

Supporting Information

Highly Efficient Regio-Selective Ring-Opening Nucleophilic Fluorination of Aziridines and Azetidines: Access to β - or γ -Fluorinated Amino Acid Derivatives

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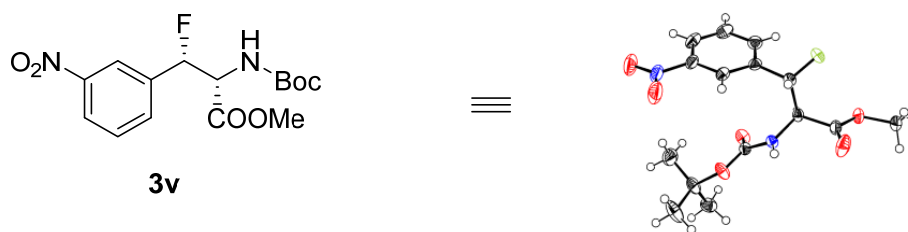
1. Conditions Optimization

Table S1. Conditions Optimization^a

1a		3a			
Entry	NuF Reagent	Solvent	Time (h)	Cover. (%)	Yield ^b (%)
1	CsF	CH ₂ Cl ₂	24	60	< 10
2	KHF ₂	CH ₂ Cl ₂	36	10	trace
3	TBAF	CH ₂ Cl ₂	24	0	-
4	Et ₃ N.HF	CH ₂ Cl ₂	24	50	N.D
5	Py.HF ^c	CH ₂ Cl ₂	0.5	100	94 (90 ^d)
6	Py.HF	Et ₂ O	5	100	90
7	Py.HF	Tol	0.5	100	78
8	Py.HF	THF	9	60	50
9	Py.HF	1,4-dioxane	9	65	56
10	Py.HF	acetone	9	40	35
11	Py.HF	CHCl ₃	0.5	100	92
12 ^e	Py.HF	CH ₂ Cl ₂	0.5	32	28
13 ^e	Py.HF	CH ₂ Cl ₂	9	85	82
14 ^f	Py.HF	CH ₂ Cl ₂	0.5	100	90
15 ^g	Py.HF	CH ₂ Cl ₂	0.5	100	94

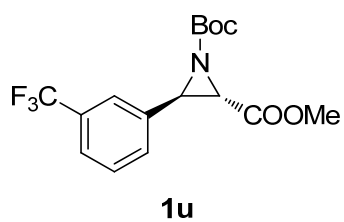
^aUnless noted, the reaction was conducted with **1a** (0.2 mmol), NuF reagent (3.0 equiv) and 2.0 mL solvent at room temperature for the specified period of time. ^bYields were determined by ¹⁹F NMR using benzotrifluoride as an internal standard. ^cPy.HF = hydrogen fluoride-pyridine (70% HF in pyridine). ^dIsolated yield. ^ePy.HF was added 1.5 equiv. ^fPy.HF was added 5.0 equiv. ^gThe reaction was carried out under nitrogen. TBAF = tetrabutylammonium fluoride, N.D. = not detected.

2. X-ray Crystallographic Data of 3v

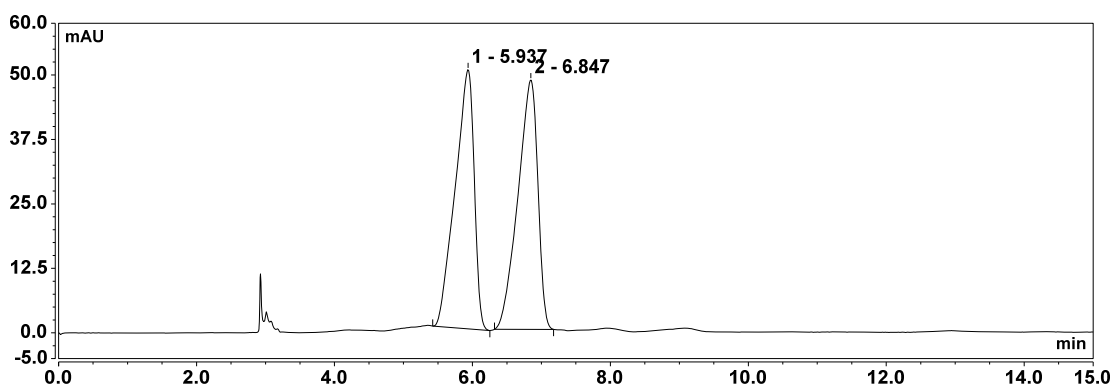


CCDC number	1887355
Empirical formula	C ₁₅ H ₁₉ FN ₂ O ₆
Formula weight	343.32
Temperature/K	150.00
Crystal system	orthorhombic
Space group	P 1 21 1
a/Å	5.1030(1)
b/Å	24.0246(3)
c/Å	13.7034(2)
α/°	90.0
β/°	98.635
γ/°	90.0
Volume/Å ³	1660.96(5)
Z	4
ρ _{calc} /g/cm ³	1.369
μ/mm ⁻¹	0.970
F(000)	720.0
Crystal size/mm ³	0.06 × 0.29 × 0.16
Radiation	MoKα(λ = 1.54184)
2θ range for data collection/°	3.68 to 73.32
Index ranges	-4 ≤ h ≤ 6, -29 ≤ k ≤ 29, -16 ≤ l ≤ 16
Reflections collected	12270
Independent reflections	6438 [R _{int} = 0.0331, R _{sigma} = 0.0395]
Data/restraints/parameters	6438/7/454
Goodness-of-fit on F ²	1.054
Final R indexes [I ≥ 2σ (I)]	R1 = 0.0360, wR2 = 0.0933
Final R indexes [all data]	R1 = 0.0375, wR2 = 0.0954
Largest diff. peak/hole / e Å ⁻³	0.152/-0.199

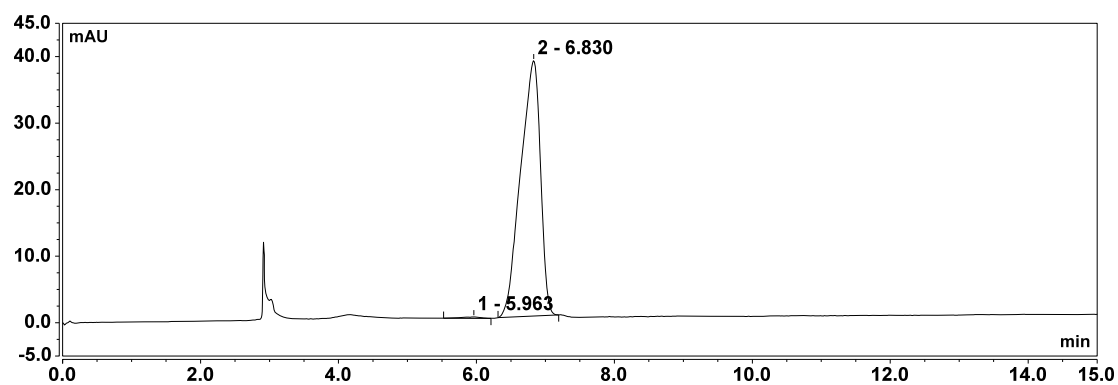
3. HPLC Results



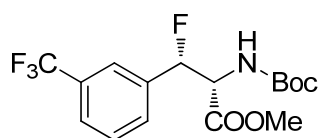
phenomenex Cellulose-1 5um 250x4.6mm (*n*-Hexane/*i*-PrOH = 97/3, flow rate = 1.0 mL/min, λ = 254 nm, 25 °C)



Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	5.937	16.0179	50.25	49.73
2	6.847	16.1914	48.39	50.27

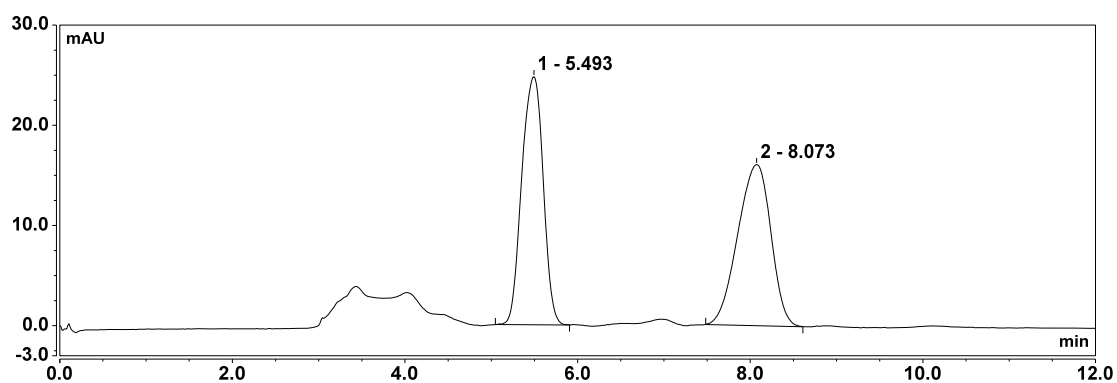


Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	5.963	0.0711	0.22	0.55
2	6.830	12.7965	38.38	99.45

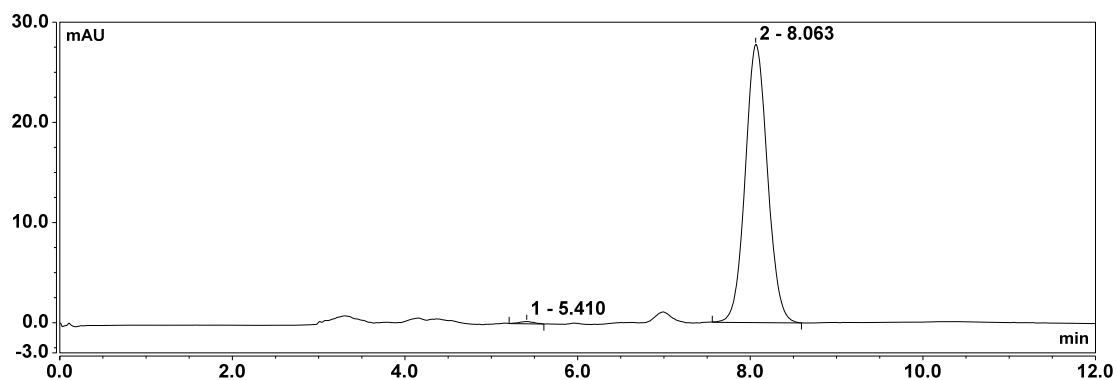


3u

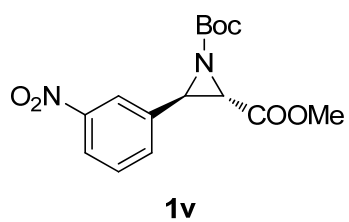
Chiral AD-H Column (*n*-Hexane/*i*-PrOH = 90/10, flow rate = 1.0 mL/min, λ = 254 nm, 25 °C)



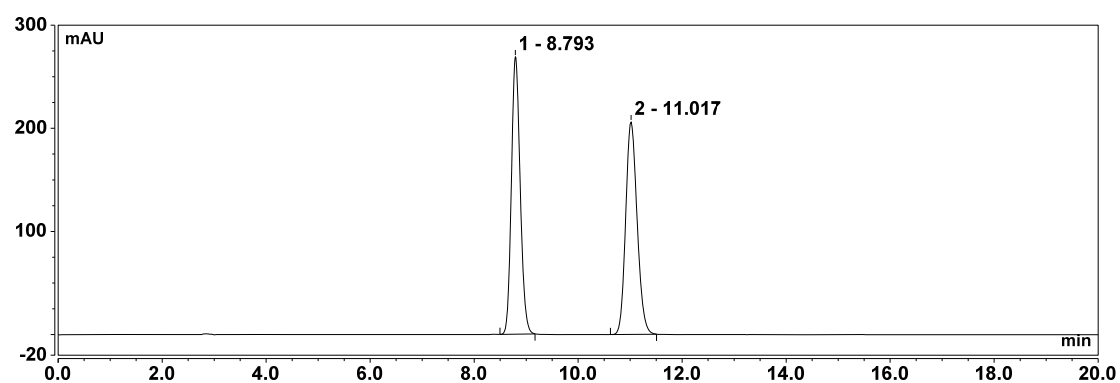
Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	5.493	7.1322	24.76	49.95
2	8.073	7.1463	16.08	50.05



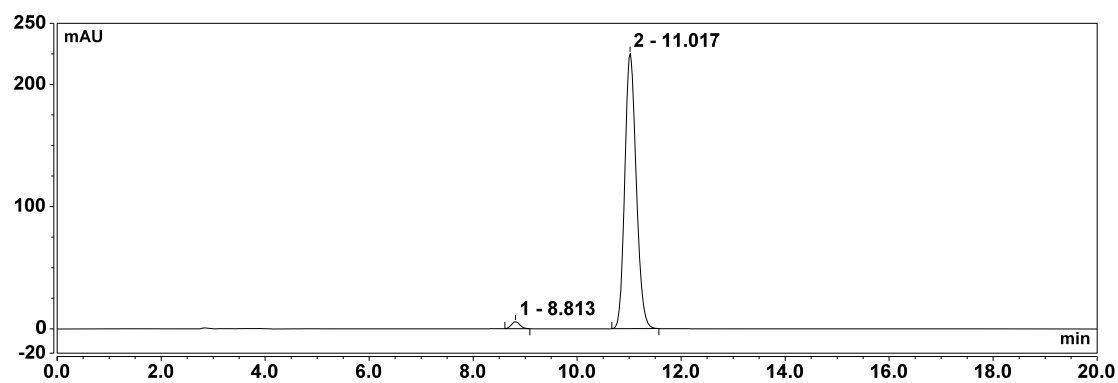
Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	5.410	0.0436	0.24	0.53
2	8.063	8.2684	27.78	99.47



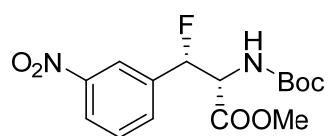
phenomenex Cellulose-1 5um 250x4.6mm (*n*-Hexane/*i*-PrOH = 90/10, flow rate = 1.0 mL/min, λ = 254 nm, 25 °C)



Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	8.793	52.6095	269.03	50.05
2	11.017	52.4941	206.30	49.95

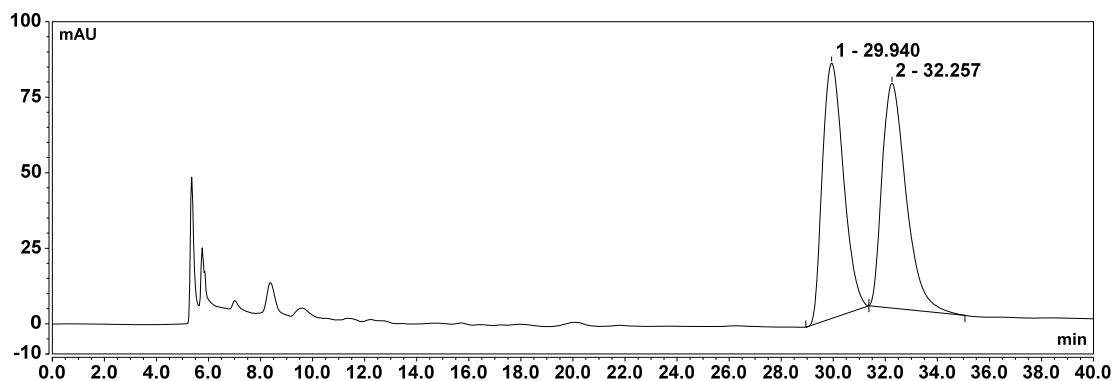


Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	8.813	1.0779	5.68	1.85
2	11.017	57.3313	225.21	98.15

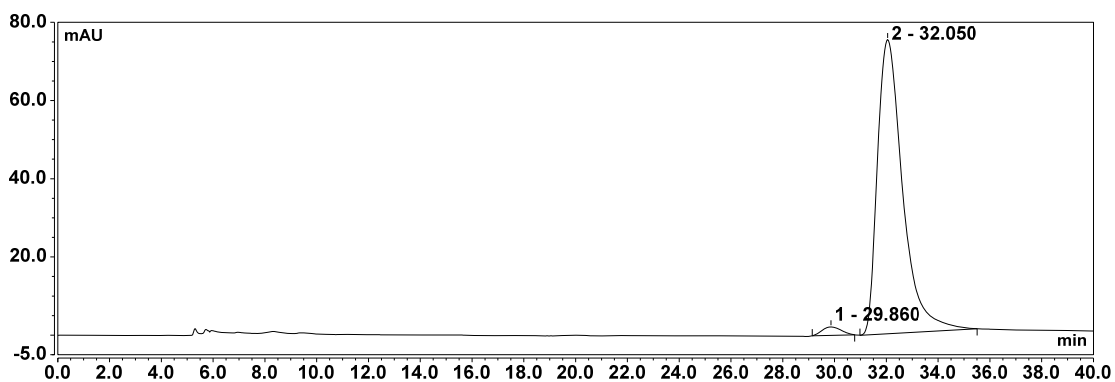


3v

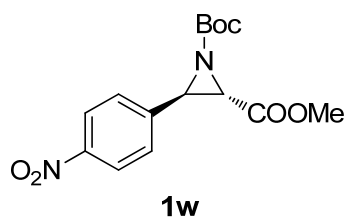
phenomenex Cellulose-1 5um 250x4.6mm (*n*-Hexane/*i*-PrOH = 95/5, flow rate = 0.5 mL/min, λ = 254 nm, 25 °C)



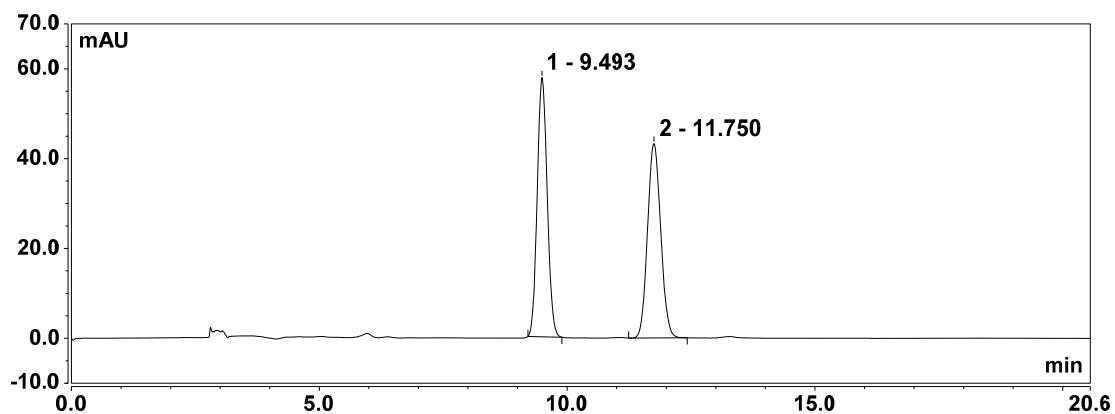
Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	29.940	79.1795	84.50	49.76
2	32.257	79.9410	74.46	50.24



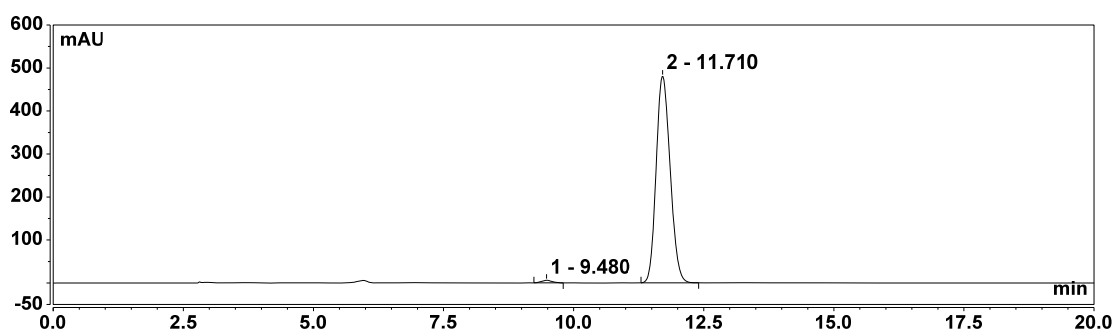
Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	29.860	1.8325	2.17	2.11
2	32.050	84.9753	75.22	97.89



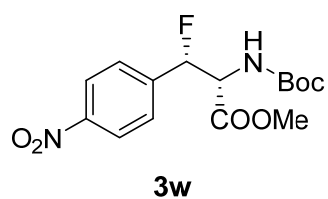
phenomenex Cellulose-1 5um 250x4.6mm (*n*-Hexane/*i*-PrOH = 90/10, flow rate = 1.0 mL/min, λ = 254 nm, 25 °C)



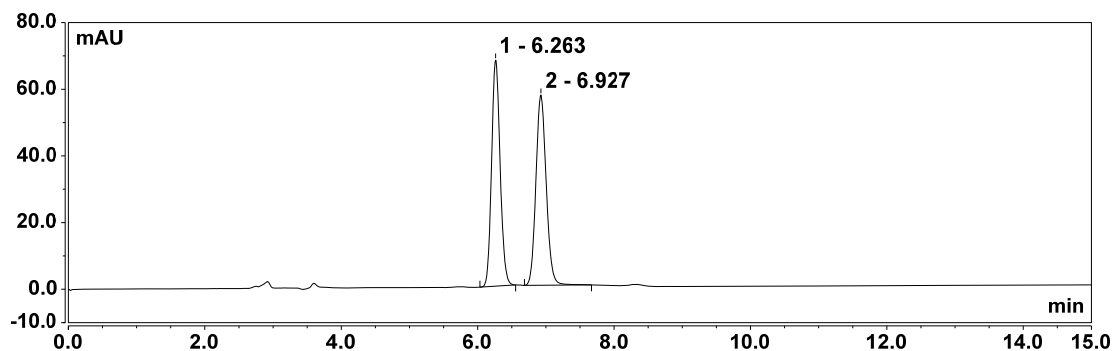
Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	9.493	13.4856	57.75	50.26
2	11.750	13.3471	43.35	49.74



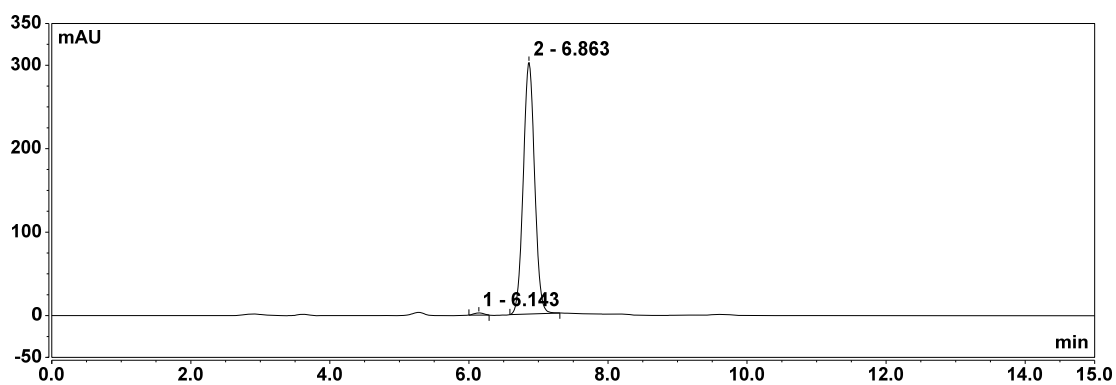
Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	9.480	1.3857	6.12	0.92
2	11.710	149.6401	480.92	99.08



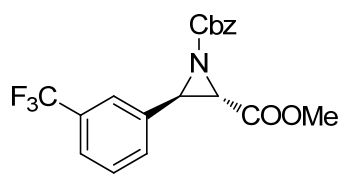
phenomenex Cellulose-1 5um 250x4.6mm (*n*-Hexane/*i*-PrOH = 80/20, flow rate = 1.0 mL/min, λ = 254 nm, 25 °C)



Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	6.263	9.8607	67.91	49.70
2	6.927	9.9809	57.07	50.30

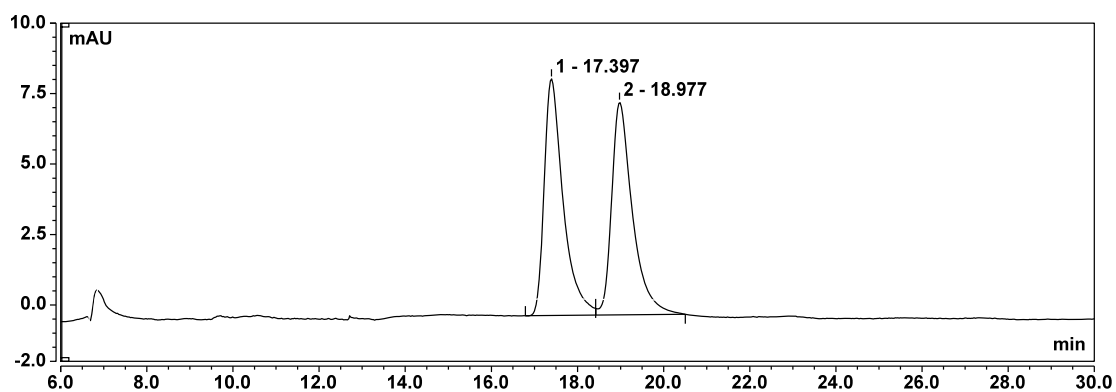


Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	6.143	0.4127	2.71	0.72
2	6.863	56.9436	301.61	99.28

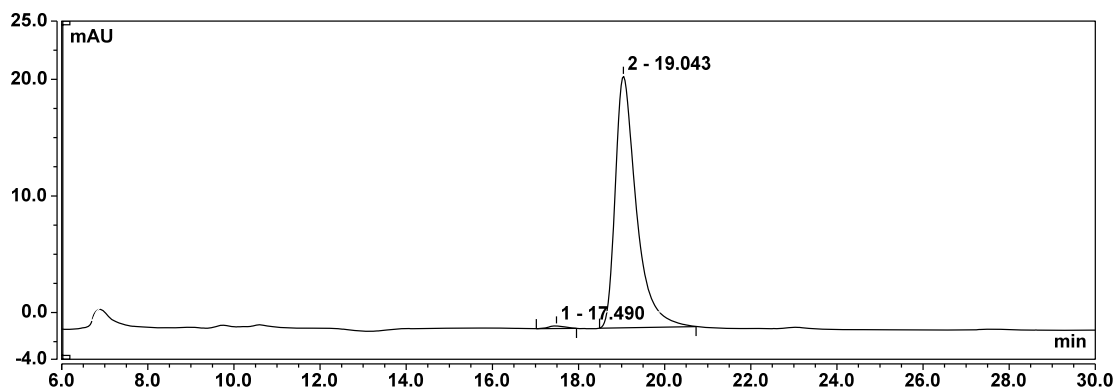


1x

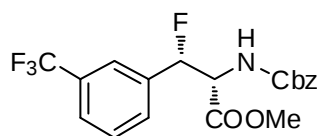
phenomenex Cellulose-1 5um 250x4.6mm (*n*-Hexane/*i*-PrOH = 95/5, flow rate = 0.5 mL/min, λ = 254 nm, 25 °C)



Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	17.397	4.2819	8.40	50.37
2	18.977	4.2191	7.55	49.63

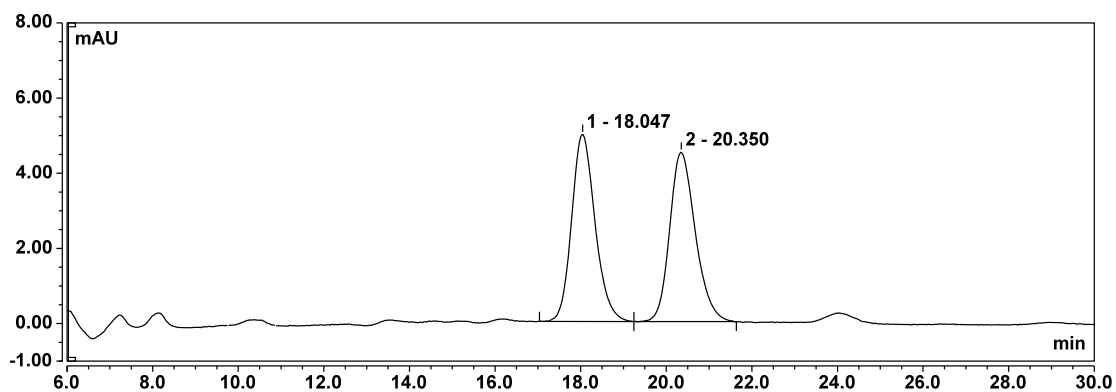


Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	17.490	0.0966	0.23	0.78
2	19.043	12.2387	21.57	99.22

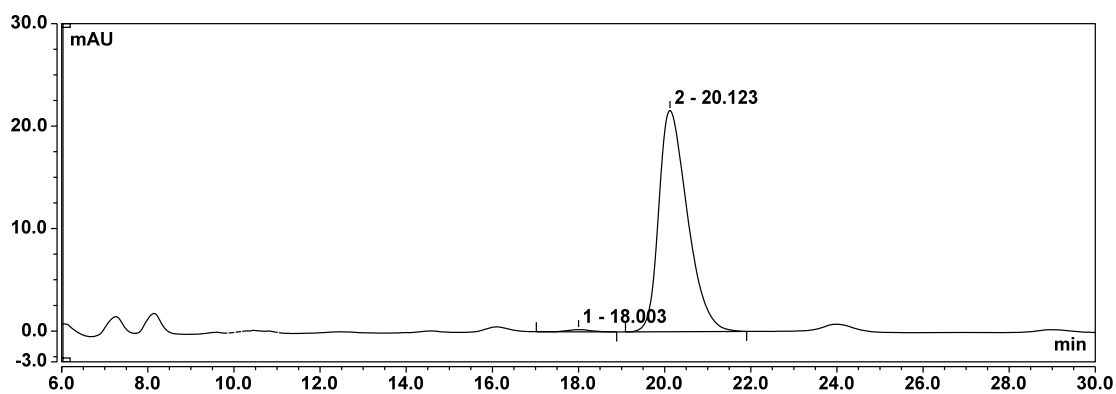


3x

Chiral AD-H Column (*n*-Hexane/*i*-PrOH = 90/10, flow rate = 0.8 mL/min, λ = 254 nm, 25 °C)



