

**Oxazoline amino acid bioconjugates:
One-pot synthesis and analysis of supramolecular interactions**

Marija Bakija,^a Berislav Perić^a and Srećko I. Kirin^{a,*}

^a Ruđer Bošković Institute, Bijenička c. 54, HR-10000 Zagreb, Croatia

* Author to whom the correspondence should be addressed. (E-mail: Srecko.Kirin@irb.hr)

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1. Compounds overview, syntheses schemes and spectroscopic characterization data

Chart S1. Bis-amino acids **2**, mixed derivatives **3** and bis-amino alcohols **4** derived from isophthalic (**m**) and terephthalic (**p**) acid synthesized in this paper, using achiral amino alcohols and chiral amino acids.

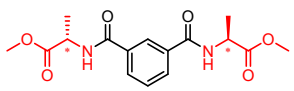
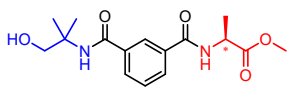
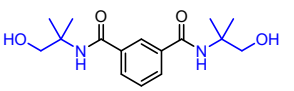
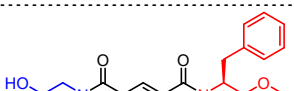
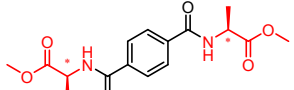
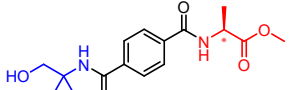
	2	3	4
m1	 2 _{m1}	 3 _{m1}	 4 _{m1}
m2	not isolated	 3 _{m2}	not isolated
p1	 2 _p	 3 _p	not isolated

Chart S2. Mixed derivatives **3** and bis-amino alcohols **4** derived from isophthalic (**m**) acid synthesized in this paper, using chiral amino alcohols and chiral amino acids.

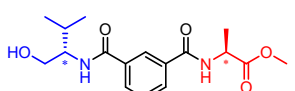
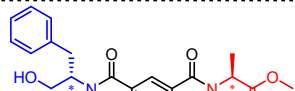
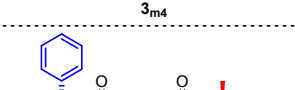
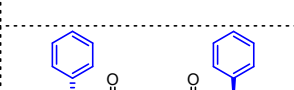
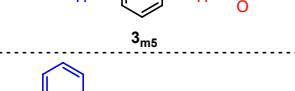
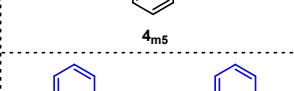
	2	3	4
m3	not isolated	 3 _{m3}	not isolated
m4	not isolated	 3 _{m4}	not isolated
m5	not isolated	 3 _{m5}	 4 _{m5}
m6	not isolated	 3 _{m6}	 4 _{m5}

Chart S3. Bis-amino acids **2**, mixed derivatives **3** and bis-amino alcohols **4** derived from 1,4-naphthalic (**n₁**), 1,5-naphthalic (**n₃**), 2,6-naphthalic (**n₄**), 2,7-naphthalic (**n₅**) or 9,10-anthracene (**a₁**) diacids, using achiral amino alcohols, achiral (**n₂**) and chiral amino acids synthesized in this paper.

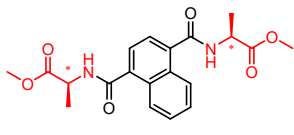
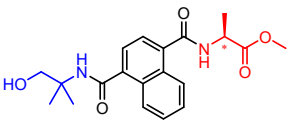
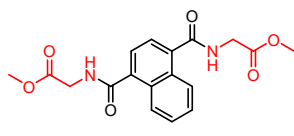
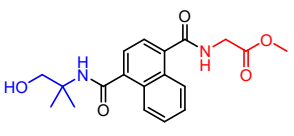
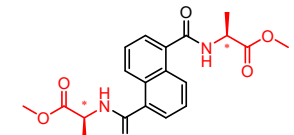
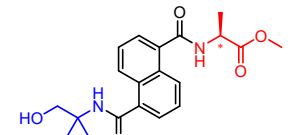
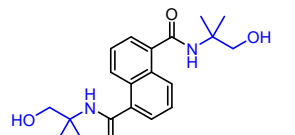
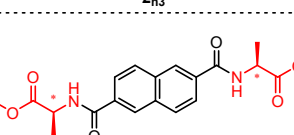
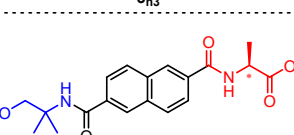
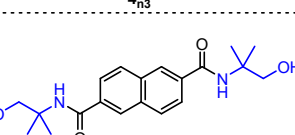
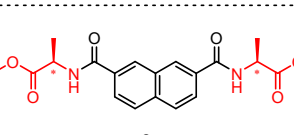
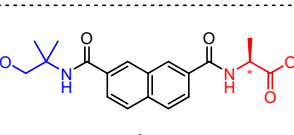
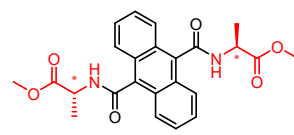
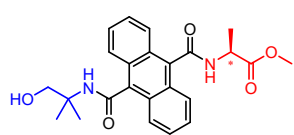
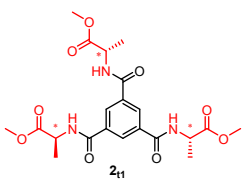
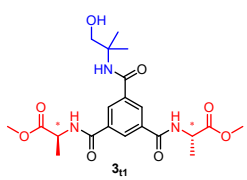
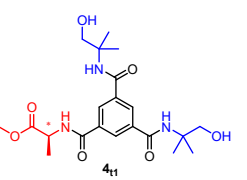
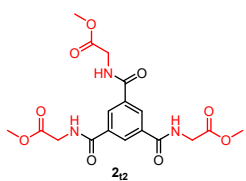
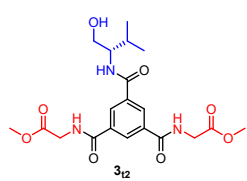
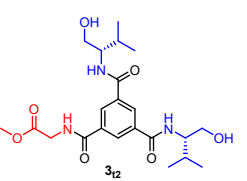
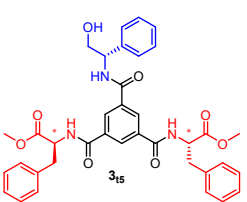
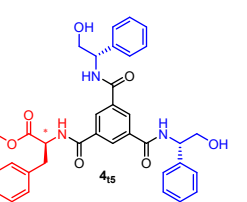
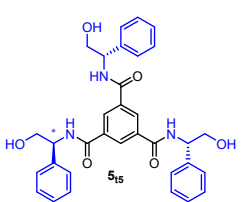
	2	3	4
n₁	 2 _{n1}	 3 _{n1}	not isolated
n₂	 2 _{n2}	 3 _{n2}	not isolated
n₃	 2 _{n3}	 3 _{n3}	 4 _{n3}
n₄	 2 _{n4}	 3 _{n4}	 4 _{n4}
n₅	 2 _{n5}	 3 _{n5}	not isolated
a₁	 2 _a	 3 _a	not isolated

Chart S4. Tris-amino acids **2**, mixed derivatives **3**, bis-amino alcohols **4** and tris-amino alcohols derived from trimesic acid (**t**), using achiral and chiral amino alcohols, as well as achiral and chiral amino acids synthesized in this paper.

	2	3	4	5
t1				not isolated
t2				not isolated
t5	not isolated			

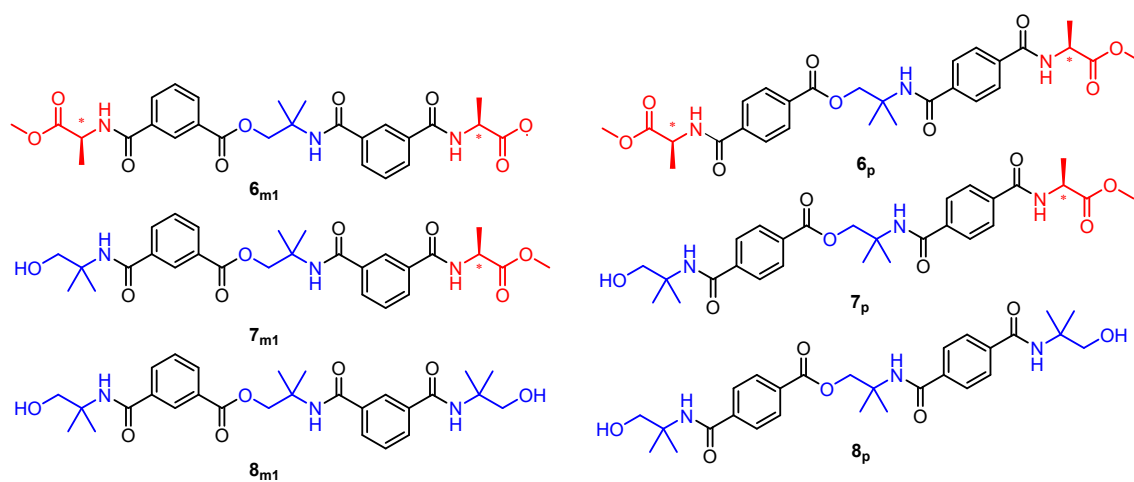
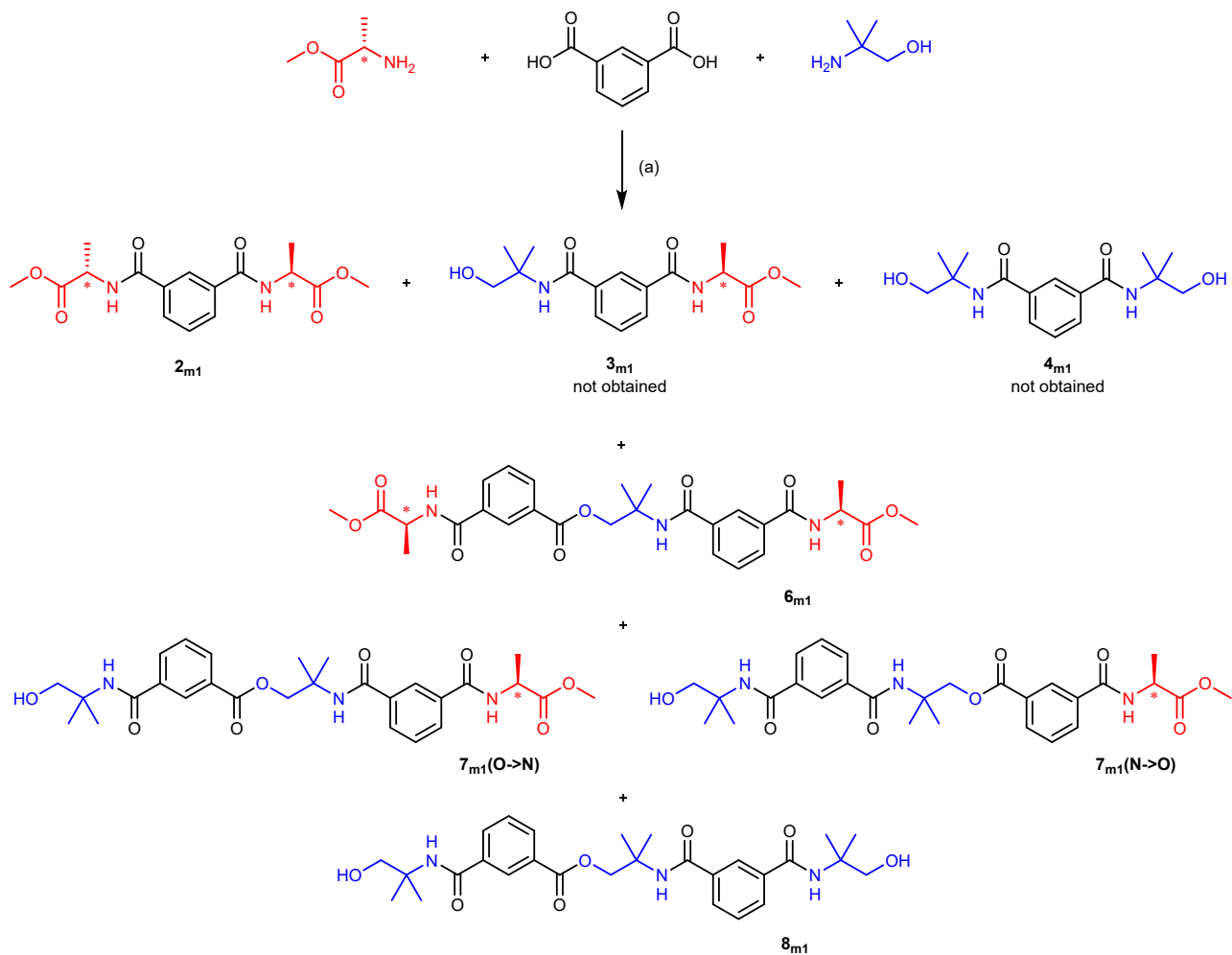
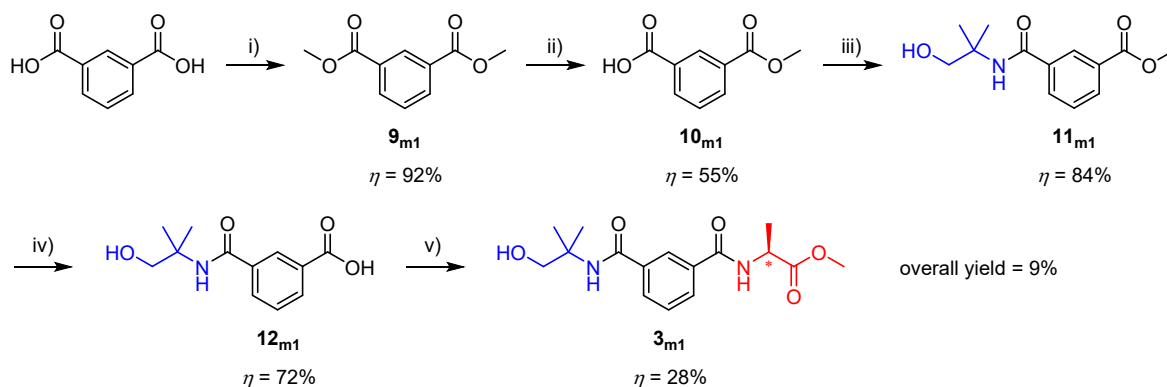


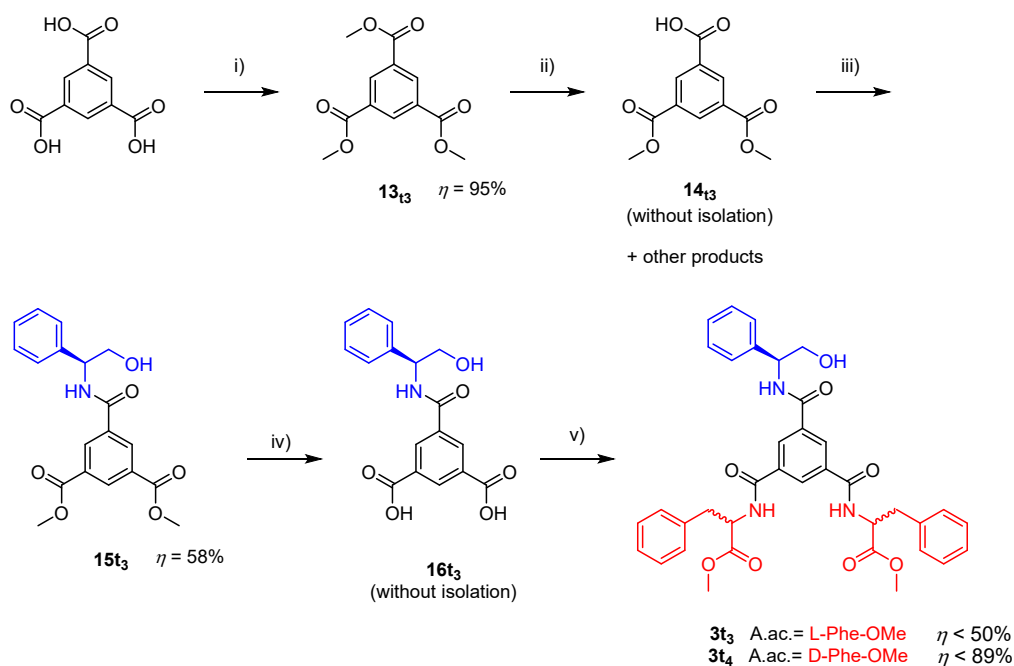
Chart S5. Ester-amide byproducts obtained in reaction 1 and 7 (tables 1 and 2 respectively).



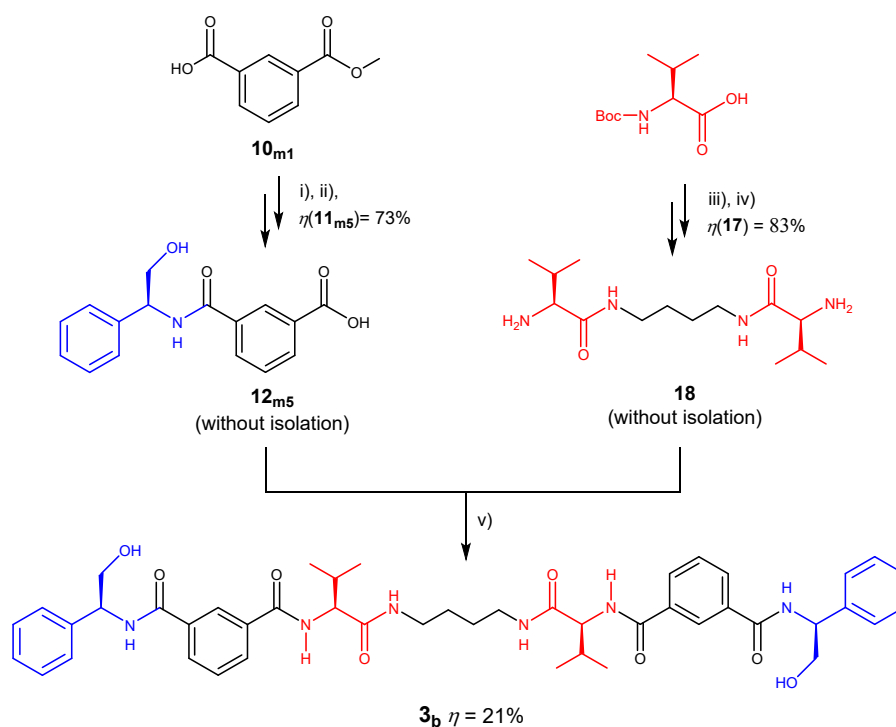
Scheme S1. Reaction conditions (a). TBTU/HOBT, DIPEA, DCM, 2 days (Table 1, reaction 1).



Scheme S2. Reaction conditions. i) H₂SO₄ (conc.), MeOH, 75 °C, stirring overnight; ii) MeOH : H₂O (2 : 1), NaOH, microwave, 50 W, 150 °C, 20 min, HCl (conc.); iii) AMP, HATU, DIPEA, DCM, 1 day; iv) MeOH:H₂O (2:1), NaOH, microwave, 50 W, 150 °C, 20 min, HCl (conc.); v) H-Ala-OMe, HATU, DIPEA, DCM, 1 day.



Scheme S3. Reaction conditions. i) H_2SO_4 (conc.), MeOH, 75 °C, stirring overnight; ii) MeOH : H_2O (2 : 1), NaOH, microwave, 50 W, 150 °C, 20 min, HCl (conc.); iii) (S)-(+)-phenylglycinol, HATU, DIPEA, DCM, 1 day; iv) MeOH: H_2O (2:1), NaOH, microwave, 50 W, 150 °C, 20 min, HCl (conc.); v) H_2N -L-Ala-OMe, HATU, DIPEA, DCM, 1 day.



Scheme S4. Reaction conditions. i) Phg[#], TBTU/HOBt, DIPEA, DCM, 1 day; ii) MeOH: H_2O (2:1), NaOH, microwave, 50 W, 150 °C, 20 min, HCl (conc.); iii) Boc-Val-OH, 1,4-diaminobutane, TBTU/HOBt, DIPEA, DCM, 1 day; iv) TFA : DCM = 1 : 1, 2h, r.t., DIPEA; v) HATU, DIPEA, DCM, 1 day.

1.1. Spectroscopic characterization of compounds 2-15

Ala-*m*C₆H₄-Ala (2_{m1}). Reactions 1, 4 and 5. $M_r(C_{16}H_{20}N_2O_6) = 336.13$. ESI-MS (m/z): 337.3 ($M + H^+$), 673.6 ($2M + H^+$). Crystals suitable for single-crystal x-ray diffraction obtained from dichloromethane/water mixture after one month. ¹H NMR (600 MHz, CDCl₃) δ /ppm: 8.22 (s, 1H), 7.95 (d, $J = 7.7$ Hz, 2H), 7.52 (t, $J = 7.7$ Hz, 1H), 6.87 (d, $J = 7.0$ Hz, 2H), 4.82 (quin., $J = 7.2$ Hz, 2H), 3.80 (s, 6H), 1.54 (d, $J = 7.2$ Hz, 6H). ¹³C NMR (151 MHz, CDCl₃) δ /ppm: 172.88, 165.24, 133.20, 129.57, 128.06, 124.42, 51.75, 47.76, 17.33.

Ala-*m*C₆H₄-AMP (3_{m1}). Reactions 3, 4 and 5. $M_r(C_{16}H_{20}N_2O_6) = 322.15$. ESI-MS (m/z): 323.1 ($M + H^+$), 645.2 ($2M + H^+$), 667.2 ($2M + Na^+$). ¹H NMR (600 MHz, CDCl₃) δ /ppm: 8.11 (s, 1H), 7.87 (d, $J = 7.8$ Hz, 2H), 7.45 (t, $J = 7.7$ Hz, 1H), 7.04 (d, $J = 7.1$ Hz, 1H), 6.38 (s, 1H), 4.80 (quin., $J = 7.2$ Hz, 1H), 4.48 (t, $J = 6.2$ Hz, 1H), 3.81 (s, 3H), 3.76 – 3.67 (m, 2H), 1.53 (d, $J = 7.2$ Hz, 3H), 1.43 (s, 5H). ¹³C NMR (151 MHz, CDCl₃) δ /ppm: 174.11, 167.64, 166.15, 135.54, 133.75, 130.55, 130.07, 129.01, 125.34, 70.31, 56.76, 52.84, 48.79, 24.63, 24.59, 18.22.

AMP-*m*C₆H₄-AMP (4_{m1}). Reaction 6. ¹H NMR (600 MHz, DMSO) δ /ppm: 8.46 – 8.41 (m, 1H), 7.89 (dd, $J = 7.5$, 1.6 Hz, 3H), 7.29 (t, $J = 7.5$ Hz, 2H), 3.34 (s, 4H), 1.15 (s, 13H). ¹³C NMR (151 MHz, DMSO) δ /ppm: 170.24, 137.45, 130.39, 130.05, 126.56, 67.43, 53.58, 23.19. ESI-MS spectrum was recorded, but the results suggested there was a problem with recording this compound.

Ala-*m*C₆H₄-AMP-*m*C₆H₄-Ala (6_{m1}). Reaction 1. $M_r(C_{40}H_{46}N_4O_{12}) = 555.22$. ESI-MS (m/z): 556.4 ($M + H^+$), 1111.9 ($2M + H^+$). ¹H NMR (600 MHz, CDCl₃) δ /ppm: 8.45 (s, 1H), 8.19 – 8.08 (m, 2H), 8.03 – 7.96 (m, 1H), 7.90 – 7.80 (m, 2H), 7.48 (t, $J = 7.8$ Hz, 1H), 7.42 (t, $J = 7.7$ Hz, 1H), 7.07 – 6.96 (m, 2H), 6.64 (s, 1H), 4.82 – 4.71 (m, 2H), 4.57 (s, 2H), 3.76 (t, $J = 5.0$ Hz, 6H), 1.58 (d, $J = 3.3$ Hz, 6H), 1.52 – 1.46 (m, 6H). ¹³C NMR (151 MHz, CDCl₃) δ /ppm: 173.70, 173.66, 166.56, 166.28, 166.24, 165.91, 132.75, 132.14, 130.3, 130.33, 130.15, 129.09, 129.08, 128.31, 125.37, 70.44, 54.63, 52.73, 48.75, 48.73, 24.32, 24.27, 18.43.

AMP-*m*C₆H₄-AMP-*m*C₆H₄-Ala (7_{m1}). Reaction 1. $M_r(C_{28}H_{35}N_3O_8) = 541.24$. ESI-MS (m/z): 542.4 ($M + H^+$), 1105.9 ($2M + Na^+$). ¹H NMR (600 MHz, CDCl₃) δ /ppm: 8.48 (s, 1H), 8.18 (d, $J = 7.8$ Hz, 1H), 8.07 (s, 1H), 8.02 (d, $J = 7.8$ Hz, 1H), 7.90 – 7.86 (m, 2H), 7.56 – 7.47 (m, 2H), 6.93 (d, $J = 7.2$ Hz, 1H), 6.66 (s, 1H), 6.38 (s, 1H), 4.79 (quin., $J = 7.2$ Hz, 1H), 4.58 (s, 2H), 3.79 (s, 3H), 3.70 (s, 2H), 1.61 – 1.59 (m, 6H).

AMP-*m*C₆H₄-AMP-*m*C₆H₄-AMP (8_{m1}). Reaction 1. $M_r(C_{28}H_{37}N_3O_7) = 527.26$. ESI-MS (m/z): 528.4 ($M + H^+$), 1077.9 ($2M + Na^+$). ¹H NMR (600 MHz, CDCl₃) δ /ppm: 8.33 (s, 1H), 8.08 (d, $J = 7.7$ Hz, 1H), 8.00 (s, 1H), 7.90 (d, $J = 7.8$ Hz, 1H), 7.78 (d, $J = 7.7$ Hz, 1H), 7.73 (d, $J = 7.7$ Hz, 1H), 7.43 (t, $J = 7.7$ Hz, 1H), 7.34 (t, $J = 7.7$ Hz, 1H), 6.76 (s, 1H), 6.60 (d, $J = 2.2$ Hz, 2H), 4.68 (s, 2H), 4.53 (s, 2H), 3.66 – 3.60 (m, 4H), 1.55 (s, 6H), 1.39 – 1.36 (m, $J = 4.0$ Hz, 12H).

Ala-*p*C₆H₄-Ala (2_p). Reaction 7. $M_r(C_{16}H_{20}N_2O_6) = 336.13$. ESI-MS (m/z): 337.1 ($M + H^+$, 55%) ¹H NMR (300 MHz, CDCl₃) δ /ppm: 7.86 (s, 4H), 6.87 (d, $J = 7.1$ Hz, 2H), 4.80 (quin., $J = 7.2$ Hz, 2H), 3.80 (s, 6H), 1.54 (d, $J = 7.2$ Hz, 6H). ¹³C NMR (151 MHz, CD₃OD) δ /ppm: 174.71, 169.18, 138.04, 128.70, 52.82, 50.17, 38.88, 17.19.

AMP-*p*C₆H₄-Ala (3_n). Reaction 7. $M_r(C_{16}H_{20}N_2O_6) = 322.15$. ESI-MS (m/z) obtained from 3_{p1} and 7_{p1} mixture: 323.1 ($M + H^+$, 33%), 345.1 ($M + Na^+$, 78%), 667.3 ($2M + Na^+$, 32%). ¹H NMR (300 MHz, CDCl₃) δ /ppm obtained from 3_{n1} and 7_{n1} mixture: 7.95 – 7.60 (m, 4H), 6.83 (d, 1H), 6.29 (s, 1H), 4.80 (quin., 1H), 4.34 (s, 1H), 3.81 (s, 3H), 3.71 (s, 2H), 1.54 (d, 3H), 1.44 (s, 6H). ¹³C NMR (75 MHz, CD₃OD) δ /ppm obtained

from **3_{p1}** and **7_{p1}** mixture: 174.72, 169.20, 166.91, 137.56, 134.07, 130.62, 128.77, 128.61, 128.52, 70.11, 55.17, 52.82, 50.17, 24.63, 17.19.

AMP-*p*C₆H₄-AMP-*p*C₆H₄-Ala (7_n**).** Reaction 7. $M_r(C_{78}H_{35}N_3O_8) = 541.24$. ESI-MS (m/z) obtained from **3_p** and **6_p** mixture: 564.2 (M + Na⁺), 1105.5 (2M + Na⁺, 5%). ¹H NMR (300 MHz, CDCl₃) δ /ppm obtained from **3_{n1}** and **7_{n1}** mixture: 8.13 – 8.00 (m, 2H), 7.95 – 7.60 (m, 6H), 6.83 (s, 1H), 6.53 (d, 1H), 6.29 (s, 1H), 4.80 (quin., 1H), 4.59 (s, 2H), 4.34 (s, 1H), 3.81 (s, 3H), 3.71 (s, 2H), 1.60 (s, 6H), 1.54 (d, 3H), 1.44 (s, 6H).

ETA-*m*C₆H₄-Phe (3_{m2}**).** Reaction 8. ¹H NMR (600 MHz, CDCl₃) δ /ppm: 8.09 (t, $J = 1.9$ Hz, 1H), 7.91 (ddd, $J = 7.7, 1.8, 1.1$ Hz, 1H), 7.74 – 7.69 (m, 1H), 7.40 (t, $J = 7.7$ Hz, 1H), 7.34 – 7.24 (m, 3H), 7.20 – 7.12 (m, 2H), 6.92 (d, $J = 7.3$ Hz, 2H), 5.07 (dt, $J = 7.6, 6.0$ Hz, 1H), 3.84 (dd, $J = 5.9, 2.7$ Hz, 2H), 3.79 (s, 3H), 3.67 – 3.59 (m, 2H), 3.49 (t, $J = 1.9$ Hz, 2H), 3.26 (ddd, $J = 54.3, 13.9, 6.0$ Hz, 2H), 2.87 (s, 1H). ¹³C NMR (151 MHz, CD₃OD) δ /ppm: 173.55, 169.61, 169.41, 138.43, 136.19, 135.64, 131.43, 131.29, 130.22, 129.81, 129.54, 127.90, 127.46, 61.55, 55.96, 52.79, 43.63, 38.18.

Val[#]-*m*C₆H₄-Ala (3_{m3}**).** Reaction 9. ¹H NMR (300 MHz, CDCl₃) δ /ppm: 8.15 (d, $J = 1.9$ Hz, 1H), 7.83 (dd, $J = 7.8, 1.7$ Hz, 2H), 7.40 (t, $J = 7.7$ Hz, 1H), 7.19 (d, $J = 7.4$ Hz, 1H), 6.78 (d, $J = 8.8$ Hz, 1H), 4.80 (s, 1H), 4.09 – 3.94 (m, 1H), 3.81 (s, 5H), 2.96 (t, $J = 5.6$ Hz, 1H), 2.00 (dt, $J = 13.8, 7.0$ Hz, 1H), 1.55 (d, $J = 7.2$ Hz, 3H), 1.02 (dd, $J = 6.8, 4.5$ Hz, 6H).

Phe[#]-*m*C₆H₄-Ala (3_{m4}**).** Reaction 10. ¹H NMR (300 MHz, CDCl₃) δ /ppm: 8.07 (t, $J = 1.8$ Hz, 1H), 7.78 (ddt, $J = 26.5, 7.8, 1.4$ Hz, 2H), 7.46 – 7.19 (m, 5H), 7.12 (d, $J = 7.4$ Hz, 1H), 6.79 (d, $J = 7.9$ Hz, 1H), 4.81 (t, $J = 7.3$ Hz, 1H), 4.42 (dd, $J = 11.0, 5.9$ Hz, 1H), 3.89 – 3.64 (m, 5H), 3.08 – 2.95 (m, 2H), 1.54 (d, $J = 7.2$ Hz, 3H).

Phg[#]-*m*C₆H₄-Ala (3_{m5}**).** Reaction 11. ¹H NMR (300 MHz, CDCl₃) δ /ppm: 8.18 (s, 1H), 7.80 (d, $J = 6.3$ Hz, 2H), 7.56 – 7.21 (m, 8H), 5.35 (s, 1H), 4.82 (s, 1H), 4.01 (s, 2H), 3.82 (s, 3H), 3.36 (t, $J = 6.3$ Hz, 1H).

(Phg[#])₂-*m*C₆H₄ (4_{m5}**).** Reaction 11. ¹H NMR (300 MHz, CDCl₃) δ /ppm: 8.03 (s, 1H), 7.82 (d, $J = 7.7$ Hz, 2H), 7.48 – 7.11 (m, 12H), 6.85 (t, $J = 7.7$ Hz, 1H), 5.30 (q, $J = 4.6$ Hz, 2H), 4.44 (s, 2H), 3.96 (d, $J = 19.7$ Hz, 4H).

Phe[#]-*m*C₆H₄-Gly-Val-Phe-OMe (3_{m6}**).** Reaction 12. ¹H NMR (300 MHz, CDCl₃) δ /ppm: 8.22 (s, 1H), 7.84 (dt, $J = 26.5, 7.9$ Hz, 2H), 7.53 (s, 1H), 7.47 – 7.11 (m, 5H), 7.07 (d, $J = 7.0$ Hz, 2H), 6.79 (s, 1H), 5.32 (d, $J = 13.5$ Hz, 1H), 4.78 (d, $J = 6.8$ Hz, 1H), 4.30 (t, $J = 7.3$ Hz, 1H), 4.22 – 3.83 (m, 5H), 3.65 (s, 3H), 3.01 (d, $J = 7.6$ Hz, 2H), 2.07 (q, $J = 6.8$ Hz, 1H), 0.97 – 0.79 (m, 6H).

Ala-1,4-Nph-Ala (2_{n1}**).** Reaction 13. $M_r(C_{70}H_{27}N_7O_6) = 386.15$. ESI-MS (m/z): 387.2 (M + H⁺, 50%), 409.0 (M + Na⁺, 39%), 773.3 (2M + H⁺, 78%), 795.1 (2M + Na⁺, 49%). ¹H NMR (600 MHz, CDCl₃) δ /ppm: 8.31 (dd, $J = 6.5, 3.3$ Hz, 2H), 7.62 – 7.53 (m, 4H), 6.62 (d, $J = 7.4$ Hz, 2H), 4.89 (quin., $J = 7.2$ Hz, 2H), 3.82 (s, 6H), 1.58 (d, $J = 7.2$ Hz, 6H). ¹³C NMR (151 MHz, CD₃OD) δ /ppm: 174.67, 171.86, 137.48, 131.64, 128.35, 126.79, 125.27, 52.87, 50.14, 17.09.

AMP-1,4-Nph-Ala (3_{n1}**).** Reaction 13. $M_r(C_{70}H_{24}N_7O_5) = 372.14$. ESI-MS (m/z): 373.2 (M + H⁺), 395.1 (M + Na⁺, 22%), 745.3 (2M + H⁺, 83%), 767.2 (2M + Na⁺, 26%). ¹H NMR (600 MHz, CDCl₃) δ /ppm: 8.31 – 8.23 (m, 1H), 8.18 – 8.11 (m, 1H), 7.59 – 7.52 (m, 2H), 7.50 (d, $J = 7.2$ Hz, 1H), 7.43 (d, $J = 7.1$ Hz, 1H), 6.68 (d, $J = 7.4$ Hz, 1H), 6.28 (s, 1H), 4.86 (quin., $J = 7.2$ Hz, 1H), 4.51 (s, 1H), 3.81 (d, $J = 9.0$ Hz, 3H), 3.74 (s, 2H), 1.56 (d, $J = 6.8$ Hz, 3H), 1.45 (s, 6H). ¹³C NMR (75 MHz, CD₃OD) δ /ppm: 174.68, 172.01, 171.89, 138.96, 136.94, 131.64, 131.50, 128.26, 128.23, 126.80, 126.63, 125.41, 124.85, 68.94, 57.16, 52.86, 49.85, 24.13, 17.11.

Gly-1,4-Nph-Gly (2_{n2}). Reaction 14. $M_r(C_{18}H_{18}N_2O_6) = 358.12$. ESI-MS (m/z): 359.1 ($M + H^+$, 40%), 717.1 ($2M + H^+$, 22%). 1H NMR (300 MHz, CD_3CN) δ/ppm : 8.38 – 8.26 (m, 2H), 7.69 – 7.59 (m, 4H), 7.39 – 7.26 (m, 2H), 4.16 (d, $J = 6.0$ Hz, 4H), 3.77 (s, 6H). ^{13}C NMR (151 MHz, CD_3OD) δ/ppm : 172.37, 171.72, 137.54, 131.62, 128.45, 126.84, 125.30, 52.74, 42.30.

AMP-1,4-Nph-Gly (3_{n2}). Reaction 14. $M_r(C_{19}H_{22}N_2O_5) = 358.15$. ESI-MS (m/z): 359.1 ($M + H^+$), 717.1 ($2M + H^+$). 1H NMR (600 MHz, CD_3CN) δ 8.35 – 8.28 (m, 1H), 8.24 – 8.18 (m, 1H), 7.64 – 7.60 (m, 3H), 7.57 (d, $J = 7.1$ Hz, 1H), 7.31 (s, 1H), 6.78 (s, 1H), 4.15 (d, $J = 6.0$ Hz, 2H), 3.97 (t, $J = 6.1$ Hz, 1H), 3.76 (s, 3H), 3.67 (d, $J = 6.1$ Hz, 2H), 1.40 (s, 6H). ^{13}C NMR (75 MHz, DMSO) δ/ppm : 170.37, 168.88, 168.34, 138.00, 135.20, 129.82, 129.77, 126.79, 125.66, 124.15, 123.59, 67.18, 55.30, 51.87, 41.11, 23.64.

Ala-1,5-Nph-Ala (2_{n3}). Reaction 15. $M_r(C_{20}H_{22}N_2O_6) = 386.15$. ESI-MS (m/z): 387.2 ($M + H^+$, 24%), 773.3 ($2M + H^+$, 7%). 1H NMR (600 MHz, $CDCl_3$) δ/ppm : 8.47 (d, $J = 8.6$ Hz, 2H), 7.70 (d, $J = 6.9$ Hz, 2H), 7.55 (dd, $J = 8.4, 7.1$ Hz, 2H), 6.55 (d, $J = 7.3$ Hz, 2H), 4.95 – 4.86 (m, 2H), 3.83 (s, 6H), 1.60 – 1.57 (m, 6H). ^{13}C NMR (151 MHz, DMSO) δ/ppm : 173.14, 168.58, 134.53, 129.85, 127.31, 125.65, 125.51, 51.99, 48.24, 16.60.

AMP-1,5-Nph-Ala (3_{n3}). Reaction 15. $M_r(C_{20}H_{24}N_2O_5) = 372.17$. ESI-MS (m/z): 373.1 ($M + H^+$), 745.2 ($2M + H^+$). 1H NMR (600 MHz, $CDCl_3$) δ/ppm : 8.42 (d, $J = 8.6$ Hz, 1H), 8.34 (d, $J = 8.6$ Hz, 1H), 7.69 (dd, $J = 7.0, 1.0$ Hz, 1H), 7.59 (dd, $J = 7.0, 1.0$ Hz, 1H), 7.56 – 7.47 (m, 2H), 6.59 (d, $J = 7.4$ Hz, 1H), 6.11 (s, 1H), 4.89 (quin., $J = 7.3$ Hz, 1H), 4.59 (s, 1H), 3.82 (s, 3H), 3.77 (d, $J = 3.9$ Hz, 2H), 1.58 (d, $J = 7.2$ Hz, 3H), 1.47 (s, 6H). ^{13}C NMR (151 MHz, $CDCl_3$) δ/ppm : 172.50, 169.55, 167.97, 134.03, 133.17, 129.44, 129.26, 127.09, 127.06, 125.03, 124.97, 124.80, 124.36, 69.64, 56.15, 51.76, 47.73, 23.79, 17.46.

AMP-1,5-Nph-AMP (4_{n3}). Reaction 15. $M_r(C_{20}H_{26}N_2O_4) = 358.19$. ESI-MS (m/z): 359.1 ($M + H^+$, 59 %), 717.2 ($2M + H^+$, 31 %). 1H NMR (600 MHz, DMSO) δ/ppm : 8.19 – 8.16 (m, 2H), 7.87 (s, 2H), 7.56 – 7.53 (m, 2H), 4.91 (s, 2H), 3.60 – 3.55 (m, 4H), 1.37 (s, 12H). ^{13}C NMR (75 MHz, DMSO) δ/ppm : 168.72, 136.35, 129.80, 126.63, 125.51, 124.89, 67.32, 55.25, 23.65.

Ala-2,6-Nph-Ala (2_{n4}). Reaction 16. $M_r(C_{20}H_{22}N_2O_6) = 386.15$. ESI-MS (m/z): 387.1 ($M + H^+$, 42%), 773.1 ($2M + H^+$, 11%). 1H NMR (600 MHz, CD_3OD) δ/ppm : 8.47 (s, 2H), 8.13 – 8.04 (m, 2H), 8.03 – 7.94 (m, 2H), 4.75 – 4.61 (m, 2H), 3.77 (s, 6H), 1.55 (d, $J = 7.3$ Hz, 6H). ^{13}C NMR (151 MHz, CD_3OD) δ/ppm : 174.85, 169.79, 135.46, 134.23, 130.48, 128.82, 125.99, 52.83, 50.27, 17.26.

AMP-2,6-Nph-Ala (3_{n4}). Reaction 16. $M_r(C_{20}H_{24}N_2O_5) = 372.42$. ESI-MS (m/z): 373.1 ($M + H^+$). 1H NMR (600 MHz, CD_3OD) δ/ppm : 8.45 (s, 1H), 8.37 (s, 1H), 8.05 (t, $J = 8.3$ Hz, 2H), 7.98 – 7.95 (m, $J = 8.5, 1.7$ Hz, 1H), 7.93 – 7.89 (m, $J = 8.5, 1.7$ Hz, 1H), 4.72 – 4.64 (m, 1H), 3.77 (s, 3H), 3.75 (s, 2H), 1.55 (d, $J = 7.3$ Hz, 3H), 1.46 (s, 6H). ^{13}C NMR (151 MHz, CD_3OD) δ/ppm : 174.86, 170.29, 169.83, 136.06, 135.47, 135.17, 133.95, 130.37, 130.32, 128.78, 128.36, 126.09, 125.86, 69.19, 56.91, 52.83, 50.26, 24.09, 17.26.

AMP-2,6-Nph-AMP (4_{n4}). Reaction 16. $M_r(C_{20}H_{26}N_2O_4) = 358.19$. ESI-MS (m/z): 359.1 ($M + H^+$), 1H NMR (600 MHz, CD_3OD) δ/ppm : 8.35 (s, 2H), 8.03 (s, 2H), 7.90 (s, 2H), 3.75 (s, 4H), 1.45 (s, 12H). ^{13}C NMR (75 MHz, DMSO) δ/ppm : 166.23, 134.11, 132.84, 128.38, 126.78, 124.91, 67.18, 54.93, 23.38.

Ala-2,7-Nph-Ala (2_{n5}). Reaction 17. $M_r(C_{20}H_{22}N_2O_6) = 386.15$. ESI-MS (m/z): 387.1 ($M + H^+$), 773.2 ($2M + H^+$). Crystals suitable for single-crystal x-ray diffraction were obtained from solution in NMR tube after several months. 1H NMR (600 MHz, $CDCl_3$)

δ/ppm : 8.39 – 8.29 (m, 2H), 7.94 – 7.89 (m, 2H), 7.88 – 7.82 (m, 2H), 7.05 (d, $J = 7.2$ Hz, 2H), 4.88 (quin., $J = 7.2$ Hz, 2H), 3.83 (s, 6H), 1.59 (d, $J = 7.2$ Hz, 6H). ^{13}C NMR (151 MHz, CDCl_3) δ/ppm : 174.00, 136.16, 132.04, 131.70, 128.54, 128.28, 125.72, 52.80, 48.80, 18.61.

AMP-2,7-Nph-Ala (3_{ns}). Reaction 17. $M_r(\text{C}_{20}\text{H}_{24}\text{N}_2\text{O}_5) = 372.42$. ESI-MS (m/z): 373.1 ($M + \text{H}^+$), 745.2 ($2M + \text{H}^+$, 85%). ^1H NMR (600 MHz, CDCl_3) δ/ppm : 8.31 (s, 1H), 8.15 (s, 1H), 7.90 – 7.70 (m, 4H), 7.11 (d, $J = 7.3$ Hz, 1H), 6.59 (s, 1H), 4.89 (quin., $J = 7.2$ Hz, 1H), 4.62 (t, $J = 6.2$ Hz, 1H), 3.84 (s, 3H), 3.76 (d, $J = 6.0$ Hz, 2H), 1.59 (d, $J = 7.2$ Hz, 3H), 1.50 (s, 6H). ^{13}C NMR (151 MHz, CDCl_3) δ/ppm : 174.33, 168.45, 167.08, 135.78, 133.20, 131.39, 131.22, 128.33, 128.06, 128.00, 127.80, 125.91, 125.42, 70.80, 56.72, 52.86, 48.82, 24.57, 24.56, 18.29.

Ala-9,10-Anth-Ala (2_{a1}). Reaction 18. $M_r(\text{C}_{24}\text{H}_{24}\text{N}_2\text{O}_6) = 436.16$. ESI-MS (m/z): 437.1 ($M + \text{H}^+$, 18%), 873.3 ($2M + \text{H}^+$, 26%). ^1H NMR (300 MHz, CDCl_3) δ/ppm : 8.43 – 7.87 (m, 4H), 7.60 – 7.49 (m, 4H), 6.55 (d, $J = 7.4$ Hz, 2H), 5.06 (quin., 2H), 3.86 (s, 6H), 1.65 (d, $J = 7.2$ Hz, 6H). ^{13}C NMR (75 MHz, CD_3OD) δ/ppm : 174.63, 129.17, 128.70, 127.70, 126.90, 126.44, 52.90, 50.30, 17.00.

AMP-9,10-Anth-Ala (3_{a1}). Reaction 18. $M_r(\text{C}_{24}\text{H}_{26}\text{N}_2\text{O}_5) = 422.18$. ESI-MS (m/z): 423.1 ($M + \text{H}^+$), 845.3 ($2M + \text{H}^+$). ^1H NMR (600 MHz, CDCl_3) δ/ppm : 8.38 – 8.04 (m, 4H), 7.59 – 7.55 (m, 4H), 6.54 (s, 1H), 6.08 (s, 1H), 5.06 (quin., $J = 7.2$ Hz, 1H), 4.56 (s, 1H), 3.91 (d, $J = 6.1$ Hz, 2H), 3.86 (s, 3H), 1.65 (d, $J = 7.2$ Hz, 3H), 1.53 (s, 6H). ^{13}C NMR (151 MHz, CD_3OD) δ/ppm : 174.63, 171.96, 171.70, 135.63, 134.02, 129.21, 128.79, 128.61, 128.56, 127.81 – 127.48 (4C), 127.02, 126.64, 126.44, 126.36, 68.94, 57.60, 52.89, 50.29, 24.24, 16.99.

1,3,5-C₆H₃-(Ala-OMe)₃ (2_{t1}). Reaction 19. ^1H NMR (300 MHz, CD_3CN) δ/ppm : 8.39 (s, 3H), 7.51 (d, $J = 7.0$ Hz, 3H), 4.60 (quin., $J = 7.2$ Hz, 3H), 3.71 (s, 9H), 1.48 (d, $J = 7.3$ Hz, 9H).

AMP-1,3,5-C₆H₃-(Ala-OMe)₂ (3_{t1}). Reaction 19. ^1H NMR (300 MHz, CD_3CN) δ/ppm : 8.37 (t, $J = 1.7$ Hz, 1H), 8.32 (d, $J = 1.7$ Hz, 2H), 7.50 (d, $J = 6.8$ Hz, 2H), 6.87 (s, 1H), 4.60 (quin., $J = 7.2$ Hz, 2H), 3.71 (s, 6H), 3.62 (s, 2H), 1.48 (d, $J = 7.3$ Hz, 6H), 1.38 (s, 6H). ^{13}C NMR (151 MHz, CD_3OD) δ/ppm : 174.70, 168.94, 168.55, 137.97, 135.91, 130.34, 130.11, 68.88, 57.01, 52.85, 50.26, 24.05, 17.23.

(AMP)₂-1,3,5-C₆H₃-Ala-OMe (4_{t1}). Reaction 19. ^1H NMR (300 MHz, CD_3CN) δ/ppm : 8.30 (s, 2H), 8.25 (s, 1H), 7.49 (s, 1H), 6.86 (s, 2H), 4.59 (quin., $J = 7.3$ Hz, 1H), 3.87 (t, $J = 6.2$ Hz, 2H), 3.70 (s, 3H), 3.62 (d, $J = 6.2$ Hz, 4H), 1.48 (d, $J = 7.3$ Hz, 3H), 1.38 (s, 12H).

1,3,5-C₆H₃-(Gly-OMe)₃ (2_{t2}). Reaction 20. ^1H NMR (300 MHz, CD_3CN) δ/ppm : 8.40 (s, 3H), 7.64 (s, 3H), 4.11 (d, $J = 5.9$ Hz, 6H), 3.72 (s, 9H).

Val[#]-1,3,5-C₆H₃-(Gly-OMe)₂ (3_{t2}). Reaction 20. ^1H NMR (300 MHz, CD_3CN) δ/ppm : 8.37 (s, 3H), 7.66 (s, 2H), 7.05 (d, $J = 9.1$ Hz, 1H), 4.11 (d, $J = 5.9$ Hz, 4H), 3.99 – 3.83 (m, 1H), 3.72 (s, 6H), 3.64 (dt, $J = 9.7, 4.1$ Hz, 2H), 0.97 (dd, $J = 7.8, 6.8$ Hz, 6H).

(Val[#])₂-1,3,5-C₆H₃-Gly-OMe (4_{t2}). Reaction 20. ^1H NMR (300 MHz, CD_3CN) δ/ppm : 8.37 (s, 3H), 7.71-7.49 (m, 1H), 7.03 (d, $J = 9.0$ Hz, 2H), 4.11 (d, $J = 5.9$ Hz, 2H), 4.02 – 3.82 (m, 2H), 3.72 (s, 3H), 3.69 – 3.60 (m, 4H), 2.97 (t, $J = 6.0$ Hz, 2H), 0.98 (dd, $J = 8.1, 6.8$ Hz, 12H).

Phg[#]-1,3,5-C₆H₃-(Phe-OMe)₂ (3_{t5}). Reaction 21. ^1H NMR (300 MHz, CDCl_3) δ/ppm : 8.16 (s, 1H), 8.02 (s, 0H), 7.58 (dd, $J = 25.6, 7.9$ Hz, 3H), 7.48 – 7.14 (m, 15H), 5.40 – 5.31 (m, 1H), 5.13 – 5.00 (m, 2H), 4.07 – 3.92 (m, 1H), 3.81 (s, 3H), 3.33 – 3.12 (m, 2H).

(Phg[#])₂-1,3,5-C₆H₃-Phe-OMe (4_{t5}). Reaction 21. ¹H NMR (300 MHz, CDCl₃) δ/ppm: 8.09 (s, 1H), 8.00 (s, 2H), 7.71 (d, *J* = 7.7 Hz, 3H), 7.48 – 7.06 (m, 15H), 5.36 (q, *J* = 6.9, 5.3 Hz, 2H), 5.05 (q, *J* = 7.2 Hz, 1H), 4.55 – 4.40 (m, 2H), 4.14 – 3.99 (m, 2H), 3.85-3.75 (m, 5H), 3.33 – 3.10 (m, 2H).

(Phg[#])₃-1,3,5-C₆H₃ (5_{t5}). Reaction 21. ¹H NMR (300 MHz, CD₃OD) δ/ppm: 8.49 (s, 2H), 8.36 (s, 1H), 7.51 – 7.18 (m, 15H), 5.23 (t, *J* = 6.6 Hz, 3H), 3.87 (d, *J* = 6.8 Hz, 6H).

Linear reaction sequence 1

Dimethyl isophthalate (9_{m1}). Isophthalic acid (1 666.1 mg, 10 mmol) was dissolved in MeOH (100mL) and 2 mL of conc. H₂SO₄ was added to the solution. The reaction mixture was refluxed at 75 °C with continuous stirring overnight. MeOH was evaporated under reduced pressure. The residue was dissolved in ethyl acetate (100 mL) and washed with water (100 mL), NaHCO₃ (sat. aq, 100 mL), water (100 mL), dried over Na₂SO₄, filtered and evaporated under reduced pressure to yield the crude white product. No further purification was required. Yield: 1 792.8 mg (9.23 mmol, 92%), white powder. ¹H NMR (300 MHz, CDCl₃) δ/ppm: 8.69 (t, *J* = 1.5 Hz, 1H), 8.23 (dd, *J* = 7.8, 1.7 Hz, 2H), 7.53 (t, *J* = 7.8 Hz, 1H), 3.95 (s, 6H).

HOOC-*m*C₆H₄-COOMe (10_{m1}). Dimethyl isophthalate (9_{m1}) (1 791.8 mg, 9.23 mmol) was dissolved in MeOH (20 mL) by using an ultrasonic bath. Aqueous solution of NaOH (368.0 mg, 9.23 mmol in 10 mL of distilled water) was added to the mixture and the reaction mixture heated in a CEM Microwave Reactor for 20 min (150 W, 50 °C). To the aqueous residue HCl (284 μL conc., 9.23 mmol in 10 mL of distilled water) was added, and the white precipitate extracted with ethyl acetate (3 x 40 mL). Combined organic extracts were washed with citric acid (10% aq, 100 mL), NaCl (sat. aq. 100 mL), dried over Na₂SO₄, filtered and evaporated under reduced pressure. Chromatography: 38 g of silica gel, 3% MeOH in DCM. Yield: 912.2 mg (5.06 mmol, 55%), white powder. ¹H NMR (300 MHz, CDCl₃) δ/ppm: 8.81 – 8.74 (m, 1H), 8.34 – 8.26 (m, 2H), 7.59 (t, *J* = 7.8 Hz, 1H), 3.97 (s, 3H).

AMP-*m*C₆H₄-COOMe (11_{m1}). HOOC-*m*C₆H₄-COOMe (10_{m1}) (360.3 mg, 2.0 mmol), HATU (836.5 mg, 2.2 mmol), DIPEA (1. 360 mL, 8.0 mmol), AMP (222.9 mg, 2.5 mmol). NMR spectrum showed significant content of tetramethyl urea (mass ratio *w*(product) = 84 %). The compound was used without further purification. Yield: 421,5 mg (as calculated from NMR spectrum, 1.68 mmol, 84%), colorless oil. ¹H NMR (300 MHz, CDCl₃) δ/ppm: 8.33 (s, 1H), 8.17 (d, *J* = 7.8 Hz, 1H), 8.00 (d, *J* = 7.8 Hz, 1H), 7.54 (t, *J* = 7.8 Hz, 1H), 6.23 (s, 1H), 3.95 (s, 3H), 3.72 (s, 2H), 1.44 (s, 7H).

AMP-*m*C₆H₄-COOH (12_{m1}). AMP-*m*C₆H₄-COOMe (11_{m1}) (421.5 mg, 1.68 mmol) was dissolved in MeOH (20 mL) by using an ultrasonic bath. Aqueous solution of NaOH (80.0 mg, 2.00 mmol in 10 mL of distilled water) was added to the mixture and the reaction mixture heated in a CEM Microwave Reactor for 20 min (150 W, 50 °C). To the aqueous residue HCl (62 μL conc., 2.0 mmol in 10 mL of distilled water) was added, and the white precipitate extracted with ethyl acetate (3 x 40 mL). Combined organic extracts were washed with citric acid (10% aq, 100 mL), NaCl (sat. aq. 100 mL), dried over Na₂SO₄, filtered and evaporated under reduced pressure. NMR spectrum showed significant content of tetramethyl urea and water (mass ratio *w*(product) = 72 %). The compound was used without further purification. Yield: 387.0 mg (as calculated from NMR spectrum, 1.63 mmol, 97%) ¹H NMR (300 MHz, CD₃OD) δ 8.43 (s, 1H), 8.17 (d, *J* = 7.8 Hz, 1H), 8.00 (d, *J* = 7.9 Hz, 1H), 7.57 (t, *J* = 7.8 Hz, 1H), 3.73 (s, 2H), 1.44 (s, 6H).

AMP-*m*C₆H₄-Ala (3_{m1}). AMP-*m*C₆H₄-COOH (11_{m1}) (387.0 mg, 1.63 mmol), HATU (762.47 mg, 2.0 mmol), DIPEA (1. 360 mL, 8.0 mmol), AMP (222.9 mg, 2.5 mmol)

Chromatography: 30 g of silica gel, 3% MeOH in DCM. Yield: 147.3 mg (0.46 mmol, 28%; overall yield: 9%), colorless oil. Recorded spectra were analogous to the previously obtained spectra of the **3_{m1}** compound.

Linear reaction sequence 2

1,3,5-C₆H₃-(COOMe)₃ (13_{t3}). Trimesic acid (1 441.7 mg, 6.9 mmol) was dissolved in MeOH (100mL) and 2 mL of conc. H₂SO₄ was added to the solution. The reaction mixture was refluxed at 75 °C with continuous stirring overnight. MeOH was evaporated under reduced pressure. The residue was dissolved in ethyl acetate (100 mL) and washed with water (100 mL), NaHCO₃ (sat. aq, 100 mL), water (100 mL), dried over Na₂SO₄, filtered and evaporated under reduced pressure to yield the crude white product. No further purification was required. Yield: 1 638.1 mg (6.5 mmol, 95%), white solid. ¹H NMR (300 MHz, CDCl₃) δ/ppm: 8.86 (s, 3H), 3.98 (s, 9H). ¹³C NMR (151 MHz, CD₃OD) δ/ppm: 164.65, 133.21, 130.72, 51.27.

HOOC-1,3,5-C₆H₃-(COOMe)₂ (14_{t3}). 1,3,5-C₆H₃-(COOMe)₃ (**13_{t3}**) (1 638.1 mg, 6.5 mmol) was dissolved in MeOH (20 mL) by using an ultrasonic bath. Aqueous solution of NaOH (259.9 mg, 6.5 mmol in 10 mL of distilled water) was added to the mixture and the reaction mixture heated in a CEM Microwave Reactor for 20 min (150 W, 50 °C). To the aqueous residue HCl (546 μL conc., 6.50 mmol in 10 mL of distilled water) was added, and the white precipitate extracted with ethyl acetate (3 x 40 mL). Combined organic extracts were washed with citric acid (10% aq, 100 mL), NaCl (sat. aq. 100 mL), dried over Na₂SO₄, filtered and evaporated under reduced pressure. The obtained mixture could not be purified by chromatography, therefore, it was used without purification in the next step. Yield: 945.0 mg (<4.0 mmol, <51%), white powder.

Phg[#]-1,3,5-C₆H₃-(COOMe)₂ (15_{t3}). HOOC-1,3,5-C₆H₃-(COOMe)₂ (**14_{t3}**) (571.5 mg, 2.4 mmol), HATU (912.6 mg, 2.4 mmol), DIPEA (1.666 mL, 9.6 mmol), Phg[#] (329.2 mg, 2.4 mmol). Chromatography: 30 g of silica gel, 3% MeOH in DCM. Yield: 497.1 mg (1.39 mmol, 58%), white solid. ¹H NMR (300 MHz, CDCl₃) δ/ppm: 8.79 (t, J = 1.7 Hz, 1H), 8.65 (d, J = 1.6 Hz, 2H), 7.39 (d, J = 4.3 Hz, 5H), 7.09 (d, J = 7.3 Hz, 1H), 5.32 (dt, J = 7.2, 4.8 Hz, 1H), 4.04 (t, J = 5.4 Hz, 2H), 3.97 (s, 6H), 2.47 (t, J = 6.1 Hz, 1H).

Phg[#]-1,3,5-C₆H₃-(Phe-OMe)₂ (3_{t3}). Phg[#]-1,3,5-C₆H₃-(COOMe)₂ (**15_{t3}**) (556.5 mg, 1.6 mmol) was dissolved in MeOH (20 mL) by using an ultrasonic bath. Aqueous solution of NaOH 200.0 mg, 5.0 mmol in 10 mL of distilled water) was added to the mixture and the reaction mixture heated in a CEM Microwave Reactor for 20 min (150 W, 50 °C). To the aqueous residue HCl (422 μL conc., 5.0 mmol in 10 mL of distilled water) was added gradually until neutralization was achieved. Water was evaporated from the mixture and the solid suspended in the DCM. The suspension was used in the next step without further processing. HATU (1 901.2 mg, 5.0 mmol), DIPEA (2.318 mL, 12.4 mmol), Phe-OMe·HCl (1 078.4 mg, 5 mmol) Chromatography: 50 g of silica gel, 2% MeOH in DCM. Yield: 505.8 mg (0.78 mmol, 50%; overall yield: 28%), white solid. ¹H NMR spectrum of compound **3_{t3}** corresponds to ¹H NMR of compound **3_{t5}**, however high amounts of contaminant TMU and other unidentified compounds are present (¹H and ¹³C NMR spectra in the supplement).

Phg[#]-1,3,5-C₆H₃-(D-Phe-OMe)₂ (3_{t4}). Phg[#]-1,3,5-C₆H₃-(COOMe)₂ (**15_{t3}**) (504.0 mg, 1.4 mmol) was dissolved in MeOH (20 mL) by using an ultrasonic bath. Aqueous solution of NaOH 200.0 mg, 5.0 mmol in 10 mL of distilled water) was added to the mixture and the reaction mixture heated in a CEM Microwave Reactor for 20 min (150 W, 50 °C). To the aqueous residue HCl (422 μL conc., 5.0 mmol in 10 mL of distilled water) was added gradually until neutralization was achieved. Water was evaporated from the mixture and the solid suspended in the DMF. The suspension was used in the next step without further processing. COMU (2 141.3 mg, 5.0 mmol), TEA (1.868 mL, 13.4 mmol), D-Phe-OMe·HCl (1 078.4 mg, 5 mmol) Chromatography: 60 g of silica gel, 3% MeOH in DCM. Yield: 819.3 mg (<1.3 mmol, <89%; overall yield: 50%), white

solid. ^1H NMR spectrum of compound **3_{t4}** showed high amounts of unidentified contaminants which could not be purified by additional chromatography, therefore the compound was used in the next step as is (^1H and ^{13}C NMR spectra in the supplement).

One-pot products within the same series do not have the same limiting reactant. To avoid discrepancies, all yields presented in the paper were calculated with the aromatic acid **P** as the limiting reactant (Obt. η). In Table S1 maximum yields (Max η) for all one-pot products for 100% of the mass of the isophthalic acid are presented (intermediates whose limiting reactant is not the aromatic acid **P** have this maximum yield value lower than 100%).

Table S1. Obtained yield values as calculated with the mass of the aromatic acid **P** as the limiting reactant.

Reactio n	Compoun d	Obt. η / %	Max. η /%
1	2_{m1}	1	100
	6_{m1}	8	100
	7_{m1}	4	100
	8_{m1}	2	100
3	3_{m1}	17	100
4	2_{m1}	25	50
	3_{m1}	13	100
5	2_{m1}	20	50
	3_{m1}	15	100
6	4_{m1}	27	100
7	2_{p1}	7	50
	3_{p1}	<13	100
8	3_{m2}	22	100
9	3_{m3}	16	100
10	3_{m4}	26	100
11	3_{m5}	13	100
	5_{m5}	3	50
12	3_{m6}	<52	100
13	2_{n1}	8	50
	3_{n1}	36	100
13	2_{n1}	8	50
	3_{n1}	36	100

Reactio n	Compoun d	Obt. η / %	Max. η /%
14	2_{n2}	17	50
	3_{n2}	18	100
15	2_{n3}	31	50
	3_{n3}	15	100
	4_{n3}	17	50
16	2_{n4}	5	50
	3_{n4}	33	100
	4_{n4}	38	50
17	2_{n5}	18	50
	3_{n5}	23	100
18	2_{a1}	19	50
	3_{a1}	16	100
19	2_{t1}	6	18
	3_{t1}	27	75
	4_{t1}	10	77
20	2_{t2}	2	50
	3_{t2}	16	76
	4_{t2}	6	75
21	3_{t5}	5	29
	4_{t5}	26	52
	5_{t5}	5	38

Table S2. Isolated yields of oxazoline compounds **1**.

Compound	yield / %	Compound	yield / %	Compound	yield / %
1_{m1}	57	1_{n1}	42	1_{t1}	86
1_p*	22	1_{n2}	35	1_{t2}	13
1_{m2}	68	1_{n3}	42	1_{t3}	38
1_{m3}	54	1_{n4}	43	1_{t4}*	24
1_{m4}	67	1_{n5}	14	1_{t5}*	33
1_{m5}	36	1_a	87	1_{t6}	10
1_{m6}*	<47			1_b	34

*not chromatographically purified in the previous step(s).

2. NMR spectra.

Table S3. NMR shifts of amide peaks of derivatives **2** and oxazolines **1** at c ~ 6 mM.

Compound	Solvent	$\delta(\text{N-H})$ /ppm	Compound	Solvent	$\delta(\text{N-H})$ /ppm
2_{m1}	CHCl ₃	6.88	2_{n3}	CHCl ₃	7.03
1_{m1}	CHCl ₃	6.77	1_{n3}	CHCl ₃	6.89
	DCM	6.76	2_{n4}	CHCl ₃	6.55
	MeCN	7.40	1_{n4}	CHCl ₃	6.56
	DMSO	9.02	2_{n5}	CHCl ₃	7.05
1_p	CHCl ₃	6.76	1_{n5}	CHCl ₃	6.88
1_{m2}	CHCl ₃	6.63	2_a	CHCl ₃	6.55
1_{m3}	CHCl ₃	6.80	1_a	CHCl ₃	6.51
1_{m4}	CHCl ₃	6.80	2_{t1}	MeCN	7.51
1_{m5}	CHCl ₃	6.85	1_{t1}	CHCl ₃	6.95
4_{m5}	CHCl ₃	6.85	2_{t2}	MeCN	7.64
1_{m6}	CHCl ₃	7.28 (Gly) 6.65 (Val) 6.48 (Phe)	1_{t2}	MeCN	7.93, 7.62
2_{n1}	CHCl ₃	6.57	1_{t3}	CHCl ₃	6.74
1_{n1}	CHCl ₃	6.53	1_{t4}	MeCN	7.50
2_{n2}	CHCl ₃	6.94	1_{t5}	CHCl ₃	6.70
1_{n2}	CHCl ₃	6.55			

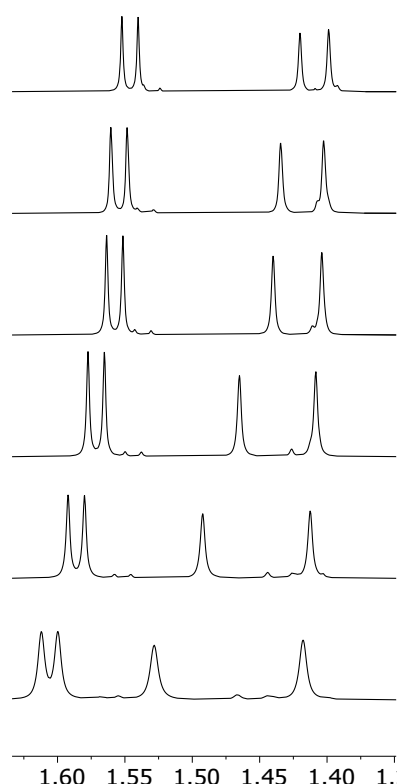
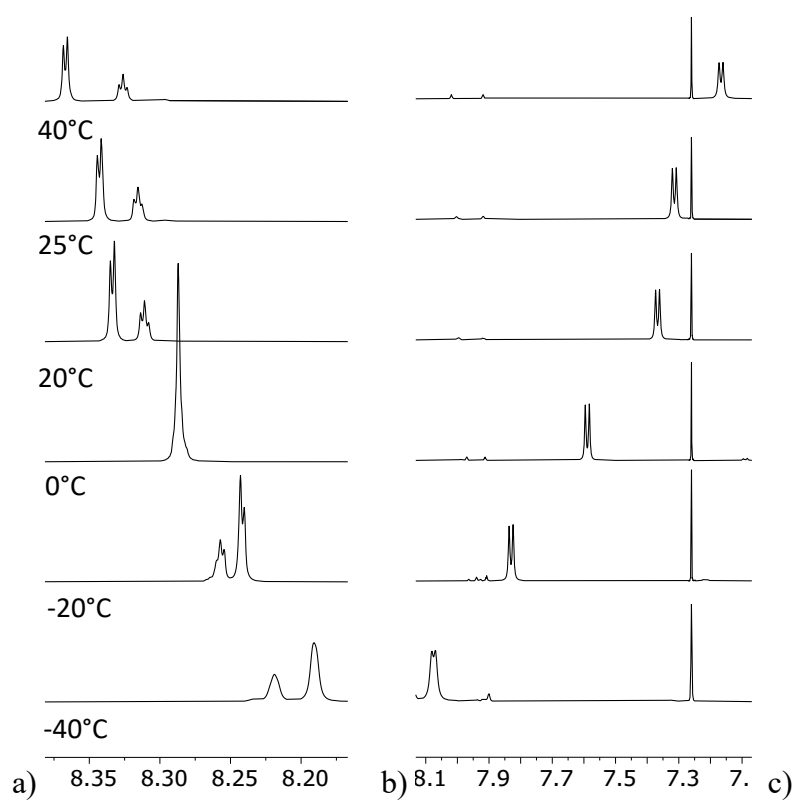
NMR spectra of selected ligands (namely **1_{m1}**, **1_p**, **1_{m5}**, **1_{m6}**, **1_{n4}**, **1_{t1}** and **1_{t5}**) were recorded in CDCl₃ and DMSO (both at *c* = 6 mM, and additionally at *c* = 60 mM in CDCl₃), and respective hydrogen bond acidity values (*A_{NMR}*) calculated from the equation:

$$A_{NMR} = 0.0065 + 0.133\Delta\delta_{DMSO-CDCl_3}$$

A_{NMR} values which are greater than 0.15 indicate no significant hydrogen bonding with the amide protons in these solutions occurs.

Table S4. Hydrogen bond acidity values (*A_{NMR}*) and $\Delta\delta_{conc.-dil}$ of amide peaks of selected ligands.

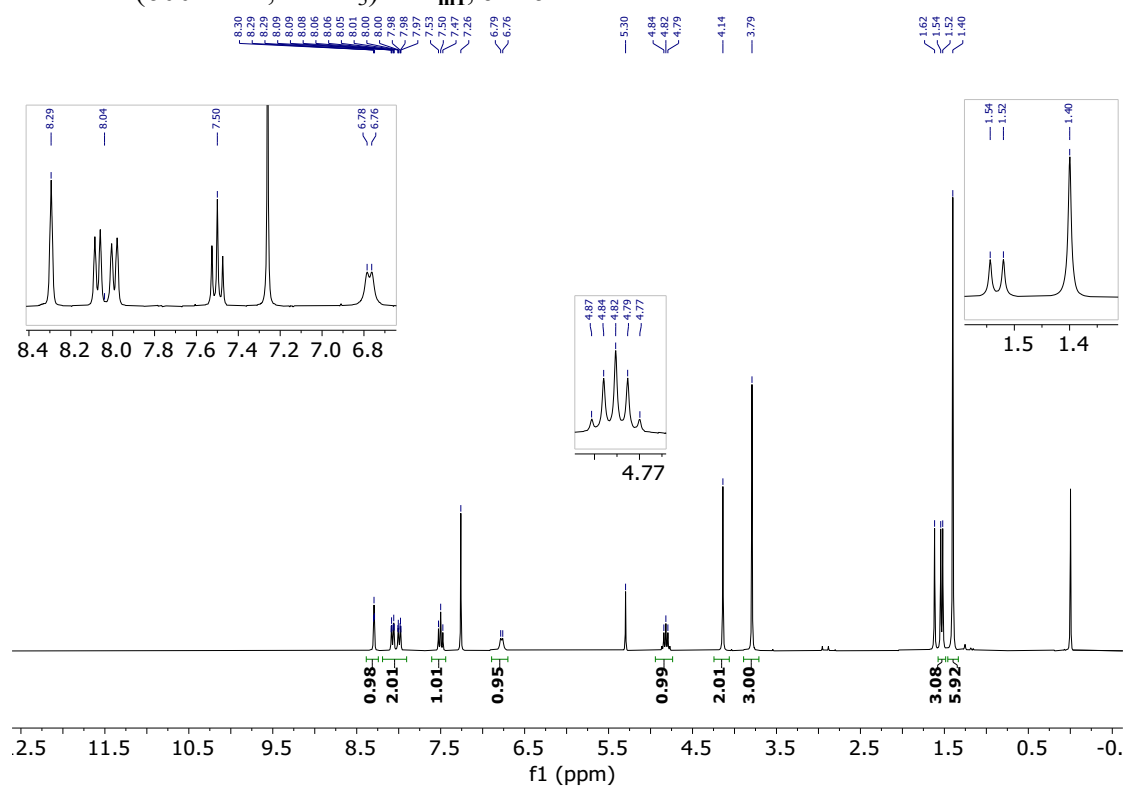
comp.	$\delta(N-H)/ppm$		$\Delta\delta_{DMSO-CDCl_3}^*$	<i>A_{NMR}</i>	CDCl ₃ (konc.)	$\Delta\delta_{conc.-dil}$
	DMSO	CDCl ₃				
1_{m1}	9.02	6.80	2.22	0.30	6.85	0.05
1_p	8.94	6.75	2.19	0.30	6.78	0.03
1_{m5}	9.04	6.79	2.25	0.31	6.87	0.08
1_{m6}	8.95(Gly)	7.28 (Gly)	1.67 (Gly)	0.23	7.82 (Gly)	0.54 (Gly)
	7.81 (Val)	6.65 (Val)	1.16 (Val)	0.16	7.03 (Val)	0.38 (Val)
	8.48 (Phe)	6.48 (Phe)	2.00 (Phe)	0.27	7.01 (Phe)	0.53 (Phe)
1_{n4}	9.01	6.88	2.13	0.29	6.94	0.06
1_{t1}	9.20	6.95	2.25	0.31	7.31	0.36
1_{t5}	9.35	6.70	2.65	0.36	6.78	0.08



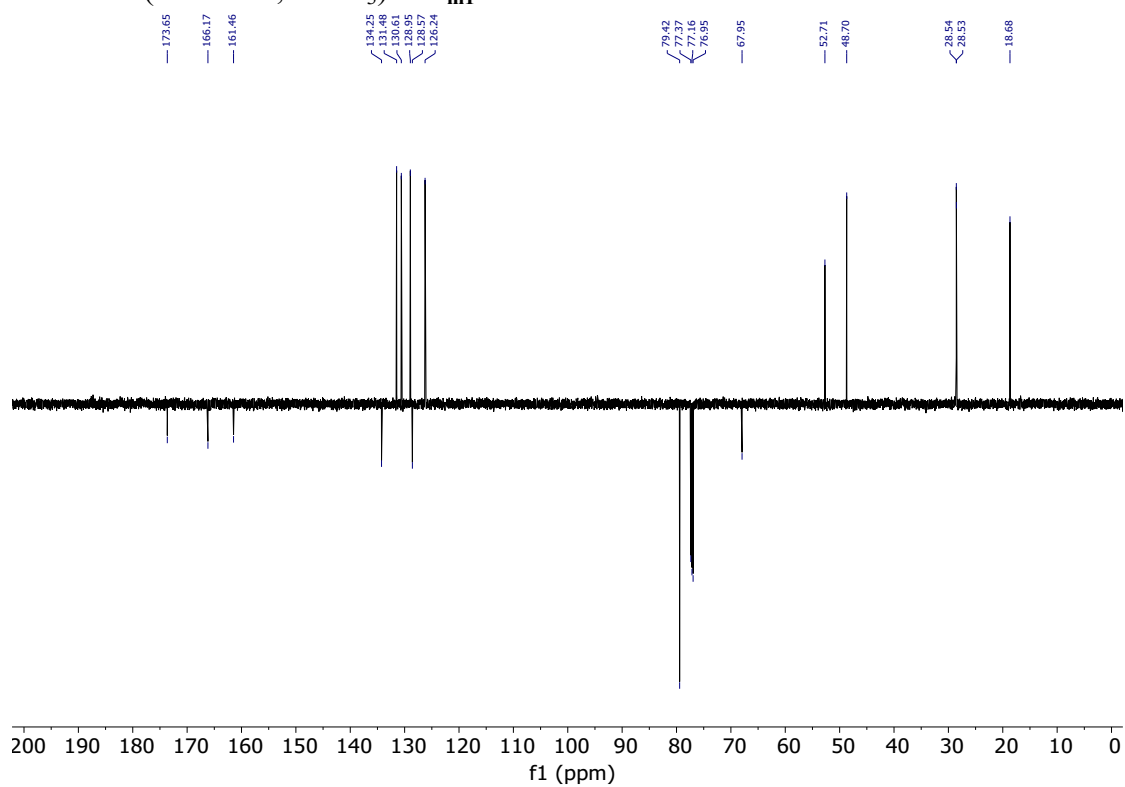
1. Figure S1. Temperature dependent ^1H NMR spectra of oxazoline $\mathbf{1}_{\text{t1}}$: a) aromatic region, b) amide region, c) aliphatic region.

2.1. Oxazolines

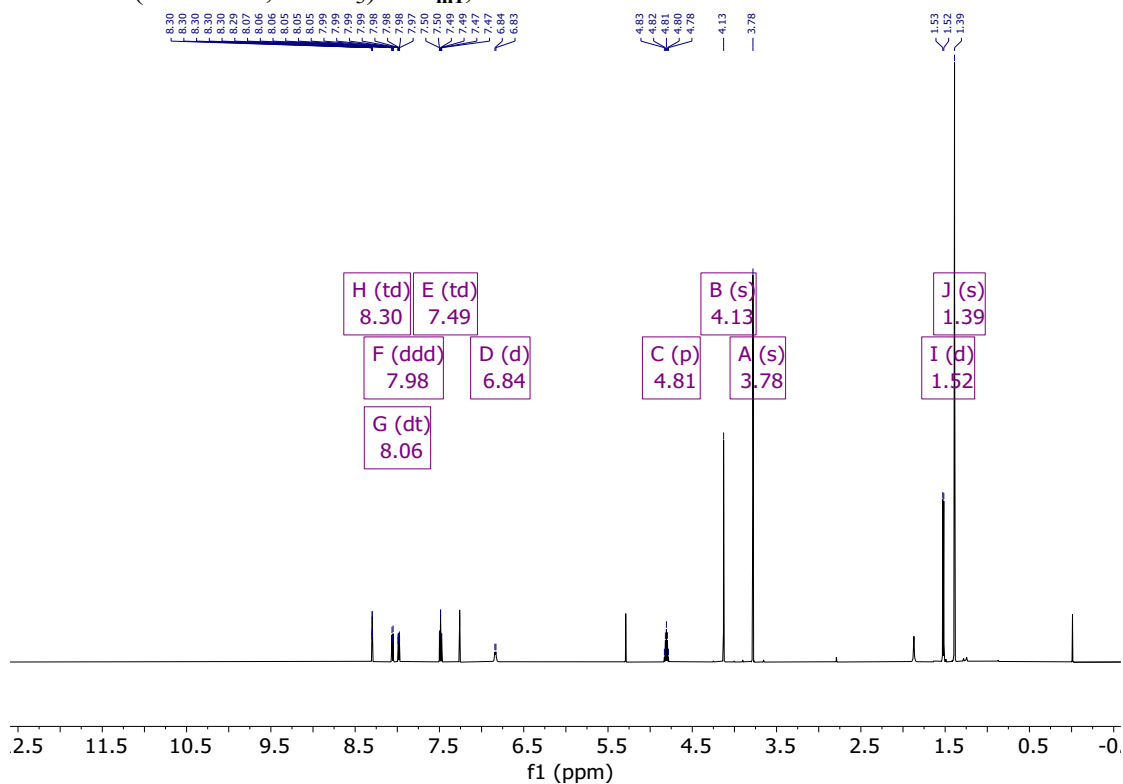
^1H NMR (600 MHz, CDCl_3) of **1_{m1}**, $c = 6$ mM.



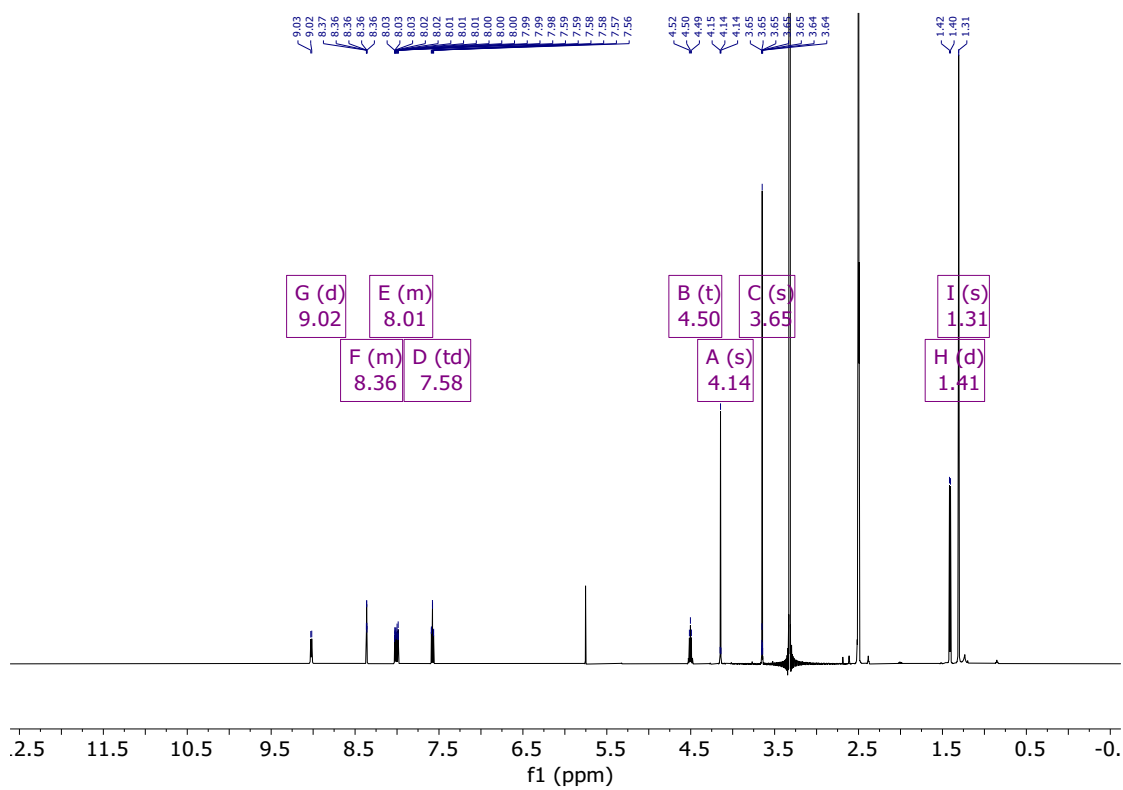
^{13}C NMR (151 MHz, CDCl_3) of **1_{m1}**.



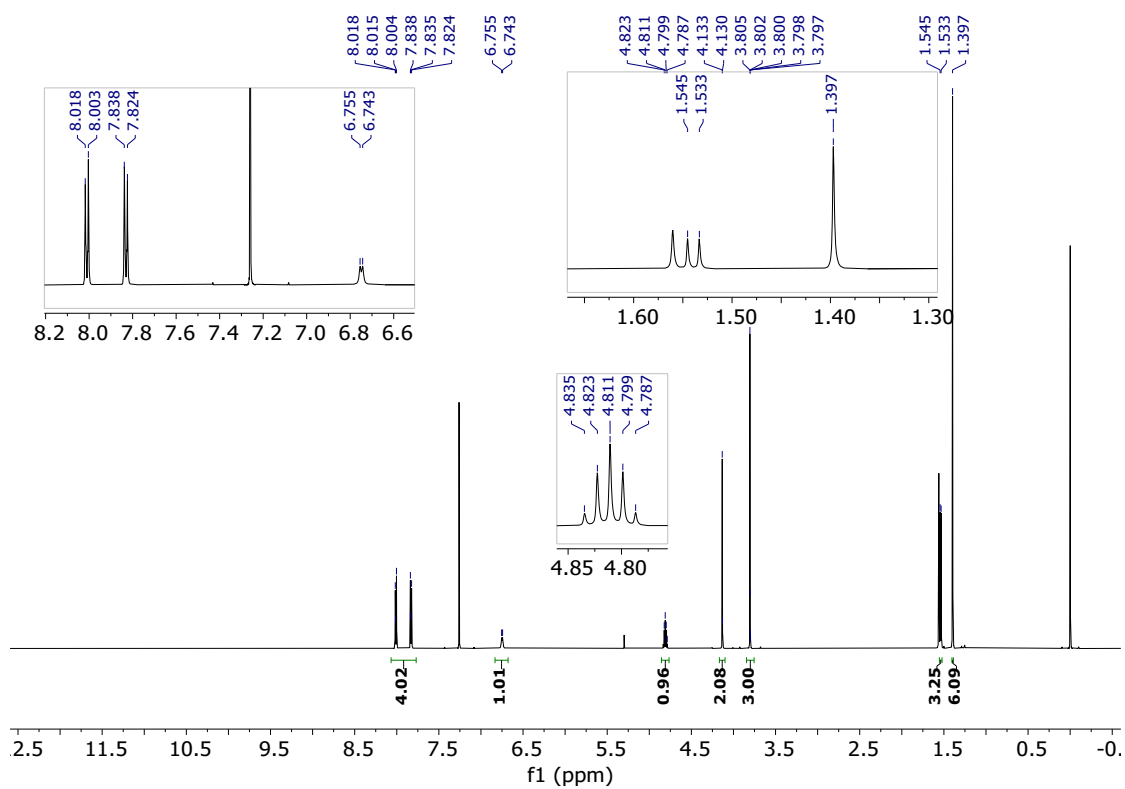
^1H NMR (600 MHz, CDCl_3) of $\mathbf{1}_{\text{m1}}$, $c = 60$ mM.



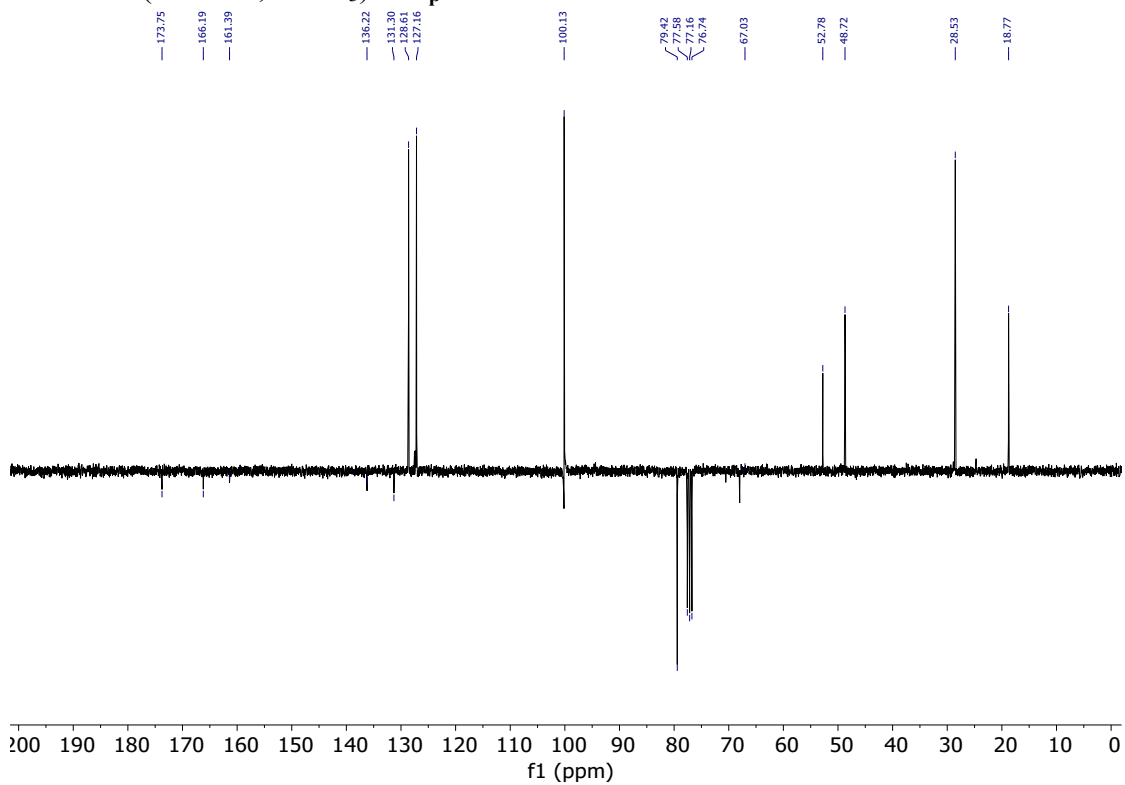
^1H NMR (600 MHz, DMSO) of $\mathbf{1}_{\text{m1}}$, $c = 6$ mM.



