

ELECTRONIC SUPPLEMENTARY INFORMATION (ESI)

Synthesis and Applications of Secondary Amine Derivatives of (+)-Dehydroabietylamine as Chiral Molecular Recognition

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1 NMR MEASUREMENT AND PROCESSING PARAMETERS

Compound characterisation

All NMR experiments were performed using Varian UNITY INOVA 500 and Varian Mercury Plus 300 instruments at 27 °C. ¹H NMR spectra were recorded with 6-16 transients, 8000-6712.5 Hz spectral width, and 1.9 s acquisition time at 500 MHz. ¹³C NMR spectra were recorded with 1876-532 transients, 31446-20000 Hz spectral width and 1-1.9 s acquisition time at 125 or 75 MHz. All 2D HSQC spectra were recorded using Varian UNITY INOVA 500 with 4 transients, 128 increments, 8000-6712.5 Hz spectral width in ¹H-dimension, 25133 Hz spectral width in ¹³C-dimension, 1.0 s relaxation delay, and 1.3 s acquisition time.

Resolution experiments

All NMR measurements were recorded using Varian UNITY INOVA 500 at 27 °C. ¹H NMR spectra were recorded with 16 transients, 8000 Hz spectral width, 5.0 s relaxation delay and 1.9 s acquisition time at 500 MHz. ¹⁹F NMR spectra were recorded with 16-32 transients, 19047 Hz spectral width, 5.0 s relaxation delay and 1.0 s acquisition time at 470 MHz. HMBC spectra were recorded using Varian UNITY INOVA 500 with 4-6 transients, 400 increments, 1.0 s relaxation delay and 0.128 s acquisition time. The spectral widths were 4247.2-8000 Hz and 30165.9-25133.5 Hz in ¹H- and ¹³C-dimensions, respectively.

DOSY Experiment (see 3.1.1)

For DOSY experiment, Bipolar Pulse Pair Simulated Echo (BPPSTE)¹ pulse sequence was used. To measure DOSY dataset, 25 different diffusion gradient amplitudes were used (range 0.5-20.0 G/cm). Gradient pulse shape was rectangular. Diffusion gradient duration was 1.0 ms, eddy current recovery delay was 0.2 ms, and diffusion time was 250.0 ms. The DOSY dataset was measured using 16 steady state scans, 4 transients, and 2.0 s relaxation delay. Spectral width in ¹H-dimension was 6982.63 Hz and the number of acquired complex data points was 7000. The data was apodized by Gaussian weighting function and zero-filled up to 16384 complex points prior to Fourier transformation. Finally, two-dimensional DOSY spectra was calculated using DOSY-macro (a monoexponential fit) incorporated in Varian VNMR 6.1C spectrometer operating software.

1 M. D. Pelta, H. Barjat, G. A. Morris, A. L. Davis, S. J. Hammond, *Magn. Reson. Chem.*, **1998**, 36, 10, 706-714

Dynamic NMR studies (see 3.1.1)

All NMR measurements were recorded using Varian UNITY INOVA 500. ¹H NMR spectra were recorded with 16-64 transients, 8000 Hz spectral width, and 1.9 s acquisition time at temperature range -65 to +50 °C.

2 CHIRAL RECOGNITION STUDIES

Solutions of **7** and **8** (22.0 mM in CDCl_3) were prepared. Compound **1-6 ab** (1.0 eq or 2.0 eq) were dissolved in 0.5 mL (1.0 eq) of prepared solution of guest and both ^1H and ^{19}F NMR spectra were recorded.

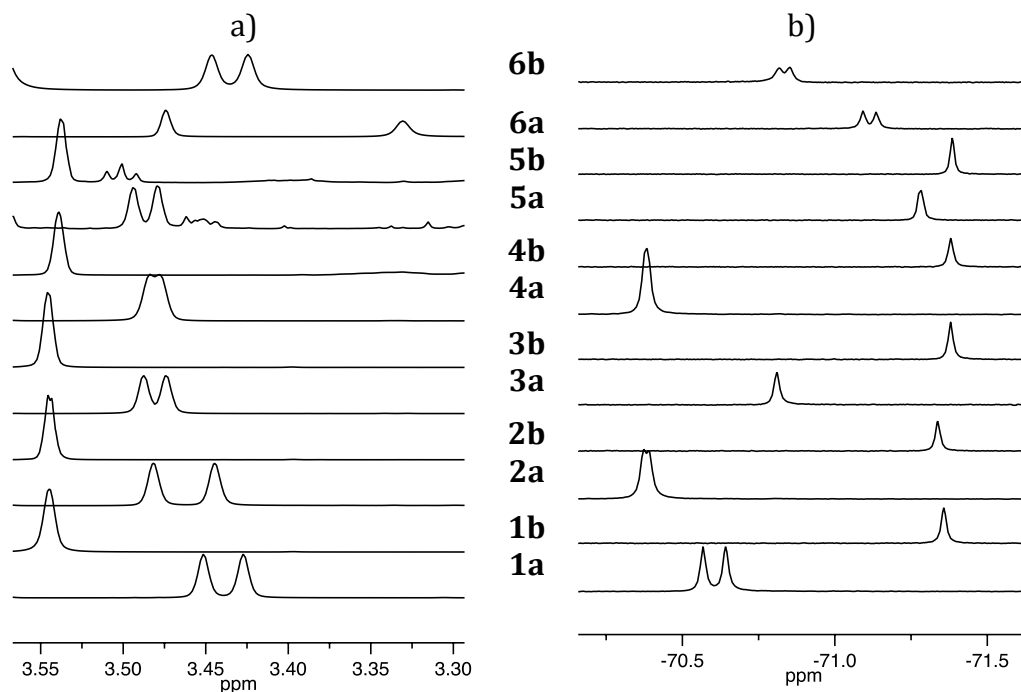


Figure 1. Chiral recognition of enantiomers of **7**. Host:guest molar ratio 1:1 a) ^1H NMR (OCH_3 peak) and b) ^{19}F NMR (CF_3 peak)

