Agonist-dependent Endocytosis of γ-Aminobutyric Acid Type A (GABA_A) Receptors Revealed by a γ2(R43Q) Epilepsy Mutation*

Severine Chaumont‡§, Caroline André‡§, David Perrais¶, Eric Boué-Grabot**‡‡, Antoine Taly‡§, and Maurice Garret†‡§

From the †Université Bordeaux, Institut de Neurosciences Cognitives et Intégratives d’Aquitaine (INCIA), UMR 5287, F-33000 Bordeaux, France; §CNRS, INCIA, UMR 5287, F-33000 Bordeaux, France; ‡Université Bordeaux, Institute for Interdisciplinary Neuroscience (IINS), UMR 5297, F-33000 Bordeaux, France; ¶CNRS, IINS, UMR 5297, F-33000 Bordeaux, France, **Université Bordeaux, Institut des Maladies Neurodégénératives, UMR 5293, F-33000 Bordeaux, France, §§CNRS, IINS, UMR 5293, F-33000 Bordeaux, France, and ¶¶CNRS, UPR 9080, Institut de Biologie Physico-Chimique, 13 Rue Pierre et Marie Curie, 75005 Paris, France

Background: Modulation of GABA_A receptors trafficking is critical for controlling inhibitory neurotransmission.

Results: A point mutation or agonist application, both affecting the GABA_A receptor extracellular domain, has an effect on receptor endocytosis.

Conclusion: Endocytosis of GABA_A receptors is linked to agonist-induced conformational changes.

Significance: This represents one of the few reports demonstrating an influence of extracellular effectors on GABAergic physiology suggesting that γ2(R43Q) trafficking in cultured hippocampal neurons and COS-7 cells can be modulated by their agonists.

GABA-gated chloride channels (GABA_ARs) trafficking is involved in the regulation of fast inhibitory transmission. Here, we took advantage of a γ2(R43Q) subunit mutation linked to epilepsy in humans that considerably reduces the number of GABA_ARs on the cell surface to better understand the trafficking of GABA_ARs. Using recombinant expression in cultured rat hippocampal neurons and COS-7 cells, we showed that receptors containing γ2(R43Q) were addressed to the cell membrane but underwent clathrin-mediated dynamin-dependent endocytosis. The γ2(R43Q)-dependent endocytosis was reduced by GABA_AR antagonists. These data, in addition to a new homology model, suggested that a conformational change in the extracellular domain of γ2(R43Q)-containing GABA_ARs increased their internalization. This led us to show that endogenous and recombinant wild-type GABA_AR endocytosis in both cultured neurons and COS-7 cells can be amplified by their agonists. These findings revealed not only a direct relationship between endocytosis of GABA_ARs and a genetic neurological disorder but also that trafficking of these receptors can be modulated by their agonist. Inhibitory transmission relies greatly on ionotropic GABA_A receptors (GABA_ARs) that are involved in many physiological functions and are the target of several drugs in wide clinical use (1). GABA_AR trafficking is modulated by a number of different mechanisms, including surface targeting, mobility, and endocytosis. Moreover, studies in physiological and pathophysiological states have revealed that the number of GABA_ARs on the cell membrane have a profound influence on GABAergic neurotransmission (1–3). Inhibitory neurotransmission is also regulated by the exchange between surface and intracellular compartments via a constitutive clathrin-mediated dynamin-dependent endocytosis pathway (4–6). This constitutive internalization is modulated by intracellular mechanisms and is altered in pathological conditions (4, 6–9).

Epilepsies are complex syndromes with multiple causes and symptoms, but it is well established that alteration or modulation of GABA neurotransmission plays an important role in the disease and its treatment (10–15). Moreover, genetic evidence has revealed a direct link between epilepsy and GABA_AR dysfunction, including trafficking alteration, supporting the hypothesis that defects in GABA_ARs lead to seizures (16–17). These mutations also offer an opportunity to obtain new insights into GABA_AR structure and function as well as clues to the role of these receptors in neurological disorders (14). For example, an R43Q mutation located in the γ2 subunit N-terminal extracellular domain is linked to childhood absence epilepsy and febrile seizure (17); heterozygous mice harboring this mutation replicate the human clinical phenotype (16). Intensive research into this mutation (18–26) has led to controversial data on its effects on GABAergic physiology suggesting that γ2(R43Q) might modify the dynamics of subunit trafficking (27).

Here, we analyzed γ2(R43Q) trafficking in cultured hippocampal neurons and COS-7 cells and revealed that receptors containing the γ2(R43Q) subunit had a shorter residence time on the plasma membrane than their wild-type counterparts.
We also showed that endocytosis of the mutated receptor was clathrin- and dynamin-dependent. However, it was surprising that a mutation in the extracellular domain (bearing binding sites for agonists and modulators) could have an influence on internalization, believed to be controlled through the intracellular domain. Moreover, endocytosis of GABA_ARs triggered by agonist exposure remains to be fully assessed (11, 28–31). Then, by using both imaging and biochemical methods, further experiments revealed that agonist exposure triggered an increase of wild-type GABA_Ar endocytosis, both on native and recombinant GABA_ARs.