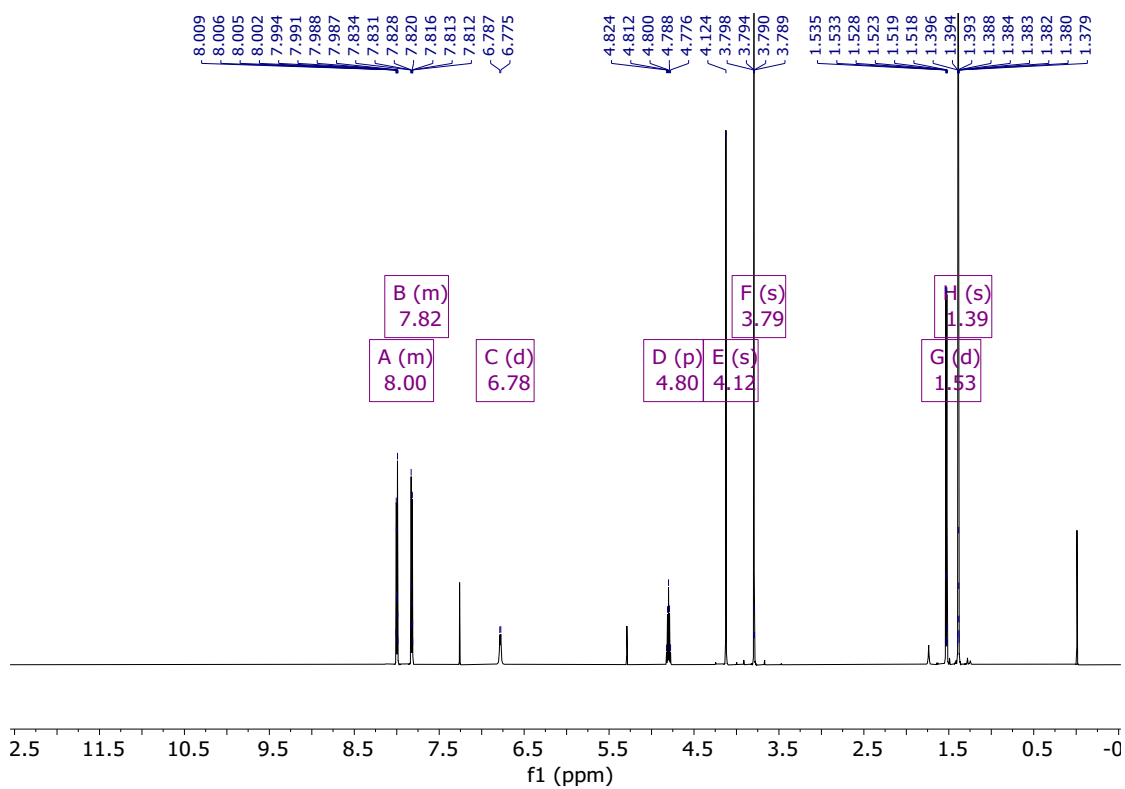
^1H NMR (300 MHz, CDCl_3) of $\mathbf{1_p}$, $c = 6$ mM.



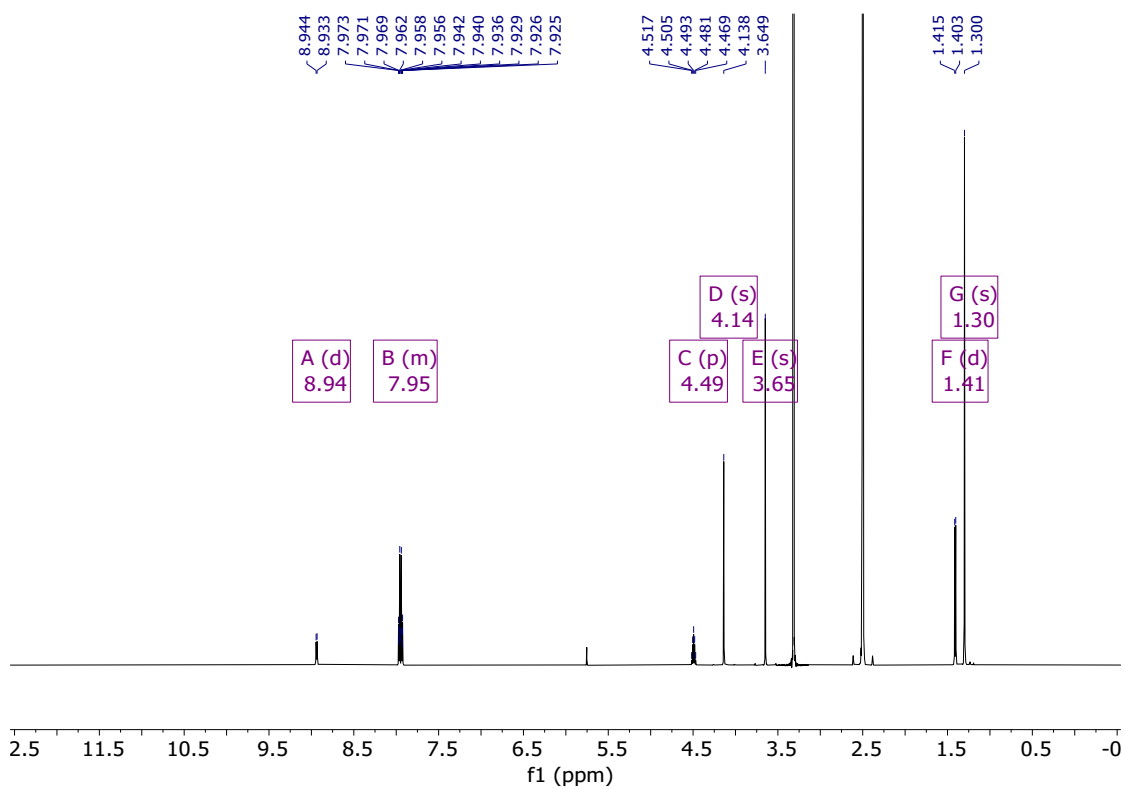
^{13}C NMR (75 MHz, CDCl_3) of $\mathbf{1_p}$.



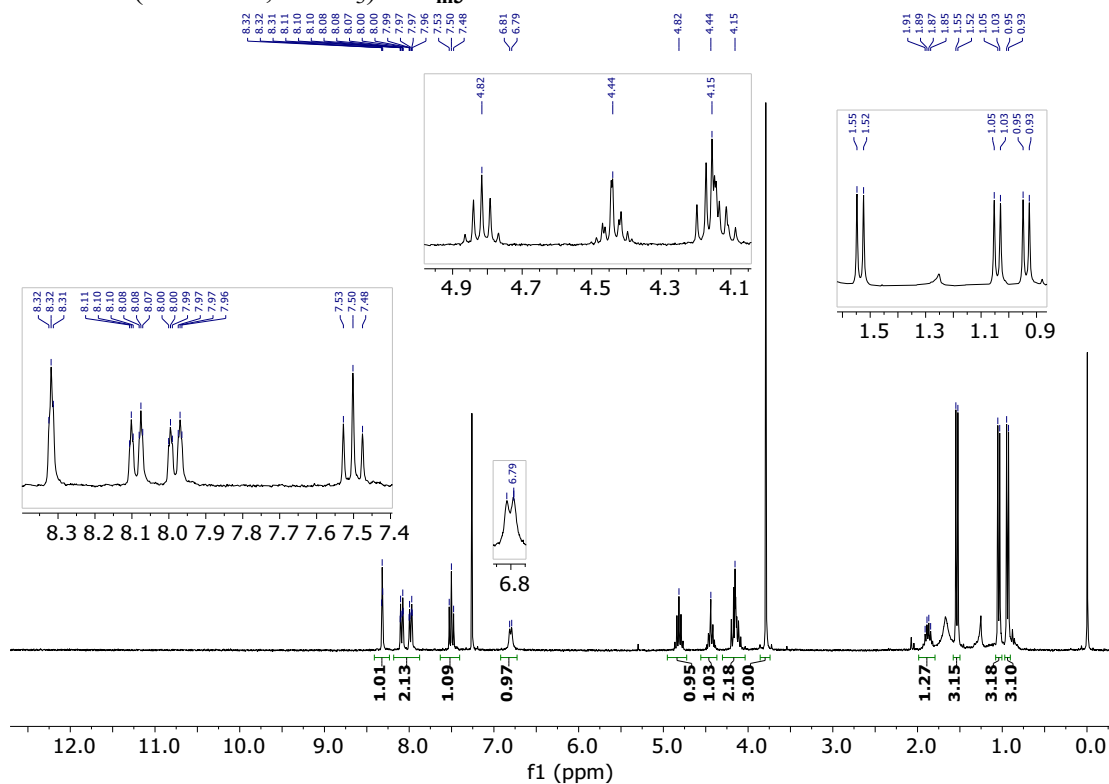
^1H NMR (600 MHz, CDCl_3) of 1_p , $c = 60$ mM.



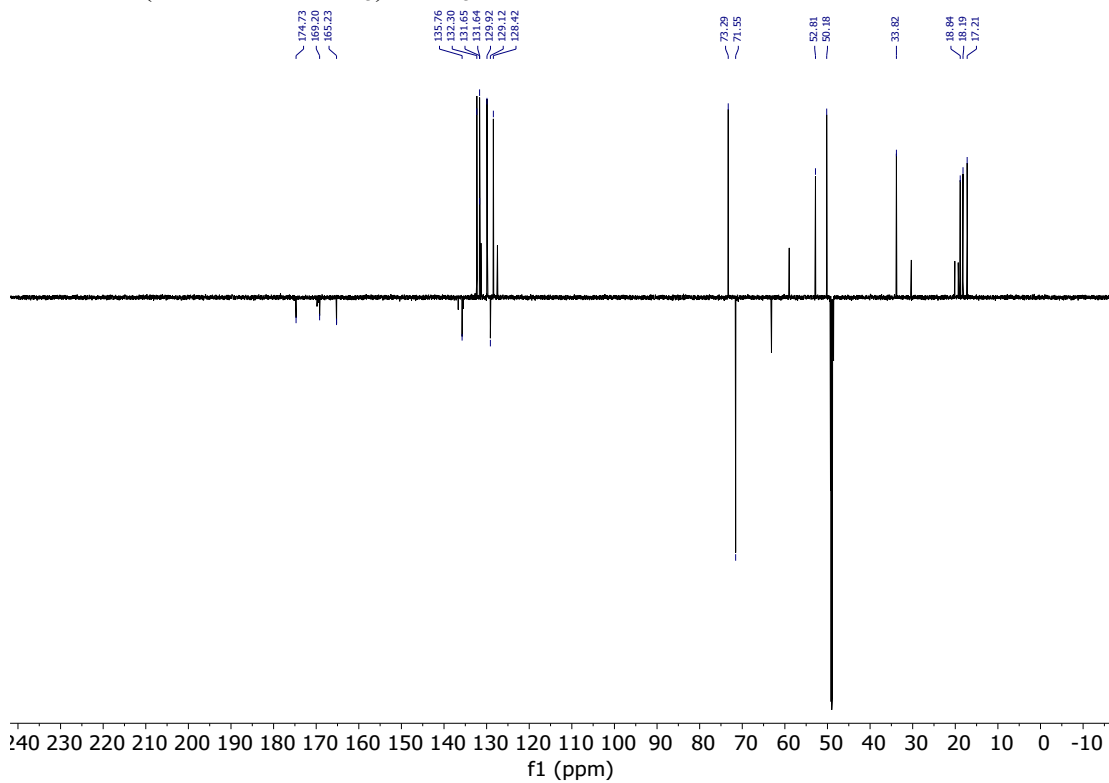
^1H NMR (600 MHz, DMSO) of 1_p , $c = 6$ mM.



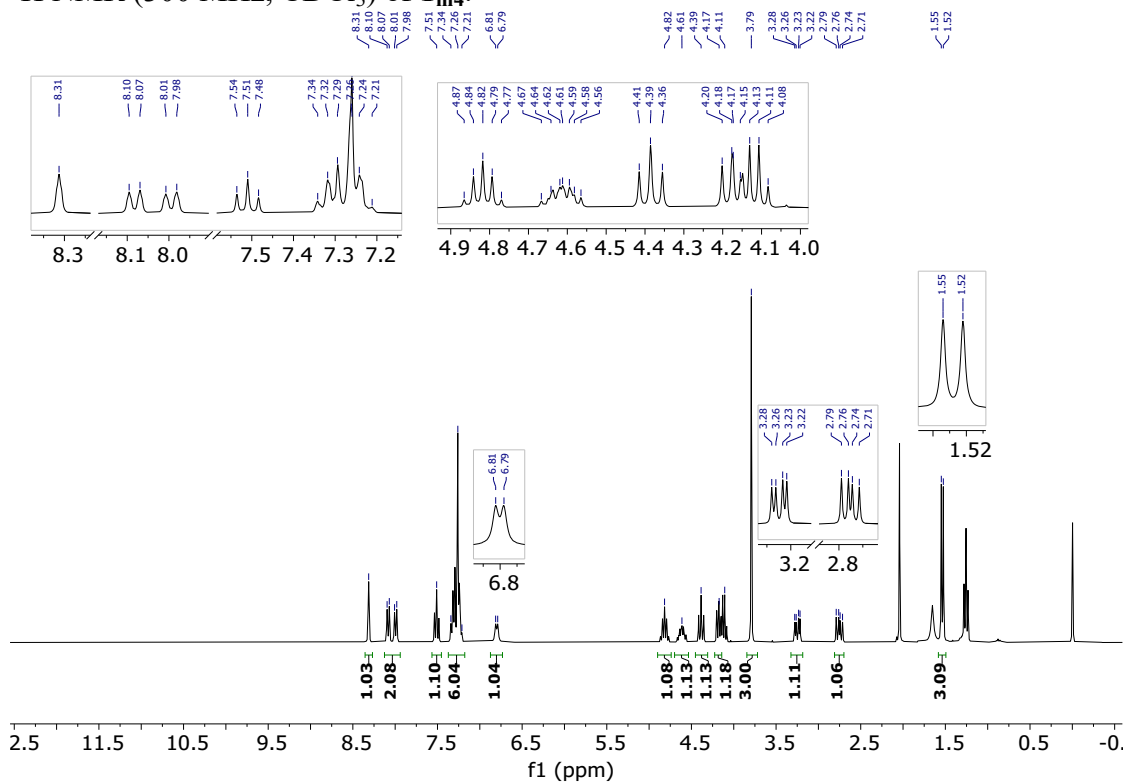
¹H NMR (300 MHz, CDCl₃) of **1m3**.



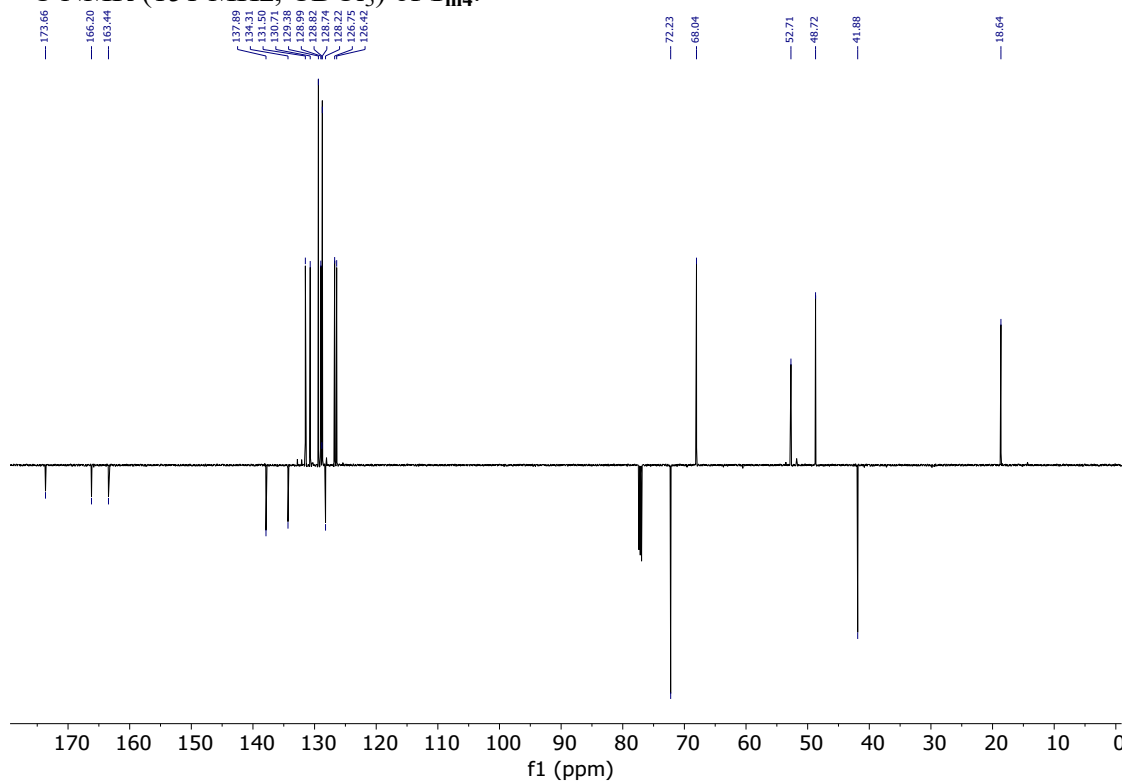
¹³C NMR (151 MHz, CDCl₃) of **1m3**.



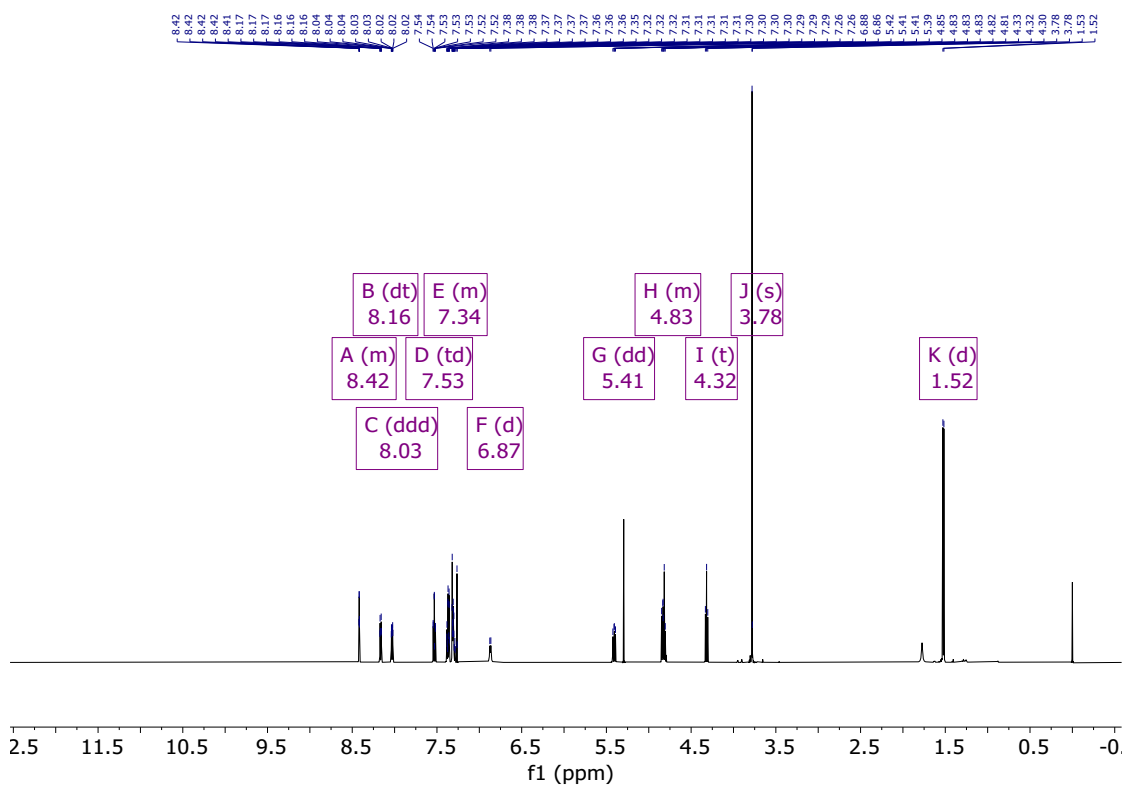
^1H NMR (300 MHz, CDCl_3) of **1m4**.



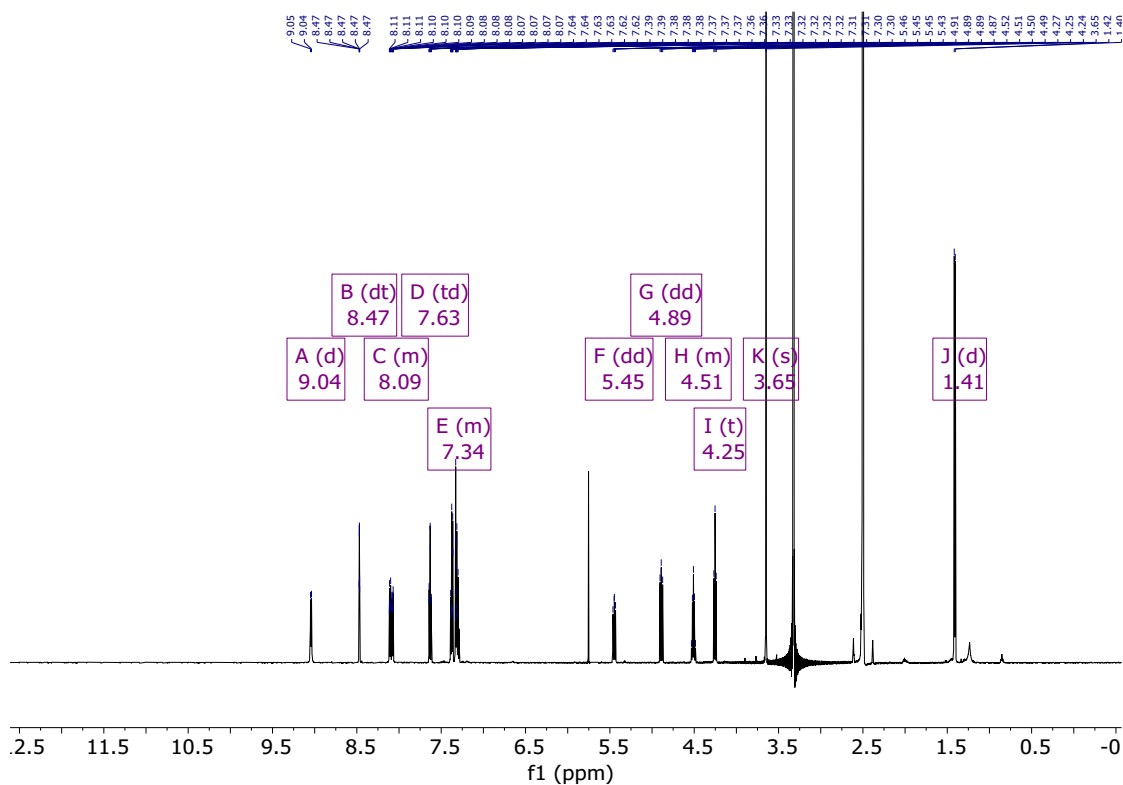
^{13}C NMR (151 MHz, CDCl_3) of **1m4**.



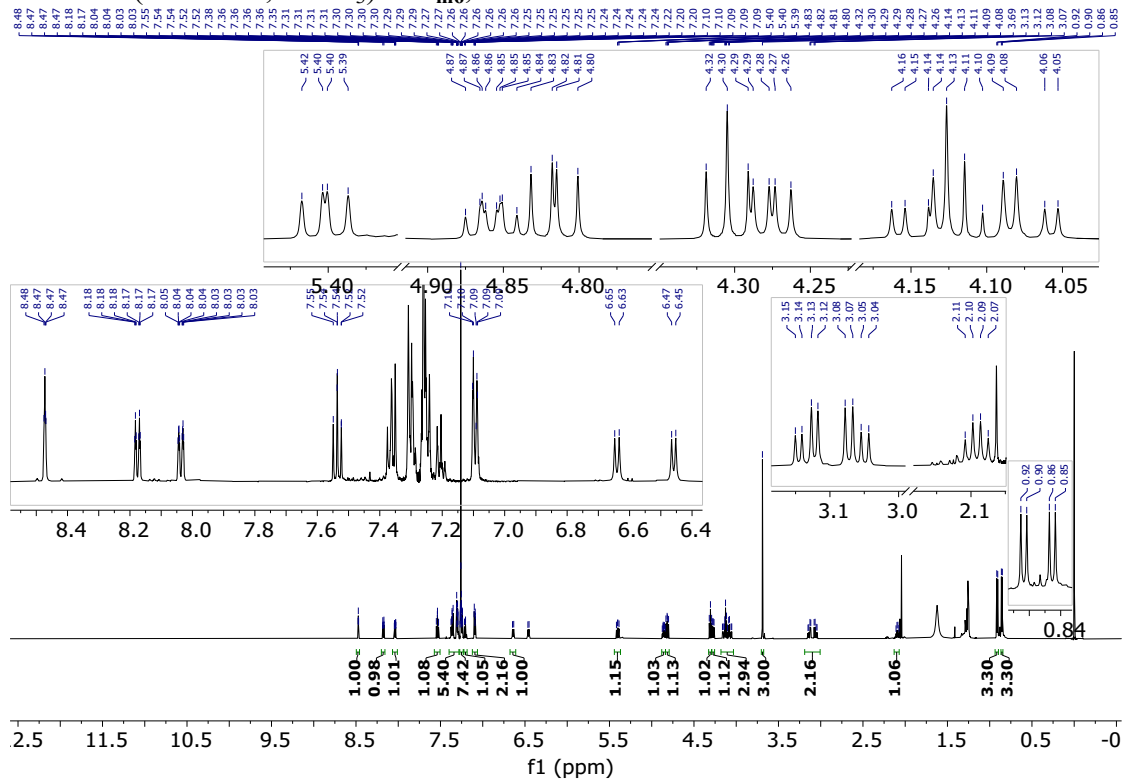
^1H NMR (600 MHz, CDCl_3) of $\mathbf{1}_{m5}$, $c = 60$ mM.



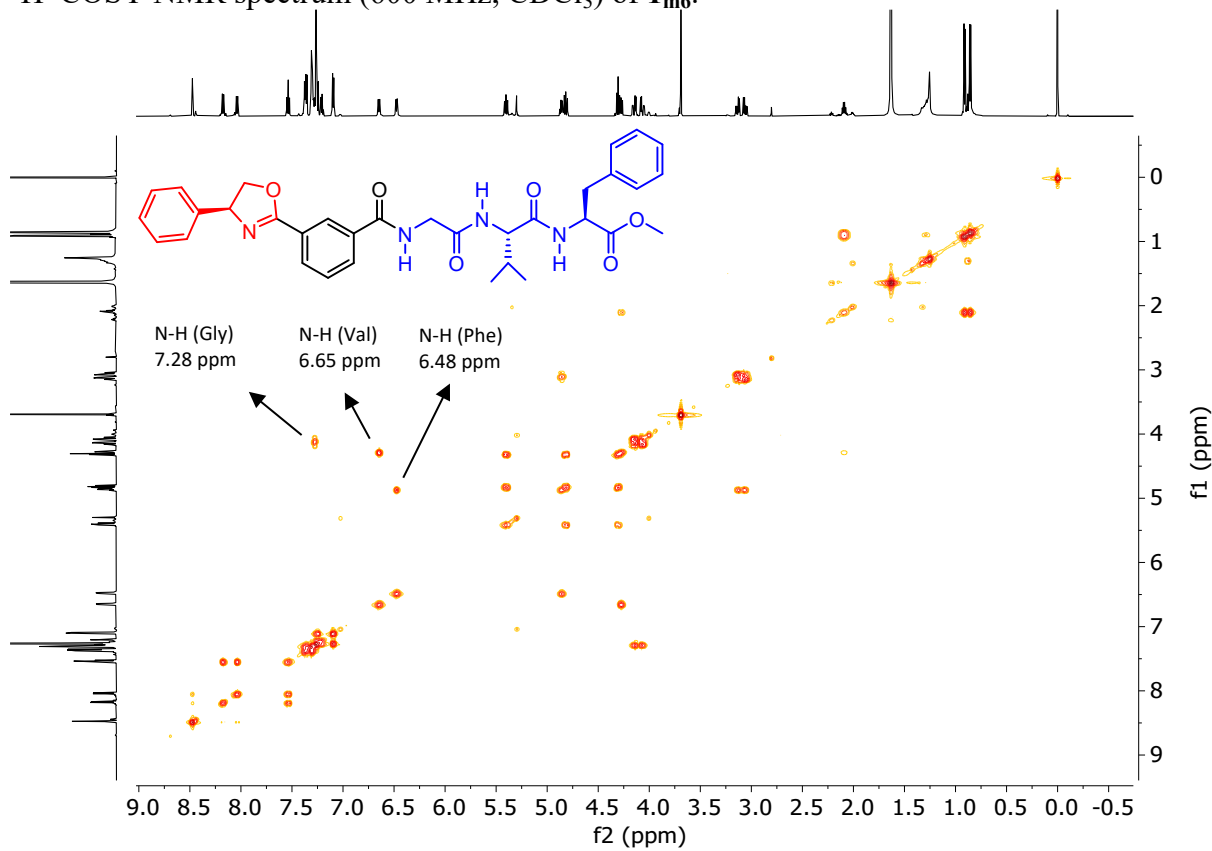
^1H NMR (600 MHz, DMSO) of $\mathbf{1}_{m5}$, $c = 6$ mM.



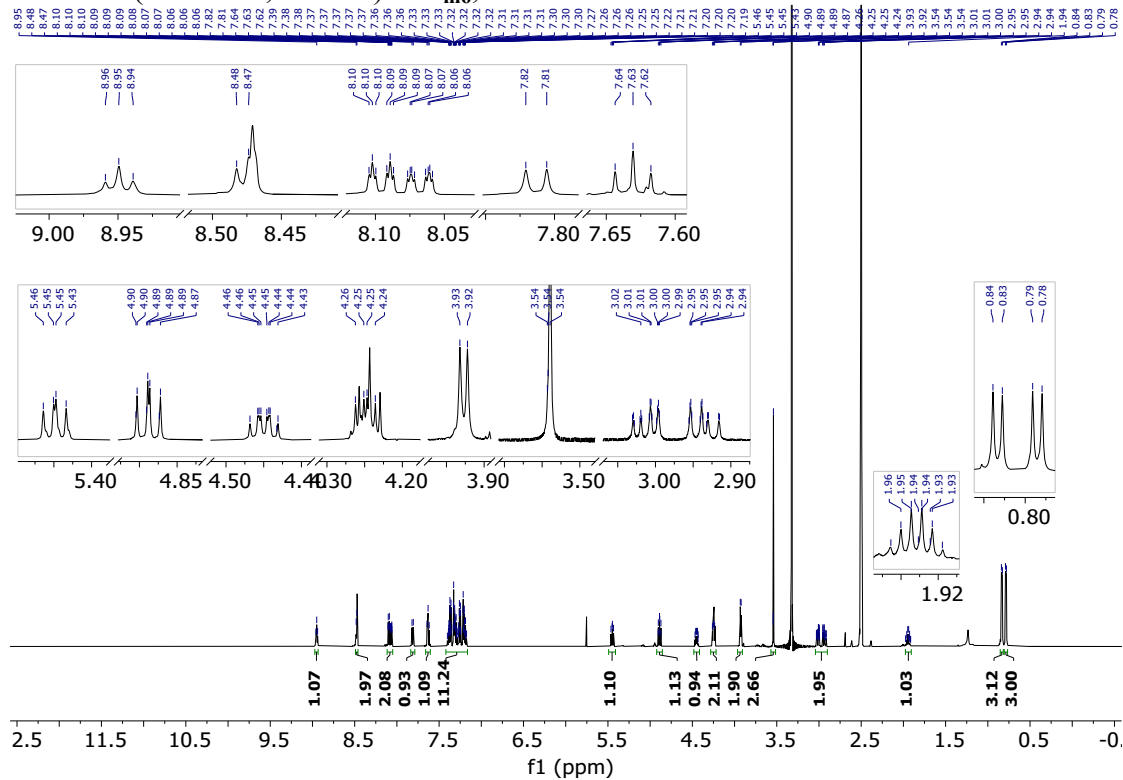
¹H NMR (600 MHz, CDCl₃) of **1_{m6}**, *c* = 6 mM.



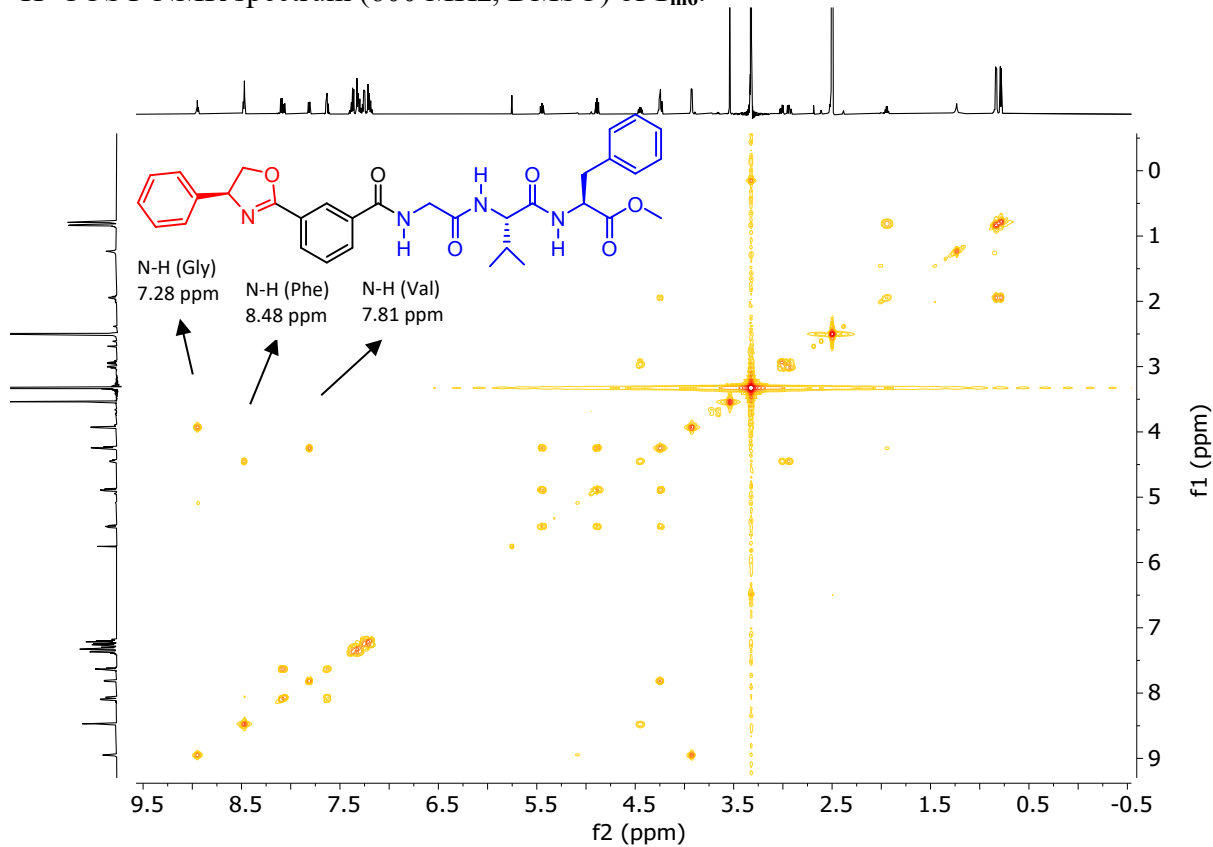
¹H COSY NMR spectrum (600 MHz, CDCl₃) of **1_{m6}**.



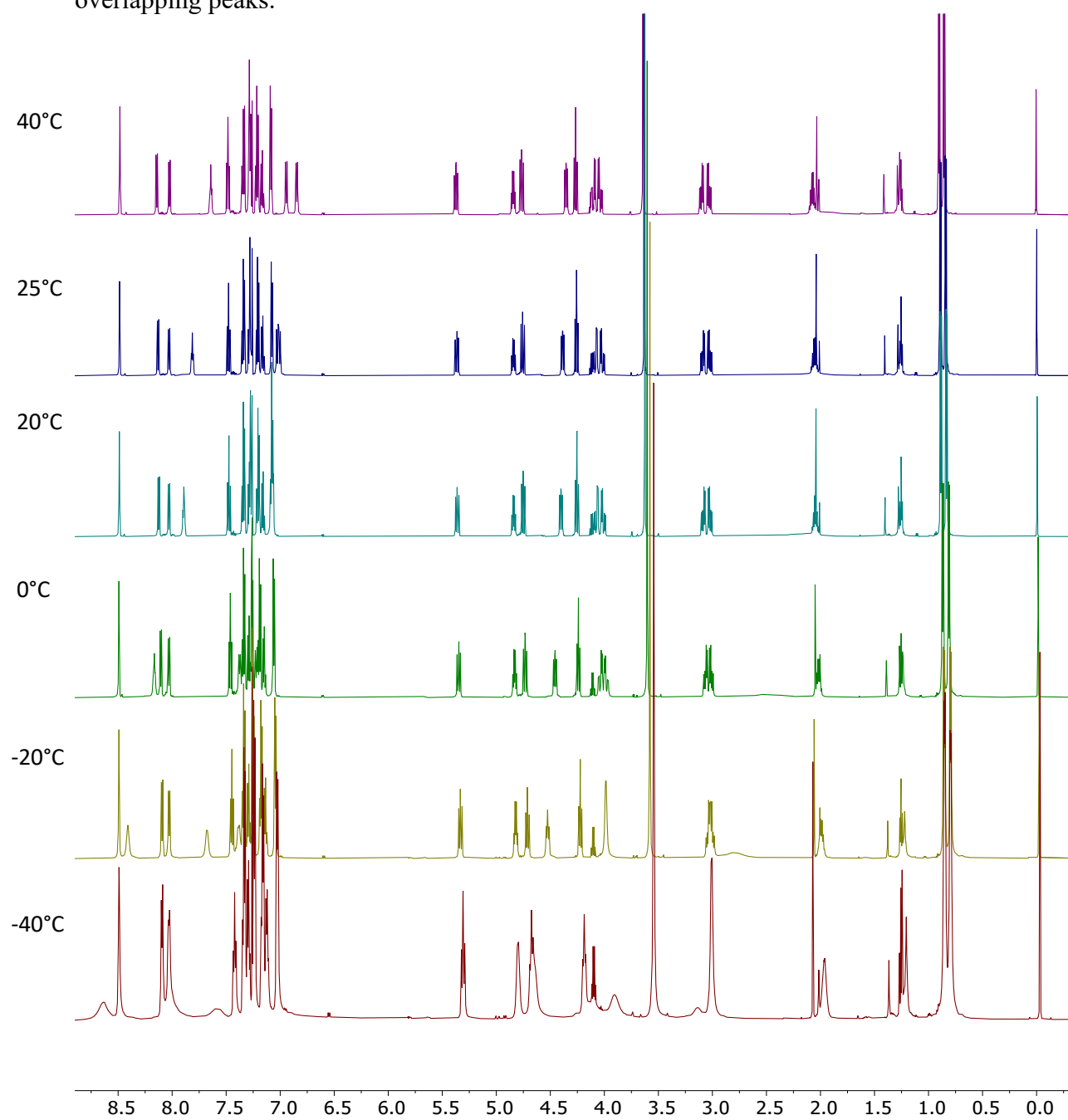
^1H NMR (600 MHz, DMSO) of $\mathbf{1}_{m6}$, $c = 6$ mM.



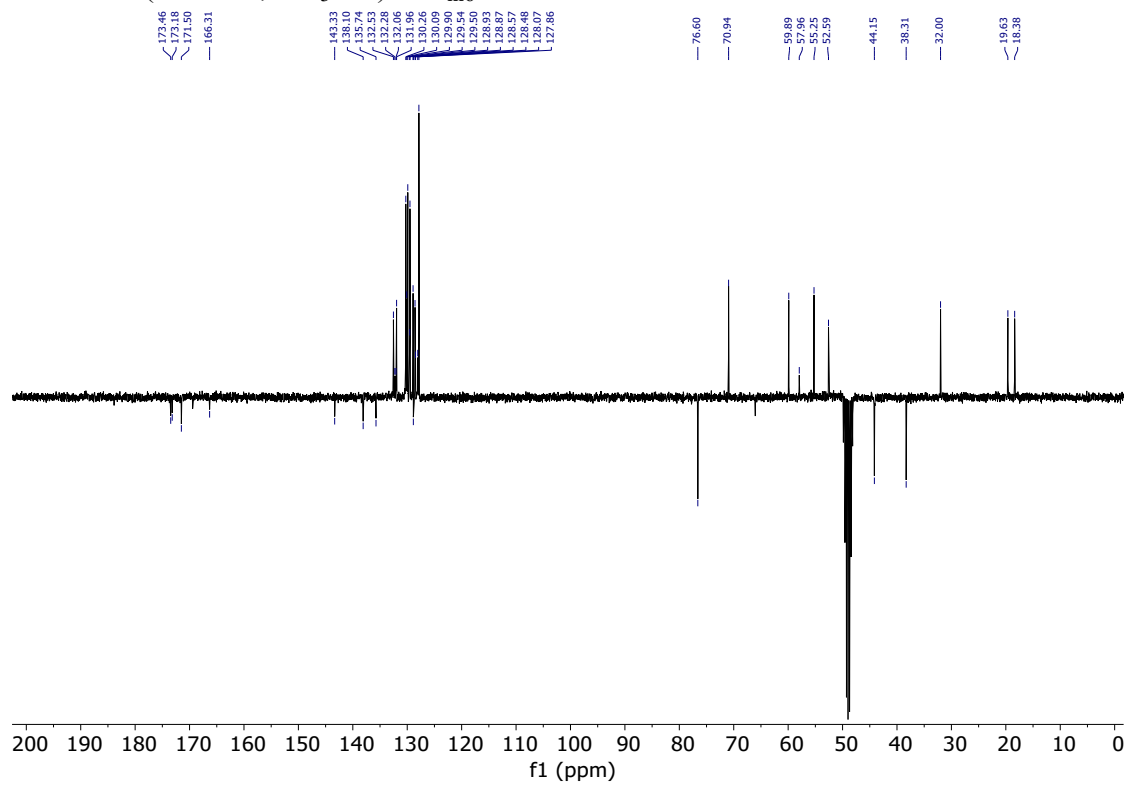
^1H COSY NMR spectrum (600 MHz, DMSO) of $\mathbf{1}_{m6}$.



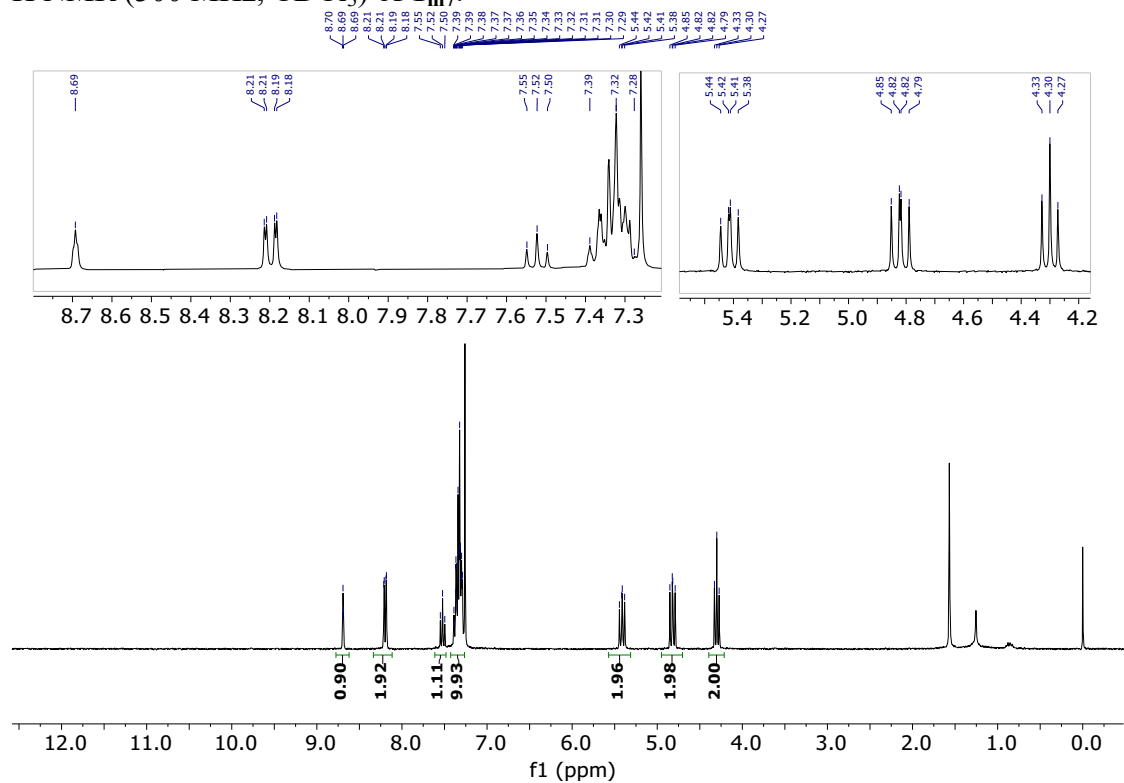
Temperature dependent ^1H NMR spectra of oxazoline $\mathbf{1}_{m6}$, at $c = 60$ mM. For each temperature a COSY spectrum was recorded as well to determine the chemical shift of overlapping peaks.



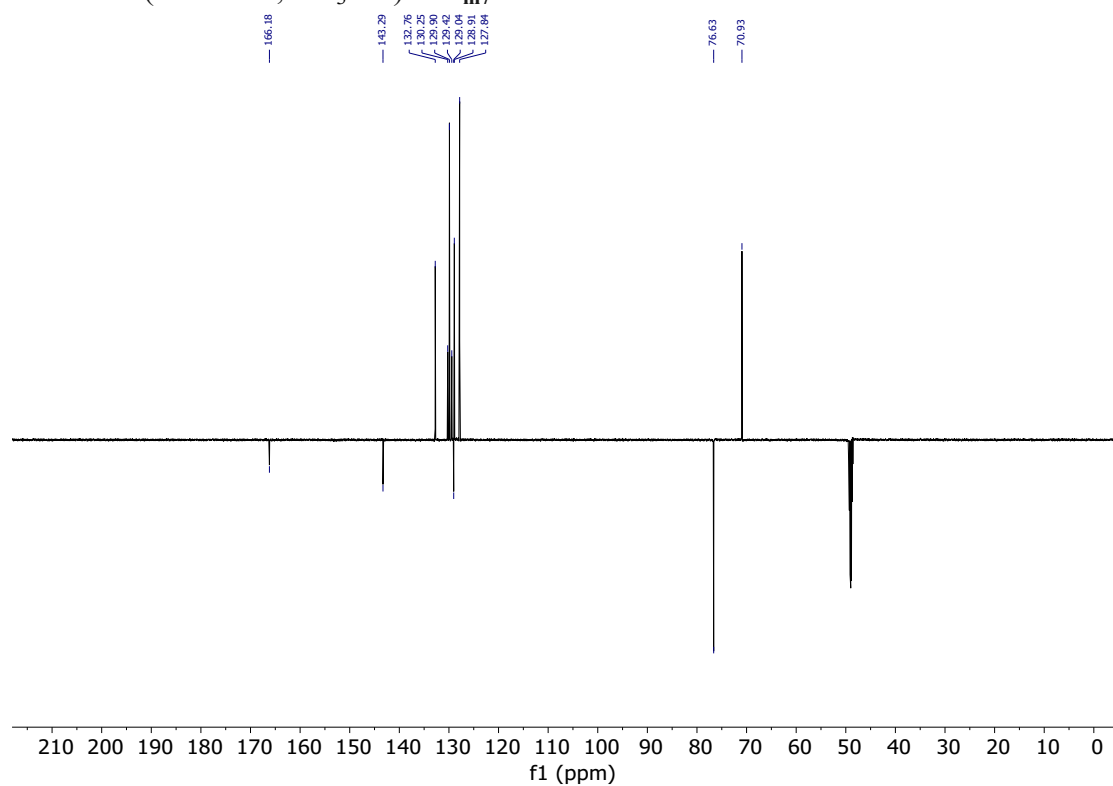
^{13}C NMR (75 MHz, CD_3OD) of **1_{m6}**.



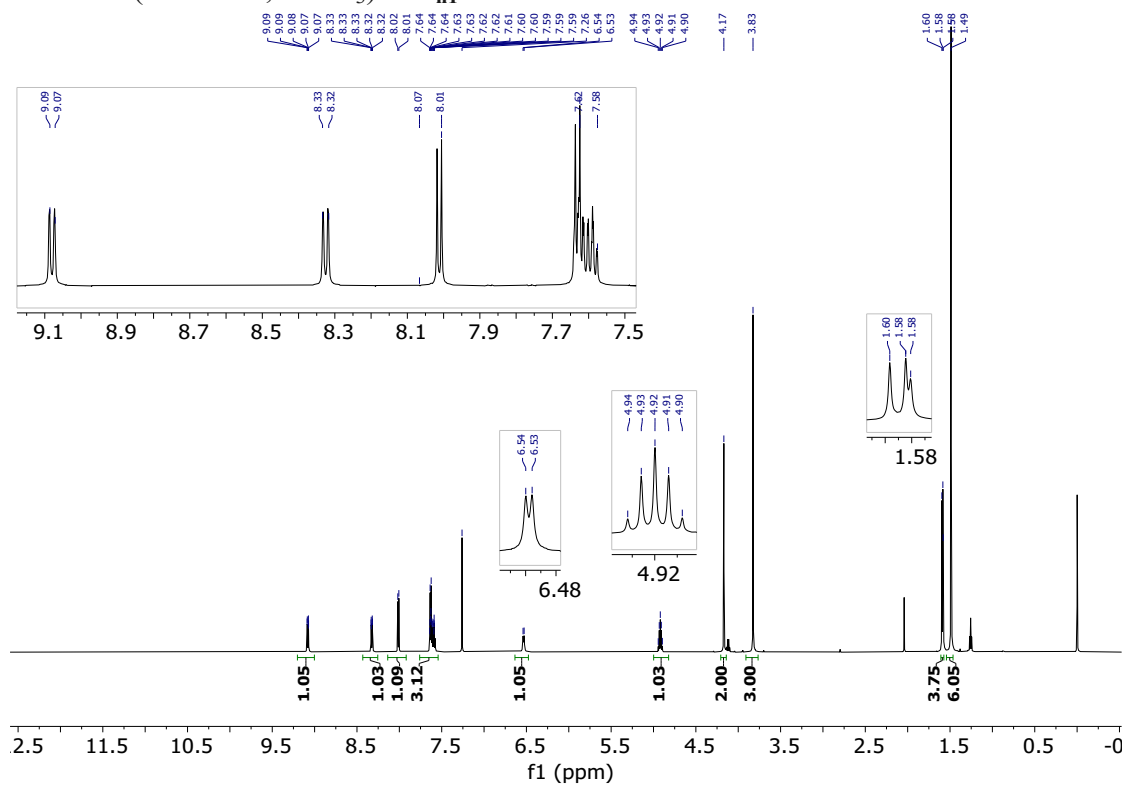
^1H NMR (300 MHz, CDCl_3) of 1_{m7} .



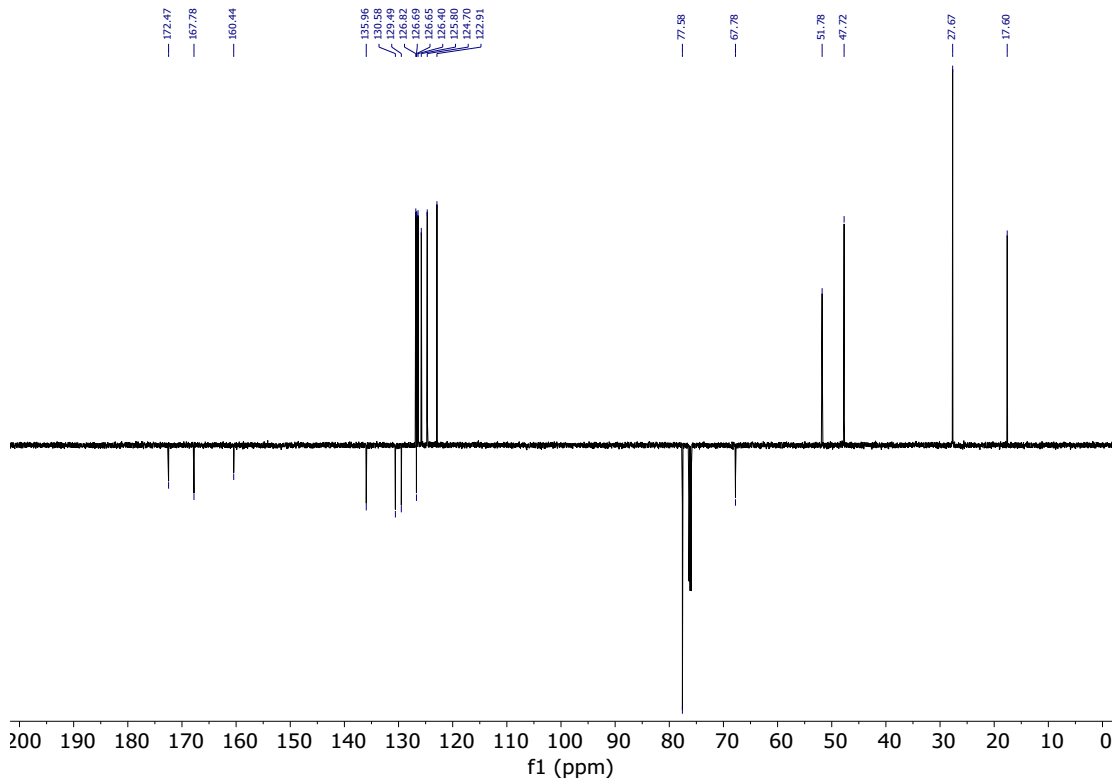
^{13}C NMR (151 MHz, CD_3OD) of 1_{m7} .



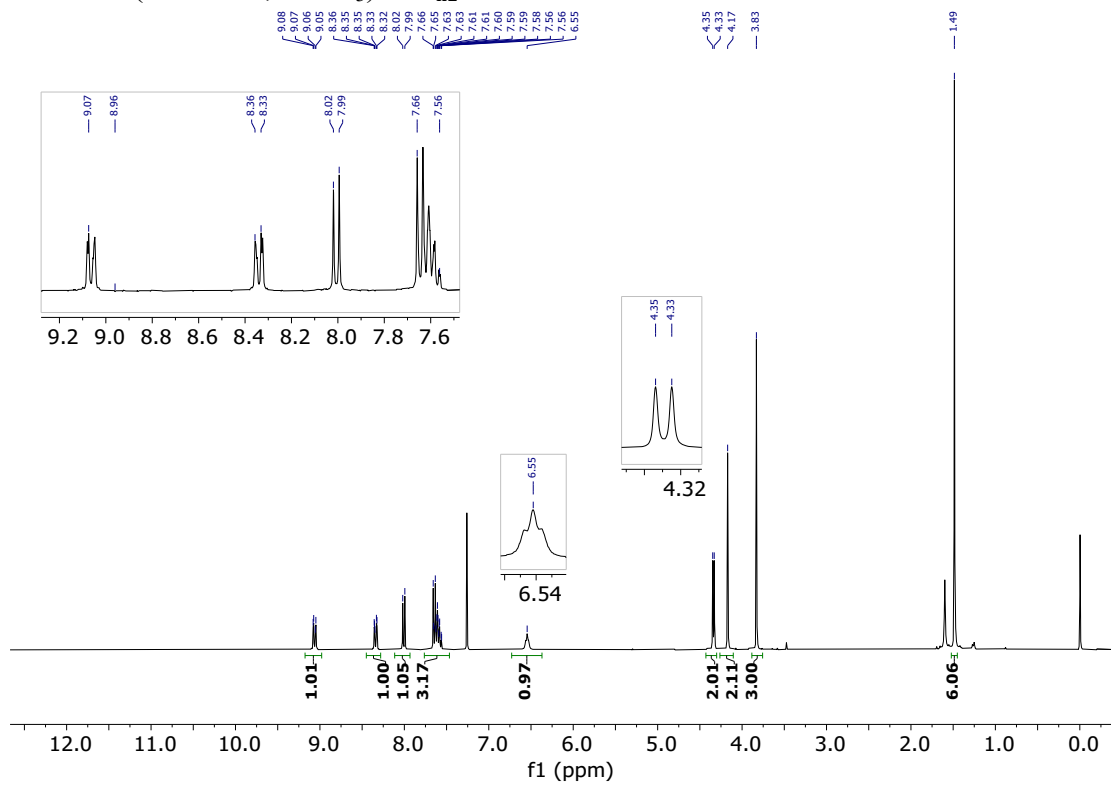
^1H NMR (600 MHz, CDCl_3) of **1n1**.



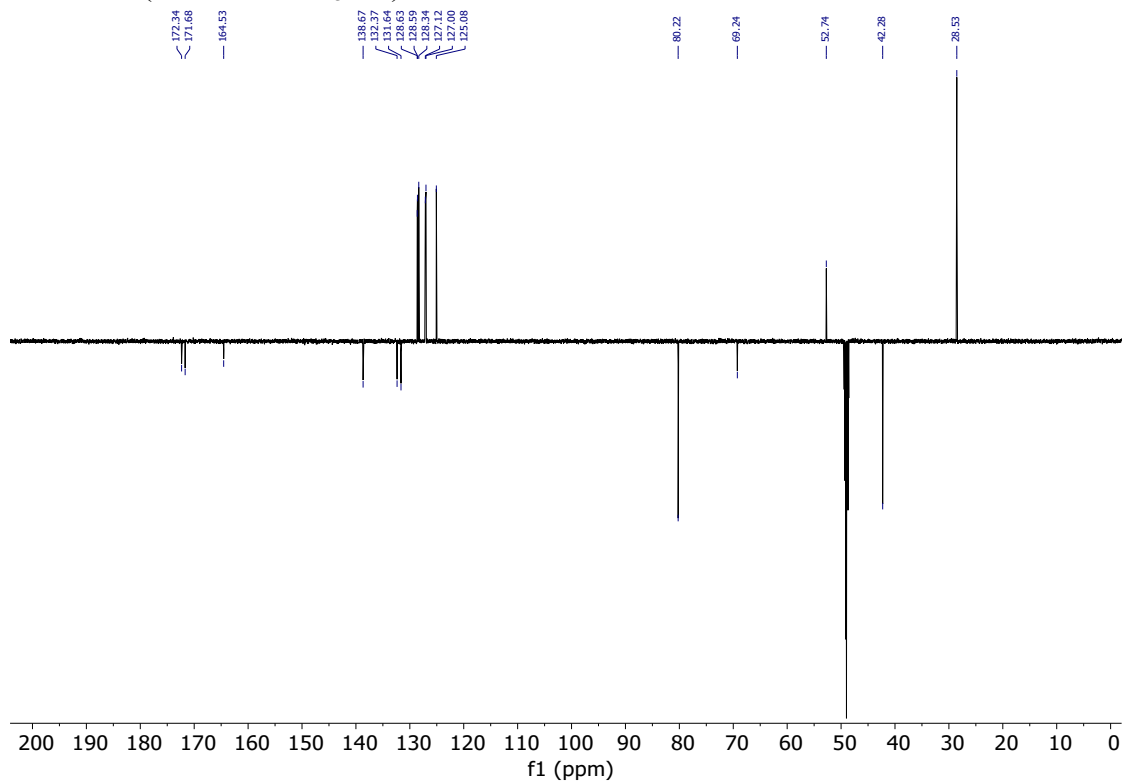
^{13}C NMR (151 MHz, CDCl_3) of **1n1**.



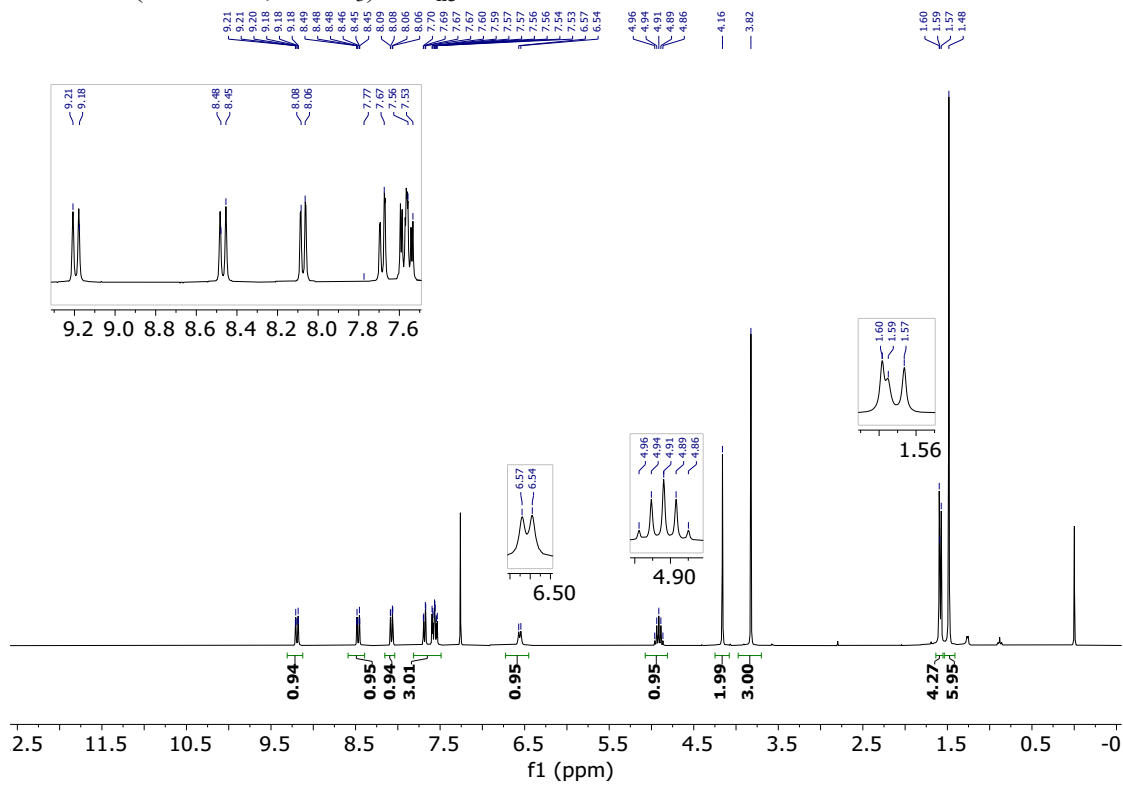
^1H NMR (300 MHz, CDCl_3) of **1n2**.



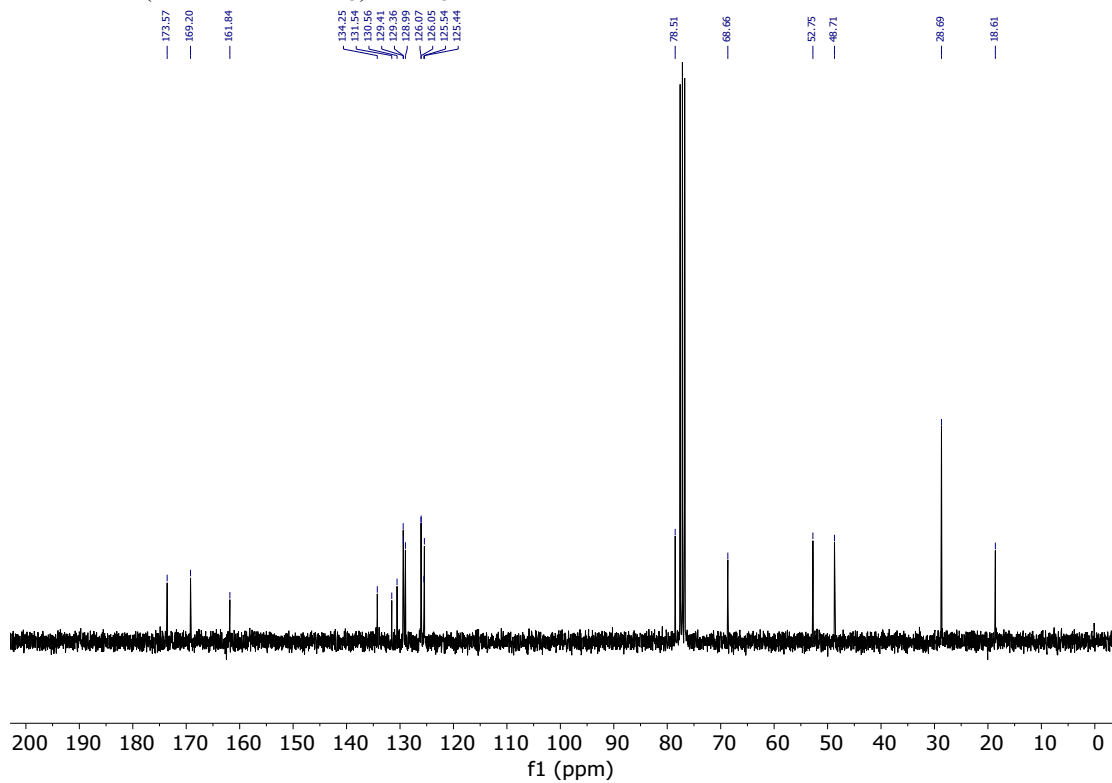
^{13}C NMR (151 MHz, CD_3OD) of **1n2**.



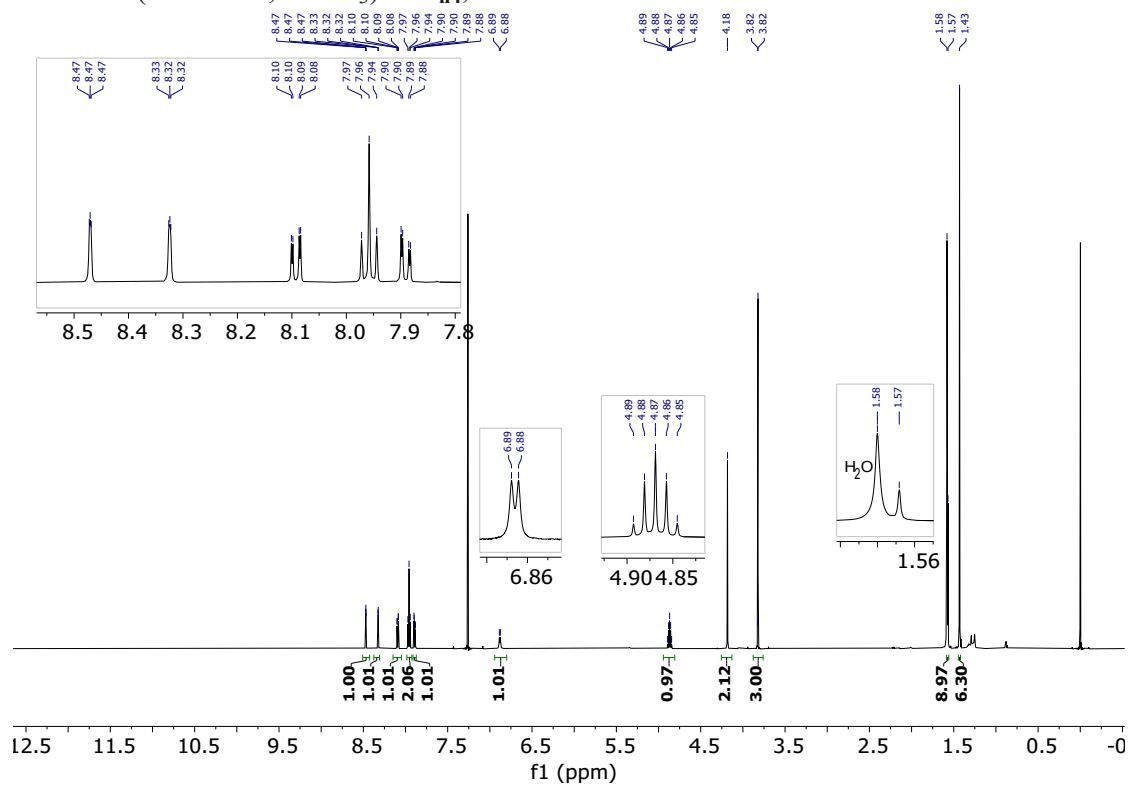
^1H NMR (300 MHz, CDCl_3) of $\mathbf{1n3}$.



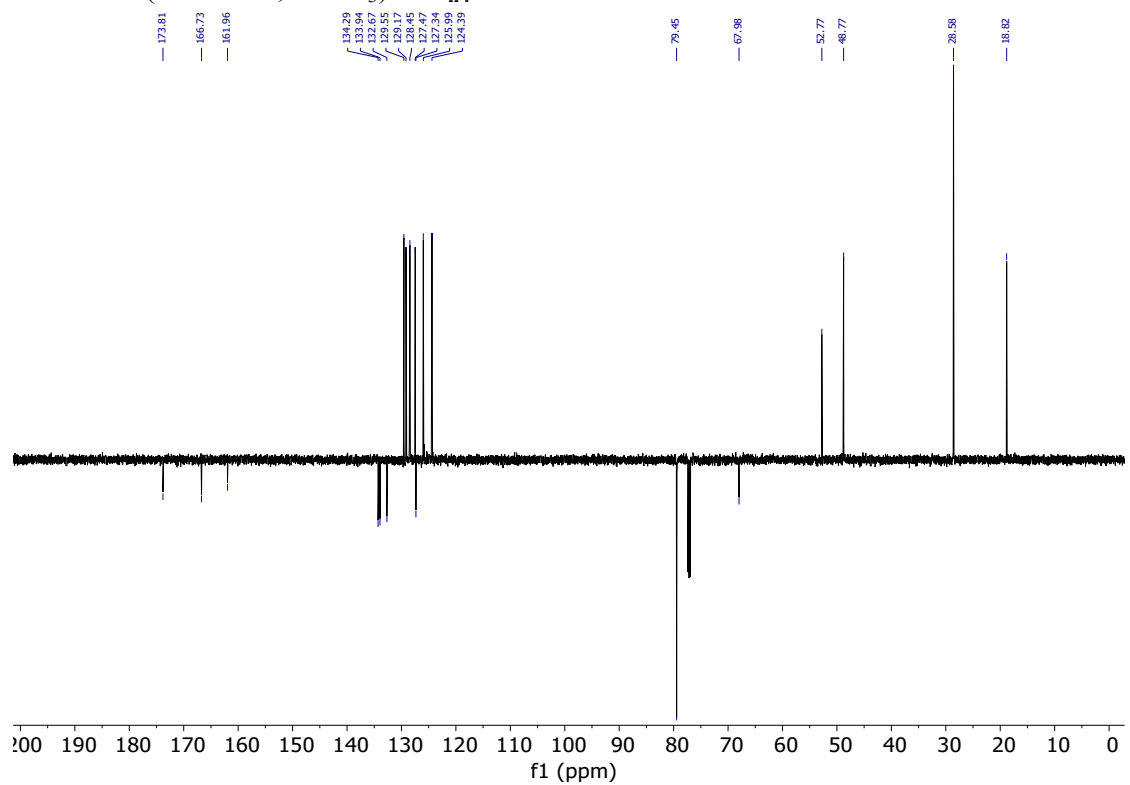
^{13}C NMR (75 MHz, CDCl_3) of $\mathbf{1n3}$.



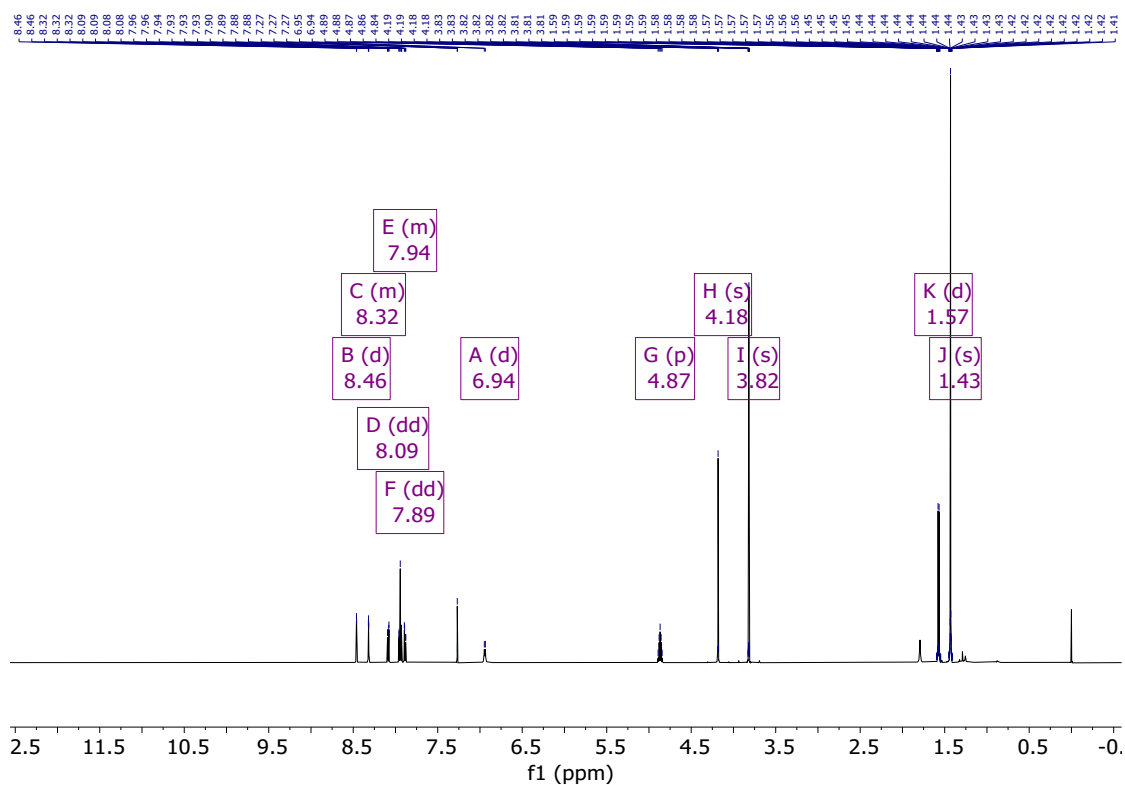
^1H NMR (600 MHz, CDCl_3) of $\mathbf{1}_{n4}$, $c = 6\text{mM}$.



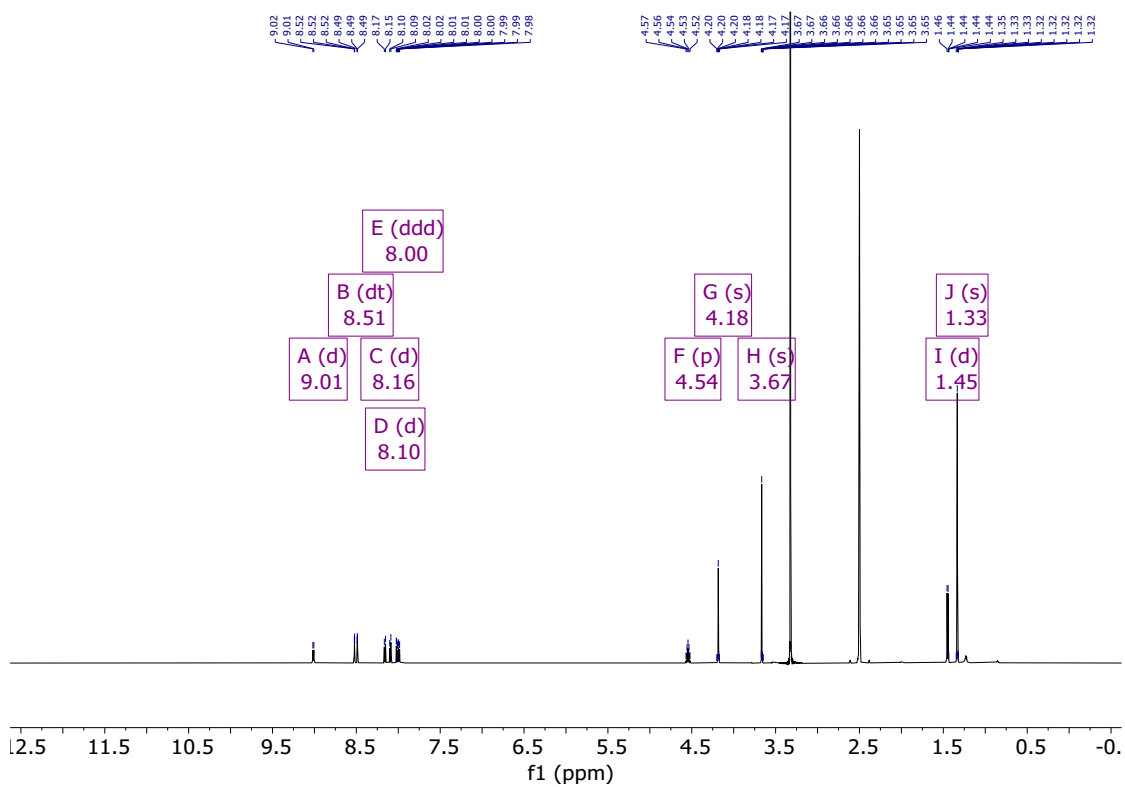
^{13}C NMR (151 MHz, CDCl_3) of $\mathbf{1}_{n4}$.



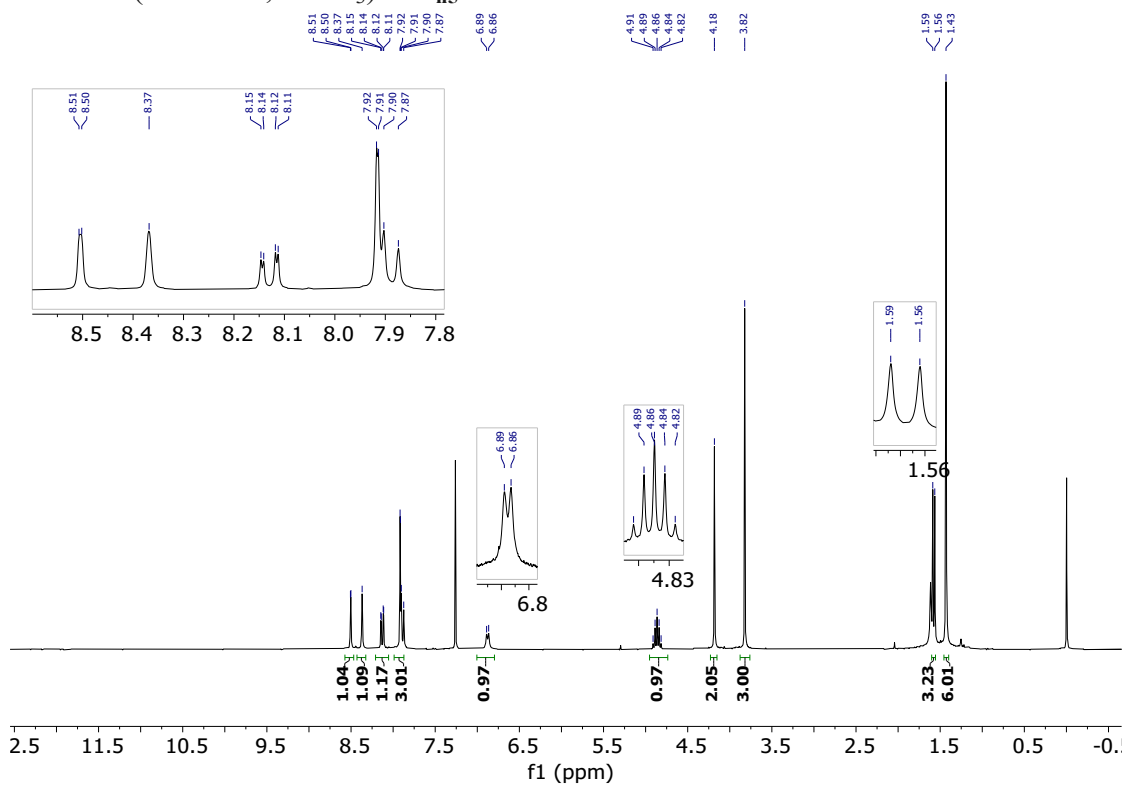
^1H NMR (600 MHz, CDCl_3) of $\mathbf{1}_{n4}$, $c = 60$ mM.



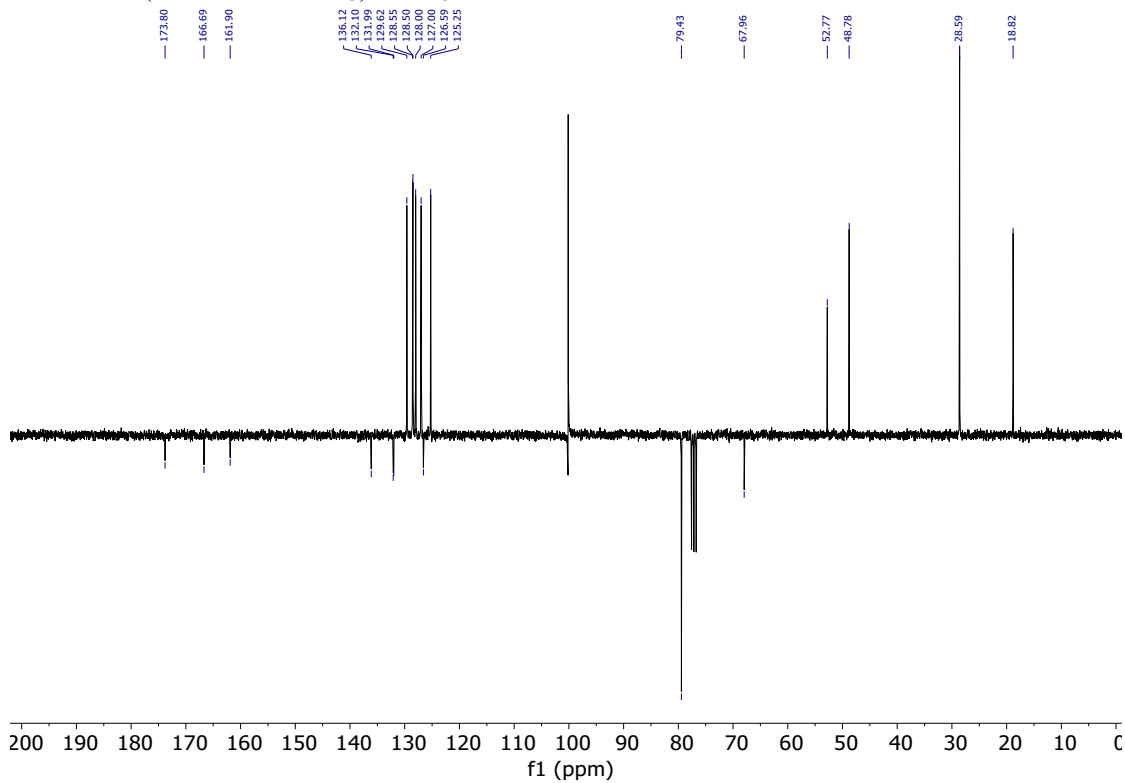
^1H NMR (600 MHz, DMSO) of $\mathbf{1}_{n4}$, $c = 6$ mM.



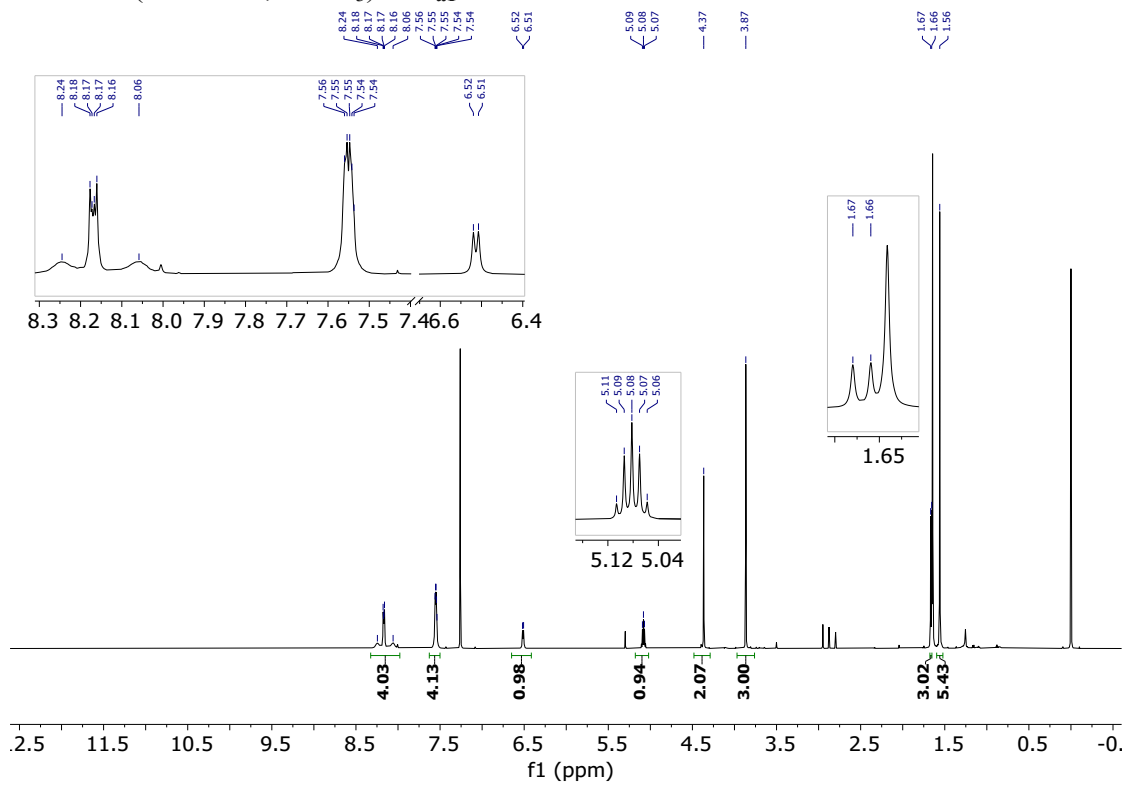
^1H NMR (300 MHz, CDCl_3) of **1n5**.