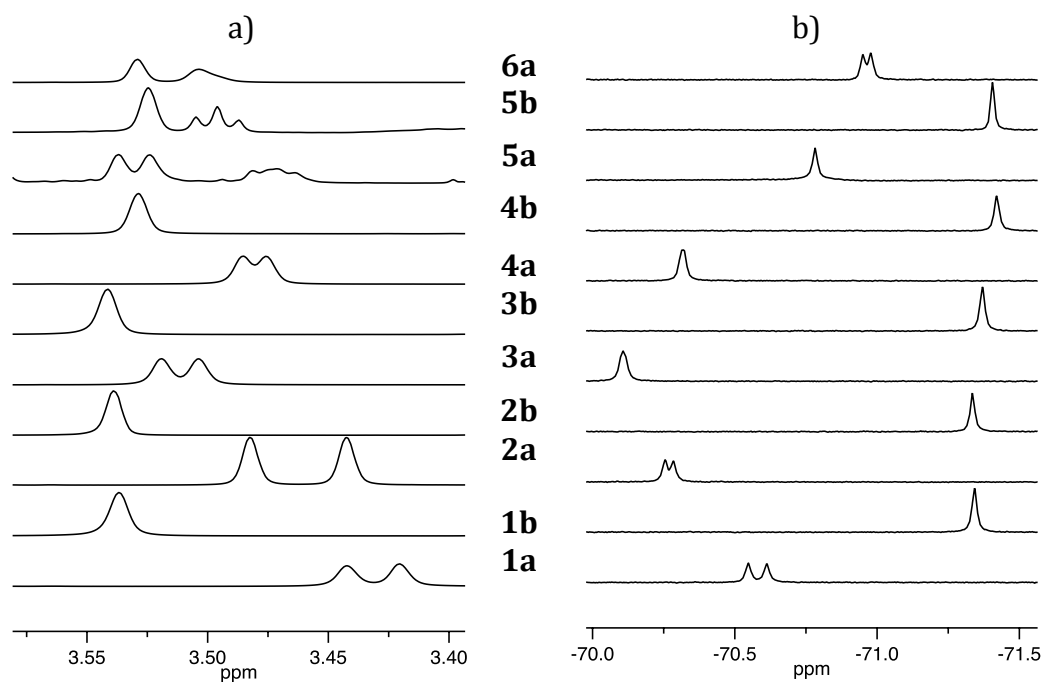


Figure 2 Chiral recognition of enantiomers of **7**. Host:guest molar ratio 2:1 a) ^1H NMR (OCH_3 peak) and b) ^{19}F NMR (CF_3 peak)

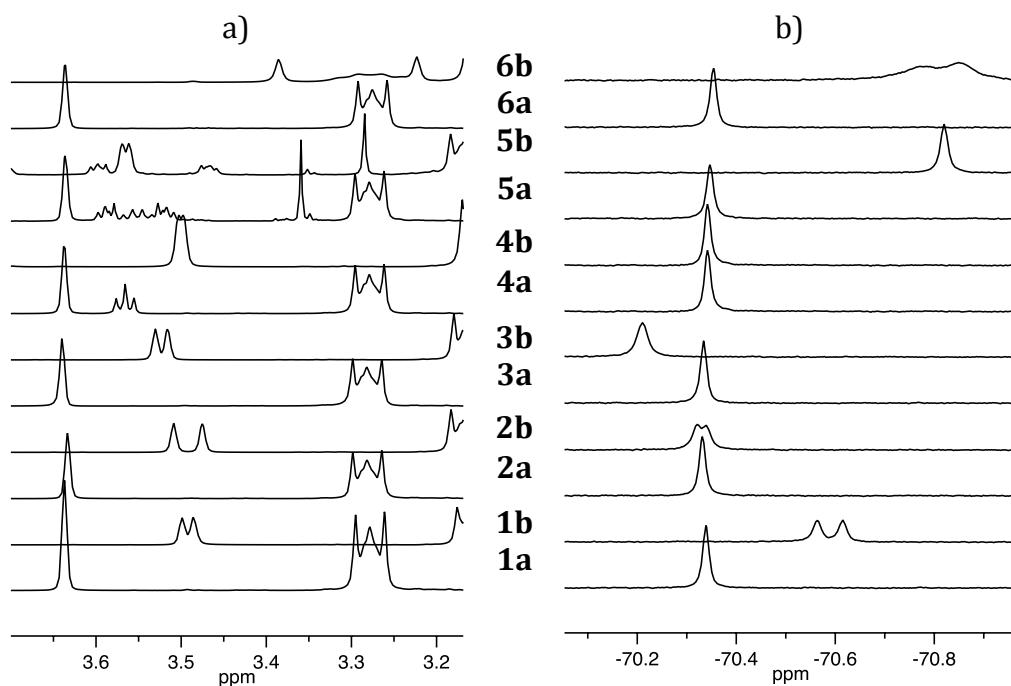


Figure 3. Chiral recognition of enantiomers of **8**. Host:guest molar ratio 1:1 a) ^1H NMR (OCH_3 peak) and b) ^{19}F NMR (CF_3 peak)

