

Supporting information

Synthesis and pharmacological evaluation of enantiomerically pure *endo*-configured KOR agonists with 2-azabicyclo[3.2.1]octane scaffold

Hendrik Jonas,^{a,b} Daniele Aiello,^b Bastian Frehland,^a Kirstin Lehmkuhl,^a Dirk Schepmann,^a Jens Köhler,^a Patrizia Diana,^b Bernhard Wunsch*^a

^a Institut für Pharmazeutische und Medizinische Chemie der Universität Münster, Corrensstraße 48, D-48149 Münster, Germany.

Tel.: +49-251-8333311; Fax: +49-251-8332144; E-mail: wunsch@uni-muenster.de

^b Dipartimento di Scienze e Tecnologie Biologiche Chimiche e Farmaceutiche, Università degli Studi di Palermo, Via Archirafi 32, I-90123 Palermo, Italy.

Content	page
1. Purity data of test compounds	S2
2. NOESY experiments	S3
3. Chiral HPLC	S8
4. Receptor binding studies	S9
5. Receptor affinity of non-racemic amines (ee ~50 %)	S13
6. References	S14
7. NMR spectra	S15

1. Purity data of test compounds determined by HPLC

Equipment: Pump: LPG-3600, degasser: DG-1210, autosampler: WPS 3000 PL, UV-detector: VWD-3400RS, interface: DIONEX UltiMate 3000, data acquisition: Chromeleon 7 (Thermo Fisher Scientific); 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 $\lambda = 210$ nm; solvents: A: demineralized 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: gradient from 90 % to 0 %; 29 - 31 min: 0 %; 31 - 31.5 min: gradient from 0 % to 90 %; 31.5 - 40 min: 90 %. Unless otherwise noted, the purity of all test compounds is higher than 95 %.

Purity data of test compounds

compound	ee	purity
(1 <i>S</i> ,5 <i>S</i> ,7 <i>R</i>)- 10a	53.6 %	99.1 %
(1 <i>S</i> ,5 <i>S</i> ,7 <i>R</i>)- 10a	99.9 %	95.1 %
(1 <i>R</i> ,5 <i>R</i> ,7 <i>S</i>)- 10a	52.8 %	97.2 %
(1 <i>R</i> ,5 <i>R</i> ,7 <i>S</i>)- 10a	96.3 %	95.7 %
(1 <i>S</i> ,5 <i>S</i> ,7 <i>R</i>)- 10b	54.3 %	99.2 %
(1 <i>S</i> ,5 <i>S</i> ,7 <i>R</i>)- 10b	99.9 %	98.0 %
(1 <i>R</i> ,5 <i>R</i> ,7 <i>S</i>)- 10b	49.8 %	98.6 %
(1 <i>R</i> ,5 <i>R</i> ,7 <i>S</i>)- 10b	98.8 %	95.1 %

2. NOESY experiments

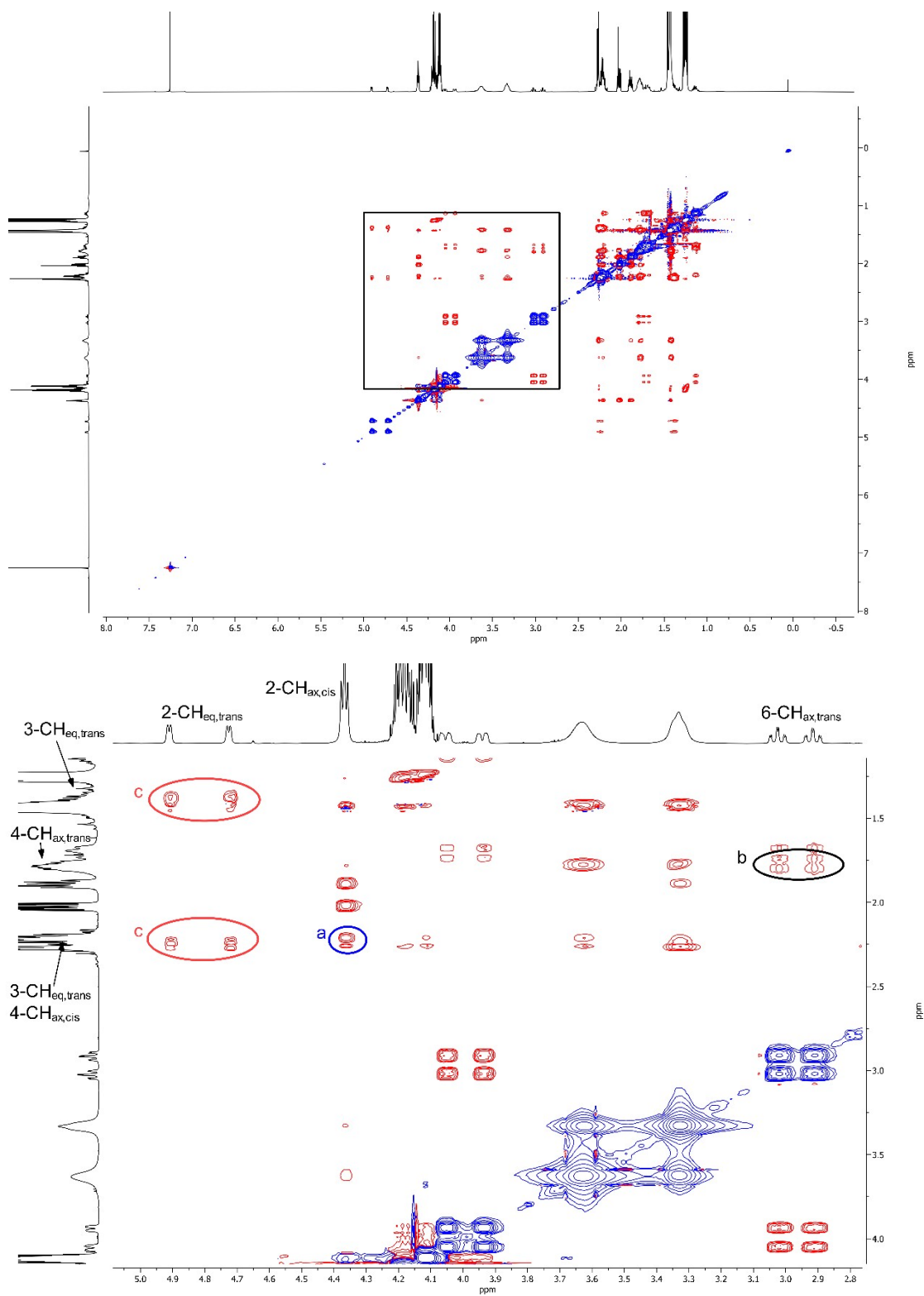
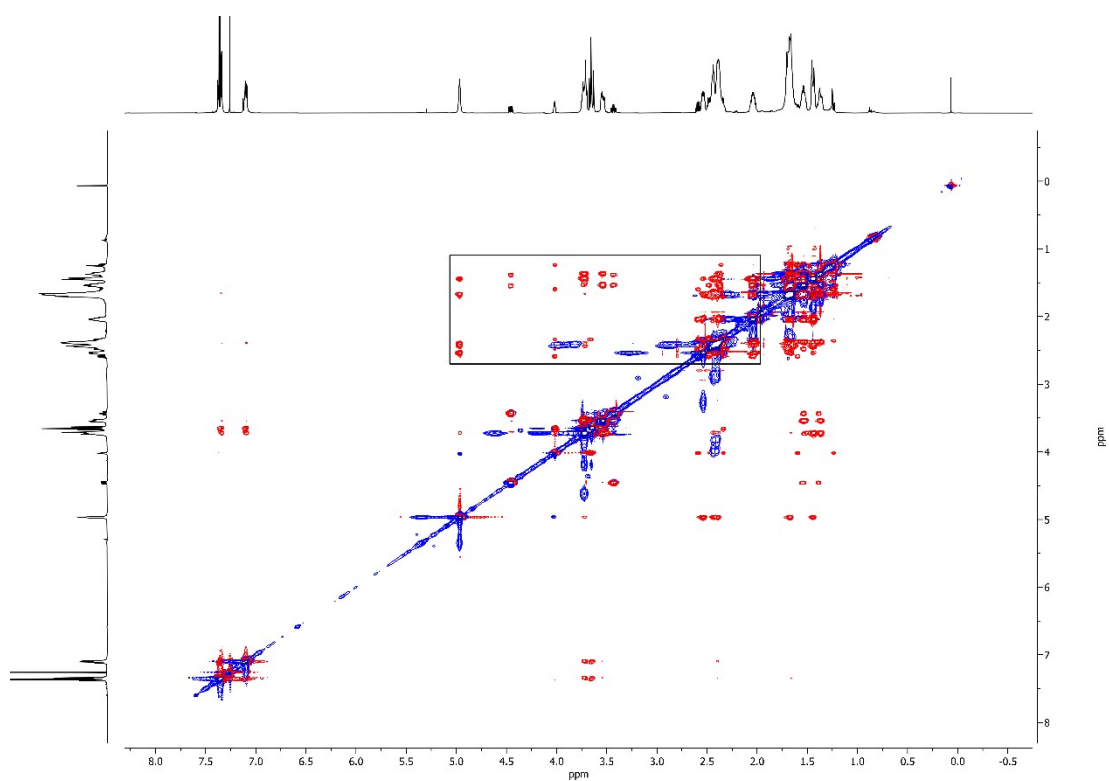


