

RESEARCH PAPER

Immuno-pharmacological characterization of group II metabotropic glutamate receptors controlling glutamate exocytosis in mouse cortex and spinal cord

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BACKGROUND AND PURPOSE

We recently proposed the existence of mGlu₃-preferring autoreceptors in spinal cord terminals and of mGlu₂-preferring autoreceptors in cortical terminals. This study aims to verify our previous conclusions and to extend their pharmacological characterization.

EXPERIMENTAL APPROACH

We studied the effect of LY566332, an mGlu₂ receptor positive allosteric modulator (PAM), and of LY2389575, a selective mGlu₃ receptor negative allosteric (NAM) modulator, on the mGlu_{2/3} agonist LY379268-mediated inhibition of glutamate exocytosis [measured as KCl-evoked release of preloaded [³H]-D-aspartate]. The mGlu₂ PAM BINA and the mGlu₃ NAM ML337, as well as selective antibodies recognizing the N-terminal of the receptor proteins, were used to confirm the pharmacological characterization of the native receptors.

KEY RESULTS

Cortical synaptosomes possess LY566332-sensitive autoreceptors that are slightly, although significantly, susceptible to LY2389575. In contrast, LY566332-insensitive and LY2389575-sensitive autoreceptors are present in spinal cord terminals. BINA and ML337 mimicked LY566332 and LY2389575, respectively, in controlling LY379268-mediated inhibition of glutamate exocytosis from both cortical and spinal cord synaptosomes. Incubation of cortical synaptosomes with anti-mGlu₂ antibody prevented the LY379268-induced inhibition of glutamate exocytosis, and this response was partially reduced by the anti-mGlu₃ antibody. Incubation of spinal cord synaptosomes with the anti-mGlu₃ antibody abolished LY379268-mediated reduction of glutamate exocytosis from these terminals, while the anti-mGlu₂ antibody was inactive. Western blot analysis and confocal microscopy data were largely consistent with these functional observations.

CONCLUSIONS AND IMPLICATIONS

We confirmed that mGlu₃-preferring autoreceptors exist in spinal cord terminals. Differently, cortical glutamatergic terminals possess mGlu₂/mGlu₃ heterodimers, whose inhibitory effect is largely mediated by mGlu₂ receptors.

Abbreviations

[³H]-D-Asp, [3H]-D-aspartate; BINA, 3'-[[2-(Cyclopentyl-2,3-dihydro-6,7-dimethyl-1-oxo-1H-inden-5-yl)oxy]methyl]-[1,1'-biphenyl]-4-carboxylic acid; LY2389575, (3S)-1-(5-bromopyrimidin-2-yl)-N-(2,4-dichlorobenzyl)pyrrolidin-3-amine methanesulfonate hydrate; LY379268, (1R,4R,5S,6R)-4-Amino-2-oxabicyclo[3.1.0]-hexane-4,6-dicarboxylic acid; LY541850, (1S,2S,4R,5R,6S)-2-amino-4-methylbicyclo[3.1.0]-hexane-2,6-dicarboxylic acid; LY566332, N-(4'-cyano-biphenyl-3-yl)-N-(3-pyridinylmethyl)-ethanesulfonamide hydrochloride; mGlu, metabotropic glutamate; ML337, [2-Fluoro-4-[2-(4-methoxyphenyl)ethynyl]phenyl][(3R)-3-hydroxy-1-piperidinyl]methanone; NAM, negative allosteric modulator; PAM, positive allosteric modulator; TRIS, Tris-(hydroxymethyl)-amino methane; t-TBS, Tris-buffered saline-Tween; VGLUT1, vesicular glutamate transporters type 1

Introduction

The group II metabotropic glutamate (mGlu) receptors comprise **mGlu₂** and **mGlu₃** receptors. The high sequence homology of the two receptors for a long time limited the possibility of distinguishing the respective contribution of mGlu₂ and mGlu₃ units to the expression of native group II mGlu receptors in the CNS (Niswender and Conn, 2010; Nicoletti *et al.*, 2011; 2015). However, there is now evidence demonstrating that mGlu₂ and mGlu₃ receptors are differently expressed in particular CNS regions (Gu *et al.*, 2008; Wright *et al.*, 2013) and in cells (neurons and astrocytes; Ohishi *et al.*, 1993; Tamaru *et al.*, 2001; Copeland *et al.*, 2017) and could have different roles in controlling synaptic transmission (Kew *et al.*, 2002; Corti *et al.*, 2007; Hanna *et al.*, 2013; Sanger *et al.*, 2013). Altogether, these observations support the existence of group II mGlu receptor subtypes.

The recent discovery of new ligands able to separate the role of the two receptors has allowed the pharmacological characterization of the native mGlu_{2/3} receptors. Consistently, we recently proposed the existence of presynaptic mGlu₃-preferring autoreceptors in spinal cord glutamate terminals and presynaptic mGlu₂-preferring autoreceptors in the cortex of adult mice (Di Prisco *et al.*, 2016a). These conclusions originated from the results obtained in release studies with **N-acetyl-aspartyl-glutamate** (NAAG), which has been shown to selectively activate the mGlu₃ receptor subtype (Neale *et al.*, 2011; Romei *et al.*, 2012; Di Prisco *et al.*, 2016a) and LY541850, a selective mGlu₂ receptor agonist with mGlu₃ receptor antagonist activity (Hanna *et al.*, 2013; Sanger *et al.*, 2013).

In this study we aimed to extend our previous work on the pharmacological characterization of the presynaptic mGlu_{2/3} autoreceptors in cortical and spinal cord synaptosomes. We investigated the effects of LY566332 and **BINA**, two positive allosteric modulators (PAMs) of the mGlu₂ receptor subtype (Rorick-Kehn *et al.*, 2005; Benneyworth *et al.*, 2007; Molinaro *et al.*, 2009; Caraci *et al.*, 2011; Sanger *et al.*, 2013), and of LY2389575 and **ML337**, which are selective mGlu₃ receptor negative allosteric modulators (NAMs; Caraci *et al.*, 2011; Wenthur *et al.*, 2013), on the **LY379268**-induced reduction of glutamate exocytosis from both synaptosomal preparations. To further characterize the respective contribution of mGlu₂ and mGlu₃ receptor proteins to the assembly of the mGlu_{2/3} autoreceptors, we applied an unconventional immunological approach. Briefly, cortical and spinal cord synaptosomes were incubated with selective antibodies raised against the outer sequences of the mGlu₂ and the mGlu₃ receptor

proteins in order to determine whether and to what extent these antibodies interfere with the LY379268-mediated inhibition of glutamate exocytosis. These antibodies were also used in Western blot and confocal microscopy analyses to demonstrate the expression of mGlu₃ and mGlu₂ receptors in spinal cord and cortical synaptosomes and to analyse their reciprocal localization. The results from functional and immunochemical studies confirm that the mGlu_{2/3} autoreceptors in spinal cord and cortical terminals differ one each other in terms of mGlu receptor composition and pharmacological profile.

Methods

Animal experiments

Mice (male, strain C57BL/6J) were obtained from Charles River (Calco, Italy) and were housed in the animal facility of DIFAR, Section of Pharmacology and Toxicology. mGlu₂ receptor knockout (mGlu₂^{-/-}) mice, mGlu₃ receptor knockout (mGlu₃^{-/-}) mice, double mGlu_{2/3} receptor knockout (mGlu_{2/3}^{-/-}) mice on a CD1 genetic background and their CD1 wild-type (WT) counterparts were kindly provided by Eli Lilly & Company (Indianapolis, IN, USA) (Linden *et al.*, 2005) and housed at IRCSS Neuromed. These mice were individually genotyped for the mGlu₂ and mGlu₃ receptor gene by PCR. All animals were kept under environmentally controlled conditions (ambient temperature = 22°C, humidity = 40%) on a 12 h light/dark cycle with food and water available *ad libitum*.

