The γ -aminobutyric acid type A receptor **(GABAAR)-associated protein GABARAP interacts with gephyrin but is not involved in receptor anchoring at the synapse**

Matthias Kneussel*, Silke Haverkamp†, Jens C. Fuhrmann*, Hongbing Wang‡, Heinz Wa¨ ssle†, Richard W. Olsen‡, and Heinrich Betz*§

Departments of *Neurochemistry and [†]Neuroanatomy, Max Planck Institute for Brain Research, Deutschordenstrasse 46, D-60528 Frankfurt/Main, Germany; and ‡Department of Molecular and Medical Pharmacology, University of California School of Medicine, Los Angeles, CA 90095-1735

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 γ -Aminobutyric acid type A receptors (GABA_ARs) are ligand-gated **chloride channels that exist in numerous distinct subunit combinations. At postsynaptic membrane specializations, different GABAAR isoforms colocalize with the tubulin-binding protein gephyrin. However, direct interactions of GABAAR subunits with** gephyrin have not been reported. Recently, the GABA_AR-associated protein GABARAP was found to bind to the γ 2 subunit of **GABAARs. Here we show that GABARAP interacts with gephyrin in both biochemical assays and transfected cells. Confocal analysis of neurons derived from wild-type and gephyrin-knockout mice revealed that GABARAP is highly enriched in intracellular compartments, but not at gephyrin-positive postsynaptic membrane specializations. Our data indicate that GABARAP–gephyrin inter**actions are not important for postsynaptic GABA_AR anchoring but may be implicated in receptor sorting and/or targeting mecha**nisms. Consistent with this idea, a close homolog of GABARAP, p16, has been found to function as a late-acting intra-Golgi transport factor.**

p16 | receptor clustering | postsynaptic density | Golgi transport | inhibitory synapse

Neuronal surface membranes contain numerous proteins that serve for intercellular communication. Among these, receptor ion channels, specialized for chemotransmission between neurons, are highly concentrated at postsynaptic densities (PSDs) apposed to the appropriate presynaptic nerve terminals. The targeting of neurotransmitter receptors to and clustering at PSDs is a complex process that requires receptor-associated proteins, cytoskeletal elements, and proteins involved in signal transduction. At excitatory synapses, PDZ domain (from postsynaptic density, disks large, zonula adherens)-mediated protein interactions generate a PSD that serves as a scaffold for glutamate receptors and regulatory enzymes involved in synaptic transmission (for reviews, see refs. 1 and 2). At inhibitory postsynaptic sites, the tubulin-binding protein gephyrin is known to play a crucial role in the synaptic localization of both glycine receptors (GlyRs) and γ -aminobutyric acid type A receptors $(GABA_ARs)$ (for a review, see ref. 3).

Gephyrin (4, 5) was originally identified by copurification with the mammalian GlyR (6, 7). Gephyrin binds with high affinity to polymerized tubulin (8), and both microtubules and actin microfilaments are implicated in GlyR localization at postsynaptic sites (3, 9). Gephyrin also interacts with the GlyR β subunit (10, 11), thus serving as a receptor-cytoskeleton linker. At developing inhibitory PSDs, formation of a gephyrin scaffold precedes GlyR synaptic clustering (12, 13), and GlyR clusters do not form upon loss of gephyrin expression, either by antisense depletion in culture (12) or by gene knockout in mice (14).

Gephyrin is also found at many GABAergic synapses (15–20). Gene depletion and knockout experiments have revealed a crucial role of gephyrin in GABAAR clustering (21–23). Mice deficient for the GABA_AR subunit γ 2 display a strong reduction of postsynaptic gephyrin and GABA_AR clusters, and the synaptic staining for the GABA_AR α 2 and γ 2 subunits observed in cultured cortical neurons is reduced upon antisense oligonucleotide treatment (21). Neurons from gephyrin knockout mice show a total loss of postsynaptic GABA_AR γ 2 and α 2 immunoreactivities, providing conclusive genetic evidence that gephyrin is essential for the postsynaptic localization of certain GABA_ARs (23). However, biochemical evidence for an association of gephyrin with $GABA_AR$ subunits is lacking at present (10), and cotransfection studies have reported only weak colocalization of the β 3 but not other $GABA_AR$ subunits with gephyrin (24).