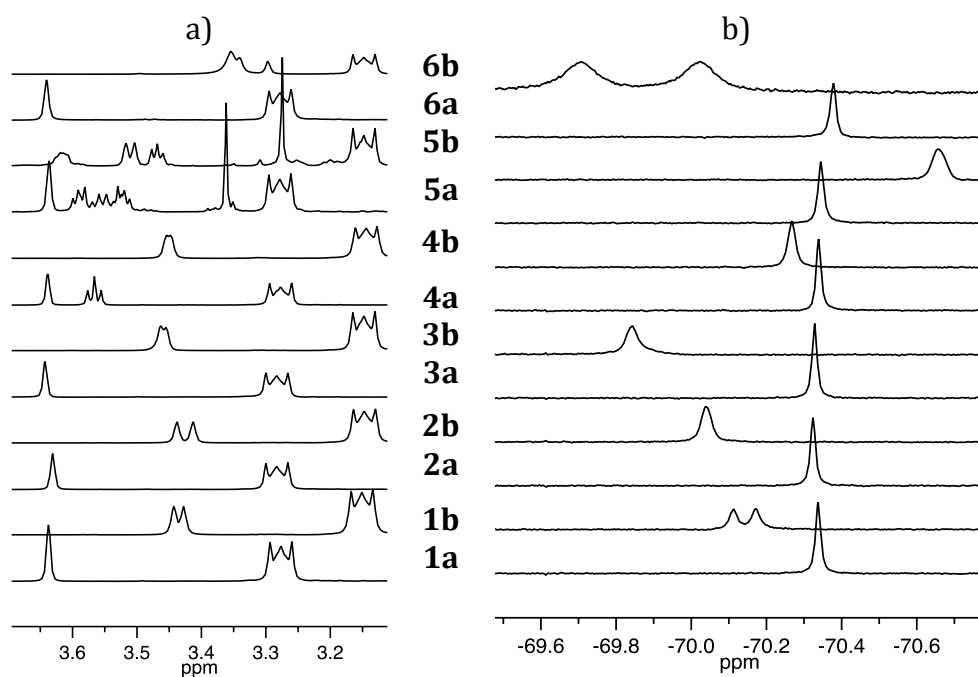


Figure 4. Chiral recognition of enantiomers of **8**. Host:guest molar ratio 2:1 a) ^1H NMR (OCH_3 peak) and b) ^{19}F NMR (CF_3 peak)

HMBC spectra from sample containing **6a** and **7**, as well as from sample containing **6b** and **8** were recorded. From spectra was noticed that **6a** and **6b** are able to resolve the chiral carbon of the used model compound (Fig 5).

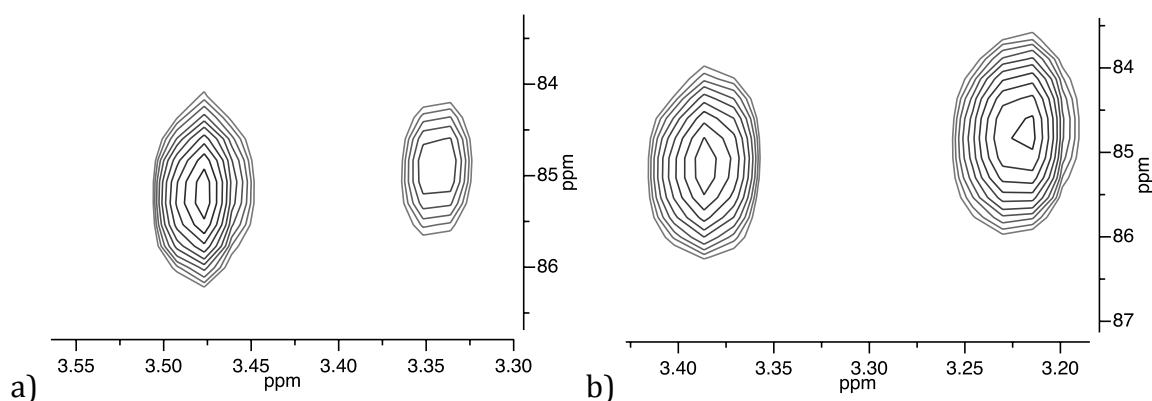


Figure 5. Separation of chiral carbon of *S* and *R* enantiomers (in HMBC spectra) of a) **7** with **6a** ($\Delta\delta$ 0.28 ppm, 35.3 Hz) and b) **8** with **6b** ($\Delta\delta$ 0.42 ppm, 52.5 Hz) in guest:host ratio 1:1.

3 TITRATION

3.1 Titration of **7** with **6a**

Titration solutions containing the guest, **7** (2.0 mM) and host, **6a** (46.6 mM) were prepared. In NMR tube 0.5 mL of guest solution was measured and titrated by 1.0-5.0 μ L doses of host solution. Both ^1H and ^{19}F NMR spectra were recorded. Concentration of guest is assumed to remain constant during the titration.

Table 1. Experimental data from titration of **7** with **6a**

c of 6a (mM)	V of 6a (mL)	^1H NMR				^{19}F NMR			
		δ of <i>S</i> (ppm)	δ of <i>R</i> (ppm)	$\Delta\delta_S$ of <i>S</i> (ppm)	$\Delta\delta_R$ of <i>R</i> (ppm)	δ of <i>S</i> (ppm)	δ of <i>R</i> (ppm)	$\Delta\delta_S$ of <i>S</i> (ppm)	$\Delta\delta_R$ of <i>R</i> (ppm)
0	0	3.5335	3.5335	0	0	-71.053	-71.053	0	0
0.093013972	0.001	3.495	3.521	0.0385	0.0125	-71.074	-71.074	0.021	0.021
0.231840796	0.0025	3.443	3.502	0.0905	0.0315	-71.09	-71.09	0.037	0.037
0.461386139	0.005	3.362	3.473	0.1715	0.0605	-71.124	-71.124	0.071	0.071
0.688669951	0.0075	3.289	3.446	0.2445	0.0875	-71.135	-71.172	0.082	0.119
0.91372549	0.01	3.23	3.423	0.3035	0.1105	-71.148	-71.221	0.095	0.168
1.003131115	0.011	3.219	3.419	0.3145	0.1145	-71.147	-71.231	0.094	0.178
1.0921875	0.012	3.225	3.424	0.3085	0.1095	-71.143	-71.231	0.09	0.178
1.180896686	0.013	3.241	3.433	0.2925	0.1005	-71.137	-71.219	0.084	0.166
1.357281553	0.015	3.279	3.453	0.2545	0.0805	-71.123	-71.196	0.07	0.143
1.792307692	0.02	3.378	3.491	0.1555	0.0425	-71.095	-71.147	0.042	0.094
2.219047619	0.025	3.442	3.513	0.0915	0.0205	-71.074	-71.117	0.021	0.064
2.637735849	0.03	3.477	3.523	0.0565	0.0105	-71.061	-71.101	0.008	0.048
3.048598131	0.035	3.497	3.528	0.0365	0.0055	-71.055	-71.093	0.002	0.04
3.451851852	0.04	3.509	3.532	0.0245	0.0015	-71.05	-71.086	-0.003	0.033
3.847706422	0.045	3.518	3.533	0.0155	0.0005	-71.045	-71.083	-0.008	0.03
4.236363636	0.05	3.524	3.534	0.0095	-0.0005	-71.043	-71.08	-0.01	0.027
4.618018018	0.055	3.535	3.535	-0.0015	-0.0015	-71.043	-71.077	-0.01	0.024
4.992857143	0.06	3.535	3.535	-0.0015	-0.0015	-71.04	-71.075	-0.013	0.022

