

Supporting Information

Development of Carborane-Based Halogenated Naphthyridinone-Analogues as Cannabinoid Receptor Type 2 (CB₂R) Ligands

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1 Experimental Section – Synthesis of compounds **1** and **E3_{o,m,p}**

The synthesis of compound **1** starts with the literature-known halogenation of aldehyde **E4** with bromine or iodine.^[1, 89] The work-up for the iodination was as following: sodium thiosulfate pentahydrate was dissolved in water (10%-aqueous solution) and sodium hydroxide, until pH 7 was reached, were added to the reaction mixture. After extraction with EtOAc (four times), drying over MgSO₄ and filtration, the semi-crude was purified by column chromatography (*n*-hexane/EtOAc, 4:1 → 1:1, (v/v)). In the second step, a Knoevenagel condensation was performed yielding **E6**,^[1] followed by the alkylation of N1 with *para*-fluorobenzyl chloride as published by Lucchesi *et al.*^[1] The basic ester hydrolysis of **E7** with LiOH·H₂O in THF/MeOH/H₂O (2:1:1, v/v/v) at 60 °C^[90] led to compound **1**. The synthesis of the aminocarboranes **E3_{o,m,p}** has been reported for the *ortho*-carborane derivative by Nie *et al.*^[100] and for the *meta*- and *para*-carborane derivatives by Scholz *et al.*^[101] and Choi and Byun^[102].

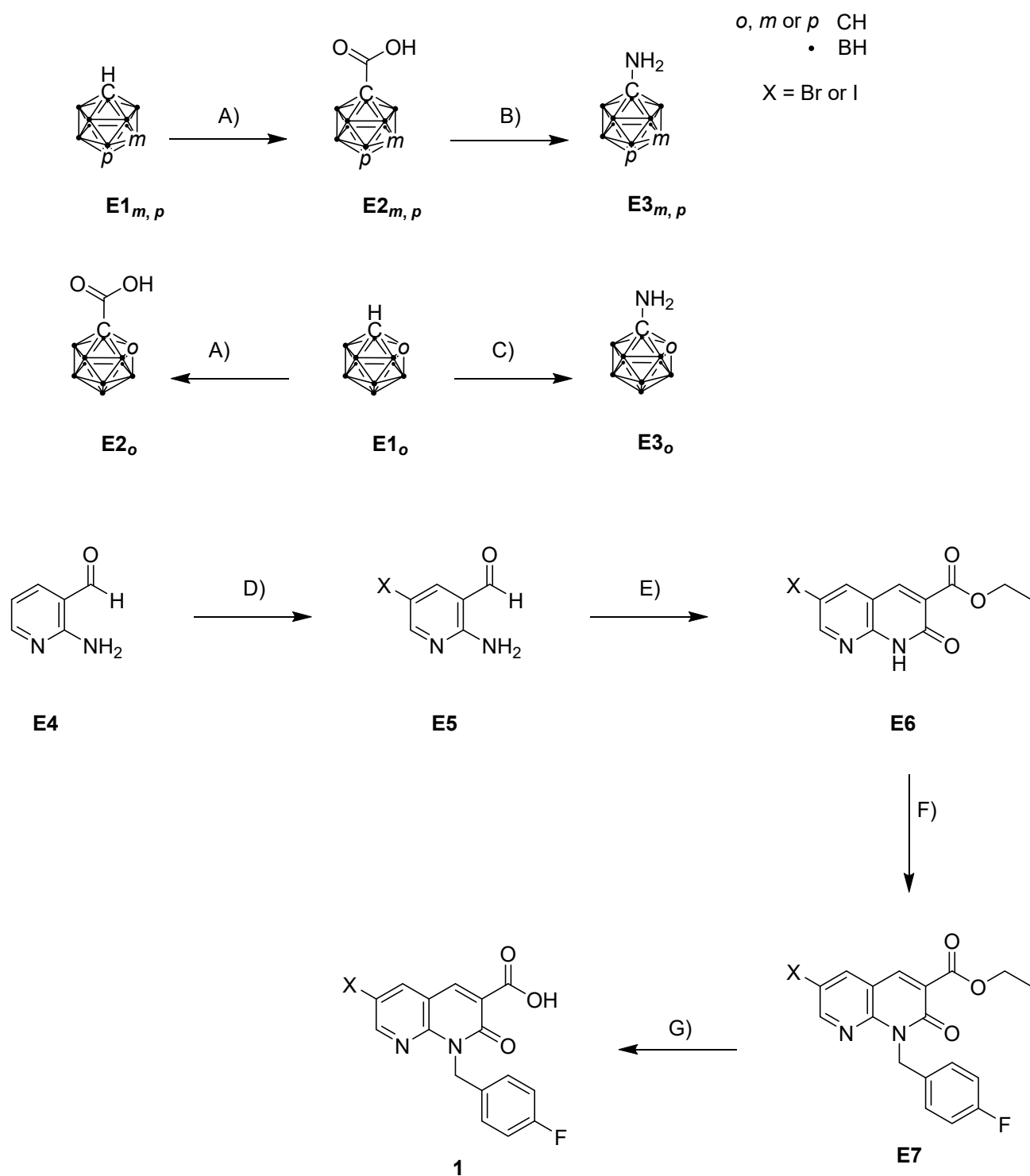


Figure S1. Synthesis of target compounds **2**_{*o,m,p*}, **3**_{*o,m,p*}, **4**, and **5**. Reagents and conditions: for **E3**_{*m,p*}: (A) (i) *n*-butyllithium, Et₂O, CO₂, -78°C → RT, 16-21 h (ii) HCl and B) (i) 4-DMAP, diphenylphosphoryl azide, NEt₃, *tert*-BuOH, reflux, 17 h (ii) TFA, CH₂Cl₂, RT, 1 d; for **E3**_{*o*}: C) *n*-butyllithium, benzyl azide, Et₂O, RT, 3h (ii) glacial acetic acid, 90°C, 2h; D) for X = Br: Br₂, AcOH, RT, 1-5 d; for X = I: HIO₄, I₂, CH₃COOH/H₂O/H₂SO₄ (117:5.8:1, v/v/v), 80°C, 19.5 h; E) diethylmalonate, piperidine, EtOH, reflux, 18-21.5 h, F) *para*-fluorobenzyl chloride, Cs₂CO₃, DMF, 50-61°C, 18.5 h; G) (i) LiOH·H₂O, THF/MeOH/H₂O, 2:1:1 (v/v/v), 60°C, 2.5-3 h, (ii) HCl.

2 NMR spectra of compounds $2_{o,m,p}$, $3_{o,m,p}$, 4, and 5

Compound 2_o

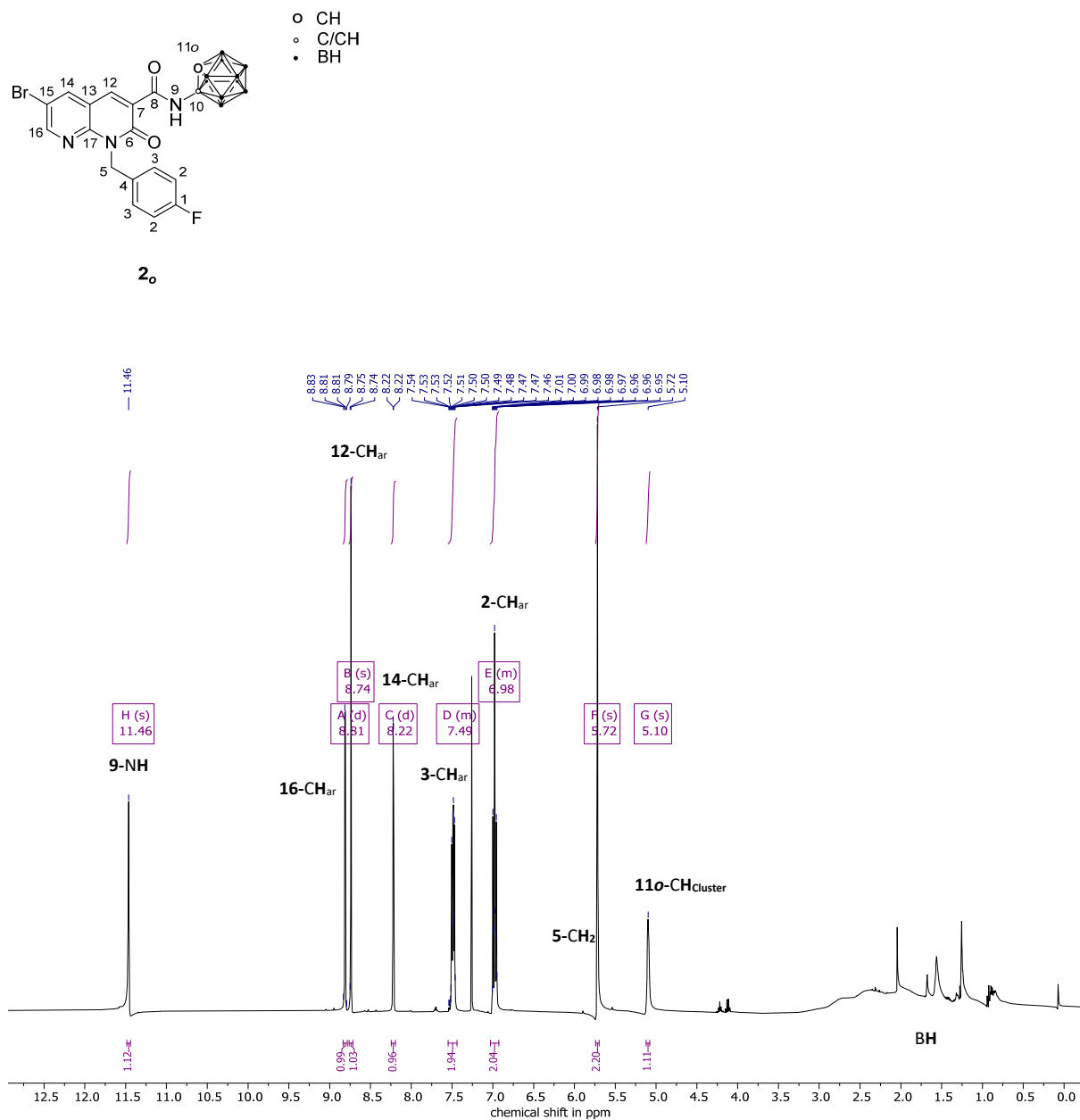


Figure S2. ^1H NMR spectrum of compound 2_o in CDCl_3 .

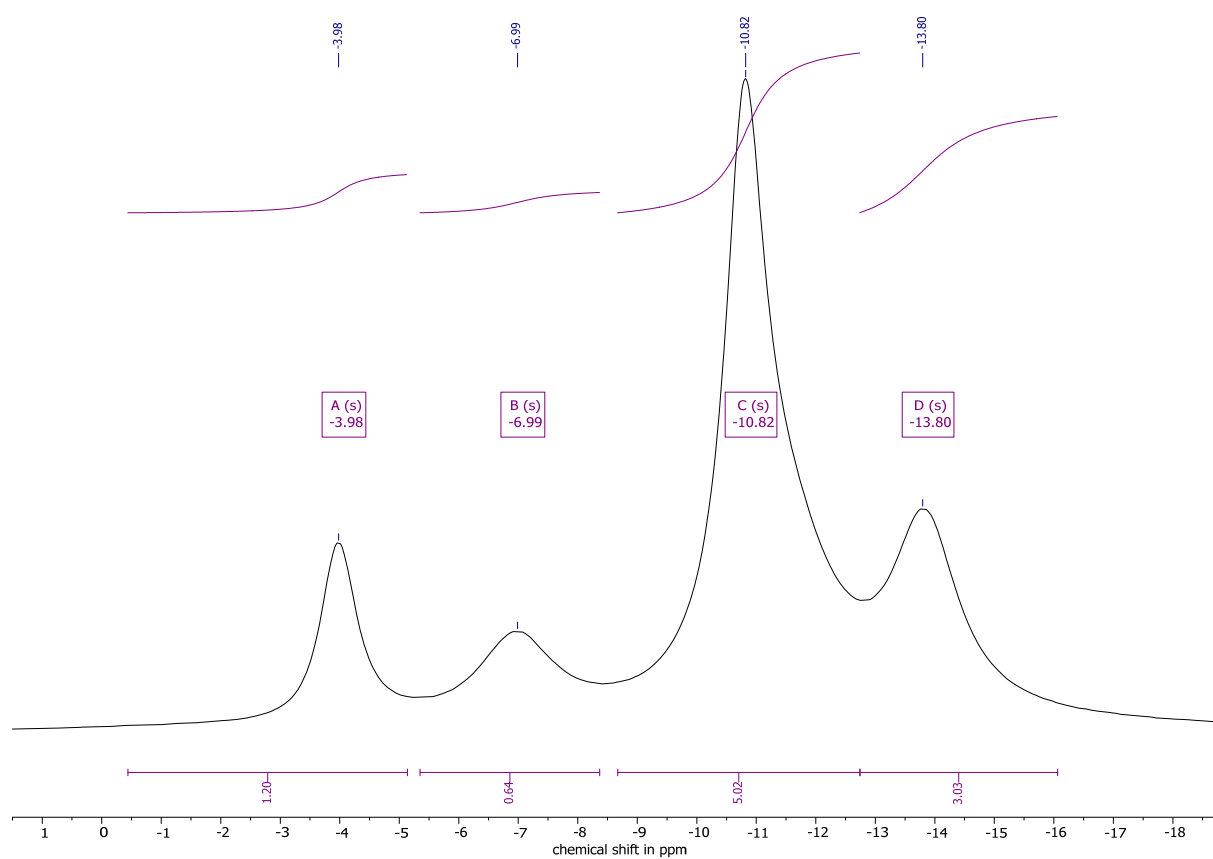


Figure S3. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **2o** in CDCl_3 .

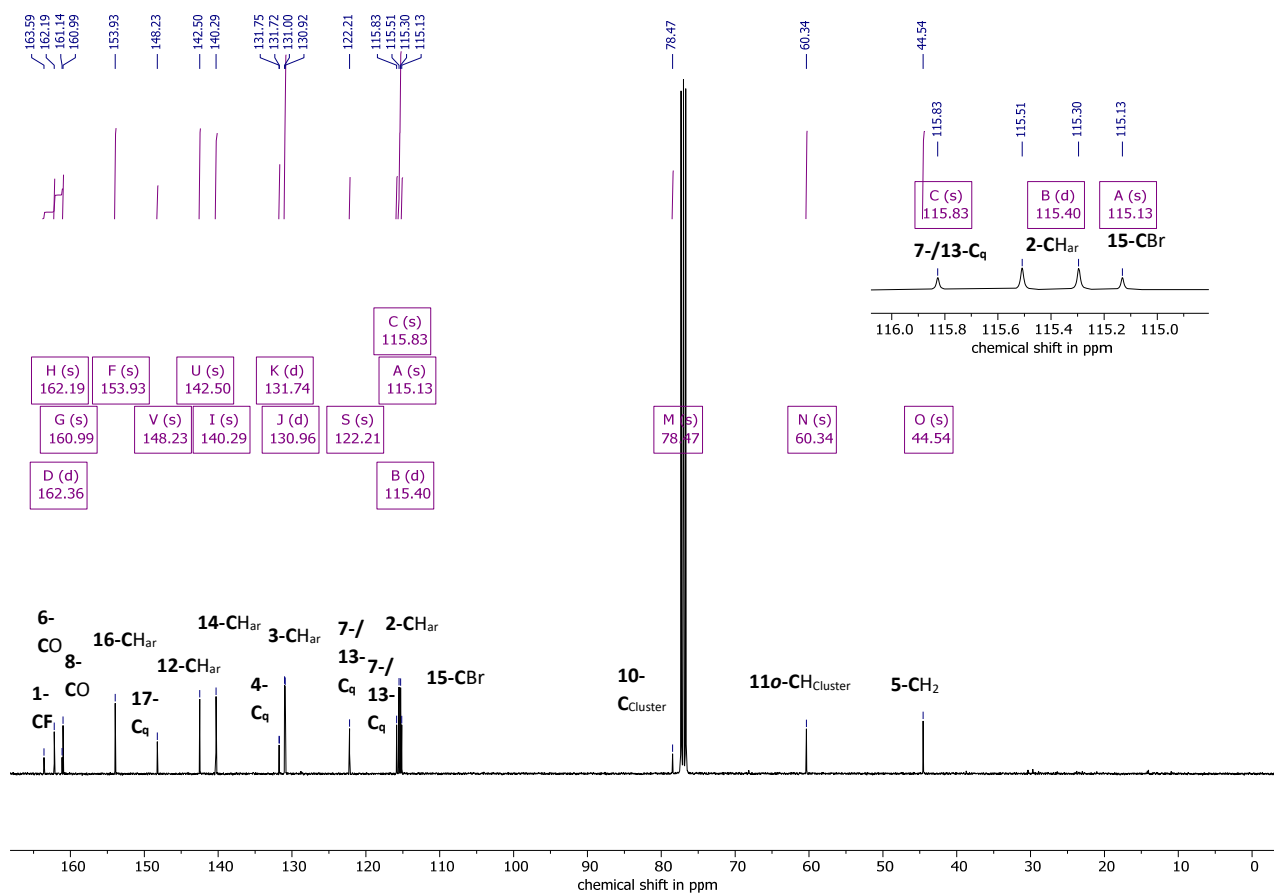


Figure S4. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **2o** in CDCl_3 .

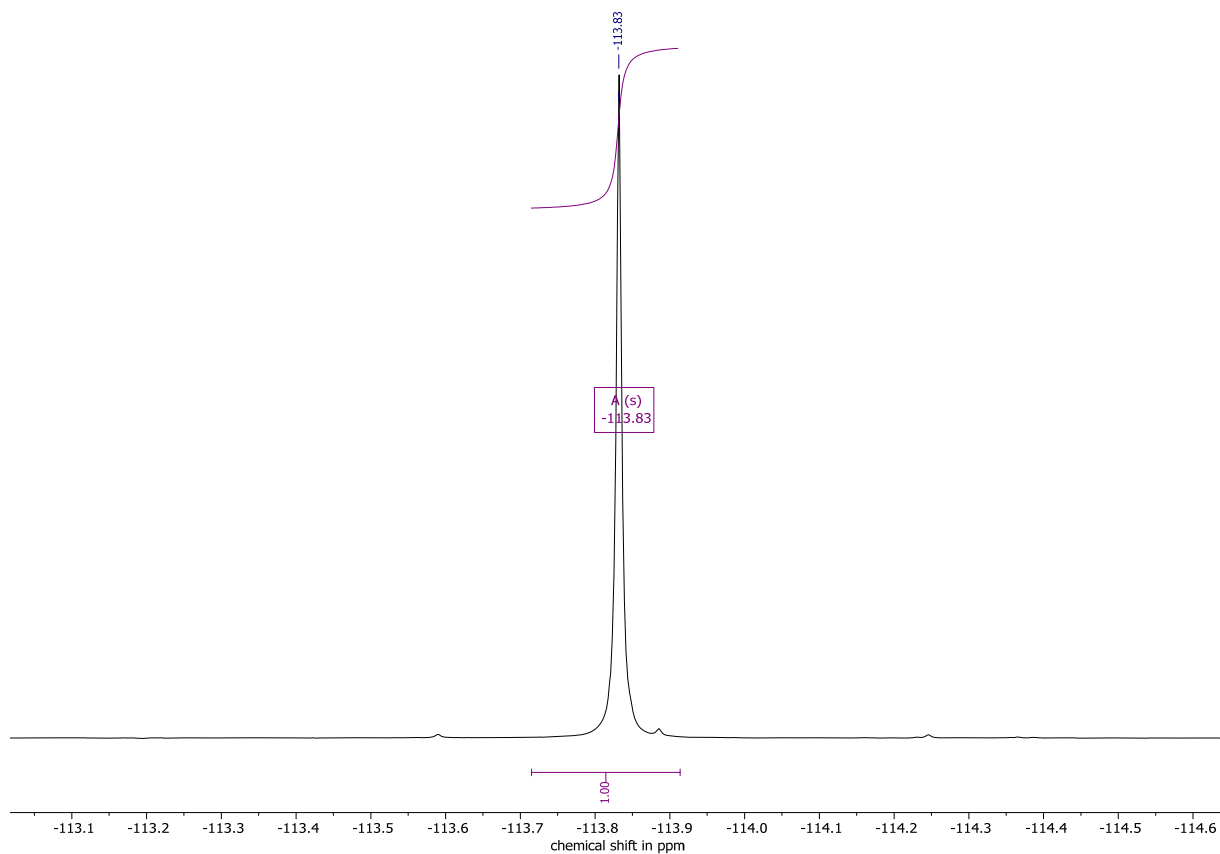


Figure S5. $^{19}\text{F}\{^1\text{H}\}$ NMR spectrum of compound **2o** in CDCl_3 .

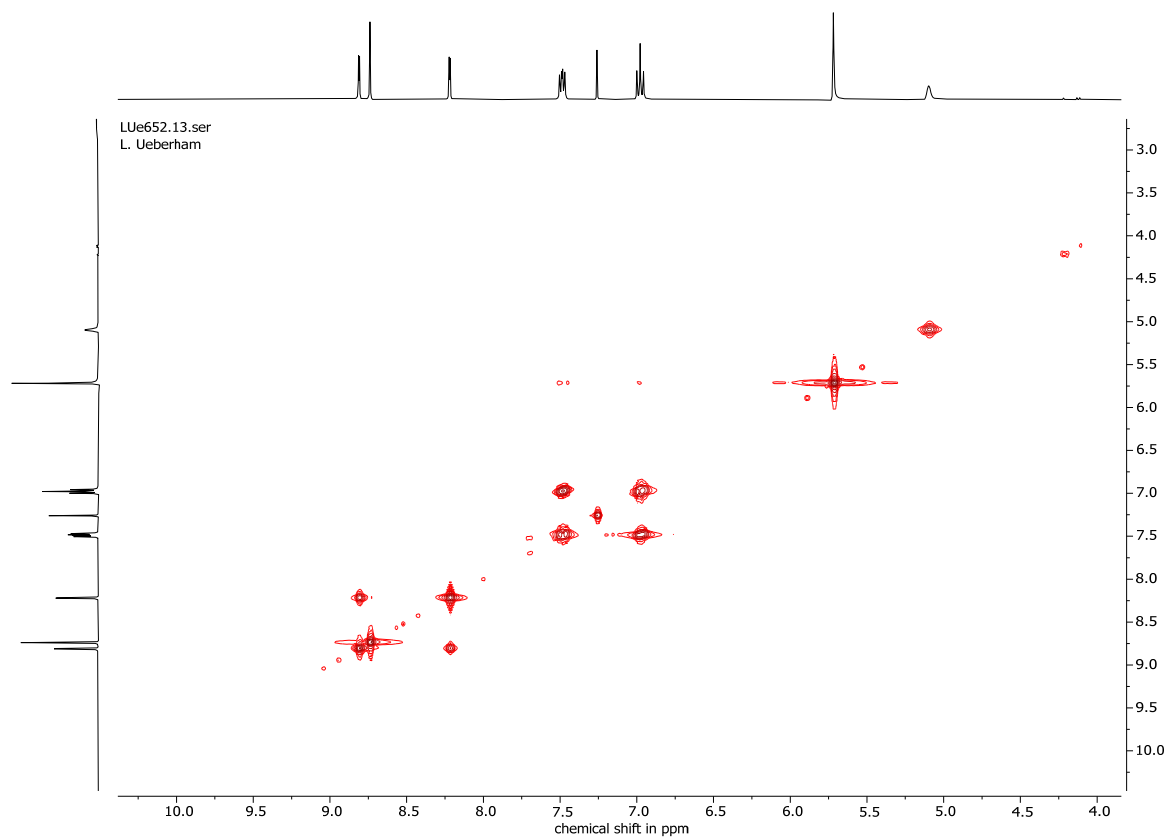


Figure S6. COSY (^1H , ^1H) NMR spectrum of compound **2o** in CDCl_3 (magnified spectrum, zoomed in).

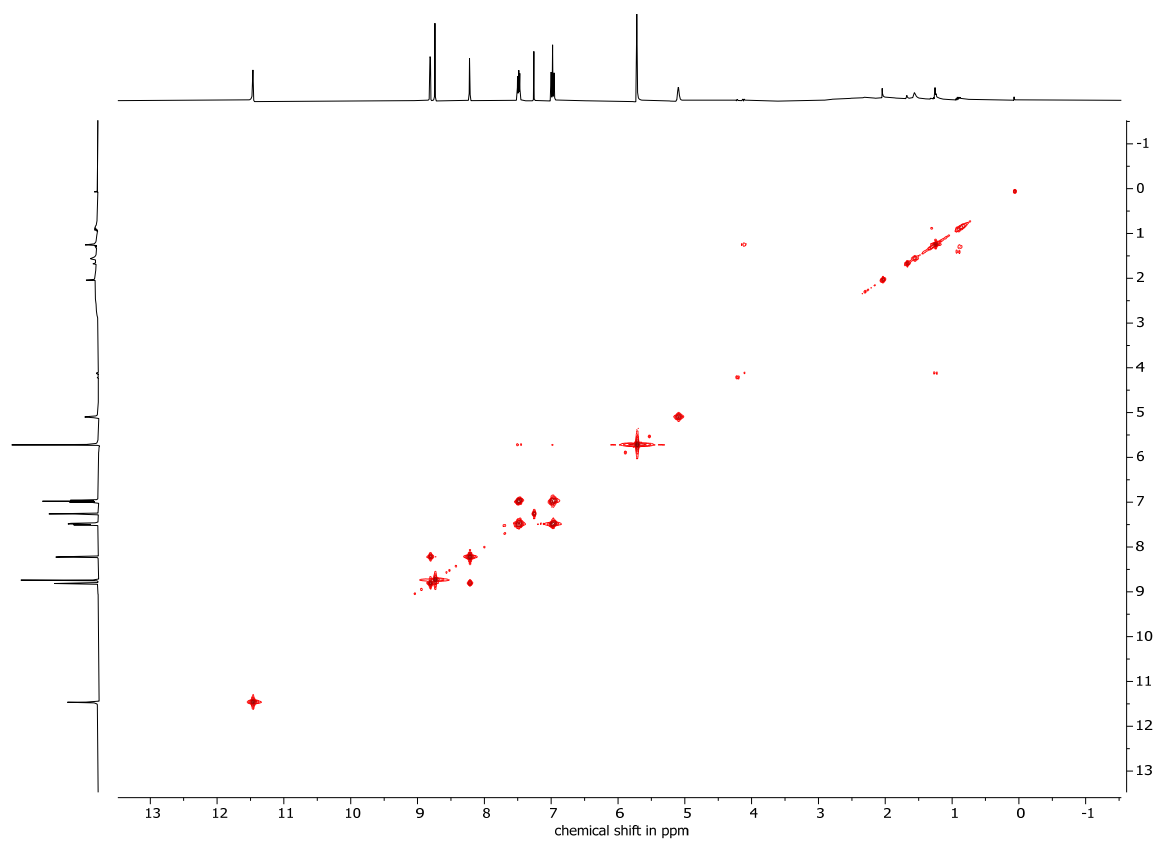


Figure S7. COSY (^1H , ^1H) NMR spectrum of compound **2o** in CDCl_3 full spectrum.

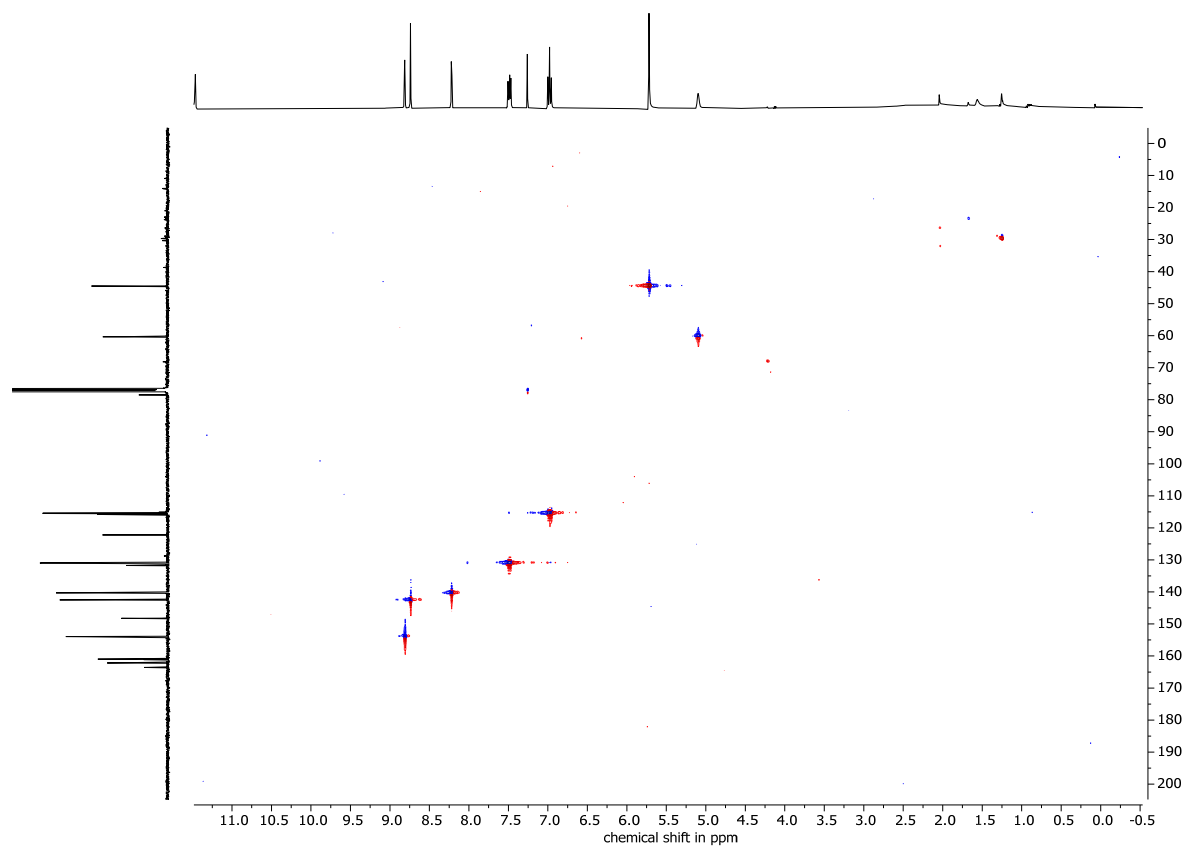


Figure S8. HSQC (^1H , ^{13}C) NMR spectrum of compound **2o** in CDCl_3 .

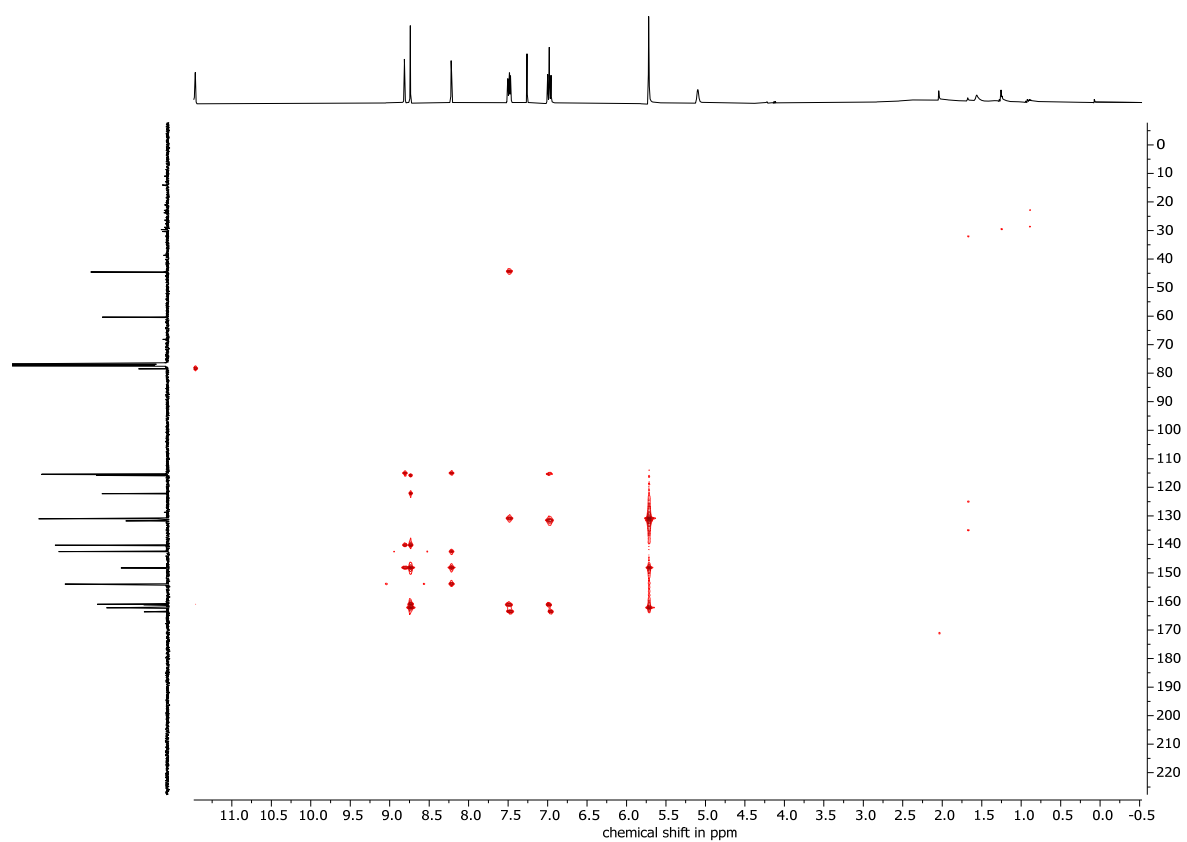


Figure S9. HMBC (^1H , ^{13}C) NMR spectrum of compound **2o** in CDCl_3 .

Compound **2_m**

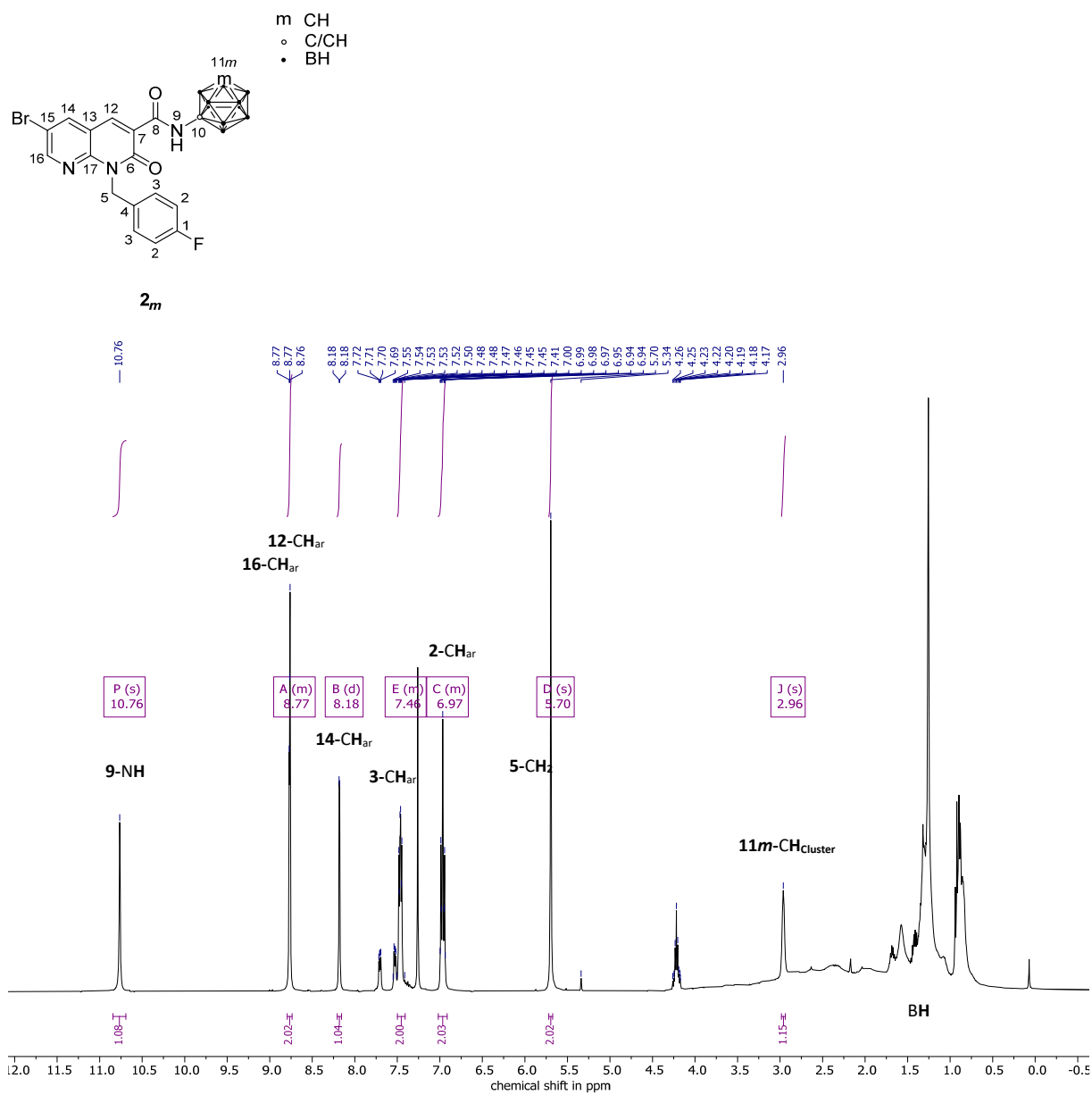


Figure S10. ¹H NMR spectrum of compound **2_m** in CDCl₃.