Recently, the GABA_AR γ 2 subunit was shown to bind the GABAAR-associated protein GABARAP, a 14-kDa polypeptide with sequence similarity to light chain 3 of microtubuleassociated proteins (MAPs) 1A and 1B (25). Consistent with a potential receptor anchoring function, GABARAP harbors a tubulin-binding motif encompassing amino acids 1–36 (25, 26). In cultured cortical neurons, GABARAP displayed punctate immunoreactivity in both cell somata and neurites and colocalized with $GABA_ARs$ (25). To clarify whether $GABARAP$ may serve as a linker between GABA_ARs and gephyrin, we have now analyzed its interaction with gephyrin by genetic and biochemical methods. Moreover, we used immunocytochemistry to study the codistribution of these proteins in spinal cord and retinal sections. Our data show that GABARAP binds to gephyrin but is not found at GABAergic synapses and is localized intracellularly. These observations and GABARAP's close homology to p16, a late-acting intra-Golgi trafficking factor, suggest a role for GABARAP in intracellular receptor transport.

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Abbreviations: PSD, postsynaptic density; GABAAR, y-aminobutyric acid type A receptor; GABARAP, GABAAR-associated protein; GlyR, glycine receptor; GST, glutathione *S*-transferase; EST, expressed sequence tag; ER, endoplasmic reticulum.

[§]To whom reprint requests should be addressed at: Max Planck Institute for Brain Research, Deutschordenstrasse 46, D-60528 Frankfurt/Main, Germany. E-mail: betz@mpihfrankfurt.mpg.de.

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Fig. 1. (*A*) Sequence alignment of GABARAP, p16, and their homologs from *Caenorhabditis elegans*, *Saccharomyces cerevisiae,* and *Arabidopsis thaliana*. The protein p16, recently described as a bovine late-acting intra-Golgi transport factor (30), shares 57.3% amino acid identity with GABARAP. Identities between GABARAP and its homologs are 78.9% (*C. elegans*), 54.7% (*S. cerevisiae*), and 54.2% (*A. thaliana*), respectively. (*B*) Phylogenetic tree of GABARAP, p16, and homolog proteins. Note that mammalian p16 is more distant from GABARAP than its homolog from *C. elegans*.

Materials and Methods

Binding Experiments. The GABARAP cDNA and a GABARAP deletion mutant encoding amino acids 36-117 (GABARAP/36-117) were subcloned into pGEX-2T (Amersham Pharmacia) to generate glutathione *S*-transferase (GST) fusion proteins. Plasmids were transformed for isopropyl β -D-thiogalactopyranoside (IPTG)induced protein production into *Escherichia coli* DH5 α or BL21. Adult rat brains (3 g wet weight) were homogenized in 30 ml of homogenization buffer [100 mM NaCl/10 mM Tris·HCl (pH 7.5)/5 mM EDTA/10 mM MgCl₂/0.5% Nonidet P-40/1% Triton X-100, including protease inhibitors] at 4°C for 60 min. After centrifugation at $10,000 \times g$ for 30 min, 2.4 ml of the resulting supernatants was incubated with glutathione-agarose charged with 5 ^mg of GST, GST-GABARAP, or GST-GABARAPy36–117. After 2 h at 4°C, the beads were washed four times with homogenization buffer. Bound proteins were eluted with 5 mM glutathione $/50$ mM Tris (pH 8.0), separated by SDS/PAGE, and analyzed by Western blotting with anti-gephyrin (Transduction Laboratories, Lexington, KY, 1:250), as described (23).

Cell Culture and Transfection. PC12 cells were cultured in Dulbecco's minimal essential medium supplemented with 10% (vol/vol) fetal calf serum and 5% (vol/vol) horse serum and seeded onto fibronectin-coated glass coverslips 1 day before transfection. Transfection was performed with 800 ng of DNA for 15 min at room temperature, using Lipofectamine Plus Reagent (Life Technologies, Eggenstein, Germany). Cells were then washed in phosphate-buffered saline and cultured in OptiMEM (Life Technologies) for 3–5 h at 37°C. Then the medium was exchanged for standard culture medium (see above), and cells were cultured for another 18 h at 37°C in an

atmosphere of 5% $CO₂/95%$ air before being processed for immunostaining.

Cortical neurons were prepared as low-density cultures in coculture with astrocyte feeder layers as described (23). Cells were cultured for 7 days before being analyzed.

