We examined the interaction between ephrins and metabotropic glutamate (mGlu) receptors in the developing brain and cultured neurons. EphrinB2 coimmunoprecipitated with mGlu1a receptors, in all of the brain regions examined, and with mGlu5 receptors in the corpus striatum. In striatal slices, activation of ephrinB2 by a clustered form of its target receptor, EphB1, amplified the mGlu receptor-mediated stimulation of polyphosphoinositide (PI) hydrolysis. This effect was abolished in slices treated with mGlu1 or NMDA receptor antagonists but was not affected by pharmacological blockade of mGlu5 receptors. An interaction among ephrinB2, mGlu1 receptor, and NMDA was supported by the following observations: (1) the NR1 subunit of NMDA receptors coimmunoprecipitated with mGlu1a receptors and ephrinB2 in striatal lysates; (2) clustered EphB1 amplified excitatory amino acid-stimulated PI hydrolysis in cultured granule cells grown under conditions that favored the expression of mGlu1a receptors; and (3) clustered EphB1 amplified the enhancing effect of mGlu receptor agonists on NMDA toxicity in cortical cultures, and its action was sensitive to mGlu1 receptor antagonists. Finally, fluorescence resonance energy transfer and coclustering analysis in human embryonic kidney 293 cells excluded a physical interaction between ephrinB2 and mGlu1a (or mGlu5 receptors). A functional interaction between ephrinB and mGlu1 receptors, which likely involves adaptor or scaffolding proteins, might have an important role in the regulation of developmental plasticity.

Key words: mGlu receptors; ephrinB; postnatal development; polyphosphoinositide hydrolysis; cerebellar granule cells; NMDA toxicity

Introduction

The evidence that ephrin (Eph) receptors associate and interact with NMDA receptors at synaptic sites (see below) provides an example of how molecules that are classically involved in developmental patterning and cell–cell communication (Kullander and Klein, 2002) may regulate excitatory neurotransmission and synaptic plasticity. Eph receptors and their ligand ephrins are subdivided into two classes named A and B. With few exceptions, A- and B-subclass ephrins bind to EphA and EphB, respectively, thus generating a bidirectional signaling (Holland et al., 1996; Kullander and Klein, 2002; Himanen et al., 2004). Although Ephs constitute the largest family of tyrosine kinase receptors, ephrin ligands signal through the engagement of adaptor or scaffolding molecules. The cytoplasmic tail of ephrinB is phosphorylated when the EphB receptor is presented by neighboring cells or as soluble fusion protein (Bruckner et al., 1997; Parker et al., 2004), thereby recruiting proteins containing Src homology 2 (SH2)/SH3 or postsynaptic density-95/Discs large/zona occludens-1 (PDZ) domains (Torres et al., 1998; Bruckner et al., 1999; Cowan and Henkemeyer, 2001; Palmer et al., 2002). Dalva et al. (2000) first showed a direct association between the N-terminal domains of EphB2 and the NMDA receptor 1 (NR1) subunit of NMDA receptors. This interaction is promoted by the presence of ephrinB acting in “trans” (i.e., either exogenously added or present on a different cell). In cultured cortical neurons, activation of EphB by ephrinB2 potentiates NMDA receptor-dependent function, suggesting a mechanism whereby activity-dependent and -independent signals converge in the regulation of synaptic plasticity (Takasu et al., 2002). Although in the above studies EphB receptors are shown to act postsynaptically and ephrinB5 presynaptically, Grunwald et al. (2004) have recently shown that the Eph/ephrin system is used in an inverted manner in the hippocampus, where postsynaptic ephrinB2 and NMDA receptors interact in the induction of long-term forms of synaptic plasticity.