Figure S1: NOESY spectrum of diesters *cis*-**12** and *trans*-**12** in the ratio 3:1. Due to rotational isomerism along the Boc group, two sets of signals are seen for *trans*-**12**. Top:

Complete NOESY spectrum. Bottom: Magnification of the region of interest. A Cross peak between 4-H (2.16 – 2.18 ppm) and 2-H (4.37 ppm, mark a) for *cis*-**12**, and two cross peaks between 4-H (1.76 – 1.84 ppm) and 6-H (2.92 ppm and 3.02 ppm, mark b) and 2-H (4.72 ppm and 4.91 ppm) and 3-H_{ax} (1.34 – 1.43 ppm) as well as 3-H_{eq} (2.21 – 2.29 ppm, mark c) for *trans*-**12** can be detected.



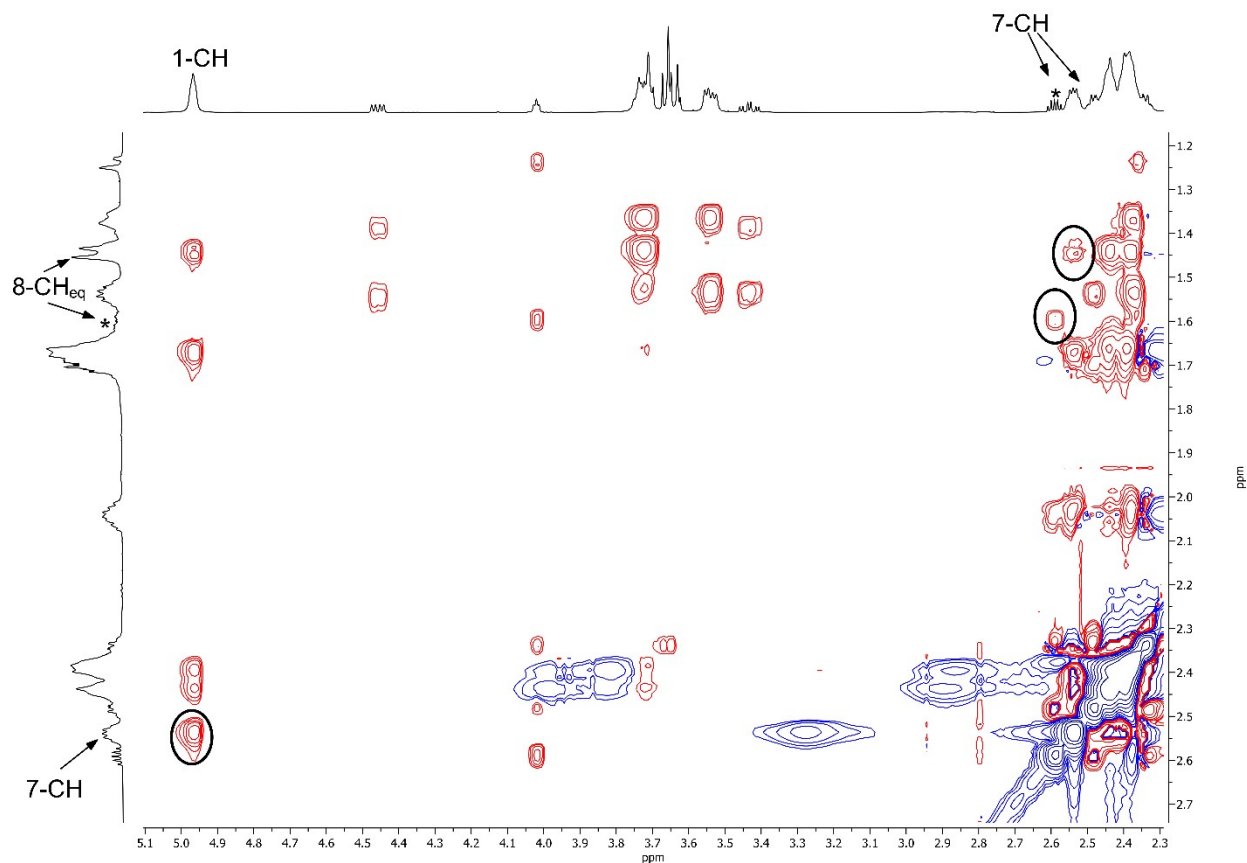


Figure S2: NOESY spectrum of bicyclic amine **10a**. Top: Complete NOESY spectrum. Bottom: Magnification of the region of interest. Cross peaks between 7-H (2.51 – 2.57 ppm) and 8-H_{eq} (1.41 – 1.46 ppm) and 7-H and 1-CH (4.97 – 5.02 ppm) for the major rotational isomer, and a cross peak between 7-H* (2.58 – 2.62 ppm) and 8-H_{eq}* (1.58 – 1.62 ppm) for the minor rotational isomer can be detected. The corresponding cross peaks are marked in black.

S6

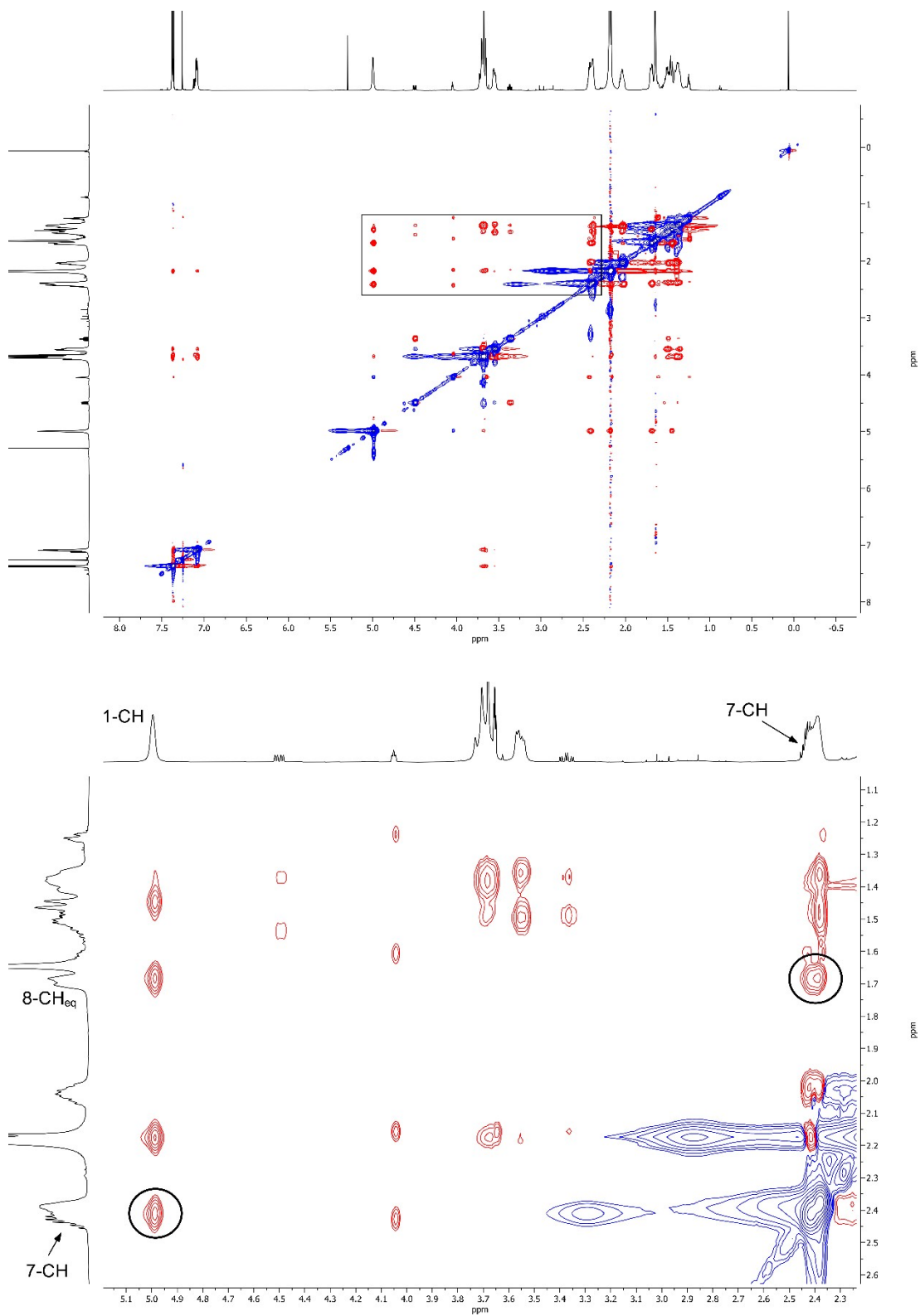


Figure S3: NOESY spectrum of bicyclic amine **10b**. Top: Complete NOESY spectrum. Bottom: Magnification of the region of interest. A Cross peak between 7-H (2.36 – 2.46