Cryostat Sections of Spinal Cord and Retina Tissue. Spinal tissue of embryonic day 19.5 mice was cut into 5-mm blocks and fixed in 4% (wt/vol) paraformaldehyde for 10 min, followed by a short wash in phosphate-buffered saline. For cryoprotection, the sections were incubated in increasing concentrations 10% (wt/vol), 20% (wt/vol), 30% (wt/vol) + 0.01% sodium azide] of sucrose at 4°C for 1 h each. Cryostat sections were refixed for 5 min in 4% (wt/vol) paraformaldehyde and processed for immunofluorescence. For retina sections, tissue of adult (8–10 weeks) mice was prepared and processed for immunofluorescence as described (18).

Antibodies. For Western blotting, a monoclonal anti-gephyrin antibody (Dianova, Hamburg, Germany) was used at a dilution of 1:250. For double-labeling experiments, the primary antibodies GABARAP 6402 (25) (1:100), anti-GABA_AR γ 2 subunit (27) (1:2000), anti-bassoon (28) (1:8000), and anti-gephyrin (1:100) were used.

Immunocytochemistry and Confocal Microscopy. Coverslips carrying cortical neurons or spinal cord sections were prepared, cultured, and processed for immunofluorescence as described (23). Confocal microscopy was performed with a TCS-SP confocal laserscanning microscope (Leica) equipped with the image software Leica TCS-NT (version 1.6.551).

Table 1. Yeast two-hybrid analysis of protein–protein interactions

Protein tested	GABARAP	p16
$GABA_\text{A}R_{\text{V}}1/350-444$	$^+$	
$GABA_\text{A}R\gamma$ 2/351-445	$^+$	
$GABA$ _A R _y 3/333-446		
$GABAAR0/327-429$		
Gephyrin, full-length		
Gephyrin/2-152		
Gephyrin/153-348	$^+$	
Gephyrin/349-736		

GABARAP and p16 bait constructs were used in binary interaction assays with GABAAR subunit and gephyrin sequences. Positive interactions revealed by $lacZ$ and $LEU2$ expression are marked with a $+$, whereas $-$ denotes white cells.

Quantification of Immunoreactive Colocalization. Three micrographs were taken from each double-labeled section with the $100\times$ objective, using red and green fluorescence filters, and printed at a final magnification of $1,500\times$. The immunofluorescent puncta of the micrographs were transferred onto tracing paper. Micrograph images of double-labeled sections were superimposed at their correct position for counting the number of colocalized puncta. As a control for random overlap, images were also superimposed after a 180° turn to determine the number of coincidentally superimposed puncta. For each measurement, between 850 and 1,700 puncta, taken from at least three sections, were sampled (29).

Yeast Two-Hybrid Experiments. Bait and prey plasmids were cotransfected with the *lacZ* reporter plasmid pSH18–34 into yeast strain EGY48 and assayed for *lacZ* and *LEU2* expression. The cDNAs of the proteins to be analyzed were cloned into the plasmids pGILDA or pJG4–5 (Origene, Rockville, MD). The following amino acid residues, encoded by the partial cDNA fragments, were used in the binary interaction assays: rat

Fig. 2. GABARAP interacts with gephyrin *in vitro*. GST-GABARAP or GST-GABARAP/36–117 (4 μ g each) was bound to glutathione-agarose and incubated with brain homogenate. After washing, bound proteins were eluted with glutathione, separated by SDS/PAGE, and probed with anti-gephyrin. Brain homogenate input (lane 1), GST-GABARAP (lane 2), and GST-GABARAP/36-117 (lane 3) interact with gephyrin, whereas GST alone (lane 4) does not. The GABARAP sequences fused to GST are schematically given (bars) above the gel lanes; putative interaction domains are indicated by arrows.

Fig. 3. Colocalization of GABARAP and gephyrin upon heterologous expression in PC12 cells. Singly expressed GABARAP protein is located at the plasma membrane (*A*), whereas singly expressed gephyrin is diffusely distributed with some enrichment at submembranous compartments (*B*). Coexpression of GABARAP and gephyrin leads to a recruitment of gephyrin to GABARAP-rich loci (C). (Scale bars, 10 μ m.)