Group I metabotropic glutamate receptors (mGlu1 and mGlu5 receptors) interact with NMDA receptors and are involved in the regulation of developmental plasticity. Expression of these receptors is developmentally regulated (Nicoletti et al., 1986a; Schoepf and Johnson, 1989; Minakami et al., 1995; Roman et al., 1996; Casabona et al., 1997) and is temporally related to critical times of experience-dependent synaptic modifications (Dudek and Bear, 1989). Group I mGlu receptors are cross-linked with NMDA receptors through a chain of anchoring proteins (Tu et al., 1999), and their activation amplifies NMDA cur-
proteins were resuspended in SDS-bromophenol blue buffer containing protein A-Sepharose beads (Sigma-Aldrich). For Western blot analysis, 3 mM EGTA, 1.5 mM MgCl2, 10% glycerol) containing 1 mM benzamidine, 3,3'-diaminobenzidine tetrachloride (Vector Laboratories). Color development was achieved by incubating the cells in PBS and then incubated for 1 h with the ABC Elite reagent (Vector Laboratories). Mouse ephrinB2 cDNA (kindly provided by Dr. David Wilkin-10 receptors. The interaction was continued for 60 min and then terminated by the addition of methanol:chloroform:water (1:1:1). The amount of [3H]inositol monophosphate (InsP) accumulated during the reaction was measured as described previously (Nicoletti et al., 1986a).

Measurement of agonist-stimulated polyphosphoinositide hydrolysis in cultured cerebellar granule cells. Primary cultures of cerebellar granule cells were prepared from 8-d-old rats, as described previously (Nicoletti et al., 1986b), and grown onto 35 mm Nunc Petri dishes in basal Eagle’s medium (Invitrogen, Milan, Italy) containing 10% fetal calf serum, 10% heat-inactivated horse serum, 0.05 mg/ml gentamycin, and 25 mM (K25) or 10 mM (K10) Ca2+ (added as CaCl2). Cytosine arabinofuranoside (10 μM) was added 16–18 h after plating to avoid the replication of non-neuronal cells. Cultures grown in K10 medium were used after 4 DIV, whereas cultures grown in K25 medium were used at 9 DIV. Western blot analysis of mGlu1a receptor, mGlu5 receptor, or ephrinB2 was performed as described above. For the stimulation of polyphosphoinositide (PI) hydrolysis, cultures were incubated overnight with [3H]inositol (2 μCi/dish), washed, and incubated for 30 min at 37°C under constant oxygenation. Receptor agonists were added, and the incubation was continued for 30 min. Clustered EphB1/Fc or MK-801 was added 5 min before receptor agonists. [3H]InsP formation was measured as described above.

Preparation of mixed cultures of mouse cortical cells and assessment of excitotoxic neuronal death. Cortical cultures containing both neurons and astrocytes were prepared from fetal mice at 14–16 d of gestation, as described previously (Rose et al., 1992). Briefly, dissociated cortical cells were plated onto 15 mm multiwell vessels (Falcon Primaria, Lincoln Park, NJ) on a layer of confluent astrocytes with minimum essential medium (MEM)–Eagle’s salts supplemented with 5% heat-inactivated horse serum, 5% fetal bovine serum, glutamine, 0.05 mg/ml gentamycin, and 25 mM (K25) or 10 mM (K10) Ca2+ (added as CaCl2). Cytosine arabinofuranoside (10 μM) was added 3–5 DIV. Cultures were exposed to 10 μM cytosine arabinofuranoside for 1–3 days before treatment with 1–3 mM QA or 1–3 mM quisqualate, 1 μM NBQX, and dizocilpine (MK-801) for the detection of GFAP. Photographs were taken at 20× using an inverted microscope.

Measurement of polyphosphoinositide hydrolysis in brain slices. Slices (350 × 350 μm) were prepared from the corpus striatum or cerebral cortex of P6–7 rats, as described previously (Nicoletti et al., 1986a), and incubated at 37°C under constant oxygenation for 45 min in Krebs-Henseleit buffer, pH 7.4. Forty microliters of gravity packed slices were transferred to vials containing 1 μCi of [3H]inositol (specific activity, 10 Ci/μmol; Amersham Biosciences) to label inositol phospholipids. At the end of this incubation, 10 μl LiCl was added, followed 10 min later by mGlu receptor agonists or carbamylcholine. When present, mGlu receptor antagonists and/or EphB1/Fc were added 5–10 min before receptor agonists. The incubation was continued for 60 min and then terminated by the addition of methanol:chloroform:water (1:1:1). The amount of [3H]inositol monophosphate (InsP) accumulated during the reaction was measured as described previously (Nicoletti et al., 1986a).
by these restriction enzymes (Roche, Vienna, Austria). Ligation was performed with the Fast-Link Ligation kit (Epicerent Technologies, Madison, WI). Sequence of the obtained construct was confirmed by sequencing. The plasmids encoding mGlu1 and mGlu5 receptors tagged with cyan fluorescent protein (CFP) on the C termini were kindly provided by Dr. Laurent Fagni (Centre CNRS-INSERM de Pharmacologie-Endocrinologie, Montpellier, France). Human dopamine D2 receptor was cloned into pECPF vector (Clontech), as described by Schmid et al. (2001). Preparation of the human serotonin transporter (hSERT) tagged with CFP and yellow fluorescent protein (YFP) (CFP-hSERT-YFP) has been described previously (Just et al., 2004).

