the presence of both vesicular structures (crossed arrows) and amorphous material of medium electron-density (empty arrows). Double-label EM immunogold with the anti-GABAergic synaptic markers gephyrin (empty arrowheads) and γ 2 GABA_ARs (filled arrowheads) showed that the immunogold particles were associated with the amorphous structures of medium electron-density (empty arrows, Fig. 2D–F), reminiscent of the ones found in the “One Triton PSD” fraction (empty arrows, Fig. 1D and E), which were also immunolabeled with antibodies to GABA_ARs and gephyrin. Some electron-dense structures with type-I PSDs morphology (filled arrows, Fig. 2F) were also present in fraction 7, although they were not immunolabeled with anti-gephyrin or anti-GABA_ARs antibodies. The presence of some type-I PSDs in fraction 7 was consistent with the presence of some PSD-95 in this fraction, as shown by immunoblotting (Fig. 2B). The EM immunogold and the immunoblot data support the notion that in the “One Triton PSD” fraction, I) there are GABA_ARs that contain α and β but little or no γ 2 subunit that are associated to DRMs and lipid rafts, and II) there are GABA_ARs that contain α , β and γ 2 subunits associated with gephyrin that are present in the type-II GABAergic postsynaptic complex.

A pool of the GABA_ARs present in synaptosomal plasma membranes is localized in DRMs

To further confirm the association of a pool of GABA_ARs with DRMs and lipid rafts, a synaptosomal plasma membrane fraction (SPM) was extracted with 1% Triton X-100 at 4°C for 30 minutes, a more stringent extraction than the one called for in the procedure for the preparation of the “One Triton PSD” fraction, which was done with 0.5% Triton X-100 at 4°C for 15 minutes. Immunoblots showed that a significant amount of GABA_ARs and gephyrin were present in both the Triton-soluble (TS) and Triton-insoluble (TI) fractions (Fig. 3A). The lipid raft markers Thy-1, Caveolin-1 and Flotillin-1 mainly concentrated in the TI fraction, while TrsfR mainly concentrated in the TS fraction. To find out whether some GABA_ARs from the TI fraction were associated to DRMs, we isolated the low-density DRM fraction, which is enriched in lipid rafts, after incubation of the SPM fraction with 1% Triton X-100 at 4°C followed by flotation by centrifugation in a discontinuous sucrose gradient, as described in the

methods section. Compared to TI, the DRM fraction showed substantial enrichment of α and β GABA_AR subunits, as well as of the lipid raft markers Thy-1, Caveolin-1 and Flotillin-1 (Fig. 3A). However, DRM showed considerably less γ 2 GABA_AR subunit, gephyrin and TrsfR than TI (Fig. 3A). The specific [³H]MUS and [³H]FNZ binding to GABA_ARs indicated that the GABA_AR subunits present in these fractions were incorporated into assembled GABA_AR (Fig. 3B). The specific [³H]MUS binding activity of DRM (1.75 ± 0.07 pmol/mg protein) was significantly higher than that of TI (1.12 ± 0.06 pmol/mg protein, $p = 0.0009$) or TS (0.55 ± 0.08 pmol/mg protein, $p = 0.0001$) fractions. The difference in [³H]MUS binding between TI and TS was also significant ($p = 0.0014$). However, [³H]FNZ binding activity of DRM (1.55 ± 0.30 pmol/mg protein) was smaller than that of TI (2.25 ± 0.17 pmol/mg protein, $p = 0.004$) and similar to that of TS (1.47 ± 0.15 pmol/mg protein, $p = 0.72$). Regarding recovery, 32% of [³H]MUS and 11% of [³H]FNZ binding in SPM was recovered in the DRM fraction. These binding and immunoblot results show that the GABA_ARs in the DRM fraction, enriched in lipid raft markers, are mainly formed by α and β subunits and bind [³H]MUS. Fewer GABA_ARs in this fraction also contain the γ 2 subunit and consequently DRM shows less [³H]FNZ binding than the TI fraction.

Cholesterol depletion disrupts the association of GABA_ARs with lipid rafts

Saponin is a detergent that sequesters cholesterol. It has frequently been used to disrupt lipid rafts (Cerneus et al. 1993, Becher et al. 2001). We treated the SPM fraction with 0.5% saponin before extraction with 1% Triton X-100 at 4°C and fractionation. Fig. 4 shows that when SPM was not treated with saponin (control, C), the α 1 and β 3 GABA_ARs subunits, as well as the lipid raft markers Thy-1, Flotillin-1 and Caveolin-1, were found in the low-density fractions (3 and 4). There was very little γ 2 GABA_AR subunit and no TrsfR could be detected in the low-density fractions 3 and 4. It is worth noticing that gephyrin also showed some association with fraction 4 of Fig. 2 and in the DRM fraction of Fig. 3, indicating that there is some association of gephyrin with DRMs and lipid rafts (Fig. 4), although the bulk of gephyrin concentrated at the heavier 9–12 fractions. After treatment of SPM with saponin (S), the α 1 and β 3 GABA_AR subunits and gephyrin (and the very small amount of γ 2 subunit seen in the control) were no longer present in fractions 3 or 4. Moreover, all the proteins, including the lipid raft markers (i.e. Thy-1, Flotillin-1 and Caveolin-1) migrated to the heavier density fractions 7–12 (Fig. 4). These experiments further support the notions that I) the Triton X-100 insoluble GABA_ARs present in the low-density fractions are associated with lipid rafts; and II) the majority of the GABA_ARs associated to lipid rafts contain α and β subunits, but relatively few contain γ 2.

In cultured hippocampal neurons, most GABA_AR clusters that contain the γ 2 subunit are not associated to lipid rafts

Others have shown that in hippocampal cultures, β 2/3-containing GABA_AR clusters and gephyrin clusters are resistant to 0.5% Triton X-100 extraction at 4°C (Allison et al. 2000). We have now investigated whether the resistance of GABA_AR clusters to Triton X-100 extraction is in part due to their association with lipid rafts and whether it depends on the subunit composition.

Live cultured hippocampal neurons (19–22 DIV) were incubated with an antibody to an extracellular epitope of the γ 2 subunit, to label the cell-surface GABA_ARs under conditions that produce no apparent antibody internalization (Levi et al. 2004). Cell permeabilization with 0.25% Triton X-100 was done after cell fixation (TX-AF). This was followed by incubation with the anti-gephyrin antibody to label the intracellular gephyrin. Under these conditions, Triton X-100 permeabilizes the cells but it does not extract the fixed GABA_ARs. In other cultures, neurons that have been surface-labeled with the anti- γ 2 GABA_AR antibody were extracted and permeabilized with 0.5% Triton X-100 at 4°C before fixation (TX-BF). After

fixation, the cells were incubated with the anti-gephyrin antibody. As shown in Fig. 5A, E, M and N, Triton X-100 at 4°C did not significantly extract the $\gamma 2$ -containing GABA_AR ($\gamma 2$ -GABA_AR) clusters from the surface of dendrites as shown by analysis of the cluster number (57.0 ± 1.7 in TX-BF compared with 59.2 ± 2.5 in TX-AF, $p = 0.48$) or the cluster size ($0.087 \pm 0.003 \mu\text{m}^2$ in TX-BF compared with $0.089 \pm 0.003 \mu\text{m}^2$ in TX-AF, $p = 0.76$).

The number and size of gephyrin clusters were also unaffected by Triton X-100 extraction at 4°C (50.7 ± 1.6 in TX-BF compared to 52.0 ± 2.2 in TX-AF, $p = 0.62$ and $0.119 \pm 0.004 \mu\text{m}^2$ in TX-BF compared with $0.106 \pm 0.007 \mu\text{m}^2$ in TX-AF, $p = 0.08$), as shown in Fig. 5B, F, M and N. Moreover, quantification of the co-localization of $\gamma 2$ and gephyrin clusters showed that in TX-AF neurons, $77.2 \pm 1.6 \%$ of $\gamma 2$ clusters co-localized with gephyrin and $87.8 \pm 1.2 \%$ of gephyrin clusters co-localized with $\gamma 2$. After extraction with Triton X-100 at 4°C (TX-BF), the co-localization of $\gamma 2$ and gephyrin clusters remained unchanged: $76.1 \pm 1.7 \%$ ($p = 0.60$) of $\gamma 2$ clusters were co-localized with gephyrin and $85.6 \pm 1.5 \%$ ($p = 0.26$) of gephyrin clusters were co-localized with $\gamma 2$. Thus, the number and size of $\gamma 2$ -GABA_AR clusters, gephyrin clusters and their co-localization remained unchanged by Triton X-100 extraction at 4°C.

