

Supplementary Material

Deconstruction – Reconstruction: Analysis of the Crucial Structural Elements of GluN2B-Selective, Negative Allosteric NMDA Receptor Modulators with 3-Benzazepine Scaffold

Nadine Ritter^{a,b} Marvin Korff^{b,c} Alexander Markus^c Dirk Schepmann^c
Guiscard Seebom^{a,b} Julian A. Schreiber^a Bernhard Wünsch^{b,c}

^aCellular Electrophysiology and Molecular Biology, Institute for Genetics of Heart Diseases (IfGH), Department of Cardiovascular Medicine, University Hospital Münster, Münster, Germany, ^bGRK 2515, Chemical biology of ion channels (Chembion), Westfälische Wilhelms-Universität Münster, Münster, Germany, ^cInstitute of Pharmaceutical and Medicinal Chemistry, Westfälische Wilhelms-Universität Münster, Münster, Germany

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1. Chemistry, General

Unless otherwise noted, moisture sensitive reactions were conducted under dry nitrogen. CH₂Cl₂ was distilled over CaH₂. Thin layer chromatography (tlc): Silica gel 60 F₂₅₄ plates (Merck). Flash chromatography (fc): Silica gel 60, 40–64 μm (Merck); parentheses include: diameter of the column (d), length of the column (l), fraction size (v), eluent. Melting point: Melting point apparatus Mettler Toledo MP50 Melting Point System, uncorrected. MS: microTOF-Q II (Bruker Daltonics); APCI, atmospheric pressure chemical ionization. IR: FT-IR spectrophotometer MIRacle 10 (Shimadzu) equipped with ATR technique. Nuclear magnetic resonance (NMR) spectra were recorded on Agilent 600-MR (600 MHz for ¹H, 151 MHz for ¹³C) or Agilent 400-MR spectrometer (400 MHz for ¹H, 101 MHz for ¹³C); δ in ppm related to tetramethylsilane and measured referring to CHCl₃ (δ = 7.26 ppm (¹H NMR) and δ = 77.2 ppm (¹³C NMR)) and CHD₂OD (δ = 3.31 ppm (¹H NMR) and δ = 49.0 ppm (¹³C NMR)); coupling constants are given with 0.5 Hz resolution; the assignments of ¹³C and ¹H NMR signals were supported by 2-D NMR techniques where necessary. HPLC: Merck Hitachi Equipment; UV detector: L-7400; autosampler: L-7200; pump: L-7100; degasser: L-7614; column: LiChrospher[®] 60 RP-select B (5 μm); LiChroCART[®] 250-4 mm cartridge; flow rate: 1.0 mL/min; injection volume: 5.0 μL; detection at λ = 210 nm; solvents: A: water with 0.05% (v/v) trifluoroacetic acid; B: acetonitrile with 0.05% (v/v) trifluoroacetic acid; gradient elution: (A%): 0-4 min: 90%, 4-29 min: 90 → 0%, 29-31 min: 0%, 31-31.5 min: 0 → 90%, 31.5-40 min: 90%. The purity of all test compounds determined by this method was higher than 95 %.

2. Receptor binding studies

2.1. Materials

Guinea pig brains, rat brains and rat livers were commercially available (Harlan-Winkelmann, Borcheln, Germany). Pig brains were a donation of the local slaughterhouse (Coesfeld, Germany). The recombinant L(tk-) cells stably expressing the GluN2B receptor were obtained from Prof. Dr. Dieter Steinhilber (Frankfurt, Germany). Homogenizers: Elvehjem Potter (B. Braun Biotech International, Melsungen, Germany) and Soniprep[®] 150 (MSE, London, UK). Centrifuges: Cooling centrifuge model Eppendorf 5427R (Eppendorf, Hamburg, Germany) and High-speed cooling centrifuge model Sorvall[®] RC-5C plus (Thermo Fisher Scientific, Langenselbold, Germany). Multiplates: standard 96 well multiplates (Diagonal, Muenster, Germany). Shaker: self-made device with adjustable temperature and tumbling speed (scientific workshop of the institute). Harvester: MicroBeta[®] FilterMate 96 Harvester. Filter: Printed Filtermat Typ A and B. Scintillator: Meltilex[®] (Typ A or B) solid state scintillator. Scintillation analyzer: MicroBeta[®] Trilux (all Perkin Elmer LAS, Rodgau-Jügesheim, Germany).