Cell culture and transfections. Human embryonic kidney 293 (HEK293) cells were grown in DMEM containing 10% fetal bovine serum, and 50 mg/L gentamicin on 10 cm diameter cell culture dishes at 37°C in an atmosphere of 5% CO2, 95% air. One day before transfection, cells were replated to obtain subconfluent cultures either on poly-D-lysine-covered glass coverslips (22 mm in diameter and placed into 6-well plates; 3 × 10^5 cells/well plate). Transient transfections were performed using the standard calcium phosphate precipitation method.

Fluorescence resonance energy transfer microscopy. Fluorescence microscopy was performed using a Zeiss (Thornwood, NY) Axiovert 200M inverted epifluorescence microscope equipped with a CoolSNAP fx cooled CCD camera (Photometrics; Roper Scientific; Tucson, AZ). The fluorescence filter sets were purchased from Chroma (Chroma Technology, Brattleboro, VT) [CFP filter set: excitation, 436 nm; dichroic mirror, 455 nm; emission, 480 nm; YFP filter set: excitation, 500 nm; dichroic mirror, 515 nm; emission, 535 nm; fluorescence resonance energy transfer (FRET) filter set: excitation, 436 nm; dichroic mirror, 455 nm; emission, 535 nm]. Coverslips with attached cells were mounted in the microscope chamber and put on the microscope stage. Images of cells with CFP- and YFP-tagged proteins were acquired through corresponding filter channels. To measure donor recovery after acceptor photobleaching (DRAP), we acquired a donor (CFP) image before (Ib) and after (Ia) photobleaching using the YFP setting for 90 s (excitation, 500 nm; dichroic mirror, 525 nm; emission, 535 nm). DRAP was quantified by FRET efficiency (E) as described by Miyawaki and Tsien (2000) according to the following equation: E = (Ia − Ib)/Ia.

Results

Coimmunoprecipitation of ephrinB2 with group I mGlu receptors in brain tissue

Expression of ephrinB2 was detected in the corpus striatum, cerebral cortex, hippocampus, and cerebellum of P6–7 rats (Fig. 1A). Lysates of brain tissue were immunoprecipitated with anti-ephrinB2 antibodies and then immunoblotted with mGlu1a, mGlu5, or mGlu2/3 receptor antibody. The mGlu1a receptor was detected in immunoprecipitates from all brain regions (Fig. 1B), whereas the mGlu5 receptor was mainly detected in immunoprecipitates from the corpus striatum (Fig. 1C). The mGlu5 receptor was also detected in ephrinB2 immunoprecipitates from the striatum of P6 wild-type mice but not in immunoprecipitates from mGlu5 knock-out mice (Fig. 1D). The mGlu2/3 receptor proteins were not detected in ephrinB2 immunoprecipitates (Fig. 1F). Neither mGlu1a nor mGlu5 receptors were detected in immunoprecipitates of ephrinA1 (Fig. 1G,H). We also examined the coimmunoprecipitation among the NR1 subunit of NMDA receptors, the mGlu1 receptor, the mGlu5 receptor, and ephrinB2 in the developing rat striatum. We could detect the NR1 subunit in ephrinB2, mGlu1, and mGlu5 receptor immunoprecipitates (Fig. 1E). We extended the study to RGS3, which is known to interact with ephrinB2 through its PDZ domain (Su et al., 2004), and Homer proteins, which are the prototypical scaffolding proteins interacting with mGlu1a and mGlu5 receptors (Brakeman et al., 1997). In control brain tissue, RGS3 was detected as a band at ~90 kDa, as expected. RGS3 could be detected in ephrinB2, mGlu1a, and mGlu5 receptor immunoprecipitates from the corpus striatum, cerebral cortex, and cerebellum (Fig. 2A). Homer proteins were detected using a pan-Homer polyclonal antibody. In control brain tissue, the antibody labeled two bands at ~30 and 43–45 kDa corresponding to the molecular size of Homer-1a and Homer-1b/c, respectively, and additional bands of higher molecular size. The immunoreactive band corresponding to Homer-1b/c was detected in ephrinB2 immunoprecipitates from the cortex, hippocampus, striatum, and cerebellum of P6–7 rats (Fig. 2B).