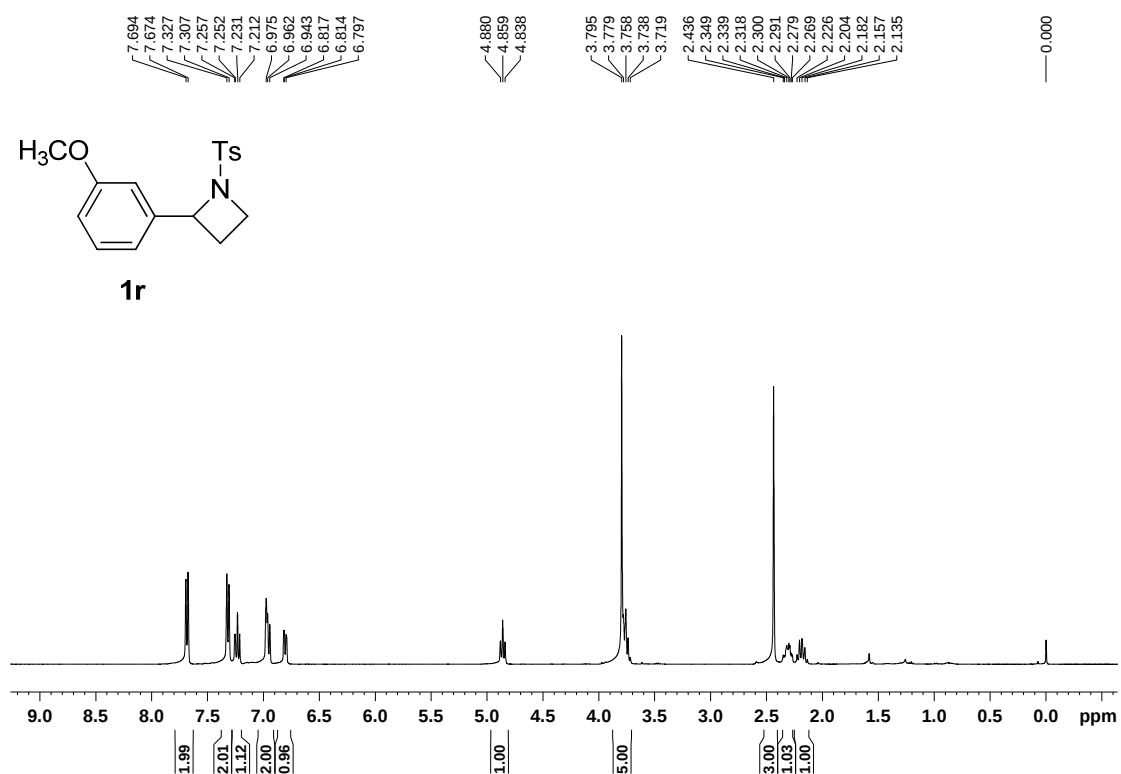
Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	18.047	3.1581	4.98	50.18
2	20.350	3.1348	4.51	49.82



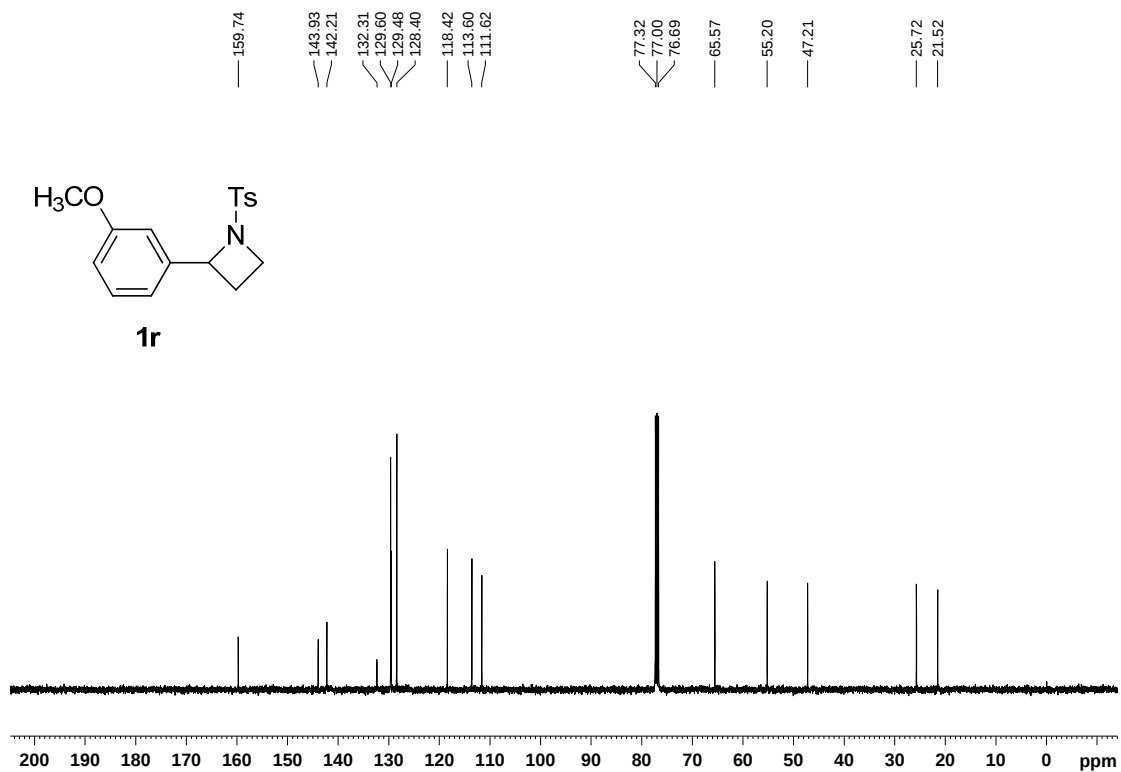
Peak #	RetTime [min]	Area mAU * min	Height [AU]	Area %
1	18.003	0.1340	0.22	0.80
2	20.123	16.6378	21.60	99.20