Mice were killed by cervical dislocation followed by decapitation and the cortex and/or the spinal cord rapidly removed. The experimental procedures were in accordance with the European legislation (European Communities Council Directive of 24 November 1986, 86/609/EEC) and were approved by the Italian Ministry of Health (DDL 26/2014 and previous legislation; protocol number no. 50/2011-B). Experiments were performed following the Guidelines for Animal Care and Use of the National Institutes of Health. Animal studies are reported in compliance with the ARRIVE guidelines (Kilkenny *et al.*, 2010; McGrath and Lilley, 2015).

Sixty-two C57BL/6 mice were used in the release experiments with selective positive and negative allosteric modulators (39 mice) and with the anti-mGlu₂ and anti-mGlu₃ selective antibodies (23 mice). In line with the 3Rs rules (replacement, refinement and reduction), both cortical and spinal cord synaptosomes were prepared from each animal, and

one preparation was used for release experiments, while the other one, for Western blot analyses or confocal microscopy. Nine genetically modified mice (mGlu2^{-/-} mice, mGlu3^{-/-} mice and mGlu2/3^{-/-} mice, three mice for each group) and three WT CD1 mice were used to confirm the specificity of anti-mGlu₂ and anti-mGlu₃ antibodies.

Preparation of synaptosomes

Mouse cortical and spinal cord purified synaptosomes were prepared as previously described (Musante *et al.*, 2011). Briefly, the tissue was homogenized in 10 volumes of 0.32 M sucrose, buffered to pH 7.4 with Tris-(hydroxymethyl)-amino methane (TRIS; final concentration 0.01 M) using a glass/Teflon tissue grinder (clearance 0.25 mm); the homogenate was centrifuged at 1000× *g* for 5 min to remove nuclei and debris, and the supernatant was gently stratified on a discontinuous Percoll gradient (6, 10 and 20% v/v⁻¹ in Tris-buffered sucrose) and centrifuged at 33 500× *g* for 5 min. The layer between 10 and 20% Percoll (synaptosomal fraction) was subsequently collected and washed by centrifugation. Synaptosomes were resuspended in a physiological solution with the following composition (mM): NaCl, 140; KCl, 3; MgSO₄, 1.2; CaCl₂, 1.2; NaH₂PO₄, 1.2; NaHCO₃, 5; HEPES, 10; glucose, 10; pH 7.2–7.4.

Release experiments

Synaptosomes were incubated for 15 min at 37°C in a rotary water bath in the presence of [³H]-D-aspartate ([³H]-D-Asp; f.c.: 50 nM). In order to study the impact of anti-mGlu antibodies on the LY379268-mediated inhibition of glutamate exocytosis, synaptosomes were incubated for 30 min in the presence of the following antibodies: polyclonal rabbit anti-mGlu₃ (1:1000) or monoclonal mouse anti-mGlu₂ (1:1000); in these experiments, the radioactive tracer was added at *t* = 15 min of incubation. Identical portions of the synaptosomal suspensions were layered on microporous filters at the bottom of parallel thermostated chambers in a Superfusion System (Raiteri *et al.*, 1974, Pittaluga, 2016; Ugo Basile, Comerio, Varese, Italy).

Synaptosomes were transiently (90 s) exposed, at *t* = 39 min, to high KCl-containing medium (12 mM KCl for cortical synaptosomes or 15 mM KCl for spinal cord synaptosomes, NaCl substituting for an equimolar concentration of KCl; Zucchini *et al.*, 2013) in the absence or in the presence of the agonist LY379268. Allosteric modulators were always added concomitantly with the agonist. Fractions were collected as follows: two 3 min fractions (basal release), one before (*t* = 36–39 min) and one after (*t* = 45–48 min) a 6 min fraction (*t* = 39–45 min; evoked release). The collected fractions and the superfused synaptosomes were measured for radioactivity.

The amount of radioactivity released into each superfusate fraction was expressed as a percentage of the total radioactivity. The K⁺-evoked overflow was estimated by subtracting the neurotransmitter content in the first and the third fractions collected (basal release, b1 and b3) from that in the 6 min fraction collected during and after the depolarization pulse (evoked release, b2). The effect of agonists/antagonists is expressed as percentage of the KCl-evoked overflow of tritium observed in the absence of receptor agonists and antagonists (% of control). In the figures

demonstrating the impact of agonists and antagonists on [³H]-D-Asp release, data are always presented as the mean ± SEM of *n* (number within brackets under the bar) independent determinations obtained in separate experiments run in triplicate (three superfusion chambers for each experimental condition) on different days.

Immunoblotting analysis

The cortical and spinal cord purified synaptosomes were lysed in ice-cold lysis buffer (150 mM NaCl, 50 mM Tris, 1% Triton X-100, protease inhibitors, pH 8.0) and quantified for protein content. Samples were boiled for 5 min at 95°C in SDS-PAGE loading buffer. Proteins were then separated by SDS 7.5% PAGE and transferred onto PVDF membranes. Membranes were incubated for 1 h at room temperature in Tris-buffered saline-Tween (t-TBS: 0.02 M Tris, 0.150 M NaCl and 0.05% Tween 20), containing 5% (w/v-1) non-fat dried milk and then probed with mouse monoclonal anti-mGlu₂ receptor antibody (1:1000 for cortical synaptosomes and 1:500 for spinal cord synaptosomes) or rabbit polyclonal anti-mGlu₃ receptor antibody (1:1000) and mouse anti-β-tubulin antibody (1:800) overnight at 4°C. After extensive washes in t-TBS, membranes were incubated for 1 h at room temperature with appropriate horseradish peroxidase-linked secondary antibodies (1:20 000).

In the experiments carried out to verify the specificity of the anti-mGlu₂ and anti-mGlu₃ antibodies, mouse cerebral cortices (from WT and knockout mice) were dissected out and homogenized at 4°C in Tris-HCl pH 7.5, 10 mM; NaCl, 150 mM; SDS 0.1%, EDTA, 5 mM and complete TM protease cocktail tablets. Proteins (20 μg) from supernatants were separated by 8% SDS-PAGE and transferred on immuno-blot PVDF membranes. Membranes were incubated with mouse anti-mGlu₂ (1:1000, 1 h at room temperature, in t-TBS), rabbit anti-mGlu₃ receptor (1:1000, 1 h at room temperature, in t-TBS) and mouse anti-β-tubulin (1:800, overnight at 4°C) antibodies and then for 1 h with the appropriate peroxidase-coupled secondary antibody. Immunoblots were visualized with an enhanced chemiluminescence plus Western blotting detection system. β-tubulin was used as an internal control.