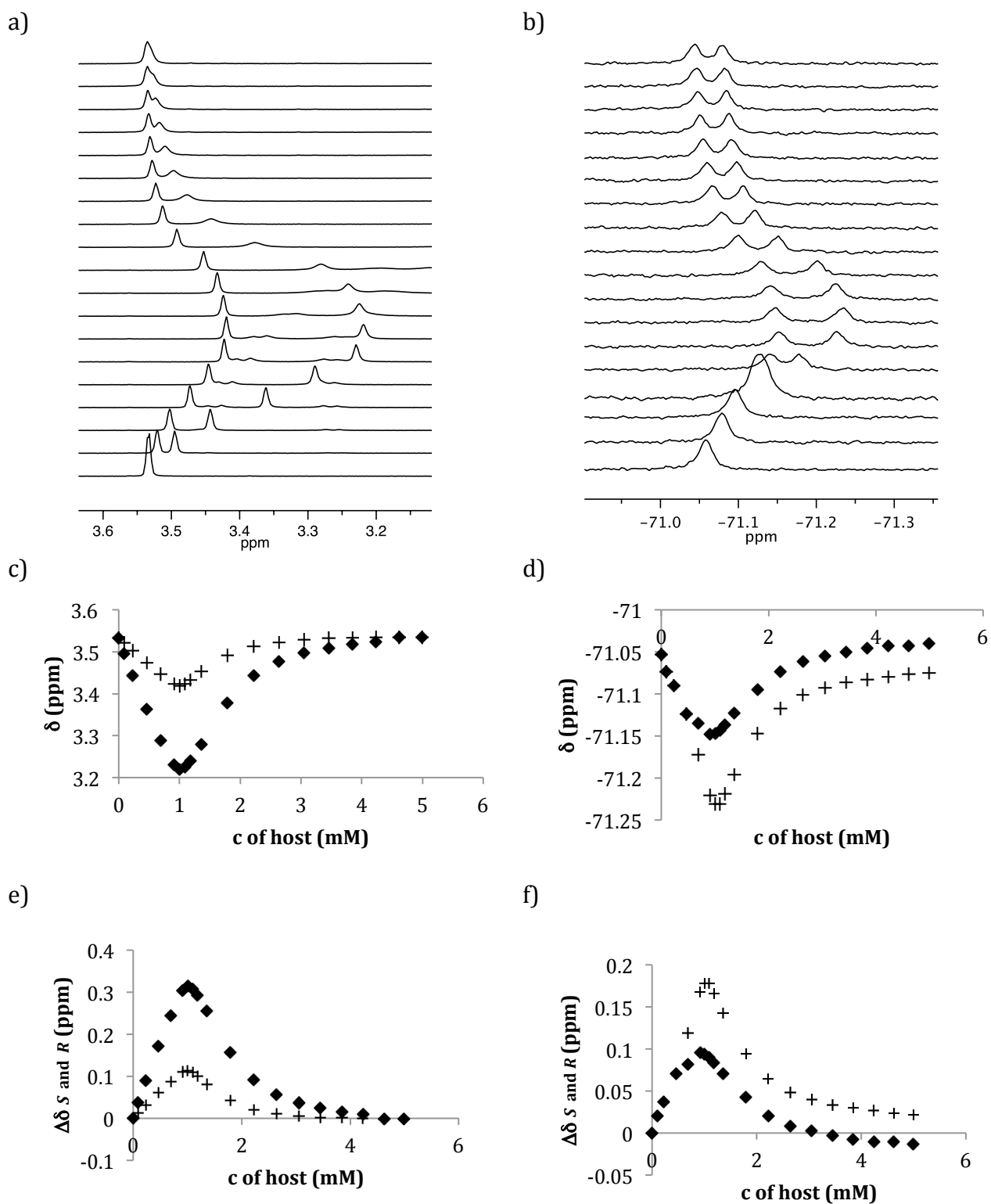


Figure 6. Spectra from titration of **7** with **6a** a) ^1H NMR and b) ^{19}F NMR. The change of chemical shifts of *S* and *R* enantiomers of **7** as a function of the concentration of **6a** c) ^1H NMR and d) ^{19}F NMR. The chemical shift change of the *S* and *R* enantiomers ($\Delta\delta_S$ and $\Delta\delta_R$) of **7**, as a function of the concentration of **6a** e) ^1H NMR and f) ^{19}F NMR. (+ *R* enantiomer and ◆ *S* enantiomer)

To study the complexation Job's plot was calculated from the titration results (Fig 7). At the point where the molar ratio is 0.5:1 Job's plot indicated complexation 0.5:1 (both *S* and *R* enantiomer has the maximum at point at 0.333 at the Job's plot).