4. NMR Spectra of Compounds

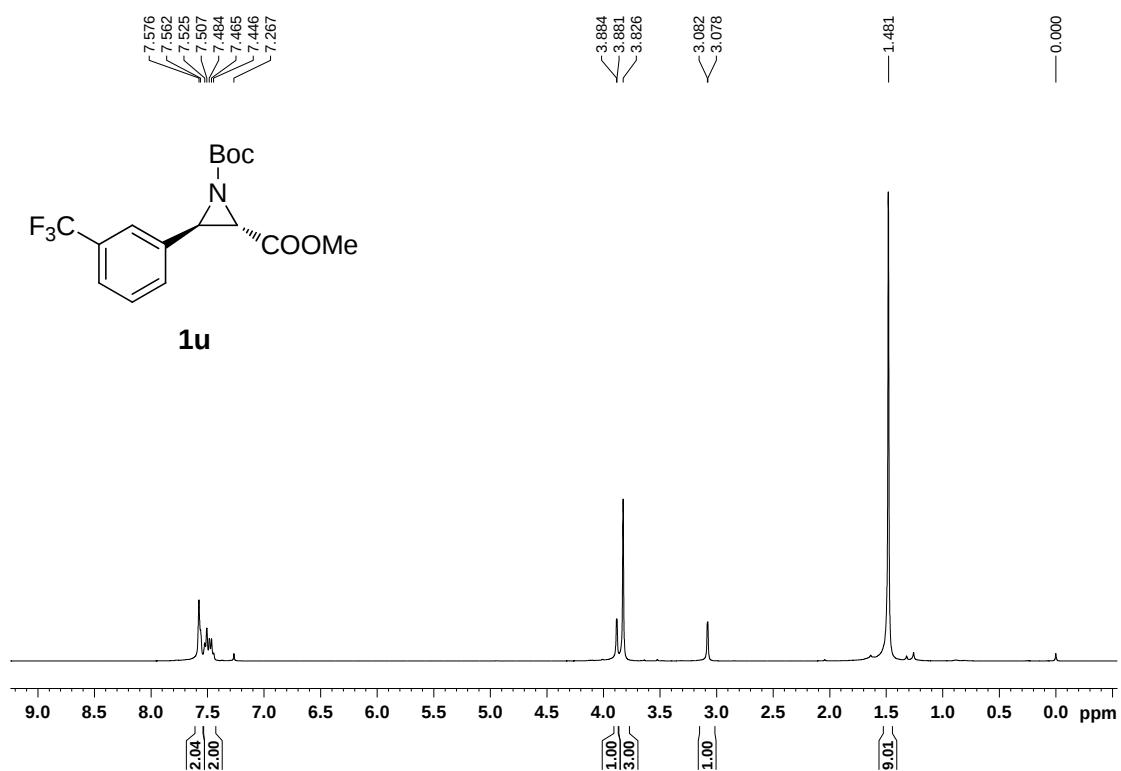
^1H NMR spectrum of **1r**



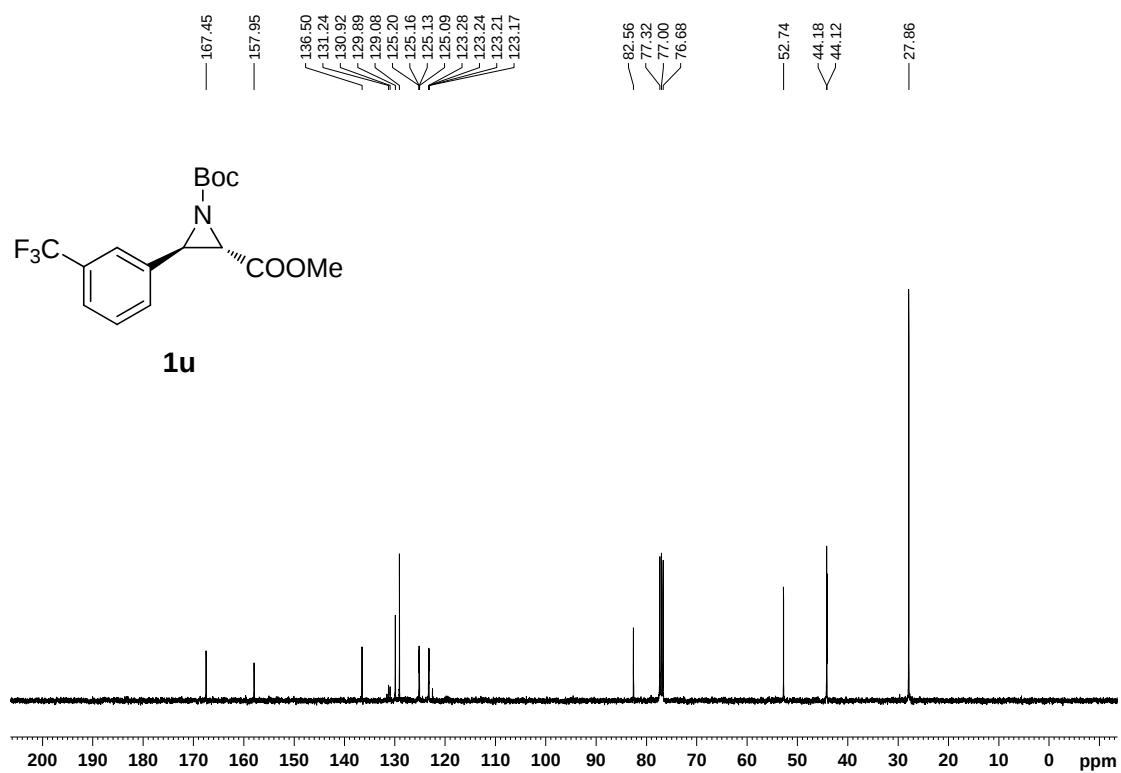
^{13}C NMR spectrum of **1r**



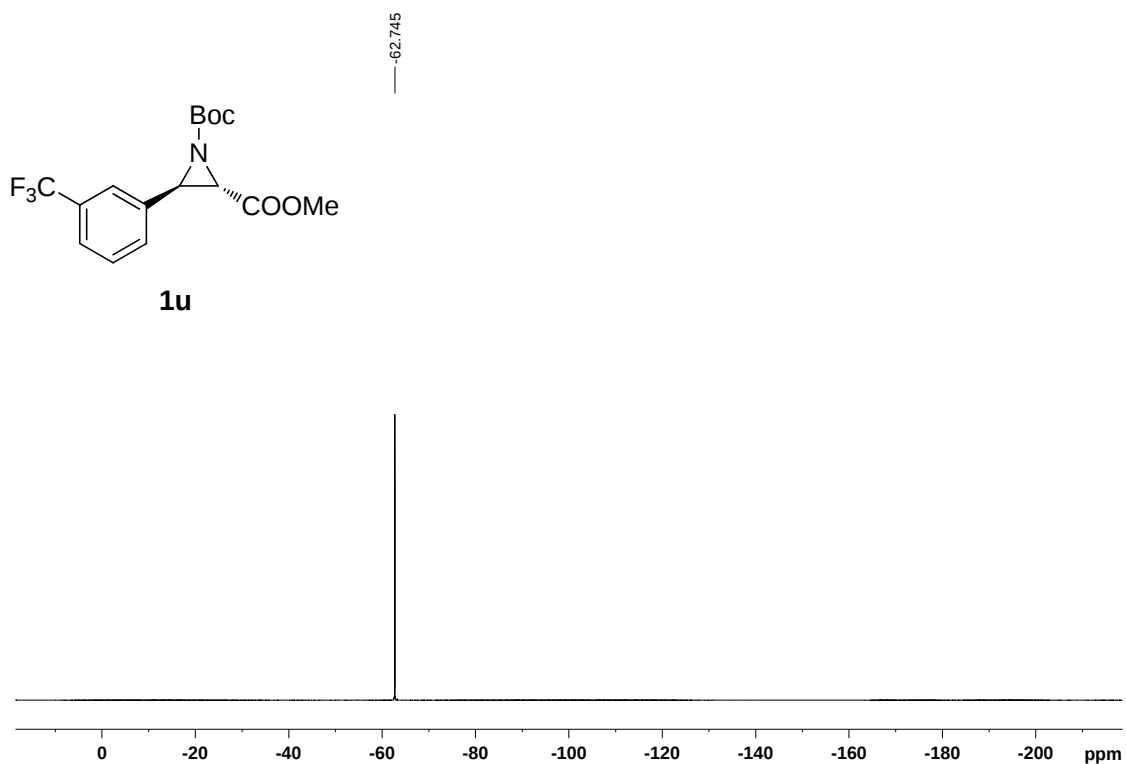
¹H NMR spectrum of **1u**



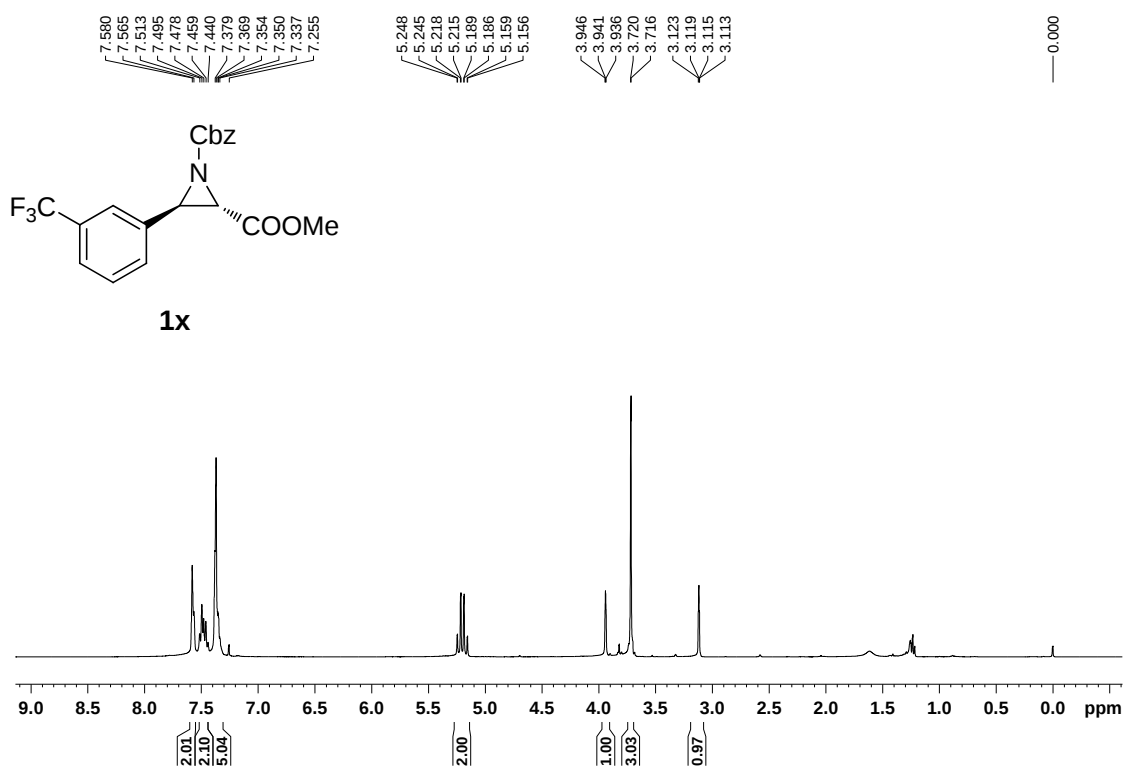
¹³C NMR spectrum of **1u**



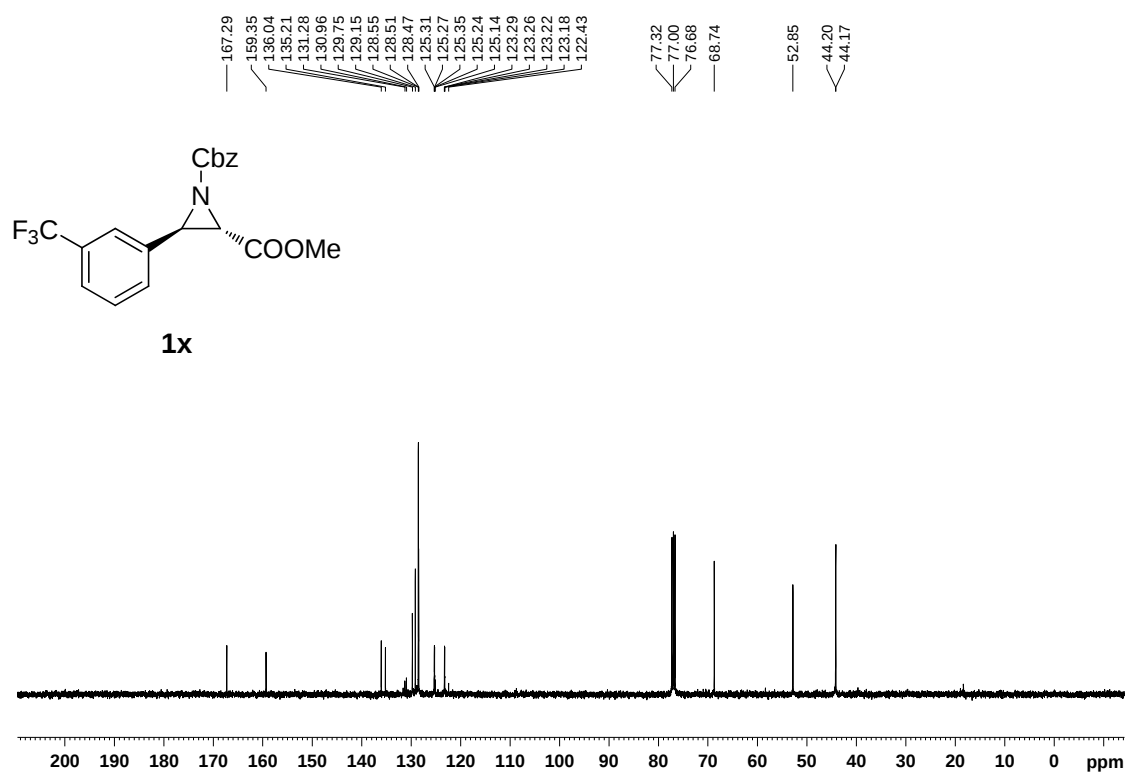
^1F NMR spectrum of **1u**



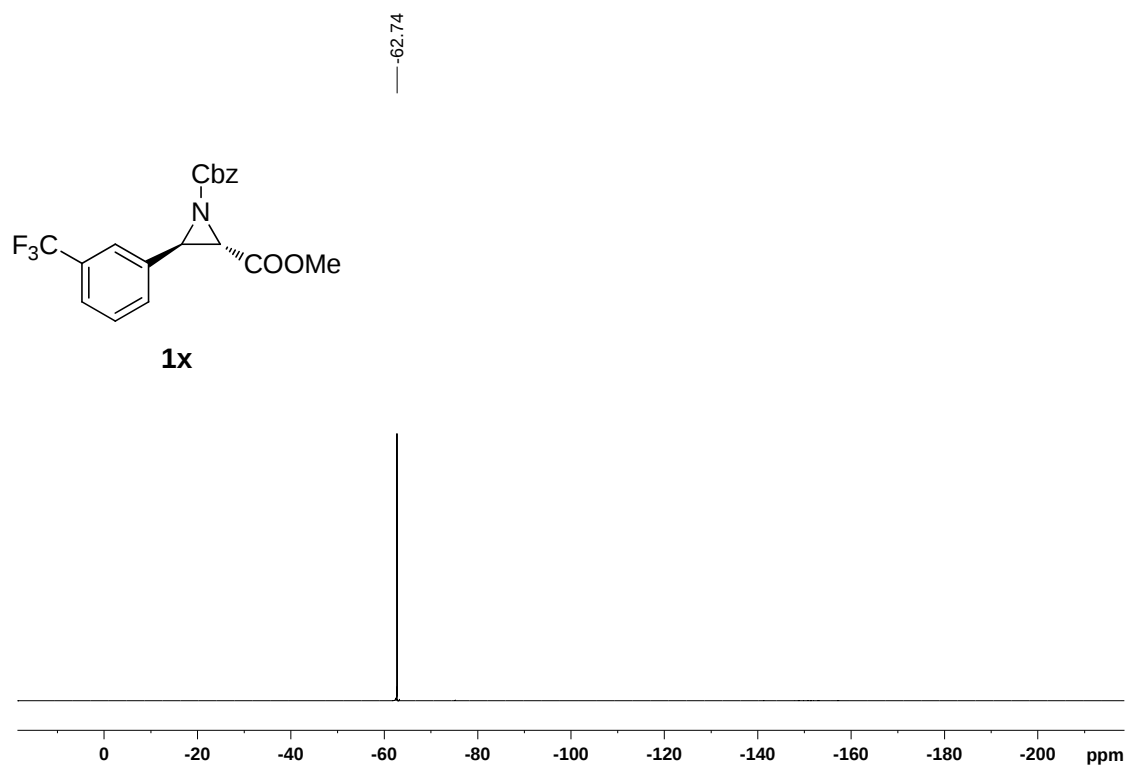
^1H NMR spectrum of **1x**



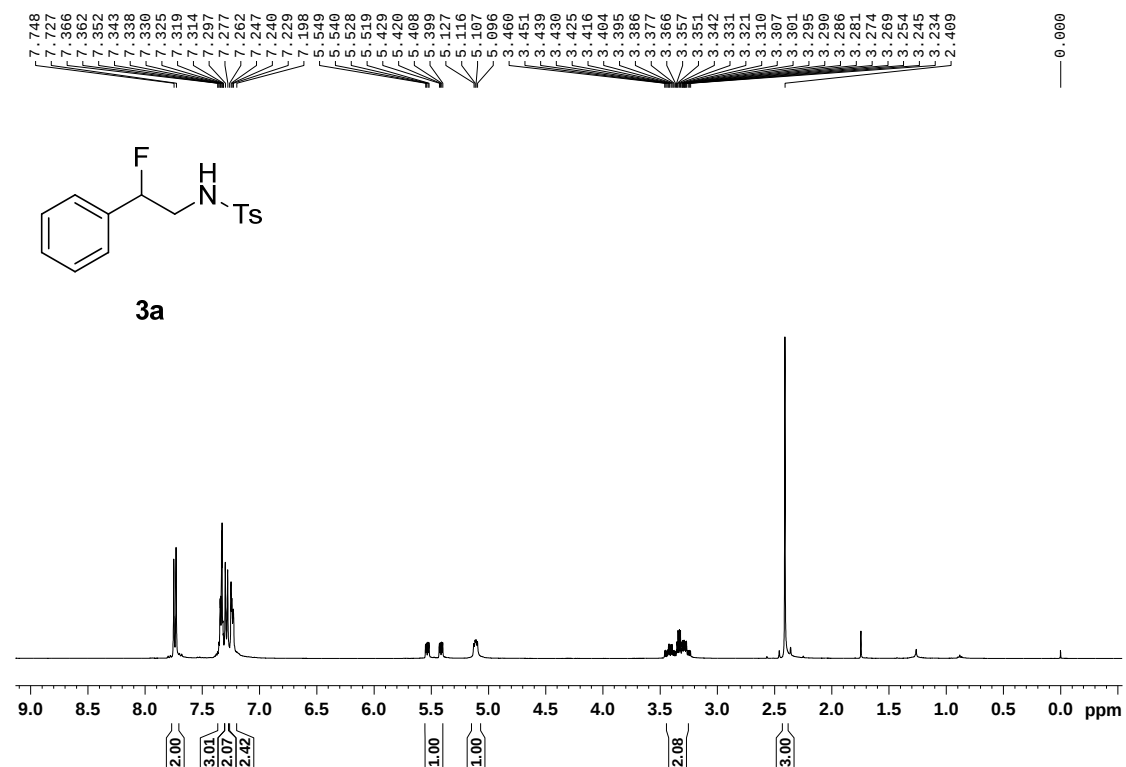
^{13}C NMR spectrum of **1x**



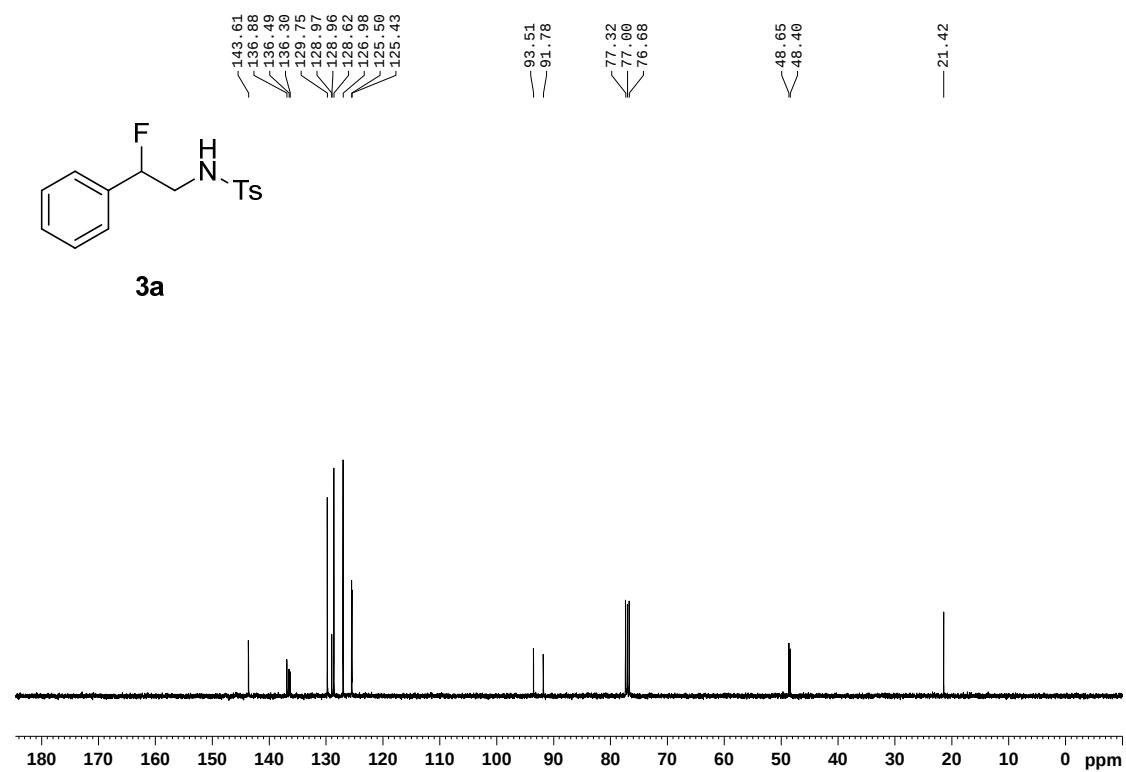
^1F NMR spectrum of **1x**



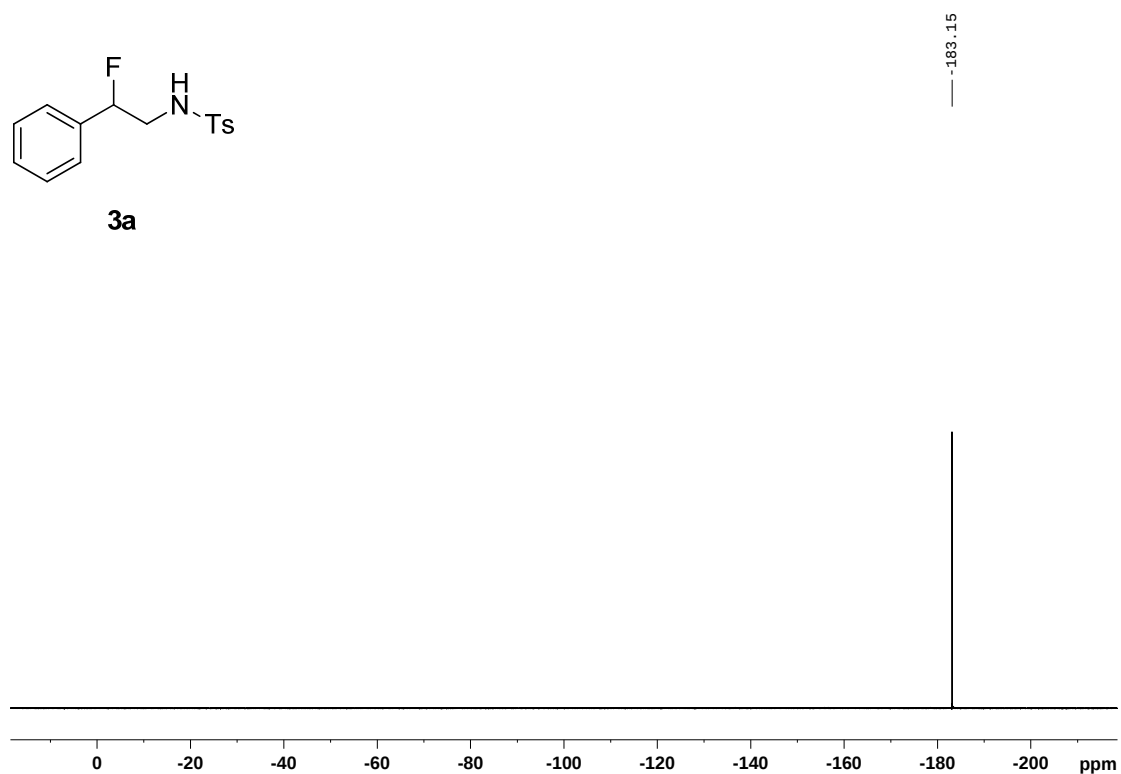
¹H NMR spectrum of **3a**



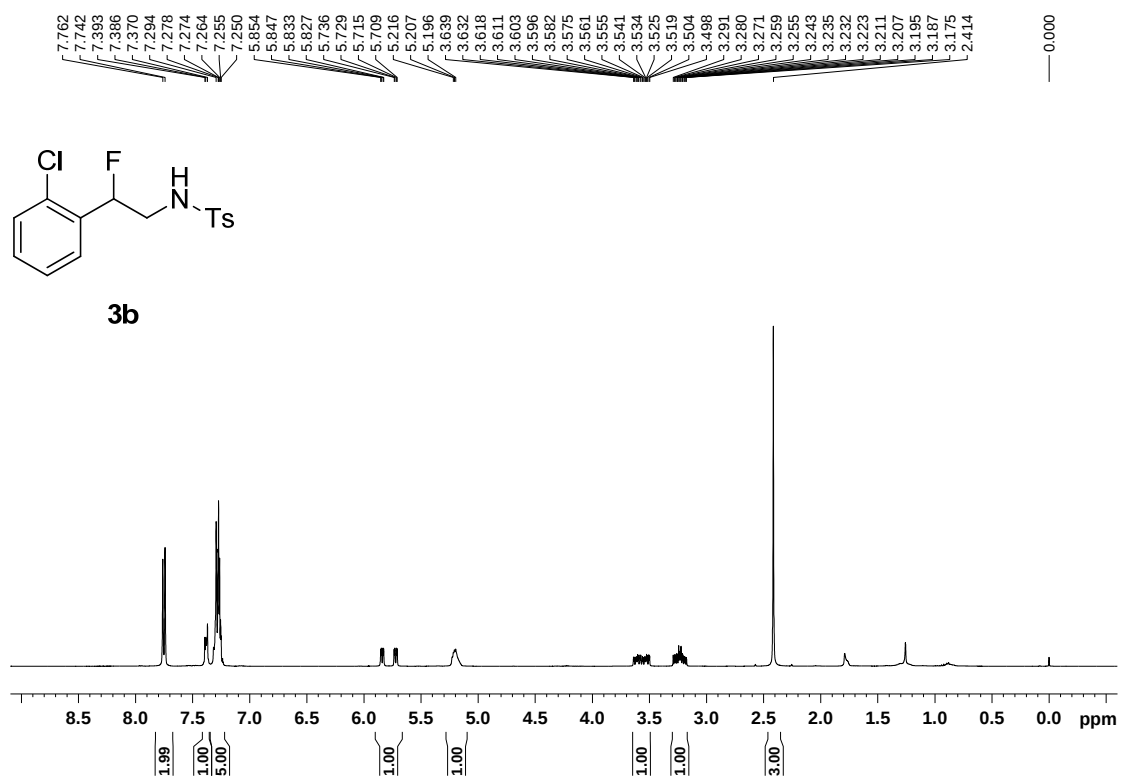
¹³C NMR spectrum of **3a**



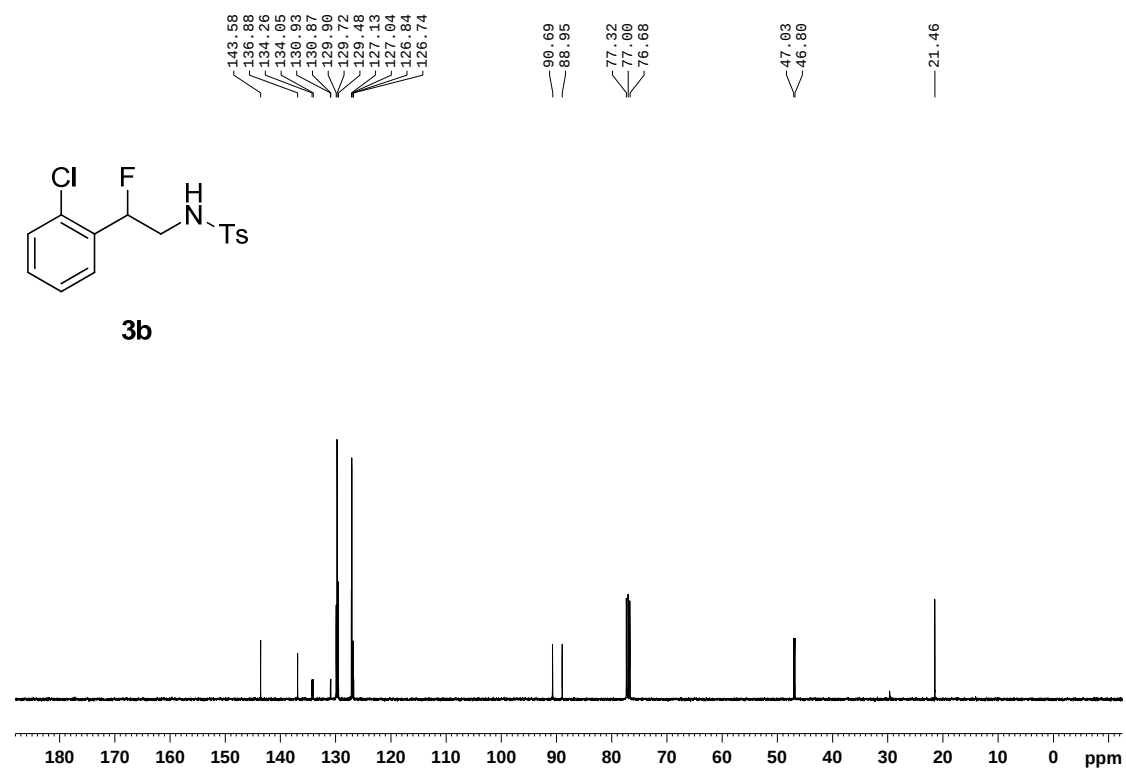
^{19}F NMR spectrum of **3a**



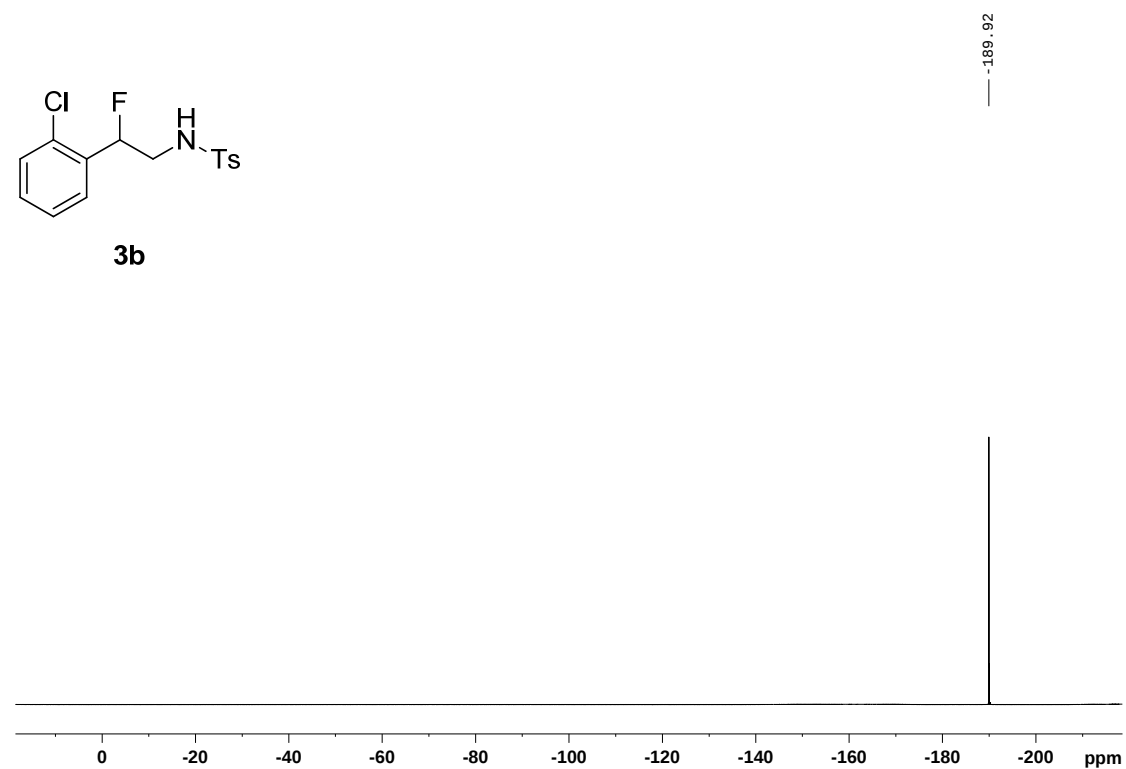
^1H NMR spectrum of **3b**



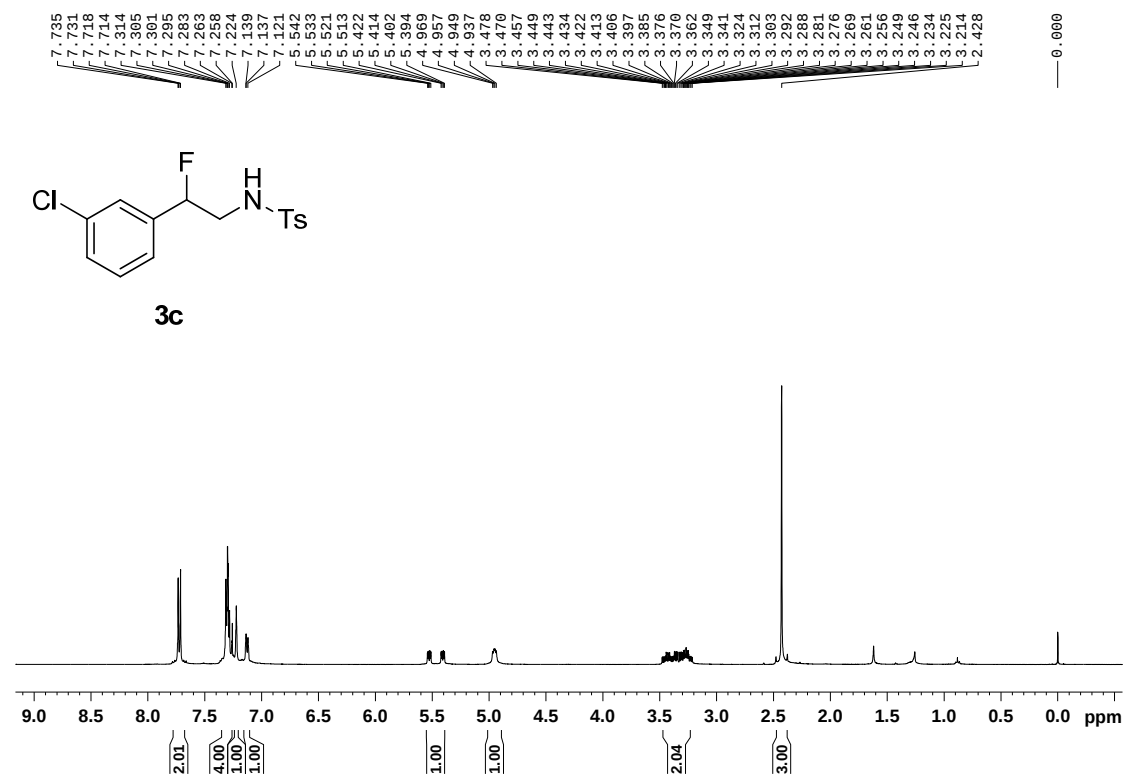
^{13}C NMR spectrum of **3b**



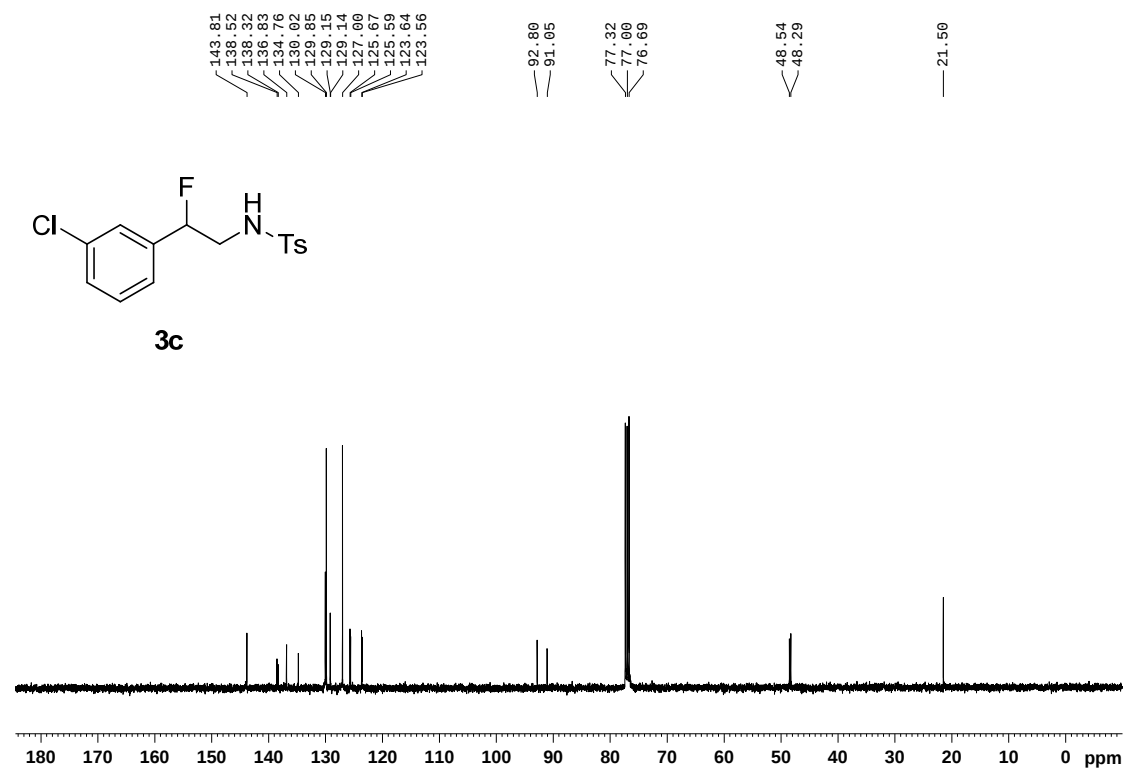
^{19}F NMR spectrum of **3b**



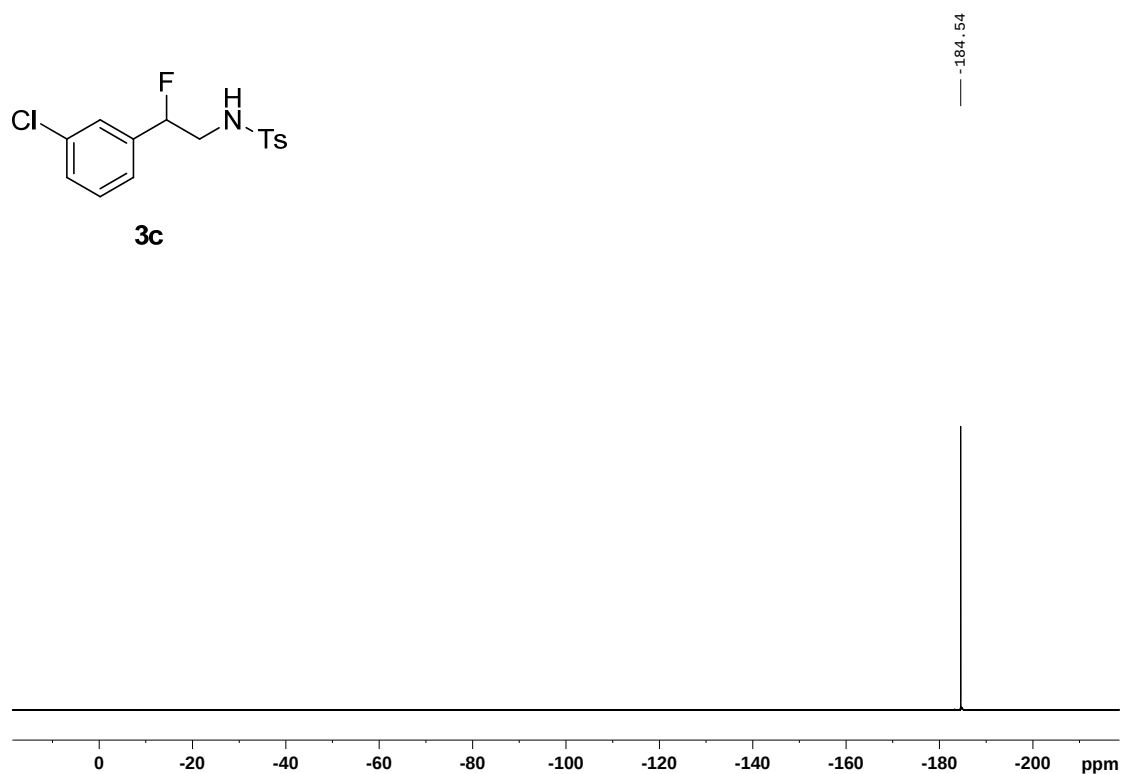
¹H NMR spectrum of **3c**



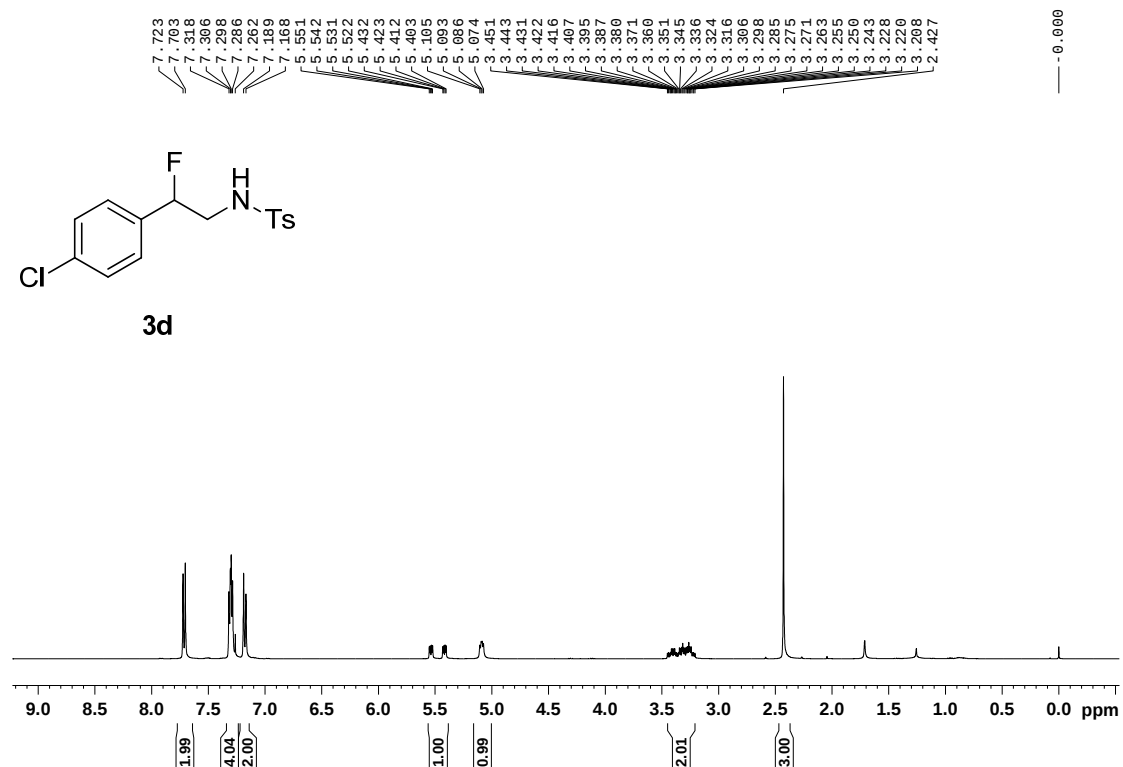
¹³C NMR spectrum of **3c**



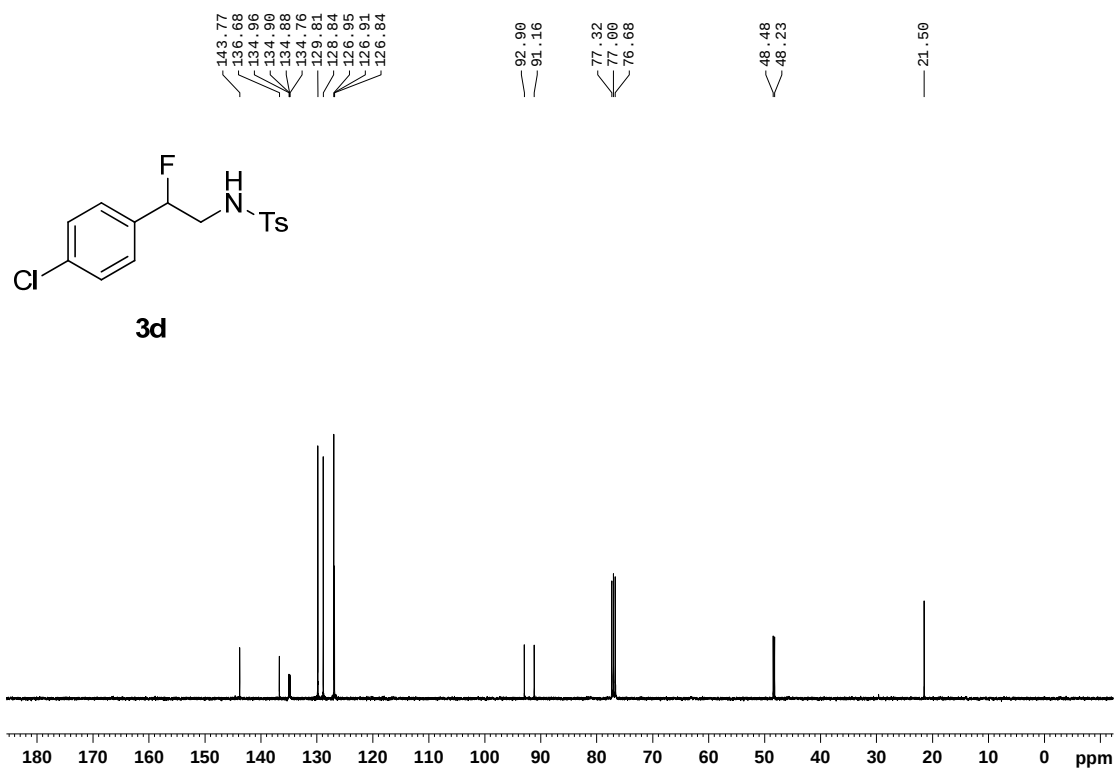
^{19}F NMR spectrum of **3c**



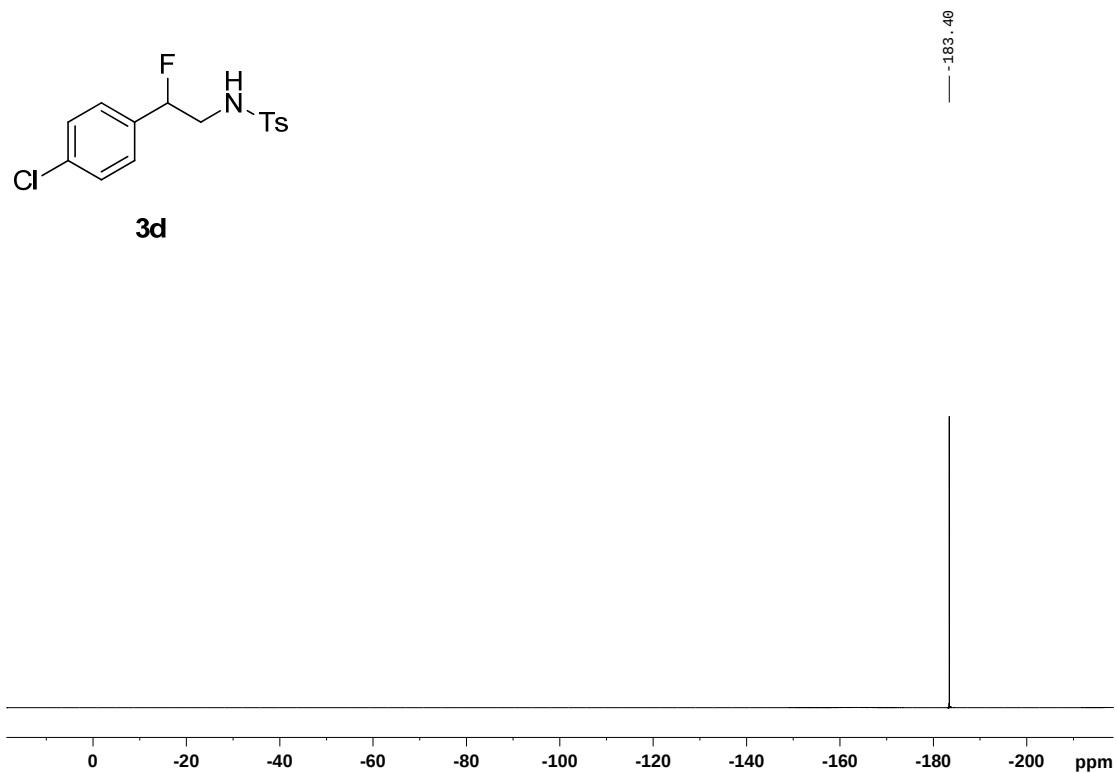
^1H NMR spectrum of **3d**



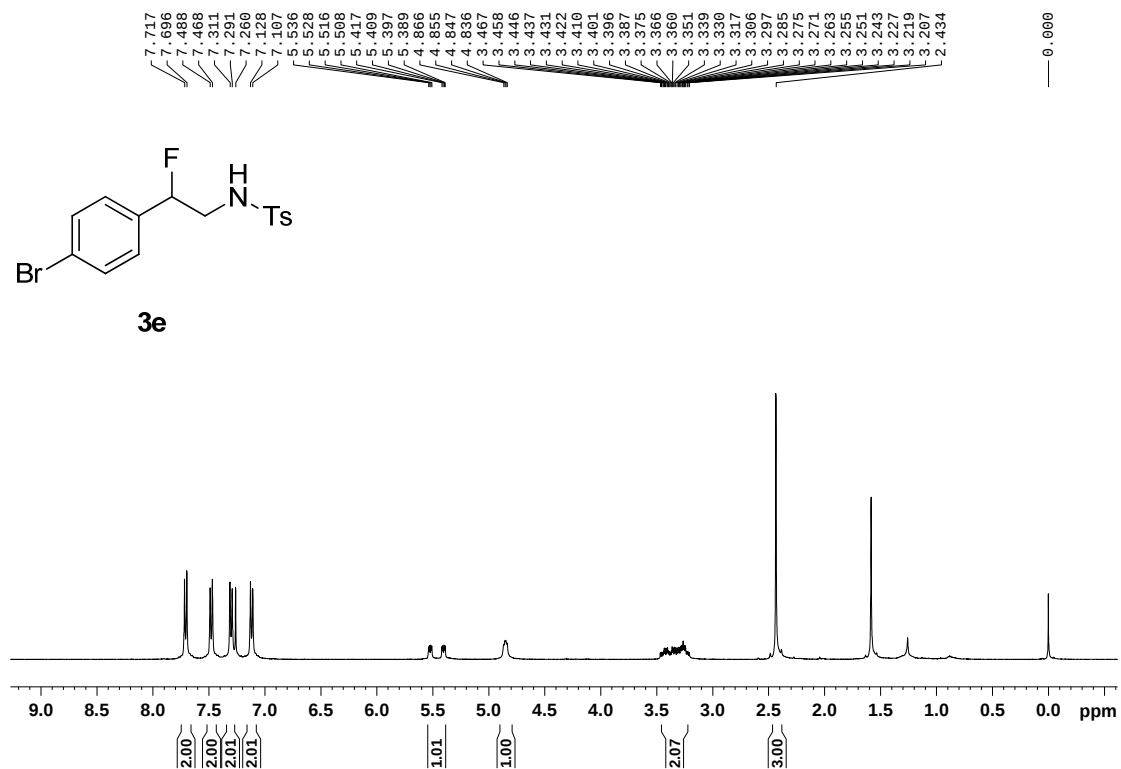
¹³C NMR spectrum of **3d**



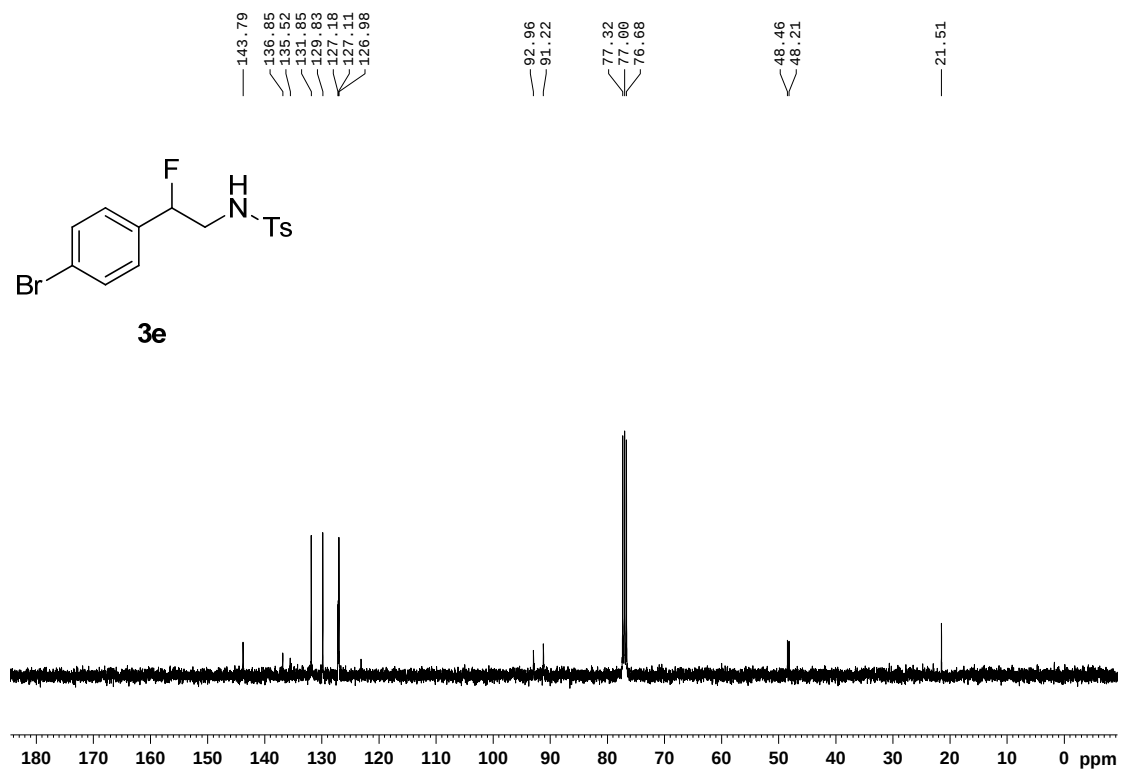
¹⁹F NMR spectrum of **3d**



¹H NMR spectrum of **3e**



¹³C NMR spectrum of **3e**



3e

Chemical structure of **3e**: BrC1=CC=C(C(F)CNc2ccc(S(=O)(=O)C)cc2)C=C1

¹H NMR spectrum (CDCl₃) of compound **3e**. The x-axis represents the chemical shift in ppm, ranging from 0 to 200. The spectrum shows a multiplet at 7.2-7.4 ppm (aromatic protons), a doublet at 6.8 ppm (NH), a multiplet at 4.2-4.4 ppm (CH₂), and a singlet at 3.0 ppm (F).