Confocal microscopy and co-localization

Mouse cortical and spinal cord synaptosomes were fixed with 2% paraformaldehyde for 15 min, permeabilized with 0.05% Triton X-100 PBS for 5 min and incubated with the following primary antibodies: mouse anti-mGlu₂ receptor (1:1000), rabbit anti-mGlu₃ receptor (1:1000) and guinea pig anti-vesicular glutamate transporters type 1 (VGLUT1; 1:500). After extensive washes, synaptosomes were incubated for 1 h at room temperature with the following antibodies: donkey anti-rabbit AlexaFluor-488 and goat anti-guinea pig AlexaFluor-633 (1:1000 both, confocal analysis aimed at identifying co-localization of mGlu₃ receptor and VGLUT1 proteins), with donkey anti-mouse AlexaFluor-488 and goat anti-guinea pig AlexaFluor-633 (1:1000 both, confocal analysis aimed at identifying co-localization of mGlu₂ receptor and VGLUT1 proteins), with donkey anti-rabbit AlexaFluor-488 with goat anti-mouse AlexaFluor-633 (1:1000 both, confocal analysis aimed at identifying co-localization of mGlu₂ and mGlu₃ receptor proteins). Synaptosomes were then applied to coverslips (Musante *et al.*, 2008a). Fluorescence images

(512 × 512 × 8 bit) were then visualized by use of a six-channel Leica TCS SP5 laser-scanning confocal microscope, equipped with 458, 476, 488, 514, 543 and 633 nm excitation lines, through a plan-apochromatic oil immersion objective 63X/1.4NA. Light collection configuration was optimized according to the combination of chosen fluorochromes. Sequential channel acquisition was performed to avoid crosstalk. A Leica 'LAS AF' software package was used for image acquisition, storage and visualization. The quantitative estimation of co-localized proteins was performed as described previously (Musante *et al.*, 2008a; Summa *et al.*, 2013), by calculating the 'co-localization coefficients' (Manders *et al.*, 1993). They express the fraction of co-localizing molecular species in each component of a dual-colour image and are based on the Pearson's correlation coefficient, a standard procedure for matching one image with another in pattern recognition. If two molecular species are co-localized, the overlay of their spatial distributions has a correlation value higher than what would be expected by chance alone. Costes *et al.* (2004) developed an automated procedure to evaluate the correlation between the green and red channels with a significance level >95%. The same procedure automatically determines an intensity threshold for each colour channel based on a linear least-square fit of the green and red intensities in the image's 2D correlation cytofluorogramme. Costes's approach was carried out by macro routines integrated as plugins (WCIF Colocalization Plugins, Wright Cell Imaging Facility, Toronto Western Research Institute, Canada) in the ImageJ 1.51p software (Wayne Rasband, NIH, USA).

Calculations and statistical analysis

Sigma plot 10 data analysis and graphing software package were used for data handling/statistics and for graph drawing. ANOVA was performed followed by Dunnett's test or Newman-Keuls multiple comparisons test, as appropriate; direct comparisons were performed by Student's *t*-test. *Post hoc* tests were done only if *F* value was significant. Data were considered significant if *P* < 0.05. Experiments were carried out to quantify the effects of antagonists alone and of pretreatment of the synaptosomes with antibodies on the KCl-evoked release of [³H]-D-Asp. In accord with our previous results (Di Prisco *et al.*, 2016a) and consistent with Curtis *et al.* (2015), at least *n* = 5 replicates were carried out for each experimental condition. The data and statistical analysis comply with the recommendations on experimental design and analysis in pharmacology (Curtis *et al.*, 2015).

Chemicals

[^{2,3}-³H]-D-Asp (specific activity 11.3 Ci·mmol⁻¹) was from Perkin Elmer (Boston, MA, USA). Mouse anti-β-tubulin, horseradish peroxidase-conjugated anti-mouse and anti-rabbit secondary antibodies were from Sigma (Milan, Italy). LY379268, LY566332, ML337 and BINA were purchased from Tocris Bioscience (Bristol, UK). The Western blotting detection system was purchased from GeHealthcare (Italy). Mouse anti-mGlu₂ receptor antibody was from Abcam (Cambridge, UK), while rabbit anti-mGlu₃ receptor antibody was from Alomone Labs (Jerusalem, Israel). Guinea pig anti-VGLUT1 was from Millipore Corporation, (Billerica, MA, USA). Donkey anti-rabbit AlexaFluor-488, goat anti guinea pig

AlexaFluor-633, donkey anti-mouse AlexaFluor-488 and goat anti-mouse AlexaFluor-633 were from Life Technologies Corporation, (Carlsbad, CA, USA). LY2389575 was kindly provided by Dr Monn and Dr McKinzie (Ely Lilly, Indianapolis, IN, USA).

Nomenclature of targets and ligands

Key protein targets and ligands in this article are hyperlinked to corresponding entries in <http://www.guidetopharmacology.org>, the common portal for data from the IUPHAR/BPS Guide to PHARMACOLOGY (Southan *et al.*, 2016), and are permanently archived in the Concise Guide to PHARMACOLOGY 2015/16 (Alexander *et al.*, 2015).

Results

Effects of LY566332 and of LY2389575 on the LY379268-induced inhibition of [³H]-D-Asp exocytosis from cortical glutamatergic synaptosomes

Purified synaptosomes isolated from the cortex of adult male mice were preloaded with [³H]-D-Asp and exposed to a mild (12 mM KCl) depolarizing stimulus (Grilli *et al.*, 2004; Di Prisco *et al.*, 2012) in the absence or in the presence of the mGlu_{2/3} receptor agonist LY379268. Previous studies showed that 3 nM LY379268 causes a significant, submaximal, inhibition of the [³H]-D-Asp exocytosis (Figure 1, see also Di Prisco *et al.*, 2016a). LY566332 (1 μM), unable on its own to modify the 12 mM KCl-evoked [³H]-D-Asp overflow, significantly enhanced the inhibitory effect of LY379268 on tritium exocytosis. A lower concentration of the mGlu₂ receptor PAM (0.1 μM) slightly, although not significantly, modified the

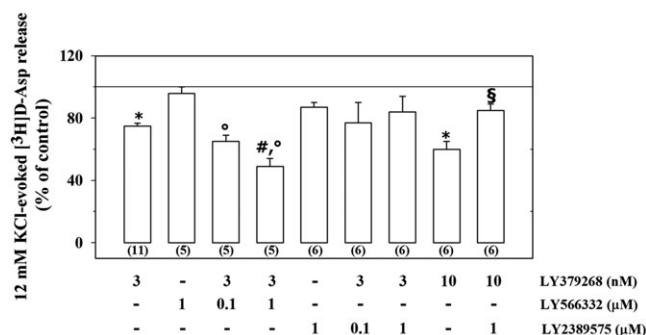


Figure 1

Effects of the mGlu₂ PAM LY566332 and of the mGlu₃ NAM LY2389575 on the inhibition of the 12 mM KCl-evoked [³H]-D-Asp overflow elicited by the mGlu_{2/3} agonist LY379268 in mouse cortical synaptosomes. The 12 mM KCl-evoked release of tritium over the basal release amounted to 1.39 ± 0.36 nCi. Results are expressed as a percentage of the 12 mM KCl-evoked tritium overflow (% of control). Data are the mean ± SEM of *n* (number within brackets) experiments. **P* < 0.05 versus the 12 mM KCl-evoked tritium overflow; °*P* < 0.05 versus the 12 mM KCl/1 μM LY566332-evoked tritium overflow; #*P* < 0.05 versus the 12 mM KCl/3 nM LY379268-evoked tritium overflow; §*P* < 0.05 versus the 12 mM KCl/10 nM LY379268-evoked tritium overflow.

LY379268-mediated inhibition of the 12 mM KCl-evoked [³H]-D-Asp overflow (Figure 1).

The impact of LY2389575 on the 12 mM KCl-evoked [³H]-D-Asp overflow from cortical synaptosomes was also quantified. The inhibition of the 12 mM KCl-evoked release of tritium elicited by LY379268 (3 nM) in the presence of LY2389575 (0.1–1 μM) did not significantly differ from that caused by LY379268 alone, nor was it significantly modified when compared to control. The mGlu₃ receptor NAM (1 μM), however, significantly modified the inhibitory effect elicited by a higher concentration (10 nM) of LY379268 on the 12 mM KCl-evoked release of tritium from cortical synaptosomes. LY2389575 failed to modify significantly, on its own, the 12 mM KCl-evoked release of [³H]-D-Asp (Figure 1).

