

Supporting Information for

**Rationalizing Hydrogen Bond Solvation with Kamlet-Taft LSER and
Molecular Torsion Balances**

Bright U. Emenike, * Arzu Sevimler, Amiel Farshadmand, and Armando J. Roman
Department of Chemistry and Physics, State University of New York, 223 Store Hill Road,
Old Westbury, NY 11568, USA

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General Experimental

All reactions were carried out under an atmosphere of nitrogen in oven-dried glassware with magnetic stirring unless otherwise indicated. Reagents were purchased from commercial sources and used without further purification. Solvents for chemical reactions were dried. Purification of the reaction products was carried out by flash chromatography using silica gel 40-63 μm (230-400 mesh) unless otherwise stated. Reactions were monitored by ^1H NMR and/or thin-layer chromatography. Visualization was accomplished with UV light, staining with 5% KMnO_4 followed by heating or with p-anisaldehyde in EtOH solution. NMR was recorded using a 400 MHz Bruker spectrometer. Data are reported as (s = singlet, d = doublet, t = triplet, q = quartet, m = multiplet; integration; coupling constant(s) in Hz). Melting points were measured with a Gallenkamp melting point apparatus.

Measurement of interaction energies (ΔG) and error analysis

The folded/unfolded ratios were measured by proton NMR spectra at room temperature ($\sim 25^\circ\text{C}$). The peak areas corresponding to the ortho methyl proton in the folded and unfolded states were integrated. The folding energies were estimated from the equation: $\Delta\text{G} = -RT\ln K = -RT\ln[\text{folded}]/[\text{unfolded}]$.

The error in the folding energies was estimated to be 0.03 kcal/mol based on a conservative estimate of the NMR measured folded/unfolded ratio of $\pm 5\%$. Multiple linear regression was performed with Excel, and the result (with errors included) are as follows:¹

$$\delta(\ln K) = \delta K/K = 0.05$$

$$\delta(\Delta\text{G}) = RT[\delta(\ln K)] = 0.03 \text{ kcal/mol}$$

Synthesis of Balance 1

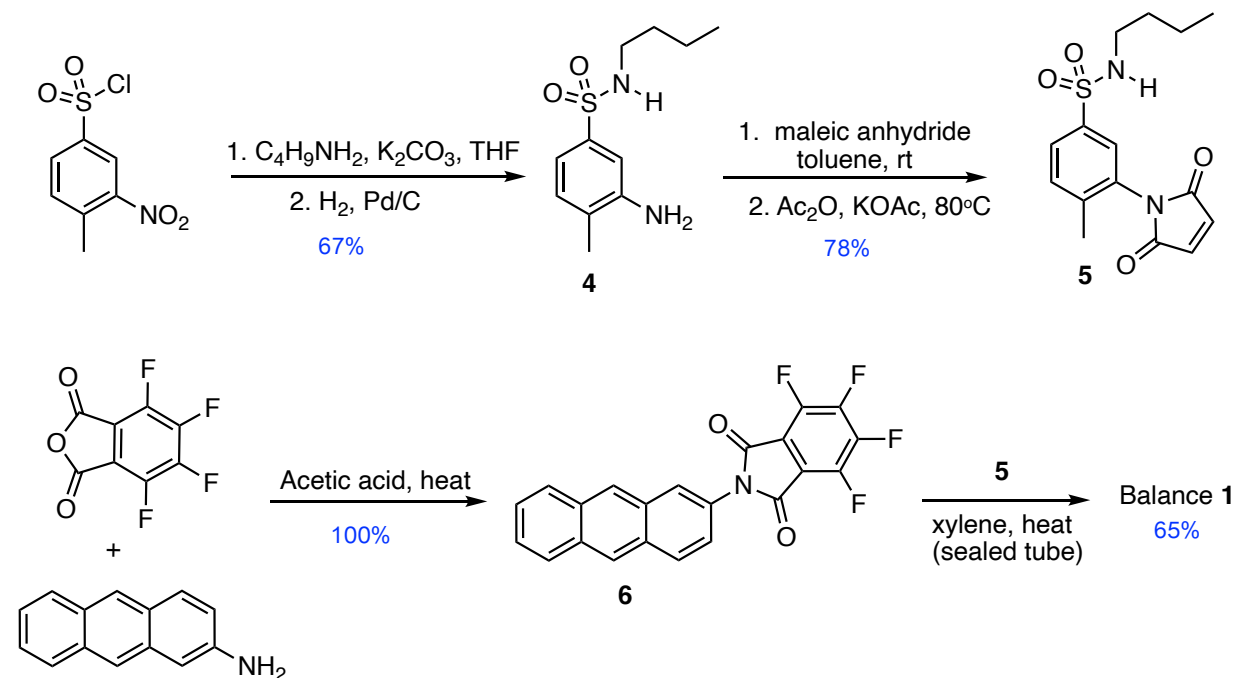


Figure S1. Synthetic scheme

Preparation and characterization of 3-amino-N-butyl-4-methylbenzenesulfonamide (**4**)

In a 100 ml round-bottom flask, 4-methyl-3-nitrobenzenesulfonyl chloride (1.5 g, 6.38 mmol) was dissolved in 20 ml of 50% (v/v) aqueous THF. To the mixture, K_2CO_3 (1.32g, 9.57 mmol) and butylamine (0.7g, 9.57 mmol) were added sequentially at room temperature. The reaction flask was then heated to reflux for 2 hours, and the reaction progress was monitored by TLC. After completion, the reaction was cooled to room temperature and the volume was reduced to half by removing solvent under reduced pressure. Next, 20 ml of water was added, and the mixture was extracted three times using 20 mL portions of ethyl acetate. The combined organic layers were washed with brine and water, dried over anhydrous $MgSO_4$, and concentrated to yield the crude intermediate. The crude intermediate (dissolved in 15 mL ethyl acetate) was then added to 150 mg of 10% Pd/C and the reaction was stirred overnight under normal atmospheric pressure of hydrogen gas. The reaction mixture was filtered, and the concentrated product was purified by flash column chromatography using mixtures of ethyl acetate and hexane to yield 1.04 g of 3-amino-N-butyl-4-methylbenzenesulfonamide (**4**).

Proton NMR:

^1H NMR (400 MHz, CDCl_3) δ 7.15 (t, $J = 8.4$ Hz, 3H), 4.96 (t, $J = 6.1$ Hz, 1H), 3.97 (s, 2H), 2.88 (q, $J = 6.8$ Hz, 2H), 2.18 (d, $J = 3.4$ Hz, 3H), 1.42 (m, 2H), 1.26 (m, 2H), 0.83 (t, $J = 7.3$ Hz, 3H)

Carbon NMR:

$^{13}\text{C}\{1\text{H}\}$ NMR (101 MHz, CDCl_3) δ 145.5, 137.9, 130.8, 127.1, 116.6, 112.6, 42.9, 31.5, 19.7, 17.5, 13.6

Mass spec: ESI(MS) for $\text{C}_{11}\text{H}_{18}\text{N}_2\text{O}_2\text{S}$ ($\text{M}+\text{Na}$) $^+$; calcd 265.0987, found 265.0980

Preparation and characterization of N-butyl-3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-4-methylbenzenesulfonamide (5)

A mixture of 3-amino-*N*-butyl-4-methylbenzenesulfonamide (1.2 g, 4.95 mmol) and maleic anhydride (0.5 g, 5.1 mmol) was prepared in 10 ml of toluene and was stirred overnight at room temperature. The toluene was removed by decantation, and acetic anhydride (20 ml) and potassium acetate (0.15 g) were added to the mixture. The reaction was heated for 30 mins, and progress was monitored by TLC. The acetic anhydride was removed under reduced pressure, and 20 ml of water was carefully added. The aqueous mixture was extracted three times using 15 ml portions of diethyl ether, and the combined extracts were washed with brine and water before being dried with anhydrous MgSO_4 . The product was purified by column chromatography to yield 1.24 g (78%) of N-butyl-3-(2,5-dioxo-2,5-dihydro-1H-pyrrol-1-yl)-4-methylbenzenesulfonamide (5).

Proton NMR:

^1H NMR (400 MHz, CDCl_3) δ 7.82 (q, $J = 3.3$, 1H), 7.66 (d, $J = 1.9$ Hz, 1H), 7.49 (d, $J = 8.1$ Hz, 1H), 6.92 (s, 2H), 4.79 (t, $J = 6.1$ Hz, 1H), 2.97 (q, $J = 6.8$ Hz, 2H), 2.25 (s, 3H), 1.46 (m, $J = 4.9$ Hz, 2H), 1.32 (m, $J = 7.4$ Hz, 2H), 0.87 (t, $J = 7.3$, 3H)

Carbon NMR:

$^{13}\text{C}\{1\text{H}\}$ NMR (101 MHz, CDCl_3) δ 168.9, 142.0, 138.7, 134.6, 131.9, 130.8, 127.8, 127.6, 43.0, 31.6, 19.7, 18.3, 13.5

Mass spec: ESI(MS) for $\text{C}_{15}\text{H}_{18}\text{N}_2\text{O}_4\text{S}$ ($\text{M}+\text{Na}$) $^+$; calcd 345.0885, found 345.0880

Preparation and characterization of 2-(anthracen-2-yl)-4,5,6,7-tetrafluoroisoindoline-1,3-dione (**6**).

A mixture containing 2-aminoanthracene (0.386 g, 2mmol) and pentafluorophthalic anhydride (0.44 g, 2mmol) in 15 ml acetic acid was then heated and refluxed overnight. The reaction was cooled, and water was added to it. The desired product was isolated by filtration. The filtrate was washed several times, first with water, then with a small amount of cold acetone, and finally with hexane. The resulting dried product was found to be pure and was used without further purification. The yield of the product (**6**) was 0.78 g.

Proton NMR:

^1H NMR (400 MHz, DMSO- d_6) δ 7.43 (m, 3H), 8.04 (m, 3H), 8.40 (m, 3H)

Carbon NMR:

$^{13}\text{C}\{1\text{H}\}$ NMR (101 MHz, DMSO- d_6) δ 169.9, 136.4, 132.1, 131.9, 130.9, 129.3, 128.3, 128.9, 128.9, 128.5, 128.1, 126.3, 126.2, 125.5, 125.4, 121.4, 114.4

Mass spec: CI(MS) for $\text{C}_{22}\text{H}_9\text{NO}_2\text{F}_4$ (M) $^+$; calcd 395.0569, found 395.0574

Diels-Alder: Preparation and characterization of Balance 1

A mixture of **5** (0.2 g, 0.85 mmol) and **6** (0.28 g, 0.71 mmol) in 1 ml xylene was heated in a sealed reaction tube for two days. The cooled reaction was then purified by column chromatography to yield 0.33g of balance **1**.

Proton NMR:

^1H NMR (400 MHz, CDCl_3) δ 7.69 (q, $J = 3.3$ Hz, 1H), 7.53 (d, $J = 1.9$ Hz), 7.49 (m, $J = 2.7$ Hz, 3H), 7.39 (q, $J = 3.3$ Hz, 1H), 7.33 (d, 2H), 6.14 (d, $J = 1.8$ Hz, 1H), 5.27 (q, $J = 4.0$ Hz, 1H), 4.99 (d, $J = 1.7$ Hz, 2H), 3.50 (t, $J = 1.5$ Hz, 2H), 2.98 (m, $J = 6.6$ Hz, 1H), 2.79 (m, $J = 6.1$ Hz, 1H), 1.49 (m, $J = 6.3$ Hz, 2H), 1.37 (q, $J = 7.4$ Hz, 2H), 0.93 (t, $J = 7.3$ Hz, 3H)

Carbon NMR:

$^{13}\text{C}\{1\text{H}\}$ NMR (101 MHz, CDCl_3) δ 175.4, 174.9, 162.3, 142.5, 140.7, 140.3, 140.2, 139.6, 139.3, 138.7, 131.6, 130.9, 129.3, 128.4, 127.7, 127.4, 127.3, 126.3, 126.2, 125.6, 124.6, 122.8, 113.4, 47.1, 47.1, 46.0, 45.6, 45.3, 45.1, 43.0, 31.6, 31.3, 19.7, 18.0, 17.0, 13.6, 13.5

Mass spec: ESI(MS) for $\text{C}_{37}\text{H}_{27}\text{N}_3\text{O}_6\text{SF}_4$ (M+Na) $^+$; calcd 740.1449, found 740.1450

Melting point: 239 – 241°C

Preparation and characterization of Balance 2

A mixture of anthracene (0.18g, 1 mmol) and **5** (0.282 g, 1.2 mmol) in 1 ml xylene was heated in a sealed reaction tube overnight. The cooled reaction was then purified by column chromatography to yield 0.395g (96%) of balance **2**.

Proton NMR:

^1H NMR (400 MHz, CDCl_3) δ 7.69 (m, $J = 2.5$ Hz, 1H), 7.33 (m, 9H), 5.94 (d, $J = 1.8$ Hz, 1H), 4.92 (s, 2H), 4.36 (dt, $J = 5.8$, 1H), 3.45 (q, $J = 1.7$, 2H), 2.91 (m, $J = 6.7$ Hz, 1.5H), 2.15 (s, 2H), 1.37 (m, 4H), 1.14 (s, 1.7H), 0.93 (t, $J = 7.3$ Hz, 1.5 H), 0.87 (t, $J = 7.3$ Hz, 1.5H)

Carbon NMR:

$^{13}\text{C}\{1\text{H}\}$ NMR (101 MHz, CDCl_3) δ 175.6, 175.5, 141.7, 141.1, 141.0, 139.3, 138.6, 138.6, 138.3, 131.7, 131.5, 131.3, 127.6, 127.4, 127.1, 126.9, 126.9, 126.8, 125.5, 125.2, 124.3, 47.3, 45.9, 45.3, 43.0, 31.6, 31.5, 19.8, 19.7, 17.9, 16.7, 13.6, 13.6

Mass spec: ESI(MS) for $\text{C}_{29}\text{H}_{28}\text{N}_2\text{O}_4\text{S}$ ($\text{M}+\text{Na}^+$); calcd 523.1, found 523.2

Melting point: 122 – 124 °C

Table 1. Summary of solvent parameters

solvents	Kamlet-Taft parameters				solvent polarity			Hunter parameters	
	α	β	π^*	δ	$E_T(30)$	μ	ϵ	α_s	β_s
CCl_4	0	0.1	0.58	0.5	32.4	0	2.2	1.4	0.6
toluene	0	0.11	0.54	1	33.9	0.31	2.4	1	2.1
benzene	0	0.1	0.59	1	34.3	0	2.3	1.1	1.6
chloroform	0.2	0.1	0.58	0.5	39.1	1.15	4.8	2.2	1.3
CH_2Cl_2	0.13	0.1	0.82	0.5	40.7	1.14	8.9	1.7	1.1
acetic acid	1.12	0.45	0.64	0	55.2	1.78	6.2	3.7	4.9
acetone	0.08	0.43	0.71	0	42.2	2.69	21	1.2	5.7
acetonitrile	0.19	0.4	0.75	0	45.6	3.5	36.6	1.5	5.1
methanol	0.98	0.66	0.6	0	55.4	2.87	32.6	2.8	4.8
pyridine	0	0.64	0.87	1	40.5	2.37	12.3	1.2	7.2
ethanol	0.86	0.75	0.54	0	51.9	1.66	24.6	2.7	5.2
DMF	0	0.69	0.88	0	43.8	3.9	36.7	2.2	7.4
DMSO	0	0.76	1	0	45.1	4.1	47	0.8	8.9
<i>t</i> -butanol	0.42	0.93	0.41	0	43.7	1.31	10.9	2.7	5.7

Symbols and definition:

α = Kamlet-Taft parameter describing hydrogen bond donor ability

β = Kamlet-Taft parameter describing hydrogen bond acceptor ability
 π^* = Kamlet-Taft parameter describing dipolarity-polarizability
 δ = correction factor: 0.5 for chlorinated solvent, 1.0 for aromatic solvents
 $E_T(30)$ = Reichardt's solvent-polarity parameter
 μ = solvent dipole moment in Debye
 ϵ = relative permittivity (dielectric constant)
 α_s = Hunter solvent hydrogen-bond donor parameter
 β_s = Hunter hydrogen-bond acceptor parameter

Table 2. Summary of experimental measurements and predicted values

solvents	% folded		equilibrium constants		folding energy (kcal/mol)		H-bond energy (kcal/mol)		
	1	2	$K_{eq}(1)$	$K_{eq}(2)$	$\Delta G(1)$	$\Delta G(2)$	expt.	(Kamlet-Taft)	(Hunter)
CCl ₄	82	46	4.56	0.85	-0.90	0.09	-0.99	-0.87	-1.02
toluene	84	52	5.25	1.08	-0.98	-0.05	-0.93	-1.02	-0.85
benzene	85	53	5.67	1.13	-1.03	-0.07	-0.96	-1.01	-0.96
chloroform	83	50	4.88	1.00	-0.94	0.00	-0.94	-0.90	-0.84
CH ₂ Cl ₂	82	53	4.56	1.13	-0.90	-0.07	-0.83	-0.71	-0.92
acetic acid	50	49	1.00	0.96	0.00	0.02	-0.02	-0.11	0.17
acetone	34	40	0.52	0.67	0.39	0.24	0.15	0.05	0.29
acetonitrile	39	42	0.64	0.72	0.26	0.19	0.07	0.00	0.16
methanol	39	51	0.64	1.04	0.26	-0.02	0.29	0.32	0.13
pyridine	31	44	0.45	0.79	0.47	0.14	0.33	0.34	0.30
ethanol	35	51	0.54	1.04	0.37	-0.02	0.39	0.48	0.22
DMF	19	42	0.23	0.72	0.86	0.19	0.67	0.73	0.69
DMSO	9	35	0.10	0.54	1.37	0.37	1.00	0.97	0.99
<i>t</i> -butanol	64	39	1.78	0.64	-0.34	0.26	-0.61	0.85	0.37

Plots of experimental H-bond (ΔG_{H-bond} in kcal/mol) with solvent polarity parameters

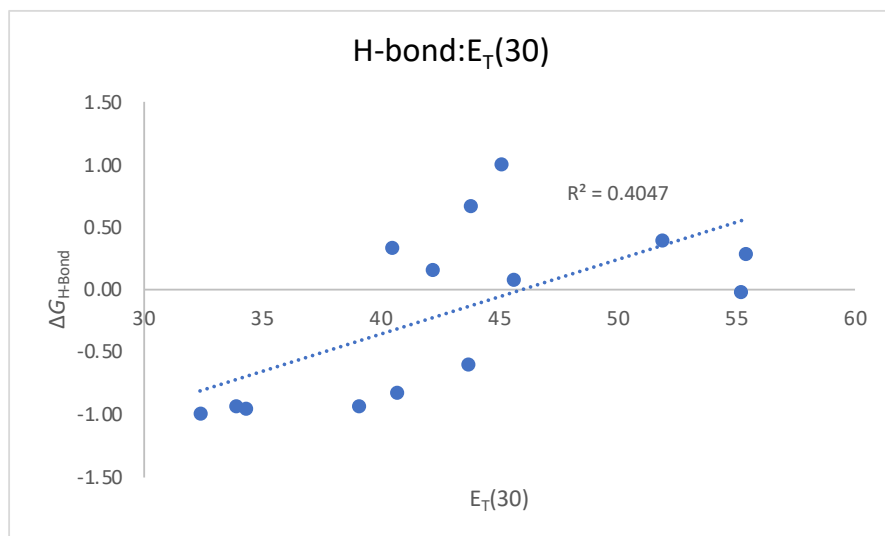


Figure S2 A plot of ΔG_{H-bond} in kcal/mol against $E_T(30)$ solvent polarity parameter

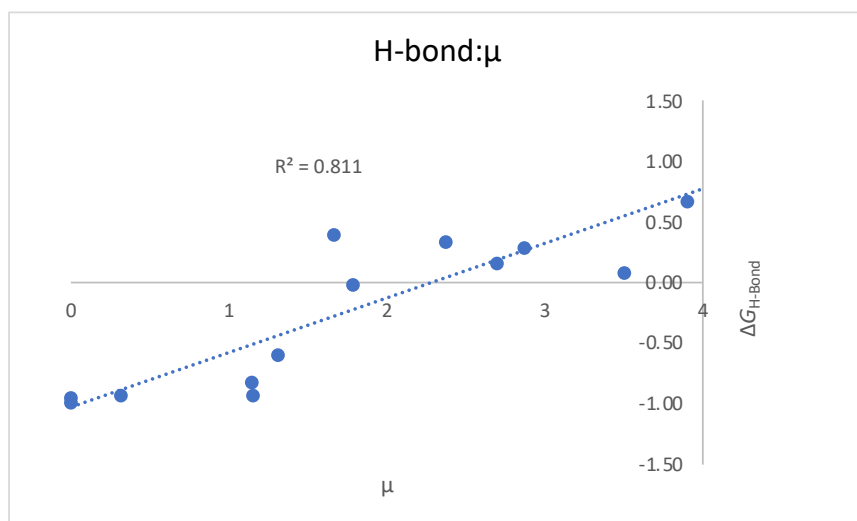


Figure S3 A plot of ΔG_{H-bond} in kcal/mol against solvents' dipole moment (in Debye)

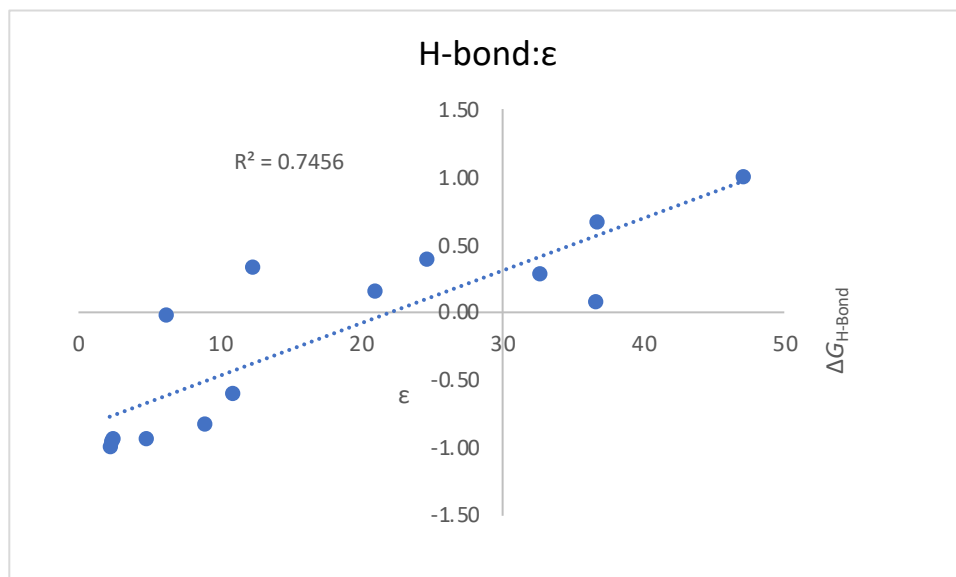


Figure S4 A plot of $\Delta G_{\text{H-bond}}$ (in kcal/mol) against solvents' dielectric constant solvent

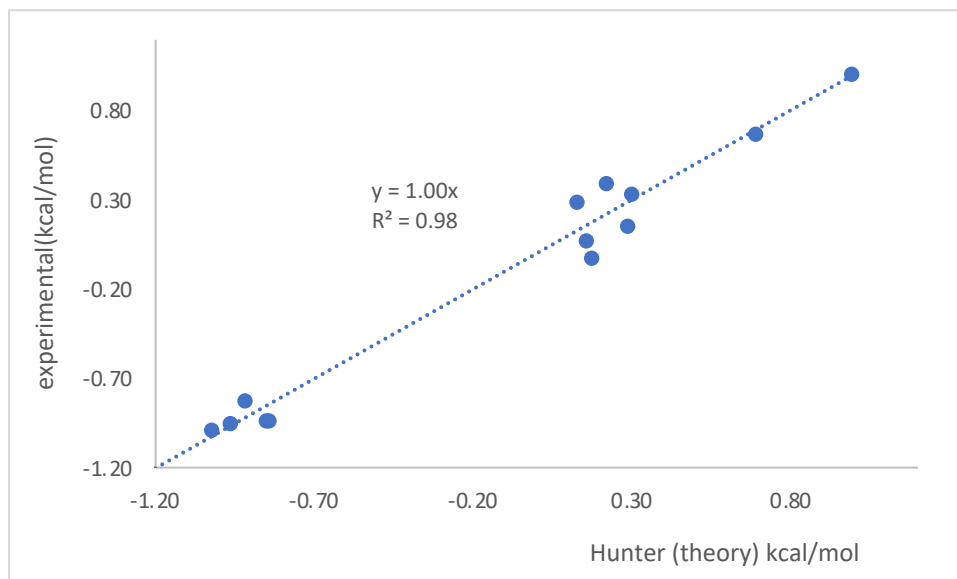


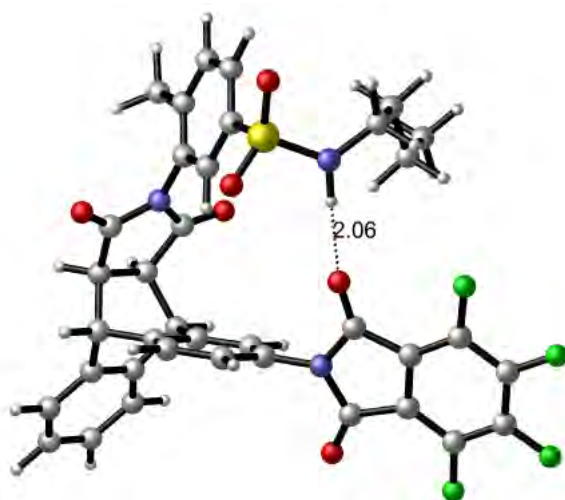
Figure S5. Correlating the experimental $\Delta G_{\text{H-bond}}$ (in kcal/mol) with Hunter solvation parameters (including π^* dipolarity-polarization term)

$$\Delta G_{\text{H-Bond}} = -1.0 + 0.03\alpha_s + 0.28\beta_s - 0.06\pi^* - 0.32\delta$$