EXPERIMENTAL PROCEDURES

Antibodies and Reagents—Antibodies were raised in rabbits against the α1 GABA_A subunit N terminus (Alomone Labs), the Myc tag (Upstate, Charlottesville, VA), membrin (1:1000; Synaptic Systems), calreticulin (1:500; Upstate Biotechnology, Inc.), or EEA1 (Sigma–Aldrich). Antibodies were raised in mice against the α1 intracellular loop (Neuromab) or the Myc tag (Roche Applied Science). Secondary antibodies were as follows: Alexa Fluor-568 goat anti-rabbit, Alexa Fluor-647 goat anti-rabbit, Alexa Fluor-488 goat anti-mouse (1:1000; Molecular Probes), and FITC-coupled anti-mouse antibody (1:200; Chemicon).

DNA Constructs—α1, β2, β3, and γ2S GABA_AγRs subunits were subcloned in the pcDNA3 vector (Invitrogen), using constructs that were available from previous studies (32, 33). The Myc epitope (MEQKLISEEDLNE, repeated 6 times) was inserted between amino acids 4 and 5 of the γ2 subunit (22). As described previously, an insertion within this domain does not modify the functional properties of GABA- or glutamate-gated channels (32, 34–36). Point mutations were constructed using the QuikChange site-directed mutagenesis system (Stratagene). All constructs were verified by automatic dideoxy DNA sequencing (Genome Express, Meylan, France). Endoplasmic reticulum–Golgi intermediate compartment was revealed with ERGIC-GFP (37), kindly provided by Jochen Lang (Bordeaux, France).

Cell Culture and Transfection—COS-7 and HEK 293 cells were maintained in Dulbecco’s modified Eagle’s medium (Invitrogen) supplemented with 10% fetal calf serum (Eurobio). Embryonic hippocampal neurons were obtained from E18 rat embryos, as described earlier (22). COS-7 and HEK 293 cells were transfected using the FuGENE 6 reagent (Roche Applied Science), according to the manufacturer’s specifications, with equal amounts of α1, β, and γ2 subunit cDNAs (and GFP for electrophysiology experiments on HEK 293 cells) (0.3 μg/well in 24-well plates). Cells were incubated with cDNAs for 24 h before analysis (22). Hippocampal neurons were transfected in vitro at 7–11 days, using the Lipofectamine 2000 reagent (Invitrogen) according to the manufacturer’s specifications. The cells were analyzed 24–60 h after transfection (22).

Immunocytochemistry—For living cell surface labeling, COS-7 cells and hippocampal neurons were incubated with antibodies at room temperature for 20 min in Dulbecco’s modified Eagle’s medium or Neurobasal medium supplemented with 10 mM HEPES, respectively. Receptors on the surface were labeled with antibodies raised in rabbits against either the α1 GABA_A subunit N-terminal domain or the Myc tag. Sera were diluted 1:500 (anti-α1) or 1:200 (anti-Myc) in medium. After incubation, cells were washed quickly by dipping coverslips in medium and fixed for 10 min in phosphate-buffered saline (PBS) containing 4% sucrose and 4% paraformaldehyde preheated to 37 °C, washed in PBS, and blocked in 0.3% bovine serum albumin and 50 mM glycine (in PBS) for 15 min. Cells were washed in PBS containing 0.3% bovine serum albumin. After cell permeabilization using 0.3% Triton X-100, intracellular tagged γ2 subunits were detected by incubating the cells with a mouse anti-Myc 9E10 antibody (1:1000; Roche Applied Science) for 2 h. Intracellular α1 subunits were detected with a mouse anti-α1 (1:1000). Polyclonal and monoclonal antibodies were detected using an Alexa Fluor-568-coupled anti-rabbit antibody and an Alexa Fluor-488-coupled anti-mouse antibody (1:1000), respectively. Endoreticulum and cis-Golgi staining were revealed with polyclonal anti-calreticulin (1:500) and antimonin (1:1000), respectively. The endoplasmic reticulum–Golgi intermediate compartment was revealed with ERGIC-GFP (37).

Internalization—Transfected neurons or COS cells were incubated in culture medium containing mouse monoclonal Myc antibody (1:200) at 37 °C for 30 min. The medium also contained GABA_AγR agonists or antagonists as required. In the case of neurons, a mixture of inhibitors (6-cyano-7-nitroquinoxalene-2,3-dione (10 μM), d-2-amino-5-phosphonovaleric acid (50 μM), and tetrodotoxin (1 μM)) was added, so as to prevent activity-dependent modulation of GABA_AγR trafficking (7). This mixture was added 5 min before the labeling experiments and was present throughout, both in control experiments and in GABA_AγR agonist/antagonist-treated neurons. Surface labeling was carried out at 20 °C using an anti-mouse secondary antibody coupled to Alexa Fluor-647 for 30 min (38); alternatively, labeled receptors remaining on the COS-7 cell surface were acid-washed for 2 min at 20 °C with culture medium adjusted at pH 2.5 (39, 40). Cells were then fixed using 4% paraformaldehyde in 4% sucrose and permeabilized in Triton X-100. Total protein content was assessed by incubating with a polyclonal antibody directed at the Myc tag. Internalized protein was then revealed by incubating with anti-mouse antibody coupled to Alexa Fluor-488 at room temperature for 1 h, whereas total protein was revealed using anti-rabbit antibody coupled to Alexa Fluor-568. For native α1 expressed in hippocampal neurons, cells were incubated in culture medium containing an antibody made in rabbits directed at the N-terminal domain (extracellular) of α1 (1:500) at 37 °C for 30 min. Surface labeling was carried as above using an anti-rabbit secondary antibody. Cells were then fixed and permeabilized as above. Total α1 subunit content was assessed by incubating with a mouse antibody directed at the Myc tag or at the native α1 at room temperature for 20 min. Internalized protein was then revealed by incubating cells with anti-mouse antibody coupled to Alexa Fluor-488 at room temperature for 1 h, whereas total protein was revealed using anti-rabbit antibody coupled to Alexa Fluor-568.