Functional interaction between ephrinB2 and group I mGlu receptors in brain slices

We assessed the activity of group I mGlu receptors by measuring the stimulation of PI hydrolysis in slices prepared from the corpus striatum or the cerebral cortex of P6–7 rats. Of the three agonists classically used in this assay (i.e., 1S,3R-ACPD, quisqualate, and DHPG), we excluded 1S,3R-ACPD, because stimulation of PI hydrolysis by this drug is partially mediated by the activation of mGlu2/3 receptors (Genazzani et al., 1994; Schoepf et al., 1996). In striatal slices, quisqualate induced a concentration-dependent stimulation of [3H]InsP formation, with an apparent
EC50 value of ~100 nM (Fig. 3A). Both mGlu1 and mGlu5 receptors contributed to this response, as shown by the inhibitory action of LY367385 (a competitive mGlu1 receptor antagonist) and MPEP (a noncompetitive mGlu5 receptor antagonist) (Fig. 3B). To activate ephrinB2, we incubated the slices with a chimeric EphB1/Fc receptor clustered with an anti-Fc IgG. Clustered EphB1/Fc (0.5 μg/ml) added to striatal slices increased the potency of quisqualate in stimulating [3H]InsP formation without affecting the efficacy of the drug (Fig. 3A). Clustered EphB1/Fc failed to induce changes in the PI response to the muscarinic cholinergic receptor agonist charbamylcholine (Table 1). No potentiation of quisqualate-stimulated PI hydrolysis was observed after application of a nonclustered form of EphB1/Fc (0.5 μg/ml; [3H]InsP formation expressed as dpm/mg protein: basal, 2700 ± 160; clustered EphA1/Fc, 2440 ± 85; quisqualate, 300 nM, 5300 ± 440; quisqualate plus EphB1/Fc, 5050 ± 115), an Eph member that does not bind to ephrinB2 (Pasquale, 2004). We examined the amplification of the PI response by clustered EphB1/Fc in the presence of a battery of glutamate receptor antagonists. Clustered EphB1/Fc still potentiated the action of quisqualate in the presence of MPEP but became ineffective in slices treated with LY367385. The AMPA receptor antagonist, NBQX (10 μM), had no effect on quisqualate-stimulated PI hydrolysis in both the absence and presence of clustered EphB1/Fc; in contrast, the NMDA receptor antagonist MK-801 (1 μM) did not affect the action of quisqualate alone but abolished the amplification of the PI response to quisqualate induced by clustered EphB1/Fc (Fig. 3B).

We extended the analysis to DHPG, which behaves as a selective mGlul/5 receptor agonist (for review, see Schoepp et al., 1999). Similarly to that observed with quisqualate, addition of clustered EphB1/Fc to striatal slices also increased the potency of DHPG in stimulating PI hydrolysis (Fig. 3C).
Table 1. EphB1/Fc selectively amplifies the mGlu receptor-mediated stimulation of PI hydrolysis in striatal slices

<table>
<thead>
<tr>
<th></th>
<th>Controls</th>
<th>Clustered EphB1/Fc (0.5 μg/ml)</th>
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<tbody>
<tr>
<td>Basal</td>
<td>3500 ± 68</td>
<td>3200 ± 130</td>
</tr>
<tr>
<td>Quisqualate</td>
<td>6800 ± 320</td>
<td>8500 ± 240*</td>
</tr>
<tr>
<td>Ch 100 μM</td>
<td>7100 ± 140</td>
<td>7920 ± 450</td>
</tr>
<tr>
<td>Ch 1 mV</td>
<td>15,000 ± 620</td>
<td>14,500 ± 850</td>
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Values are means ± SEM of 9–12 determinations. *p < 0.05 (Student’s t test) versus the respective basal values.