Cholesterol depletion was done by incubating the cultures with 5mM methyl- β -cyclodextrin (M β CD) after live cell incubation with the antibodies and prior to the extraction with Triton X-100 at 4°C before fixation (TX-BF). Cholesterol depletion had no significant effect on the insolubility of $\gamma 2$ -GABA_AR clusters in Triton X-100 at 4°C, either in the number (55.7 ± 1.2) compared to the TX-BF (57.0 ± 1.7 , $p = 0.51$) or to the TX-AF (59.2 ± 2.5 , $p = 0.51$) or in the size ($0.094 \pm 0.003 \mu\text{m}^2$) compared to the TX-BF ($0.087 \pm 0.003 \mu\text{m}^2$, $p = 0.09$) or to the TX-AF ($0.089 \pm 0.003 \mu\text{m}^2$, $p = 0.15$), as shown in Fig. 5A, E, I, M and N. Neither cholesterol depletion by M β CD had an effect on the insolubility of gephyrin clusters in Triton X-100 at 4°C, nor significantly changed the number of clusters (49.2 ± 1.4) compared to TX-BF (50.7 ± 1.6 , $p = 0.46$), or compared to TX-AF (52.0 ± 2.2 , $p = 0.26$), or the size ($0.116 \pm 0.003 \mu\text{m}^2$) compared to the TX-BF ($0.119 \pm 0.004 \mu\text{m}^2$, $p = 0.48$) or to the TX-AF ($0.106 \pm 0.007 \mu\text{m}^2$, $p = 0.18$), of gephyrin clusters, as shown in Fig. 5B, F, J, M and N. Treatment with M β CD prior to extraction with Triton X-100 at 4°C did not affect the co-localization of $\gamma 2$ and gephyrin clusters ($76.0 \pm 1.7 \%$ of $\gamma 2$ clusters co-localized with gephyrin, $p = 0.56$ and $86.1 \pm 1.4 \%$ of gephyrin clusters co-localized with $\gamma 2$, $p = 0.51$), compared with TX-BF ($76.1 \pm 1.7 \%$ of $\gamma 2$ clusters co-localized with gephyrin and $85.6 \pm 1.5 \%$ of gephyrin clusters co-localized with $\gamma 2$).

The persistent insolubility of the $\gamma 2$ -GABA_AR or gephyrin clusters in Triton X-100 at 4°C even after cholesterol depletion indicates that these clusters are not associated with lipid rafts. The resistance of $\gamma 2$ -GABA_AR and gephyrin clusters to Triton X-100 extraction is likely due to their association with the cytoskeleton and scaffold proteins, as in a way being reminiscent of the insolubility in Triton X-100 of the GABA_ARs and gephyrin that are present at the GABAergic type-II PSD complex in brain synapses, as shown above.

Some $\alpha 1$ -GABA_AR clusters in cultured hippocampal neurons are associated with lipid rafts

We did similar live-cell incubation experiments with an anti- $\alpha 1$ GABA_AR subunit antibody that recognizes an extracellular epitope. The observed $\alpha 1$ -GABA_AR clusters resisted solubilization in Triton X-100 at 4°C. Quantification showed that there was no significant difference between TX-BF and TX-AF neurons in the number (57.5 ± 2.0 vs. 58.3 ± 2.3 , $p = 0.79$) and size ($0.092 \pm 0.003 \mu\text{m}^2$ vs. $0.095 \pm 0.003 \mu\text{m}^2$, $p = 0.49$) of the $\alpha 1$ -GABA_AR clusters, as shown in Fig. 5C, G, M and N. Nevertheless, when cultures were treated with M β CD to deplete cholesterol from the membrane, a significant number of $\alpha 1$ -GABA_AR clusters became solubilized by Triton X-100 at 4°C, as shown by the reduction in the number of surface $\alpha 1$ -GABA_AR clusters after M β CD treatment (40.6 ± 1.8), compared to TX-BF or TX-AF cells

(57.5 ± 2.0 and 58.3 ± 2.3 respectively, $p < 0.0001$ in both cases), as shown in Fig. 5C, G, K, and M. The cluster size was not significantly affected in M β CD ($0.084 \pm 0.004 \mu\text{m}^2$) when compared to TX-BF ($0.092 \pm 0.003 \mu\text{m}^2$, $p = 0.095$), but decreased when compared to a TX-AF ($0.095 \pm 0.003 \mu\text{m}^2$, $p = 0.018$) as shown in Fig. 5C, G, K and N. Thus, approximately 30% of the $\alpha 1$ -GABA_AR clusters present on the membrane surface were extracted by Triton X-100 at 4°C after M β CD treatment, indicating that these clusters are associated with lipid rafts. Extraction of the cultures with 0.5% Triton X-100 at 37°C before fixation, condition in which lipid rafts are solubilized (Hering et al. 2003), led to a similar reduction in the number of $\alpha 1$ -GABA_AR clusters (not shown).

To test the efficiency of lipid rafts disruption by the M β CD treatment in these cultures, neurons were labeled with an antibody to Thy-1, a surface protein associated with lipid rafts, in double-label experiments with anti- $\alpha 1$. Thy-1 clusters were found on the soma and dendrites for TX-AF condition (Fig. 5D). When extraction with Triton X-100 at 4°C was done before fixation (TX-BF), the size of the Thy-1 clusters increased ($0.236 \pm 0.01 \mu\text{m}^2$, Fig. 5H and N) when compared to TX-AF ($0.111 \pm 0.005 \mu\text{m}^2$, $p < 0.0001$ Fig. 5D and N), apparently due to the coalescence of the smaller Thy-1 clusters induced by the detergent, while the number of Thy-1 clusters significantly decreased (83.8 ± 2.5 for TX-BF vs. 97.6 ± 4.0 for TX-AF, $p = 0.0044$) as shown in Fig. 5D, H and M. Compared to TX-BF neurons, cholesterol depletion with M β CD led to the solubilization by Triton X-100 at 4°C of the majority of Thy-1 clusters, as shown by a dramatic decrease in number (17.59 ± 1.05 , $p < 0.0001$ Fig. 5H, L and M) and size ($0.122 \pm 0.008 \mu\text{m}^2$, $p < 0.0001$ Fig. 5H, L and N) of the Thy-1 clusters when compared with TX-BF or TX-AF. These results show that I) the solubilization behavior of Thy-1 clusters in Triton X-100 at 4°C is the one expected for a protein associated with lipid rafts; and II) that the M β CD treatment is effective in disrupting lipid rafts in these cultures. The results also validate the assay used in Figs. 5 and 6 to study the association of clustered GABA_AR to lipid rafts.

Taken together, these results indicate that I) surface $\gamma 2$ -GABA_AR and gephyrin clusters resist extraction by Triton X-100 at 4°C. These clusters are not associated with lipid rafts; and II) about 30% of surface $\alpha 1$ -GABA_AR clusters are associated with lipid rafts. These clusters do not contain the $\gamma 2$ subunit and are not associated with gephyrin (see below).