Summary of computational single-point calculations performed at the M06-2X/6-31+G* using conformers optimized at the B3LYP/6-31G+* level

Model System	Conformer	Total Energy		Relative Energies		Imaginary Frequency
		(a.u)	(kcal/mol)	(a.u)	(kcal/mol)	
1	folded	-2838.454735	-1781158.73	-0.005403	-3.39	0
1	unfolded	-2838.449332	-1781155.34	0	0.00	0
2	folded	-1929.919842	-1211044.00	-0.00071	-0.45	0
2	unfolded	-1929.919132	-1211043.55	0	0.00	0

**The cartesian coordinates for the optimized conformer at the B3LYP/6-31+G* level:
Balance 1 folded conformer**

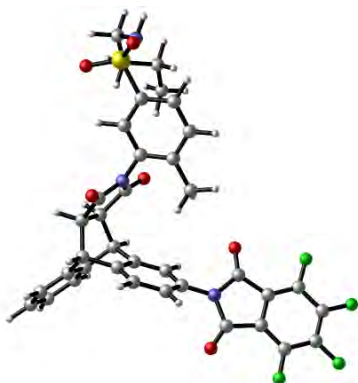


C-C_R	2.11643	-0.14443	4.79692
C-C_R	1.88722	1.19886	4.44644
C-C_3	1.3801	-1.13306	3.90498
H-H_	1.56594	-2.17578	4.17346
C-C_3	0.96519	1.35808	3.24639

C-C_R	1.75722	-0.82154	2.4669
C-C_R	2.36443	0.06371	-0.094
C-C_R	2.25629	-1.72246	1.52871
C-C_R	1.54546	0.52488	2.11649
C-C_R	1.84227	0.97218	0.83384
C-C_R	2.56997	-1.27558	0.24056
H-H_	2.40241	-2.76721	1.78983
H-H_	1.67839	2.00776	0.54928
H-H_	2.96407	-1.96493	-0.49971
C-C_3	-0.14356	-0.78554	4.07396
H-H_	-0.43237	-0.98649	5.11033
C-C_3	-0.39262	0.68309	3.66051
H-H_	-0.84283	1.27292	4.46565
C-C_R	-1.03791	-1.61507	3.1617
C-C_R	-1.39106	0.60824	2.50883
O-O_R	-1.15784	-2.82092	3.16388
O-O_R	-1.84013	1.54225	1.87548
N-N_R	-1.7186	-0.73833	2.29461
C-C_R	-2.58762	-1.19684	1.24597
C-C_R	-4.21132	-2.18153	-0.79068
C-C_R	-3.9781	-1.01191	1.34384
C-C_R	-2.00888	-1.85545	0.16191
C-C_R	-2.82707	-2.33991	-0.85745
C-C_R	-4.76917	-1.5178	0.30264
H-H_	-4.84117	-2.59414	-1.57227
N-N_R	2.69999	0.52412	-1.41112
C-C_R	3.80046	1.36967	-1.68835
C-C_R	1.98426	0.18345	-2.56856
O-O_R	4.57211	1.81731	-0.87239
O-O_R	0.99215	-0.51518	-2.60777
C-C_R	3.77948	1.57254	-3.16982
C-C_R	3.27214	1.61323	-5.89502
C-C_R	4.61687	2.30456	-3.99221
C-C_R	2.69215	0.86039	-3.69737
C-C_R	2.42782	0.87226	-5.05541
C-C_R	4.35361	2.32096	-5.36981
H-H_	0.8067	2.40094	2.96195
S-S_3+4	-2.06869	-3.18734	-2.26395

O-O_2	-3.13958	-3.95283	-2.91442
O-O_2	-0.8506	-3.8169	-1.74236
N-N_3	-1.56143	-2.05423	-3.35601
H-H_	-0.69151	-1.59198	-3.10163
C-C_3	-2.41327	-1.46468	-4.39271
C-C_R	2.48332	2.22612	5.17361
H-H_	2.30987	3.26498	4.90098
C-C_R	3.31659	1.90926	6.25543
H-H_	3.79059	2.70636	6.82218
C-C_R	3.545	0.5756	6.60262
H-H_	4.19538	0.33662	7.43997
C-C_R	2.94119	-0.4587	5.87396
H-H_	3.12067	-1.49722	6.14389
H-H_	-3.12705	-2.23606	-4.69504
H-H_	-1.75962	-1.28002	-5.25531
C-C_3	-3.14901	-0.16898	-4.01528
H-H_	-3.81912	0.07886	-4.85261
H-H_	-3.7948	-0.34972	-3.14557
C-C_3	-2.22912	1.02663	-3.73511
H-H_	-1.57348	1.18616	-4.6044
H-H_	-1.56938	0.79255	-2.88952
C-C_3	-3.00269	2.31564	-3.43169
H-H_	-3.64325	2.60316	-4.27565
H-H_	-2.32334	3.15264	-3.22929
H-H_	-3.64627	2.19202	-2.55186
F-F_	1.40309	0.21193	-5.60076
F-F_	3.04401	1.64773	-7.20962
F-F_	5.13836	3.02218	-6.1904
F-F_	5.66091	2.99503	-3.53071
H-H_	-0.93537	-2.00849	0.12413
H-H_	-5.84862	-1.40185	0.35966
C-C_3	-4.606	-0.30878	2.52096
H-H_	-5.69532	-0.40711	2.49489
H-H_	-4.36055	0.75987	2.5163
H-H_	-4.25379	-0.72574	3.47216

The cartesian coordinates for the optimized conformer at the B3LYP/6-31G+* level:
Balance 1 unfolded conformer

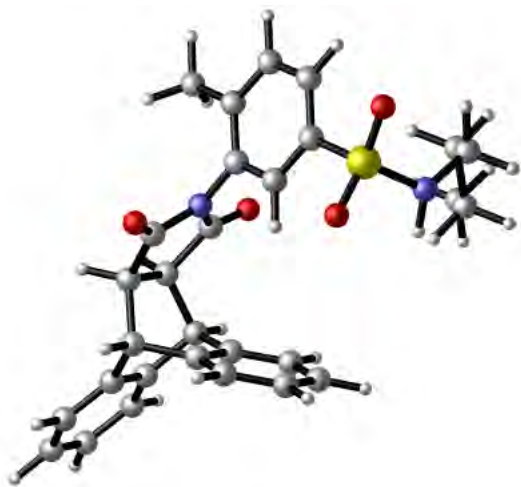


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C-C_R	-2.32589	3.46543	1.09893
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H-H_	-2.76417	2.83653	-2.21716
C-C_3	-1.46268	2.21367	1.19474
C-C_R	-2.77058	1.30285	-0.66068
C-C_R	-3.40222	-1.06649	0.65875
C-C_R	-3.55635	0.34657	-1.29926
C-C_R	-2.28558	1.06372	0.63803
C-C_R	-2.60051	-0.11503	1.30573
C-C_R	-3.87228	-0.84516	-0.63895
H-H_	-3.92767	0.52199	-2.30567
H-H_	-2.23069	-0.29952	2.30903
H-H_	-4.49368	-1.58931	-1.12464
C-C_3	-0.80261	2.72603	-1.21403
H-H_	-0.5003	3.70473	-1.60256
C-C_3	-0.2662	2.47784	0.21448
H-H_	0.30052	3.33217	0.599
C-C_R	-0.12961	1.66814	-2.08323
C-C_R	0.71124	1.31591	0.07809
O-O_R	-0.29415	1.48667	-3.27032
O-O_R	1.36969	0.81004	0.96353
N-N_R	0.75439	0.93202	-1.27189
C-C_R	1.68742	-0.03439	-1.78431

C-C_R	3.57869	-1.8269	-2.77348
C-C_R	2.8127	0.45006	-2.45338
C-C_R	1.47595	-1.41019	-1.59714
C-C_R	2.44472	-2.28776	-2.11018
C-C_R	3.75632	-0.45189	-2.93481
H-H_	2.94986	1.51568	-2.6038
H-H_	4.31129	-2.52342	-3.16891
N-N_R	-3.74638	-2.27919	1.33742
C-C_R	-5.06296	-2.79006	1.42672
C-C_R	-2.81022	-3.11015	1.99208
O-O_R	-6.05771	-2.28477	0.95768
O-O_R	-1.61691	-2.91682	2.06817
C-C_R	-4.94821	-4.05848	2.20882
C-C_R	-4.17507	-6.28209	3.67361
C-C_R	-5.91453	-4.97115	2.5945
C-C_R	-3.60175	-4.24811	2.54936
C-C_R	-3.20077	-5.35359	3.27899
C-C_R	-5.51534	-6.09295	3.33571
H-H_	-1.09692	2.01873	2.20579
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H-H_	0.27086	-1.65661	0.18088
H-H_	0.22905	-3.03091	-0.92983
H-H_	-0.66144	-1.54918	-1.3172
H-H_	2.30063	-3.35772	-1.98442
S-S_3+4	5.22845	0.16336	-3.76188
O-O_2	5.01844	1.59691	-4.00751
O-O_2	5.55727	-0.77048	-4.84829
N-N_3	6.46413	-0.0512	-2.62328
H-H_	7.21107	-0.55189	-3.09915
C-C_3	6.92357	1.09794	-1.80969
C-C_R	-2.6301	4.34086	2.13744
H-H_	-2.25271	4.15483	3.14079
C-C_R	-3.43096	5.46268	1.88023
H-H_	-3.67681	6.14625	2.68865
C-C_R	-3.9135	5.70314	0.59171
H-H_	-4.53491	6.57363	0.39837
C-C_R	-3.60284	4.82433	-0.4558
H-H_	-3.97863	5.01357	-1.4591

H-H_	6.90527	2.0183	-2.40545
H-H_	7.97049	0.88102	-1.56616
C-C_3	6.12477	1.28905	-0.51661
H-H_	6.50136	2.2049	-0.03742
H-H_	5.07284	1.48738	-0.76191
C-C_3	6.21283	0.11891	0.4717
H-H_	7.27096	-0.07531	0.7054
H-H_	5.82979	-0.79027	-0.00812
C-C_3	5.44136	0.38301	1.77079
H-H_	5.82406	1.27561	2.28305
H-H_	5.5314	-0.46209	2.46375
H-H_	4.37337	0.53979	1.57597
F-F_	-1.9304	-5.57319	3.62419
F-F_	-3.82531	-7.3595	4.38032
F-F_	-6.42078	-6.9932	3.72527
F-F_	-7.20638	-4.83201	2.2905

**The cartesian coordinates for the optimized conformer at the B3LYP/6-31G+* level:
Balance 2 folded conformer**

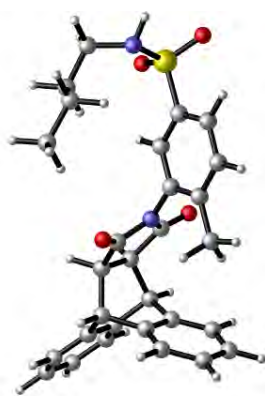


C-C_R	2.80322	0.28408	3.75049
C-C_R	2.51264	1.60568	3.36364
C-C_3	2.0954	-0.7601	2.90056
H-H_	2.32974	-1.78548	3.19743
C-C_3	1.56408	1.68953	2.17717

C-C_R	2.43278	-0.47687	1.44604
C-C_R	2.95248	0.35403	-1.16531
C-C_R	2.95909	-1.38412	0.52822
C-C_R	2.1527	0.84626	1.05765
C-C_R	2.40883	1.26267	-0.24707
C-C_R	3.22294	-0.96272	-0.78207
H-H_	3.16194	-2.41029	0.82541
H-H_	2.19053	2.28436	-0.55026
H-H_	3.63647	-1.66494	-1.50091
C-C_3	0.56105	-0.47182	3.08917
H-H_	0.29541	-0.66002	4.13437
C-C_3	0.24458	0.97634	2.64774
H-H_	-0.20503	1.56883	3.45144
C-C_R	-0.31286	-1.35778	2.21173
C-C_R	-0.78051	0.837	1.52817
O-O_R	-0.38282	-2.56747	2.23501
O-O_R	-1.29205	1.73573	0.89061
N-N_R	-1.05212	-0.52722	1.34586
C-C_R	-1.91825	-1.03412	0.32034
C-C_R	-3.53698	-2.0683	-1.69834
C-C_R	-3.31321	-0.89231	0.43317
C-C_R	-1.3325	-1.67964	-0.76793
C-C_R	-2.14901	-2.18835	-1.7768
C-C_R	-4.10122	-1.42188	-0.59857
H-H_	-4.16219	-2.4927	-2.47734
H-H_	1.35557	2.71633	1.86679
S-S_3+4	-1.39007	-2.98823	-3.20462
O-O_2	-2.45041	-3.75836	-3.86197
O-O_2	-0.13578	-3.59035	-2.7418
N-N_3	-0.91603	-1.81007	-4.28944
H-H_	-0.04941	-1.36553	-4.00185
C-C_3	-1.86689	-0.99785	-5.0617
C-C_R	3.07514	2.6779	4.05165
H-H_	2.85195	3.70006	3.75272
C-C_R	3.93708	2.42964	5.12919
H-H_	4.38376	3.26294	5.66548
C-C_R	4.22743	1.11779	5.51146
H-H_	4.8994	0.93118	6.34515

C-C_R	3.65634	0.03797	4.82317
H-H_	3.88349	-0.98367	5.12065
H-H_	-2.76699	-1.60603	-5.18859
H-H_	-1.4332	-0.87218	-6.06293
C-C_3	-2.20926	0.37704	-4.46652
H-H_	-2.99882	0.8163	-5.09434
H-H_	-2.64053	0.25232	-3.46482
C-C_3	-1.0243	1.35157	-4.40389
H-H_	-0.58485	1.45238	-5.40755
H-H_	-0.23424	0.94156	-3.75781
C-C_3	-1.41946	2.73593	-3.87473
H-H_	-2.17535	3.2038	-4.51846
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H-H_	-0.25476	-1.79945	-0.81431
H-H_	-5.18277	-1.33619	-0.53037
C-C_3	-3.94997	-0.20728	1.61585
H-H_	-5.03569	-0.34114	1.60218
H-H_	-3.73919	0.86865	1.60552
H-H_	-3.57282	-0.61	2.56338
H-H_	3.1666	0.67845	-2.18067

**The cartesian coordinates for the optimized conformer at the B3LYP/6-31G+* level:
Balance 2 unfolded conformer**



C-C_R	-3.79856	2.6579	0.39431
C-C_R	-3.30906	2.42857	1.69468
C-C_3	-3.34528	1.60867	-0.61284

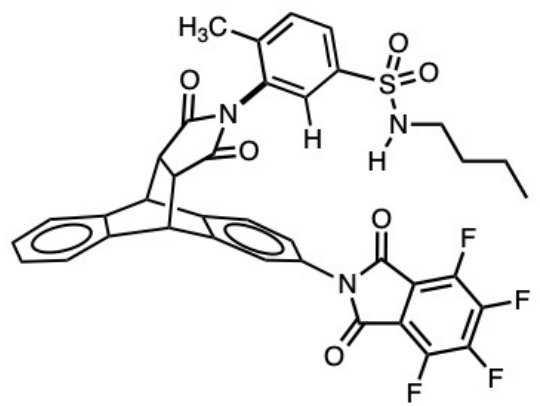
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C-C_R	-3.73018	0.24856	-0.05432
C-C_R	-4.31831	-2.13989	1.26672
C-C_R	-4.49682	-0.72179	-0.69692
C-C_R	-3.24456	0.02138	1.24636
C-C_R	-3.53399	-1.17133	1.90698
C-C_R	-4.79411	-1.91758	-0.02869
H-H_	-4.86534	-0.54786	-1.70542
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C-C_3	-1.77885	1.69677	-0.61264
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H-H_	-0.66914	2.30177	1.19683
C-C_R	-1.09534	0.65004	-1.48512
C-C_R	-0.27054	0.28138	0.68
O-O_R	-1.24376	0.47937	-2.67578
O-O_R	0.37793	-0.23682	1.56707
N-N_R	-0.22021	-0.09588	-0.67086
C-C_R	0.70722	-1.06602	-1.18397
C-C_R	2.5861	-2.86829	-2.18177
C-C_R	1.83545	-0.58742	-1.85239
C-C_R	0.48756	-2.44136	-1.0018
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H-H_	-0.71725	-2.66333	0.77325
H-H_	-0.75974	-4.05698	-0.32198
H-H_	-1.65371	-2.58081	-0.72161
H-H_	1.29817	-4.39346	-1.39835
S-S_3+4	4.24793	-0.88457	-3.16401
O-O_2	4.04676	0.55127	-3.40278
O-O_2	4.57012	-1.81542	-4.25493
N-N_3	5.48376	-1.11247	-2.02721

H-H_	6.22822	-1.61226	-2.508
C-C_3	5.94969	0.03064	-1.2087
C-C_R	-3.62268	3.30683	2.72812
H-H_	-3.24402	3.1297	3.73268
C-C_R	-4.43504	4.4192	2.46539
H-H_	-4.68797	5.10433	3.27039
C-C_R	-4.91981	4.64793	1.17559
H-H_	-5.54972	5.51126	0.97731
C-C_R	-4.59975	3.76648	0.133
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C-C_3	5.23591	-0.95535	1.06852
H-H_	6.29324	-1.15478	1.30142
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C-C_3	4.46548	-0.69423	2.36884
H-H_	4.85124	0.19479	2.88492
H-H_	4.55293	-1.54262	3.05822
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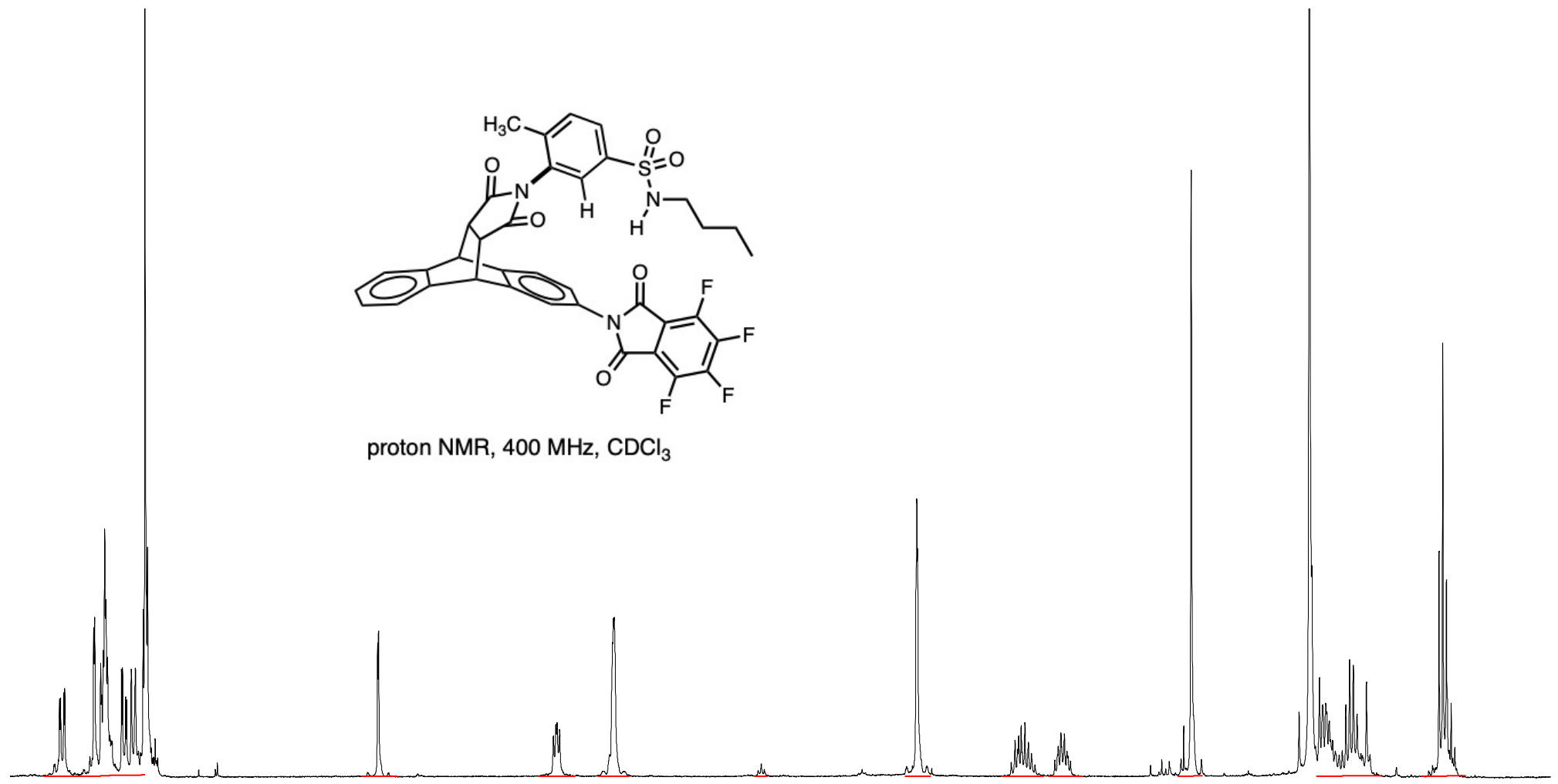
References

1. Gardarsson, H.; Schweizer, W. B.; Trapp, N.; Diederich, F., Structures and Properties of Molecular Torsion Balances to Decipher the Nature of Substituent Effects on the Aromatic Edge-to-Face Interaction. *Chemistry – A European Journal* **2014**, *20* (16), 4608-4616.