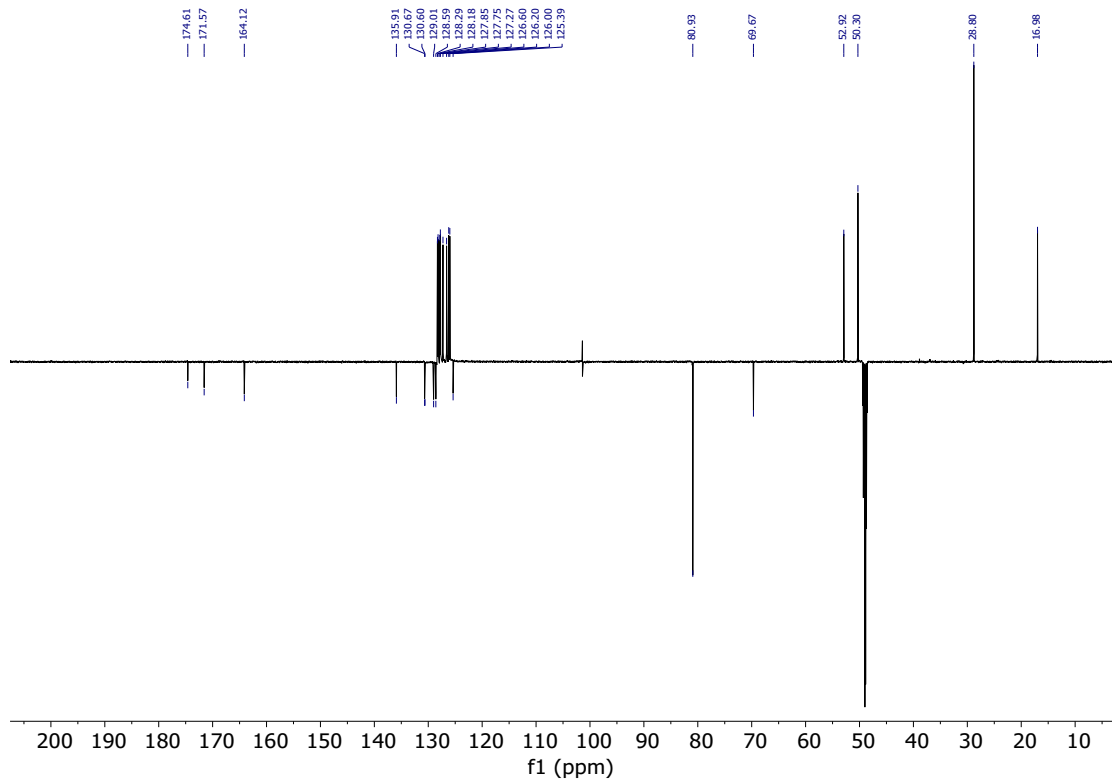
^{13}C NMR (75 MHz, CDCl_3) of **1n5**.



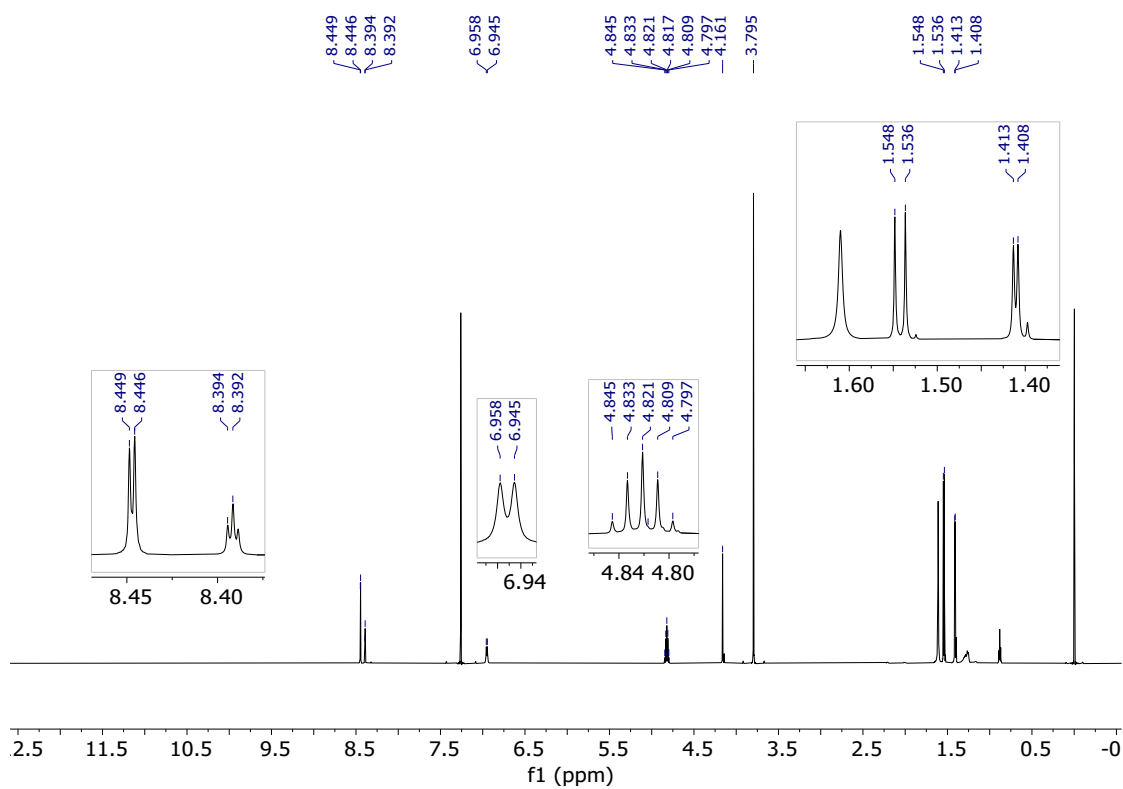
^1H NMR (600 MHz, CDCl_3) of **1a1**.



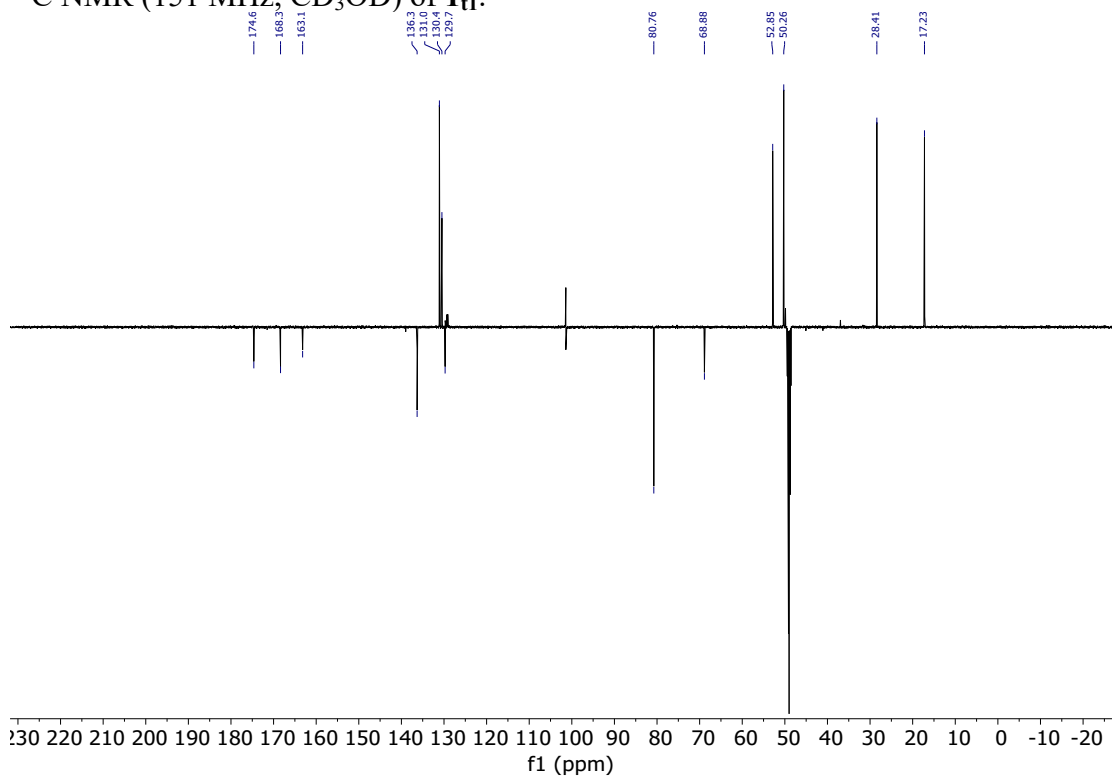
^{13}C NMR (151 MHz, CDCl_3) of **1a1**.



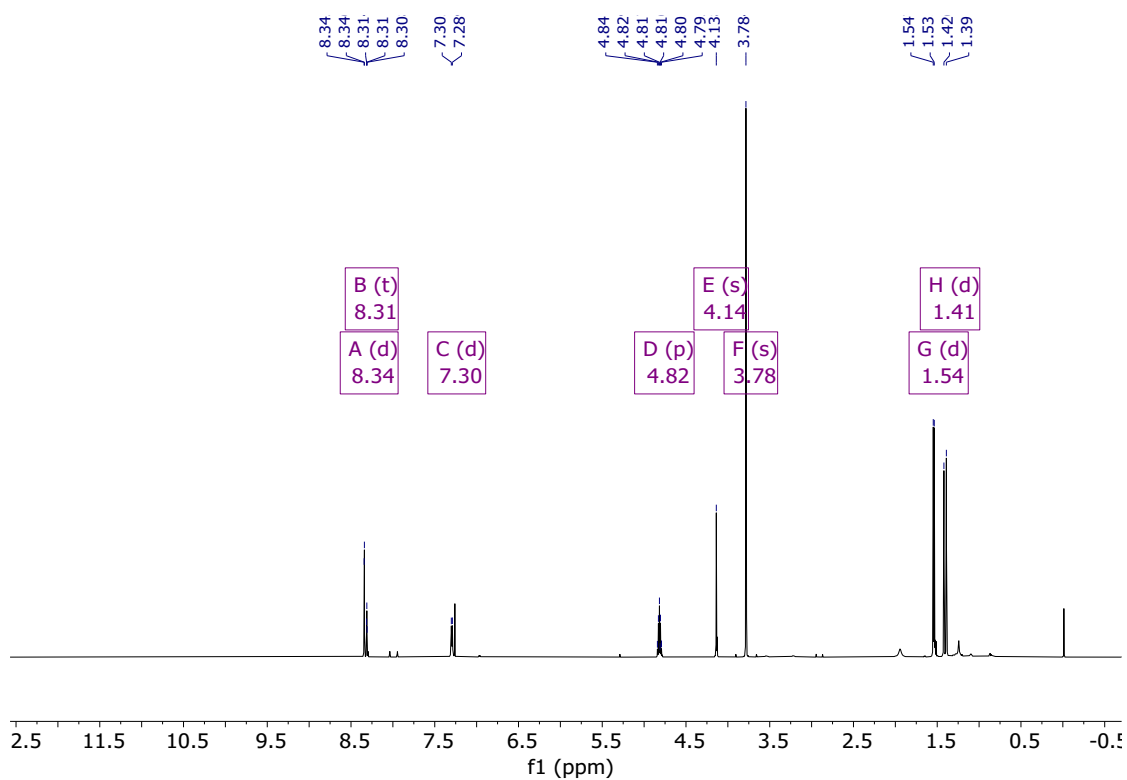
^1H NMR (600 MHz, CDCl_3) of $\mathbf{1}_{\text{tl}}$, $c = 6 \text{ mM}$



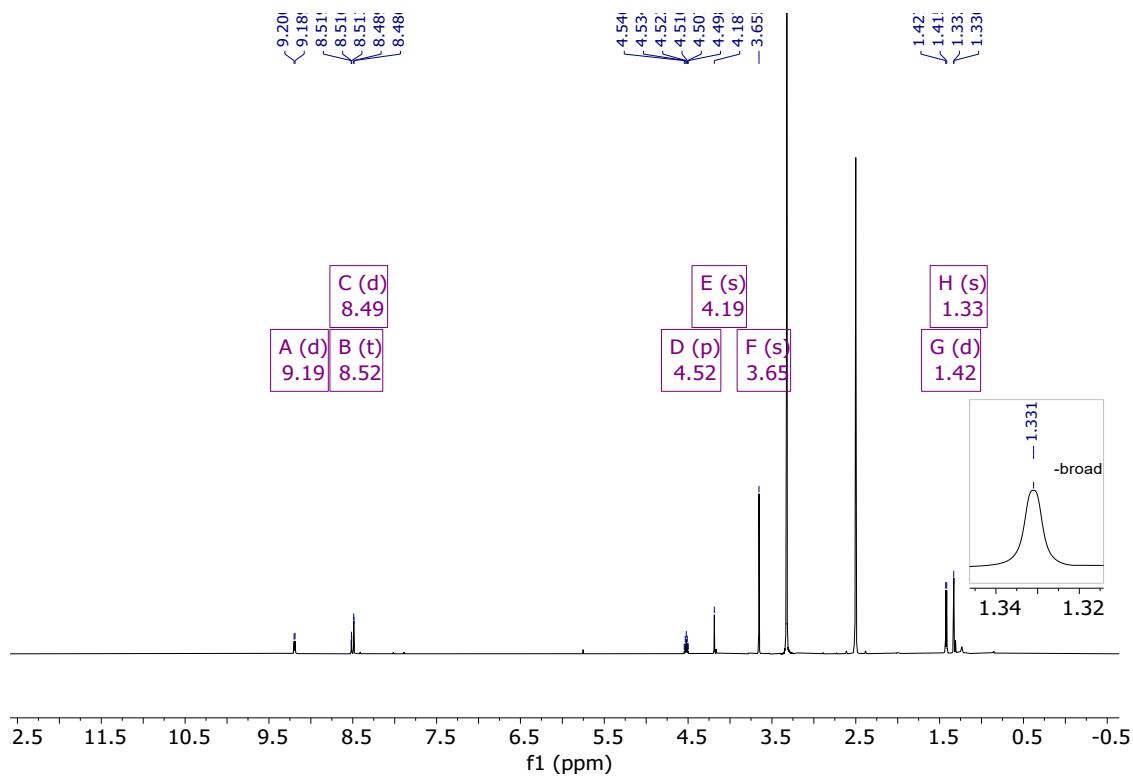
^{13}C NMR (151 MHz, CD_3OD) of $\mathbf{1}_{\text{tl}}$.



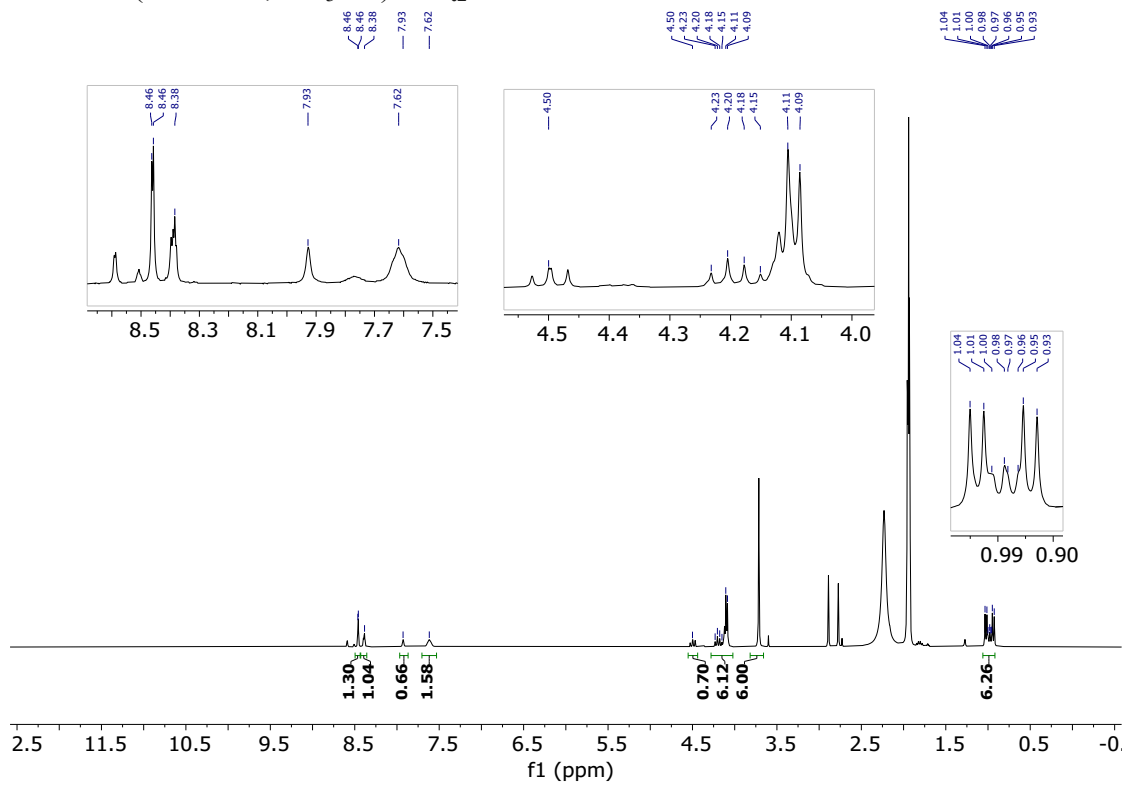
^1H NMR (600 MHz, CDCl_3) of $\mathbf{1}_{\text{t1}}$, $c = 60 \text{ mM}$



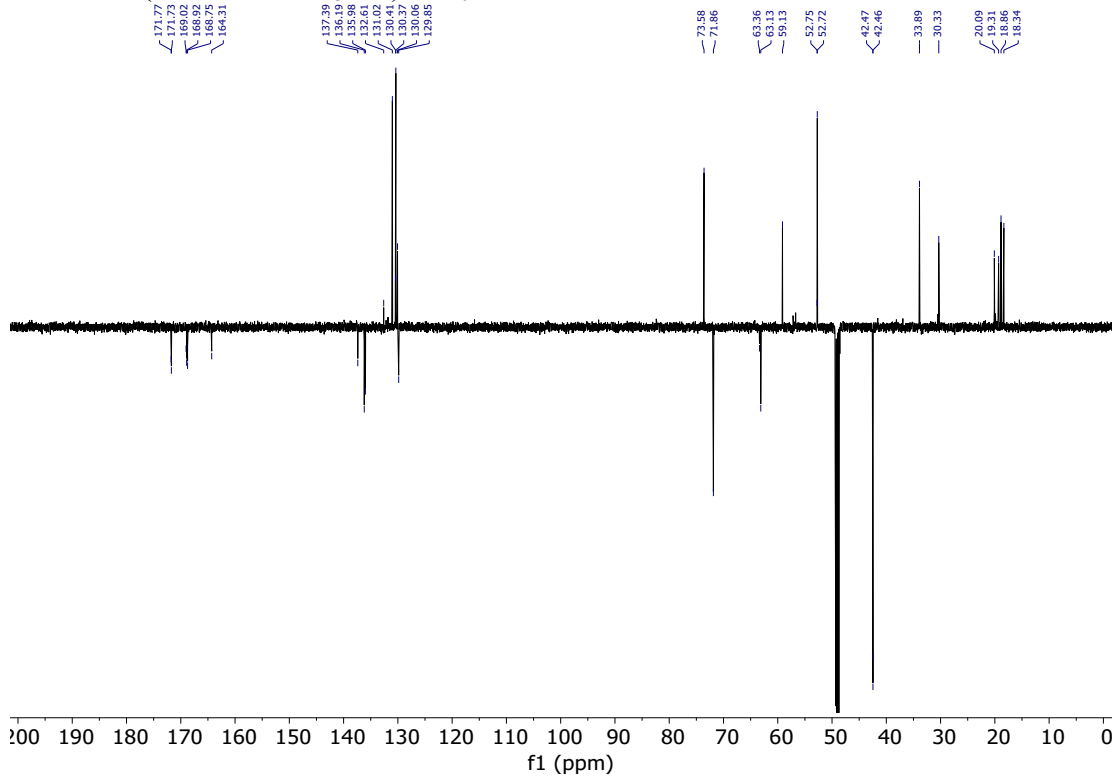
^1H NMR (600 MHz, DMSO) of $\mathbf{1}_{\text{t1}}$, $c = 6 \text{ mM}$



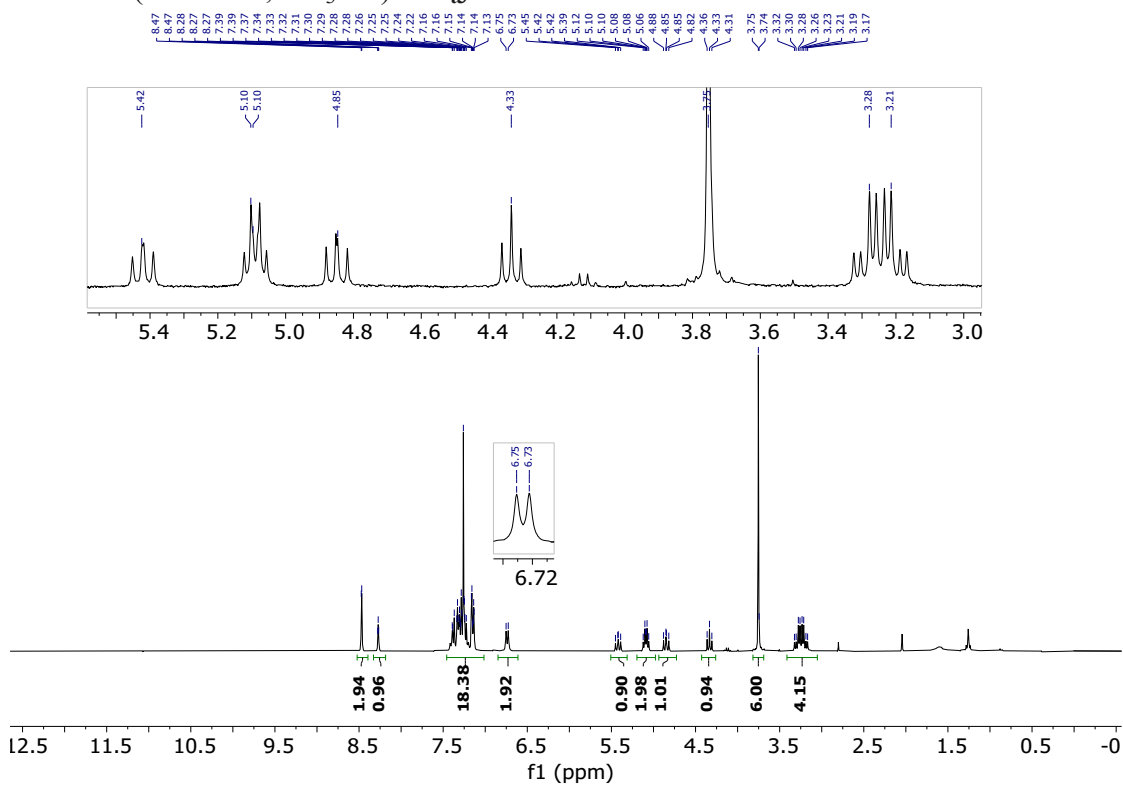
^1H NMR (300 MHz, CD_3CN) of $\mathbf{1}_2$.



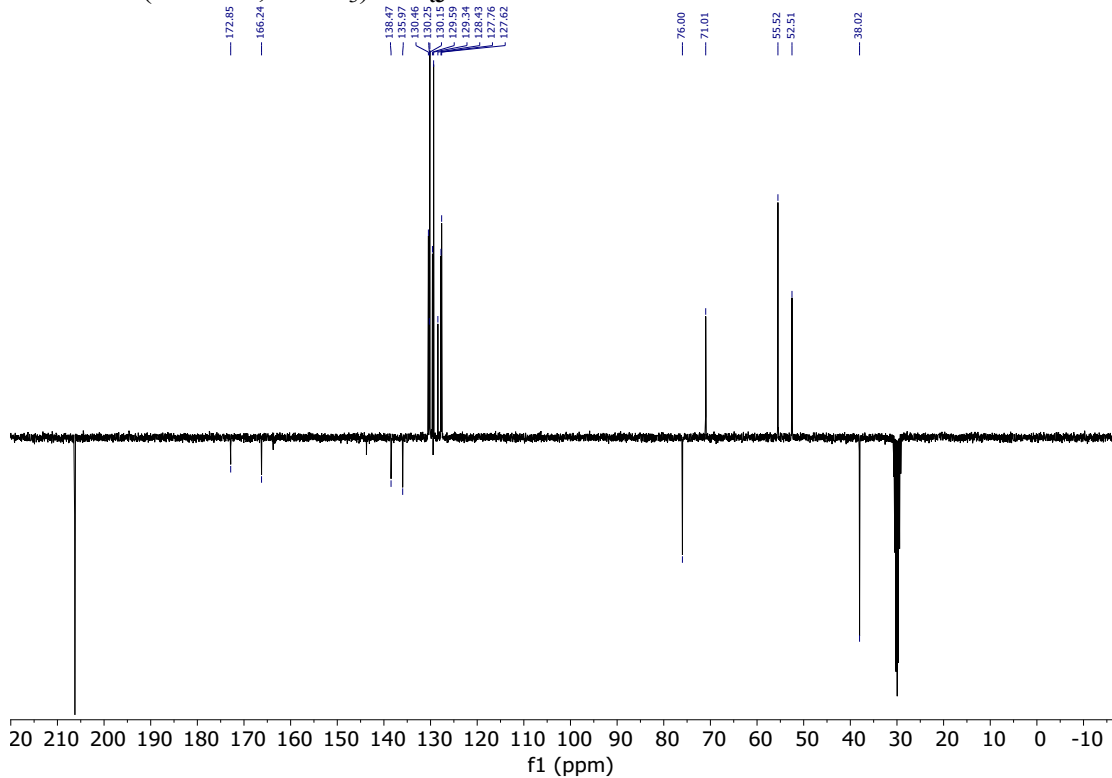
^{13}C NMR (151 MHz, CD_3OD) of $\mathbf{1}_2$.



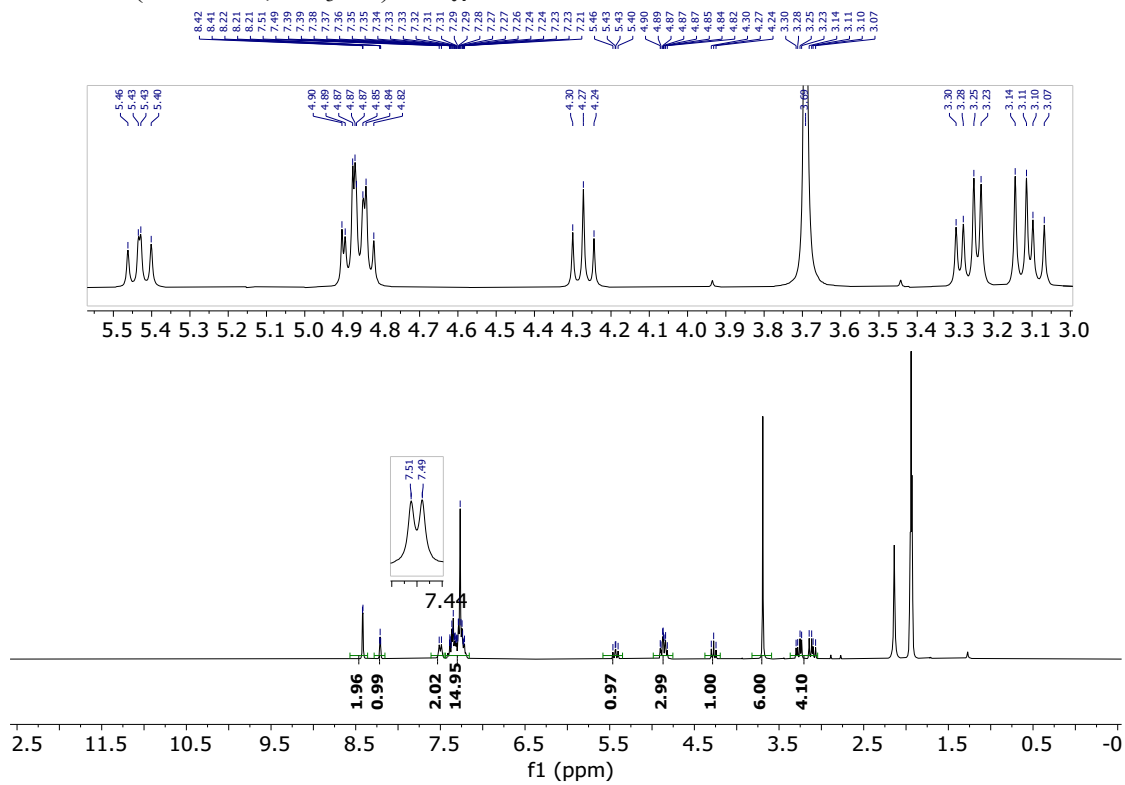
^1H NMR (300 MHz, CD_3CN) of **1₃**.



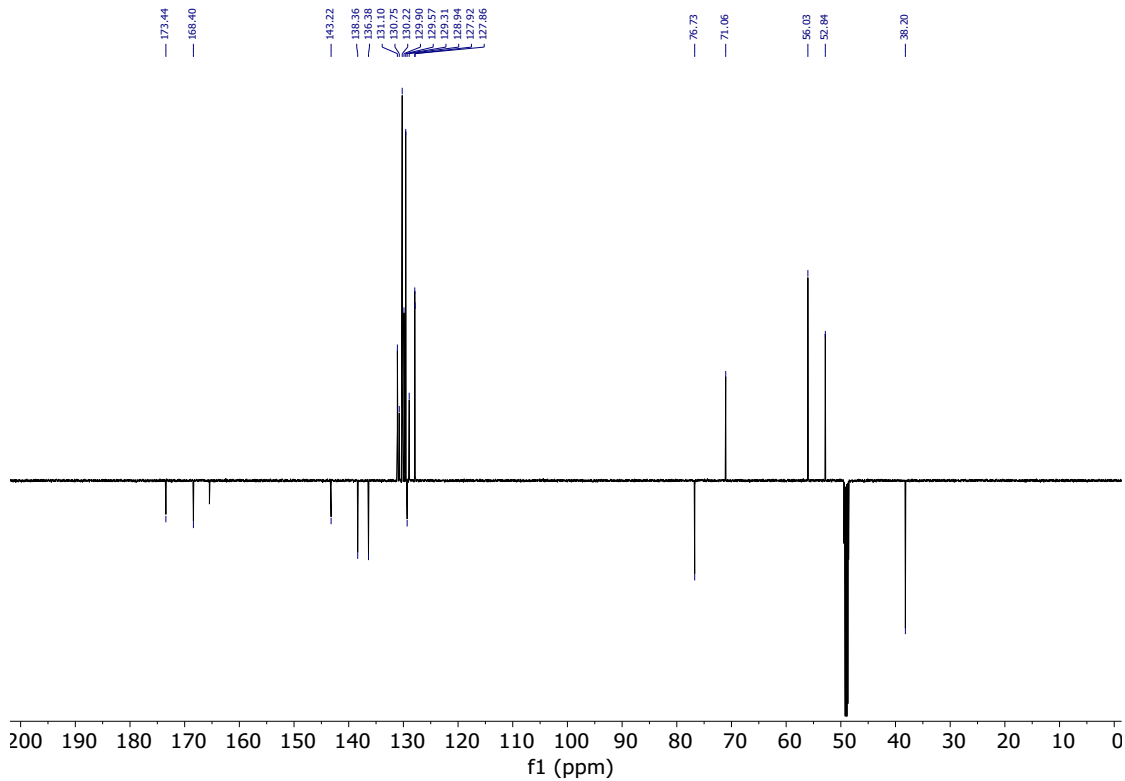
^{13}C NMR (75 MHz, CDCl_3) of **1₃**.



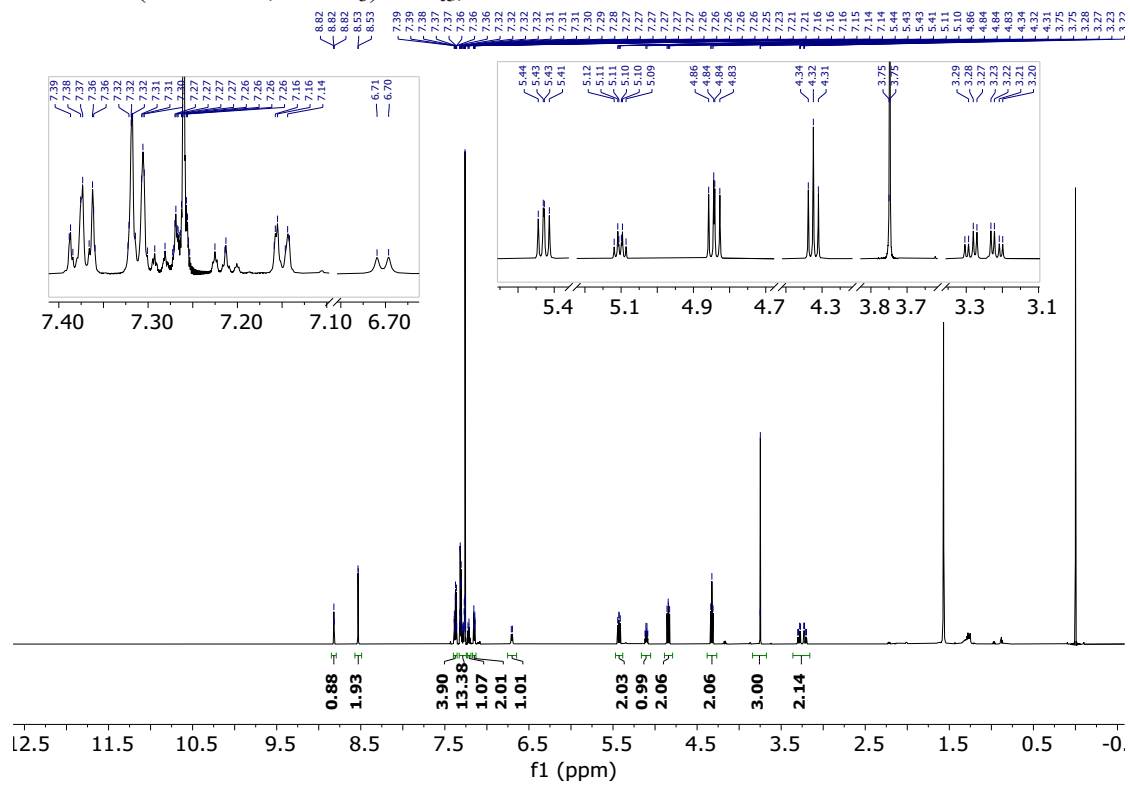
^1H NMR (300 MHz, CD_3CN) of **1_{t4}**.



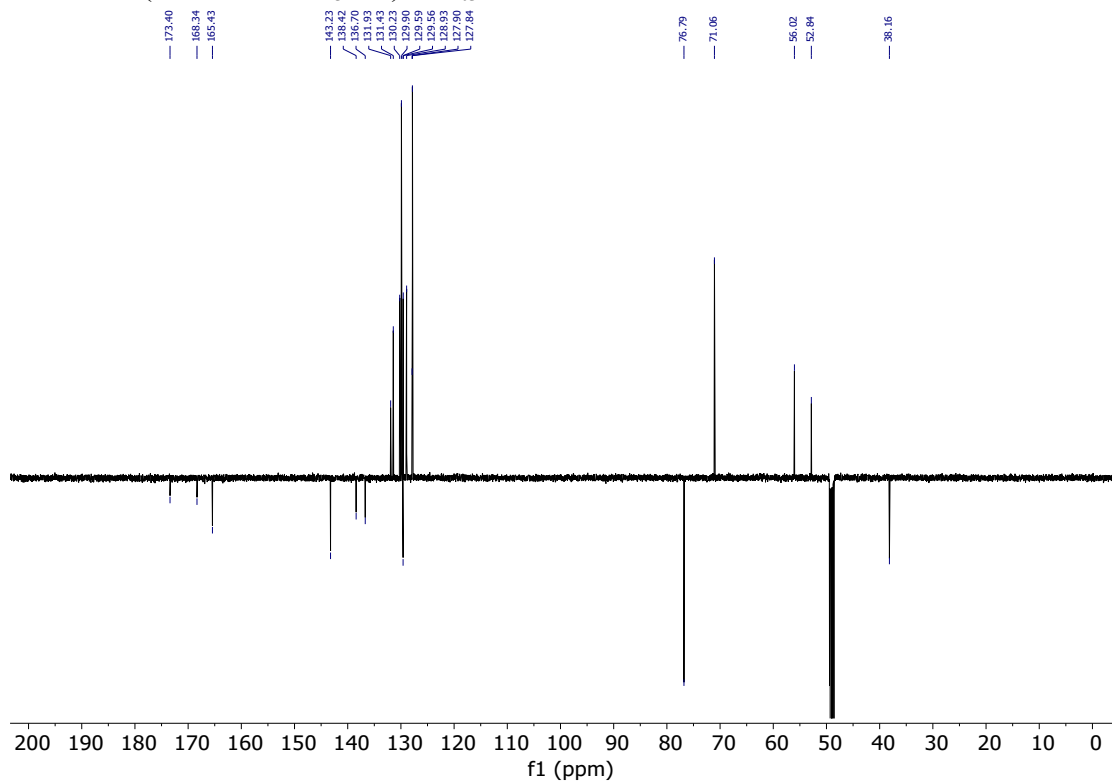
^{13}C NMR (151 MHz, CD_3OD) of **1_{t4}**.



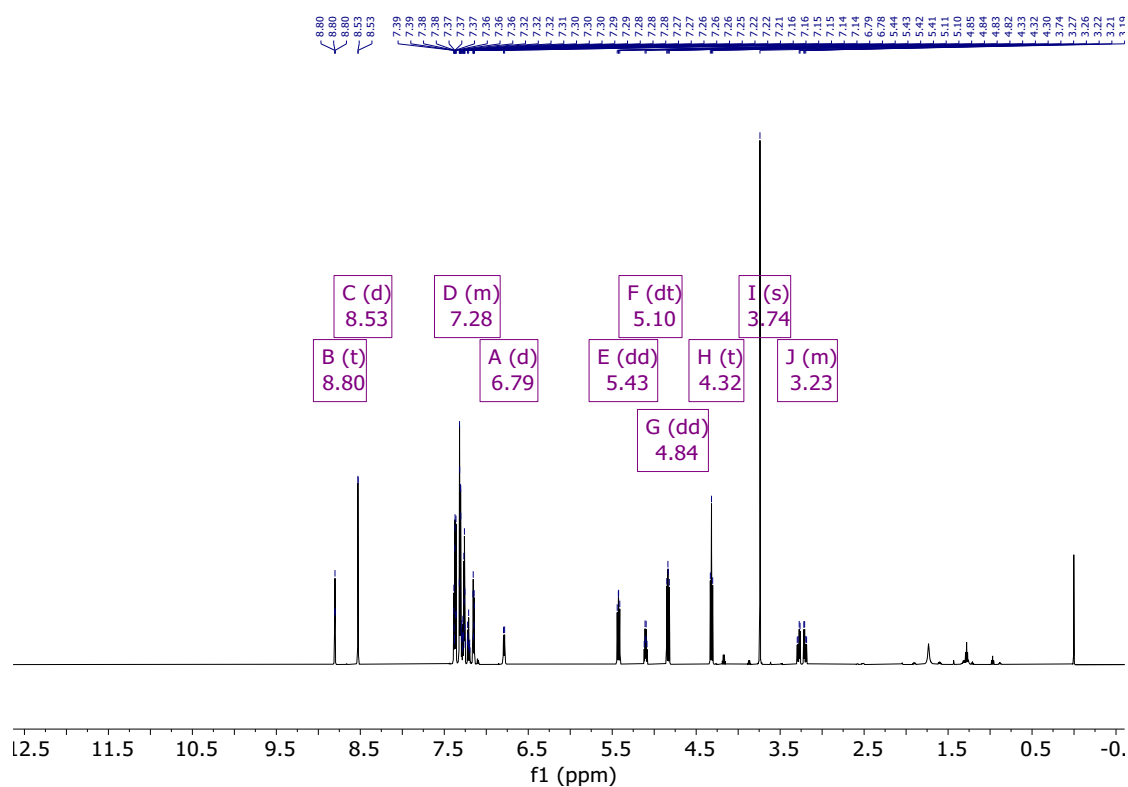
^1H NMR (600 MHz, CDCl_3) of $\mathbf{1}_{\mathbf{t5}}$, $c = 6$ mM.



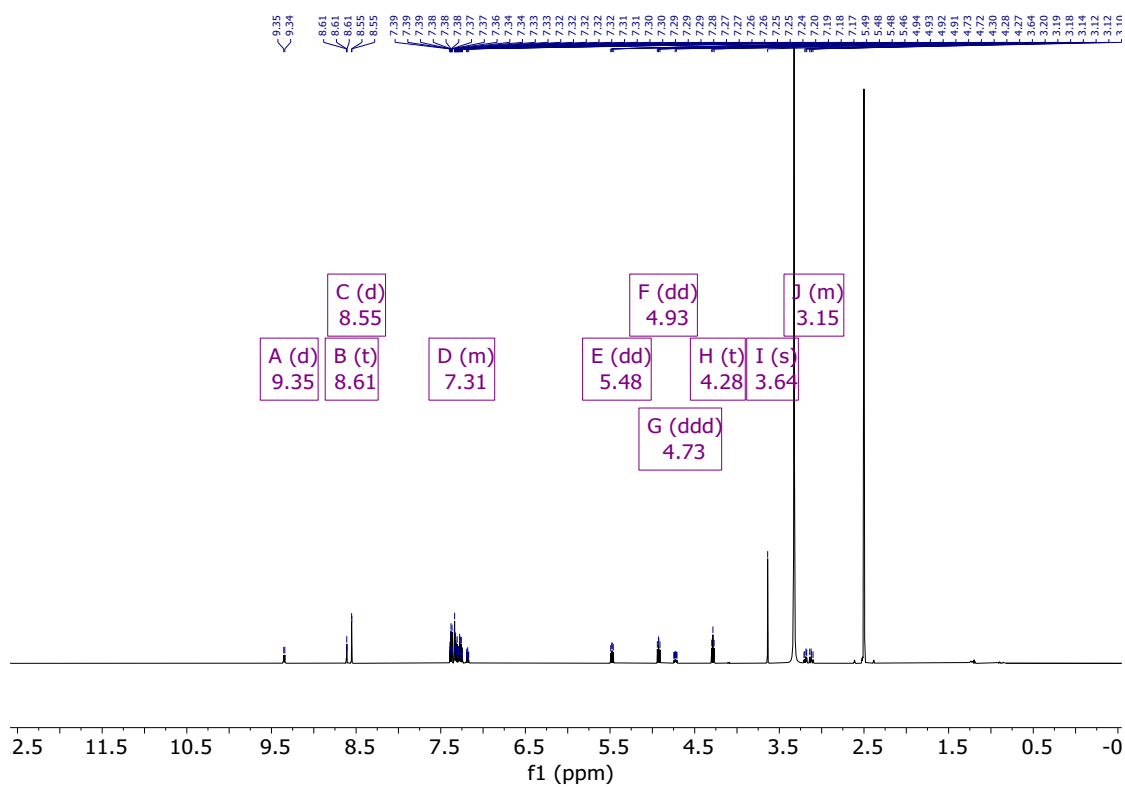
^{13}C NMR (151 MHz, CD_3OD) of $\mathbf{1}_{\mathbf{t5}}$.



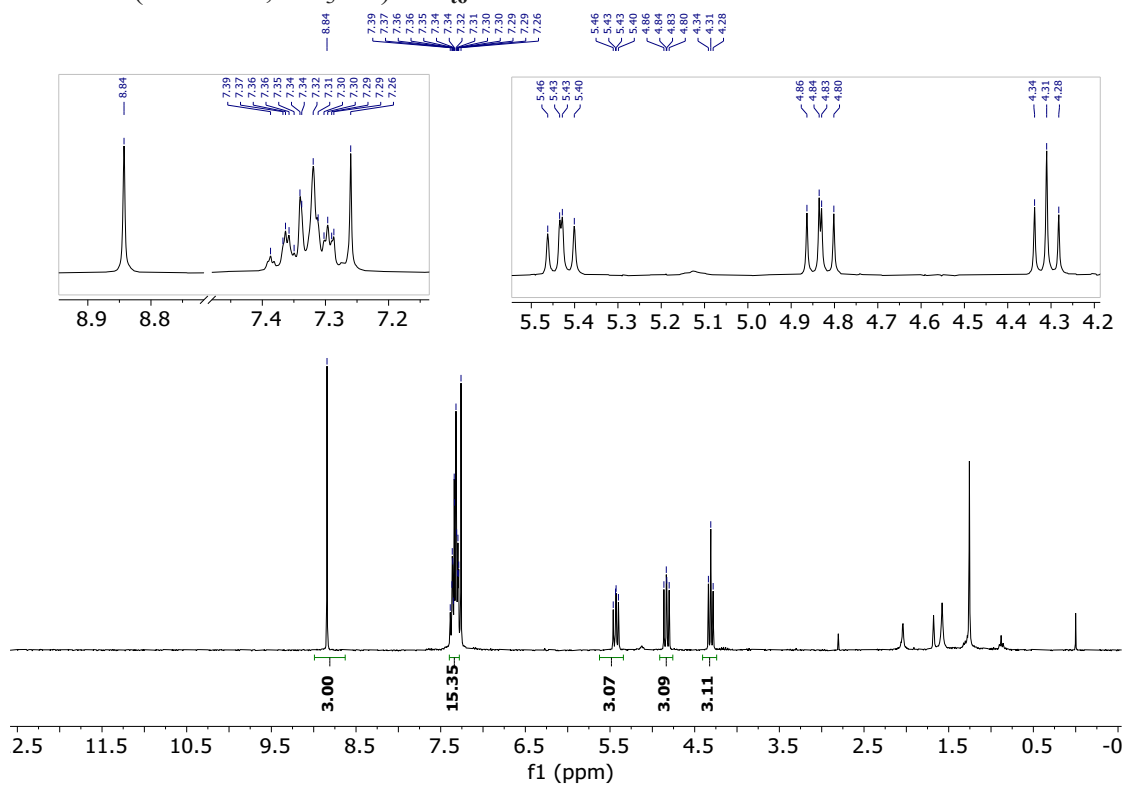
^1H NMR (600 MHz, CDCl_3) of $\mathbf{1}_{\text{t5}}$, $c = 60$ mM.



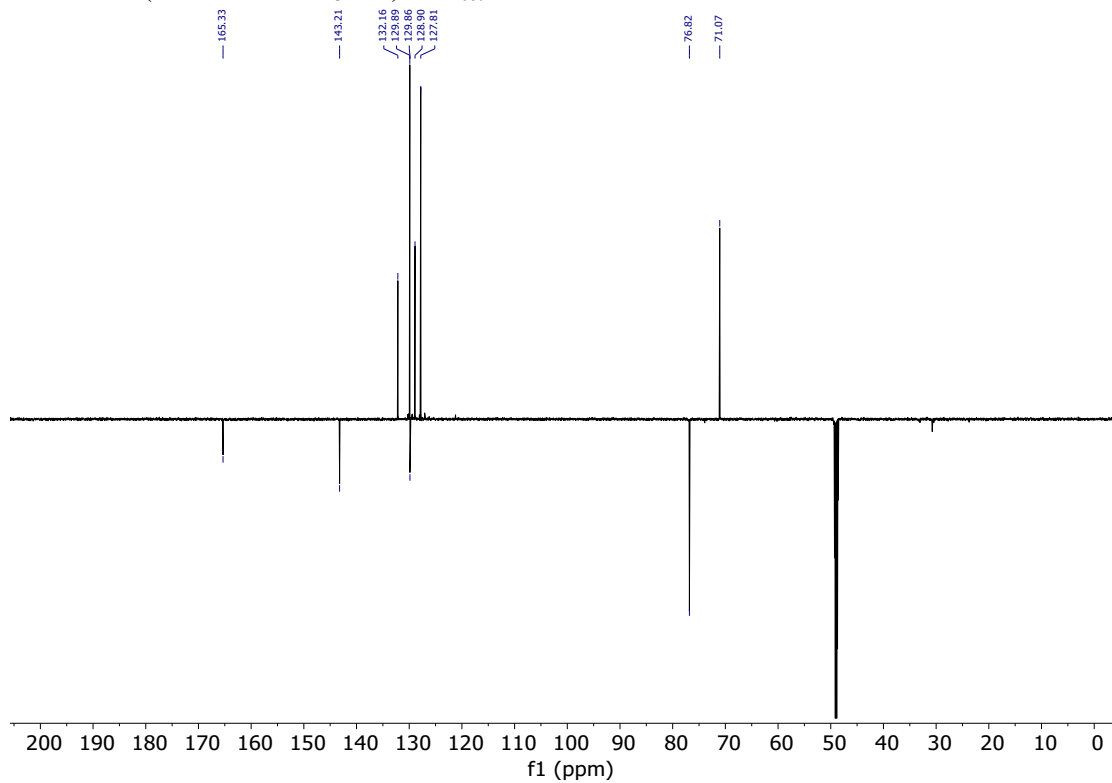
^1H NMR (600 MHz, DMSO) of $\mathbf{1}_{\text{t5}}$, $c = 6$ mM.



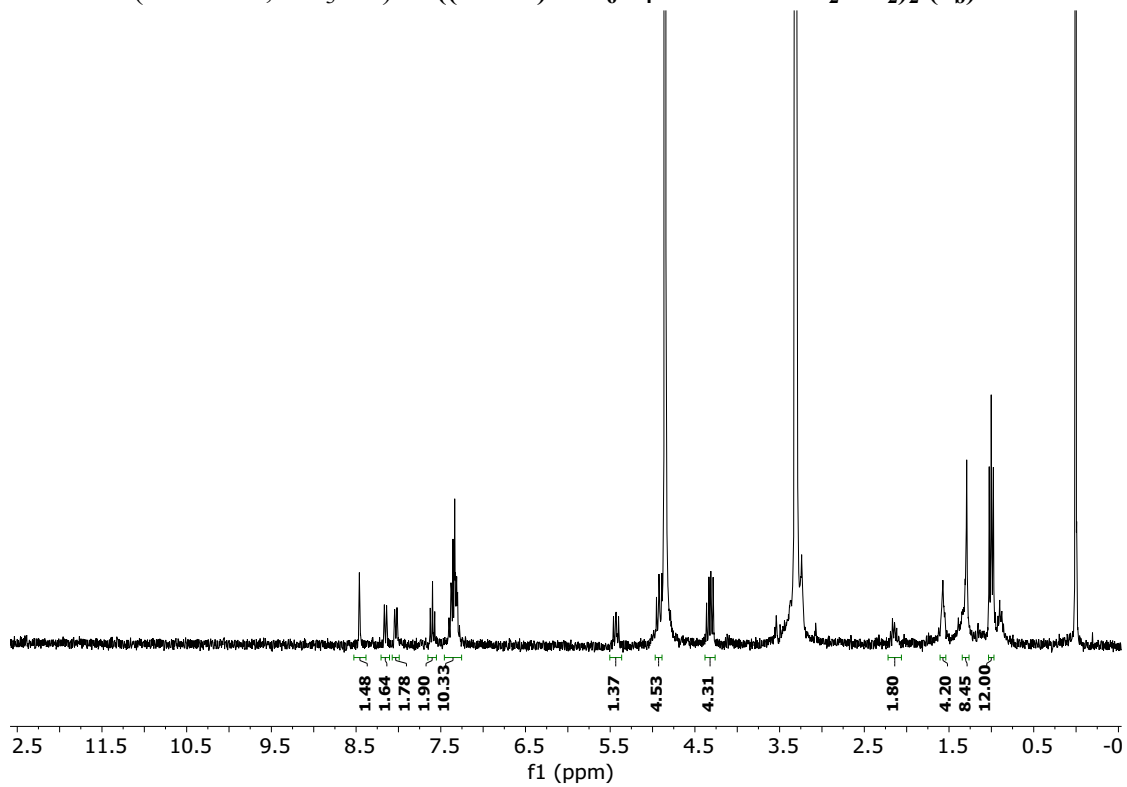
^1H NMR (300 MHz, CD_3CN) of **1_{t6}**.



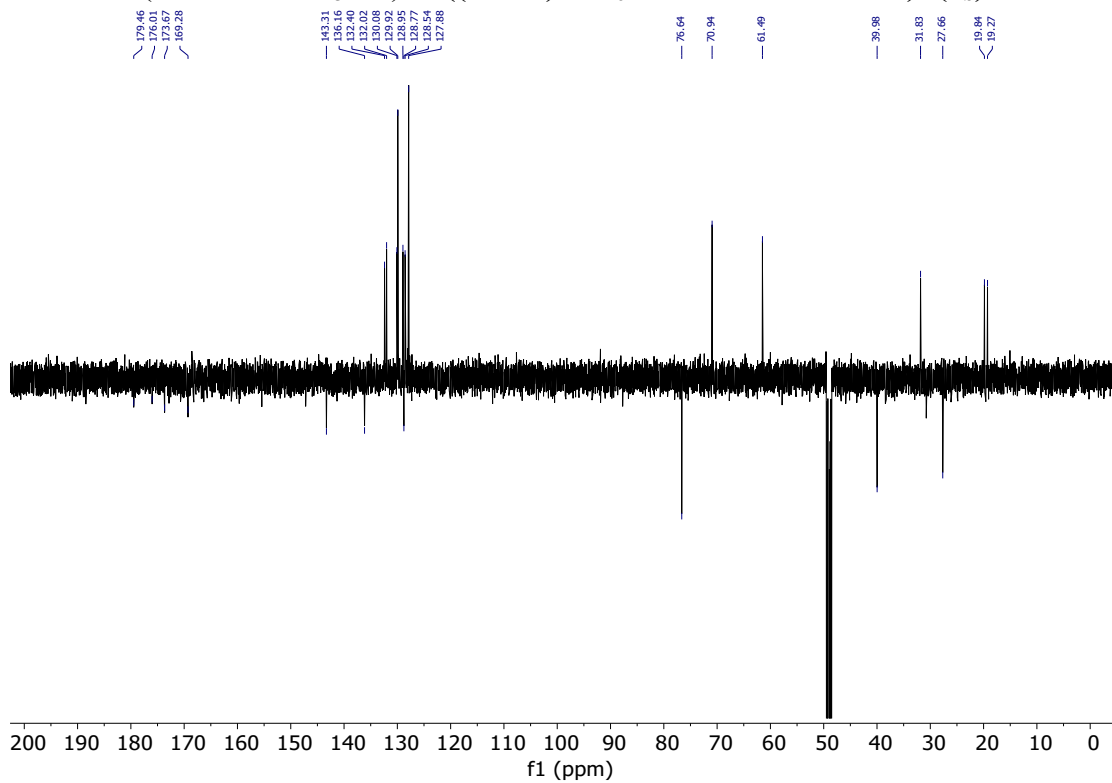
^{13}C NMR (151 MHz, CD_3OD) of **1_{t6}**.



^1H NMR (300 MHz, CD_3OD) of $((\text{Ph-ox})\text{-}m\text{C}_6\text{H}_4\text{-Val-NH-CH}_2\text{CH}_2)_2$ (**1b**).

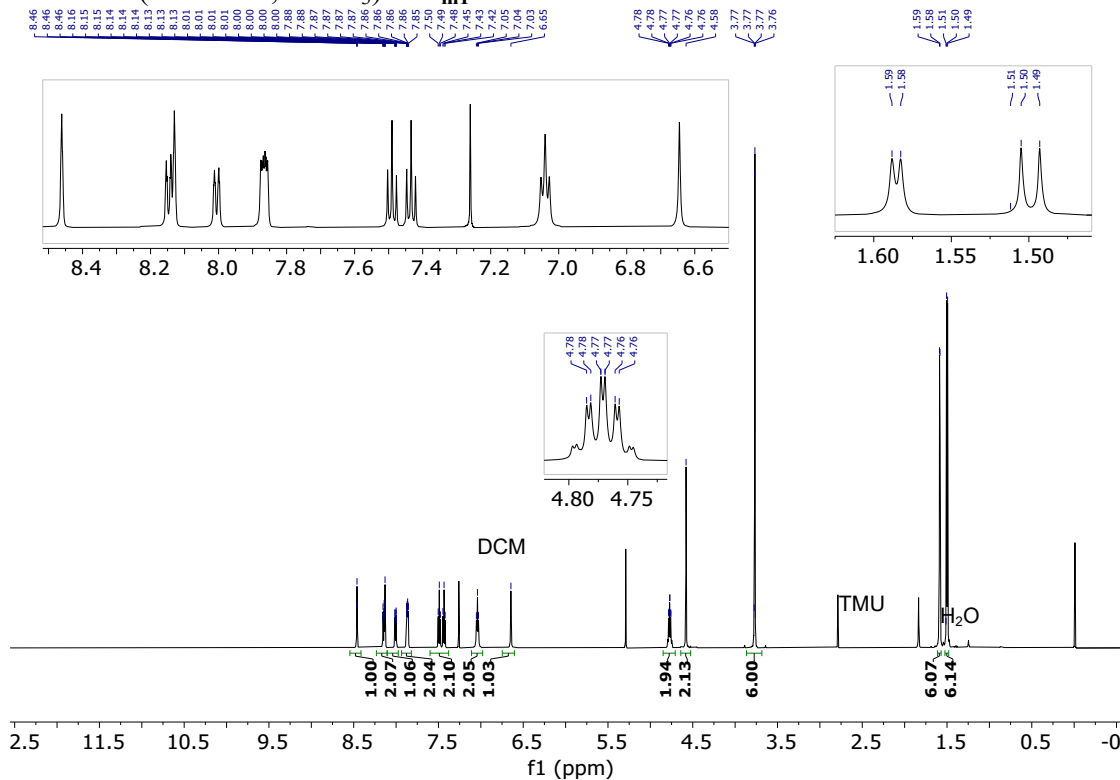


^{13}C NMR (151 MHz, CD_3OD) of $((\text{Ph-ox})\text{-}m\text{C}_6\text{H}_4\text{-Val-NH-CH}_2\text{CH}_2)_2$ (**1b**).

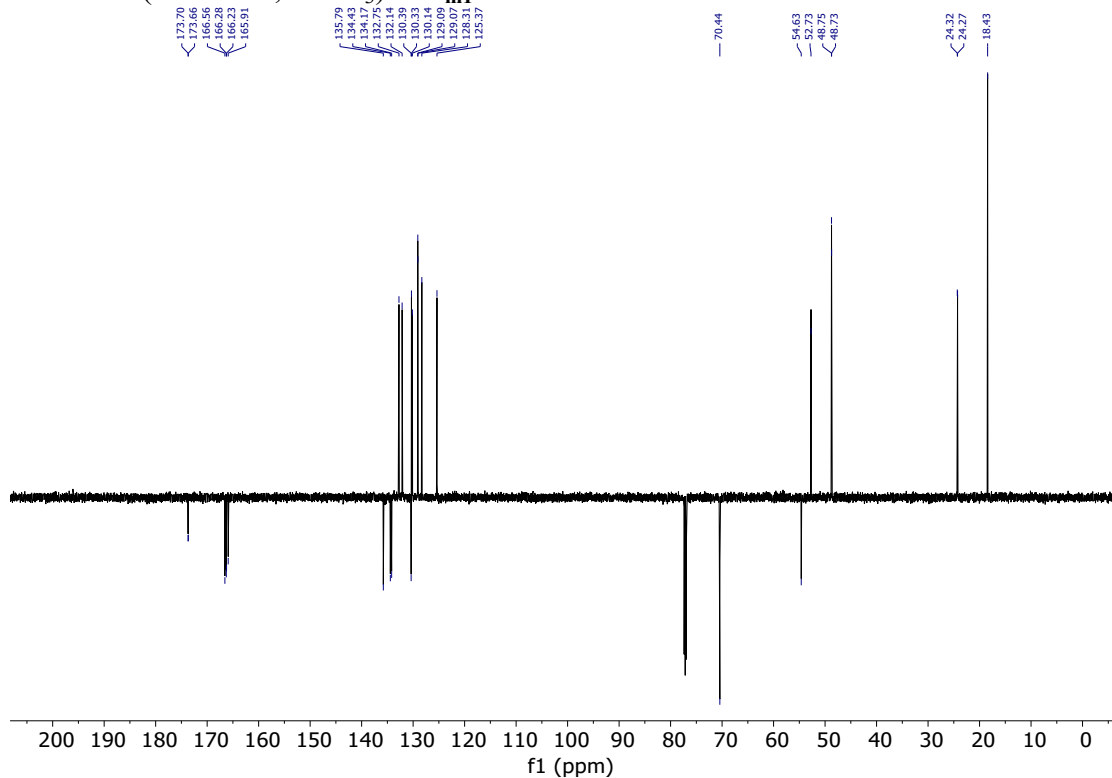


2.2. Reaction 1

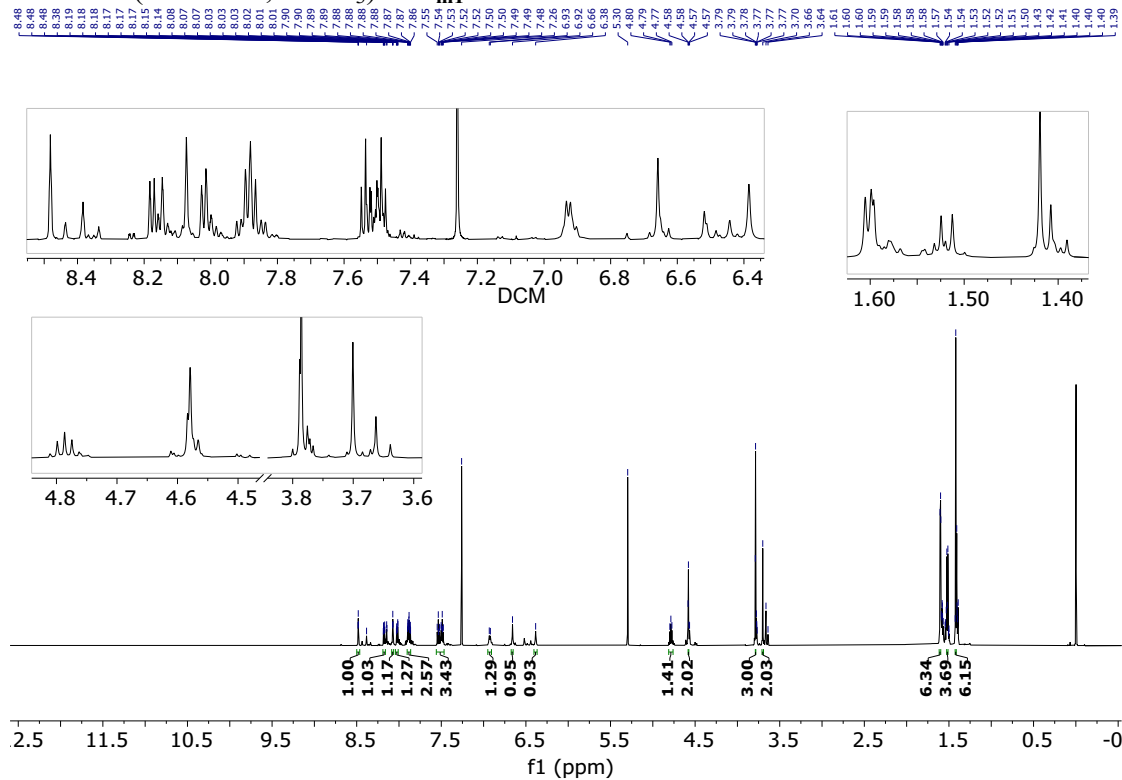
^1H NMR (600 MHz, CDCl_3) of $\mathbf{6m1}$.



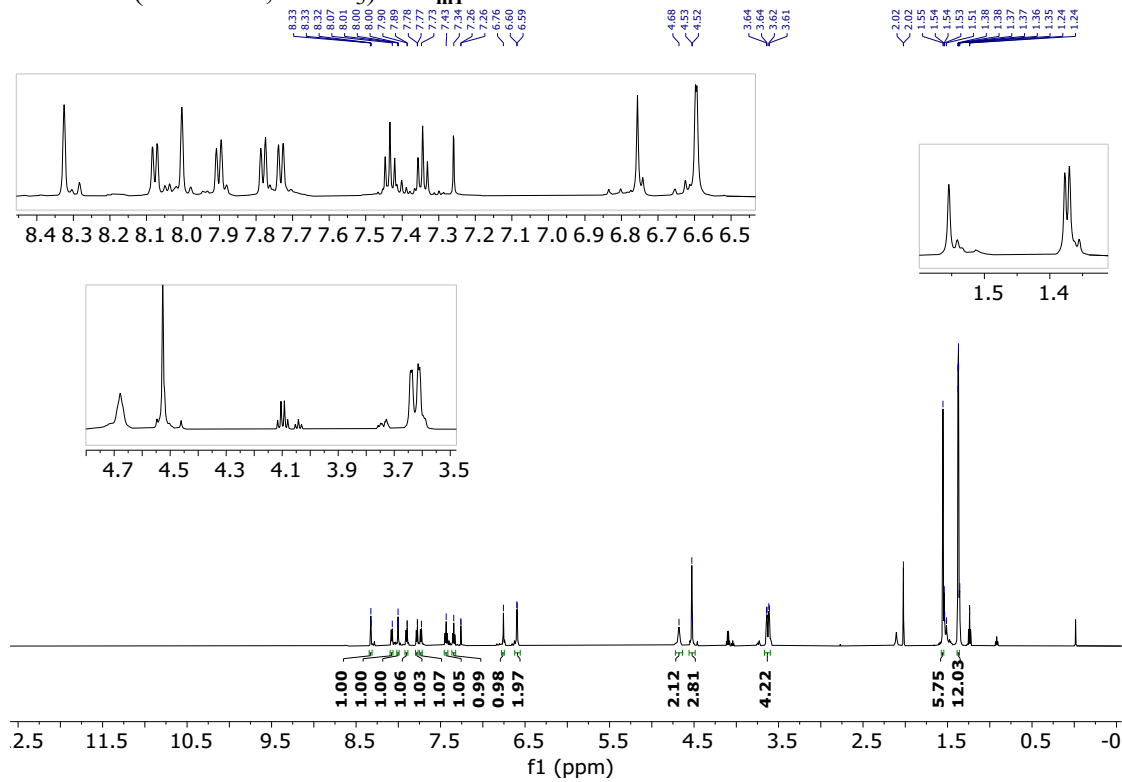
^{13}C NMR (151 MHz, CDCl_3) of $\mathbf{6m1}$.



^1H NMR (600 MHz, CDCl_3) of **7m1**.

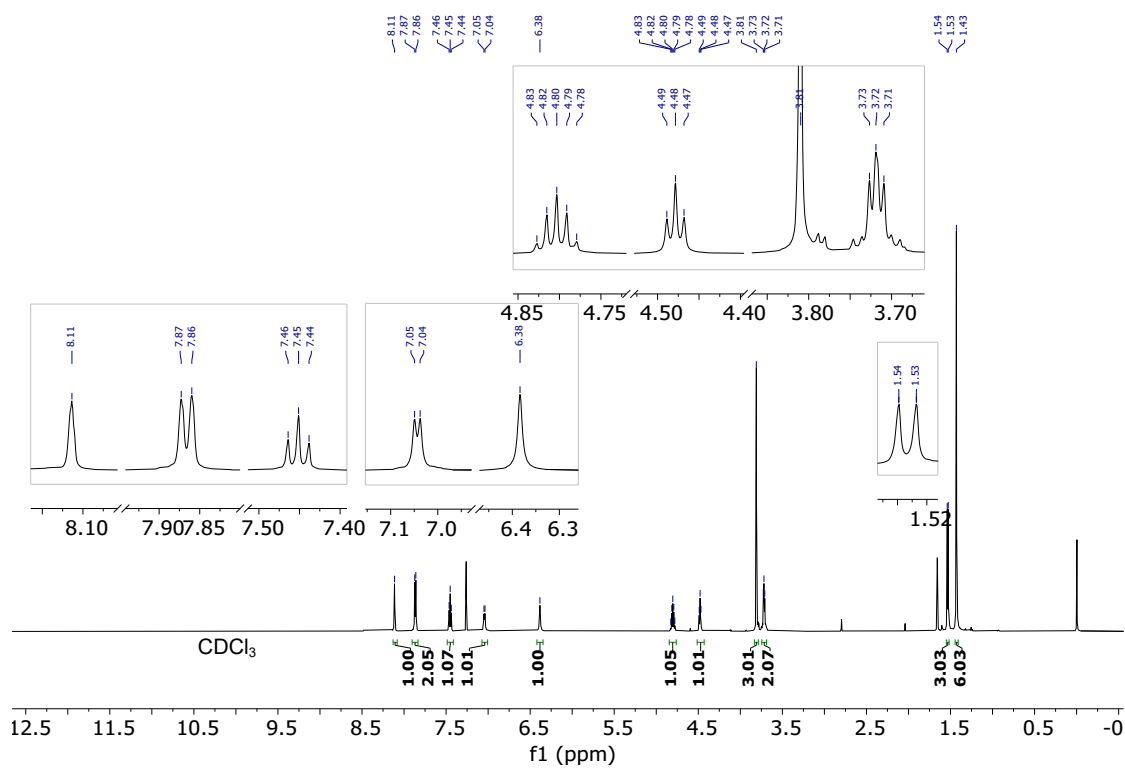


^1H NMR (600 MHz, CDCl_3) of **8m1**.

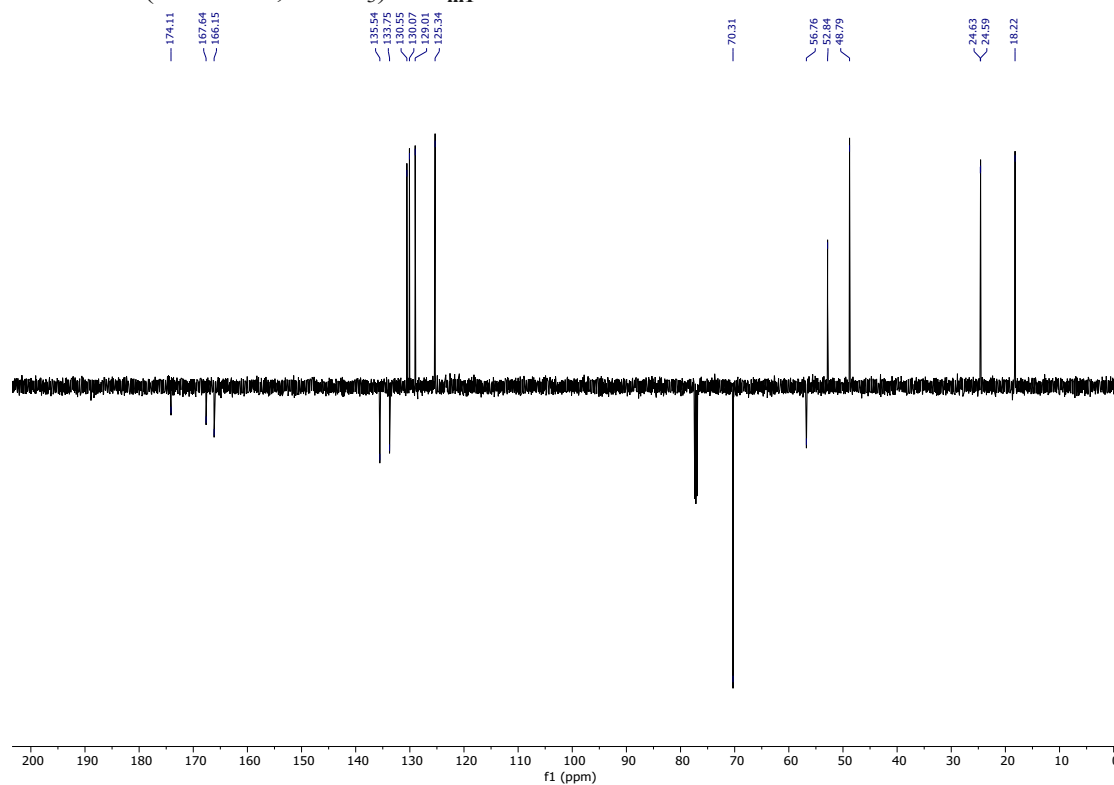


2.3. Reaction 5

^1H NMR (600 MHz, CDCl_3) of **3_{m1}**.

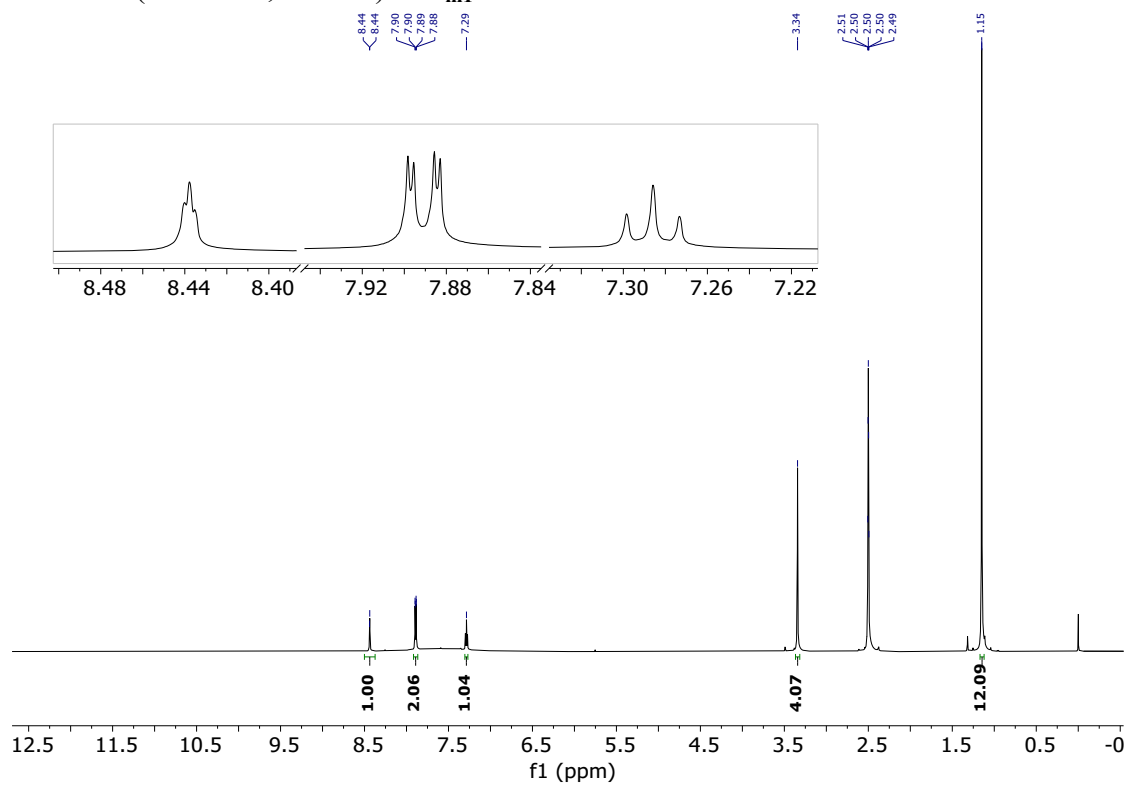


^{13}C NMR (151 MHz, CDCl_3) of **3_{m1}**.

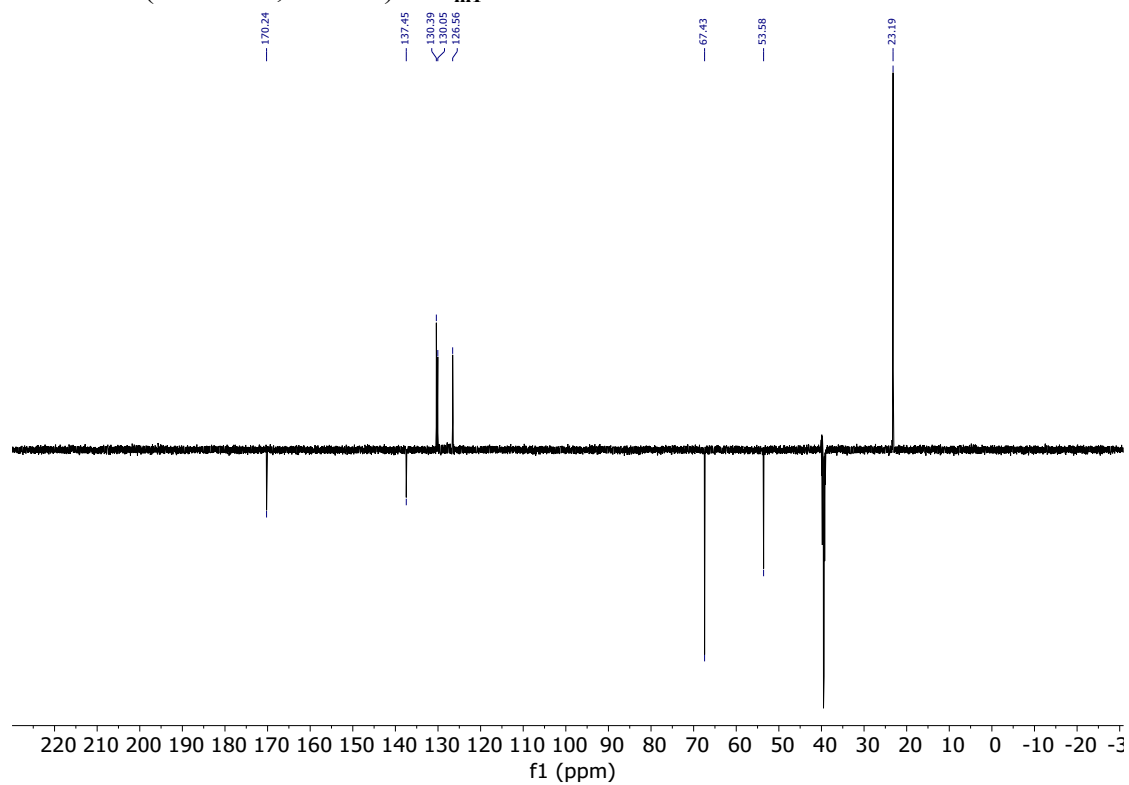


2.4. Reaction 6

^1H NMR (600 MHz, DMSO) of **4_{m1}**.

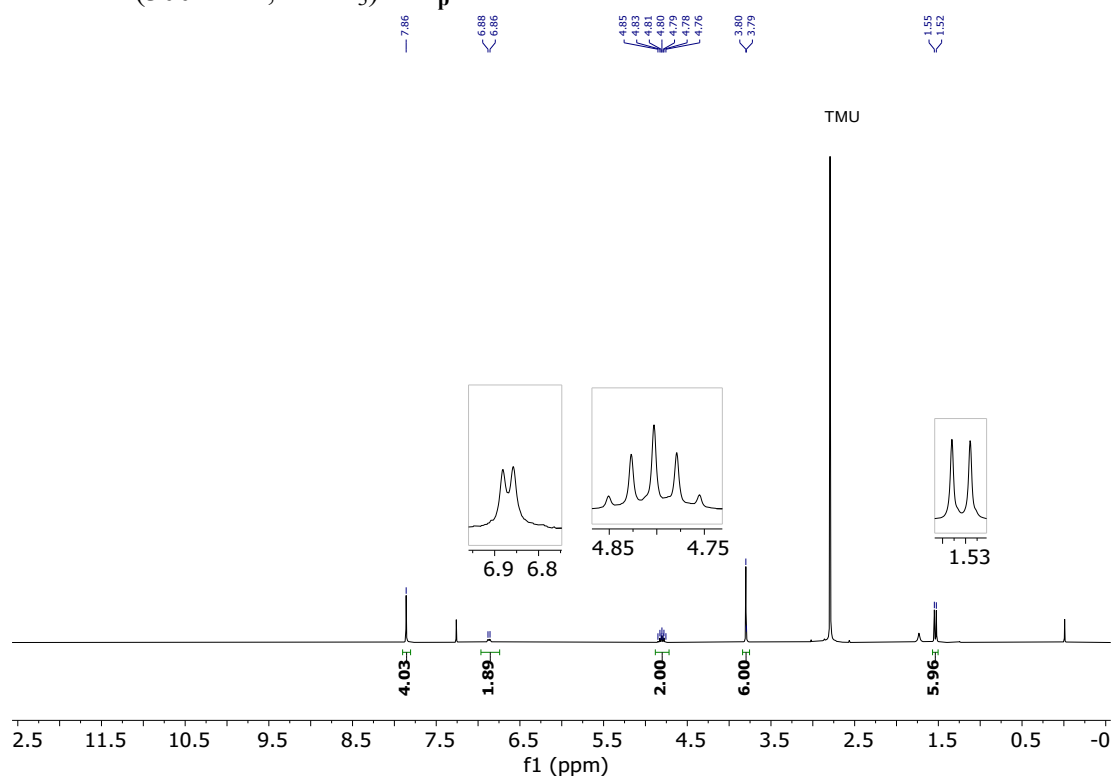


^{13}C NMR (151 MHz, DMSO) of **4_{m1}**.

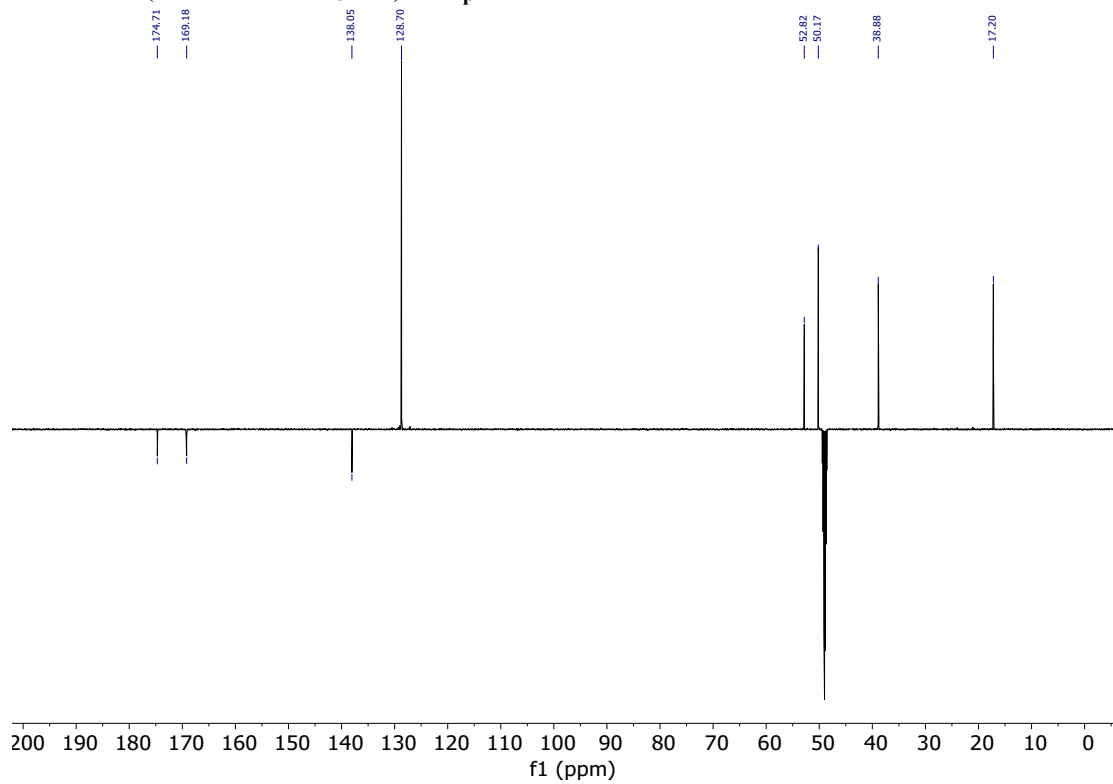


2.5. Reaction 7

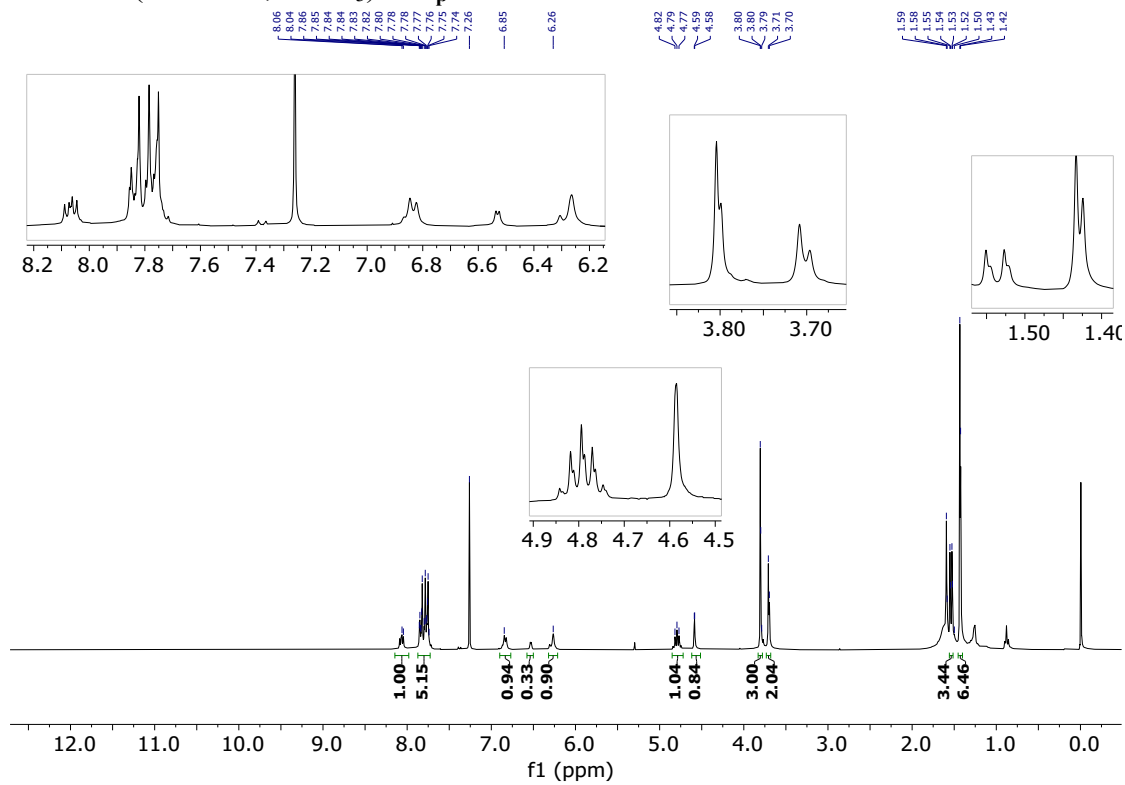
^1H NMR (300 MHz, CDCl_3) of **2p**.