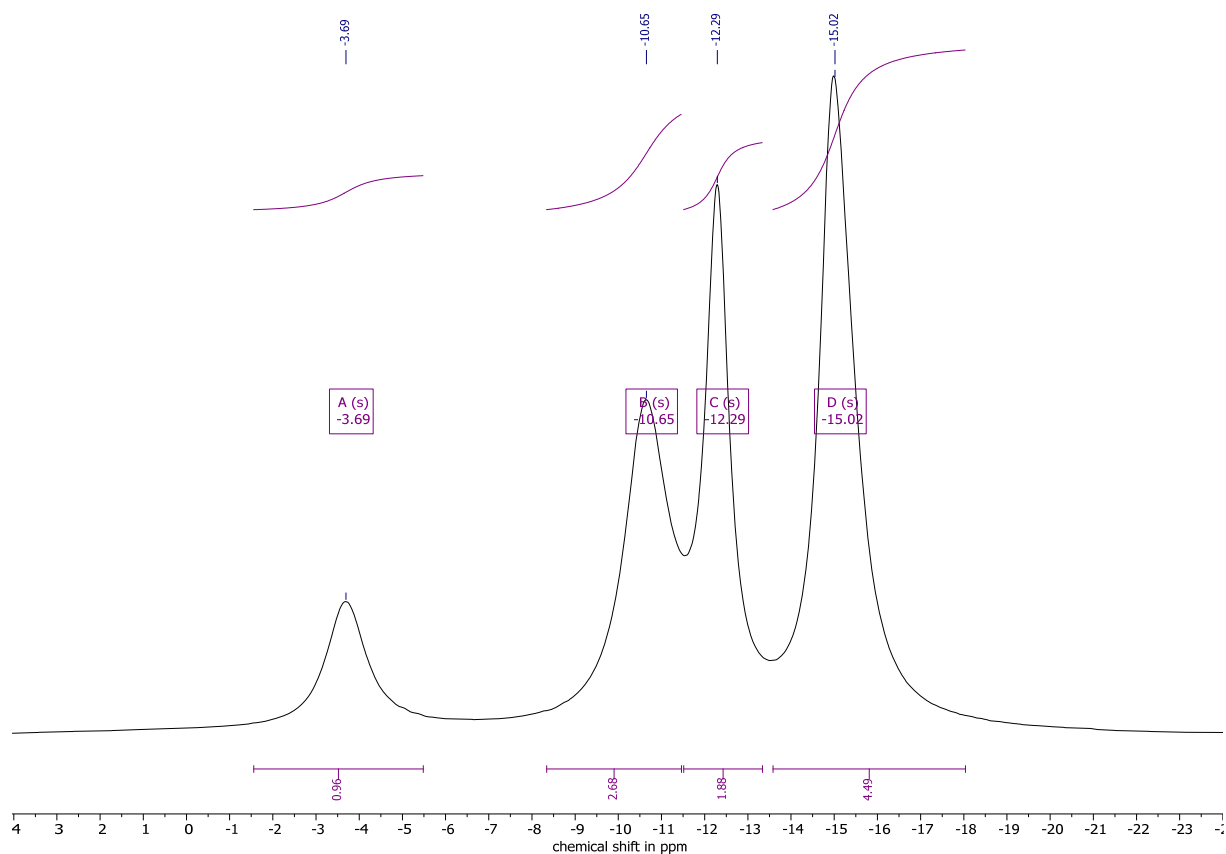


Figure S11. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **2_m** in CDCl_3 .

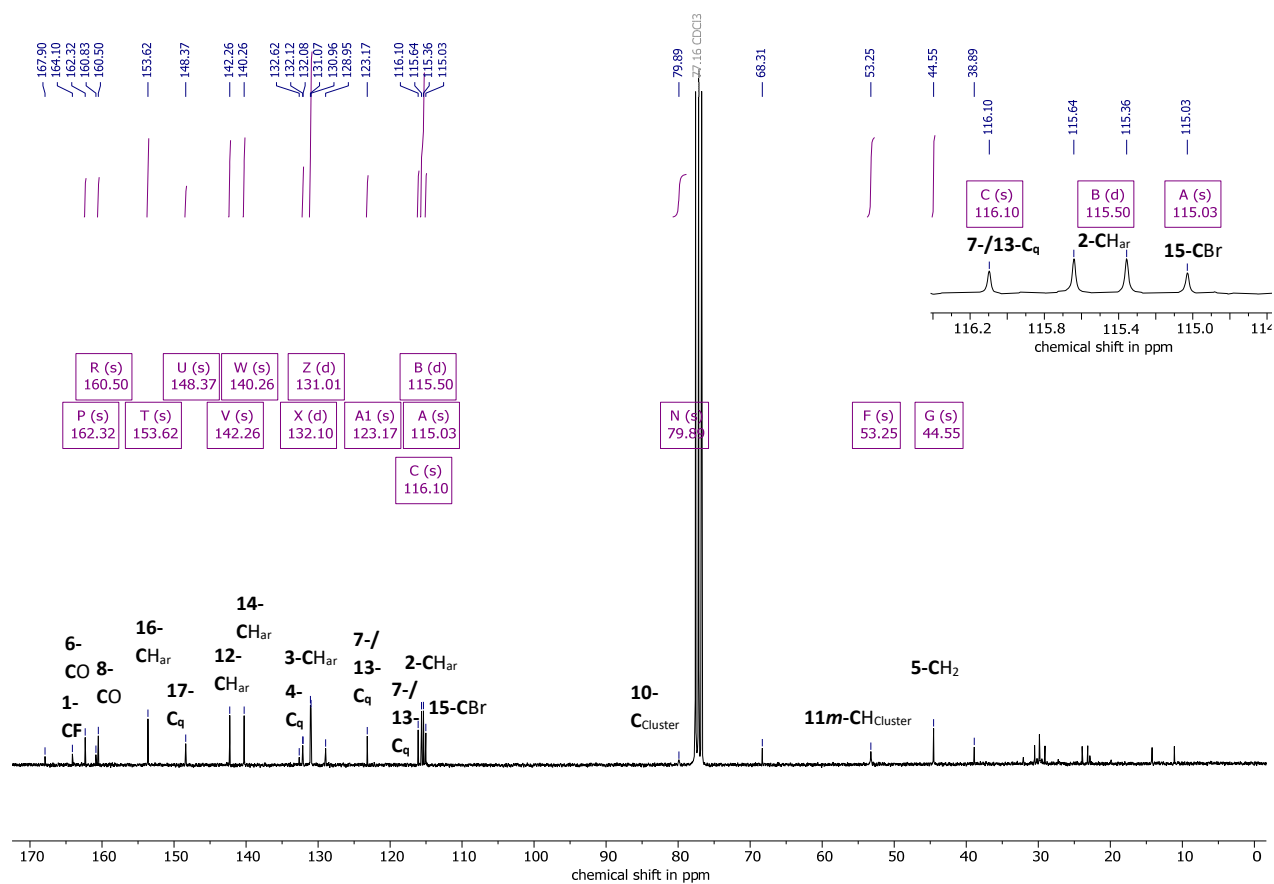


Figure S12. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **2_m** in CDCl_3 .

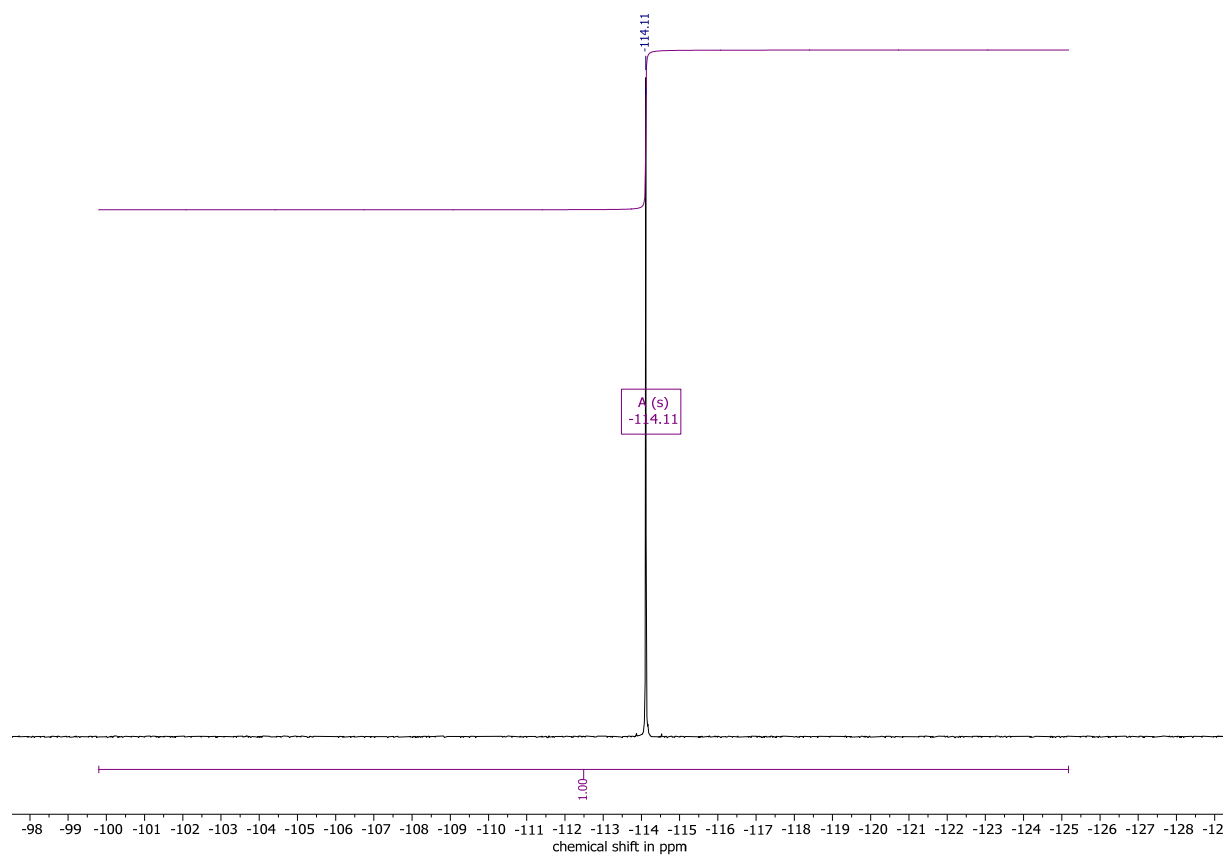


Figure S13. $^{19}\text{F}\{^1\text{H}\}$ NMR spectrum of compound **2_m** in CDCl_3 .

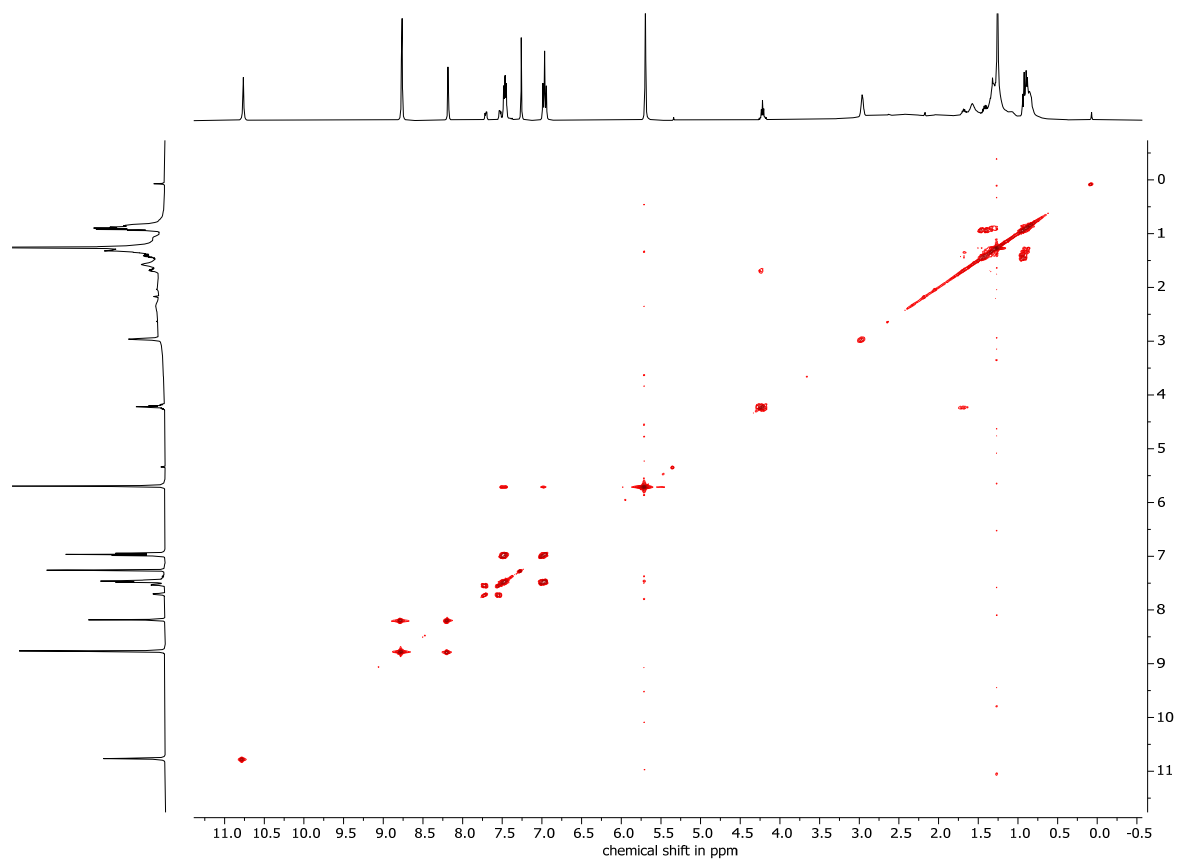


Figure S14. COSY (^1H , ^1H) NMR spectrum of compound **2_m** in CDCl_3 .

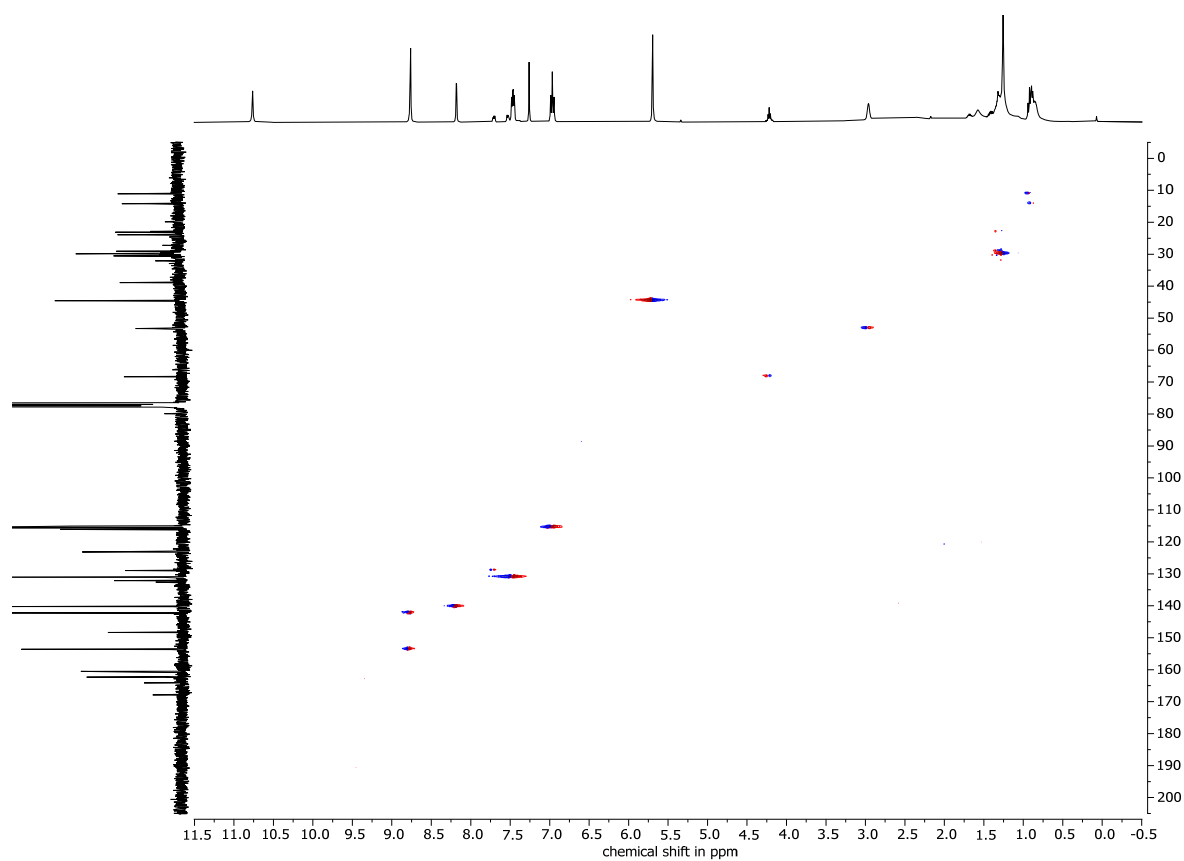


Figure S15. HSQC (¹H, ¹³C) NMR spectrum of compound **2_m** in CDCl₃.

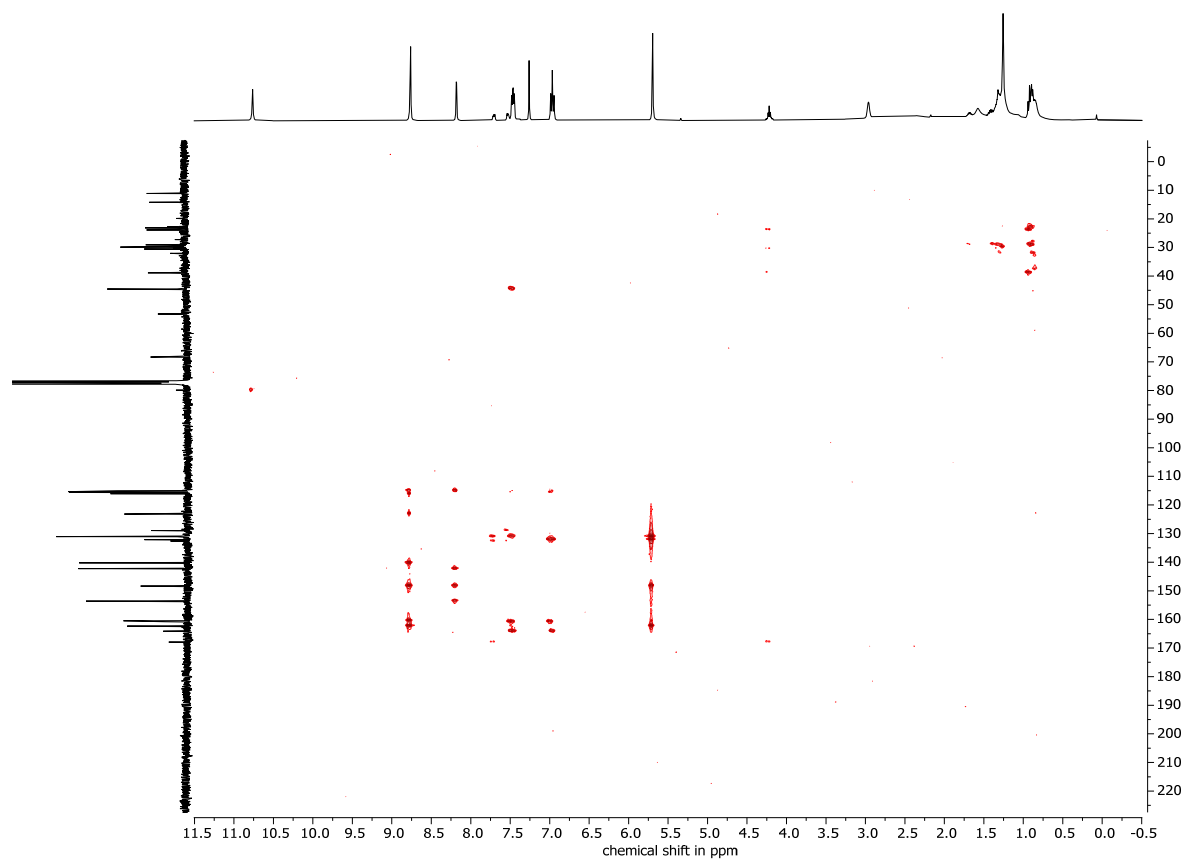


Figure S16. HMBC (¹H, ¹³C) NMR spectrum of compound **2_m** in CDCl₃.

Compound **2_p**

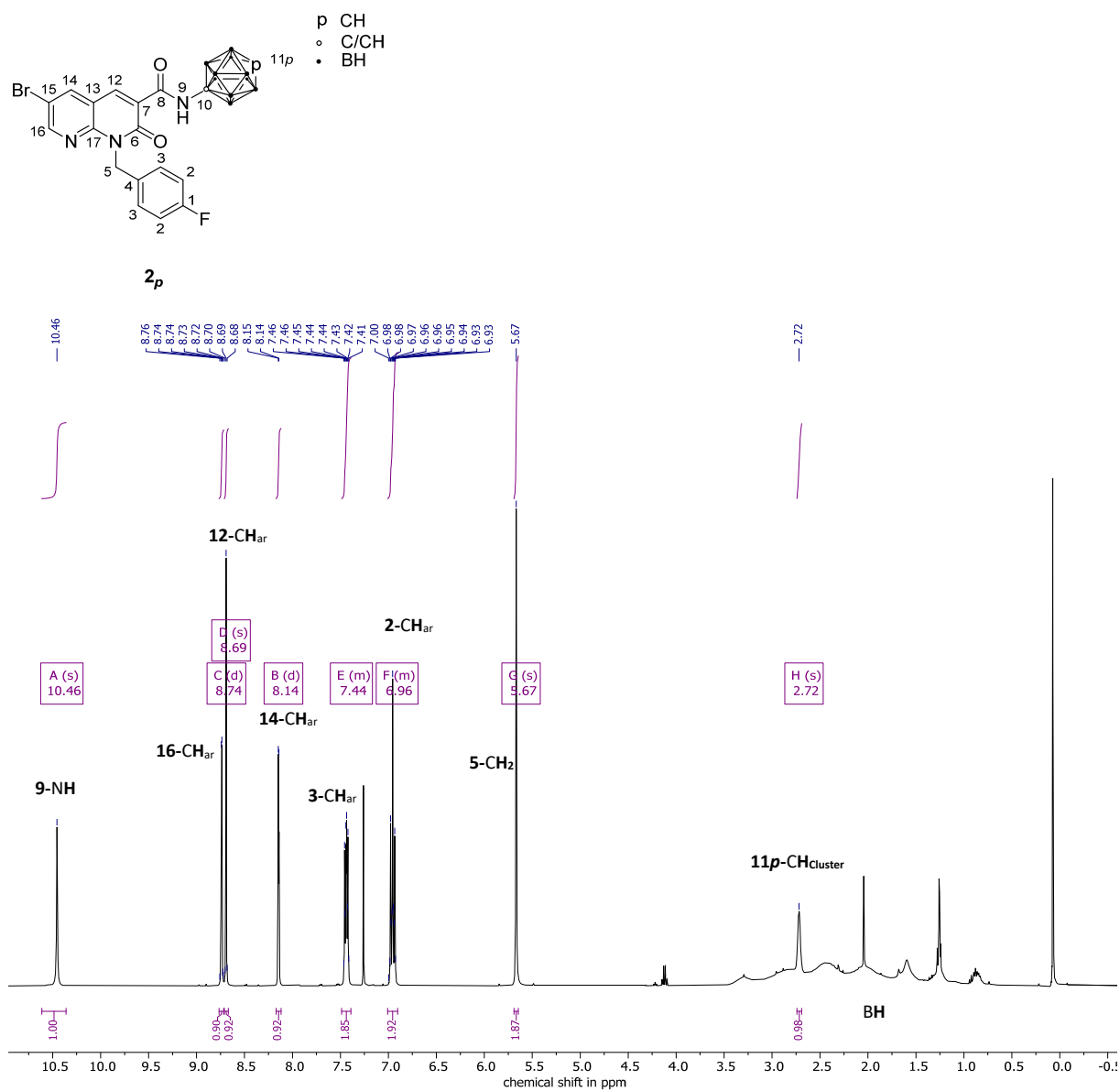


Figure S17. ¹H NMR spectrum of compound **2_p** in CDCl₃.

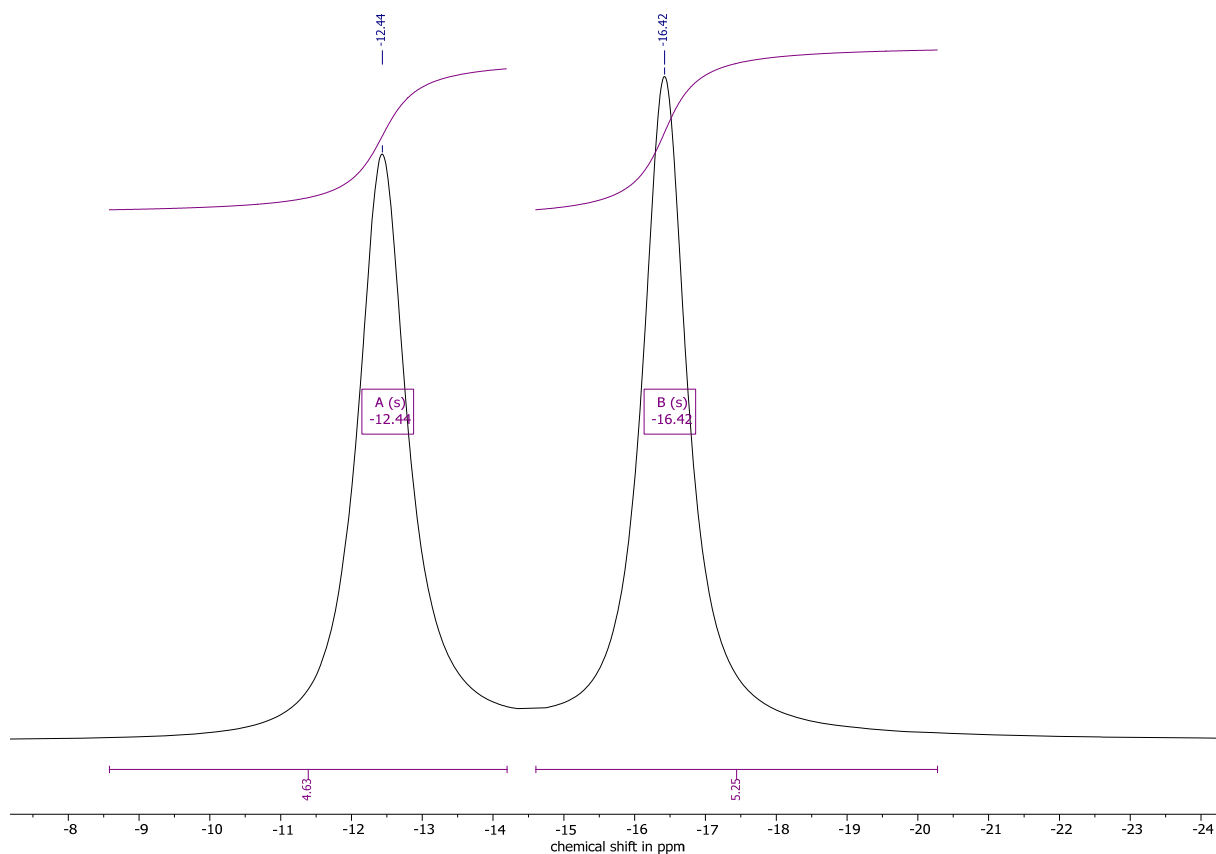


Figure S18. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **2p** in CDCl_3 .

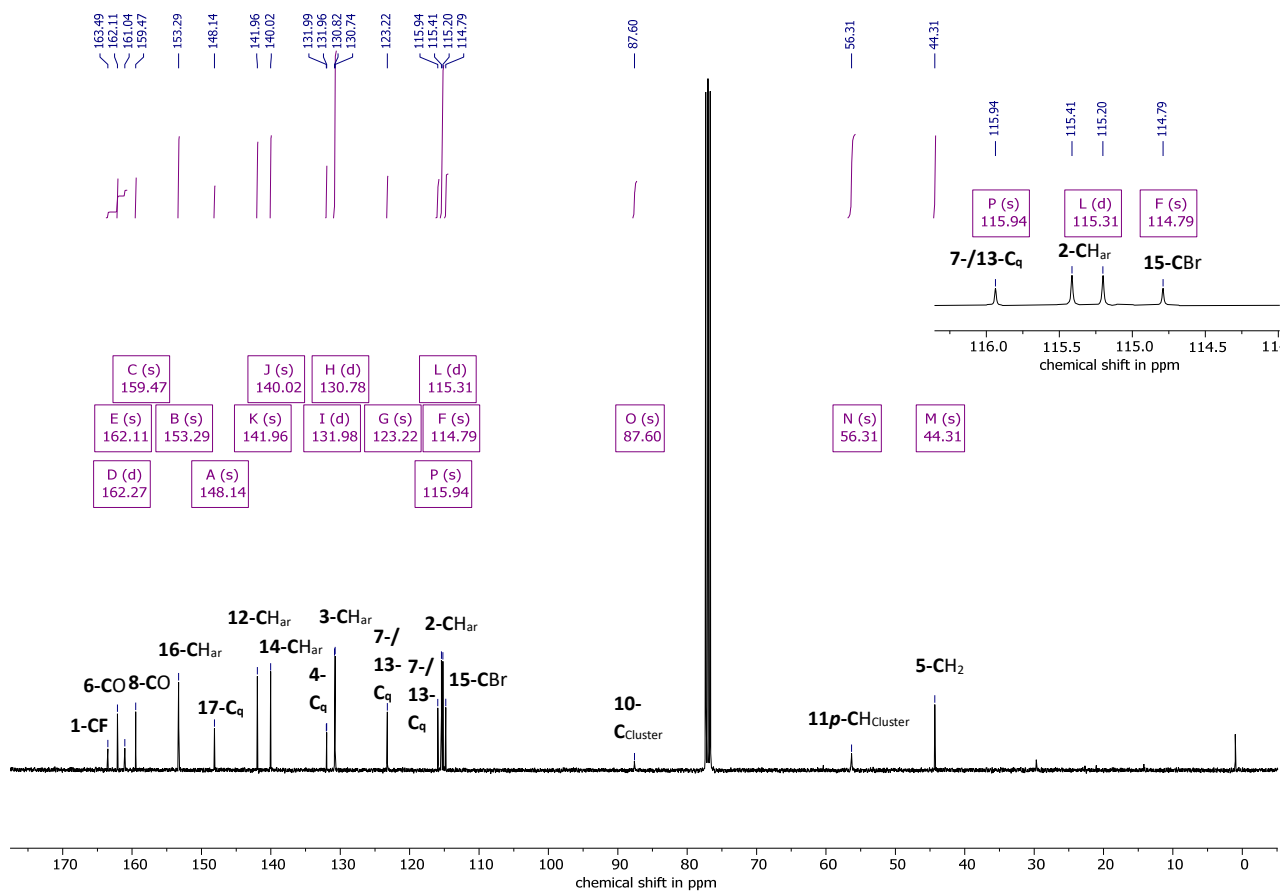


Figure S19. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **2p** in CDCl_3 .

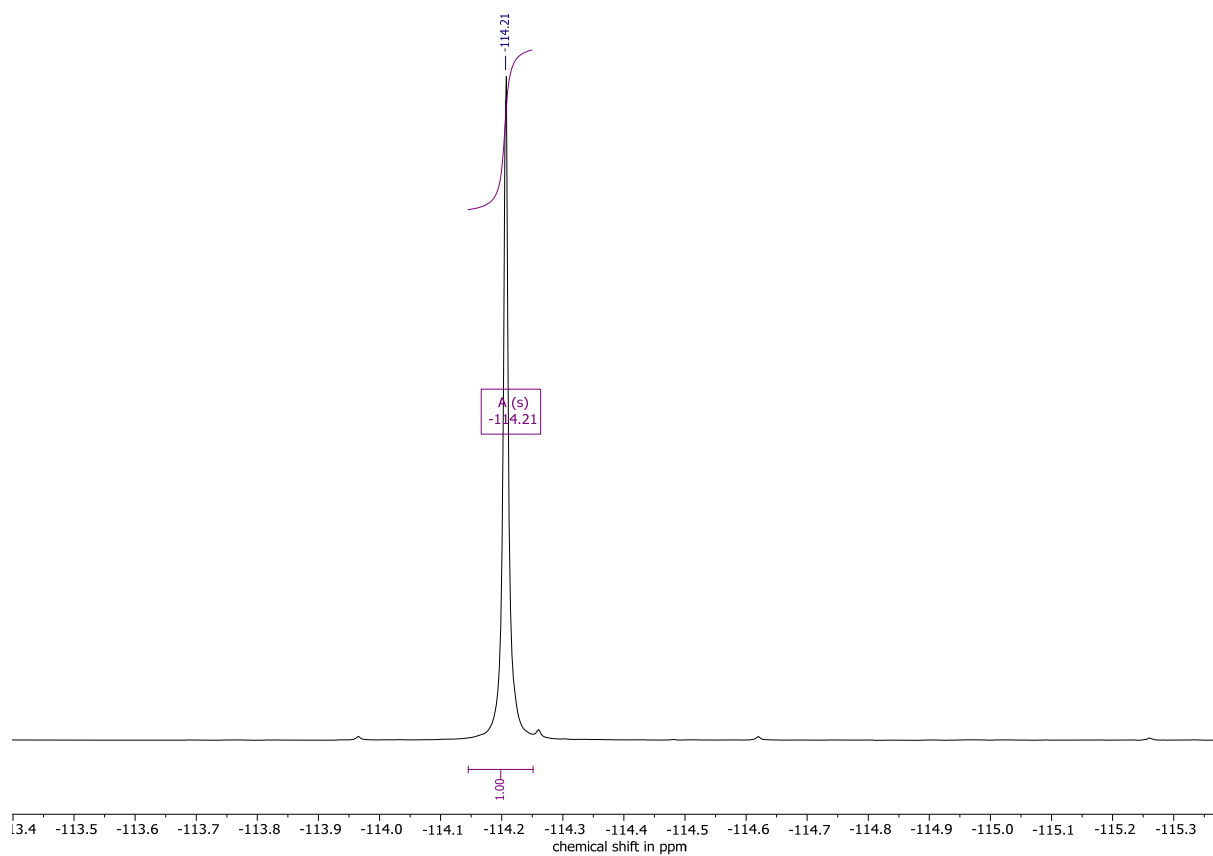


Figure S20. $^{19}\text{F}\{^1\text{H}\}$ NMR spectrum of compound **2p** in CDCl_3 .