ppm) and 8-H_{eq} (1.66 – 1.72 ppm) and a cross peak between 7-H (2.36 – 2.46 ppm) and 1-H (4.97 – 5.02 ppm) for the minor rotational isomer can be detected. The corresponding cross peaks are marked in black.

3. Chiral HPLC

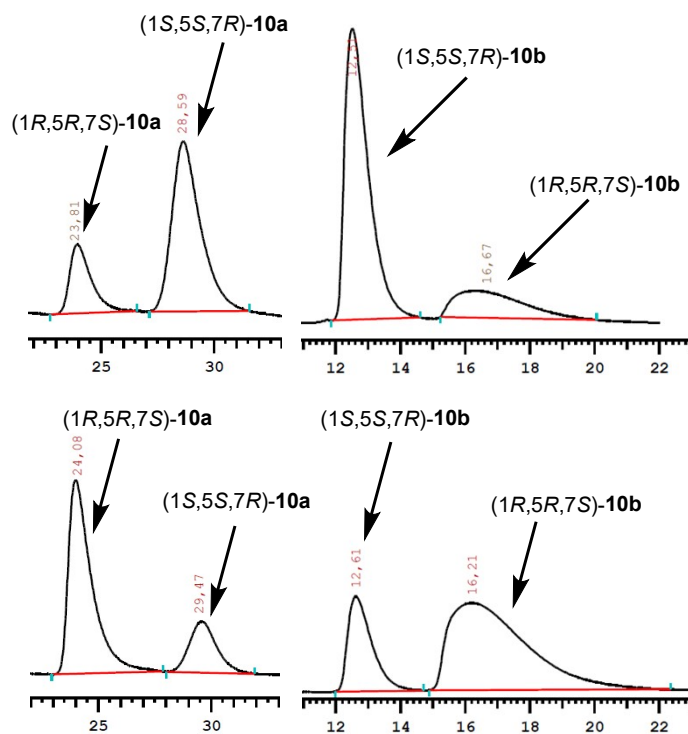


Figure S4: HPLC chromatogram before the separation of enantiomers **10a,b**. Column: Daicel Chiralpak IA; eluent: *iso*-hexane/ EtOH = 9:1 + 0.1 % diethylamine + 0.1 % formic acid (**10a**), *iso*-hexane/ *iso*-propanol = 95:5 + 0.1 % diethylamine (**10b**), isocratic elution; flow rate: 1.0 mL/min; detection λ = 275 nm. The addition of formic acid led to an inversion of the signals.

4. Receptor binding studies

4.1. Materials

Guinea pig brains, rat brains and rat livers were commercially available (Harlan-Winkelmann, Borcheln, 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).

4.2. Preparation of membrane homogenates from guinea pig brain¹⁻³

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.

4.3. Preparation of membrane homogenates from rat brain⁴⁻⁶

5 rat brains (species: Sprague Dawley rats) 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.

4.4. Preparation of membrane homogenates from rat liver¹⁻³

2 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.

4.5. Protein determination

The protein concentration was determined by the method of Bradford,⁷ modified by Stoscheck.⁸ 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).

4.6. 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 $^{\circ}\text{C}$. The solid scintillator was melted on the dried filtermats at a temperature of 95 $^{\circ}\text{C}$ 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.⁹ The K_i values are given as mean value \pm SEM from three independent experiments.

4.7. Performance of the KOR assay⁴⁻⁶

The assay was performed with the radioligand [^3H]U-69,593 (55 Ci/mmol, BIOTREND). The thawed guinea pig brain membrane preparation (about 100 μg of the protein) was incubated with various concentrations of test compounds, 1 nM [^3H]U-69,593, and TRIS-MgCl₂-buffer (50 mM TRIS, 8 mM MgCl₂, pH 7.4) at 37 $^{\circ}\text{C}$. The non-specific binding was determined with 10 μM unlabeled U-69,593. The K_d value of U-69,593 is 0.69 nM.

4.8. Performance of the MOR assay⁴⁻⁶

The assay was performed with the radioligand [^3H]DAMGO (51 Ci/mmol, Perkin Elmer). The thawed guinea pig brain membrane preparation (about 100 μg of the protein) was incubated with various concentrations of test compounds, 3 nM [^3H]DAMGO, and TRIS-MgCl₂-buffer (50 mM TRIS, 8 mM MgCl₂, pH 7.4) at 37 $^{\circ}\text{C}$. The non-specific binding was determined with 10 μM unlabeled naloxone. The K_d value of DAMGO is 0.57 nM.

4.9. Performance of the DOR assay⁴⁻⁶

The assay was performed with the radioligand [³H]DPDPE (69 Ci/mmol, BIOTREND). The thawed rat brain membrane preparation (about 75 µg of the protein) was incubated with various concentrations of test compounds, 3 nM [³H]DPDPE, and TRIS-MgCl₂-buffer (50 mM TRIS, 8 mM MgCl₂, pH 7.4) supplemented with SIGMAFAST® protease inhibitor mix (Sigma Aldrich Biochemicals, Hamburg, Germany; 1 tablet dissolved in 100 mL of buffer) at 37 °C. The non-specific binding was determined with 10 µM unlabeled morphine. The *K_d* value of DPDPE is 0.65 nM.

4.10. Performance of the σ_1 receptor assay¹⁻³

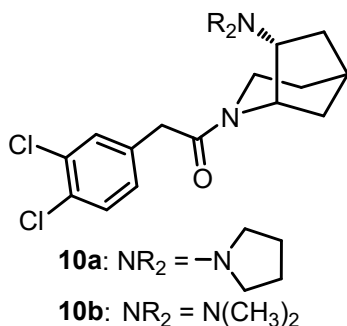
The assay was performed with the radioligand [³H]-(+)-pentazocine (22.0 Ci/mmol; Perkin Elmer). The thawed membrane preparation of guinea pig brain (about 100 µg of the protein) was incubated with various concentrations of test compounds, 2 nM [³H]-(+)-pentazocine, and TRIS buffer (50 mM, pH 7.4) at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

4.11. Performance of the σ_2 receptor assay¹⁻³

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.¹¹

5. Receptor affinity of non-racemic amines (ee ~50 %)

Table S11: Affinities of non-racemic 2-azabicyclo[3.2.1]octanes **10aa** and **10b** (ee ~50 %).