3f

Chemical structure of **3f** is shown above the spectrum. The spectrum displays peaks corresponding to the structure, with integration values and chemical shifts (ppm) indicated.

Chemical Shift (ppm)	Integration
7.741, 7.721, 7.711, 7.701, 7.690, 7.681, 7.671, 7.274, 7.273, 7.272, 7.213, 7.048, 7.065, 7.066	1.99, 1.98, 2.00, 1.98
5.549, 5.540, 5.528, 5.519, 5.519, 5.430, 5.420, 5.410, 5.400, 5.135, 5.123, 5.115, 5.104	1.00, 0.97
3.441, 3.432, 3.421, 3.411, 3.406, 3.397, 3.385, 3.376, 3.369, 3.360, 3.349, 3.340, 3.333, 3.332, 3.325, 3.320, 3.313, 3.311, 3.304, 3.299, 3.290, 3.284, 3.278, 3.270, 3.264, 3.258, 3.243, 3.235, 3.223, 2.425	2.02, 3.00

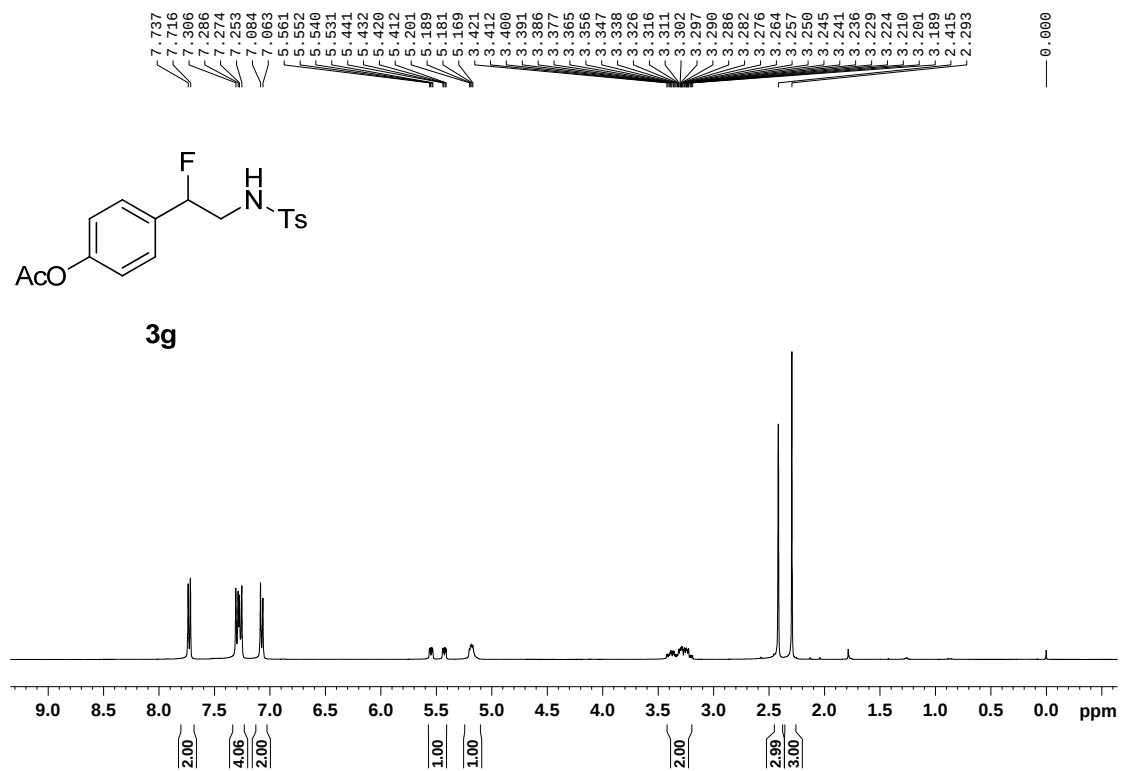
CC(F)(c1ccc(F)cc1)CNc2ccc(cc2)S(=O)(=O)c3ccccc3

3f

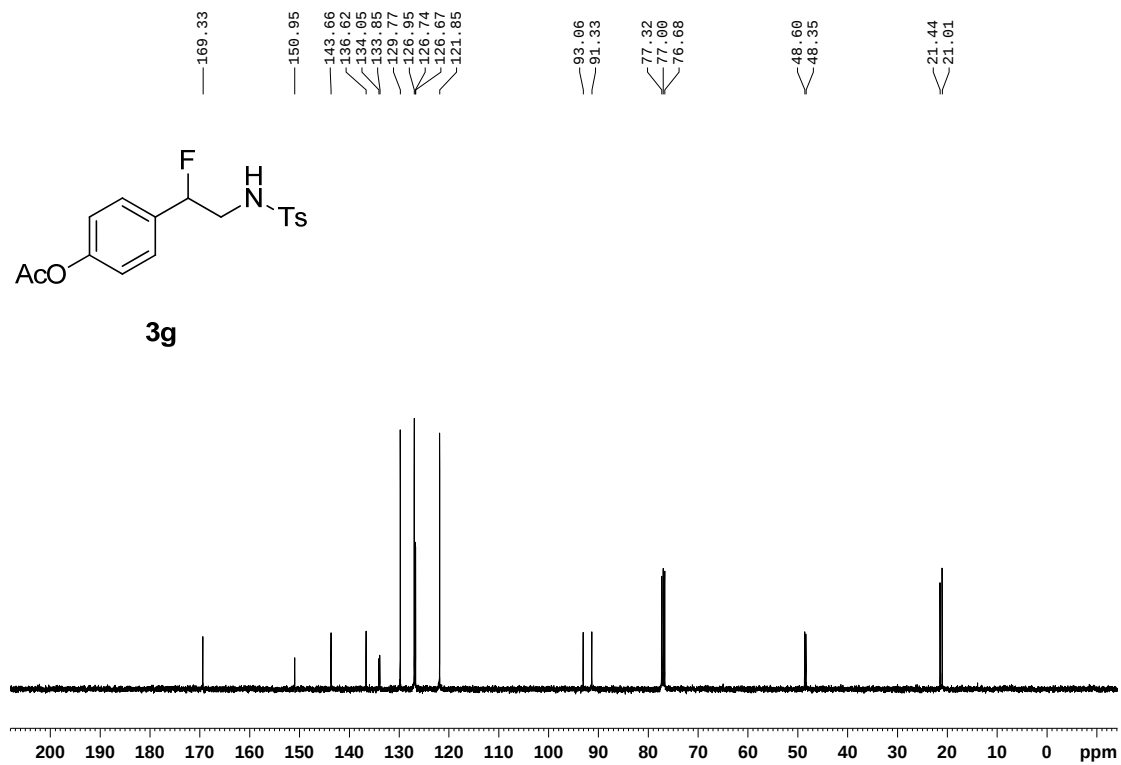
164.19
 164.17
 161.73
 161.70
 143.74
 136.73
 132.34
 132.31
 132.14
 132.11
 129.80
 127.57
 127.51
 127.49
 127.42
 126.97
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 77.00
 76.68
 48.56
 48.31
 21.47

[illegible]