In hippocampal cultures there is a pool of $\alpha 1$ -GABA_AR associated with lipid rafts that is diffusely distributed at the neuronal surface and that can be induced to cluster by antibody capping

We and others have shown that when living cells are incubated with a primary anti-GABA_AR antibody to an external epitope, it leads to the clustering (one-step capping) of the normally non-clustered GABA_AR that are present on the cell surface (Brunig et al. 2002; Levi et al. 2004; Christie et al. 2006). The capping of the non-clustered GABA_AR is further enhanced by sequential incubations of the live cells with the primary and secondary antibodies before fixation (two-step capping, Levi et al. 2004; Christie et al. 2006). When the cells were fixed, followed by permeabilization with Triton X-100 before incubation with the primary antibodies (TX-AF, no capping) in double-label immunofluorescence experiments, all the observed $\alpha 1$ -GABA_AR and gephyrin clusters (Fig. 6A–C) are the ones that are normally present in those neurons in the absence of antibody capping. These clusters have been thoroughly documented by us and others (Craig et al. 1996; Christie et al. 2002a, b; Christie and De Blas 2003). When the live cells were incubated with the anti- $\alpha 1$ antibody followed by fixation and treatment with Triton X-100 (one-step capping TX-AF, Fig. 6D and V), the number of $\alpha 1$ clusters (58.1 ± 2.0) was significantly higher than when TX-AF cells were not subjected to capping (48.4 ± 2.1 , $p = 0.0014$). Two-step capping induced the formation of a considerably larger number of $\alpha 1$ -

GABA_AR clusters (141.5 ± 7.5 , $p < 0.0001$, Fig. 6M and V) compared to no capping or one-step capping in the TX-AF cells.

The number of gephyrin clusters did not significantly change in TX-AF cells from no capping (47.8 ± 2.2 , Fig. 6B and V) to one-step capping with anti- $\alpha 1$ (51.3 ± 2.0 , $p = 0.23$, Fig. 6E and V). A very small increase in gephyrin cluster number was detected in two-step capping with anti- $\alpha 1$ (55.2 ± 2.7 , $p = 0.03$, Fig. 6N and V). In TX-AF cells the number of $\alpha 1$ -GABA_AR clusters colocalizing with gephyrin clusters (or the number of gephyrin clusters colocalizing with $\alpha 1$ -GABA_AR clusters, which is the same, Fig. 6V) remained unchanged in no-capping (40.1 ± 2.1) when compared with one-step capping with anti- $\alpha 1$ (39.3 ± 1.9 , $p = 0.79$), or with two-step capping (39.9 ± 3.2 , $p = 0.96$). Therefore the increased number of $\alpha 1$ -GABA_AR clusters induced by one-step or two-step antibody capping of the $\alpha 1$ -GABA_ARs that are normally diffusely distributed at the neuronal surface is not accompanied by simultaneous gephyrin capping (Levi et al. 2004; Christie et al. 2006).

Lipid rafts can be induced to coalesce by antibodies recognizing proteins associated with lipid rafts (Holowka et al. 2005). We have investigated whether the $\alpha 1$ -GABA_AR clusters associated with lipid rafts described above under one-step capping conditions (Fig. 5C, G, K) are the $\alpha 1$ -GABA_AR clusters normally existing in these neurons or the clusters whose formation was induced by one-step capping with anti- $\alpha 1$. We also tested the solubility of the clusters in Triton X-100 at 4°C before fixation after one-step and two-step capping. After one-step capping of live cells, extraction with Triton X-100 at 4°C before fixation (TX-BF) did not change the number of $\alpha 1$ -GABA_AR clusters (58.2 ± 2.1 vs. 58.1 ± 2.0 , $p = 0.96$) or the number of $\alpha 1$ -GABA_AR clusters colocalizing with gephyrin (40.3 ± 1.9 vs. 39.3 ± 1.9 , $p = 0.73$), when compared to one-step capping of live cells in which extraction with Triton X-100 was done after fixation (TX-AF, Fig. 6D–I and W). Although the number of $\alpha 1$ -GABA_AR clusters in the two-step capping experiments was considerably larger than in the one-step capping, there was no significant difference between two-step capping TX-AF and TX-BF in the number of $\alpha 1$ -GABA_AR clusters (141.5 ± 7.5 vs. 139.3 ± 5.5 respectively, $p = 0.82$) or the number of $\alpha 1$ -GABA_AR clusters colocalizing with gephyrin (39.9 ± 3.2 vs. 40.0 ± 2.1 respectively, $p = 0.98$), as shown in Fig. 6M–R and W. These experiments show that both the original $\alpha 1$ -GABA_AR clusters and the ones induced by one-step or two-step capping resist solubilization by Triton X-100 at 4°C.

After one-step capping (Fig. 6J–L and W) or two-step capping (Fig. 6S–U and W), treatment of the culture with M β CD for cholesterol depletion prior to Triton X-100 extraction before fixation, led to a significant reduction in the number of $\alpha 1$ -GABA_AR clusters compared to the corresponding TX-BF cultures not treated with M β CD (43.2 ± 1.8 vs. 58.2 ± 2.1 , $p < 0.0001$ for one-step capping and 57.0 ± 2.3 vs. 139.3 ± 5.5 , $p < 0.0001$ for two-step capping, respectively). The results (Fig. 6W) also showed that I) the original $\alpha 1$ -GABA_AR clusters, which are present in the absence of capping, co-localized with gephyrin and remained insoluble in Triton X-100 at 4°C after cholesterol depletion by M β CD and II) M β CD treatment facilitated the solubilization in Triton X-100 of the majority (93%) of the $\alpha 1$ -GABA_AR clusters whose formation was induced by one-step or two-step capping with anti- $\alpha 1$. In contrast, and in similar experiments, many (84%) of the $\gamma 2$ GABA_ARs whose cluster formation was induced by two step-capping with anti- $\gamma 2$ remained insoluble in Triton X-100 at 4°C even after cholesterol depletion by M β CD (not shown).

These experiments show that I) most of the $\alpha 1$ -GABA_AR clusters induced by one-step or two-step capping with the anti- $\alpha 1$ antibody are associated with lipid rafts, but are not associated with gephyrin; II) most of the γ -GABA_ARs whose clustering is induced by one-step or two-step capping with the anti- $\gamma 2$ antibody are not associated with lipid rafts and III) the normally existing $\alpha 1$ -GABA_AR or $\gamma 2$ -GABA_AR clusters that are present in the cultured hippocampal

neurons, are not associated to lipid rafts. Many of these clusters co-localize with gephyrin clusters.

DISCUSSION

We have shown by EM immunogold that after brain homogenization, synaptosomes derived from GABAergic synapses, which contain postsynaptic GABA_ARs and gephyrin and that have Gray's type-II synaptic morphology, are present in the classical synaptosomal fraction. We have also found that the "One Triton insoluble PSD" fraction, which is derived from the synaptosomal fraction and is enriched in glutamatergic type-I PSDs (Hunt et al. 1996 and Kennedy 1997), is also enriched in GABA_ARs and gephyrin, which are proteins associated with GABAergic type-II synapses. These results support and expand the notion put forward by Matus et al. 1980 that the detergent-insoluble PSD fraction contains not only type-I PSDs but also type-II PSDs. They based their hypothesis in that the detergent-insoluble PSD fraction showed [³H]GABA and [³H]Muscimol binding. Our EM immunogold studies of the "One Triton PSD" fraction show that the anti-GABA_AR and anti-gephyrin antibodies immunolabel amorphous structures of lower electron-density than those of type-I PSDs. We hypothesize that these structures correspond to type-II PSDs. This notion was confirmed after sedimentation of the "One Triton PSD" fraction on a continuous sucrose density gradient, which led to the partial separation of the GABAergic type-II PSDs from the glutamatergic type-I PSDs, as shown by the separation of two peaks, one of a heavier density ($\rho = 1.18\text{--}1.23$ g/ml) enriched in PSD-95, a marker of the glutamatergic type-I PSDs, and another one of a lighter density ($\rho = 1.10\text{--}1.15$ g/ml) enriched in gephyrin and $\gamma 2$ -GABA_ARs, markers of the GABAergic type-II PSDs. The EM immunogold of the $\rho = 1.10\text{--}1.15$ g/ml peak fraction shows that the GABA_AR and gephyrin immunolabeling concentrates in amorphous structures similar to those immunolabeled with anti-gephyrin and anti-GABA_AR subunit antibodies in the "One Triton PSD" fraction. We propose that these structures correspond to type-II GABAergic PSDs.