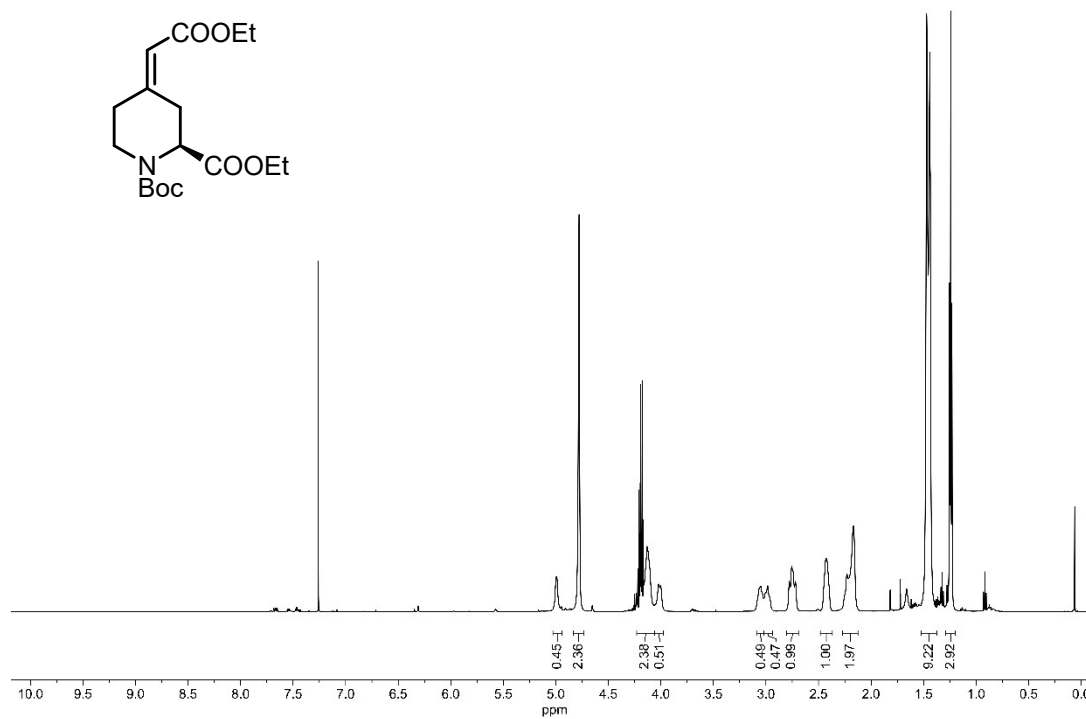
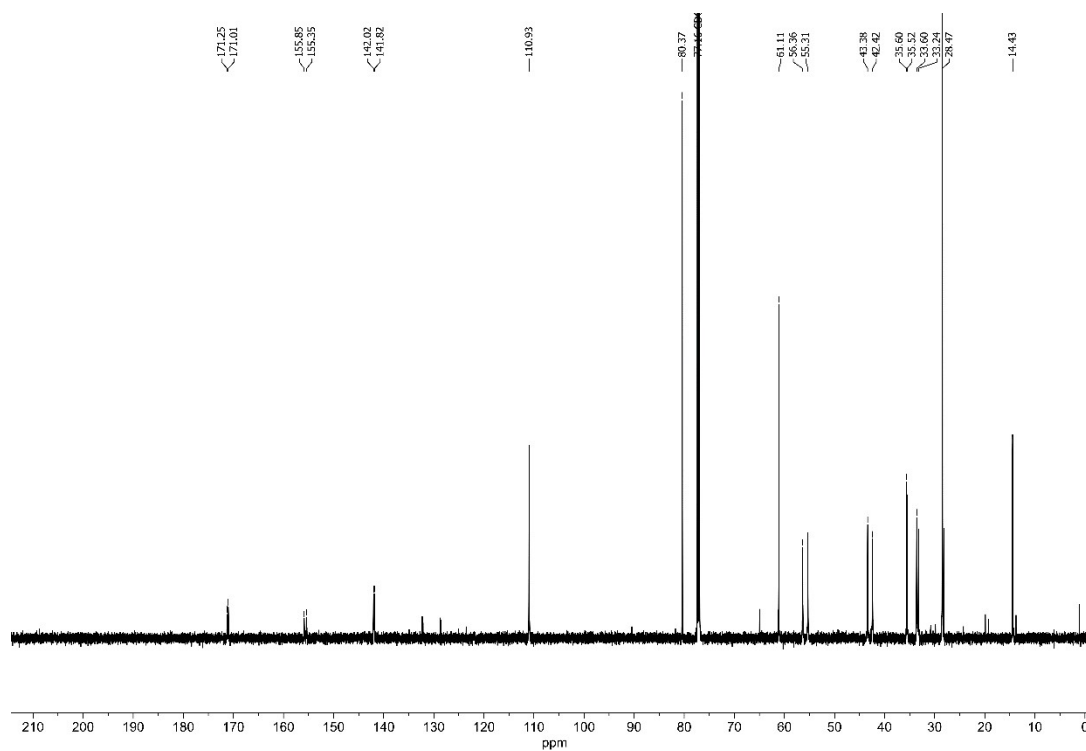
compd.	NR ₂	ee [%]	<i>K_i</i> ± SEM [nM] (n = 3) ^{a)}				
			KOR	MOR	DOR	σ ₁	σ ₂
			[³ H] U-69,593	[³ H] DAMGO	[³ H] DPDPE	(+)- [³ H]pentazocine	[³ H] DTG
10a		53.6	12 ± 2	8%	3200	7 ± 4	310
<i>ent</i> - 10a		52.8	53 ± 5	0%	1800	20 ± 4	243
10b	N(CH ₃) ₂	54.3	18 ± 2	10%	0%	76 ± 6	17%
<i>ent</i> - 10b		49.8	49 ± 18	5%	565	32 ± 10	366

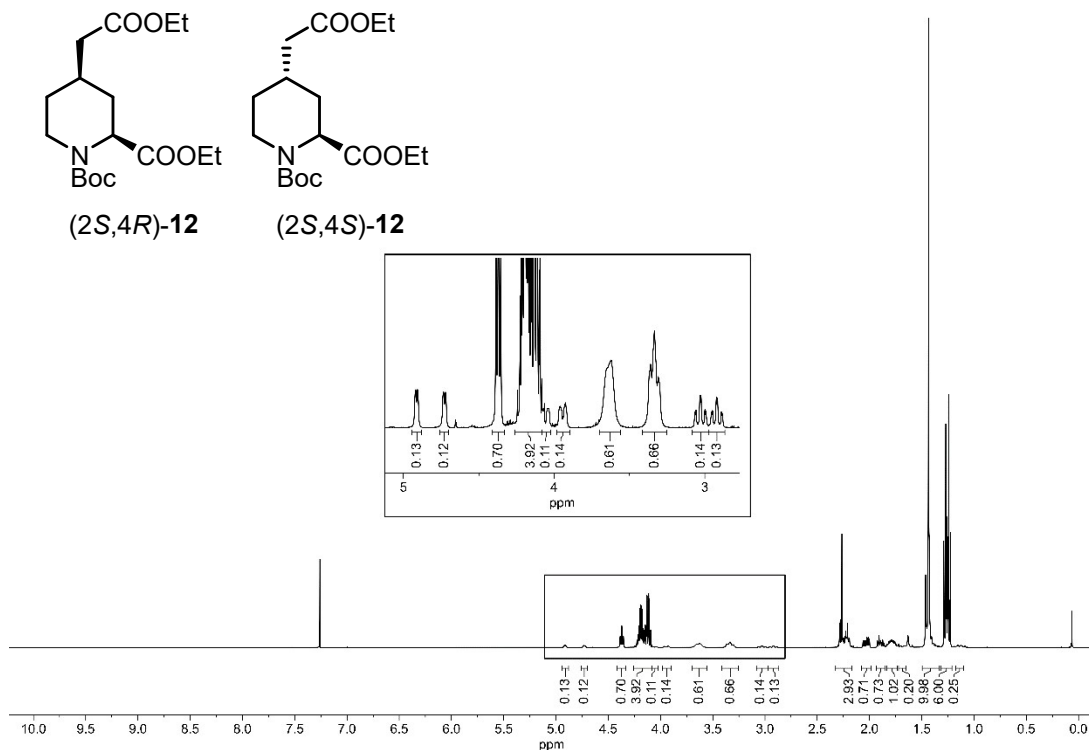
^{a)} A value in % reflects the inhibition of the radioligand binding at a test compound concentration of 1 μM. *K_i* values without SEM values represent the mean of two experiments (n = 2), *K_i* values with SEM values represent the mean of three experiments (n = 3).