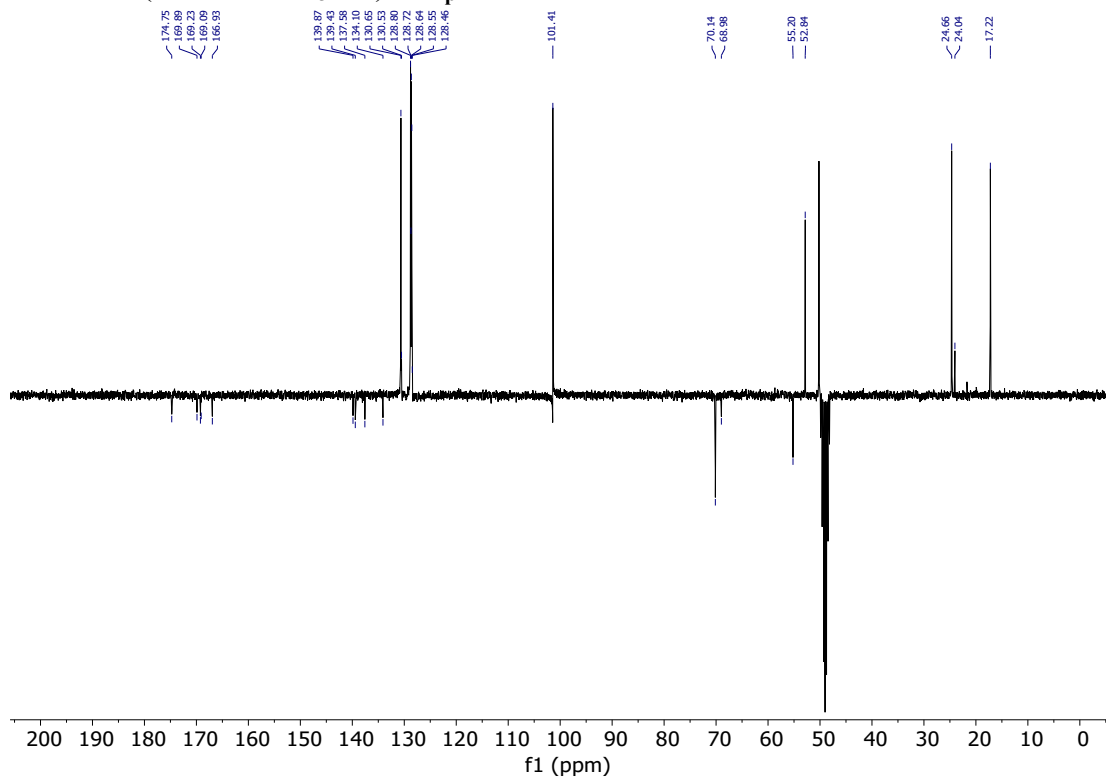
^{13}C NMR (151 MHz, CD_3OD) of **2p**.



^1H NMR (300 MHz, CDCl_3) of **3_p**.

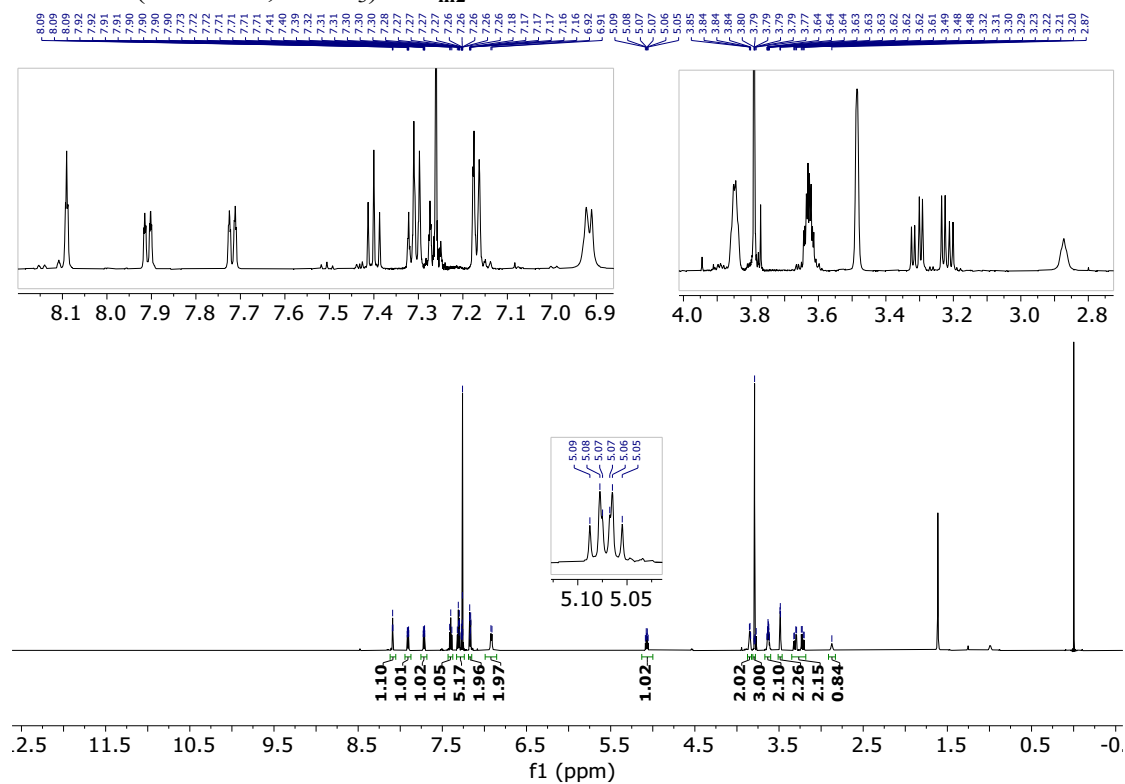


^{13}C NMR (75 MHz, CD_3OD) of **3_p**.

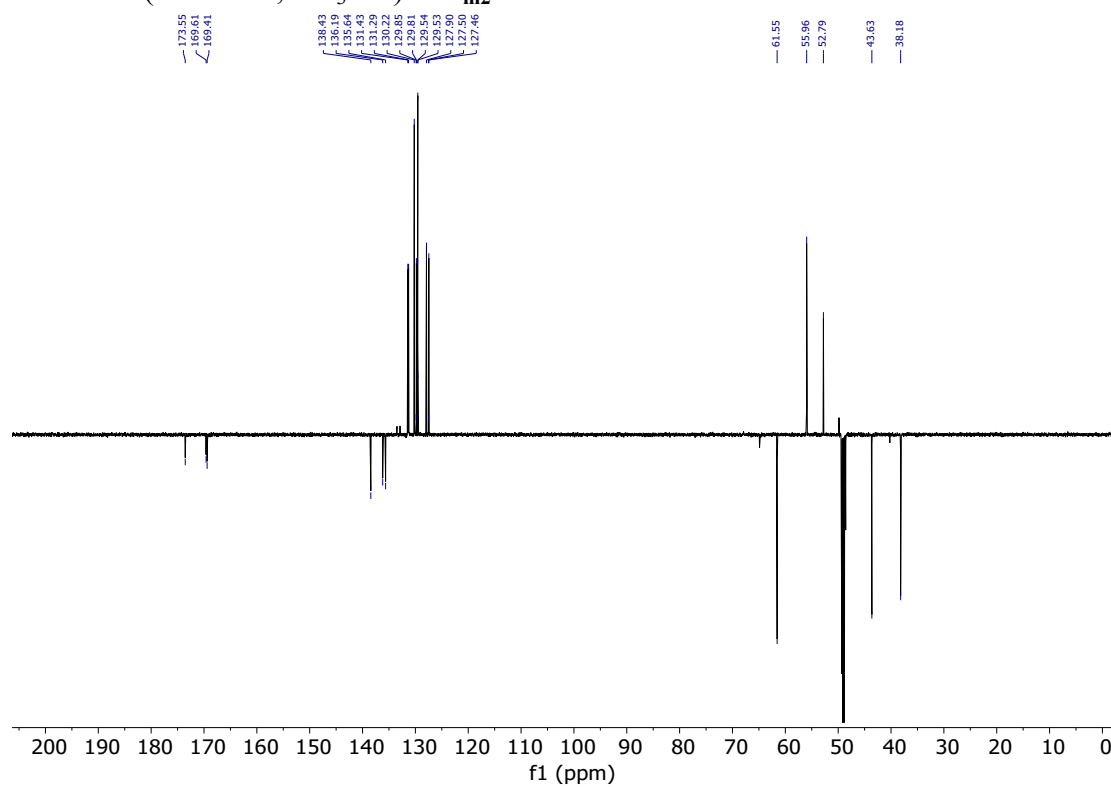


2.6. Reaction 8

^1H NMR (600 MHz, CDCl_3) of $3_{\text{m}2}$.

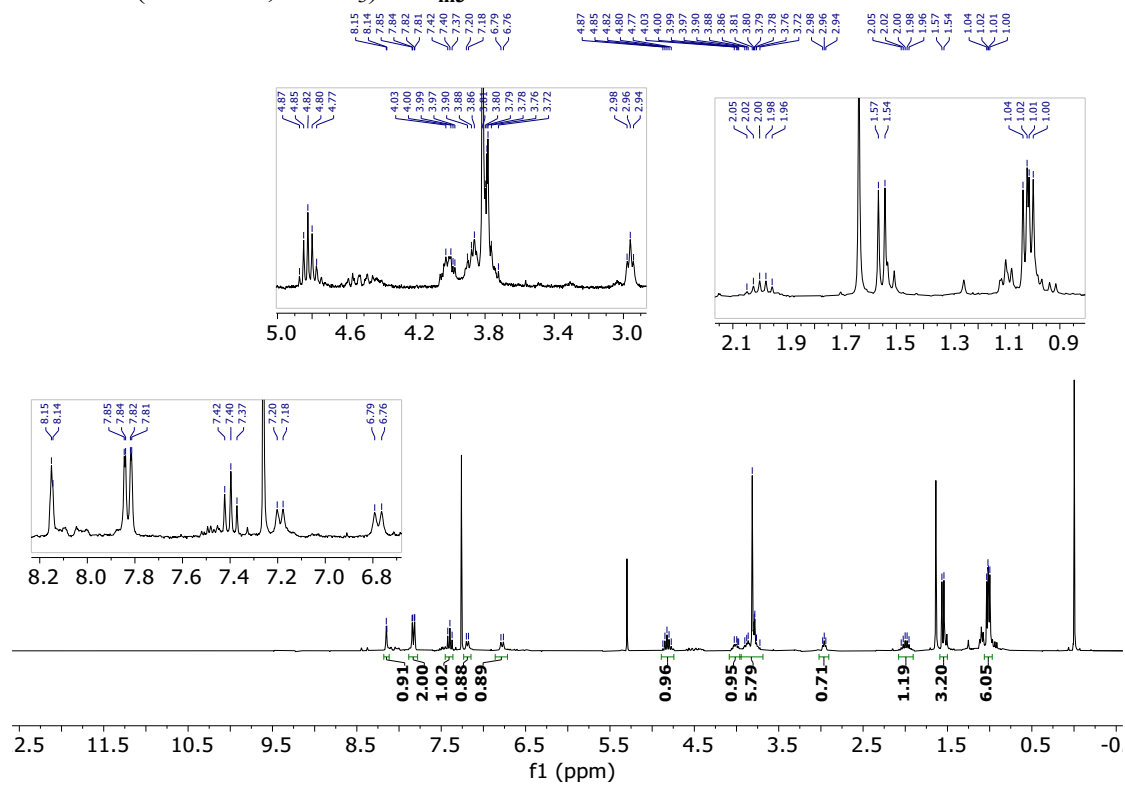


^{13}C NMR (151 MHz, CD_3OD) of $3_{\text{m}2}$.



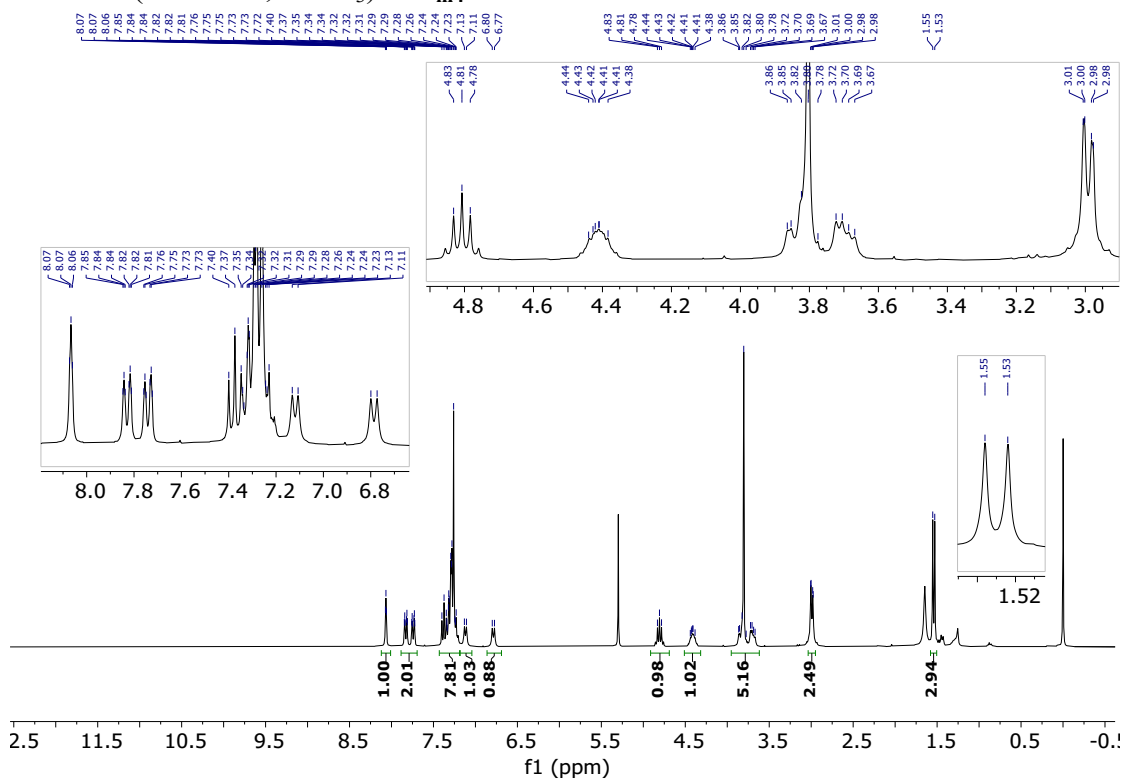
2.7. Reaction 9

^1H NMR (600 MHz, CDCl_3) of **3m3**.



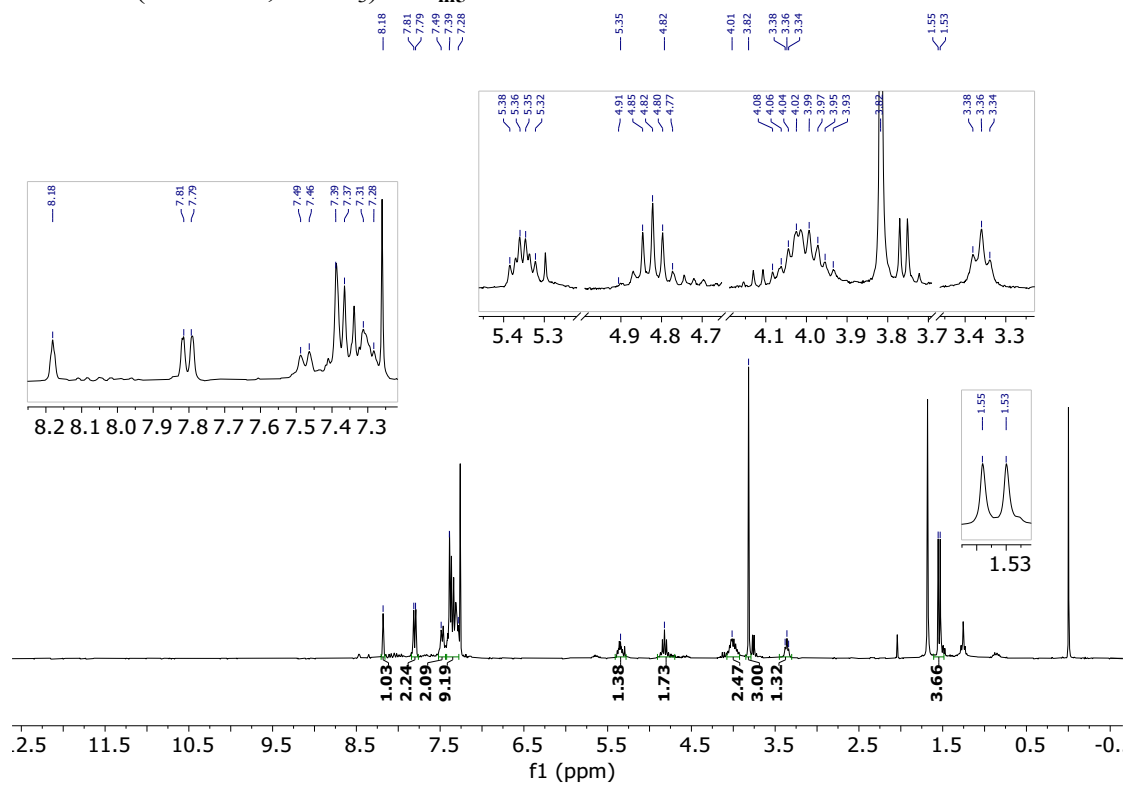
2.8. Reaction 10

^1H NMR (300 MHz, CDCl_3) of $\mathbf{3m4}$.

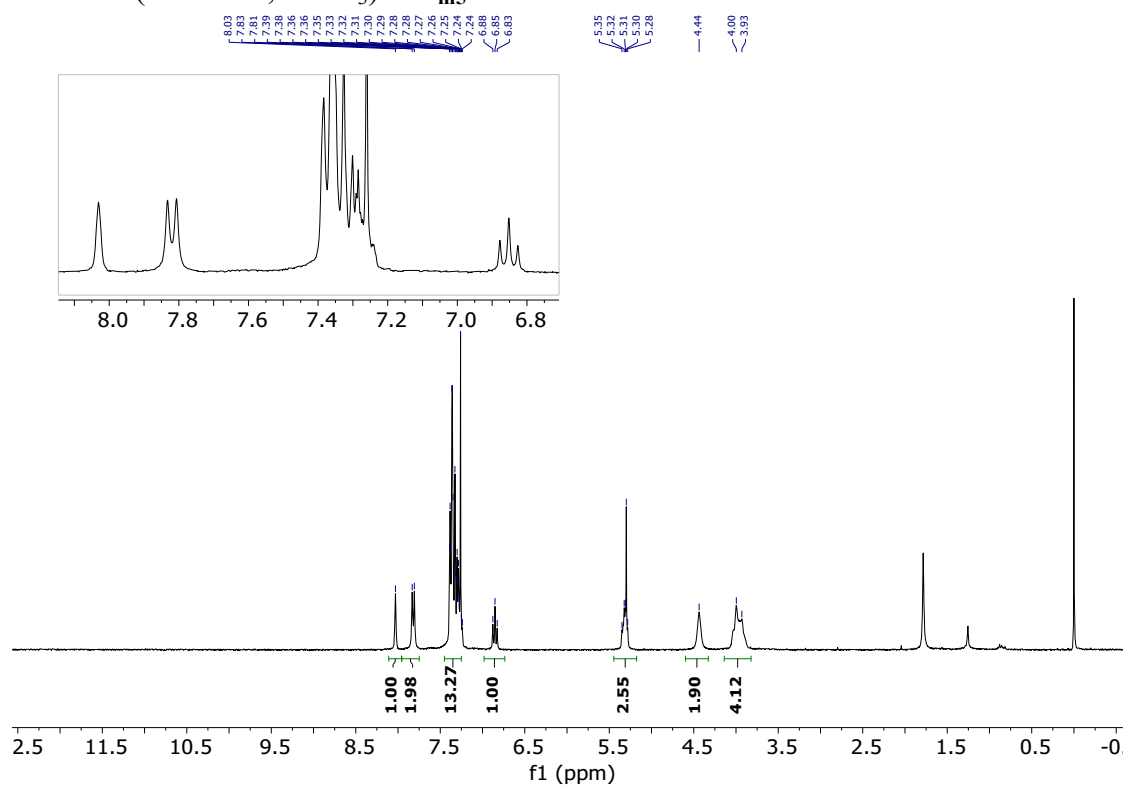


2.9. Reaction 11

^1H NMR (300 MHz, CDCl_3) of **3_{m5}**.

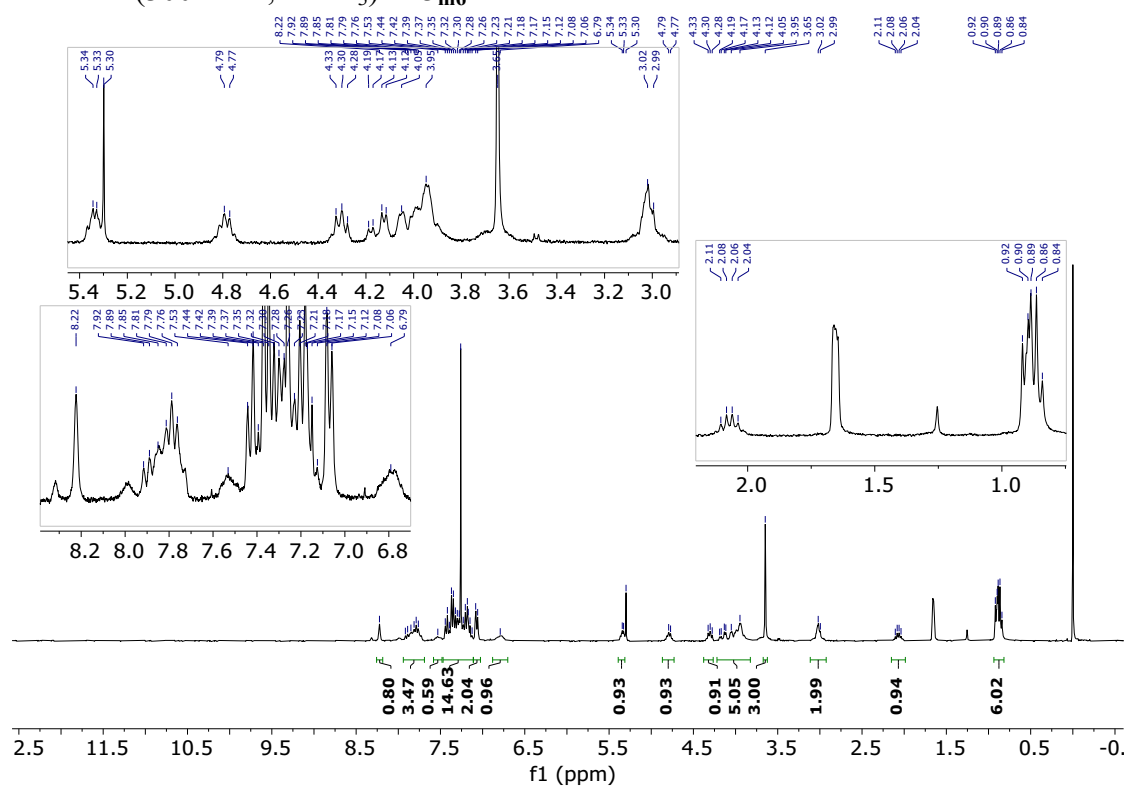


^1H NMR (300 MHz, CDCl_3) of **4_{m5}**.



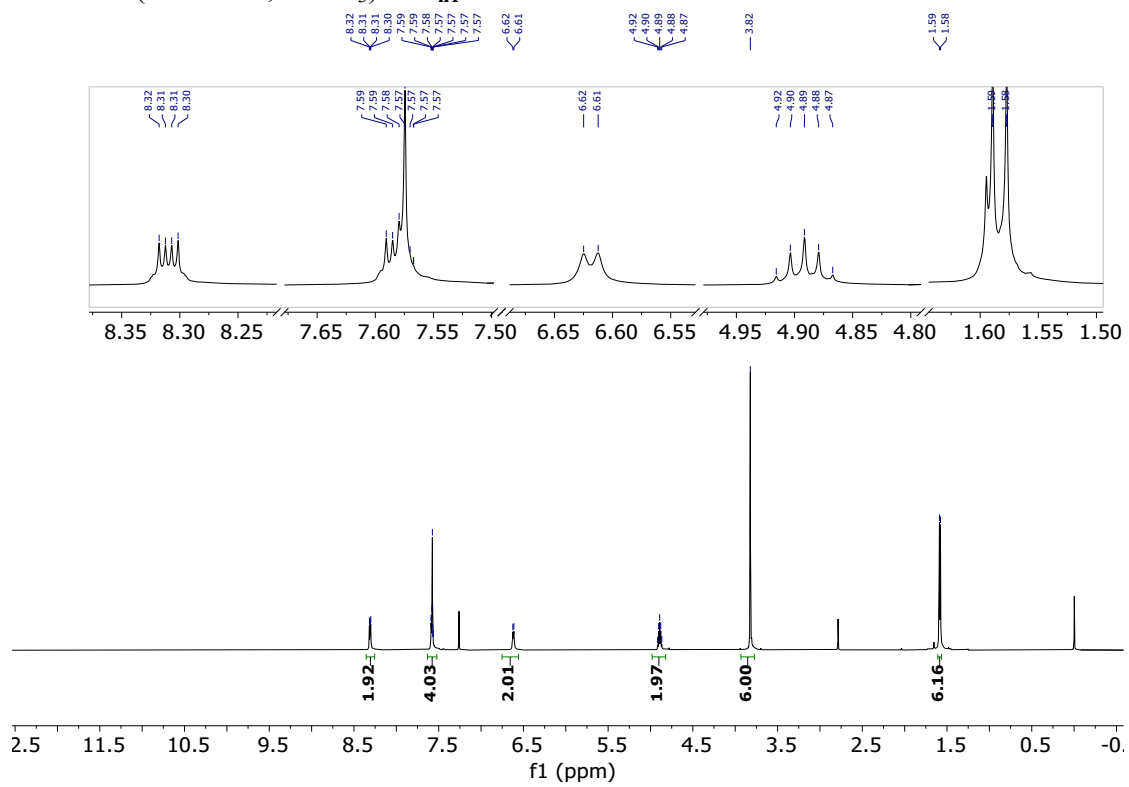
2.10. Reaction 12

^1H NMR (300 MHz, CDCl_3) of **3_{m6}**.

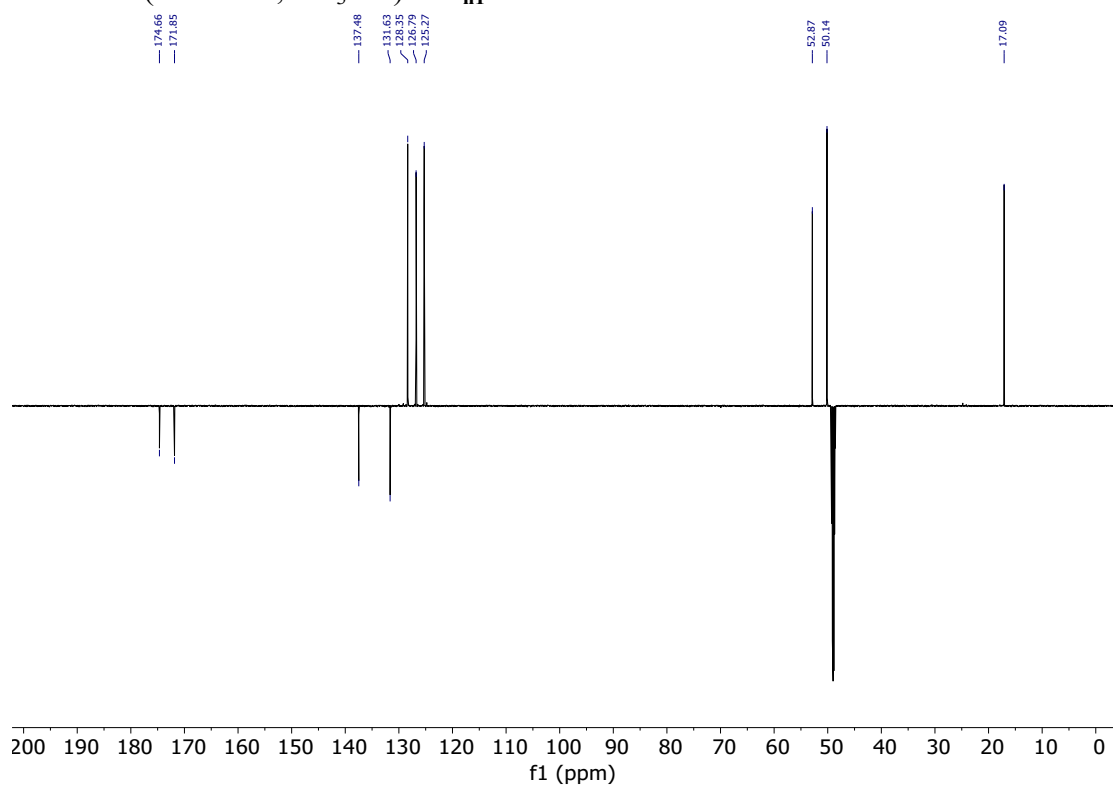


2.11. Reaction 13

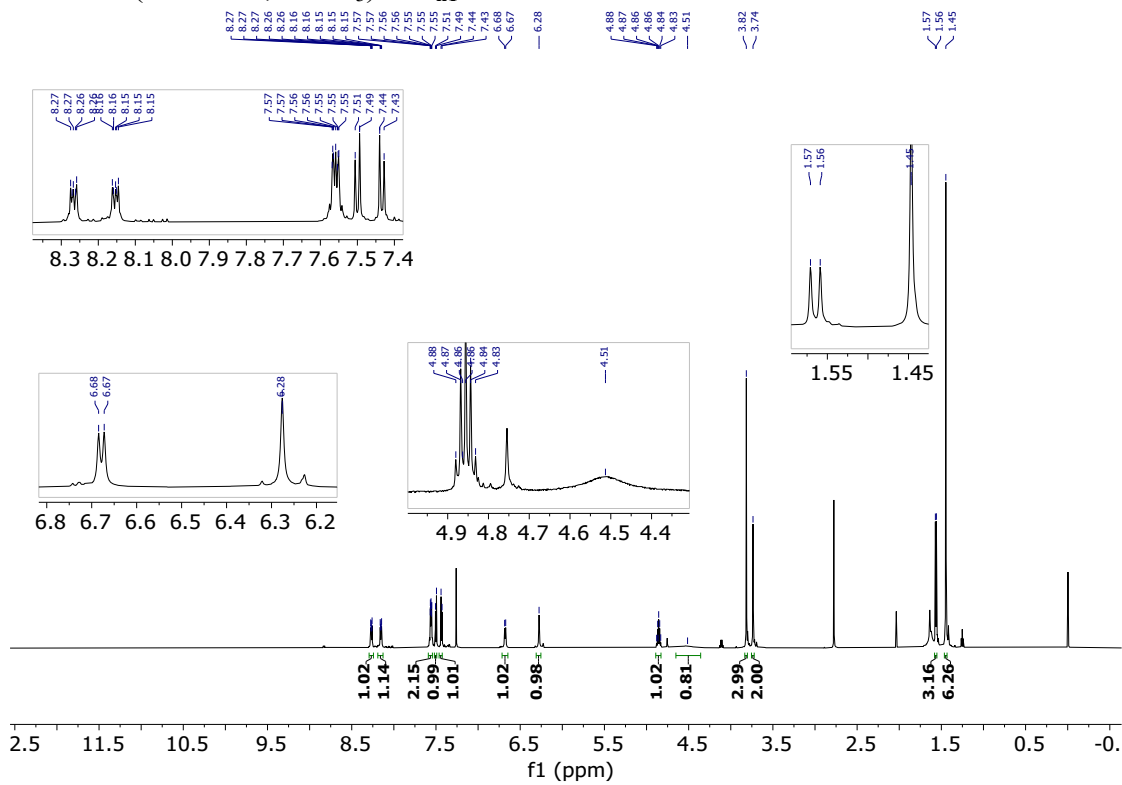
^1H NMR (600 MHz, CDCl_3) of **2_{n1}**.



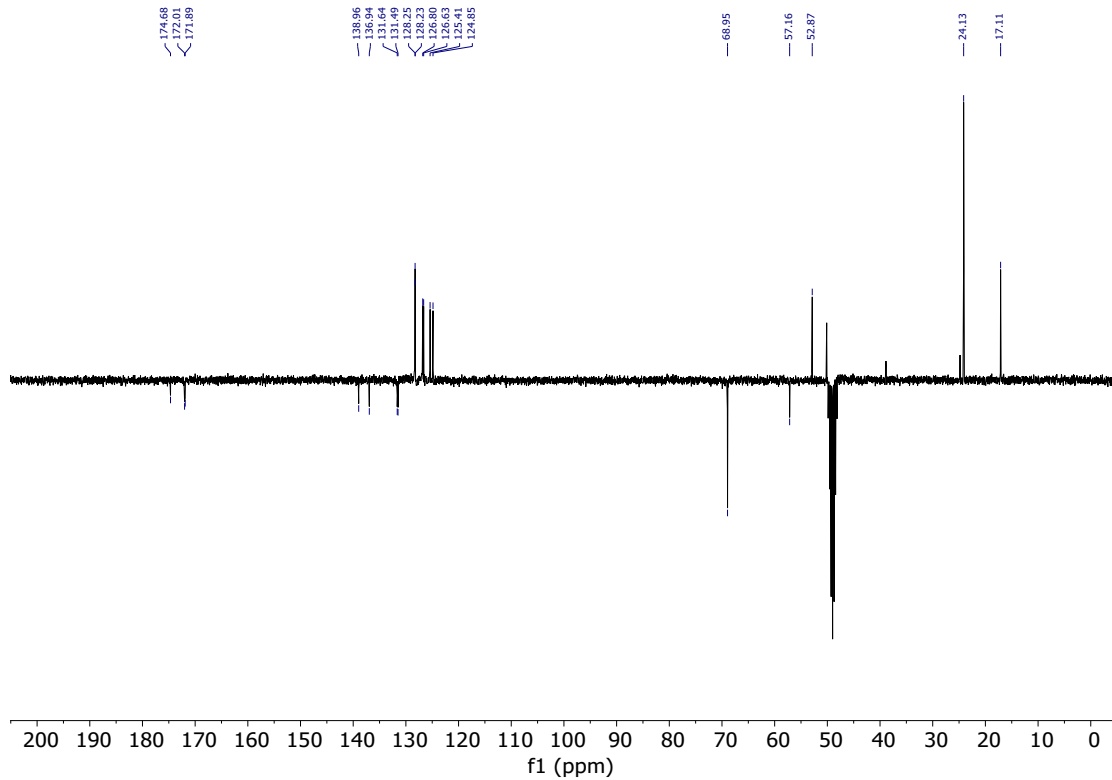
^{13}C NMR (151 MHz, CD_3OD) of **2_{n1}**.



^1H NMR (600 MHz, CDCl_3) of **3_{n1}**.

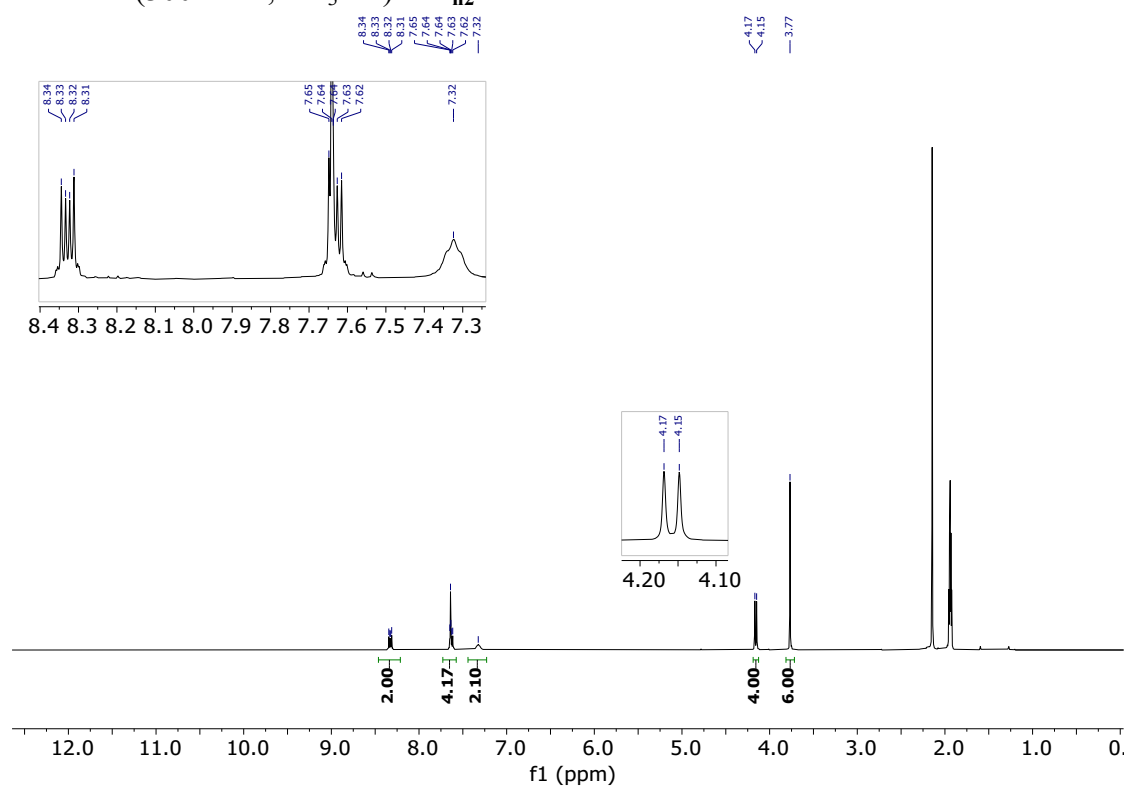


^{13}C NMR (75 MHz, CD_3OD) of **3_{n1}**.

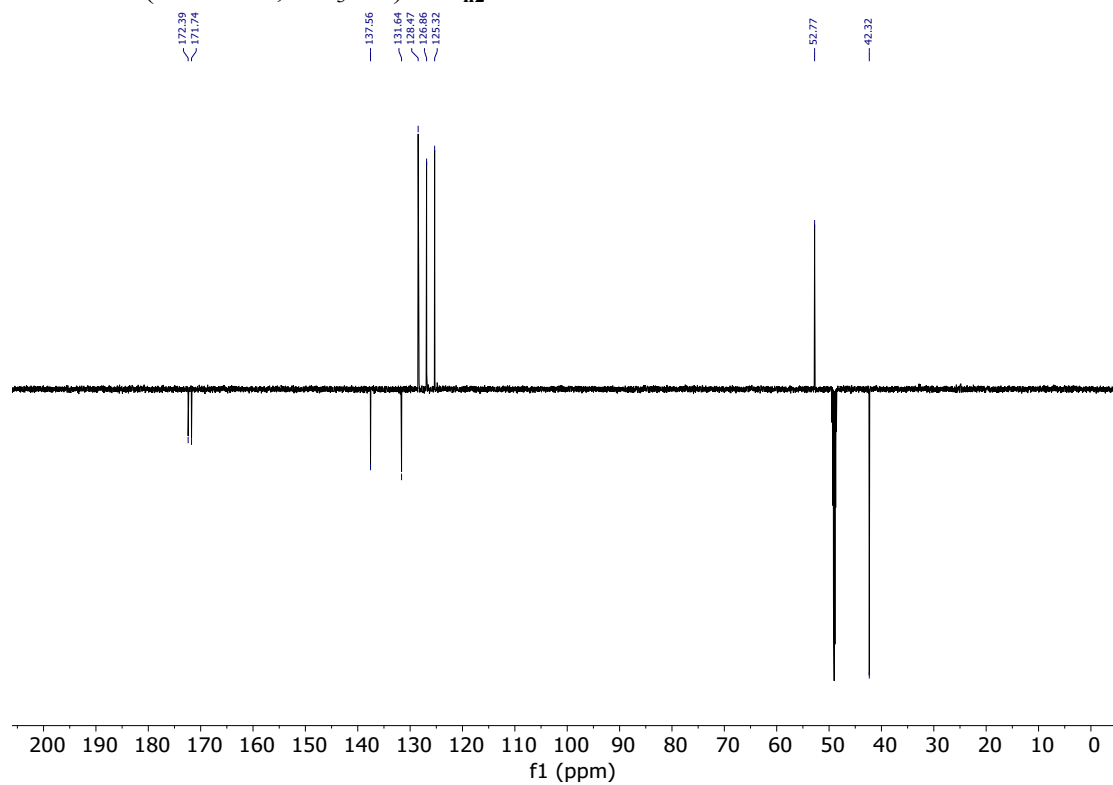


2.12. Reaction 14

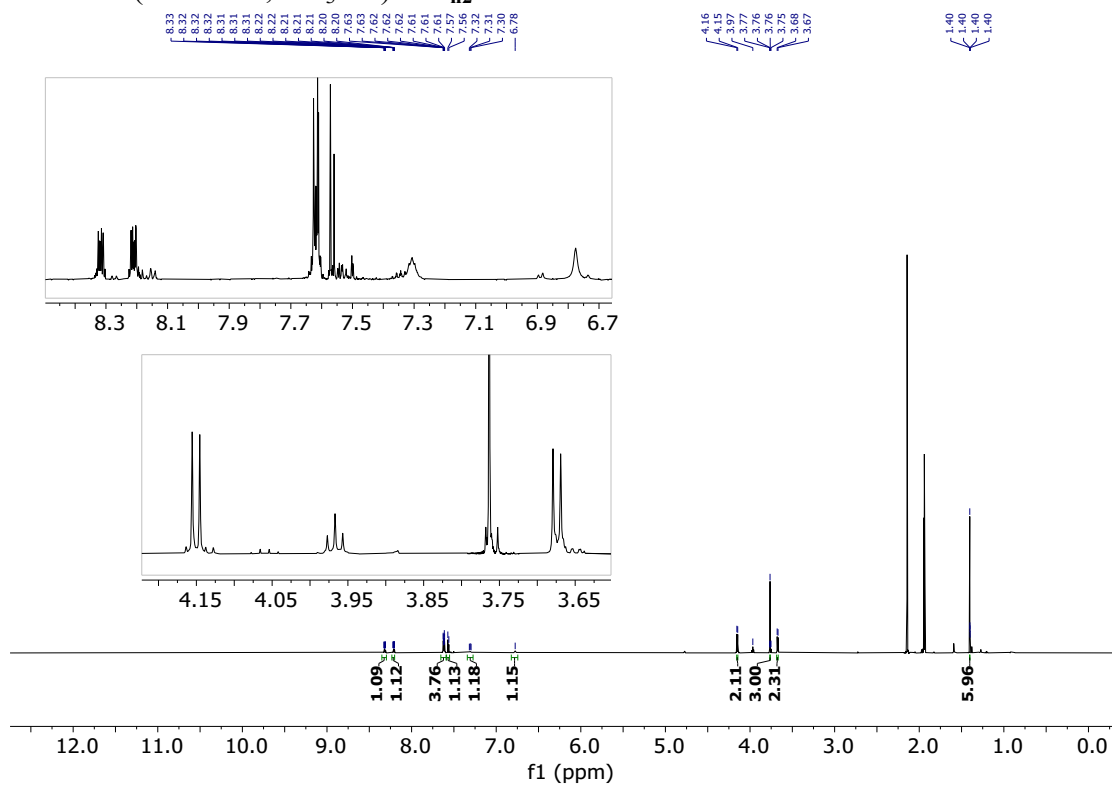
^1H NMR (300 MHz, CD_3CN) of 2_{n2} .



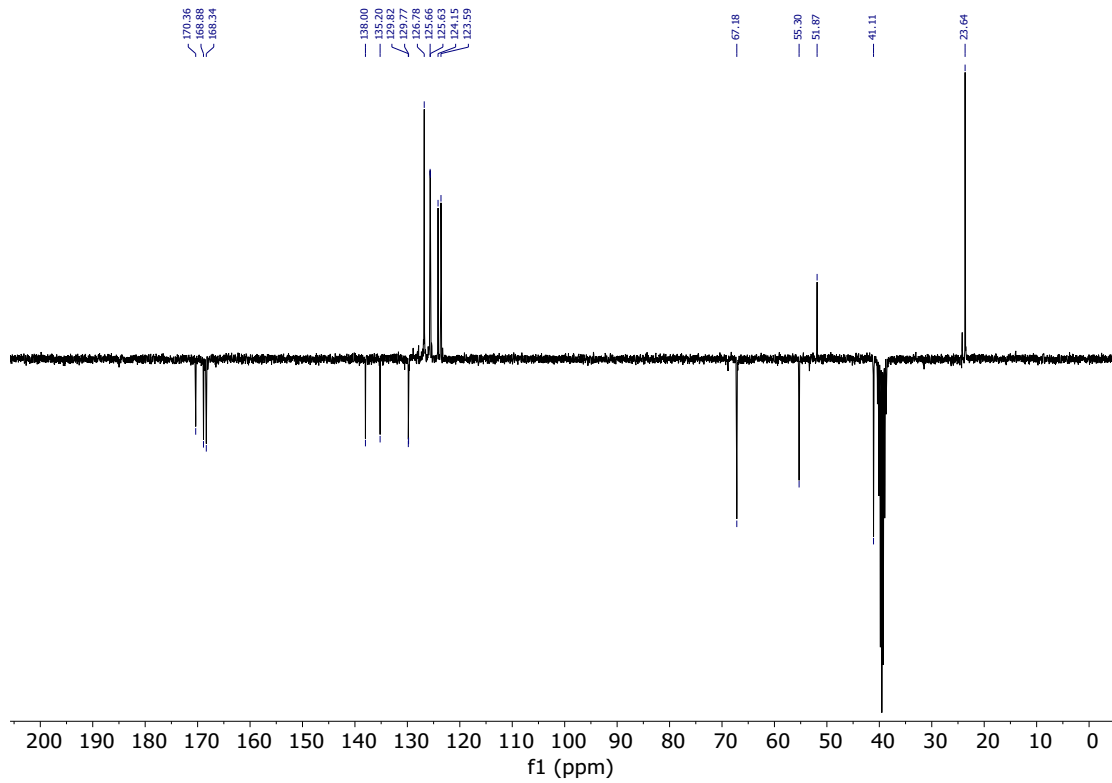
^{13}C NMR (151 MHz, CD_3OD) of 2_{n2} .



^1H NMR (600 MHz, CD_3CN) of 3_{n2} .

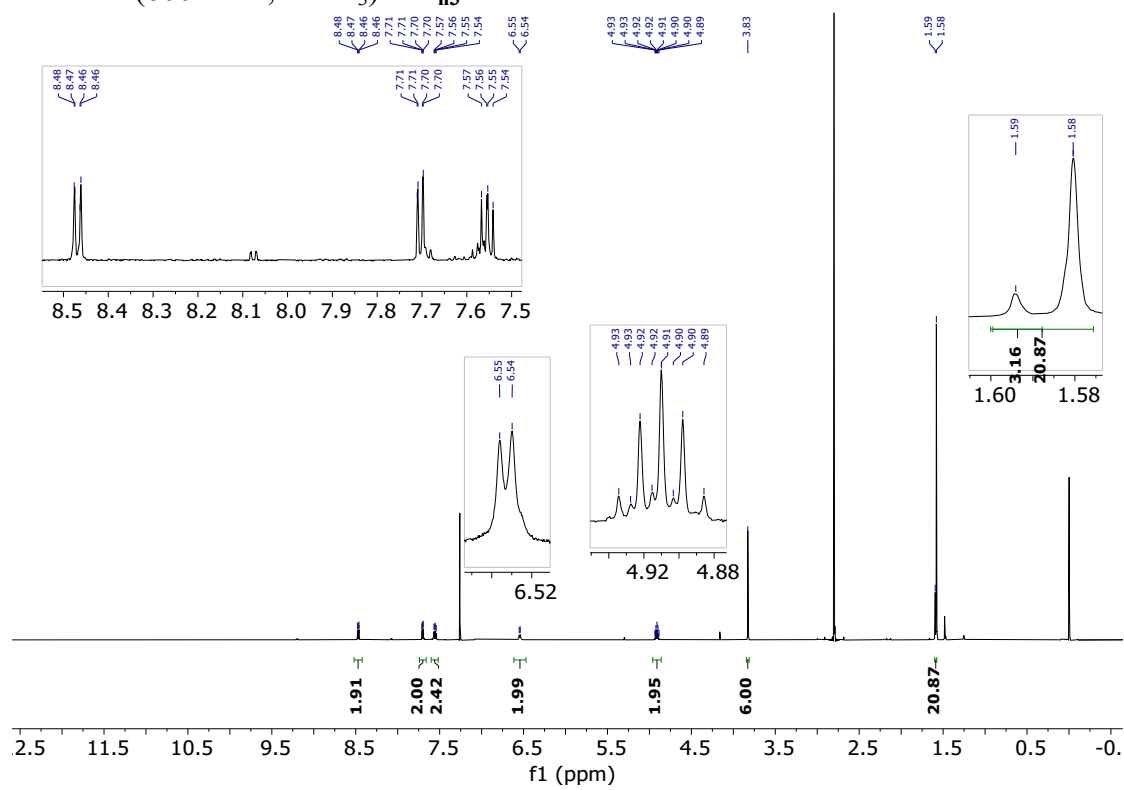


^{13}C NMR (75 MHz, DMSO) of 3_{n2} .

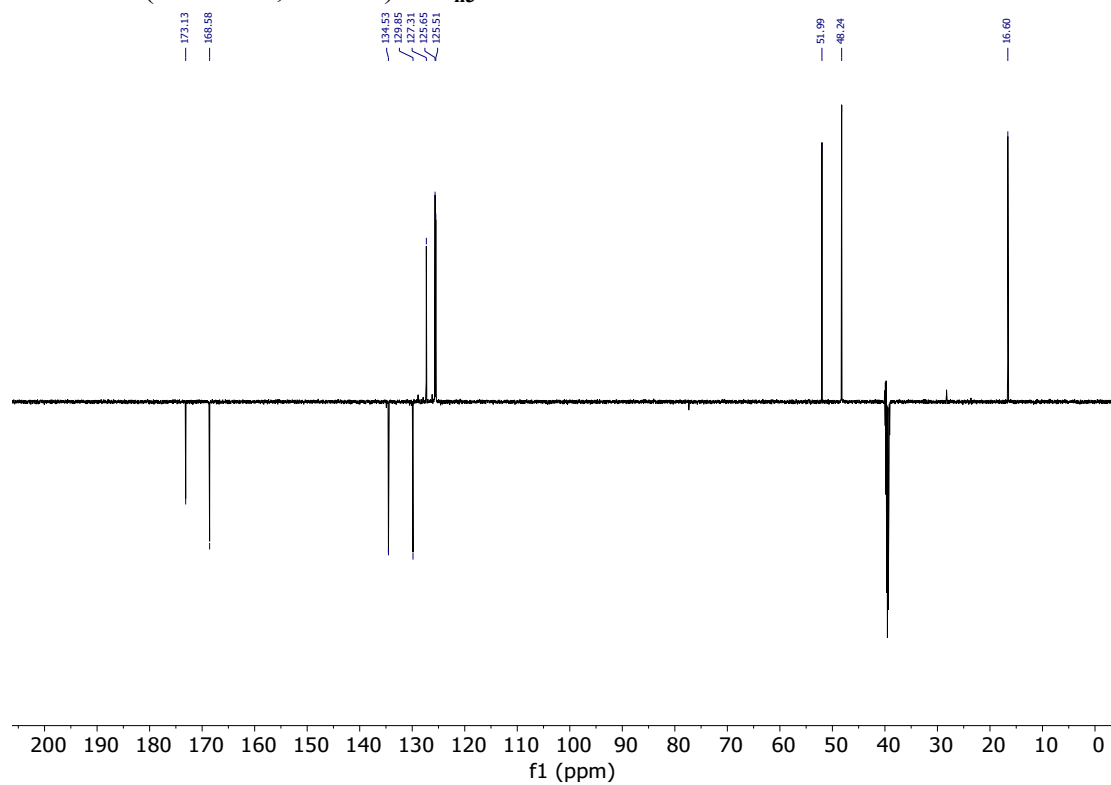


2.13. Reaction 15

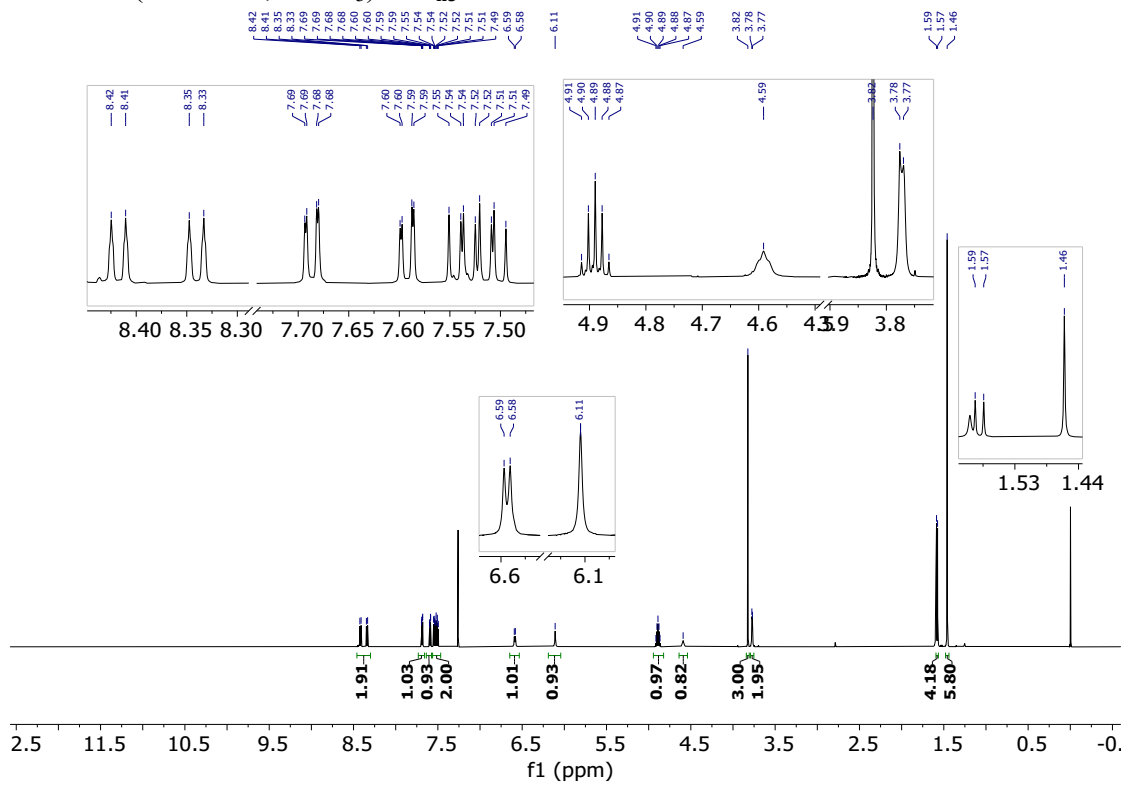
^1H NMR (600 MHz, CDCl_3) of **2_{n3}**.



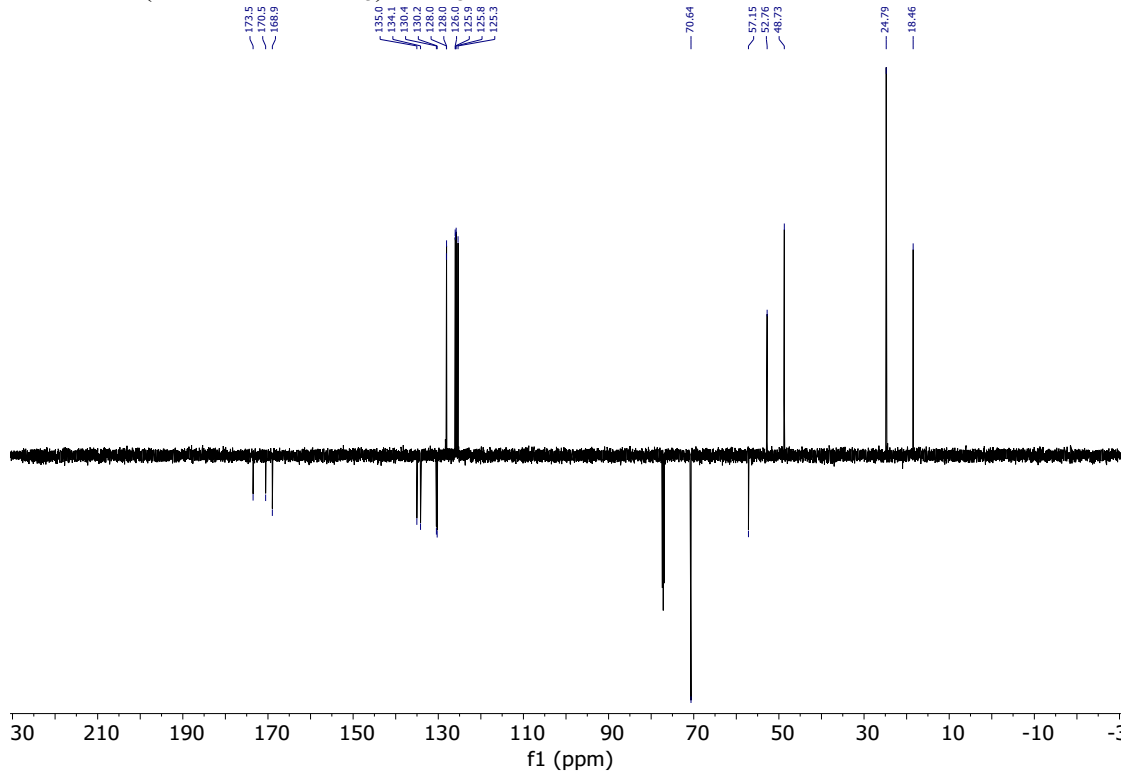
^{13}C NMR (151 MHz, DMSO) of **2_{n3}**.



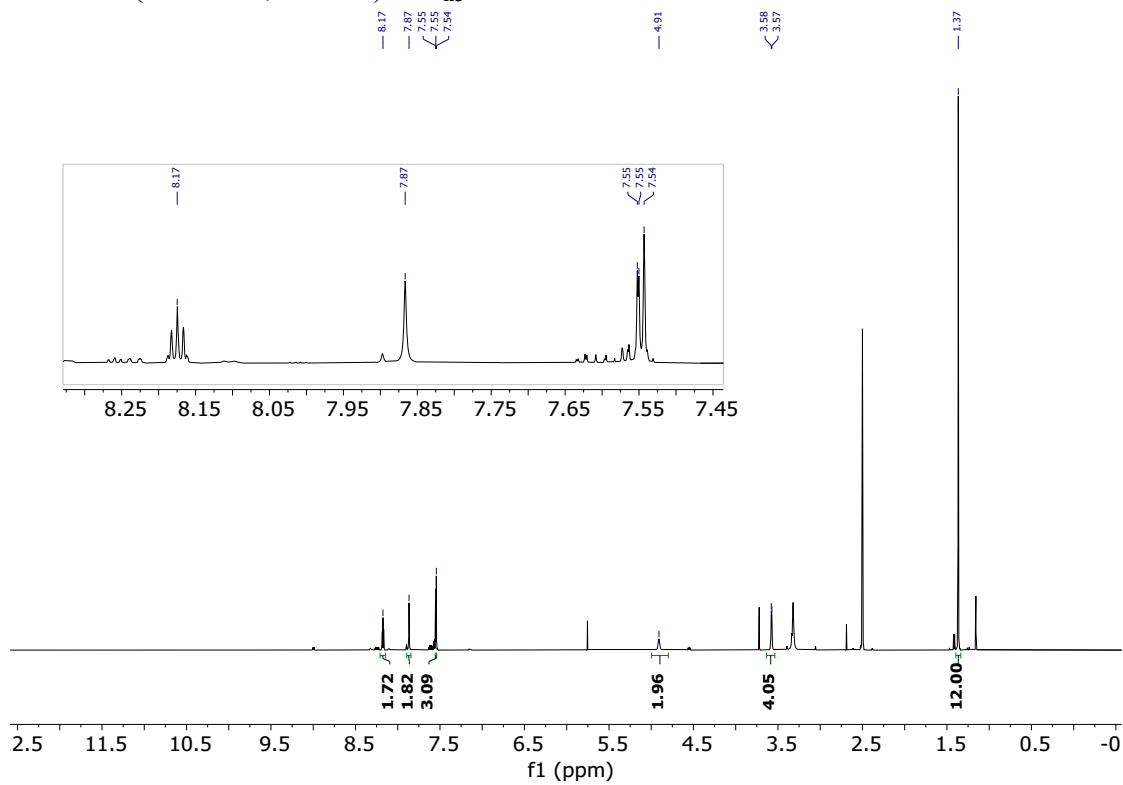
^1H NMR (600 MHz, CDCl_3) of **3_{n3}**.



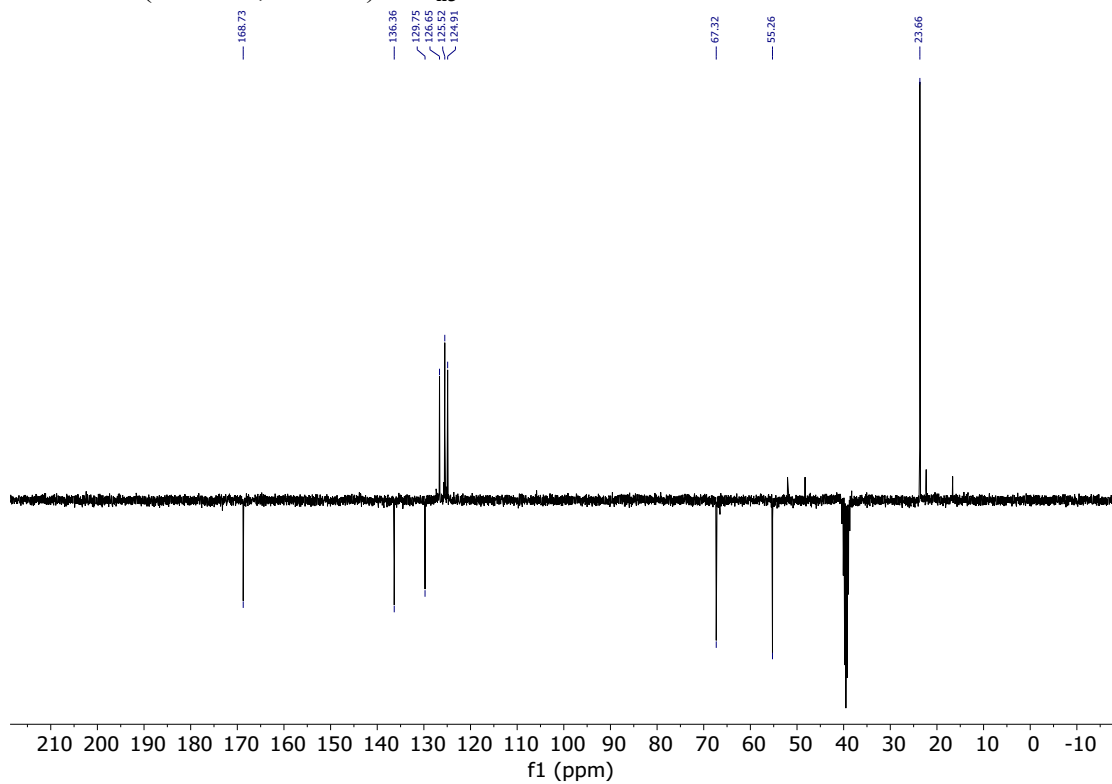
^{13}C NMR (151 MHz, CDCl_3) of **3_{n3}**.



^1H NMR (600 MHz, DMSO) of 4_{n3} .

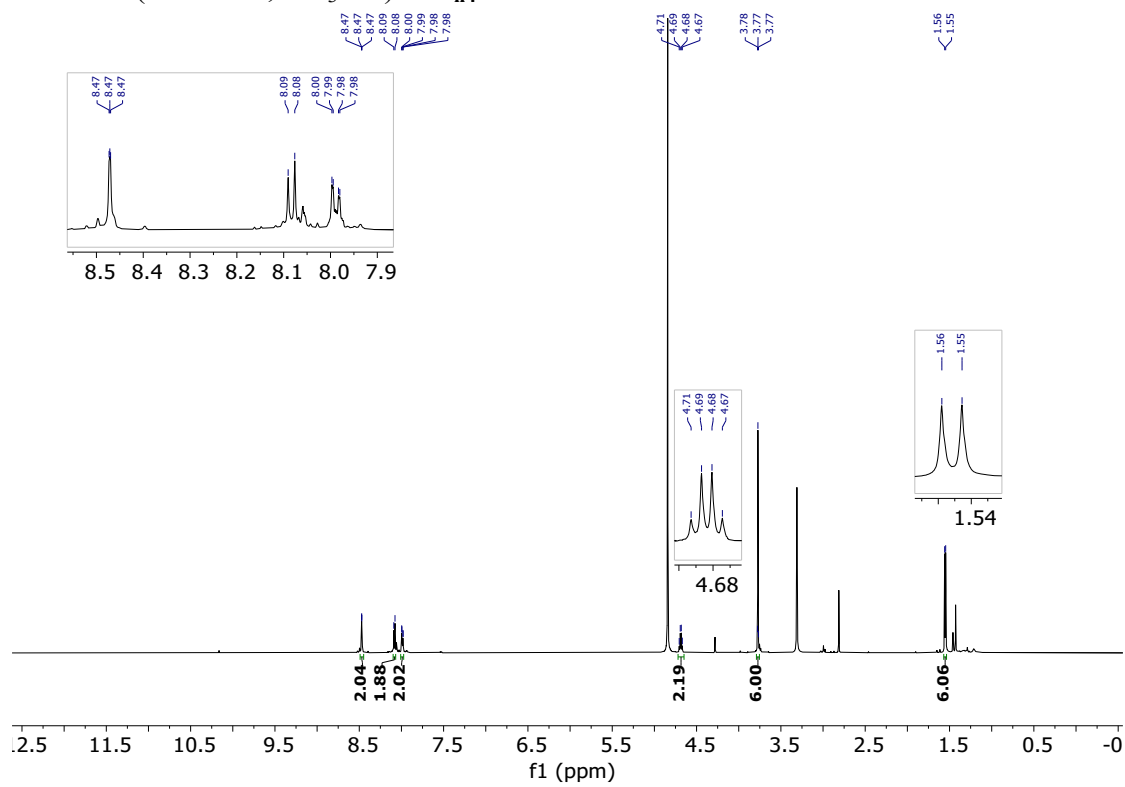


^{13}C NMR (75 MHz, DMSO) of 4_{n3} .

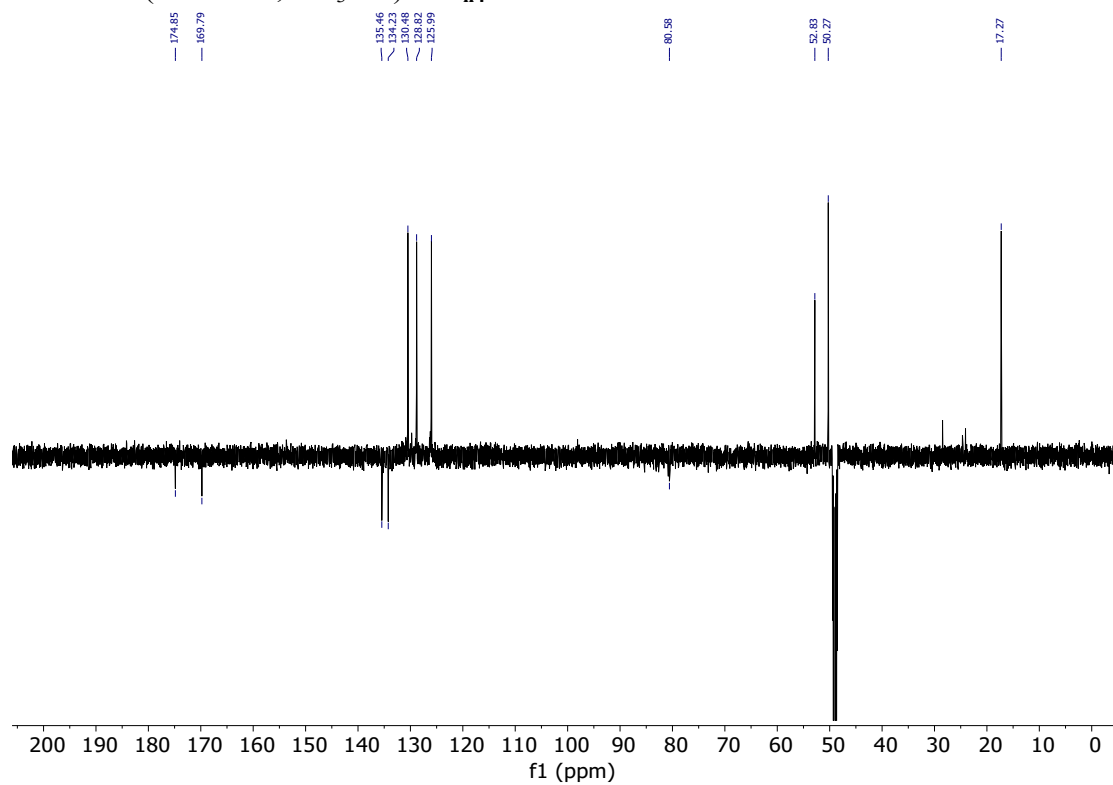


2.14. Reaction 16

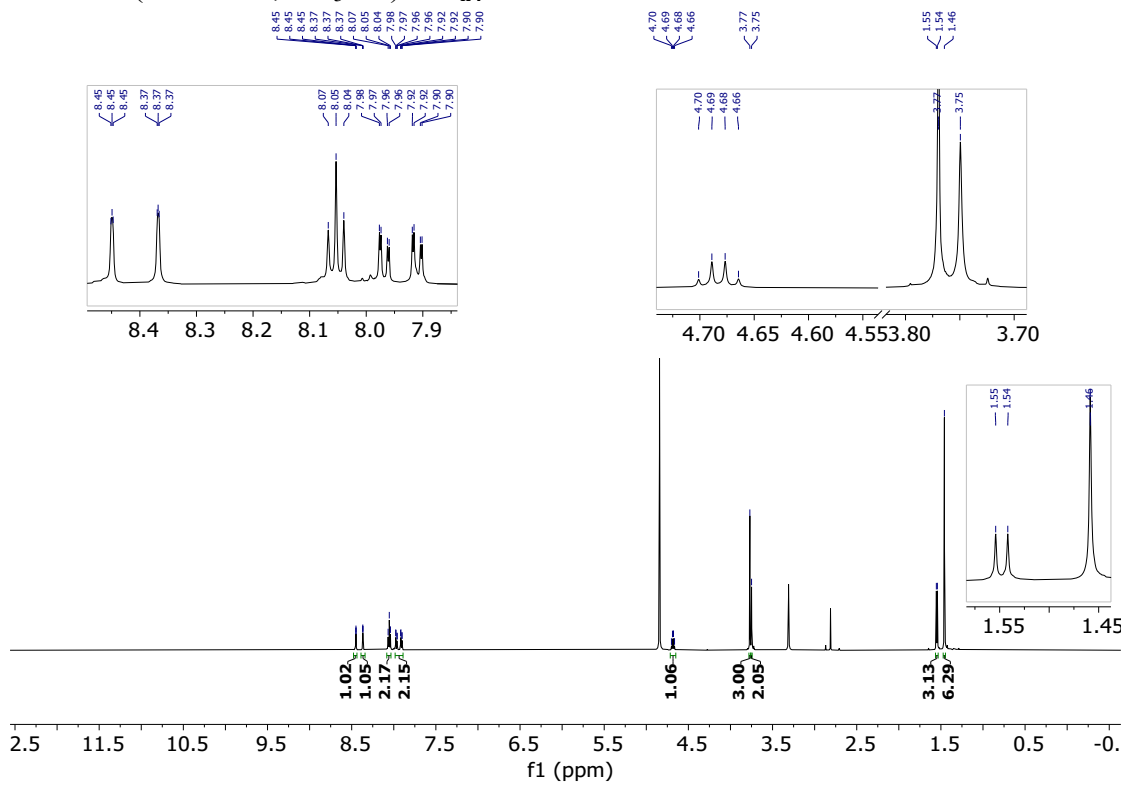
^1H NMR (600 MHz, CD_3OD) of 2_{n4} .



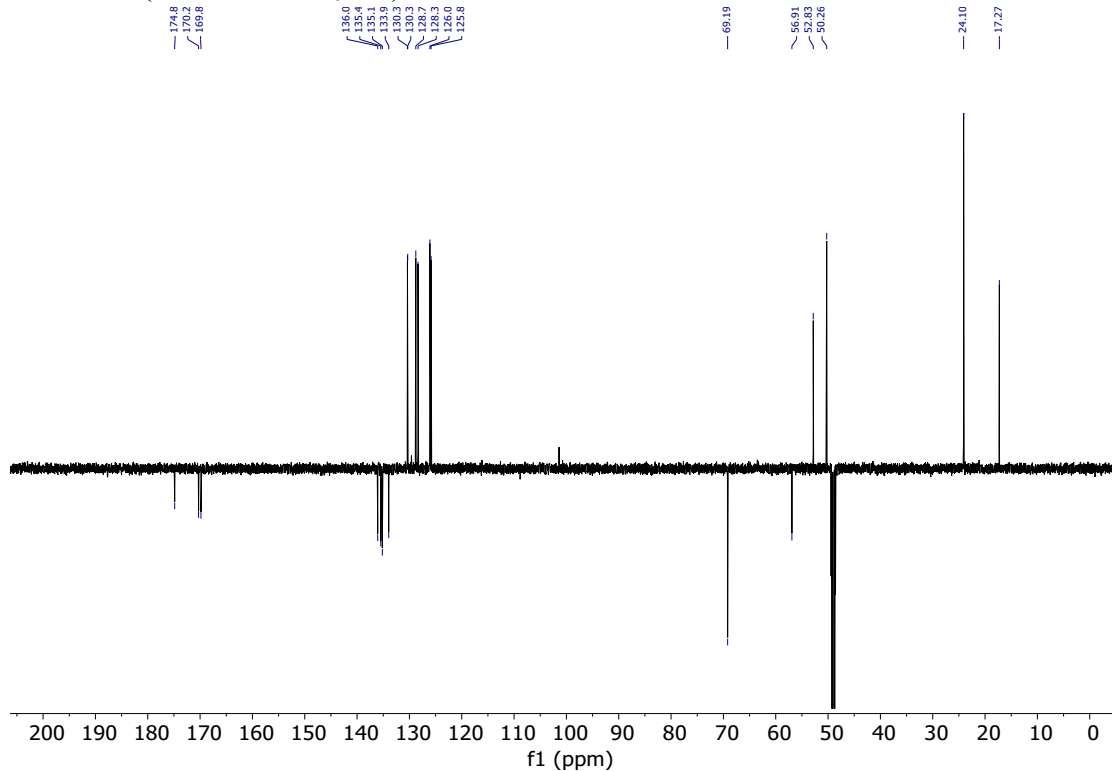
^{13}C NMR (151 MHz, CD_3OD) of 2_{n4} .



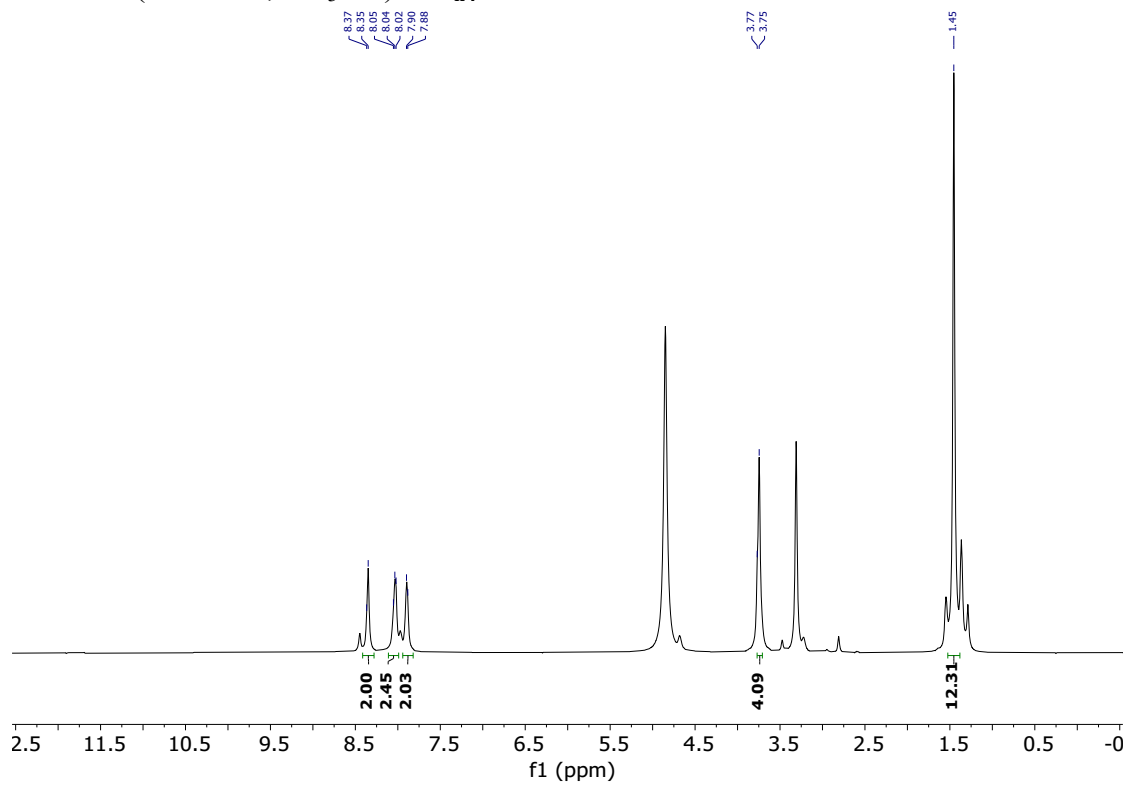
^1H NMR (600 MHz, CD_3OD) of 3_{n4} .



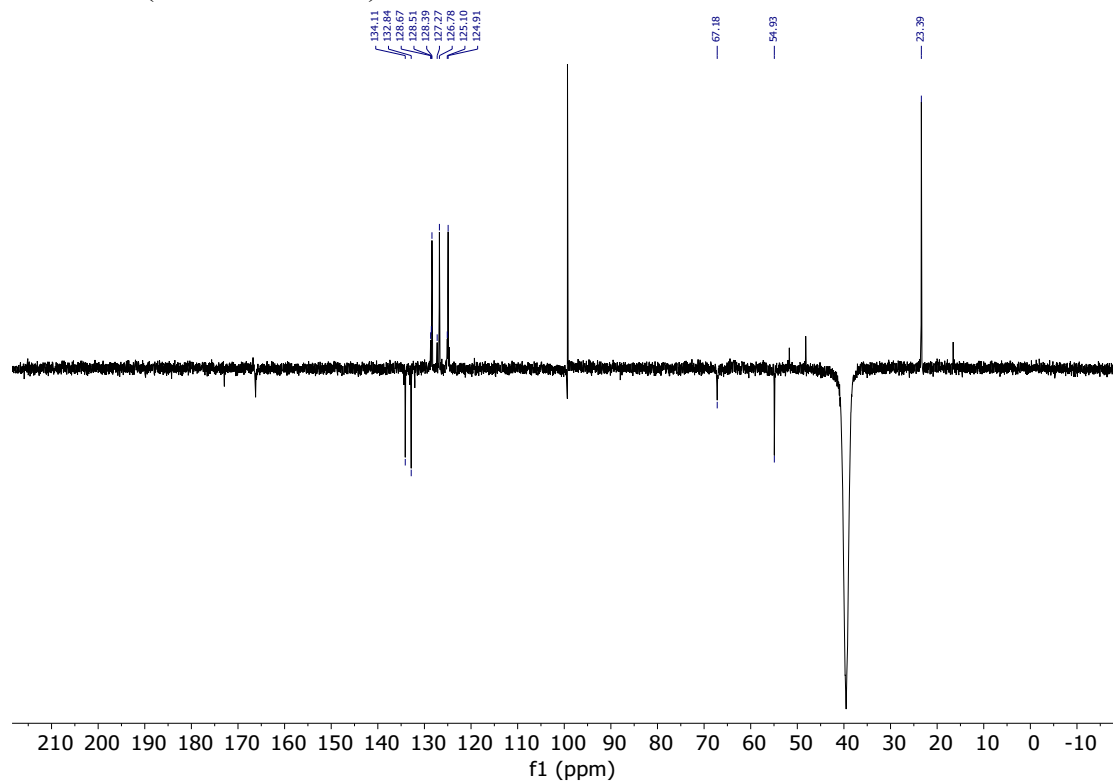
^{13}C NMR (151 MHz, CD_3OD) of 3_{n4} .



^1H NMR (600 MHz, CD_3OD) of 4_{n4} .

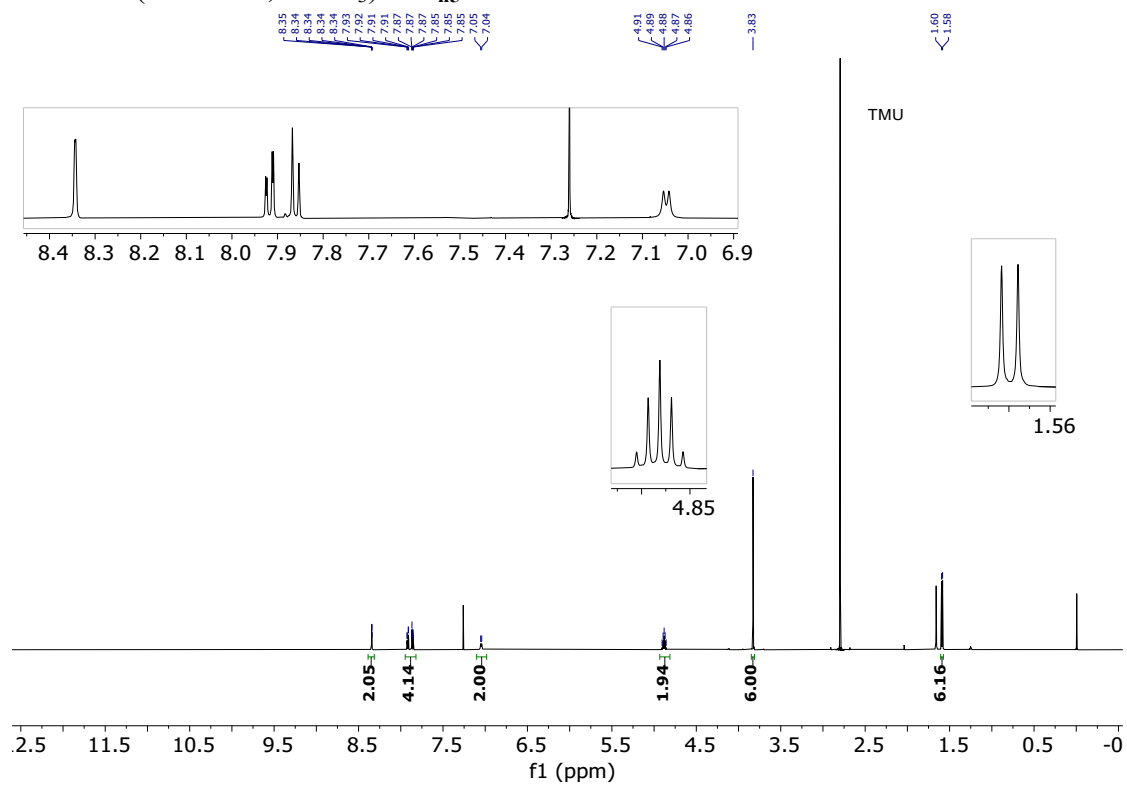


^{13}C NMR (75 MHz, DMSO) of 4_{n4} .

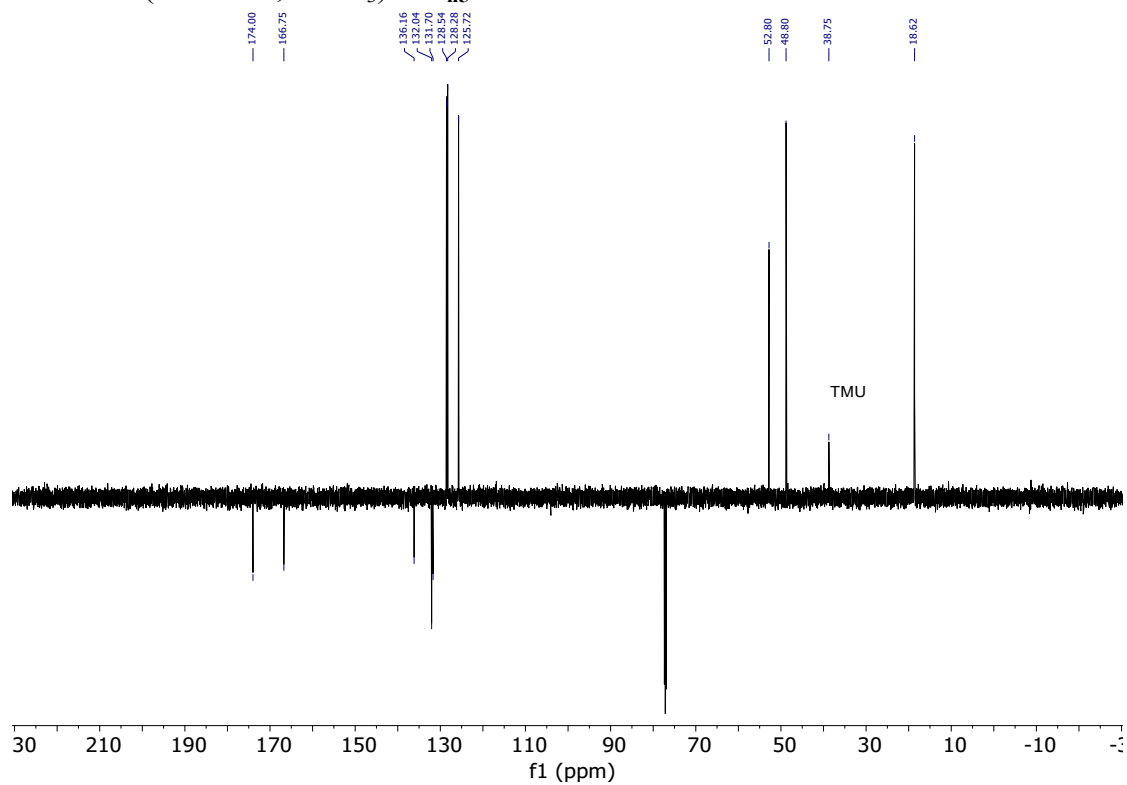


2.15. Reaction 17

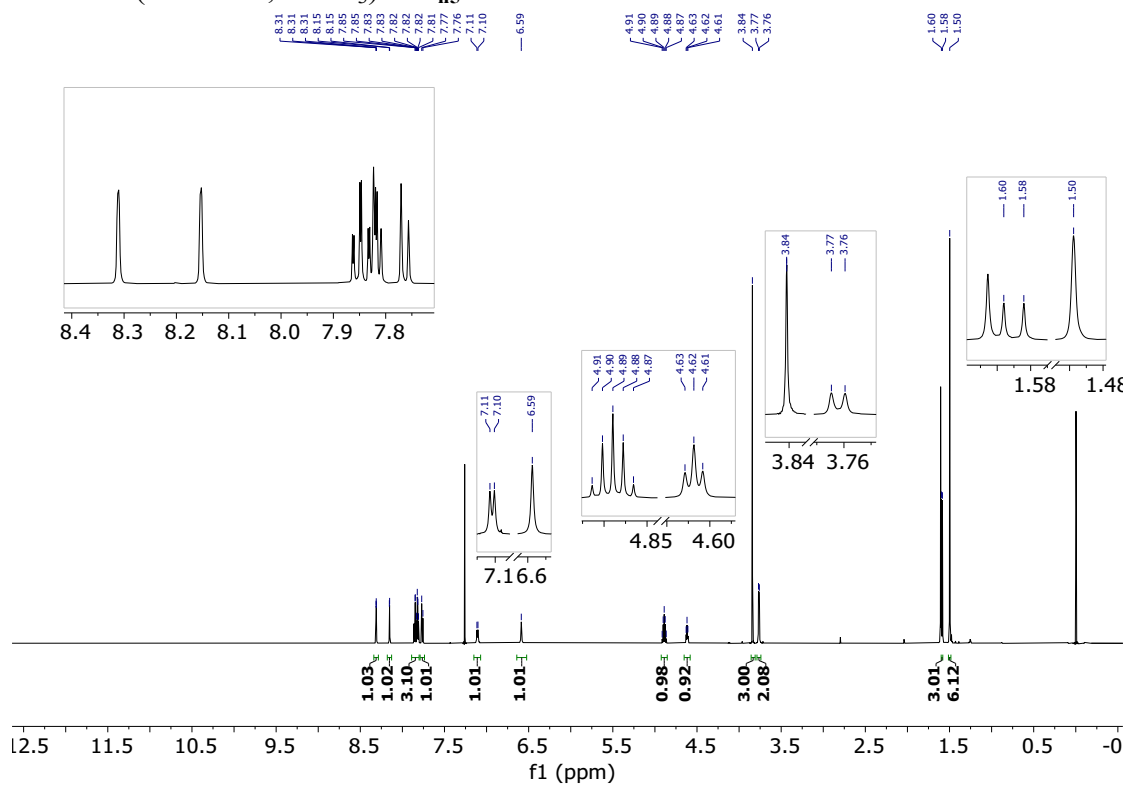
^1H NMR (600 MHz, CDCl_3) of **2_{n5}**.