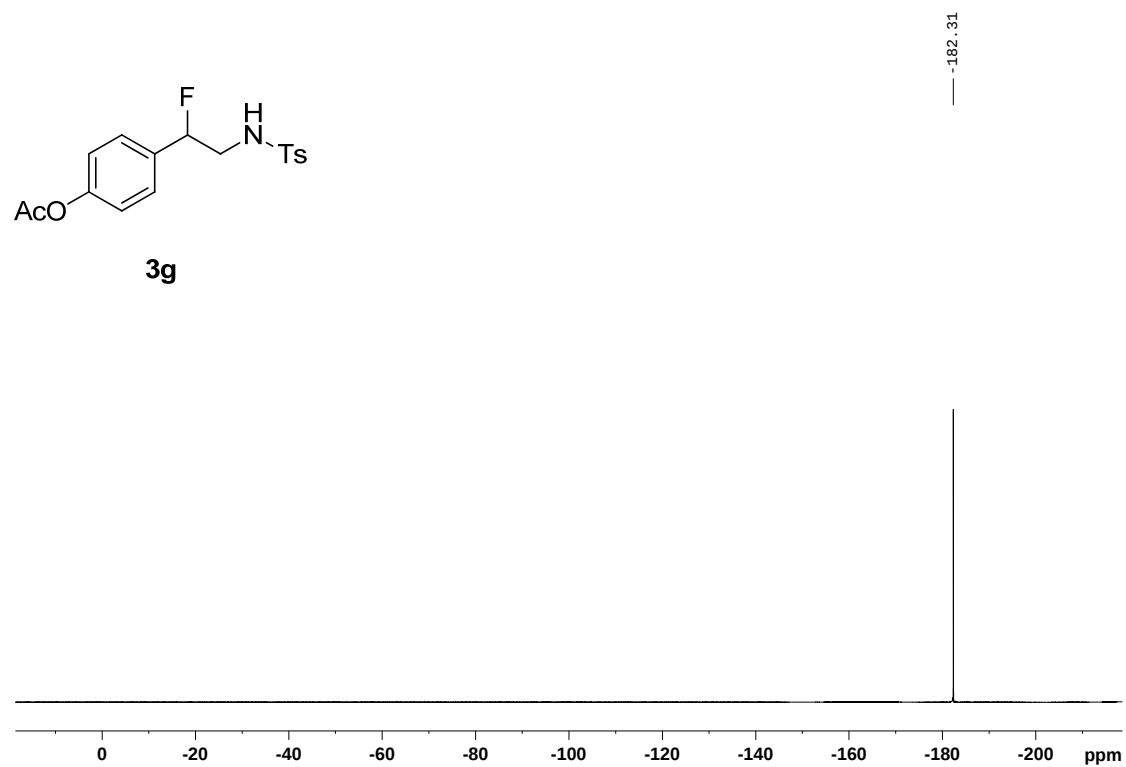
¹H NMR spectrum of **3g**



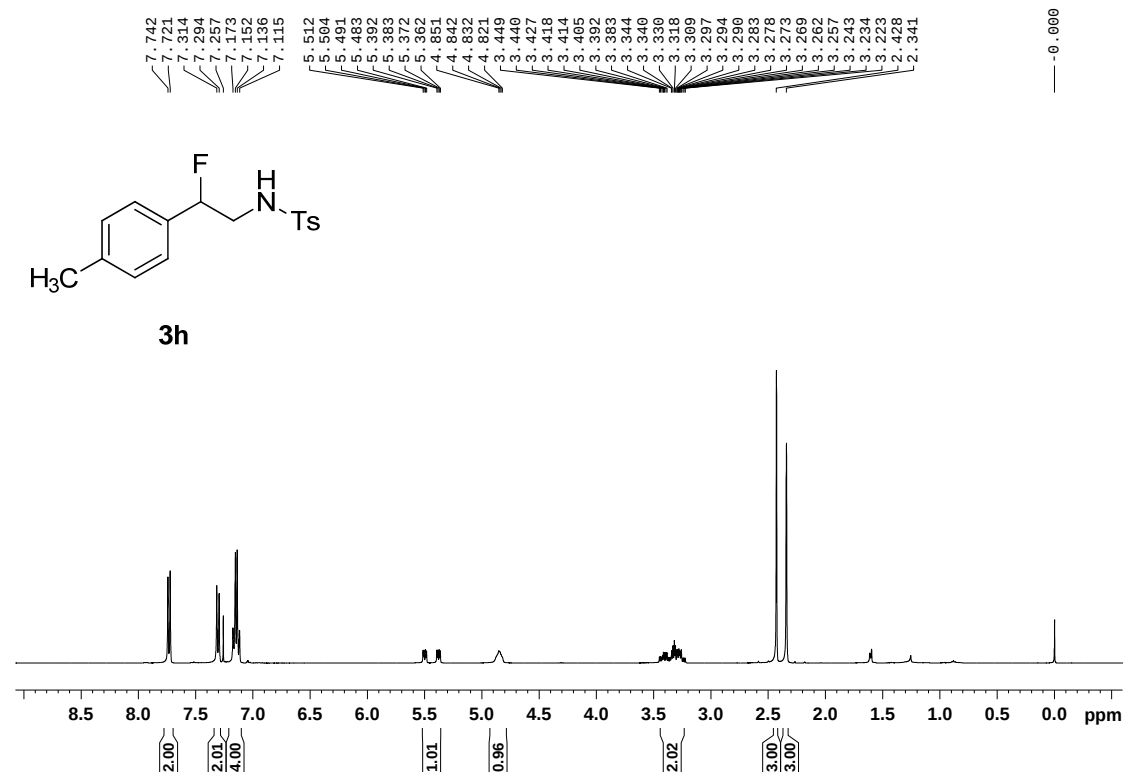
¹³C NMR spectrum of **3g**



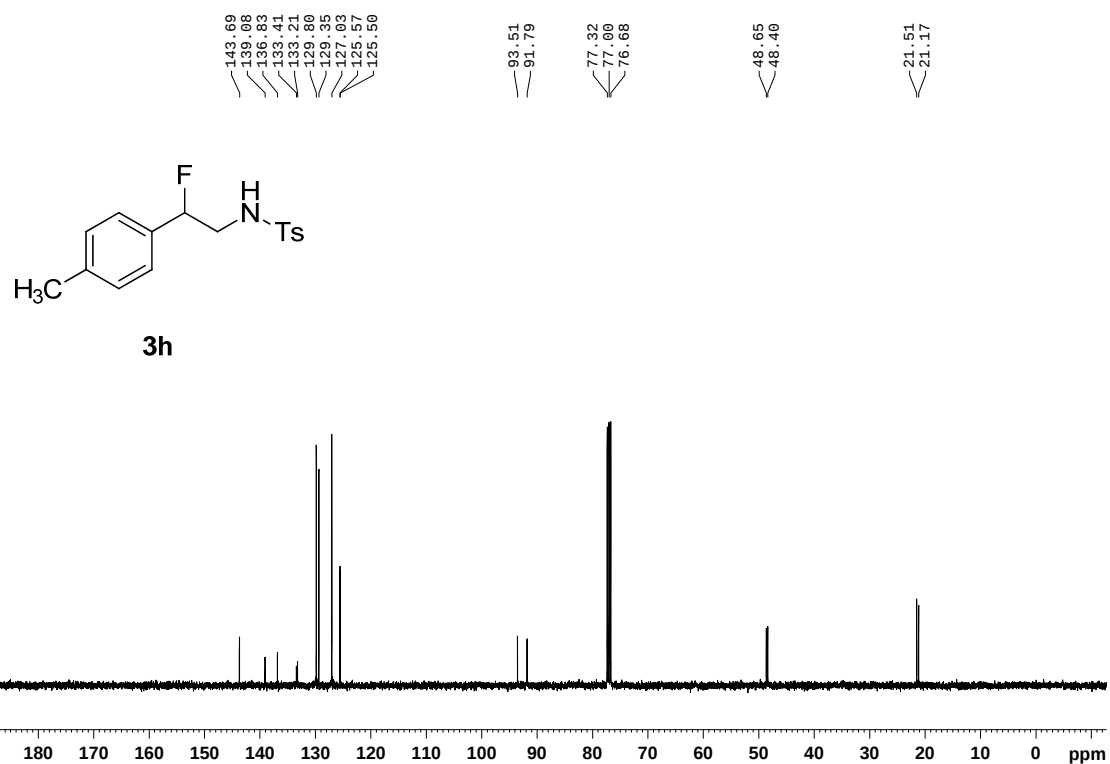
^{19}F NMR spectrum of **3g**



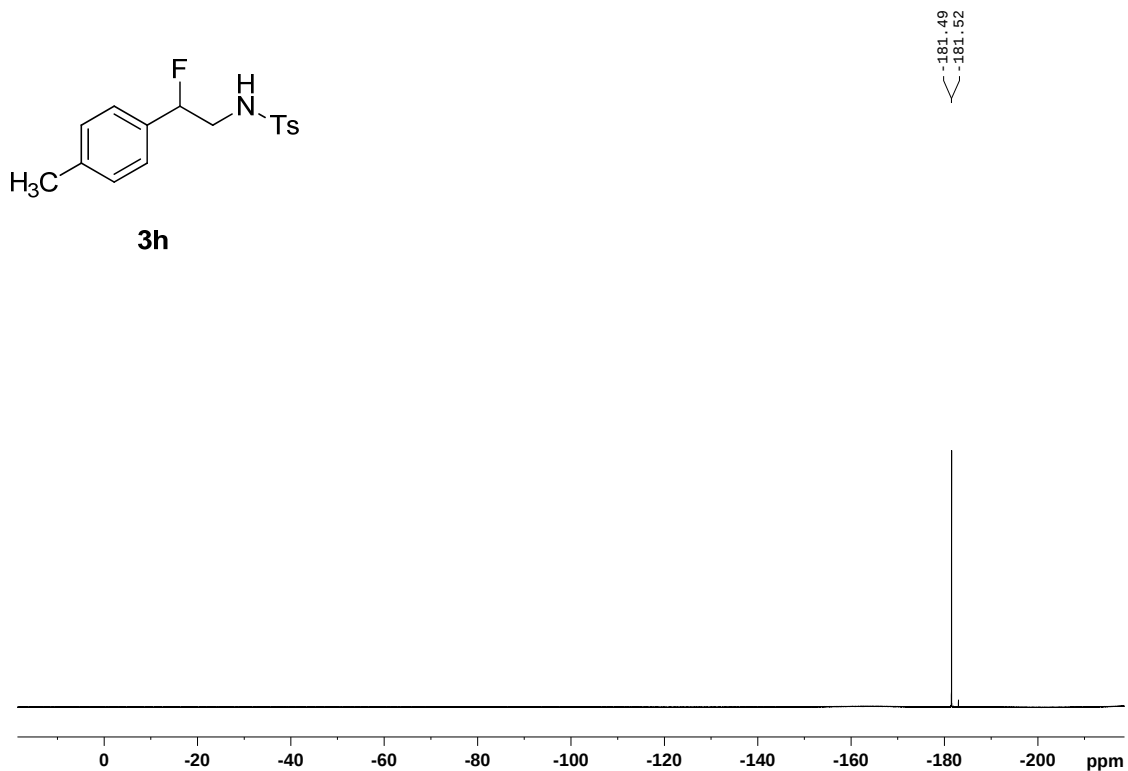
^1H NMR spectrum of **3h**



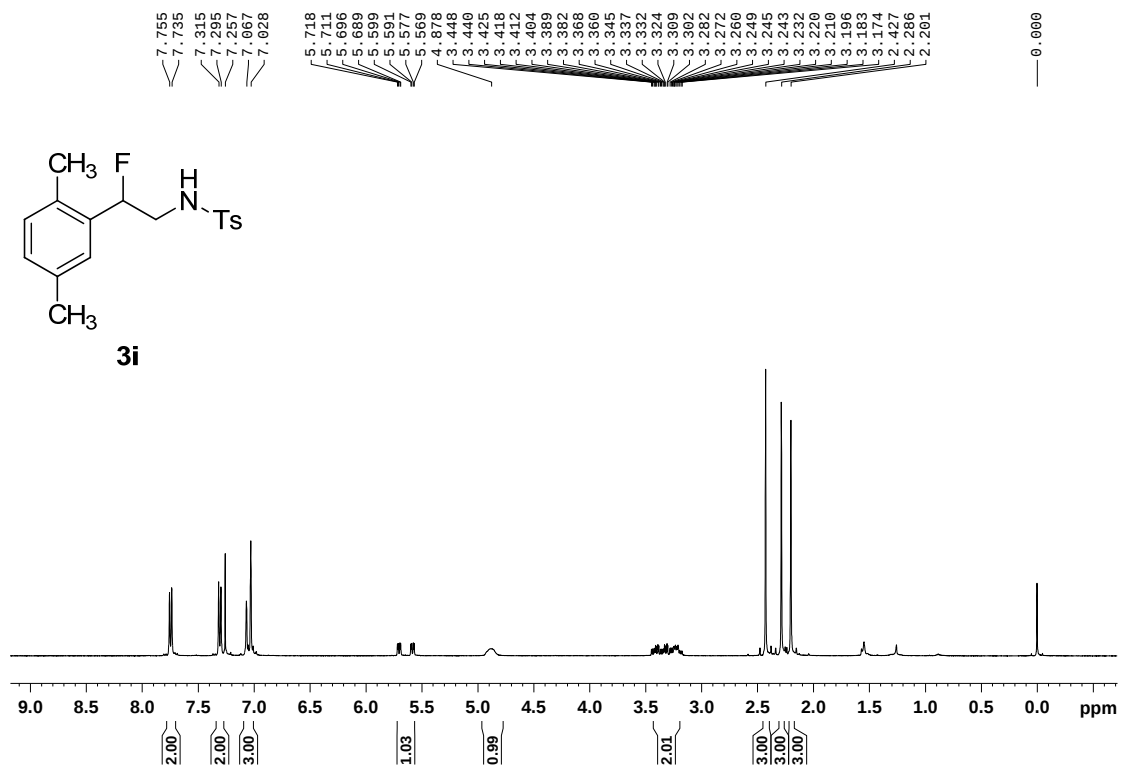
¹³C NMR spectrum of **3h**



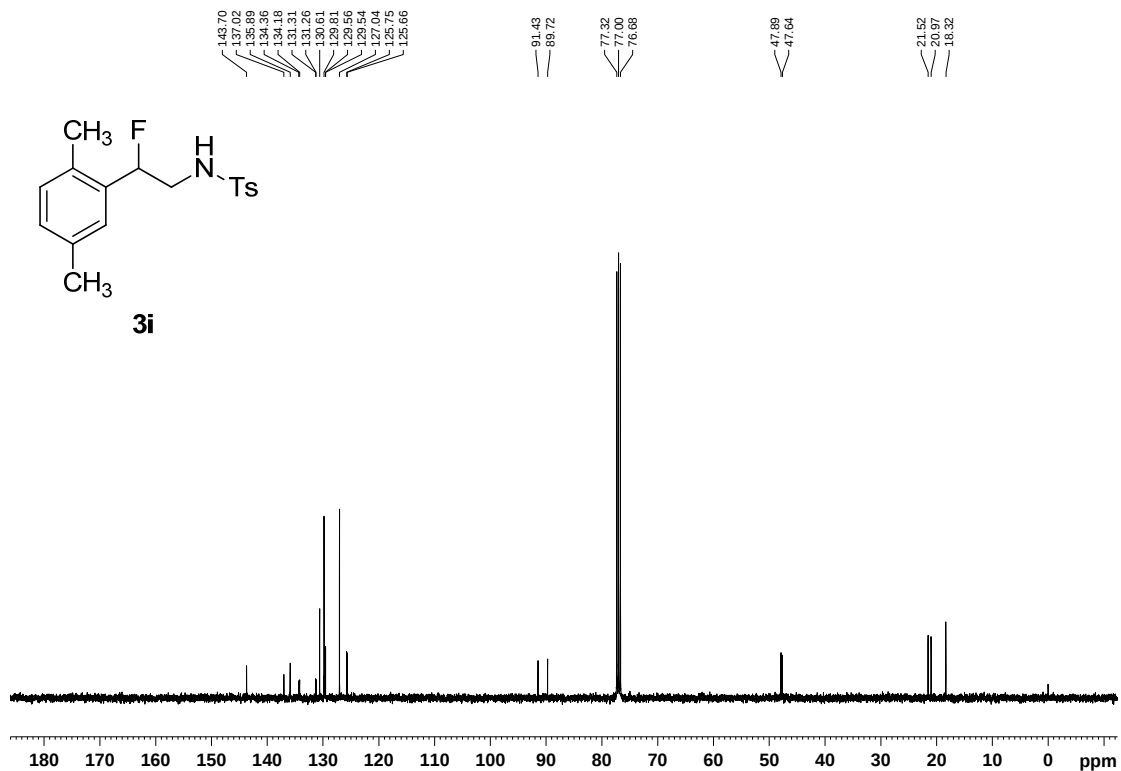
¹⁹F NMR spectrum of **3h**



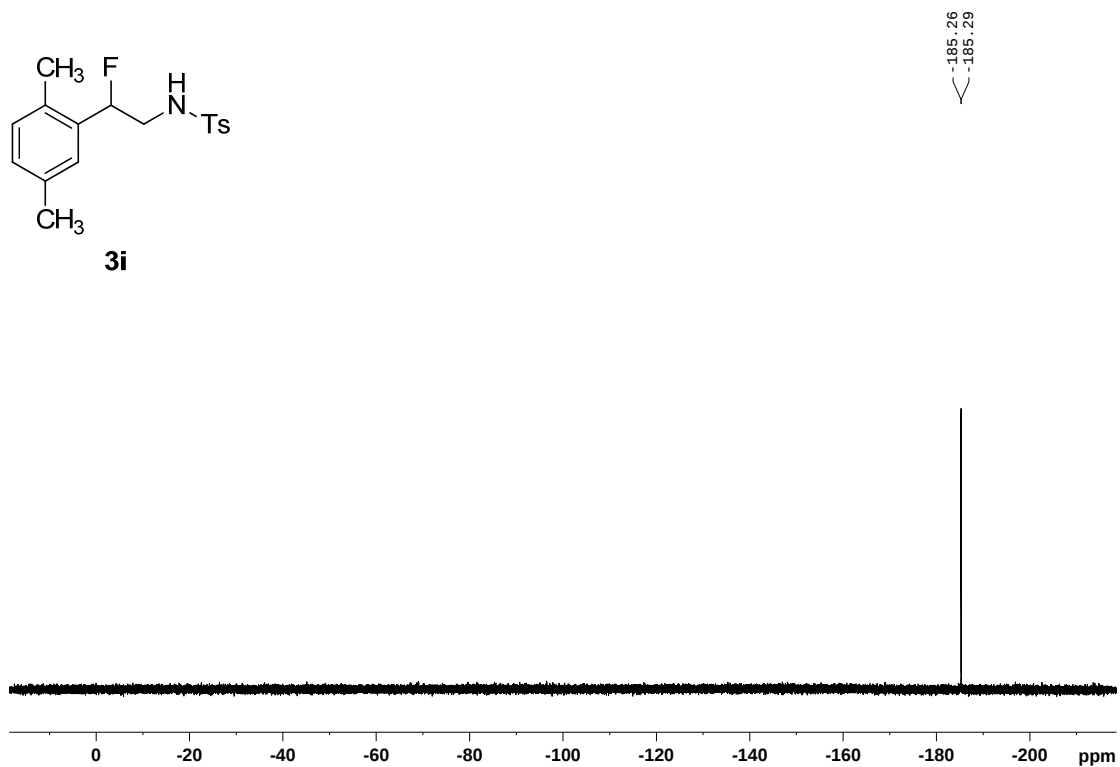
¹H NMR spectrum of **3i**



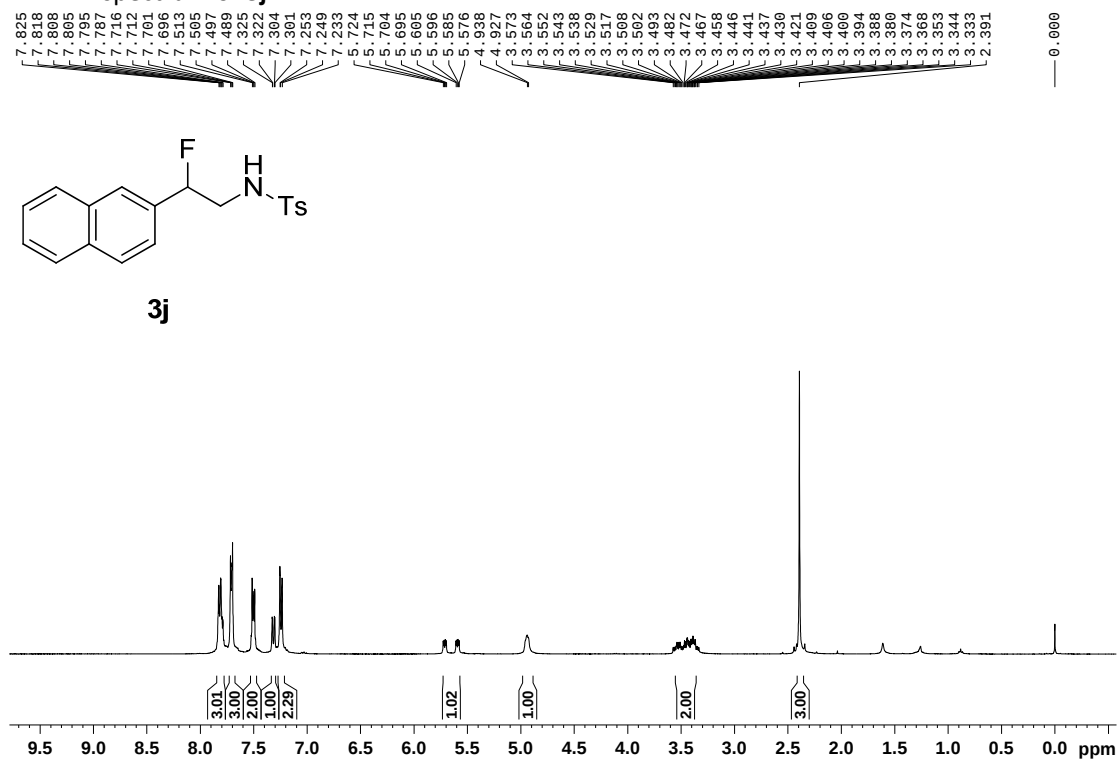
¹³C NMR spectrum of **3i**



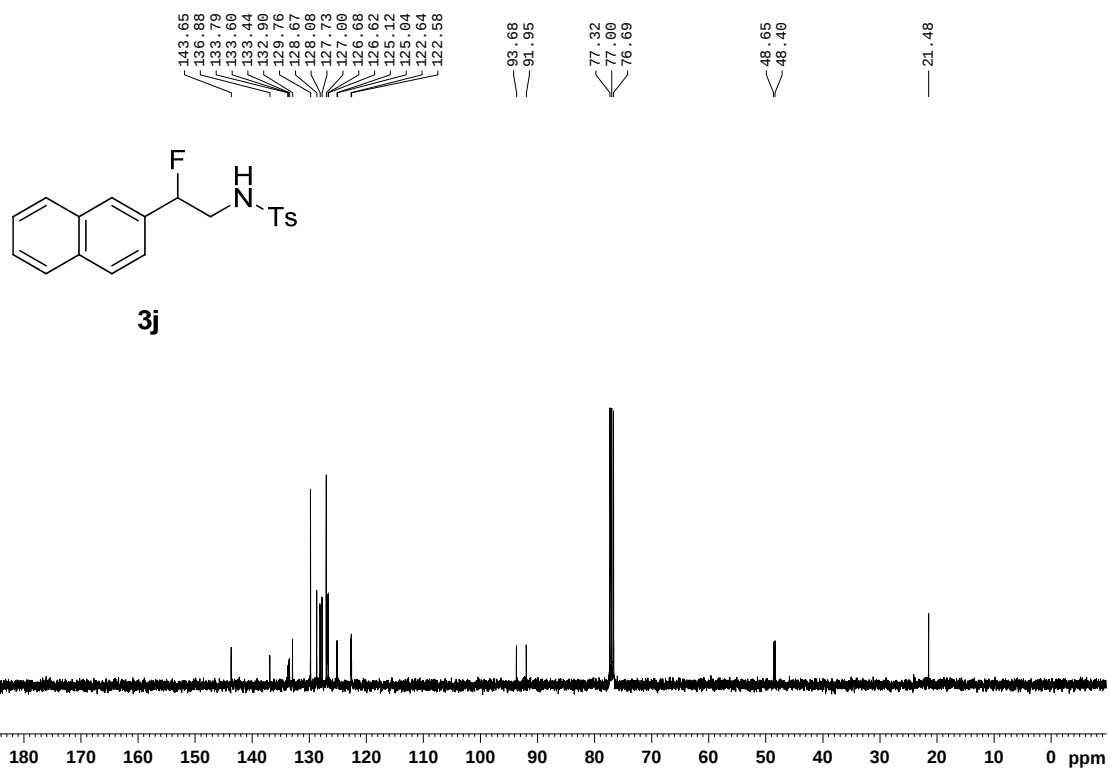
^{19}F NMR spectrum of **3i**



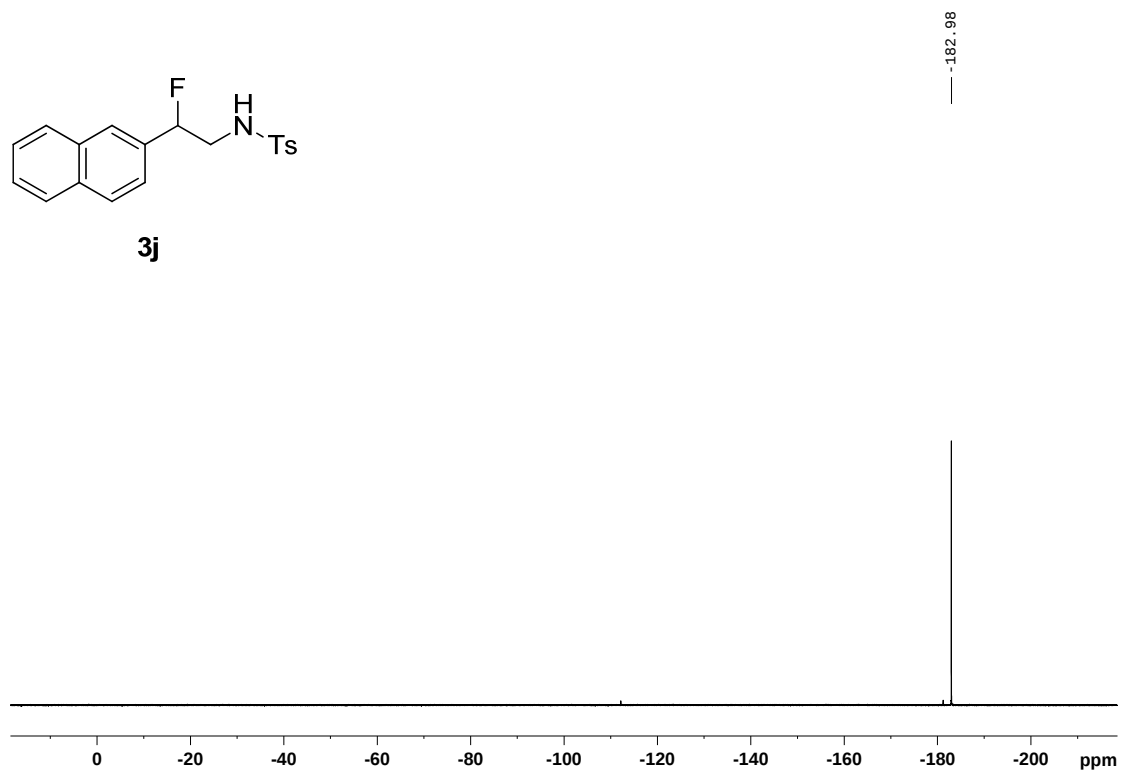
^1H NMR spectrum of **3j**



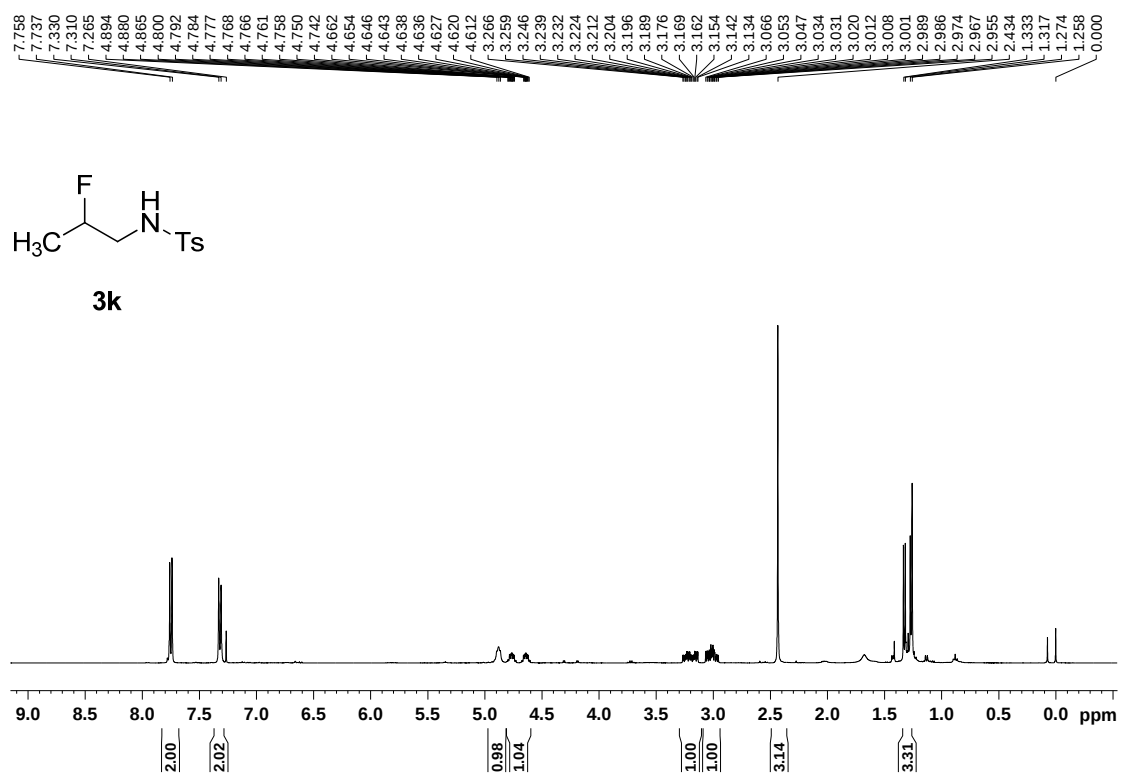
¹³C NMR spectrum of **3j**



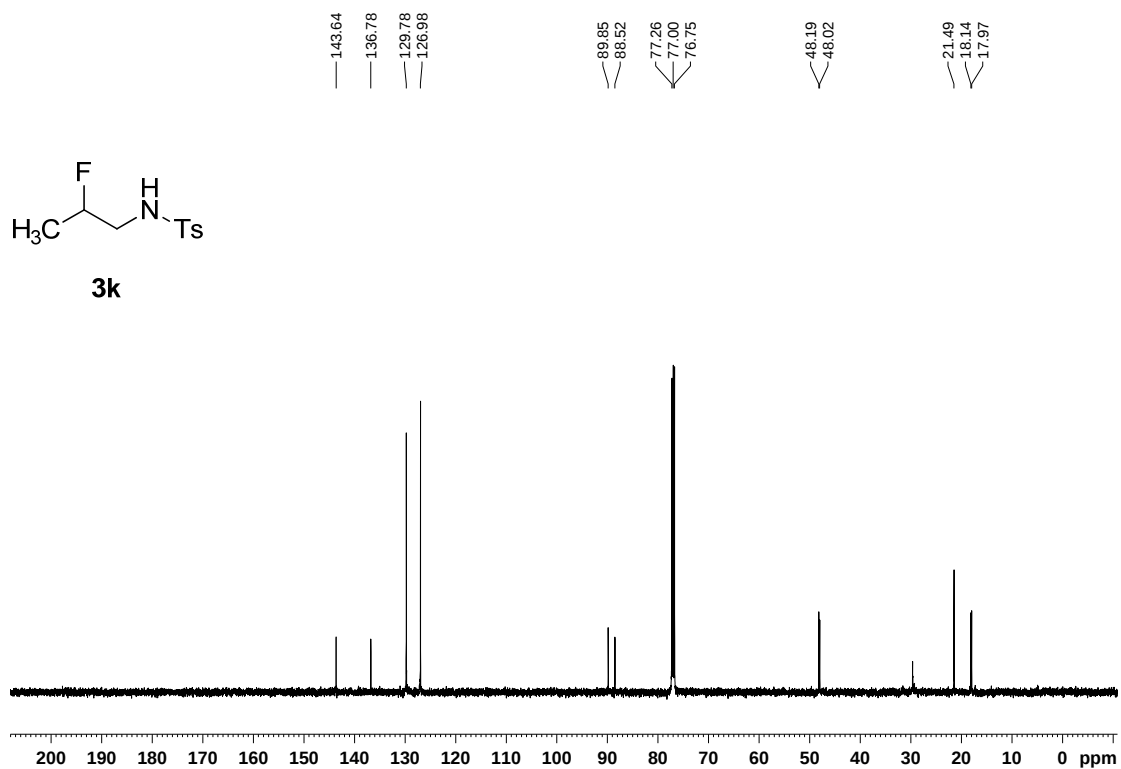
¹⁹F NMR spectrum of **3j**



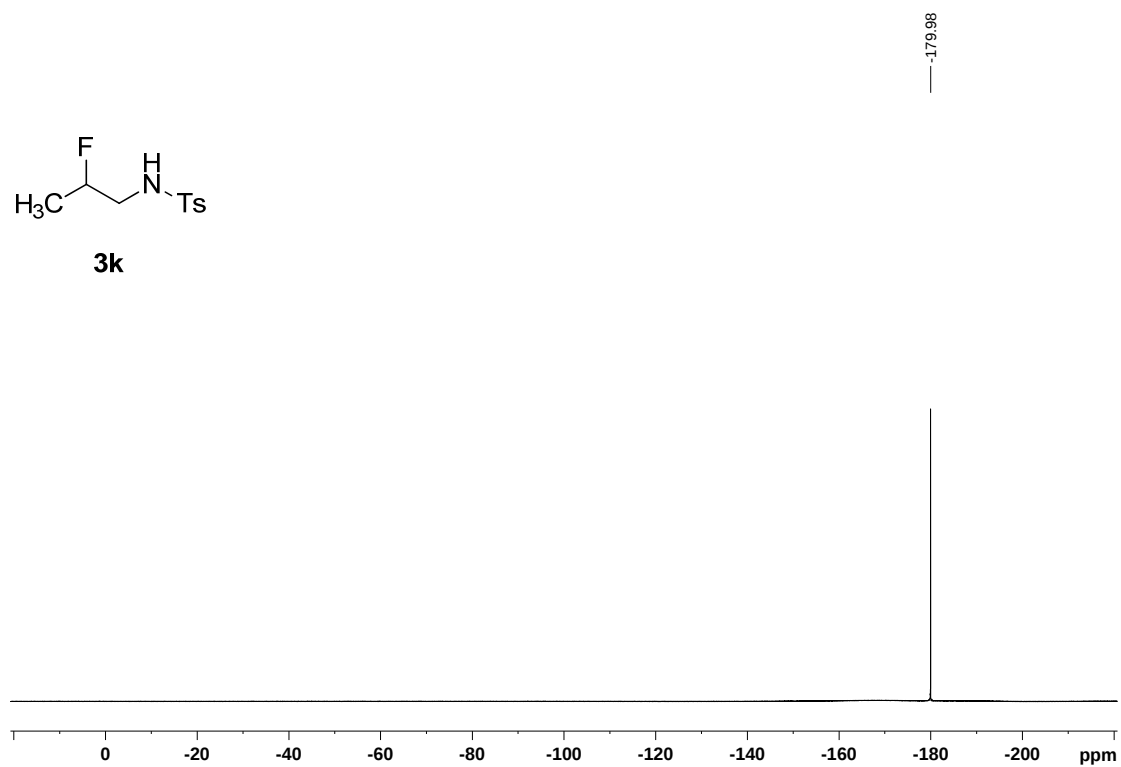
¹⁹H NMR spectrum of **3k**