6. References

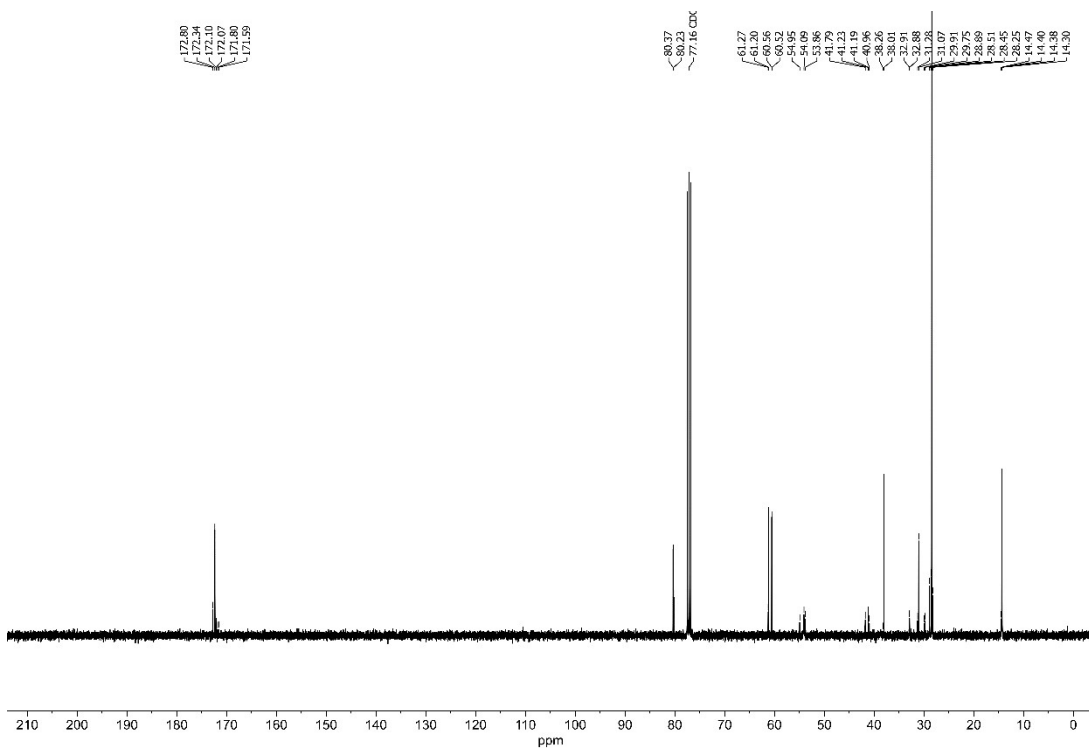
1. Hasebein, P.; Frehland, B.; Lehmkuhl, K.; Fröhlich, R.; Schepmann, D.; Wünsch, B. Synthesis and pharmacological evaluation of like- and unlike-configured tetrahydro-2-benzazepines with the α -substituted benzyl moiety in the 5-position. *Org. Biomol. Chem.* **2014**, *12* (29), 5407–5426.
2. Meyer, C.; Neue, B.; Schepmann, D.; Yanagisawa, S.; Yamaguchi, J.; Würthwein, E.-U.; Itami, K.; Wünsch, B. Improvement of σ_1 receptor affinity by late-stage C-H bond arylation of spirocyclic lactones. *Bioorg. Med. Chem.* **2013**, *21* (7), 1844–1856.
3. Miyata, K.; Schepmann, D.; Wünsch, B. Synthesis and σ receptor affinity of regioisomeric spirocyclic furopyridines. *Eur. J. Med. Chem.* **2014**, *83*, 709–716.
4. Wittig, C.; Schepmann, D.; Soeberdt, M.; Daniliuc, C. G.; Wünsch, B. Stereoselective Synthesis of Conformationally Restricted KOR Agonists Based on the 2,5-diazabicyclo[2.2.2]octane Scaffold. *Org. Biomol. Chem.* **2017**, *15* (31), 6520–6540.
5. Geiger, C.; Zelenka, C.; Lehmkuhl, K.; Schepmann, D.; Englberger, W.; Wünsch, B. Conformationally Constrained K Receptor Agonists: Stereoselective Synthesis and Pharmacological Evaluation of 6,8-Diazabicyclo[3.2.2]nonane Derivatives. *J. Med. Chem.* **2010**, *53*, 4212–4222.
6. Kracht, D.; Rack, E.; Schepmann, D.; Fröhlich, R.; Wünsch, B. Stereoselective Synthesis and Structure–affinity Relationships of Bicyclic κ Receptor Agonists. *Org. Biomol. Chem.* **2010**, *8*(1), 212–225.
7. Bradford, M. M. A Rapid and Sensitive Method for the Quantitation of Microgram Quantities of Protein Utilizing the Principle of Protein-Dye Binding. *Anal. Biochem.* **1976**, *72*, 248–254.
8. Stoscheck, C. Quantitation of Protein. *Methods Enzymol.* **1990**, *182*, 50–68.
9. Yung-Chi, C.; Prusoff, W. H. Relationship between the Inhibition Constant (KI) and the Concentration of Inhibitor Which Causes 50 per Cent Inhibition (I50) of an Enzymatic Reaction. *Biochem. Pharmacol.* **1973**, *22* (23), 3099–3108.
10. DeHaven-Hudkins, D. L.; Fleissner, L. C.; Ford-Rice, F. Y. Characterization of the Binding of ($[^3\text{H}]$)-Pentazocine to Sigma Recognition Sites in Guinea Pig Brain. *Eur. J. Pharmacol.* **1992**, *227* (4), 371–378.
11. Mach, R. H.; Smith, C. R.; Childers, S. R. Ibogaine Possesses a Selective Affinity for Sigma 2 Receptors. *Life Sci.* **1995**, *57* (4), 57–62.

7. NMR spectra

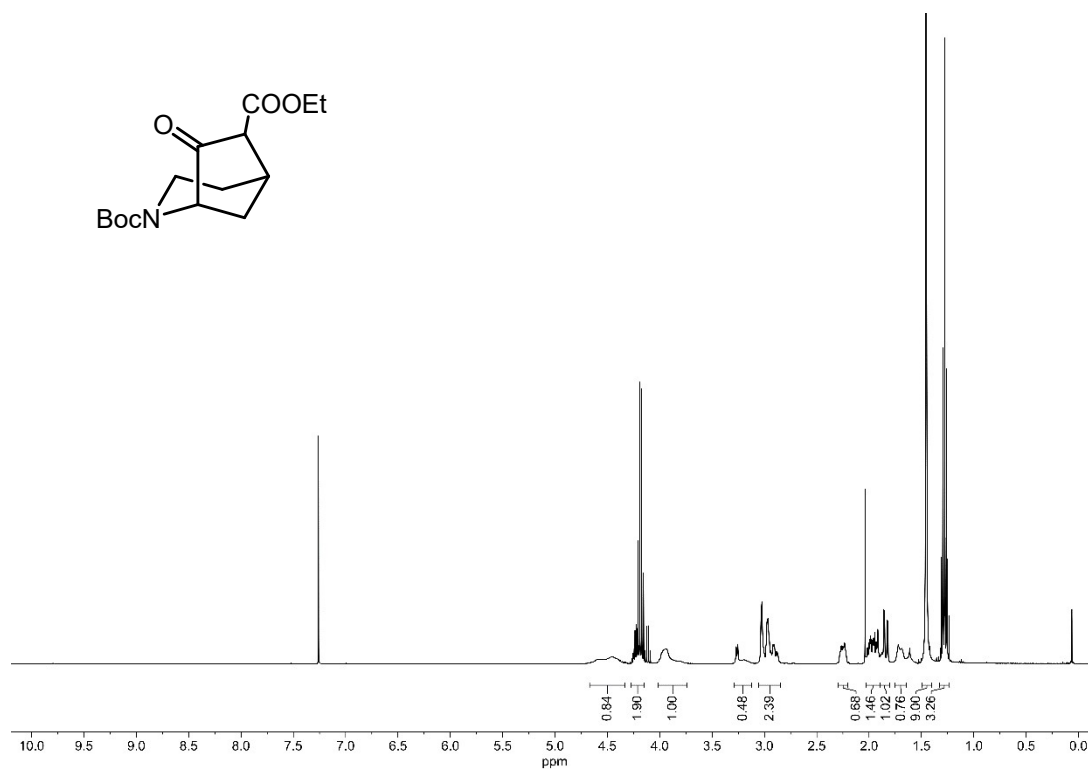
¹H NMR spectrum (CDCl₃) of **14**.¹³C NMR spectrum (CDCl₃) of **14**.