2.2. Cell culture and preparation of membrane homogenates from GluN2B cells

Mouse L(tk-) cells stably transfected with the dexamethasone-inducible eukaryotic expression vectors pMSG GluN1a, pMSG GluN2B (1:5 ratio) were grown in Modified Earl's Medium (MEM) containing 10 % of standardized FBS Superior (Biochrom AG, Berlin, Germany). The expression of the NMDA receptor at the cell surface was induced after the cell density of the adherent growing cells had reached approximately 90 % of confluency. For the induction, the original growth medium was replaced by growth medium containing 4 μ M dexamethasone and 4 μ M ketamine (final concentration). After 24 h, the cells were rinsed with phosphate buffered saline solution (PBS, Biochrom AG, Berlin, Germany), harvested by mechanical detachment and pelleted (10 min, 1,200 x g).

For the binding assay, the cell pellet was resuspended in PBS solution and the number of cells was determined using a Scepter[®] cell counter (MERCK Millipore, Darmstadt, Germany). Subsequently, the cells were lysed by sonication (4 °C, 6 x 10 s cycles with breaks of 10 s). The resulting cell fragments were centrifuged with a high performance cool centrifuge (23,500 x g, 4 °C). The supernatant was discarded and the pellet was

resuspended in a defined volume of PBS yielding cell fragments of approximately 500,000 cells/mL. The suspension of membrane homogenates was sonicated again (4 °C, 2 x 10 s cycles with a break of 10 s) and stored at -80 °C.

2.3. Preparation of membrane homogenates from guinea pig brain

5 guinea pig brains were homogenized with the potter (500-800 rpm, 10 up and down strokes) in 6 volumes of cold 0.32 M sucrose. The suspension was centrifuged at 1,200 x g for 10 min at 4 °C. The supernatant was separated and centrifuged at 23,500 x g for 20 min at 4 °C. The pellet was resuspended in 5-6 volumes of buffer (50 mM TRIS, pH 7.4) and centrifuged again at 23,500 x g (20 min, 4 °C). This procedure was repeated twice. The final pellet was resuspended in 5-6 volumes of buffer and frozen (-80 °C) in 1.5 mL portions containing about 1.5 mg protein/mL.

2.4. Preparation of membrane homogenates from rat liver

Two rat livers were cut into small pieces and homogenized with the potter (500-800 rpm, 10 up and down strokes) in 6 volumes of cold 0.32 M sucrose. The suspension was centrifuged at 1,200 x g for 10 min at 4 °C. The supernatant was separated and centrifuged at 31,000 x g for 20 min at 4 °C. The pellet was resuspended in 5-6 volumes of buffer (50 mM TRIS, pH 8.0) and incubated at rt for 30 min. After the incubation, the suspension was centrifuged again at 31,000 x g for 20 min at 4 °C. The final pellet was resuspended in 5-6 volumes of buffer and stored at -80 °C in 1.5 mL portions containing about 2 mg protein/mL.

2.5. Protein determination

The protein concentration was determined by the method of Bradford,¹ modified by Stoscheck.² The Bradford solution was prepared by dissolving 5 mg of Coomassie Brilliant Blue G 250 in 2.5 mL of EtOH (95 %, v/v). 10 mL deionized H₂O and 5 mL phosphoric acid (85 %, m/v) were added to this solution, the mixture was stirred and filled to a total volume of 50 mL with deionized water. The calibration was carried out using bovine serum albumin as a standard in 9 concentrations (0.1, 0.2, 0.4, 0.6, 0.8, 1.0, 1.5, 2.0 and 4.0 mg /mL). In a 96 well standard multiplate, 10 µL of the calibration solution or 10 µL of the membrane receptor preparation were mixed with 190 µL of the Bradford solution, respectively. After 5 min, the UV absorption of the protein-dye

complex at $\lambda = 595$ nm was measured with a plate reader (Tecan Genios[®], Tecan, Crailsheim, Germany).