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0.908
0.885



proton NMR, 400 MHz, CDCl₃



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1.5
1.0
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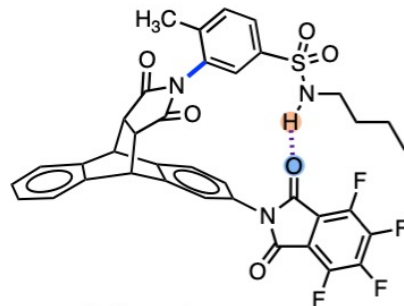
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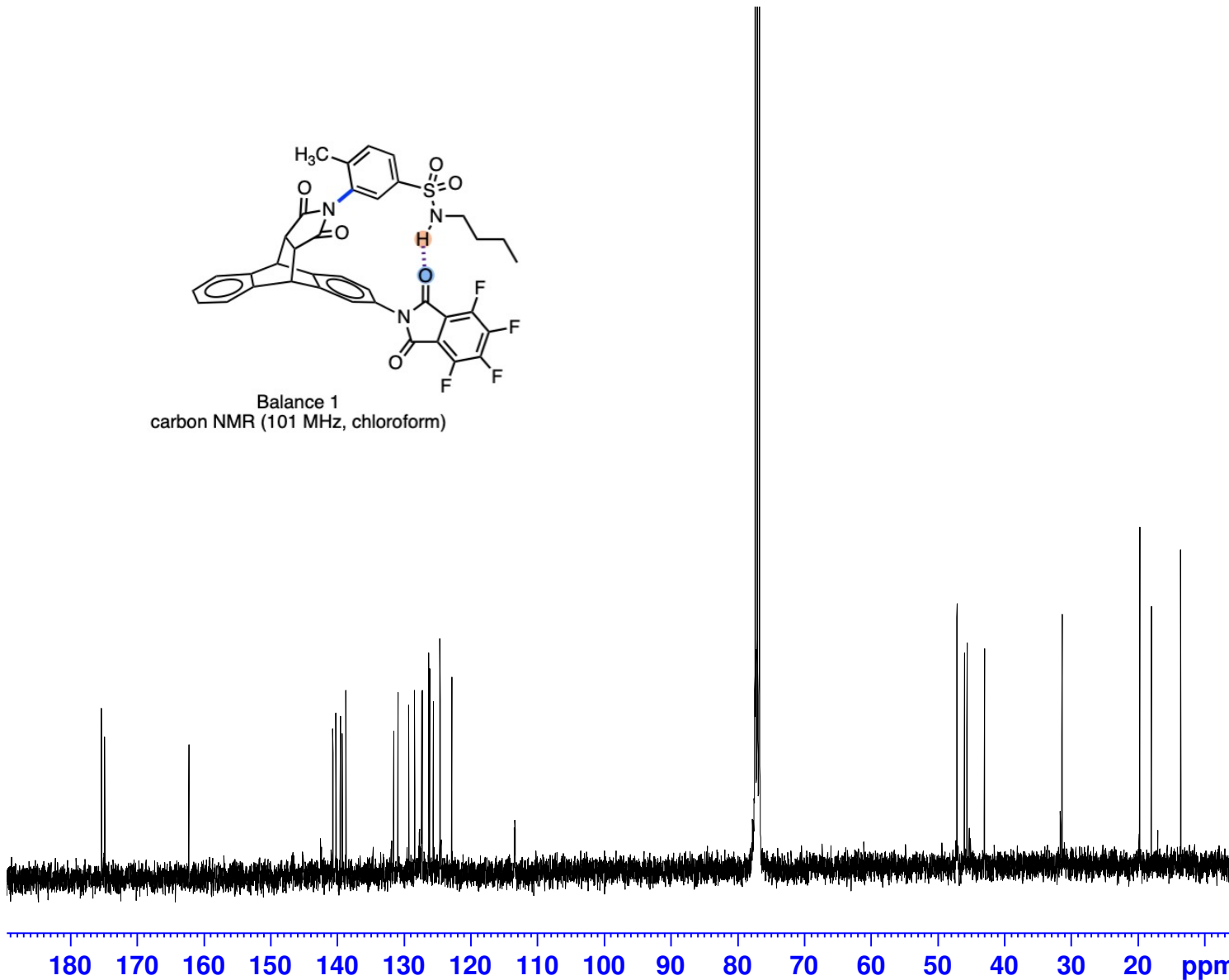
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122.82
113.36

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47.07
45.96
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42.97
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Balance 1
carbon NMR (101 MHz, chloroform)



Current Data Parameters
NAME controljan142023balanc
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20230114
Time 19.48
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zgpg30
TD 65536
SOLVENT CDC13
NS 2883
DS 4
SWH 24038.461 Hz
FIDRES 0.366798 Hz
AQ 1.3631488 sec
RG 2050
DW 20.800 usec
DE 6.50 usec
TE 300.0 K
D1 2.0000000 sec
D11 0.0300000 sec
TD0 1

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NUC1 13C
P1 7.50 usec
PLW1 75.0000000 W

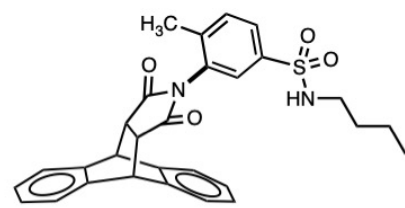
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NUC2 1H
CPDPRG[2] waltz16
PCPD2 90.00 usec
PLW2 18.0000000 W
PLW12 0.19220001 W
PLW13 0.09667500 W

F2 - Processing parameters
SI 32768
SF 100.6127690 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40

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7.443
7.435
7.430
7.422
7.394
7.385
7.381
7.372
7.363
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7.344
7.286
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Current Data Parameters
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EXPNO 1
PROCNO 1

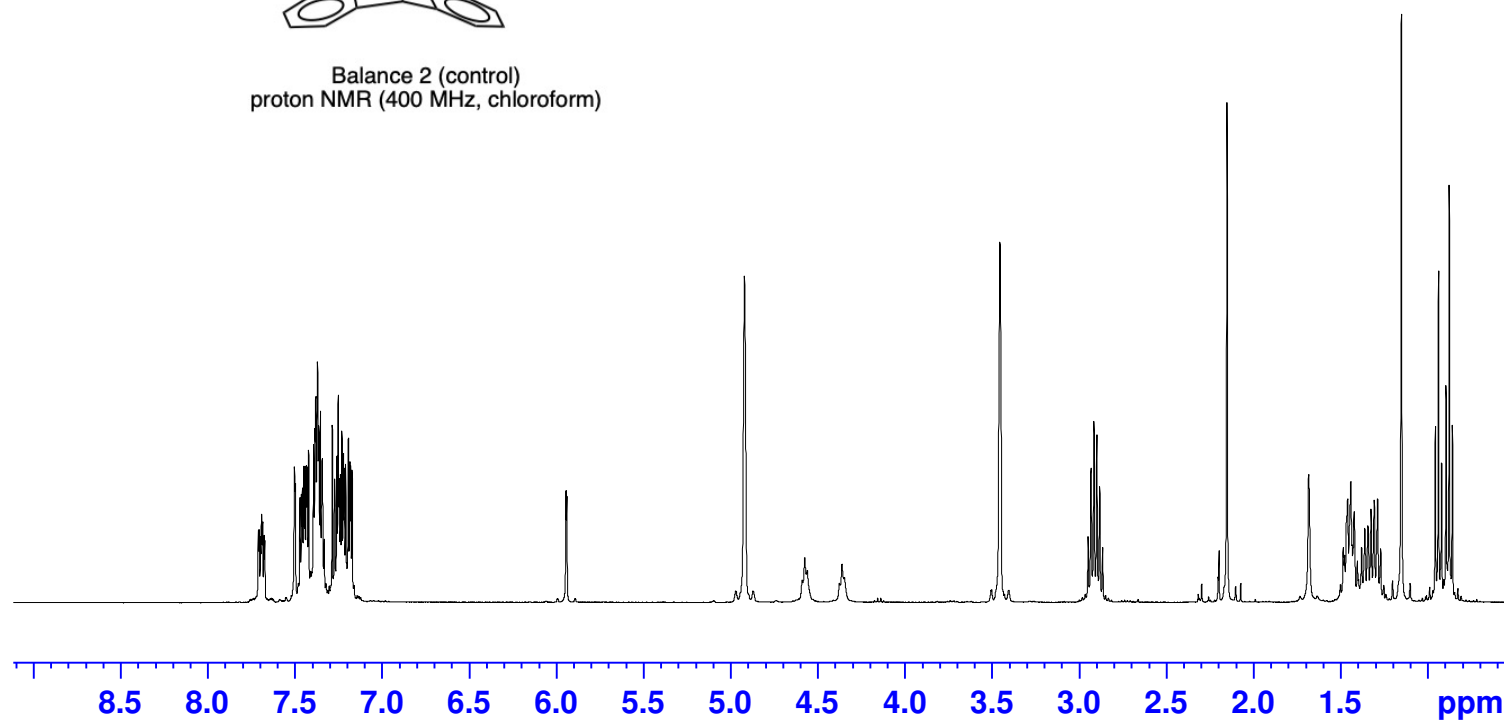
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DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 144
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1



Balance 2 (control)
proton NMR (400 MHz, chloroform)

===== CHANNEL f1 =====
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NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



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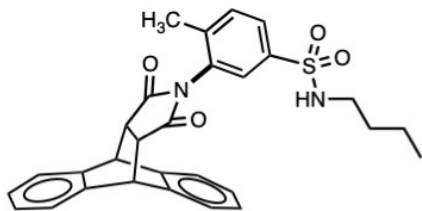
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124.31

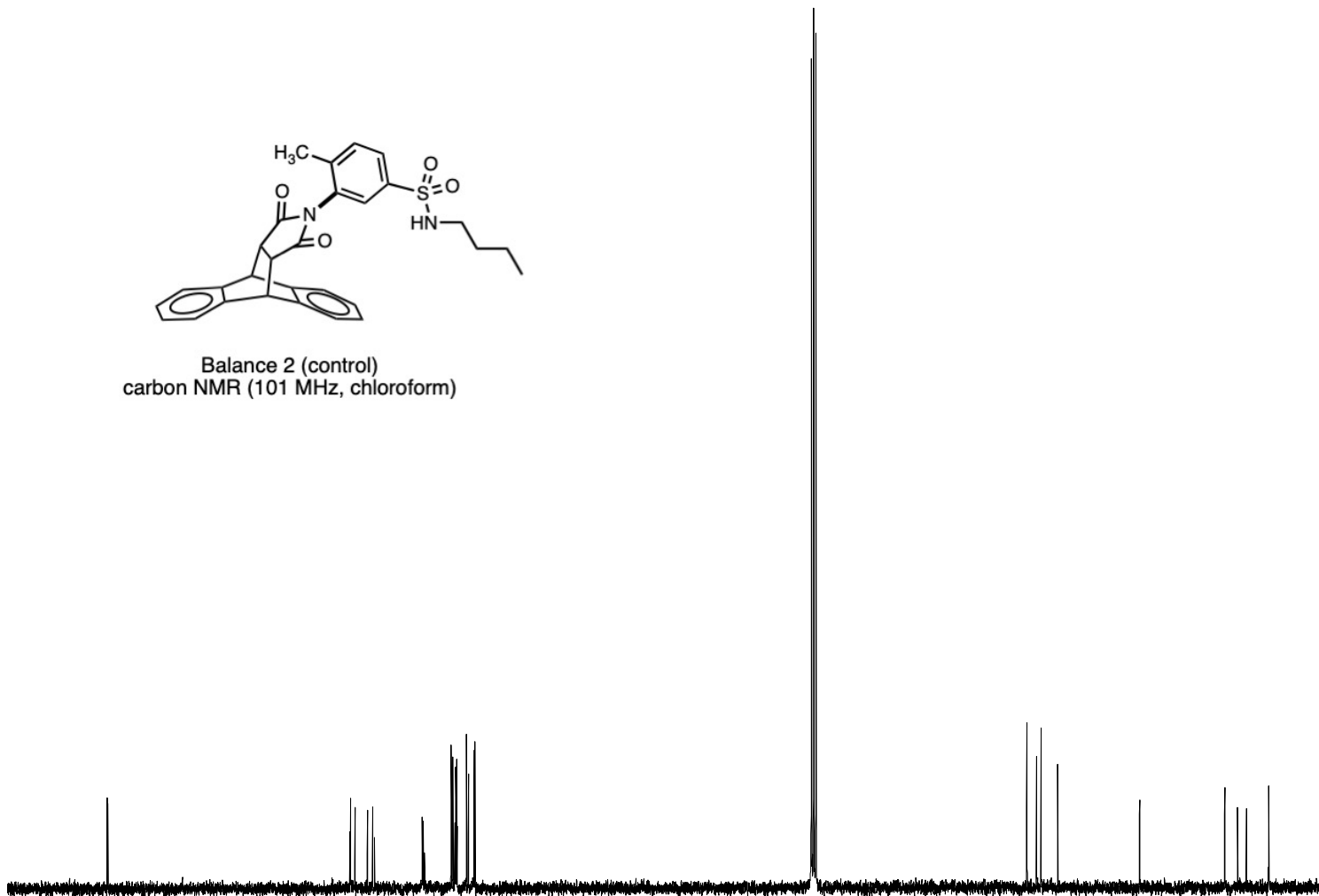
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16.68
13.61
13.55



Balance 2 (control)
carbon NMR (101 MHz, chloroform)



Current Data Parameters
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EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
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PULPROG zgpg30
TD 65536
SOLVENT CDC13
NS 383
DS 4
SWH 24038.461 Hz
FIDRES 0.366798 Hz
AQ 1.3631488 sec
RG 2050
DW 20.800 usec
DE 6.50 usec
TE 300.0 K
D1 2.00000000 sec
D11 0.03000000 sec
TDO 1

==== CHANNEL f1 =====
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NUC1 13C
P1 7.50 usec
PLW1 75.00000000 W

==== CHANNEL f2 =====
SFO2 400.1316005 MHz
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CPDPRG[2] waltz16
PCPD2 90.00 usec
PLW2 18.00000000 W
PLW12 0.19220001 W
PLW13 0.09667500 W

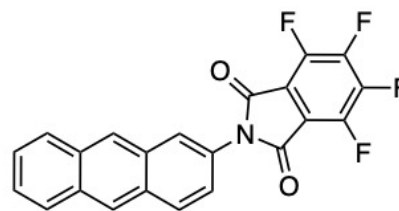
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SI 32768
SF 100.6127690 MHz
WDW EM
SSB 0
LB 1.00 Hz
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8.403

8.035
8.016

7.569
7.528
7.506
7.501
7.481
7.463
7.445
7.431



6

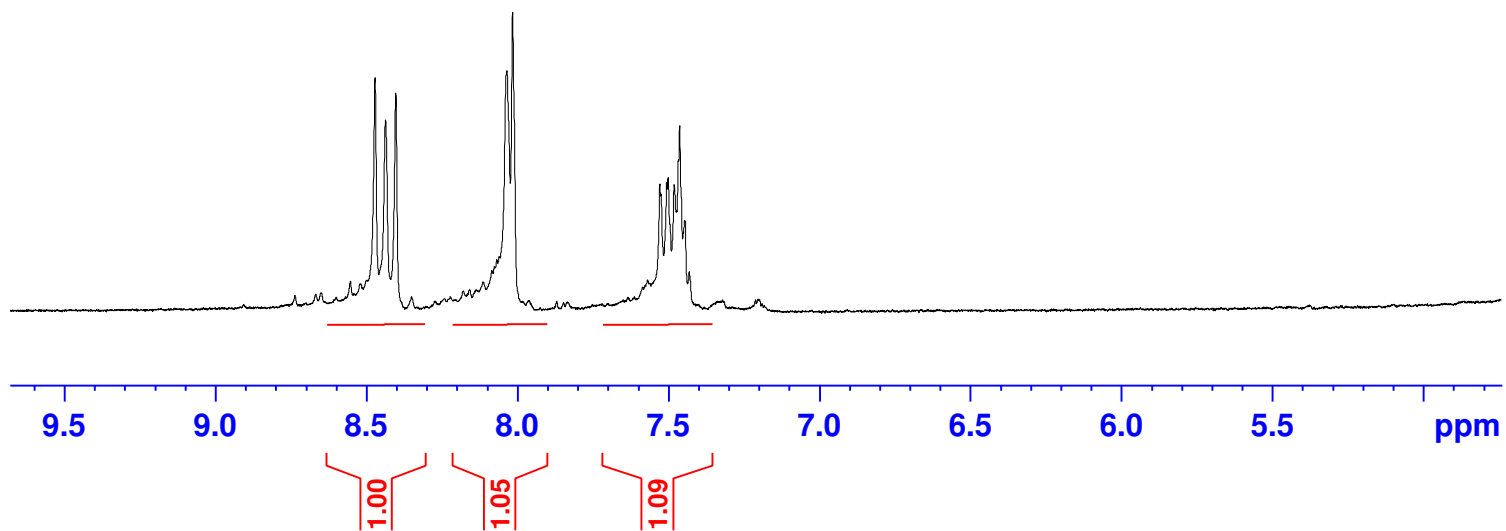
proton NMR (400 MHz, DMSO-d6)

Current Data Parameters
NAME controljan1162023balance1PROTON
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20230116
Time 16.36
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 32
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

===== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

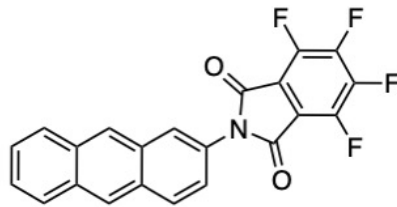
F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



169.88

136.40
132.12
131.99
130.85
129.31
128.93
128.50
128.13
126.31
126.21
125.53
125.38
121.39

114.41



6

Carbon NMR (101 MHz, DMSO-d6)

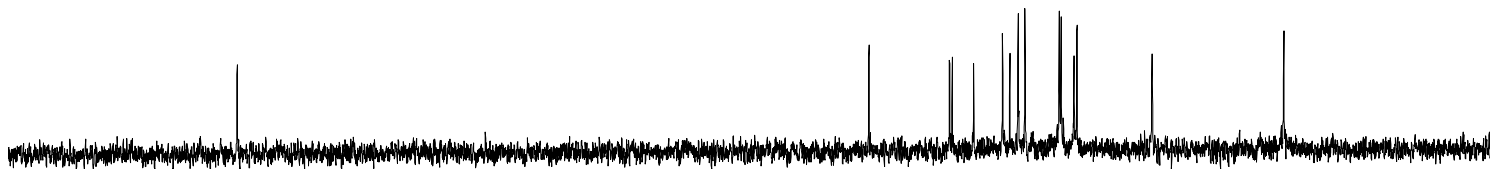
Current Data Parameters
NAME controljan1162023balance1CARBON
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20230116
Time 19.18
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zgpg30
TD 65536
SOLVENT DMSO
NS 2782
DS 4
SWH 24038.461 Hz
FIDRES 0.366798 Hz
AQ 1.3631488 sec
RG 2050
DW 20.800 usec
DE 6.50 usec
TE 300.0 K
D1 2.00000000 sec
D11 0.03000000 sec
TD0 1

===== CHANNEL f1 =====
SFO1 100.6228293 MHz
NUC1 13C
P1 7.50 usec
PLW1 75.00000000 W

===== CHANNEL f2 =====
SFO2 400.1316005 MHz
NUC2 1H
CPDPRG[2] waltz16
PCPD2 90.00 usec
PLW2 18.00000000 W
PLW12 0.19220001 W
PLW13 0.09667500 W

F2 - Processing parameters
SI 32768
SF 100.6127690 MHz
WDW EM
SSB 0
LB 1.00 Hz
GB 0
PC 1.40



180 175 170 165 160 155 150 145 140 135 130 125 120 115 110 ppm

7.837
7.832
7.817
7.812
7.662
7.658
7.499
7.479
6.917

4.809
4.794
4.779

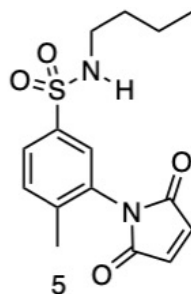
3.000
2.983
2.967
2.950
2.248
1.499
1.482
1.474
1.463
1.456
1.444
1.426
1.355
1.337
1.318

Current Data Parameters
NAME controljan142023proton2
EXPNO 1
PROCNO 1

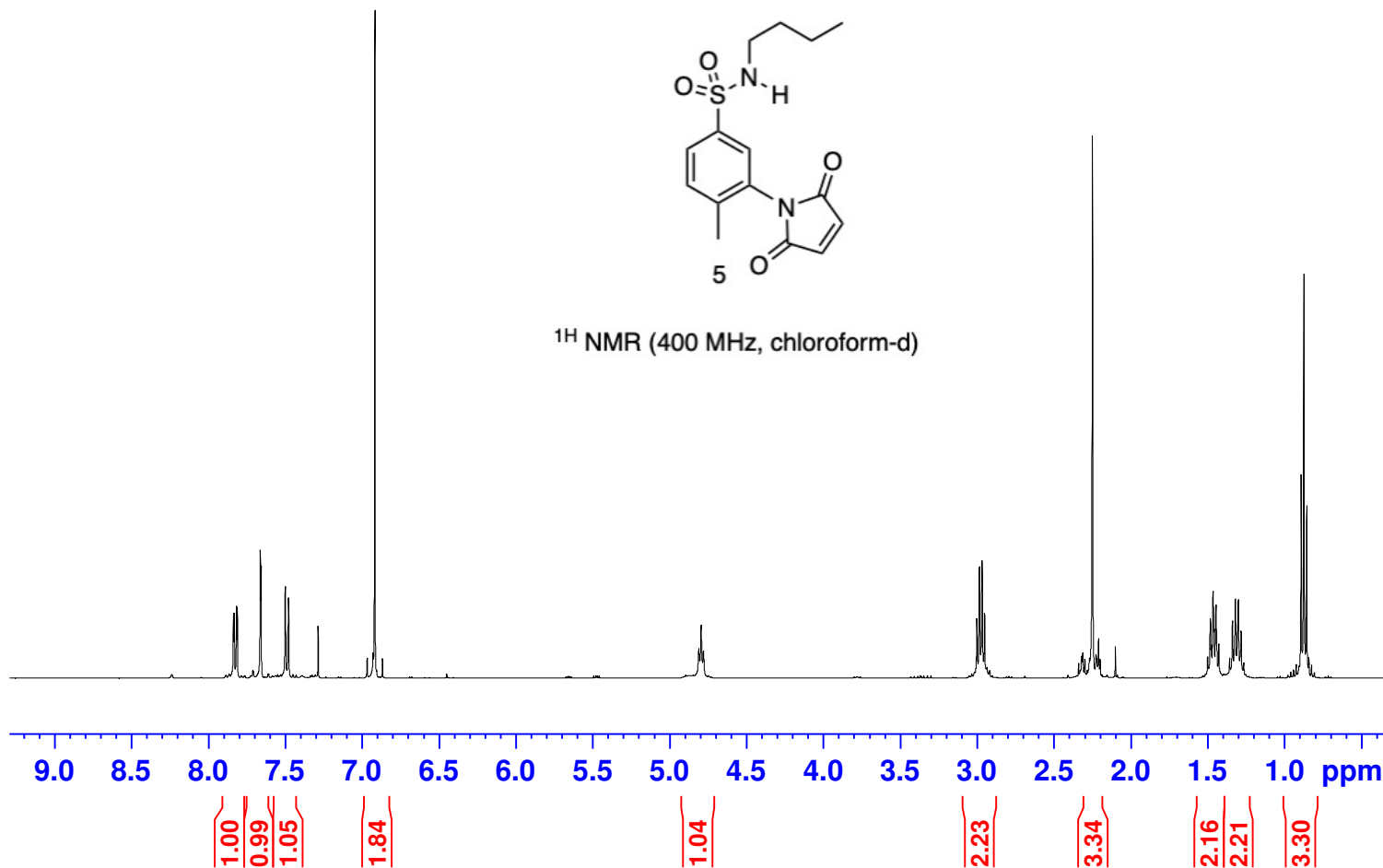
F2 - Acquisition Parameters
Date_ 20230114
Time 16.14
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT CDC13
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 114
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

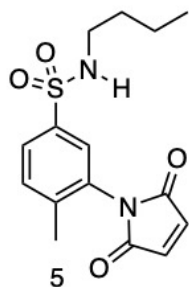
==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

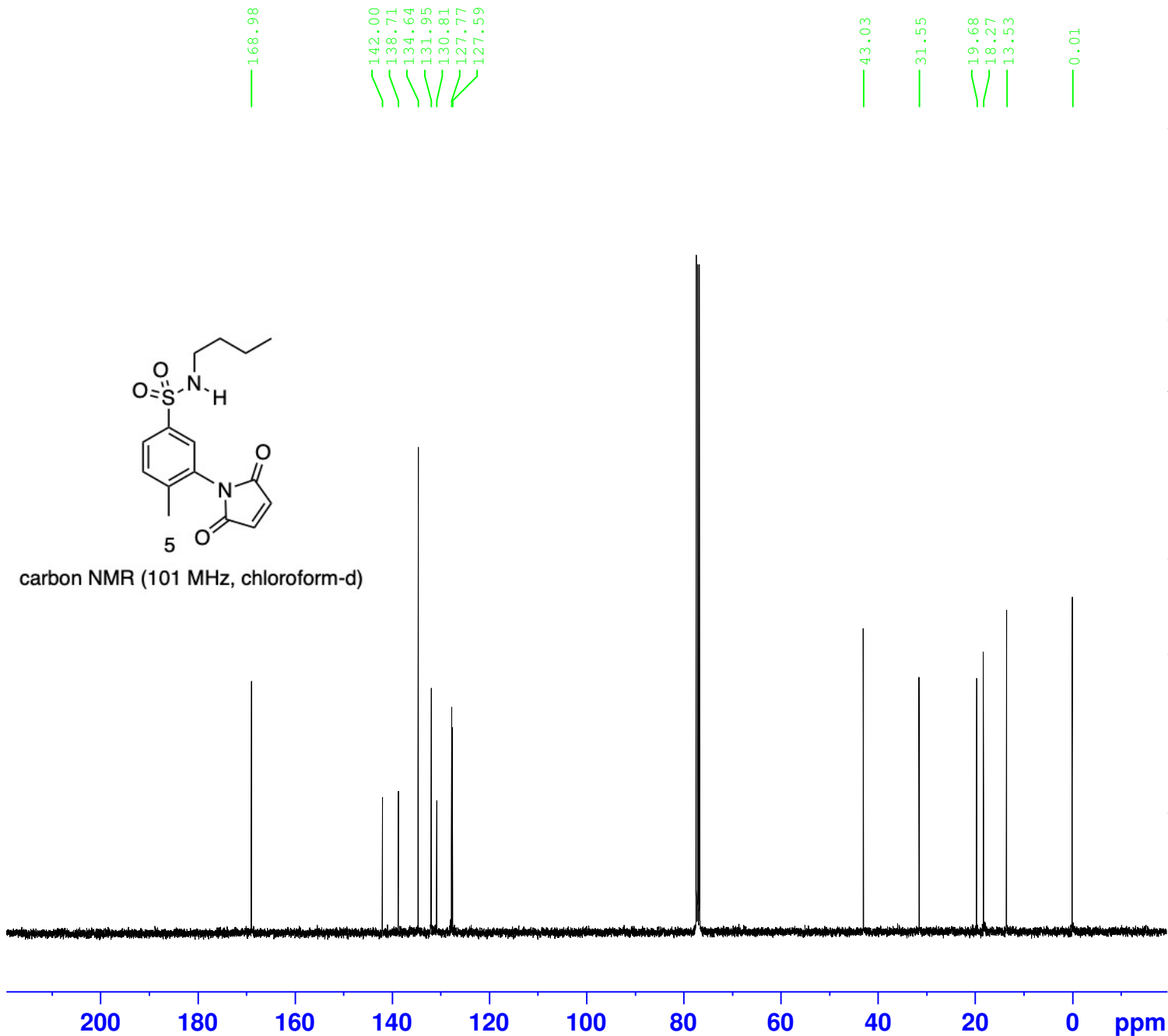


¹H NMR (400 MHz, chloroform-d)





carbon NMR (101 MHz, chloroform-d)