Figure 4. Clustered EphB1/Fc fails to amplify mGlu receptor-mediated PI hydrolysis in cortical slices from P6–7 rats. Note that the PI response to quisqualate (A) or DHPG (B) was attenuated by MPEP but not by the mGlu1 receptor antagonists CPCCOEt and LY367385. Values are means ± SEM of 12 determinations. #p < 0.05 (one-way ANOVA plus Fisher’s PLSD) compared with the respective values obtained with quisqualate or quisqualate plus clustered EphB2/Fc without receptor antagonists.

Figure 5. Western blot analysis of mGlu1a receptor, mGlu5 receptors, and ephrinB2 in cultured cerebellar granule cells grown in medium containing 25 mM K⁺ (K25) at 9 DIV or in medium containing 10 mM K⁺ (K10) at 4 DIV is shown. Expression in the rat cerebellum (CB) or striatum (STR) is also shown.

In cortical slices, the stimulation of PI hydrolysis by quisqualate or DHPG was antagonized by the mGlu5 receptor antagonist, MPEP, but it was not affected by the mGlu1 receptor antagonists LY367385 or CPCCOEt. Using this preparation, we could not observe any effect of clustered EphB1/Fc on excitatory amino acid-stimulated PI hydrolysis (Fig. 4A, B).

Interaction between ephrinB2 and group I mGlu receptors in cultured cerebellar granule cells

We examined the effect of clustered EphB1/Fc on group I mGlu receptor signaling in cultured cerebellar granule cells at 9 DIV grown in medium containing 25 mM K⁺ (K25). These cultures expressed ephrinB2, high levels of mGlu1a receptors, and low levels of mGlu5 receptors (Fig. 5). The addition of clustered EphB1/Fc did not stimulate PI hydrolysis, per se, but enhanced the stimulation produced by low concentrations of quisqualate (100 nM). No potentiation was observed when cultures were stimulated by higher concentrations of quisqualate (1 or 100 μM). Clustered EphB1/Fc also amplified the PI response to 10 μM glutamate but not to carbachol (10 μM) (Table 2). Knowing that a component of the PI response to glutamate in granule cells is mediated by the activation of NMDA receptors (Nicoletti et al., 1986b), we examined [³H]InsP₃ formation in the presence of the NMDA receptor antagonist MK-801 (1 μM). Clustered EphB1/Fc failed to potentiate the PI response to glutamate in the presence of MK-801, indicating that activation of NMDA receptors is necessary for the modulation of mGlu receptors by ephrinB/EphB (Table 2). We also used young cultured granule cells (4 DIV) grown in 10 mM K⁺ (K10), which expressed mGlu5 receptors and very low levels of mGlu1 receptors (Fig. 5) (Copani et al., 1998). In these cultures, stimulation of PI hydrolysis by 1 μM quisqualate was not affected by clustered EhsB1/Fc (Table 3), supporting the specificity of ephrinB/EphB for cells expressing mGlu1 receptors.

Interaction between ephrinB2 and group I mGlu receptors in mixed cultures of cortical cells: study on NMDA toxicity