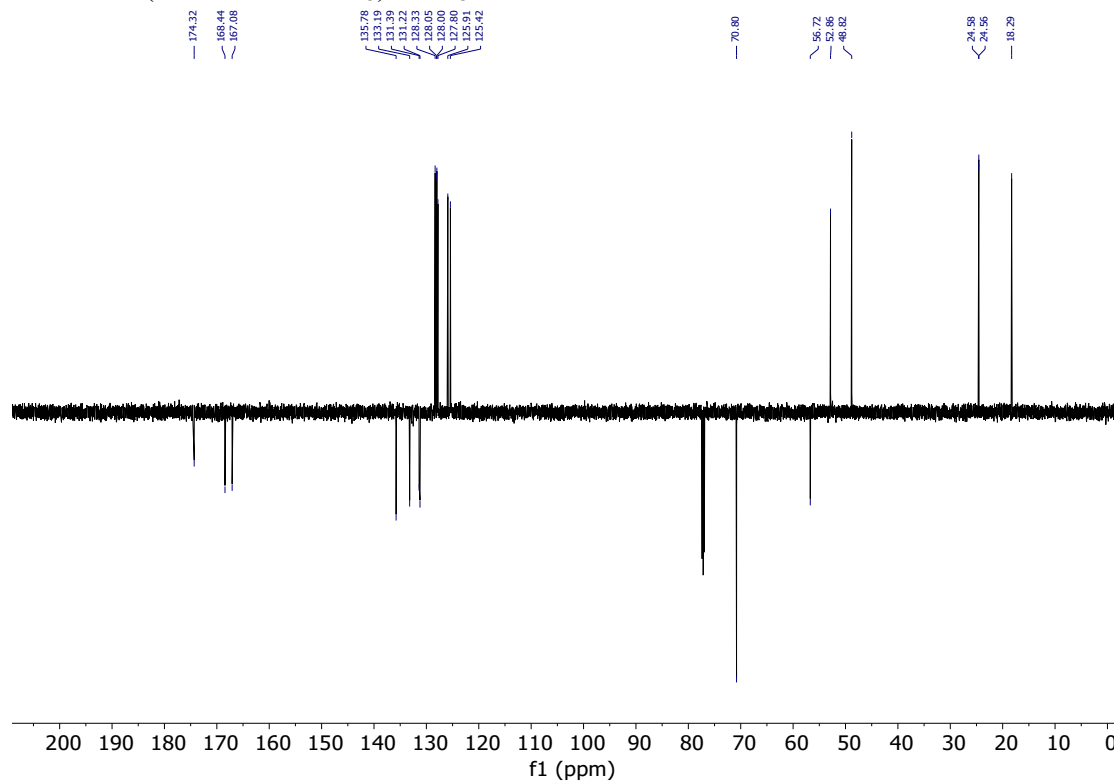
^{13}C NMR (151 MHz, CDCl_3) of **2_{n5}**.



^1H NMR (600 MHz, CDCl_3) of 3_{n5} .

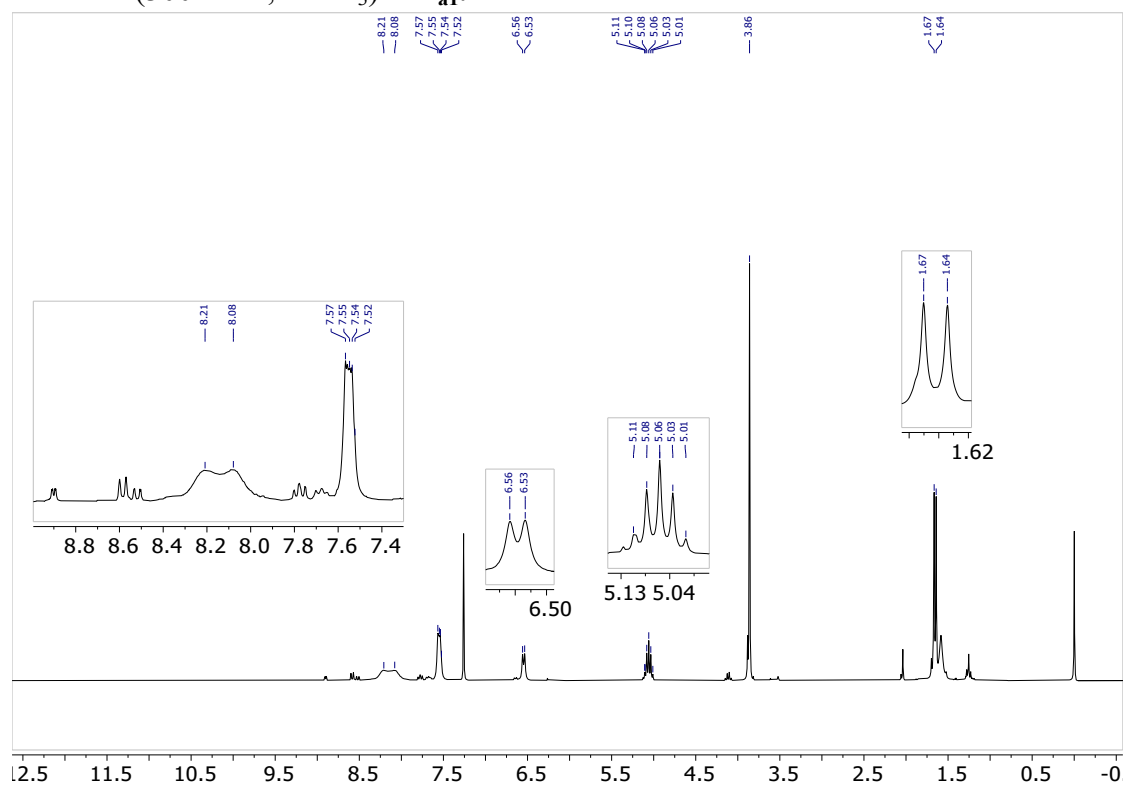


^{13}C NMR (151 MHz, CDCl_3) of 3_{n5} .

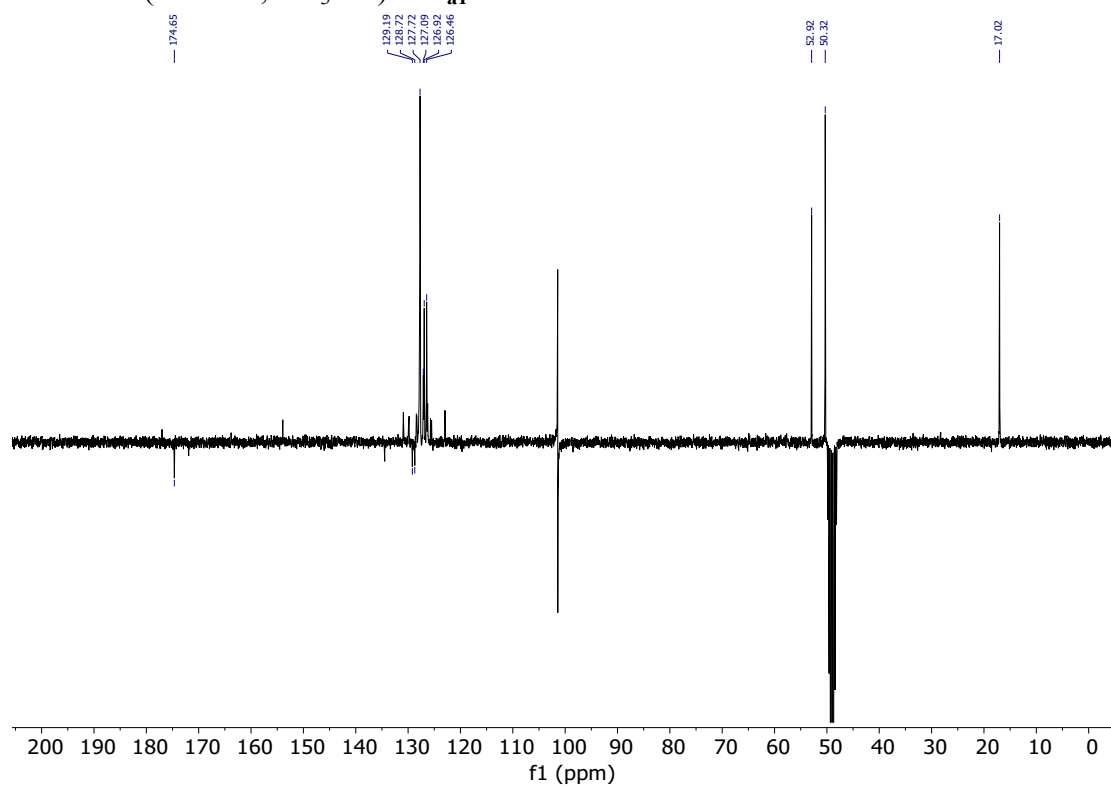


2.16. Reaction 18

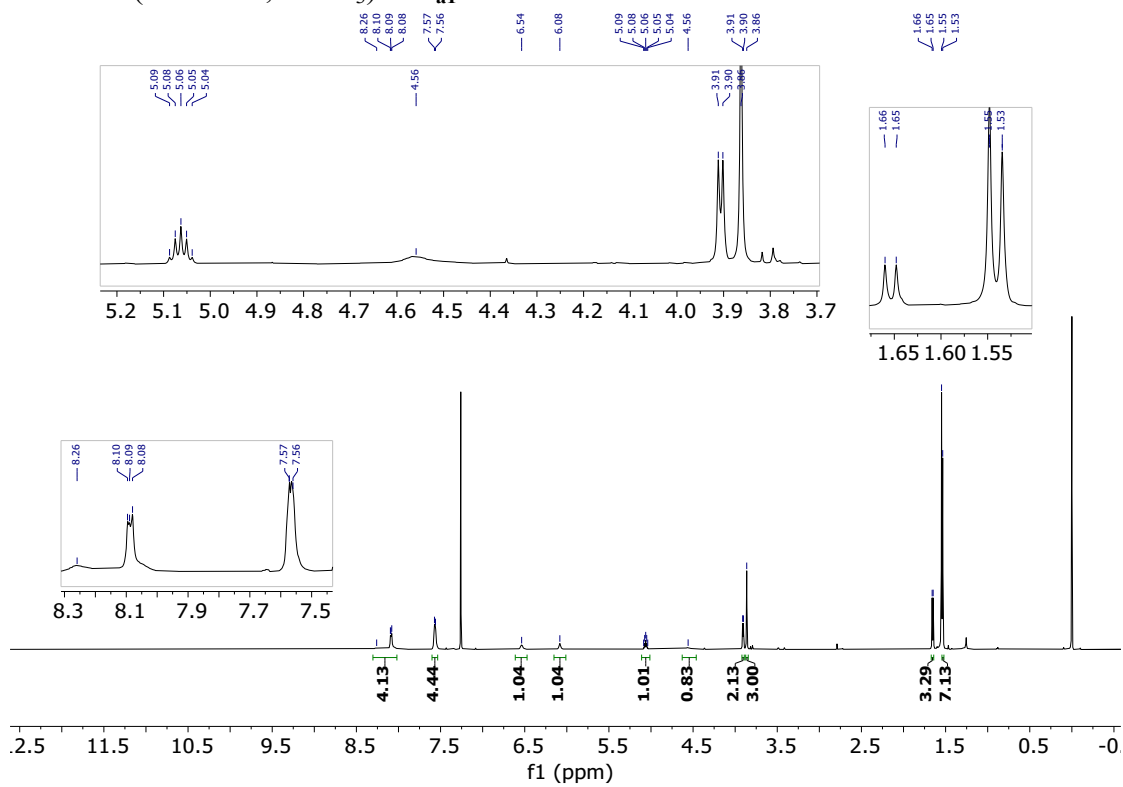
^1H NMR (300 MHz, CDCl_3) of **2a1**.



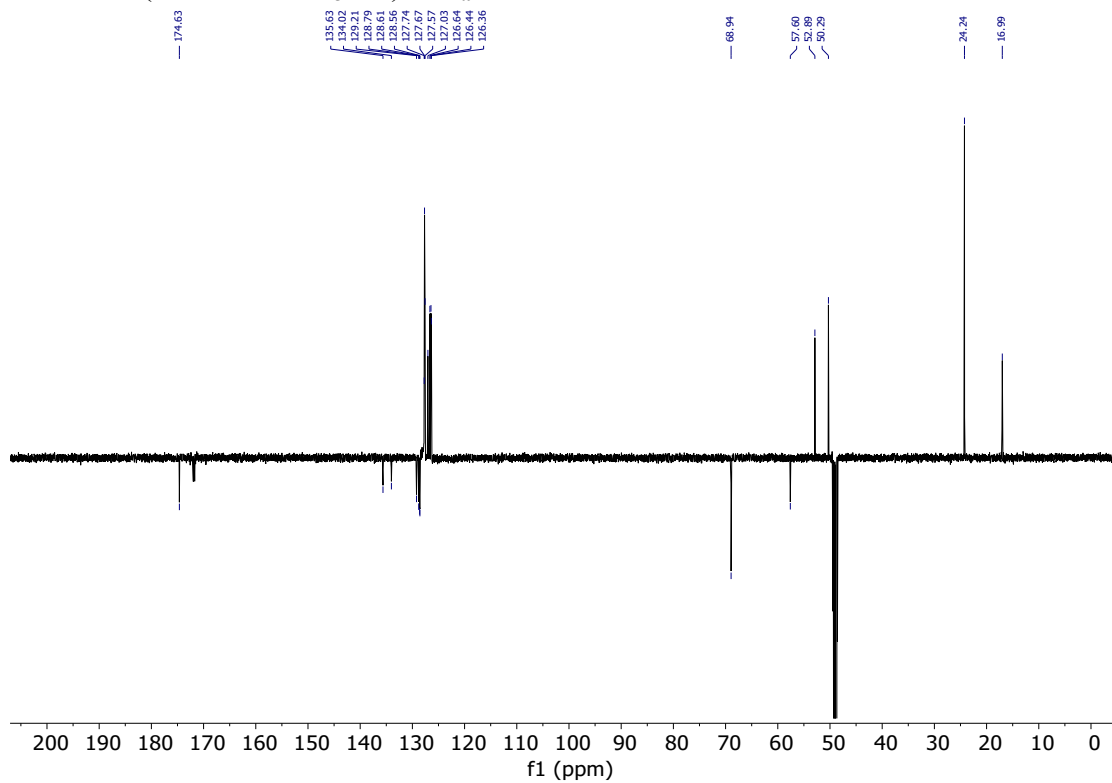
^{13}C NMR (75 MHz, CD_3OD) of **2a1**.



^1H NMR (300 MHz, CDCl_3) of **3a1**.

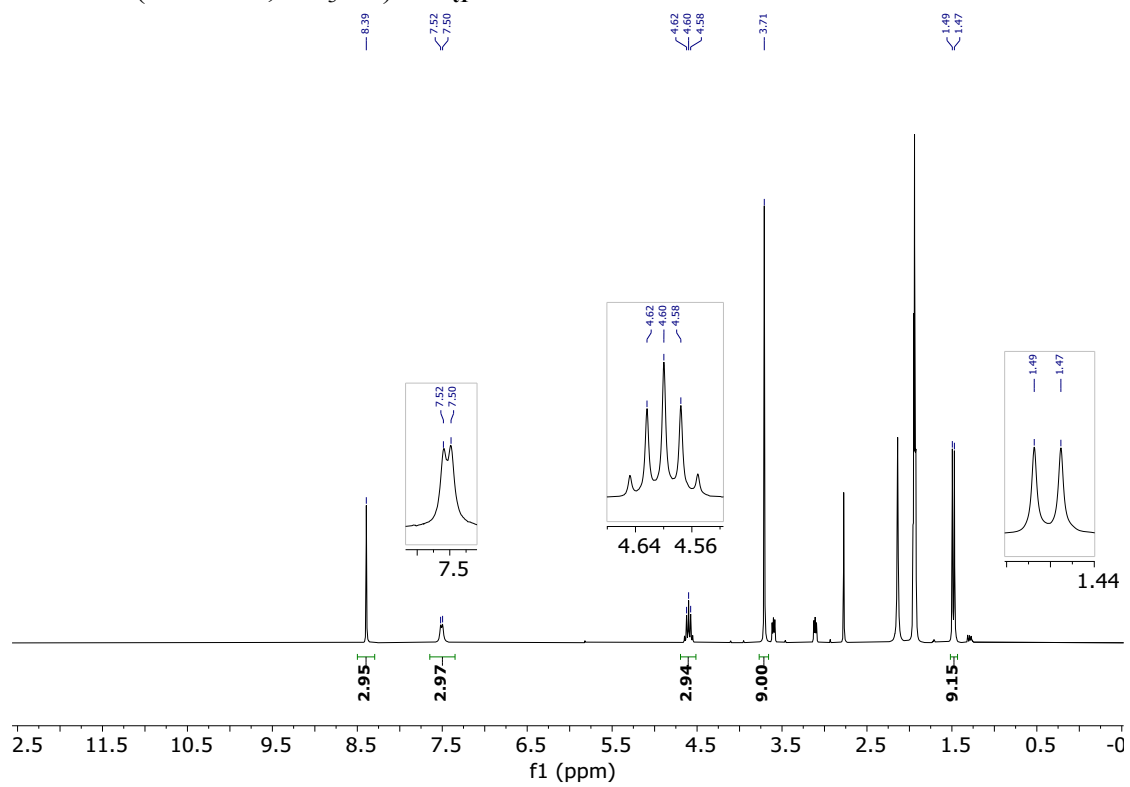


^{13}C NMR (151 MHz, CD_3OD) of **3a1**.

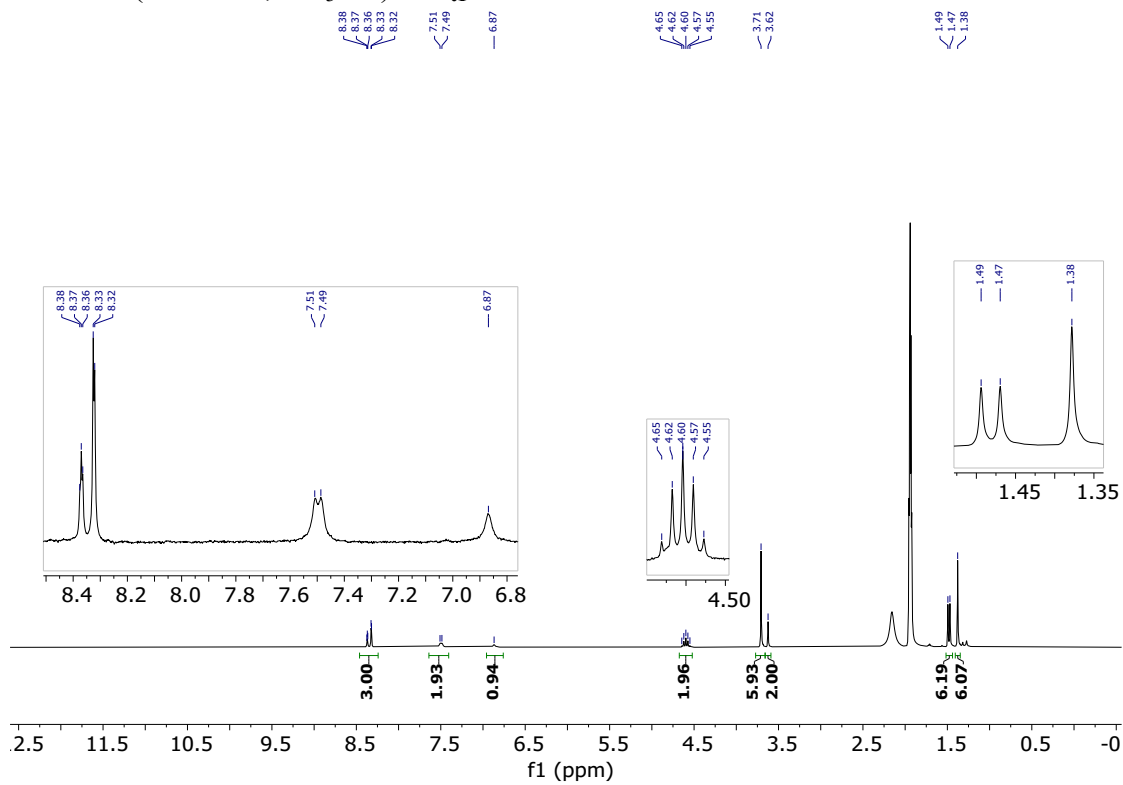


2.17. Reaction 19

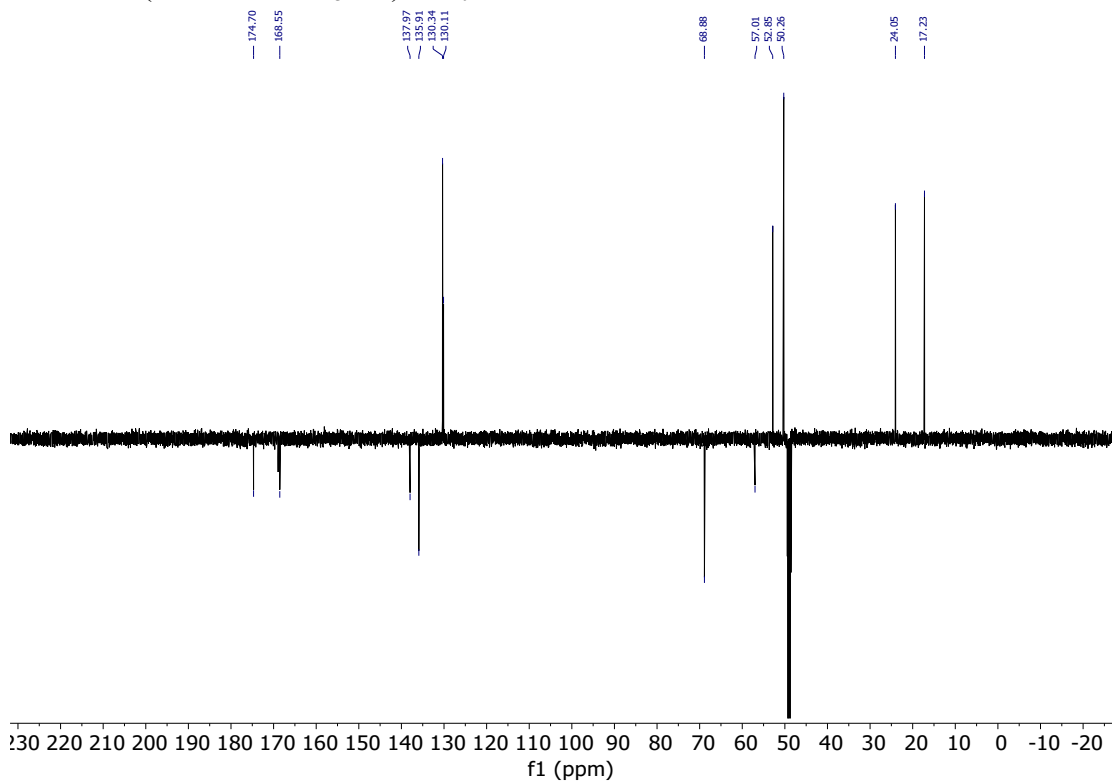
^1H NMR (300 MHz, CD_3CN) of **211**.



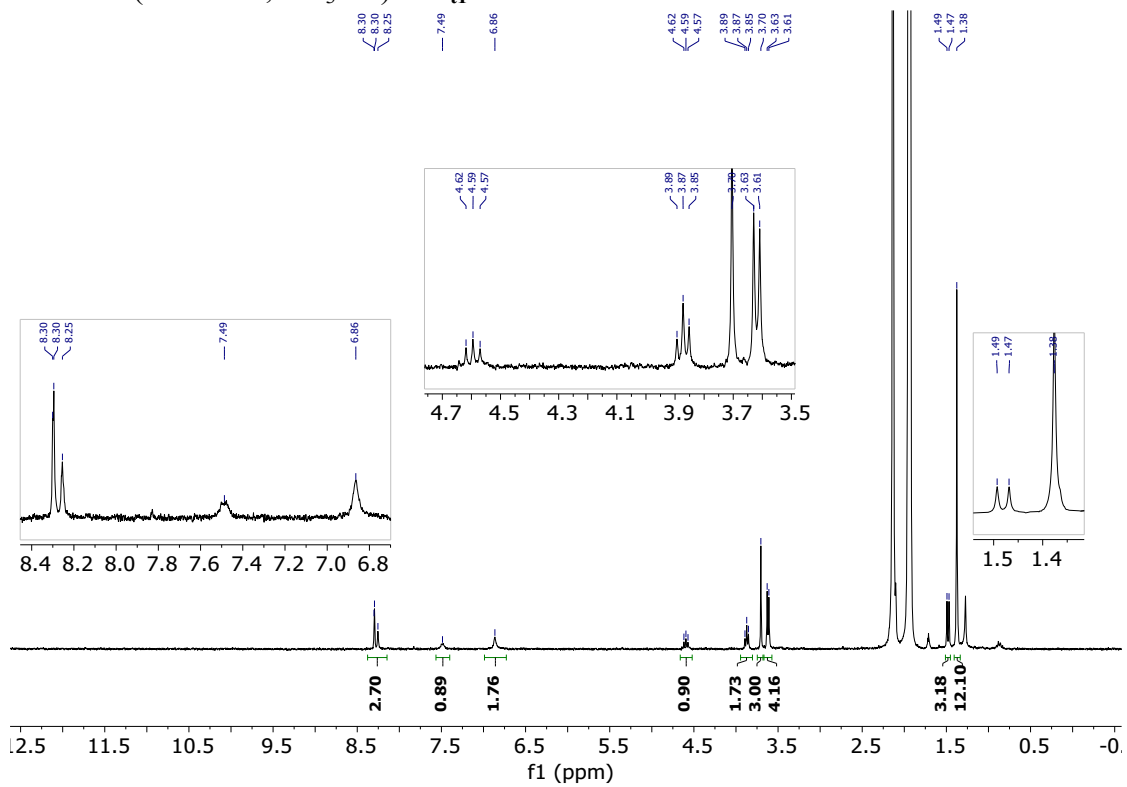
^1H NMR (300 MHz, CD_3CN) of **3_{tl}**.



^{13}C NMR (151 MHz, CD_3OD) of **3_{tl}**.

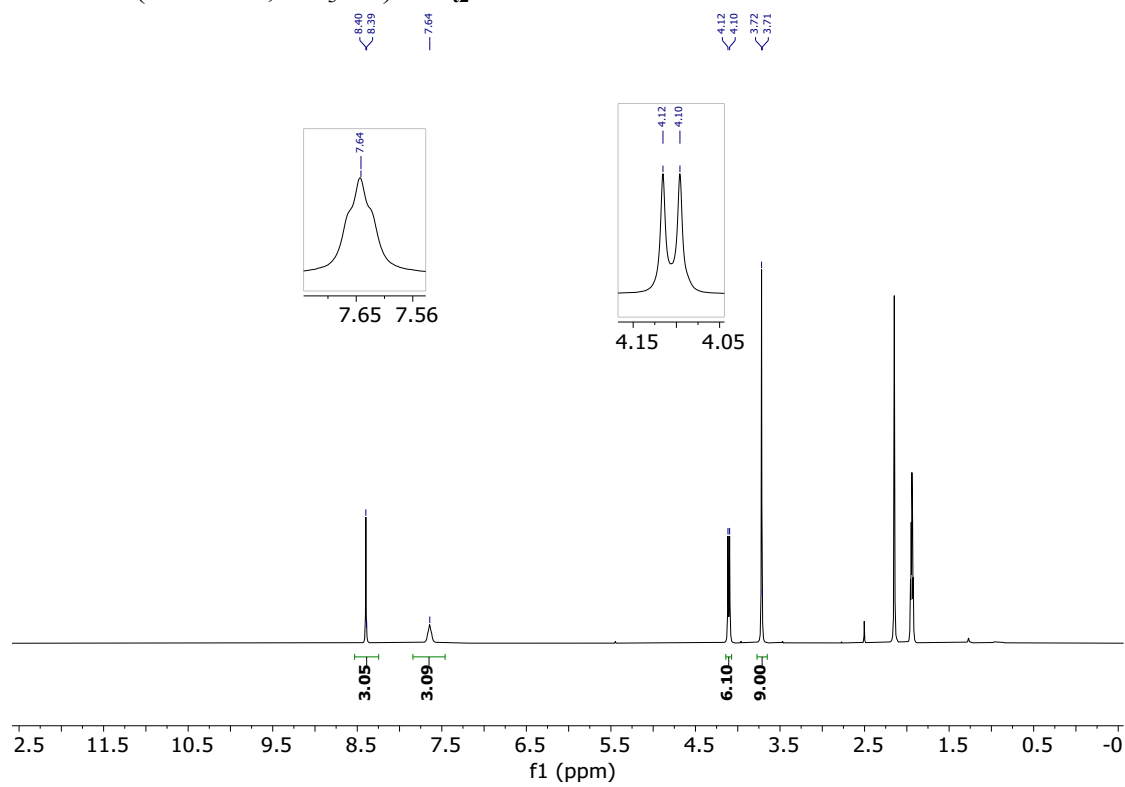


^1H NMR (300 MHz, CD_3CN) of **4₁₁**.

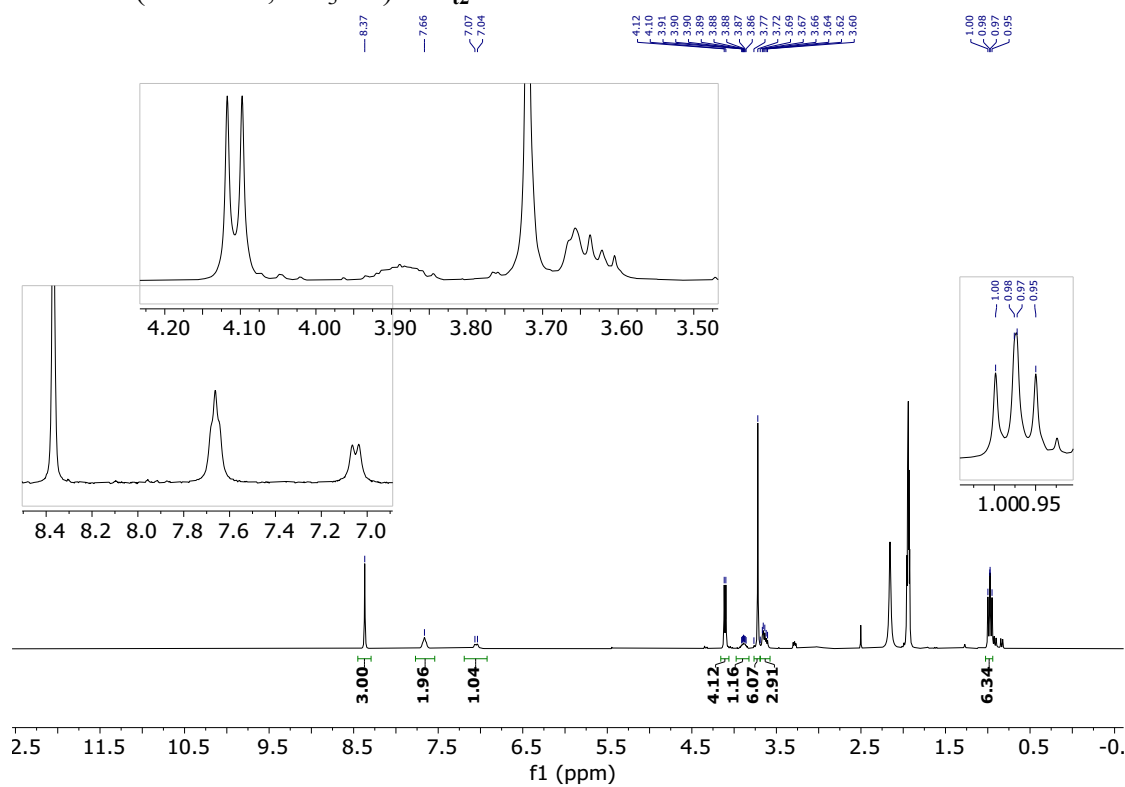


2.18. Reaction 20

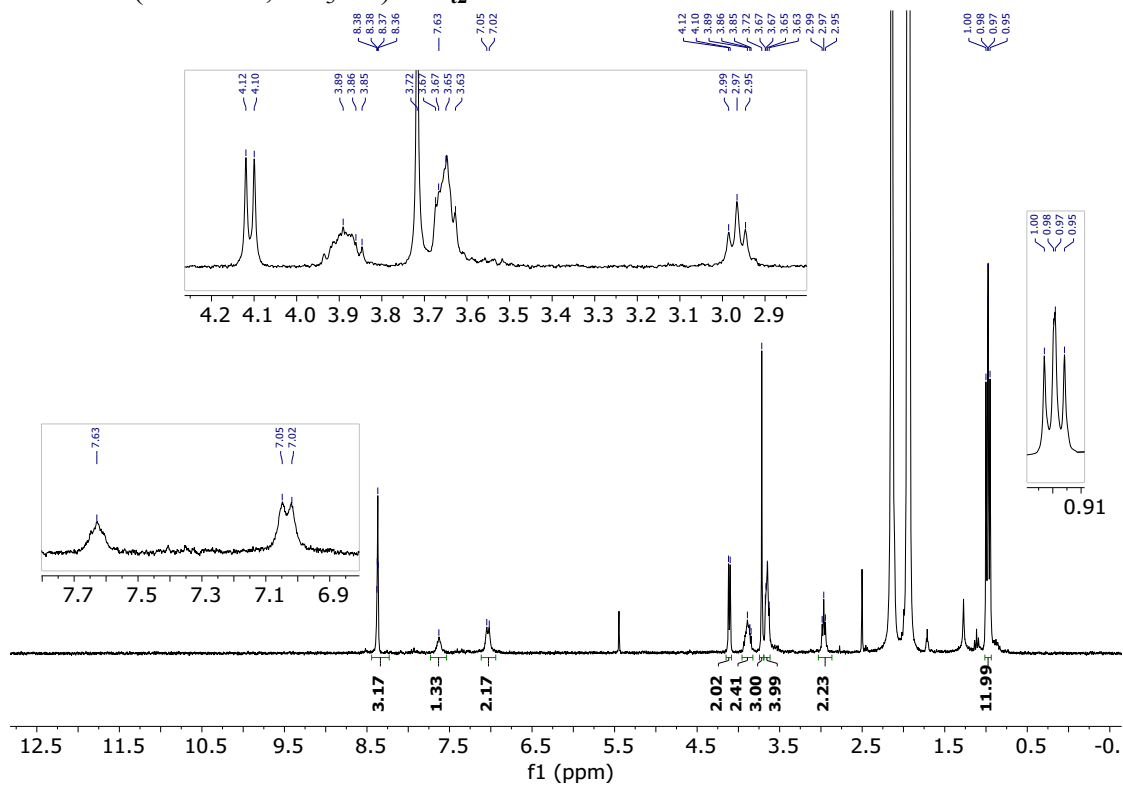
^1H NMR (300 MHz, CD_3CN) of **2**₁₂.



^1H NMR (300 MHz, CD_3CN) of **3**₁₂.

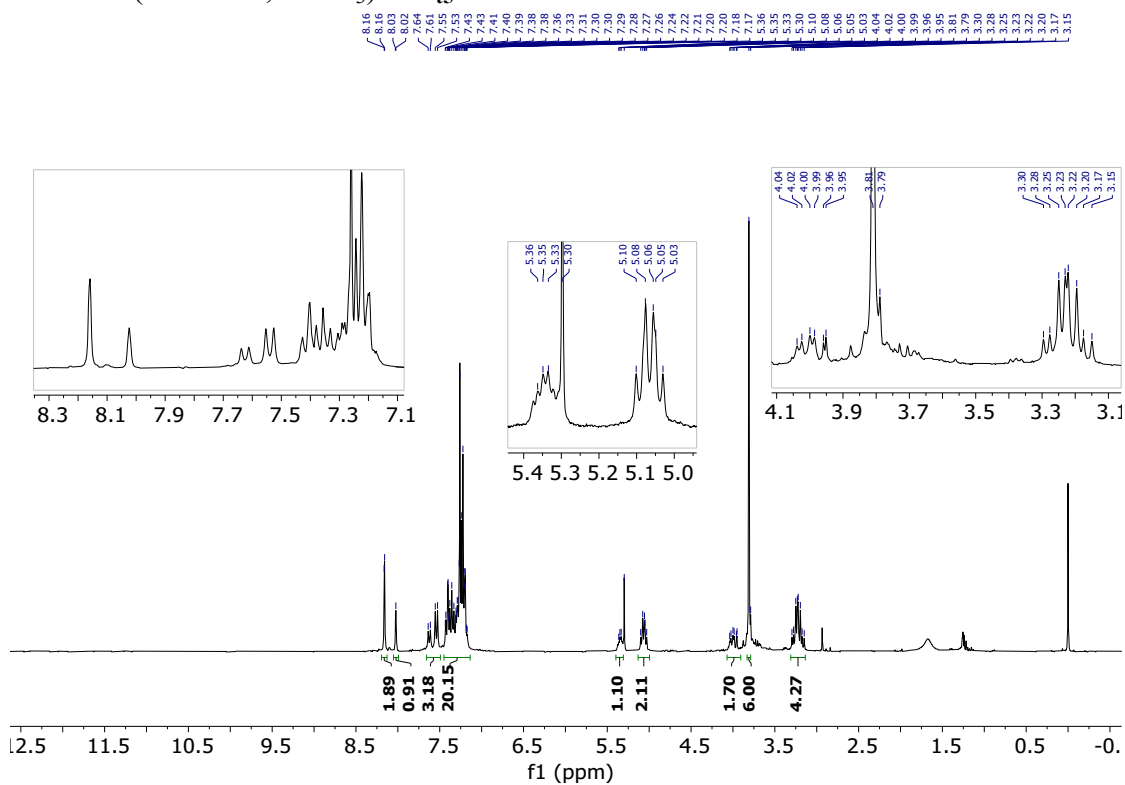


^1H NMR (300 MHz, CD_3CN) of 4_{12} .

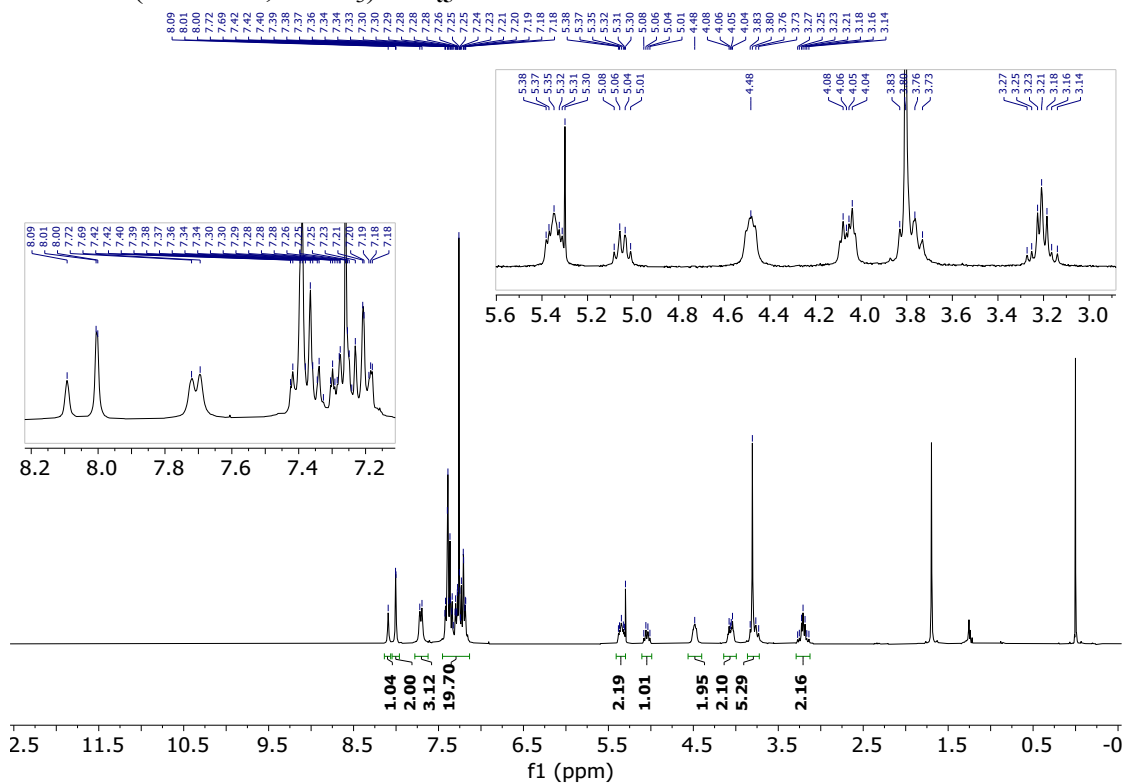


2.19. Reaction 21

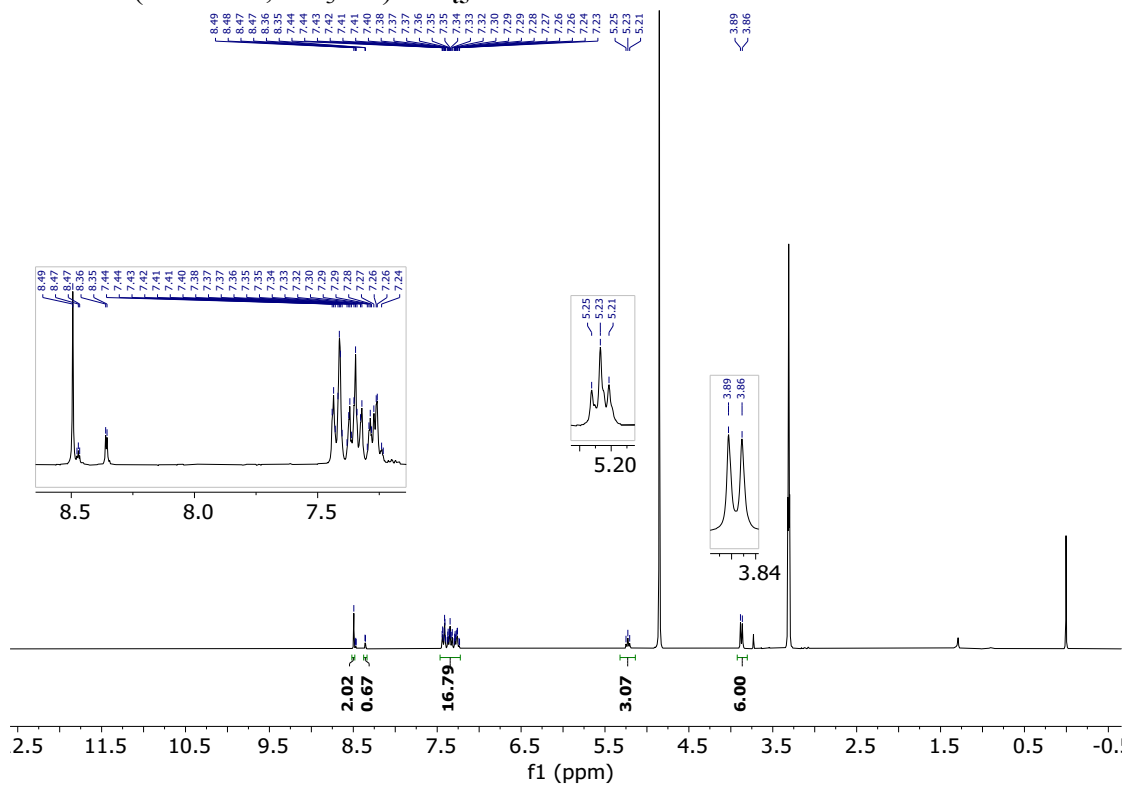
^1H NMR (300 MHz, CDCl_3) of **3₁₅**.



^1H NMR (300 MHz, CDCl_3) of **4₁₅**.

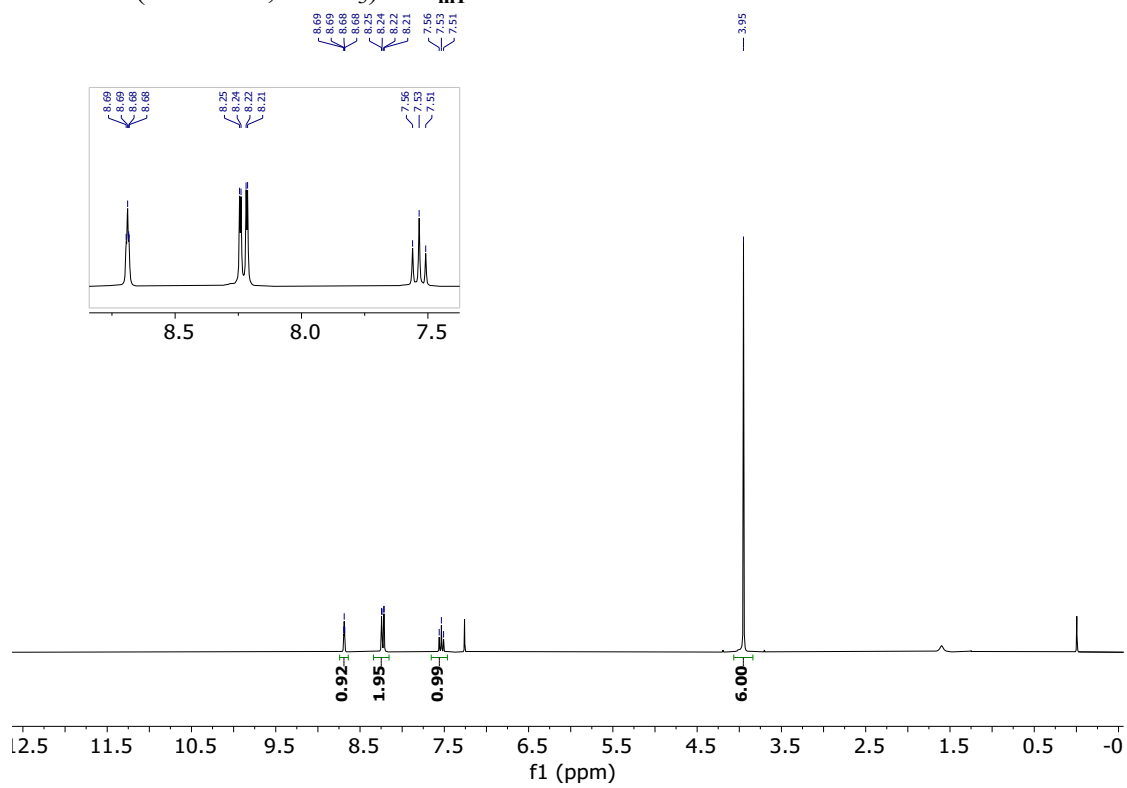


¹H NMR (300 MHz, CD₃OD) of **5_{t5}**.

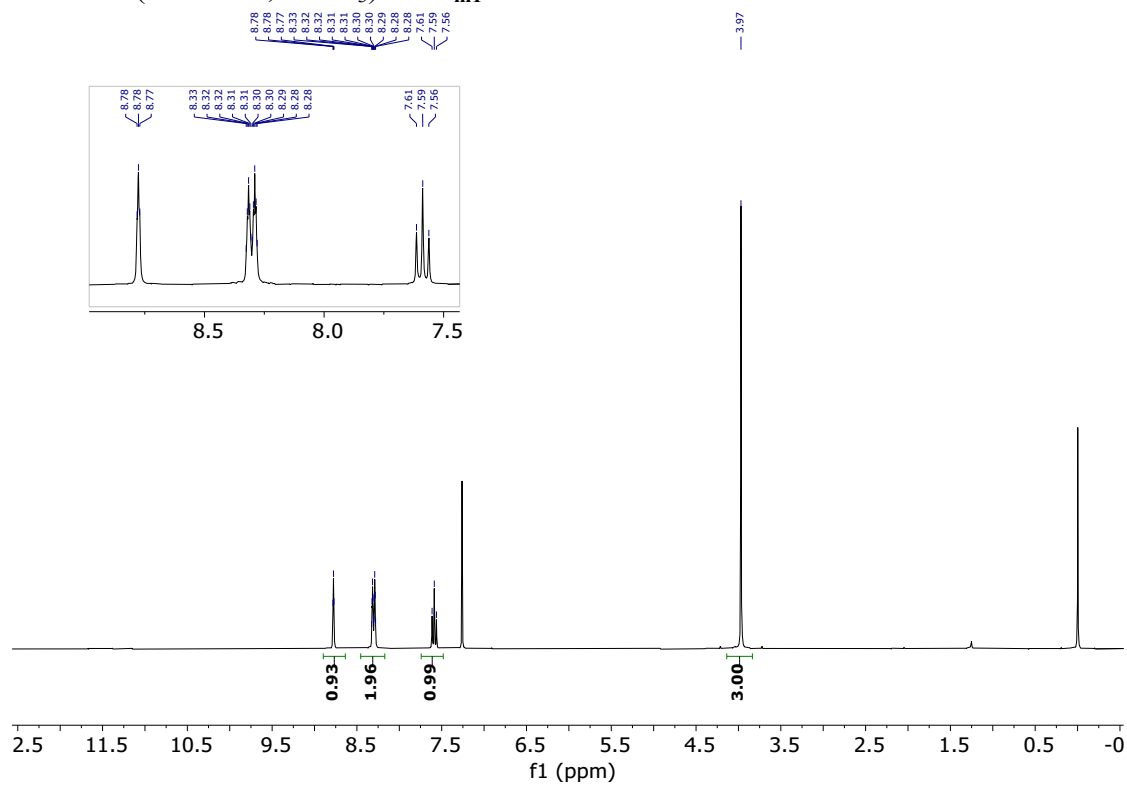


2.20. Linear reaction sequence 1

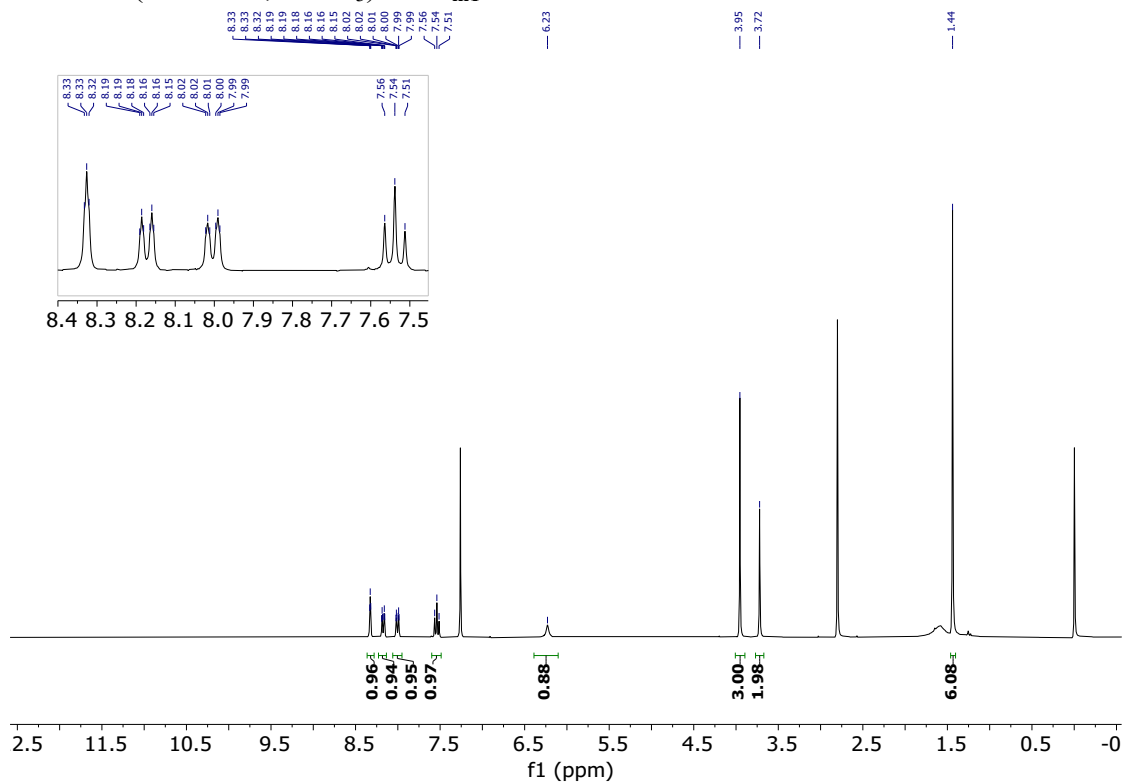
^1H NMR (300 MHz, CDCl_3) of **9_{m1}**.



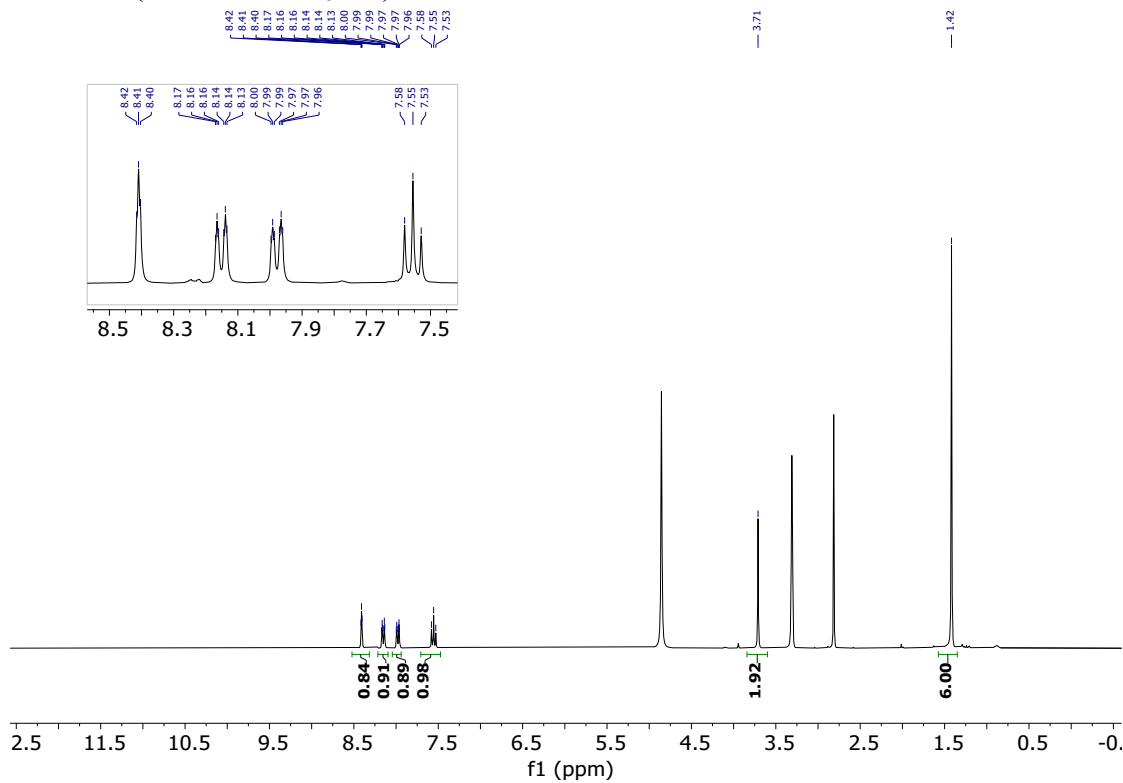
^1H NMR (300 MHz, CDCl_3) of **10_{m1}**.



^1H NMR (300 MHz, CDCl_3) of **11**_{m1}.

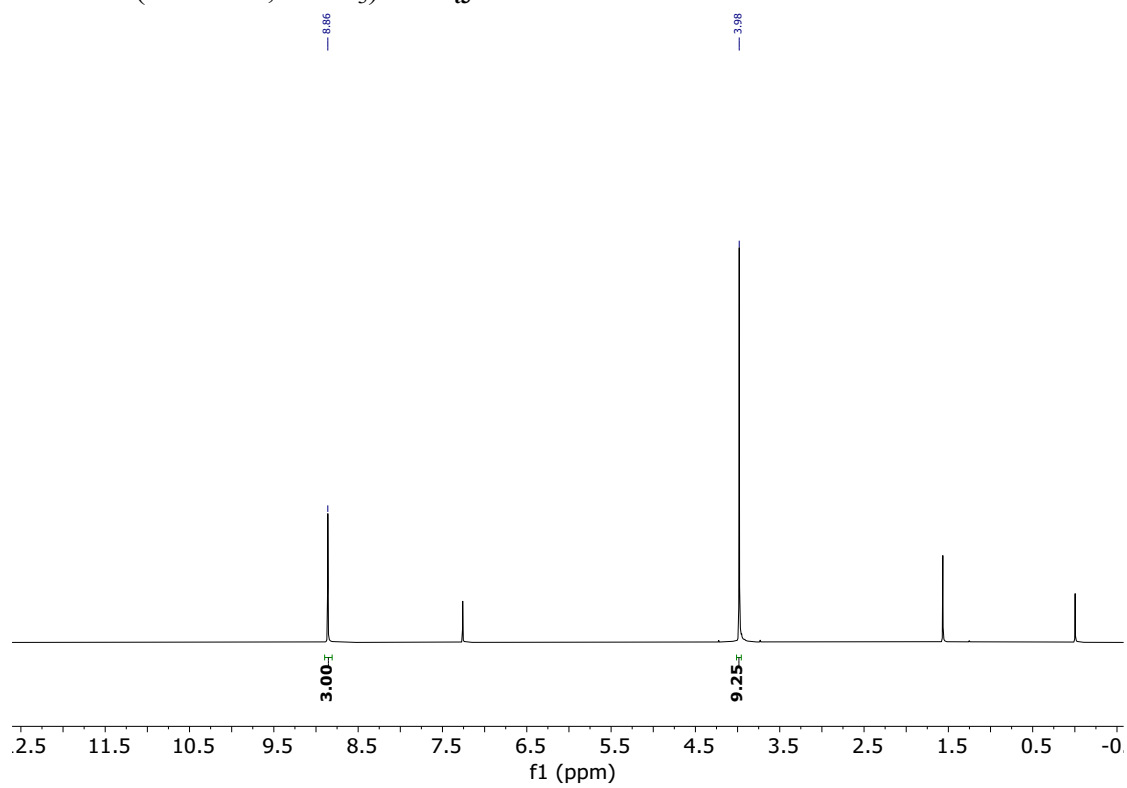


^1H NMR (300 MHz, CD_3OD) of **12**_{m1}.

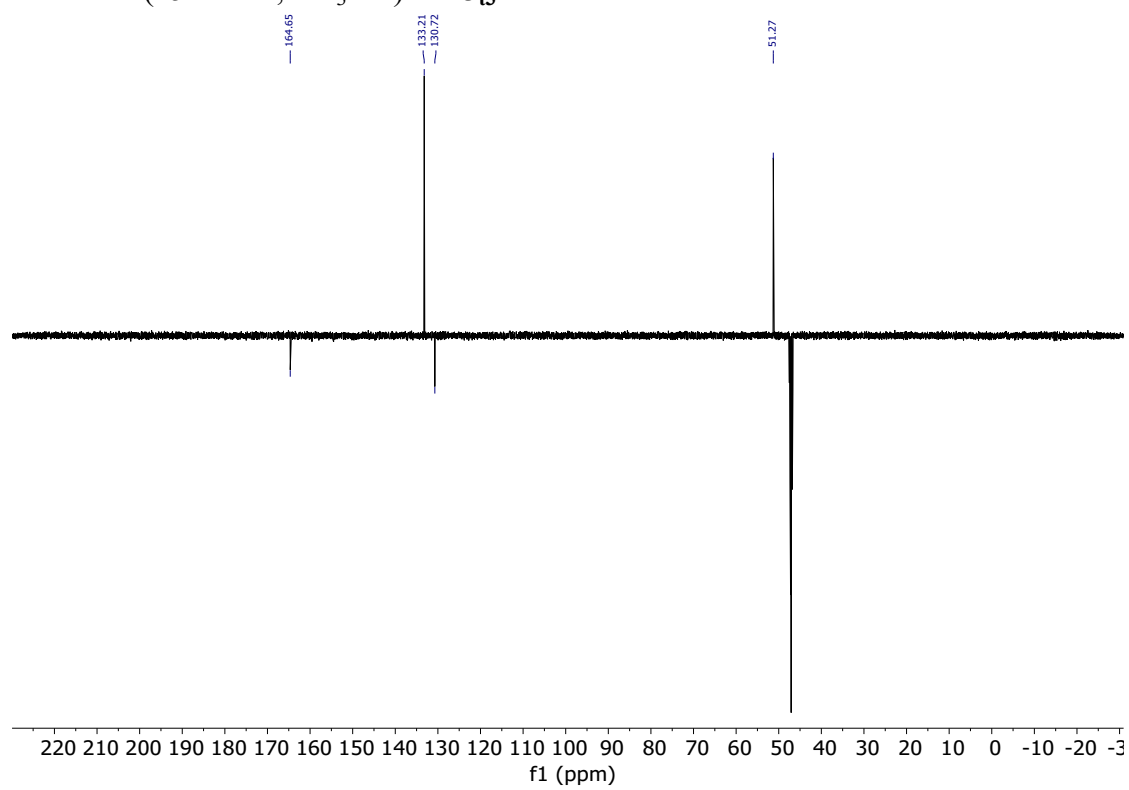


2.21. Linear reaction sequence 2

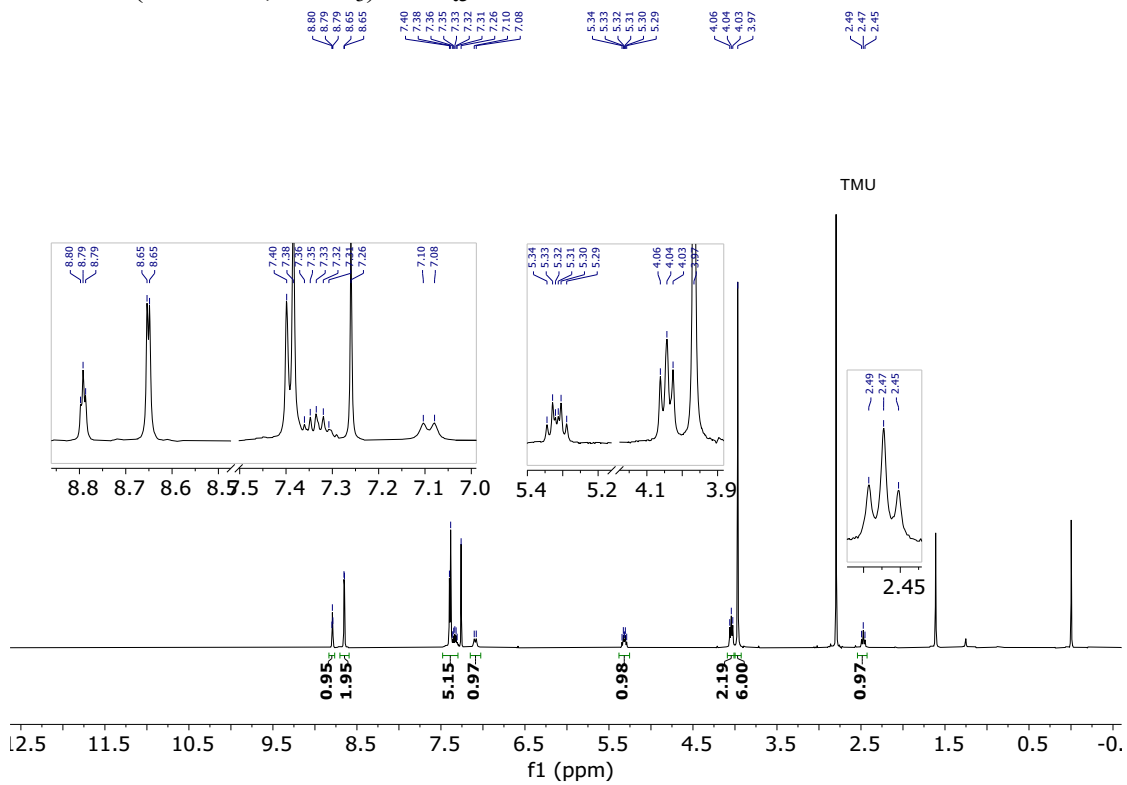
^1H NMR (300 MHz, CDCl_3) of **13**_{t3}.



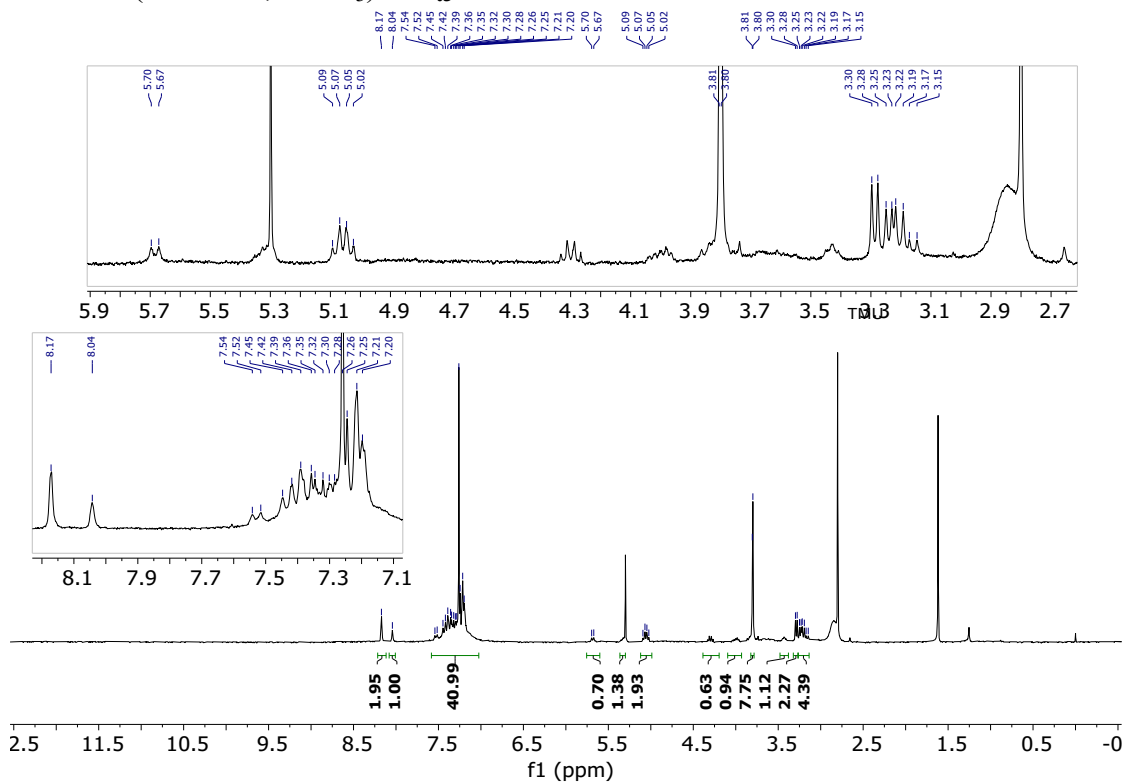
^{13}C NMR (151 MHz, CD_3OD) of **13**_{t3}.



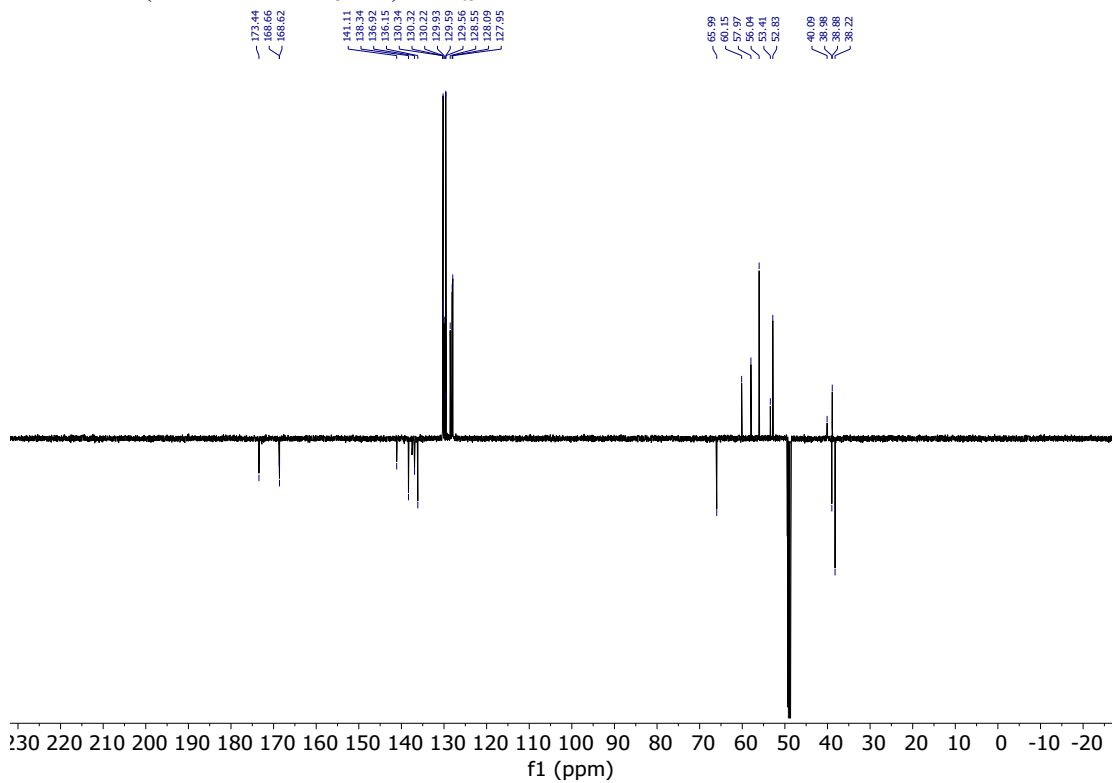
^1H NMR (300 MHz, CDCl_3) of **15t3**.



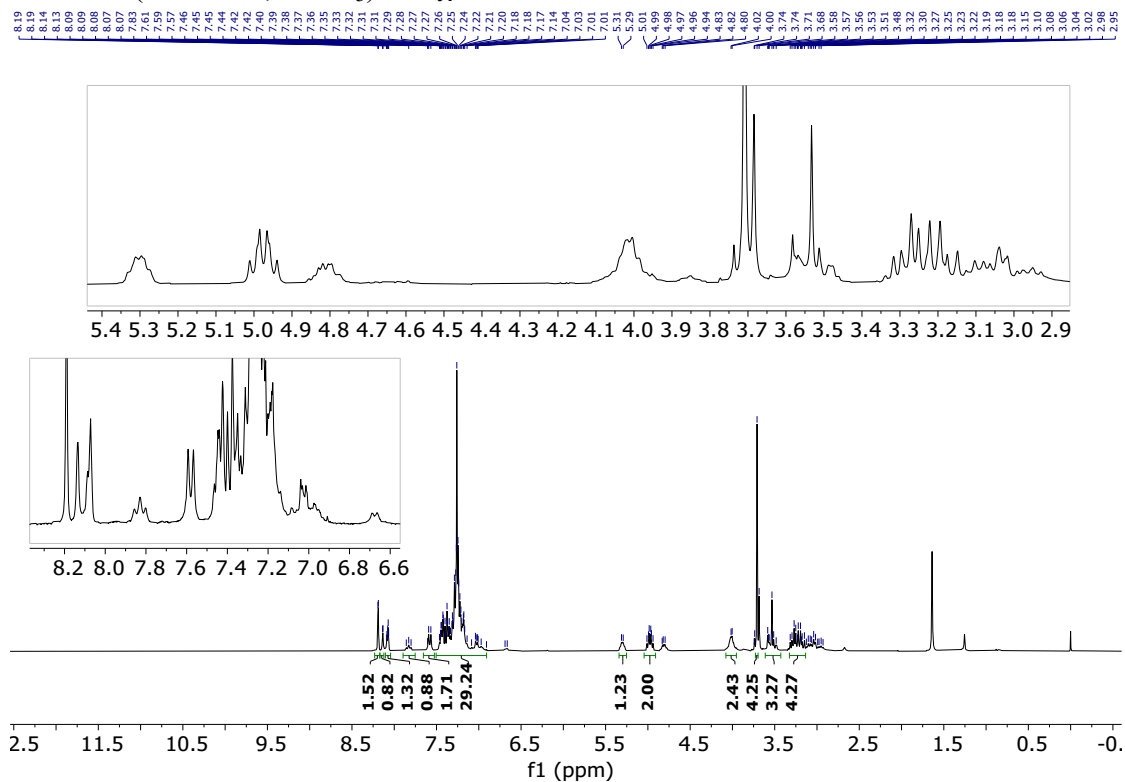
^1H NMR (300 MHz, CDCl_3) of **3**₁₃.



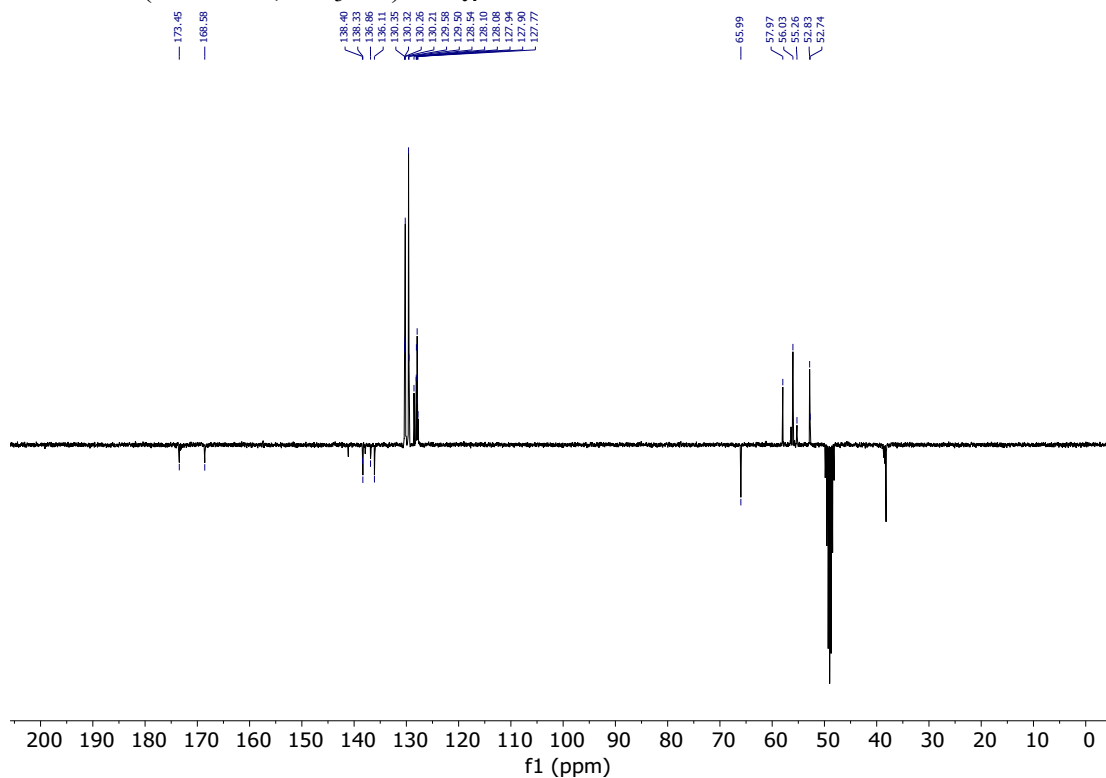
^{13}C NMR (151 MHz, CD_3OD) of **3**₁₃.



¹H NMR (300 MHz, CDCl₃) of **3**_{t4}.

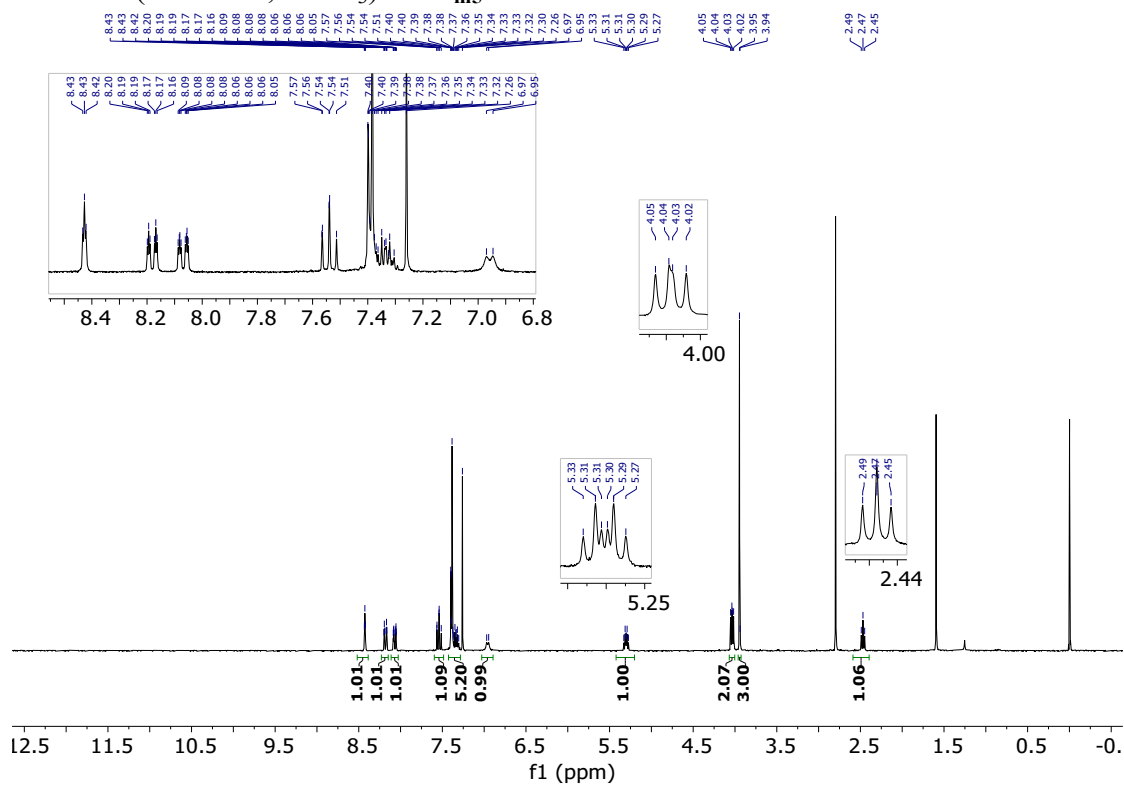


¹³C NMR (151 MHz, CD₃OD) of **3**_{t4}.

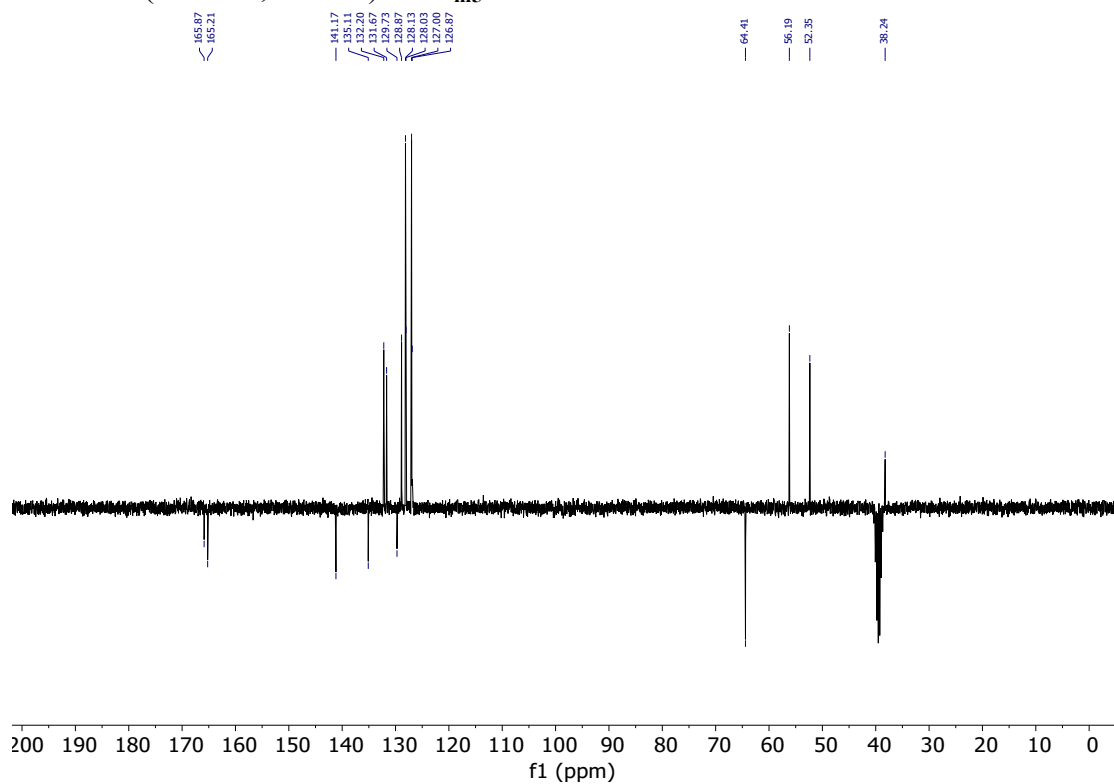


2.22. Synthesis of derivative **3b**

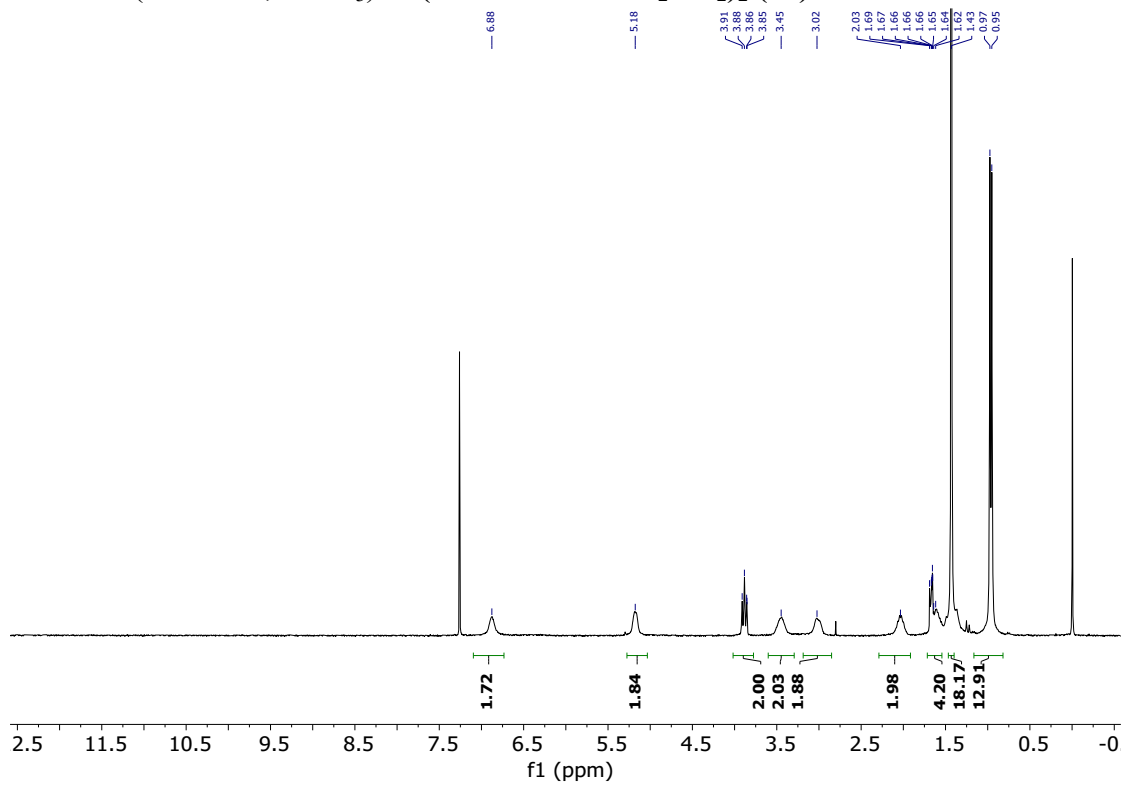
^1H NMR (300 MHz, CDCl_3) of **11m5**.



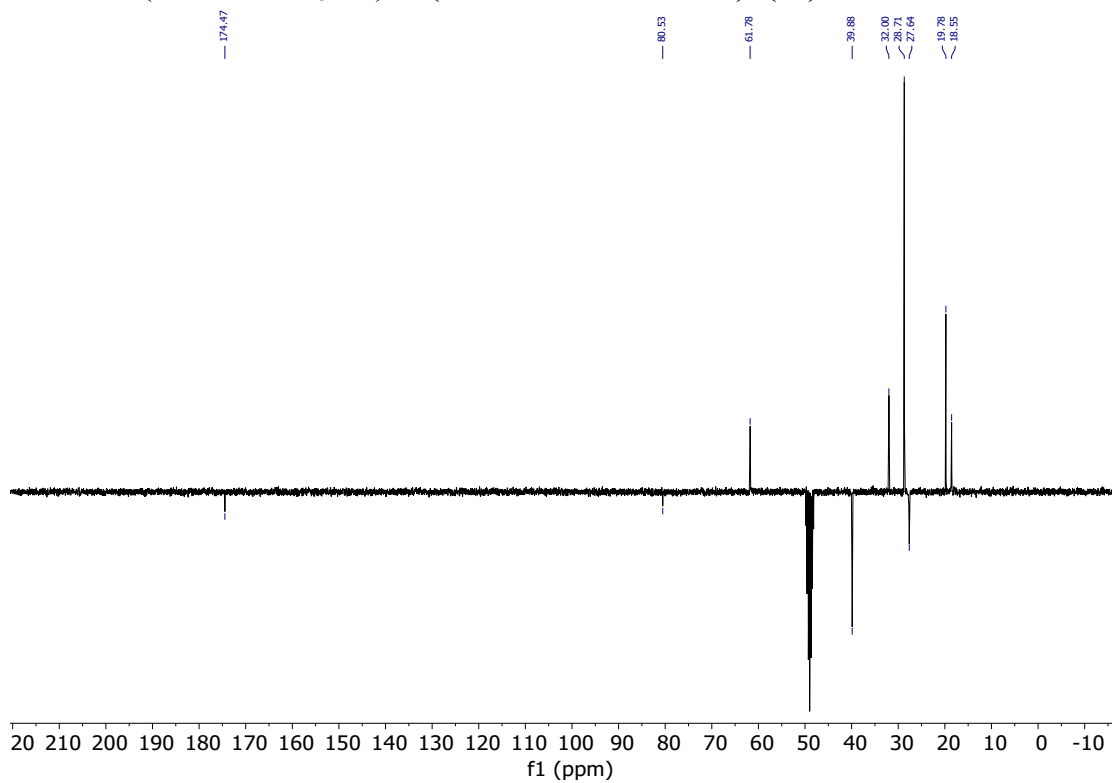
^{13}C NMR (75 MHz, DMSO) of **11m5**.