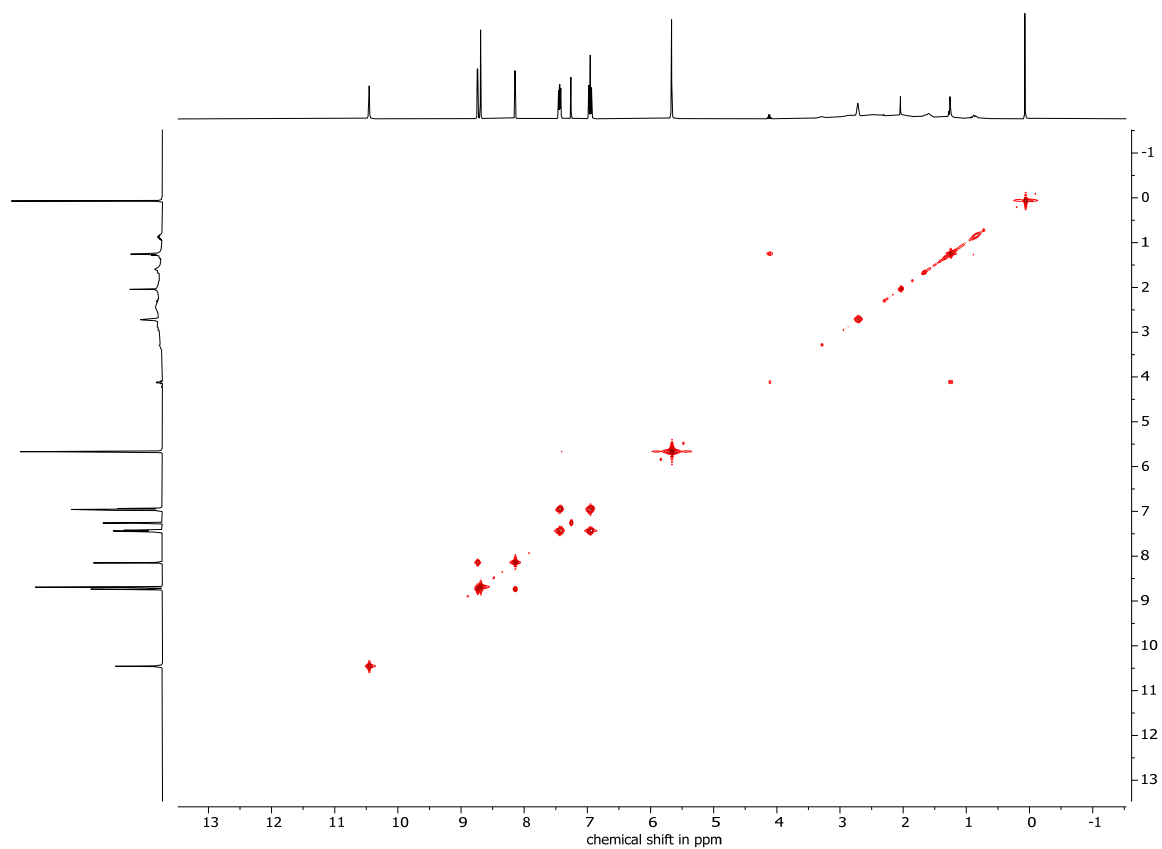


Figure S21. COSY (^1H , ^1H) NMR spectrum of compound **2p** in CDCl_3 .

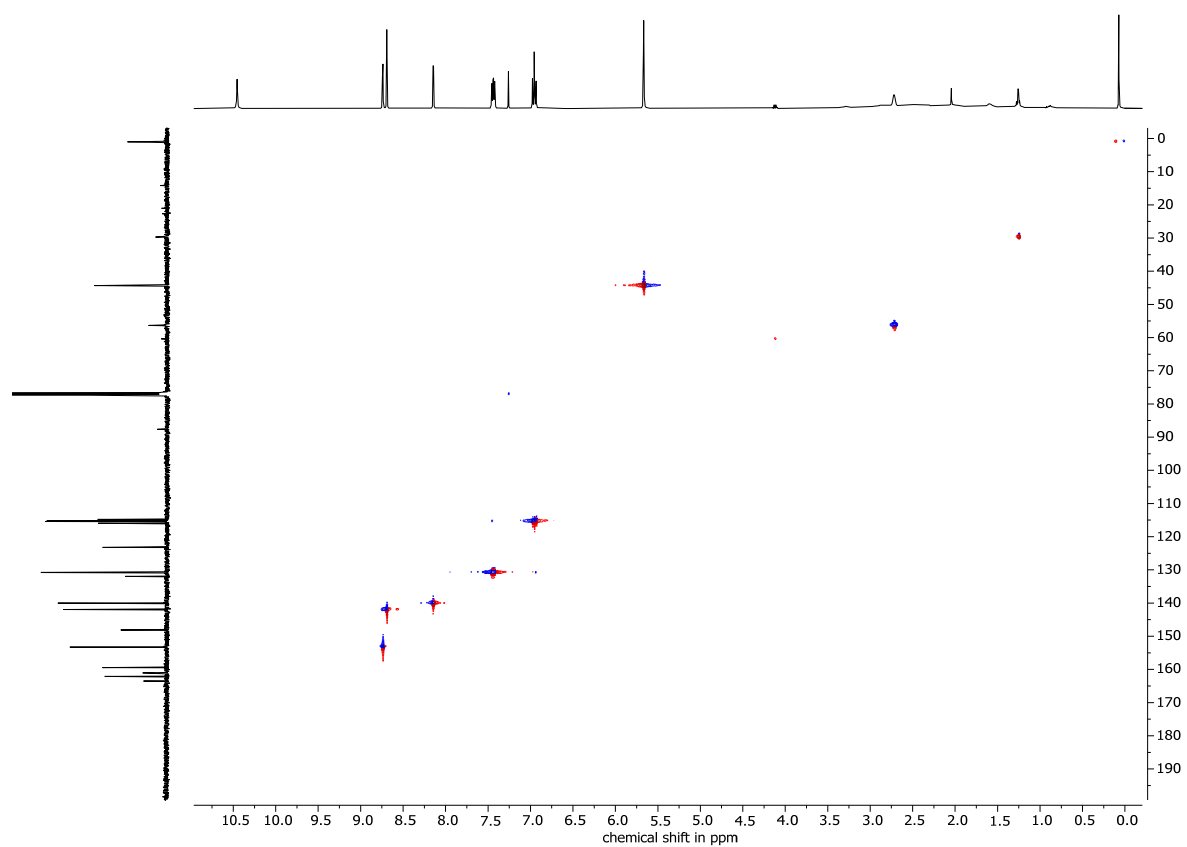


Figure S22. HSQC (^1H , ^{13}C) NMR spectrum of compound **2p** in CDCl_3 .

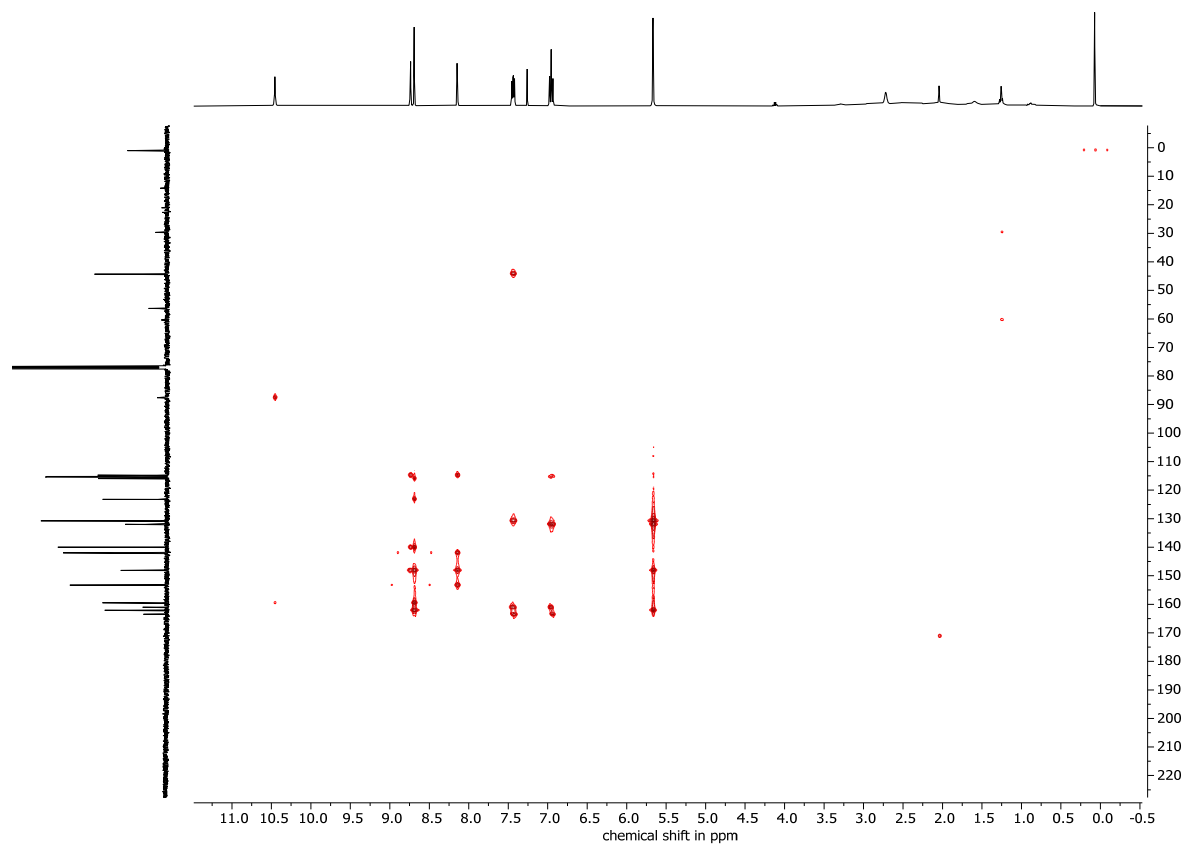
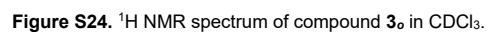


Figure S23. HMBC (^1H , ^{13}C) NMR spectrum of compound **2p** in CDCl_3 .



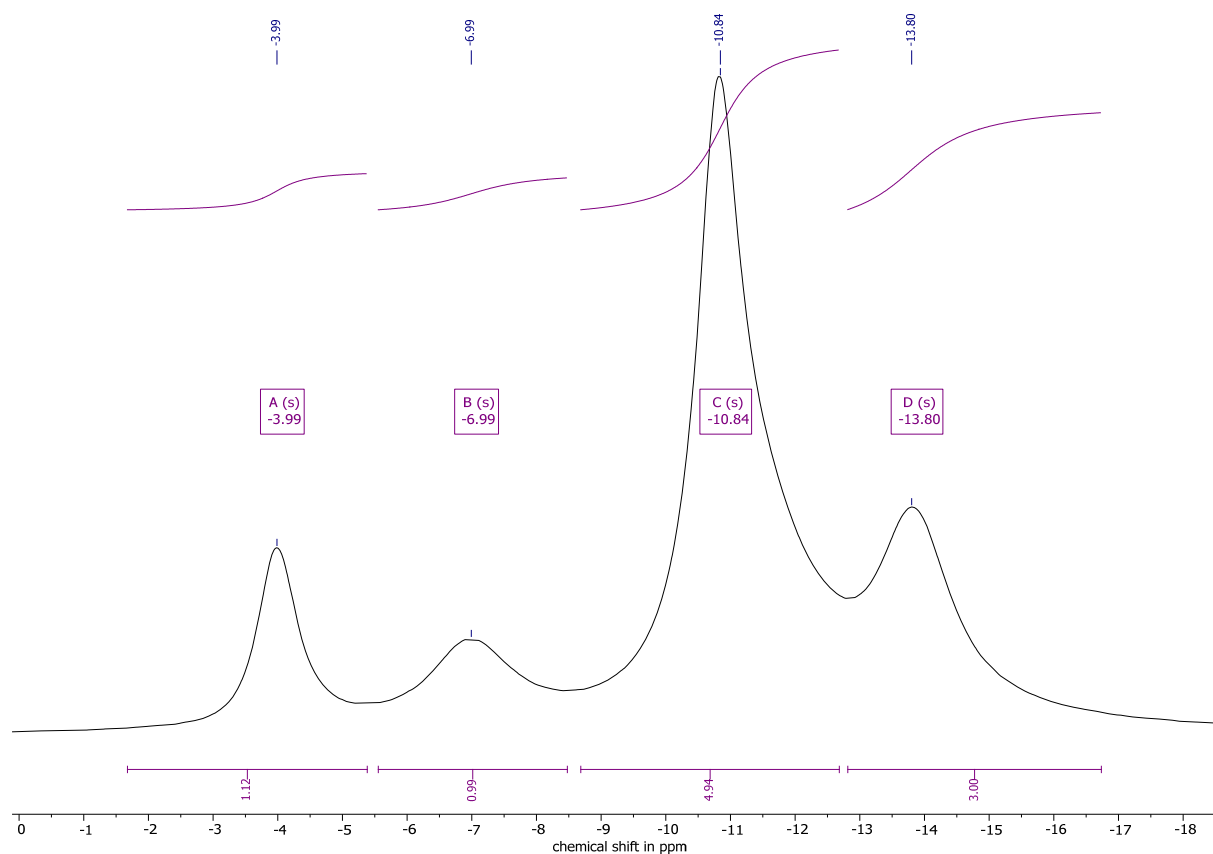


Figure S25. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **3o** in CDCl_3 .

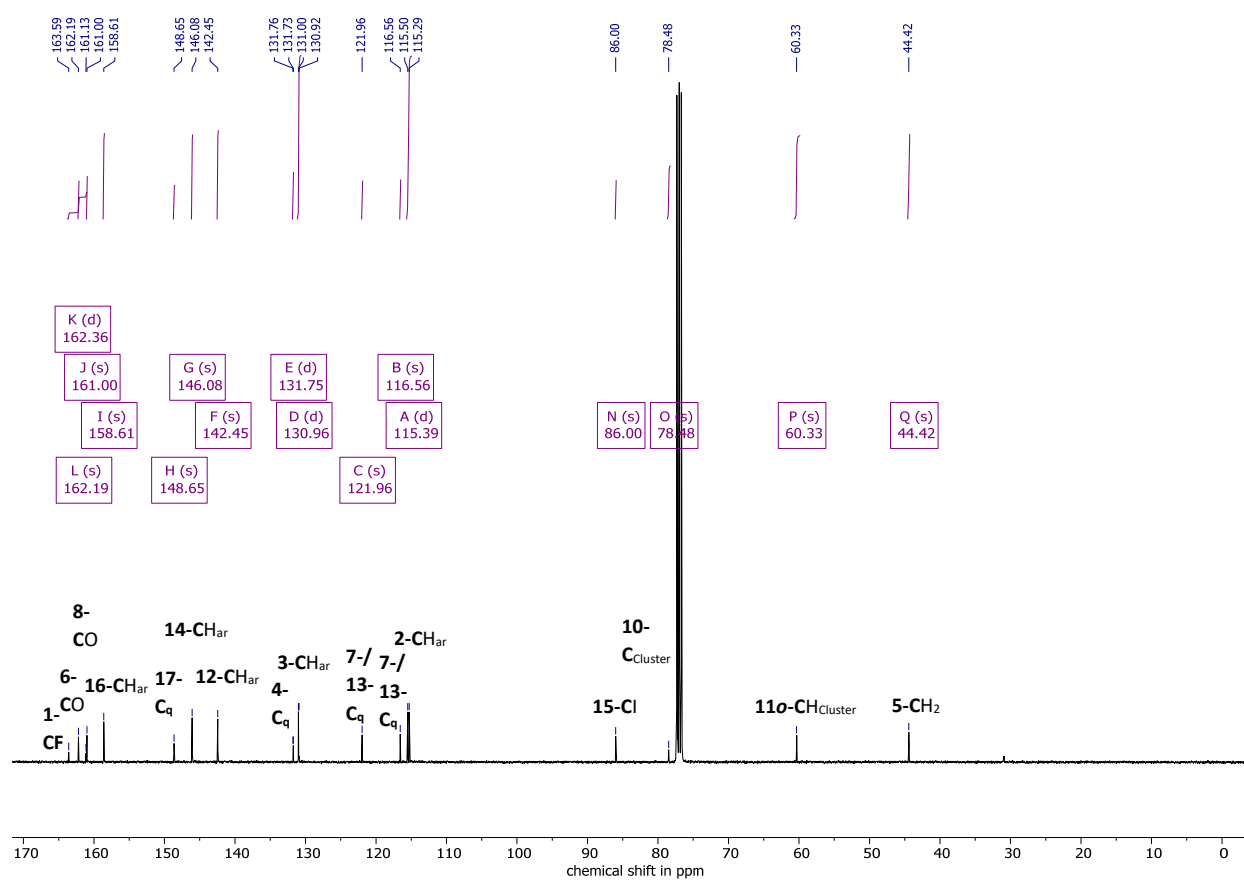


Figure S26. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **3o** in CDCl_3 .

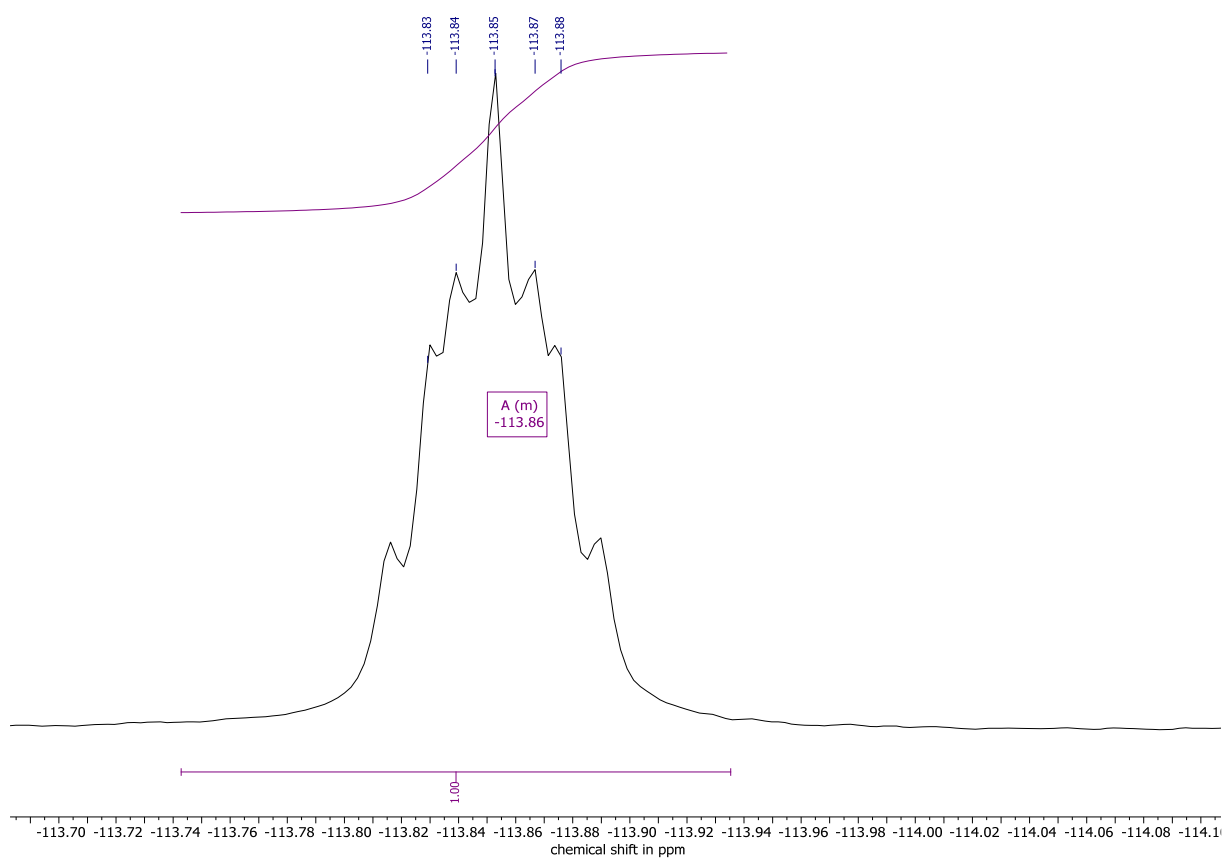


Figure S27. $^{19}\text{F}\{^1\text{H}\}$ NMR spectrum of compound **3o** in CDCl_3 .

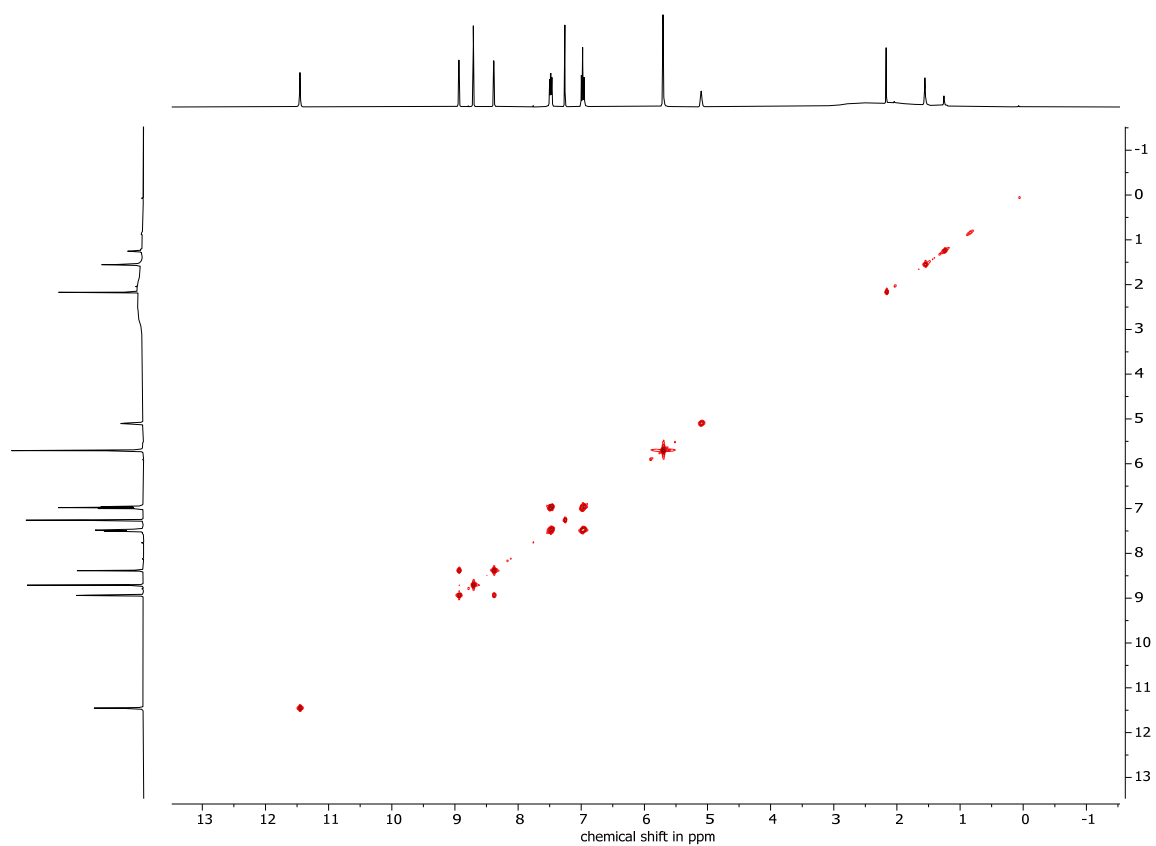


Figure S28. COSY (^1H , ^1H) NMR spectrum of compound **3o** in CDCl_3 .

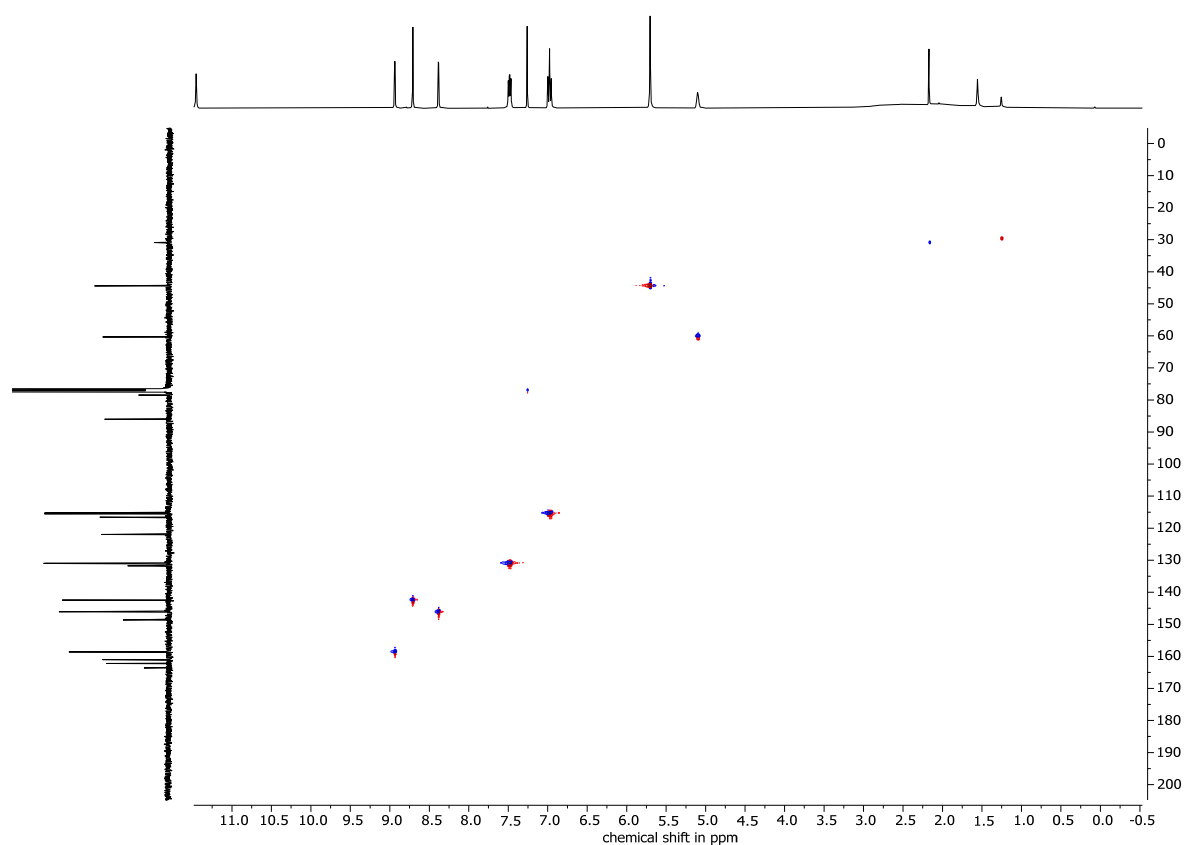


Figure S29. HSQC (¹H, ¹³C) NMR spectrum of compound **3o** in CDCl₃.

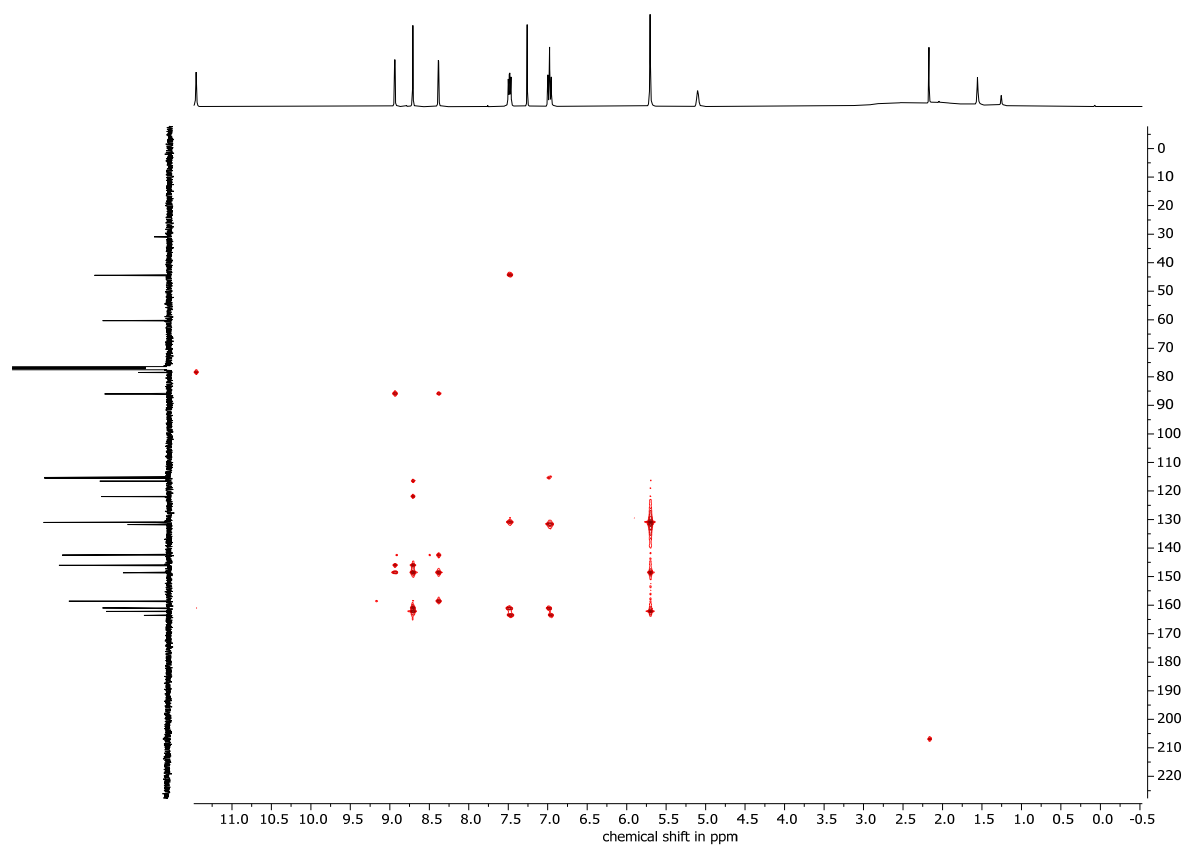


Figure S30. HMBC (¹H, ¹³C) NMR spectrum of compound **3o** in CDCl₃.

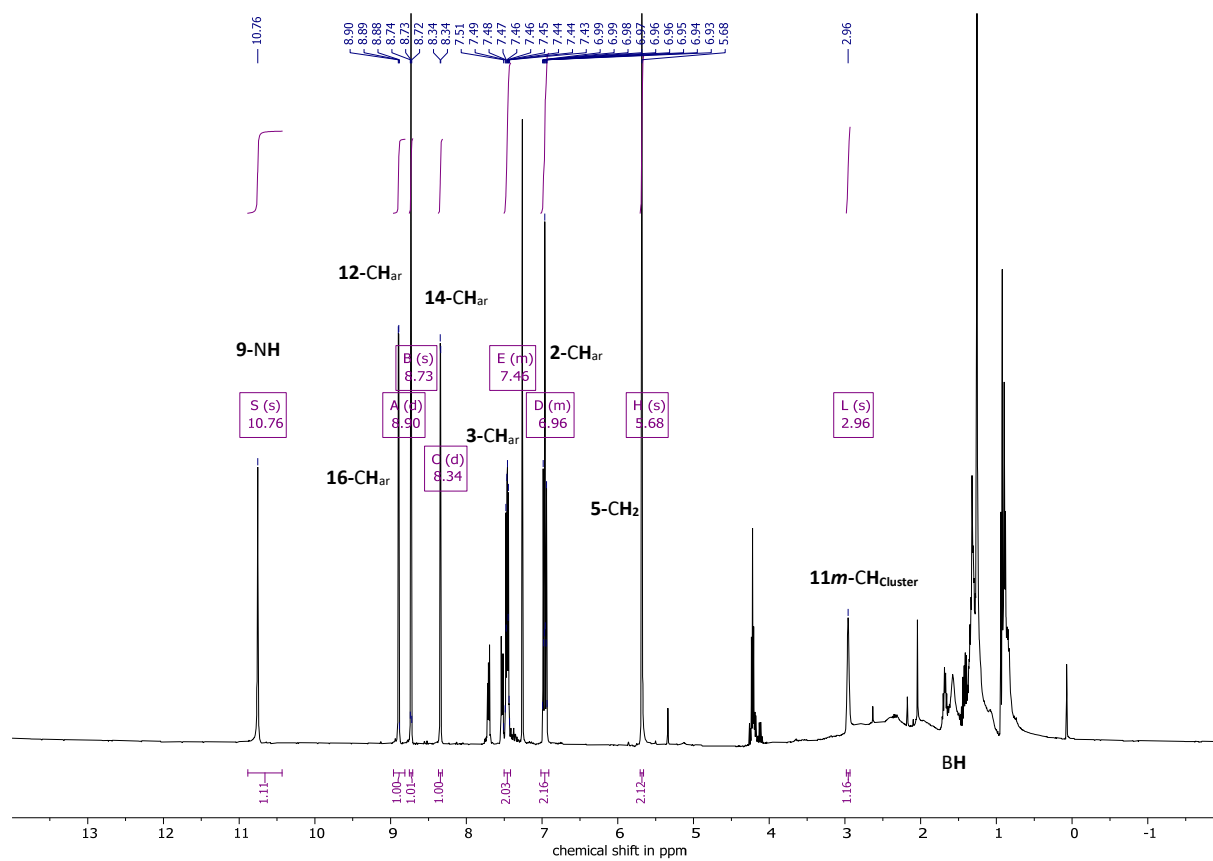


Figure S31. ^1H NMR spectrum of compound **3_m** in CDCl_3 .

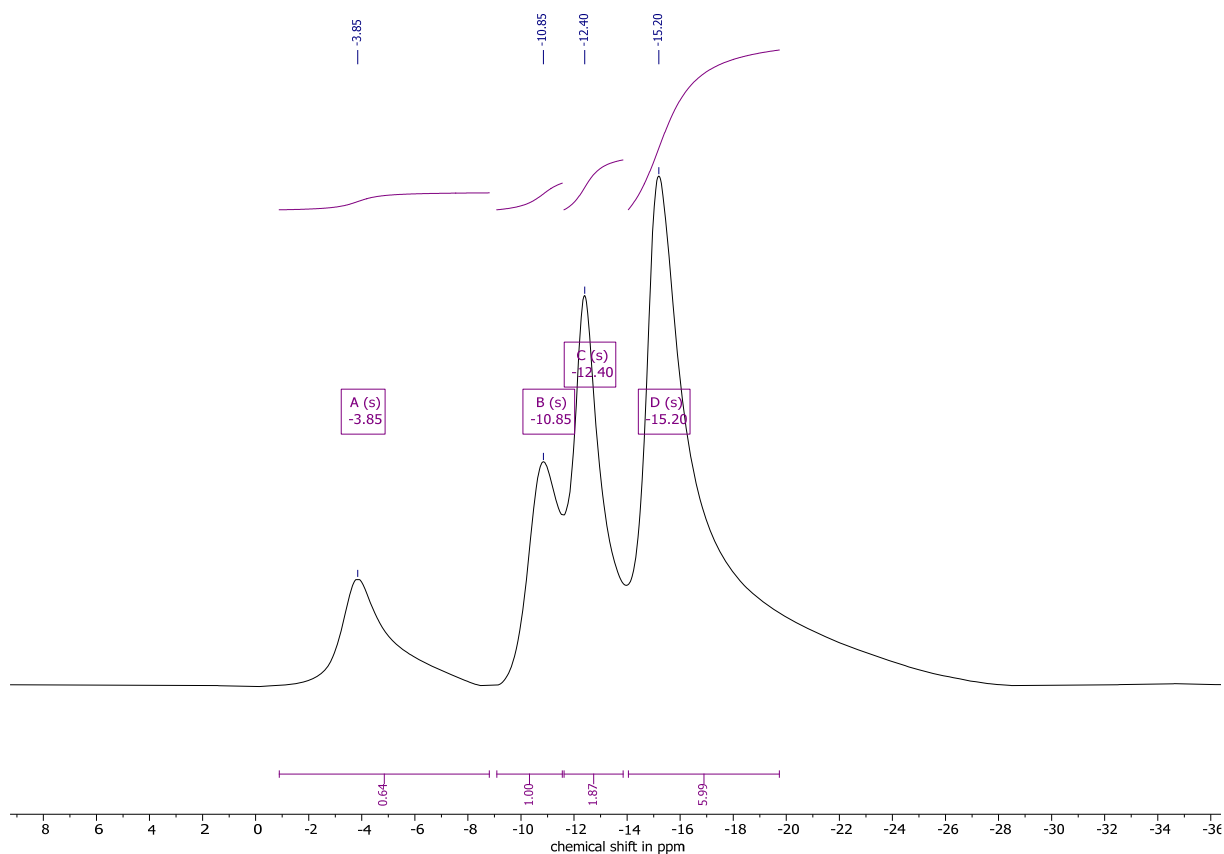


Figure S32. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **3_m** in CDCl_3 .