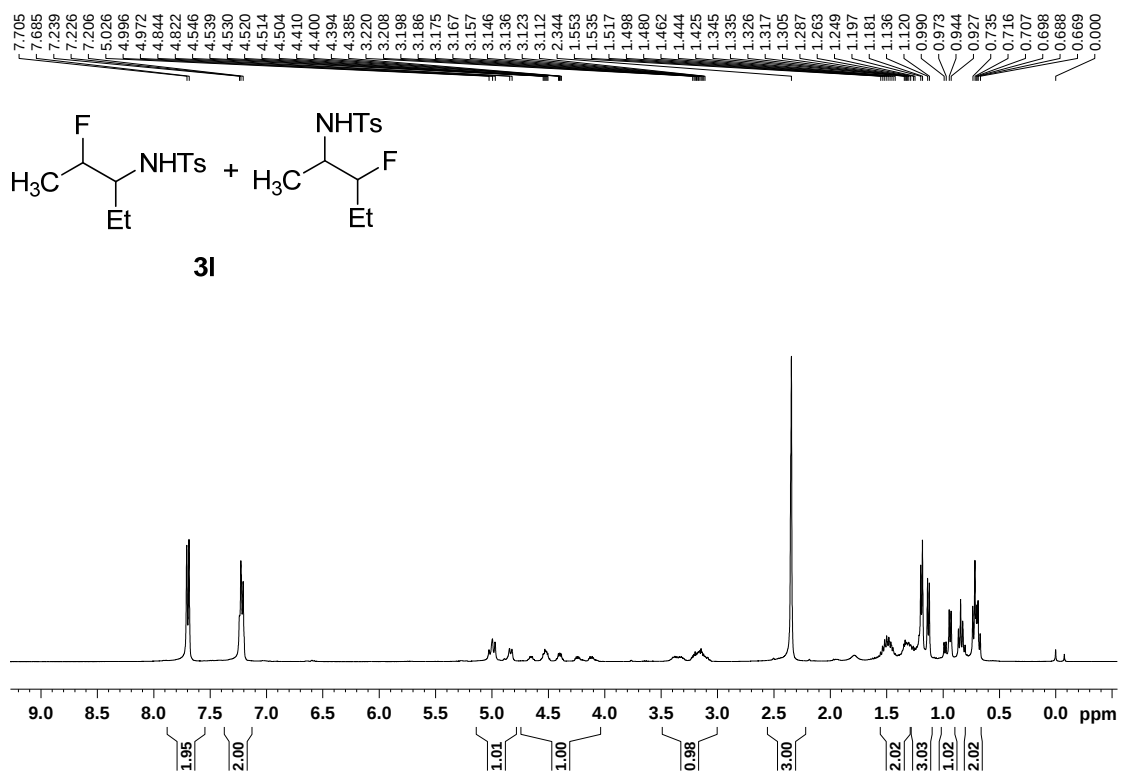
¹³C NMR spectrum of **3k**



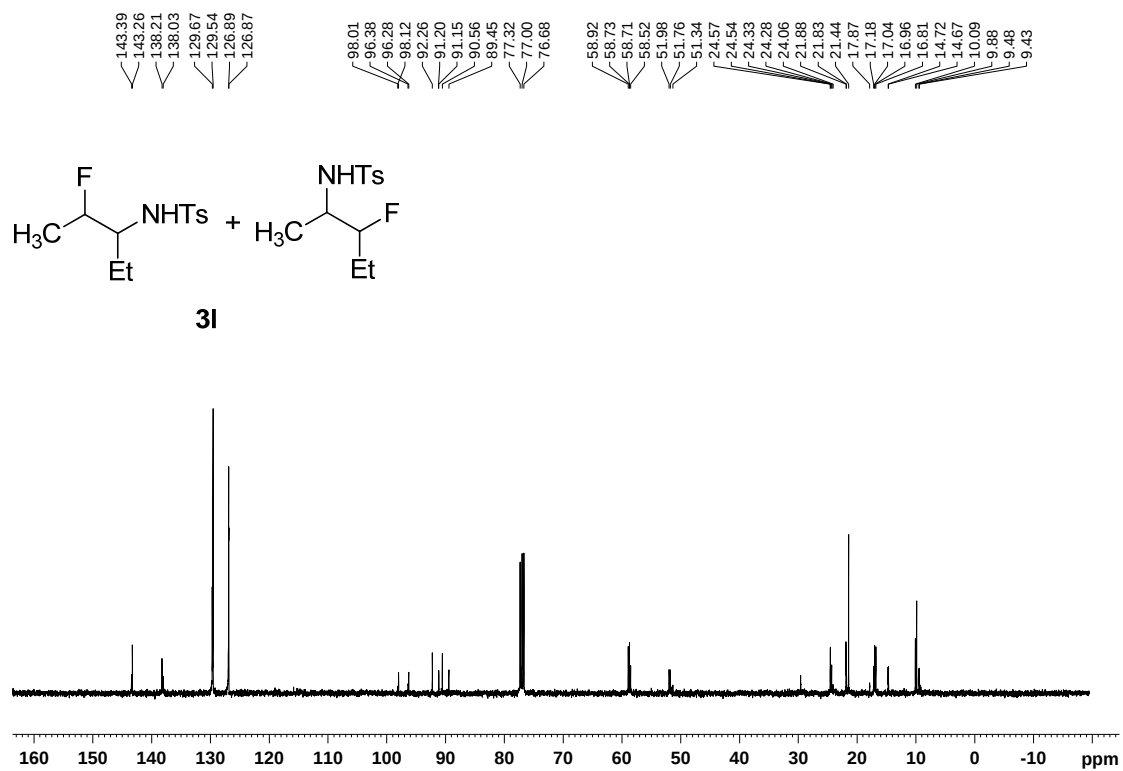
¹⁹F NMR spectrum of **3k**



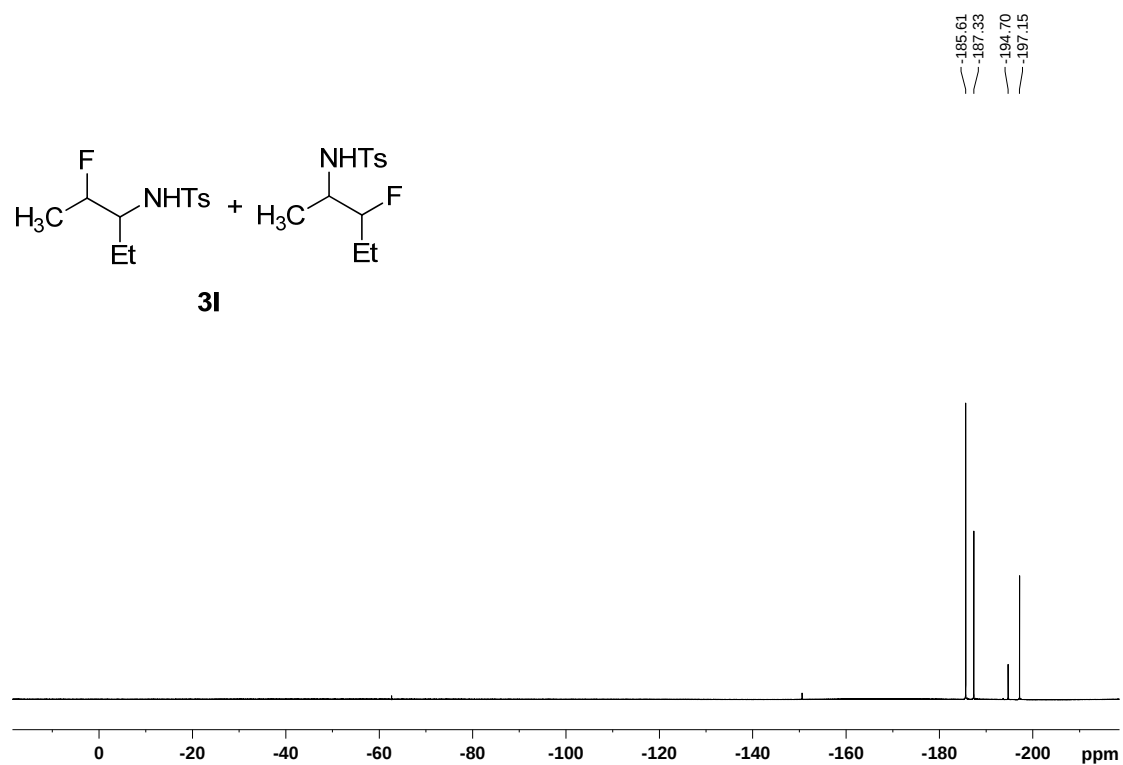
¹H NMR spectrum of **3l**



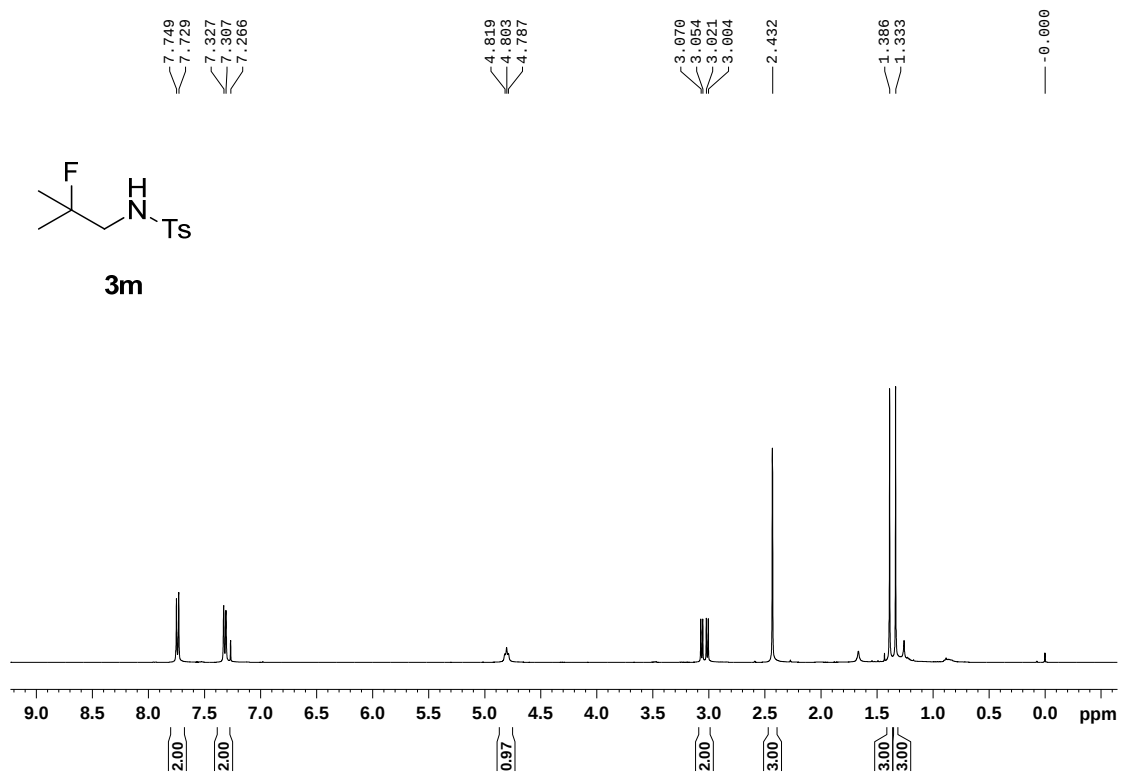
¹³C NMR spectrum of **3I**



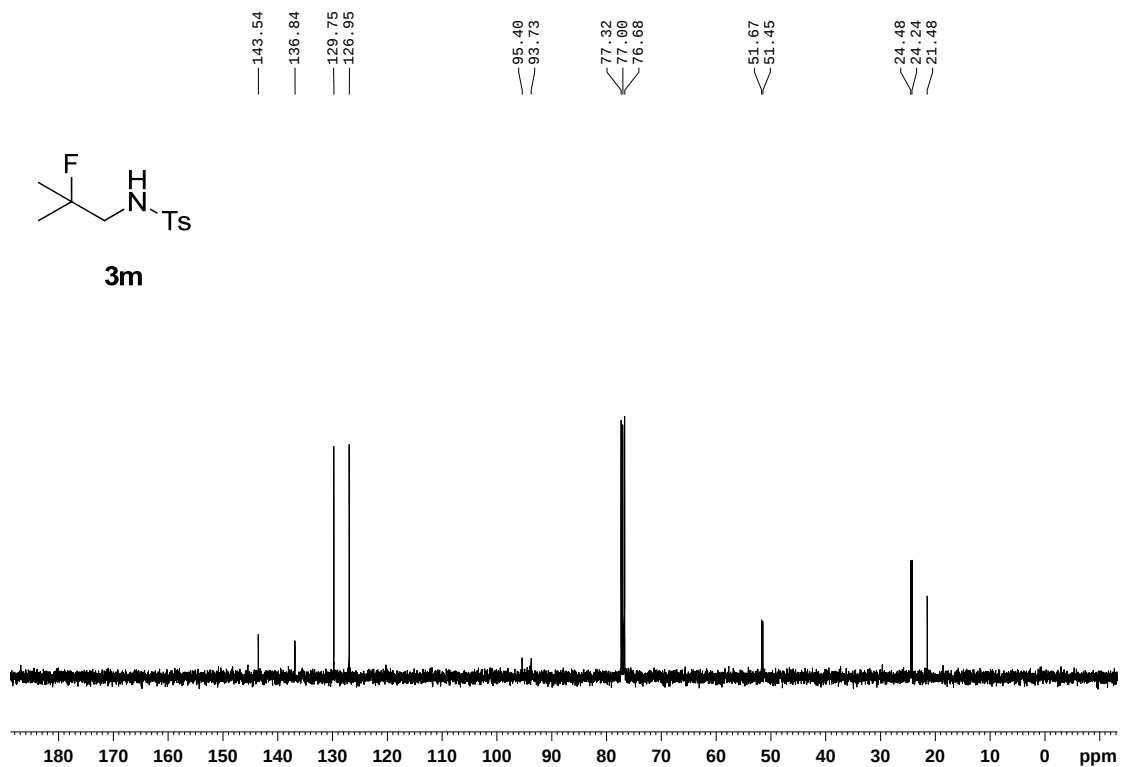
¹⁹F NMR spectrum of **3I**



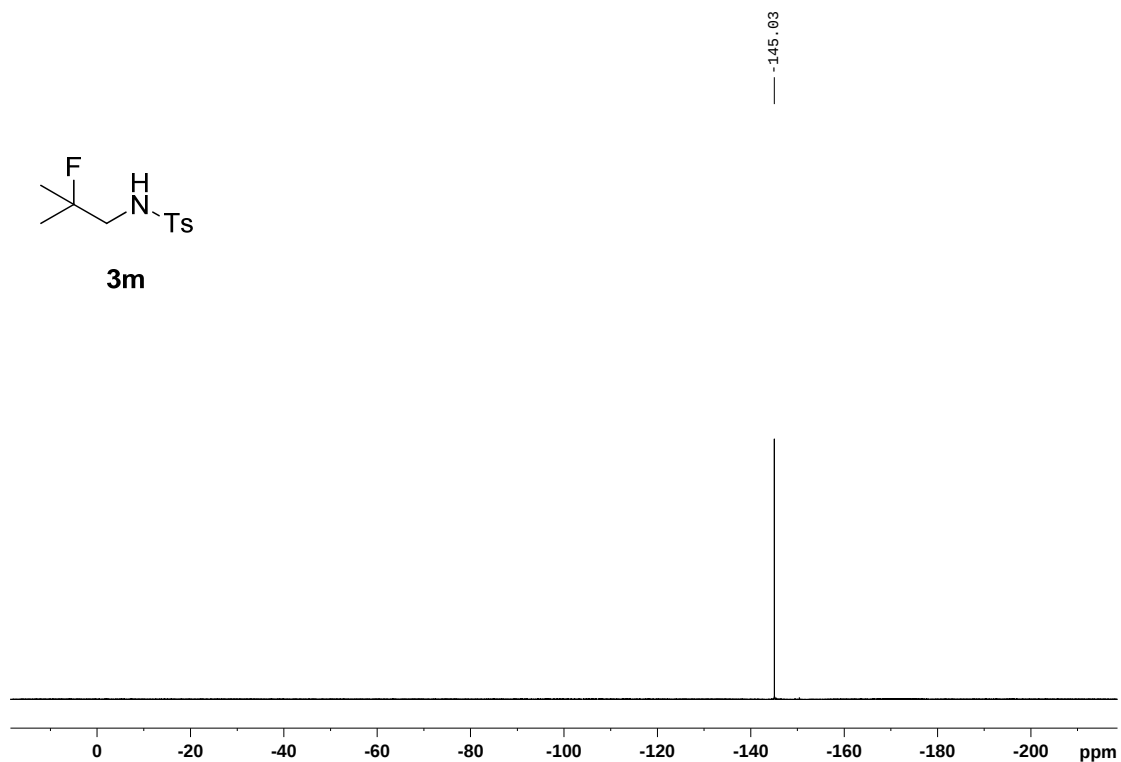
¹H NMR spectrum of **3m**



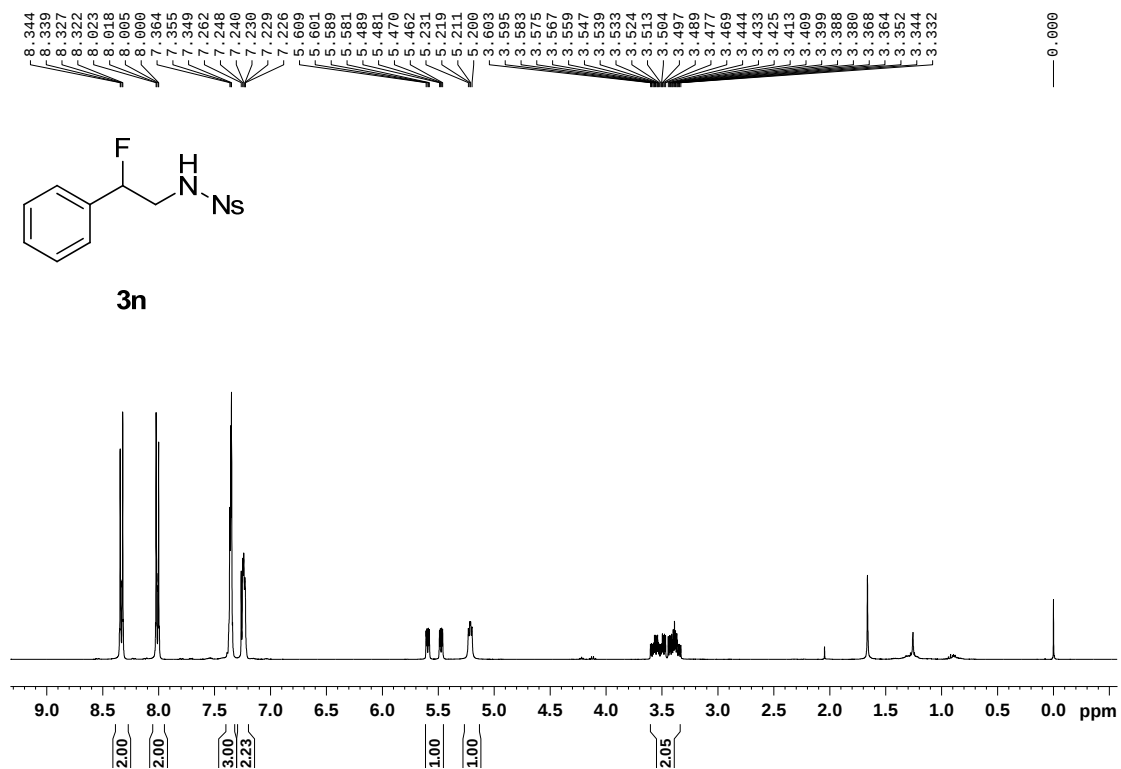
¹³C NMR spectrum of **3m**



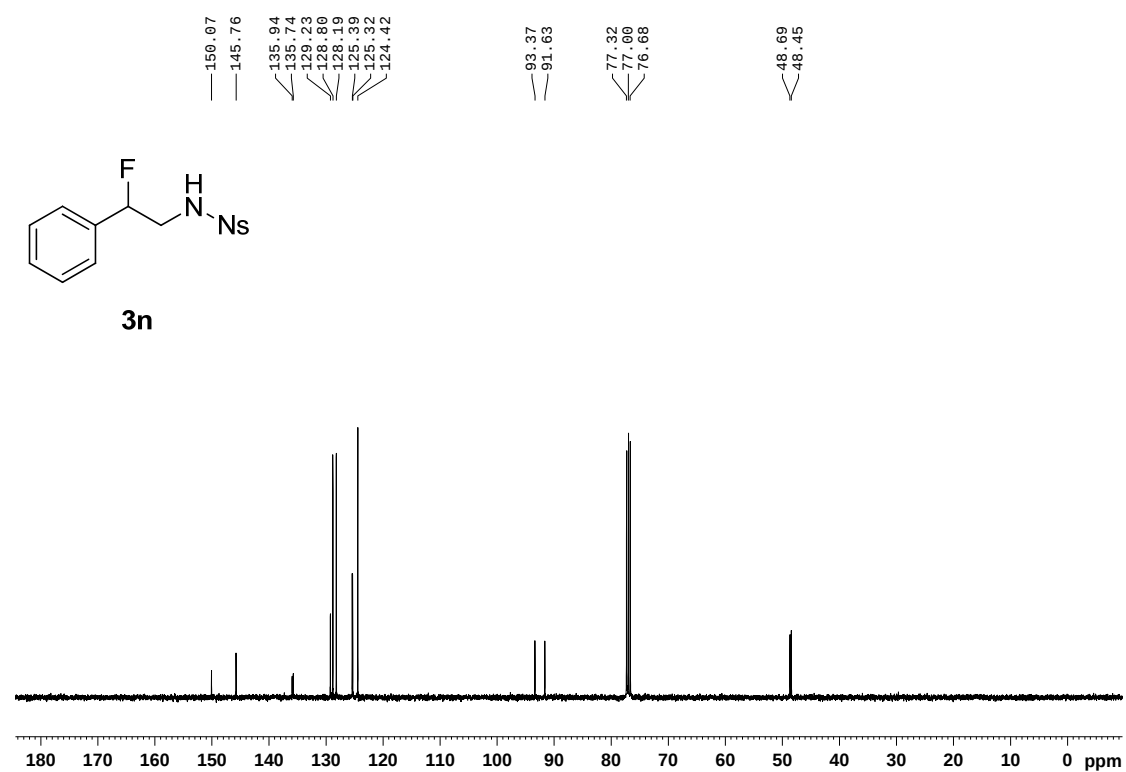
^{19}F NMR spectrum of **3m**



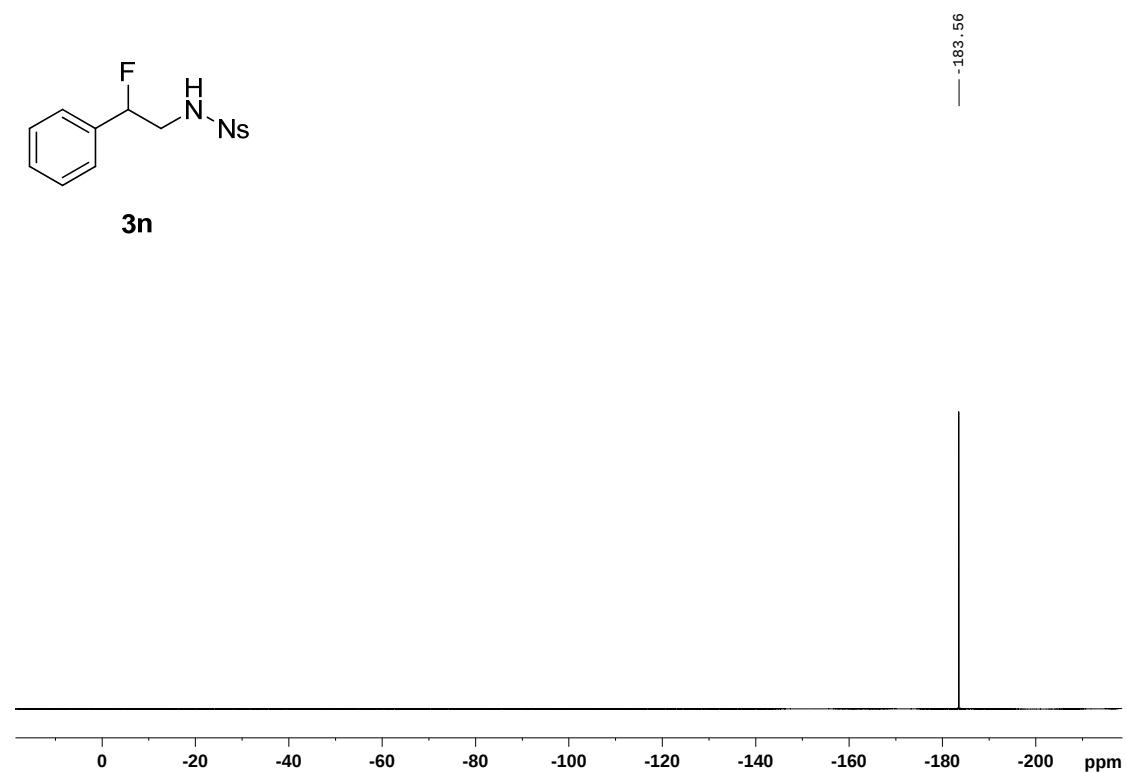
^1H NMR spectrum of **3n**



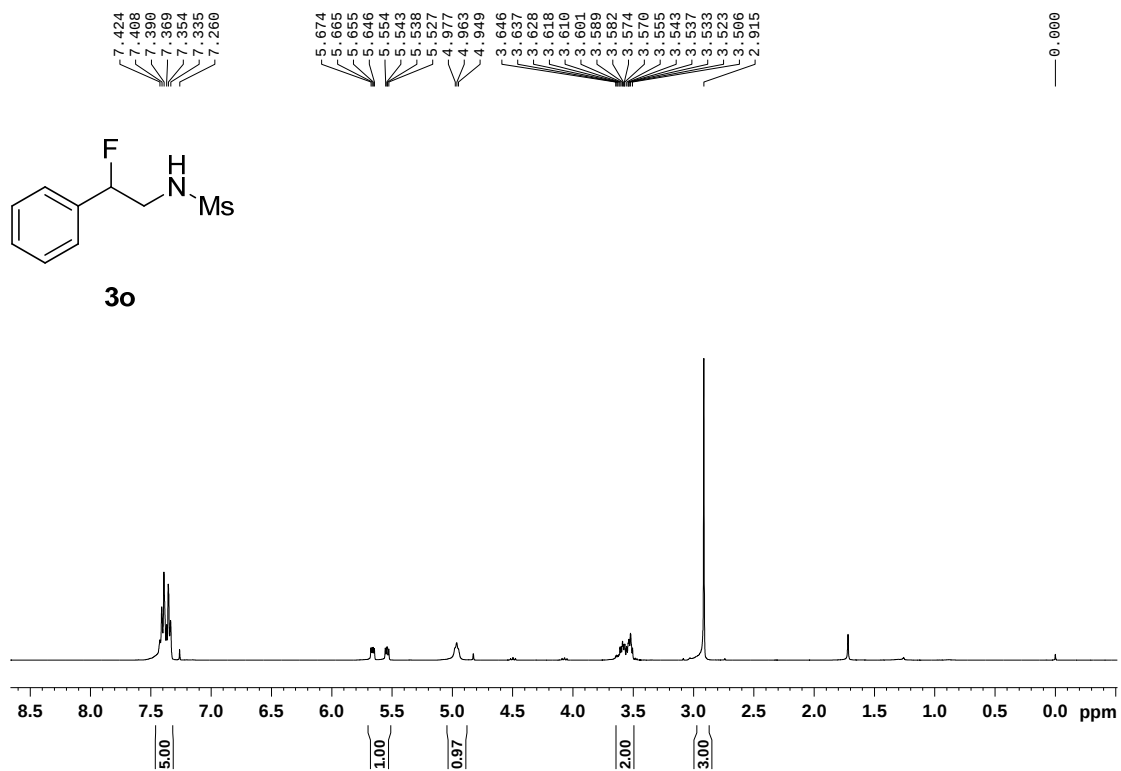
¹³C NMR spectrum of **3n**



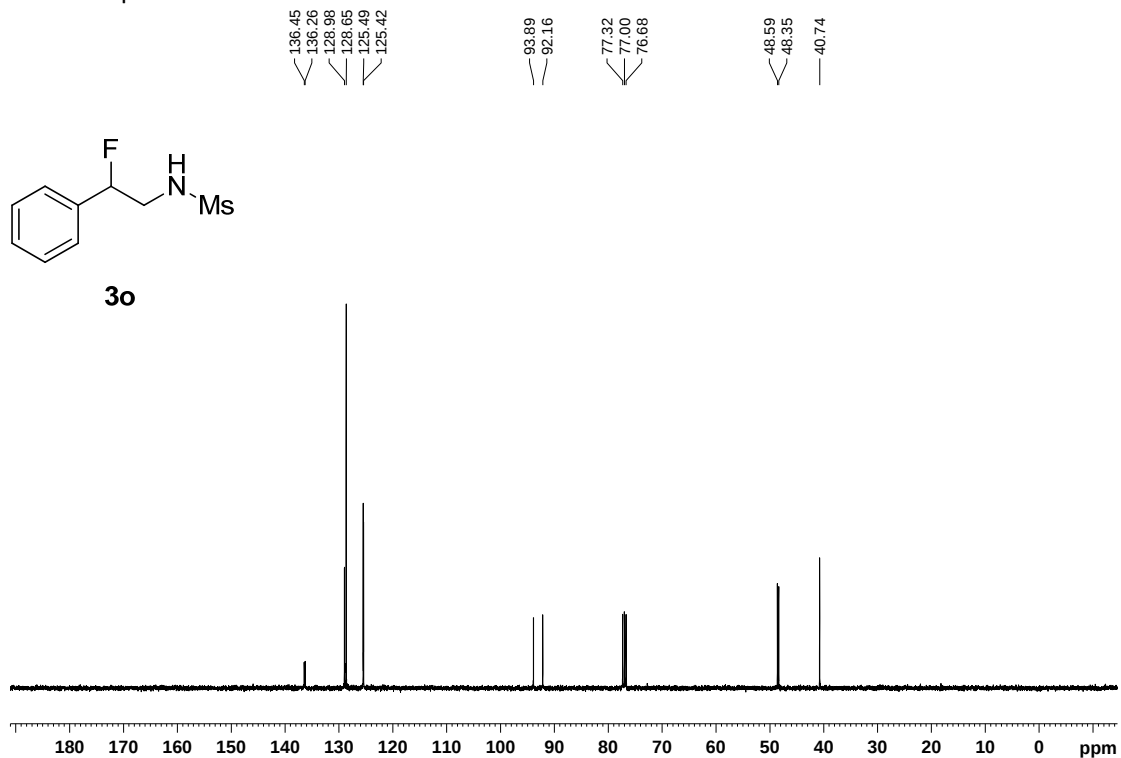
¹³F NMR spectrum of **3n**



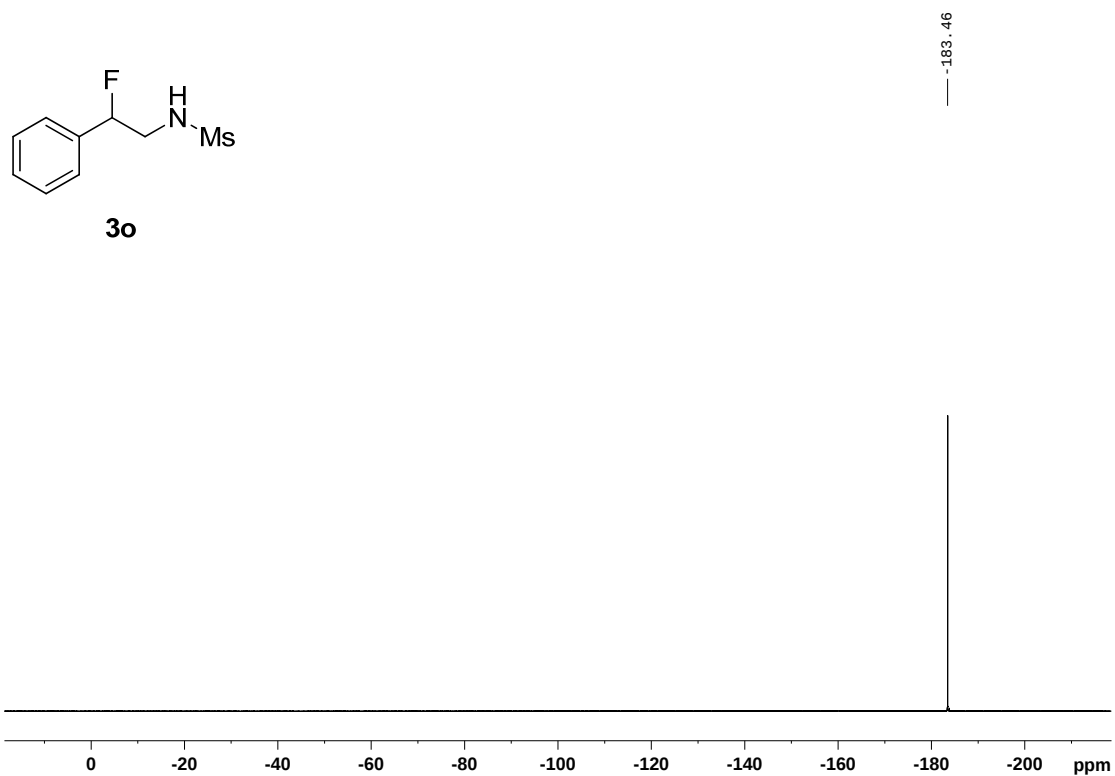
¹H NMR spectrum of **3o**



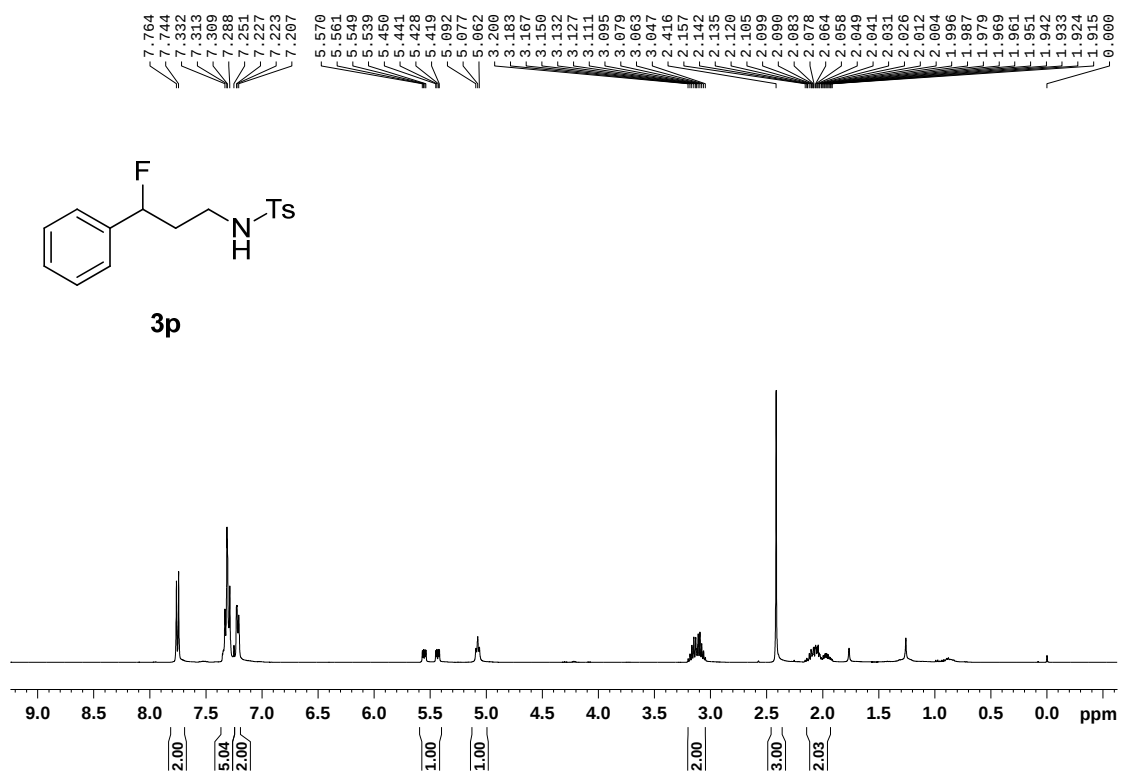
¹³C NMR spectrum of **3o**