We extended the study of the interaction between ephrinB and group I mGlu receptors to mixed cultures of mouse cortical cells. These cultures are widely used for the study of excitotoxic neuronal death and carry the advantage of maintaining the physiological interplay between neurons and astrocytes. Using this model, we (and others) have shown that activation of either mGlu1 receptor or mGlu5 receptor amplifies NMDA toxicity (Bruno et al., 1995, 2000, 2001; Buisson and Choi, 1995; Battaglia et al., 2001). Western blot analysis confirmed the presence of mGlu1 and mGlu5 receptors in these cultures and also showed the expression of ephrinB2 (Fig. 6A). Immunohistochemistry showed that both mGlu1a receptors and ephrinB2 were expressed by neurons in mixed cortical cultures, whereas no expression was found in the underlying monolayer of astrocytes identified as GFAP” cells (Fig. 6B). This contrasts with the widespread expression of mGlu5 receptors, which were also found in astrocytes (data not shown). The lack of ephrinB2 and mGlu1a receptors in glial cells was confirmed using pure cultures of mouse cortical astrocytes (Fig. 6B). Mixed cultures were challenged with a 10 min pulse of 30 μM NMDA for the induction of excitotoxic neuronal death. As expected, NMDA toxicity was amplified by the mGlu1/5 receptor agonist quisqualate (100 nM or 1 μM) or DHPG (3, 30, or 300 μM). Clustered EphB1/Fc (0.5 μg/ml) applied during the toxic pulse further amplified the enhancing effect of quisqualate or DHPG on NMDA toxicity, producing only small effects on its own. In addition, quisqualate alone has no discernible effect on cell survival even when combined with clustered EphB1 (Fig. 6C,D; Table 4) (data not shown). Nonclustered EphB1/Fc was inactive (data not shown). To examine which group I mGlu receptor subtype was positively modulated by clustered EphB1/Fc, we treated the cultures with the mGlu5 receptor antagonist MPEP or with the mGlu1 receptor antagonists CPCCOEt or LY367385. These three antagonists were similarly effec-
tive in reducing NMDA toxicity both in the absence and in the presence of quisqualate or DHPG (Bruno et al., 2000). However, clustered EphB1/Fc could still amplify responses to quisqualate or DHPG when mGlu5 receptors were antagonized by MPEP (3 μM) but became inactive when mGlu1 receptors were blocked by CPCCOEt (10 μM) or LY367385 (1 μM) (Table 4).

FRET analysis of the interaction between group I mGlu receptors and ephrinB2 in transfected HEK293 cells

To address the question of whether mGlu1 and mGlu5 directly associate with ephrinB2, we used FRET microscopy on transiently transfected HEK293 cells. We fused ephrinB2 cDNA via its C terminus in frame with YFP. Plasmids encoding the obtained construct and mGlu1-CFP and mGlu5-CFP were transiently coexpressed in HEK293 cells. We were unable to observe an increase in fluorescence of donor (CFP) after acceptor photobleaching in both cases (mGlu1-CFP plus ephrinB2-YFP and mGlu5-CFP plus ephrinB2-YFP) (Fig. 7A), which is indicative of little or no direct interaction within the pairs of proteins. FRET relies on the distance between the fluorophores and their relative orientation. An interaction may escape detection if the two fluorophores are separated by a distance that is larger than the Förster distance (50–100 Å in the case of YFP and CFP) and if their relative orientation is unfavorable (e.g., because of rational constraints). We therefore used an independent method to assess the interaction between mGlu1/5-CFP and ephrinB2-YFP chimeras. For that purpose, we coexpressed both proteins in HEK293 cells and applied the preclustered EphB1-Fc for 20 min at 37°C. EphrinB2-YFP was clustered into punctate aggregates in cell membrane and internalized. However, both mGlu1-CFP and mGlu5-CFP did not redistribute together with ephrinB2, which would be expected provided the receptors directly associate with the latter. Similar experiments were also performed in the presence of 50 μM DHPG, which induced internalization of mGlu1 and mGlu5 receptors; the presence of the mGlu receptor agonist did not facilitate coclustering (Fig. 7B). This indicates once again that the interaction between mGlu1 or mGlu5 and ephrinB2 is unlikely to be direct.

**Discussion**

We moved from the evidence that both the Eph/ephrin system and group I mGlu receptors interact with NMDA receptors and are involved in the regulation of synaptic plasticity during development and in adulthood. Group I mGlu receptors, mGlu1 and mGlu5 receptors, are both coupled to Gq-proteins, and their activation stimulates PI hydrolysis with ensuing intracellular Ca2+ release and activation of protein kinase C (De Blasi et al., 2001). We found here that ephrinB2 coimmunoprecipitates with mGlu1a receptors in different regions of rats at P6–7. Coimmunoprecipitation between ephrinB2 and mGlu5 receptors was mainly detected in the corpus striatum. The specificity of these data was strengthened by the lack of mGlu1a or mGlu5 receptors in ephrinA1 immunoprecipitates. Interestingly, ephrinB2, mGlu1a, and mGlu5 receptors also communoprecipitated with the NR1 subunit of NMDA receptors. This suggests that ephrins associate with mGlu receptors and that association with NMDA receptors is not restricted to EphB receptors (for review, see Palmer and Klein, 2003) but extends to ephrinB ligands. The latter possibility has been raised by Gruwal et al. (2004), who showed that postsynaptic ephrinB2 and NMDA receptors interact in the induction of long-term potentiation in the hippocampus. It is noteworthy that group I mGlu receptors are also involved in the regulation of hippocampal synaptic plasticity (Bortolotto et al., 1999).