Quantitative Analysis of Fluorescence Signals—Fluorescence microscopy was performed using a Zeiss Axioplan 2 microscope, with a ×63, 1.4 numerical aperture oil immersion lens.
**γ2(R43Q) and GABA<sub>R</sub> Trafficking**

Quantification of fluorescence signals and background subtraction were performed using ImageJ (National Institutes of Health). For each image acquired, background levels were determined using the surface and intracellular signals measured in neighboring non-transfected cells and subtracted from the values obtained in transfected cells. Numerical data are presented as mean ± S.E., and statistical significance was assessed using one-way analysis of variance (Origin, Originlab Corp.) (significance level, < 0.05). Confocal microscopy was performed using an upright Leica DMR TCS SPZ AOBS, with a ×63, 1.4 numerical aperture Leica HPCL Fluoar oil objective. Colocalization was quantified using a plugin for ImageJ designed by F. Levet and C. Poujol (BIC (Bordeaux Imaging Center), Bordeaux, France). Briefly, two images, one containing GABA<sub>R</sub> subunit labeling and one containing the labeling for a cellular compartment, were thresholded in the same way. The plugin calculates the percentage of pixels containing γ2 subunit labeling that also contain specific labeling for a cellular compartment. The percentage of colocalization was normalized for total γ2 and γ2(R43Q) immunoreactivity, respectively. Analyses were performed in parallel cultures, blind to experimental conditions.

Quantification of surface clusters or intracellular punctate labeling, blind to experimental conditions, was performed using ImageJ (National Institutes of Health). Threshold was applied to the images, and the number as well as the area of surface clusters or internalized particles were measured using the particle analyzer module of ImageJ. For COS-7 cells, the whole cell was counted. For neurons, an area of 10-μm length along a dendrite was counted. For all experiments, total protein expression was assessed by antibody labeling after permeabilization of the cells and was measured for the same area to allow normalization of the values. To calculate fluorescence ratios, a stack was created for each cell in ImageJ with the image corresponding to the surface and total labeling. This allows us to draw the outline of the cell and measure the average surface and total fluorescence for the same area.

**Biotinylation Assays** — Biotinylation experiments were performed essentially as described previously (36, 38). COS-7 cells were transfected in 6-well plates (2 wells/condition) and were incubated 24 h post-transfection. Cells were then washed two times with PBS, pH 8.0, incubated with 1 mg/ml EZ-Link Sulfo-NHS-SS-Biotin (Pierce) in PBS for 30 min at 4 °C, washed three times with PBS, and scraped in lysis buffer containing 25 mm HEPES, 150 mM NaCl, 1% Triton X-100, and a mix of protease inhibitors (Roche Applied Science). After centrifugation, the supernatant was immunoprecipitated with 50 μl of immunopure immobilized streptavidin-beaded agarose overnight at 4 °C and washed extensively. Surface and total proteins were separated on SDS-PAGE and revealed by Western blotting using anti-Myc antibodies at a 1:1000 dilution. Quantification of Western blots was performed using ImageJ (National Institutes of Health) or with the Chemi Doc XRS+ under the control of the Image Lab software (Bio-Rad). To assess γ2 internalization, plates were returned to 37 °C for 30 min after biotinylation to allow endocytosis. Cells were then exposed to 50 mM MESNA, which cleaved biotin from proteins remaining on the surface. A sample was kept at 4 °C (instead of 37 °C) to control to ensure that the cleavage with MESNA was complete. Samples were analyzed by Western blot as above.

**Electrophoretic and Western Blot Analyses** — COS-7 cells were homogenized in buffer containing 20 mM HEPES, 0.15 mM EDTA, and 10 mM KCl, pH 8, supplemented with a mixture of protease inhibitors (Roche Applied Science). The buffer was then adjusted to 12% sucrose, and after four more strokes, the cells were centrifuged at 2000 rpm for 3 min to remove genomic DNA. The supernatant was centrifuged at 15,000 rpm for 30 min. The pellet was recovered, and cell membranes were solubilized with 15 strokes in a buffer containing 20 mM Tris-HCl, 0.15 mM EDTA, 150 mM NaCl, 2% Triton X-100, and 0.5% deoxycholate, pH 8, supplemented with a mixture of protease inhibitors, and then incubated for 45 min. The sample was centrifuged for 45 min at 15,000 rpm. The supernatant was supplemented with loading buffer and analyzed as described (22).