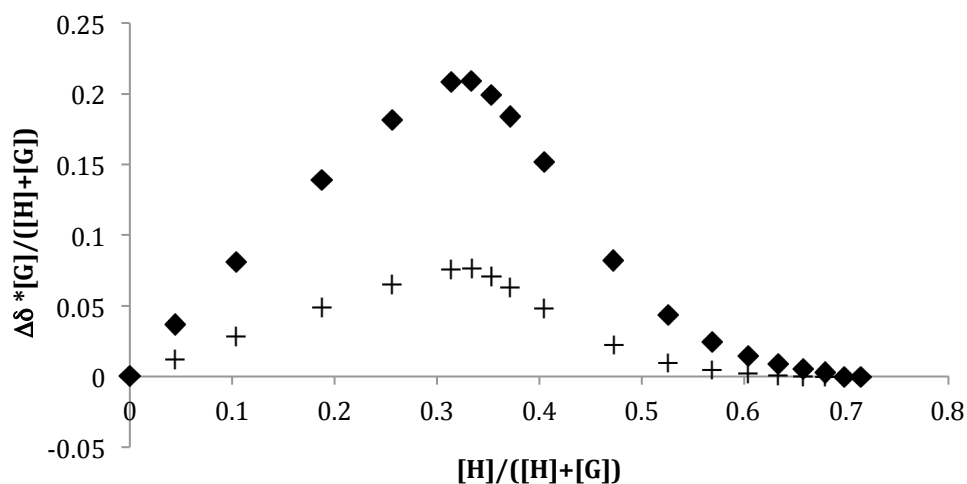


Figure 7. Job's plot from titration results (based on results from ^1H NMR) (+ *R* enantiomer and ◆ *S* enantiomer).

3.1.1 The effect of increasing concentration of **6a**

During the titration (as well as while performing chiral recognition studies) the methoxy peak signal from *S* enantiomer of **7** was noted to broaden. This was presumed to originate from strong binding of *S* enantiomer to host **6a** leading to intermediate exchange between free and bound forms.

To observe the free and bound forms of *S* enantiomer dynamic NMR study was performed (Fig 8). Sample (in CD_2Cl_2) containing only *S*-**7** (0.5 mL, 2.0 M, 1.0 eq) and host **6a** (0.22 μL , 46.6 M, 1.0 eq) was cooled down to $-65\text{ }^\circ\text{C}$. To maximise the temperature range CD_2Cl_2 (min $-95\text{ }^\circ\text{C}$) was used as solvent instead of CDCl_3 ($-63\text{ }^\circ\text{C}$). Also the effect of increasing temperature to the peak width was studied. Sample containing **7** (0.5 mL, 2.0 M, 1.0 eq) and host **6a** (0.22 μL , 46.6 M, 1.0 eq) in CDCl_3 was heated up to $+50\text{ }^\circ\text{C}$.

At temperature $-65\text{ }^\circ\text{C}$ the separation of the free and bound forms of *S* enantiomer is clearly detected (Fig. 8 a)). Temperature has notable effect on the shape of peak of *S* enantiomer (Fig 8 b)). The line width increases while temperature decreases (i.e. decreased exchange rate between free and bound form) and oppositely decreases at higher temperature (i.e. increased exchange rate between free and bound form). The line width of *R* enantiomer remains virtually unaffected throughout the studied temperature range. The result thus indicates higher affinity of *S* enantiomer with the host compared to *R* enantiomer.

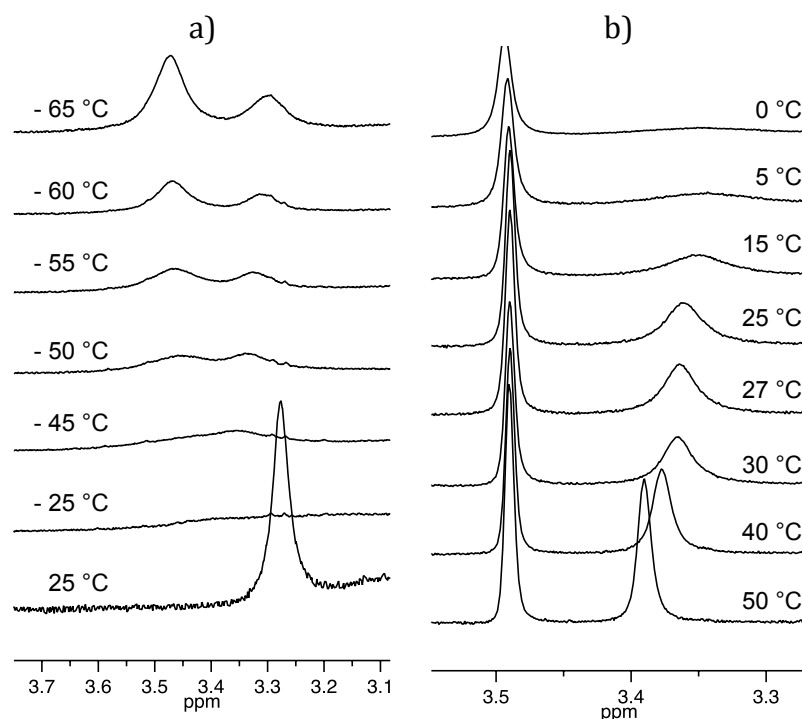


Figure 8. a) Dynamic NMR experiment in CD_2Cl_2 with enantiopure *S*-7 (-65 °C to +25 °C) and b) dynamic NMR experiment in CDCl_3 with **7** (0 °C to 50 °C)

Stronger binding also increases apparent molar weight of *S* enantiomer compare to *R* enantiomer. This was observed in DOSY experiment (Fig 9) where *S* enantiomer displays smaller diffusion coefficient.