2.6. General procedures for the binding assays

The test compound solutions were prepared by dissolving approximately 10 μmol (usually 2-4 mg) of test compound in DMSO so that a 10 mM stock solution was obtained. To obtain the required test solutions for the assay, the DMSO stock solution was diluted with the respective assay buffer. The filtermats were presoaked in 0.5 % aqueous polyethylenimine solution for 2 h at rt before use. All binding experiments were carried out in duplicates in the 96 well multiplates. The concentrations given are the final concentration in the assay. Generally, the assays were performed by addition of 50 μL of the respective assay buffer, 50 μL of test compound solution in various concentrations (10^{-5} , 10^{-6} , 10^{-7} , 10^{-8} , 10^{-9} and 10^{-10} mol/L), 50 μL of the corresponding radioligand solution and 50 μL of the respective receptor preparation into each well of the multiplate (total volume 200 μL). The receptor preparation was always added last. During the incubation, the multiplates were shaken at a speed of 500-600 rpm at the specified temperature. Unless otherwise noted, the assays were terminated after 120 min by rapid filtration using the harvester. During the filtration, each well was washed five times with 300 μL of water. Subsequently, the filtermats were dried at 95 °C. The solid scintillator was melted on the dried filtermats at a temperature of 95 °C for 5 min. After solidifying of the scintillator at rt, the trapped radioactivity in the filtermats was measured with the scintillation analyzer. Each position on the filtermat corresponding to one well of the multiplate was measured for 5 min with the [³H]-counting protocol. The overall counting efficiency was 20 %. The IC_{50} values were calculated with the program GraphPad Prism[®] 3.0 (GraphPad Software, San Diego, CA, USA) by non-linear regression analysis. Subsequently, the IC_{50} values were transformed into K_i values using the equation of Cheng and Prusoff.³ The K_i values are given as mean value \pm SEM from three independent experiments.

2.7. Performance of the binding assays

2.7.1. Ifenprodil binding site of GluN2B subunit containing NMDA receptors

The competitive binding assay was performed with the radioligand [³H]ifenprodil (60 Ci/mmol; BIOTREND, Cologne, Germany). The thawed cell membrane preparation from the transfected L(tk-) cells (about 20 μg protein) was incubated with various

concentrations of test compounds, 5 nM [³H]ifenprodil, and TRIS/EDTA-buffer (5 mM TRIS/1 mM EDTA, pH 7.5) at 37 °C. The non-specific binding was determined with 10 μM unlabeled ifenprodil. The K_d value of ifenprodil is 7.6 nM.⁵

2.7.2. σ_1 receptor assay

The assay was performed with the radioligand [³H]-(+)-pentazocine (22.0 Ci/mmol; Perkin Elmer). The thawed membrane preparation of guinea pig brain (about 100 μg of the protein) was incubated with various concentrations of test compounds, 2 nM [³H]-(+)-pentazocine, and TRIS buffer (50 mM, pH 7.4) at 37 °C. The non-specific binding was determined with 10 μM unlabeled (+)-pentazocine. The K_d value of (+)-pentazocine is 2.9 nM.⁶

2.7.3. σ_2 receptor assay

The assays were performed with the radioligand [³H]di-*o*-tolyguanidine (specific activity 50 Ci/mmol; ARC, St. Louis, MO, USA). The thawed rat liver membrane preparation (about 100 μg protein) was incubated with various concentrations of the test compound, 3 nM [³H]di-*o*-tolyguanidine and buffer containing (+)-pentazocine (500 nM (+)-pentazocine in TRIS buffer (50 mM TRIS, pH 8.0)) at rt. The non-specific binding was determined with 10 μM non-labeled di-*o*-tolyguanidine. The K_d value of di-*o*-tolyguanidine is 17.9 nM.⁷

3. Molecular biology and oocyte preparation

3.1. Data analysis

The obtained data were analyzed using Ana (Dr. Michael Pusch, Genova, Italy) and statistics were performed via OriginPro 2016 (OriginLab Corporation, Northampton, USA). The inhibitory effect of each compound was calculated using the following equation;

$$\text{inhibition} = 1 - \frac{I_c - I_h}{I_a - I_h}$$

where I_h describes the current without agonists; I_a represents the steady-state current with the agonists present; I_c is defined as the steady-state after agonist and compound bound. All dose-response curves were fitted to the following logistic equation;

$$y = \frac{A1 - A2}{1 + \left(\frac{x}{x_0}\right)^p} + A2$$

A1 describes the minimal inhibition of a compound and was set to 0. A2 represents the maximal inhibition of a compound; p is the slope of the curve; x_0 is defined as the concentration at half-maximal inhibition and x is the tested concentration, respectively.

Table 1: Inhibitory effects of the compounds **8**, **9** and **10** towards the GluN1-1a/GluN2B expressing oocytes. IC_{50} values, hill coefficients and A2 values (level of maximal inhibition) derived from the fitted curves.

compound	$IC_{50} \pm SE$ (nM)	$h \pm SE$	$A2 \pm SE$ (%)	n
8	574 ± 95	1.6 ± 0.5	86 ± 4	7
9	778 ± 142	1.6 ± 0.5	86 ± 5	8
10	496 ± 316	1.0 ± 0.2	92 ± 4	6

The data were recorded according to reference⁸.

4. References

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