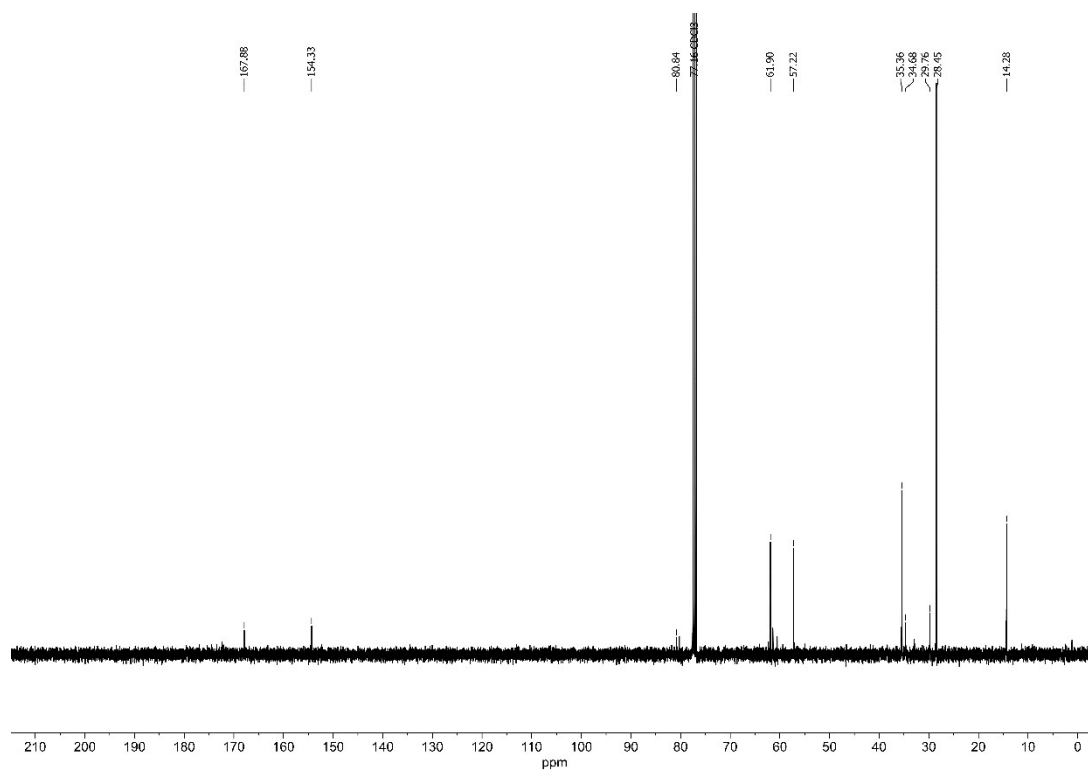
¹H NMR spectrum (CDCl₃) of (2*S*,4*R*)-**12** (major) and (2*S*,4*S*)-**12** (minor). The diastereomers were obtained in a 3:1 ratio. Due to rotational isomerism along the Boc group, the minor diastereomer shows two sets of signals.



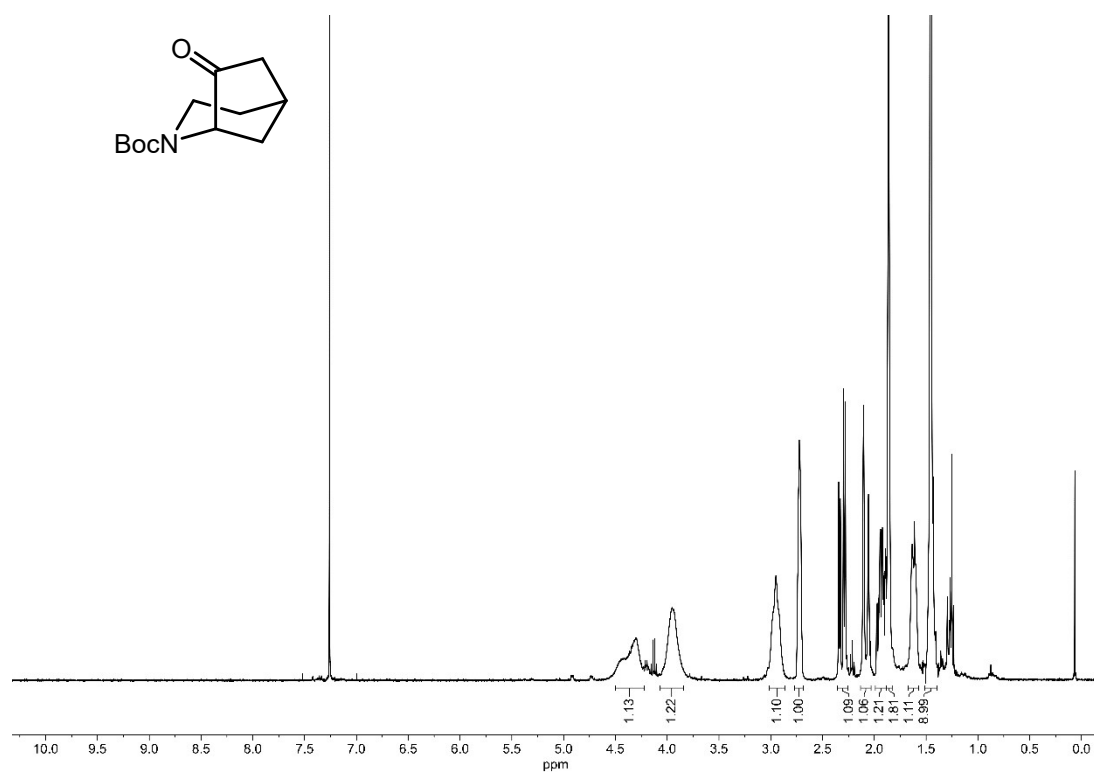
¹³C NMR spectrum (CDCl₃) of (2*S*,4*R*)-**12** (major) and (2*S*,4*S*)-**12** (minor).



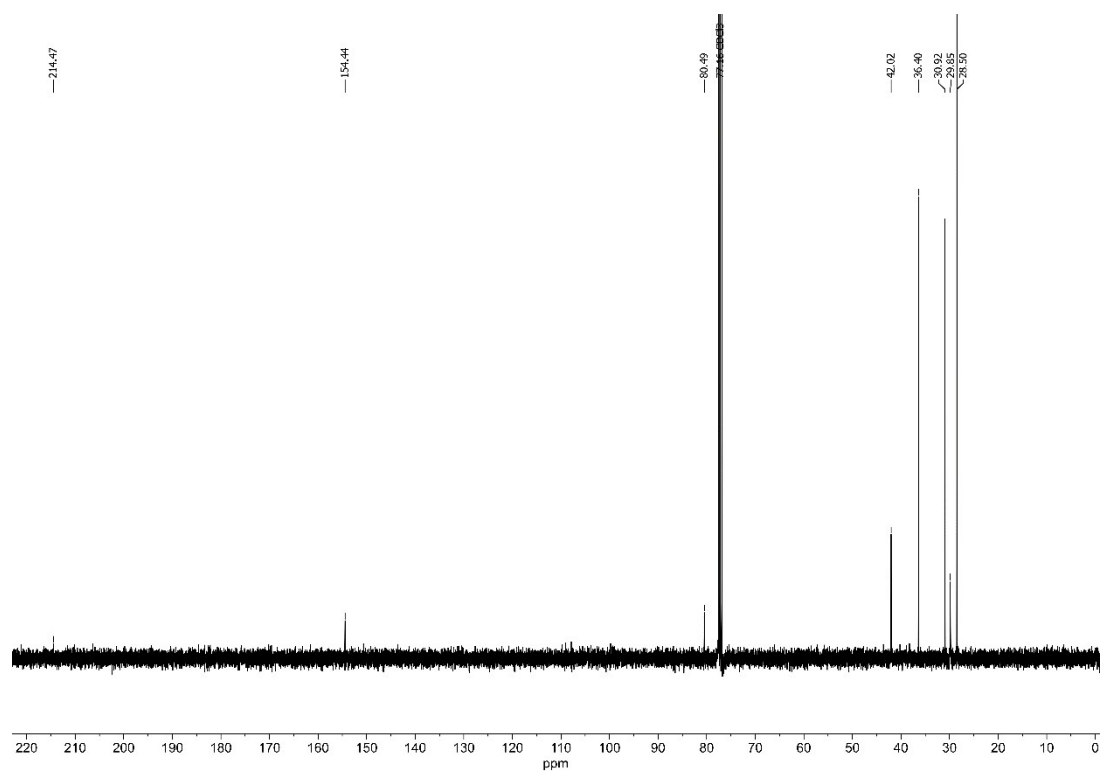
^1H NMR spectrum (CDCl_3) of **15**.