Current Data Parameters
 NAME controljan142023carbon2
 EXPNO 1
 PROCNO 1

F2 - Acquisition Parameters
 Date_ 20230114
 Time 16.50
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zgpg30
 TD 65536
 SOLVENT CDCl3
 NS 561
 DS 4
 SWH 24038.461 Hz
 FIDRES 0.366798 Hz
 AQ 1.3631488 sec
 RG 2050
 DW 20.800 usec
 DE 6.50 usec
 TE 300.0 K
 D1 2.00000000 sec
 D11 0.03000000 sec
 TD0 1

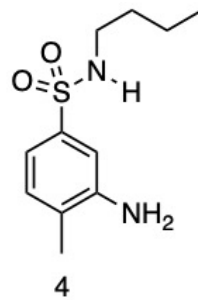
===== CHANNEL f1 =====
 SFO1 100.6228293 MHz
 NUC1 13C
 P1 7.50 usec
 PLW1 75.00000000 W

===== CHANNEL f2 =====
 SFO2 400.1316005 MHz
 NUC2 1H
 CPDPRG[2] waltz16
 PCPD2 90.00 usec
 PLW2 18.00000000 W
 PLW12 0.19220001 W
 PLW13 0.09667500 W

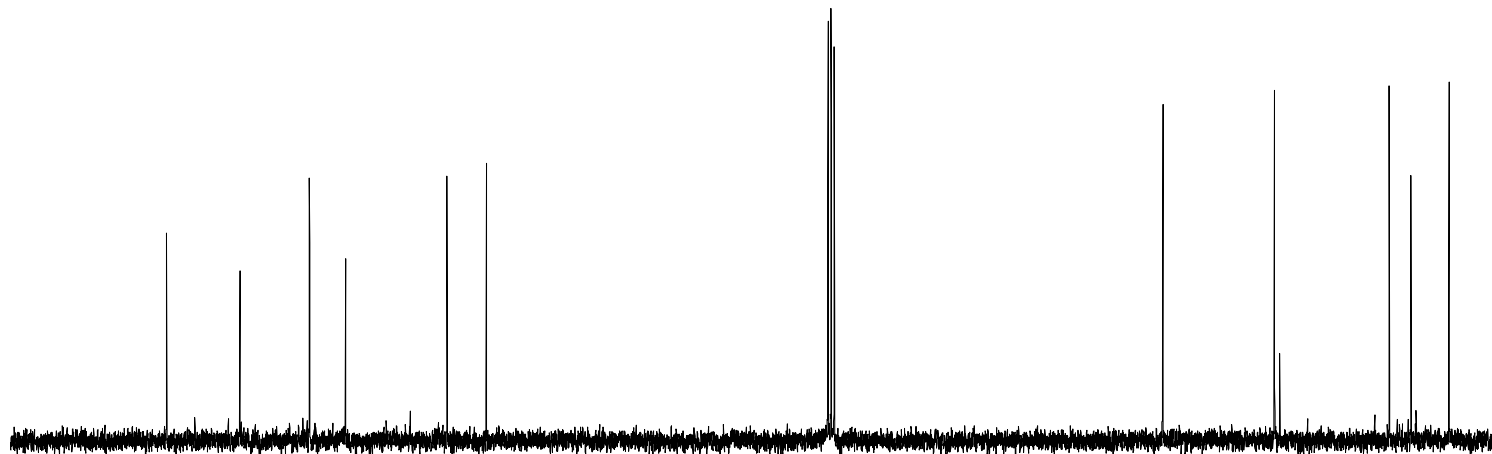
F2 - Processing parameters
 SI 32768
 SF 100.6127690 MHz
 WDW EM
 SSB 0
 LB 1.00 Hz
 GB 0
 PC 1.40

145.48
137.94
130.78
127.07
116.64
112.59

42.99
31.50
19.72
17.47
13.56



carbon NMR (101 MHz, chloroform-d)



150 140 130 120 110 100 90 80 70 60 50 40 30 20 ppm

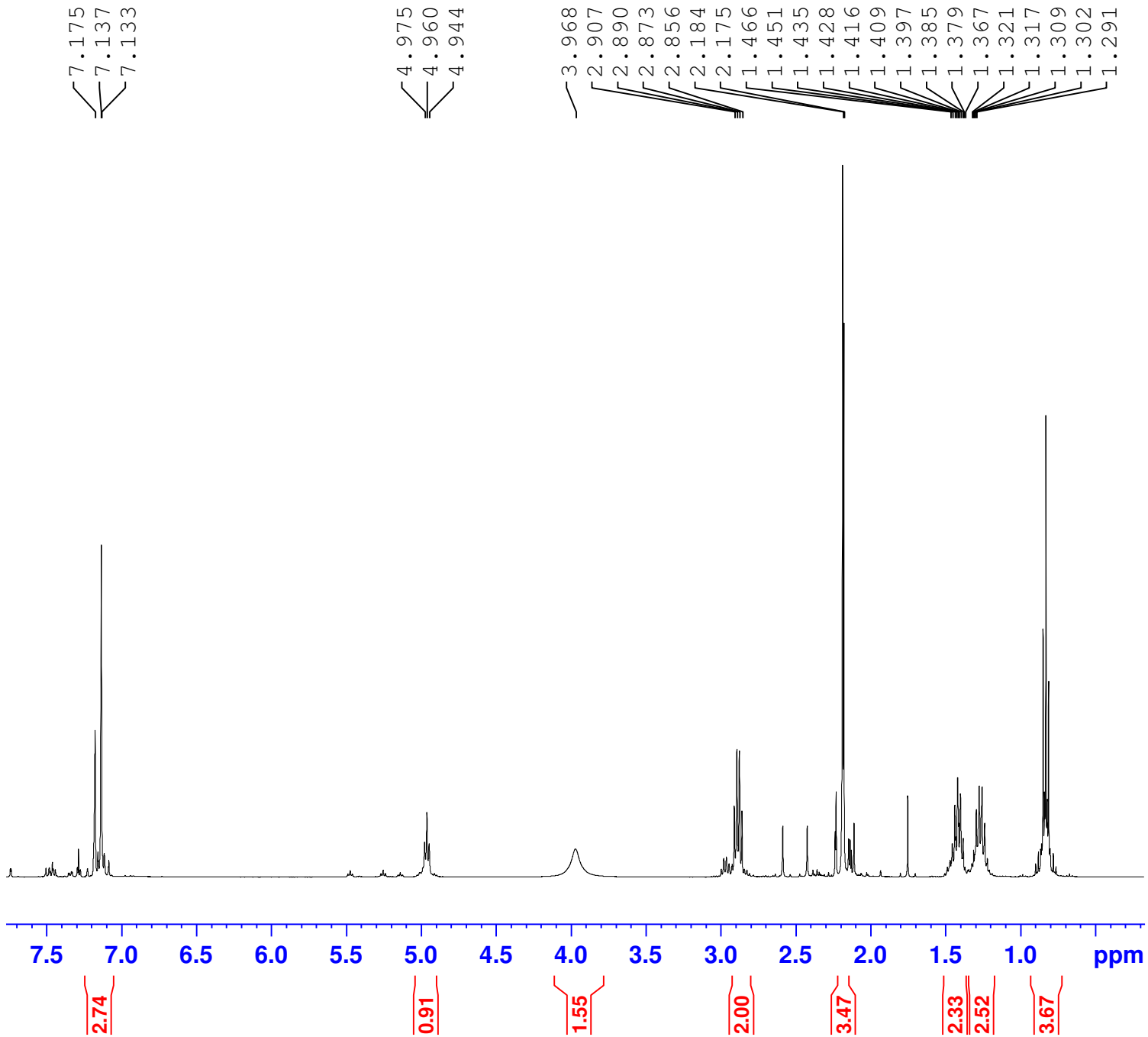
```
Current Data Parameters
NAME      N-butyl(3-methyl-4-amino)benzenesulfanamide-carbon
EXPNO     1
PROCNO    1

F2 - Acquisition Parameters
Date_     20230124
Time      19.37
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   zgpg30
TD         65536
SOLVENT   CDCl3
NS         25
DS         4
SWH        24038.461 Hz
FIDRES     0.366798 Hz
AQ         1.3631488 sec
RG         2050
DW         20.800 usec
DE         6.50 usec
TE         300.0 K
D1         2.00000000 sec
D11        0.03000000 sec
TD0        1

----- CHANNEL f1 -----
SFO1      100.6228293 MHz
NUC1       13C
P1         7.50 usec
PLW1       75.00000000 W

----- CHANNEL f2 -----
SFO2      400.1316005 MHz
NUC2       1H
CPDPRG2    waltz16
PCPD2      90.00 usec
PLW2       18.00000000 W
PLW12      0.19220001 W
PLW13      0.09667500 W

F2 - Processing parameters
SI         32768
SF         100.6127690 MHz
WDW        EM
SSB        0
LB         1.00 Hz
GB         0
PC         1.40
```



```

Current Data Parameters
NAME      N-butyl(3-methyl-4-amino)benzenesulfanamide-prot
EXPNO     1
PROCNO    1

F2 - Acquisition Parameters
Date_     20230124
Time      19.34
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   zg30
TD         65536
SOLVENT   CDC13
NS         7
DS         2
SWH        8012.820 Hz
FIDRES     0.122266 Hz
AQ         4.0894465 sec
RG         71.8
DW         62.400 usec
DE         6.50 usec
TE         298.0 K
D1         1.00000000 sec
TDO        1

----- CHANNEL f1 -----
SFO1      400.1324710 MHz
NUC1      1H
P1         9.30 usec
PLW1      18.00000000 W

F2 - Processing parameters
SI         65536
SF         400.1300000 MHz
WCM        EM
SSB        0
LB         0.30 Hz
GB         0
PC         1.00
  
```

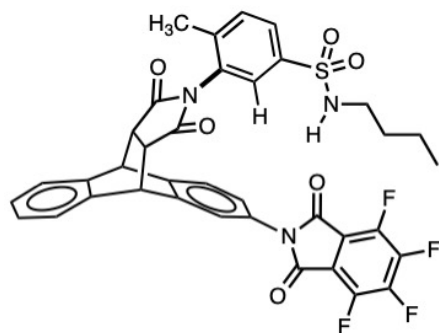


Current Data Parameters
NAME conc-study-9.9mMOLAR-toluene
EXPNO 1
PROCNO 1

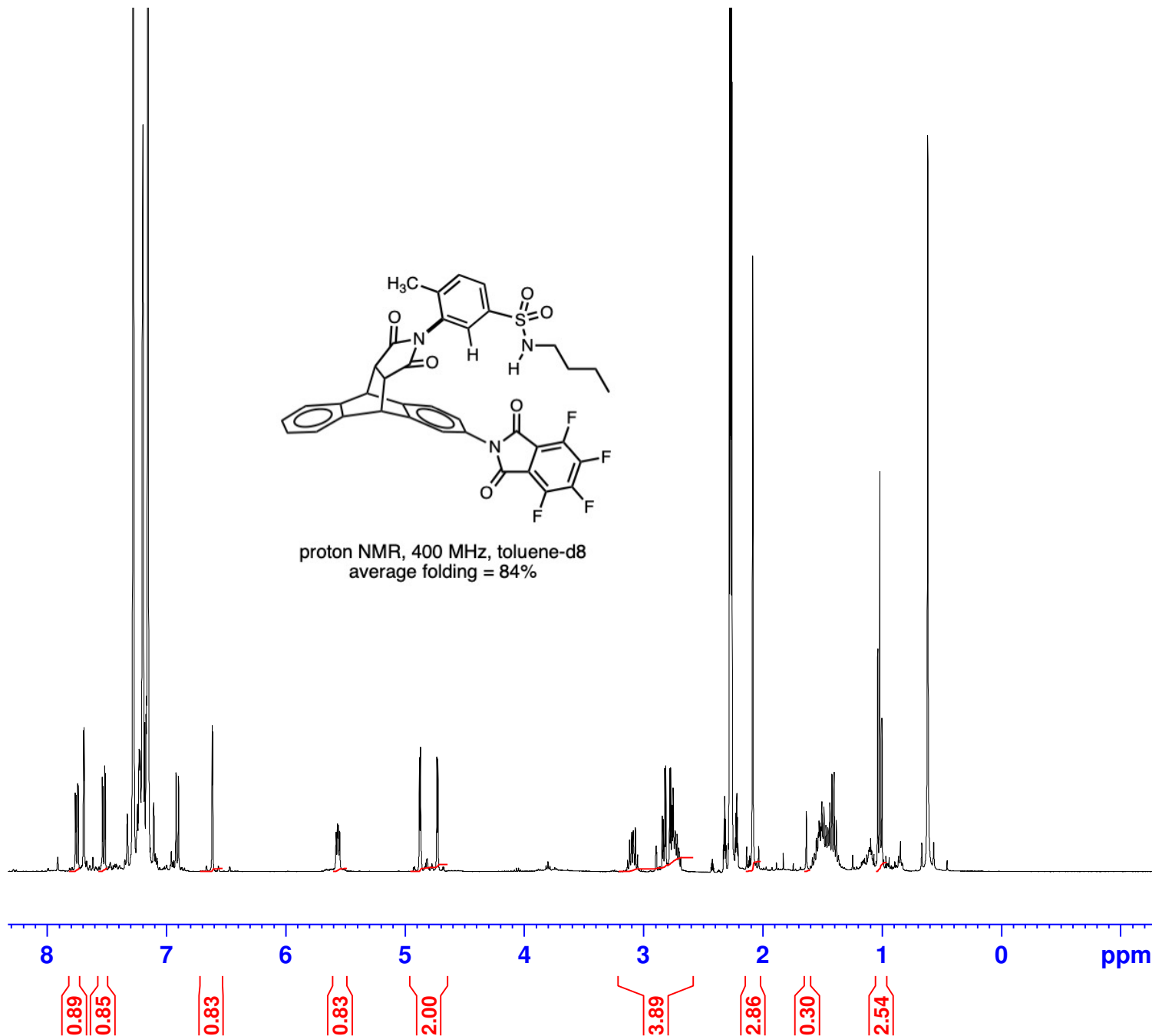
F2 - Acquisition Parameters
Date_ 20211013
Time 18.38
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT Tol
NS 800
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 512
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1299439 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



proton NMR, 400 MHz, toluene-d8
average folding = 84%

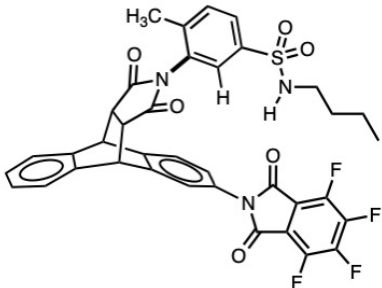


7.502
7.471
7.451
7.442
7.419
7.405
7.397
7.376
7.370
7.308
7.288
7.195
7.175
7.164
7.153
7.129
6.062

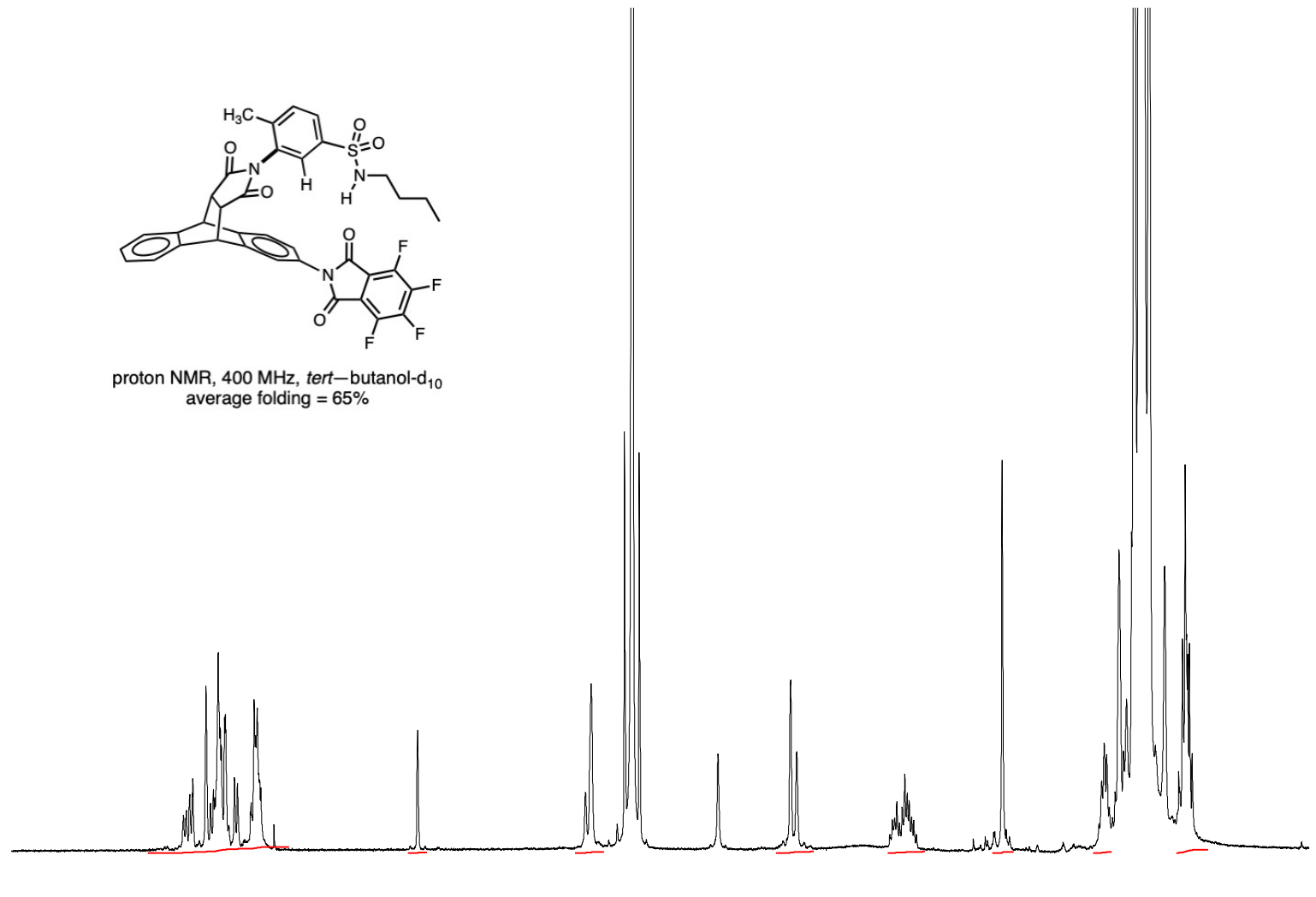
4.922
4.883

3.525
3.484
2.835
2.818
2.804
2.786
2.768
2.750
2.733
2.720
2.703
2.689
2.088

0.862
0.844
0.826
0.814
0.796



proton NMR, 400 MHz, *tert*-butanol-*d*₁₀
average folding = 65%



8.5 8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 ppm

9.15 0.65 2.00 2.06 2.03 1.85 1.87 5.19

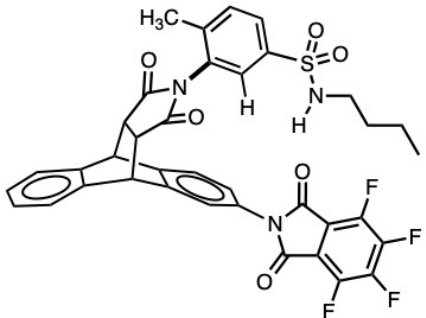
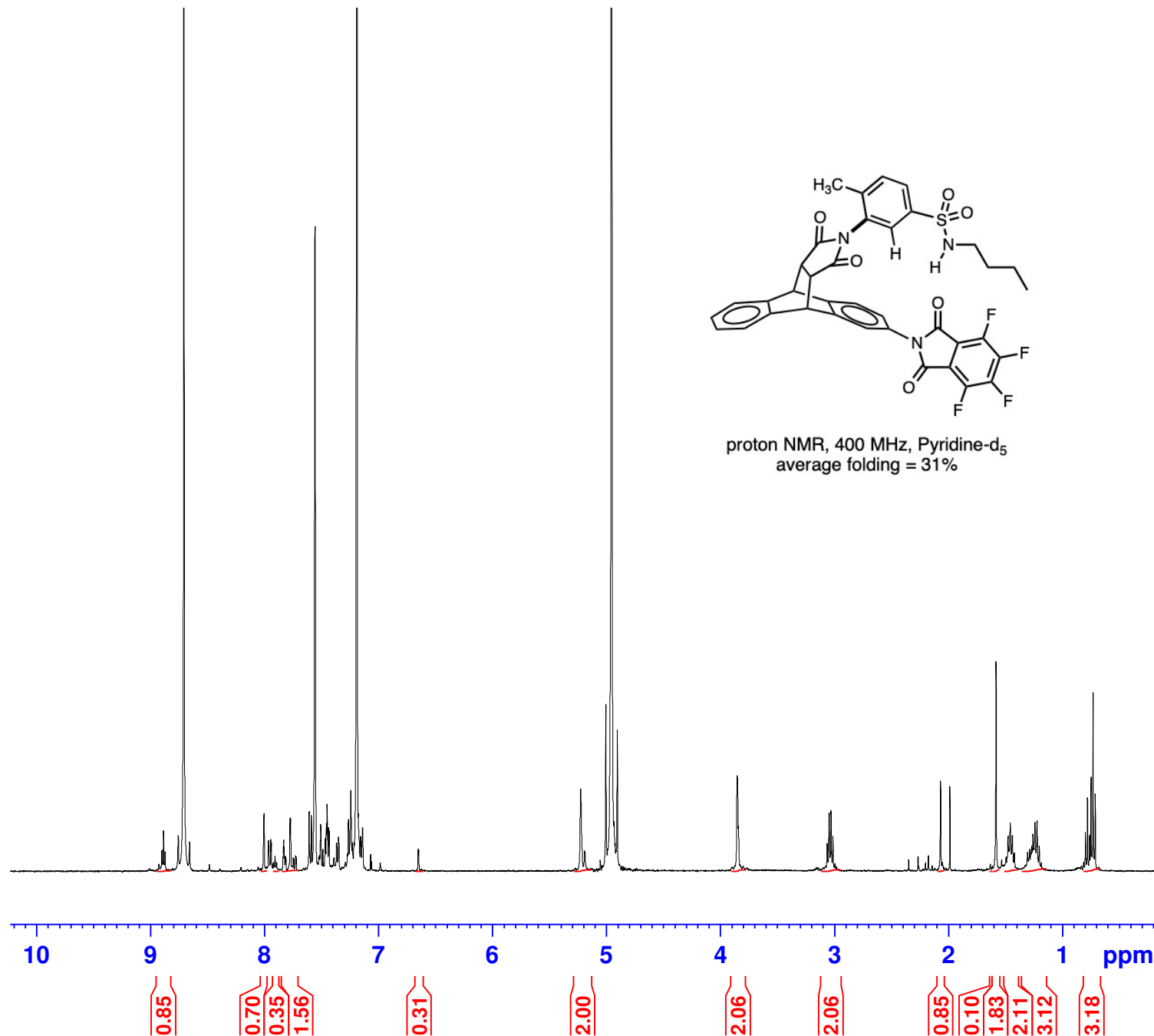
```

Current Data Parameters
NAME      conc-study~9.9mMOLAR-tertbutanol1
EXPNO     1
PROCNO    1

F2 - Acquisition Parameters
Date_     20221107
Time      15.19
INSTRUM   spect
PROBHD    5 mm PABBO BB-
PULPROG   zg30
TD         65536
SOLVENT   EtOH
NS         320
DS         2
SWH        8012.820 Hz
FIDRES     0.122266 Hz
AQ         4.0894465 sec
RG         256
DW         62.400 usec
DE         6.50 usec
TE         298.0 K
D1         1.00000000 sec
TDO        1

===== CHANNEL f1 =====
SFO1      400.1324710 MHz
NUC1       1H
P1         9.30 usec
PLW1      18.00000000 W

F2 - Processing parameters
SI         65536
SF         400.1300000 MHz
WDW        EM
SSB        0
LB         0.30 Hz
GB         0
PC         1.00
  
```



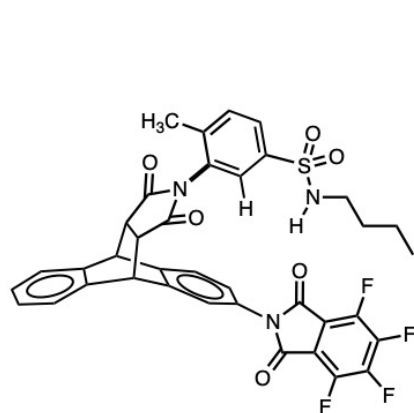
proton NMR, 400 MHz, Pyridine-d₅
average folding = 31%