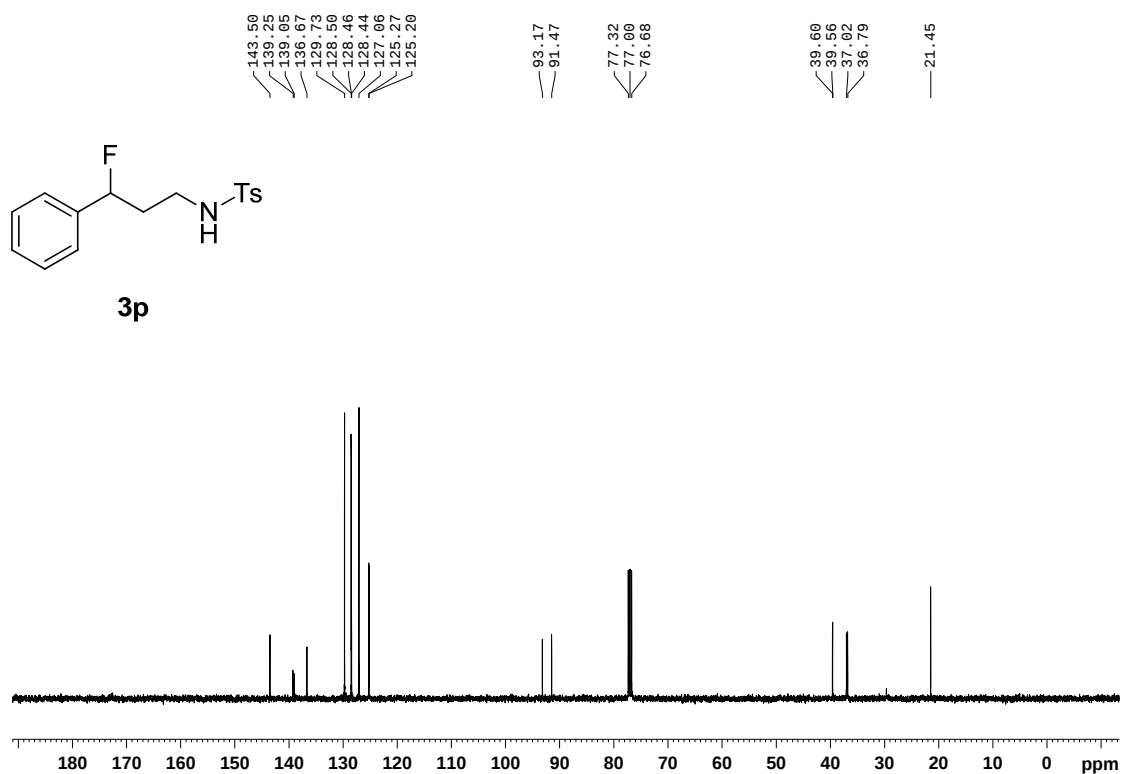
^{19}F NMR spectrum of **3o**



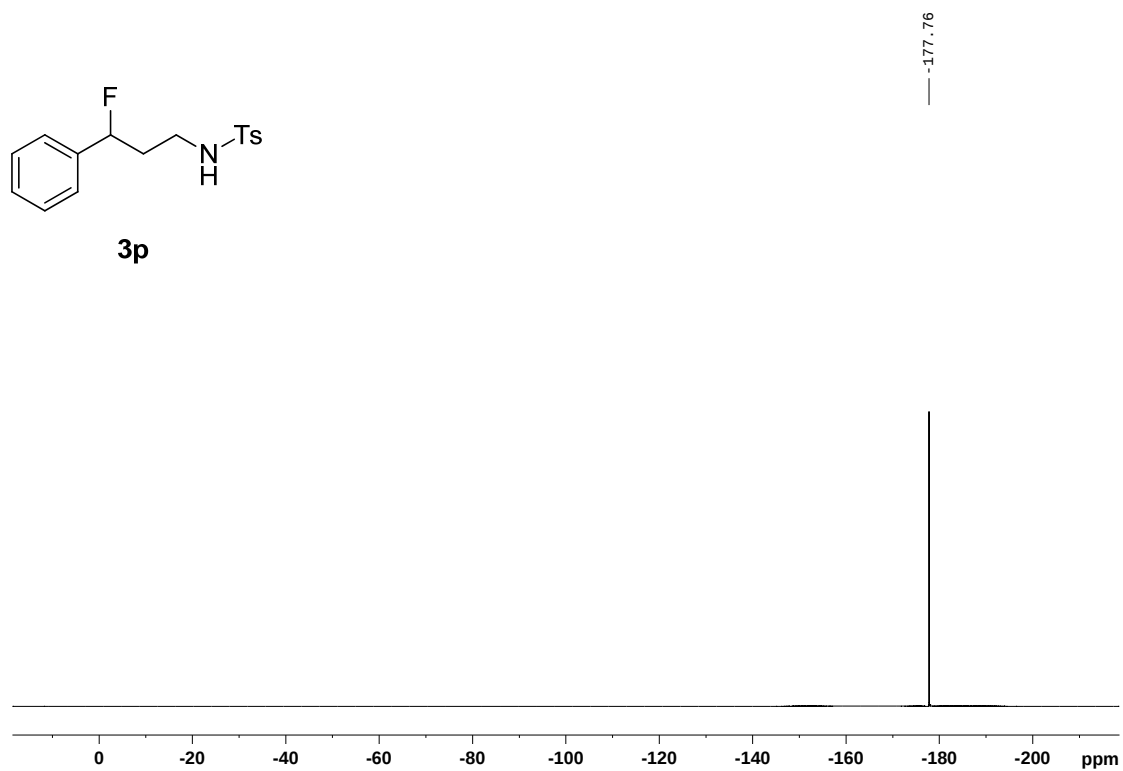
^1H NMR spectrum of **3p**



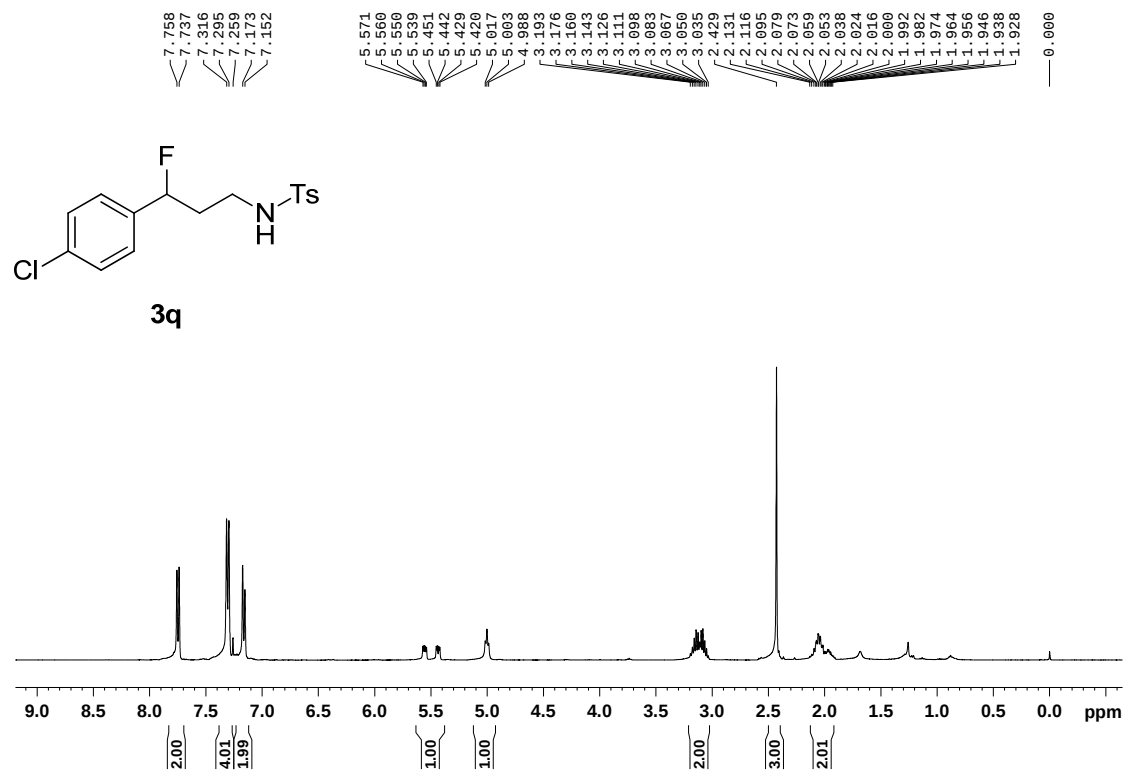
¹³C NMR spectrum of **3p**



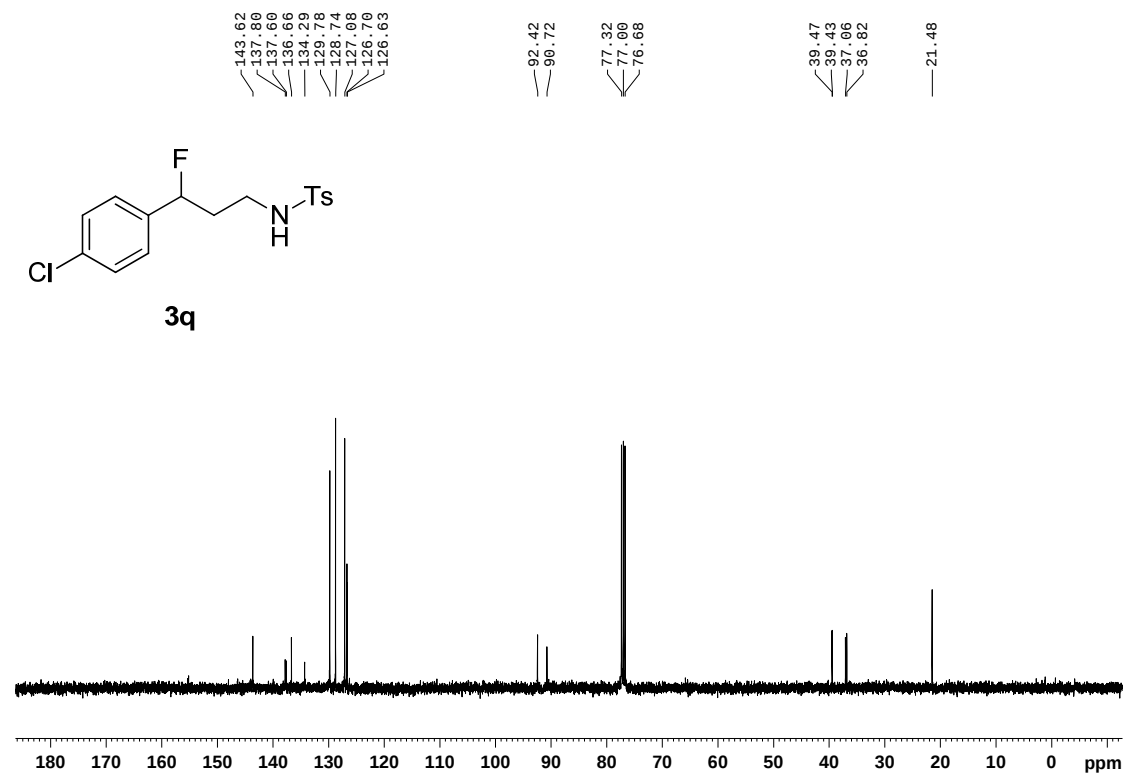
¹⁹F NMR spectrum of **3p**



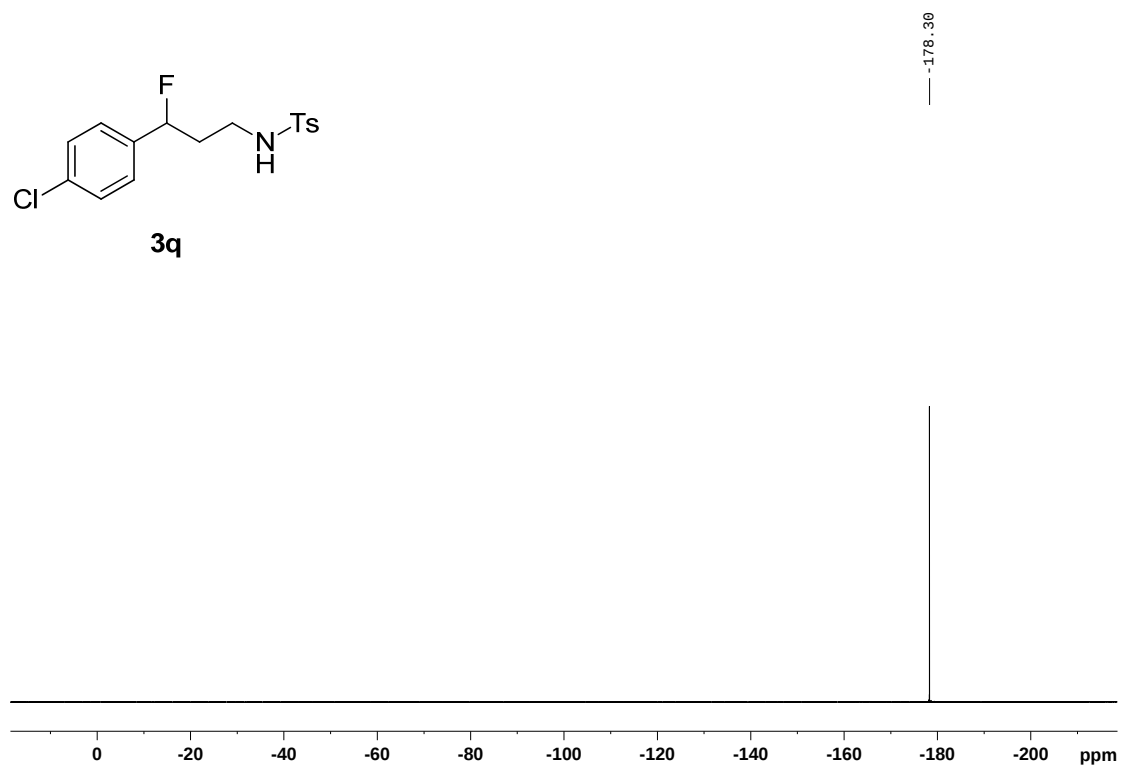
¹H NMR spectrum of **3q**



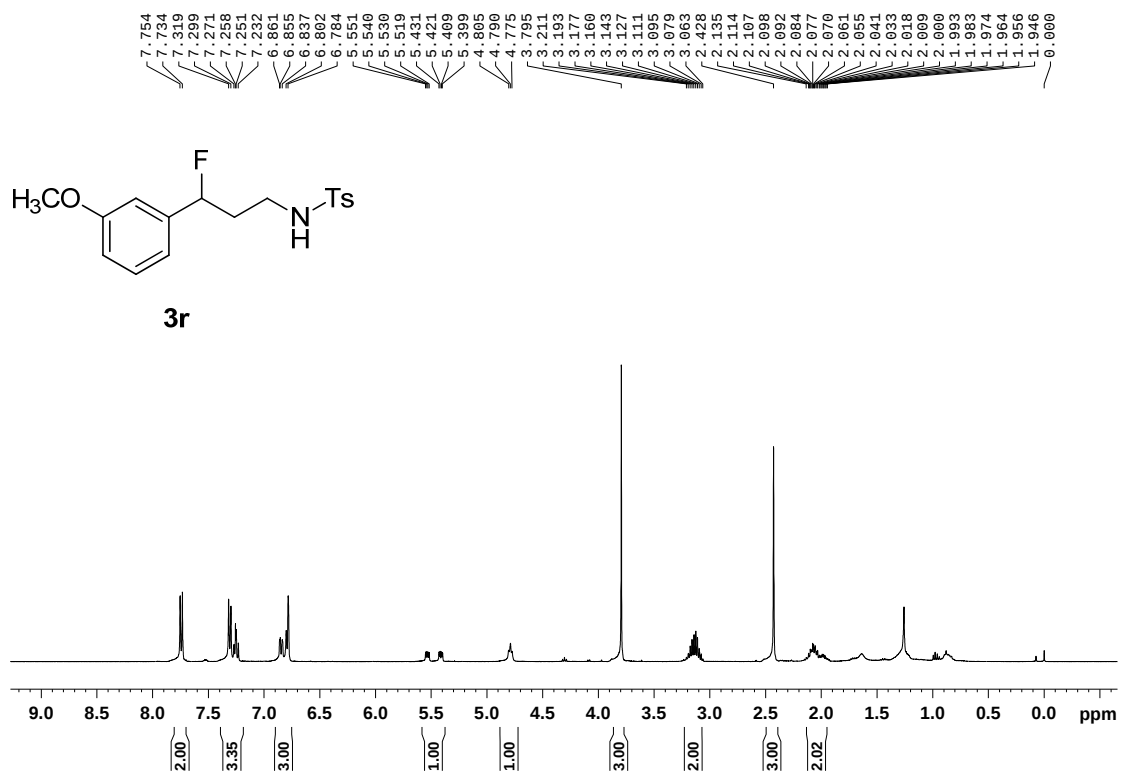
¹³C NMR spectrum of **3q**



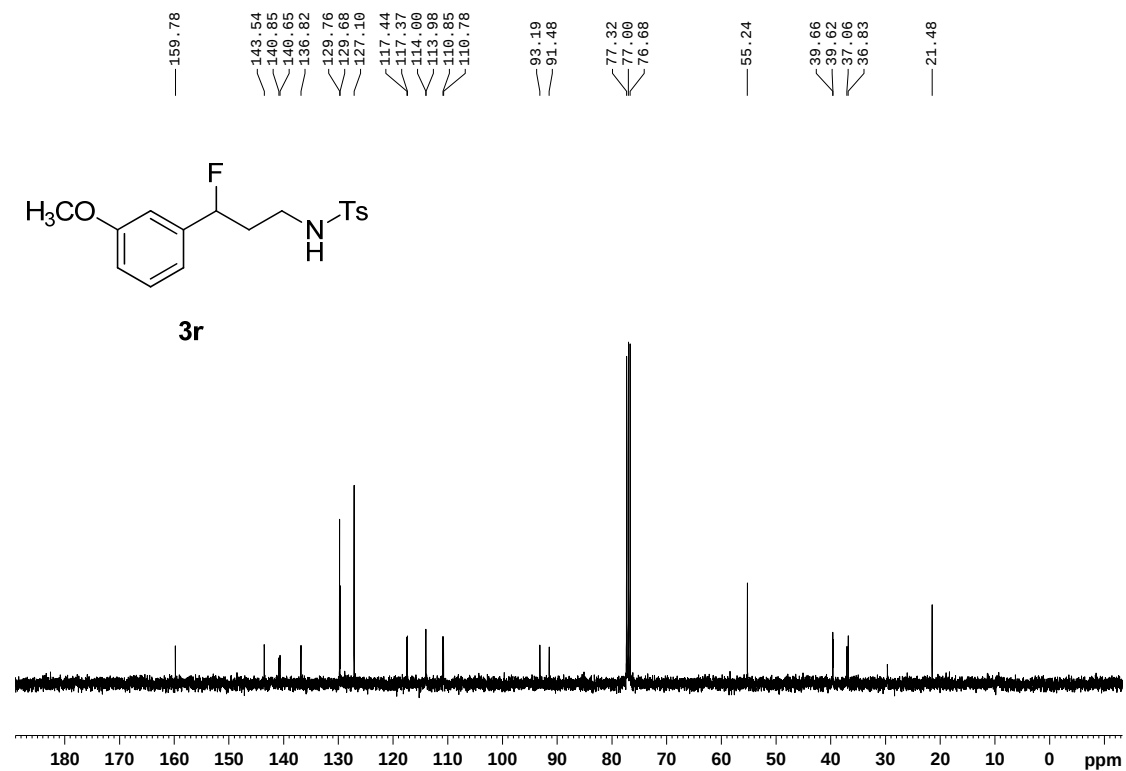
^{19}F NMR spectrum of **3q**



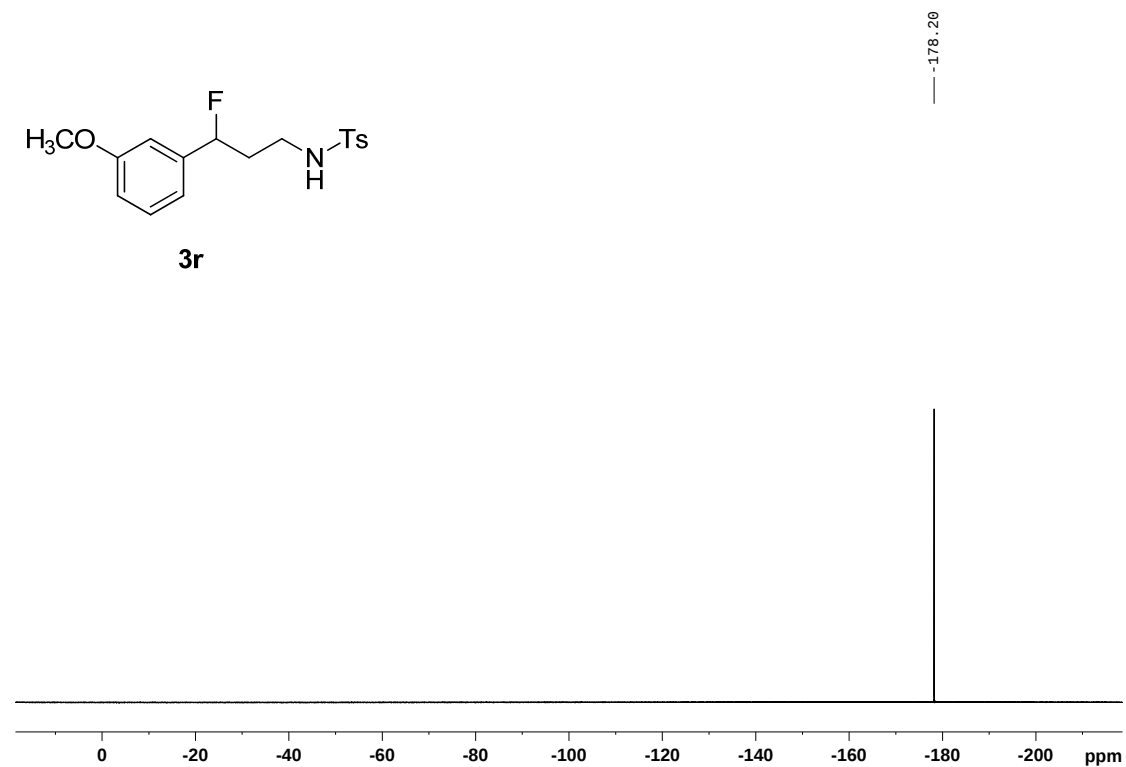
^1H NMR spectrum of **3r**



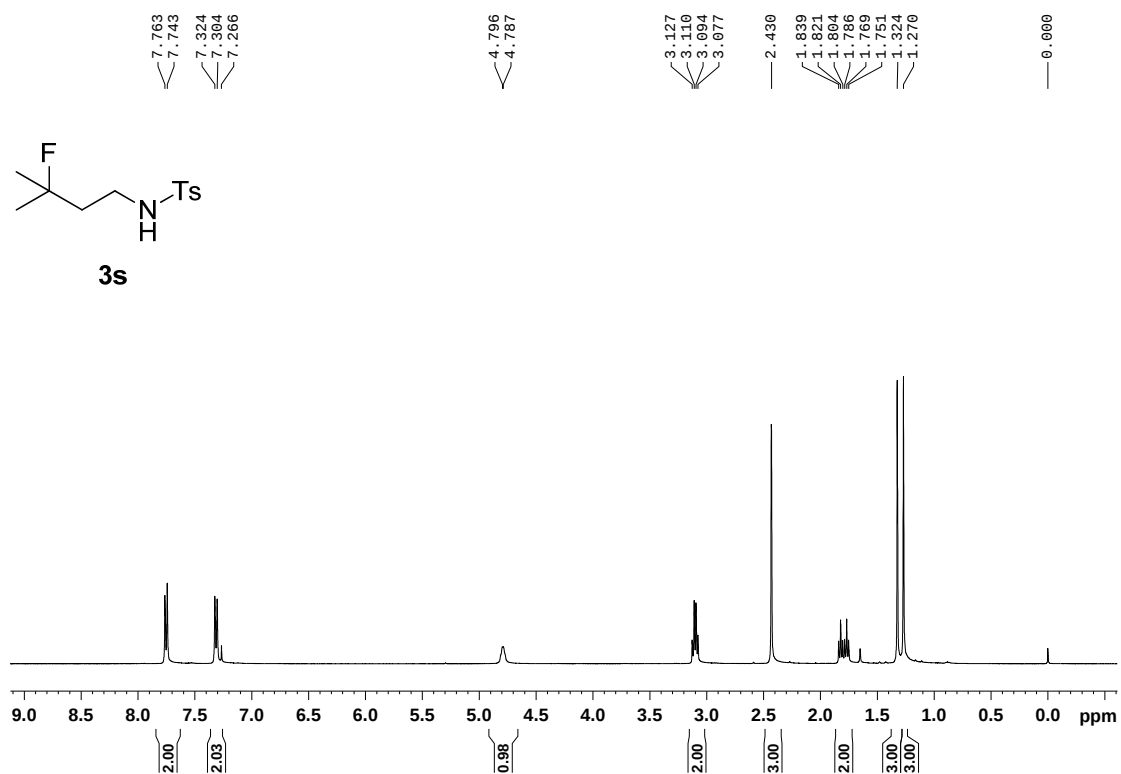
¹³C NMR spectrum of **3r**



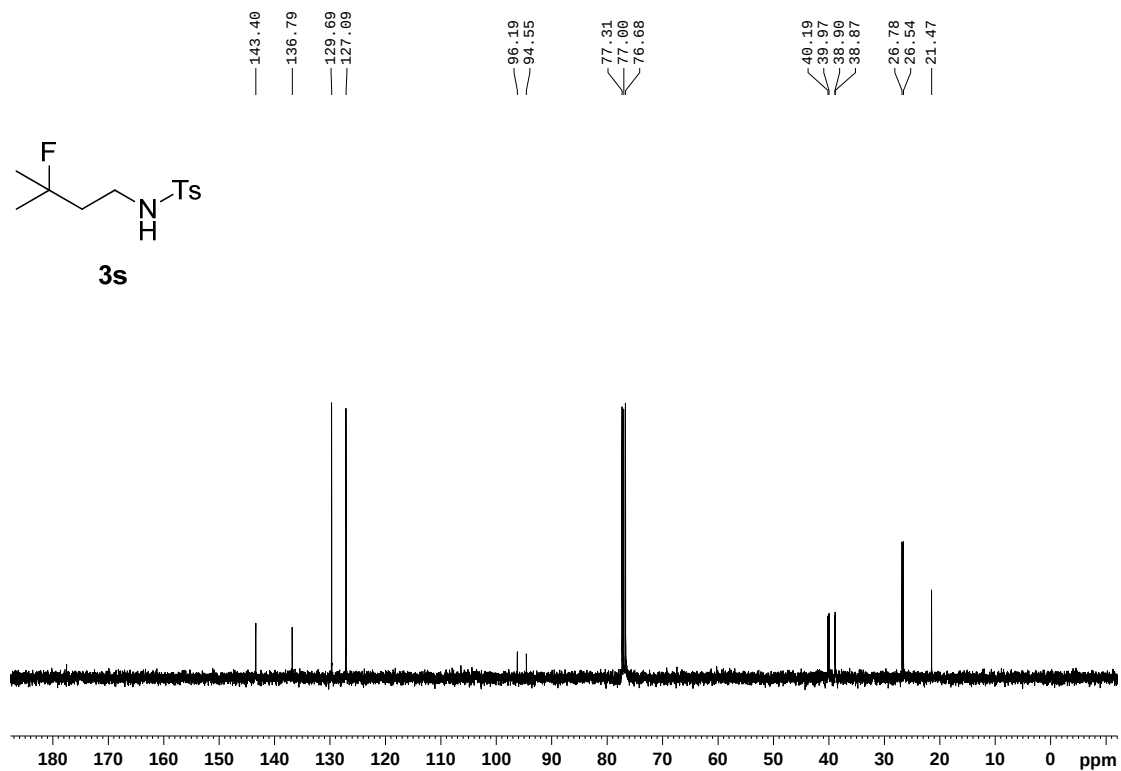
¹⁹F NMR spectrum of **3r**



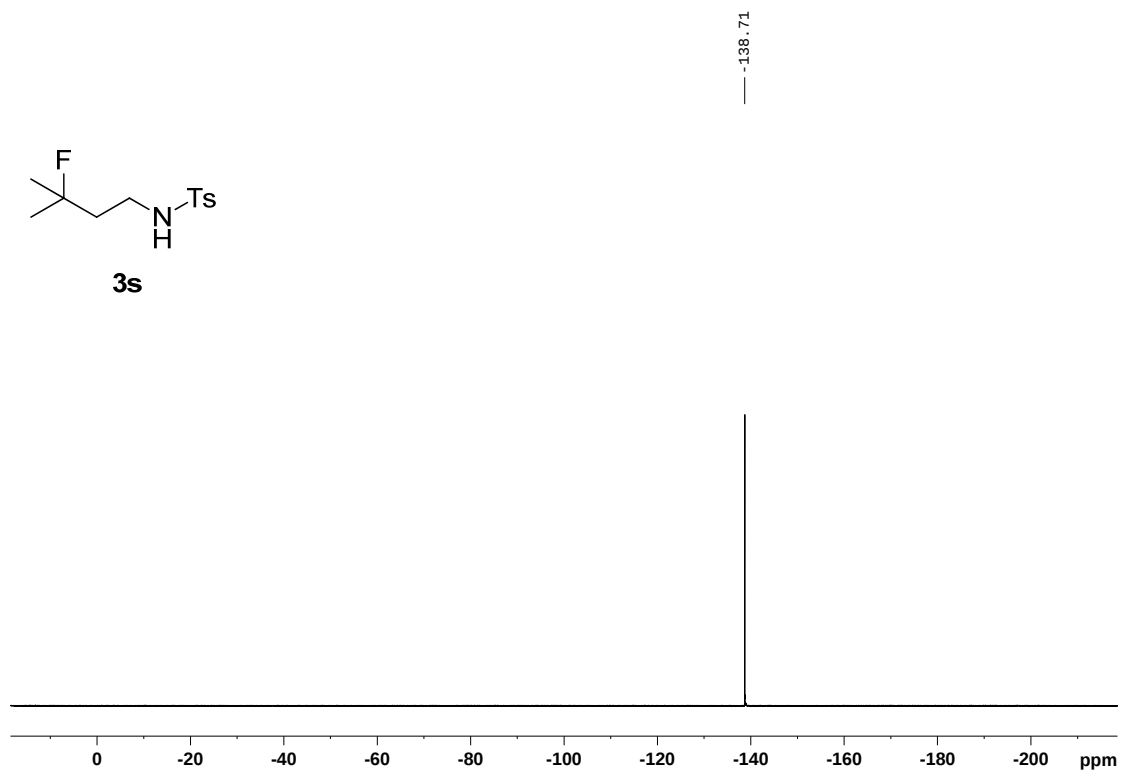
¹H NMR spectrum of **3s**



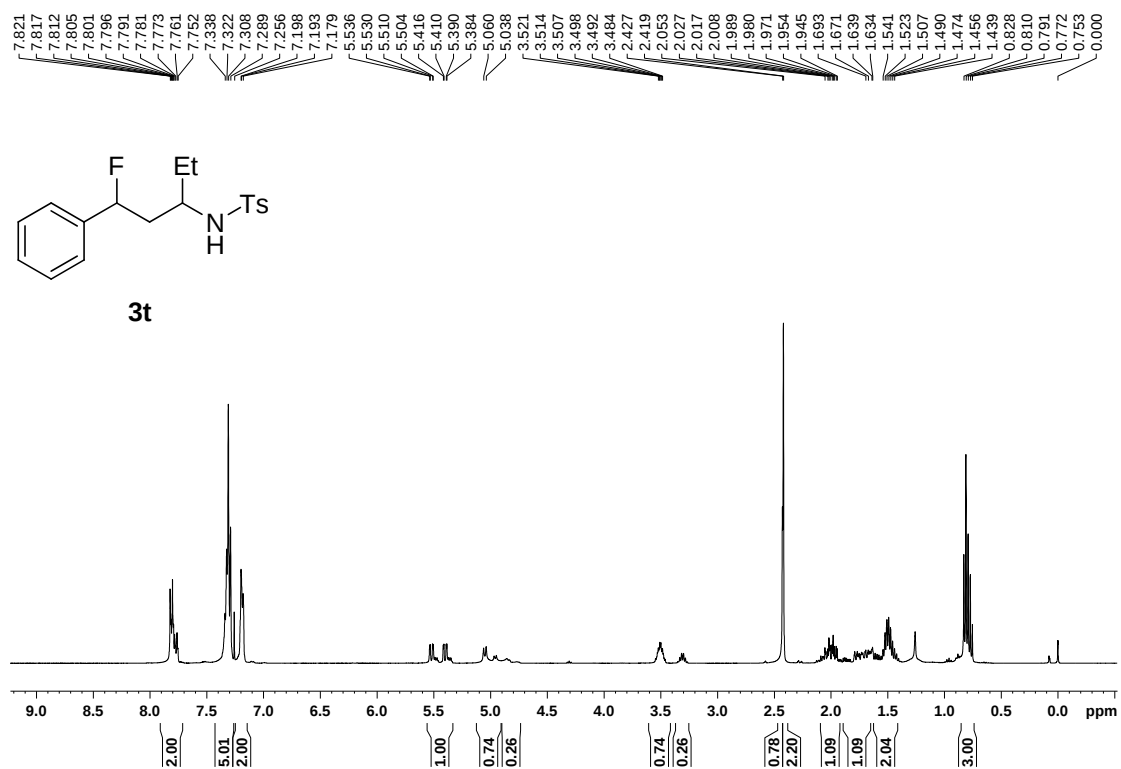
¹³C NMR spectrum of **3s**



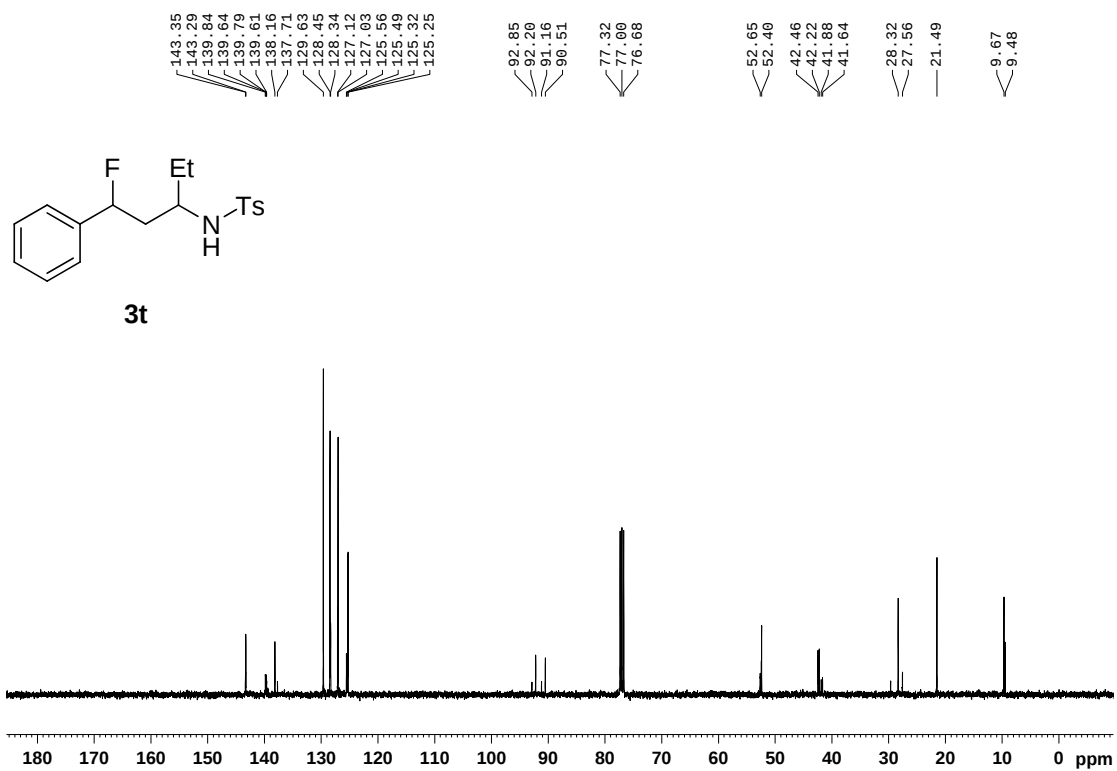
¹⁹F NMR spectrum of **3s**