This interpretation is consistent with the known concentration and co-localization of gephyrin and $\gamma 2$ -GABA_ARs in the type-II postsynaptic GABAergic complexes both in brain and in cultured hippocampal neurons (Sassoe-Pognetto et al. 2000; Christie et al. 2002a; Craig et al. 1996; Giustetto et al. 1998; Kneussel et al. 1999; Christie et al. 2002b; van Rijnsoever et al. 2005). It has been shown that postsynaptic gephyrin is involved in the clustering of some postsynaptic GABA_ARs (Essrich et al. 1998; Kneussel et al. 1999, 2001; Christie et al. 2002a; Levi et al. 2004; Jacob et al. 2005), by restricting the lateral mobility of GABA_ARs in the membrane (Jacob et al. 2005), and that the $\gamma 2$ subunit is essential for the postsynaptic clustering of GABA_ARs and gephyrin, as shown by studies on the $\gamma 2$ knockout mouse (Essrich et al. 1998; Schweizer et al. 2003) and by $\gamma 2$ RNA interference studies (Li et al. 2005b).

The gradient fractions 6 and 7 (Fig. 2), which are enriched in $\gamma 2$ -GABA_ARs and gephyrin but not in PSD-95, should be valuable for identifying various protein components of the GABAergic type-II PSDs by proteomics. There is a limited knowledge on the protein components and the signal transduction mechanisms in type-II PSDs (Moss and Smart 2001; Owens and Kriegstein 2002). Proteomic approaches have been successfully used to identify various protein components of the purified type-I PSDs (Kennedy 1997; Walikonis et al. 2000; Yoshimura et al. 2004; Peng et al. 2004; Jordan et al. 2004; Collin et al. 2006).

A second pool of Triton X-100-insoluble GABA_ARs is present in the "One Triton PSD". This very low-density ($\rho = 1.06\text{--}1.09$ g/ml) pool of GABA_ARs is associated to DRMs and the lipid raft markers Caveolin-1, Thy-1 and Flotillin-1, but not to transferrin receptor. To the best of our knowledge, this is the first time that two pools of Triton X-100 -insoluble GABA_ARs in the rat brain have been identified in the "One Triton PSD" fraction: one associated with DRMs (and probably to lipid rafts), and another one associated with type-II PSDs. Thus, an important

consideration for those who use the “One Triton PSD” fraction in their research is that not all the Triton X-100-insoluble proteins that are present in this fraction are associated to PSDs. Some of these proteins are associated instead to the DRMs that are also present in the “One Triton PSD” fraction. It has been previously reported that some GABA_ARs in cultured rat cerebellar granule cells are associated to DRMs (Dalskov et al. 2005).

Our results show that the majority of the forebrain GABA_ARs associated with DRMs have α and β subunits, but little or no $\gamma 2$, and are largely not associated with gephyrin. Although most of the δ -GABA_ARs in the forebrain are also associated to DRMs, they represent a relatively small proportion of the α and β -GABA_ARs present in the forebrain DRMs. This notion is supported by I) the weak intensity of the δ protein band observed in immunoblots with three different anti- δ antibodies and II) although the δ subunit is highly expressed in rat cerebellum and thalamus, in the rat cerebral cortex (the main component of our forebrain preparation) there is very little expression of the δ subunit while there is high expression of the α , β and $\gamma 2$ subunits (Fritschy and Mohler 1995; Pirker et al. 2000). Thus, many of the GABA_ARs in the DRMs from forebrain contain α and β but not $\gamma 2$ or δ subunits while some contain α , β and δ subunits. The subunit composition of the GABA_ARs in the DRMs contrasts with that of the synaptic GABA_ARs associated with type-II PSDs, which have the $\gamma 2$ subunit (plus α and β) and are associated with gephyrin. The absence of the $\gamma 2$ and gephyrin suggests that the majority of GABA_ARs associated with DRMs (and lipid rafts) are non-synaptic, since as indicated above, the $\gamma 2$ subunit is essential for the postsynaptic localization of GABA_ARs (Essrich et al. 1998; Schweizer et al. 2003; Li et al. 2005b), and gephyrin concentrates at GABAergic synapses and is involved in the postsynaptic clustering of some GABA_ARs. Moreover, the δ -GABA_ARs, which we have shown to be associated with DRMs, are extrasynaptic and involved in tonic inhibition (Farrant and Nusser 2005; Nusser and Mody 2002; Stell et al. 2003). The difference in subunit composition between the GABA_ARs in type-II PSDs and the GABA_ARs in DRMs translates into different radioligand binding properties. Although both types of GABA_ARs show [³H]MUS, non-synaptic GABA_ARs present in the DRM fraction show low [³H]FNZ binding, which agrees with the low content of $\gamma 2$ subunit in the DRM fraction. In contrast, the synaptic GABA_ARs present in type-II PSDs contain the $\gamma 2$ subunit and show high [³H]FNZ binding. The DRM fraction is enriched in caveolin. Thus, some lipid raft-associated GABA_ARs might be endocytosed by caveolae and caveosomes. Caveolae are morphologically identified as flask-like invaginations of the plasma membrane that are enriched in lipid rafts and in the coat protein caveolin (Brown and London 1998, Masserini et al. 1999). It is also worth noticing that although the majority of gephyrin and $\gamma 2$ GABA_AR subunit are associated to type-II PSDs, there are some gephyrin and $\gamma 2$ associated to DRMs (in the DRM fraction of Fig. 3 and in the low-density fraction of Fig. 4). We do not know yet whether a small proportion of the $\gamma 2$ -GABA_ARs and gephyrin present at GABAergic synapses is associated to lipid rafts, or whether they represent $\gamma 2$ -GABA_ARs and gephyrin associated to lipid rafts that are trafficking in and out the GABAergic synapses.

There is a third pool of GABA_ARs that we have not considered in this study. This is a pool of GABA_ARs that can be solubilized by Triton X-100 at 4°C and that has been the object of many biochemical and pharmacological studies in the literature (and in Fig. 3). The GABA_ARs that are present in this Triton X-100-soluble (TS) fraction have α , β and $\gamma 2$ subunits and show [³H]MUS and [³H]FNZ binding (Fig. 3), indicating that they are fully assembled receptors. The binding activity recovered in the TS fraction was 45% of the [³H]MUS and 51% of the [³H]FNZ binding activity present in the SPM fraction. The solubility of the GABA_ARs in Triton X-100 at 4°C indicates that they are not associated to lipid rafts and suggests that they might not be associated to the type-II PSDs and to the GABA_ARs that are present in the Triton X-100-insoluble (TI) fraction. The GABA_ARs in the TS fraction might represent extrasynaptic membrane receptors with lateral mobility that can eventually be anchored at the postsynaptic

complexes and become part of the immobilized synaptic pool (Choquet and Triller 2003; Thomas et al. 2005).

In hippocampal cultures, the normal $\gamma 2$ -GABA_AR and gephyrin clusters are not associated with lipid rafts since they cannot be solubilized by Triton X-100 at 4°C even after cholesterol depletion. These results with cultures agree with the results obtained with brain fractions in the sense that the majority of brain $\gamma 2$ -GABA_AR and gephyrin are not present in DRMs. Moreover, the majority (74%) of the $\alpha 1$ -GABA_AR clusters, that are normally present in the cultured hippocampal neurons, are not associated to lipid rafts, since they are resistant to solubilization with Triton X-100 at 4°C even after cholesterol depletion. The majority of these $\alpha 1$ -GABA_AR clusters colocalize with gephyrin clusters. However, the remaining 26% of Triton-insoluble $\alpha 1$ -GABA_AR clusters become solubilized by Triton X-100 when cholesterol is depleted, indicating that they are associated to lipid rafts. These $\alpha 1$ -GABA_AR clusters do not co-localize with gephyrin clusters.