**Electrophysiology** — Brightly fluorescent isolated HEK 293 cells were selected for recording. Cells were bathed in a solution containing 150 mM NaCl, 2 mM KCl, 2 mM MgCl<sub>2</sub>, 2 mM CaCl<sub>2</sub>, 10 mM glucose, 10 mM HEPES, equilibrated to pH 7.4 with NaOH. Cells were recorded in whole cell mode and placed under the flow of a theta tube pulled to a final opening of ~100 μm mounted on a piezoelectric translator (Physik Instrumente). Currents were evoked by applications of 100 μM GABA for 5 s every minute at ~80 mV and recorded at a sampling frequency of 2 kHz by an EPC10 amplifier (HEKA). GABA was exchanged for gabazine (100 μM) and applied to the cell after 3 min, necessary for complete exchange. Thereafter, the medium was exchanged again for GABA, and we observed complete recovery of the response to the agonist. Data were analyzed with Igor 5 (Wavemetrics).

**Model Building** — The sequences of the human α1, β2, and γ2 GABA receptor subunits were retrieved from the Ligand Gated Ion Channel database (41). The model of the α1β2γ2 receptor was constructed by homology modeling using the structure of the glutamate-gated chloride channel as a template (Protein Data Bank code 3RIF) (42) and sequences alignments obtained with T-coffee (43). Homology modeling was performed with Modeiler version 9.5 (44) using default settings. 100 models were prepared, and the best model according to the Discrete Optimized Protein Energy function (DOPE) was selected. Figures were prepared with PyMOL (77).

**RESULTS**

γ2(R43Q) Displays Increased Clathrin-mediated and Dynamin-dependent Endocytosis — To analyze the impact of the R43Q substitution on intracellular trafficking, the colocalization of wild-type or mutated γ2 with markers of intracellular compartments was assessed in transfected, cultured hippocampal neurons (Fig. 1, A and B). Increased retention in the endoplasmic reticulum (CaR colocalization: 38.3 ± 2.5% for γ2(R43Q)-containing subunits versus 26.7 ± 2.3% for wild-type receptors, n = 31 and 27, respectively; p < 0.005) confirmed previous findings in heterologous cells (HEK293 (24)). However, importantly, γ2(R43Q) colocalization with Golgi apparatus markers did not differ significantly from that of the wild-type subunit, showing that the mutated subunit was also present on the route to the cell membrane (membrin: 19.3 ± 2.4% for the wild-type versus...
15.2 ± 1.6% for the γ2(R43Q)-containing receptor, n = 35 and 38, respectively; ERGIC: 18.6 ± 1.7% for the wild-type versus 13.4 ± 1.9% for the γ2(R43Q)-containing receptor, n = 28 and 34, respectively). We therefore checked whether the absence of surface labeling was due to fast endocytosis mechanisms, as proposed to explain the exclusion of sodium channels from somatic domains (45). When living neurons, expressing γ2(R43Q), were incubated at 37 °C with antibodies directed against the tagged extracellular N-terminal γ2-domain, intracellular punctate labeling was detected (Fig. 1C), suggesting receptor internalization. This was confirmed by the increased colocalization of γ2/γ4(R34Q) with EEA1, an early endosome marker (Fig. 1, D and E) (22.8 ± 4.7% for the wild-type versus 39.5 ± 4.1% for the γ2(R43Q)-containing receptor, n = 14 and 16, respectively, p < 0.005).

Because clathrin-mediated, dynamin-dependent endocytosis is the major neuronal GABA_A_R internalization mechanism (4, 6), we tested whether γ2(R43Q) internalization was driven by a similar pathway. When dynamin was inhibited by incubation with 80 μM dynasore (46), γ2(R43Q) was detected at the plasma membrane of transfected COS-7 cells (Fig. 2A). The ratio of γ2(R43Q) subunit labeled on the cell surface versus γ2(R43Q) labeled within intracellular compartments was increased from 0.42 ± 0.2 to 0.78 ± 0.18 (Fig. 2B). COS-7 cells were then transfected with a β2(LL/AA) mutant; this mutation within the β2 intracellular domain reduces the interaction of GABA_A_R with the AP2 complex (6). In this case, the signal ratio between surface and intracellular γ2(R43Q) increased from 0.18 ± 0.08 to 0.53 ± 0.1 (Fig. 2, C and D), showing that γ2(R43Q) was present on the cell surface when co-expressed with α1 and β2(LL/AA). Taken together, these data showed that the γ2 subunit containing the R43Q mutation increased endocytosis of GABA_A_Rs, hindering their detection on the cell surface.