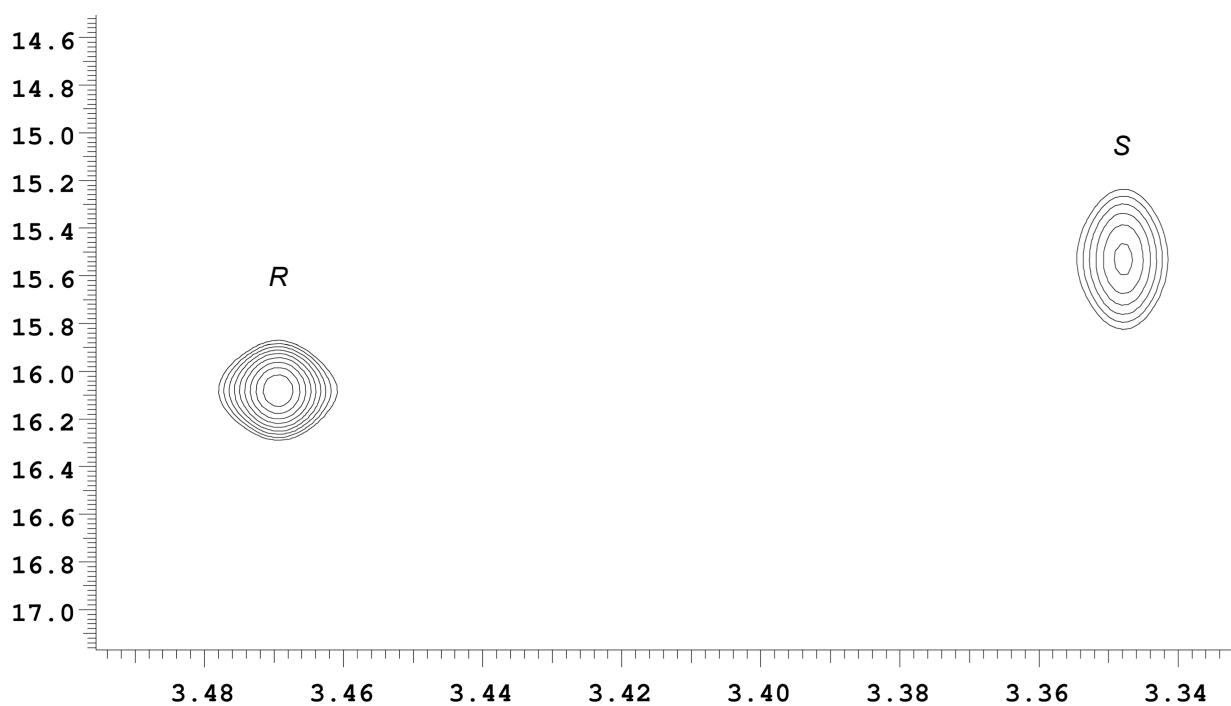


Figure 9. DOSY experiment of sample containing **7** and **6a** in 1:1 composition ($D_S = 15.5 \times 10^{-10} \text{ m}^2/\text{s}$ and $D_R = 16.1 \times 10^{-10} \text{ m}^2/\text{s}$).

To find if the complex formation occur in the same manner with pure enantiomer, *S*-7 was titrated with **6a** (Fig. 10). As in the case of racemic **7** the maximum separation was obtained at point where the host guest ratio was 1:2. When the concentration of host was increased the chemical shift change decreased as in the case of **7**. Also the peak width was observed to increase when the 1:2 host:guest ratio was reached.

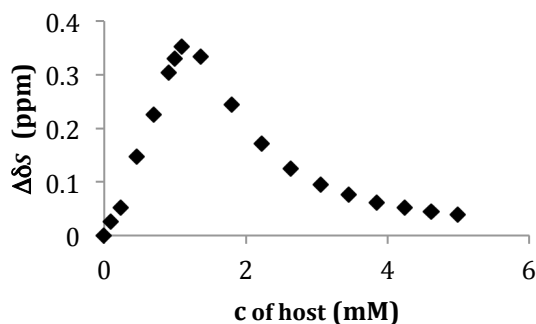


Figure 10. Chemical shift change of *S*-7 in the function of concentration.

The chemical shift of *S*-enantiomer in racemic mixture (0.315 ppm) was smaller than in titration containing only *S*-enantiomer (0.352 ppm). This might be caused by formation of complex *SHR* in racemic mixture instead of *SHS* as with pure *S*-7. It is also possible that some amount of *SHS* (as well as *RHR*) is formed with *SHR* when racemic **7** is titrated. Based on these assumptions the complexation pattern could be presumed to go in the manner illustrated in the Figure 11.

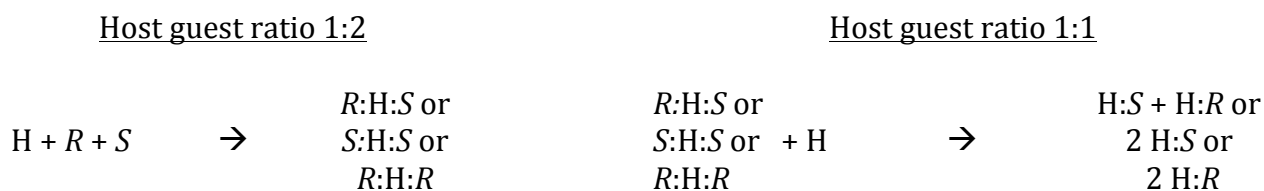


Figure 11. Possible complexation pattern formation in the titration of **7** with **6a**.

3.2 Titration of **8** with **6b**

Due to solubility problems of **6b** to CDCl₃, titration was performed in opposite manner compared to **6a**. Titration solutions containing the **6b**, host (2.0 mM) and guest **8** (46.6 mM) were prepared. In NMR tube 0.5 mL of host solution (1.0 eq) was measured and titrated by 2.5-5.0 μL doses of guest solution. Both ¹H and ¹⁹F NMR spectra were recorded. Concentration of host (**6b**) is presumed to remain constant during titration.