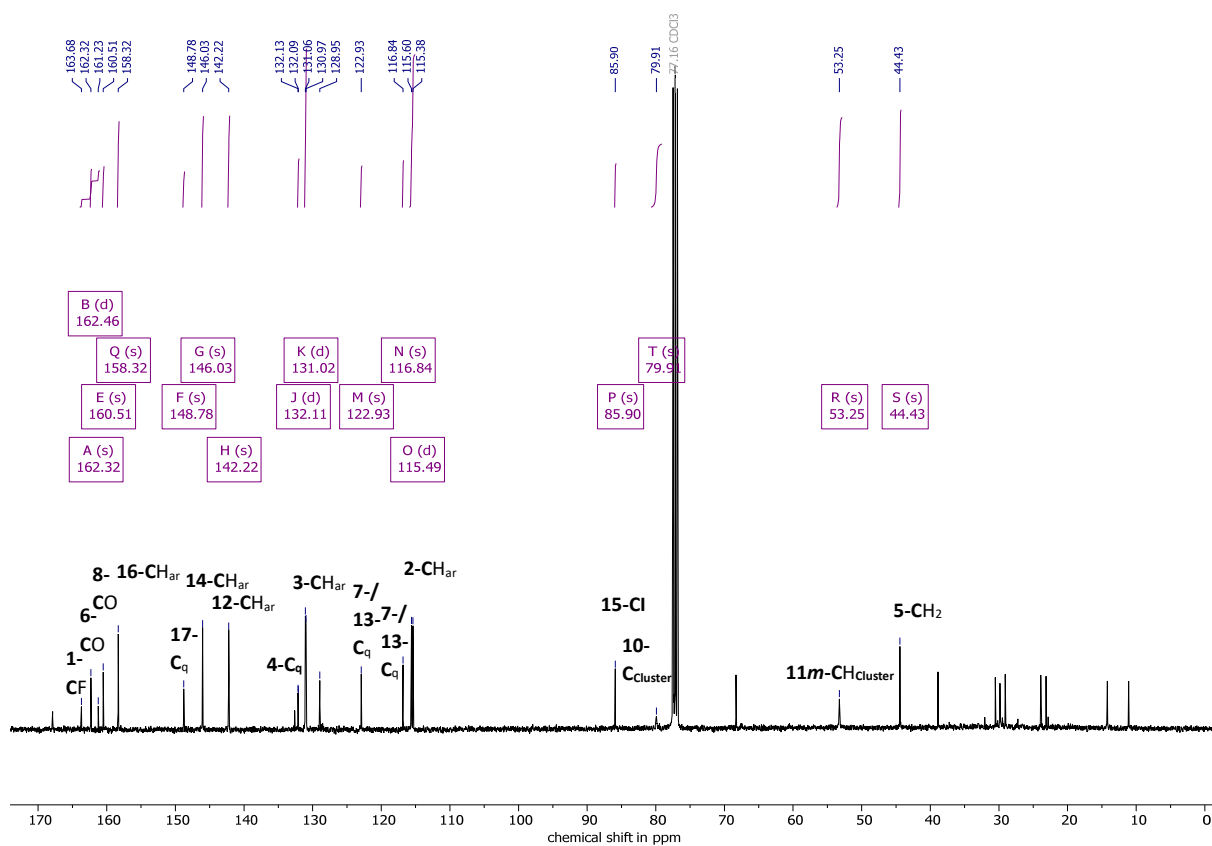


Figure S33. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **3_m** in CDCl_3 .

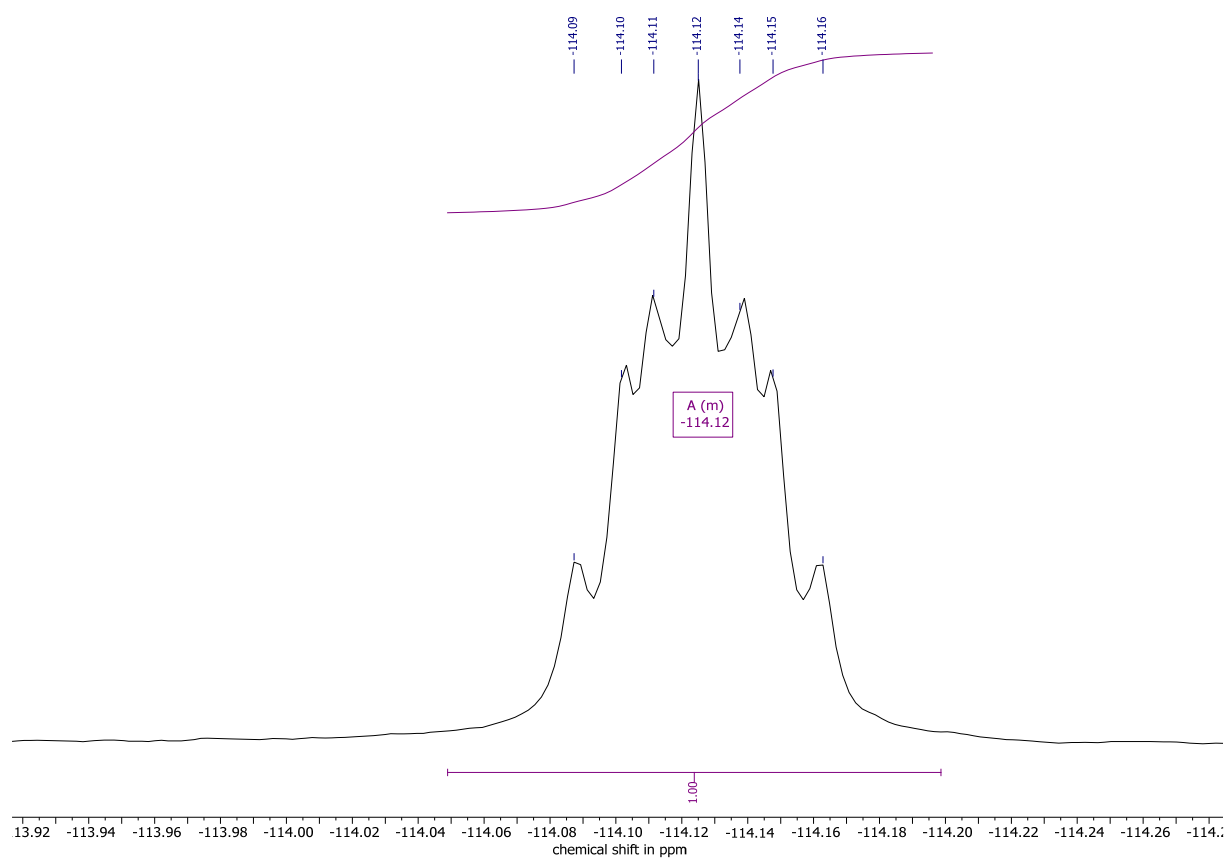


Figure S34. $^{19}\text{F}\{^1\text{H}\}$ NMR spectrum of compound **3_m** in CDCl_3 .

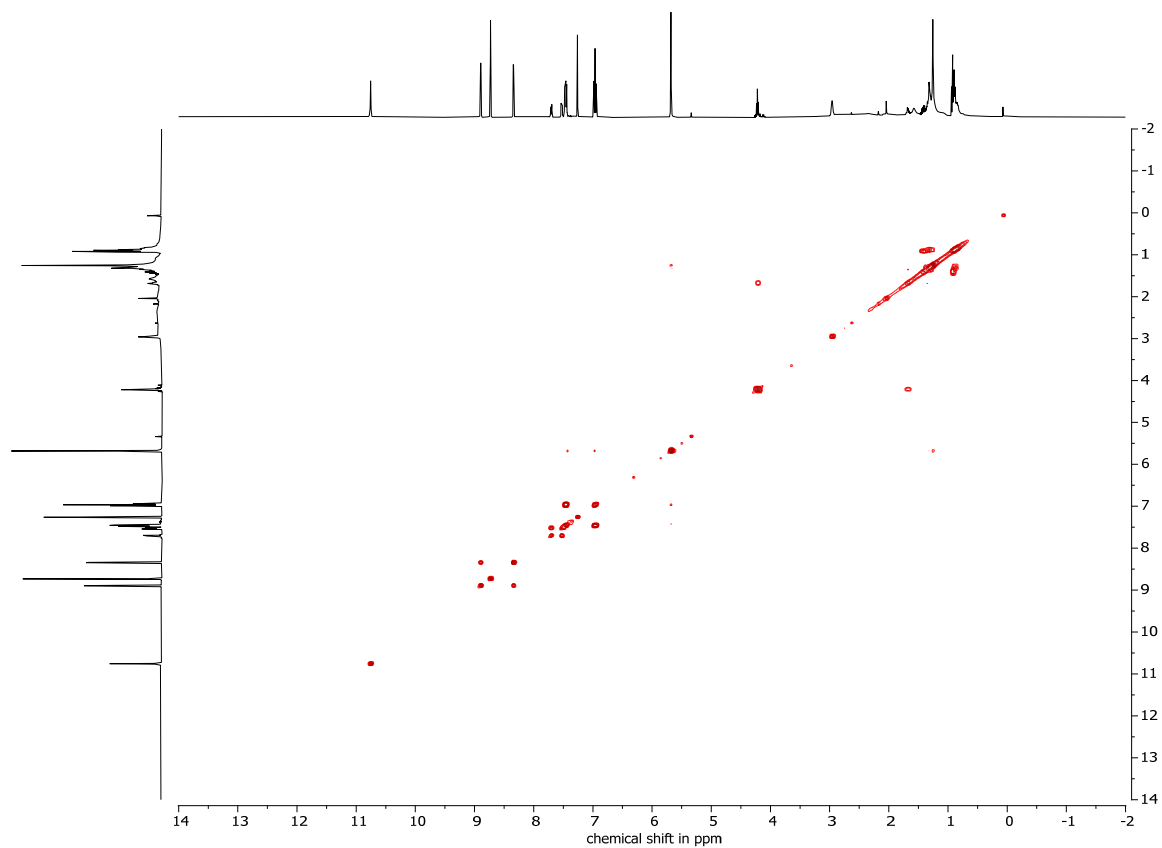


Figure S35. COSY (^1H , ^1H) NMR spectrum of compound **3_m** in CDCl_3 .

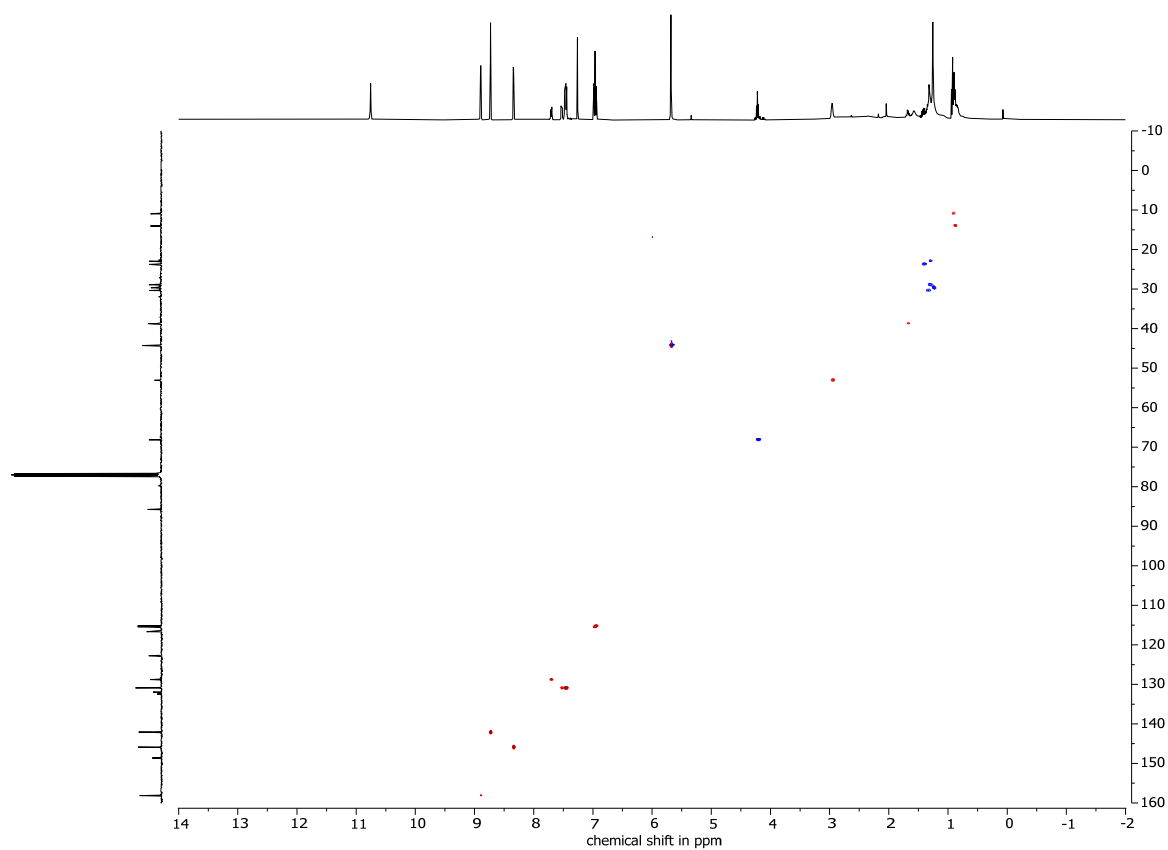


Figure S36. HSQC (^1H , ^{13}C) NMR spectrum of compound **3_m** in CDCl_3 .

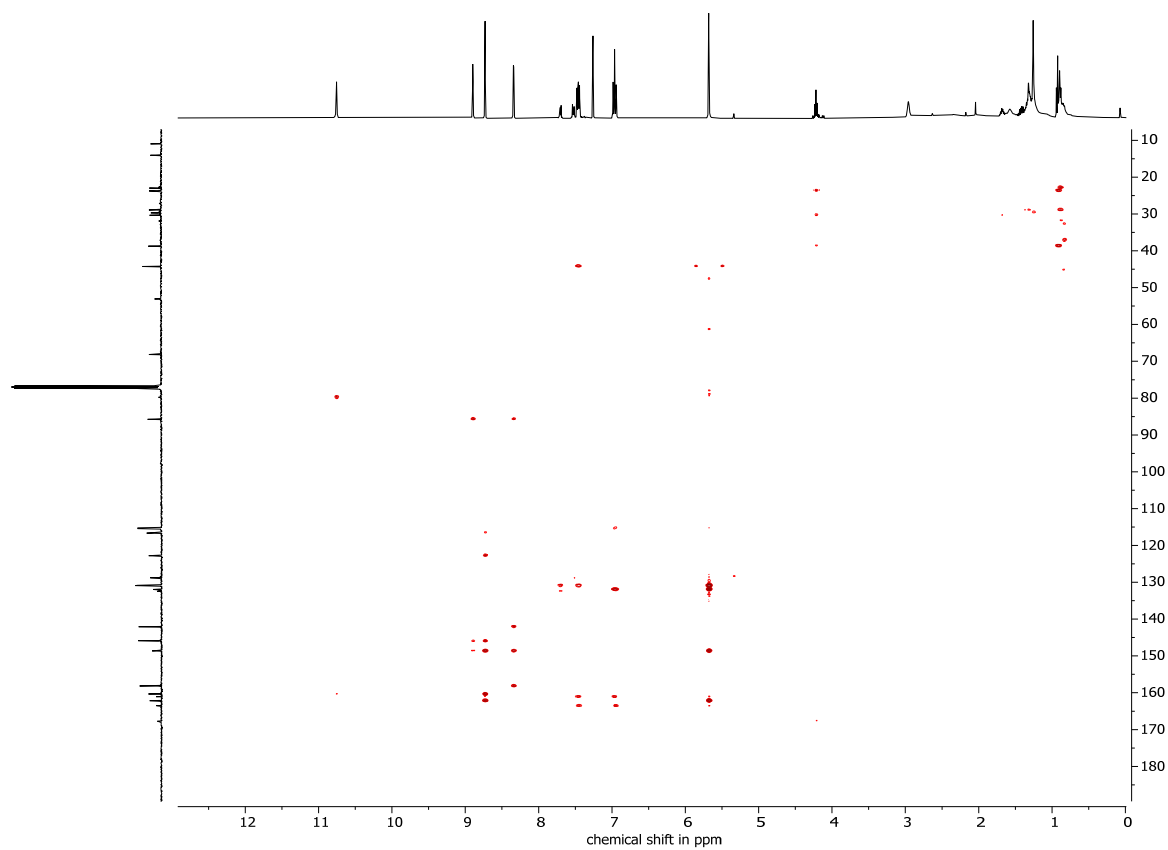
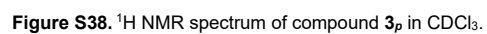


Figure S37. HMBC (^1H , ^{13}C) NMR spectrum of compound **3_m** in CDCl_3 .



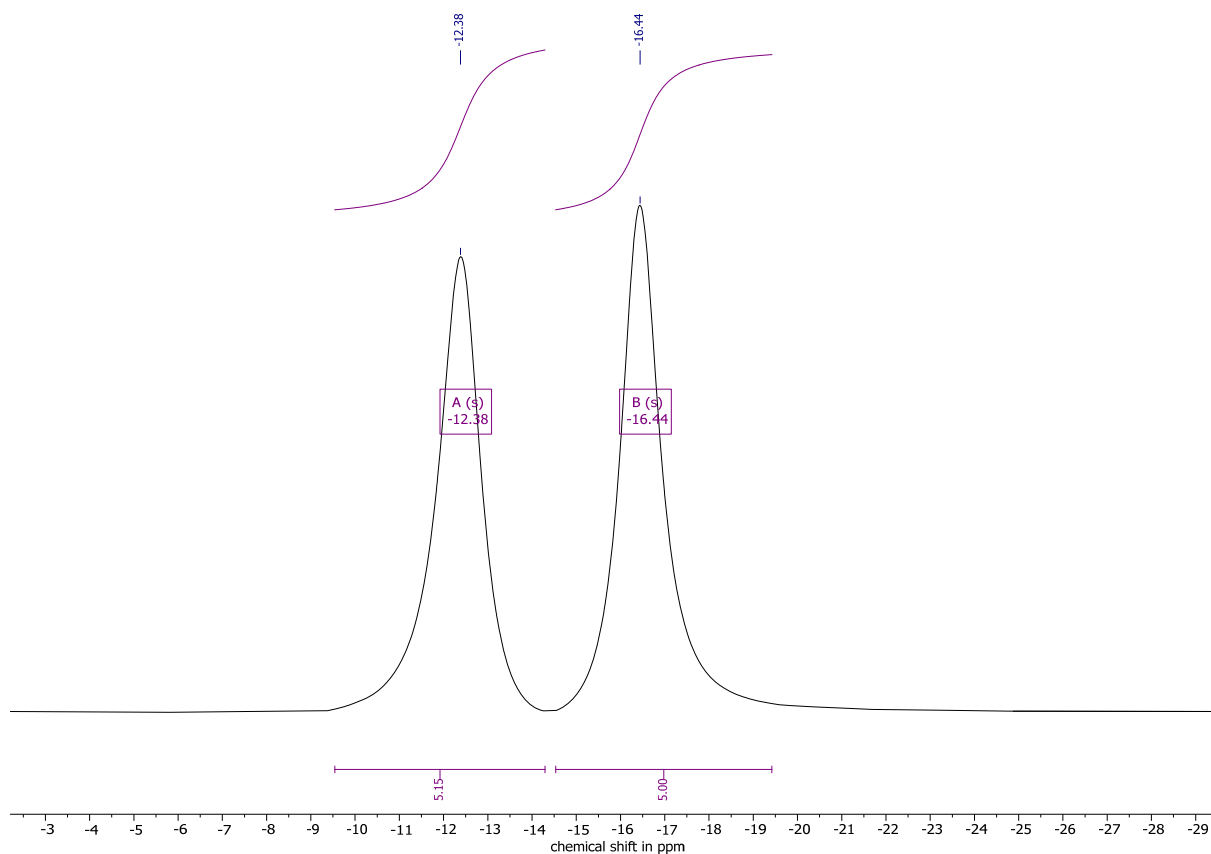


Figure S39. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **3p** in CDCl_3 .

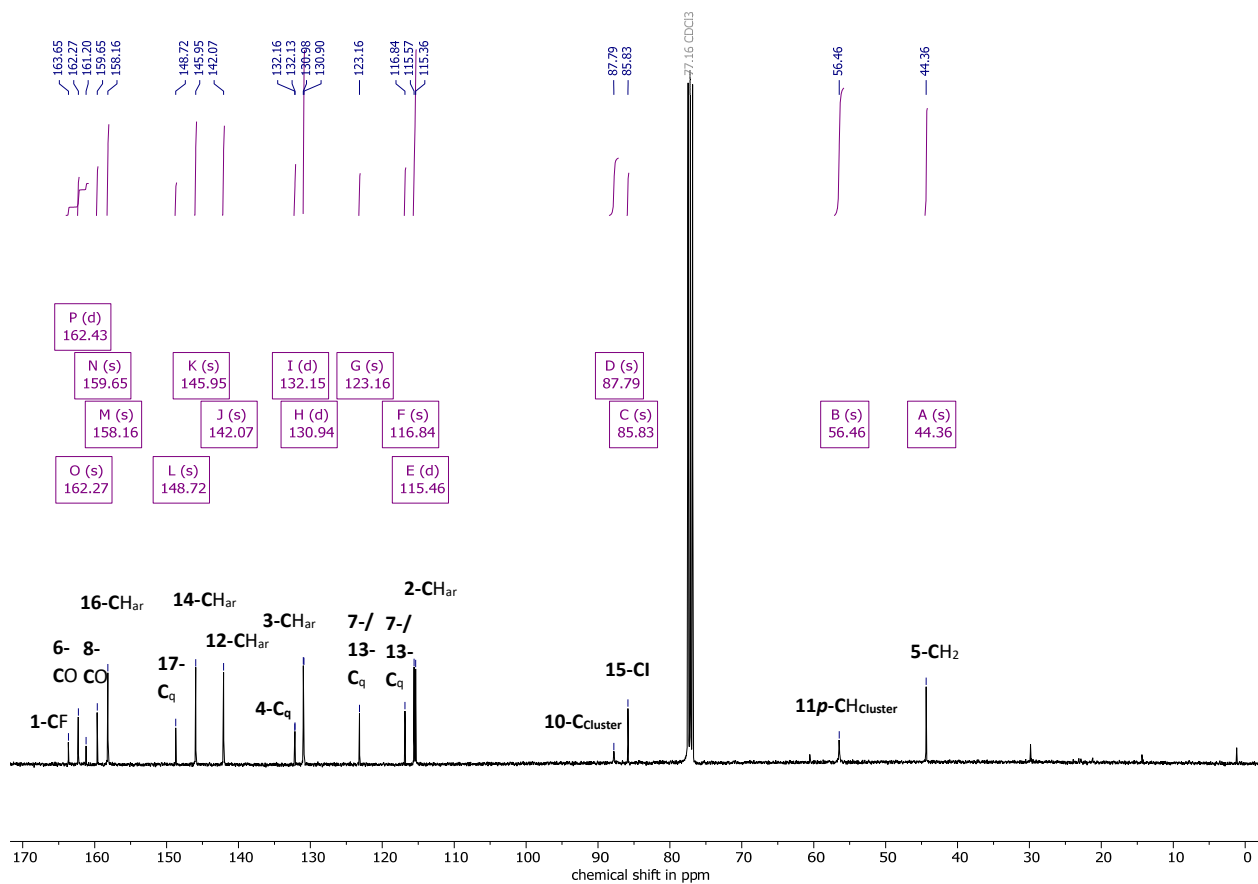


Figure S40. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **3p** in CDCl_3 .

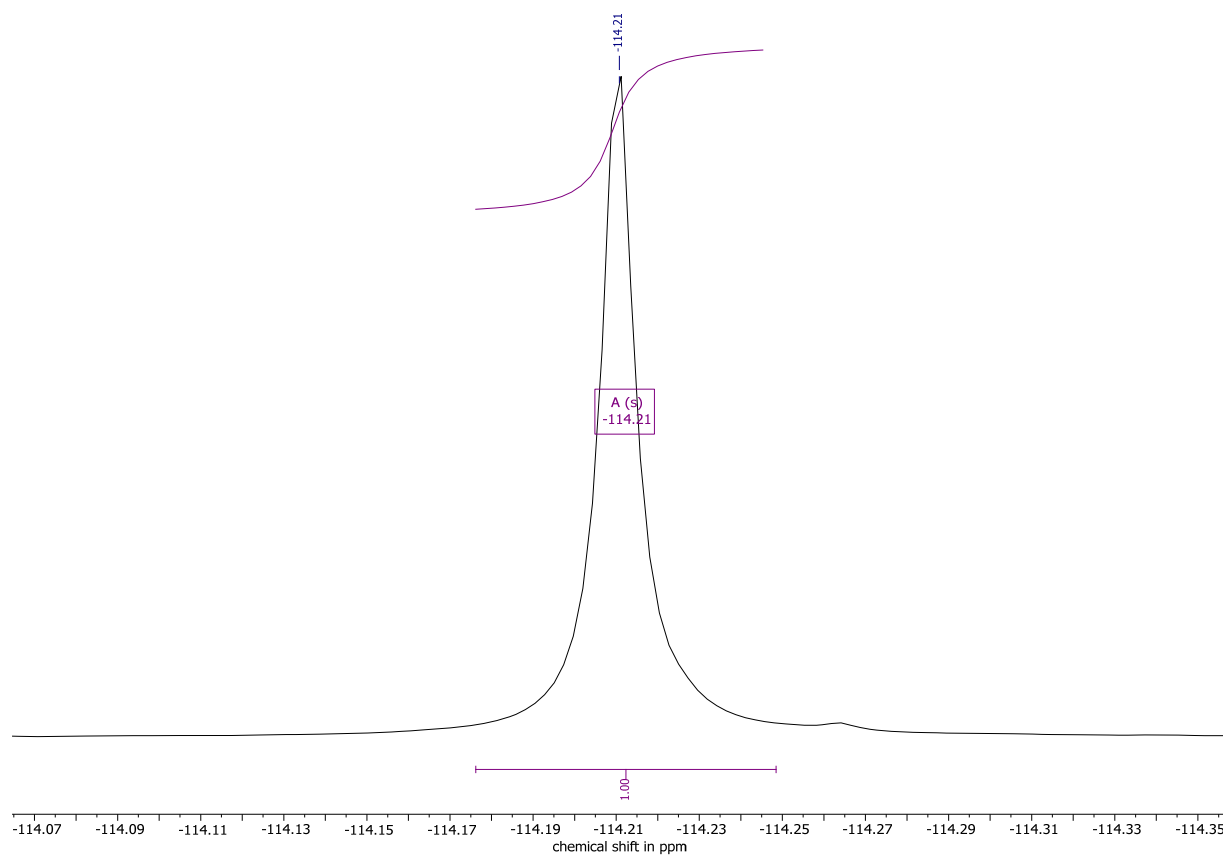


Figure S41. $^{19}\text{F}\{^1\text{H}\}$ NMR spectrum of compound **3p** in CDCl_3 .

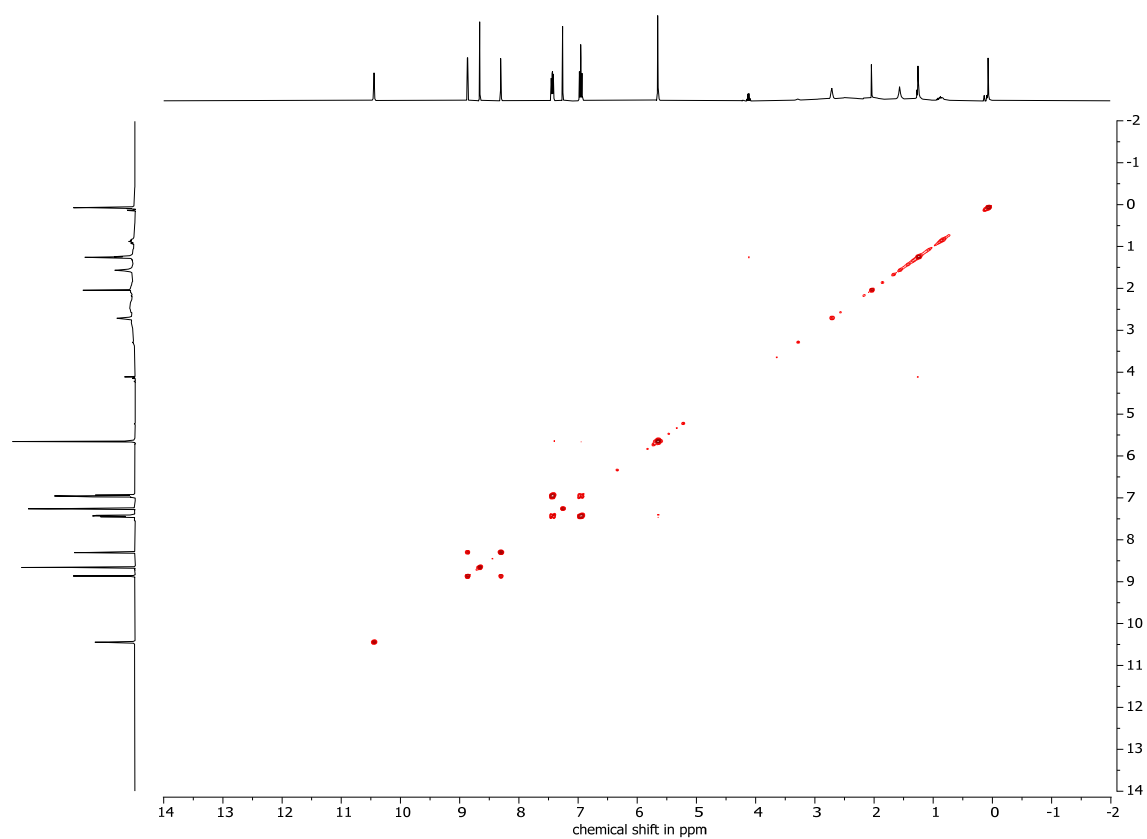


Figure S42. COSY (^1H , ^1H) NMR spectrum of compound **3p** in CDCl_3 .

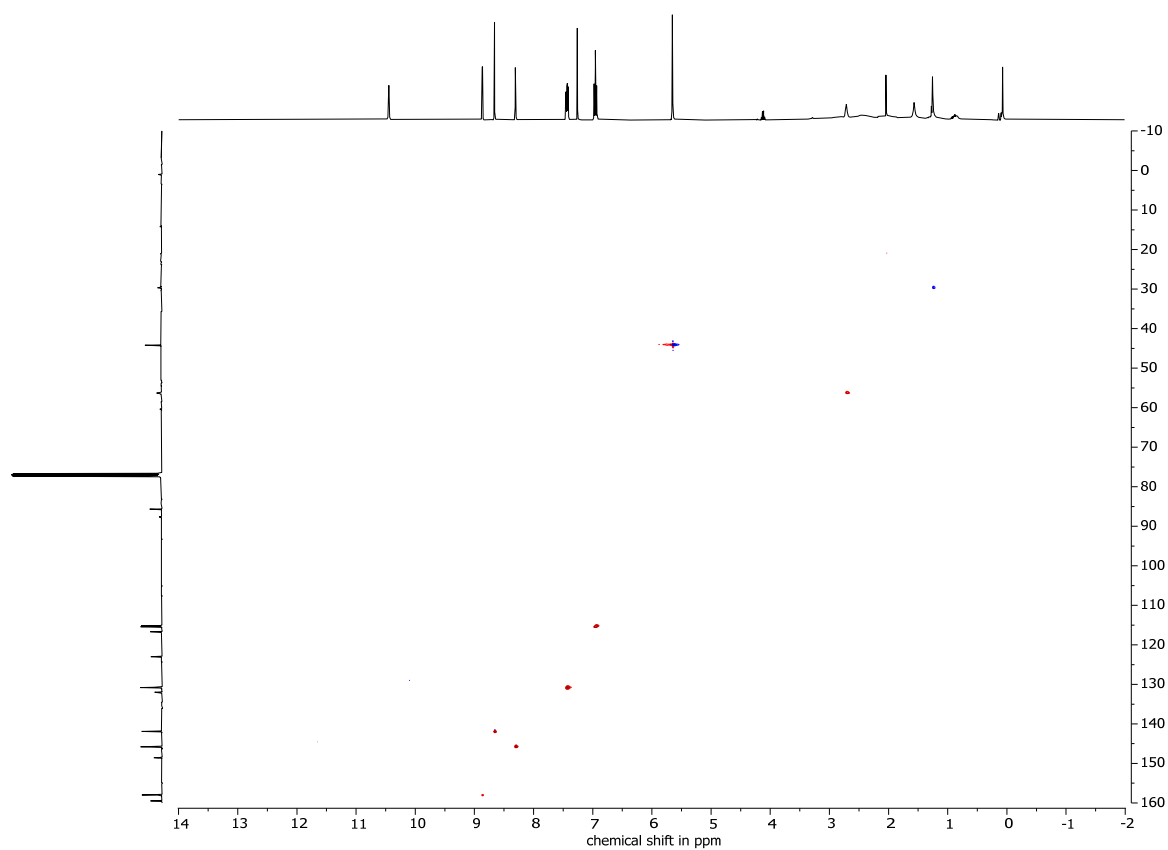


Figure S43. HSQC (^1H , ^{13}C) NMR spectrum of compound **3p** in CDCl_3 .

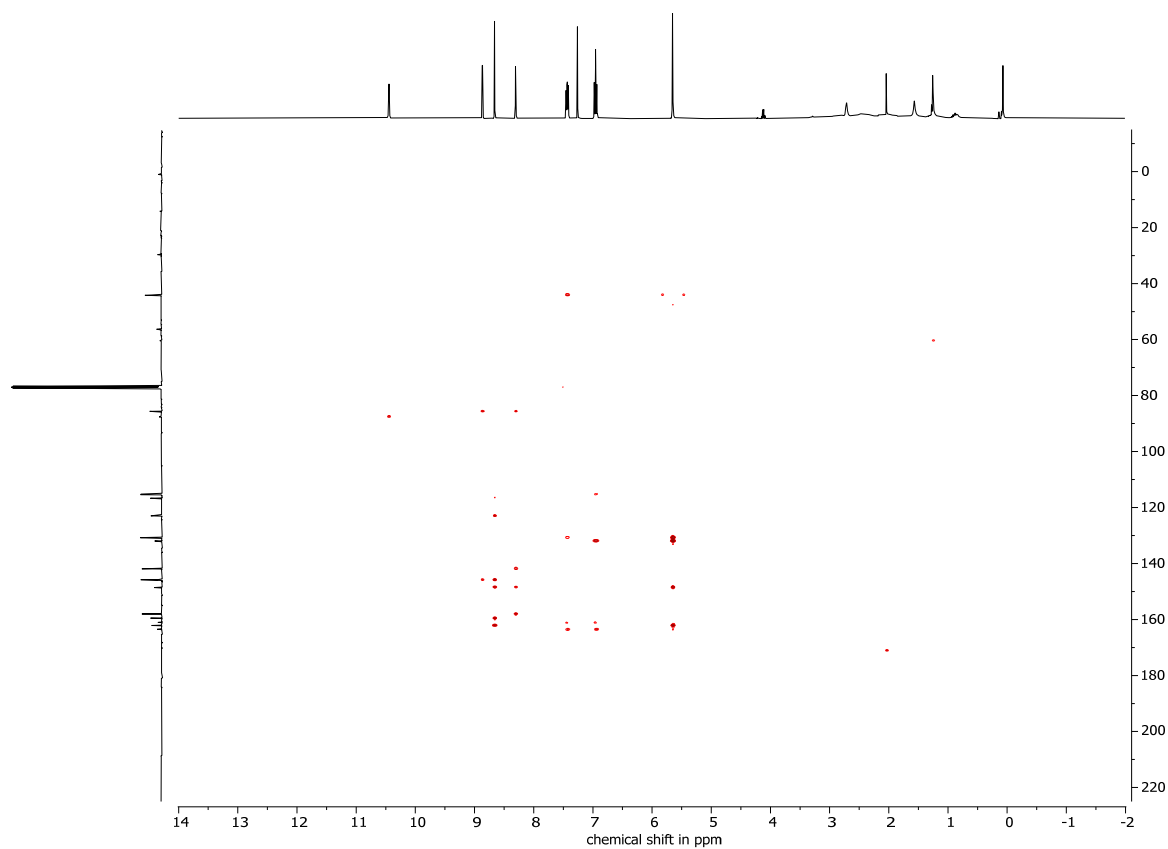
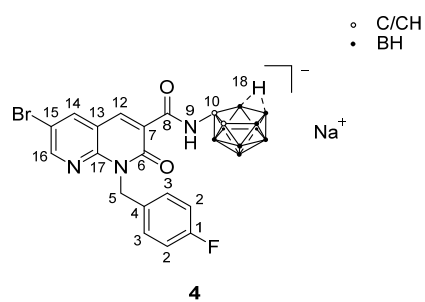


Figure S44. HMBC (^1H , ^{13}C) NMR spectrum of compound **3p** in CDCl_3 .