Based on FRET and redistribution analysis in HEK293 cells, we conclude that a direct interaction between ephrinB2 and mGlu1 or mGlu5 receptors is unlikely to occur. However, an interaction at the C terminus could have been masked by the large fluorescent protein tags or, alternatively, requires additional proteins that are not present in HEK293 cells. We found that ephrinB2 communoprecipitates with Homer proteins in brain tissue. These proteins interact with the C-terminus domain of mGlu1 or mGlu5 receptors (Brakeman et al., 1997; Xiao et al., 2000) and regulate receptor targeting (Ango et al., 2000, 2002) and coupling to other receptors, adaptors, and signaling proteins (Tu et al., 1999; Kammermeier et al., 2000). The use of a pan-Homer antibody revealed that ephrinB2 coimmunoprecipitated with putative Homer-1b/c but not with the shorter isoform, Homer-1a. Interestingly, only the long isoforms of Homer (such as Homer-1b and -1c) can be involved in the formation of multiprotein complexes (Xiao et al., 2000). Long isoforms of Homer mediate physical interaction between group I mGlu and NMDA receptors through additional anchoring proteins (Tu et al., 1999).

We examined whether a functional interaction exists between group I mGlu receptors and ephrinB2 using three different models as follows: (1) brain slices prepared from the striatum of P6–7 rats; (2) primary cultures of rat cerebellar granule cells; and (3)
mixed cultures of mouse cortical cells. We focused on striatal slices because of the established role for the ephrinB2/EphB1 system in the development of the nigro-striatal pathway and in drug-induced striatal plasticity (Yue et al., 1999; Halladay et al., 2000). EphrinB2 activated by clustered EphB1/Fc increased the potency of mGlu receptor agonists in enhancing PI hydrolysis in striatal slices, suggesting an increased ligand affinity at mGlu receptors or an increased efficiency of receptor signaling. Potentiation was abolished by mGlu1 (but not mGlu5) receptor antagonists, indicating that activated ephrinB2 specifically amplified responses mediated by mGlu1 receptors. Interestingly, NMDA receptor blockade by MK-801, which did not affect the PI response by itself (Nicoletti et al., 1986a,b), inhibited the amplifying activity of activated ephrinB2. This discloses an unexpected NMDA component in the PI response to mGlu receptor agonists, which becomes unmasked after activation of ephrinB2. Although the underlying mechanism is unclear, we suggest the following models. It is possible that activated ephrinB2 relieves the Mg\(^{2+}\) blockade of the NMDA channel (the slice incubation buffer contained 1.2 mM Mg\(^{2+}\)), thus allowing the activation of NMDA receptors by the endogenous glutamate. Ca\(^{2+}\) influx would then amplify mGlu1 receptor signaling by preventing receptor desensitization (Alagarsamy et al., 1999). Alternatively, activated ephrinB2 might promote a cascade of reactions by primarily facilitating mGlu1 receptors. Activation of protein kinase C would then relieve the Mg\(^{2+}\) blockade of the NMDA receptor (Chuang et al., 2000), thus allowing Ca\(^{2+}\) influx and a additional amplification of mGlu1 receptor signaling. These hypotheses do not exclude that other protein partners are involved. PDZ-containing regulators of G-protein signaling (RGS) proteins are likely candidates, because they are known to bind ephrinB (Lu et al., 2001) and potentially interact with both mGlu1a and NMDA receptors through their PDZ domain. RGS proteins accelerate the GTPase activity of the \(z\) subunit of the G-proteins, thus inhibiting signal transduction at G-protein-coupled receptors (Dohlman and Thorner, 1997). At least two RGS proteins (i.e., RGS2 and RGS4) inhibit mGlu1a receptor signaling (Saugstad et al., 1998; Kammermeier et al., 1999). The PDZ-RGS, RGS3, coimmunoprecipitated with both ephrinB2 and mGlu1a receptors, raising the possibility that both mGlu1a and NMDA receptors act in cis on the same membrane domain, presumably at the postsynaptic site of striatal neurons, where all these proteins are predominantly found in the first week of postnatal life (Yue et al., 1999; Smith et al., 2000). This is somehow in contrast with studies showing that it is rather the EphB receptor that interacts in cis with NMDA receptors on the postsynaptic membrane (Dalva et al., 2000) but may be consistent with the recent finding by Grunwald et al. (2004) (see above).