Current Data Parameters
 NAME emenikemodella-inpyridine3mg
 EXPNO 1
 PROCNO 1

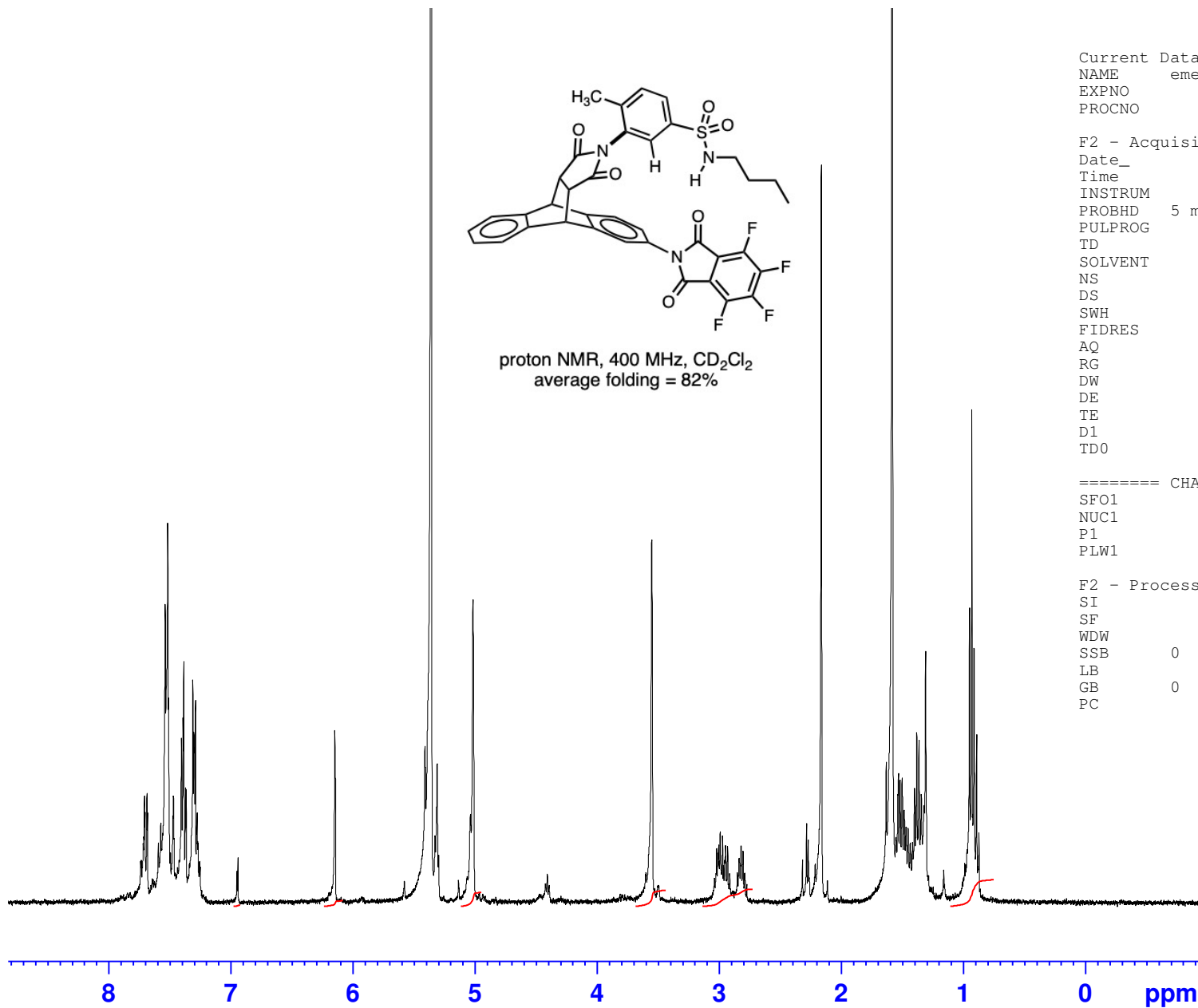
F2 - Acquisition Parameters
 Date_ 20221031
 Time 17.43
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zg30
 TD 65536
 SOLVENT Pyr
 NS 16
 DS 2
 SWH 8012.820 Hz
 FIDRES 0.122266 Hz
 AQ 4.0894465 sec
 RG 287
 DW 62.400 usec
 DE 6.50 usec
 TE 298.0 K
 D1 1.00000000 sec
 TD0 1

===== CHANNEL f1 =====
 SFO1 400.1324710 MHz
 NUC1 1H
 P1 9.30 usec
 PLW1 18.00000000 W

F2 - Processing parameters
 SI 65536
 SF 400.1300000 MHz
 WDW EM
 SSB 0
 LB 0.30 Hz
 GB 0
 PC 1.00



proton NMR, 400 MHz, CD₂Cl₂
average folding = 82%



0.17

0.82

2.00

2.29

2.44

3.78

Current Data Parameters

NAME emenikemodella-indichloromethane3mg
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters

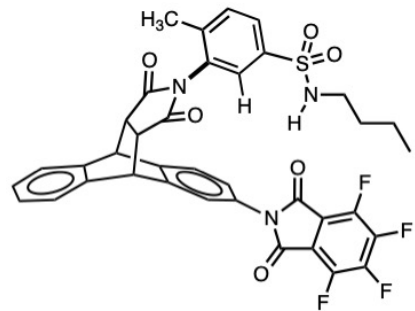
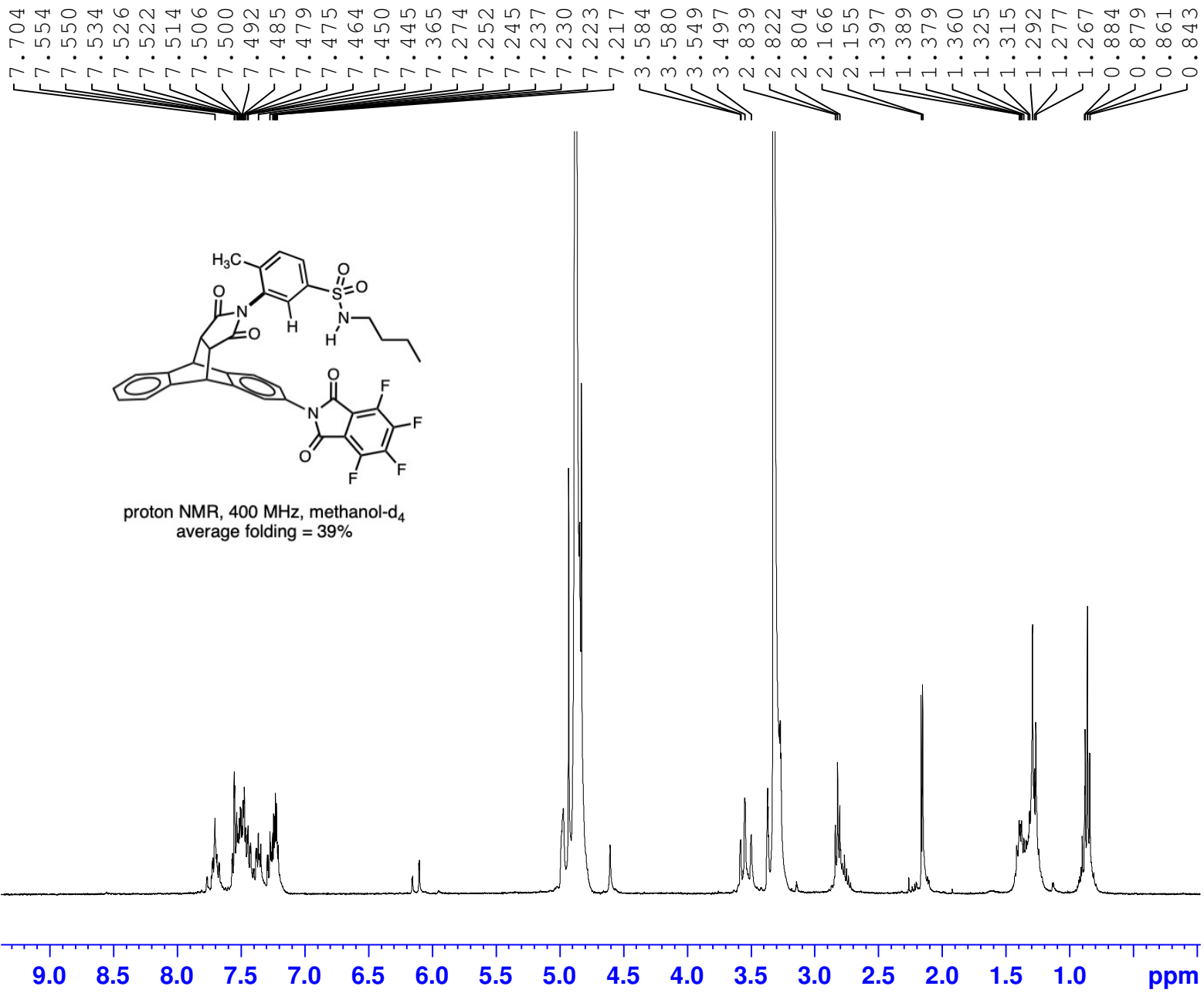
Date_ 20221031
Time 16.38
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT CD2Cl2
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 575
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====

SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters

SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



proton NMR, 400 MHz, methanol-d₄
average folding = 39%

Current Data Parameters
 NAME conc-study-9.9mMOLAR-MeOH-1
 EXPNO 1
 PROCNO 1

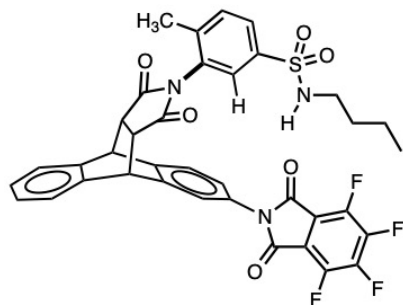
F2 - Acquisition Parameters
 Date_ 20211009
 Time 17.48
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zg30
 TD 65536
 SOLVENT MeOD
 NS 103
 DS 2
 SWH 8012.820 Hz
 FIDRES 0.122266 Hz
 AQ 4.0894465 sec
 RG 512
 DW 62.400 usec
 DE 6.50 usec
 TE 298.0 K
 D1 1.00000000 sec
 TD0 1

==== CHANNEL f1 =====
 SFO1 400.1324710 MHz
 NUC1 1H
 P1 9.30 usec
 PLW1 18.00000000 W

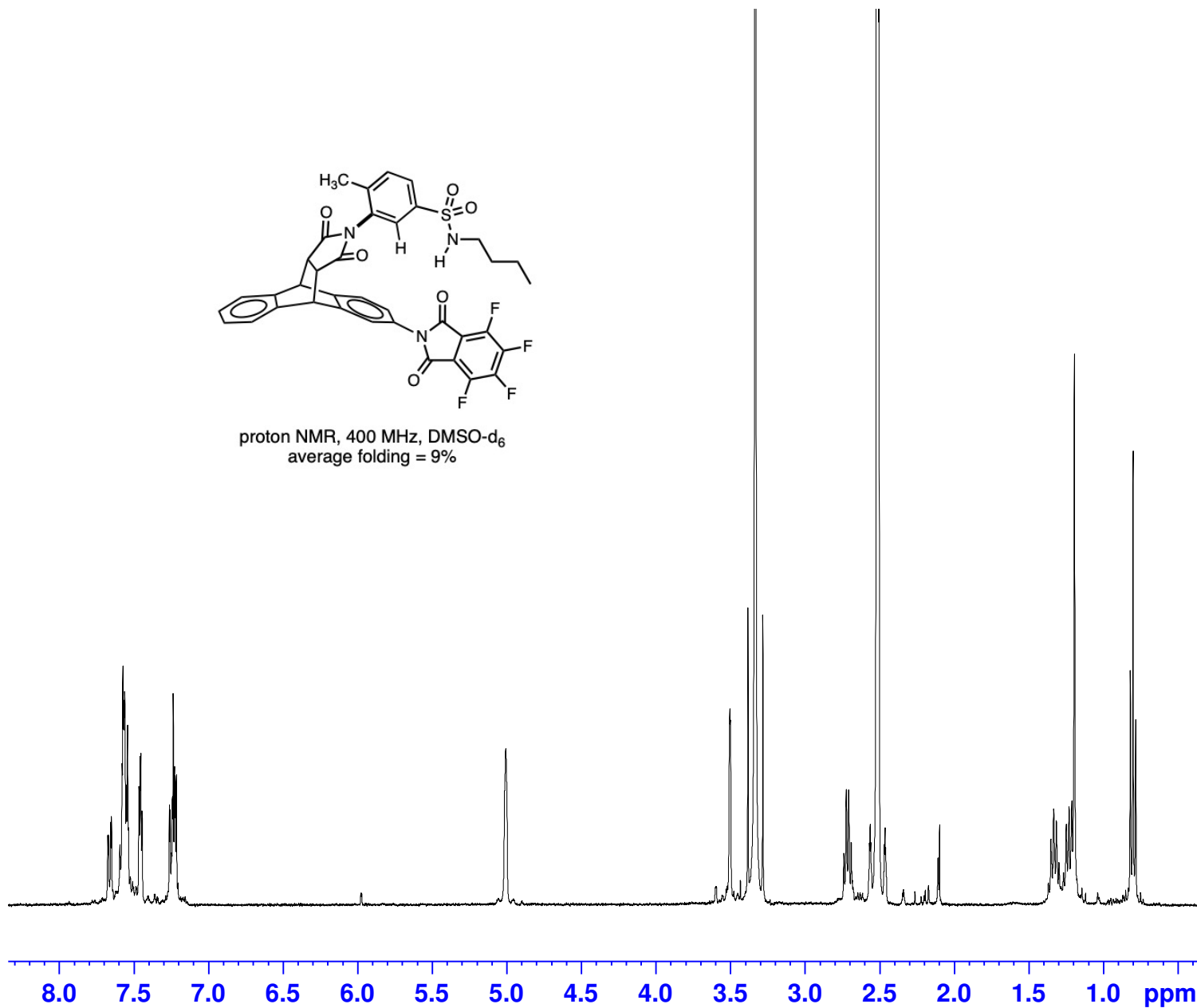
F2 - Processing parameters
 SI 65536
 SF 400.1300033 MHz
 WDW EM
 SSB 0
 LB 0.30 Hz
 GB 0
 PC 1.00

9.0 8.5 8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 ppm

9.09 0.39 2.01 2.13 0.30 0.74 6.37 3.00



proton NMR, 400 MHz, DMSO-d₆
average folding = 9%



8.00

3.07

0.09

2.00

1.94

1.98

6.83

3.12

Current Data Parameters
NAME emenikemodella-indmso3
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20221031
Time 16.20
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 322
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.0000000 sec
TD0 1

==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.0000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

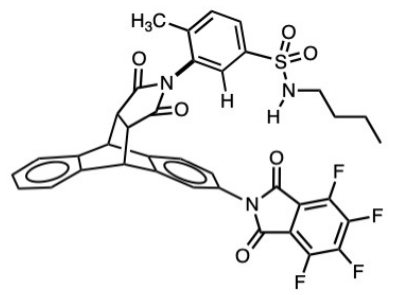
8.075
7.772
7.767
7.752
7.747
7.669
7.665
7.634
7.613
7.600
7.589
7.556
7.551
7.489
7.472
7.459
7.370
7.365
7.350
7.345
7.266
7.258
7.253
7.245
6.173
5.075
2.973
2.968
2.959
2.873
2.853
2.836
2.819
2.801
2.796
2.790
1.435
1.417
1.398
1.380
1.284
1.266
1.249
1.171
0.875
0.862
0.806

Current Data Parameters
NAME conc-study-9.9mMOLAR-dmf-1
EXPNO 1
PROCNO 1

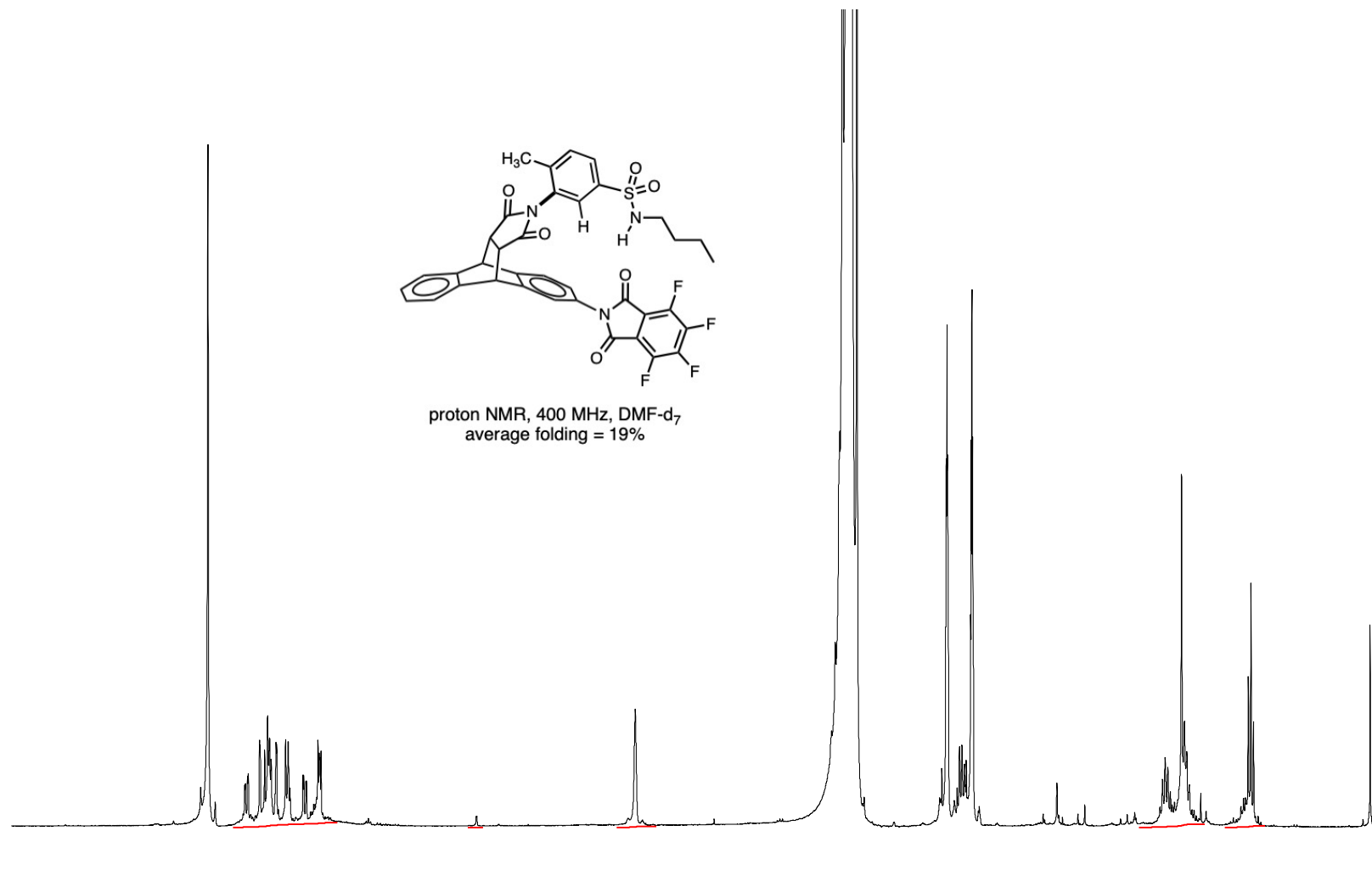
F2 - Acquisition Parameters
Date_ 20211006
Time 17.19
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT DMF
NS 220
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 203
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300033 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

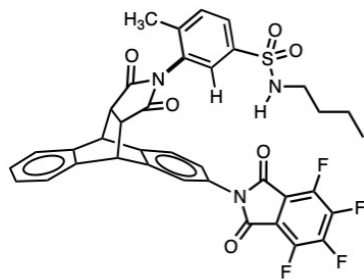


proton NMR, 400 MHz, DMF-d₇
average folding = 19%

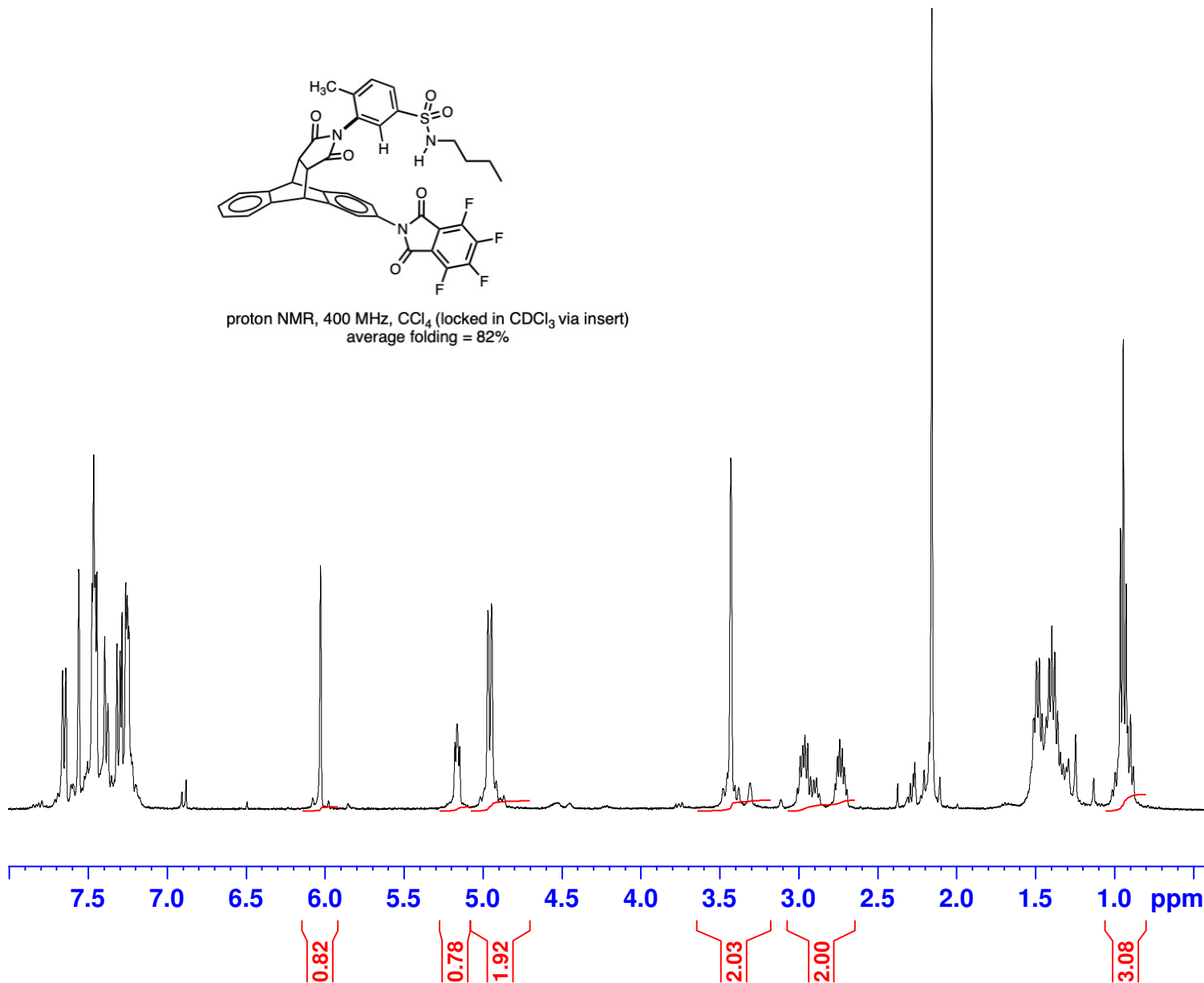


9.0 8.5 8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 ppm

10.22
0.19
2.00
7.37
3.20



proton NMR, 400 MHz, CCl₄ (locked in CDCl₃ via insert)
average folding = 82%

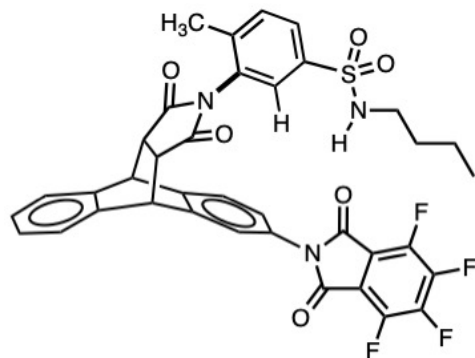


Current Data Parameters
NAME balance-sulfanamide-CCl4
EXPNO 1
PROCNO 1

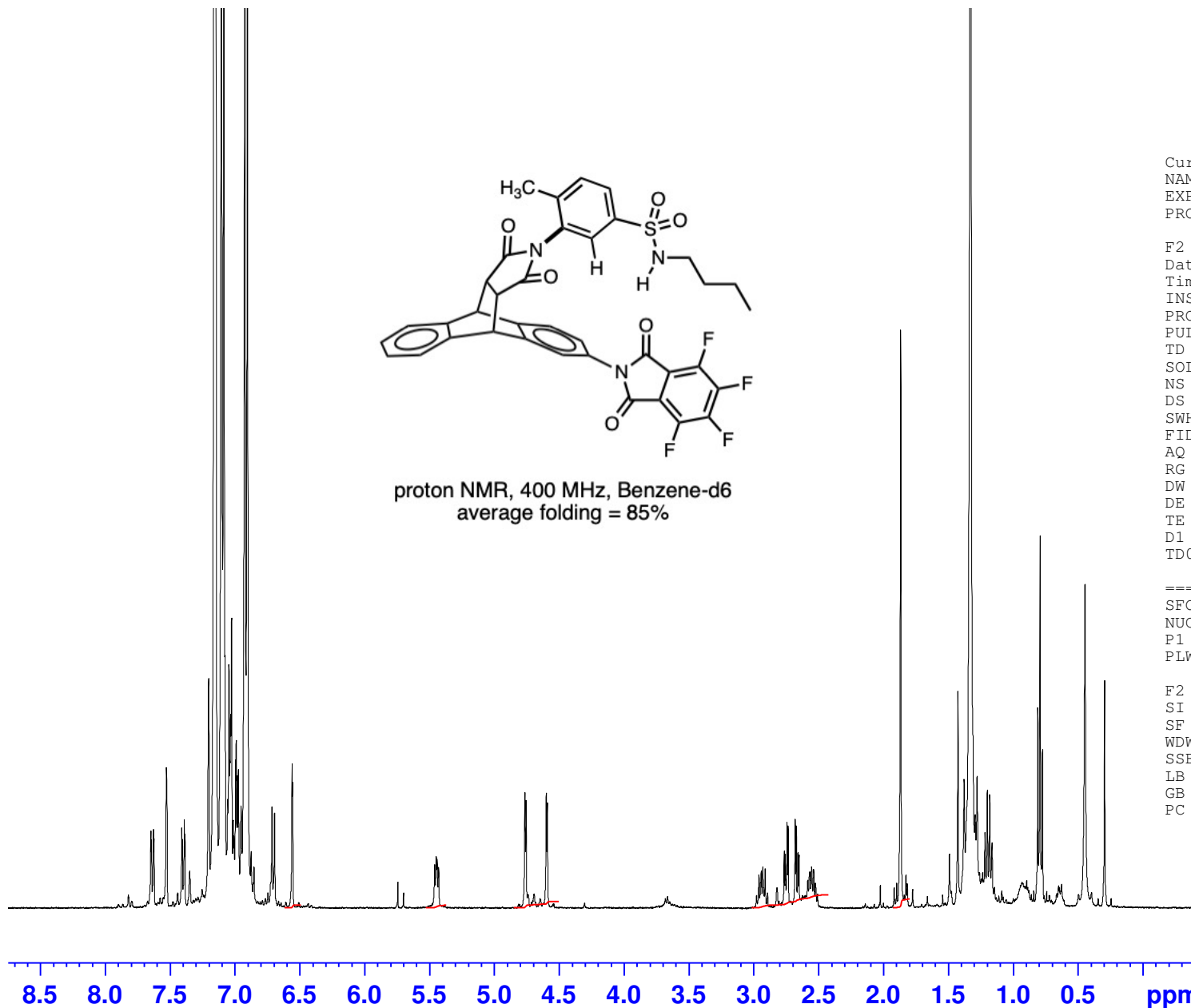
F2 - Acquisition Parameters
Date_ 20221224
Time 15.58
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT CDCl3
NS 48
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 322
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



proton NMR, 400 MHz, Benzene-d6
average folding = 85%



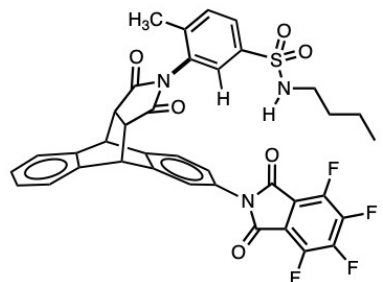
Current Data Parameters
NAME balance-sulfanamide-benzene
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20221224
Time 15.25
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT C6D6
NS 69
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 256
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