Effects of LY566332 and of LY2389575 on the LY379268-induced inhibition of [³H]-D-Asp exocytosis from spinal cord glutamatergic synaptosomes

Exposure of spinal cord synaptosomes to 15 mM KCl-enriched solution causes the Ca²⁺-dependent release of preloaded [³H]-D-Asp from these terminals (Di Prisco *et al.*, 2013; 2016a). The concomitant presence of LY379268 (30 pM) significantly reduced this release (Figure 2; see also Di Prisco *et al.*, 2013; 2016a). LY566332 (0.1–1 μM) failed to significantly modify the 30 pM LY379268-induced inhibition of tritium exocytosis (Figure 2). Experiments were performed to investigate the impact of this compound on the tritium release evoked by 15 mM KCl in the presence of a lower (1 pM) concentration of the mGlu_{2/3} agonist. Again, the mGlu₂ PAM was devoid of activity (Figure 2). However, the inhibitory effect elicited by LY379268 (30 pM) was totally abolished when LY2389575 (1 μM) was added concomitantly (Figure 2). A lower concentration (0.1 μM) of the mGlu₃ NAM slightly, albeit not significantly, affected the 15 mM KCl/30 pM

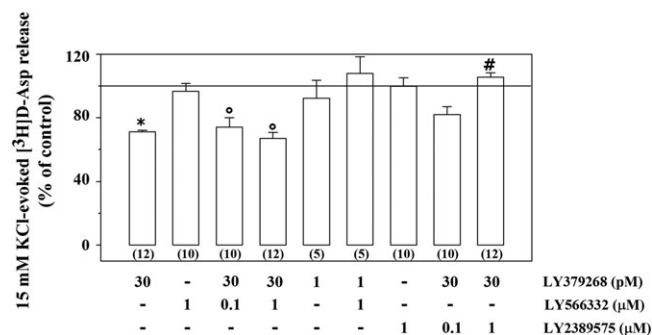


Figure 2

Effects of the mGlu₂ PAM LY566332 and of the mGlu₃ NAM LY2389575 on the inhibition of the 15 mM KCl-evoked [³H]-D-Asp overflow elicited by the mGlu_{2/3} agonist LY379268 in mouse spinal cord synaptosomes. The 15 mM KCl-evoked release of tritium over the basal release amounted to 4.23 ± 0.79 nCi. Results are expressed as a percentage of the 15 mM KCl-evoked tritium overflow (% of control). Data are the mean ± SEM of *n* (number within brackets) experiments. **P* < 0.05 versus the 15 mM KCl-evoked tritium overflow; ^o*P* < 0.05 versus the 15 mM KCl/1 μM LY566332-evoked tritium overflow; [#]*P* < 0.05 versus the 15 mM KCl/30 pM LY379268-evoked tritium overflow.

LY379268-evoked release of [³H]-D-Asp. The mGlu₃ receptor NAM, on its own, did not modify the 15 mM KCl-evoked release of tritium from spinal cord synaptosomes (Figure 2).

Effects of BINA and of ML337 on the LY379268-induced inhibition of [³H]-D-Asp exocytosis from cortical and spinal cord glutamatergic synaptosomes

To confirm the pharmacological profile of the mGlu_{2/3} autoreceptors under study, experiments were carried out to evaluate the impact of the mGlu₂ PAM BINA and mGlu₃ NAM ML337 on glutamate exocytosis in the absence and in the presence of LY379268 from both synaptosomal preparations. The concomitant addition of the mGlu₂ PAM BINA (1 μM) to the depolarizing stimulus (12 mM KCl) did not significantly modify the [³H]-D-Asp exocytosis from cortical synaptosomes. However, the release of tritium elicited by 12 mM KCl/3 nM LY379268/1 μM BINA was significantly lower than that elicited by 12 mM KCl/3 nM LY379268 and by 12 mM KCl/1 μM BINA. The mGlu₃ NAM ML337 also significantly modified the 12 mM KCl/10 nM LY379268-evoked release of [³H]-D-Asp, leaving unchanged the 12 mM KCl-induced [³H]-D-Asp exocytosis (Figure 3A). In contrast, in spinal cord synaptosomes, BINA (1 μM) failed to affect the 15 mM KCl-evoked release of [³H]-D-Asp in the absence and in the presence of 3 pM LY379268. Moreover, ML337 did not significantly change the 15 mM KCl-evoked release of tritium but almost totally reversed the inhibitory effect elicited by 10 pM LY379268 on the 15 mM KCl-evoked exocytosis of the radioactive tracer (Figure 3B).

Expression of mGlu₂ and mGlu₃ receptor proteins in mouse cortical and spinal cord synaptosomes

To assess the expression of mGlu₂ and mGlu₃ receptor proteins in mouse cortical and spinal cord synaptosomes, selective antibodies recognizing amino acid sequences within the outer NH₂ terminus of these receptor proteins were used. Their specificity was confirmed in Western blot analysis on cortical homogenates from WT, mGlu₂^{-/-}, mGlu₃^{-/-} and mGlu_{2/3}^{-/-} mice. Both receptor antibodies recognized proteins with a mass of about 200 kDa in the brain homogenate from the WT mice, which is consistent with the presence of the mGlu₂ and the mGlu₃ receptor dimers. The anti-mGlu₂ antibody failed to recognize the band corresponding to 200 kDa in lysates from mGlu₂^{-/-} and from mGlu_{2/3}^{-/-} mice. Moreover, the anti-mGlu₃ antibody did not recognize the protein with comparable mass in lysates from mGlu₃^{-/-} and from mGlu_{2/3}^{-/-} mice (Figure 4). Both receptor antibodies recognized proteins with a mass of about 100 kDa in the brain homogenates from the mice belonging to the four different strains. In particular, the anti-mGlu₃ antibody recognized a band at 100 kDa largely conserved in the four lysates (Figure 4B), consistent with the presence of non-specific immunoreactivity. In contrast, the band at about 100 kDa recognized by the anti-mGlu₂ antibody has a higher intensity in the WT and in the mGlu₃^{-/-} lysates when compared to mGlu₂^{-/-} and mGlu_{2/3}^{-/-} lysates, suggesting the presence of a monomeric form of the mGlu₂ receptor protein (Figure 4A).