GABA_AR γ 1/350–444, rat GABA_AR γ 2S/351–445, rat GABA_AR γ 3/333–446, rat GABA_AR δ /327–429, rat gephyrin/2–733, rat gephyrin/2–152, rat gephyrin/153–348, rat gephyrin/349–736, GABARAP/37–117, and p16/37–117.

Results

GABARAP Is Homologous to p16, a Protein That Functions in Intra-Golgi Transport and Has Homologs in Different Invertebrate Species. Database searches revealed a bovine homolog (GenBank accession no. AF020262; ref. 30), termed p16, that shares 57.3% amino acid identity with GABARAP (Fig. 1). The p16 protein was described as a late-acting 16-kDa intra-Golgi transport factor and was originally purified from bovine brain cytosol (30). The human p16 protein is termed ganglioside expression factor 2 (GenBank accession no. NP009216, unpublished) and shares 100% amino acid identity with the bovine protein. Additional homology searches revealed related sequences (Fig. 1) from *Caenorhabditis elegans* (GenBank accession no. Q09490), *Saccharomyces cerevisiae* (Genbank accession no. CAA84899), and *Arabidopsis thaliana* (GenBank accession no. AAD24645); the last has been classified as a putative microtubule-associated

Fig. 4. GABARAP immunoreactivity in embryonic day 19.5 spinal cord sections derived from wild-type $(+/+)$ and gephyrin-knockout $(-/-)$ mice. Anti-GABARAP stains punctate structures in spinal cells of wild-type tissue; in addition, a diffuse staining is seen (*A*). Gephyrin appears in synaptic clusters of wild-type tissue (*B*, and see ref. 14). In tissue sections derived from gephyrindeficient mice, GABARAP immunoreactivity is unaltered (*C*), whereas gephyrin immunoreactivity is lost (D). (Scale bar, 10 μm.)

protein (31). These observations suggest a widespread function of p16 in many organisms that is not restricted to the nervous system, a finding that is consistent with GABARAP's ubiquitous mRNA expression (25). Searching mouse expressed sequence tag (EST) databases with the nucleotide sequence of p16 identified ESTs obtained from various tissues. These included hypothalamus (GenBank accession no. AA968244), liver (Gen-Bank accession no. AI528168), kidney (GenBank accession no. AW318968), pancreas (GenBank accession no. AV056273), mammary gland (GenBank accession no. AA822987), lymph node (GenBank accession no. AA275333), tongue (GenBank accession no. AV082088), stomach (GenBank accession no. AV076945), and diaphragm (GenBank accession no. AA065794), suggesting that p16 is a ubiquitous protein with essential functions in all cells. Interestingly, ESTs corresponding to the p16 nucleotide sequence were also obtained from blastocysts (GenBank accession no. AA795530), indicating expression during early embryonic development.

GABARAP, But Not p16, Binds Gephyrin. In the yeast two-hybrid system, GABARAP gave strong positive signals for an interaction with the cytoplasmic loop regions of the GABAAR subunits γ 1 and γ 2 but not of the γ 3 and δ subunits (Table 1). To investigate whether GABARAP might also bind to the receptoranchoring protein gephyrin, we analyzed its interaction with either full-length gephyrin or partial gephyrin sequences, corresponding to amino acids 2–152, 153–348, or 349–736 (Table 1). Whereas full-length gephyrin did not give any β -galactosidase signal indicative of protein interaction, the fragment corresponding to amino acids 153–348 showed significant interaction (Table 1). In contrast, p16 did not interact with any of the analyzed GABA_AR subunits (γ 1, γ 2, γ 3, and δ), or with gephyrin or its fragments (Table 1).

To further investigate the GABARAP–gephyrin interaction *in vitro*, we fused GABARAP to GST. Bacterially expressed

Fig. 5. GABARAP staining of cultured cortical neurons derived from wildtype $(+/+)$, heterozygous $(+/-)$, and homozygous $(-/-)$ gephyrinknockout mice. A punctate distribution of GABARAP (25) is seen throughout the cytoplasm of neurons derived from wild-type (*A*), heterozygous (*B*), and homozygous (*C*) gephyrin-knockout mice. Individual confocal sections are shown. (Scale bar, 10 μ m.)

fusion protein was incubated with rat brain homogenate. After washing, bound protein was eluted, subjected to SDS/PAGE, and probed with anti-gephyrin. Full-length GABARAP as well as a truncated protein containing amino acids 36–117 displayed efficient binding of the 93-kDa protein gephyrin (Fig. 2, lanes 2 and 3). In contrast, GST alone did not interact with gephyrin (Fig. 2, lane 4). Thus GABARAP, which binds the $GABA_AR$ subunit γ 2 (25), also binds gephyrin.