In cultured hippocampal neurons, there is also a pool of extrasynaptic $\alpha 1$ -GABA_ARs that are diffusely distributed at the neuronal surface and that can be induced to form clusters by capping with a primary antibody that recognizes an extracellular epitope of the GABA_AR (one-step capping), or by the sequential incubation of live neurons with the primary antibody and the fluorophore-labeled secondary antibody (two-step capping). Antibody-induced capping has been used by us and others to reveal diffuse non-clustered GABA_ARs on the surface of cultured neurons (Levi et al. 2004; Christie et al. 2006). The pool of diffuse GABA_ARs that can be clustered by one-step or two-step capping induced by anti- $\alpha 1$ is associated to lipid rafts since it resists extraction with Triton X-100 at 4°C, but it becomes solubilized by this detergent after cholesterol depletion. The antibody-induced clustering of the $\alpha 1$ -GABA_ARs does not induce the simultaneous co-clustering of gephyrin, as we and others have previously observed, following GABA_AR capping induced with antibodies to other GABA_AR subunits (Levi et al. 2004; Christie et al. 2006), indicating that the GABA_ARs in the non-clustered state are not associated with gephyrin. These results also support the notion that in the absence of association with gephyrin, GABA_ARs show high lateral mobility in the membrane (Jacob et al. 2005), which allows the capping by the antibody of the diffuse membrane GABA_ARs.

Thus, the studies with rat forebrain fractions and hippocampal cultures support the hypothesis that there are two pools of assembled GABA_ARs that resist Triton X-100 solubilization at 4°C. One pool is associated with gephyrin, contains the $\gamma 2$ subunit (plus α and β subunits), shows high specific activity of both [³H]MUS and [³H]FNZ bindings and co-purifies with type-II PSDs of the GABAergic synapses. This synaptic pool forms a Triton X-100 insoluble protein complex containing GABA_ARs, gephyrin, the postsynaptic cytoskeleton and scaffold proteins among others. These synaptic GABA_ARs are not associated to DRMs or lipid rafts. There is another pool of GABA_ARs in the forebrain that is also insoluble in Triton X-100 but that is associated to DRMs and lipid rafts. These GABA_ARs do not contain the $\gamma 2$ subunit but the majority contain α and β subunits but not δ and some contain α , β and δ subunits. These GABA_ARs show high [³H]MUS binding but low [³H]FNZ binding and show little or no association with gephyrin. These GABA_ARs are non-synaptic, many are diffusely distributed through the neuronal surface and can be induced to form clusters by anti- $\alpha 1$ antibody-induced capping.

The association of some GABA_ARs with lipid rafts might have functional relevance since cholesterol modulates GABA_ARs (Sooksawate and Simmonds 2001). It is thought that the synaptic GABA_ARs that are present in type-II GABAergic synapses are involved in phasic inhibition while the diffuse extrasynaptic GABA_ARs are involved in tonic inhibition (Luscher and Keller 2004; Kullmann et al. 2005; Farrant and Nusser 2005; Mody 2005). Our results are consistent with the notion that the GABA_ARs associated to lipid rafts in the forebrain are

extrasynaptic and are involved in tonic inhibition since: I) some of the forebrain GABA_ARs associated with lipid rafts contain the δ subunit and it has been shown that δ -GABA_ARs are extrasynaptic and involved in tonic inhibition (Farrant and Nusser 2005; Nusser and Mody 2002; Stell et al. 2003); II) other forebrain GABA_ARs present in lipid rafts contain α and β subunits but not γ 2 or δ . It has recently been reported the existence of extrasynaptic GABA_ARs that contain only α and β subunits and that are involved in tonic inhibition (Mortensen and Smart 2006).

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Abbreviation used

DRM	detergent resistant membrane
EM	electron microscopy
FNZ	flunitrazepam
GABA_AR	γ -aminobutyric acid type-A receptor
Geph	gephyrin
mAb	monoclonal antibody
MβCD	methyl- β -cyclodextrin
MUS	muscimol
PSD	postsynaptic density
PSD-95	postsynaptic density protein 95
SDS-PAGE	sodium dodecyl sulfate-polyacrylamide gel electrophoresis
SPM	synaptosomal plasma membrane fraction
SYN	synaptosomal fraction
TI	Triton X-100 insoluble fraction
TS	Triton X-100 soluble fraction

TX-AF

Triton X-100 extraction after fixation

TX-BF

Triton X-100 extraction before fixation

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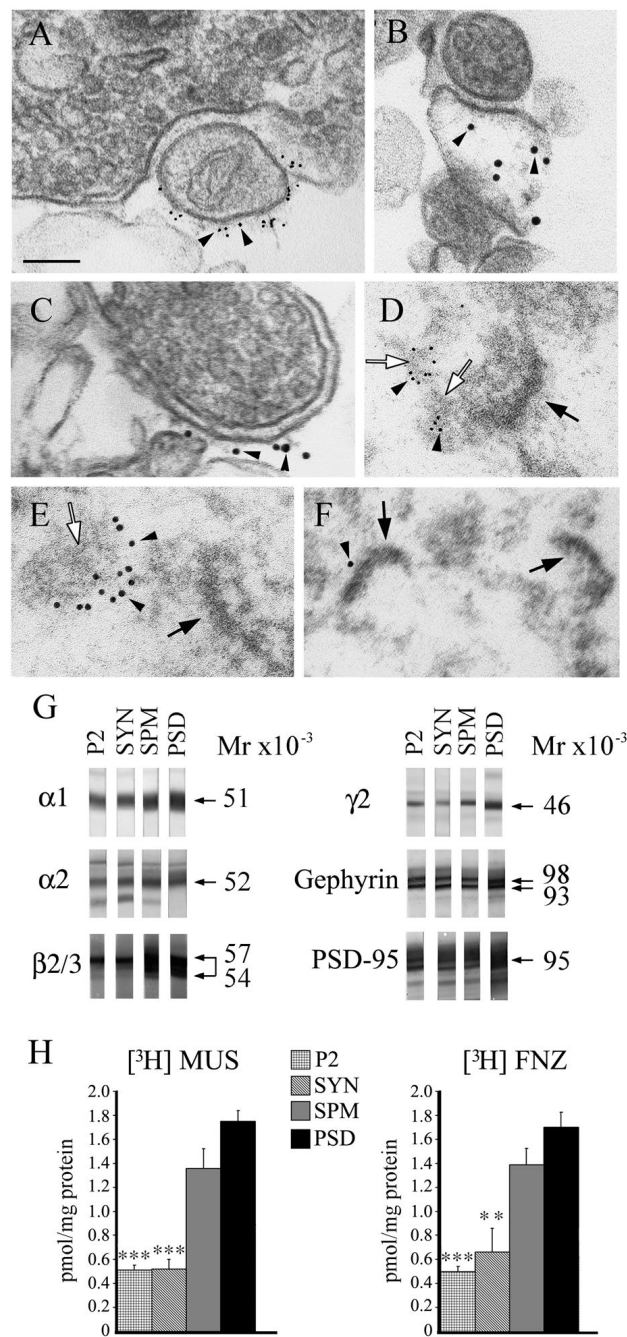
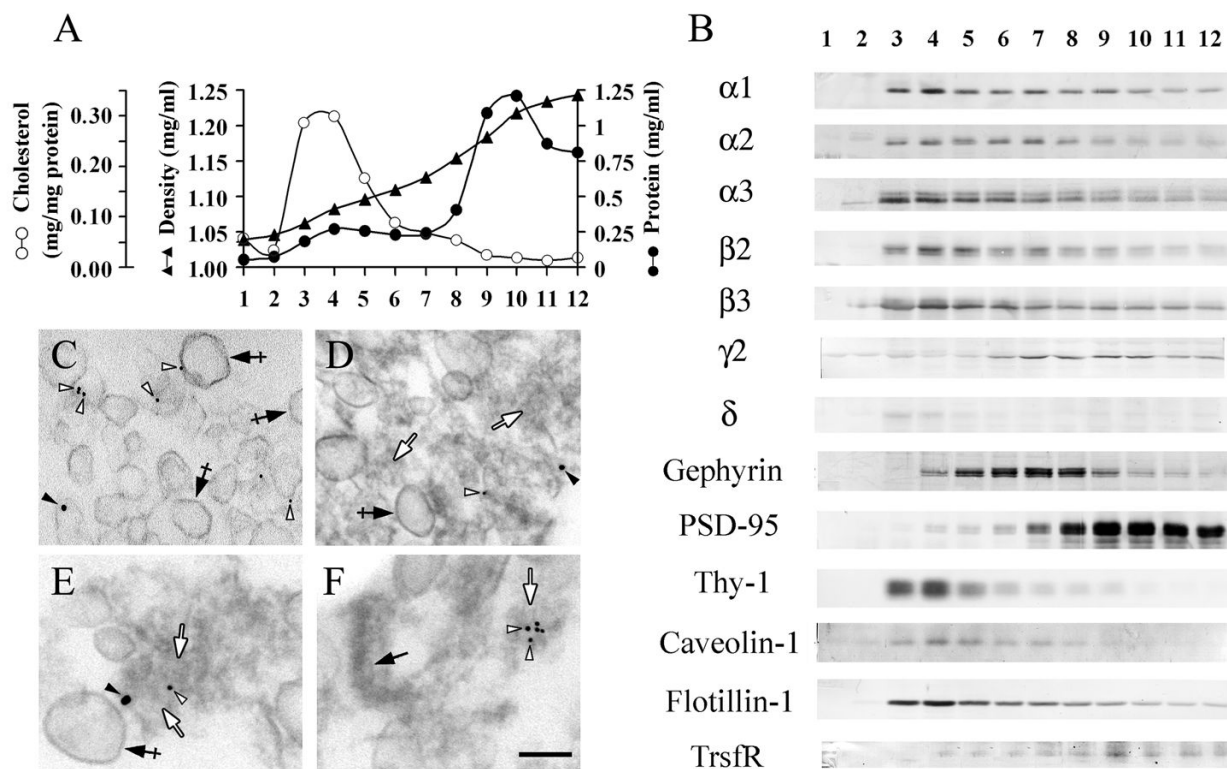