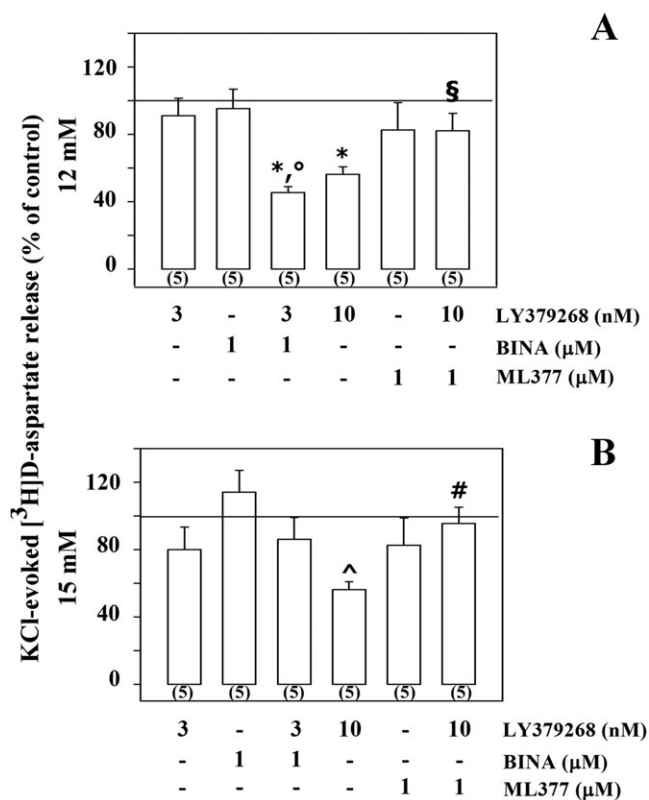


Figure 3

Effects of the mGlu₂ PAM BINA and of the mGlu₃ NAM ML337 on the inhibition of the KCl-evoked [³H]-D-Asp overflow elicited by the mGlu_{2/3} agonist LY379268 in mouse cortical (A) and spinal cord (B) synaptosomes. (A) The 12 mM KCl-evoked release of tritium from cortical synaptosomes amounted to 1.75 ± 0.42 nCi. (B) The 15 mM KCl-evoked release of tritium from cortical synaptosomes amounted to 4.76 ± 0.63 nCi. Results are expressed as a percentage of the KCl-evoked tritium overflow (% of control). Data are the mean \pm SEM of *n* (number within brackets) experiments. **P* < 0.05 versus the 12 mM KCl-evoked tritium overflow; ^o*P* < 0.05 versus the 12 mM KCl/1 μM BINA-evoked tritium overflow; §*P* < 0.05 versus the 12 mM KCl/10 nM LY379268-evoked tritium overflow; ^*P* < 0.05 versus the 15 mM KCl-evoked tritium overflow; #*P* < 0.05 versus the 15 mM KCl/10 pM LY379268-evoked tritium overflow.

Immunochemical studies showed that the anti-mGlu₂ receptor antibody recognized a clear mGlu₂ immunoreactivity in the lysates of the synaptosomes from the cortex of WT mice at about 200 kDa, which is consistent with the presence of the dimeric form of the receptor (Figure 5A). Also mGlu₂ immunopositivity was detectable, although to a lesser extent, in spinal cord synaptosomal lysates (Figure 5B). The anti-mGlu₃ antibody recognized a band with an apparent mass of about 200 kDa in both cortical and spinal cord synaptosomal lysates (Figure 5A,B).

mGlu₂ and mGlu₃ receptor proteins co-localize with VGLUT1 protein in mouse cortical and spinal cord synaptosomes

Confocal microscopy was carried out to investigate whether mGlu₂ and mGlu₃ receptors co-localize with VGLUT1, here used as a selective marker of glutamate terminals. A diffuse mGlu₂-immunoreactivity (Figure 6A, green) was observed in VGLUT1-positive (Figure 6B, red) cortical synaptosomes (Figure 6C, yellow, merge, $91 \pm 4\%$, *n* = 12). The mouse cortical synaptosomal preparation also showed a significant colocalization of mGlu₃ receptors and VGLUT1 (Figure 6D, green; E, red; F, yellow, merge, $74 \pm 4\%$, *n* = 12). Finally, a significant percentage of mGlu₂ positive particles (Figure 6G, green) was also reactive for mGlu₃ receptors (Figure 6H, red; I, yellow, merge, $83 \pm 4\%$, *n* = 12), while the expression of mGlu₂ immunoreactivity in mGlu₃ receptor-positive particles was less pronounced (Figure 6I, yellow, merge $71 \pm 3\%$, *n* = 12).

In spinal cord synaptosomes, mGlu₂ receptor immunoreactivity (Figure 7A, green) and mGlu₃ receptor immunoreactivity (Figure 7D, green) were observed in VGLUT1-immunopositive (Figure 7B,E, red) particles (Figure 7C, yellow, merge, 55.7 ± 3.1 of mGlu₂ receptor localization in VGLUT1-positive cells; Figure 7F, yellow, merge, 46.5 ± 7.2 of mGlu₃ receptor localization in VGLUT1-positive cells). Finally, mGlu₃ receptor immunopositive particles showed a low immunoreactivity for mGlu₂ receptor signalling (Figure 7I, merge, yellow, 33.1 ± 6.2 of mGlu₂ receptor immunoreactivity in mGlu₃ receptor immunoreactive particles), while a larger co-localization of mGlu₃ receptor immunosignal with mGlu₂ receptor-positive particles was detected (Figure 7I,

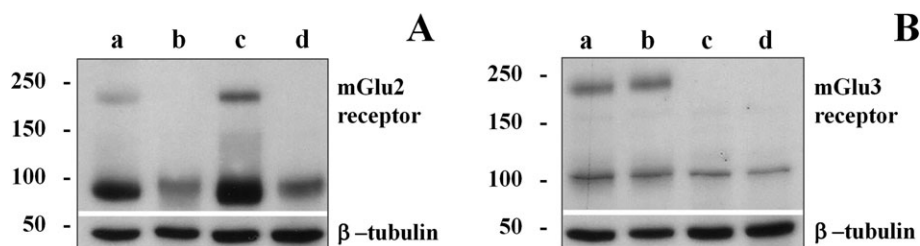


Figure 4

Western blot analysis of mGlu₂ (A) and mGlu₃ (B) receptors in mouse cortical homogenates of wild-type mice (panel a), mGlu₂ receptor knockout mice (mGlu₂^{-/-}, panel b), mGlu₃ receptor knockout mice (mGlu₃^{-/-}, panel c) and double mGlu₂ and mGlu₃ receptor knockout mice (mGlu_{2/3}^{-/-}, panel d). The figure shows a representative Western blot.

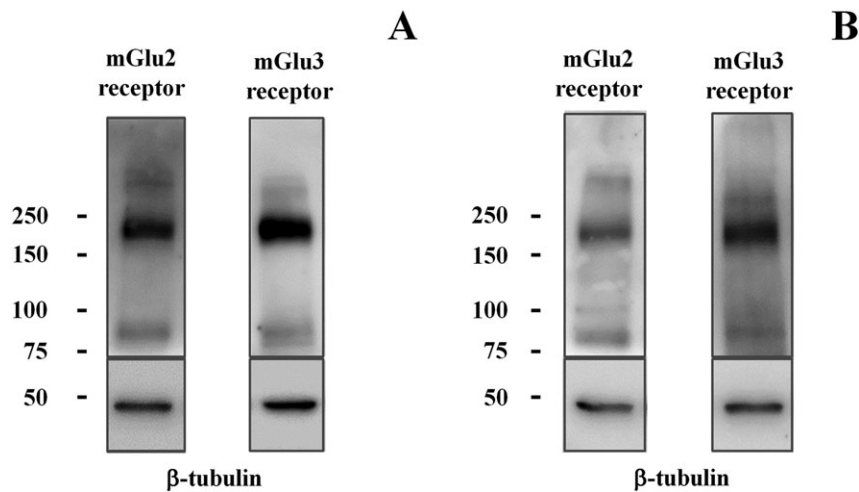


Figure 5

Western blot analysis reveals the presence of mGlu₂ receptor and mGlu₃ receptor protein dimers in mouse cortical (A) and spinal cord (B) synaptosomes (10 µg protein for cortical lysates and 20 µg protein for spinal cord lysates). The figure shows a representative blot of five (cortex) to seven (spinal cord) analyses carried out on different days.

yellow, merge, 52.1 ± 9.2 mGlu₃ receptor immunoreactivity in mGlu₂ receptor-immunopositive particles).

mGlu₂ and mGlu₃ receptor antibodies prevent the LY379268-induced inhibition of the 12 mM KCl-evoked [³H]-D-Asp exocytosis from cortical glutamatergic synaptosomes