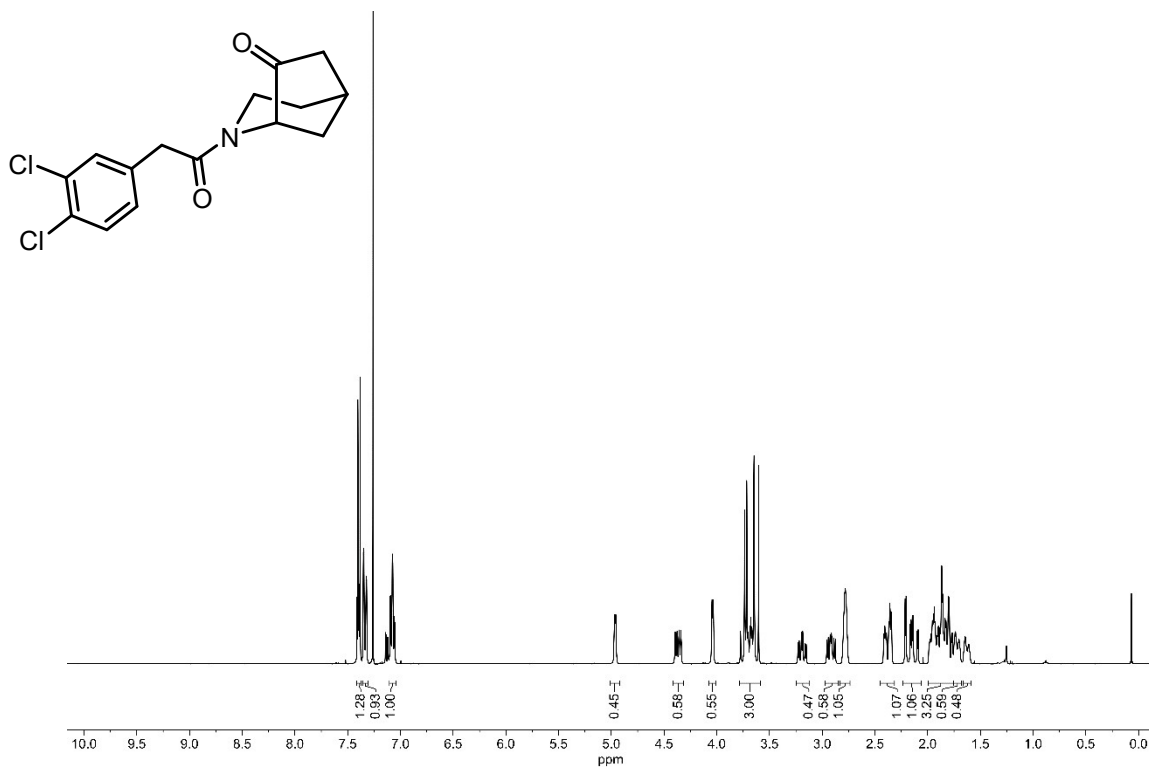
^{13}C NMR spectrum (CDCl_3) of **15**.



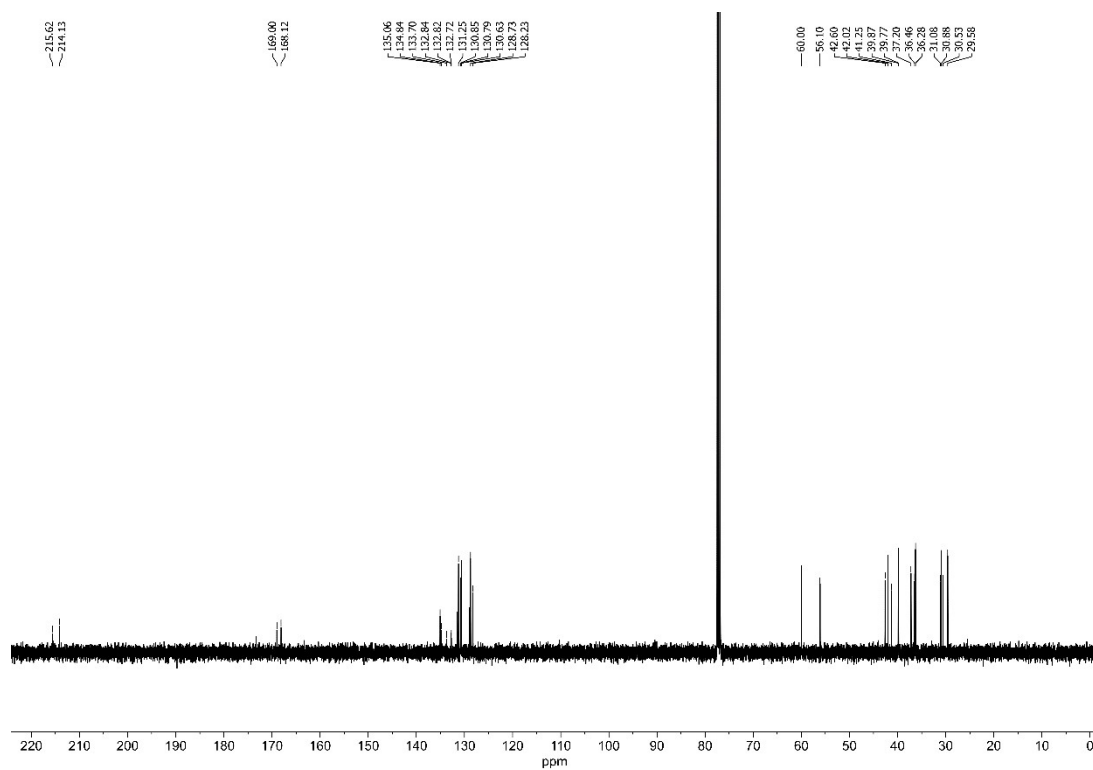
¹H NMR spectrum (CDCl₃) of **16**.



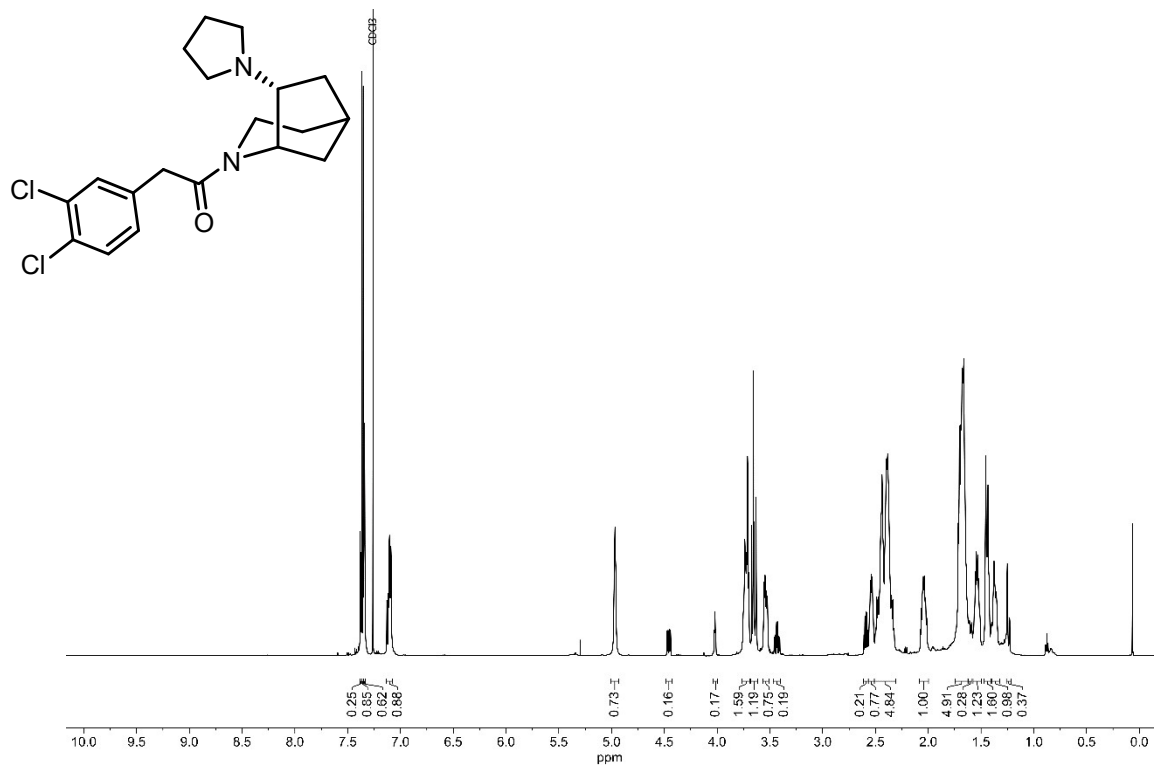
¹³C NMR spectrum (CDCl₃) of **15**.