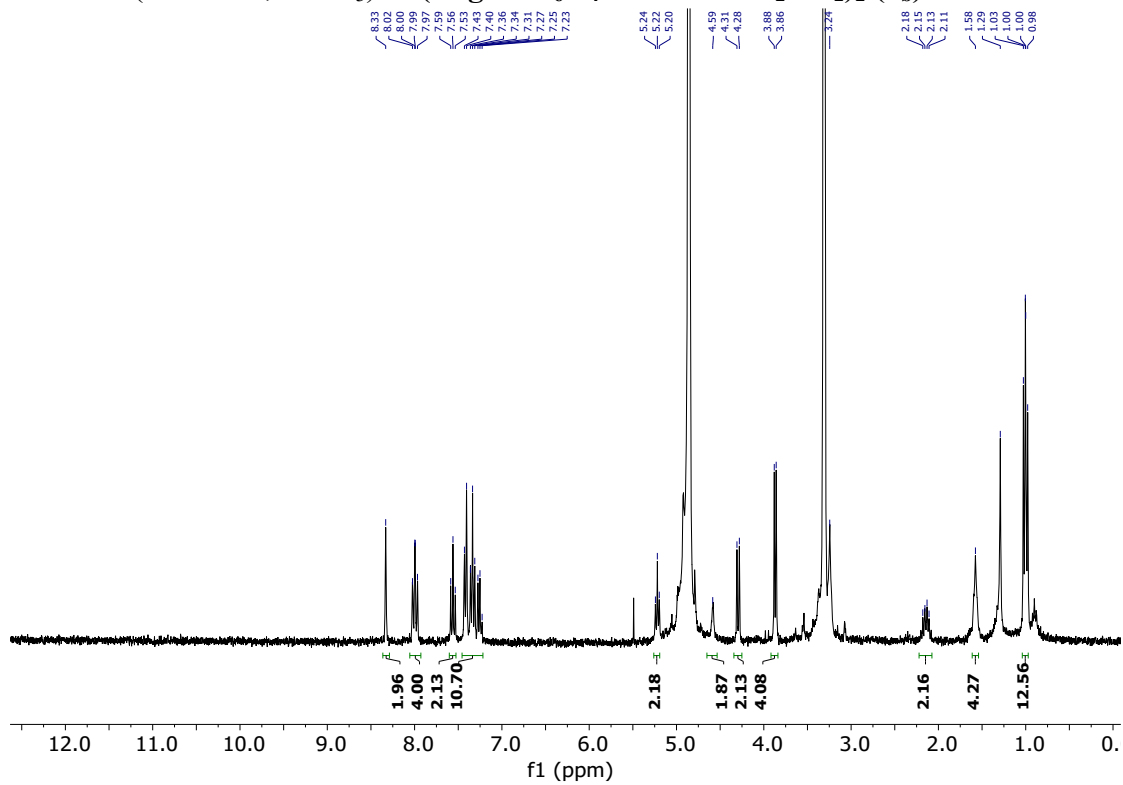
^1H NMR (300 MHz, CDCl_3) of **(Boc-Val-NH-CH₂CH₂)₂** (17).



^{13}C NMR (75 MHz, CD_3OD) of **(Boc-Val-NH-CH₂CH₂)₂** (17).



^1H NMR (300 MHz, CDCl_3) of **(Phg[#]-*m*C₆H₄-Val-NH-CH₂CH₂)₂ (3_b)**.



3. CD spectra.

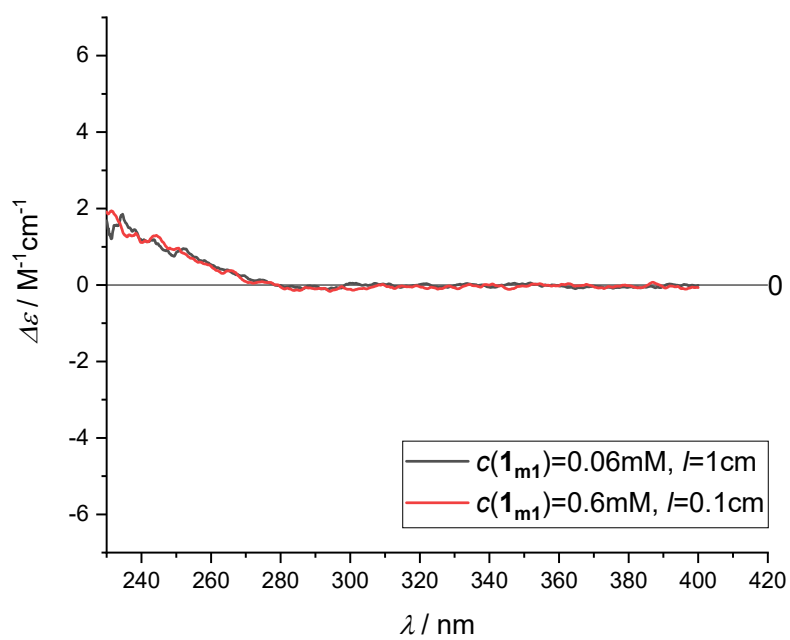


Figure S2. CD spectra (DCM) of $\mathbf{1}_{m1}$ solutions.

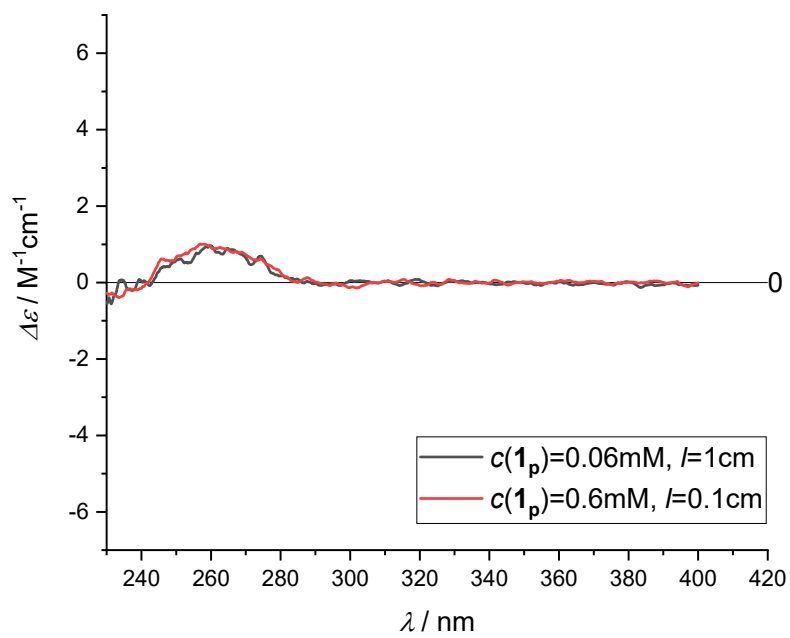


Figure S3. CD spectra (DCM) of $\mathbf{1}_p$ solutions.

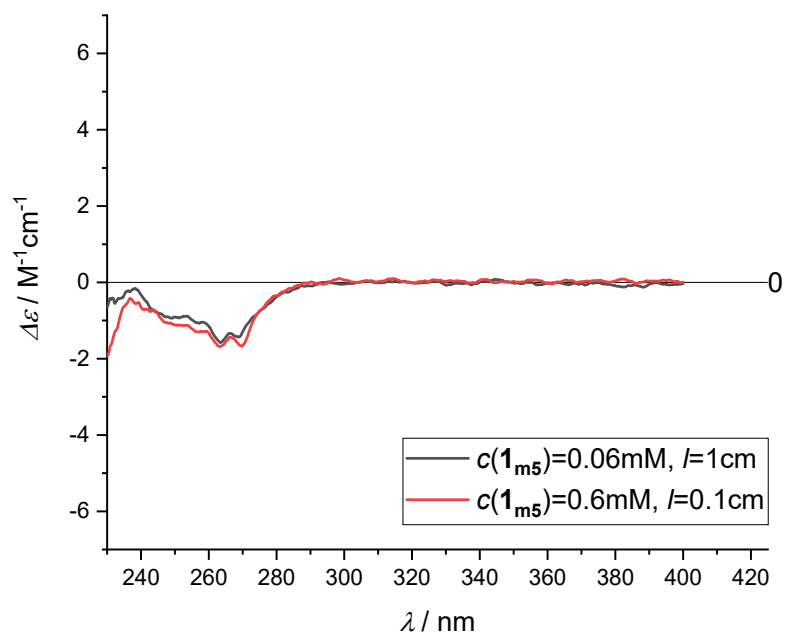


Figure S4. CD spectra (DCM) of 1_{m5} solutions.

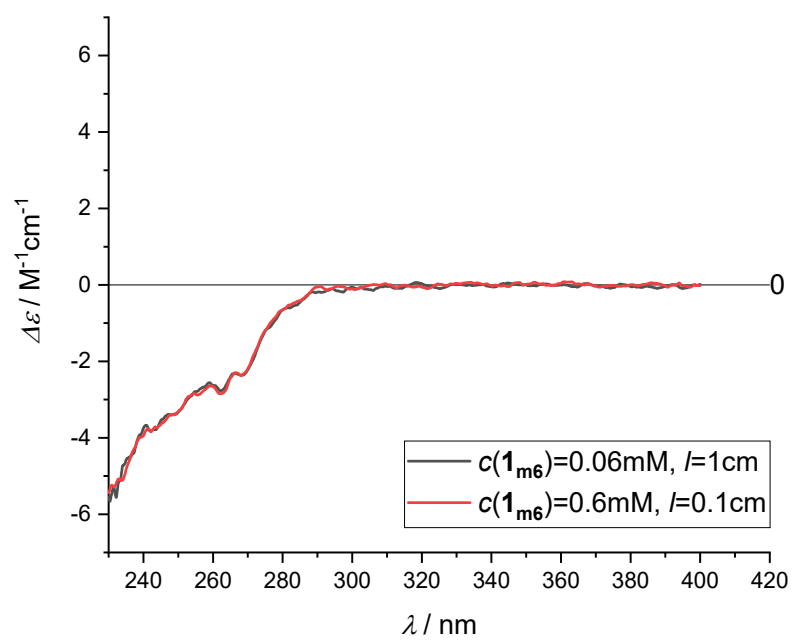


Figure S5. CD spectra (DCM) of 1_{m6} solutions.

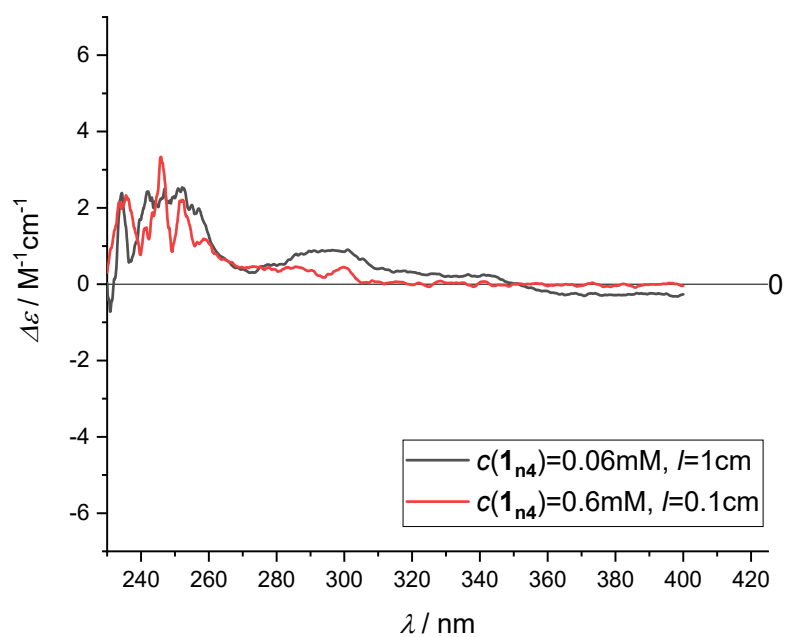


Figure S6. CD spectra (DCM) of $\mathbf{1}_{n4}$ solutions.

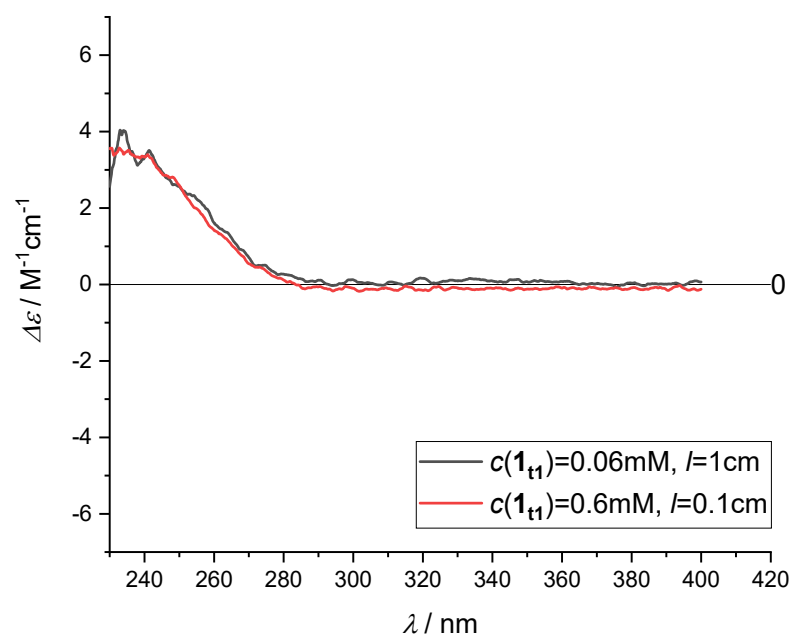


Figure S7. CD spectra (DCM) of $\mathbf{1}_{t1}$ solutions.

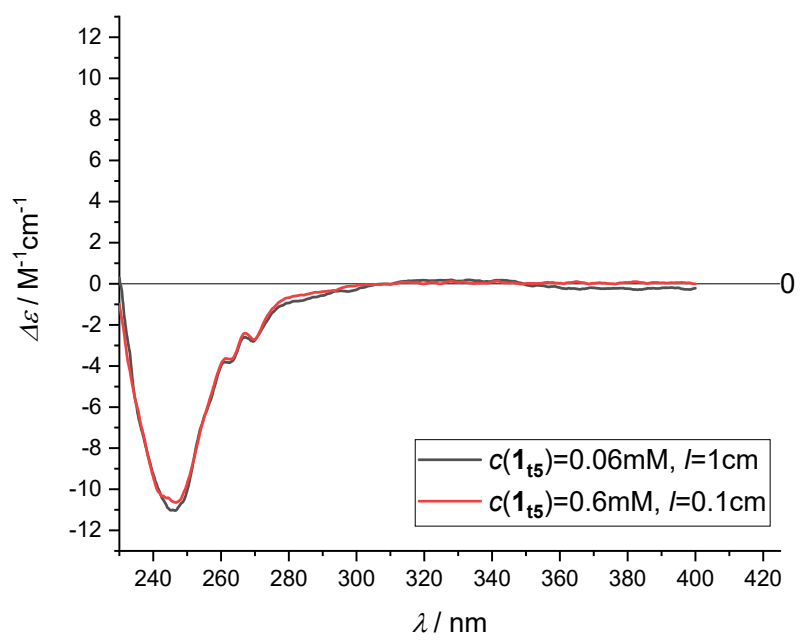
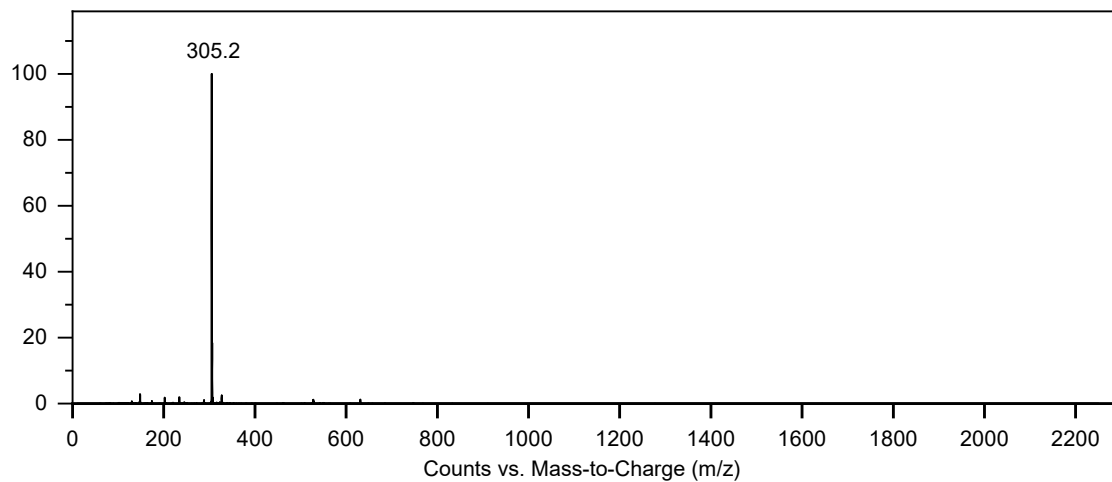


Figure S8. CD spectra (DCM) of 1_{t5} solutions.

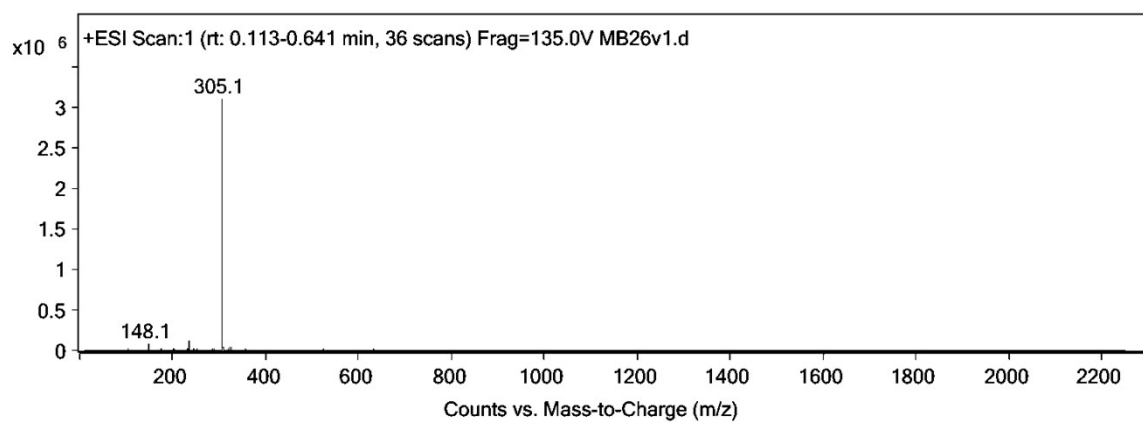
4. Mass spectra.

4.1. Oxazolines

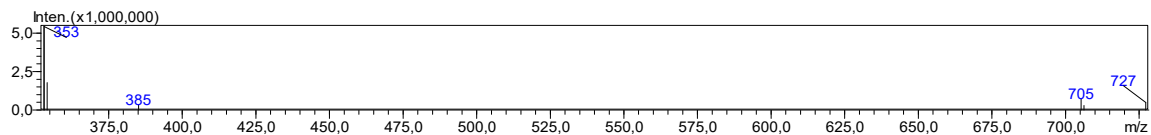
ESI-MS of **1_{m1}**.



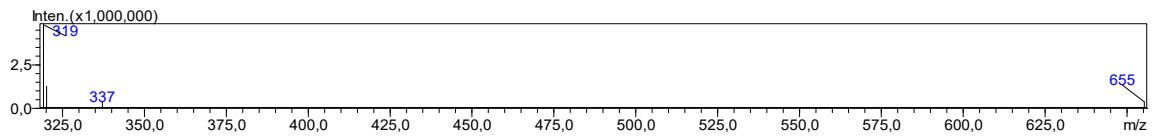
ESI-MS of **1_p**.



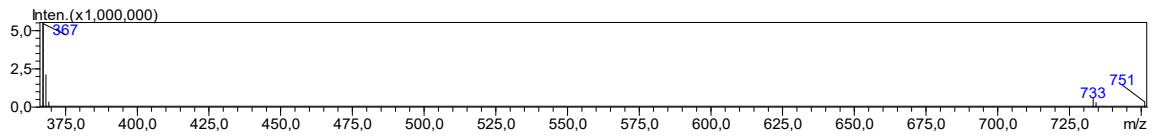
ESI-MS of **1_{m2}**.



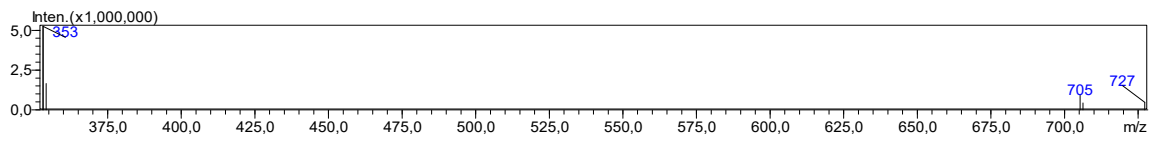
ESI-MS of **1_{m3}**.



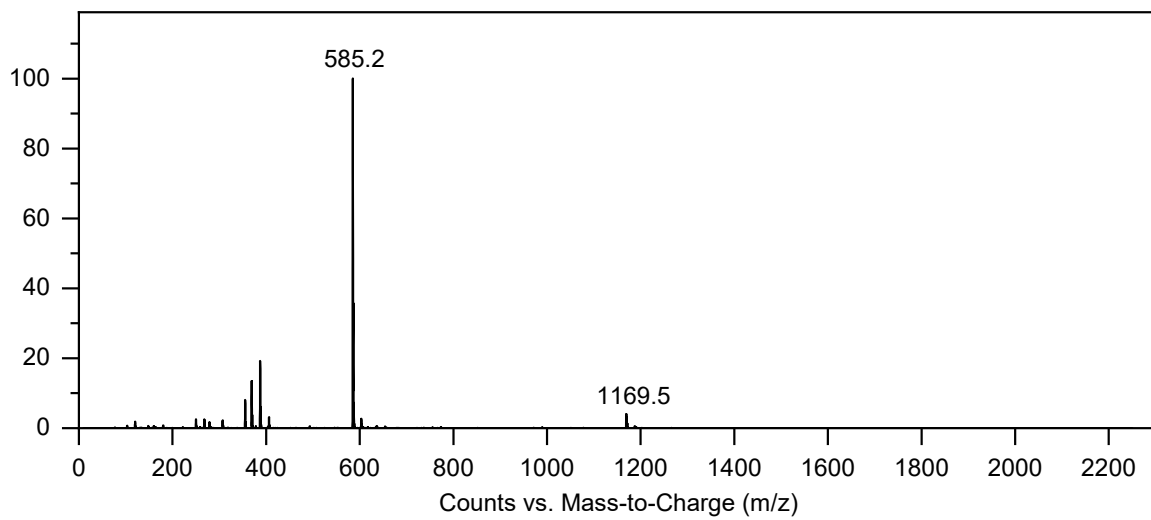
ESI-MS of **1_{m4}**.



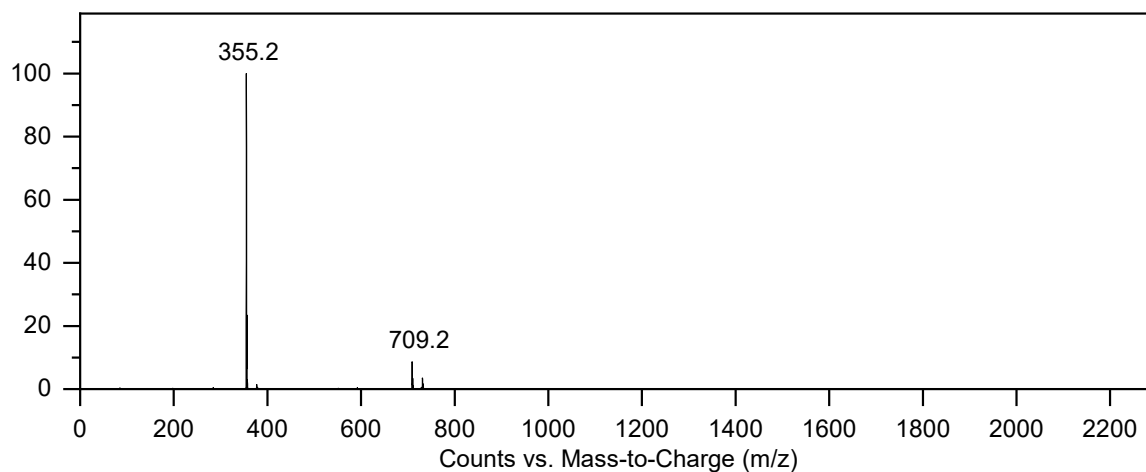
ESI-MS of **1_{m5}**.



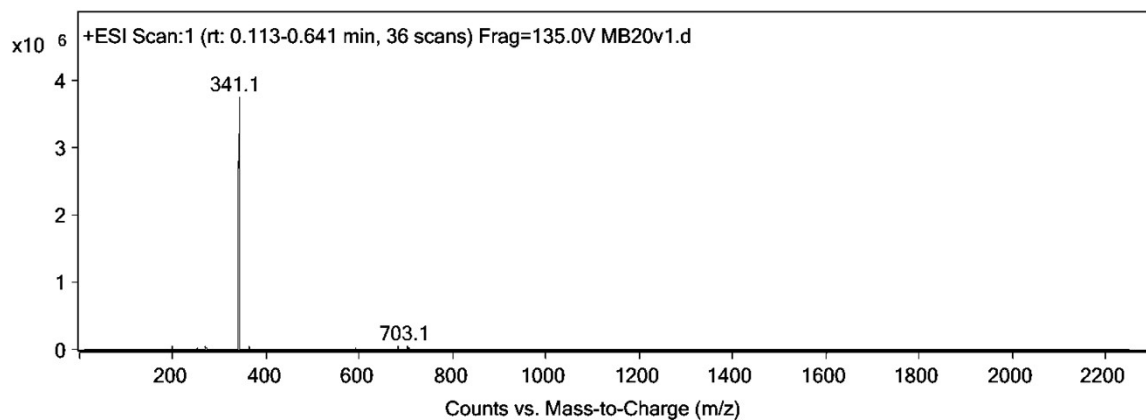
ESI-MS of **1_{m6}**.



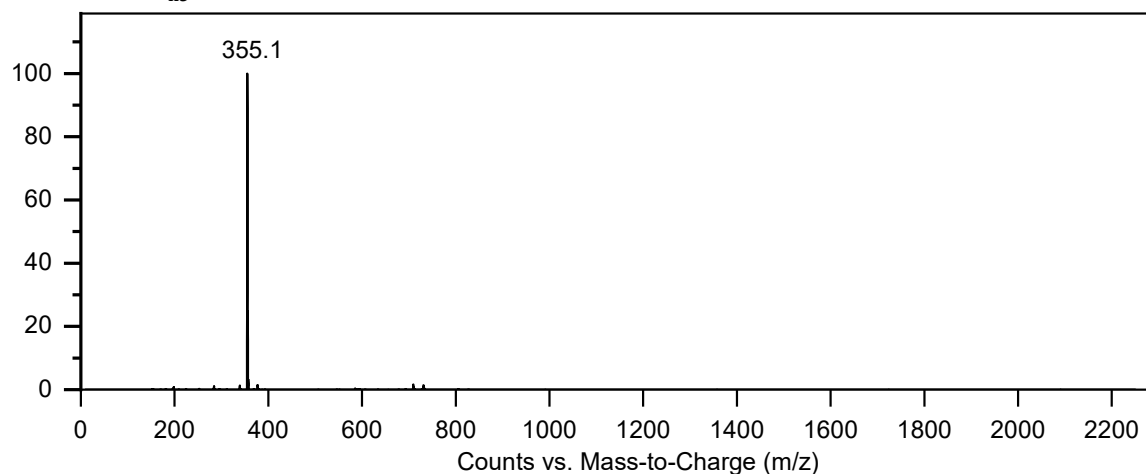
ESI-MS of **1_{n1}**.



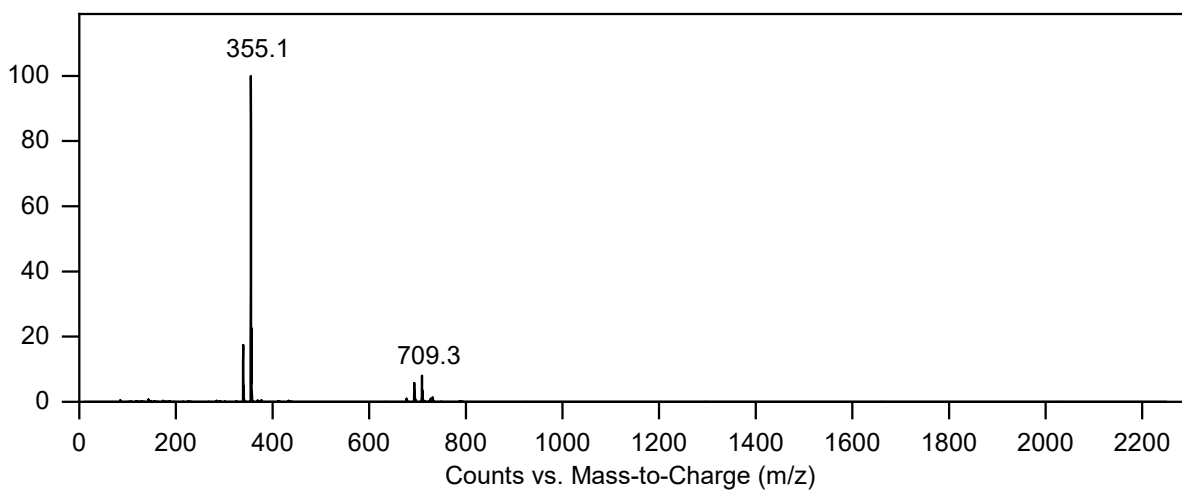
ESI-MS of **1_{n2}**.



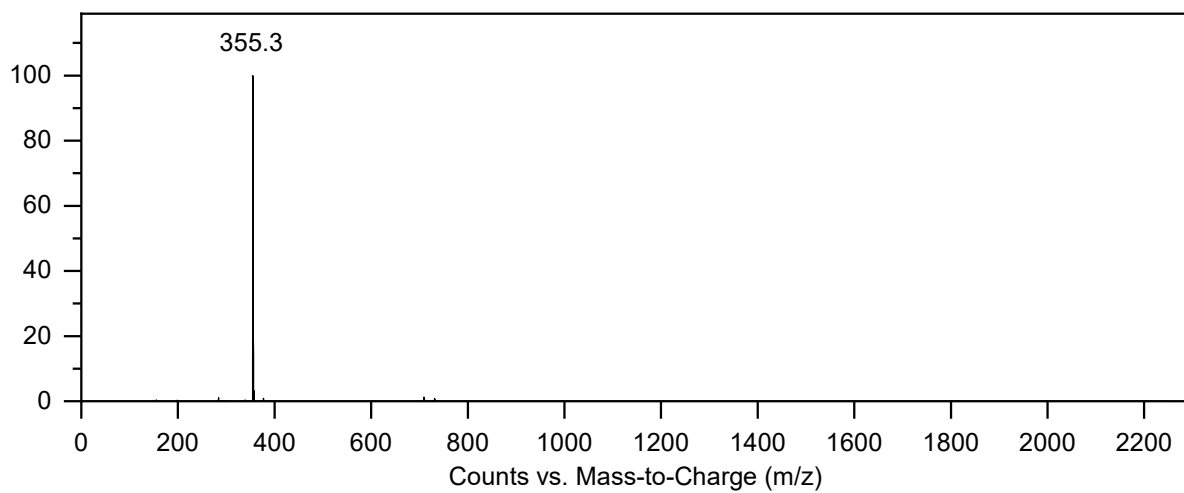
ESI-MS of **1_{n3}**.



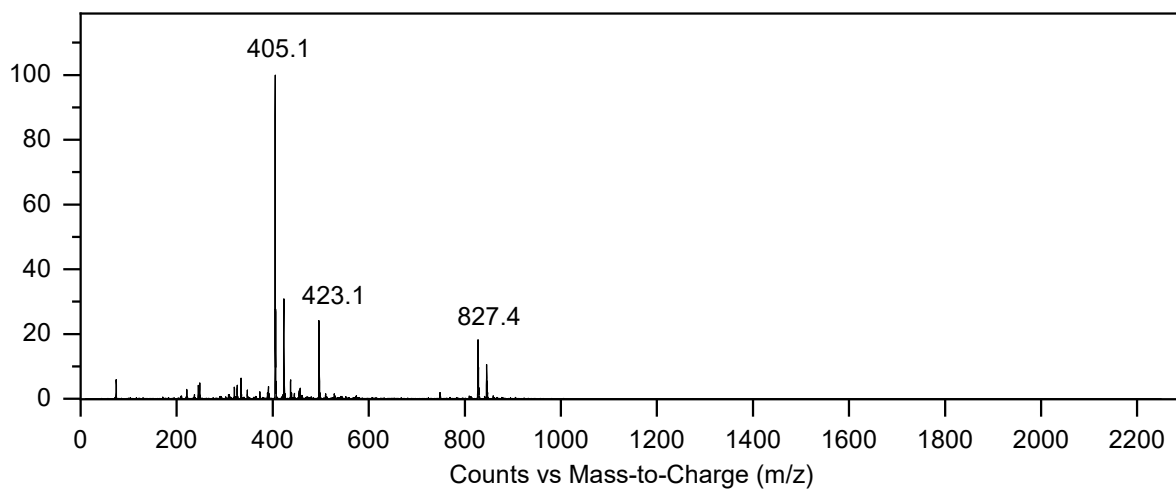
ESI-MS of **1_{n4}**.



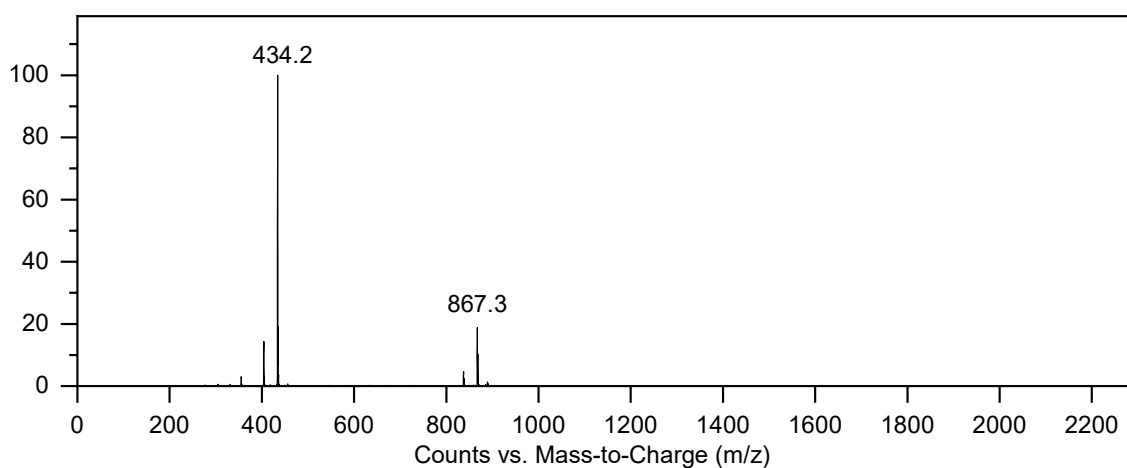
ESI-MS of **1_{n5}**.



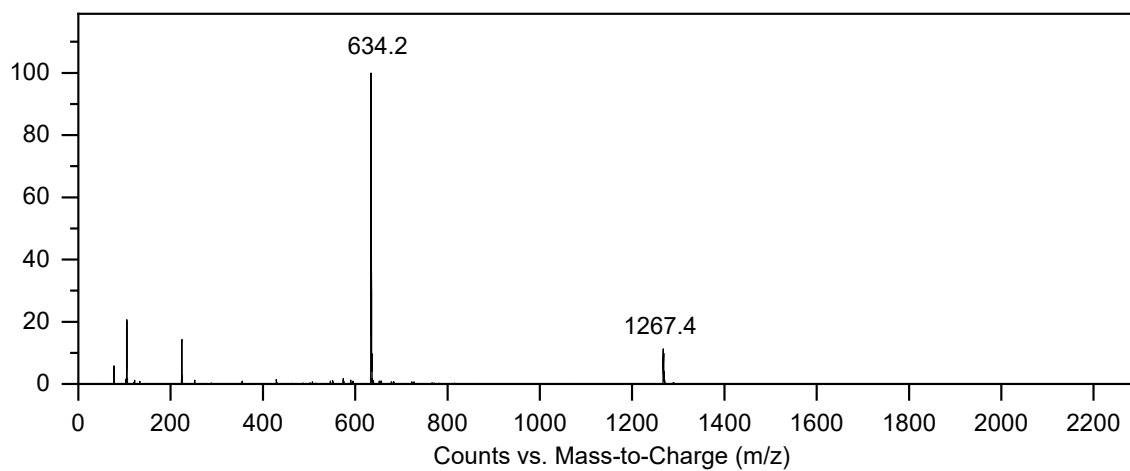
ESI-MS of **1_a**.



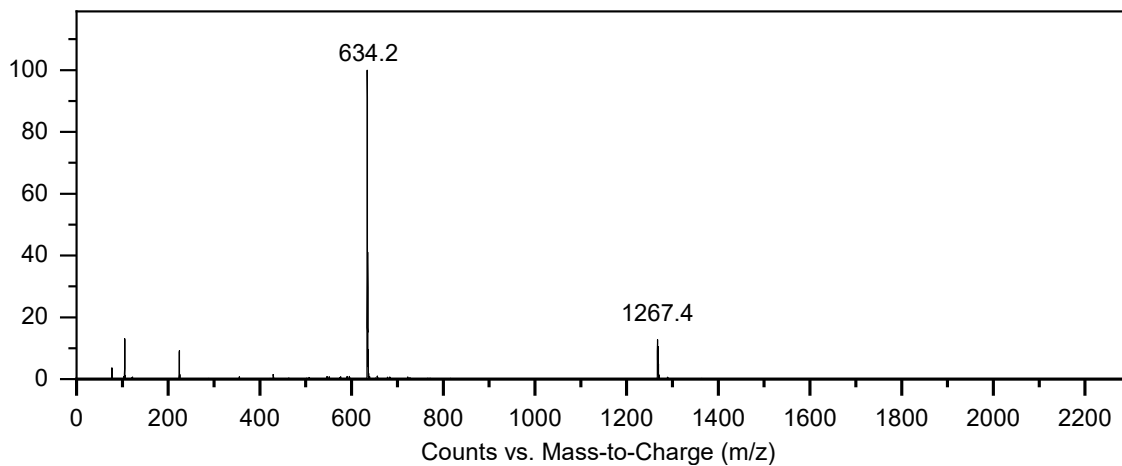
ESI-MS of **1_{t1}**.



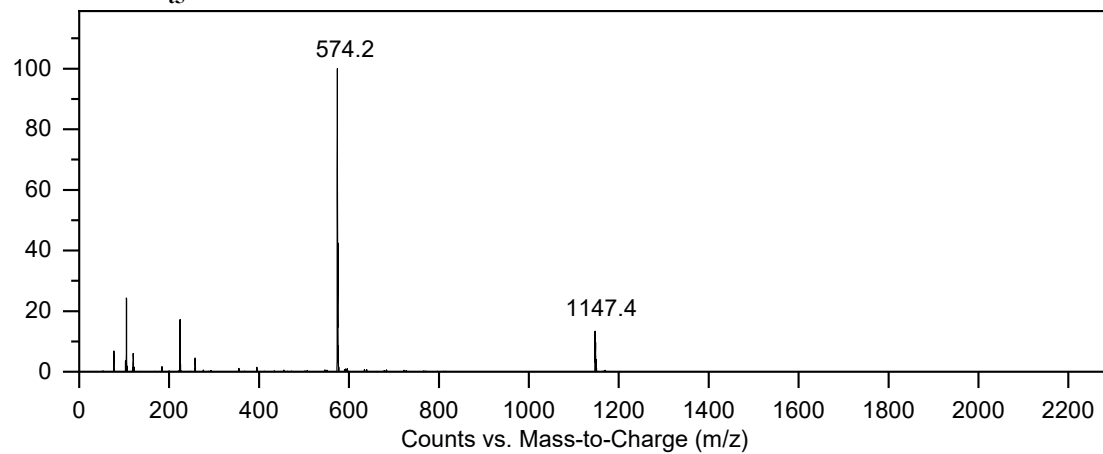
ESI-MS of **1_{t3}**.



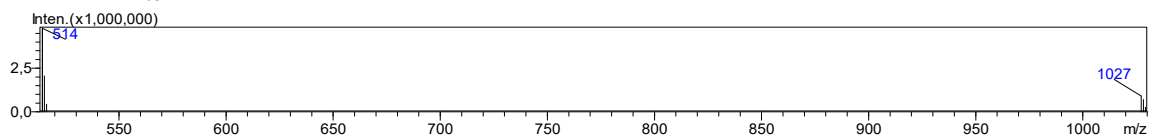
ESI-MS of **1_{t4}**.



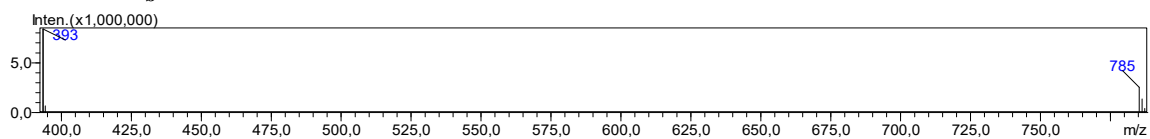
ESI-MS of **1_{t5}**.



ESI-MS of **1_{t6}**.

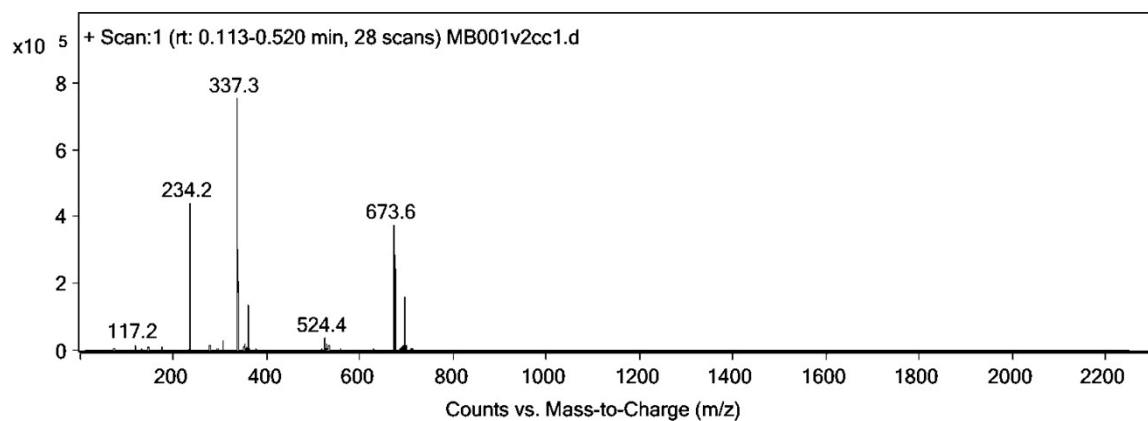


ESI-MS of **1_b**.

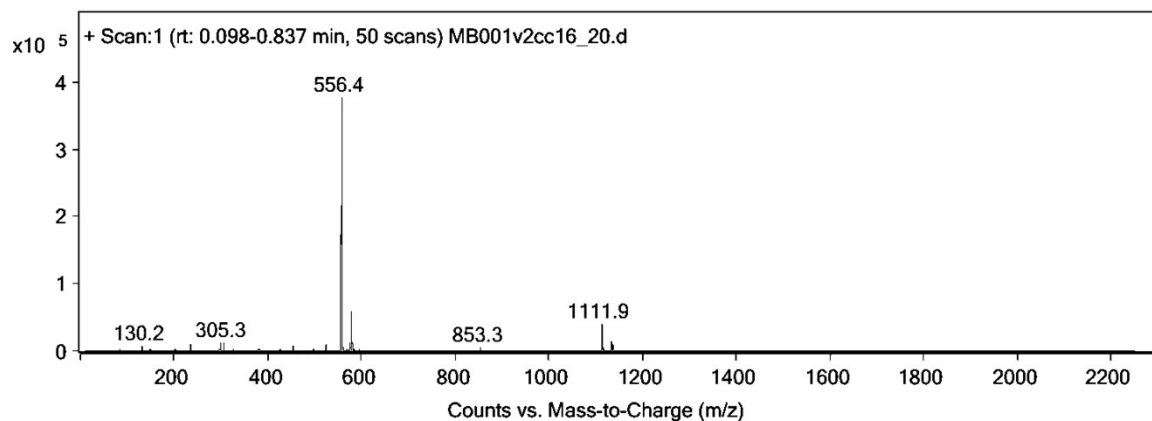


4.2. Reaction 1

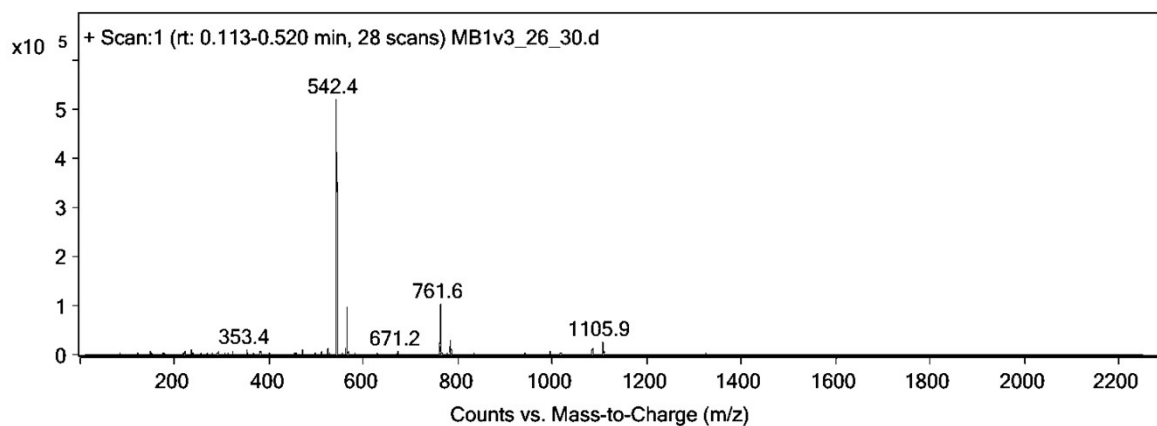
ESI-MS of **2**_{m1}.



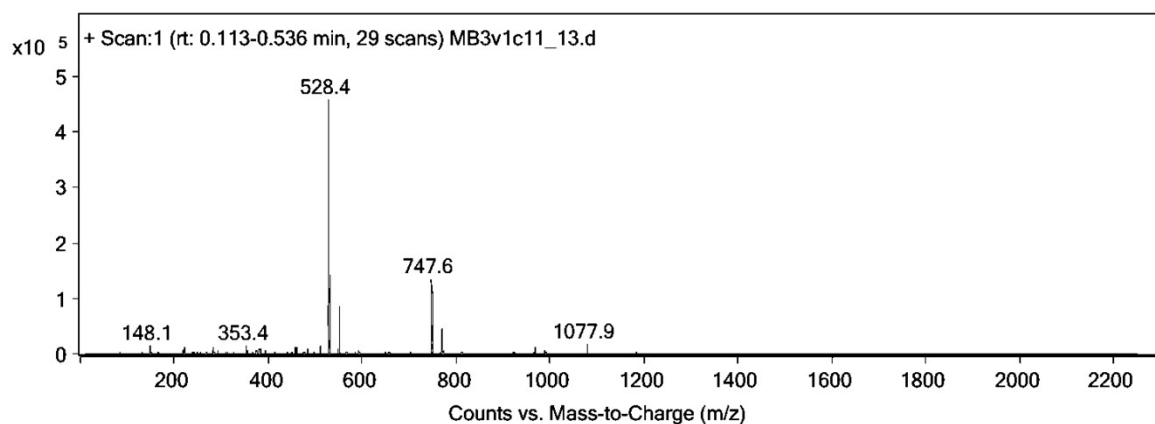
ESI-MS of **6**_{m1}.



ESI-MS of **7**_{m1}.

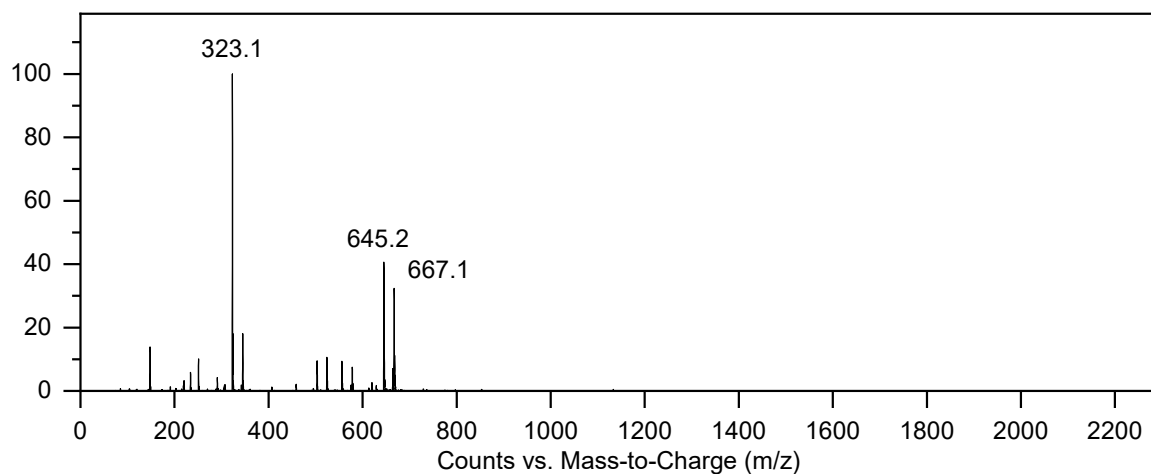


ESI-MS of **8**_{m1}.



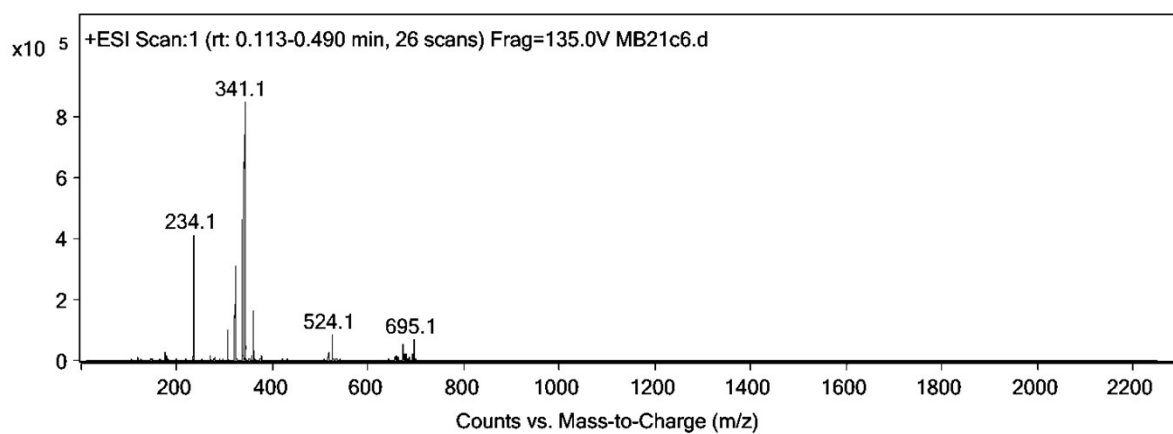
4.3. Reaction 5

ESI-MS of **3**_{m1}.



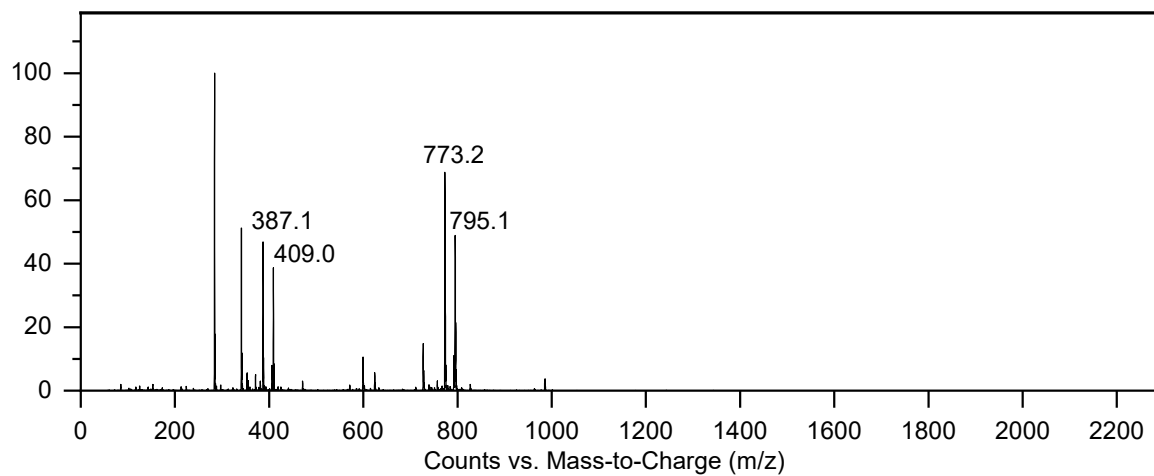
4.4. Reaction 7

ESI-MS of **2**_p.

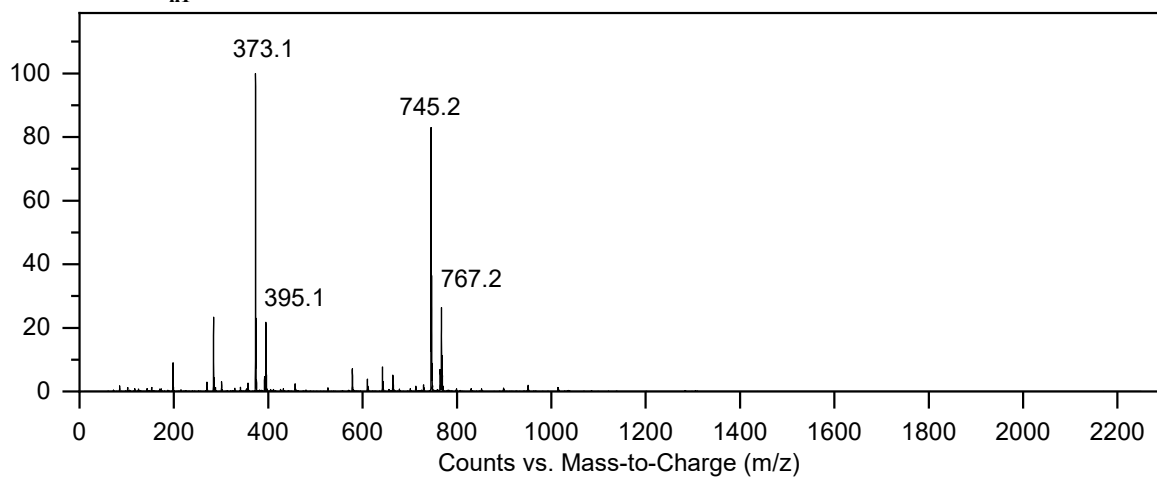


4.5. Reaction 13

ESI-MS of **2_{n1}**.

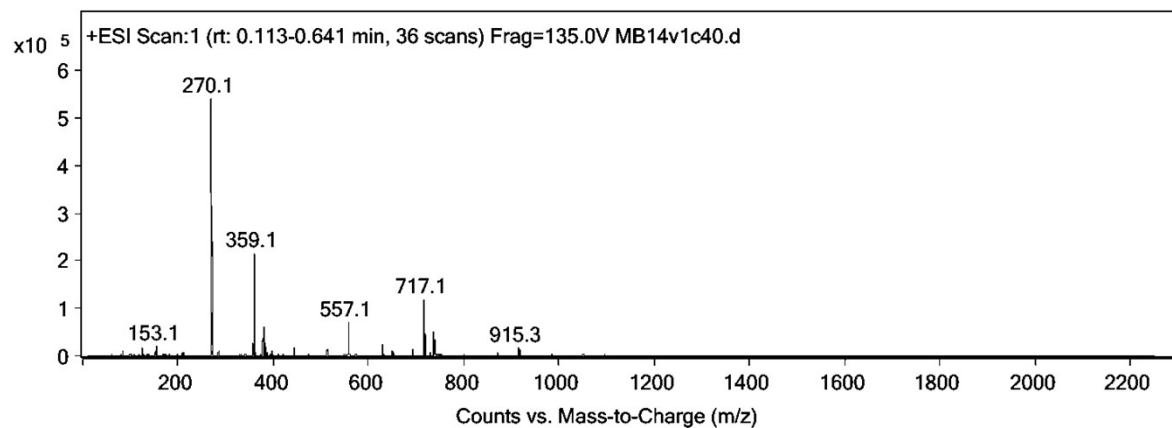


ESI-MS of **3_{n1}**.

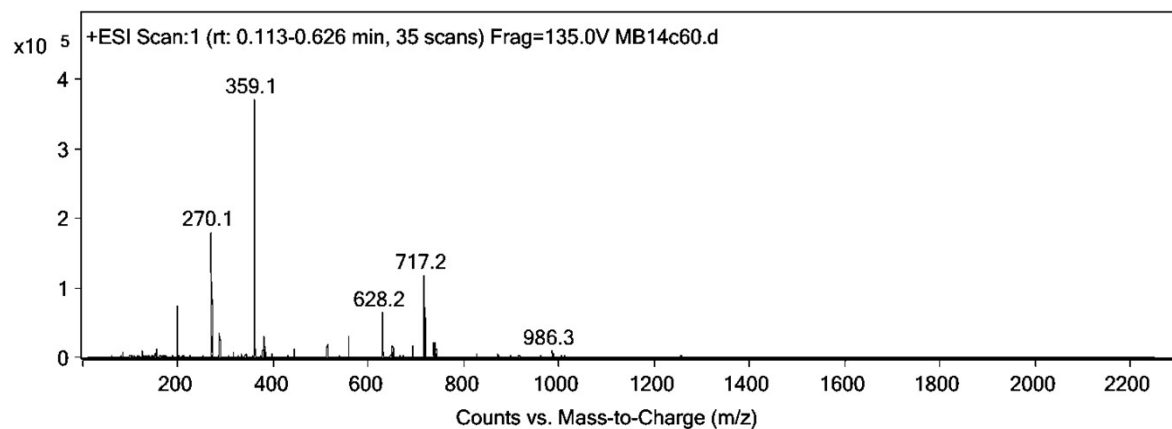


4.6. Reaction 14

ESI-MS of **2**_{n2}.

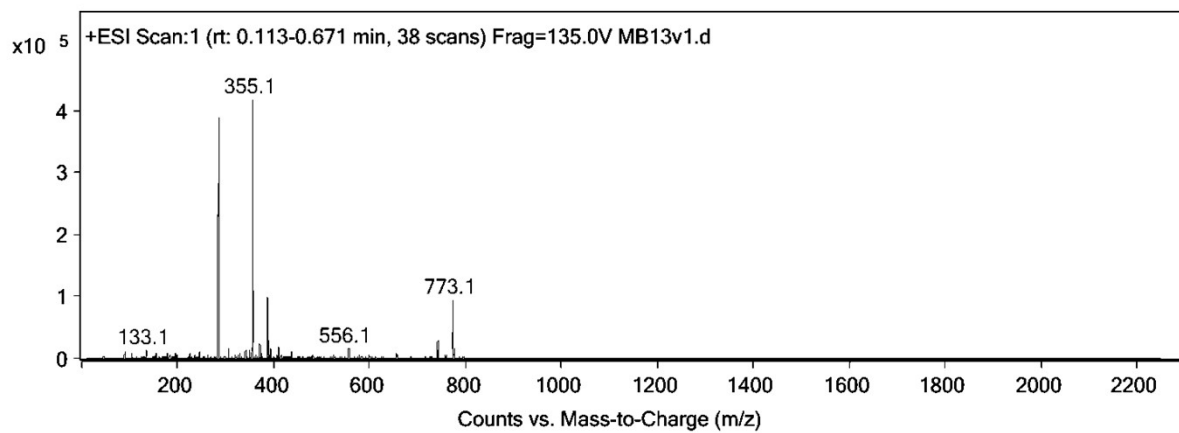


ESI-MS of **3**_{n2}.

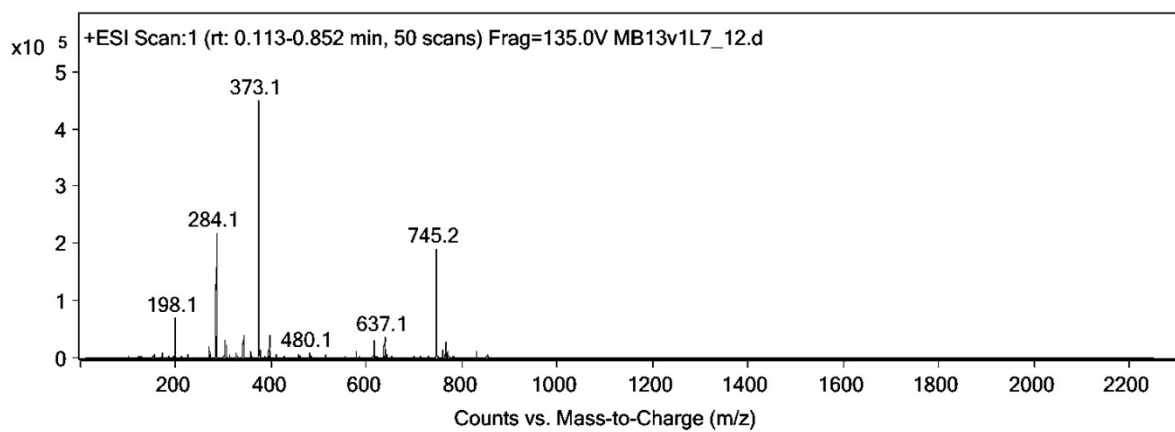


4.7. Reaction 15

ESI-MS of **2**_{n3}.

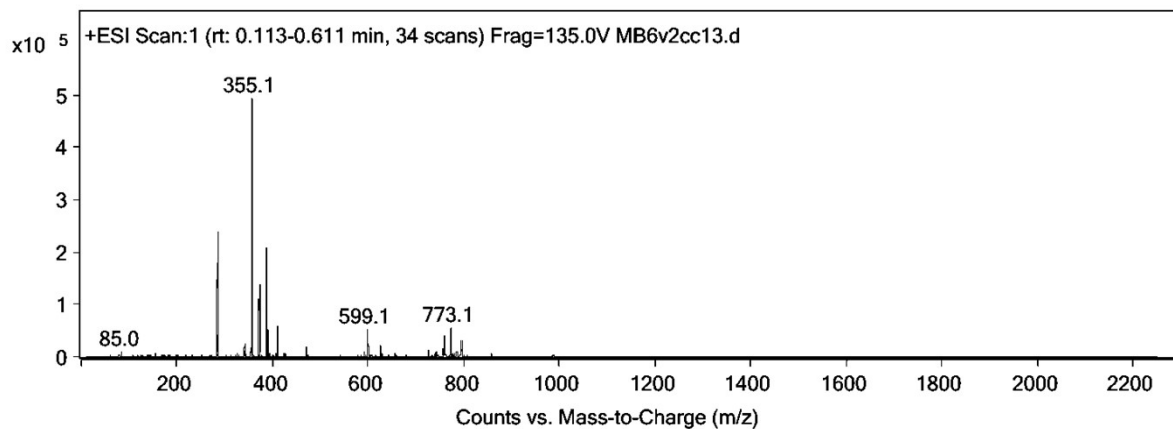


ESI-MS of **3**_{n3}.

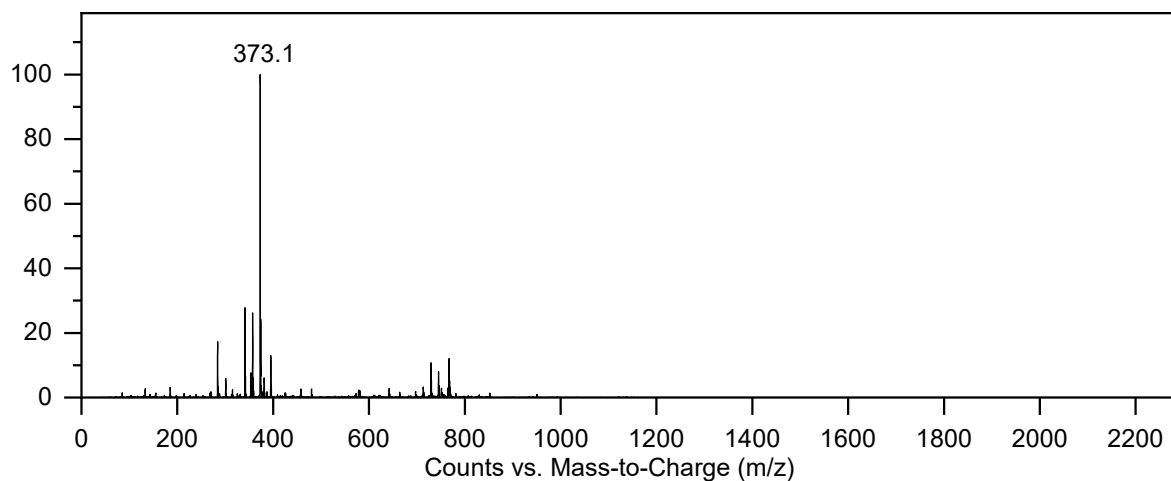


4.8. Reaction 16

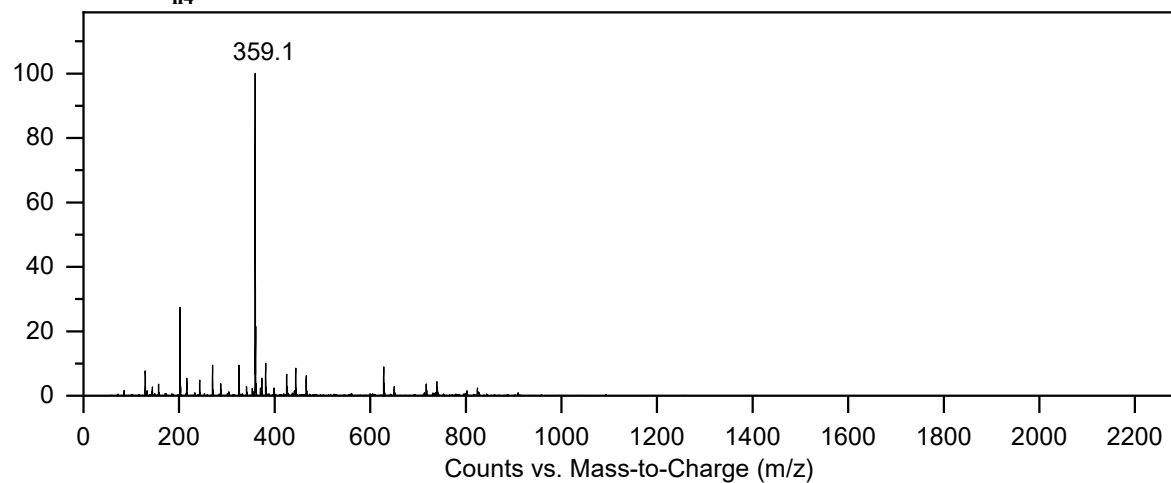
ESI-MS of **2**_{n4}.



ESI-MS of **3**_{n4}.

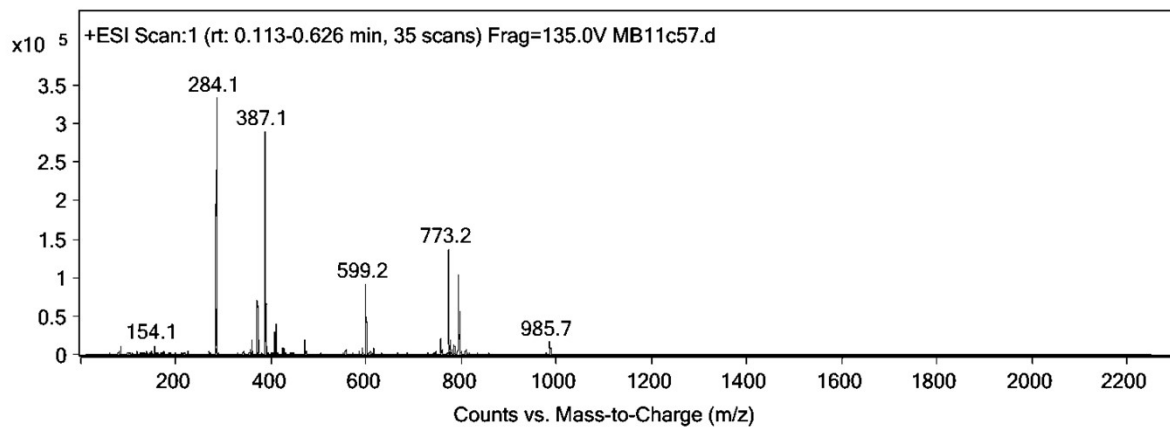


ESI-MS of **4**_{n4}.

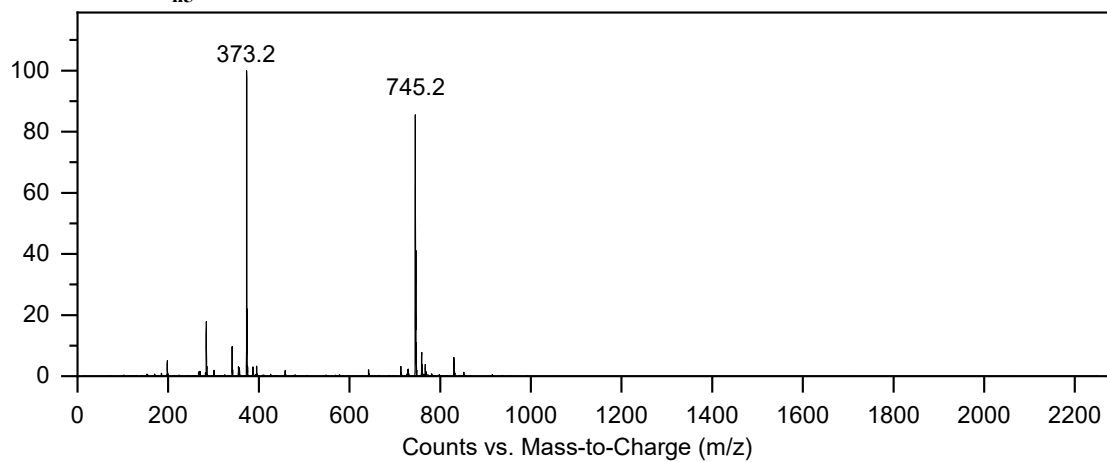


4.9. Reaction 17

ESI-MS of **2**_{n5}.

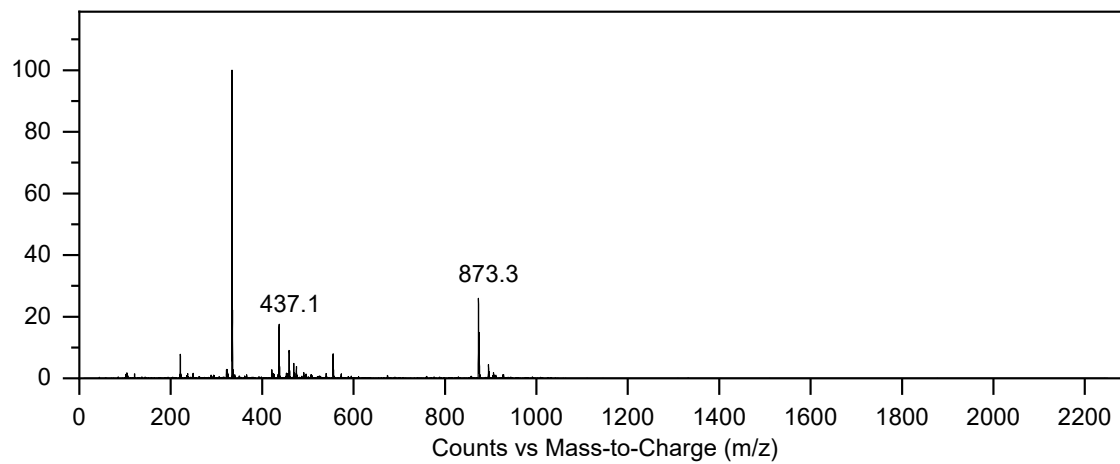


ESI-MS of **3**_{n5}.

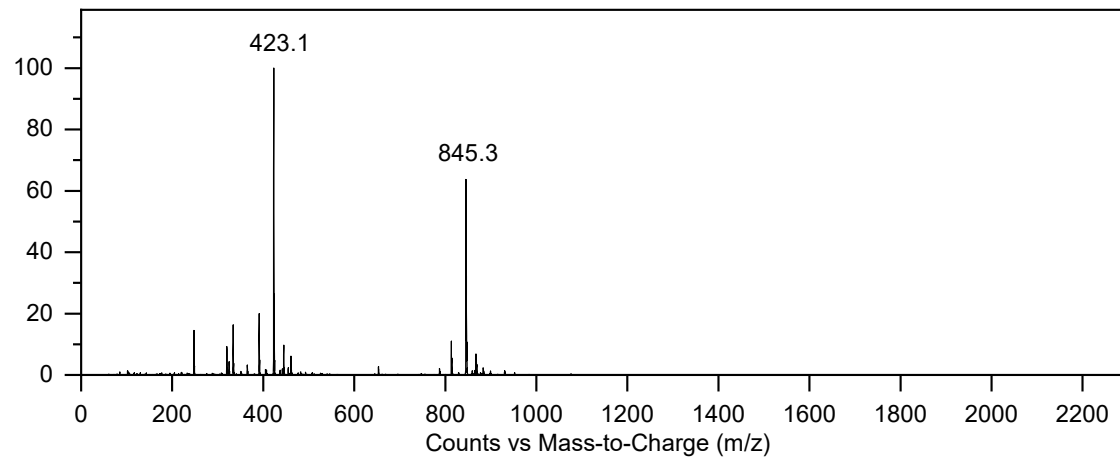


4.10. Reaction 18

ESI-MS of **2_{a1}**.



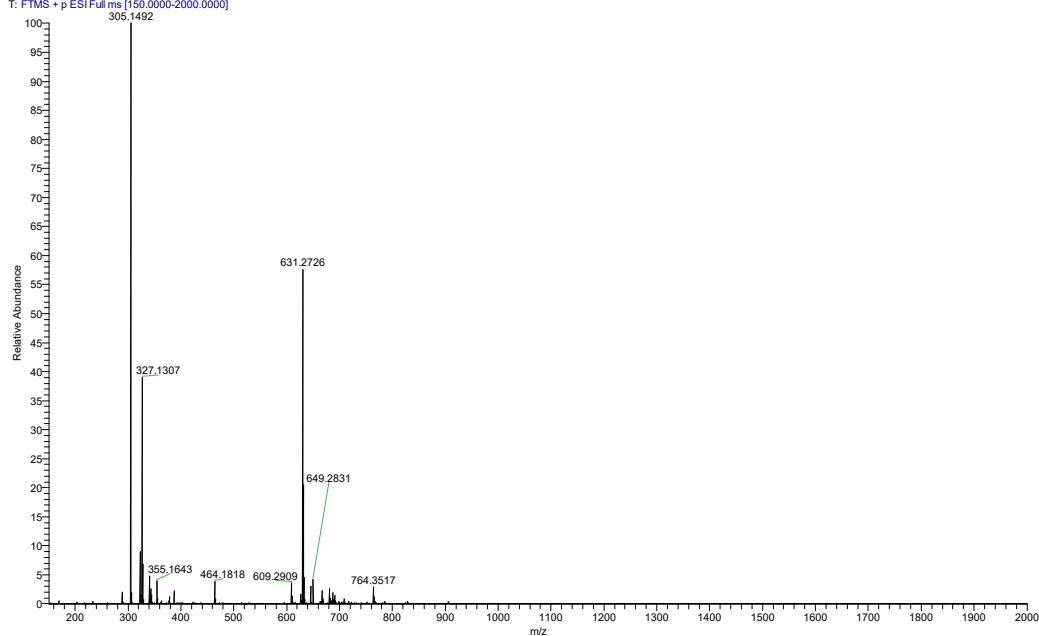
ESI-MS of **3_{a1}**.



5. HRMS

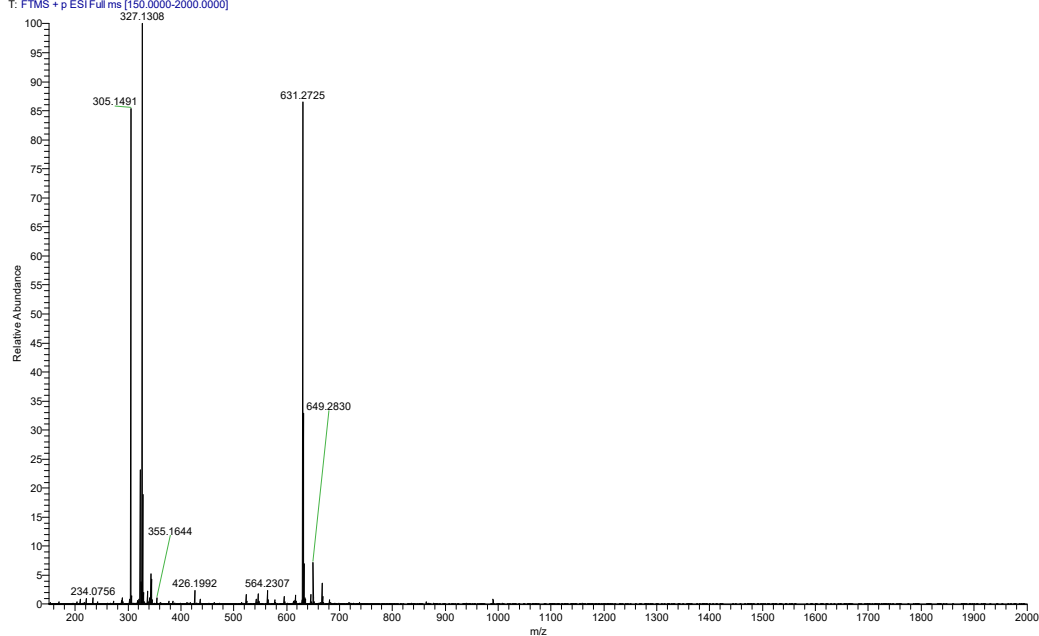
ESI-HRMS of **1_m**.

MB25v3 #1-48 RT: 0.00-0.25 AV: 48 NL: 8.08E7
T: FTMS + p ESI Full ms [150.0000-2000.0000]

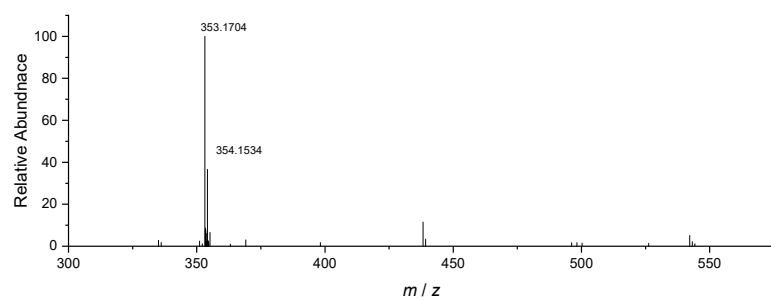


ESI-HRMS of **1_p**.

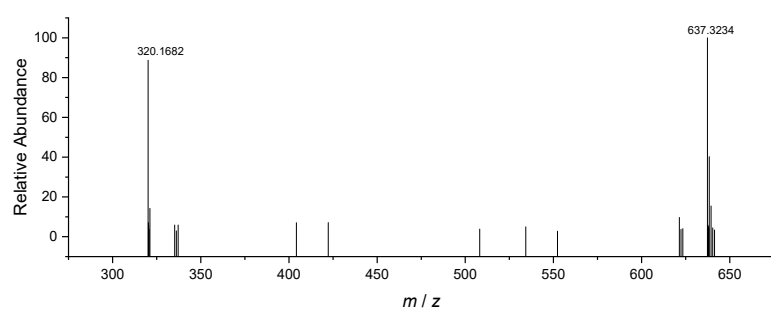
MB26v1 #1-44 RT: 0.00-0.25 AV: 44 NL: 8.62E6
T: FTMS + p ESI Full ms [150.0000-2000.0000]



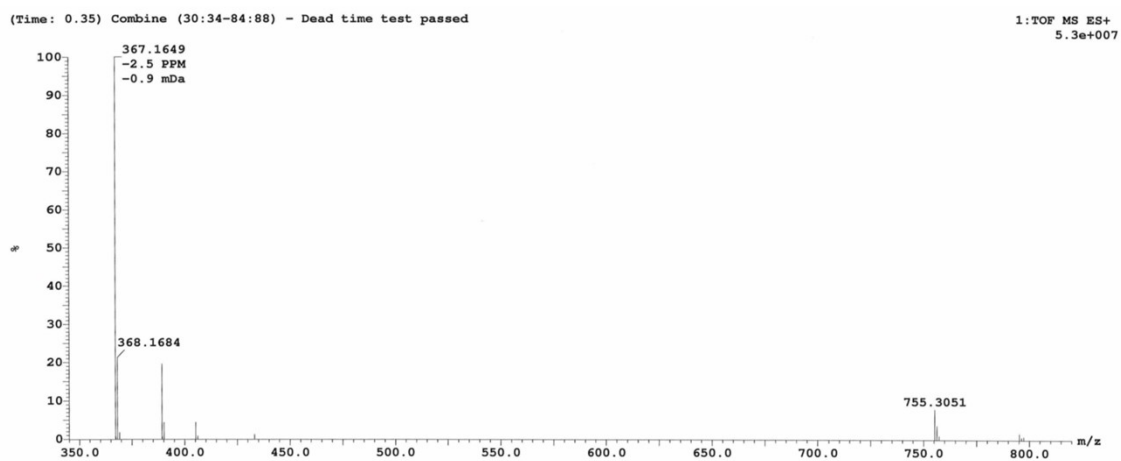
MALDI-HRMS of **1_{m2}**.



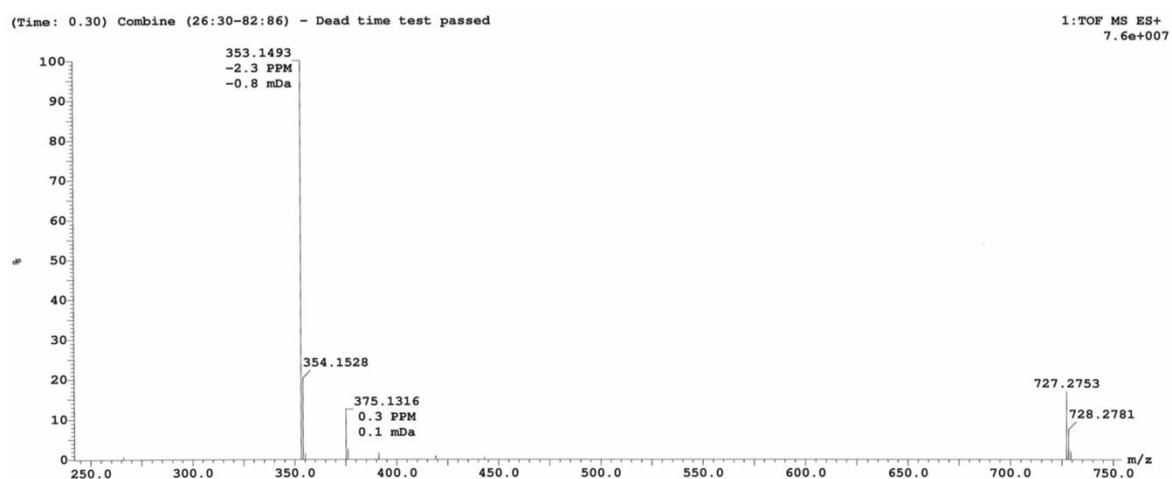
MALDI-HRMS of **1_{m3}**.



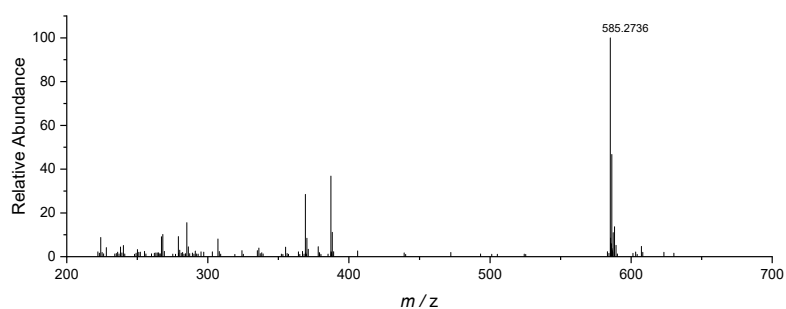
ESI-TOF-HRMS of **1_{m4}**.



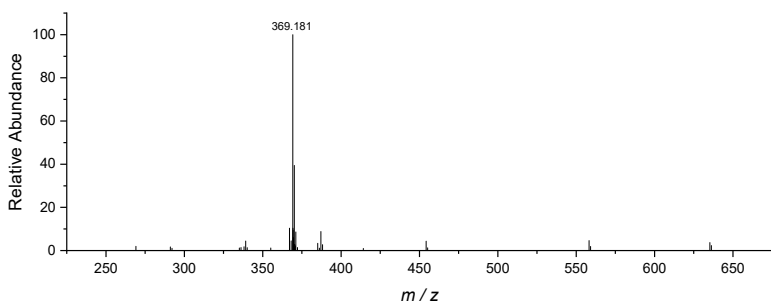
ESI-TOF-HRMS of **1_{m5}**.



MALDI-HRMS of **1_{m6}**.

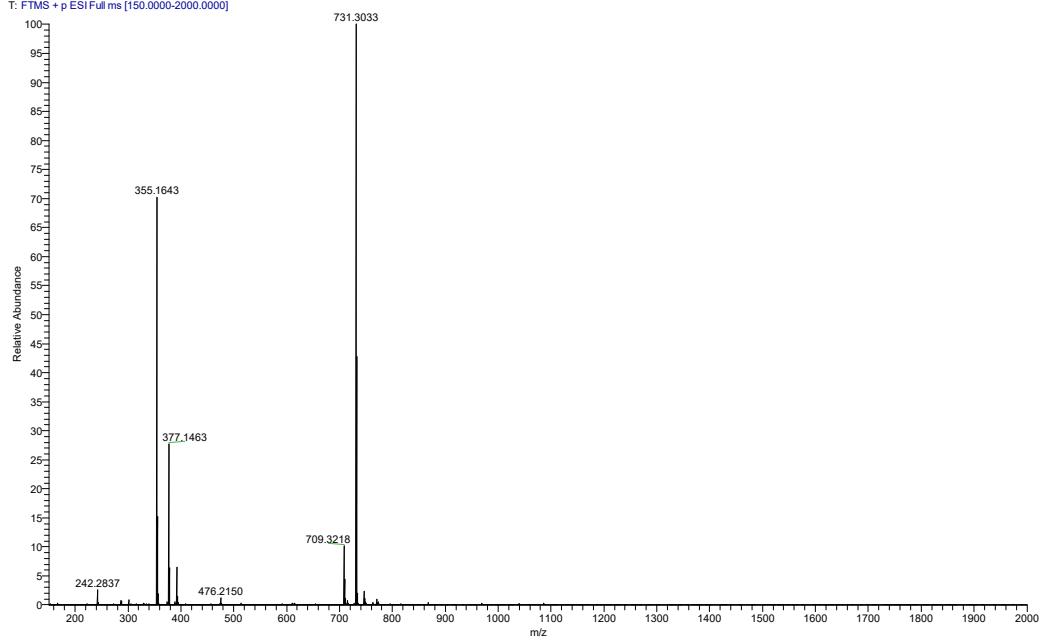


MALDI-HRMS of **1_{m7}**.