Fig. 1. GABA_ARs are present in various subcellular brain fractions. The P2, SYN, SPM, PSD, were prepared as described in Materials and Methods. **A–F** show EM immunogold of synaptosomal (A–C) and “One Triton PSD” (D–F) fractions with rabbit anti-α1 GABA_AR subunit (A and D, arrowheads), mAb to gephyrin (B, C and E, arrowheads) and mAb to PSD-95 (F, arrowhead). Filled arrows indicate type-I PSDs, which have high electron-density. Empty arrows point to structures of medium electron-density immunolabeled with antibodies to α1 or gephyrin, likely corresponding to type-II PSDs. The secondary goat anti-rabbit IgG and anti-mouse IgG antibodies were conjugated to 6 nm and 12 nm diameter colloidal gold particles respectively. Scale bar = 100 nm. **G:** Immunoblots of various brain fractions with rabbit anti-α1, rabbit anti-

$\alpha 2$, rabbit anti- $\gamma 2$, and rabbit anti-gephyrin antisera, mAb to $\beta 2/3$ and mAb to PSD-95. Same amount of total protein from each fraction (10 μ g/lane) was used for SDS-PAGE. The immunoreactivity of the specific protein band was blocked by the corresponding antigenic peptide (data not shown). **H:** [3 H]MUS and [3 H]FNZ binding to various brain fractions. Specific activity values (pmol/mg protein) are mean \pm S.E.M of at least three experiments each done in triplicate. *** $p < 0.001$; ** $p < 0.01$. The p values are for the various fractions compared to PSD.

**Fig. 2.**

In the "One Triton PSD" fraction, some GABA_ARs are associated with DRMs and some are associated with GABAergic type-II PSDs. Fractionation of "One Triton PSD" after centrifugation on a continuous 0.32–2.0 M sucrose density gradient. **A**: Distribution of cholesterol and protein in the density gradient. **B**: Immunoblots of the various fractions with antibodies to GABA_ARs subunits ($\alpha 1$, $\alpha 2$, $\alpha 3$, $\beta 2$, $\beta 3$, $\gamma 2$, δ), Gephyrin, PSD-95, transferrin receptor (TrsfR) and the lipid raft markers Thy-1, Caveolin-1 and Flotillin-1. The same amount of total protein (4 μ g/lane) was used for SDS-PAGE. Fractions 1 and 12 correspond to the top and bottom of the gradient respectively. **C**: Double-labeled EM Immunogold of pooled gradient fractions 3 and 4 with rabbit anti- $\alpha 1$ and mAb anti- $\beta 2/3$. Crossed arrows show the DRM vesicles present in this fraction. Some of these structures are immunolabeled with rabbit anti- $\alpha 1$ (filled arrowhead) and/or mAb anti- $\beta 2/3$ (empty arrowheads). **D–F**: Double-label EM immunogold of fraction 7 with rabbit anti- $\gamma 2$ and mAb to gephyrin. Filled arrow in F indicates a type-I PSD that has high electron-density. Empty arrows in D–F point to amorphous structures of medium electron-density. Some of these structures are immunolabeled with rabbit anti- $\gamma 2$ (filled arrowheads) and/or mAb to gephyrin (empty arrowheads). Crossed arrows in D and E indicate vesicular membranes. The secondary goat anti-rabbit IgG and goat anti-mouse IgG antibodies were conjugated to 18 nm and 10 nm diameter colloidal gold particles respectively. Scale bar represents 150 nm in C and D and 100 nm in E and F.