GABARAP and Gephyrin Colocalize in PC12 Cells. To unravel whether GABARAP and gephyrin interact in intact cells, we expressed both proteins in the neuron-like cell line PC12. GABARAP immunoreactivity was largely found at the plasma membrane (Fig. 3*A*), as revealed by confocal microscopy. Untransfected cells also displayed some endogenous GABARAP immunoreactivity, which again was mainly located at the membrane (not shown). This localization suggests the existence of membrane binding sites for GABARAP in PC12 cells. In contrast, single expression of gephyrin generated a largely diffuse cytoplasmic staining (Fig. 3*B*). Coexpression of both GABARAP and gephyrin caused a recruitment of cytoplasmic gephyrin to

Subcellular Distribution of GABARAP in Neurons of Wild-Type and Gephyrin-Knockout Mice. To unravel whether the interaction of GABARAP and gephyrin seen *in vitro* and in transfected cells may result in a synaptic colocalization of these proteins, we analyzed spinal cord sections and cultured cortical neurons from wild-type and gephyrin-deficient mice for GABARAP immunoreactivity. In spinal cord sections, the punctate distribution of GABARAP was less prominent than it was in cultured cortical neurons (25) but was still detectable (Fig. 4). Notably, sections derived from wild-type $(+/+)$ and homozygous gephyrinknockout mutants $(-/-)$ did not significantly differ in the intensity and/or number of GABARAP immunoreactive puncta (Fig. 4) or show colocalization with gephyrin (not shown). Similarly, in cultured cortical neurons derived from wild-type and mutant mice, GABARAP puncta were not significantly different in number, size, and location between the different genotypes (Fig. 5), suggesting that GABARAP functions prior to or independently of gephyrin. More importantly, confocal images revealed that the majority of GABARAP puncta were seen in intracellular compartments, which according to their location represented putative endoplasmic reticulum (ER) and Golgi structures (Fig. 5). These data suggest that, in contrast to what is seen upon expression in transfected PC12 cells, in neurons GABARAP predominantly functions intracellularly.

Quantitative Analysis of GABARAP, GABA AR, and Gephyrin Immunoreactivity in Retina. To quantitatively evaluate whether GABA_AR, GABARAP, and gephyrin colocalize at synaptic sites, we used retina sections, in which gephyrin is known to be exclusively synaptic, as revealed by electron microscopy (18). Such sections were stained for GABARAP, gephyrin, the GABA_AR γ 2 subunit, and the presynaptic protein bassoon (28). Notably, GABARAP and gephyrin immunoreactivities failed to show any significant colocalization above randomized values (Fig. 6 *B* and E). Furthermore, none of the \rm{GABA} _AR γ 2 subunit (Fig. 6 C and *E*) or the bassoon (Fig. 6 *D* and *E*) immunoreactive puncta colocalized with GABARAP, whereas 64% of GABA_AR γ 2 subunit immunoreactive puncta colocalized with synaptic gephyrin (Fig. 6 *A* and *E*). These data indicate that GABARAP is not clustered at inhibitory synaptic sites, suggesting that GABARAP's interactions with the GABA_AR γ 2 subunit and gephyrin may be important for cellular functions other than receptor anchoring.

Fig. 6. Cofocal micrographs of vertical sections through the inner plexiform layer (IPL) of double-immunostained mouse retinae. Selected areas of the micrographs (frames) are shown at higher magnification, to the right. (*A*) Gephyrin (red) and the γ 2 subunit of the GABA_AR (green) are aggregated in synaptic hot spots, which are often colocalized. (*B*) GABARAP (red) shows diffuse, punctate distribution in the IPL but is not clustered with gephyrin (green) in synaptic hot spots. The white arrows point to the cell bodies of ganglion cells and show the expression of GABARAP in the cytoplasm. (*C*) GABARAP (red) and the γ 2 subunit of the GABA_AR (green) appear not to be aggregated within the same hot spots. (*D*) The presynaptic cytomatrix protein bassoon (green), which is clustered at both excitatory and inhibitory synapses, is not colocalized with GABARAP. (Scale bar, 10 μ m.) (*E*) Quantifications of the colocalizations at their correct superpositions and at random superpositions. Only in the case of the γ 2 subunit of the GABA_AR and gephyrin was a significant colocalization of puncta observed.