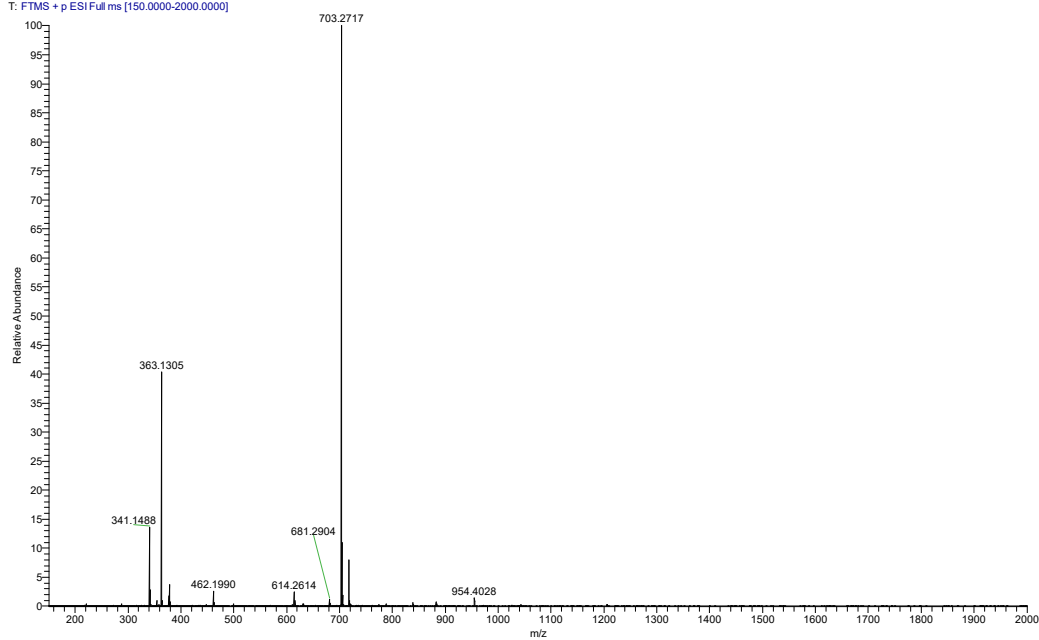
ESI-HRMS of **1_{n1}**.

MB9v1 #1-44 RT: 0.00-0.25 AV: 44 NL: 1.44E7
T: FTMS + p ESI Full ms [150.0000-2000.0000]



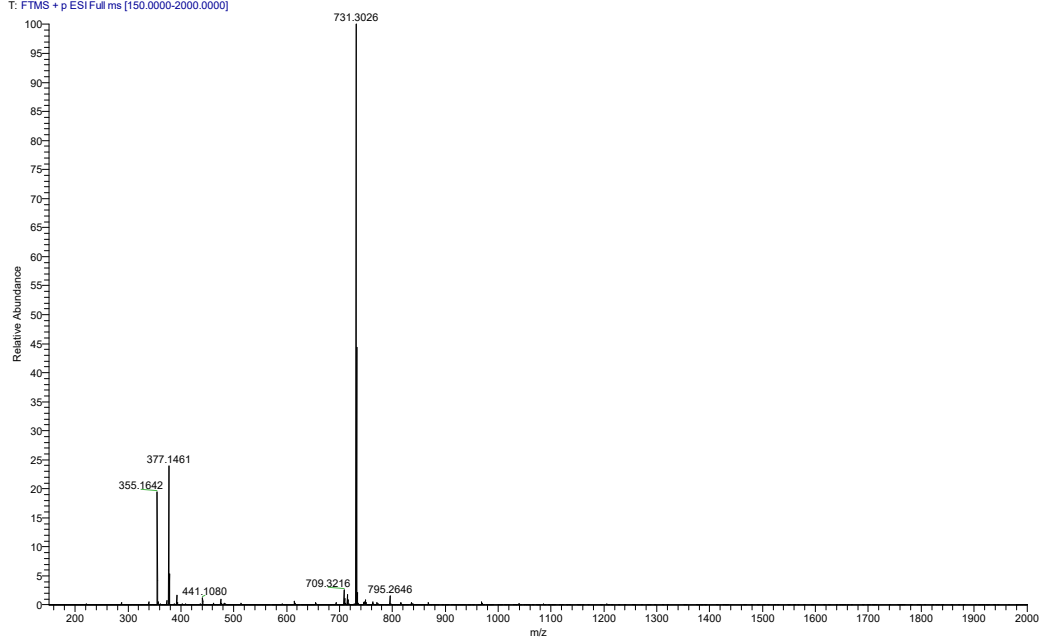
ESI-HRMS of **1_{n2}**.

MB20c44 #1-44 RT: 0.00-0.25 AV: 44 NL: 1.45E7
T: FTMS + p ESI Full ms [150.0000-2000.0000]



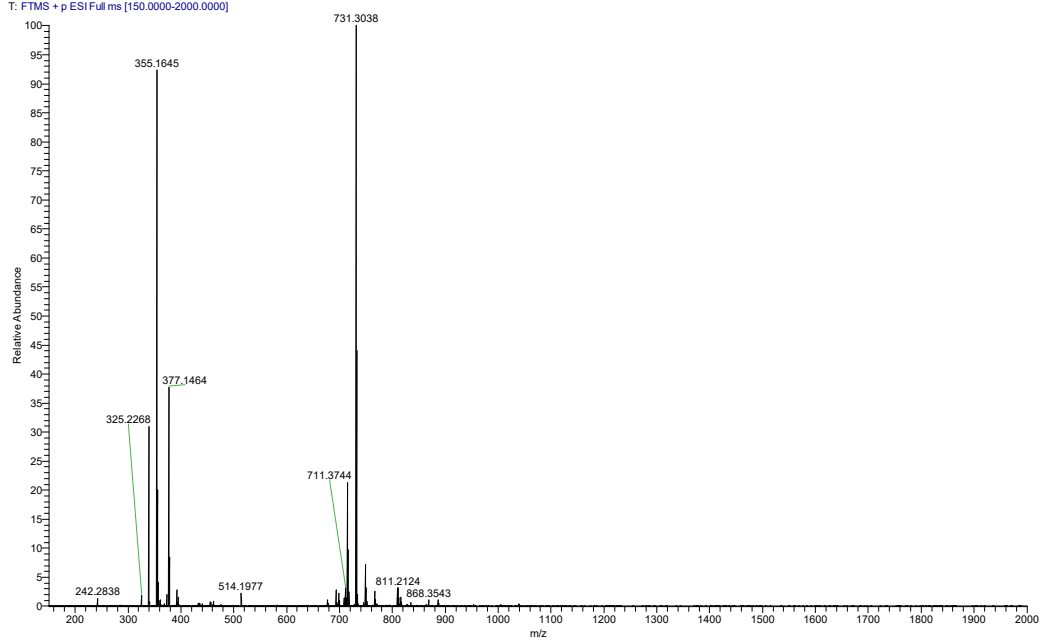
ESI-HRMS of 1_{n3} .

MB19v1 #1-45 RT: 0.00-0.25 AV: 45 NL: 2.58E7
T: FTMS + p ESI Full ms [150.0000-2000.0000]



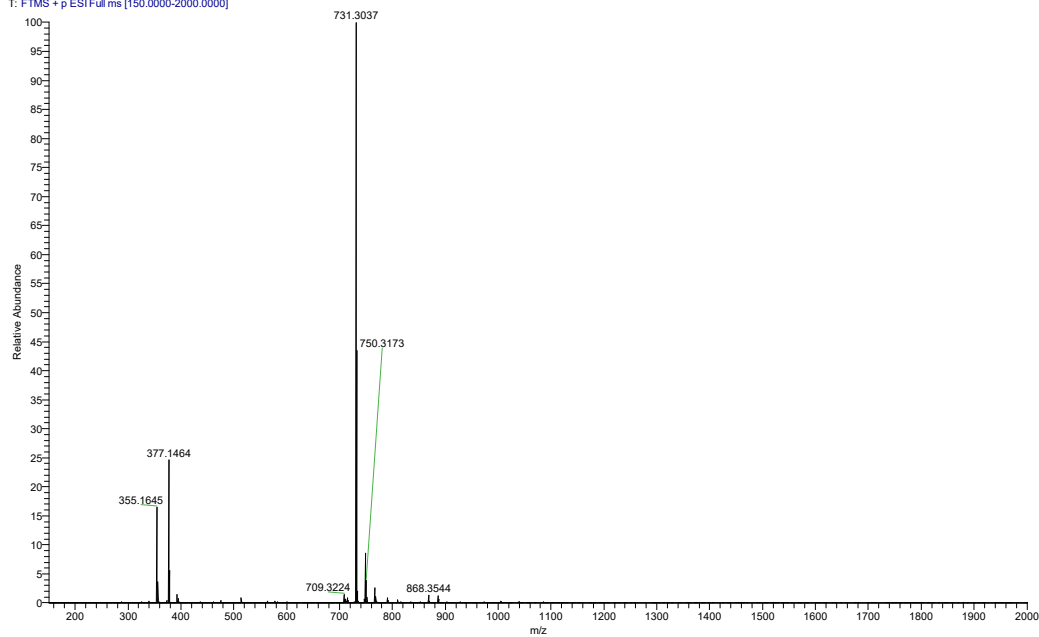
ESI-HRMS of 1_{n4} .

MB10v1 #1-43 RT: 0.00-0.25 AV: 43 NL: 7.11E6
T: FTMS + p ESI Full ms [150.0000-2000.0000]



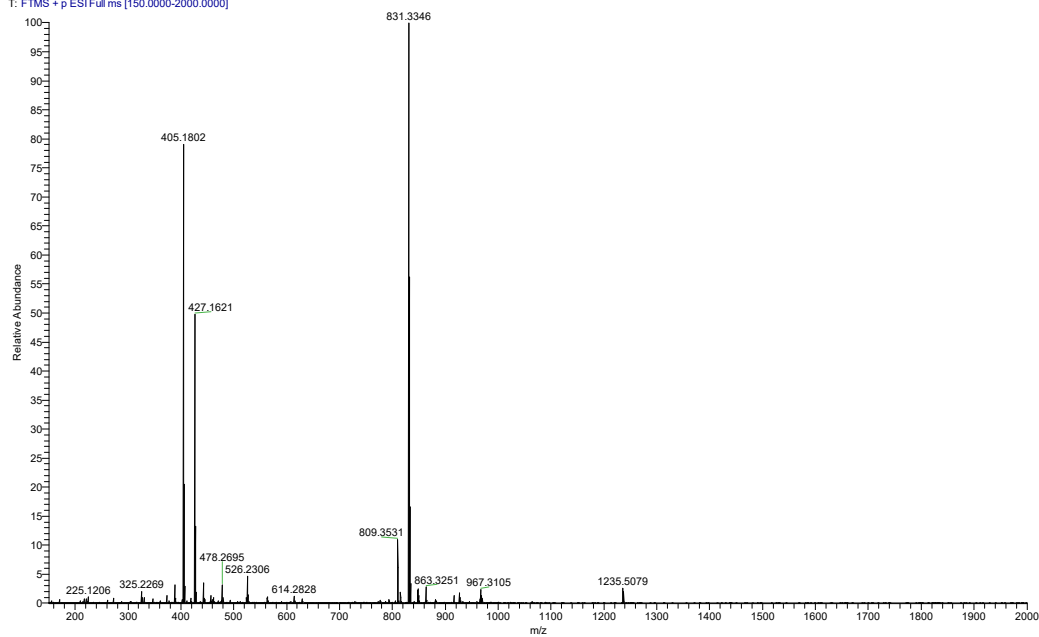
ESI-HRMS of **1_{n5}**.

MB15v2 #1-44 RT: 0.00-0.25 AV: 44 NL: 1.65E7
T: FTMS + p ESI Full ms [150.0000-2000.0000]

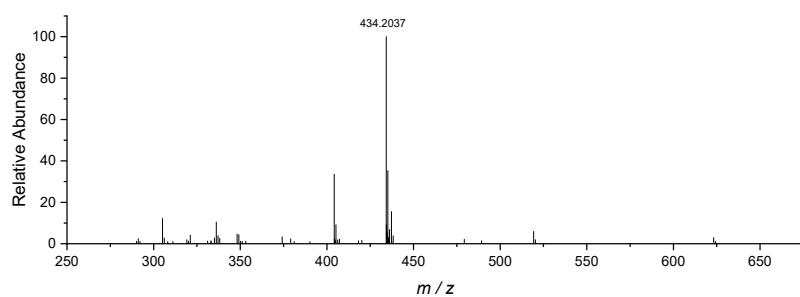


ESI-HRMS of **1_{a1}**.

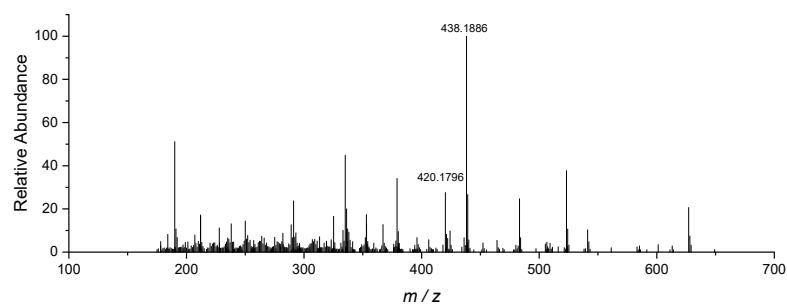
MB118 #1-47 RT: 0.00-0.25 AV: 47 NL: 1.95E7
T: FTMS + p ESI Full ms [150.0000-2000.0000]



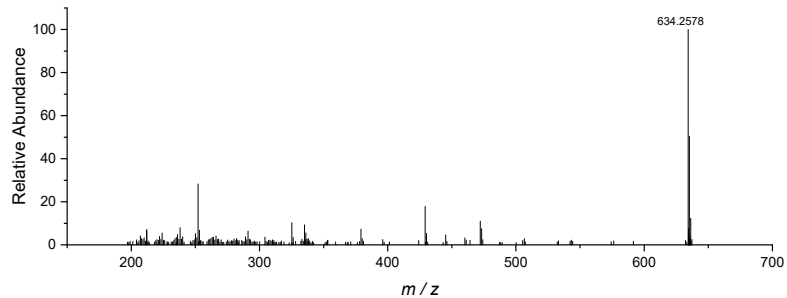
MALDI-HRMS of **1_{t1}**.



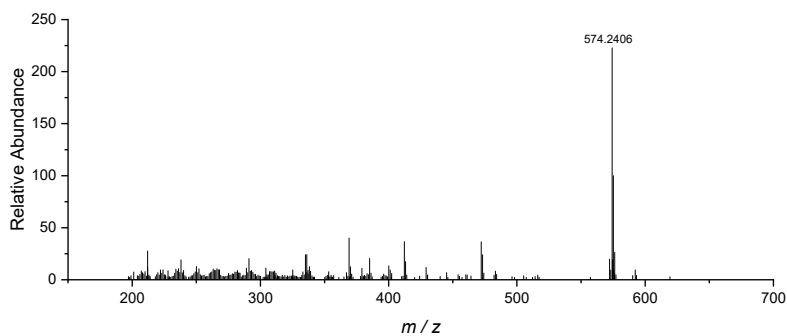
MALDI-HRMS of **1_{t2}**.



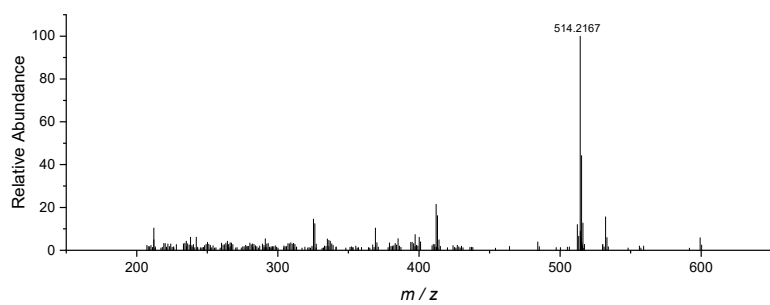
MALDI-HRMS of **1_{t3}**.



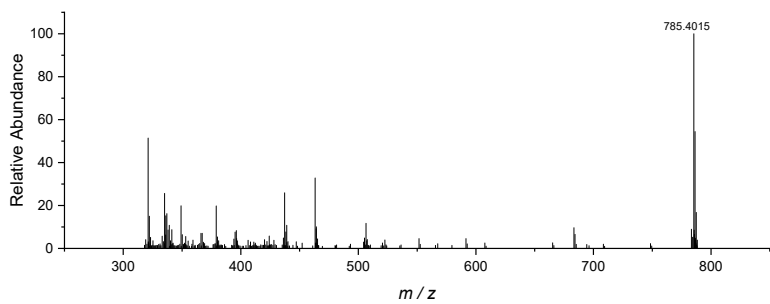
MALDI-HRMS of **1_{t5}**.



MALDI-HRMS of **1_{t6}**.

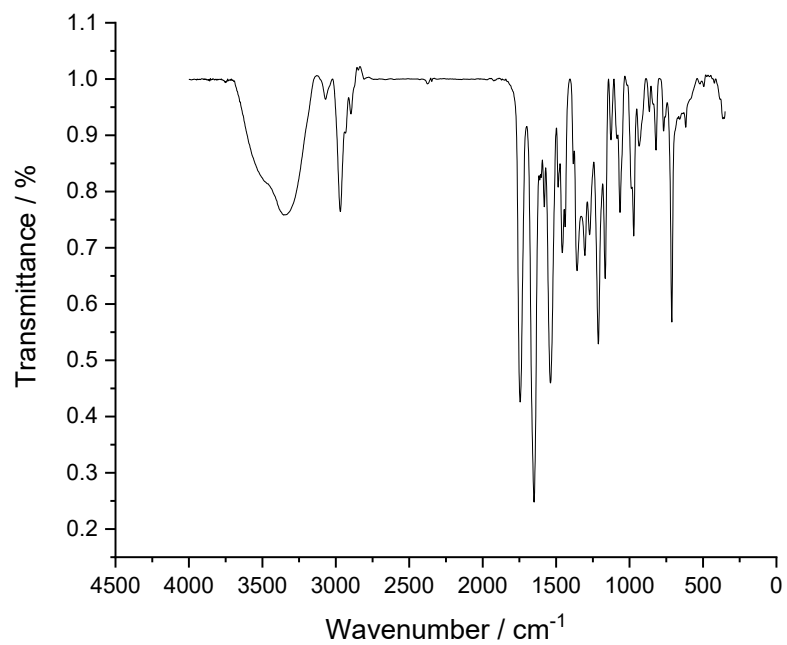


MALDI-HRMS of **1_b**.

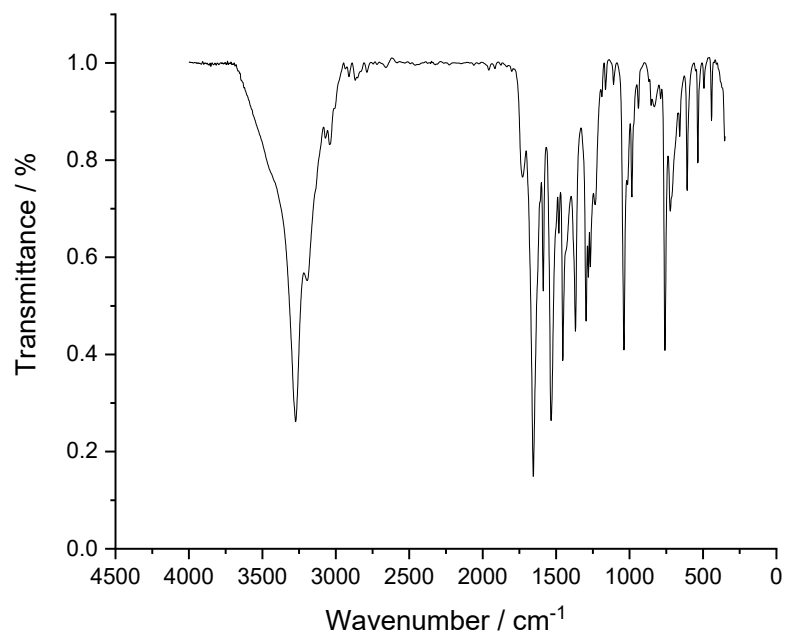


6. IR spectra.

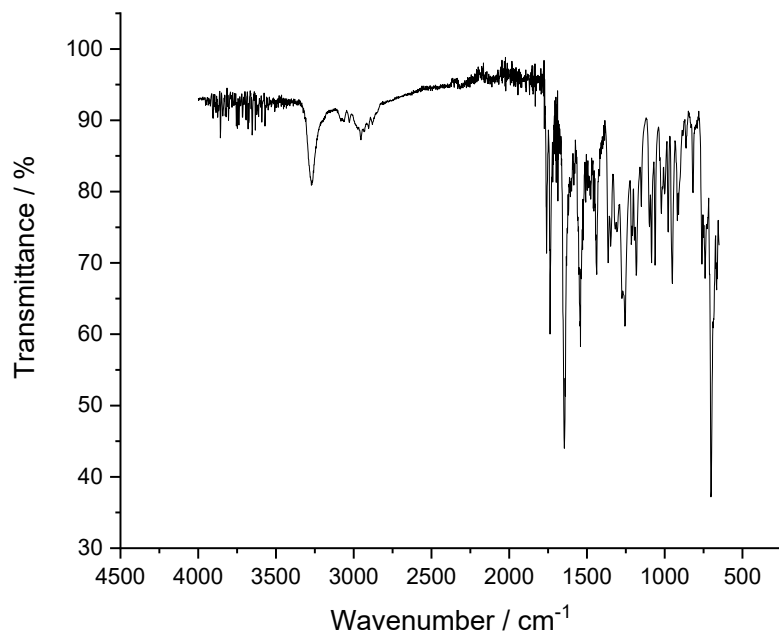
IR (KBr) of **1_{m1}**.



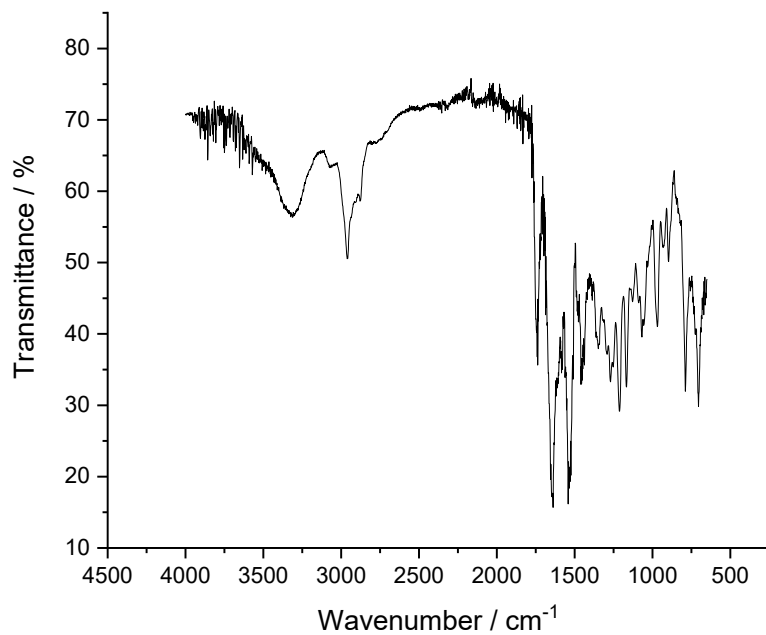
IR (KBr) of **1_{p1}**.



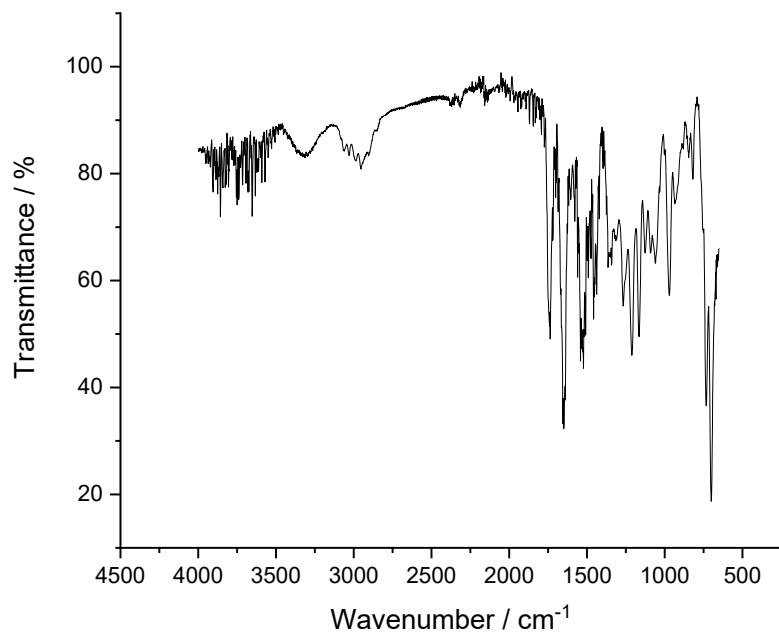
IR (ATR) of **1_{m2}**.



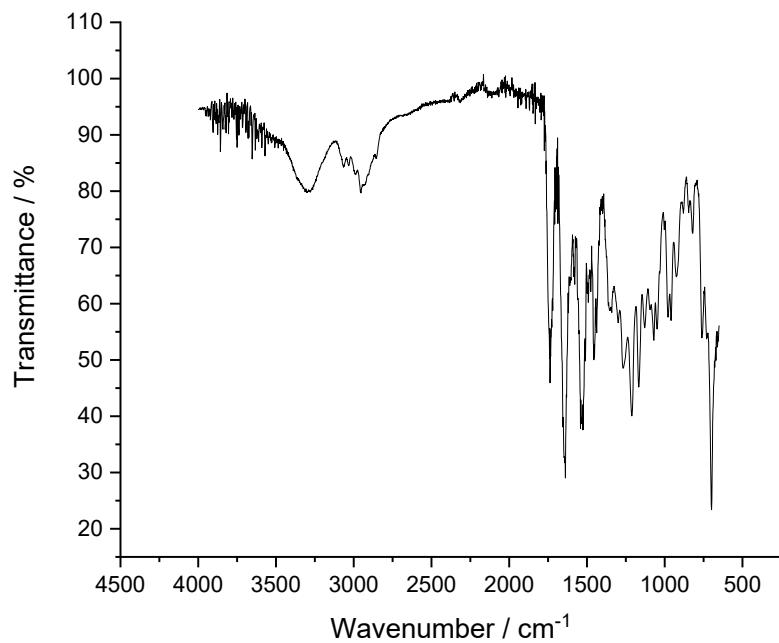
IR (ATR) of **1_{m3}**.



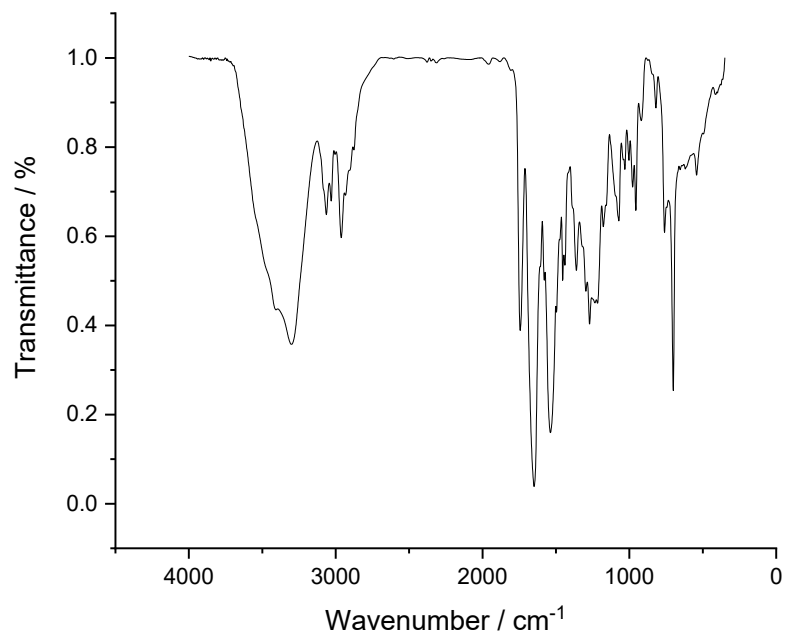
IR (ATR) of **1_{m4}**.



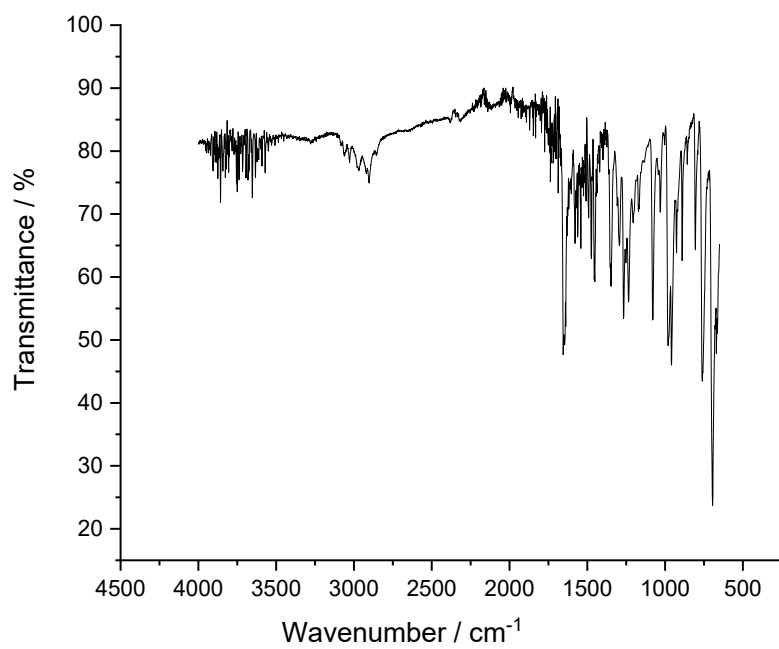
IR (ATR) of **1_{m5}**.



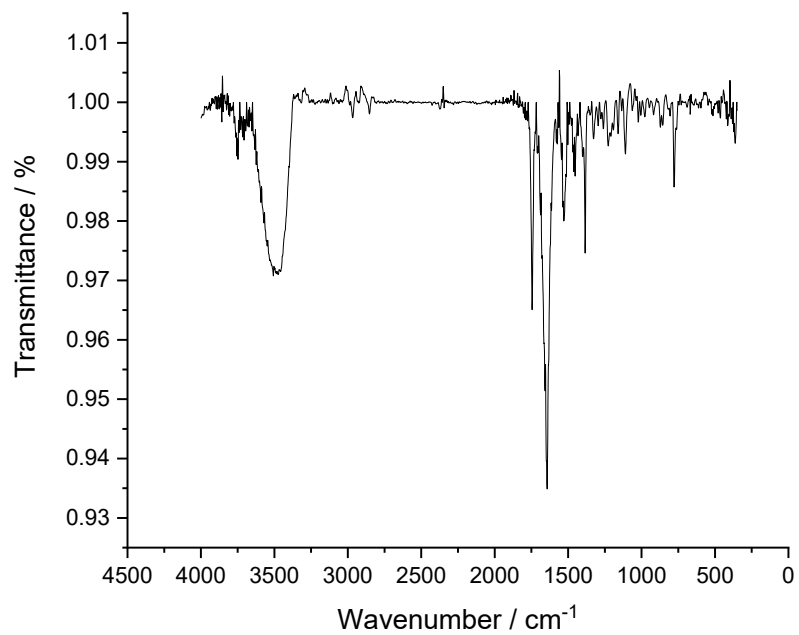
IR (KBr) of **1_{m6}**.



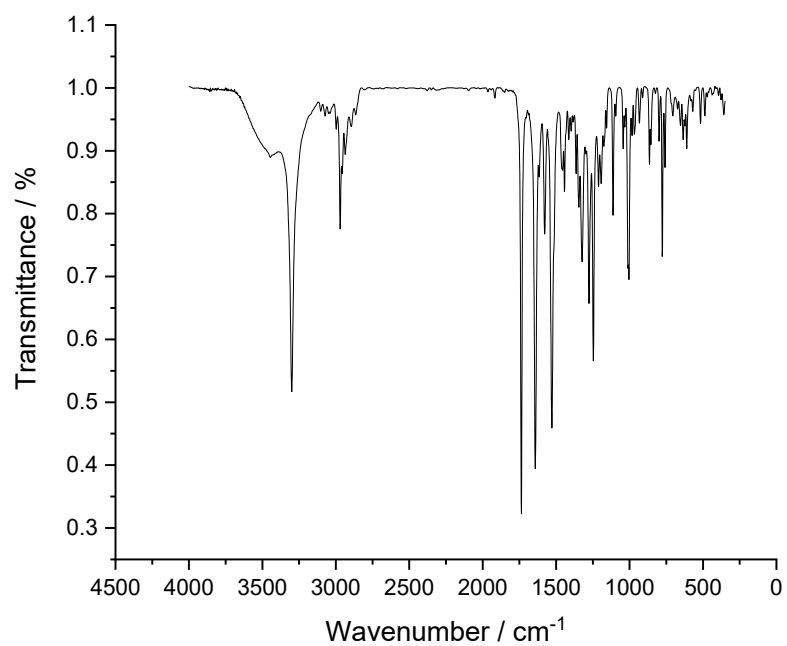
IR (ATR) of **1_{m7}**.



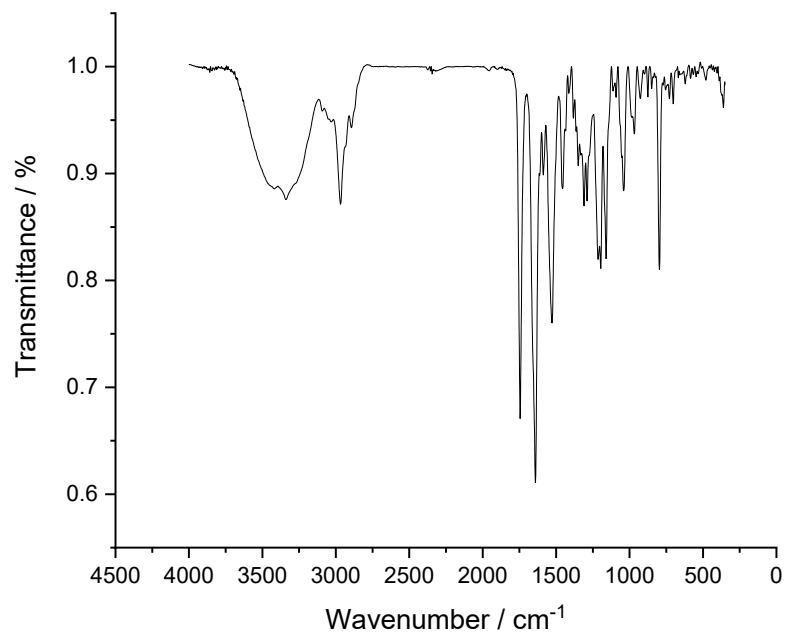
IR (KBr) of **1_{n1}**.



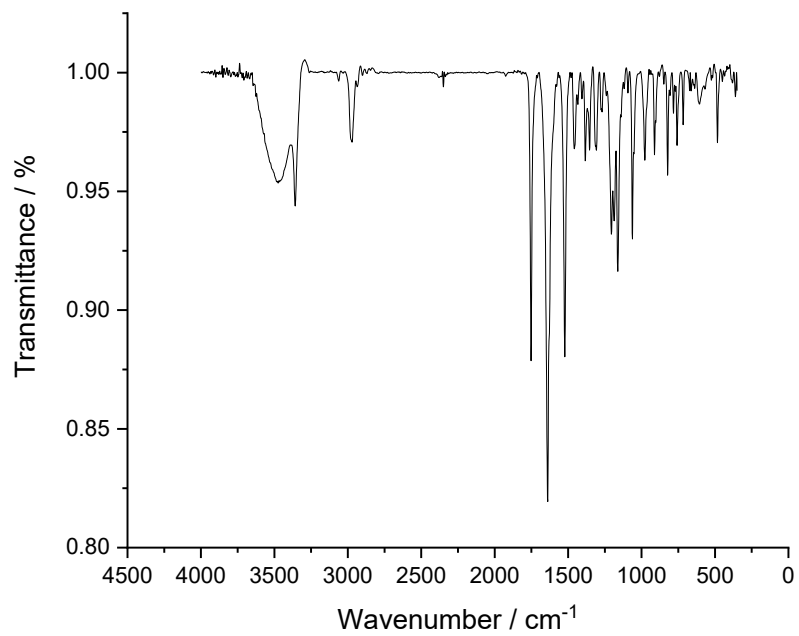
IR (KBr) of **1_{n2}**.



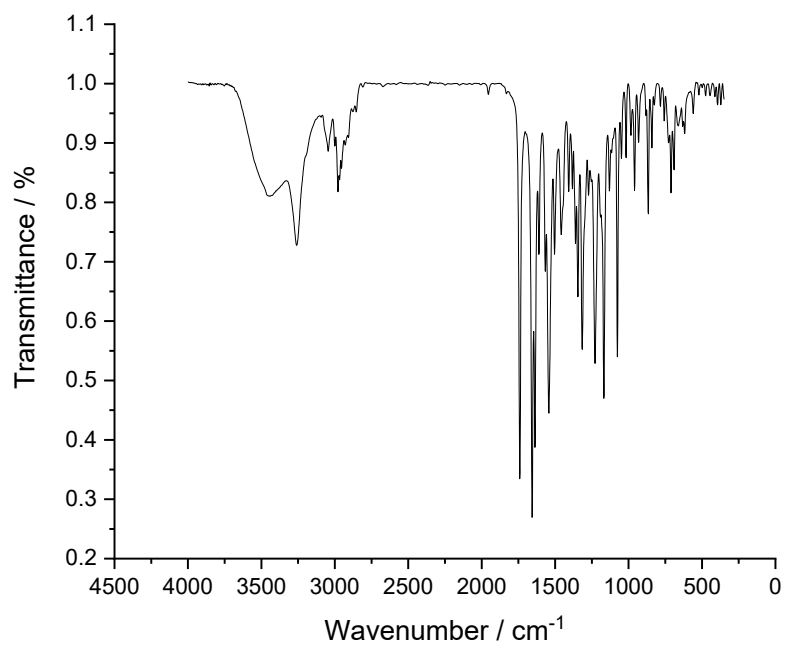
IR (KBr) of **1_{n3}**.



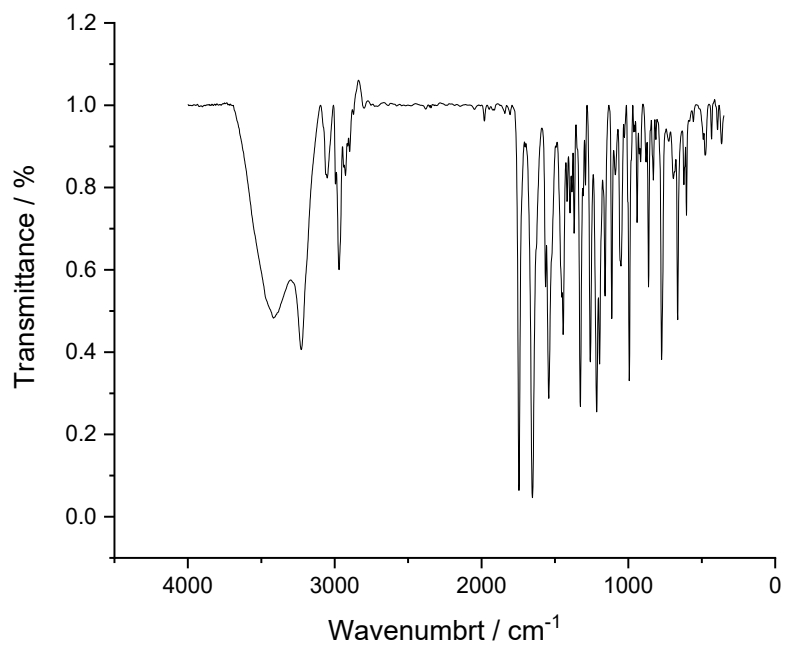
IR (KBr) of **1_{n4}**.



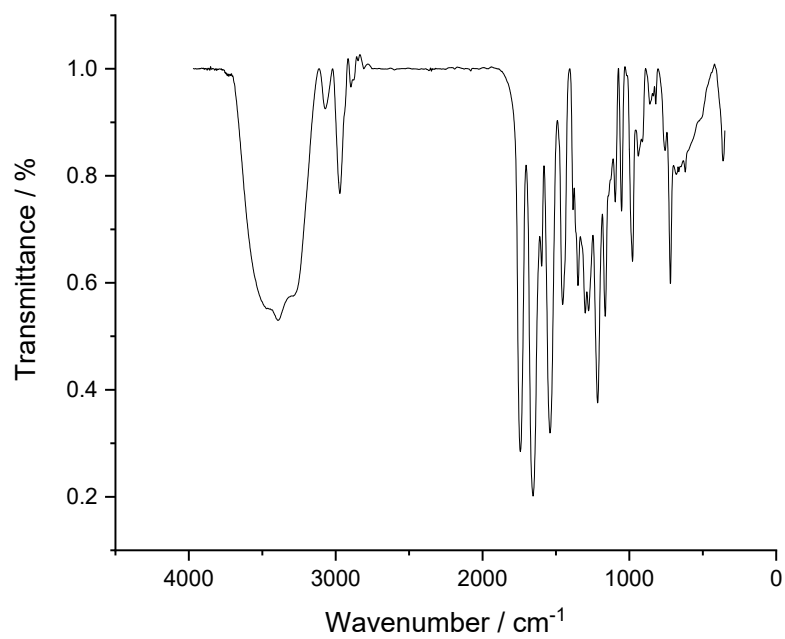
IR (KBr) of **1_{n5}**.



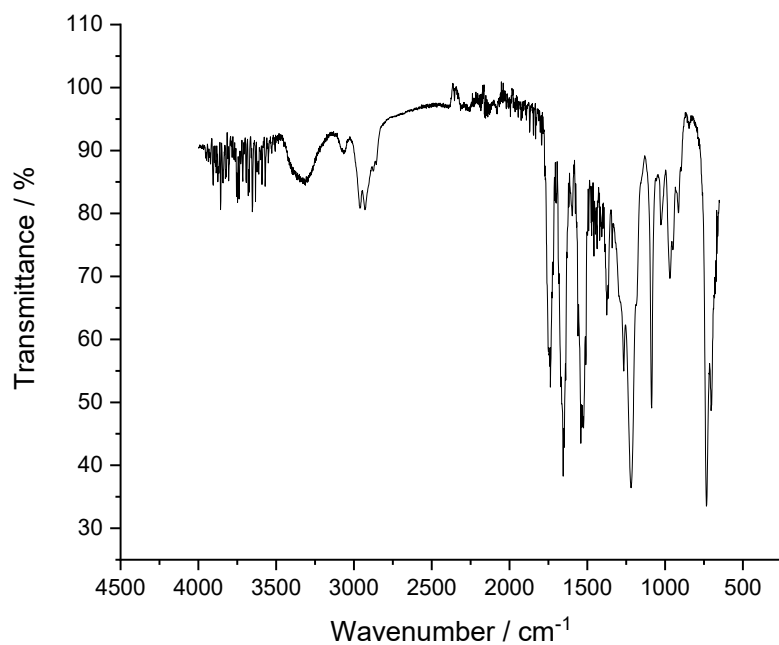
IR (KBr) of **1_{a1}**.



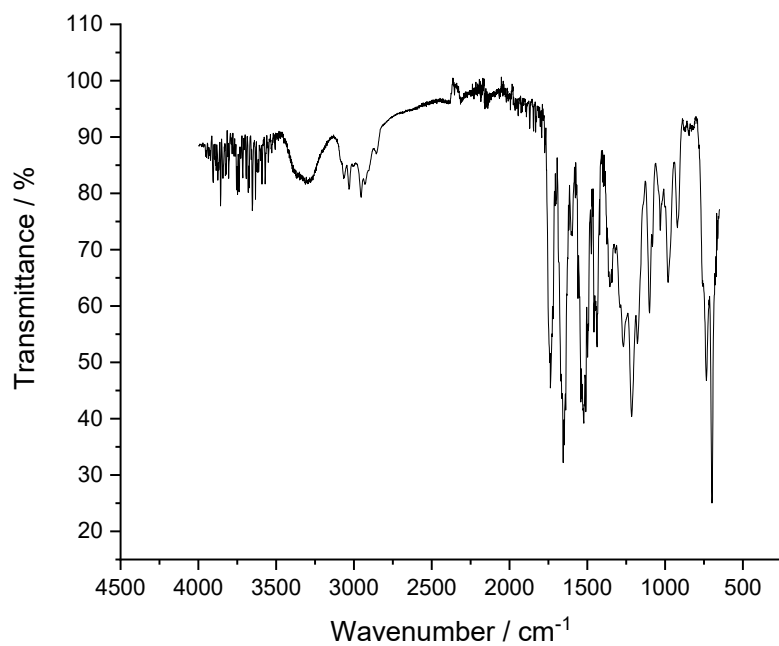
IR (KBr) of **1_{1t}**.



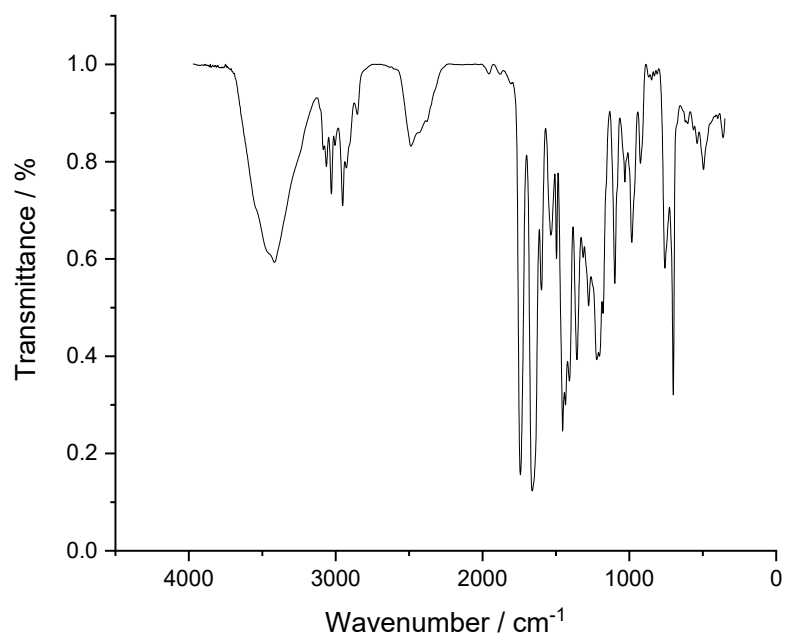
IR (ATR) of **1₂**.



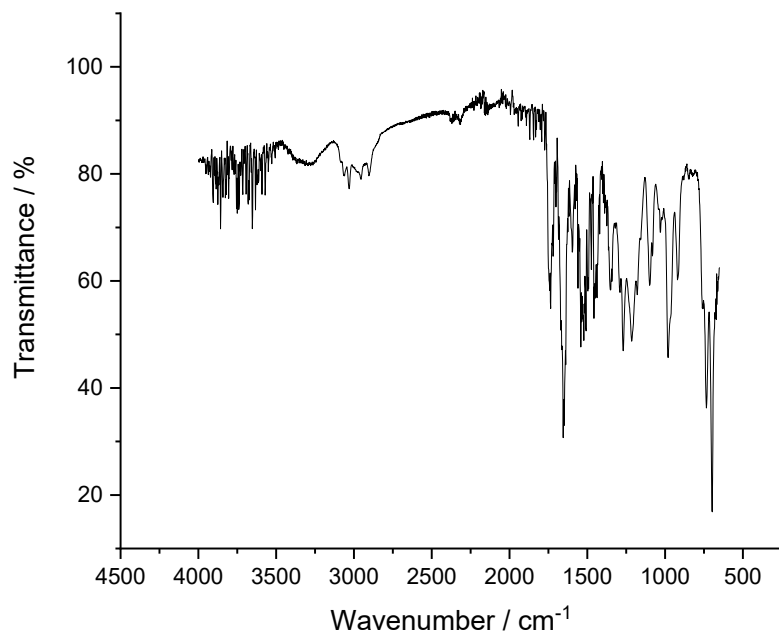
IR (ATR) of **1₃**.



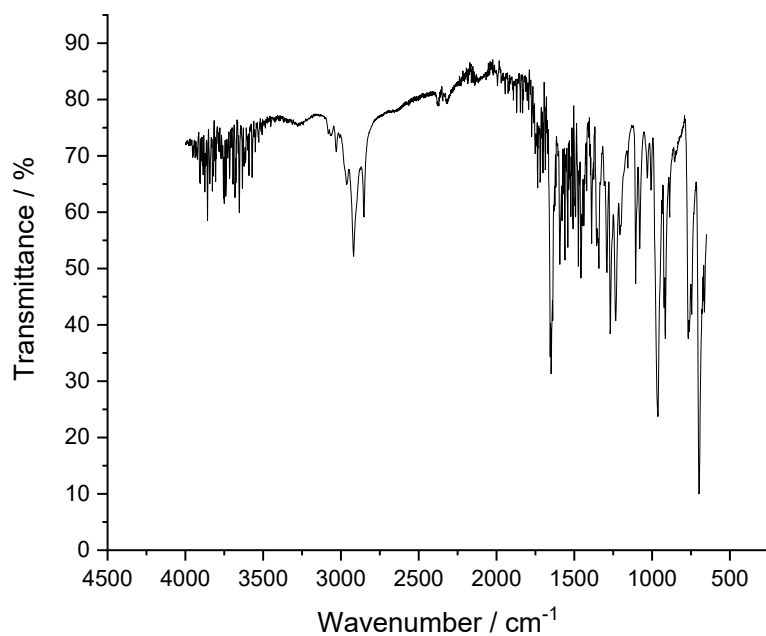
IR (KBr) of **1₄**.



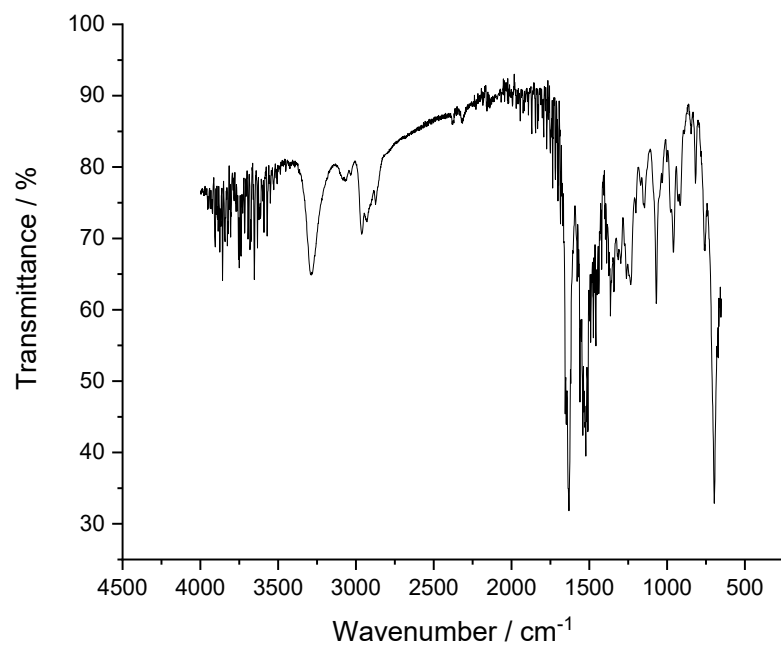
IR (ATR) of **1_{t5}**.



IR (ATR) of **1_{t6}**.



IR (ATR) of **1_b**.



7. X-ray single crystal structures.

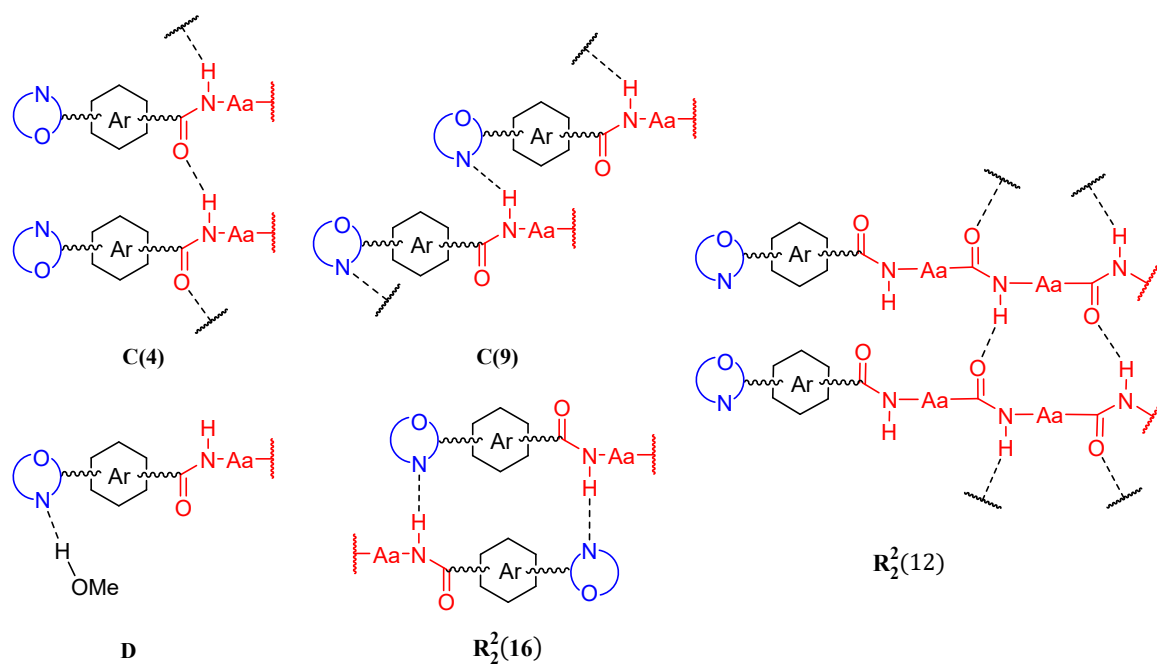


Figure S9. Types of hydrogen bonding interactions that occur in obtained crystal structures. Details are given in Table S5.

Table S5. Types of graph set motifs that appear in each oxazoline compound.

Oxazoline	Types of hydrogen bonding
1_{p1}	C(9)
1_{m6}[*]	C(4)C(4)D R₂²(12)[*]
1_{n2}	C(4)
1_{n4}	C(4)
1_{n5}	C(4)
1_{a1}	C(9)
1_{t4}	C(4) R₂²(16)

^{*} **R₂²(12)** is a binary graph set motif

Structures of similar oxazoline compounds with motifs disclosed above have been reported previously in literature. These structures can be divided into four categories according to unitary motifs which appear in the structure: oxazoline compounds in which oxazoline rings participate in a) a chain motif¹⁻⁸ or b) in a ring motif⁹⁻¹³ as well as oxazoline compounds in which oxazoline rings do not participate but have c) a chain motif^{11,14-20} or d) a ring motif.^{12,21} In all of these reported structures, only the oxazoline nitrogen atom acts as a hydrogen bonding acceptor. There are only two reported supramolecular structures (which do not contain a metal atom) in which the oxazoline oxygen atom acts as the hydrogen bonding acceptor.^{13,22}

The molecular structure of the oxazoline bioconjugates is dominated by six dihedral angles α , φ , θ , ϕ , ψ , and χ (Figure S10). Angle α is defined by $N_{ox}=C_{ox}-C_{a1}=O_{a1}$ dihedral angles (Figure S10, a). Values of α give information about relative directionality between the oxazoline and the amide double bond. Angles φ and θ , defined by $N_{ox}=C_{ox}-C_{Ar1}=C_{Ar2}$ and $O_{a1}=C_{a1}-C_{Ar4}=C_{Ar3}$ atoms, respectively, give insight in coplanarity of the oxazoline and amide C=O double bonds with the central aromatic unit (Figure S10, b and c). Angles ϕ and ψ^* are the Ramachandran dihedral angles of the amino acid residues, defined by $C_{a1}-N_{a1}-C_{\alpha}-C_{e(a2)}$ and $N_{a1}-C_{\alpha}-C_{e(a2)}-O_e(N_{a2})$ atom (Figure S10, d and e). Angles χ are defined by angles between two planes of amide and amide, or amide and ester carbonyl bonds (Figure S10, f). If a molecule has more than one χ value, then the successive values of χ are reported in order from N-terminus to C-terminus. Angles χ near 0° indicate a parallel orientation, values of 90° indicate a perpendicular orientation and values near 180° indicate antiparallel orientation. The experimental data of all defined angles for the 7 oxazolines are collected in Table S6. Data from crystal structures of similar compounds, amino-acid-aromatic and oxazoline-aromatic conjugates from the crystal base are collected in Table S6 (see below).

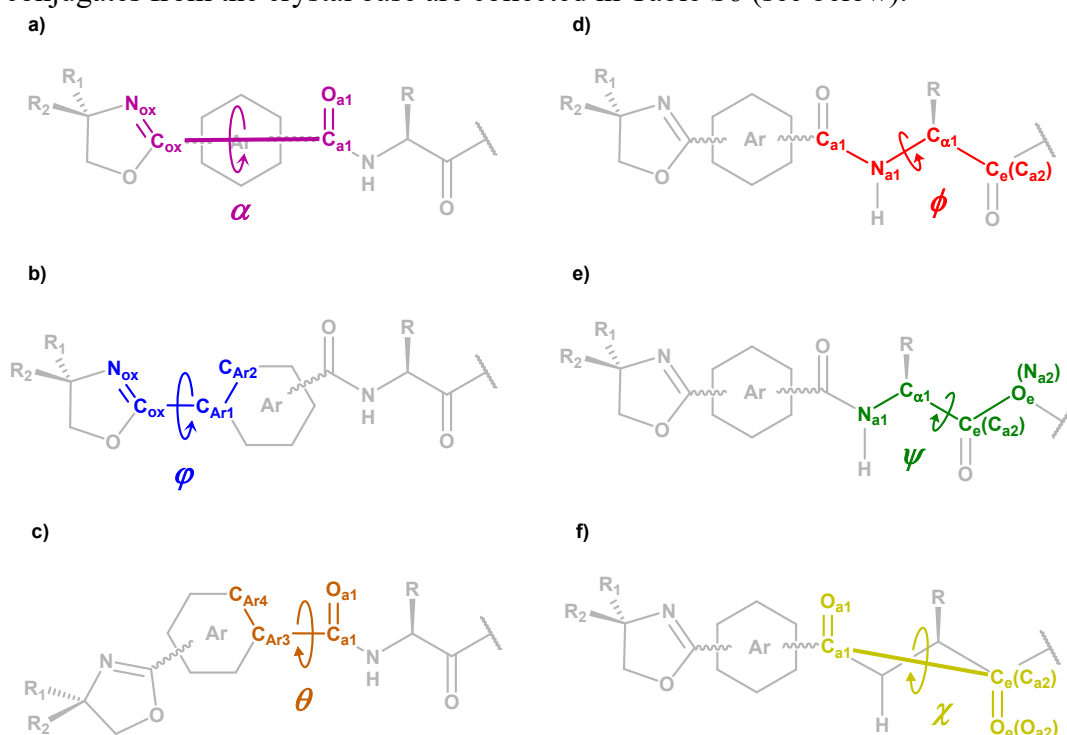


Figure S10. Characteristic dihedral angles: a) α , b) ϕ , c) θ , d) φ , e) ψ and f) χ of oxazoline compounds as defined in the text above.

Table S6. Dihedral angles as defined in Figure S10, for reported oxazoline structures and examples of structurally most similar compound from the literature.

Compound	$\alpha / ^\circ$	$\varphi / ^\circ$	$\theta / ^\circ$	$\phi / ^\circ$	$\psi / ^\circ$	$\chi / ^\circ$
1_p	-44	-12	-31	-85	-175	86
1_{m6}	140	-5	-21	-84 (Gly) -132 (Val) 37 (Phe)	172 (Gly) 127 (Val) -179 (Phe)	87 (Gly-Val) -126 (Val-Phe) -153 (Phe-ester)
1_{n2}	-76	22	-21, -47	-62	-31	78
1_{n4}	-50	-12	-30	-73	158	83
1_{n5}	2	-14	-16	-75	151	89
1_a	-17	-93	80	53	42	76
1_{t4}*	-41 (Phe ₁) 136 (Phe ₂)	-7	-21 (Phe ₁) -26 (Phe ₂)	95(Phe ₁) 61(Phe ₂)	-169 (Phe ₁) -141 (Phe ₂)	80 (Phe ₁) 79 (Phe ₂)
2_{m1}	-62	-	-34	-77	156	-112
2_{n5}	-61	-	-34	-77	158	-111
4_{m5}	-0.71	-	-42, 41	-147, -121	56, 50	-
3_{t3}	-101, - 115, 136	-	29, 35, 14	-114(A.al), -75, -115	71, 151, - 174	-116, 68
<i>p</i>C₆H₄(Gly-OMe)₂²³	180	-	-28, 28	65, -65	-152, 152	106, -106
2-Nph-Phe-OMe²⁴	-	-	25	-70	-37	11
2-Nph-Val-OMe²⁴	-	-	29	-72	138	-122
9,10-Anth(Phe-OMe)₂²⁵	-178	-	-67, 66	66, -90	-153, 164	106, -115
9,10-Anth(Val-OMe)₂²⁵	153	-	-95, 68	-88, -75	141, 137	-126, -119
1,3,5-C₆H₃-(Phe-OMe)₃²⁶	122,123, 142	-	158, - 29, 169	88, 98, 102	21, -32, -45	-14, -119 -128
1,3,5-C₆H₃-(Gly-OMe)₃²⁶	-88,-89, - 84	-	-	-	-	-
(Phg-ox)<i>m</i>C₆H₄CN²⁷	-167	-173	-	-	-	-
(ETA-ox)₂<i>m</i>C₆H₄²⁸	-169	178, 19	-	-	-	-

carbonyl is separated from the aromatic ring by the amide nitrogen.

7.1 Oxazolines

Table S7. Experimental data for the X-ray diffraction studies

Compound	1_{p1}	1_{m6}	1_{n2}	1_{n4}
Formula	C ₁₆ H ₂₀ N ₂ O ₄	C ₃₃ H ₃₆ N ₄ O ₆ ·Cl O	C ₁₉ H ₂₀ N ₂ O ₄	C ₂₀ H ₂₂ N ₂ O ₄
<i>F_w</i> (g mol ⁻¹)	304.34	616.70	340.37	354.39
Crystal system	Tetragonal	Monoclinic	Monoclinic	Monoclinic
Space group	<i>P</i> 4 ₃ (No. 78)	<i>P</i> 2 ₁ (No. 4)	<i>P</i> 2 ₁ / <i>c</i> (No. 14)	<i>C</i> 2 (No. 5)
<i>a</i> (Å)	7.30610(10)	4.85790(10)	14.4335(2)	22.688(4)
<i>b</i> (Å)	7.30610(10)	21.1948(4)	13.2892(2)	5.2784(6)
<i>c</i> (Å)	29.5083(6)	15.9374(3)	9.4990(2)	16.099(2)
<i>α</i> (°)	90	90	90	90
<i>β</i> (°)	90	91.905(2)	105.099(2)	105.082(17)
<i>γ</i> (°)	90	90	90	90
<i>V</i> (Å ³)	1575.13(5)	1640.04(6)	1759.10(5)	1861.6(5)
<i>Z</i>	4	2	4	4
<i>D_{calc}</i> (g cm ⁻³)	1.283	1.249	1.285	1.264
<i>F</i> (000)	648	656	720	752
Instrument	XtaLAB	Xcalibur	XtaLAB	Xcalibur
Radiation (Å)	1.54184	1.54184	1.54184	1.54184
Temperature (K)	293(2)	293(2)	293(2)	293(2)
Reflections collected	5610	8050	12736	2235
Independent reflections	2524	4801	3615	1615
<i>R_{init}</i>	0.0253	0.0320	0.0349	0.0307
Reflections observed	2375	4108	3143	1209
Parameters	207	423	233	243
<i>R</i> ₁ [<i>I</i> > 2σ(<i>I</i>)] ^[a]	0.0376	0.0504	0.0528	0.0549
<i>wR</i> ₂ (all data) ^[b]	0.0930	0.1596	0.1550	0.1352
Goof, <i>S</i> ^[c]	1.105	1.169	1.063	1.030
Maximum/minimum electron density (e Å ³)	0.152/-0.195	0.247/-0.228	0.306/-0.392	0.168/-0.135

^[a] $R_1 = \frac{\sum ||F_o| - |F_c||}{\sum |F_o|}$. ^[b] $wR_2 = \{\frac{\sum [w(F_o^2 - F_c^2)^2]}{\sum [w(F_o^2)^2]}\}^{1/2}$. ^[c] $S = \{\frac{\sum [w(F_o^2 - F_c^2)^2]}{(n - p)}\}^{1/2}$ where *n* is number of reflections and *p* is the total number of parameters refined

Table S7. Experimental data for the X-ray diffraction studies (continuation)

Compound	1_{n5}	1_{a1}	1_{t4}
Formula	C ₂₀ H ₂₂ N ₂ O ₄	C ₂₄ H ₂₄ N ₂ O ₄	C ₃₇ H ₃₅ N ₃ O ₇
<i>F_w</i> (g mol ⁻¹)	354.39	404.45	633.68
Crystal system	Orthorhombic	Orthorhombic	Tetragonal
Space group	<i>P</i> 2 ₁ 2 ₁ 2 ₁ (No. 11)	<i>P</i> 2 ₁ 2 ₁ 2 (No. 18)	<i>P</i> 4 ₁ 2 ₁ 2 (No. 92)
<i>a</i> (Å)	5.1830(1)	13.2430(4)	18.8070(2)
<i>b</i> (Å)	5.5120(1)	20.7980(6)	18.8070(2)
<i>c</i> (Å)	64.5994(13)	7.7513(2)	19.0641(5)
<i>α</i> (°)	90	90	90
<i>β</i> (°)	90	90	90
<i>γ</i> (°)	90	90	90
<i>V</i> (Å ³)	1845.52(6)	2134.92(10)	6743.0(2)
<i>Z</i>	4	4	8
<i>D_{calc}</i> (g cm ⁻³)	1.275	1.258	1.248
<i>F</i> (000)	752	856	2672
Instrument	XtaLAB	XtaLAB	Xcalibur
Radiation (Å)	1.54184	1.54184	1.54184
Temperature (K)	293(2)	293(2)	293(2)
Reflections collected	14023	8620	14949
Independent reflections	3887	4022	5899
<i>R_{init}</i>	0.0353	0.0318	0.0326
Reflections observed	3813	3596	4958
Parameters	243	279	434
<i>R</i> ₁ [<i>I</i> > 2σ(<i>I</i>)] ^[a]	0.0867	0.0468	0.0419
<i>wR</i> ₂ (all data) ^[b]	0.2865	0.1288	0.1103
Goof, <i>S</i> ^[c]	1.161	1.083	1.038
Maximum/minimum electron density (e Å ⁻³)	0.498/-0.350	0.218/-0.176	0.138/-0.109

^[a] $R_1 = \frac{\sum ||F_o| - |F_c||}{\sum |F_o|}$. ^[b] $wR_2 = \{\frac{\sum [w(F_o^2 - F_c^2)^2]}{\sum [w(F_o^2)^2]}\}^{1/2}$. ^[c] $S = \{\frac{\sum [w(F_o^2 - F_c^2)^2]}{(n - p)}\}^{1/2}$ where *n* is number of reflections and *p* is the total number of parameters refined

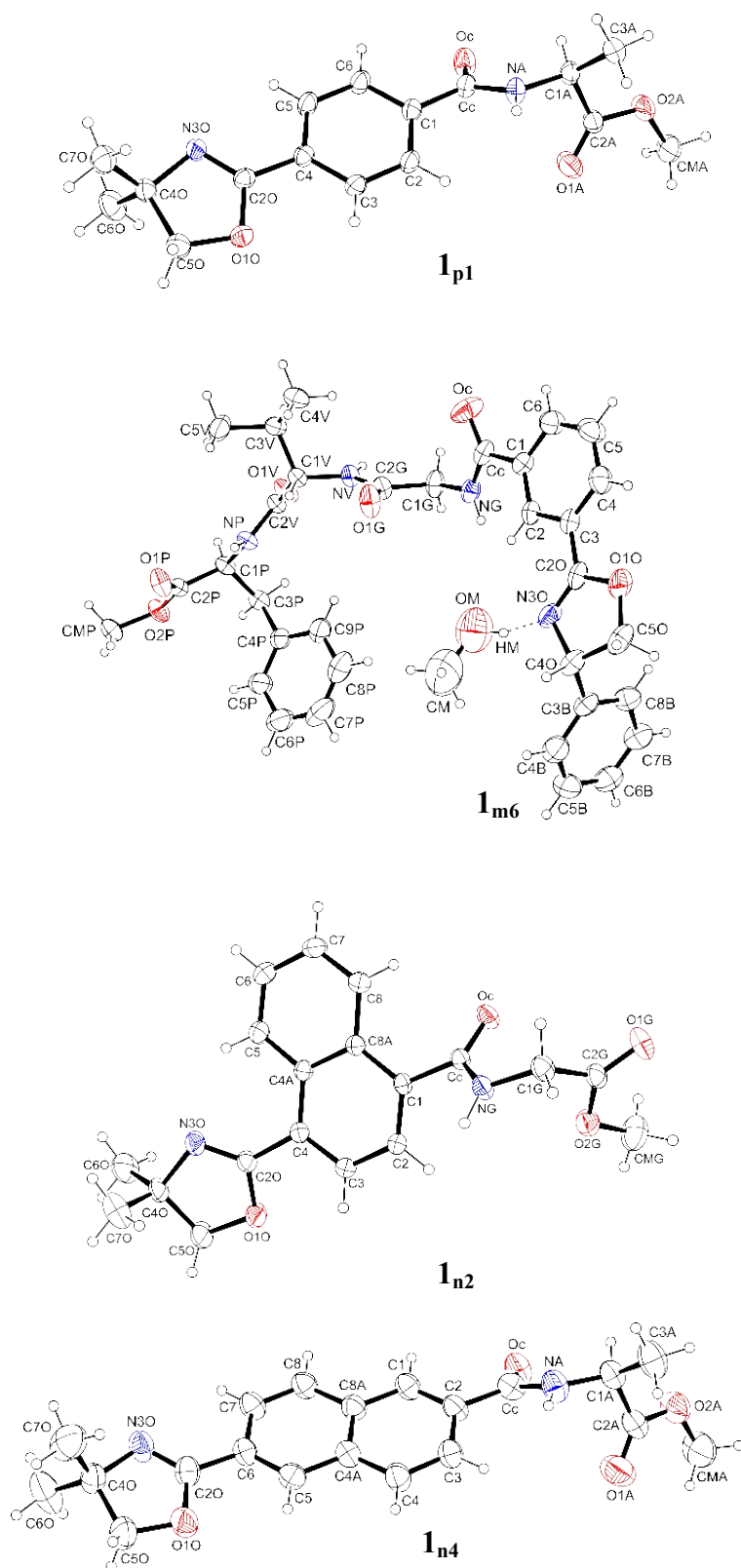
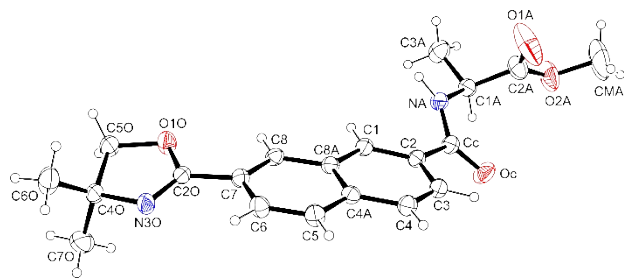
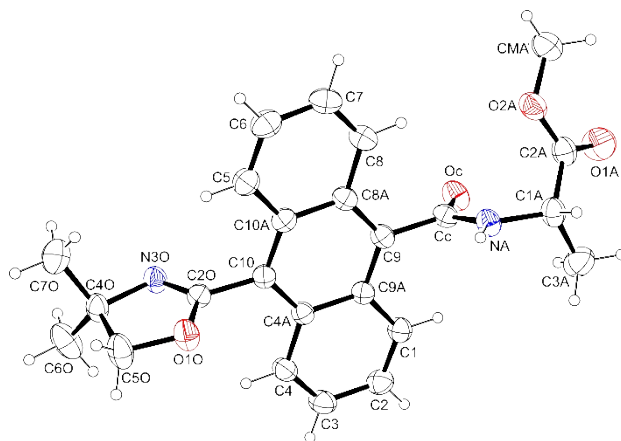


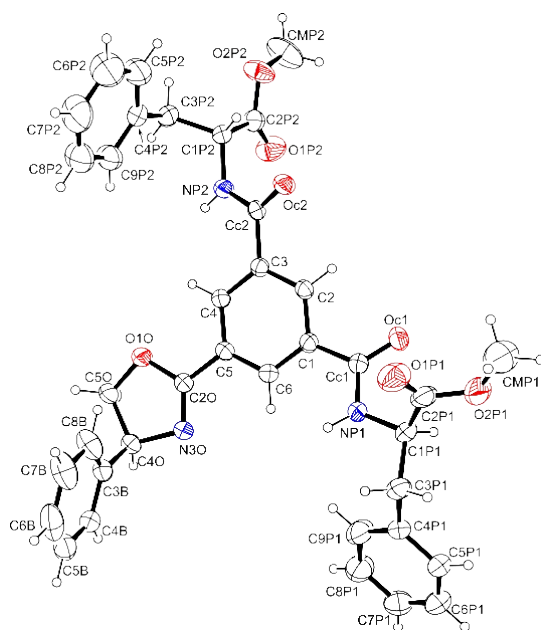
Figure S11. ORTEP-III drawings (Farrugia)²⁹ for SCXRD determined structures with complete atom numbering schemes and 30 % ellipsoid probability level.



1_{n5}



1_{a1}



1_{t4}

Figure S11. ORTEP-III drawings (Farrugia)²⁹ for SCXRD determined structures with complete atom numbering schemes and 30 % ellipsoid probability level. (continuation).

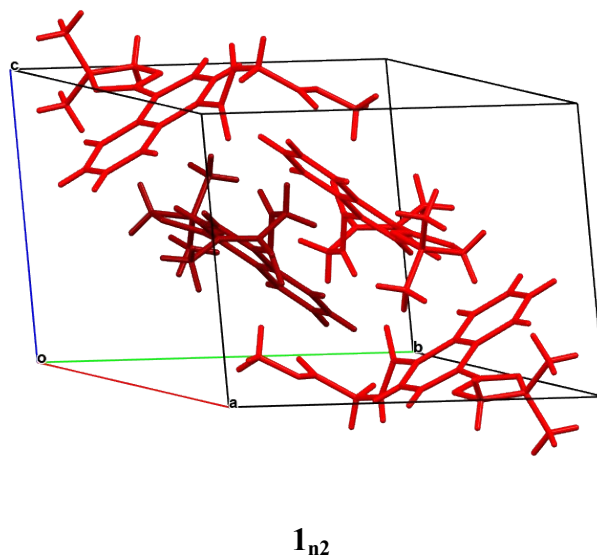
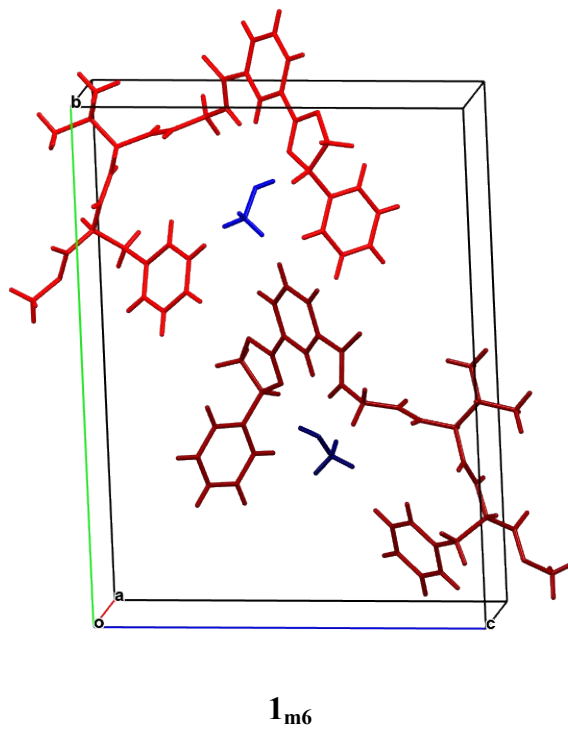
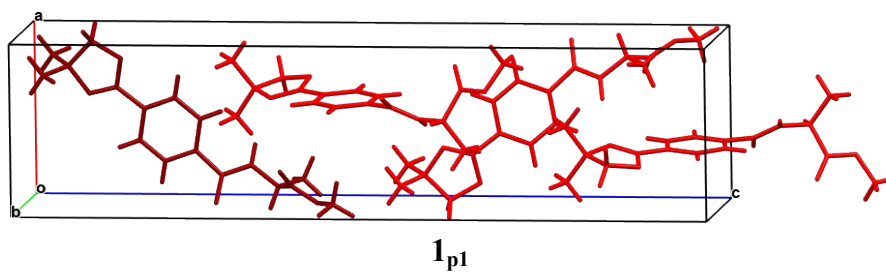


Figure S12. Crystal packings for SCXRD determined structures. Solvent methanol molecules in 1_{m6} are shown in blue.

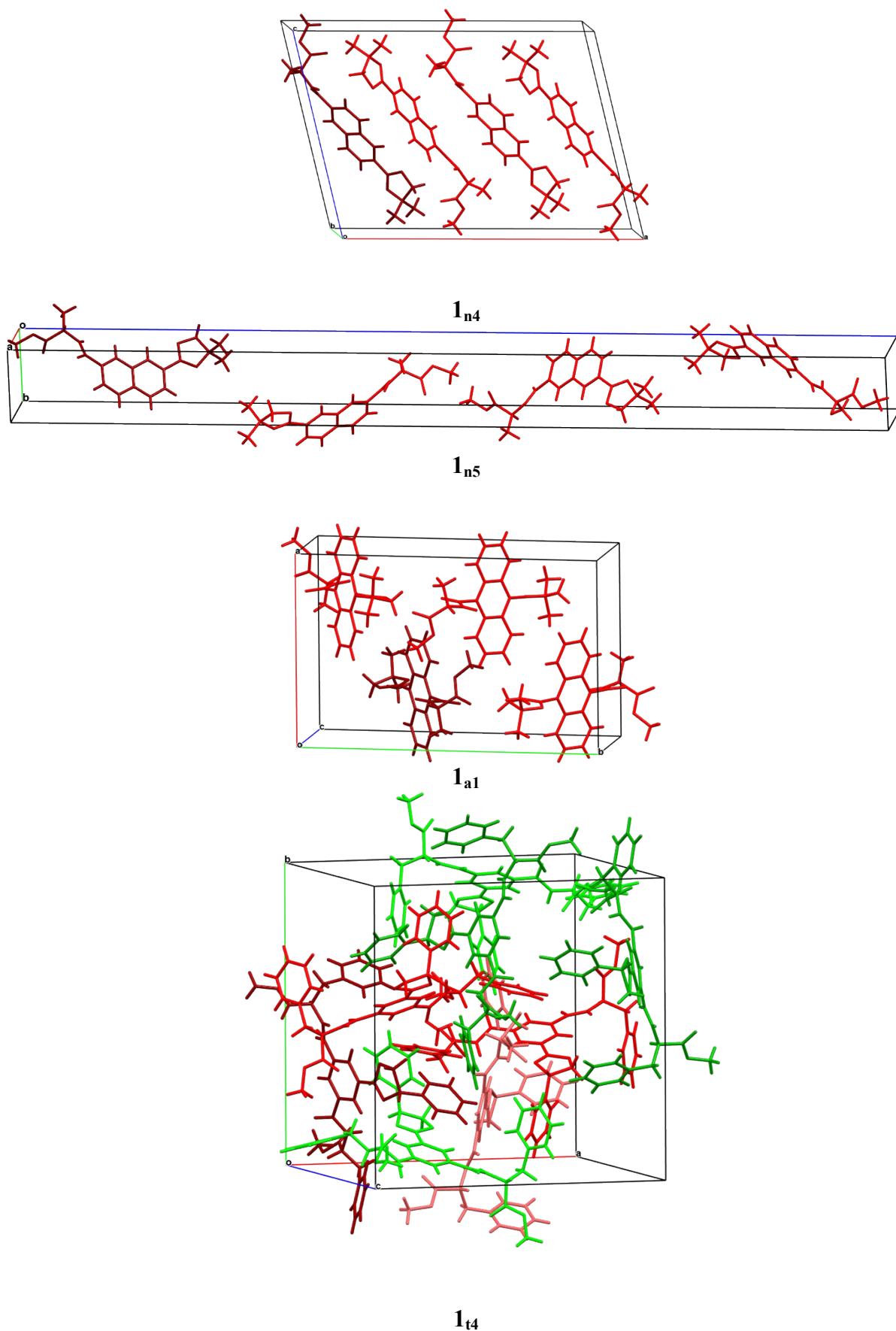


Figure S12. Crystal packings for SCXRD determined structures. (continuation)

7.2. Precursors

Table S8. Experimental data for the X-ray diffraction studies.

Compound	2_{m1}	4_{m5}	2_{n5}	3_{t3}
Formula	C ₁₆ H ₂₀ N ₂ O ₆	C ₂₄ H ₂₄ N ₂ O ₄	C ₂₀ H ₂₂ N ₂ O	C ₃₇ H ₃₇ N ₃ O ₈
<i>F_w</i> (g mol ⁻¹)	336.34	404.45	386.39	651.69
Crystal system	Monoclinic	Monoclinic	Monoclinic	Tetragonal
Space group	C2 (No. 5)	C2 (No. 5)	C2 (No. 5)	P4 ₃ 2 ₁ 2 (No. 96)
<i>a</i> (Å)	6.4358(2)	20.892(2)	6.5044(4)	18.8640(2)
<i>b</i> (Å)	8.0804(3)	4.9926(3)	8.0195(6)	18.8640(2)
<i>c</i> (Å)	16.4380(6)	21.517(2)	18.7137(1)	19.6976(4)
<i>α</i> (°)	90	90	90	90
<i>β</i> (°)	98.198(3)	111.246(12)	95.141(6)	90
<i>γ</i> (°)	90	90	90	90
<i>V</i> (Å ³)	846.10(5)	2091.8(4)	972.22(11)	7009.4(2)
<i>Z</i>	2	4	2	8
<i>D_{calc}</i> (g cm ⁻³)	1.320	1.284	1.320	1.235
<i>F</i> (000)	356	856	408	2755
Instrument	Xcalibur	Xcalibur	Xcalibur	Xcalibur
Radiation (Å)	1.54184	1.54184	1.54184	1.54184
Temperature (K)	293(2)	293(2)	293(2)	293(2)
Reflections collected	1358	11420	1845	22636
Independent reflections	994	3874	1343	7226
<i>R_{init}</i>	0.0195	0.0849	0.0258	0.0411
Reflections observed	980	2974	1261	4601
Parameters	116	281	134	443
<i>R₁</i> [<i>I</i> > 2σ(<i>I</i>)] ^[a]	0.0483	0.0671	0.0416	0.0852
<i>wR₂</i> (all data) ^[b]	0.1483	0.1950	0.1413	0.2769
Goof, <i>S</i> ^[c]	1.165	1.060	1.179	1.048
Maximum/minimum electron density (e Å ⁻³)	0.279/-0.176	0.217/-0.222	0.292/-0.2	0.295/-0.194

^[a] $R_1 = \sum ||F_o| - |F_c|| / \sum |F_o|$. ^[b] $wR_2 = \{\sum [w(F_o^2 - F_c^2)^2] / \sum [w(F_o^2)^2]\}^{1/2}$. ^[c] $S = \{\sum [w(F_o^2 - F_c^2)^2] / (n - p)\}^{1/2}$ where n is number of reflections and p is the total number of parameters refined

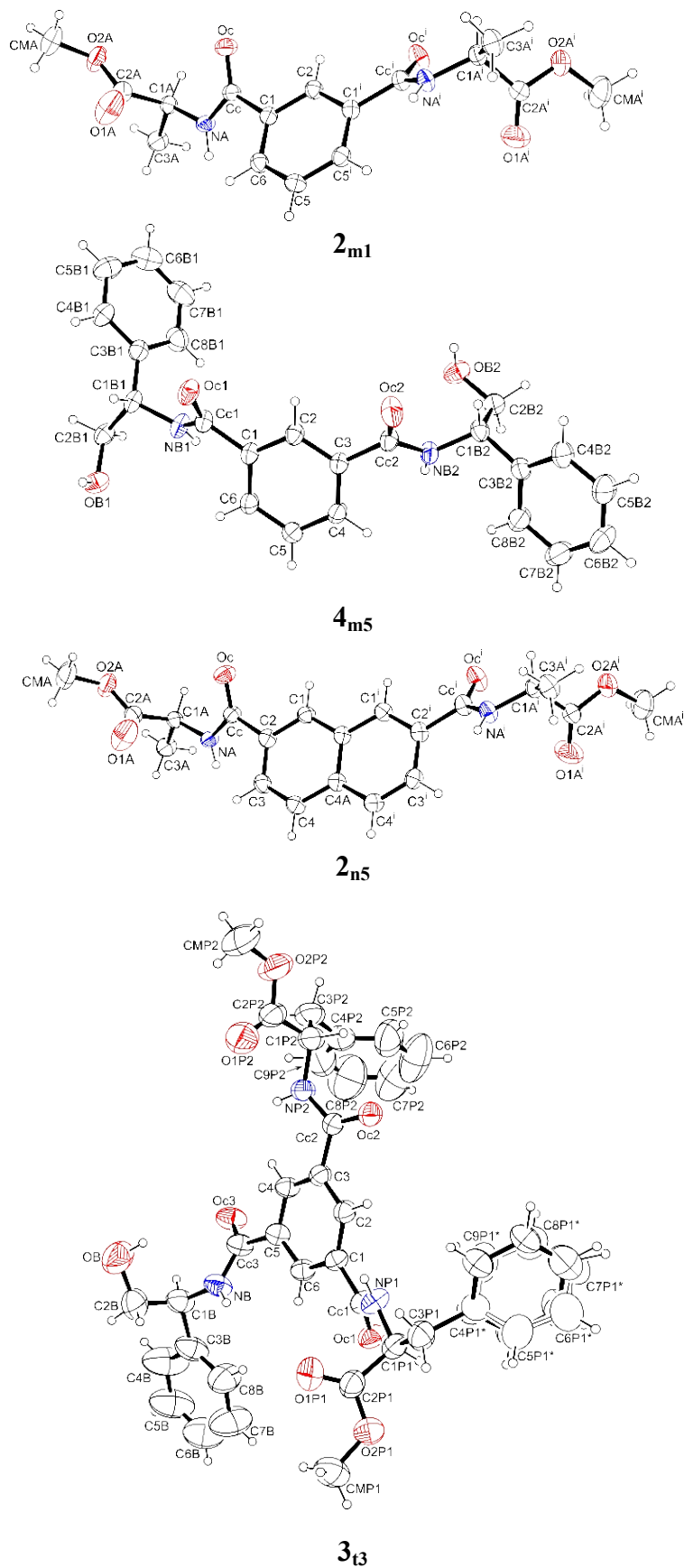
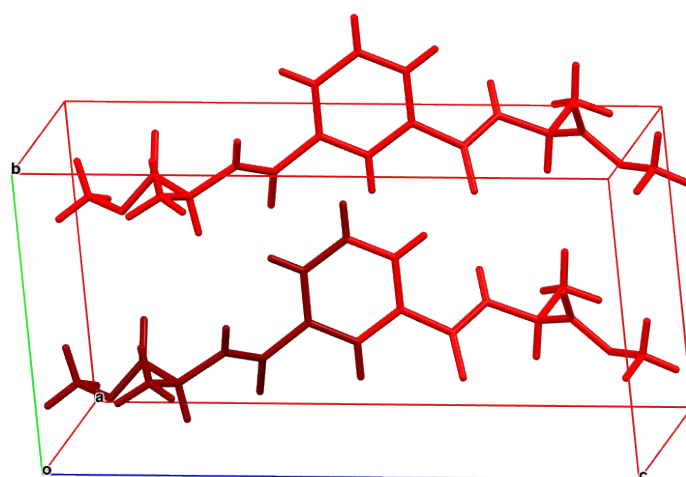
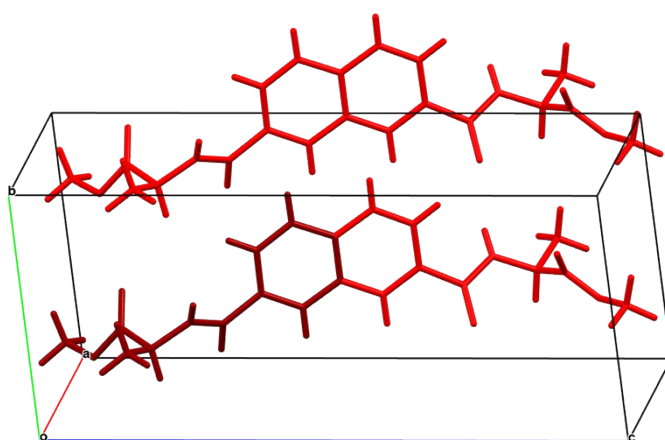


Figure S13. ORTEP-III drawings (Farrugia)²⁹ for SCXRD determined structures with complete atom numbering schemes and 30 % ellipsoid probability level.