7.714
7.709
7.694
7.689
7.563
7.543
7.528
7.524
7.520
7.514
7.512
7.506
7.455
7.442
7.437
7.410
7.263
7.255
7.251
7.246
7.242
7.232
7.226
5.000
3.570
3.566
3.539
3.535
3.531
2.830
2.814
1.386
1.383
1.365
1.296
1.291
1.273
1.271
1.246
0.882
0.863
0.861
0.843
0.825



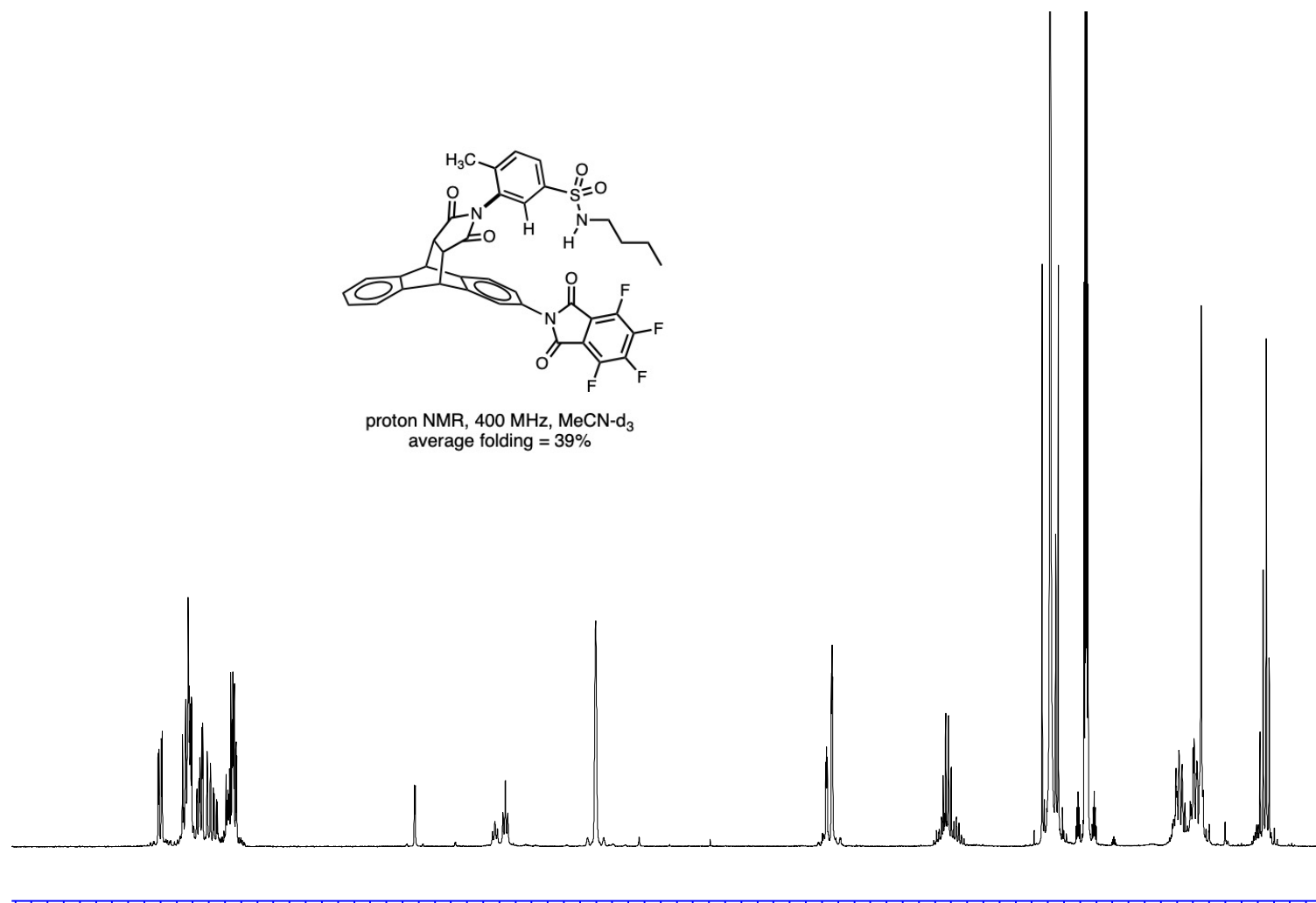
proton NMR, 400 MHz, MeCN-d₃
average folding = 39%

Current Data Parameters
NAME conc-study-9.9mMOLAR-MECN
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20210925
Time 15.34
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT CD3CN
NS 120
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 512
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TDO 1

===== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

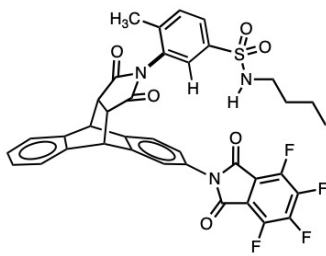
F2 - Processing parameters
SI 65536
SF 400.1300033 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



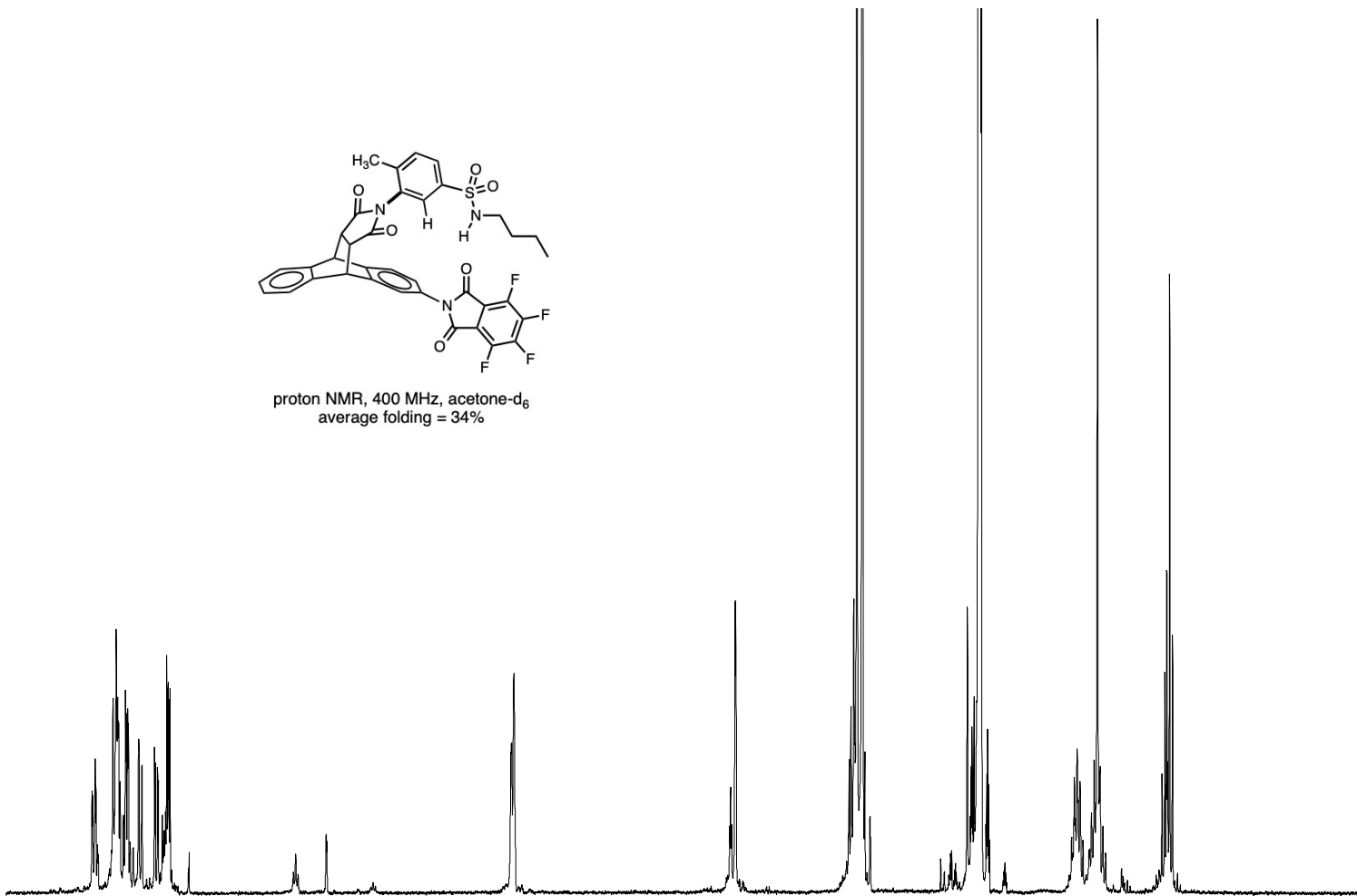
8.5 8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 ppm

9.08
0.39
0.32
0.59
1.88
2.00
2.03
6.03
2.91

7.734
7.729
7.714
7.709
7.600
7.580
7.571
7.565
7.556
7.520
7.515
7.506
7.502
7.436
7.415
7.335
7.330
7.315
7.310
7.264
7.257
7.248
7.243
7.235
5.057
5.041
3.629
3.626
3.623
1.465
1.447
1.440
1.428
1.355
1.338
1.316
1.306
1.299
0.905
0.886
0.874
0.868
0.856
0.837



proton NMR, 400 MHz, acetone-d₆
average folding = 34%



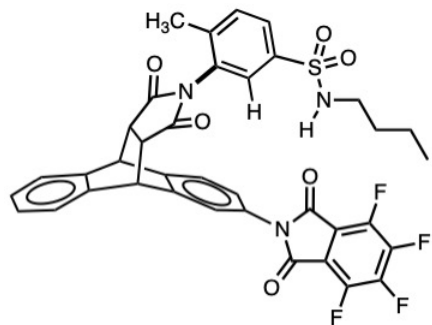
8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 0.5 ppm

Current Data Parameters
NAME emenikemodella-inacetone3mg
EXPNO 1
PROCNO 1

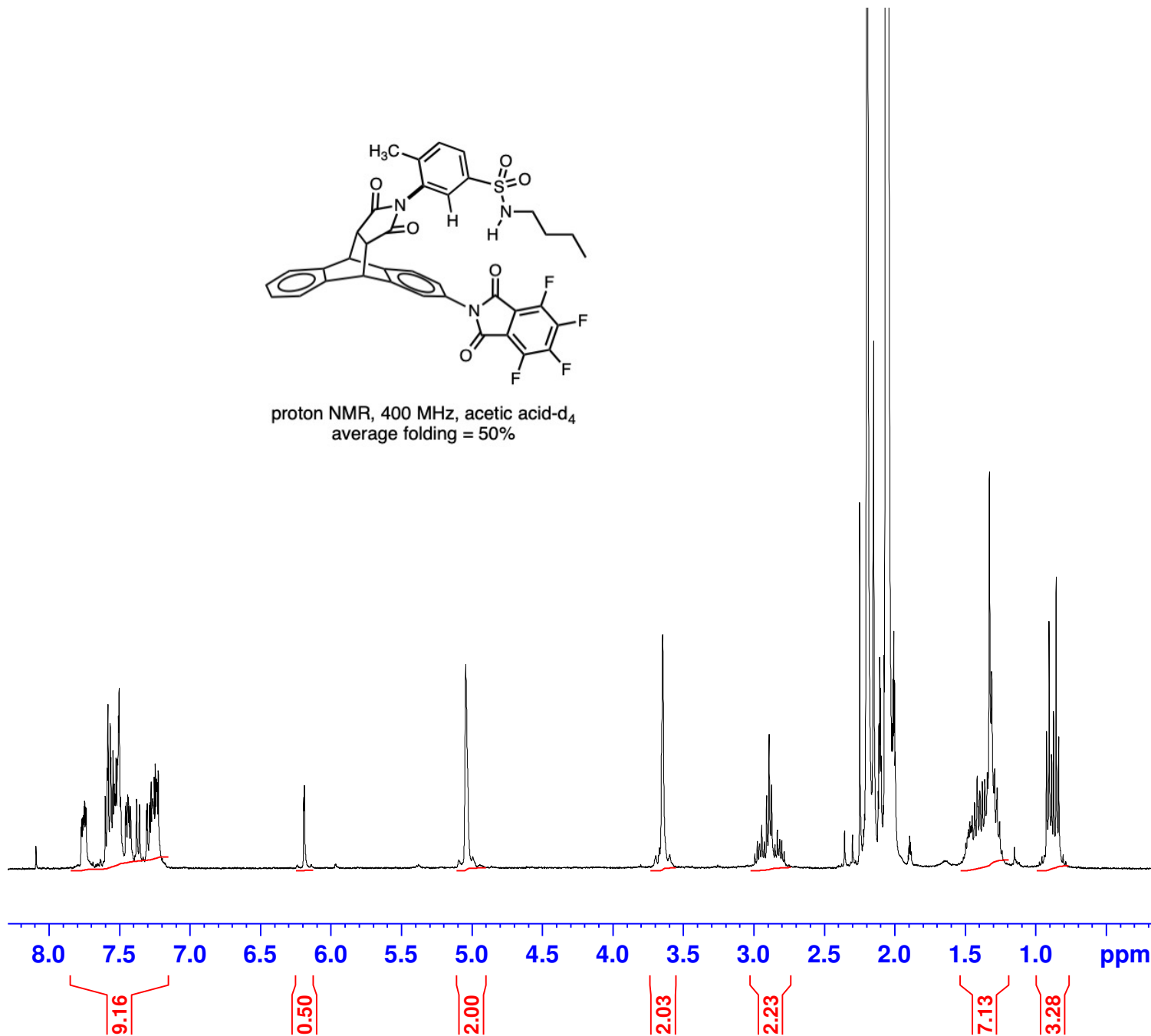
F2 - Acquisition Parameters
Date_ 20221031
Time_ 15.37
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT Acetone
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 362
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



proton NMR, 400 MHz, acetic acid-d₄
average folding = 50%



Current Data Parameters
NAME conc-study-9.9mMOLAR-aceticacid
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters

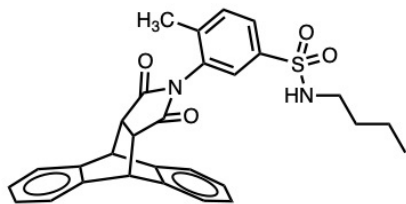
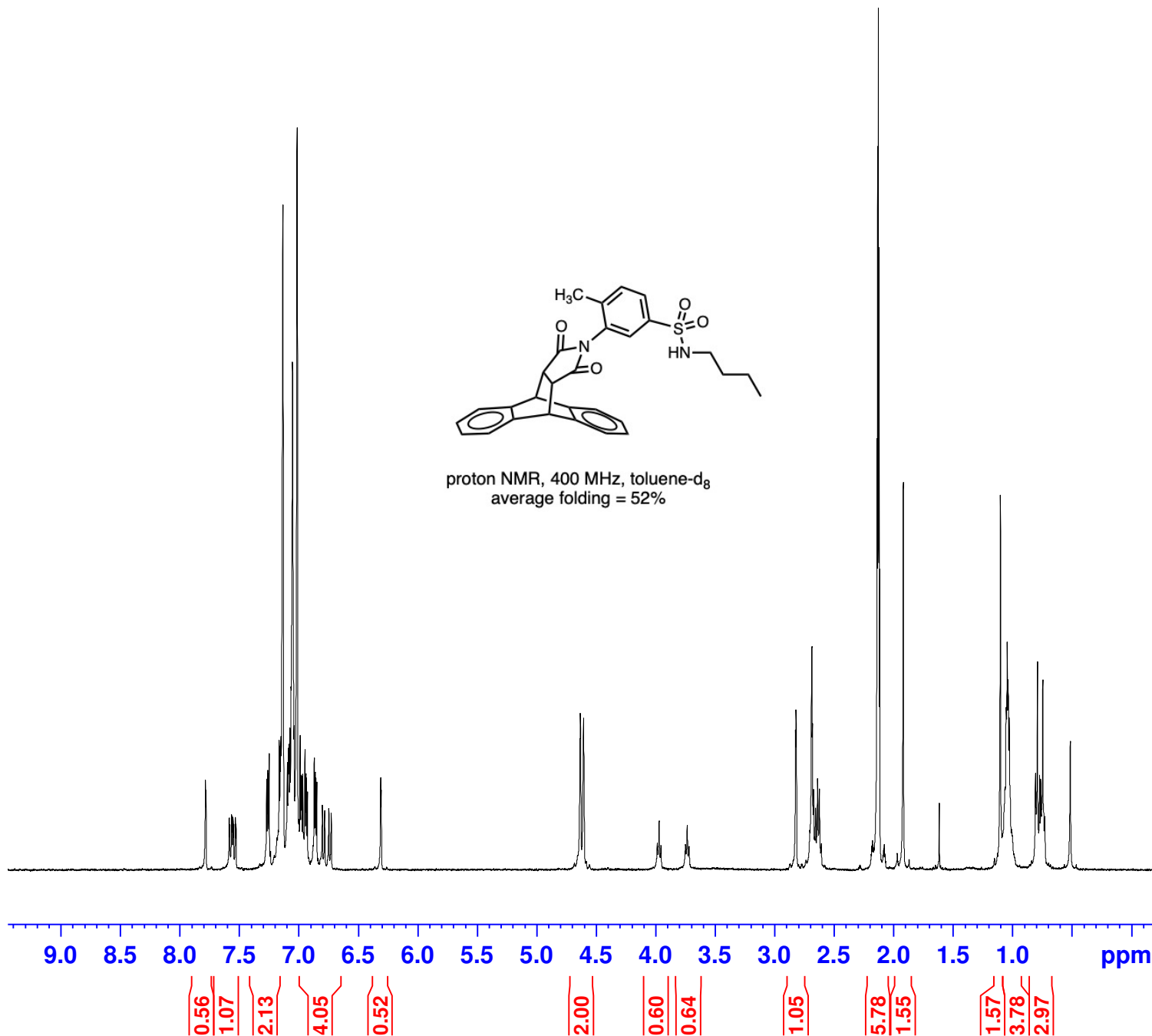
Date_ 20211009
Time 13.13
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT Acetic
NS 150
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 575
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

===== CHANNEL f1 =====

SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters

SI 65536
SF 400.1300033 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

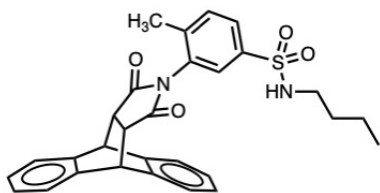


Current Data Parameters
 NAME controlnov19balanceintoluene
 EXPNO 1
 PROCNO 1

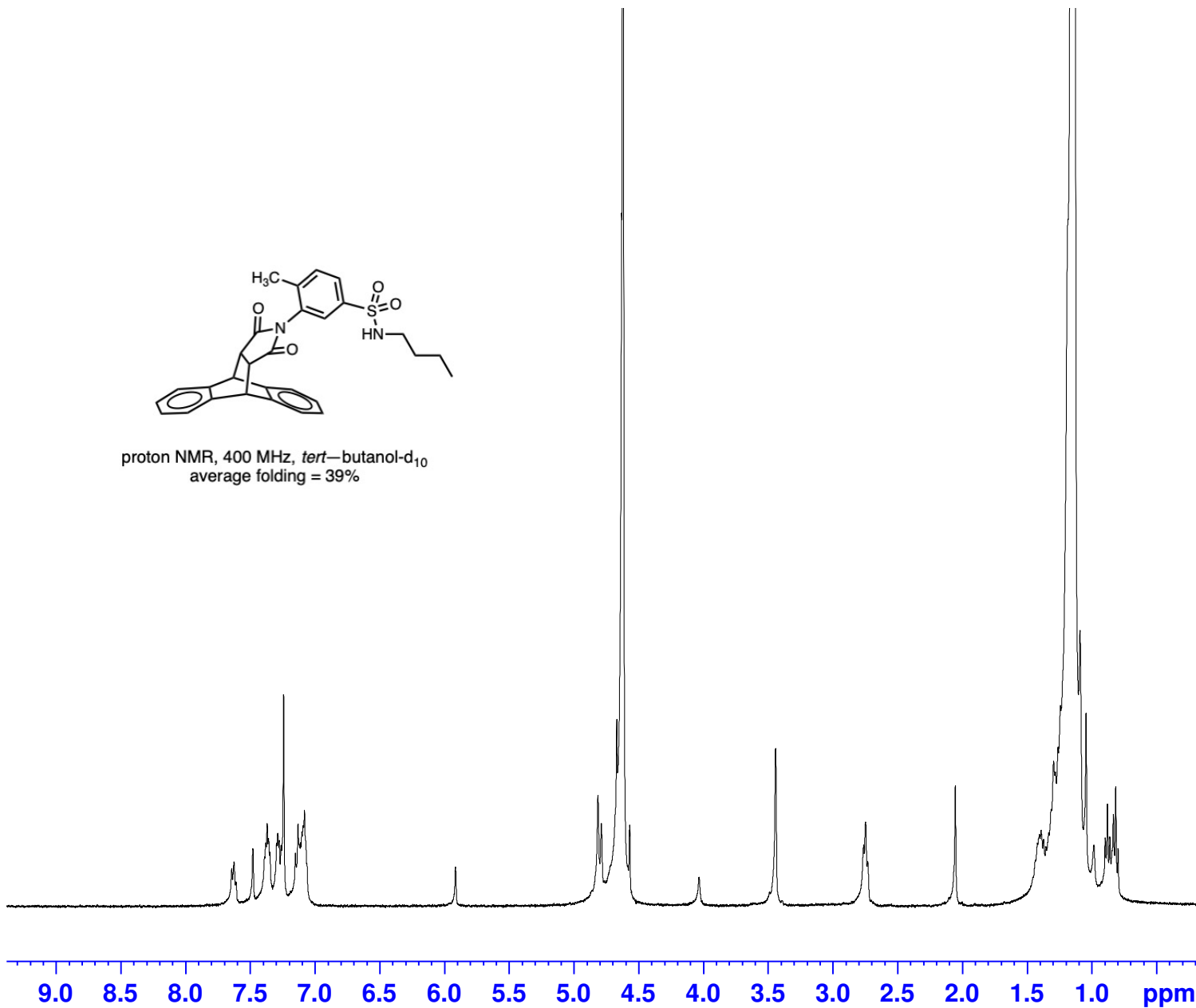
F2 - Acquisition Parameters
 Date_ 20221119
 Time 16.20
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zg30
 TD 65536
 SOLVENT Tol
 NS 16
 DS 2
 SWH 8012.820 Hz
 FIDRES 0.122266 Hz
 AQ 4.0894465 sec
 RG 322
 DW 62.400 usec
 DE 6.50 usec
 TE 298.0 K
 D1 1.00000000 sec
 TD0 1