γ2(R43Q) Endocytosis Is Inhibited by GABA_A_R Antagonists—This increased internalization of a ligand-gated channel, resulting from a mutation within the extracellular domain of the molecular complex, was surprising. This GABA_A_R domain contains binding sites for agonists and allosteric modulators, whereas the intracellular domain mediates interactions with trafficking factors (1, 4, 6, 47). We therefore tested whether GABA_A_R ligands interfered with endocytosis. Incubating transfected neurons (Fig. 3, A and B) or COS-7 cells (Fig. 3, C and D) with two different antagonists (i.e. picrotoxin and gabazine) for 1 h significantly increased γ2(R43Q) surface expression, whereas neither GABA_A_R agonist nor antagonists altered surface expression levels of receptors containing wild-type γ2 subunit. The area of clusters on the surface of neurons transfected with γ2 was 2.23 ± 0.23 mm²/10 μm in control conditions and 2.50 ± 0.19 or 2.36 ± 0.16 in the presence of gabazine (100 μM) or picrotoxin (100 μM), respectively (Fig. 3B). The area of clusters on the surface of neurons transfected with γ2(R43Q) was 0.08 ± 0.02 mm²/10 μm in control conditions and increased to 0.79 ± 0.15 and 0.79 ± 0.12 in the presence of gabazine (100 μM) or picrotoxin (100 μM), respectively (Fig. 3B). In COS-7 cells transfected with α1-, β2-, and γ2 subunits, the surface/intra ratio were 3.8 ± 0.5 in control conditions and 4 ± 0.9 or 3.3 ± 0.4 in the presence of gabazine (100 μM) or picrotoxin (100 μM), respectively (Fig. 3B). In COS-7 cells transfected with α1, β2, and γ2(R43Q) subunits, the surface/intra ratio increased from 0.3 ± 0.1 to 1.27 ± 0.2 for gabazine and 1.41 ± 0.23 for picrotoxin (Fig. 3D). Muscimol (10 μM), a GABA_A_R agonist, had no detectable effect, compared with the control experiments (Fig. 3, B and D). Cell surface biotinylation...
on α1-, β2-, and γ2-transfected COS-7 cells was not modified by 100 μM gabazine + picrotoxin treatment (107 ± 4%), whereas experiments on α1-, β2-, and γ2(R43Q)-transfected cells confirmed that antagonist treatment increased significantly the surface fraction of γ2(R43Q) (213 ± 29% of control, p < 0.05, three independent experiments; Fig. 3, E and F).

The fact that picrotoxin and gabazine are both allosteric GABA_AR antagonists (48–50) indicated that, compared with their wild-type counterparts, the equilibrium between states in γ2(R43Q)-containing GABA_ARs shifted away from the resting state toward the active or desensitized state. Our new homology model based on a glutamate-gated chloride channel (42) suggests that the Arg-43 residue of the γ2 subunit is connected to Tyr = 174 and Glu-178 from the loop B and to Asp-84 and Arg-86 of the β2 subunit via polar interactions (Fig. 4, B–D). In addition, this new model shows that γ2Arg-43 and β2Asp-84/Arg-86 are on loops (Allo1 and Allo2) identified as involved in the motion that opens the channel pore (51). This prompted us to test whether this alteration in the equilibrium between states favored an open channel conformation. We thus co-expressed wild-type or γ2(R43Q) constructs with α1 and β2 subunits in HEK cells. In α1-, β2-, and γ2(R43Q)-transfected cells, maximum GABA-gated currents were 16% (p < 0.01) of currents from cells expressing wild-type receptors (Fig. 4E), in agreement with previous findings (19, 23, 52). Gabazine, an allosteric antagonist, did not cause a significant change in the current base line during whole cell voltage clamp recordings (Fig. 4F) in HEK cells co-expressing α1-, β2-, and either wild-type or γ2(R43Q) subunits (n = 7 and 15, respectively), showing that γ2(R43Q)-containing receptors had no constitutive activity.

Collectively, our data suggested that γ2(R43Q) mutation triggered GABA_AR endocytosis through a structural change in the extracellular domain. Because agonist exposure triggers a long range conformational change (53, 54), our finding on mutated receptors opened the interesting possibility that endocytosis of wild-type GABA_ARs may be modulated by agonists. Our data did not yet indicate significant alteration of the surface level of wild-type γ2-containing receptors in the presence of agonists (Fig. 3). We thus decided to examine directly the impact of agonists on the internalized fraction of GABA_ARs.

GABA_AR Agonist Enhances Receptor Endocytosis—To test a possible link between GABA-induced conformational change and endocytosis, suggested by our studies on γ2(R43Q), we directly quantified internalization by measuring the level of
endogenous α1-containing receptors internalized during agonist application on cultured neurons (Fig. 5A). In control and muscimol experiments, excitatory activity was blocked with 6-cyano-7-nitroquinolxene-2,3-dione, d-2-amino-5-phosphonovaleric acid, and tetrodotoxin in the medium. Neurons were incubated with an antibody directed at the extracellular N-terminal domain of α1 subunit for 30 min at 37 °C. Labeled receptors remaining at the cell surface were labeled with saturating concentration of Alexa Fluor-647 secondary antibodies, internalized receptors were labeled with Alexa Fluor-488 secondary antibodies (Fig. 5A, green), whereas total α1 subunits were labeled with a monoclonal antibody and Alexa Fluor-568 secondary antibodies (Fig. 5A, red). Quantification of the area of punctate labeling of internalized receptors showed a significant increase when neurons were incubated with 10 μM muscimol (the area of intracellular punctate labeling/10 μm increased to 3.14 ± 0.36 μm² from 1.62 ± 0.17 in control condition). This increase was abolished when muscimol was co-applied with 100 μM gabazine or picrotoxin (1.68 ± 0.15 or 1.89 ± 0.08 μm², respectively). Application of gabazine or picrotoxin alone had no effect (1.69 ± 0.19 or 1.61 ± 0.16 μm², respectively) (Fig. 5B).