2_{m1}



2_{n5}

Figure S14. Crystal packings for SCXRD determined structures. For reason of clarity, symmetry equivalent molecules are drawn in different shades of red and green.

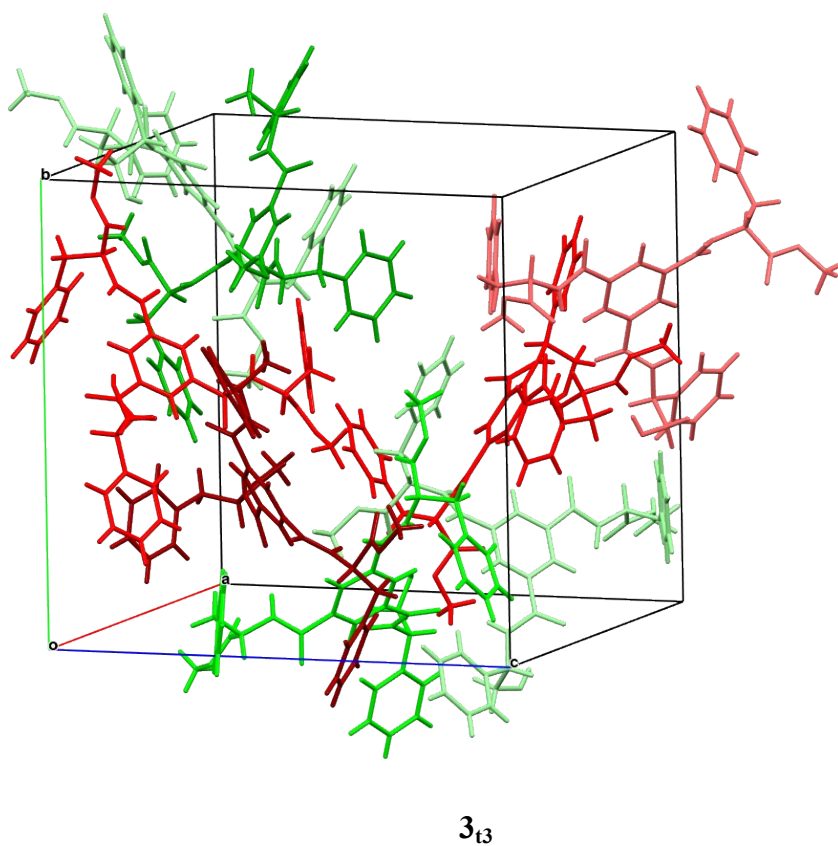
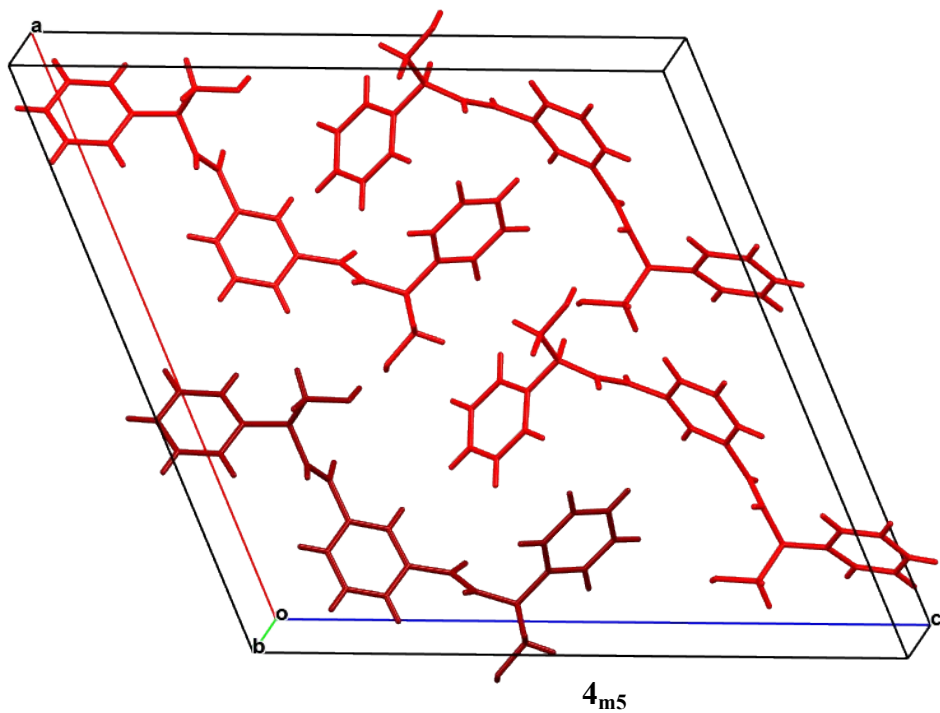


Figure S15. Crystal packings for SCXRD determined structures. For reason of clarity, symmetry equivalent molecules are drawn in different shades of red and green.

7.3. Hydrogen bonds

Table S9. Geometry parameters (\AA , $^\circ$) for hydrogen bonds in SCXRD determined structures.

<i>D-H</i> ⋯ <i>A</i> , type	<i>D-H</i>	<i>H</i> ⋯ <i>A</i>	<i>D</i> ⋯ <i>A</i>	<i>D-H</i> ⋯ <i>A</i>
1_{p1} NA-HA⋯N3O, am⋯oxz	0.90(3)	2.24(3)	3.119(3)	166(3)
1_{m6} NV-HV⋯O1G, am⋯am NP-HP⋯O1V, am⋯am OM-HM⋯N3O, dimer with solvent	0.88(2) 0.86(2) 0.82	2.13(2) 2.28(3) 2.24	2.988(4) 3.062(4) 2.916(11)	167(4) 151(4) 139.3
1_{n2} NG-HG⋯OC, am⋯am	0.89(2)	2.01(2)	2.8208(16)	150.8(16)
1_{n4} NA-HA⋯OC, am⋯am	0.77(9)	2.46(9)	3.158(11)	152(8)
1_{n5} NA-HA⋯OC, am⋯am	0.91(6)	2.14(6)	3.050(6)	171(5)
1_{a1} NA-HA⋯N3O, am⋯oxz	0.97(4)	2.11(4)	3.054(3)	165(3)
1_{t4} NP2-HP2⋯OC2, am⋯am NP1-HP1⋯N3O, am⋯oxz	0.87(2) 0.89(2)	2.13(3) 2.29(2)	2.956(3) 3.175(3)	159(3) 173(3)
2_{m1} NA-HA⋯OC, am⋯am	0.87(4)	2.16(5)	3.019(4)	170(3)
4_{m5} NB1-HB1⋯OC1, am⋯am NB2-HB2⋯OC2, am⋯am OB1-HOB1⋯OB2, OH⋯OH OB2-HOB2⋯OB1, OH⋯OH	0.88(8) 0.96(7) 0.82 0.82	2.10(8) 1.96(7) 1.98 1.88	2.974(5) 2.911(5) 2.772(6) 2.696(6)	179(5) 173(6) 161.2 174.2
2_{n5} NA-HA⋯OC, am⋯am	0.81(4)	2.23(4)	3.020(4)	166(3)
3_{t3} NP1-HP1⋯OC2, am(1)⋯am(2) NB-HB⋯OC3, am(3)⋯am(3) NP2-HP2⋯OC1, am(2)⋯am(1) OB-HOB⋯O1P1, OH⋯O=C OB-HOB⋯NB, intramolecular	0.87(3) 0.85(3) 0.84(3) 0.82 0.82	2.01(3) 2.13(3) 2.04(4) 2.09 2.59	2.879(6) 2.967(6) 2.813(6) 2.744(10) 2.973(10)	177(6) 168(6) 153(7) 136.0 109.7

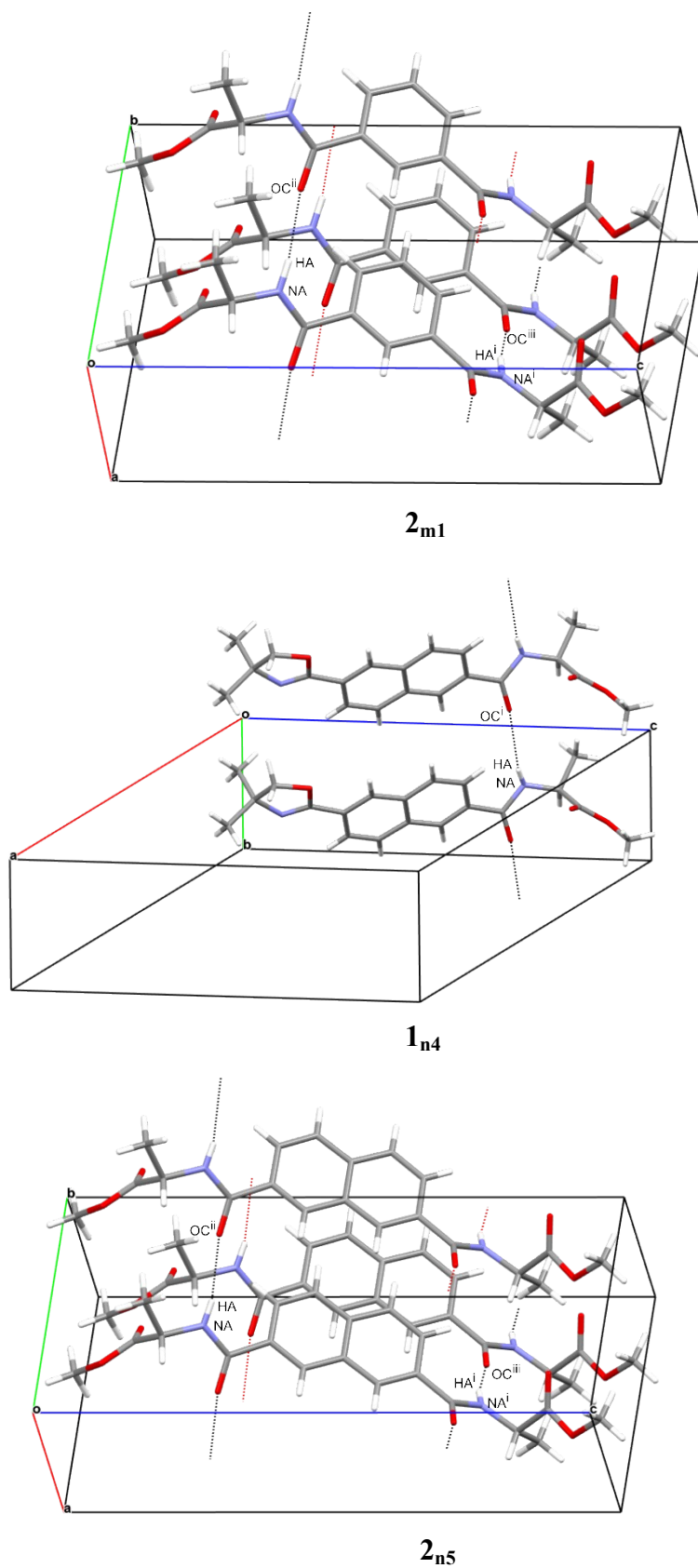
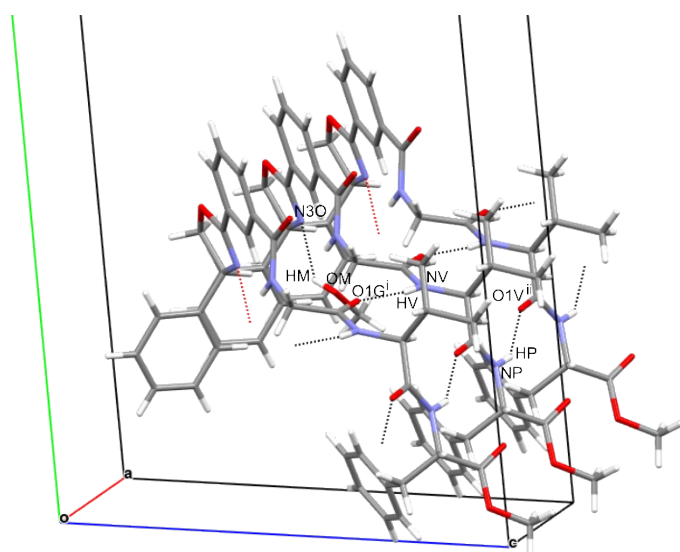
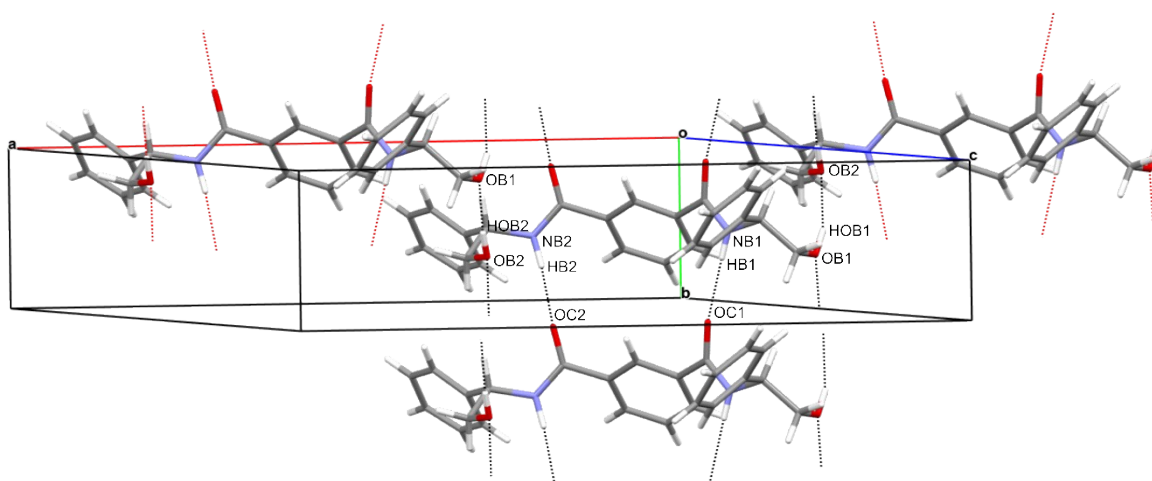


Figure S16. Hydrogen bonds (Table S9) for SCXRD determined structures. Symmetry codes: (i) x,y,z , (ii) x,y,z (iii) x,y,z for **2_{m1}** and **2_{n5}**, (i) x,y,z for **1_{n4}**.

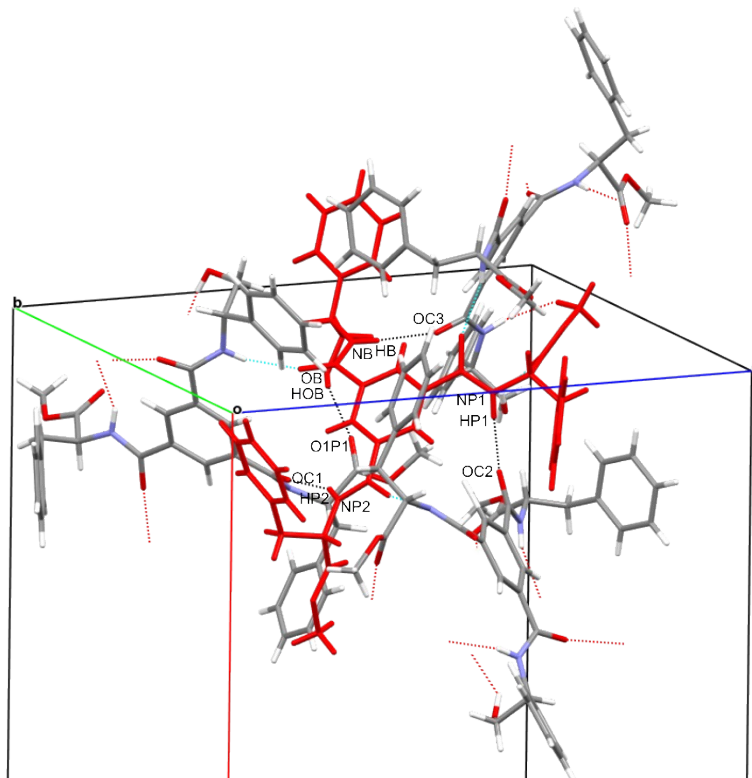


1_{m6}

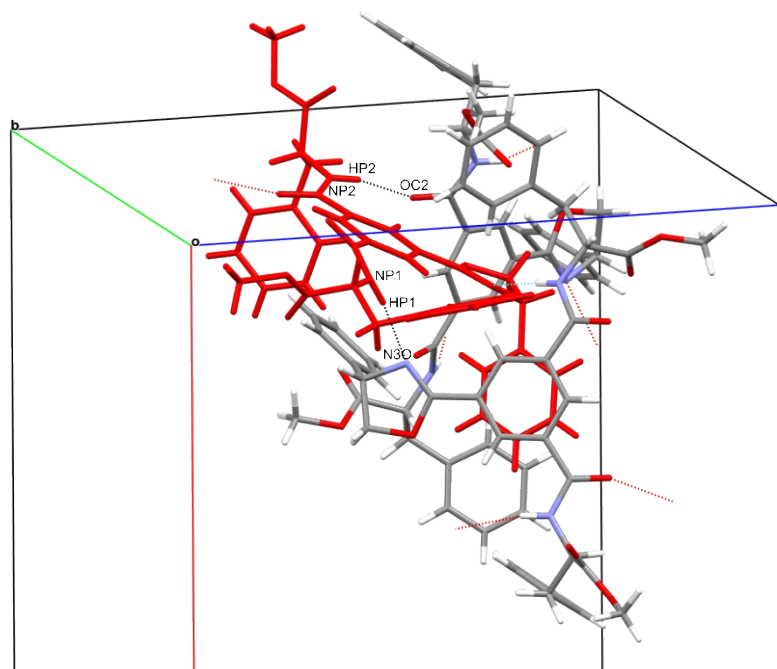


4_{m5}

Figure S17. Hydrogen bonds (Table S9) for SCXRD determined structures. Symmetry codes: (i) x,y,z, (ii) x,y,z **1_{m6}**, (i) x,y,z, (ii) x,y,z and (iii) x,y,z for **4_{m5}**.



3_{t3}



1_{t4}

Figure S18. Hydrogen bonds (Table S9) for SCXRD determined structures. Symmetry codes: (i) x,y,z , (ii) x,y,z and (iii) 3_{t3} ; (i) x,y,z and (ii) x,y,z for 1_{t4} . For reason of clarity, origin molecules are drawn in red.

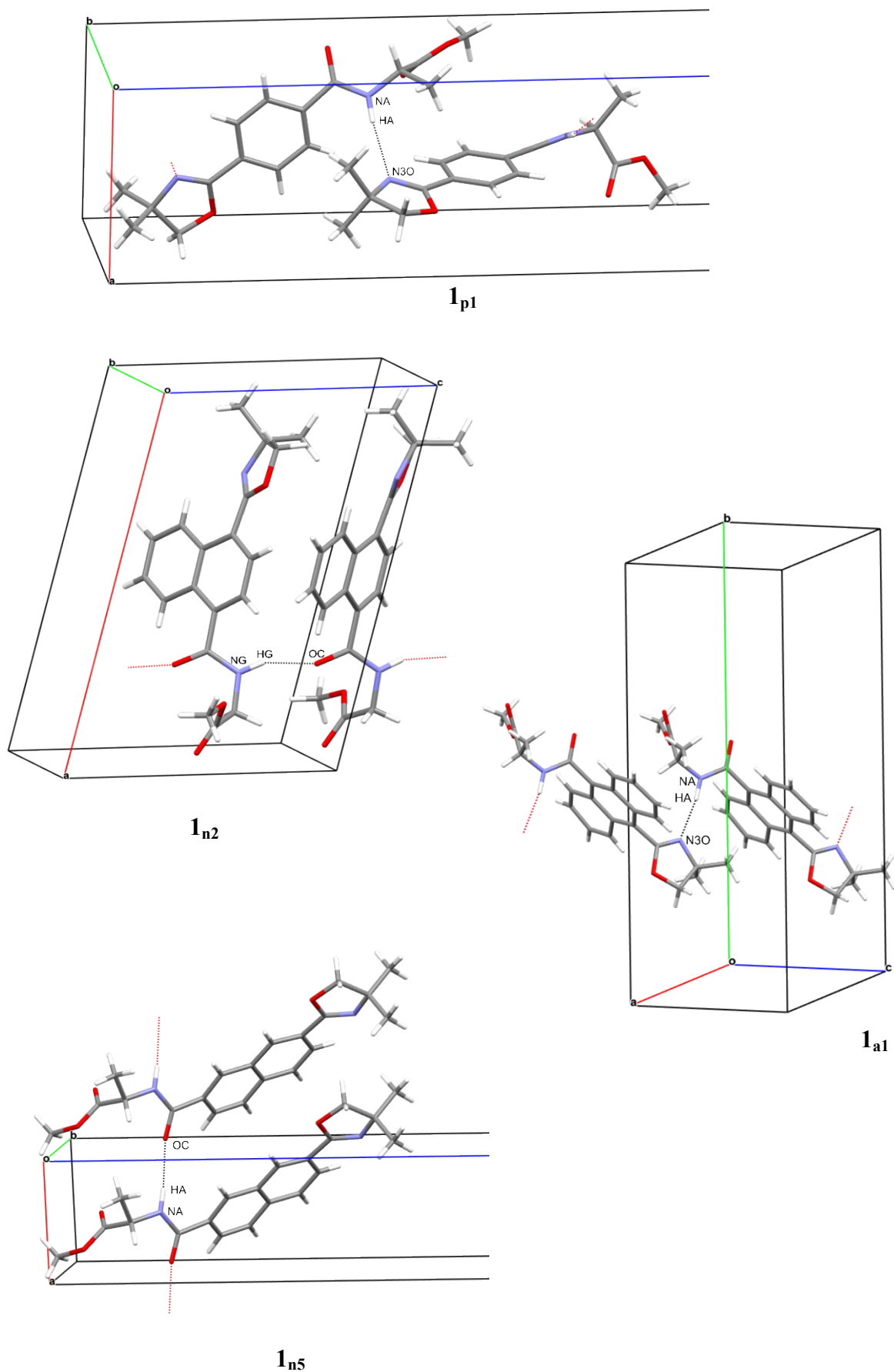


Figure S19. Hydrogen bonds (Table S9) for SCXRD determined structures. Symmetry codes: (i) x,y,z, for **1p1**; (i) x,y,z for **1n2**; (i) x,y,z for **1a1** and (i) x,y,z for **1n5**.

8. Computational calculations

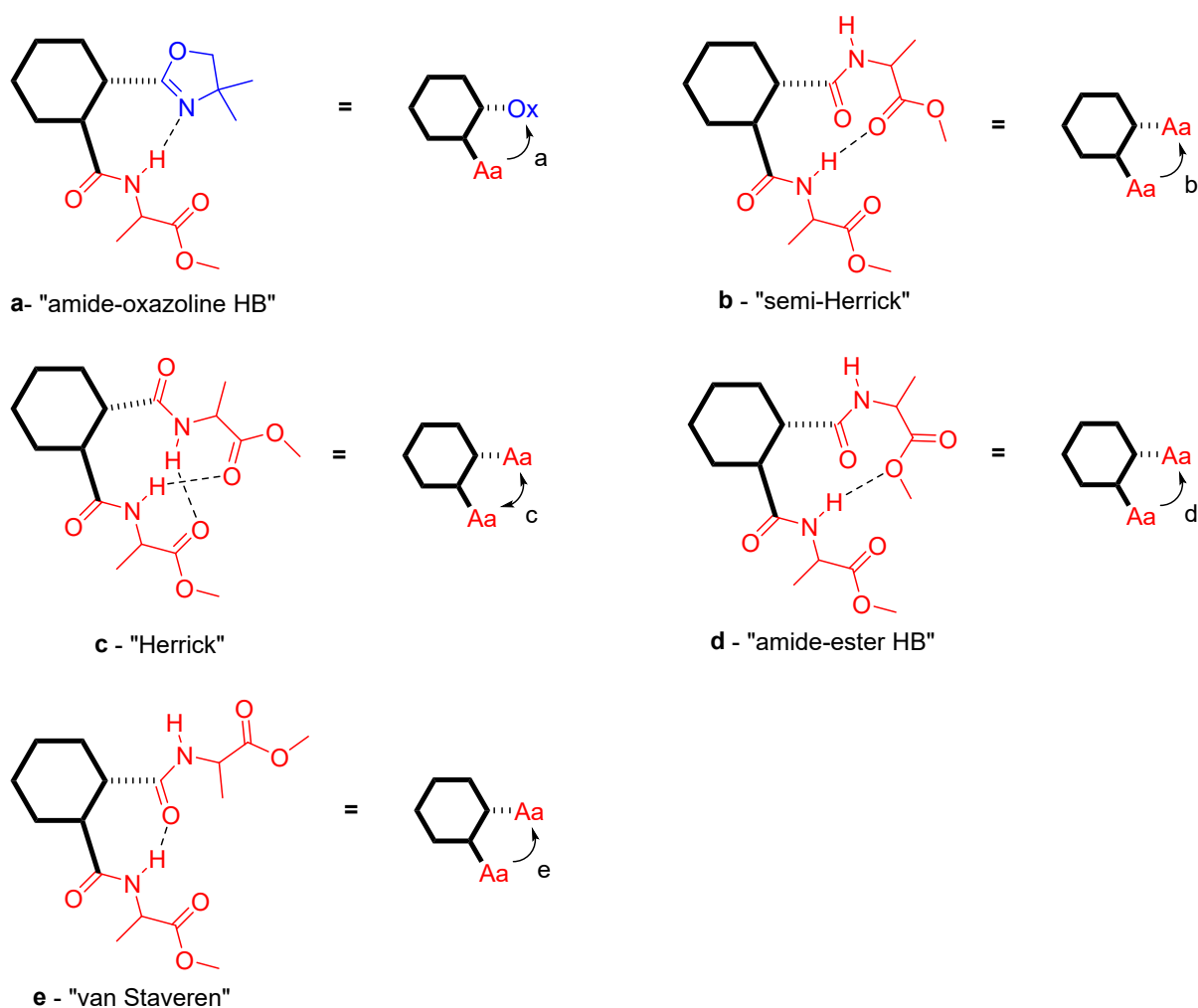


Figure S20. Types of hydrogen bonding: a-HB with oxazoline, b-semi-Herrick, c-Herrick, d-HB with the methoxy oxygen, e- van Staveren HB. The arrowhead indicates the direction of the amide proton donation (each arrowhead on an arrow represents one amide proton). BW = Boltzmann weight, Ox = oxazoline ring, Aa = amino acid, HBA-hydrogen bond acceptor, HBD-hydrogen bond donor.

'Anti' and 'syn' relative orientations of oxazoline rings in dimers are defined by making a 2D projection of the dimer vertically in relation to the benzene rings (Figure S21, bottom right). Then, the origin of the coordinate system is placed in the center of the benzene rings. We define angles $\alpha_{x1,x2}$ as the angles between two crossed lines of which the first line passes through heteroatom X1 and the origin of the coordinate system, while the second line passes through heteroatom X2 and the origin of the coordinate system. Numbers 1 and 2 in angle indices in Figure S21 are omitted as it is assumed that one indice denotes a heteroatom from one molecule from the dimer, and the other indice denotes the heteroatom from the other molecule from the dimer.

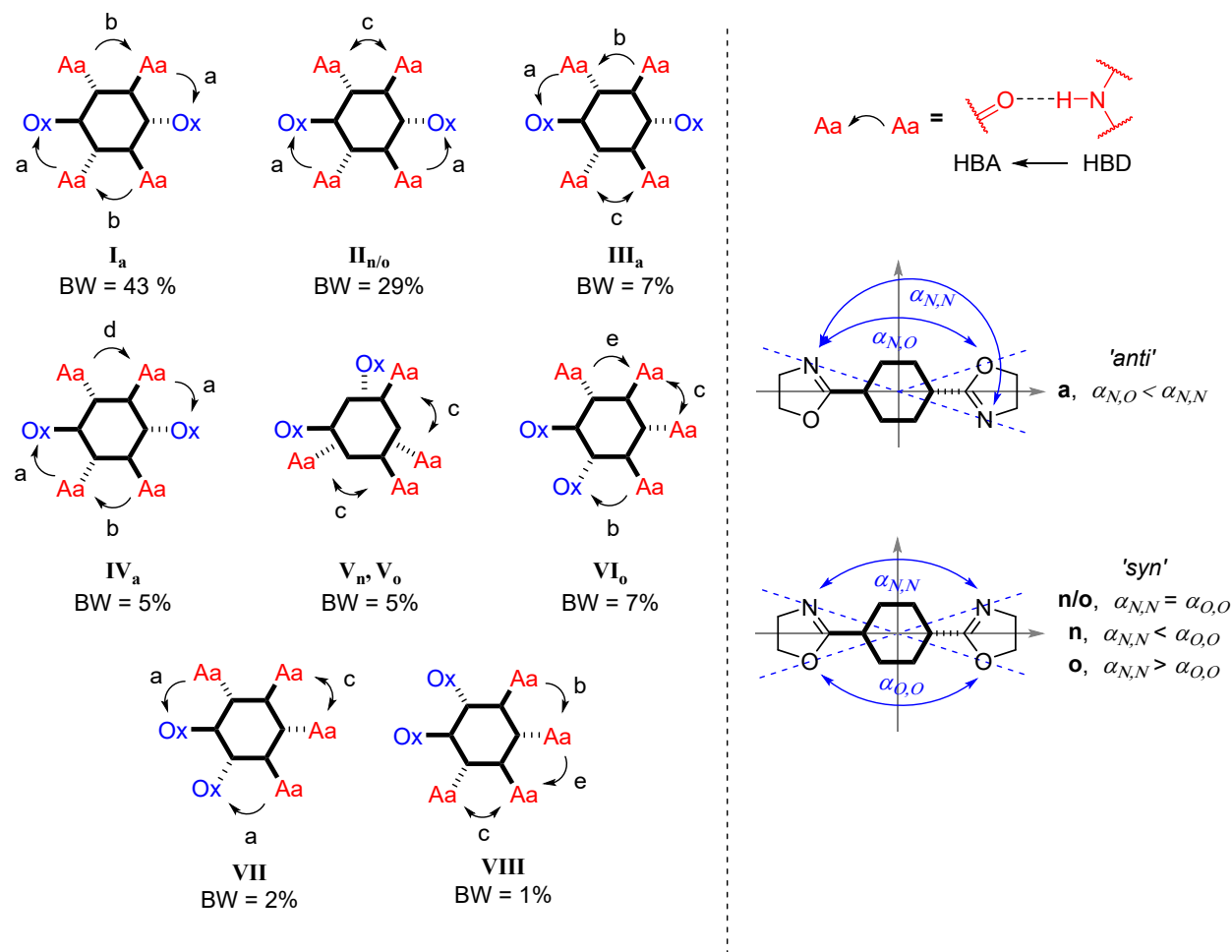


Figure S21. DFT structures of 1_{t1} dimers and corresponding hydrogen bonding with Boltzmann weight values greater than 1%

Table S10. Characteristic parameters of conformers obtained by CREST/CENSO protocol with Boltzmann weight contributions $\geq 1\%$.

Conf.	Type	Stacked oxazoline relative positions	Oxazoline relative orientation	Type of hydrogen bonded amide hydrogen atom				$\Delta G_{tot} / kcal/mol$ ₁	Boltzmann weight	BW SUM over type
				Herrick	van Staveren	semi-Herrick	oxazoline			
1	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	0	18	43
4	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	0.42	9	
6	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	0.75	5	
7	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	0.95	4	
8	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	0.98	3	
14	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	1.22	2	
18	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	1.59	1	
24	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	1.81	1	
27	I	1,4'	<i>anti</i>	/	/	++(3,6')	++(5,2')	2.02	1	
2	II	1,4'	<i>syn</i>	++(5,6')	/	/	++(3,2')	0.27	11	29
3	II	1,4'	<i>syn</i>	++(5,6')	/	/	++(3,2')	0.39	9	
9	II	1,4'	<i>syn</i>	++(5,6')	/	/	++(3,2')	1.05	3	
11	II	1,4'	<i>syn</i>	++(5,6')	/	/	++(3,2')	1.08	3	
20	II	1,4'	<i>syn</i>	++(5,6'-ester)	/	/	++(3,2')	1.63	1	
25	II	1,4'	<i>syn</i>	++(5,6')	/	/	++(3,2')	1.97	1	
28	II	1,4'	<i>syn</i>	++(5,6')	/	/	++(3,2')	2.08	1	
5	III	1,4'	<i>anti</i>	++(5,6')	/	+(3)	+(2')	0.59	7	
13	IV	1,4'	<i>anti</i>	/	/	++(3-ester,6')	++(5,2')	1.18	2	5
17	IV	1,4'	<i>anti</i>	/	/	++(3-ester,6')	++(5,2')	1.45	2	
21	IV	1,4'	<i>anti</i>	/	/	++(3-ester,6')	++(5,2')	1.67	1	
10	V	P1,3'	<i>syn(N away)</i>	++/++(3,4'/5,6')	/	/	/	1.07	3	5
19	V	P1,3'	<i>syn(N towards each other)</i>	++/++(3,4'/5,6')	/	/	/	1.6	1	
29	V	P1,3'	<i>syn(N away)</i>	++/++(3,4'/5,6'-ester)	/	/	/	2.11	1	
12	VI	P1,6'	<i>syn(N away)</i>	++(3,4')	+(2')	/	+(5)	1.14	3	7
15	VI	P1,6'	<i>syn(N away)</i>	++(3,4')	+(2')	/	+(5)	1.23	2	
16	VI	P1,6'	<i>syn(N away)</i>	++(3,4')	+(2')	/	+(5)	1.37	2	
22	VII	P1,6'	<i>syn(N away)</i>	++(3,4')	/	/	++(5,2')	1.69	1	2
26	VII	P1,6'	<i>syn(N away)</i>	++(3,4')	/	/	++(5,2')	1.99	1	
23	VIII	P1,6'	<i>anti</i>	++(3,4')	+(5)			1.75	1	1
Total BW SUM=									97	

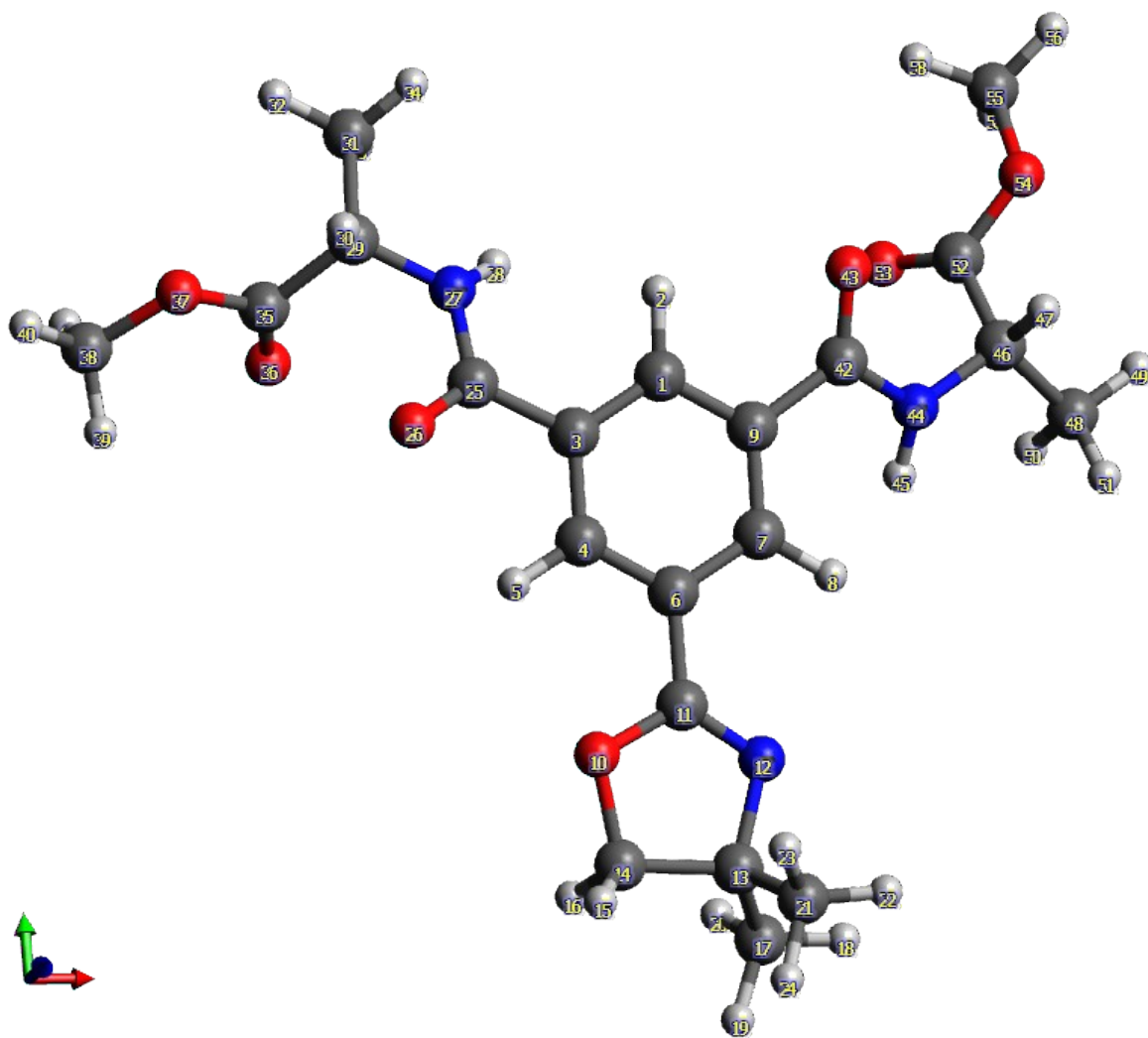


Figure S22. DFT optimized structure of lowest energy conformer of **1₁₁** (CONF1) with atom labels.

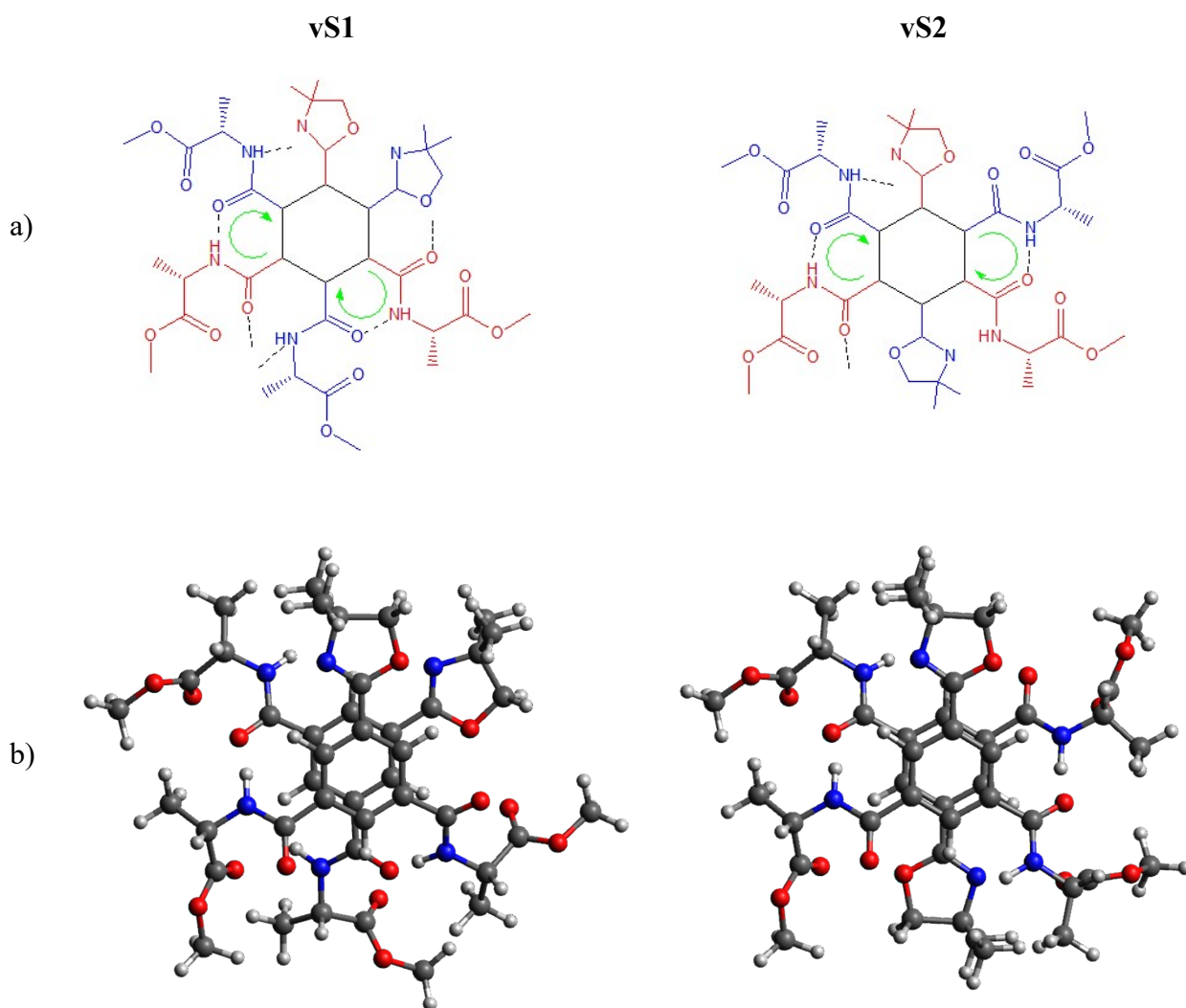


Figure S23. Schematic (a) and 3D “ball and stick” (b) representation of starting dimer models for 1_{tI} molecules (vS1 and vS2), obtained by the Avogadro software from two copies of the lowest energy conformer of 1_{tI} . Labels of atoms in one part of dimer is identical as in the monomer labelling (Figure S22), labels of the second part of dimer are accordingly increased for 58. (Atom 1 in one part is identical to atom 59 in second part of dimer, atom 2 is identical to atom 60, etc.) In order to mimic the van Staveren hydrogen bonding motifs for vS2 it was necessary to change torsion angles of two amide groups with respect to benzene rings, in particular C(1)-C(3)-C(25)-O(26) and C(65)-C(67)-C(100)-O(101) torsion angles from $\approx -150^\circ$ in CONF1 to $\approx 150^\circ$."

Table S11. Gibbs free energies (G) for monomer conformers of $\mathbf{1}_{\text{H}}$ at different temperatures.

CONF	$G [E_{\text{h}}]$ 233 K	$\Delta G [\text{kcal/mol}]$ 233 K	$G [E_{\text{h}}]$ 253 K	$\Delta G [\text{kcal/mol}]$ 253 K	$G [E_{\text{h}}]$ 273 K	$\Delta G [\text{kcal/mol}]$ 273 K
CONF1	-1506.7451897	0.00	-1506.7509090	0.00	-1506.7568827	0.00
CONF2	-1506.7447804	0.26	-1506.7505004	0.26	-1506.7564741	0.26
CONF3	-1506.7443329	0.54	-1506.7500520	0.54	-1506.7560282	0.54
CONF4	-1506.7441282	0.67	-1506.7498446	0.67	-1506.7558183	0.67
CONF5	-1506.7440714	0.70	-1506.7497902	0.70	-1506.7557630	0.70
CONF6	-1506.7438705	0.83	-1506.7495897	0.83	-1506.7555655	0.83
CONF7	-1506.7438737	0.83	-1506.7495891	0.83	-1506.7555606	0.83
CONF8	-1506.7438461	0.84	-1506.7495641	0.84	-1506.7555391	0.84
CONF9	-1506.7438008	0.87	-1506.7495135	0.88	-1506.7554789	0.88
CONF10	-1506.7436885	0.94	-1506.7494037	0.94	-1506.7553739	0.95
CONF11	-1506.7435880	1.01	-1506.7493114	1.00	-1506.7552884	1.00
CONF12	-1506.7434740	1.08	-1506.7491924	1.08	-1506.7551650	1.08
CONF13	-1506.7433286	1.17	-1506.7490587	1.16	-1506.7550463	1.15
CONF14	-1506.7433336	1.16	-1506.7490488	1.17	-1506.7550212	1.17
CONF15	-1506.7433183	1.17	-1506.7490368	1.17	-1506.7550122	1.17
CONF16	-1506.7432941	1.19	-1506.7490112	1.19	-1506.7549852	1.19
CONF17	-1506.7431828	1.26	-1506.7489051	1.26	-1506.7548807	1.26
CONF18	-1506.7429376	1.41	-1506.7486535	1.42	-1506.7546249	1.42
CONF19	-1506.7429500	1.41	-1506.7486599	1.41	-1506.7546248	1.42
CONF20	-1506.7428644	1.46	-1506.7485814	1.46	-1506.7545544	1.46
CONF21	-1506.7427883	1.51	-1506.7485189	1.50	-1506.7545056	1.49
CONF22	-1506.7424890	1.69	-1506.7482067	1.70	-1506.7541817	1.69
CONF23	-1506.7424365	1.73	-1506.7481663	1.72	-1506.7541534	1.71
CONF24	-1506.7424369	1.73	-1506.7481575	1.73	-1506.7541388	1.72
CONF25	-1506.7423019	1.81	-1506.7480287	1.81	-1506.7540121	1.80
CONF26	-1506.7422470	1.85	-1506.7479516	1.86	-1506.7539147	1.86
CONF27	-1506.7420906	1.94	-1506.7478268	1.93	-1506.7538185	1.92
CONF28	-1506.7421297	1.92	-1506.7478474	1.92	-1506.7538219	1.92
CONF29	-1506.7420525	1.97	-1506.7477827	1.96	-1506.7537689	1.95
CONF30	-1506.7418123	2.12	-1506.7475292	2.12	-1506.7535057	2.12
CONF31	-1506.7417322	2.17	-1506.7474637	2.16	-1506.7534522	2.15
CONF32	-1506.7416213	2.24	-1506.7473470	2.24	-1506.7533335	2.23
CONF33	-1506.7415279	2.30	-1506.7472467	2.30	-1506.7532217	2.30
CONF34	-1506.7415045	2.31	-1506.7472216	2.31	-1506.7531994	2.31
CONF35	-1506.7412985	2.44	-1506.7470144	2.44	-1506.7529888	2.44
CONF36	-1506.7409763	2.64	-1506.7466693	2.66	-1506.7526216	2.67
CONF37	-1506.7405839	2.89	-1506.7462834	2.90	-1506.7522415	2.91
CONF38	-1506.7404543	2.97	-1506.7461561	2.98	-1506.7521174	2.99
CONF39	-1506.7403628	3.03	-1506.7460853	3.03	-1506.7520681	3.02
CONF40	-1506.7403688	3.03	-1506.7460783	3.03	-1506.7520478	3.03
CONF41	-1506.7402343	3.11	-1506.7459260	3.13	-1506.7518759	3.14
CONF42	-1506.7400646	3.22	-1506.7457682	3.23	-1506.7517318	3.23
CONF43	-1506.7400251	3.24	-1506.7457362	3.25	-1506.7517070	3.25
CONF44	-1506.7399567	3.28	-1506.7456541	3.30	-1506.7516092	3.31
CONF45	-1506.7398280	3.36	-1506.7455551	3.36	-1506.7515418	3.35
CONF46	-1506.7399157	3.31	-1506.7456008	3.33	-1506.7515436	3.35
CONF47	-1506.7397351	3.42	-1506.7454458	3.43	-1506.7514167	3.43

Table S11(continuation). Gibbs free energies (G) for monomer conformers of $\mathbf{1}_{\text{H}}$ at different temperatures.

CONF	$G [E_{\text{h}}]$ 293 K	ΔG [kcal/mol] 293 K	$G [E_{\text{h}}]$ 298 K	ΔG [kcal/mol] 298 K	$G [E_{\text{h}}]$ 313 K	ΔG [kcal/mol] 313 K
CONF1	-1506.7631071	0.00	-1506.7647019	0.00	-1506.7695789	0.00
CONF2	-1506.7626979	0.26	-1506.7642926	0.26	-1506.7691686	0.26
CONF3	-1506.7622573	0.53	-1506.7638537	0.53	-1506.7687361	0.53
CONF4	-1506.7620453	0.67	-1506.7636413	0.67	-1506.7685222	0.66
CONF5	-1506.7619860	0.70	-1506.7635804	0.70	-1506.7684560	0.70
CONF6	-1506.7617941	0.82	-1506.7633904	0.82	-1506.7682722	0.82
CONF7	-1506.7617842	0.83	-1506.7633792	0.83	-1506.7682568	0.83
CONF8	-1506.7617671	0.84	-1506.7633633	0.84	-1506.7682448	0.84
CONF9	-1506.7616932	0.89	-1506.7632853	0.89	-1506.7681535	0.89
CONF10	-1506.7615954	0.95	-1506.7631896	0.95	-1506.7680648	0.95
CONF11	-1506.7615154	1.00	-1506.7631108	1.00	-1506.7679892	1.00
CONF12	-1506.7613881	1.08	-1506.7629827	1.08	-1506.7678585	1.08
CONF13	-1506.7612875	1.14	-1506.7628870	1.14	-1506.7677788	1.13
CONF14	-1506.7612468	1.17	-1506.7628424	1.17	-1506.7677223	1.17
CONF15	-1506.7612408	1.17	-1506.7628370	1.17	-1506.7677190	1.17
CONF16	-1506.7612121	1.19	-1506.7628079	1.19	-1506.7676886	1.19
CONF17	-1506.7611058	1.26	-1506.7627007	1.26	-1506.7675773	1.26
CONF18	-1506.7608479	1.42	-1506.7624426	1.42	-1506.7673195	1.42
CONF19	-1506.7608408	1.42	-1506.7624337	1.42	-1506.7673047	1.43
CONF20	-1506.7607796	1.46	-1506.7623749	1.46	-1506.7672537	1.46
CONF21	-1506.7607446	1.48	-1506.7623434	1.48	-1506.7672326	1.47
CONF22	-1506.7604101	1.69	-1506.7620064	1.69	-1506.7668884	1.69
CONF23	-1506.7603938	1.70	-1506.7619930	1.70	-1506.7668840	1.69
CONF24	-1506.7603767	1.71	-1506.7619759	1.71	-1506.7668677	1.70
CONF25	-1506.7602482	1.79	-1506.7618463	1.79	-1506.7667336	1.79
CONF26	-1506.7601324	1.87	-1506.7617261	1.87	-1506.7666012	1.87
CONF27	-1506.7600618	1.91	-1506.7616616	1.91	-1506.7665537	1.90
CONF28	-1506.7600495	1.92	-1506.7616456	1.92	-1506.7665268	1.92
CONF29	-1506.7600073	1.95	-1506.7616059	1.94	-1506.7664945	1.94
CONF30	-1506.7597377	2.11	-1506.7613352	2.11	-1506.7662218	2.11
CONF31	-1506.7596938	2.14	-1506.7612933	2.14	-1506.7661851	2.13
CONF32	-1506.7595766	2.22	-1506.7611770	2.21	-1506.7660727	2.20
CONF33	-1506.7594492	2.30	-1506.7610451	2.29	-1506.7659257	2.29
CONF34	-1506.7594338	2.31	-1506.7610320	2.30	-1506.7659211	2.30
CONF35	-1506.7592177	2.44	-1506.7608143	2.44	-1506.7656977	2.44
CONF36	-1506.7588291	2.68	-1506.7604204	2.69	-1506.7652883	2.69
CONF37	-1506.7584543	2.92	-1506.7600468	2.92	-1506.7649182	2.92
CONF38	-1506.7583341	3.00	-1506.7599277	3.00	-1506.7648027	3.00
CONF39	-1506.7583069	3.01	-1506.7599061	3.01	-1506.7647982	3.00
CONF40	-1506.7582731	3.03	-1506.7598690	3.03	-1506.7647508	3.03
CONF41	-1506.7580800	3.15	-1506.7596703	3.16	-1506.7645350	3.17
CONF42	-1506.7579514	3.24	-1506.7595459	3.24	-1506.7644234	3.24
CONF43	-1506.7579335	3.25	-1506.7595296	3.25	-1506.7644122	3.24
CONF44	-1506.7578180	3.32	-1506.7594095	3.32	-1506.7642773	3.33
CONF45	-1506.7577841	3.34	-1506.7593842	3.34	-1506.7642785	3.33
CONF46	-1506.7577402	3.37	-1506.7593286	3.37	-1506.7641872	3.38
CONF47	-1506.7576436	3.43	-1506.7592399	3.43	-1506.7641230	3.42

Table S12. Gibbs free energies (G) for dimer conformers of $\mathbf{1}_{II}$ at different temperatures.

CONF (dimer)	$G [E_h]$ 233 K	ΔG [kcal/mol] 233 K	$G [E_h]$ 253 K	ΔG [kcal/mol] 253 K	$G [E_h]$ 273 K	ΔG [kcal/mol] 273 K
CONF1	-1506.7451897	0.00	-1506.7509090	0.00	-1506.7568827	0.00
CONF2	-1506.7447804	0.28	-1506.7505004	0.28	-1506.7564741	0.28
CONF3	-1506.7443329	0.29	-1506.7500520	0.32	-1506.7560282	0.35
CONF4	-1506.7441282	0.42	-1506.7498446	0.42	-1506.7558183	0.42
CONF5	-1506.7440714	0.59	-1506.7497902	0.58	-1506.7557630	0.58
CONF6	-1506.7438705	0.77	-1506.7495897	0.77	-1506.7555655	0.76
CONF7	-1506.7438737	0.88	-1506.7495891	0.90	-1506.7555606	0.92
CONF8	-1506.7438461	0.92	-1506.7495641	0.94	-1506.7555391	0.96
CONF9	-1506.7438008	1.02	-1506.7495135	1.03	-1506.7554789	1.04
CONF10	-1506.7436885	0.99	-1506.7494037	1.01	-1506.7553739	1.04
CONF11	-1506.7435880	1.12	-1506.7493114	1.11	-1506.7552884	1.10
CONF12	-1506.7434740	1.16	-1506.7491924	1.15	-1506.7551650	1.15
CONF13	-1506.7433286	1.14	-1506.7490587	1.15	-1506.7550463	1.17
CONF14	-1506.7433336	1.21	-1506.7490488	1.21	-1506.7550212	1.21
CONF15	-1506.7433183	1.18	-1506.7490368	1.19	-1506.7550122	1.21
CONF16	-1506.7432941	1.31	-1506.7490112	1.32	-1506.7549852	1.34
CONF17	-1506.7431828	1.41	-1506.7489051	1.42	-1506.7548807	1.43
CONF18	-1506.7429376	1.50	-1506.7486535	1.53	-1506.7546249	1.55
CONF19	-1506.7429500	1.42	-1506.7486599	1.47	-1506.7546248	1.53
CONF20	-1506.7428644	1.69	-1506.7485814	1.67	-1506.7545544	1.66
CONF21	-1506.7427883	1.61	-1506.7485189	1.62	-1506.7545056	1.64
CONF22	-1506.7424890	1.66	-1506.7482067	1.66	-1506.7541817	1.67
CONF23	-1506.7424365	1.74	-1506.7481663	1.74	-1506.7541534	1.74
CONF24	-1506.7424369	1.74	-1506.7481575	1.76	-1506.7541388	1.78
CONF25	-1506.7423019	1.89	-1506.7480287	1.91	-1506.7540121	1.94
CONF26	-1506.7422470	1.98	-1506.7479516	1.98	-1506.7539147	1.98
CONF27	-1506.7420906	1.97	-1506.7478268	1.98	-1506.7538185	1.99
CONF28	-1506.7421297	2.06	-1506.7478474	2.06	-1506.7538219	2.07
CONF29	-1506.7420525	2.09	-1506.7477827	2.09	-1506.7537689	2.10
CONF30	-1506.7418123	2.05	-1506.7475292	2.08	-1506.7535057	2.11
CONF31	-1506.7417322	2.18	-1506.7474637	2.20	-1506.7534522	2.22
CONF32	-1506.7416213	2.25	-1506.7473470	2.25	-1506.7533335	2.25
CONF33	-1506.7415279	2.31	-1506.7472467	2.36	-1506.7532217	2.40
CONF34	-1506.7415045	2.34	-1506.7472216	2.37	-1506.7531994	2.41
CONF35	-1506.7412985	2.43	-1506.7470144	2.44	-1506.7529888	2.45
CONF36	-1506.7409763	2.44	-1506.7466693	2.46	-1506.7526216	2.48
CONF37	-1506.7405839	2.62	-1506.7462834	2.64	-1506.7522415	2.65
CONF38	-1506.7404543	2.82	-1506.7461561	2.86	-1506.7521174	2.89
CONF39	-1506.7403628	2.86	-1506.7460853	2.89	-1506.7520681	2.93
CONF40	-1506.7403688	3.11	-1506.7460783	3.17	-1506.7520478	3.23
CONF41	-1506.7402343	3.24	-1506.7459260	3.31	-1506.7518759	3.37

Table S12. Gibbs free energies (G) for dimer conformers of $\mathbf{1}_{\text{fl}}$ at different temperatures. (continuation)

CONF	G [E_{h}] 293 K	ΔG [kcal/mol] 293 K	G [E_{h}] 298 K	ΔG [kcal/mol] 298 K	G [E_{h}] 313 K	ΔG [kcal/mol] 313 K
CONF1	-1506.7631071	0.00	-1506.7647019	0.00	-1506.7695789	0.00
CONF2	-1506.7626979	0.27	-1506.7642926	0.27	-1506.7691686	0.27
CONF3	-1506.7622573	0.39	-1506.7638537	0.39	-1506.7687361	0.42
CONF4	-1506.7620453	0.42	-1506.7636413	0.42	-1506.7685222	0.42
CONF5	-1506.7619860	0.59	-1506.7635804	0.59	-1506.7684560	0.59
CONF6	-1506.7617941	0.75	-1506.7633904	0.75	-1506.7682722	0.74
CONF7	-1506.7617842	0.95	-1506.7633792	0.95	-1506.7682568	0.98
CONF8	-1506.7617671	0.98	-1506.7633633	0.98	-1506.7682448	0.99
CONF9	-1506.7616932	1.05	-1506.7632853	1.05	-1506.7681535	1.06
CONF10	-1506.7615954	1.06	-1506.7631896	1.07	-1506.7680648	1.09
CONF11	-1506.7615154	1.09	-1506.7631108	1.08	-1506.7679892	1.08
CONF12	-1506.7613881	1.14	-1506.7629827	1.14	-1506.7678585	1.15
CONF13	-1506.7612875	1.18	-1506.7628870	1.18	-1506.7677788	1.20
CONF14	-1506.7612468	1.22	-1506.7628424	1.22	-1506.7677223	1.22
CONF15	-1506.7612408	1.22	-1506.7628370	1.23	-1506.7677190	1.24
CONF16	-1506.7612121	1.36	-1506.7628079	1.37	-1506.7676886	1.38
CONF17	-1506.7611058	1.44	-1506.7627007	1.45	-1506.7675773	1.45
CONF18	-1506.7608479	1.58	-1506.7624426	1.59	-1506.7673195	1.61
CONF19	-1506.7608408	1.59	-1506.7624337	1.60	-1506.7673047	1.65
CONF20	-1506.7607796	1.64	-1506.7623749	1.63	-1506.7672537	1.62
CONF21	-1506.7607446	1.66	-1506.7623434	1.67	-1506.7672326	1.68
CONF22	-1506.7604101	1.69	-1506.7620064	1.69	-1506.7668884	1.70
CONF23	-1506.7603938	1.75	-1506.7619930	1.75	-1506.7668840	1.75
CONF24	-1506.7603767	1.80	-1506.7619759	1.81	-1506.7668677	1.83
CONF25	-1506.7602482	1.96	-1506.7618463	1.97	-1506.7667336	1.99
CONF26	-1506.7601324	1.99	-1506.7617261	1.99	-1506.7666012	2.00
CONF27	-1506.7600618	2.01	-1506.7616616	2.02	-1506.7665537	2.03
CONF28	-1506.7600495	2.07	-1506.7616456	2.08	-1506.7665268	2.09
CONF29	-1506.7600073	2.11	-1506.7616059	2.11	-1506.7664945	2.12
CONF30	-1506.7597377	2.14	-1506.7613352	2.15	-1506.7662218	2.17
CONF31	-1506.7596938	2.24	-1506.7612933	2.25	-1506.7661851	2.26
CONF32	-1506.7595766	2.25	-1506.7611770	2.25	-1506.7660727	2.25
CONF33	-1506.7594492	2.44	-1506.7610451	2.45	-1506.7659257	2.48
CONF34	-1506.7594338	2.45	-1506.7610320	2.46	-1506.7659211	2.49
CONF35	-1506.7592177	2.47	-1506.7608143	2.47	-1506.7656977	2.48
CONF36	-1506.7588291	2.51	-1506.7604204	2.52	-1506.7652883	2.54
CONF37	-1506.7584543	2.67	-1506.7600468	2.68	-1506.7649182	2.70
CONF38	-1506.7583341	2.93	-1506.7599277	2.94	-1506.7648027	2.97
CONF39	-1506.7583069	2.97	-1506.7599061	2.98	-1506.7647982	3.01
CONF40	-1506.7582731	3.29	-1506.7598690	3.30	-1506.7647508	3.35
CONF41	-1506.7580800	3.45	-1506.7596703	3.46	-1506.7645350	3.52

Table S13. Boltzmann averaged free energy G of ensemble of 47 monomers of $\mathbf{1}_{\text{H}}$ (E_{h}):

T / K	$\langle E_{\text{gas}} \rangle$	$\langle G_{\text{mRRHO}} \rangle$	$\langle G_{\text{solv}} \rangle$	$\langle G \rangle$
233	-1507.1118680	0.4091322	-0.0414216	-1506.7441574
253	-1507.1117665	0.4034176	-0.0414428	-1506.7497917
273	-1507.1116754	0.3974478	-0.0414598	-1506.7556874
293	-1507.1115927	0.3912264	-0.0414738	-1506.7618401
298	-1507.1115732	0.3896321	-0.0414769	-1506.7634180
313	-1507.1115171	0.3847563	-0.0414854	-1506.7682461

Table S14. Boltzmann averaged free energy G of ensemble of 41 dimer structures of $\mathbf{1}_{\text{H}}$ (E_{h}):

T / K	$\langle E_{\text{gas}} \rangle$	$\langle G_{\text{mRRHO}} \rangle$	$\langle G_{\text{solv}} \rangle$	$\langle G \rangle$
233	-3014.2977470	0.8458926	-0.0506042	-3013.5024586
253	-3014.2975688	0.8363840	-0.0506951	-3013.5118799
273	-3014.2974074	0.8263455	-0.0507776	-3013.5218394
293	-3014.2972638	0.8157857	-0.0508500	-3013.5323280
298	-3014.2972306	0.8130652	-0.0508666	-3013.5350319
313	-3014.2971361	0.8047120	-0.0509137	-3013.5433377

Table S15. Gibbs free energy of formation as calculated from equation $\Delta G = G(\text{dimer}) - 2 \times G(\text{monomer})$.

T / K	$\Delta G / [E_{\text{h}}]$	$\Delta G / \text{kcal mol}^{-1}$
233	0.014144	8.88
253	0.012297	7.72
273	0.010465	6.57
293	0.008648	5.43
298	0.008196	5.14
313	0.006846	4.30

Table S16. Experimental and calculated NMR parameters for monomer ensemble of 1_{t1} at different temperatures.

Part 1: ^1H NMR experimental shifts and calculated shieldings and shifts*

Lab. (^1H)	shift exp. (233 K)	shield. averaged (233 K)	shift calc. (233 K)	shift exp. (253 K)	shield. averaged (253 K)	shift calc. (253 K)	shift exp. (273 K)	shield. averaged (273 K)	shift calc. (273 K)
2	8.220	22.483	8.195	8.260	22.493	8.229	8.290	22.492	8.264
5	8.190	22.252	8.397	8.240	22.254	8.442	8.290	22.254	8.478
15	4.180	26.892	4.314	4.170	26.892	4.316	4.161	26.894	4.315
16	4.160	27.080	4.149	4.160	27.082	4.147	4.159	27.080	4.149
18	1.420	29.970	1.606	1.410	29.971	1.577	1.410	29.972	1.554
22	1.530	30.049	1.537	1.490	30.049	1.508	1.460	30.048	1.486
28	8.120	24.388	6.518	7.870	24.385	6.546	7.610	24.382	6.569
30	4.860	26.851	4.351	4.850	26.854	4.349	4.840	26.855	4.350
32	1.610	29.856	1.706	1.590	29.855	1.680	1.570	29.855	1.659
39	3.820	27.537	3.747	3.810	27.542	3.737	3.810	27.544	3.732
		MAE(C-H):	0.13881			0.13778			0.13402
		AE(N-H):	1.60189			1.32362			1.04140

Lab. (^1H)	shift exp. (293 K)	shield. averaged (293 K)	shift calc. (293 K)	shift exp. (298 K)	shield. averaged (298 K)	shift calc. (298 K)	shift exp. (313 K)	shield. averaged (313 K)	shift calc. (313 K)
2	8.310	22.492	8.299	8.320	22.492	8.308	8.330	22.491	8.325
5	8.330	22.254	8.513	8.340	22.254	8.523	8.370	22.255	8.540
15	4.150	26.987	4.240	4.150	26.987	4.239	4.140	26.987	4.241
18	1.404	29.975	1.542	1.403	29.975	1.535	1.420	29.976	1.525
22	1.440	30.044	1.479	1.434	30.044	1.472	1.399	30.043	1.464
28	7.380	24.370	6.603	7.330	24.369	6.608	7.170	24.368	6.620
30	4.830	26.849	4.364	4.830	26.849	4.364	4.820	26.849	4.366
32	1.560	29.854	1.651	1.550	29.854	1.644	1.550	29.854	1.636
39	3.800	27.544	3.737	3.800	27.544	3.735	3.790	27.545	3.734
		MAE(C-H):	0.13506			0.13488			0.13024
		AE(N-H):	0.77710			0.72172			0.54950

* Shift calc. is obtained by using linear regression equation between shield. averaged (x) and shift exp (y), see Figure S22-S27. Linear regression equation is calculated for all ^1H nuclei bonded to carbon atoms, independently for each T . MAE(C-H) is the mean absolute error for ^1H nuclei with respect to the calculated linear regression equation. AE(N-H) is absolute error for ^1H amide nuclei with respect to the calculated linear regression equation.

Part 2: Experimental and calculated ^1H - ^1H J couplings**

T (K)	2-bond		3-bond				4-bond							
	$J(15-16)$	exp.	$J(28-30)$	exp.	$J(15-16)$	exp.	$J(2-5)$	exp.	$J(5-8)$	exp.	$J(15-18)$	$J(16-18)$	$J(18-22)$	$J(28-32)$
233	-9.00	7.9	5.94	7.9	7.04	7.4	0.90		0.75		0.13	-0.10	0.24	-0.29
253	-9.00	7.8	5.84	7.6	7.13	7.4	0.89	1.7	0.75	1.7	0.13	-0.10	0.24	-0.29
273	-9.00		5.81	7.6	7.13	7.3	0.89		0.75		0.13	-0.10	0.24	-0.29
293	-9.00		5.76	7.5	7.13	7.3	0.89	1.7	0.75	1.7	0.13	-0.10	0.24	-0.28
298	-9.00		5.76	7.5	7.13	7.3	0.89	1.7	0.75	1.7	0.13	-0.10	0.24	-0.28
313	-9.00		5.74	7.5	7.13	7.2	0.89	1.7	0.75	1.7	0.13	-0.10	0.24	-0.28

** Experimentally observed J couplings and their corresponding calculated values (left from exp. values) are shaded grey.

Part 3: ^{13}C NMR experimental shifts and calculated shieldings and shifts for $T = 298\text{ K}$ ***

Lab. (^{13}C)	shield. averaged (293 K)	shift exp. (293 K)	shift calc. (293 K)
1	44.735	129.679	129.213
3	39.860	134.903	133.767
4	43.546	129.083	130.324
6	46.157	128.334	127.884
11	10.697	161.781	161.008
13	110.184	67.472	68.077
14	98.519	79.352	78.973
17	154.933	27.002	26.277
25	6.169	166.965	165.238
29	128.827	48.850	50.663
31	167.004	15.823	15.001
35	-4.962	173.230	175.635
38	127.566	51.445	51.840
		MAE:	0.99512

*** Shift calc. is obtained by using linear regression equation between shield. averaged (x) and shift exp (y), see Figure S25. Linear regression equation is calculated for all ^{13}C nuclei. MAE is the mean absolute error for ^{13}C nuclei with respect to the calculated linear regression equation.

Table S17. Experimental and calculated NMR parameters for dimer ensemble of $\mathbf{1}_{t1}$ at different temperatures.

Part 1: ^1H NMR experimental shifts and calculated shieldings and shifts*

Lab. (^1H)	shift exp. (233 K)	shield. averaged (233 K)	shift calc. (233 K)	shift exp. (253 K)	shield. averaged (253 K)	shift calc. (253 K)	shift exp. (273 K)	shield. averaged (273 K)	shift calc. (273 K)
2	8.220	22.535	8.416	8.260	22.549	8.447	8.290	22.563	8.473
5	8.190	22.961	8.011	8.240	22.957	8.055	8.290	22.953	8.095
15	4.180	26.917	4.256	4.170	26.917	4.256	4.161	26.917	4.260
16	4.160	26.990	4.187	4.160	26.994	4.183	4.159	26.993	4.187
18	1.420	29.916	1.409	1.410	29.918	1.378	1.410	29.917	1.357
22	1.530	29.714	1.602	1.490	29.714	1.574	1.460	29.720	1.548
28	8.120	22.276	8.662	7.870	22.280	8.705	7.610	22.282	8.745
30	4.860	26.407	4.741	4.850	26.406	4.747	4.840	26.409	4.752
32	1.610	29.728	1.588	1.590	29.731	1.557	1.570	29.732	1.536
39	3.820	27.415	3.784	3.810	27.415	3.779	3.810	27.415	3.778
		MAE(C-H):	0.08180			0.08488			0.08862
		AE(N-H):	0.54169			0.83490			1.13510

Lab. (^1H)	shift exp. (293 K)	shield. averaged (293 K)	shift calc. (293 K)	shift exp. (298 K)	shield. averaged (298 K)	shift calc. (298 K)	shift exp. (313 K)	shield. averaged (313 K)	shift calc. (313 K)
2	8.310	22.567	8.503	8.320	22.568	8.513	8.330	22.569	8.525
5	8.330	22.951	8.129	8.340	22.951	8.138	8.370	22.950	8.152
15	4.150	26.955	4.231	4.150	26.955	4.232	4.140	26.955	4.231
18	1.404	29.917	1.348	1.403	29.917	1.342	1.420	29.917	1.330
22	1.440	29.722	1.537	1.434	29.723	1.532	1.399	29.724	1.519
28	7.380	22.283	8.779	7.330	22.283	8.790	7.170	22.284	8.804
30	4.830	26.409	4.763	4.830	26.409	4.765	4.820	26.408	4.766
32	1.560	29.733	1.527	1.550	29.733	1.522	1.550	29.733	1.510
39	3.800	27.415	3.784	3.800	27.415	3.783	3.790	27.415	3.780
		MAE(C-H):	0.09311			0.09320			0.10213
		AE(N-H):	1.39946			1.46029			1.63419

* Shift calc. is obtained by using linear regression equation between shield. averaged (x) and shift exp (y), see Figure S22-Figure S27. Linear regression equation is calculated for all ^1H nuclei bonded to carbon atoms, independently for each T . MAE(C-H) is the mean absolute error for ^1H nuclei with respect to the calculated linear regression equation. AE(N-H) is absolute error for ^1H amide nuclei with respect to the calculated linear regression equation.

Part 2: Experimental and calculated ^1H - ^1H J couplings**

T (K)	2-bond		3-bond				4-bond							
	$J(15-16)$	exp.	$J(28-30)$	exp.	$J(15-16)$	exp.	$J(2-5)$	exp.	$J(5-8)$	exp.	$J(15-18)$	$J(16-18)$	$J(18-22)$	$J(28-32)$
233	-8.49	7.9	7.62	7.9	6.82	7.4	0.87		0.64		0.12	-0.08	0.23	-0.21
253	-8.88	7.8	7.89	7.6	7.12	7.4	0.90	1.7	0.67	1.7	0.12	-0.09	0.24	-0.21
273	-8.90		7.85	7.6	7.12	7.3	0.90		0.68		0.12	-0.09	0.24	-0.22
293	-8.90		7.85	7.5	7.12	7.3	0.90	1.7	0.68	1.7	0.12	-0.09	0.24	-0.22
298	-8.90		7.85	7.5	7.12	7.3	0.90	1.7	0.68	1.7	0.12	-0.09	0.24	-0.22
313	-8.90		7.85	7.5	7.12	7.2	0.90	1.7	0.68	1.7	0.13	-0.09	0.24	-0.22

** Experimentally observed J couplings and their corresponding calculated values (left from exp. values) are shaded grey.

Part 3: ^{13}C NMR experimental shifts and calculated shieldings and shifts for $T = 298\text{ K}$ ***

Lab. (^{13}C)	shield. averaged (293 K)	shift exp. (293 K)	shift calc. (293 K)
1	43.181	129.679	130.235
3	40.622	134.903	132.612
4	43.442	129.083	129.993
6	46.960	128.334	126.724
11	8.045	161.781	162.876
13	110.004	67.472	68.156
14	98.491	79.352	78.852
17	155.372	27.002	26.010
25	6.772	166.965	164.059
29	129.526	48.850	50.020
31	166.668	15.823	15.515
35	-6.212	173.230	176.121
38	126.685	51.445	52.660
		MAE:	1.31755

*** Shift calc. is obtained by using linear regression equation between shield. averaged (x) and shift exp (y), see Figure S27. Linear regression equation is calculated for all ^{13}C nuclei. MAE is the mean absolute error for ^{13}C nuclei with respect to the calculated linear regression equation.

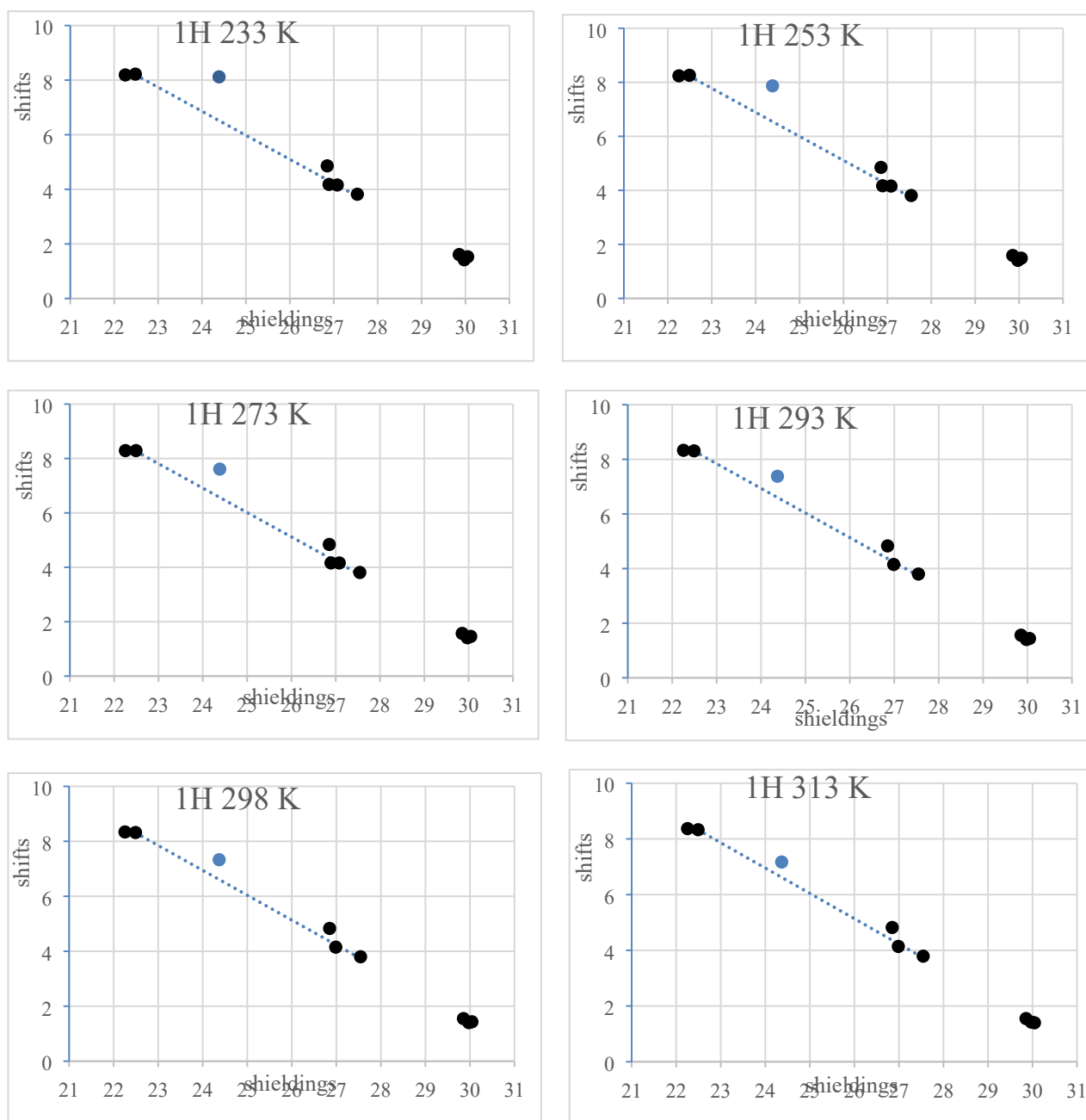


Figure S24. Linear regressions at different temperatures between ^1H calculated shieldings and ^1H experimental shifts for monomer ensemble of $\mathbf{1}_{\text{H}}$.

* ^1H nuclei bonded to carbon atoms are black, ^1H amide nucleus is blue.

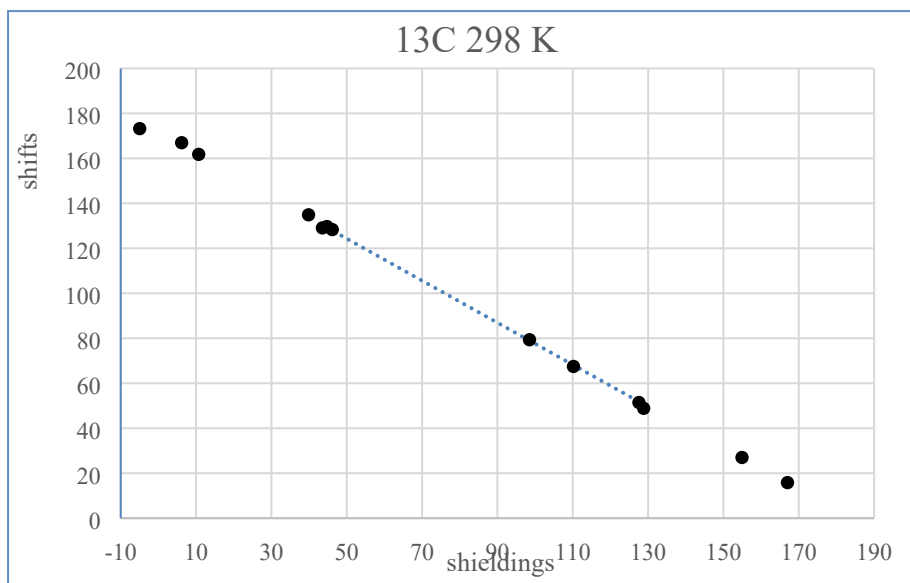


Figure S25. Linear regression at 298 K between ^{13}C calculated shieldings and ^{13}C experimental shifts for monomer ensemble of $\mathbf{1}_{\text{tt}}$.

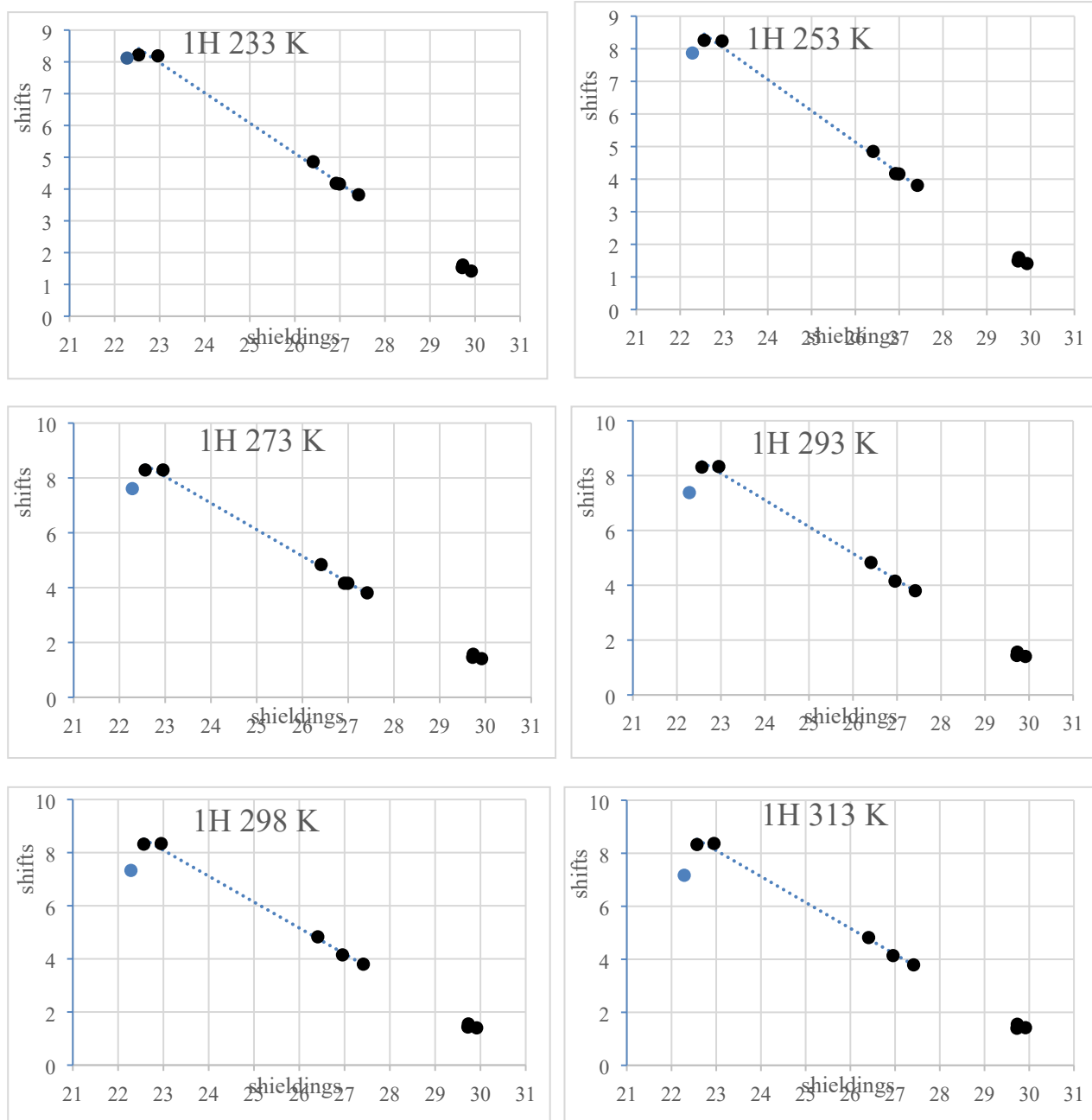


Figure S26. Linear regressions at different temperatures between ^1H calculated shieldings and ^1H experimental shifts for dimer ensemble of 11I .

* ^1H nuclei bonded to carbon atoms are black, ^1H amide nucleus is blue.

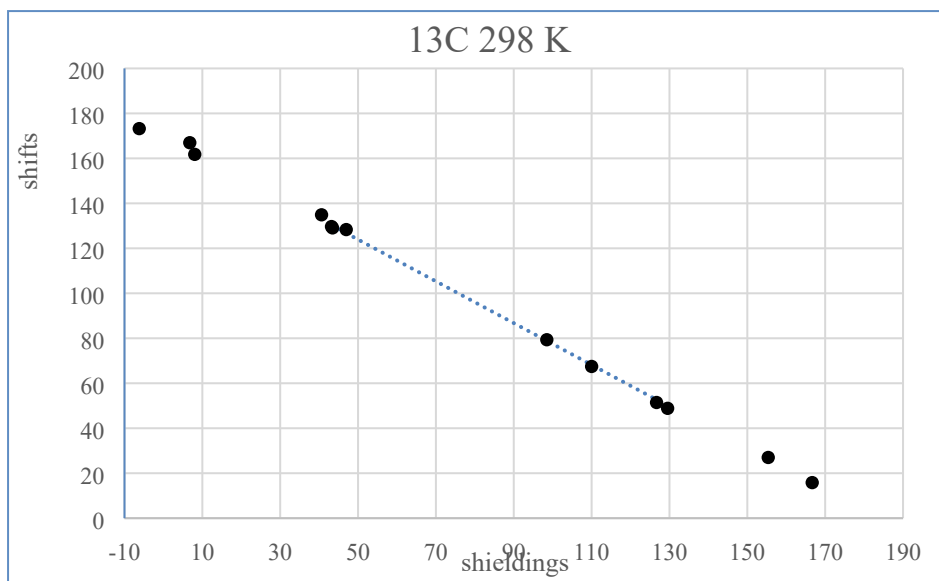


Figure S27. Linear regression at 298 K between ^{13}C calculated shieldings and ^{13}C experimental shifts for dimer ensemble of **1t1**.

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