Group I mGlu receptors might be a part of a ternary complex with ephrinB2 and NMDA receptors at postsynaptic densities, thus regulating different aspects of synaptic plasticity not only in
the striatum but also in other brain regions. The widespread nature of this interaction is suggested by results obtained with the two additional models (i.e., cultured cerebellar granule cells and mixed cultures of mouse cortical cells). Cultured cerebellar granule cells are usually grown in medium containing 25 mM K\textsubscript{+}, a condition that mimics the excitatory drive provided by mossy fibers in the intact cerebellum. If extracellular K\textsubscript{+} concentration is lowered to 25 mM, cells are viable only for 4 d and then die by apoptosis (Gallo et al., 1987). Activated ephrinB2 potentiated glutamate or quisqualate-stimulated PI hydrolysis only under growth conditions that favored the expression of mGlu1a receptors (i.e., in K25 cultures at 9 DIV) (Copani et al., 1998). Potentiation required again the activation of NMDA receptors, which here contributed to the overall response to glutamate even in the absence of clustered EphB1/Fc (Nicoletti et al., 1986c, 1987). This is not surprising, because glutamate may activate AMPA or kainate receptors expressed by mature granule cells (Condorelli et al., 1993), thus depolarizing neuronal membranes and relieving the Mg\textsuperscript{2+} blockade of the NMDA channel.

Mixed cultures of mouse cortical cells have been widely used for the characterization of how group I mGlu receptors modulate excitotoxic neuronal death. mGlu1/5 receptor agonists may either facilitate or reduce NMDA toxicity depending on the functional state of mGlu receptors (i.e., naive vs preactivated) and on the presence of astrocytes (Bruno et al., 1995, 2001; Buisson and Choi, 1995; Nicoletti et al., 1999). In contrast, mGlu1 or mGlu5 receptor antagonists are consistently neuroprotective (for review, see Bruno et al., 2001). Using an experimental protocol in which mGlu1/5 receptor agonists enhance NMDA toxicity (Bruno et al., 1995), we found that activated ephrinB2 further amplified the action of these drugs and that amplification was abrogated by mGlu1 receptor antagonists. Both ephrinB2 and mGlu1 receptors were expressed by neurons and not by astrocytes in cortical cultures, although the presence of ephrinB in astrocytes has been reported previously (Conover et al., 2000; Bundesen et al., 2003). Our data raise the possibility that ephrins are involved in processes of neurodegeneration/neuroprotection.

In conclusion, we have provided evidence for a novel form of interaction among ephrinB2, mGlu1a receptors, and NMDA receptors in the developing brain. Although the precise subcellular localization of this interaction is unknown, we suggest that all of these proteins act in cis in the same membrane domain. Interestingly, studies on striatal slices disclose an NMDA component in the PI response to mGlu receptor agonists that was not recognized in previous studies. Whether activated ephrinB2 primarily facilitates NMDA receptor activation (which in turn potentiates mGlu1a receptor signaling) or directly amplifies mGlu1a responses (thus promoting a vicious circle leading to a secondary activation of NMDA receptors and then to a further amplification of mGlu1a signaling) remains to be determined. Whatever the mechanism(s), our observations introduce a new partner (i.e., the mGlu1a receptor) in the ephrinB/EphB-NMDA receptor network. This might have important implications in the regulation of developmental plasticity, associative learning, and even in the regulation of pain threshold, which involves the ephrin/Eph system (Battaglia et al., 2003) as well as group I mGlu receptors.
receptors (Neugebauer, 2002; Varney and Gereau, 2002). It will be interesting to examine whether strategies aimed at preventing ephrinB2 activation will reduce mGlu1 receptor signaling and disrupt the physiological interplay between group I mGlu receptors and NMDA receptors. This might be relevant to the experimental treatment of neurodegenerative disorders and chronic pain.

References


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