Table 2. Experimental data from titration of **8** with **6b**

c of 8 (mM)	V of 8 (mL)	¹ H NMR ^[a]				¹⁹ F NMR ^[b]			
		δ of <i>S</i> (ppm)	δ of <i>R</i> (ppm)	Δδ _{<i>S</i>} of <i>S</i> (ppm)	Δδ _{<i>R</i>} of <i>R</i> (ppm)	δ of <i>S</i> (ppm)	δ of <i>R</i> (ppm)	Δδ _{<i>S</i>} of <i>S</i> (ppm)	Δδ _{<i>R</i>} of <i>R</i> (ppm)
0	0	0	0	0	0	0	0	0	0
0.231840796	0.0025	3.328	3.407	0.303	0.224	-70.585	-70.636	-0.029	0.022
0.461386139	0.005	3.325	3.406	0.306	0.225	-70.614	-70.676	0	0.062
0.91372549	0.01	3.329	3.406	0.302	0.225	-70.745	-70.797	0.131	0.183
1.357281553	0.015	3.241	3.408	0.39	0.223	-71.007	-71.007	0.393	0.393
1.575845411	0.0175	3.223	3.41	0.408	0.221	-71.069	-71.108	0.455	0.494
1.792307692	0.02	3.213	3.412	0.418	0.219	-71.109	-71.18	0.495	0.566
2.006698565	0.0225	3.213	3.417	0.418	0.214	-71.133	-71.219	0.519	0.605
2.219047619	0.025	3.221	3.425	0.41	0.206	-71.143	-71.236	0.529	0.622
2.429383886	0.0275	3.242	3.443	0.389	0.188	-71.127	-71.206	0.513	0.592
2.637735849	0.03	3.273	3.462	0.358	0.169	-71.105	-71.166	0.491	0.552
3.048598131	0.035	3.323	3.493	0.308	0.138	-71.061	-71.095	0.447	0.481
3.451851852	0.04	3.364	3.516	0.267	0.115	-71.024	-71.024	0.41	0.41
3.847706422	0.045	3.395	3.532	0.236	0.099	-70.986	-70.986	0.372	0.372
4.236363636	0.05	3.421	3.543	0.21	0.088	-70.951	-70.951	0.337	0.337
4.618018018	0.055	3.441	3.553	0.19	0.078	-70.922	-70.922	0.308	0.308
4.992857143	0.06	3.458	3.561	0.173	0.07	-70.896	-70.896	0.282	0.282
5.722807018	0.07	3.485	3.571	0.146	0.06	-70.854	-70.854	0.24	0.24
6.427586207	0.08	3.505	3.579	0.126	0.052	-70.823	-70.823	0.209	0.209
7.108474576	0.09	3.521	3.586	0.11	0.045	-70.796	-70.796	0.182	0.182

[a] The peak of OCH₃ in ¹H NMR is at 3.631 ppm and [b] the peak of CF₃ in ¹⁹F NMR is at -70.614 ppm when the concentration of **8** is 2.0 mM (no host added).

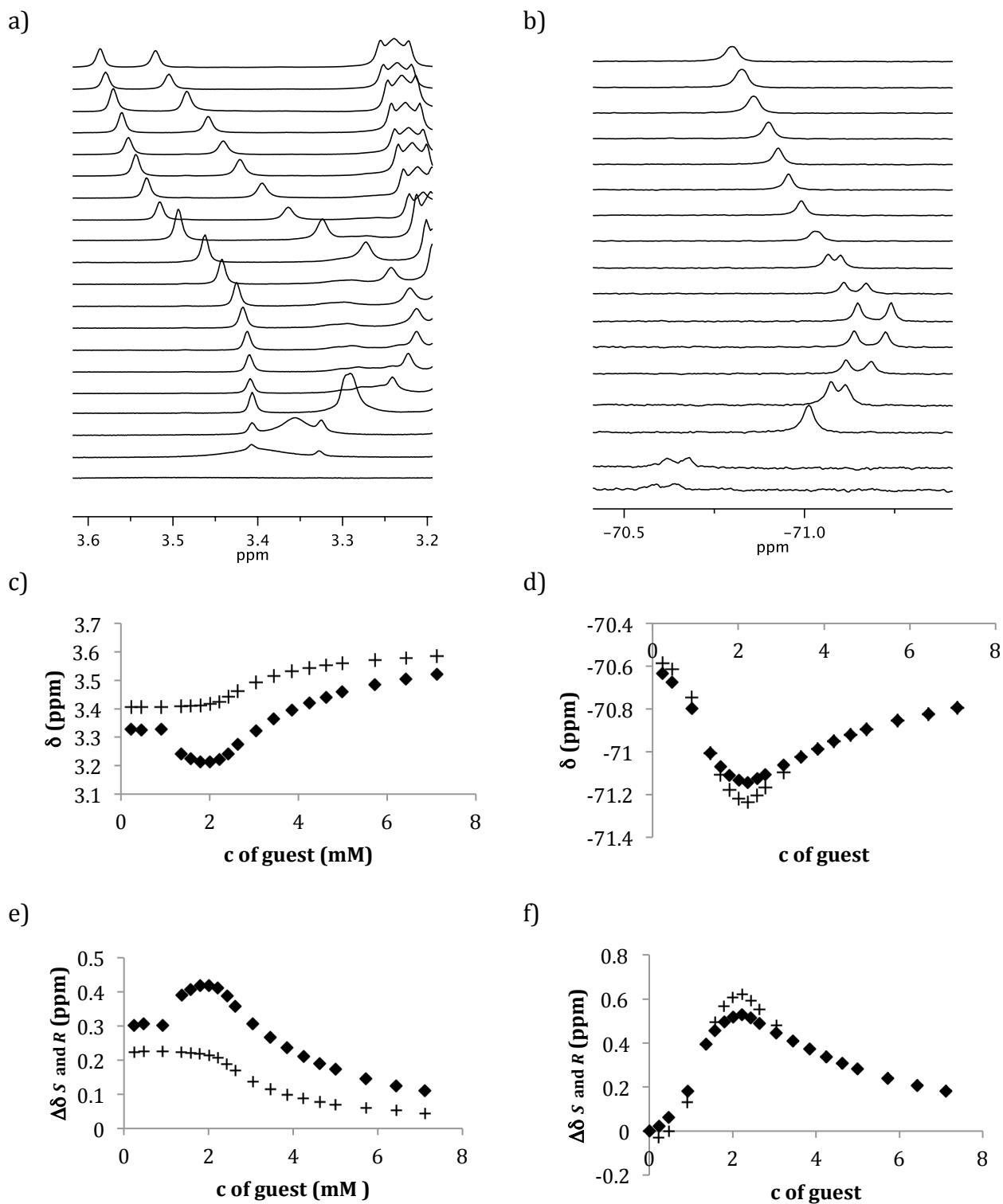


Figure 12. Spectra from titration of **8** with **6b** a) ^1H NMR and b) ^{19}F NMR. The change of chemical shifts of *S* and *R* enantiomers of **8** c) ^1H NMR and d) ^{19}F NMR. The chemical shift change of the *S* and *R* enantiomers ($\Delta\delta_S$ and $\Delta\delta_R$) of **8** e) ^1H NMR and f) ^{19}F NMR. (+ *R* enantiomer and ♦ *S* enantiomer)

To study the complexation Job's plot was calculated from the titration results (Fig. 12). At the point where the molar ratio is 1:1 Job's plot indicated complexation 1:1 (both *S* and *R* enantiomer has the maximum at point at 0.5).