Compound 4



nido-ClusterCH = 11n

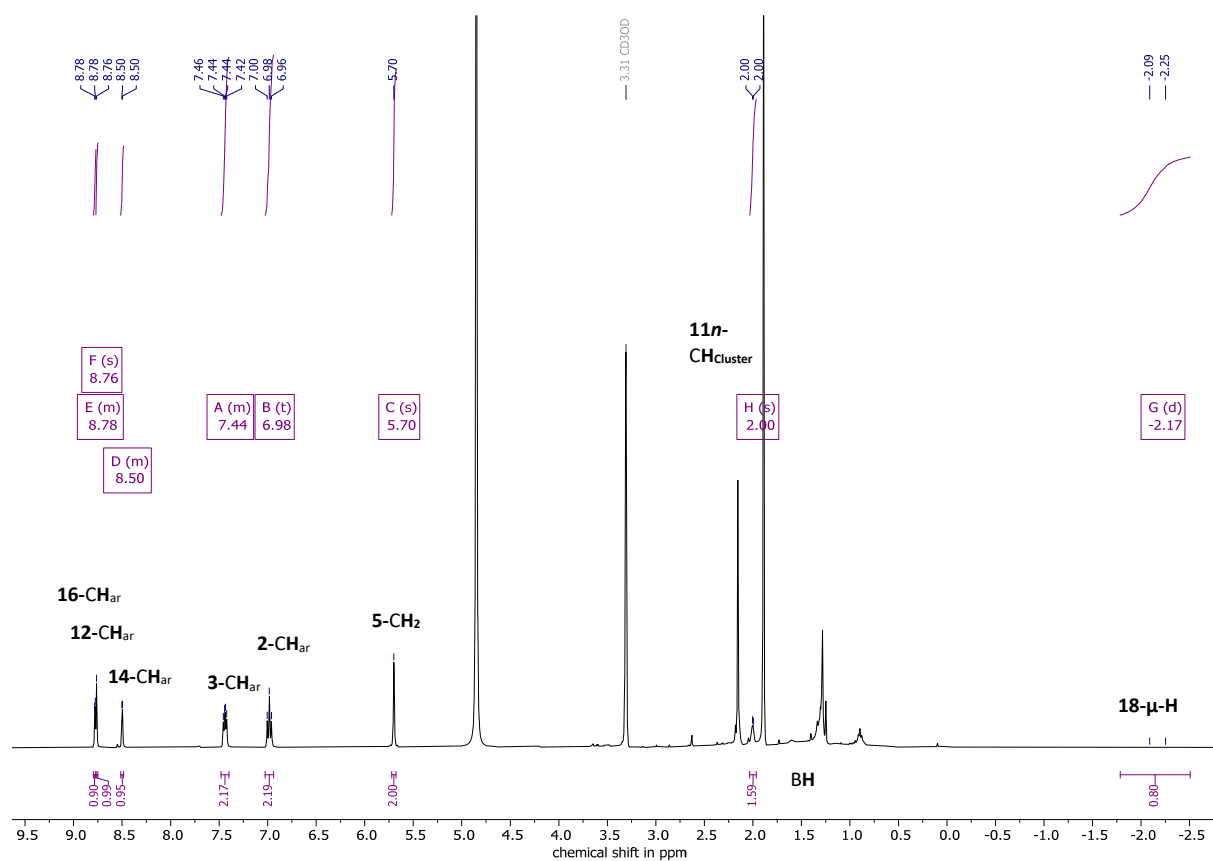


Figure S45. ¹H NMR spectrum of compound 4 in CD₃OD full spectrum.

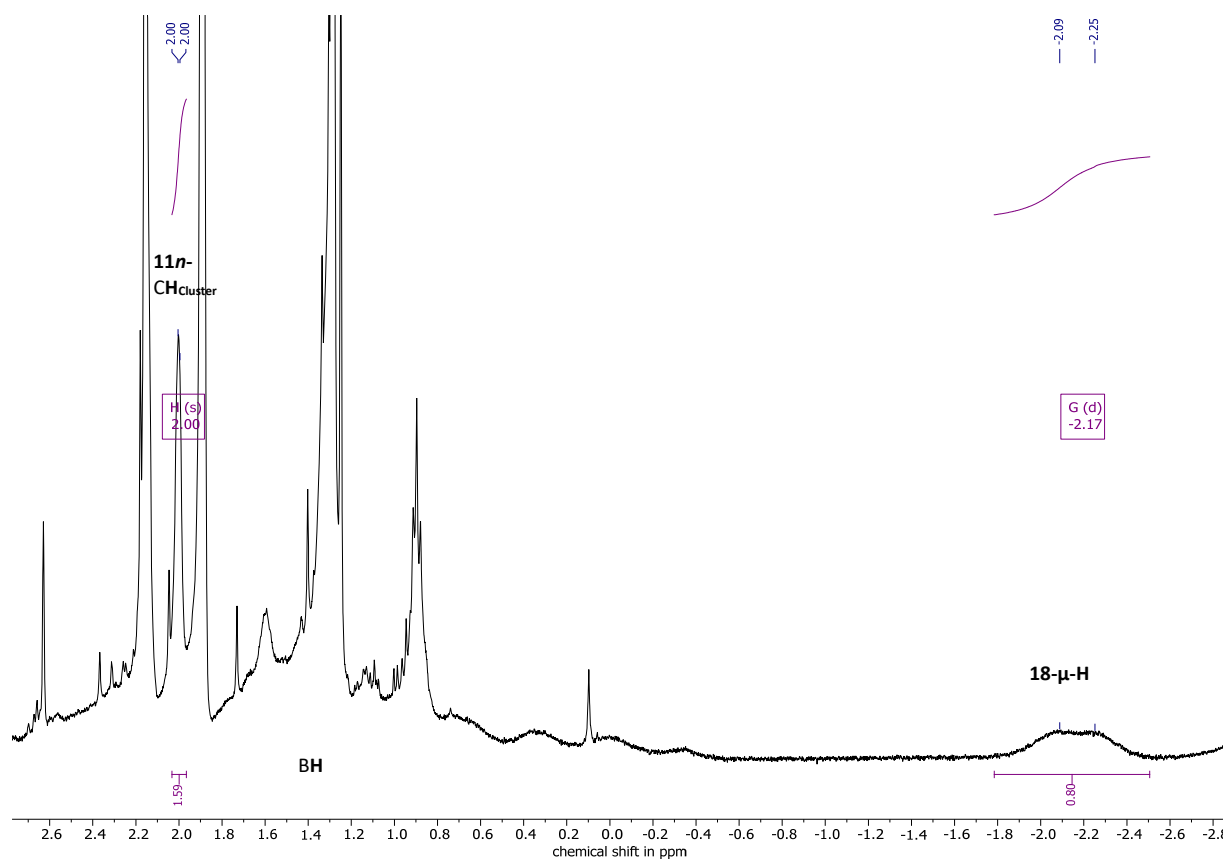


Figure S46. ^1H NMR spectrum of compound **4** in CD_3OD (magnified spectrum, zoomed in).

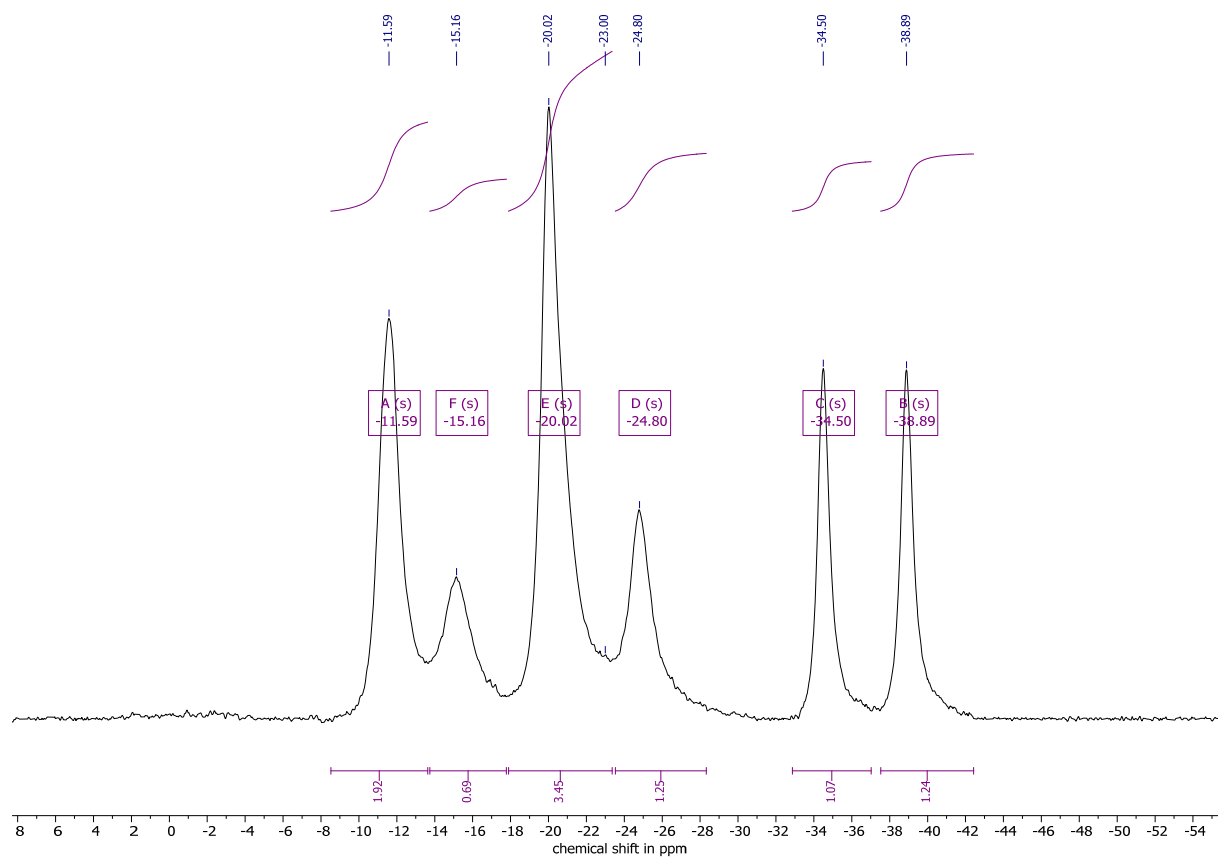


Figure S47. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **4** in CD_3OD .

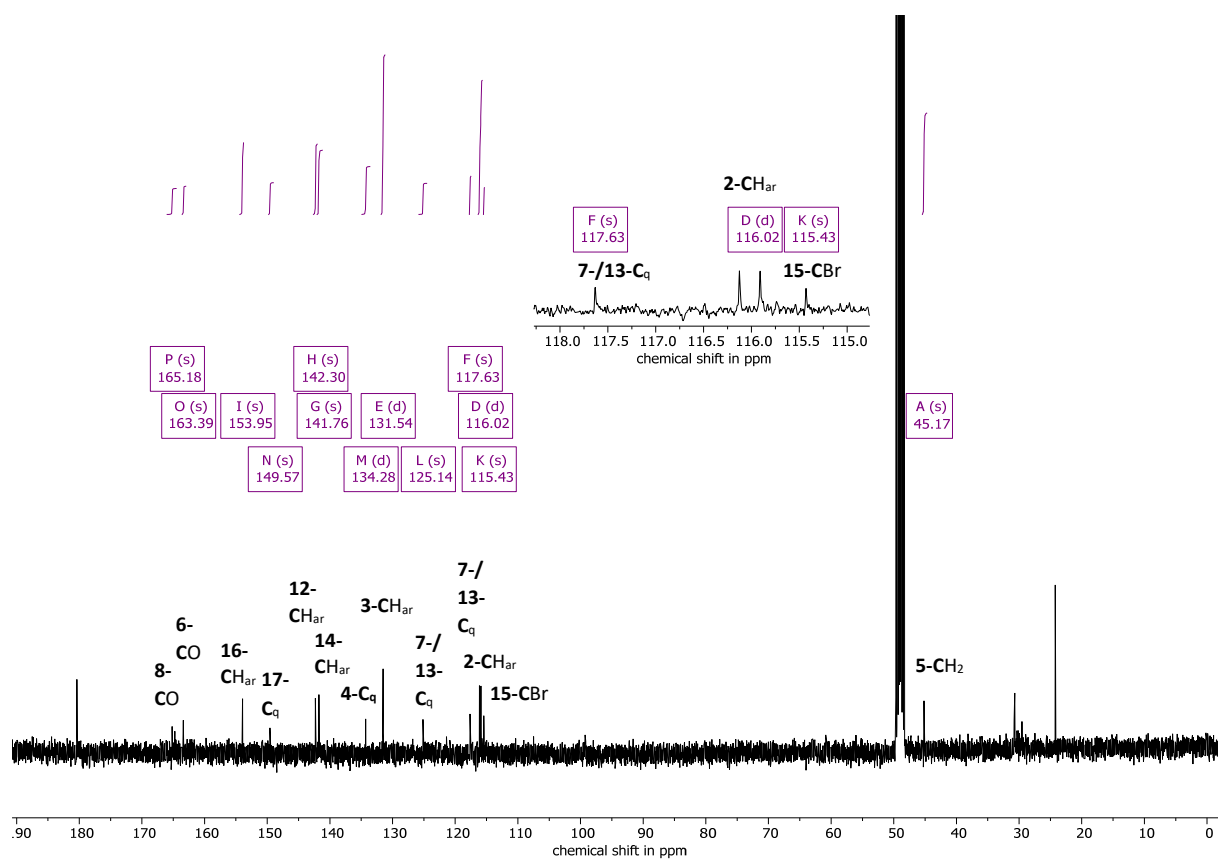


Figure S48. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **4** in CD_3OD .

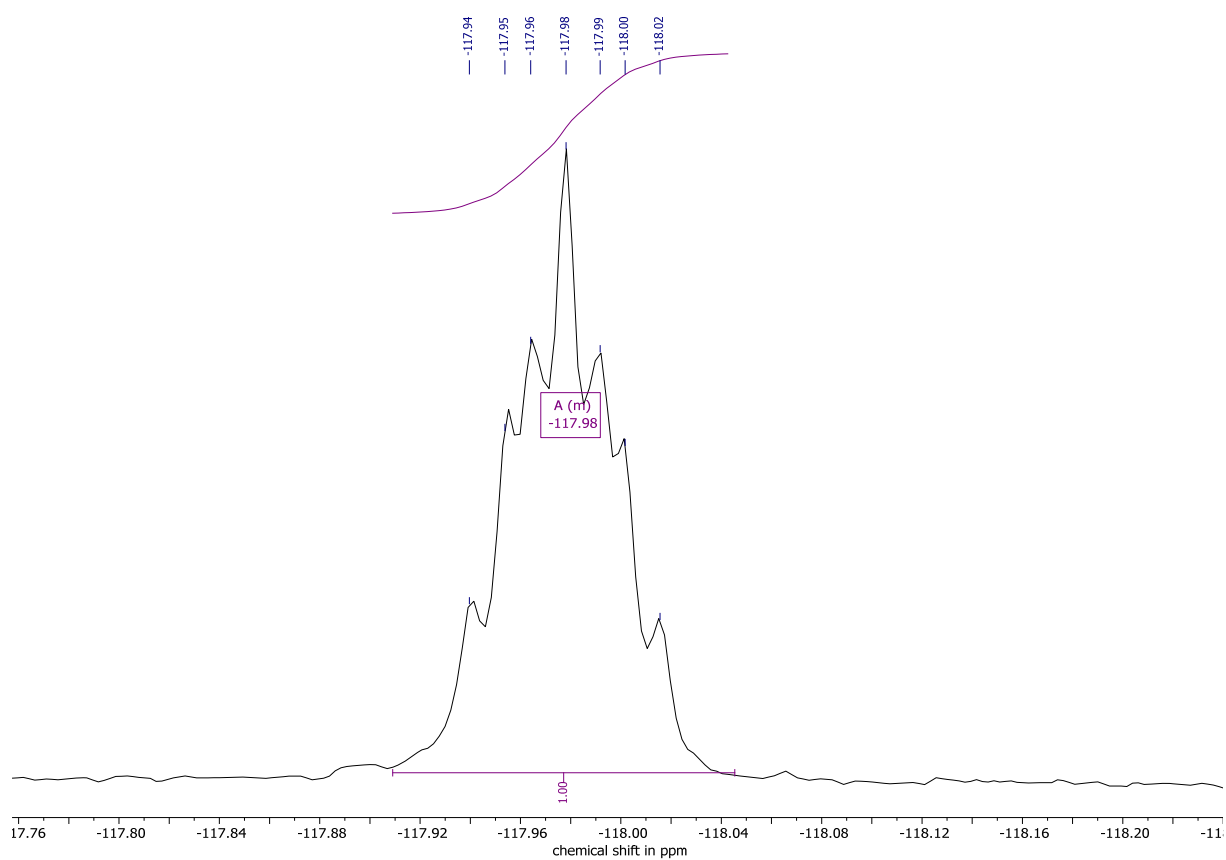


Figure S49. $^{19}\text{F}\{^1\text{H}\}$ NMR spectrum of compound **4** in CD_3OD .

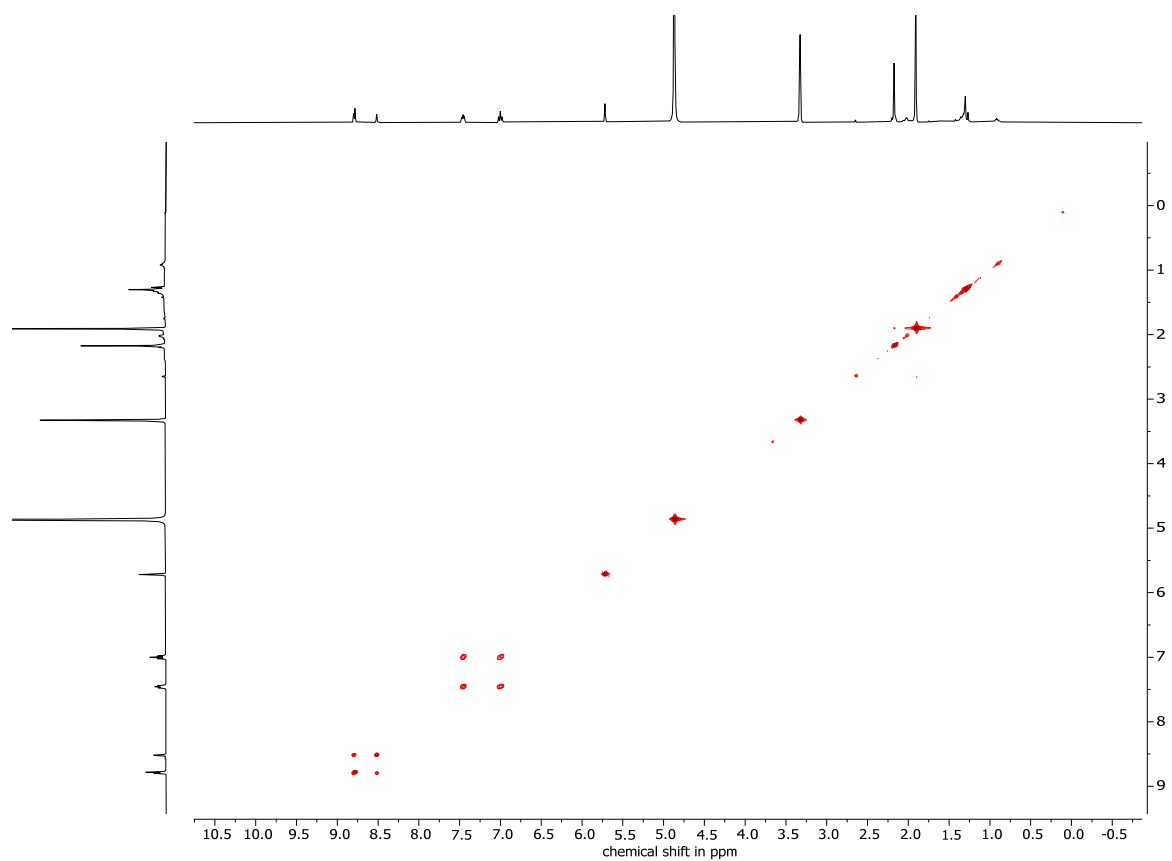


Figure S50. COSY (^1H , ^1H) NMR spectrum of compound **4** in CD_3OD .

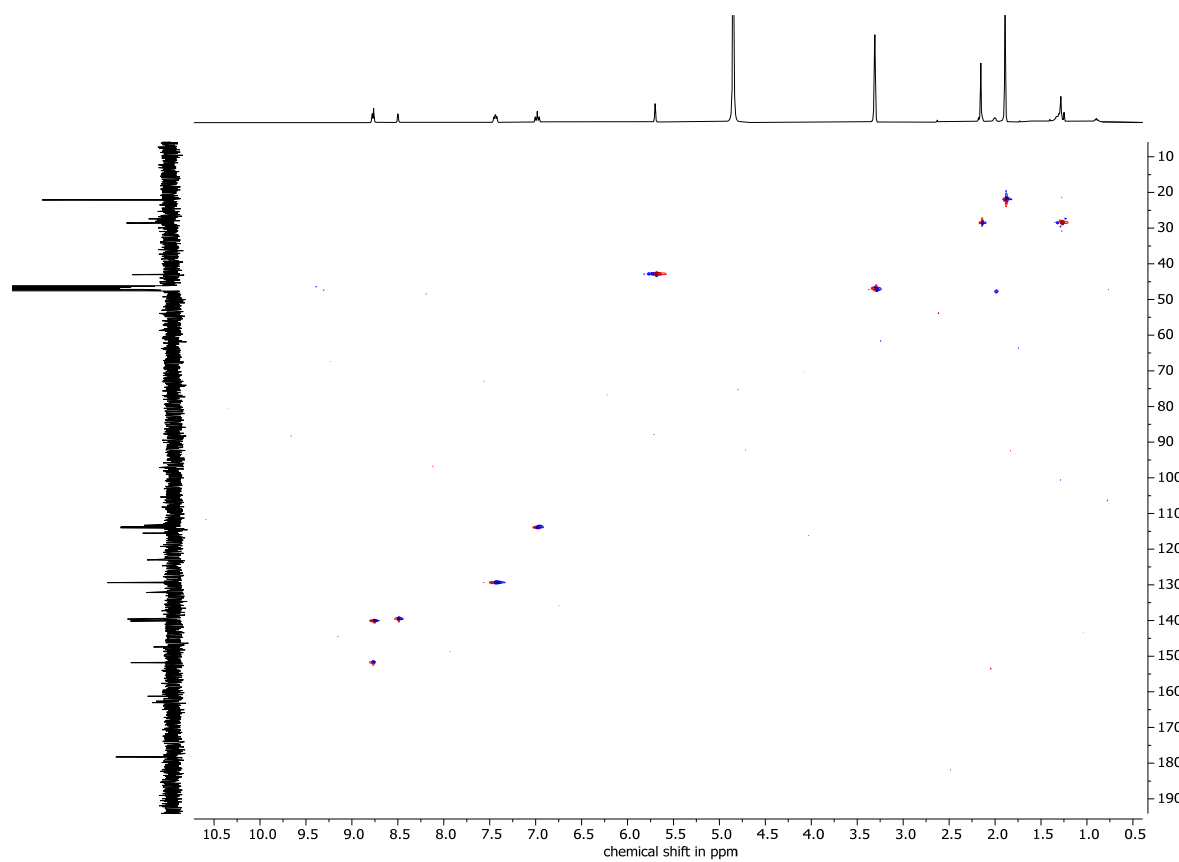


Figure S51. HSQC (^1H , ^{13}C) NMR spectrum of compound **4** in CD_3OD .

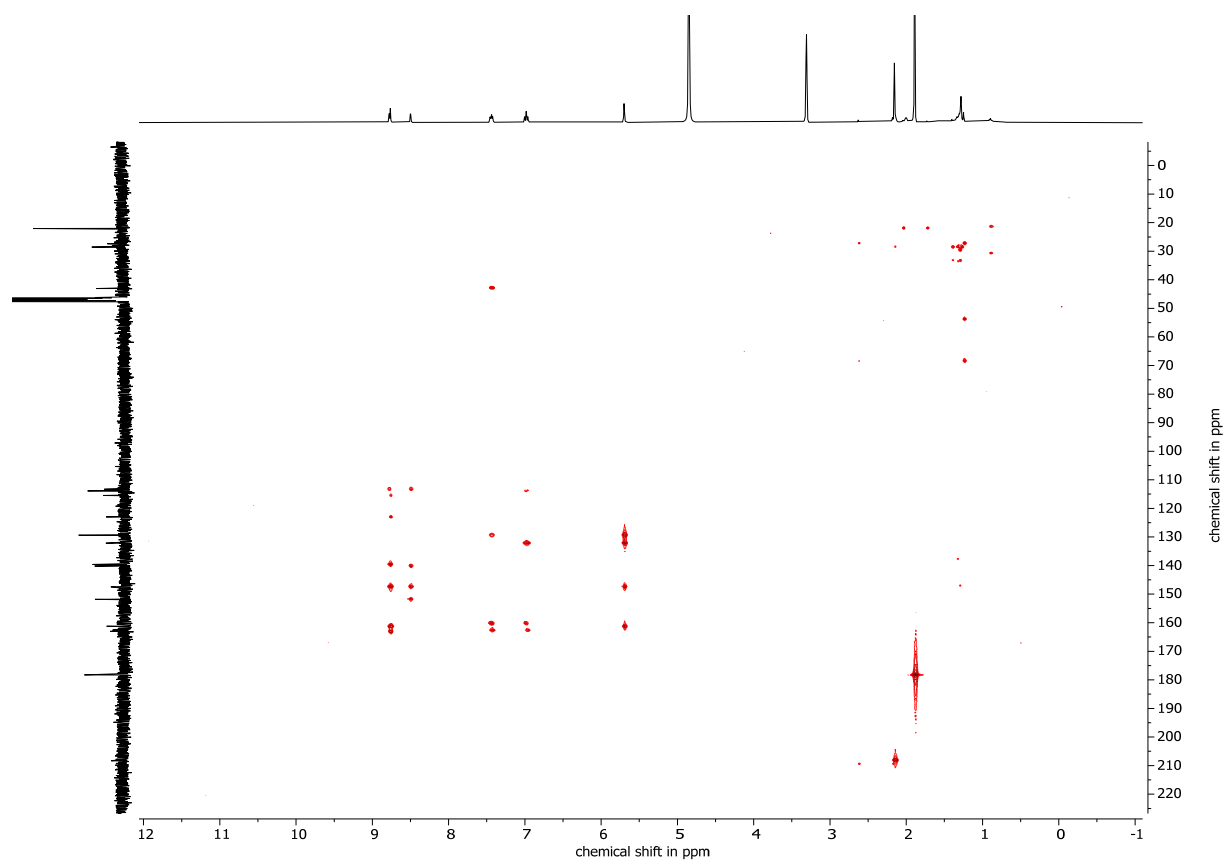
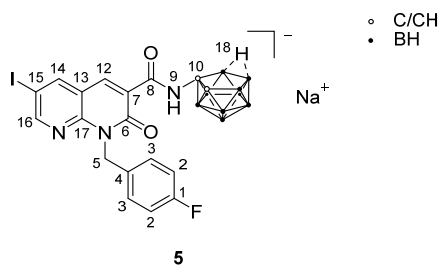


Figure S52. HMBC (^1H , ^{13}C) NMR spectrum of compound **4** in CD_3OD .

Compound **5**



nido-ClusterCH = 11*n*

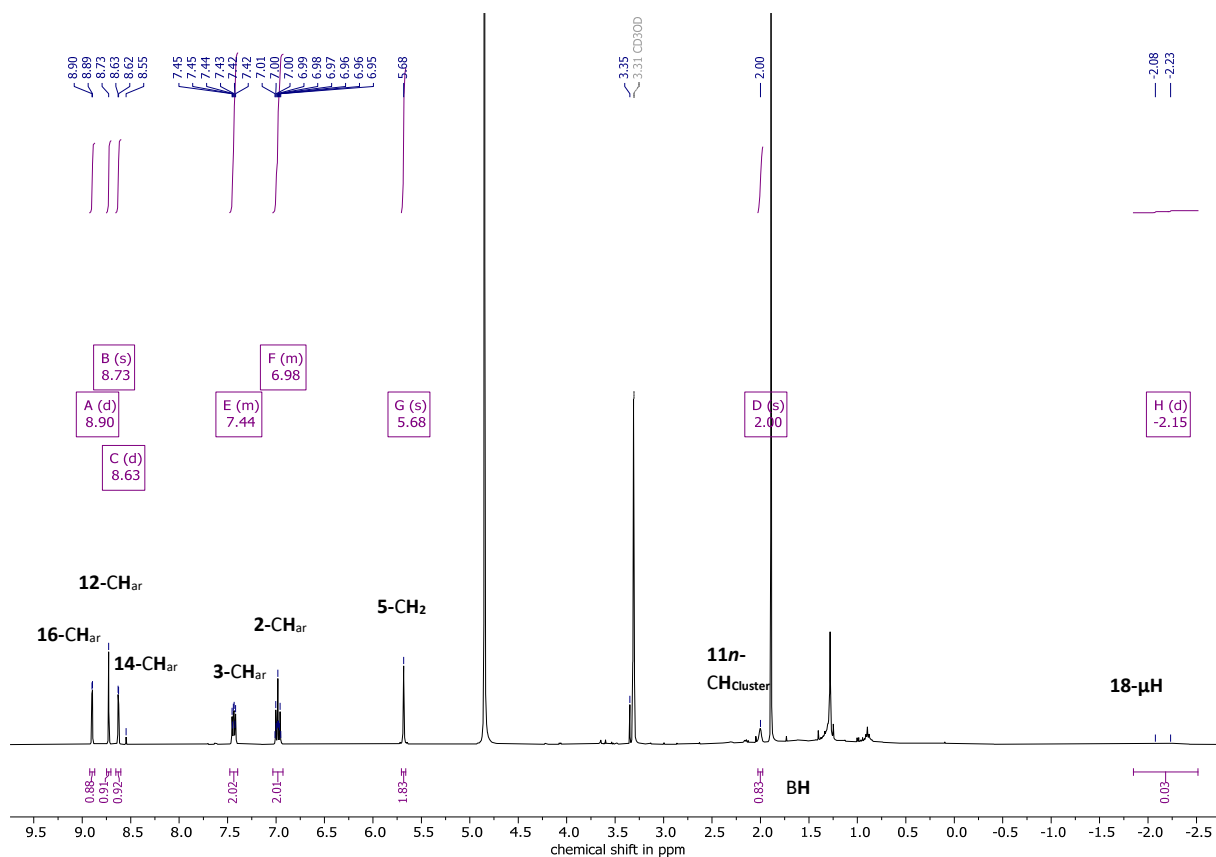


Figure S53. ^1H NMR spectrum of compound **5** in CD_3OD full spectrum.

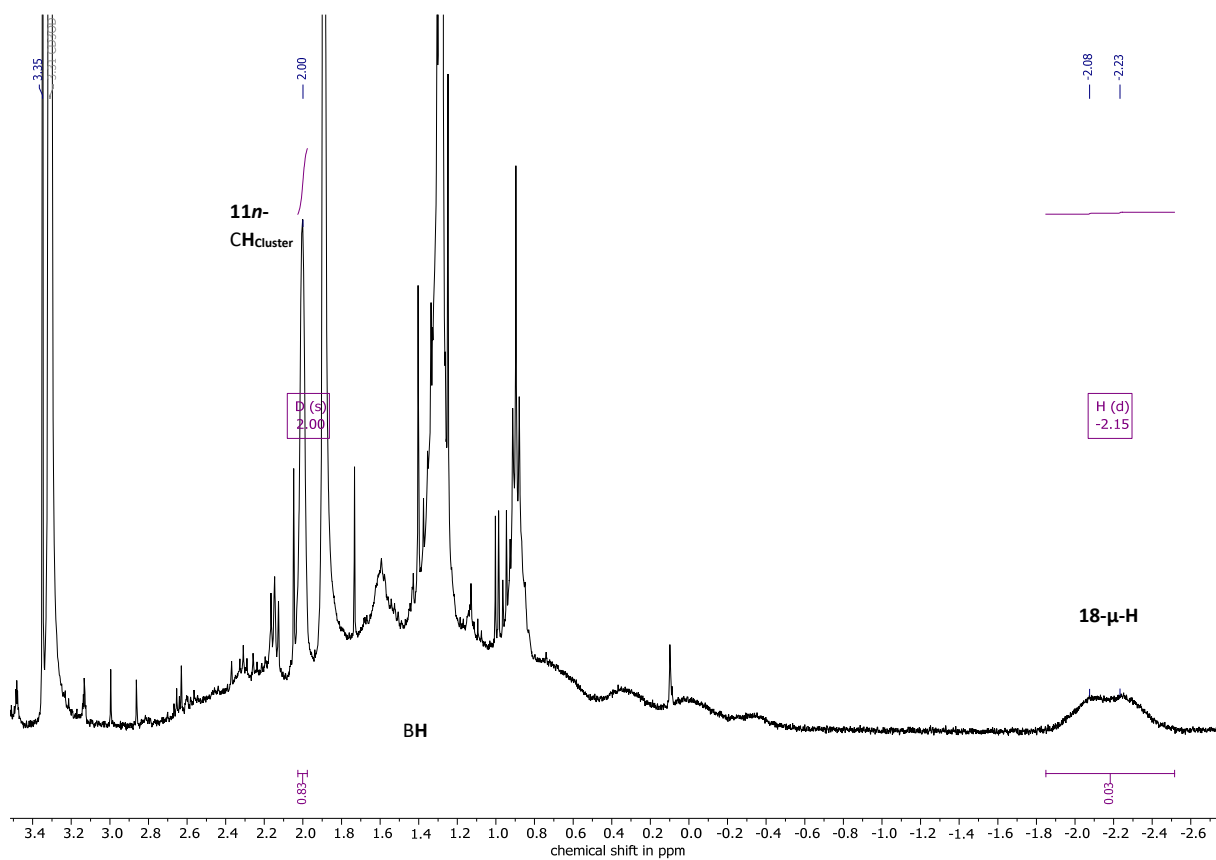


Figure S54. ^1H NMR spectrum of compound **5** in CD_3OD (magnified spectrum, zoomed in).

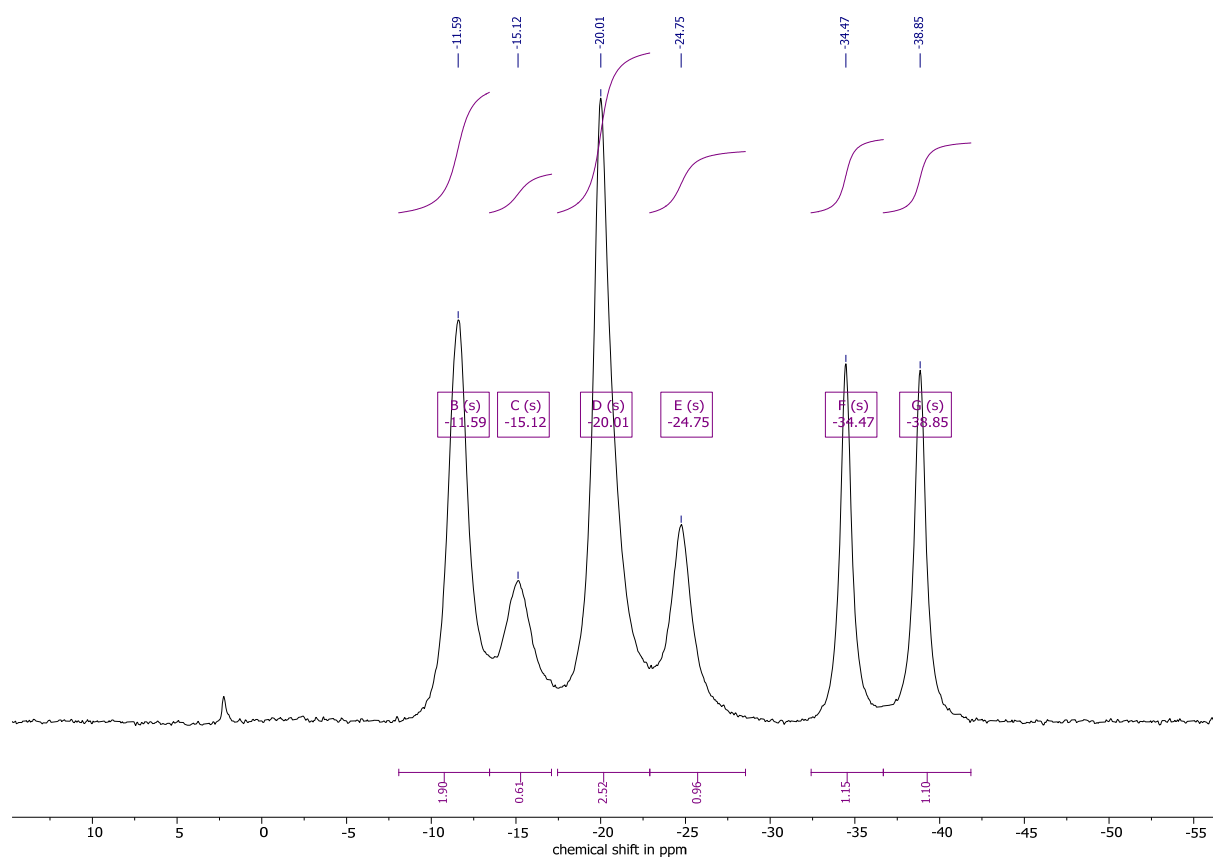


Figure S55. $^{11}\text{B}\{^1\text{H}\}$ NMR spectrum of compound **5** in CD_3OD .