Discussion

Here we have used yeast two-hybrid analysis as well as biochemical and anatomical methods to investigate whether GABARAP may mediate the specific localization of $GABA_ARs$ at gephyrinrich PSDs. In yeast, a partial gephyrin sequence, but not full-length gephyrin, gave a strong β -galactosidase signal, which is indicative of protein interaction with GABARAP. Furthermore, a recombinant GST-GABARAP fusion protein displayed high-affinity *in vitro* binding to gephyrin solubilized from brain. Upon heterologous coexpression in the neuron-like PC12 cell line, GABARAP and gephyrin were found to colocalize, indicating that these proteins can interact in a cellular environment. Because gephyrin has been shown to be essential for GABAAR clustering (21–23), these data are consistent with the proposal of Wang *et al.* (25) that GABARAP may be involved in the processes that underlie $GABA_AR$ targeting to and/or clustering at postsynaptic sites.

Despite the observed interaction of the two proteins, the comparison of spinal cord sections and cultured cortical neurons from wild-type and gephyrin-deficient mice did not reveal any differences in the number, size, or location of GABARAP immunoreactive puncta. Moreover, in wild-type retinae there was no significant colocalization of GABARAP with either the $GABA_A\overline{R}$ γ 2 subunit or gephyrin, whereas gephyrin-positive structures showed highly significant colocalization with the GABAAR. These observations indicate that GABARAP is not clustered at gephyrin-rich PSDs and is therefore unlikely to serve as a GABAAR-gephyrin linker molecule at differentiated synapses. Indeed, confocal sections of cultured cortical neurons revealed that the punctate GABARAP immunoreactivity is mainly found in intracellular compartments. According to their location, these GABARAP-positive sites are likely to represent $ER/Golgi$ structures. This location is consistent with p16, a homolog of GABARAP, functioning as a late-acting intra-Golgi transport factor in mammalian brain (30). Thus GABARAP may be implicated in intracellular receptor sorting and targeting processes that precede and/or initiate receptor clustering at the synapse. Indeed, coexpression of GABARAP and GABA_ARs in QT6 cells has been found to increase the fraction of clustered GABAARs (L. Chen, H.W., S*.* Vicini, and R.W.O., unpublished results). In any case, the absence of GABARAP from GABAer-

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gic postsynaptic sites raises the question of whether there are yet-unidentified proteins that interact with both GABA_ARs and gephyrin and thus cause their synaptic colocalization.

Intracellular vesicular protein traffic includes different steps, such as budding, targeting, docking, and fusion of vesicles with their target membranes. Each step requires a different set of cytosolic factors. Its homology to p16 protein suggests that GABARAP may participate in a step preceding insertion of GABAARs into the plasma membrane. The tubulin cytoskeleton is known to be crucial for the organization of the Golgi complex (32) and the spatial distribution of peroxisomes (33). Furthermore, microtubules provide the structural basis for organelle transport via kinesin/dynein motor proteins, and a Golgi network-associated protein was recently found to directly bind to microtubules (34). The assignment of a putative tubulin-binding site to the N-terminal region of GABARAP (25, 26) is consistent with this protein having a role in the transport of GABA_ARcontaining Golgi vesicles before vesicle fusion with the plasma membrane. The yeast two-hybrid data obtained here show that GABARAP binds different GABAAR subunits and that only GABARAP and not its homolog p16 interacts with gephyrin. These data point to a role for GABARAP in receptor trafficking processes that may involve both different $GABA_AR$ isoforms and the receptor anchoring protein gephyrin. Whether gephyrin is transported to synaptic sites in vesicle-bound form, however, is not yet solved. Furthermore, it remains to be elucidated whether gephyrin can interact with inhibitory receptor subunits already at the level of the ER/Golgi compartment. If so, plasma membrane insertion of vesicles containing preassembled receptor–gephyrin complexes may provide a GABARAP-driven mechanism to allow for postnatal growth of inhibitory PSDs.

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