Synaptosomes were incubated with anti-mGlu₂ or anti-mGlu₃ receptor antibodies and the release of preloaded [³H]-D-Asp elicited by the KCl stimulus was quantified in the absence or presence of LY379268. Incubation of cortical synaptosomes with the anti-mGlu₂ receptor antibody did not cause a significant changes to the 12 mM KCl-evoked release of [³H]-D-Asp (control synaptosomes: 1.04 ± 0.8 ; synaptosomes incubated with the anti-mGlu₂ receptor antibody: 1.10 ± 0.6 , $n = 10$, results expressed as induced overflow; mean \pm SEM) but totally abolished the LY379268-induced inhibition of the radioactive overflow (Figure 8). Similarly, incubation of cortical synaptosomes with the anti-mGlu₃ receptor antibody did not significantly modify the 12 mM KCl-evoked tritium exocytosis (synaptosomes incubated with anti-mGlu₃ receptor antibody: 1.11 ± 0.7 , $n = 8$, results expressed as induced overflow; mean \pm SEM). In contrast, LY379268 (3 nM) still significantly inhibited the 12 mM KCl-evoked release of tritium from cortical synaptosomes pre-exposed to the anti-mGlu₃ receptor antibody, although the inhibitory effect was significantly less pronounced than that observed in control synaptosomes (Figure 8).

mGlu₂ and mGlu₃ receptor antibodies prevent the LY379268-induced inhibition of the 15 mM KCl-evoked [³H]-D-Asp exocytosis from spinal cord glutamatergic synaptosomes

Incubation of synaptosomes with antibodies failed to significantly affect the 15 mM KCl-evoked release of [³H]-D-Asp

(control synaptosomes: 4.52 ± 0.72 ; synaptosomes incubated with the anti-mGlu₂ receptor antibody: 4.61 ± 0.52 ; synaptosomes incubated with the anti-mGlu₃ receptor antibody: 4.83 ± 0.39 , $n = 8$, results expressed as induced overflow; mean \pm SEM). The inhibition of the 15 mM KCl-evoked [³H]-D-Asp exocytosis elicited by LY379268 (30 pM) was unmodified in synaptosomes pre-exposed to the anti-mGlu₂ receptor antibody, but it was significantly abolished in synaptosomes pre-exposed to the anti-mGlu₃ receptor antibody (Figure 9).

Discussion

Since the 90s, novel ligands for the group II mGlu receptors have been synthesized and their activity characterized. These compounds are typified by selectivity and by nanomolar affinities for these receptors and might represent the prototype of new neuroprotective agents (Nickols and Conn, 2014; Yin and Niswender, 2014). The novel compounds include ligands displaying mixed agonist/antagonist pharmacological effects on both mGlu₂ and mGlu₃ receptors. The highly conserved sequence of the two receptor proteins makes it difficult to compare the pharmacological profile of these mixed agents, in terms of affinity, activity and efficacy at each receptor subtype, and also to identify selective *native* neuronal and glial systems to be used as cellular models for their pharmacological characterization (but see Schoepp, 2001; Hanna *et al.*, 2013; Sanger *et al.*, 2013; Durand *et al.*, 2014).

We used new molecules and established compounds to characterize the pharmacological profile of native mGlu_{2/3} autoreceptors in cortical and spinal cord terminals. We recently proposed that these native autoreceptors belong to different receptor subtypes, having a different receptor composition and pharmacological profile, and might, therefore, provide native receptor models for studying the

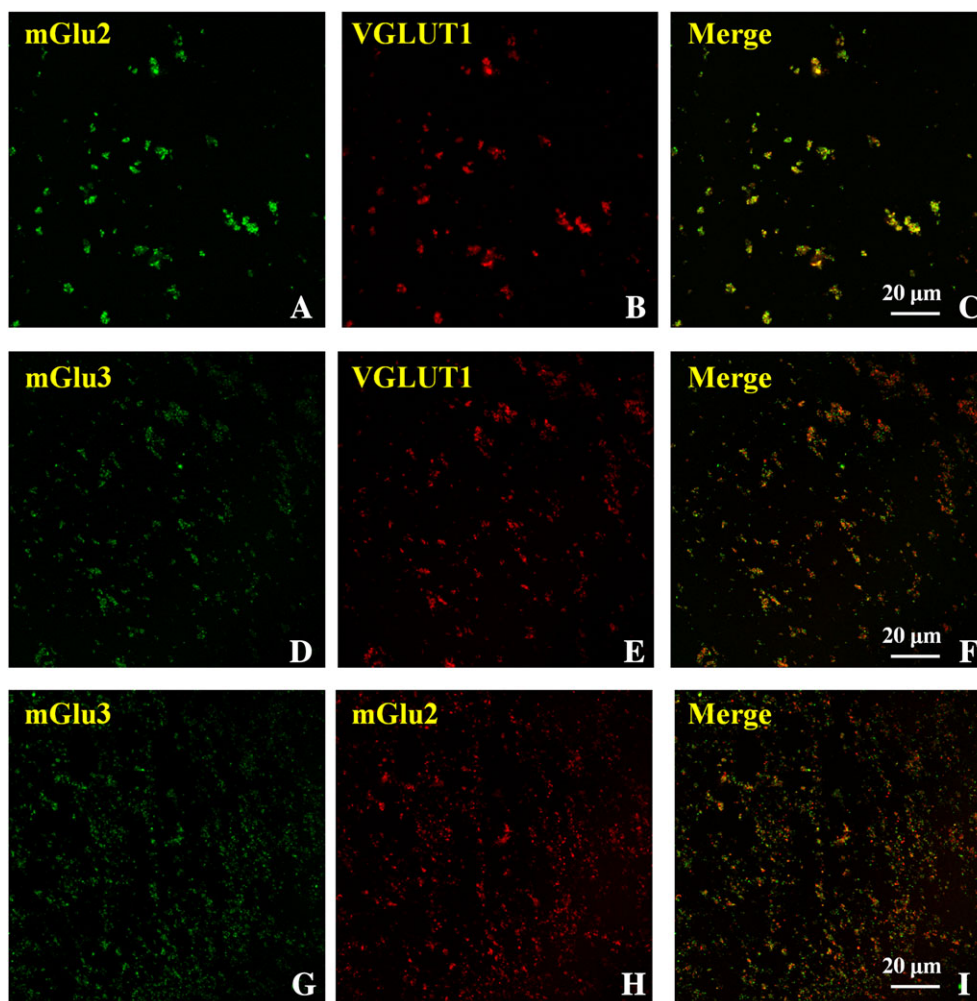


Figure 6

Confocal analysis of mGlu₂ and mGlu₃ receptor immunoreactivities in glutamatergic VGLUT1-positive cortical nerve terminals: co-localization of the two receptor proteins. The figure shows representative images of six independent experiments carried out on different days.

pharmacological profile of new drugs (Di Prisco *et al.*, 2016a). This conclusion is now further supported by the functional and immunochemical studies described in the present work.