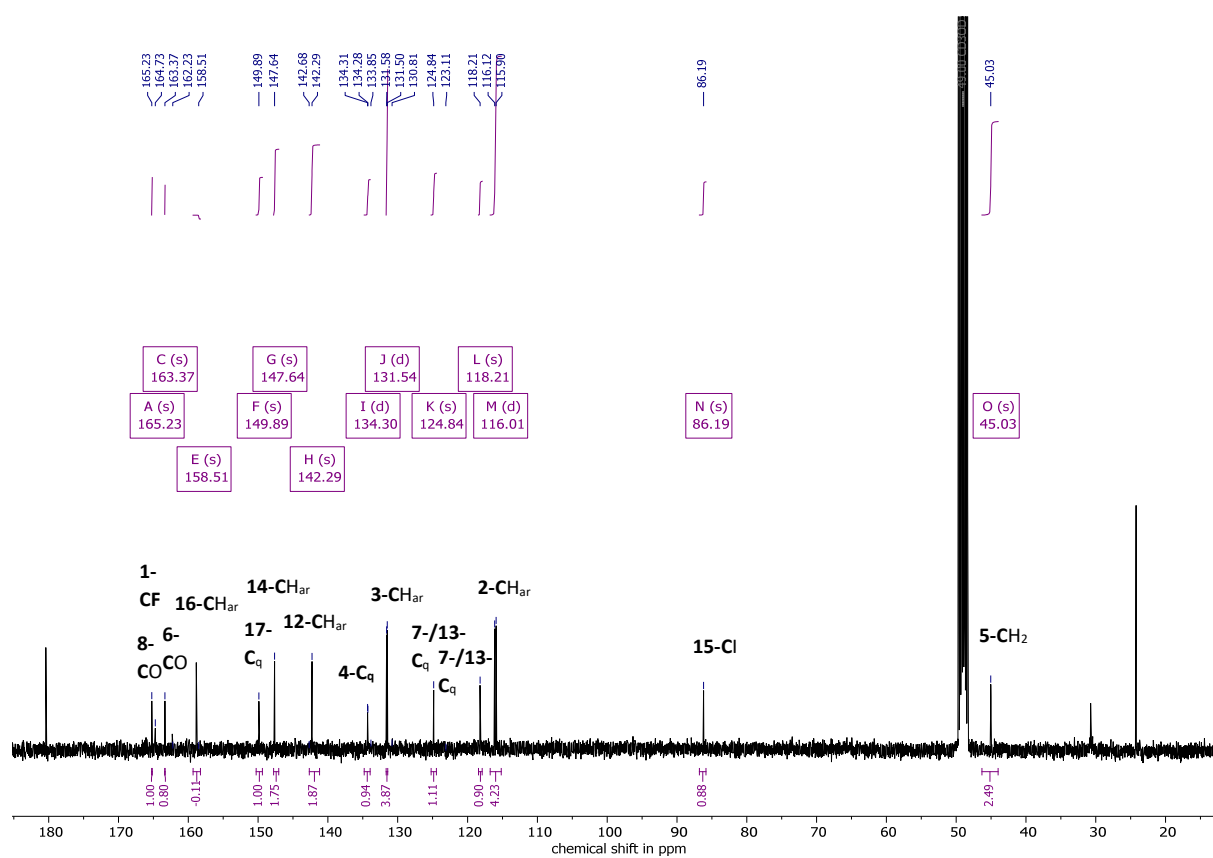


Figure S56. $^{13}\text{C}\{^1\text{H}\}$ NMR spectrum of compound **5** in CD_3OD .

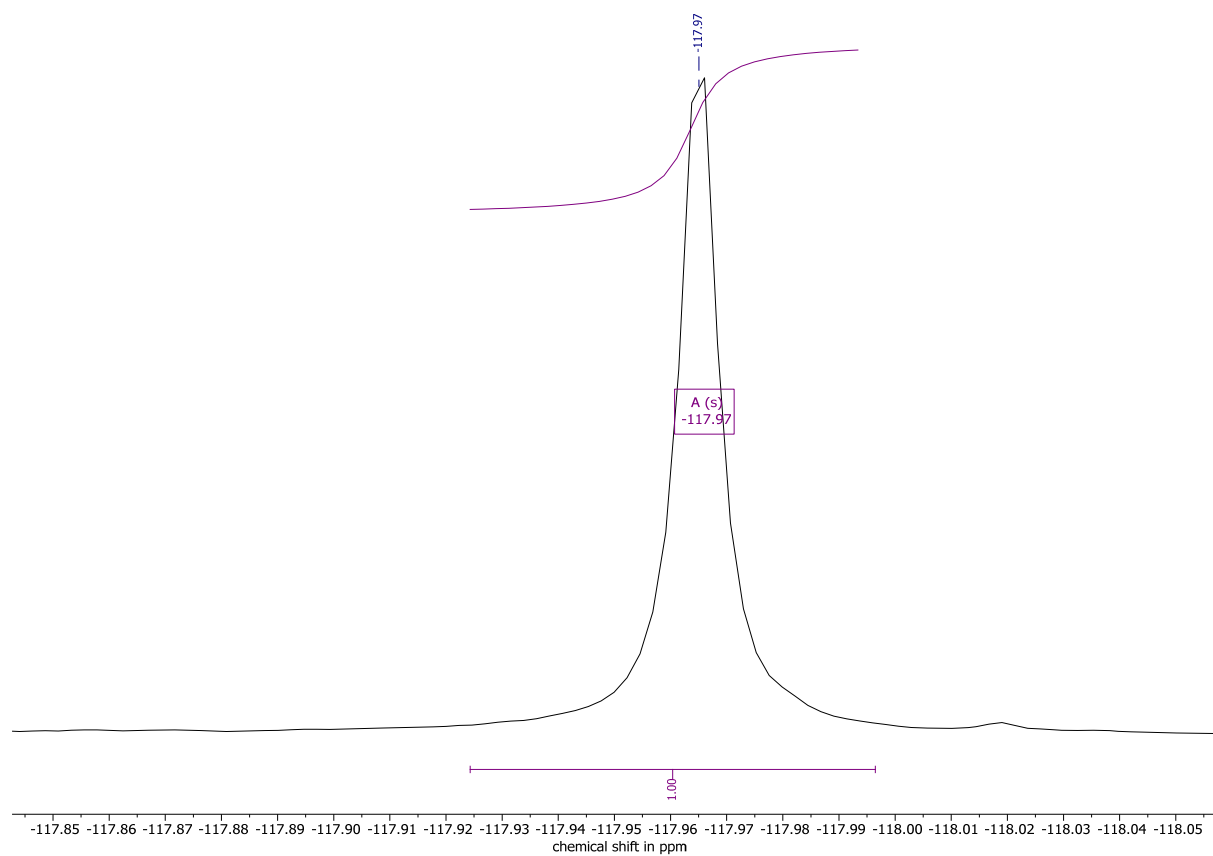


Figure S57. $^{19}\text{F}\{^1\text{H}\}$ NMR spectrum of compound **5** in CD_3OD .

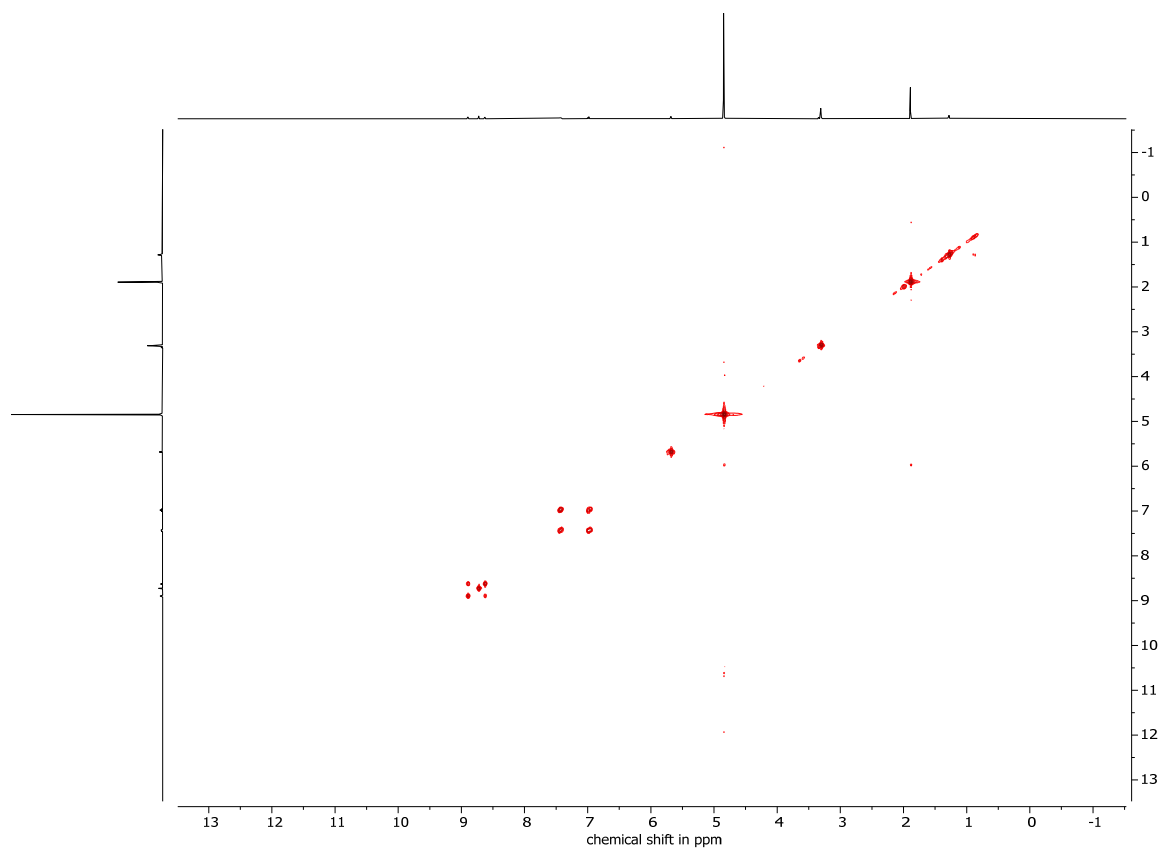


Figure S58. COSY (^1H , ^1H) NMR spectrum of compound **5** in CD_3OD .

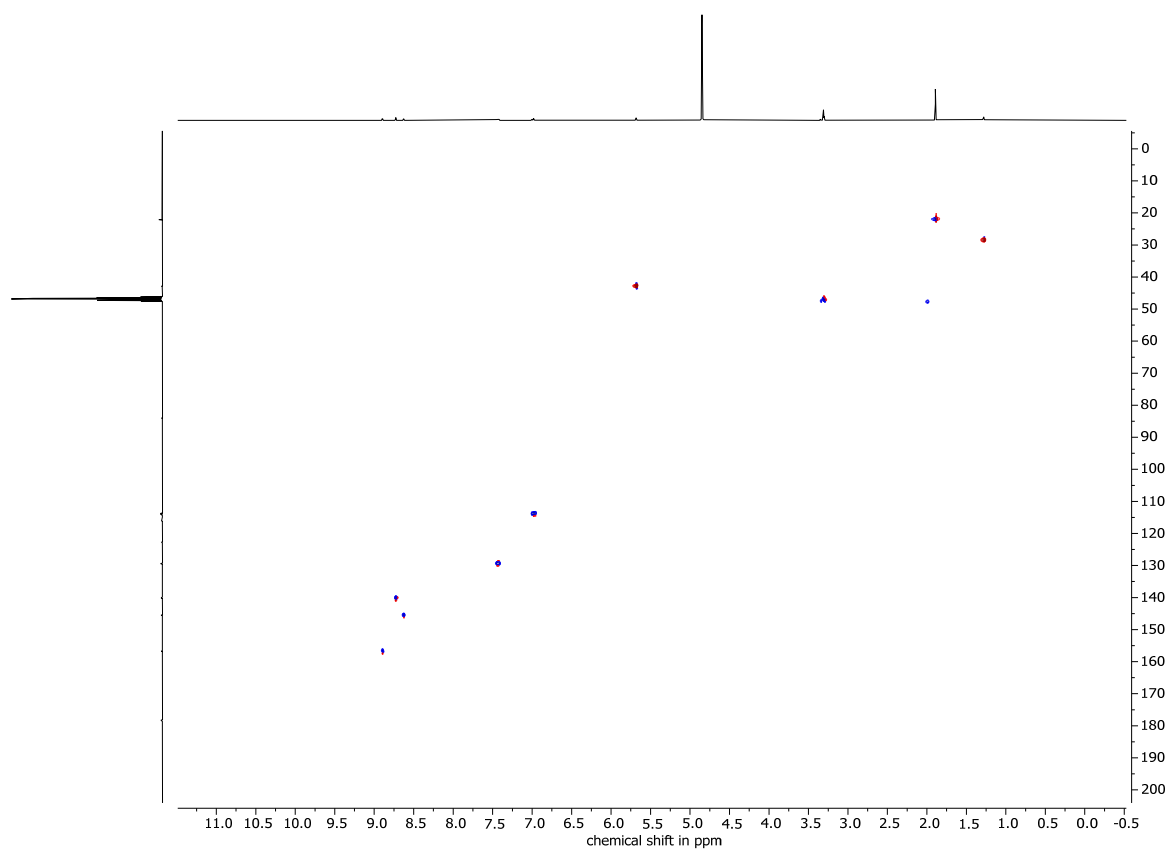


Figure S59. HSQC (^1H , ^{13}C) NMR spectrum of compound **5** in CD_3OD .

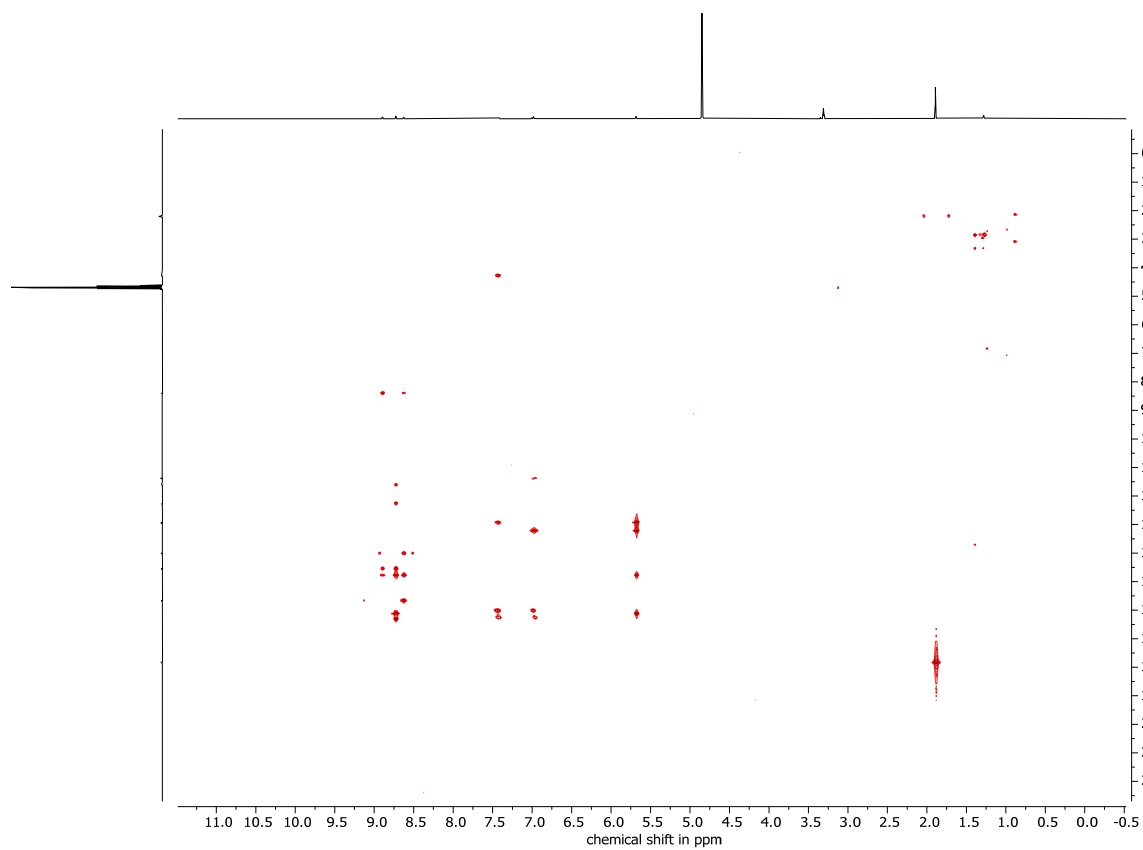


Figure S60. HMBC (^1H , ^{13}C) NMR spectrum of compound **5** in CD_3OD .

3 HR-ESI Mass Spectra of Compounds $2_{o,m,p}$, $3_{o,m,p}$, 4, and 5

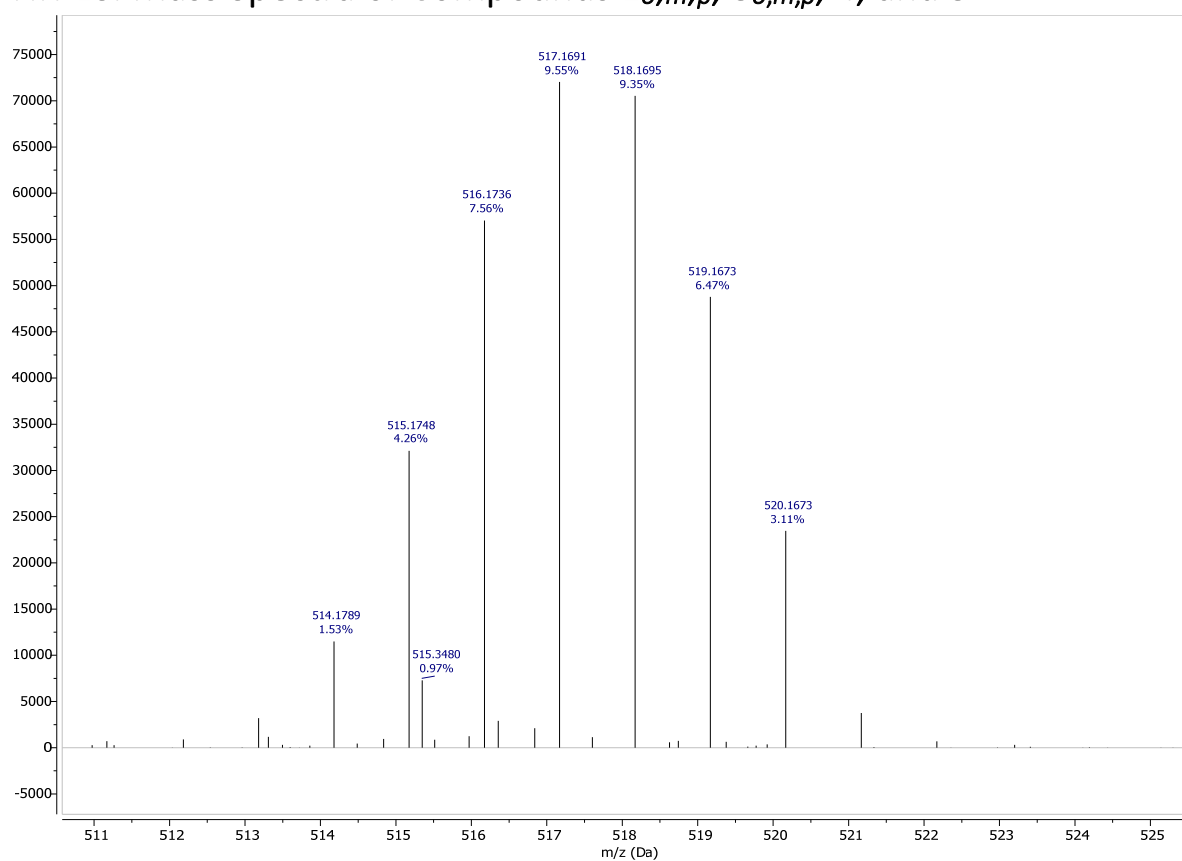


Figure S61. HRMS (ESI-) of compound 2_o in CH_3CN .

P:/HR ESI MS 2...11_01_50434.zip Injection 1 +MS profile He...LUe674_050724 MS + spectrum 0.64

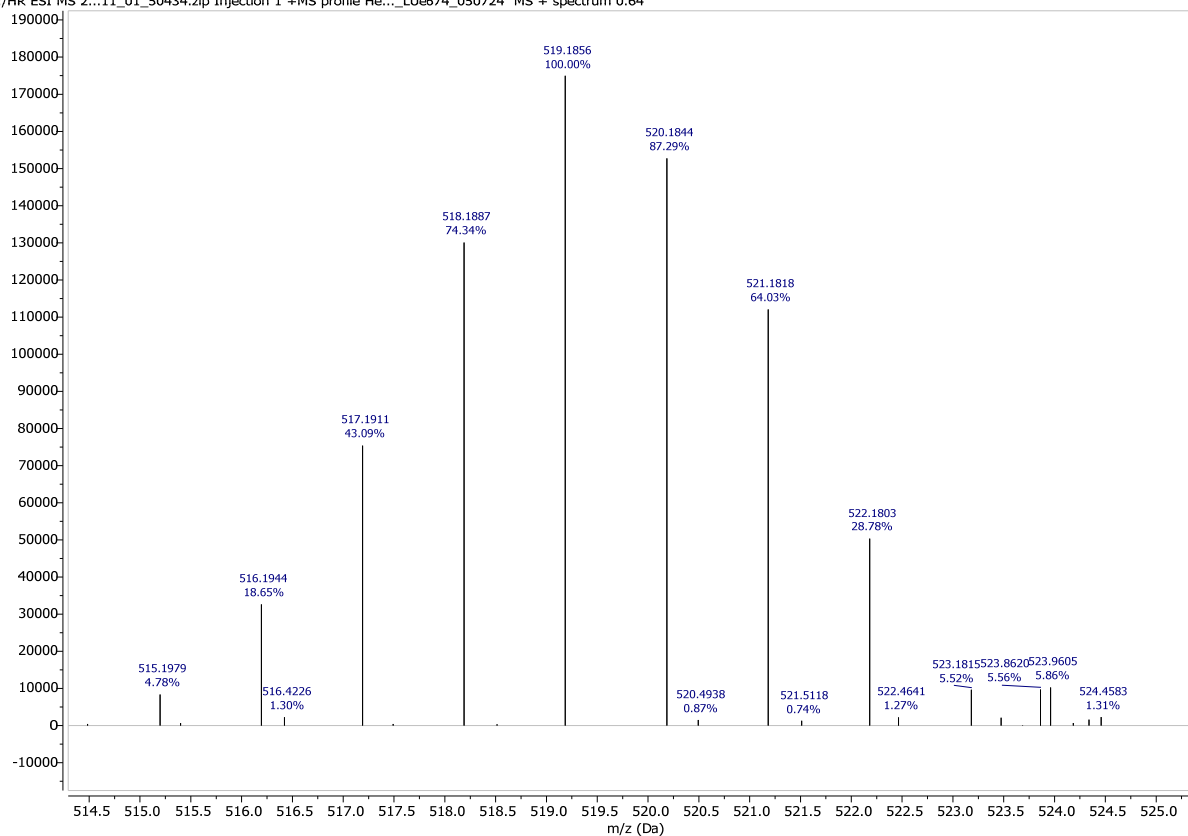


Figure S62. HRMS (ESI+) of compound 2_m in CH_3CN .

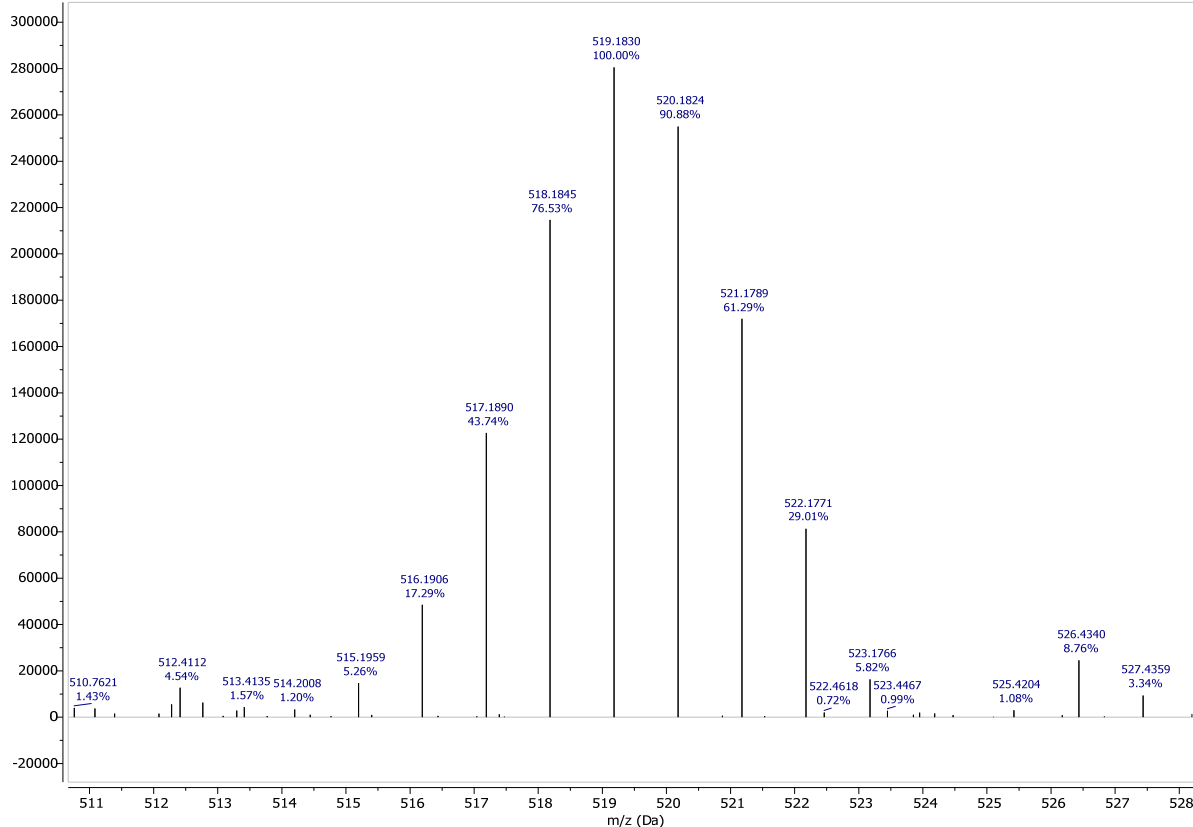


Figure S63. HRMS (ESI+) of compound **2_p** in CH₃CN.

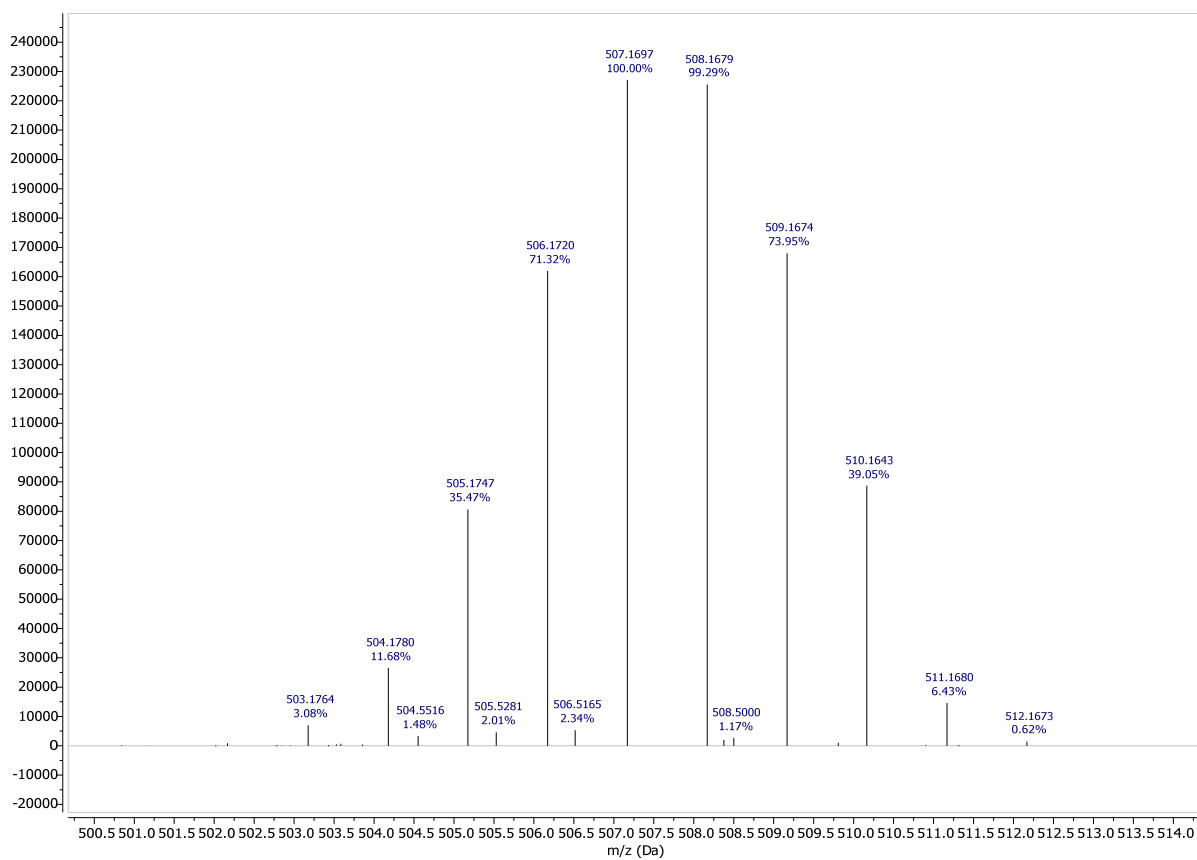


Figure S64. HRMS (ESI-) of compound **4** in MeOH.

P:/HR ESI MS 2...._6_01_50014.zip Injection 1 -MS profile He..._LUe668_170624 MS - spectrum 0.67

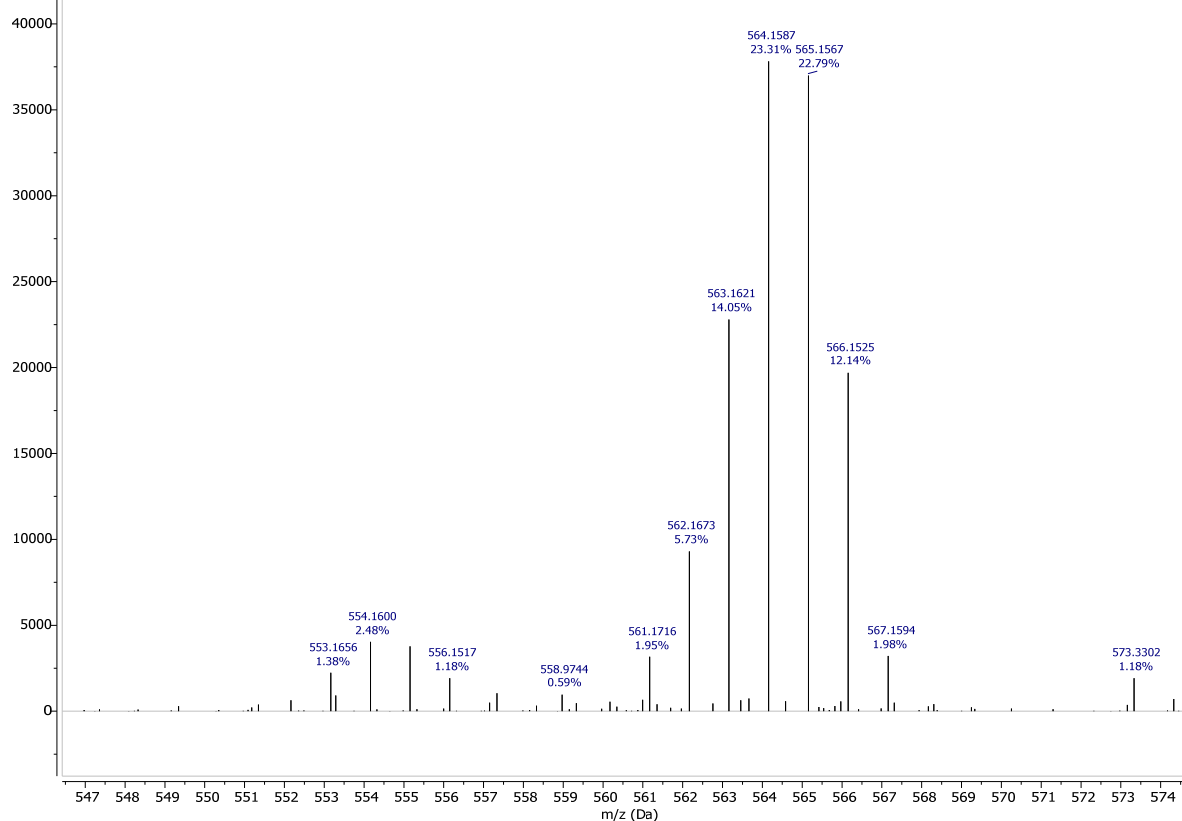


Figure S65. HRMS (ESI-) of compound **3_o** in CH₃CN.

P:/HR ESI MS 2...._5_01_50061.zip Injection 1 +MS profile He..._LUe672_170624 MS + spectrum 0.80

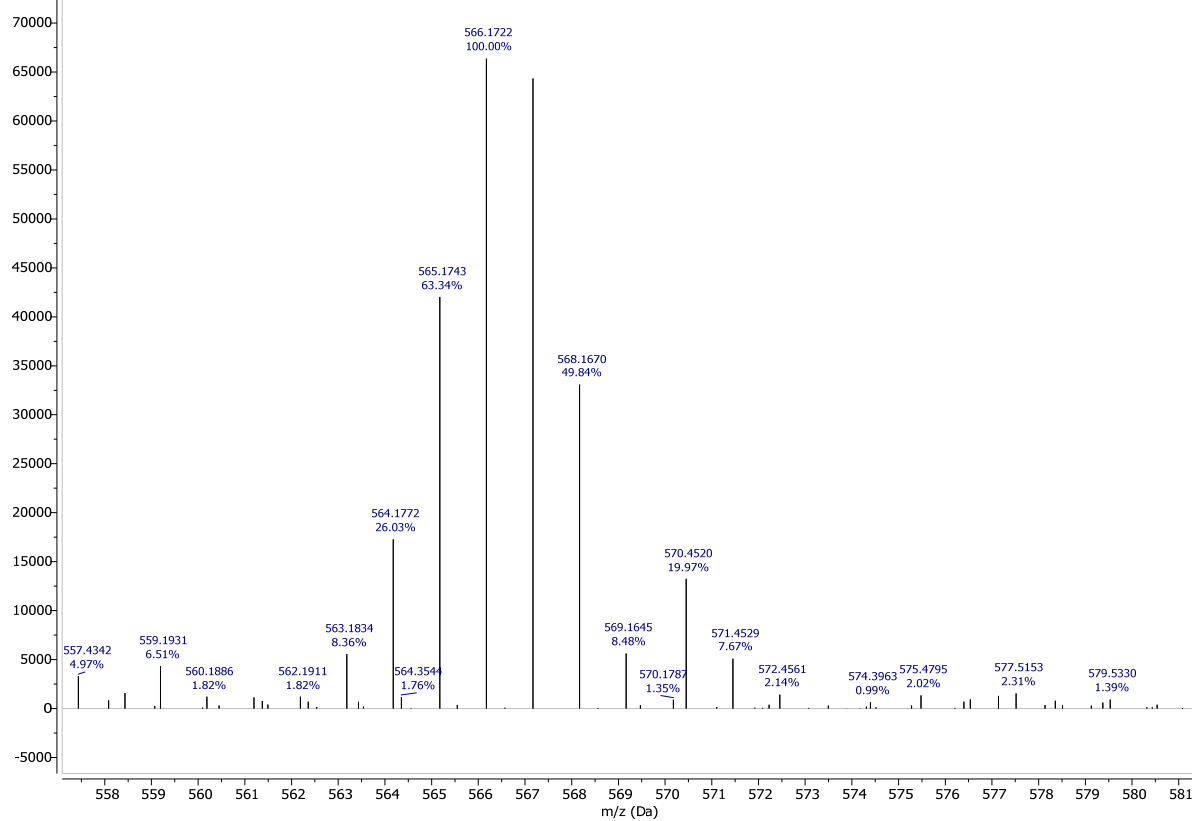


Figure S66. HRMS (ESI+) of compound **3_m** in CH₃CN.