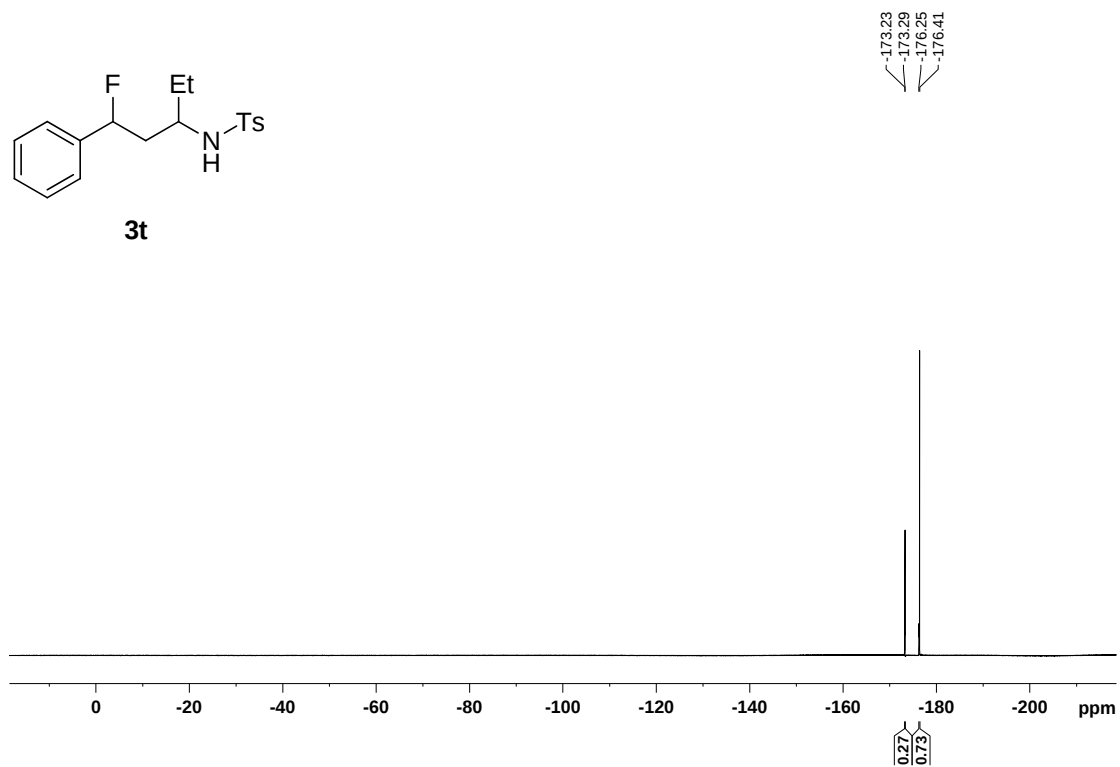
¹H NMR spectrum of **3t**



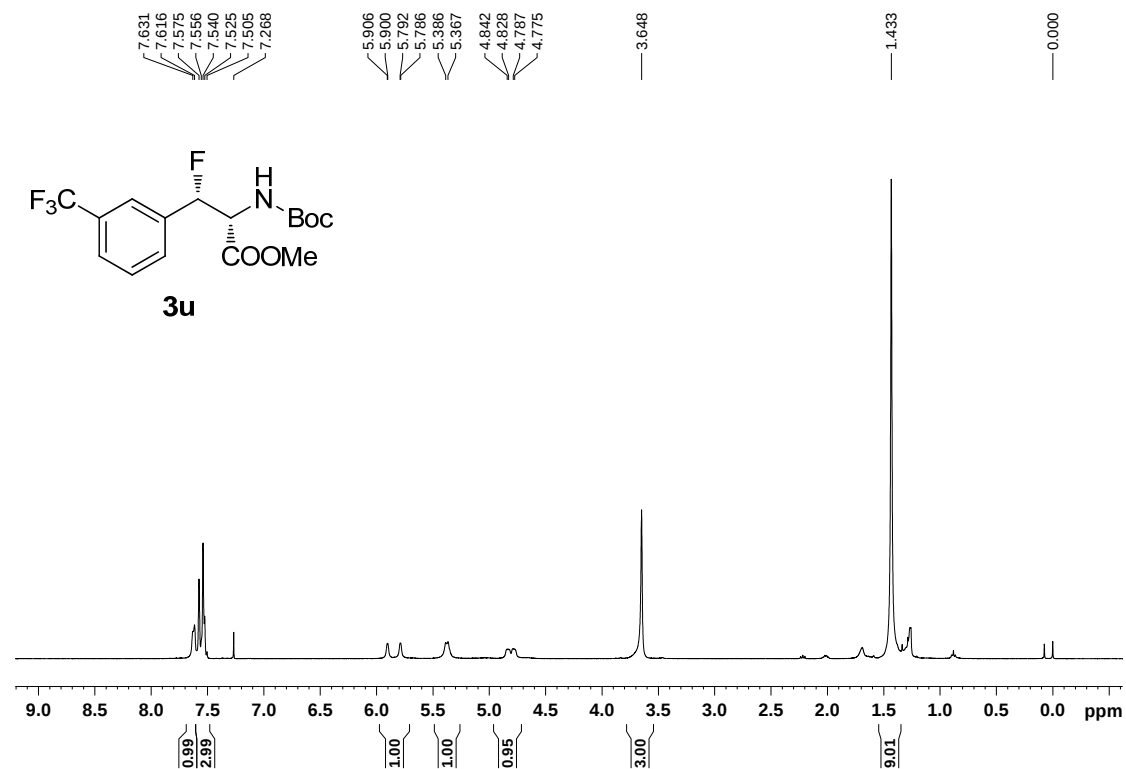
¹³C NMR spectrum of **3t**



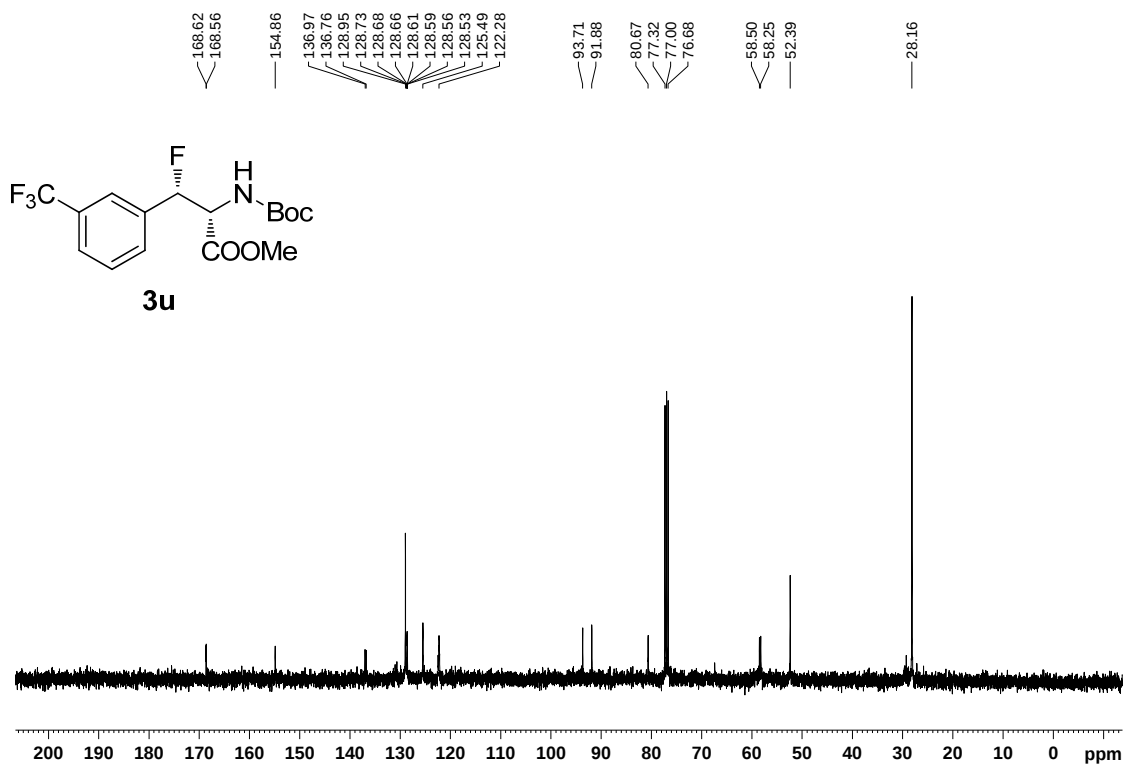
¹⁹F NMR spectrum of **3t**



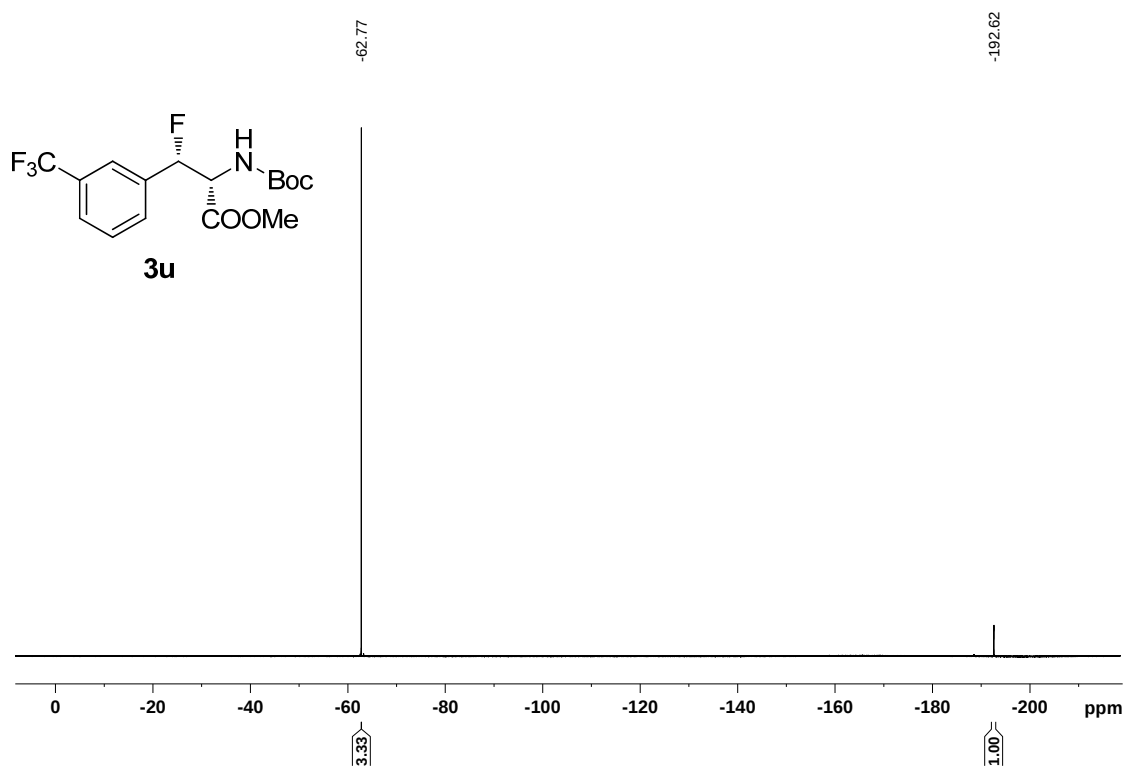
¹H NMR spectrum of **3u**



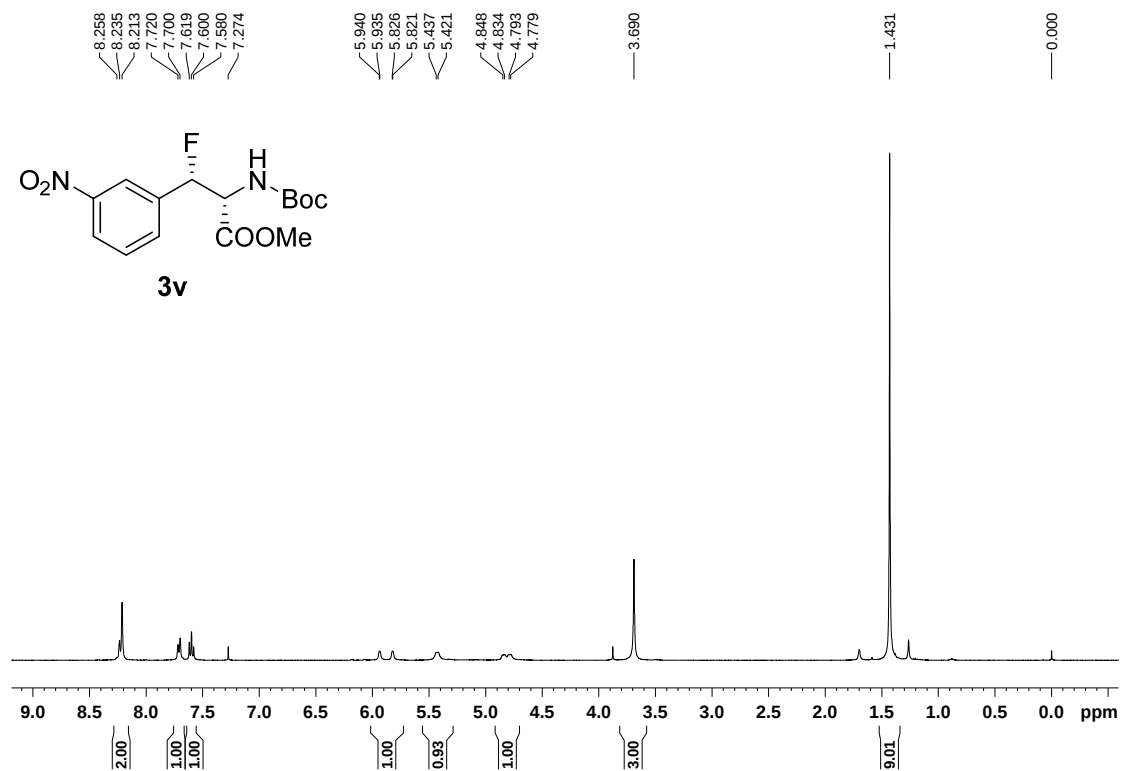
¹³C NMR spectrum of **3u**



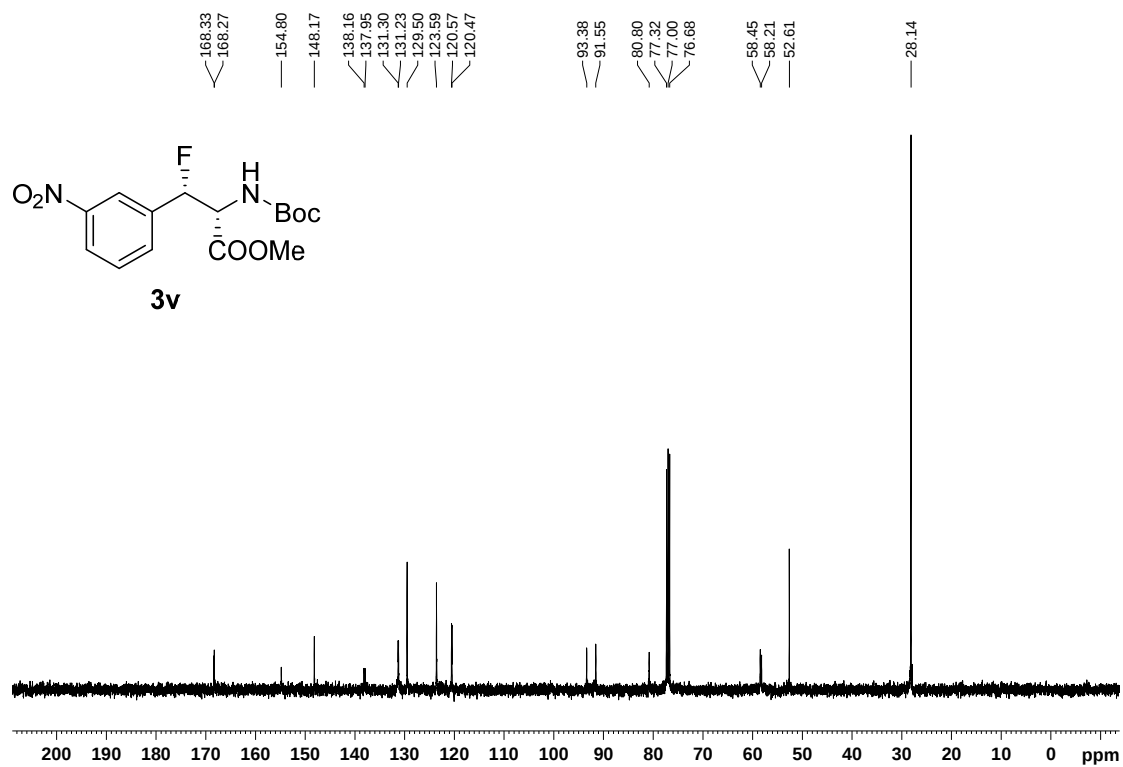
^{19}F NMR spectrum of **3u**



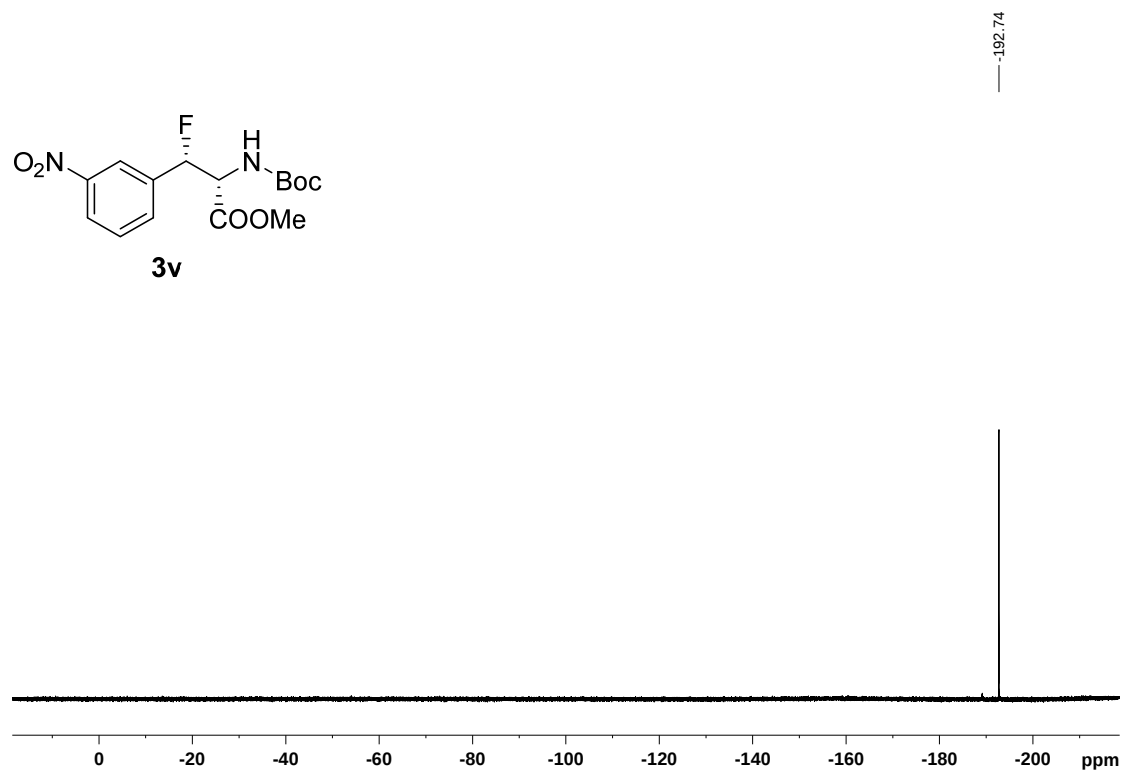
^1H NMR spectrum of **3v**



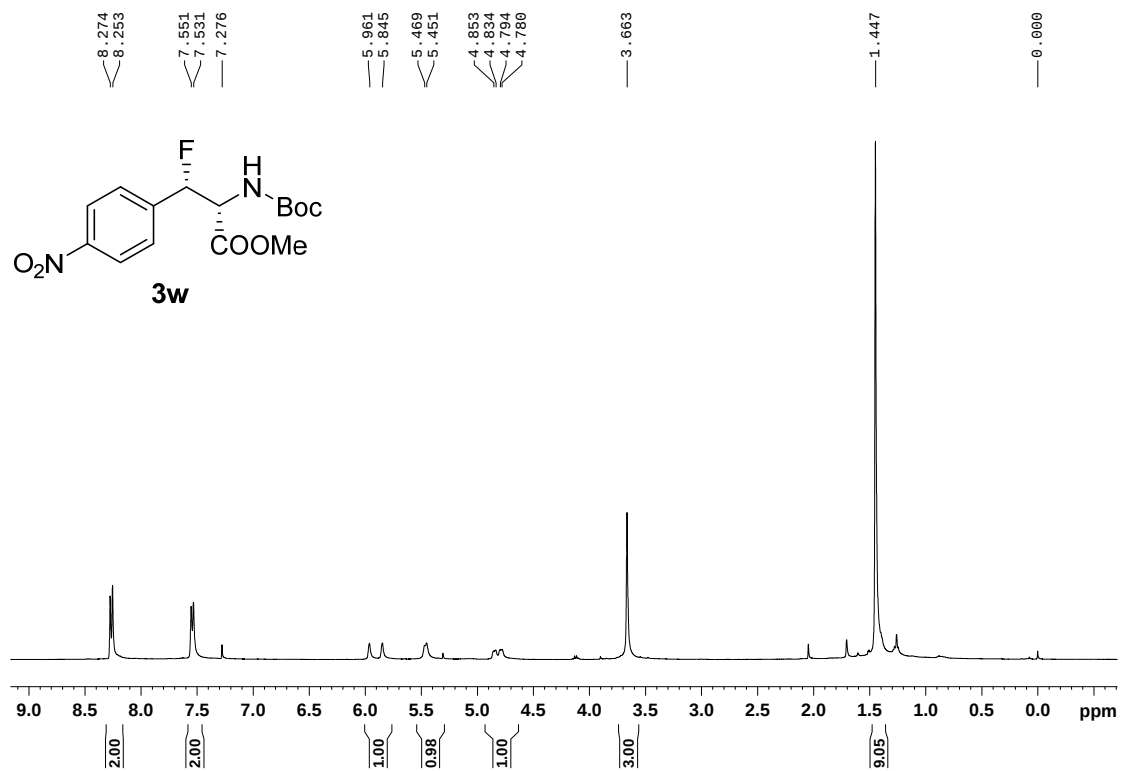
¹³C NMR spectrum of **3v**



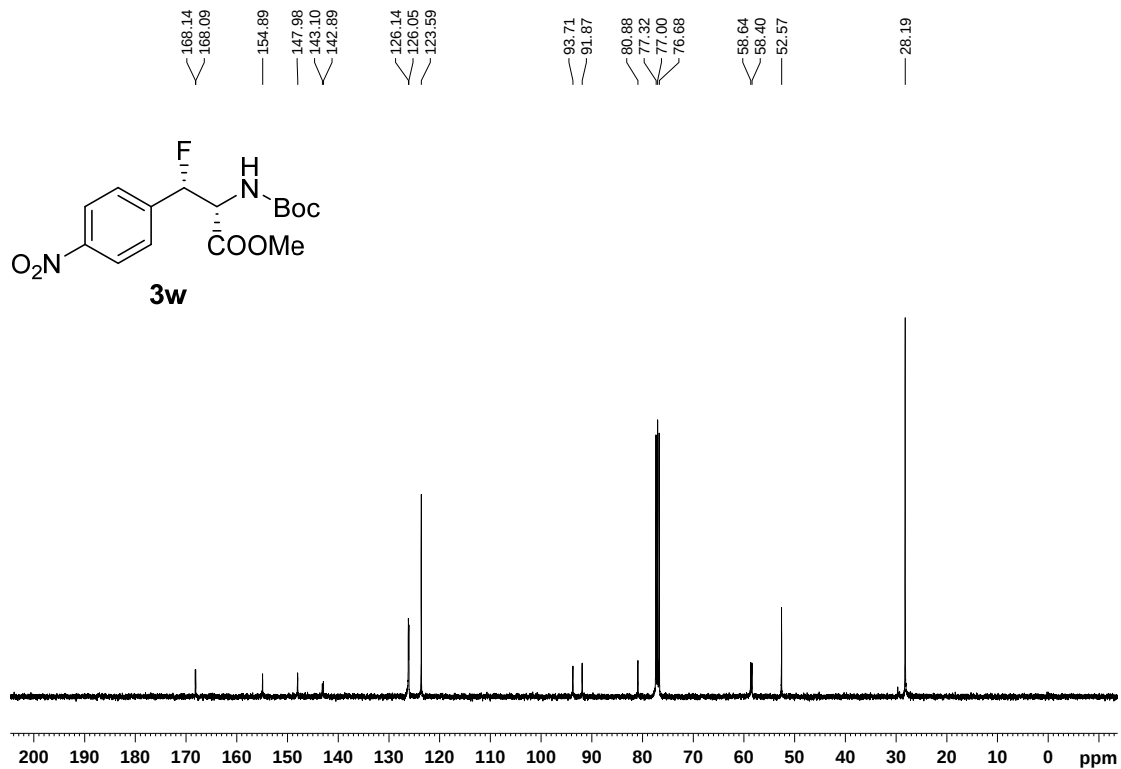
¹⁹F NMR spectrum of **3v**



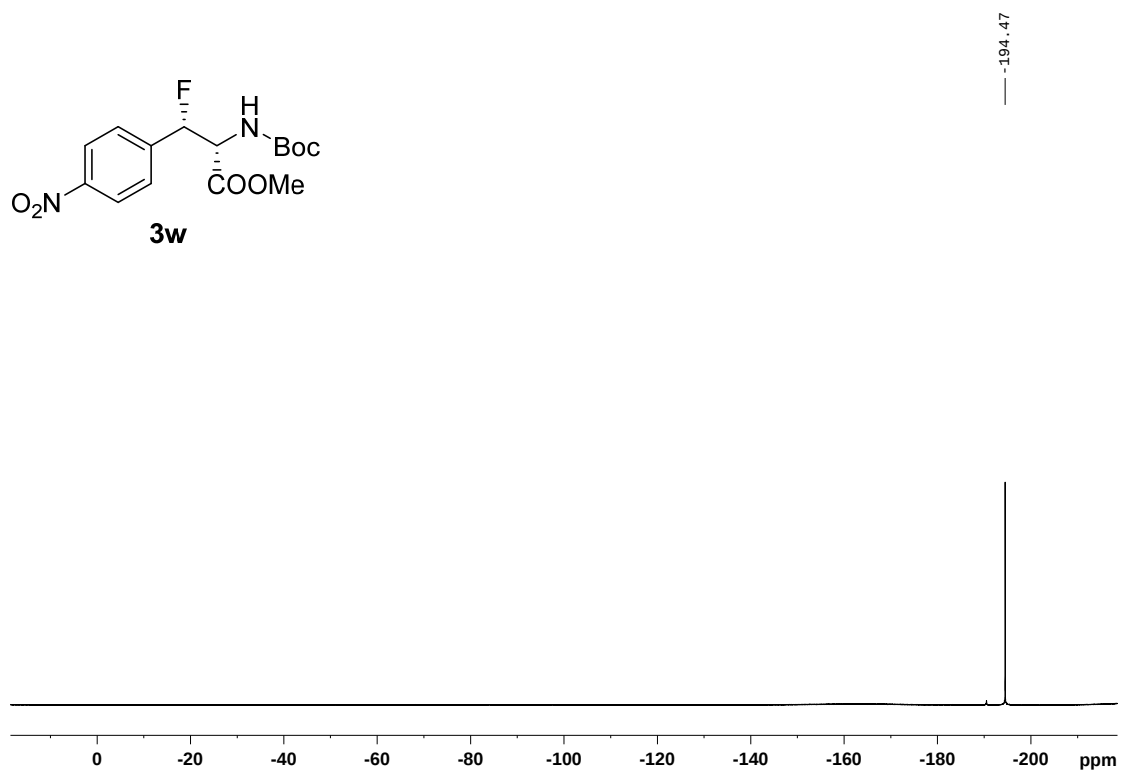
¹H NMR spectrum of **3w**



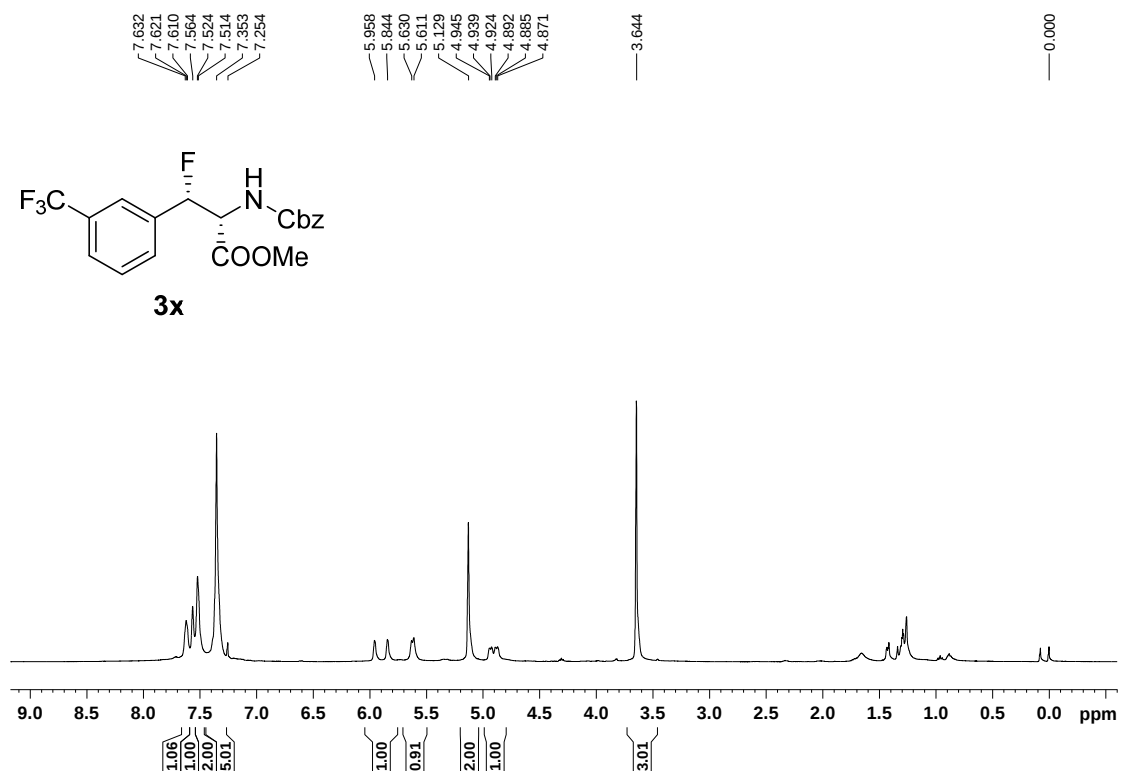
¹³C NMR spectrum of **3w**



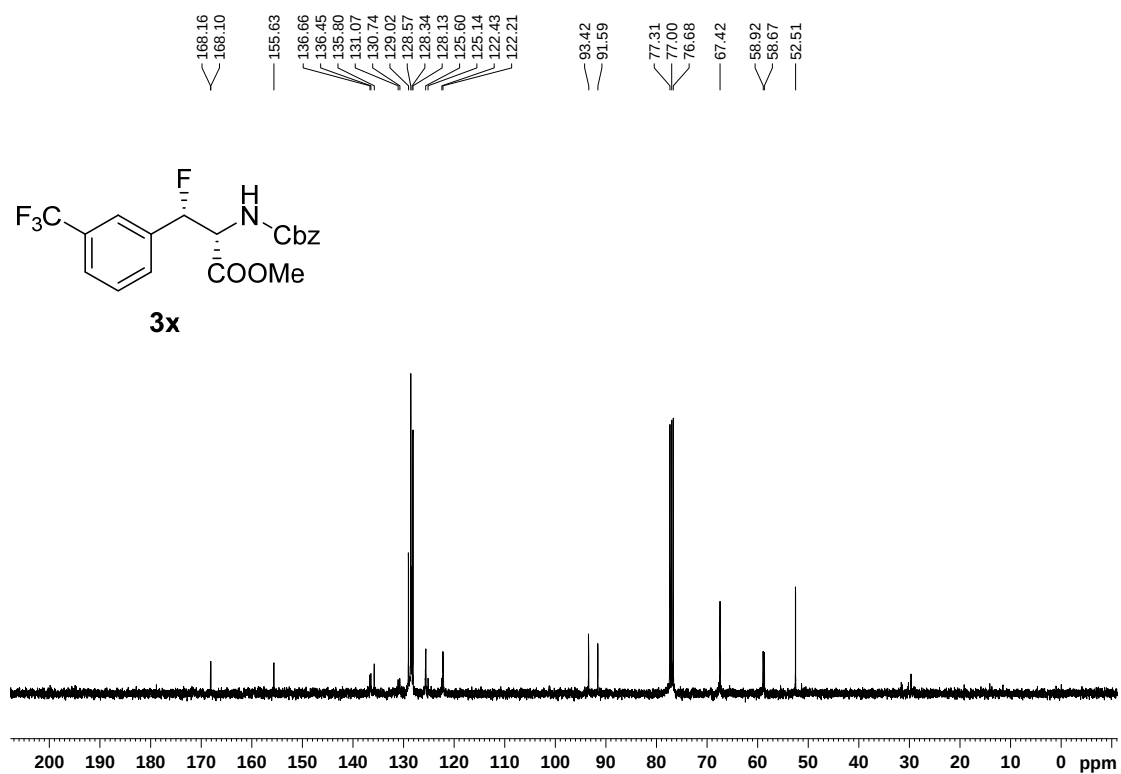
^{19}F NMR spectrum of **3w**



^1H NMR spectrum of **3x**



¹³C NMR spectrum of **3x**



¹⁹F NMR spectrum of **3x**