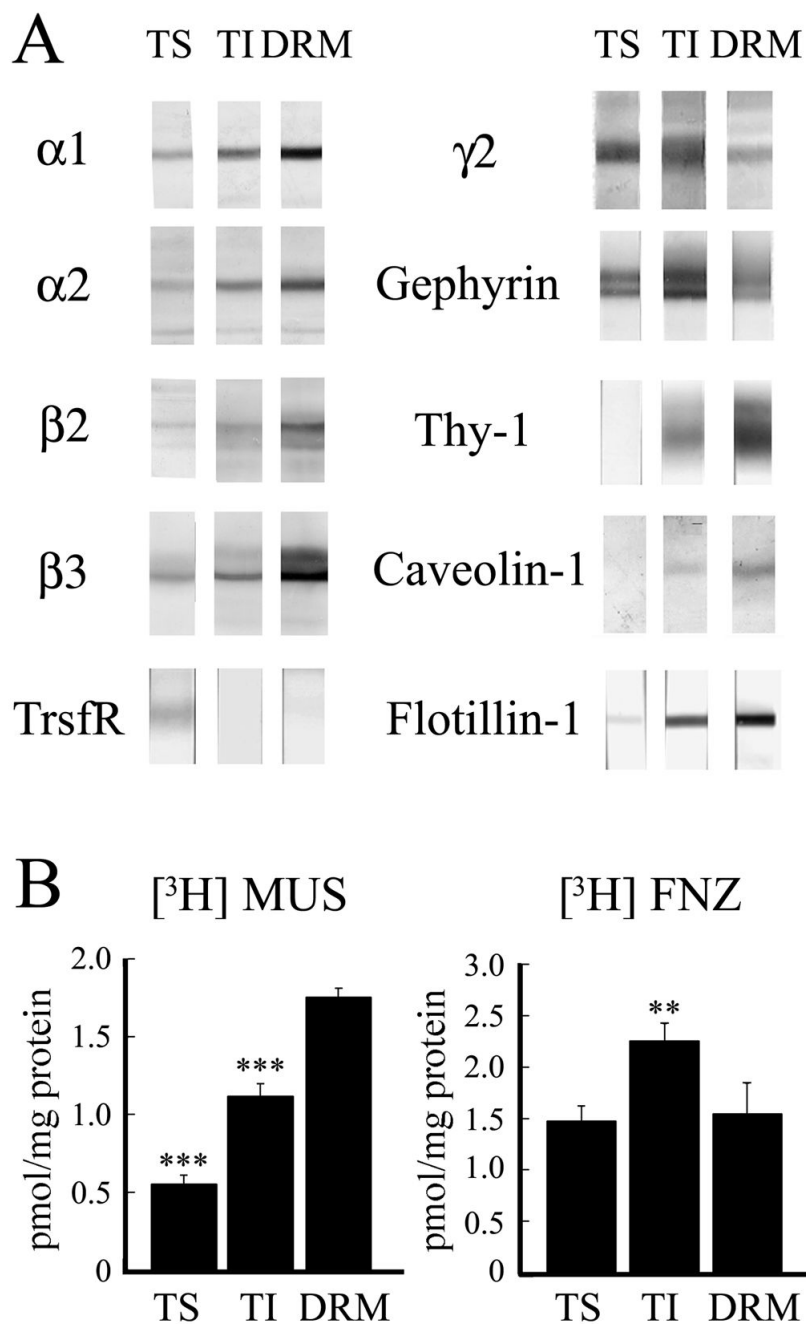


Fig. 3. Some GABA_AR in synaptosomal plasma membrane are associated with DRMs. The SPM fraction was extracted by 1% Triton X-100 at 4°C and the soluble (TS) and insoluble (TI) fractions were collected. The low-density DRM fraction was also collected after flotation in a discontinuous sucrose gradient. **A:** Immunoblots with rabbit antibodies to $\alpha 1$, $\alpha 2$, $\beta 2$, $\beta 3$, $\gamma 2$ GABA_AR subunits, gephyrin, transferrin receptor (TrsfR) and lipid raft markers Thy-1, Caveolin-1 and Flotillin-1. Same amount of total protein (4 μ g/lane) was used for SDS-PAGE. **B:** [^3H]MUS and [^3H]FNZ binding to GABA_AR in TS, TI and DRM. Specific activity values (pmol/mg protein) are mean \pm S.E.M of at least three experiments each done in triplicate. *** $p < 0.001$; ** $p < 0.01$. The p values are for the TS or TI fractions compared to DRM.

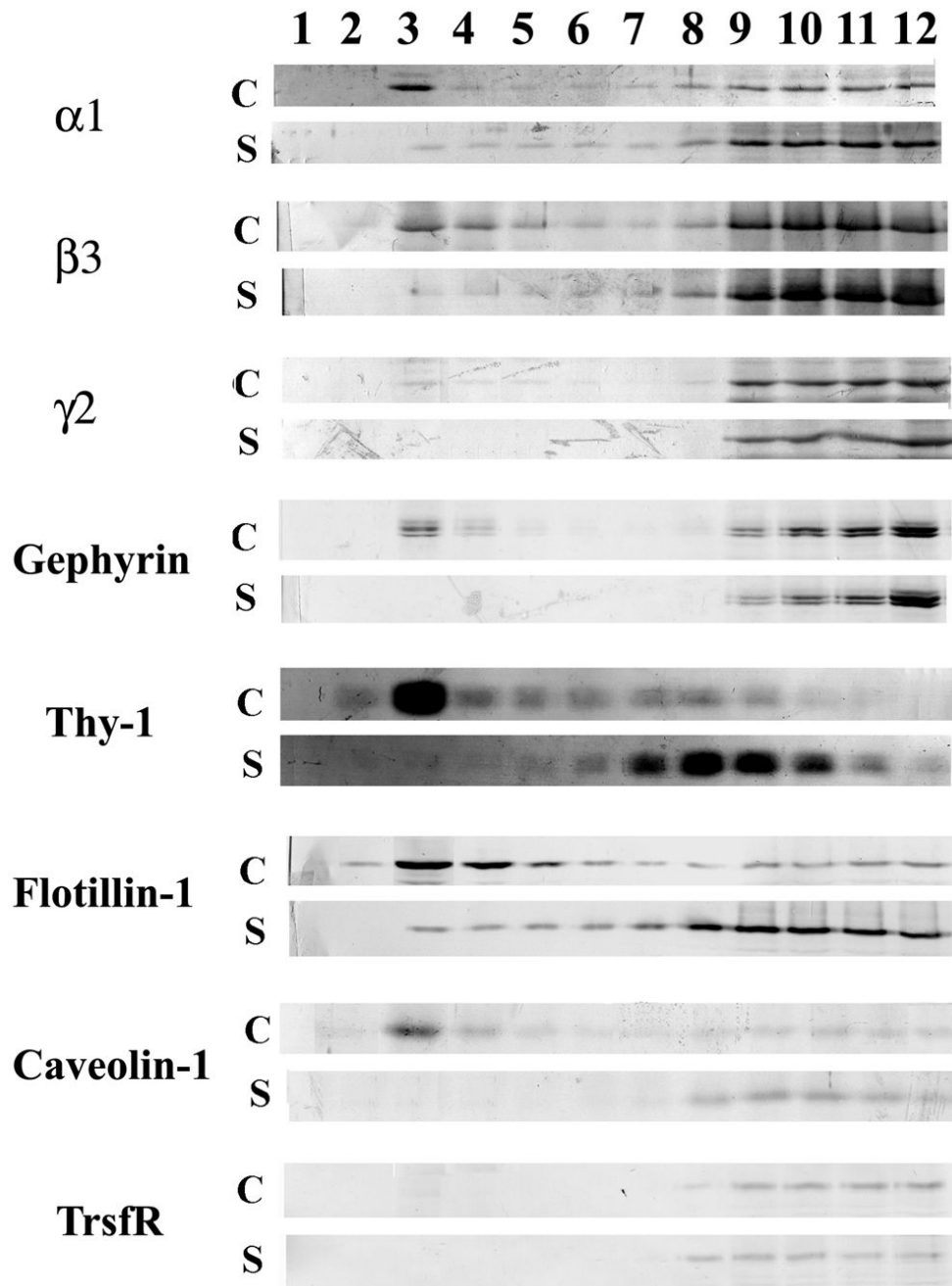


Fig. 4. Saponin treatment disrupts the association of GABA_ARs with DRMs. The SPM fraction was incubated with 0.5% saponin (S) or without saponin (C, control) followed by extraction with 1% Triton X-100 at 4°C, flotation in a sucrose density gradient and fractionation. Immunoblotting was done with rabbit antibodies to $\alpha 1$, $\beta 3$, $\gamma 2$ GABA_ARs, gephyrin, transferrin receptor (TrsfR) and the lipid raft markers Thy-1, Flotillin-1 and Caveolin-1. The same volume (12 μ l/lane) from each fraction was used for SDS-PAGE. Fractions 1 and 12 correspond to the top and bottom of the gradient respectively.