P:/HR ESI MS 2..._9_01_50065.zip Injection 1 +MS profile He..._LUe673_170624 MS + spectrum 0.59

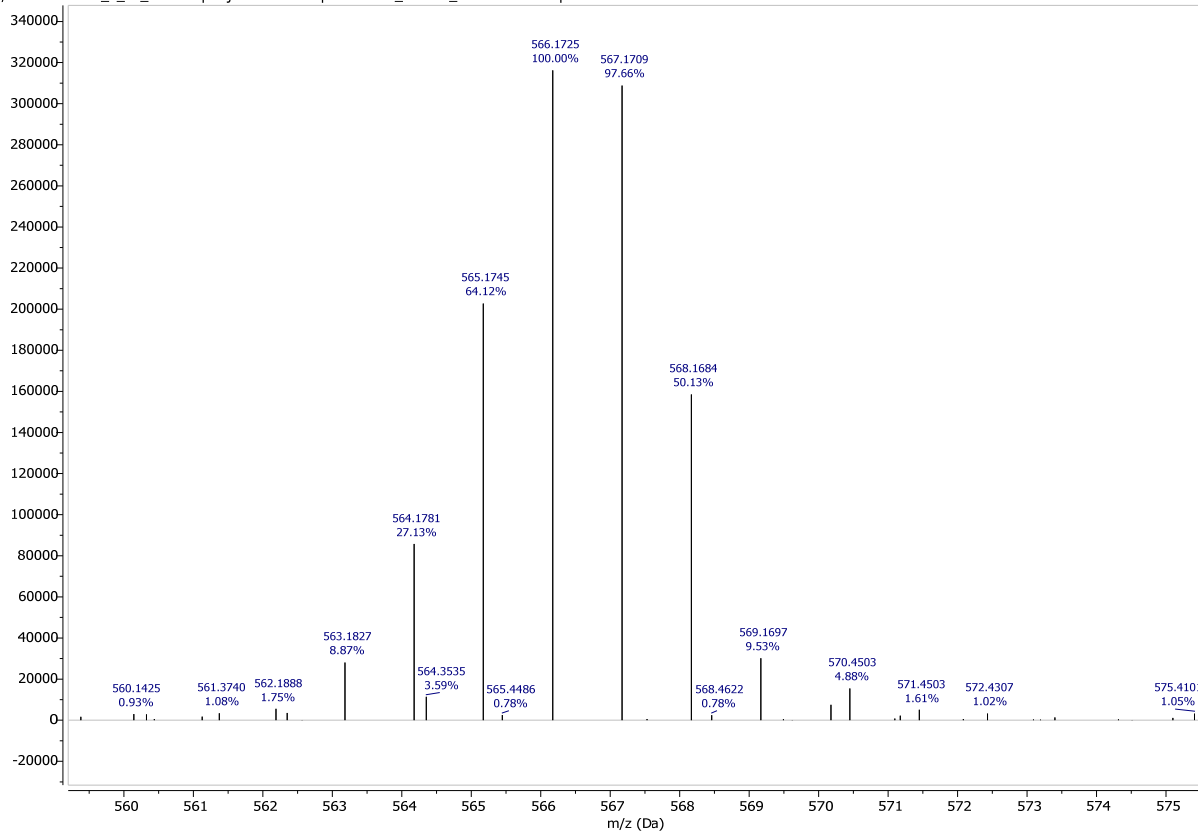


Figure S67. HRMS (ESI+) of compound **3_p** in CH₃CN.

P:/HR ESI MS 2...15_01_50873.zip Injection 1 -MS profile He..._LUe689_260724 MS - spectrum 0.42

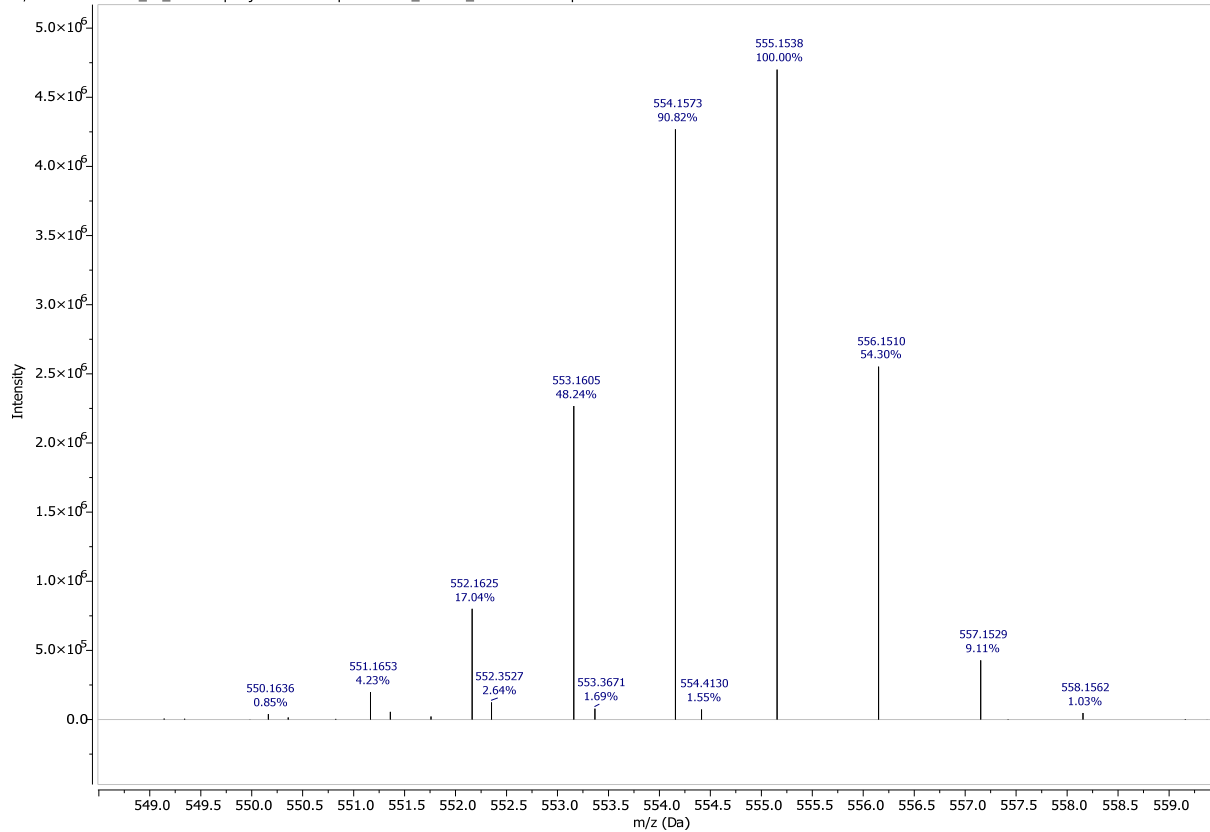


Figure S68. HRMS (ESI-) of compound **5** in MeOH.

4 Determination of HPLC Purity of Compounds **2_{o,m,p}**, **3_{o,m,p}**, **4**, and **5**

The HPLC purity was determined using HPLC-UV-MS with an RP column. The *closo*-carborane derivatives **2_{o,m,p}** and **3_{o,m,p}** were dissolved in CH₃CN, the *nido*-carborane derivatives **4** and **5** in MeOH. All compounds had a purity >95%. The double peaks or shoulders visible for the signals of the compounds are related to the column that has been used and not to the compound, since variations in gradient system could not resolve the double-peaks or shoulders. Broad signals have been observed in case of *nido*-carborane derivatives.

Compound **2_o**

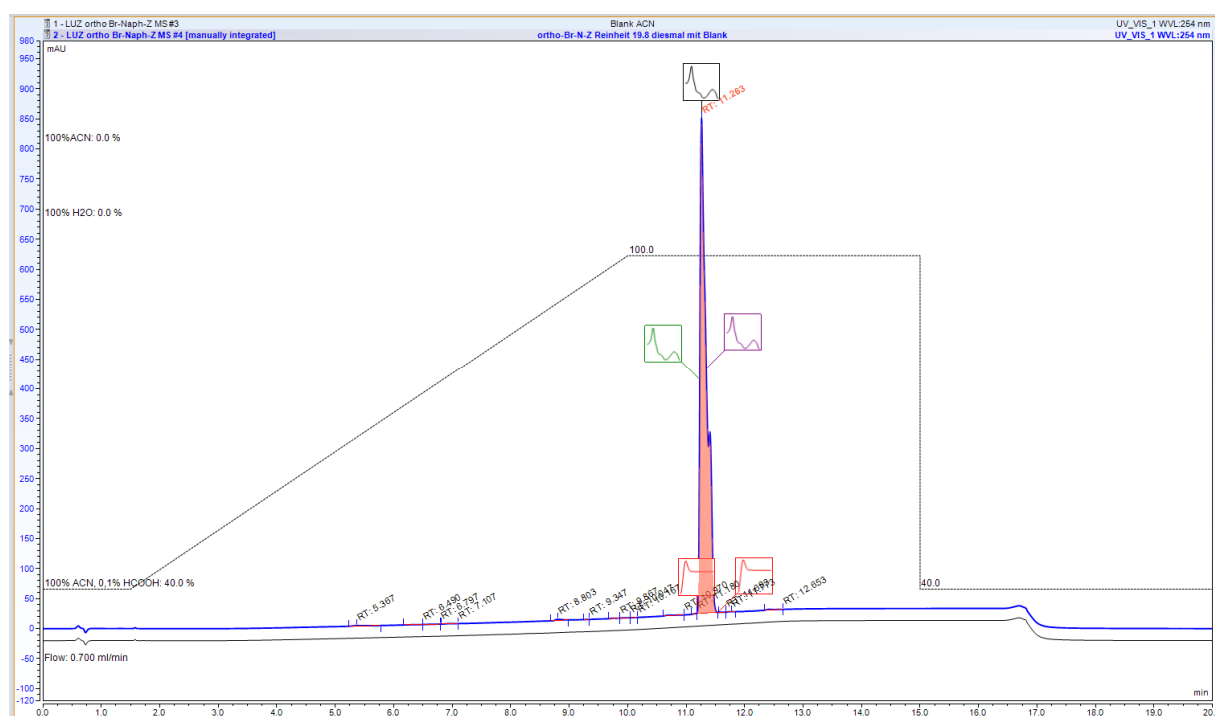


Figure S69. RP-HPLC chromatogram of blank (CH₃CN) and compound **2_o**, retention time: 11.3 min.

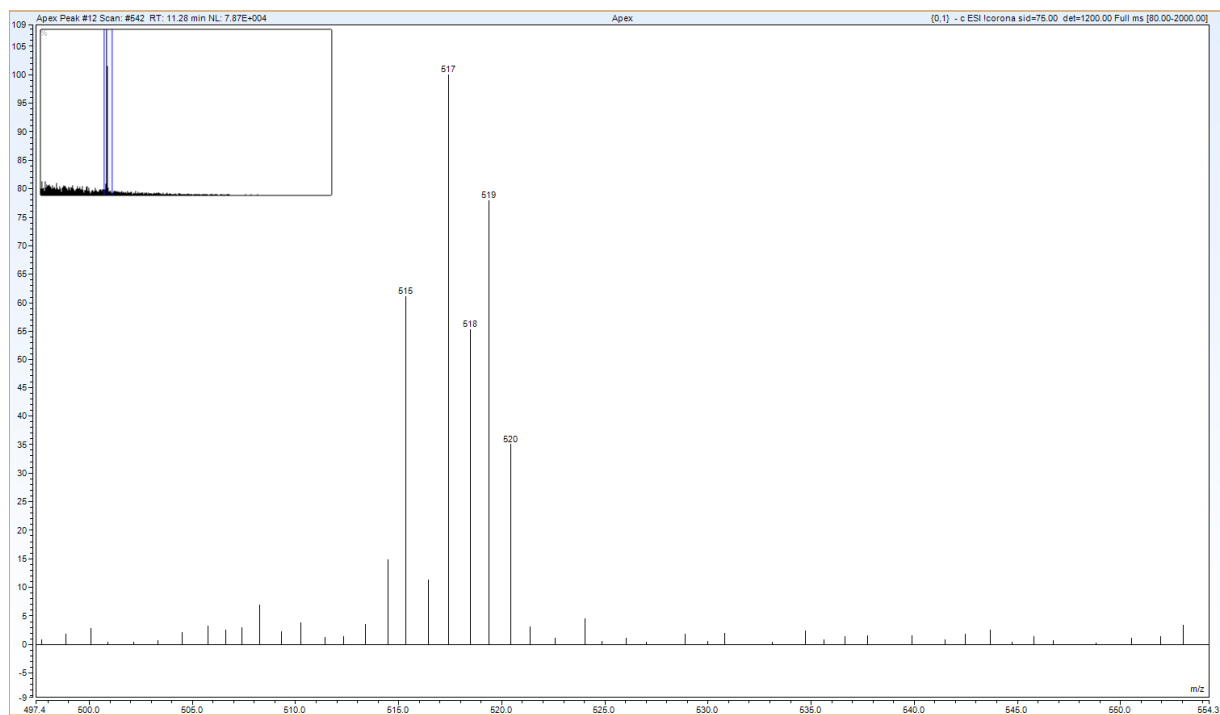


Figure S70. MS(-) of compound **2_o**, retention time: 11.3 min.

Compound **2_m**

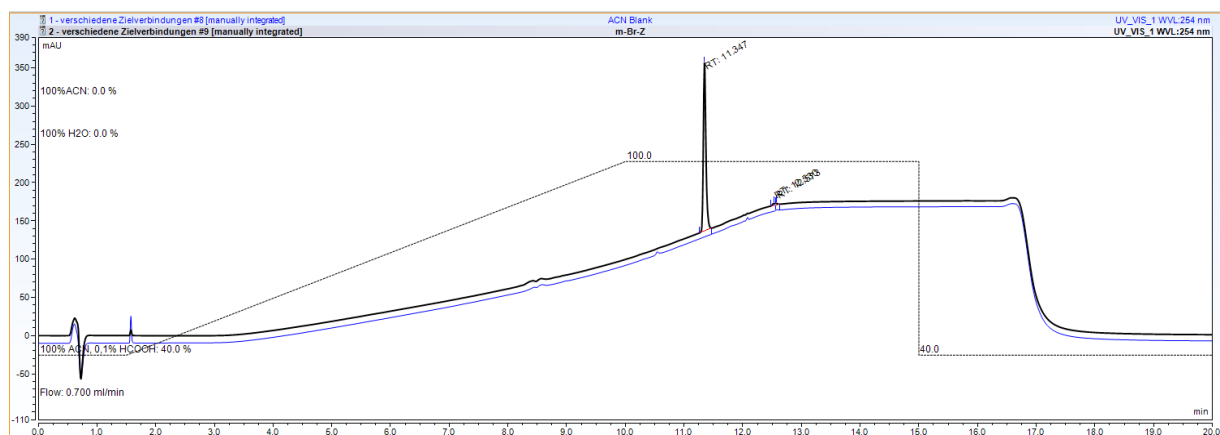


Figure S71. RP-HPLC chromatogram of blank (CH₃CN) and compound **2_m**, retention time: 11.3 min.

Compound **2_p**

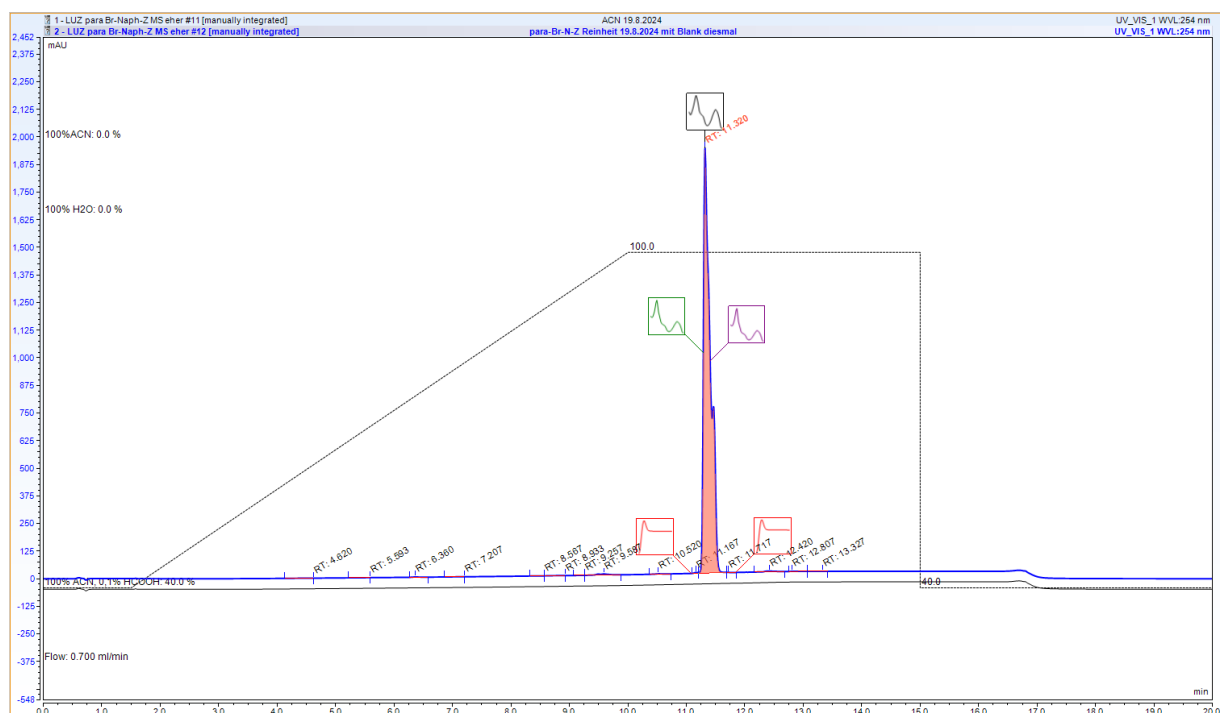


Figure S72. RP-HPLC chromatogram of blank (CH₃CN) and compound **2_p**, retention time: 11.3 min.

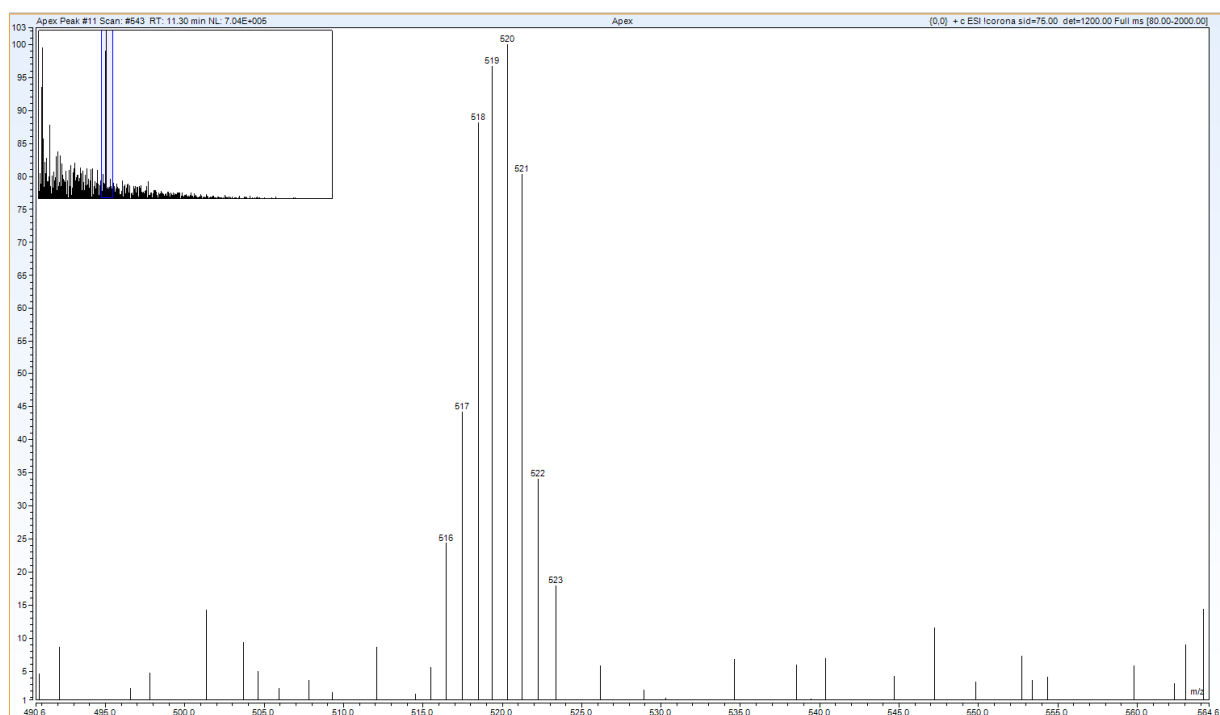


Figure S73. MS(+) of compound **2_p**, retention time: 11.3 min.

Compound **3_o**

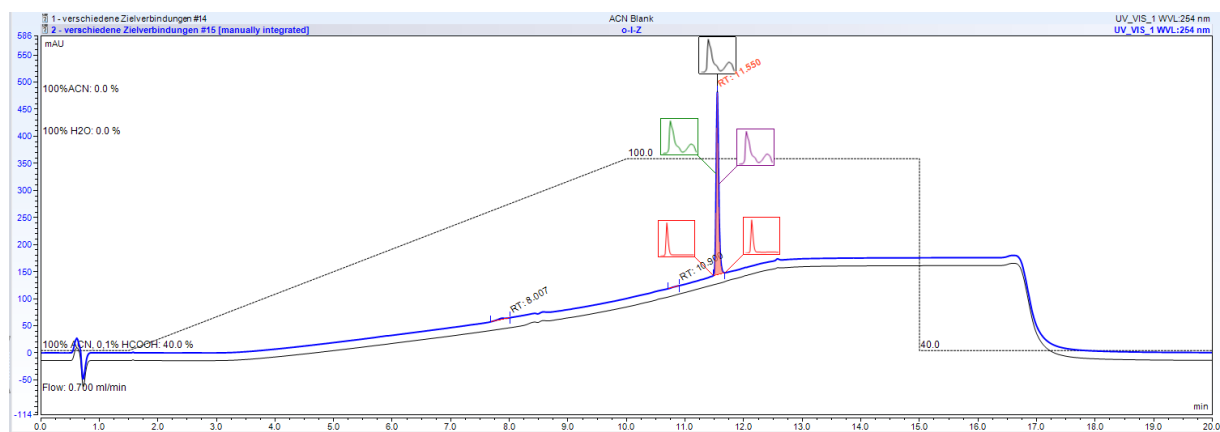


Figure S74. RP-HPLC chromatogram of blank (CH₃CN) and compound **3_o**, retention time: 11.6 min.

Compound **3_m**

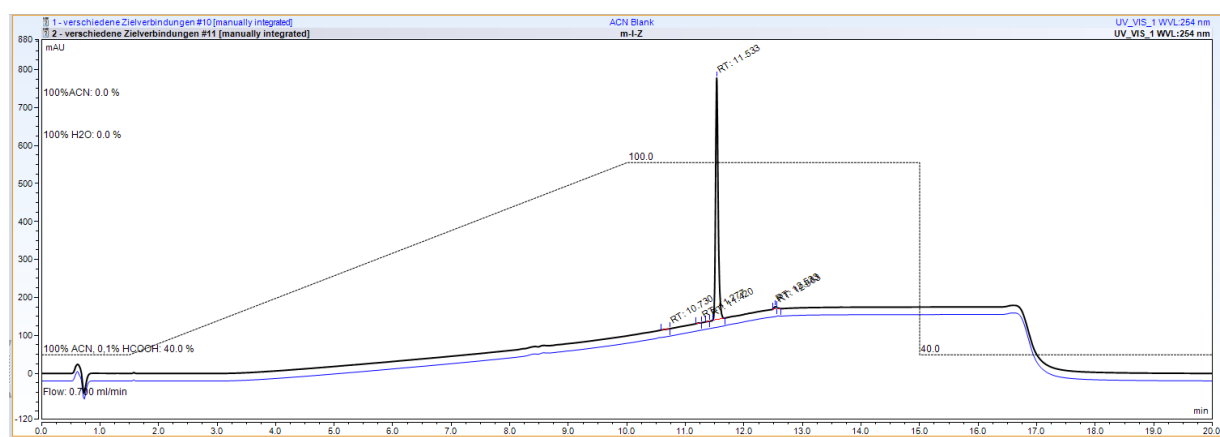


Figure S75. RP-HPLC chromatogram of blank (CH₃CN) and compound **3_m**, retention time: 11.5 min.

Compound **3_p**

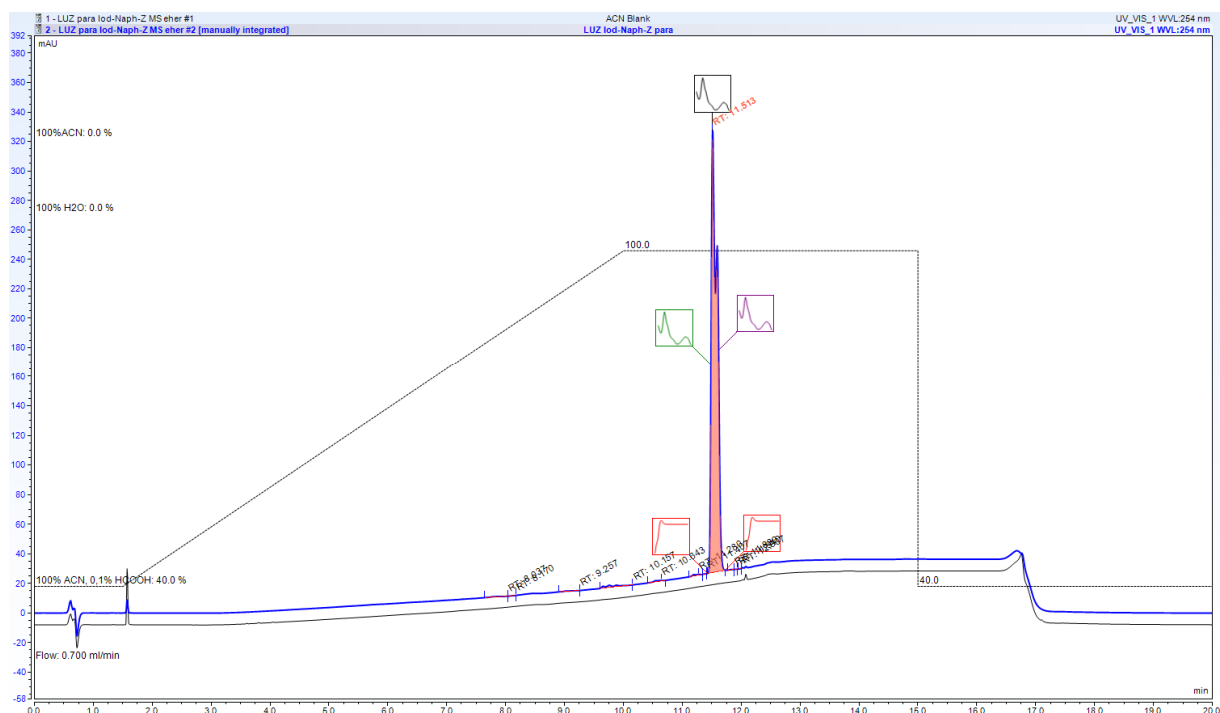


Figure S76. RP-HPLC chromatogram of blank (CH₃CN) and compound **3_p**, retention time: 11.5 min.

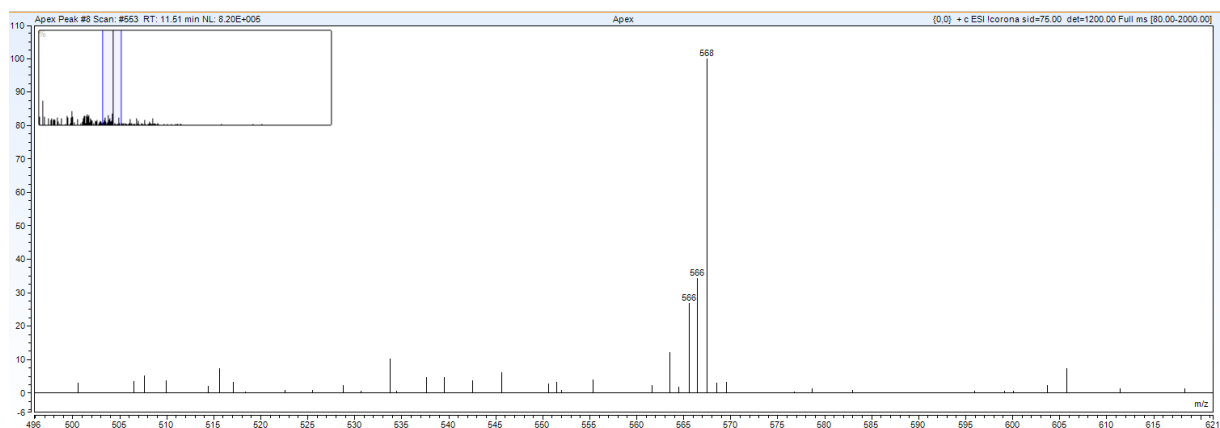


Figure S77. MS(+) of compound **3_p**, retention time: 11.5 min.

Compound 4

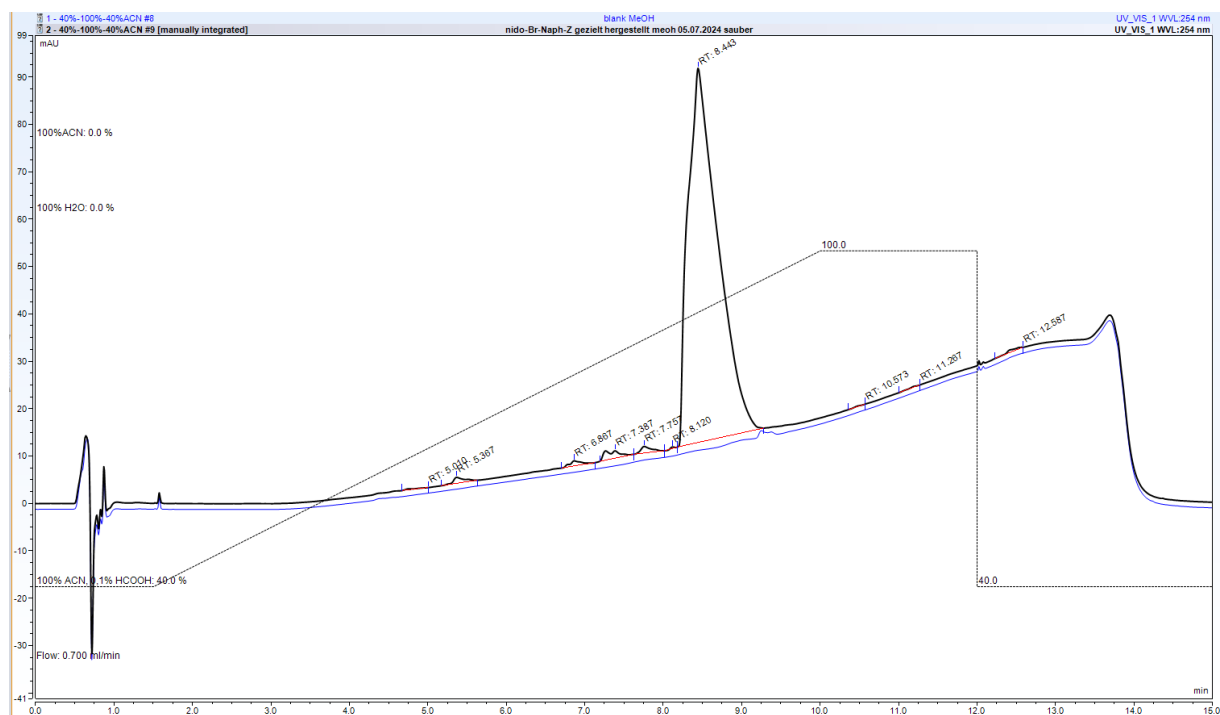


Figure S78. RP-HPLC chromatogram of blank (MeOH) and compound **4**, retention time: 8.4 min.

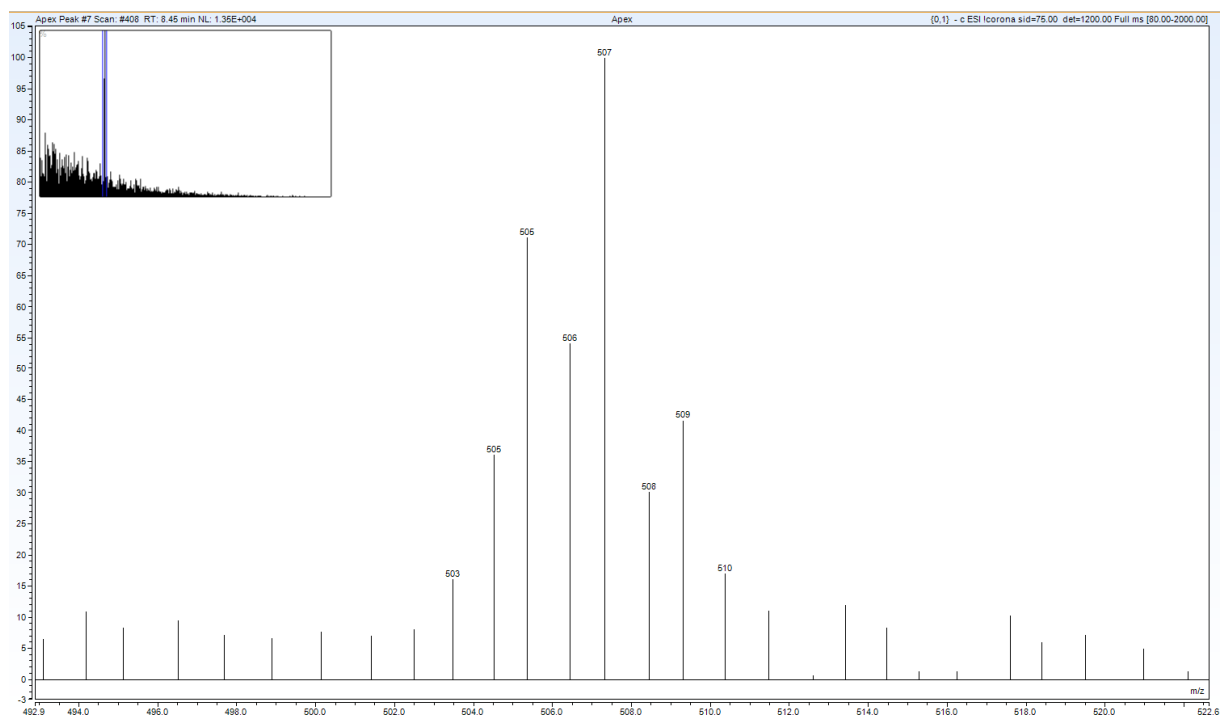


Figure S79. MS(-) of compound **4**, retention time: 8.45 min.

Compound 5

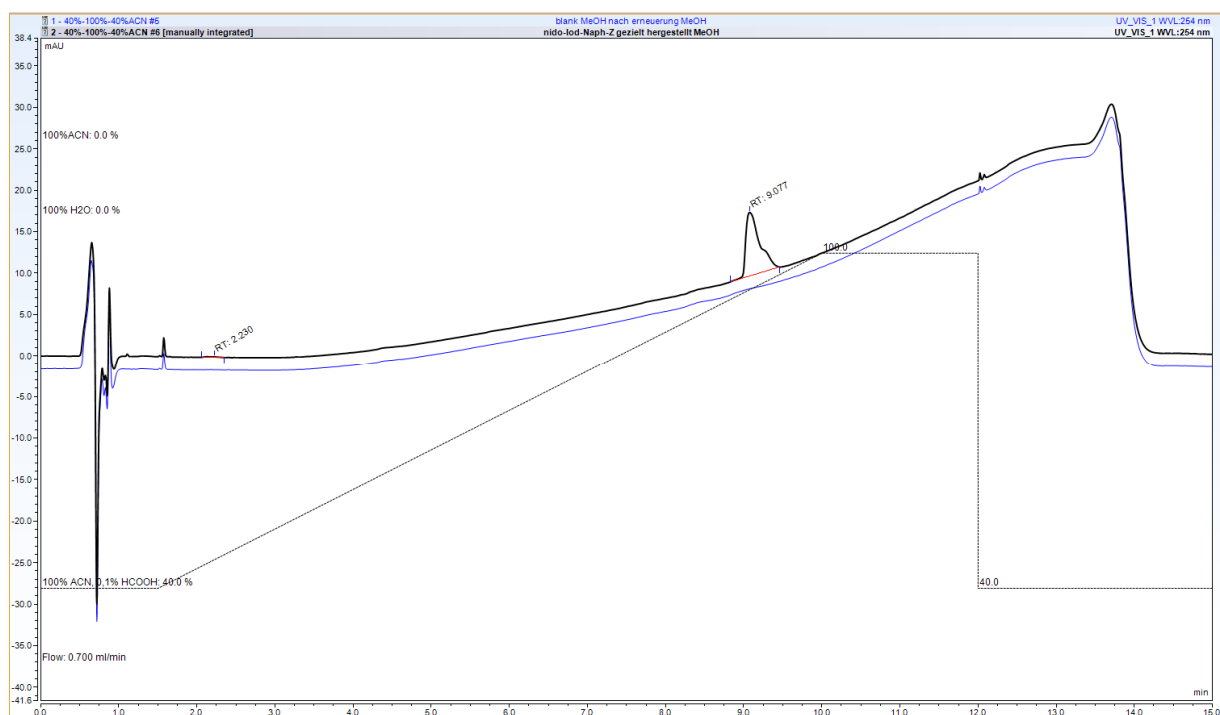


Figure S80. RP-HPLC chromatogram of blank (MeOH) and compound **5**, retention time: 9.1 min.