As a first approach, we focussed on two novel allosteric modulators able to discriminate between mGlu₂ and mGlu₃ receptors: the mGlu₂ receptor PAM LY566332 and the mGlu₃ receptor NAM LY2389575. It is known that PAMs preferentially amplify receptor-mediated events elicited by submaximal concentrations of orthosteric agonists, while NAMs impede receptor-mediated events regardless of the amount of agonist applied. To study the impact of PAM and NAM, mGlu_{2/3} receptor-mediated inhibition of glutamate exocytosis was elicited by exposing synaptosomes to concentrations of LY379268 that assure the submaximal activation of both of the two autoreceptors. When choosing the LY379268 concentrations to inhibit glutamate exocytosis from the two synaptosomal preparations, we took into consideration the strikingly different potency (about 4 order of magnitude) of LY379268 in cerebral cortex when compared to spinal cord

synaptosomes highlighted in our previous study (Di Prisco *et al.*, 2016a). A molecular explanation of this striking difference is not yet available, and future studies will be required to address this aspect. We found that LY566332 amplified the inhibitory activity of LY379268 at presynaptic cortical autoreceptors, leaving unchanged the LY379268-mediated effect in spinal cord terminals. In spinal cord synaptosomes, the positive allosteric effects of LY566332 could not be revealed even if the mGlu_{2/3} agonist was present at a lower concentration, further confirming that the mGlu₂ receptor subtype does not participate in the LY379268-mediated inhibition of glutamate exocytosis in these terminals. In contrast, LY2389575 significantly recovered the inhibitory activity exerted by LY379268 at spinal cord mGlu₃-preferring autoreceptors, while it slightly, although not significantly, modified the efficacy of a submaximal amount of LY379268 at presynaptic mGlu₂-preferring autoreceptors in cortical synaptosomes. The antagonist-like activity of the mGlu₃ NAM, however, was disclosed when cortical synaptosomes were exposed to a concentration of LY379268 eliciting the

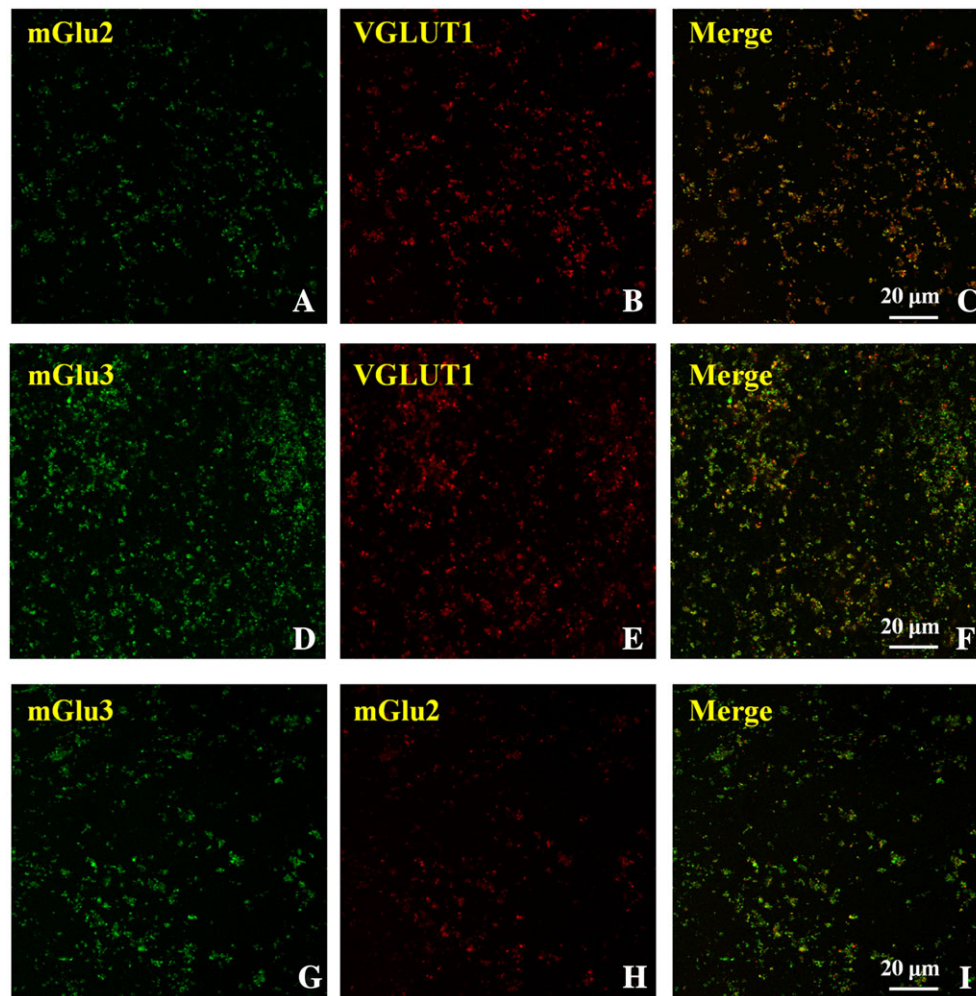


Figure 7

Confocal analysis highlighting the presence of mGlu₂ and mGlu₃ receptor immunoreactivities in glutamatergic spinal cord VGLUT1-positive nerve terminals: co-localization of the two receptor proteins. The figure shows representative images of eight independent experiments carried out on different days.

almost maximal inhibitory effect at these terminals. Comparable results were obtained when studying the impact of the mGlu₂ PAM BINA and the mGlu₃ NAM ML337. Altogether, these observations largely corroborated the hypothesis that mGlu₃ receptor subtypes account for presynaptic control of glutamate exocytosis from spinal cord terminals, while both mGlu₂ and mGlu₃ receptors participate in the control of glutamate overflow from cortical terminals.

To further support the data obtained with group II NAMs and PAMs, we approached the pharmacological characterization of these receptors by using selective anti-mGlu₂ and anti-mGlu₃ receptor antibodies as receptor ligands. Classically, antibodies have been used to characterize the subunit composition of receptors in immunochemical studies. More recently, these reagents were proposed as tools to define the pharmacological profile of receptors, in particular GPCRs (Gupta *et al.*, 2008; Musante *et al.*, 2008b and references therein, Di Prisco *et al.*, 2012, 2016b; Meregá *et al.*, 2015). Actually, the binding of an antibody to the ligand-binding pocket of the receptor or to its extraterminal region(s) causes

either activation and/or blockade of the receptor-mediated functions, mimicking the effect of an agonist or antagonist at these structures (Musante *et al.*, 2008b; Di Prisco *et al.*, 2012).

Antibodies raised against the extracellular epitope of the mGlu receptors have different effects on the LY379268-mediated inhibition of glutamate exocytosis in the two synaptosomal preparations. In particular, the anti-mGlu₂ receptor antibody impeded the inhibitory activity of LY379268 on glutamate exocytosis in cortical synaptosomes, but not in spinal cord terminals. Whereas, the LY379268-induced inhibition of glutamate exocytosis was impeded by the anti-mGlu₃ receptor antibody in spinal cord synaptosomes, and it was slightly, although significantly, reduced in cortical terminals pretreated with the antibody.

Before drawing any conclusions from these findings, these observations deserve some brief comments. Firstly, the possibility that antibody cross-reactivity might affect our results is excluded on the basis of the high selectivity of the two antibodies, revealed in the Western blot analysis carried out with cortical homogenates from mGlu₂ and/or mGlu₃

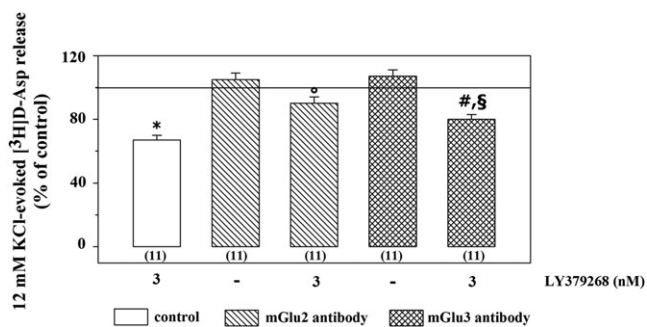


Figure 8

Effects of the incubation of cortical synaptosomes with an anti-mGlu₂ receptor or anti-mGlu₃ receptor antibody on the 12 mM KCl/LY379268-evoked release of [³H]-D-Asp. The 12 mM KCl-evoked release of tritium amounted to the following: control synaptosomes, 1.48 ± 0.43 nCi; anti-mGlu₂ antibody-incubated synaptosomes, 1.63 ± 0.27 nCi; and anti-mGlu₃ antibody-incubated synaptosomes 1.48 ± 0.29 nCi. Results are expressed as percentage of the 12 mM KCl-evoked [³H]-D-Asp release (% of control). Data are the mean ± SEM of *n* (number within brackets) experiments **P* < 0.05 versus the 12 mM KCl-evoked tritium overflow from control cortical synaptosomes; °*P* < 0.05 versus the 12 mM KCl/3 nM LY379268-evoked tritium overflow from control cortical synaptosomes; #*P* < 0.05 versus the 12 mM KCl-evoked tritium overflow from cortical synaptosomes incubated with the anti-mGlu₃ receptor antibody; §*P* < 0.05 versus the 12 mM KCl/3 nM LY379268-evoked tritium overflow from control cortical synaptosomes.