In another set of experiments, we analyzed γ2 subunit internalization in COS cells transfected with α1, β2, and tagged γ2 subunits (Fig. 6A). Average values ± S.E. from four independent experiments and 105 cells for each condition were plotted (Fig. 6B). These experiments showed that the number of internalized γ2 subunits increased during agonist treatment. The area of intracellular punctate labeling rose from 21.41 ± 3.28 μm² in the control cells to 59.70 ± 6.89 μm² when cells were treated with 100 μM GABA at 37 °C for 30 min. This increase was abolished when GABA was applied with gabazine, picrotoxin, or picrotoxin + gabazine (27.36 ± 3.29, 21.72 ± 2.82, or 21.41 ± 3.28 μm², respectively). When compared with control, gabazine or picrotoxin had no significant effect on the area of intracellular punctate labeling (36.39 ± 5.42 or 28.89 ± 4.09, respectively).

We also assessed γ2 internalization by performing a biotinylation assay on COS-7 cells transfected as above. After labeling with cleavable biotin, plates were returned to 37 °C for 30 min to allow endocytosis. Cells were then exposed to MESNA, which cleaved biotin from proteins remaining on the surface, allowing determination of intracellular biotinylated receptors. As shown in Fig. 6, C and D, application of GABA induced a significant increase of the internalization (212 ± 40% of control, n = 6, p < 0.05) that was suppressed when agonists were co-applied with GABA (99.5 ± 27%).

We next examined whether agonist-induced internalization reduced the amount of receptors on the cell surface (Fig. 6E) by surface biotinylation experiments. Comparison of the relative surface/total ratio, in the absence or the presence of GABA (Fig. 6, E and F), showed that the ligand did not change significantly the amount of receptors on the surface. Expression is normalized to 1 for control and is 1.08 ± 0.11 for GABA-treated cells and 1.06 ± 0.12 for antagonist-treated cells; n = 3 independent experiments (Fig. 6F). These data implied that GABAₐRs endocytosis may be compensated for by exocytosis or reinsertion of GABAₐRs. Indeed, inhibition of receptor recycling with monensin (55, 56) (Fig. 6G) reduced significantly the surface expression of GABAₐRs following application of 10 μM muscimol. Expression is normalized to 1 for control and is 0.722 ± 0.034 for muscimol-treated cells (80 cells for each condition, five independent experiments), suggesting that removal and insertion of GABAₐRs on the cell surface is a use-dependent process that is tightly regulated.
DISCUSSION

Modulation of surface stability of GABA<sub>2</sub>Rs is essential for regulating the physiological properties of inhibitory neurotransmission. Modification of inhibitory signaling and altered receptor trafficking has been associated with several neurological diseases, including schizophrenia, substance abuse, pain, or epilepsy (2, 11, 57–61). It is also established that GABA<sub>2</sub>Rs endocytosis is regulated through intracellular signaling pathways (1, 2, 4). Here we show that a mutation in the N-terminal domain and ligand application have both an influence on receptor endocytosis. Thus, our data uncover a mechanism that links the extracellular domain of GABA<sub>2</sub>Rs to their stability on the cell surface. It must be noted that it was possible that we had detected the traffic of recombinant γ2 subunits in

FIGURE 4. The γ2(R43Q) mutation is not associated with an open channel conformation. A, schematic diagram of an α1β2γ2 GABA<sub>2</sub>R, which illustrates the five combined subunits that form the complex, the two GABA active binding sites at the β2 and α1 interfaces (gray circles), and the benzodiazepine (BDZ; blue circle) allosteric binding site at the α1 and γ2 interface. In current models, γ2Arg-43 is at the interface with β2. B–D, GABA<sub>2</sub>R model viewed from the outside (B), the top (C), and inside (D); only the γ2 and β2 subunits are shown for clarity (γ2 in green and β2 in blue). γ2 (Arg-43, Tyr-174, and Glu-178) and β2 (Asp-84 and Arg-86) residues are represented by sticks. These residues are within loops, shown in gray (γ2 subunit) or dark blue (β2), identified as being involved in the channel-opening motion. E, GABA application on HEK cells transfected with α1β2γ2 or α1β2γ2(R43Q) activated membrane currents, whereas gabazine application (F) did not change the holding current in either type of receptor.