===== CHANNEL f1 =====
 SFO1 400.1324710 MHz
 NUC1 1H
 P1 9.30 usec
 PLW1 18.00000000 W

F2 - Processing parameters
 SI 65536
 SF 400.1300000 MHz
 WDW EM
 SSB 0
 LB 0.30 Hz
 GB 0
 PC 1.00



proton NMR, 400 MHz, *tert*-butanol-d₁₀
average folding = 39%



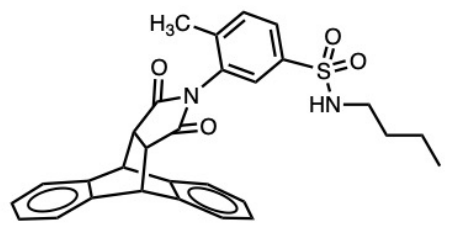
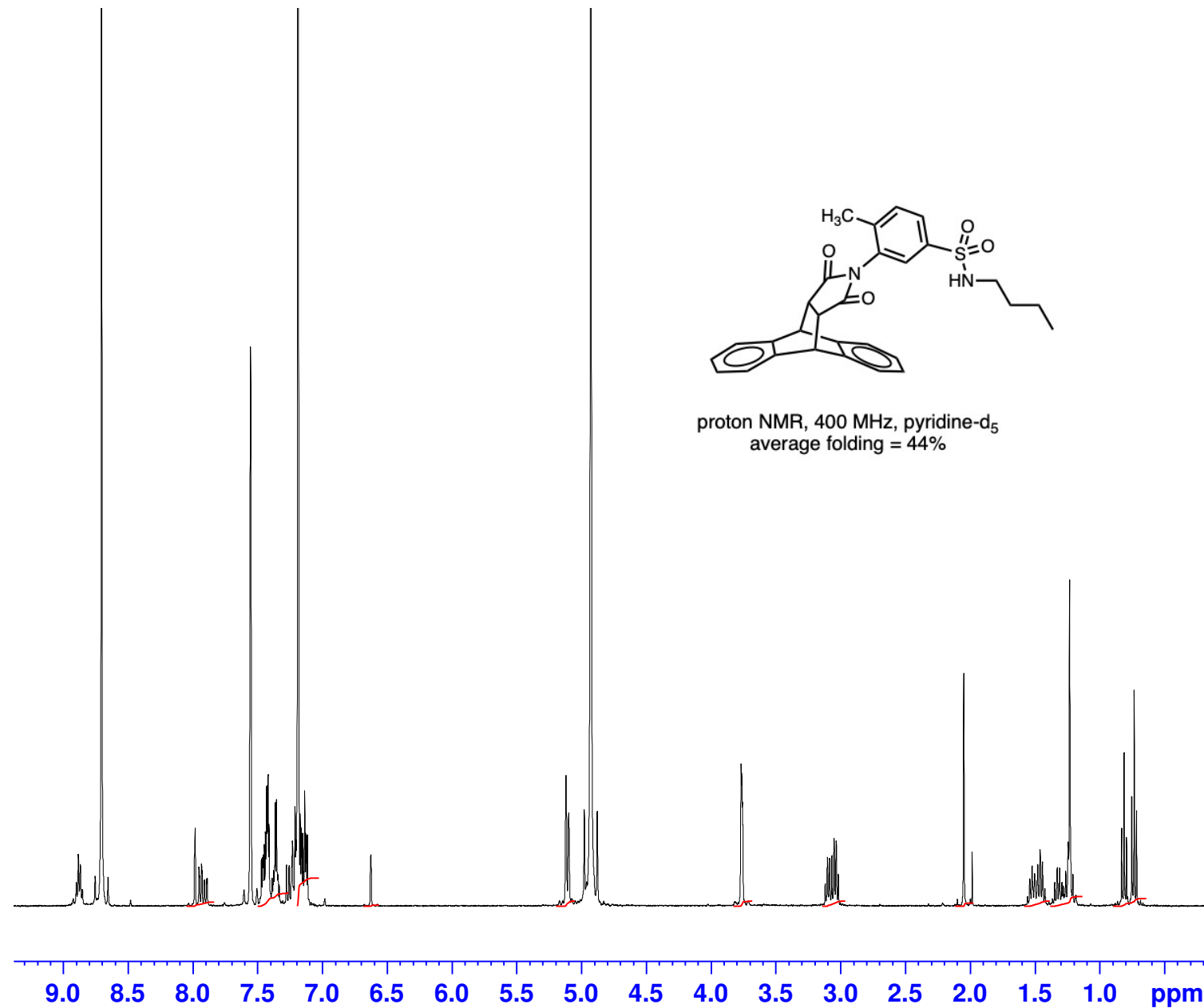
Integration values (in red): 10.35, 0.39, 0.53, 2.00, 1.98, 1.21

Current Data Parameters
NAME controlnovi9balancein-tertbutanol
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20221119
Time 18.15
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT EtOD
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 228
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====
SF01 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



proton NMR, 400 MHz, pyridine-d₅
average folding = 44%

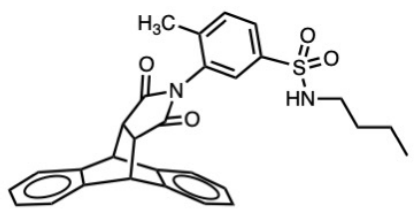
Current Data Parameters
 NAME controlnov19balanceinpyridine
 EXPNO 1
 PROCNO 1

F2 - Acquisition Parameters
 Date_ 20221119
 Time 15.40
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zg30
 TD 65536
 SOLVENT Pyr
 NS 16
 DS 2
 SWH 8012.820 Hz
 FIDRES 0.122266 Hz
 AQ 4.0894465 sec
 RG 287
 DW 62.400 usec
 DE 6.50 usec
 TE 298.0 K
 D1 1.0000000 sec
 TDO 1

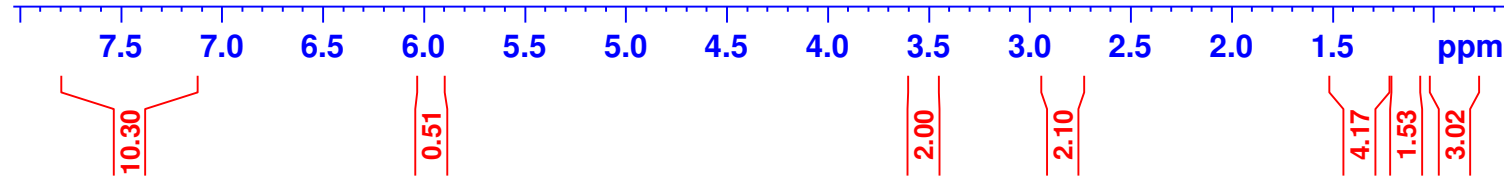
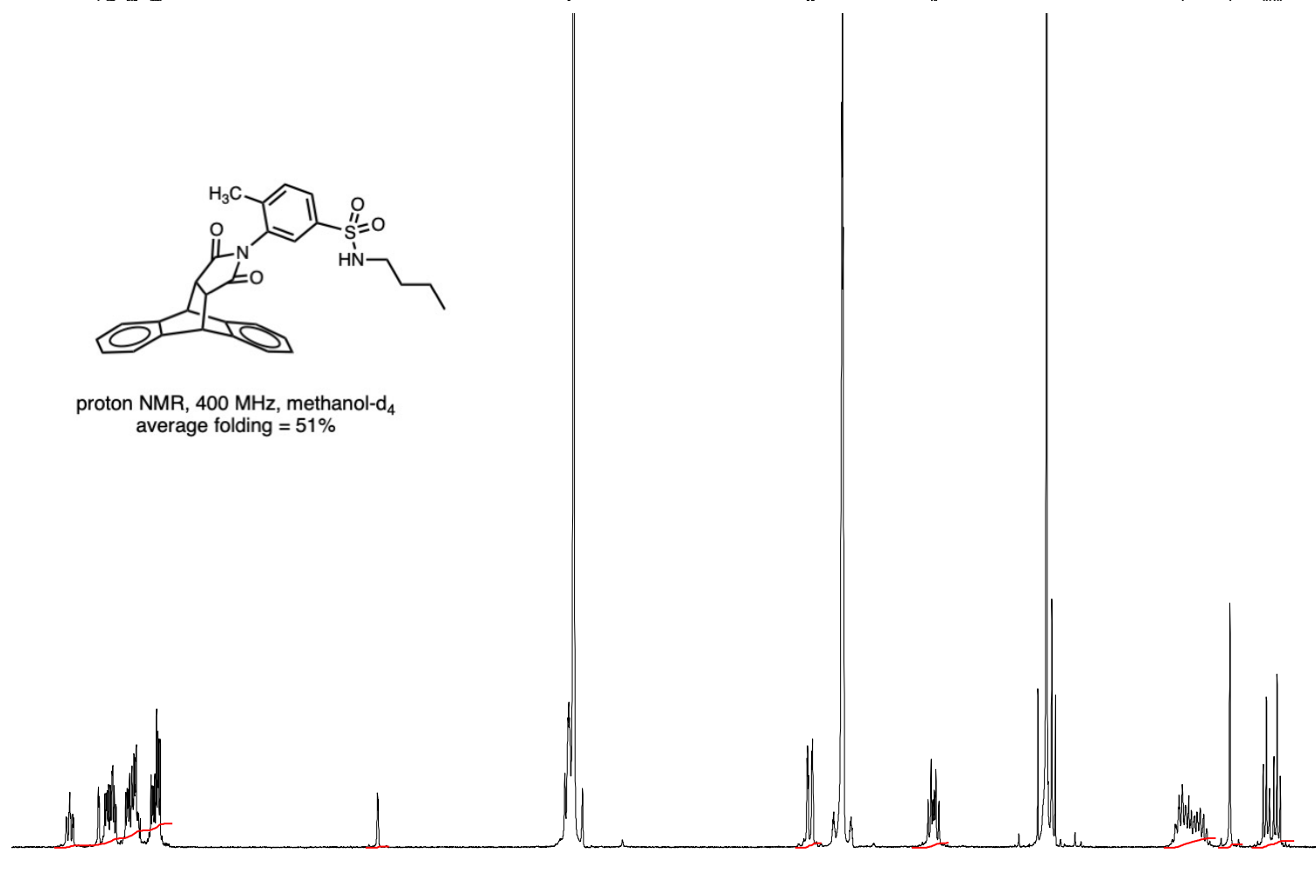
==== CHANNEL f1 =====
 SFO1 400.1324710 MHz
 NUC1 1H
 P1 9.30 usec
 PLW1 18.0000000 W

F2 - Processing parameters
 SI 65536
 SF 400.1300000 MHz
 WDW EM
 SSB 0
 LB 0.30 Hz
 GB 0
 PC 1.00

7.543
7.492
7.484
7.475
7.466
7.461
7.453
7.379
7.374
7.366
7.352
7.341
7.336
7.331
7.328
7.245
7.237
7.231
7.223
7.214
7.206
7.200
7.192
4.884
4.880
4.876
3.531
3.527
3.526
3.522
3.504
3.500
3.496
2.828
2.810
2.801
1.406
1.136
0.948
0.930
0.912
0.888
0.869
0.851



proton NMR, 400 MHz, methanol-d₄
average folding = 51%

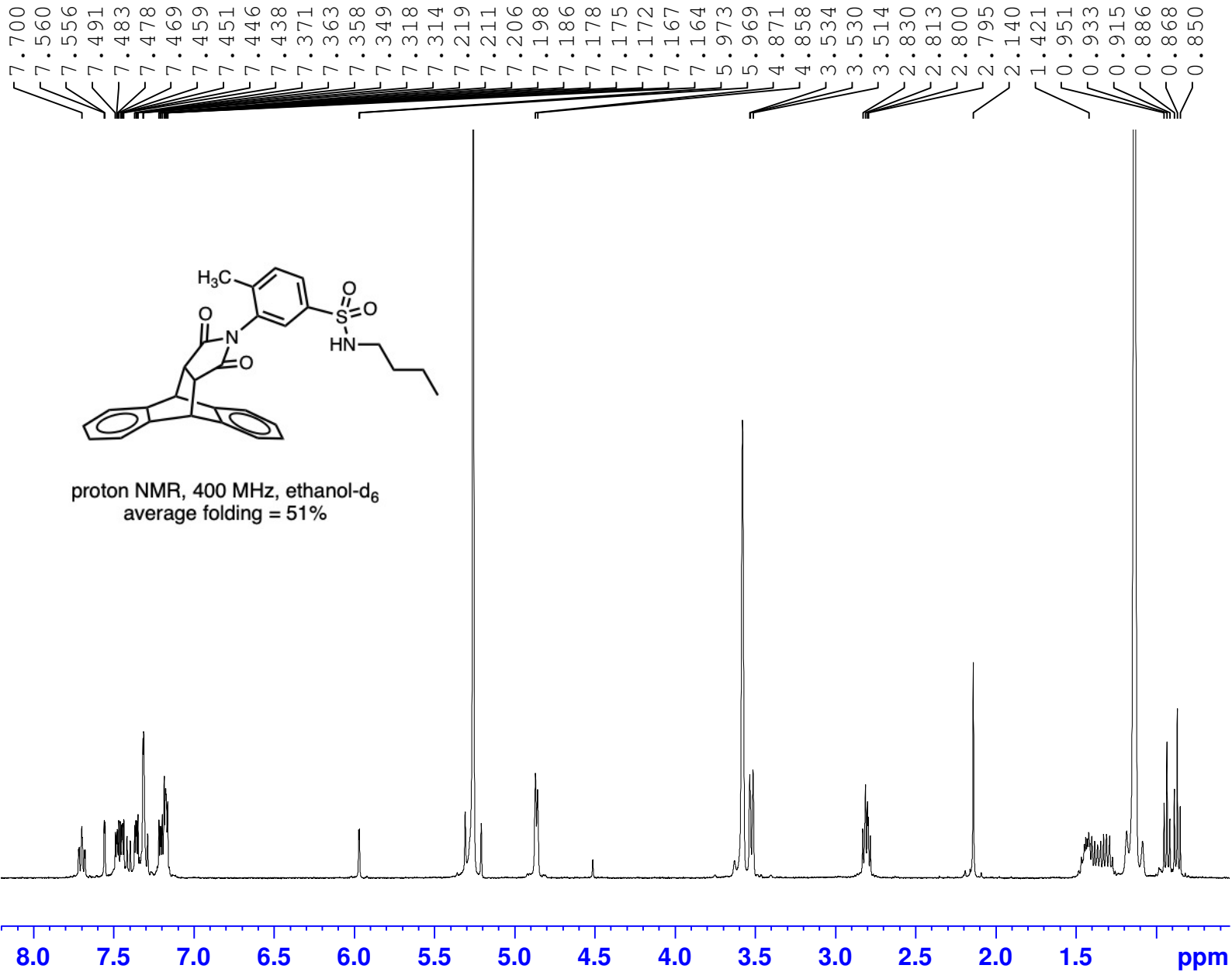


Current Data Parameters
NAME controlnov19balanceinmethanol
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20221119
Time 16.39
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT MeOD
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 322
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

===== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

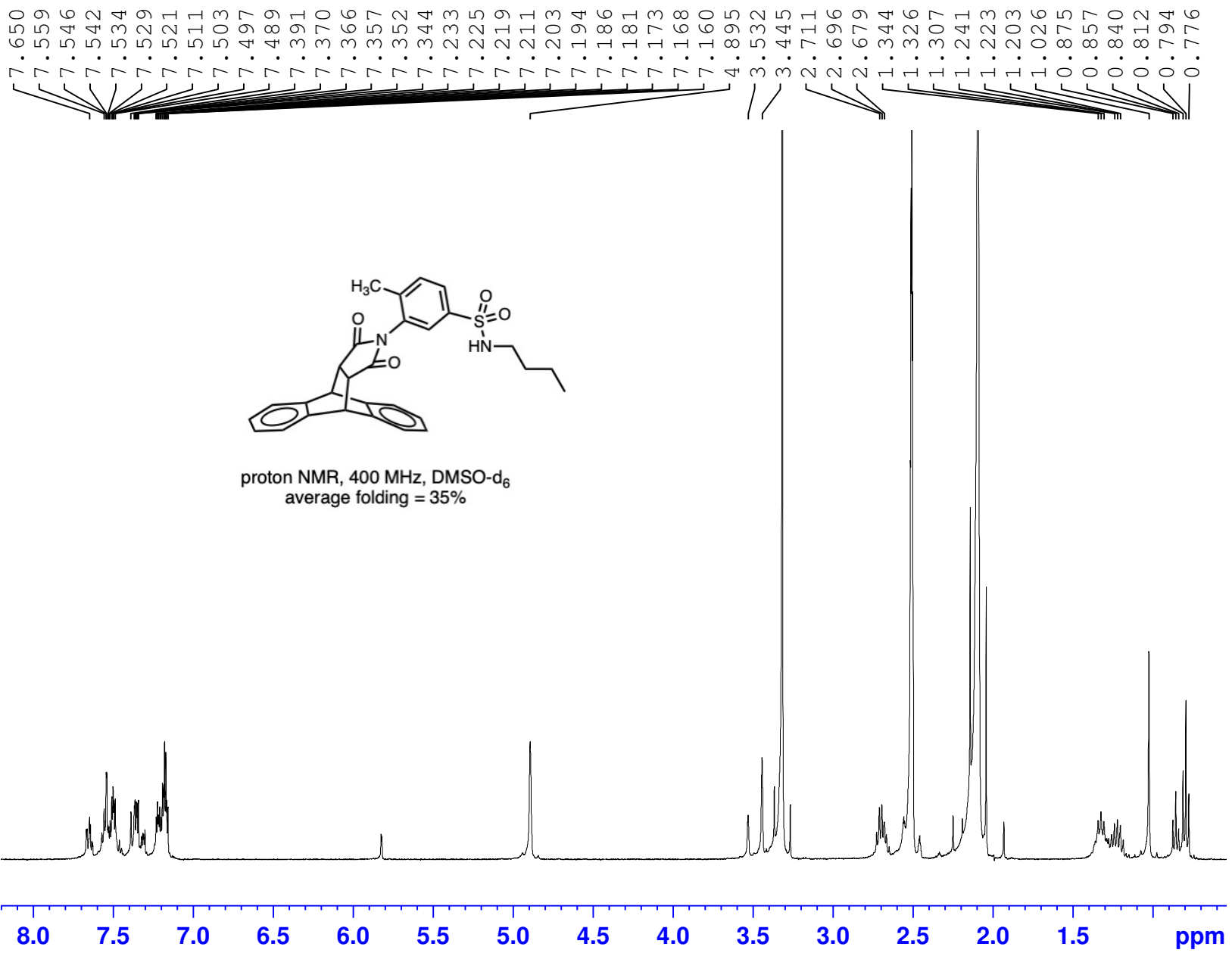


Current Data Parameters
 NAME controlnov19balancein-ethanol
 EXPNO 1
 PROCNO 1

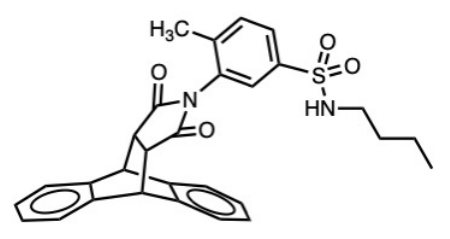
F2 - Acquisition Parameters
 Date_ 20221119
 Time 18.52
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zg30
 TD 65536
 SOLVENT EtOD
 NS 16
 DS 2
 SWH 8012.820 Hz
 FIDRES 0.122266 Hz
 AQ 4.0894465 sec
 RG 322
 DW 62.400 usec
 DE 6.50 usec
 TE 298.0 K
 D1 1.00000000 sec
 TD0 1

===== CHANNEL f1 =====
 SFO1 400.1324710 MHz
 NUC1 1H
 P1 9.30 usec
 PLW1 18.00000000 W

F2 - Processing parameters
 SI 65536
 SF 400.1300000 MHz
 WDW EM
 SSB 0
 LB 0.30 Hz
 GB 0
 PC 1.00



7.650
7.559
7.546
7.542
7.534
7.529
7.521
7.511
7.503
7.497
7.489
7.391
7.370
7.366
7.357
7.352
7.344
7.233
7.225
7.219
7.211
7.203
7.194
7.186
7.181
7.173
7.168
7.160
4.895
3.532
3.445
2.711
2.696
2.679
1.344
1.326
1.307
1.241
1.223
1.203
1.026
0.875
0.857
0.840
0.812
0.794
0.776



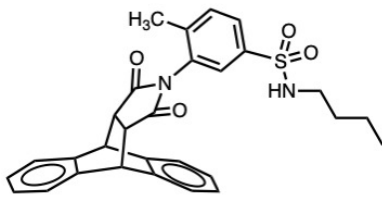
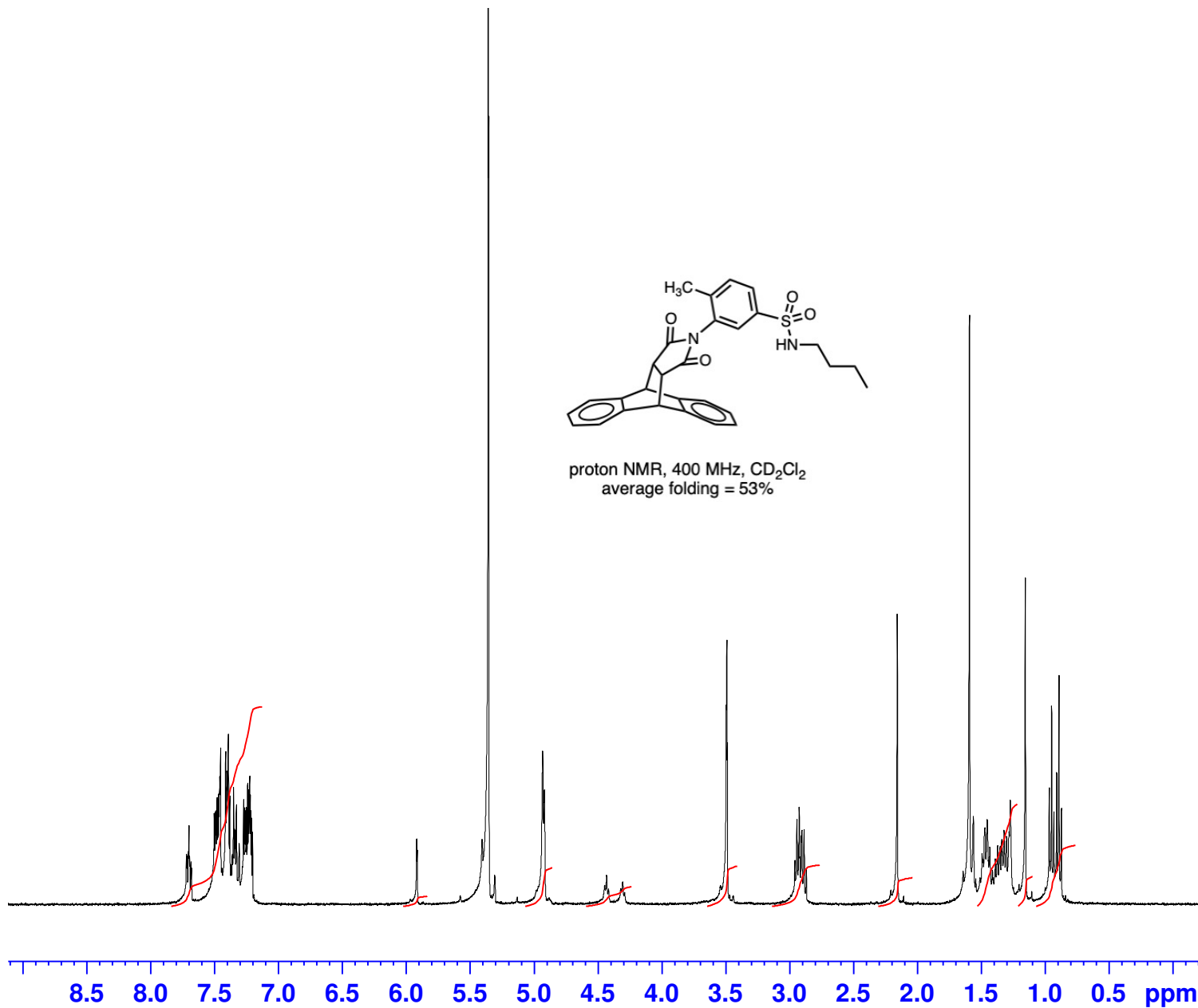
proton NMR, 400 MHz, DMSO-d₆
average folding = 35%

Current Data Parameters
NAME controlnov19balanceindms
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20221119
Time 14.33
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT DMSO
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 256
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