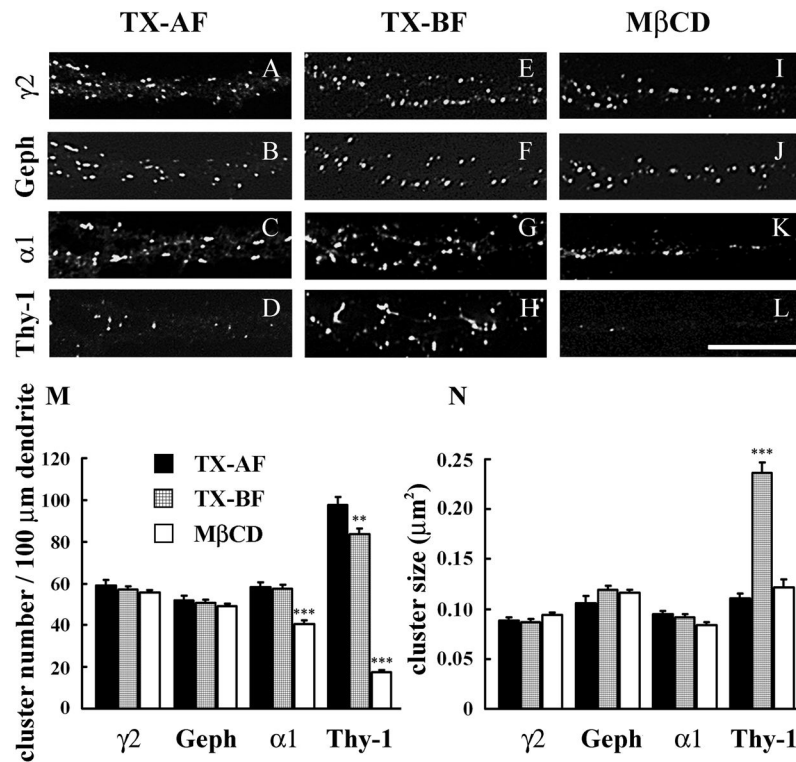
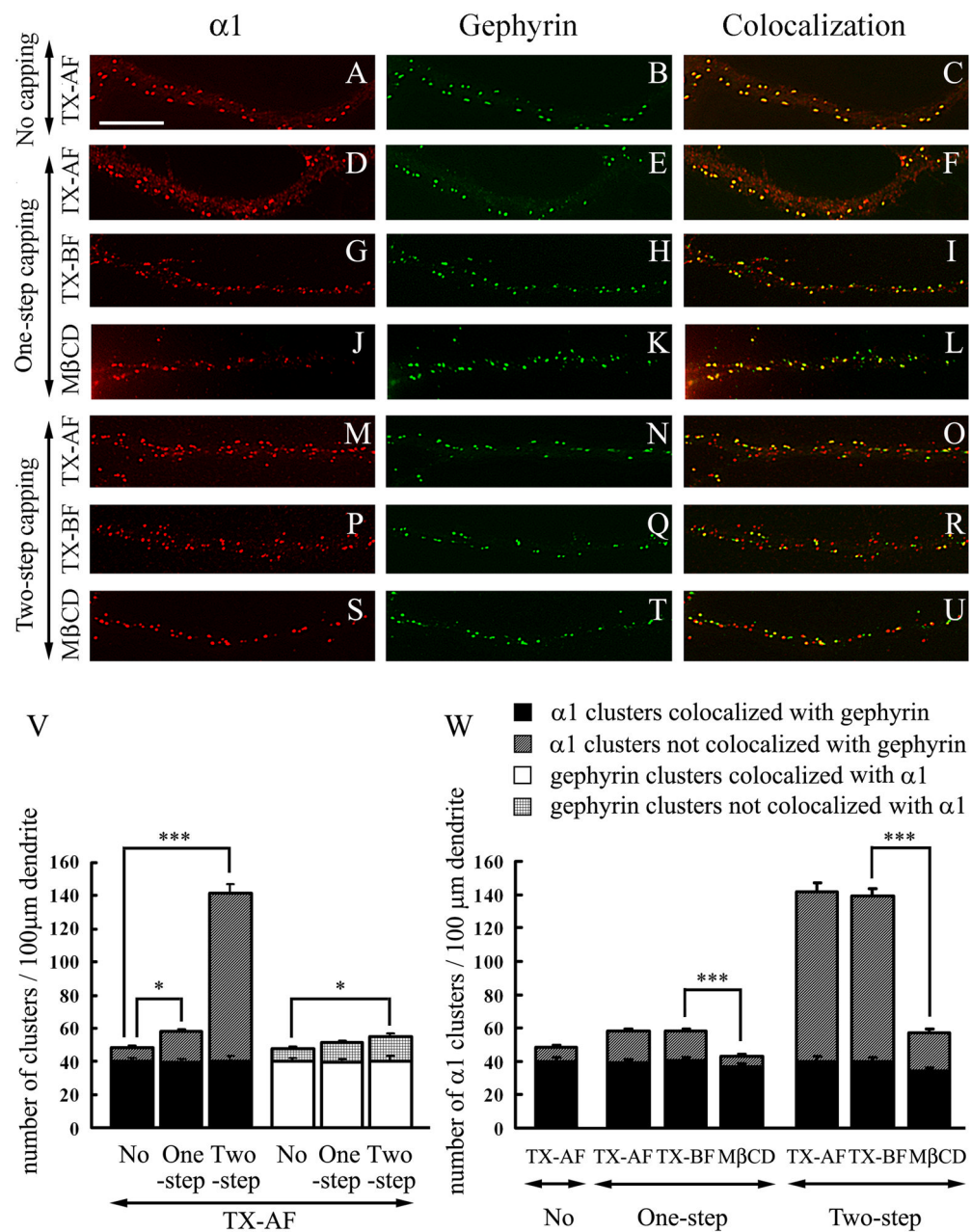


Fig. 5. In cultured hippocampal neurons some $\alpha 1$ -GABA_AR clusters, but no $\gamma 2$ -GABA_AR clusters or gephyrin clusters, are associated with lipid rafts. Double-label immunofluorescence of pyramidal cell dendrites. Live hippocampal neurons were subjected to one-step capping with anti- $\gamma 2$ or anti- $\alpha 1$ GABA_AR antibody and double-labeling with mAb to gephyrin or Thy-1 respectively. **A–D**: After one-step capping with guinea pig anti- $\gamma 2$ (A and B) or rabbit anti- $\alpha 1$ (C and D), cells were fixed followed by incubation with Triton X-100 at 4°C (Triton X-100 after fixation, TX-AF). **E–H**: After surface labeling and one-step capping with guinea pig anti- $\gamma 2$ (E and F) or rabbit anti- $\alpha 1$ (G and H), cells were treated with Triton X-100 at 4°C before fixation (TX-BF). **I–L**: After surface labeling and one-step capping with guinea pig anti- $\gamma 2$ (I and J) or rabbit anti- $\alpha 1$ (K and L), cells were treated with MβCD before Triton X-100 extraction at 4°C followed by fixation (MβCD). Double-label immunofluorescence of $\gamma 2$ -GABA_AR subunit (A, E and I) and gephyrin (B, F and J) or $\alpha 1$ -GABA_AR subunit (C, G and K) and Thy-1 (D, H and L). The secondary antibodies were Texas Red conjugated anti- guinea pig or anti- rabbit IgG and FITC conjugated anti- mouse IgG, all made in donkey. Scale bar: 10 μm. **M and N**: Quantification of cluster density and cluster size respectively. Data in M and N were collected from 30–40 dendrites from 15–20 cells of 3–4 separate experiments. *** $p < 0.001$; ** $p < 0.01$. The p values are for the TX-BF or MβCD compared to TX-AF.

**Fig. 6.**

An anti- $\alpha 1$ antibody induces the capping of $\alpha 1$ -GABA_ARs that are associated with lipid rafts. Double-label immunofluorescence of pyramidal cell dendrites. **A–C**: Hippocampal cultures were fixed, permeabilized with Triton X-100 and double-labeled with rabbit anti- $\alpha 1$ and mAb to gephyrin (no capping, TX-AF). **D–L**: Live neurons were one-step capped with anti- $\alpha 1$ antibody and double-labeled with mAb to gephyrin. **M–U**: Live neurons were two-step capped with sequential incubation of anti- $\alpha 1$ antibody and secondary antibody and then double-labeled with mAb to gephyrin. In TX-AF, one-step capping (D–F) or two-step capping (M–O) was followed by 0.25% Triton X-100 permeabilization after fixation. In TX-BF, one-step capping (G–I) or two-step capping (P–R) was followed by incubation with 0.5% Triton X-100 at 4°C before fixation. In MβCD, one-step capping (J–L) or two-step capping (S–U) was followed by

treatment with M β CD before incubation with Triton X-100 at 4°C and fixation. Panels A–U show representative dendrites. Scale bar: 10 μ m. **V and W:** Quantification of α 1-GABA_AR and gephyrin cluster density and their co-localization in hippocampal cultures in the various experimental conditions represented in panels A–U. For quantification, 30 dendrites were randomly chosen from 15–20 cells from 3 separate experiments. Values represent mean \pm SEM. *** $p < 0.001$; * $p < 0.05$.