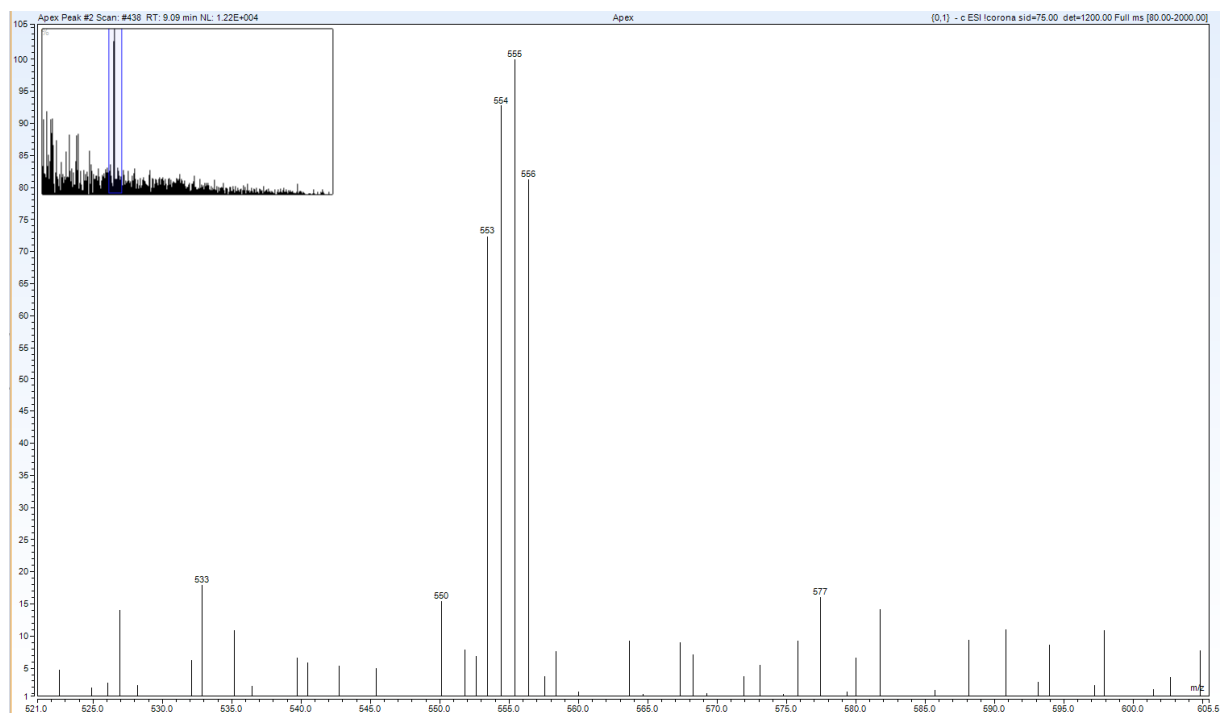


Figure S81. MS(-) of compound **5**, retention time: 9.1 min.

5 Determination of the Stability of Compounds 2_{o,m,p}, 3_{o,m,p}, 4, and 5 by HPLC

The stability measurements have been performed with an HPLC-UV-MS on an RP column. Blank samples of pure DMSO/H₂O (1:1, v/v) have been measured prior to the target compounds. The target compounds have been dissolved in DMSO/H₂O (1:1, v/v) for stability determination. The measurements have been started directly after the addition of the water. The double peaks or shoulders of the signals of the compound are related to the column that has been used. Broad signals have been observed in case of the *nido*-carborane derivatives.

Compound 2_o

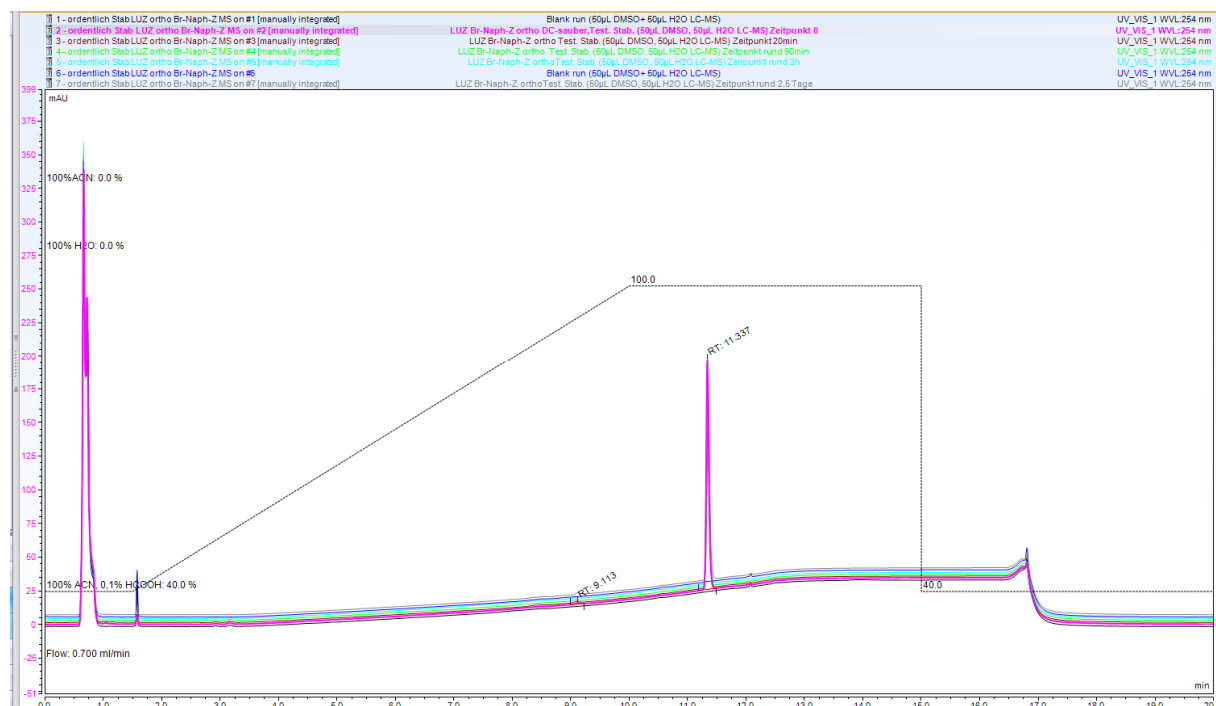


Figure S82. RP-HPLC chromatograms of blank (DMSO/H₂O, black and dark blue) and compound 2_o, retention time: 11.4 min, purity after 2.5 d: >98%.

Compound **2_m**

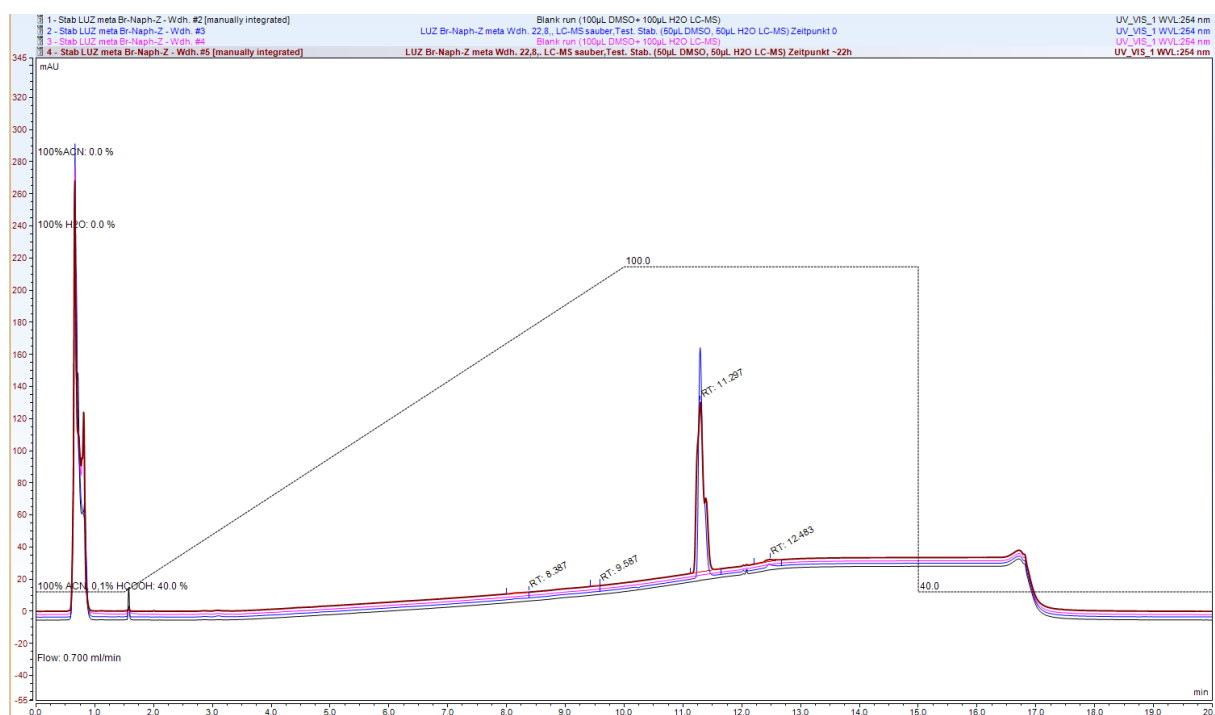


Figure S83. RP-HPLC chromatograms of blank (DMSO/H₂O, black, pink) and compound **2_m**, retention time: 11.3 min, purity after 1 d: ~98%.

Compound **2_p**

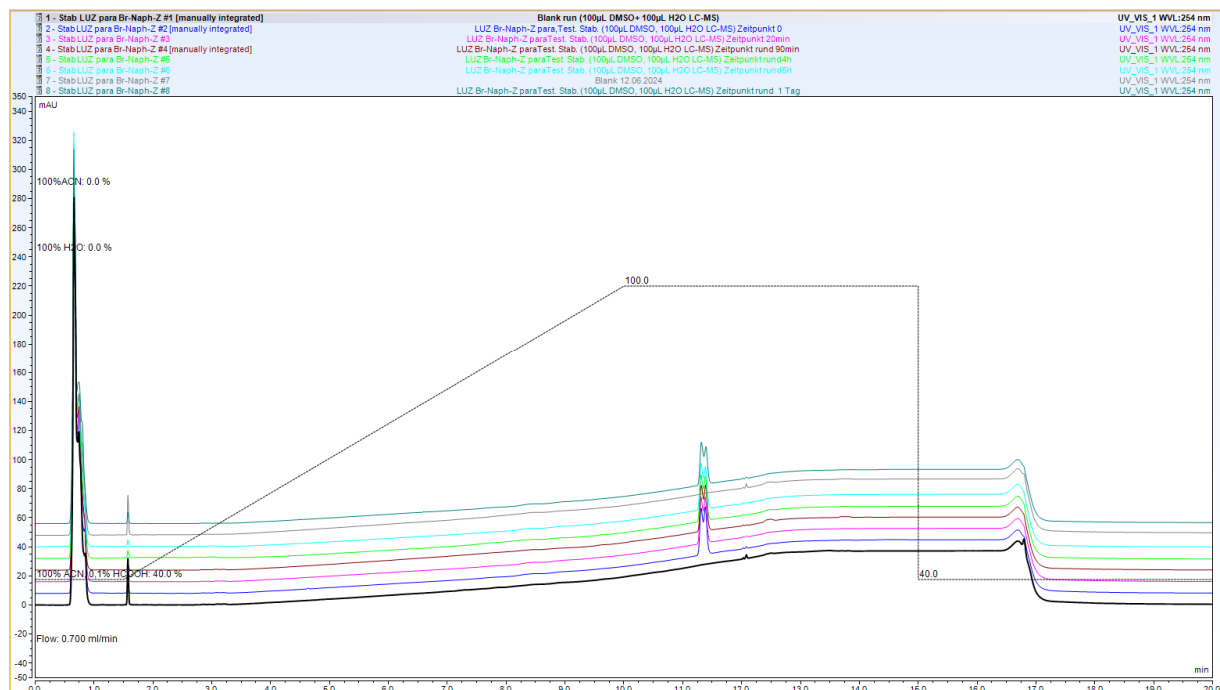


Figure S84. RP-HPLC chromatograms of blank (DMSO/H₂O, black, grey) and compound **2_p**, retention time: 11.4 min

Compound **3_o**

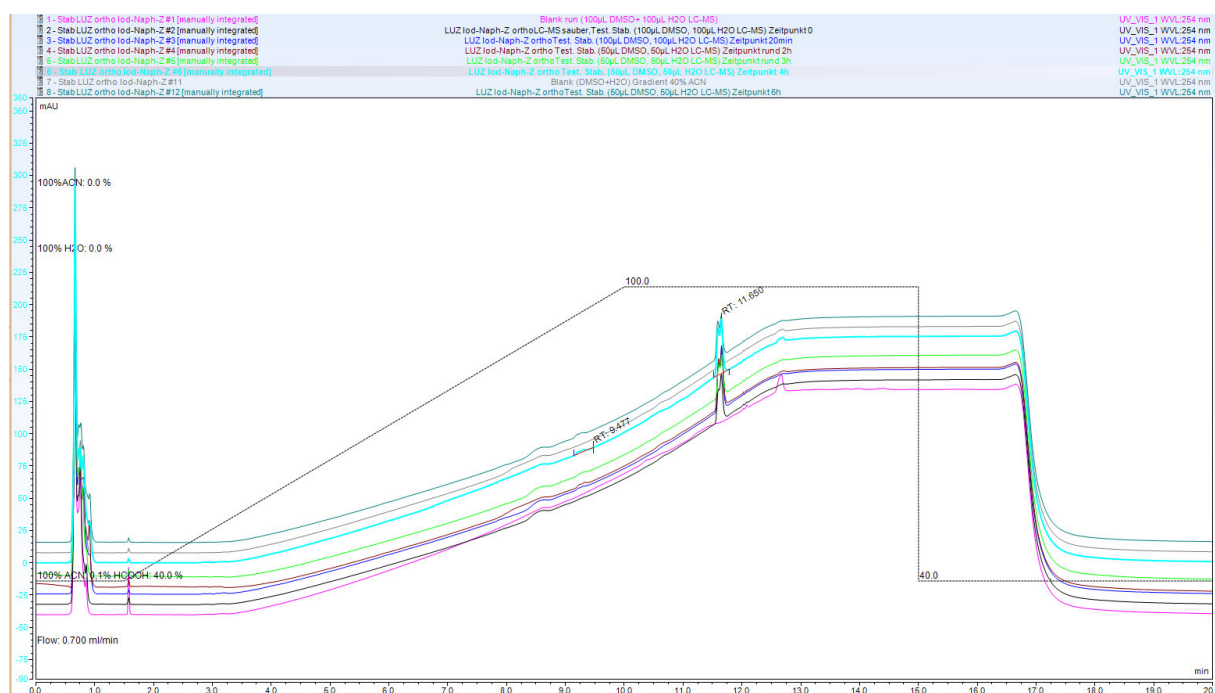


Figure S85. RP-HPLC chromatograms of blank (DMSO/H₂O, pink, grey) and compound **3_o**, retention time: 11.7 min, purity after 6 h: ~93%

Compound **3_m**

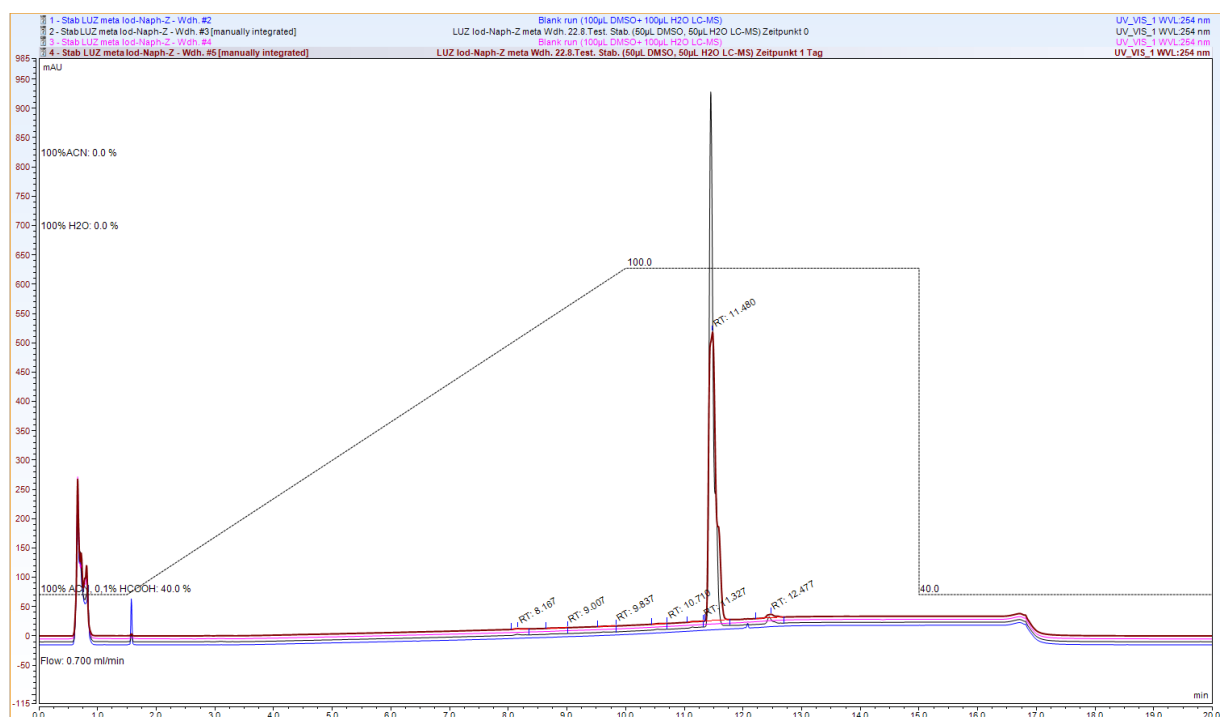


Figure S86. RP-HPLC chromatograms of blank (DMSO/H₂O, pink, blue) and compound **3_m**, retention time: 11.5 min, purity after 1 d: ~98%

Compound 3_p

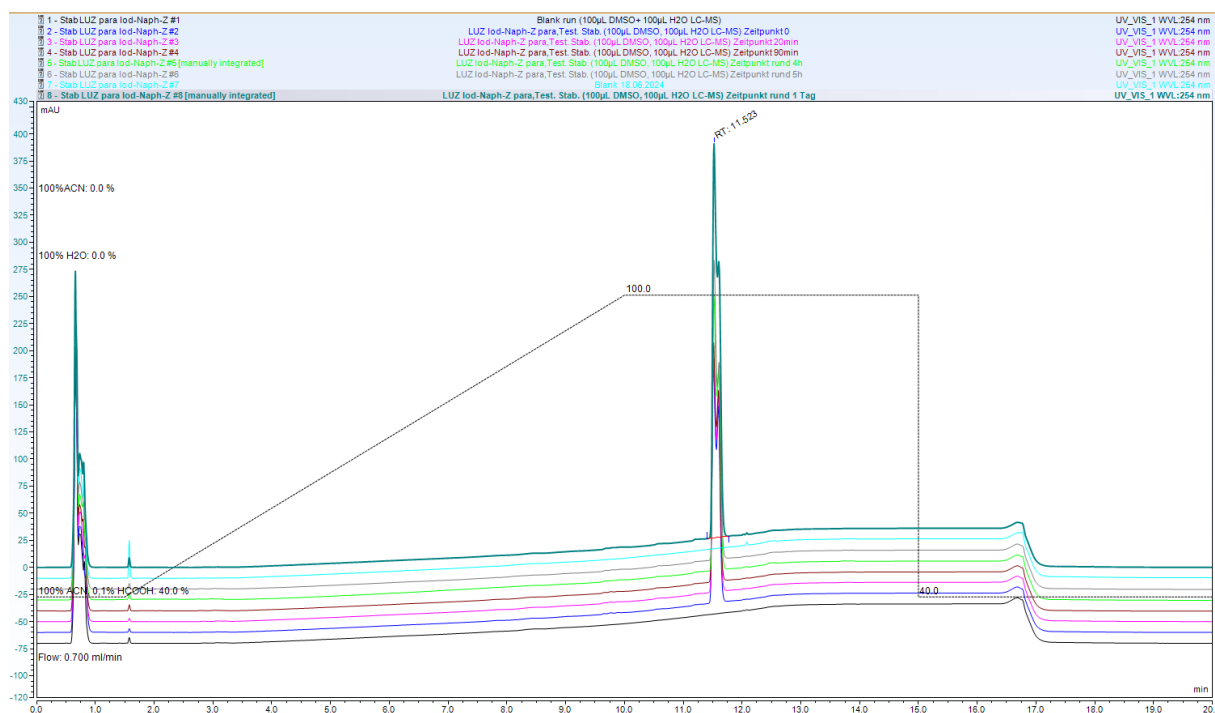


Figure S87. RP-HPLC chromatograms of blank (DMSO/H₂O, black, light blue) and compound 3_p, retention time: 11.5 min, purity after 1 d: >98%

Compound 4

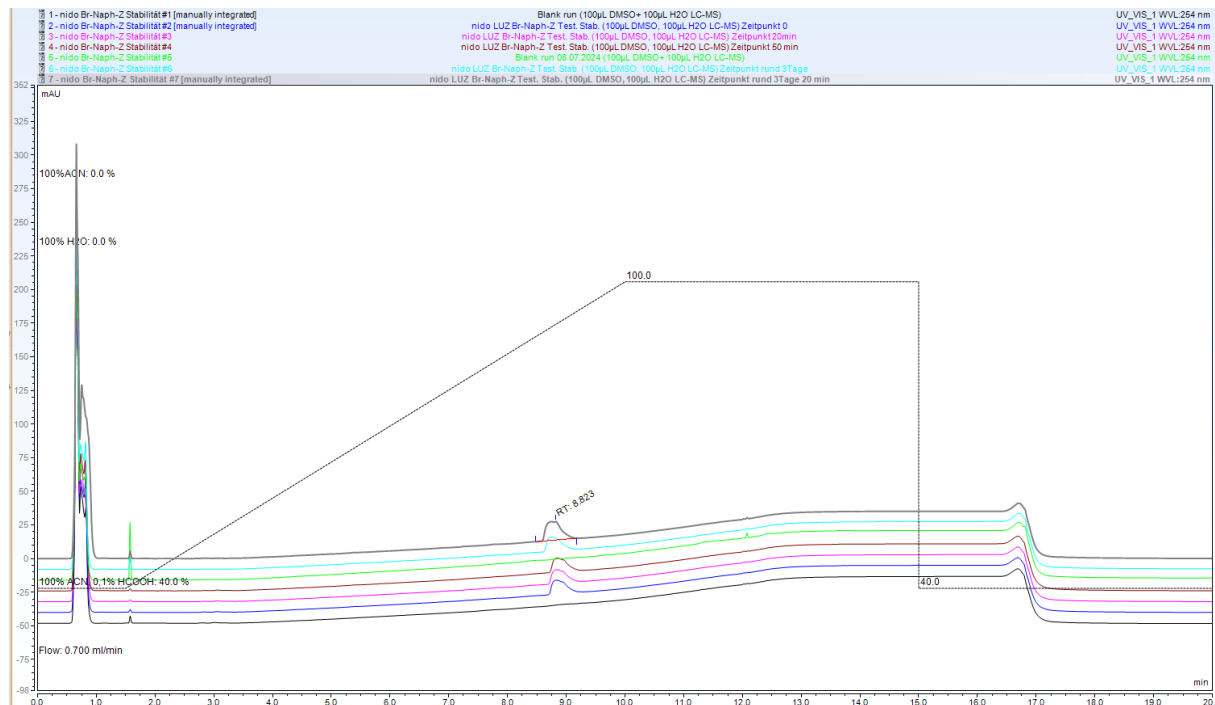


Figure S88. RP-HPLC chromatograms of blank (DMSO/H₂O, black, green) and compound 4, retention time: 8.8 min, purity after 3 d: ~98%

Compound 5

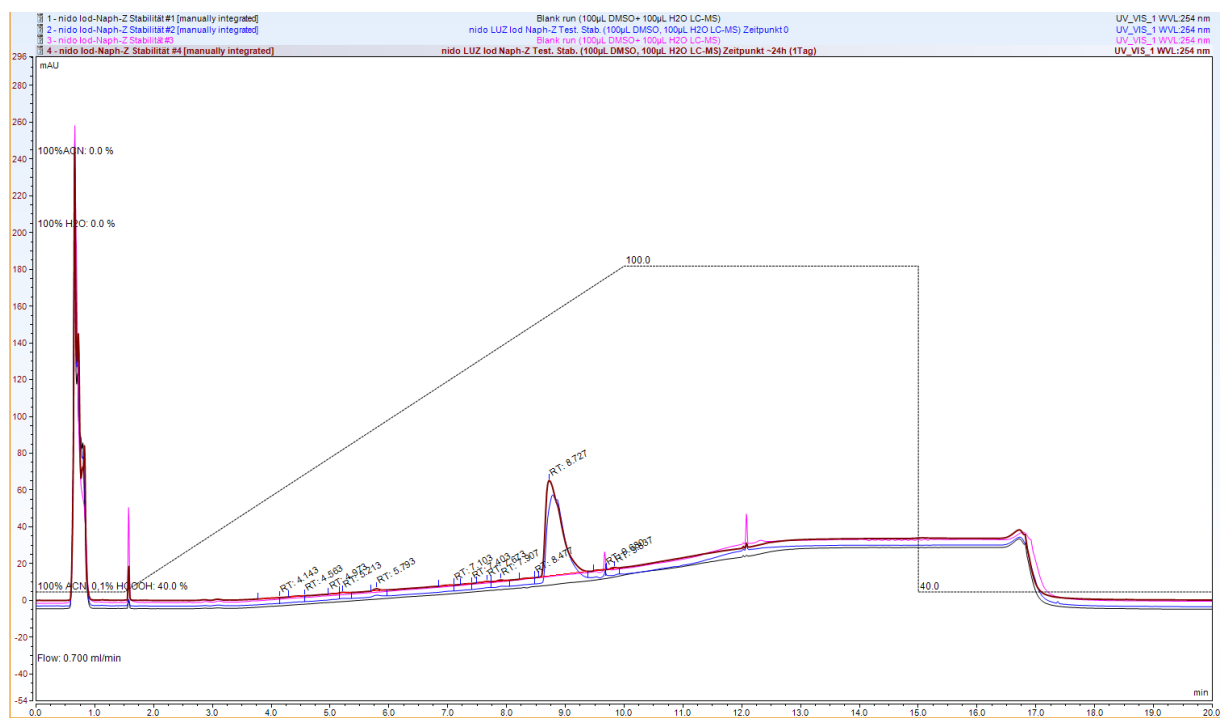


Figure S89. RP-HPLC chromatograms of blank (DMSO/H₂O, black, pink) and compound **5**, retention time: 8.7 min, purity after 1 d: >97%

6 X-ray Crystallography Data of Compounds **2_o**, **2_p**, **3_o**, and **3_m**

Table S1. Fundamental structure parameters of **2_o**, **2_p**, **3_o**, **3_m** at 190(2) K.

Compound	2_o	2_p	3_o	3_m
Empirical formula	C ₁₈ H ₂₁ B ₁₀ BrFN ₃ O ₂	C ₁₈ H ₂₁ B ₁₀ BrFN ₃ O ₂	C ₁₈ H ₂₁ B ₁₀ FIN ₃ O ₂	C ₁₈ H ₂₁ B ₁₀ FIN ₃ O ₂
Formula weight	518.39	518.39	565.38	565.38
Temperature [K]	190(2)	190(2)	190(2)	190(2)
Wavelength [pm]	71.073	71.073	71.073	71.073
Crystal system	Triclinic	Triclinic	Triclinic	Triclinic
Space group	<i>P</i> $\bar{1}$	<i>P</i> $\bar{1}$	<i>P</i> $\bar{1}$	<i>P</i> $\bar{1}$
Unit cell dimensions				
a [pm]	717.95(1)	716.05(2)	729.17(2)	729.57(2)
b [pm]	1262.93(3)	1259.81(5)	1261.01(4)	1251.29(4)
c [pm]	1441.12(3)	1438.06(5)	1438.32(4)	1435.06(4)
α [deg]	114.074(2)	114.635(3)	113.052(3)	112.654(3)
β [deg]	94.818(2)	95.020(2)	95.013(2)	94.816(2)
γ [deg]	99.527(2)	99.560(3)	100.414(3)	99.656(3)
Volume [nm ³]	1.15981(4)	1.14489(7)	1.17845(7)	1.17587(6)
Z	2	2	2	2
ρ (calculated) [Mg/m ³]	1.484	1.504	1.593	1.597
μ [mm ⁻¹]	1.804	1.828	1.390	1.393
F(000)	520	520	556	556
Crystal size [mm ³]	0.40 · 0.26 · 0.20	0.40 · 0.26 · 0.19	0.28 · 0.24 · 0.09	0.32 · 0.13 · 0.09
Θ_{Min} / Θ_{Max} [deg]	2.861 / 30.330	2.877 / 30.269	2.838 / 32.296	2.826 / 32.382
Index ranges	-10 ≤ h ≤ 10 -17 ≤ k ≤ 17 -20 ≤ l ≤ 19	-9 ≤ h ≤ 10 -17 ≤ k ≤ 17 -20 ≤ l ≤ 20	-10 ≤ h ≤ 10 -18 ≤ k ≤ 18 -20 ≤ l ≤ 21	-10 ≤ h ≤ 10 -18 ≤ k ≤ 18 -21 ≤ l ≤ 21
Reflections collected	30819	22051	21855	25123
Indp. reflections (<i>R</i> _{int})	6479 (0.0315)	6278 (0.0291)	7700 (0.0262)	7754 (0.0276)
Completeness (Θ_{Max}) [deg]	99.9 % (28.285)	99.9 % (28.285)	99.9 % (30.510)	100.0 % (30.510)
<i>T</i> _{Max} / <i>T</i> _{Min}	1.00000 / 0.93618	1.00000 / 0.88629	1.00000 / 0.98898	1.00000 / 0.91640
Restraints / Gof on <i>F</i> ²	0 / 400 1.077	0 / 400 1.030	0 / 400 1.019	0 / 400 1.041
<i>R</i> 1 / <i>wR</i> 2 (<i>I</i> > 2σ(<i>I</i>))	0.0343 / 0.0783	0.0322 / 0.0758	0.0298 / 0.0631	0.0300 / 0.0625
<i>R</i> 1 / <i>wR</i> 2 (all data)	0.0537 / 0.0867	0.0438 / 0.0819	0.0409 / 0.0679	0.0414 / 0.0675
Residual electron density [e·Å ⁻³]	0.324 / -0.416	0.416 / -0.534	0.462 / -0.556	0.616 / -0.622
Comments	† ¹	-	-	-
CCDC No ^[103]	2432537	2432538	2432539	2432540

†¹: The crystals always tend to crack at temperatures below 180 K. For this reason, all samples had been recorded at 190 K.

Table S2. Fundamental structure parameters of **2_o**, **2_p** at 297(2) K.

Compound	2_o	2_p
Empirical formula	C ₁₈ H ₂₁ B ₁₀ BrFN ₃ O ₂	C ₁₈ H ₂₁ B ₁₀ BrFN ₃ O ₂
Formula weight	518.39	518.39
Temperature [K]	297(2)	297(2)
Wavelength [pm]	71.073	71.073
Crystal system	Triclinic	Triclinic
Space group	<i>P</i> $\bar{1}$	<i>P</i> $\bar{1}$
Unit cell dimensions		
a [pm]	715.15(2)	716.53(1)
b [pm]	1268.90(6)	1267.64(3)
c [pm]	1457.16(4)	1447.18(3)
α [deg]	113.502(3)	114.284(2)
β [deg]	95.248(2)	95.478(2)
γ [deg]	98.669(3)	98.721(2)
Volume [nm ³]	1.18189(8)	1.16601(4)
Z	2	2
ρ _(calculated) [Mg/m ³]	1.457	1.476
μ [mm ⁻¹]	1.770	1.794
F(000)	520	520
Crystal size [mm ³]	0.35 · 0.24 · 0.16	0.41 · 0.27 · 0.17
Θ_{Min} / Θ_{Max} [deg]	2.812/ 29.197	2.856/ 28.703
Index ranges	-9 ≤ h ≤ 9 -16 ≤ k ≤ 17 -19 ≤ l ≤ 19	-9 ≤ h ≤ 9 -16 ≤ k ≤ 16 -19 ≤ l ≤ 19
Reflections collected	27843	27492
Indp. reflections (R_{int})	5745 (0.0292)	5440 (0.0278)
Completeness (Θ_{Max}) [deg]	99.9 % (26.375)	99.9 % (26.375)
T_{Max} / T_{Min}	1.00000 / 0.94443	1.00000 / 0.91888
Restraints / parameters	421 / 429	0 / 400
Gof on F ²	1.017	1.044
R1 / wR2 ($I > 2\sigma(I)$)	0.0373 / 0.0839	0.0375 / 0.0904
R1 / wR2 (all data)	0.0685 / 0.0973	0.0565 / 0.1004
Residual electron density [e·Å ⁻³]	0.208 / -0.510	0.474 / -0.490
Comments	† ²	-
CCDC No ^[103]	2432541	2432542

†²: The *ortho*-carborane unit is disordered on two positions with a ratio of 0.67(1):0.33(1).

7 Docking Studies of Compounds **2_{o,m,p}**, **3_{o,m,p}**, **4**, and **5**

Molecular docking was performed using AutoDockTools4^[104] with the Lamarckian Genetic Algorithm.^[105] The force-field parameters for boron atoms were manually added to the AutoDockTools4 parameter file.

The protein structure (PDB ID: 5ZTY)^[9] was obtained from the Protein Data Bank (PDB) and originally contained the ligand *N*-(adamantan-1-yl)-1-(5-hydroxypentyl)-4-methyl-5-phenyl-1*H*-pyrazole-3-carboxamide. The docking site was selected based on the position of this ligand within the binding pocket. Prior to docking, the ligand and water molecules were removed, and the protein structure was protonated using the Reduce software.^[106]

For docking, water molecules were eliminated, and non-polar hydrogen atoms were merged. The docking grid box was set with the parameters below:

58 x 44 x 54 center at (7.24, 1.919, -60.355) for **2_o**, **2_m**, **2_p**, **4**,

9.441 0.12 -59.514 center at (8.982 1.344 -54.339) for **3_o**, **3_m**, **3_p**, **5**,

aligning with the ligand-binding domain (LBD).

The following docking parameters were used:

- Number of hybrid GA-LS runs: 100
- Population size: 150
- Maximum number of energy evaluations: 25,000,000
- Top individuals surviving to the next generation: 1
- Gene mutation rate: 0.02
- Crossover rate: 0.8
- Mean of Cauchy distribution for gene mutation: 0.0
- Variance of Cauchy distribution for gene mutation: 1.0

8 Chemical Structures of WIN55212-2 and SR141716A

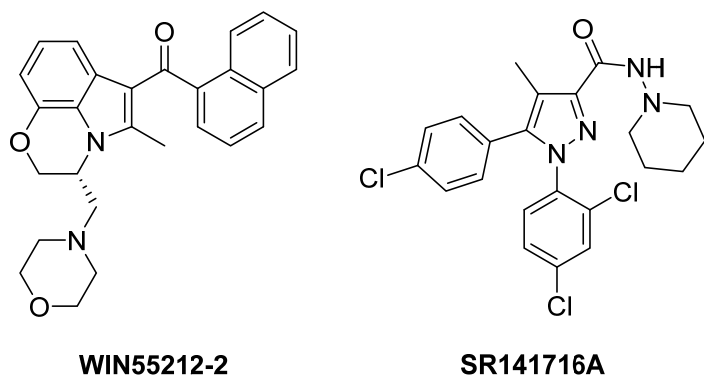


Figure S90. Chemical Structures of CB₂R agonist **WIN55212-2** and CB₂R antagonist/inverse agonist **SR141716A**.

- [1] V. Lucchesi, D.P. Hurst, D.M. Shore, S. Bertini, B.M. Ehrmann, M. Allarà, L. Lawrence, A. Ligresti, F. Minutolo, G. Saccomanni, H. Sharir, M. Macchia, V. Di Marzo, M.E. Abood, P.H. Reggio, C. Manera, *J. Med. Chem.* **2014**, *57*, 8777–8791.
- [9] X. Li, T. Hua, K. Vemuri, J.-H. Ho, Y. Wu, L. Wu, P. Popov, O. Benchama, N. Zvonok, K. Locke, L. Qu, G.W. Han, M.R. Iyer, R. Cinar, N.J. Coffey, J. Wang, M. Wu, V. Katritch, S. Zhao, G. Kunos, L.M. Bohn, A. Makriyannis, R.C. Stevens, Z.-J. Liu, *Cell* **2019**, *176*, 459-467.e13.
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