FIGURE 5. GABA<sub>2</sub>R internalization in neurons is increased by muscimol. A, internalized labeling was obtained after incubating live cultured hippocampal neurons with a polyclonal antibody directed at the extracellular domain of the native α1 subunit at 37 °C (Internalized). Total staining of the same neurons was performed after permeabilization and labeling with a monoclonal antibody directed at the intracellular domain of the native α1 subunit (total). Arrows, labeling of internalized receptors; arrowheads, labeling of the intracellular domain. Scale bar, 10 μm. B, quantitative analysis of the area of intracellular punctate labeling of internalized receptors formed by α1 subunits in dendrites of neurons incubated in culture medium (control), with muscimol, gabazine (gbz), or picrotoxin (ptx); a total of four experiments on independent neuron cultures where carried out, with the number of quantified neurons going from n = 32 to 40. **, p < 0.005. Error bars, S.E.
monomer form because it has been shown in some heterologous cells transfected with this subunit alone. However, we have previously shown that this is unlikely (22). Furthermore, in the present work, we found that the amount of \( \gamma_2(\text{R43Q}) \) detected on the cell surface increased in the presence of a \( \gamma_2 \) subunit bearing the LL/AA substitution or following treatment with gabazine, a competitive antagonist of the GABA binding site. Because \( \gamma_2(\text{LL/AA}) \) should be associated with \( \gamma_2(\text{R43Q}) \) to increase cell surface labeling and the gabazine binding site is within the interface between \( \alpha \) and \( \beta \) subunits (48), these experiments strongly suggest that \( \gamma_2(\text{R43Q}) \) detected on the cell surface is part of an oligomer also containing \( \alpha \) and \( \beta \) subunits. The same holds true for wild type \( \gamma_2 \) subunit endocytosis promoted by GABA (or muscimol), the binding site of which is at the \( \alpha/\beta \) interface (Fig. 4A).

Analysis of \( \gamma_2(\text{R43Q}) \) fate in neurons showed an increased retention in the endoplasmic reticulum, in agreement with previous findings in human embryonic kidney 293-T cells (24). Additionally, we provide evidence that \( \gamma_2(\text{R43Q}) \) is not entirely retained within the reticulum and is transported via

**FIGURE 6.** GABA\(_4\)R internalization in COS-7 cells is increased by GABA. A, COS-7 cells were transiently co-transfected with \( \alpha_1-\), \( \beta_2-\), and Myc-tagged \( \gamma_2 \) subunits. Internalized labeling (red) was obtained after incubating live cells with polyclonal antibody directed at the extracellular Myc tag at 37 °C followed by stripping of antibodies remaining on the surface. Intracellular staining (green) of the same cells was obtained after permeabilization and labeling with a monoclonal antibody directed at the same tag (white frames detailed in lower panels). Scale bar, 10 μm. B, quantification of the area of intracellular punctate labeling of internalized receptors formed by \( \gamma_2 \) subunit in cells incubated in culture medium (Ctrl) with GABA, gabazine (GBZ), or picrotoxin (PTX). ***, \( p < 0.001 \). C, COS-7 cells transfected as in A were labeled with biotin and returned to 37 °C in control medium, GABA, gabazine, or picrotoxin, as indicated. Biotin remaining on the surface after the incubation at 37 °C was removed by cleaving. Total and internalized (biotinylated) \( \gamma_2 \) subunits were detected with an anti-Myc antibody. GAPDH staining shows that intracellular proteins were not biotinylated. D, means of internalized/total expression ratio normalized to total \( \gamma_2 \) from experiments described in C. **, \( p < 0.005 \); *, \( p < 0.01 \). E, COS-7 cells transfected as in A and C and labeled with biotin. Representative Western blot shows Myc-tagged \( \gamma_2 \) in total or surface-biotinylated extracts. F, means of surface/total expression ratio normalized to total \( \gamma_2 \) from experiments described in E show that surface expression was unchanged by treatment with GABA. G, quantitative analysis of \( \alpha_1\beta_2\gamma_2 \) GABA\(_4\)Rs surface expression in COS-7 cells transiently co-transfected with \( \alpha_1-\), \( \beta_2-\), and Myc-tagged \( \gamma_2 \) subunits and incubated with monensin with or without muscimol for 30 min before surface and total immunofluorescence labeling. A value of 1 for cells incubated without muscimol was used. **, \( p < 0.005 \). Error bars, S.E.
the Golgi apparatus to the cell membrane, where mutated receptors are highly internalized via a clathrin- and dynamin-dependent mechanism. Blockade of endocytosis leads to a major increase in γ2(R43Q) surface targeting in neurons and COS-7 cells, indicating that internalization is a major mechanism for down-regulating cell surface expression of γ2(R43Q)-containing receptors.

We also show that gabazine or picrotoxin increases dramatically the surface expression of γ2(R42Q) subunit. These two GABA_2R antagonists are both negative allosteric modulators, acting at different sites (48–50, 62). The gabazine-and picrotoxin-sensitive internalization of γ2(R43Q) suggested that endocytosis could be linked to a constitutive activity of the mutated receptor. Electrophysiological recordings of γ2(R43Q)-expressing cells clearly showed that this mutated subunit did not give rise to constitutive currents. Therefore, the effect of antagonists on γ2(R43Q) endocytosis is probably related to another conformational state (e.g. the desensitized state). Interestingly, it has been shown that the γ2(R43Q) mutation favors desensitized states (52).

Consequently, our findings, showing that GABA_2R antagonists prevent γ2(R43Q) endocytosis, suggest that internalization is driven by a global conformational change. Molecular models show that the γ2Arg-43 residue is at the γ2/β2 interface in the extracellular domain, on a loop positioned above the pocket, which is homologous to the GABA binding sites. Interestingly, many mutations in nicotinic receptors linked to diseases are at the interface between receptor subunits (63); they alter the gating allosterically (i.e. from a distance) (63–64). A model indicates that γ2Arg-43 and γ2Glu-178 are connected through a bifurcated salt bridge; this model has been used to study the γ2(R43Q) mutation (22, 26, 52). One of these studies has suggested that these positions have a long range allosteric effect (52).