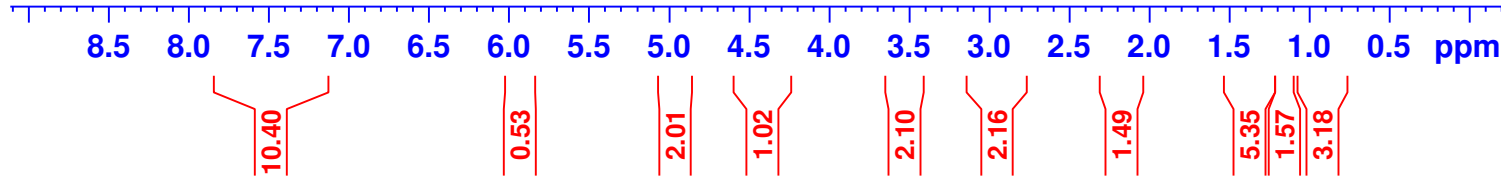
proton NMR, 400 MHz, CD₂Cl₂
average folding = 53%

Current Data Parameters
 NAME controlnov19balancein-methylenechloride
 EXPNO 1
 PROCNO 1

F2 - Acquisition Parameters
 Date_ 20221119
 Time 19.00
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zg30
 TD 65536
 SOLVENT CD2Cl2
 NS 16
 DS 2
 SWH 8012.820 Hz
 FIDRES 0.122266 Hz
 AQ 4.0894465 sec
 RG 575
 DW 62.400 usec
 DE 6.50 usec
 TE 298.0 K
 D1 1.00000000 sec
 TD0 1

===== CHANNEL f1 =====
 SFO1 400.1324710 MHz
 NUC1 1H
 P1 9.30 usec
 PLW1 18.00000000 W

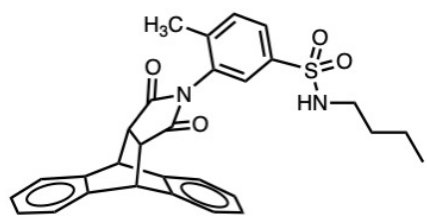
F2 - Processing parameters
 SI 65536
 SF 400.1300000 MHz
 WDW EM
 SSB 0
 LB 0.30 Hz
 GB 0
 PC 1.00



6.942
6.845
6.775
6.767
6.757
6.749
6.744
6.736
6.686
6.673
6.664
6.656
6.634
6.620
6.534
6.526
6.516
6.508
6.503
6.494
6.486
6.472
6.464
4.170
4.131
2.653
2.548
1.462
0.771
0.753
0.735
0.729
0.718
0.693
0.676
0.657
0.638
0.454
0.314
0.297
0.279
0.259
0.241
0.223

Current Data Parameters
NAME controlnov19balancein-CCl4
EXPNO 1
PROCNO 1

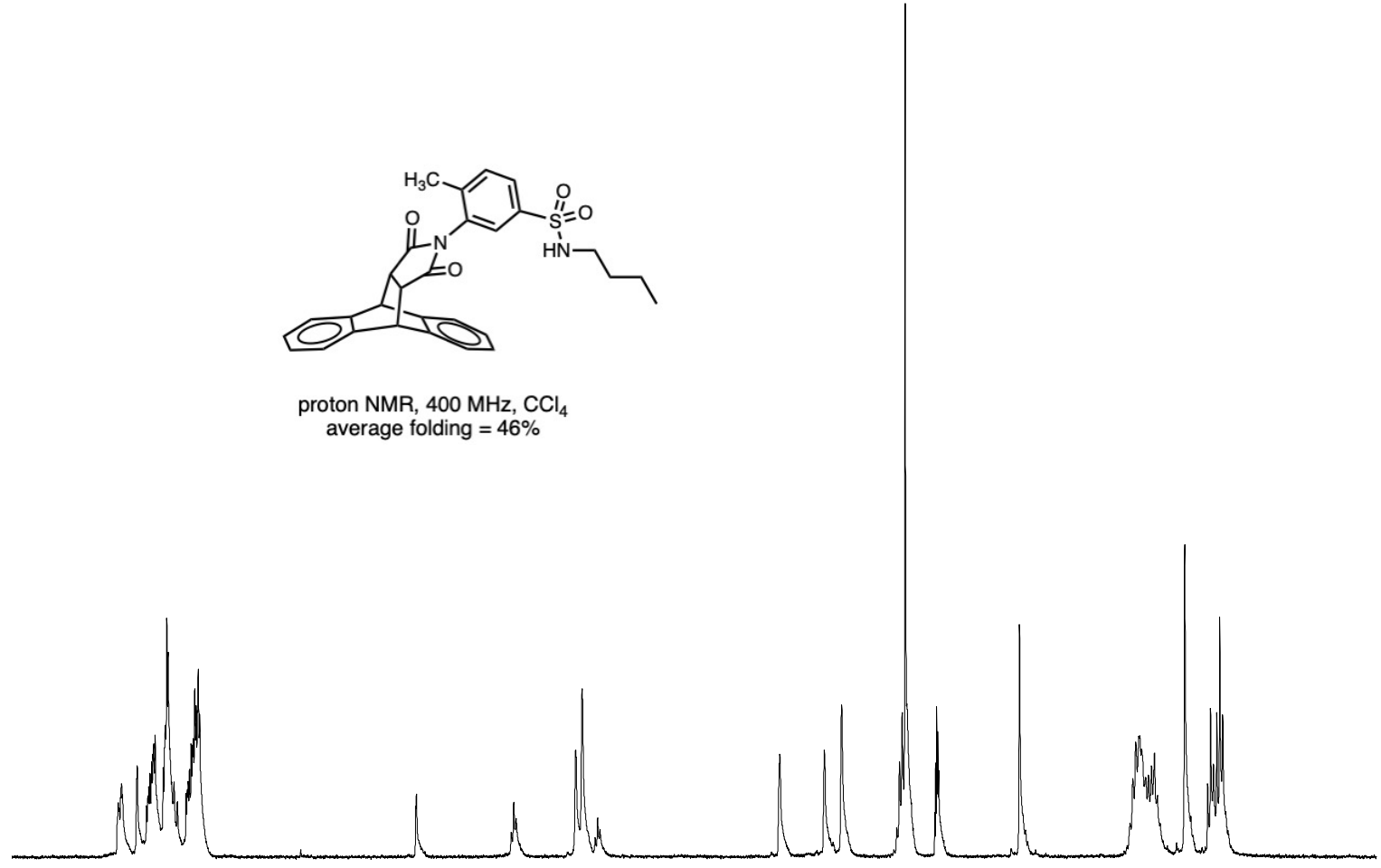
F2 - Acquisition Parameters
Date_ 20221119
Time 19.43
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT CD3CN
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 645
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1



proton NMR, 400 MHz, CCl₄
average folding = 46%

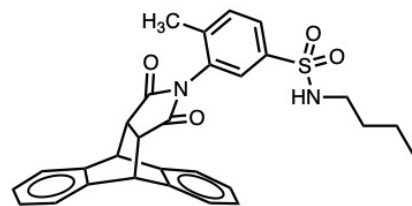
===== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

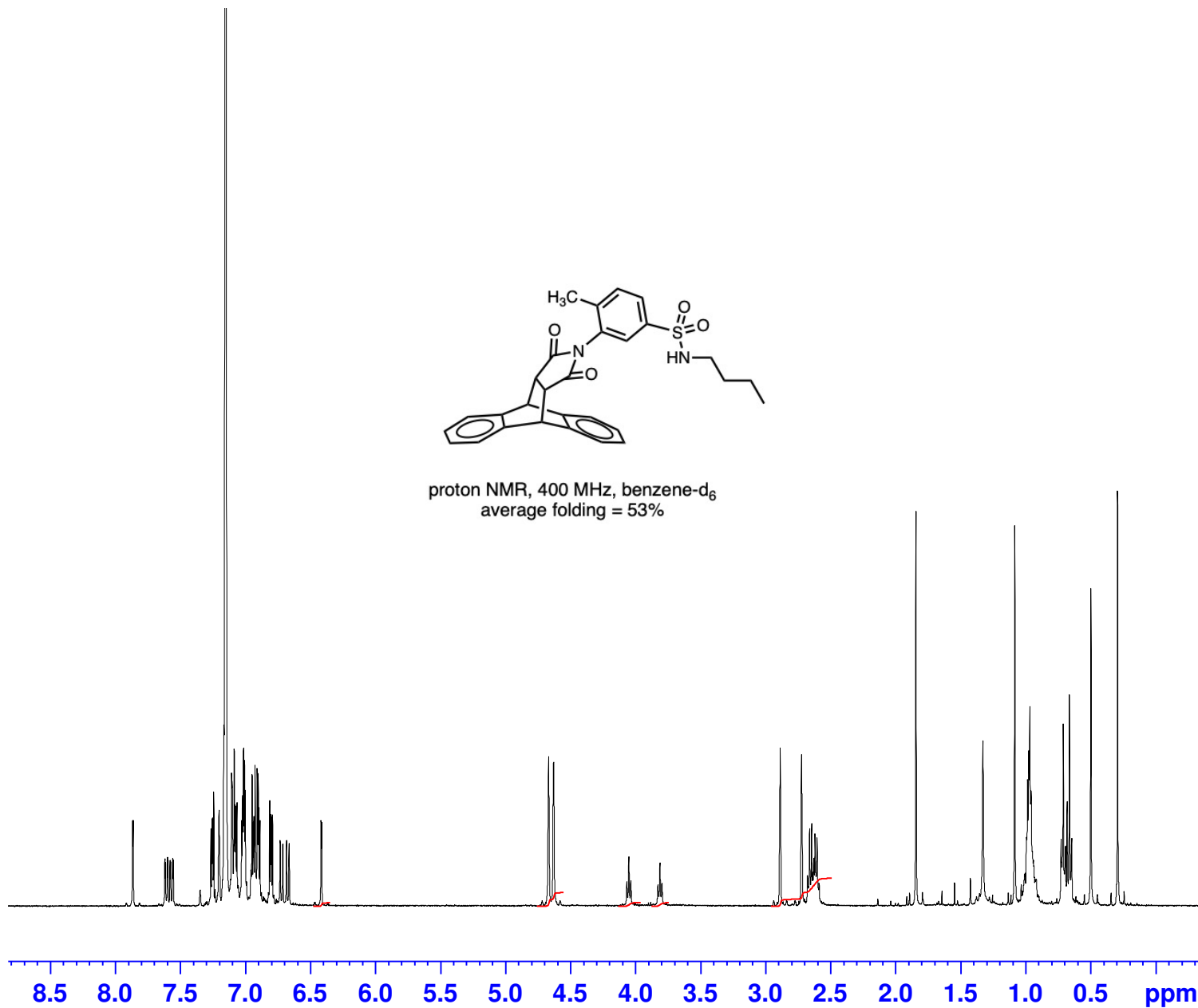


7 6 5 4 3 2 1 0 ppm

9.44 0.46 0.68 2.00 1.00 1.27 4.48 1.75 2.98



proton NMR, 400 MHz, benzene-d₆
average folding = 53%



0.53

2.00

0.55

0.53

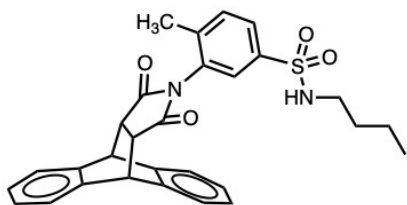
4.04

Current Data Parameters
NAME balance-sulfanamide-control-benzene
EXPNO 1
PROCNO 1

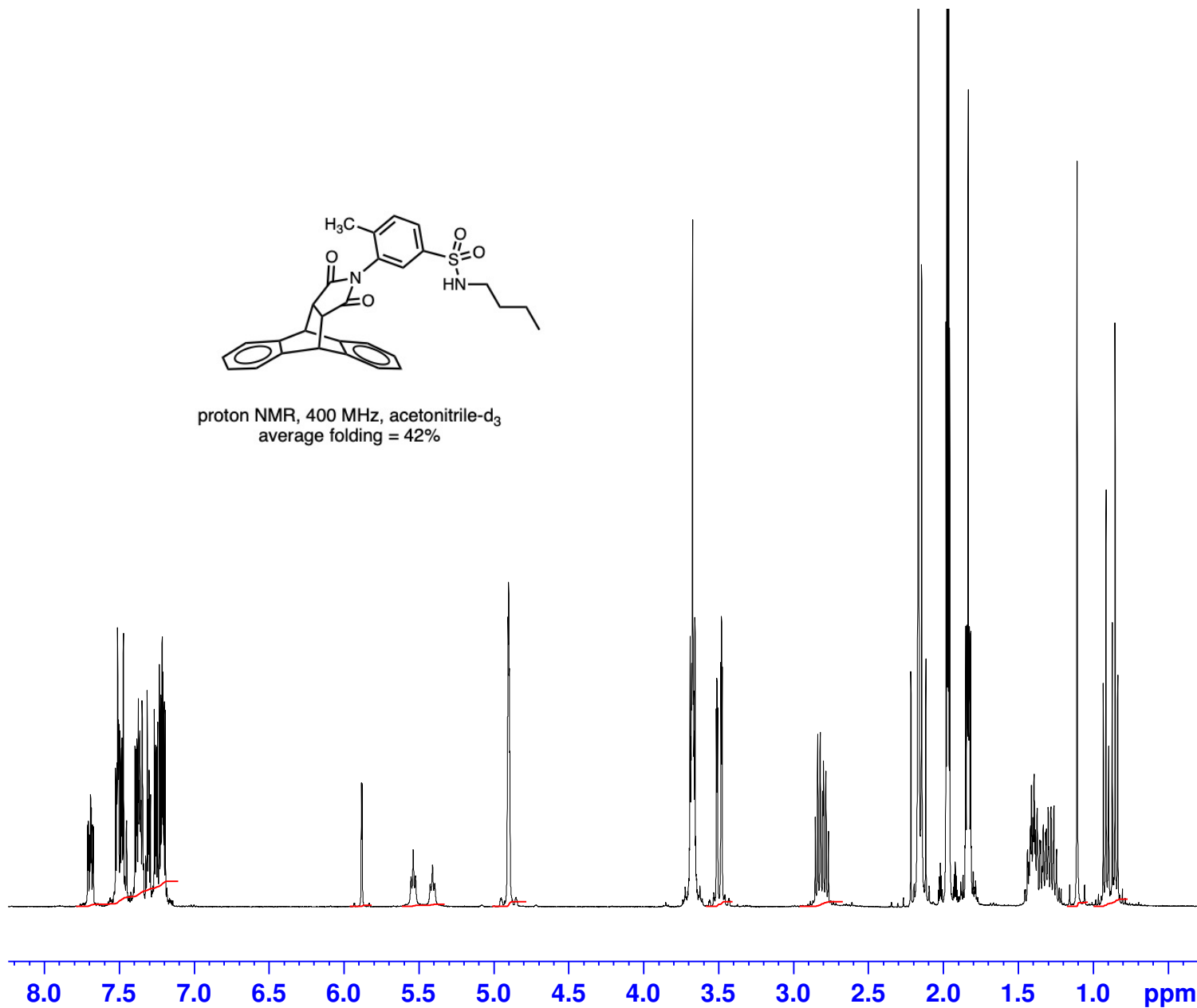
F2 - Acquisition Parameters
Date_ 20221224
Time 15.31
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT C6D6
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 256
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

==== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00



proton NMR, 400 MHz, acetonitrile-d₃
average folding = 42%



Current Data Parameters
NAME controlnov19balancein-
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20221122
Time 13.00
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT CD3CN
NS 160
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 512
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

===== CHANNEL f1 =====
SFO1 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00

8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 ppm

10.55

0.42

0.98

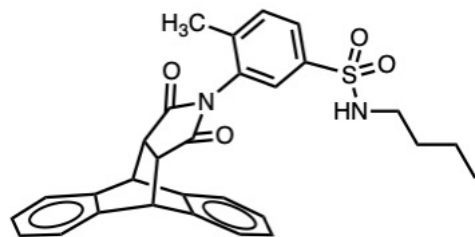
2.00

2.01

2.09

1.70

3.14



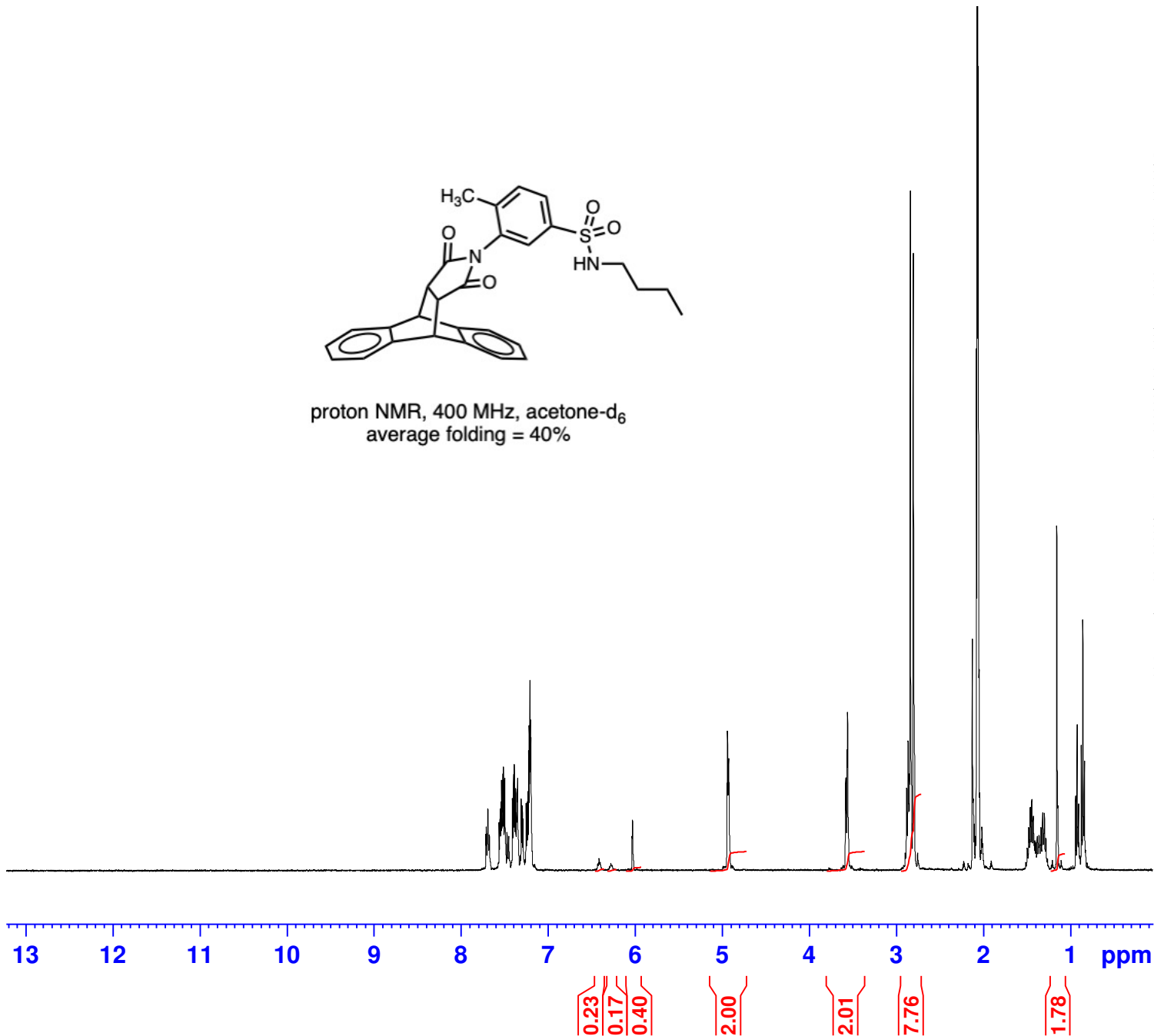
proton NMR, 400 MHz, acetone-d₆
average folding = 40%

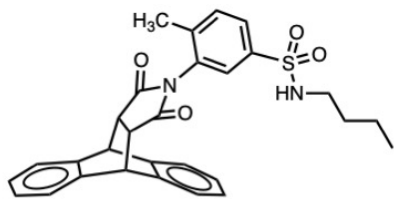
Current Data Parameters
NAME controlnov19balanceinacetone
EXPNO 1
PROCNO 1

F2 - Acquisition Parameters
Date_ 20221119
Time 15.27
INSTRUM spect
PROBHD 5 mm PABBO BB-
PULPROG zg30
TD 65536
SOLVENT Acetone
NS 16
DS 2
SWH 8012.820 Hz
FIDRES 0.122266 Hz
AQ 4.0894465 sec
RG 512
DW 62.400 usec
DE 6.50 usec
TE 298.0 K
D1 1.00000000 sec
TD0 1

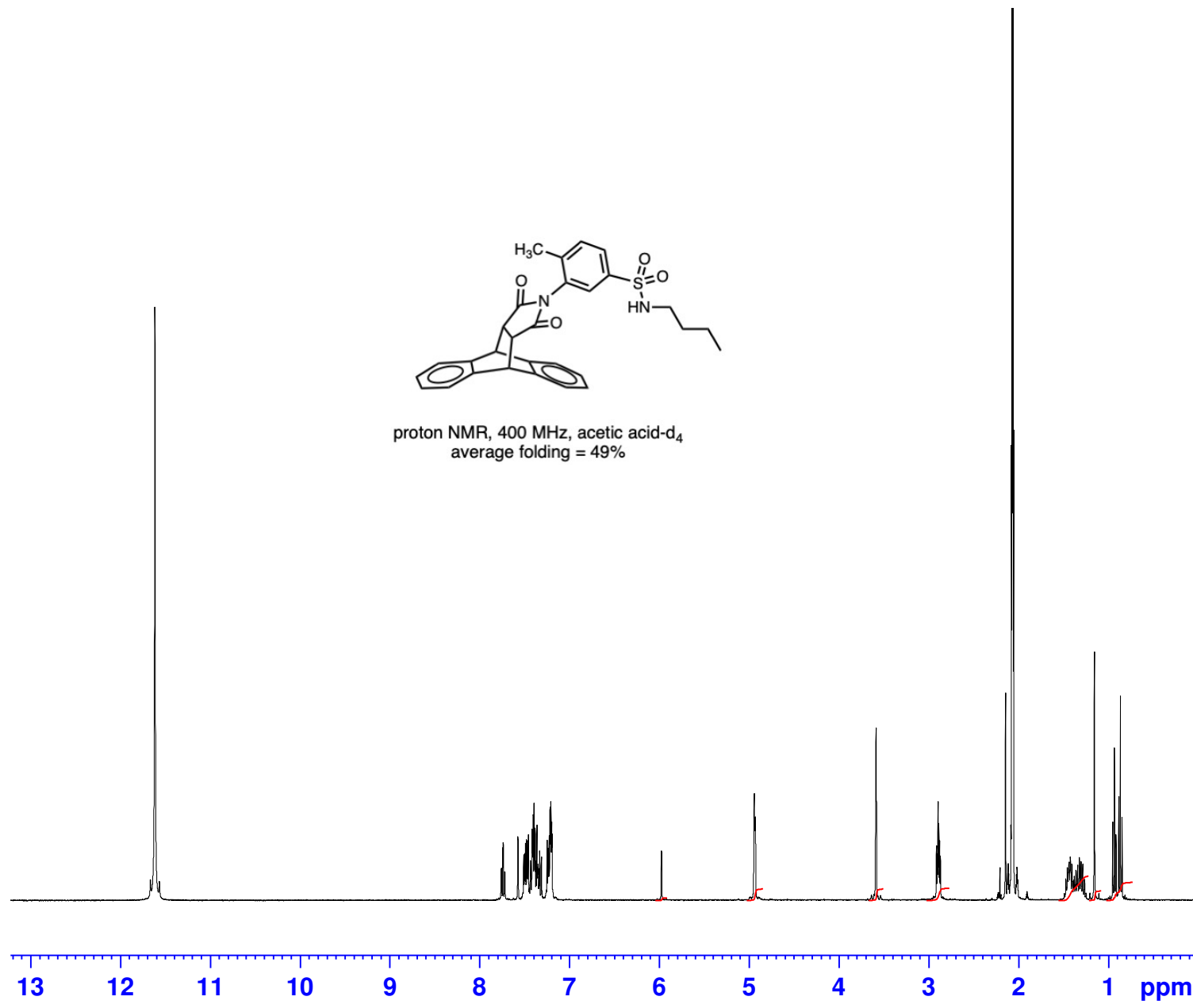
===== CHANNEL f1 =====
SF01 400.1324710 MHz
NUC1 1H
P1 9.30 usec
PLW1 18.00000000 W

F2 - Processing parameters
SI 65536
SF 400.1300000 MHz
WDW EM
SSB 0
LB 0.30 Hz
GB 0
PC 1.00





proton NMR, 400 MHz, acetic acid-d₄
average folding = 49%



Current Data Parameters
 NAME controlnov19balanceinaceticacid
 EXPNO 1
 PROCNO 1

F2 - Acquisition Parameters
 Date_ 20221119
 Time 16.32
 INSTRUM spect
 PROBHD 5 mm PABBO BB-
 PULPROG zg30
 TD 65536
 SOLVENT Acetic
 NS 16
 DS 2
 SWH 8012.820 Hz
 FIDRES 0.122266 Hz
 AQ 4.0894465 sec
 RG 322
 DW 62.400 usec
 DE 6.50 usec
 TE 298.0 K
 D1 1.00000000 sec
 TD0 1

===== CHANNEL f1 =====
 SF01 400.1324710 MHz
 NUC1 1H
 P1 9.30 usec
 PLW1 18.00000000 W

F2 - Processing parameters
 SI 65536
 SF 400.1300000 MHz
 WDW EM
 SSB 0
 LB 0.30 Hz
 GB 0
 PC 1.00