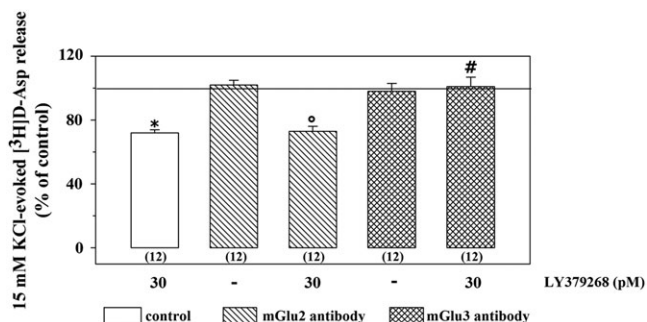


Figure 9

Effects of the incubation of spinal cord synaptosomes with anti-mGlu₂ receptor or anti-mGlu₃ receptor antibody on the 15 mM KCl/LY379268-evoked release of [³H]-D-Asp. The 15 mM KCl-evoked release of tritium amounted to the following: control synaptosomes, 4.76 ± 0.69 nCi; anti-mGlu₂ antibody-incubated synaptosomes, 3.21 ± 0.55 nCi; and anti-mGlu₃ antibody-incubated synaptosomes 3.98 ± 0.42 nCi. Results are expressed as percentage of the 15 mM KCl-evoked [³H]-D-Asp release. Data are the mean ± SEM of *n* (number within brackets) experiments **P* < 0.05 versus the 15 mM KCl-evoked tritium overflow from control spinal cord synaptosomes; °*P* < 0.05 versus the 15 mM KCl-evoked tritium overflow from spinal cord synaptosomes incubated with the anti-mGlu₂ receptor antibody; #*P* < 0.05 versus the 15 mM KCl/30 pM LY379268-evoked tritium overflow from control spinal cord synaptosomes.

receptor knockout mice. In our previous studies, we demonstrated that incubation of synaptosomes with antibodies raised against cytosolic proteins (i.e. β actin) did not affect

the spontaneous and the K⁺-evoked release of glutamate from both cortical and spinal cord terminals, ruling out the possibility that unspecific events might account for the effects observed (Musante *et al.*, 2008b; Di Prisco *et al.*, 2012; Merrega *et al.*, 2015). Secondly, the fact that both antibodies failed to modify on their own the KCl-evoked glutamate exocytosis in both synaptosomal preparations allows us to rule out the occurrence of unspecific effects and also suggest that the antibodies do not mimic mGlu_{2/3} receptor agonist(s) and/or PAM(s) at mGlu_{2/3} receptors. Rather, our results indicate that antibodies preferentially act as selective antagonist and/or NAM at these targets. Finally, even if monoclonal antibodies are thought to be the best candidates as receptor agonists/antagonists, our results demonstrate that polyclonal antibodies can also be applied to function as receptor agonists or antagonists (reviewed by Gupta *et al.*, 2008). Notably, the results obtained with the two antibodies were superimposable on those obtained with selective PAM and NAM and suggest that in spinal cord terminals, mGlu₃-preferring autoreceptors are pivotal for LY379268-induced inhibition of glutamate exocytosis, while in cortical synaptosomes, mGlu₂ autoreceptors play the main role, although the mGlu₃ receptors also participate in this effect.

Quite interestingly, the use of the selective antibodies in immunochemical analysis unveiled a scenario well in line with the functional conclusions from release studies. In fact, Western blot analysis revealed the presence of both anti-mGlu₂ and anti-mGlu₃ receptor immunoreactivities in cortical (see also Ferraguti and Shigemoto, 2006) and spinal cord synaptosomal lysates; this is compatible with the idea that synaptosomal particles originating from both CNS regions are endowed with the two mGlu receptor proteins. These results, however, do not provide information on where the receptors are located (i.e. on glutamatergic or non-glutamatergic spinal cord terminals). This conclusion is proposed on the basis of the results obtained with confocal microscopy, which was carried out to highlight the co-localization of anti-mGlu₂ and anti-mGlu₃ receptor immunoreactivities with VGLUT1, used here as a selective marker of the glutamatergic particles.

Confocal analysis suggested that within the cortical synaptosomal preparations, there are glutamatergic synaptosomes endowed with group II mGlu receptors containing the mGlu₂ receptor protein and synaptosomes possessing group II mGlu receptors containing the mGlu₃ receptor protein. Here, the high co-localization of mGlu₂ positivity with VGLUT1 when compared to mGlu₃ receptor indicates the predominant expression of mGlu₂ receptors in these terminals. However, the high co-localization of mGlu₃ receptor reactivity with the mGlu₂ receptor-positive particles suggests that most of the cortical glutamatergic terminals possessing mGlu₂-preferring autoreceptors also express co-localized mGlu₃ receptors. In contrast in the spinal cord, the largest population of glutamatergic terminals was endowed with mGlu₃ receptors, but mGlu₃ immunopositivity (i.e. of the receptor that shows the largest distribution in glutamatergic terminals) slightly co-localized with the less diffuse mGlu₂ receptor immunoreactivity, suggesting that most of the spinal cord terminals do not possess both receptor proteins.

As a whole, based on both the functional and non-functional observations described here, we confirm our

original working hypothesis that cortical terminals are endowed with mGlu₂-preferring receptors, the releasing activity of which is also influenced, but not obviously so, by co-localized mGlu₃ receptors, while spinal cord terminals possess mGlu₃-preferring receptors that control glutamate release. We propose the selective antibodies, that recognize the N-terminal of mGlu receptor protein, as new pharmacological tools for characterizing these receptors. Also we confirm that the group II mGlu autoreceptors controlling glutamate exocytosis in spinal cord terminals represent an appropriate model of native mGlu₃-preferring receptor for studying the detailed pharmacological profile of ligands at this receptor subtype.

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Author contributions

A.P. was involved in the supervision, planning and execution of the research. T.B. and G.O. were involved in animal care and release experiments. G.O. and M.V. were involved in Western blot analysis. C.U. performed confocal microscopy analysis. B.R. and G.B. analysed the specificity of anti mGlu₂ and anti-mGlu₃ receptor antibodies. G.B. and F.N. provided the transgenic mice. A.P. wrote the manuscript. A.P., G.B. and F.N. discussed the results and revised the manuscript.

Conflict of interest

The authors declare no conflicts of interest.

Declaration of transparency and scientific rigour

This Declaration acknowledges that this paper adheres to the principles for transparent reporting and scientific rigour of preclinical research recommended by funding agencies, publishers and other organisations engaged with supporting research.

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