In our new GABA_2R model derived from the glutamate-gated chloride channel (42), the Arg-43 residue of the β2 subunit is connected to Tyr-174 and Glu-178 from the loop B and to the β2 subunit via polar interactions that should be sensitive to the R43Q substitution and positioned on a loop thought to be involved in the channel pore opening motion (51). Moreover, electrophysiological recordings and kinetic analyses have shown that the long distance effects of γ2(R43Q) substitution extend as far as the transmembrane domains (52). Therefore, γ2(R43Q) mutation might have an influence on receptor endocytosis in line with the current views on pentameric ligand-gated ion channels, describing a link between extracellular, transmembrane, and intracellular domains (53, 65, 66).

Because ligand binding in the Cys-loop receptor family is followed by a whole chain of interconnections, including the extracellular domain (67), it is of interest to assess whether GABA binding may influence GABA_2R endocytosis. Although it is established that the number of surface GABA_2Rs is regulated by constitutive endocytosis and neuromodulation through intracellular signaling (5–8, 68, 69), previous findings on ligand-independent or -dependent internalization are conflicting (28–31). Several studies have investigated GABA_2 receptor internalization following agonist application (29, 30). However, data were obtained by analyzing the amount of receptors remaining at the surface after agonist application. Here, we used a different approach (i.e. quantification of internalized receptors). Altogether, biochemical and immunocytochemical analyses of internalized receptor fraction, both in neurons and in COS-7, showed an increased number of internalized receptors during agonist application (Figs. 5B and 6, B and D), whereas the surface/intracellular ratio or surface labeling (Figs. 3, B and D, and 6E) remained unchanged. These data show that an overall counting of receptors on the cell membrane may overlook an increased endocytosis.

It has been suggested that internalization of GABA_2Rs or increase in neuronal activity is accompanied by insertion of new receptors (4, 7, 70–72). Here, all of the experiments performed on neurons were conducted in the presence of tetrodotoxin and glutamate receptor inhibitors, suggesting that agonist binding endocytosis associated with the insertion of new receptors should instead represent an additional homeostasis mechanism. Knowing that internalized receptors are recycled back to the surface membrane or targeted for degradation and that endocytosis may be compensated for by surface targeting of distinct receptor subtypes, this balance between agonist-induced removal and insertion of receptors may regulate the number, but also the identity, of GABA_2Rs on the cell surface. Because the functional and pharmacological properties of GABA_2Rs depend on subunit composition, our findings showing that ligand stimulation increased endocytosis imply that this mechanism may be an important process for a fine tuning of GABAergic neurotransmission (4). It is of note that benzodiazepines (allosteric modulators of GABA_2Rs) induce a subtype-specific change via enhanced degradation rather than alterations in receptor insertion or endocytosis, thus revealing another mechanism that might regulate GABAergic neurotransmission (73).

The γ2(R43Q) mutation is directly linked to epilepsy (16, 17). Thus, our findings, revealing a direct relationship between receptor endocytosis and a neurological disorder, are in line with the emerging concept that GABA_2 trafficking deficiencies are key factors in initiating and maintaining several diseases, including epilepsy (4, 6, 15). For example, status epilepticus leads to enhanced GABA_2 endocytosis (28–30). The K289M substitution in the γ2 subunit known to be responsible for generalized epilepsy with febrile seizures plus alters the membrane diffusion of GABA_2Rs (61). It must be also noted that a shortened lifetime caused by the epilepsy mutation A322D on α1-containing GABA_2Rs has been proposed (74) (but also see Ref. 75). Interestingly, experiments on γ2(R43Q) knock-in mice and transfected neurons or COS-7 cells have shown that α1 and α3 subunit surface expression was not reduced, despite a dramatic decrease in γ2(R43Q) surface labeling (16, 22). Earlier data, together with our present findings, show that the consequence of the mutation is a complex and dynamic process, suggesting that the defect is an active phenomenon instead of a mere static retention of mutated receptors in the intracellular compartments and that γ2(R43Q)-containing receptor internalization is associated with a compensatory insertion of distinct GABA_2 subtypes.
CONCLUSION

γ2(R43Q)-containing GABA_ARs are in a conformational state that promotes internalization, providing evidence for a direct link between GABA_AR endocytosis and epilepsy. Furthermore, our data suggest that GABA_AR endocytosis is use-dependent, consistent with a model in which ligand binding induces a conformation of the receptor that is a substrate for the biochemical events leading to endocytosis (70). Because GABA is the main inhibitory neurotransmitter in the brain and GABA_ARs are the target of many drugs, this property may have important functional and pathophysiological implications and should therefore be fully characterized (1, 4). Our data suggest that the γ2(R43Q) mutant is a useful model for this purpose. Our findings also illustrate the fact that mutations offer insights not only into diseases but also receptor physiology (61, 64, 74, 76). It would be also of interest to assess whether the different allosteric drugs acting on GABA_ARs have an influence on receptor trafficking.

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