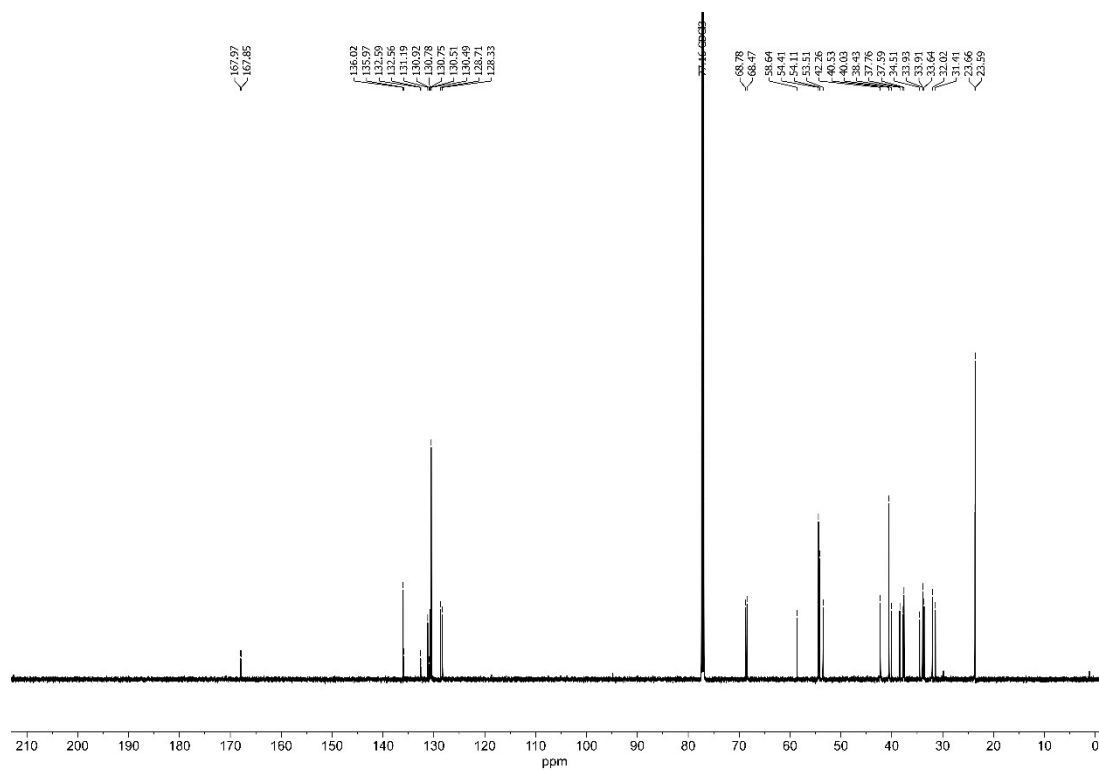
¹H NMR spectrum (CDCl₃) of **11**.



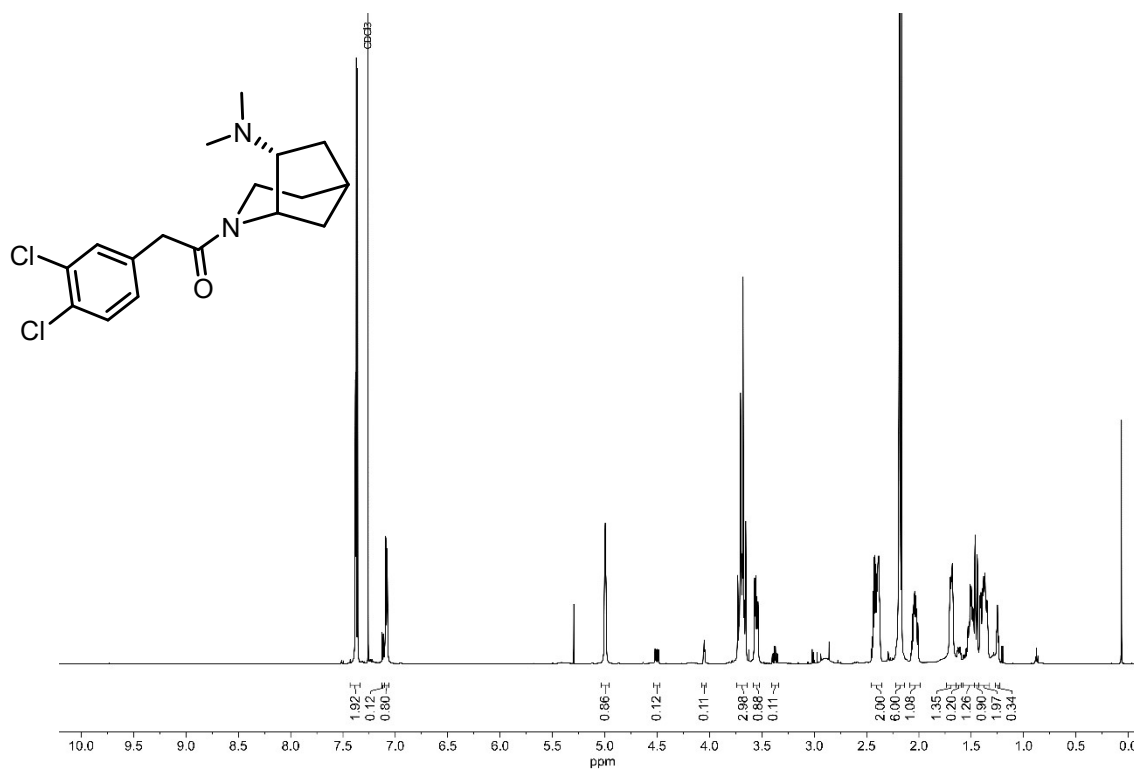
¹³C NMR spectrum (CDCl₃) of **10**.



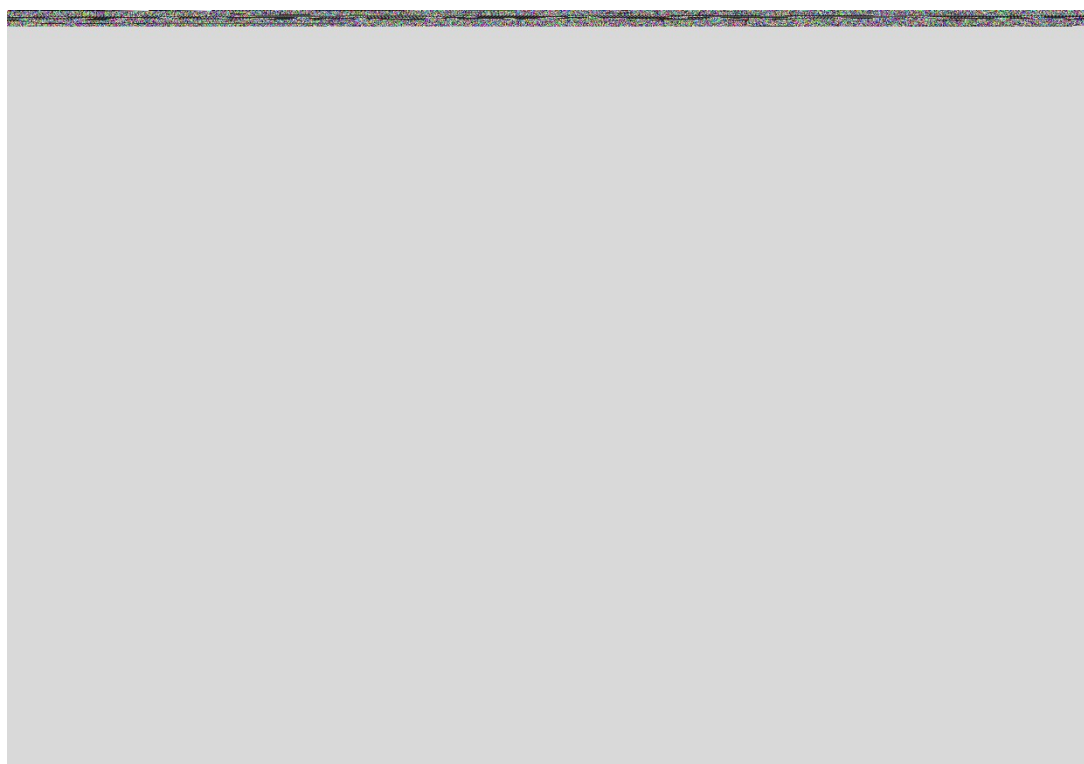
¹H NMR spectrum (CDCl₃) of **10a**.



¹³C NMR spectrum (CDCl₃) of **10a**.



¹H NMR spectrum (CDCl₃) of **10b**.



¹³C NMR spectrum (CDCl₃) of **10b**.