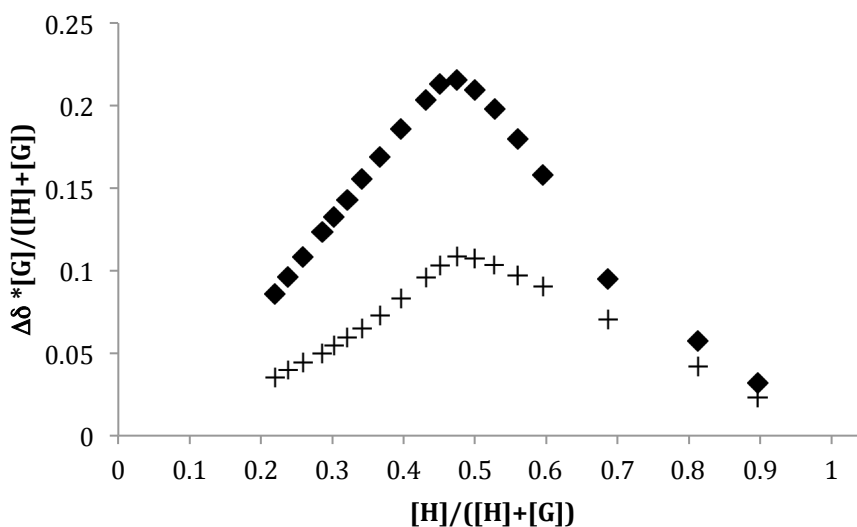


Figure 13. Job's plot from titration results (based on results from ^1H NMR) (+ *R* enantiomer and ◆ *S* enantiomer).

3.2.1 Dependence of concentration on $\Delta\delta$

During the titration of **6b** (2.0 mM) the chemical shift difference between *R* and *S* enantiomers ($\Delta\delta$) in ^{19}F NMR spectra was not as high as in enantiomeric separation experiments (22.0 mM). To investigate if the concentration had an effect on $\Delta\delta$, a sample containing host:guest in ration 2:1 (22.0 mM) was diluted by CDCl_3 (Fig. 14). Results indicate that the concentration has distinct effect on the $\Delta\delta$ at ^{19}F NMR (change of $\Delta\delta$: 0.08 ppm, 37.6 Hz), however, in the case of ^1H NMR slight increase in $\Delta\delta$ values was detected (change of $\Delta\delta$: 0.006 ppm, 3.0 Hz).

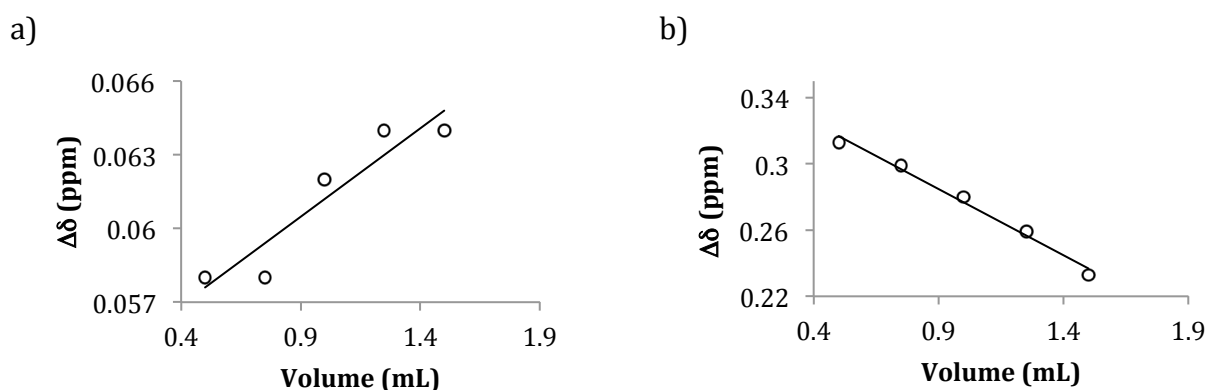


Figure 14. The effect of increased solvent volume (CDCl_3) on $\Delta\delta$ in a) ^1H NMR and b) ^{19}F NMR.

4 ENANTIOMERIC EXCESS STUDIES

4.1 Enantiomeric excess measurements of 7 with 6a

In ee% determination studies solutions of racemic **7** and *S*-**7** were prepared (2.0 mM). Mixtures of enantiomerically enriched samples were prepared in NMR tube (0.5 mL, 2.0 mM, 1.0 eq) and added **6a** (46.6 mM, 22.5 μ L, 1.0 eq) and recorded spectra. Due to overlapping of peaks of host and guest, 1.0 eq (22.5 μ L 46.6 mM) of **6a** was added instead of 0.5.

Table 3. ee% determined from ^1H NMR spectra

<i>S</i> (mL)	<i>R/S</i> (mL)	Expected <i>S/R</i>	Measured <i>S/R</i>	Area <i>S</i>	Area <i>R</i>	Expected ee% (<i>S</i>)	Measured ee% (<i>S</i>)
0	0.5	1	0.965717071	2488.509	2576.851	50	49.12797906
0.1	0.4	1.5	1.474179057	23892.418	16207.27	60	59.58255336
0.2	0.3	2.333	2.31684909	25898.506	11178.331	70	69.85090449
0.25	0.25	3	2.915923268	28472.871	9764.616	75	74.46323813
0.3	0.2	4	3.401767245	28246.343	8303.432	80	77.28185194

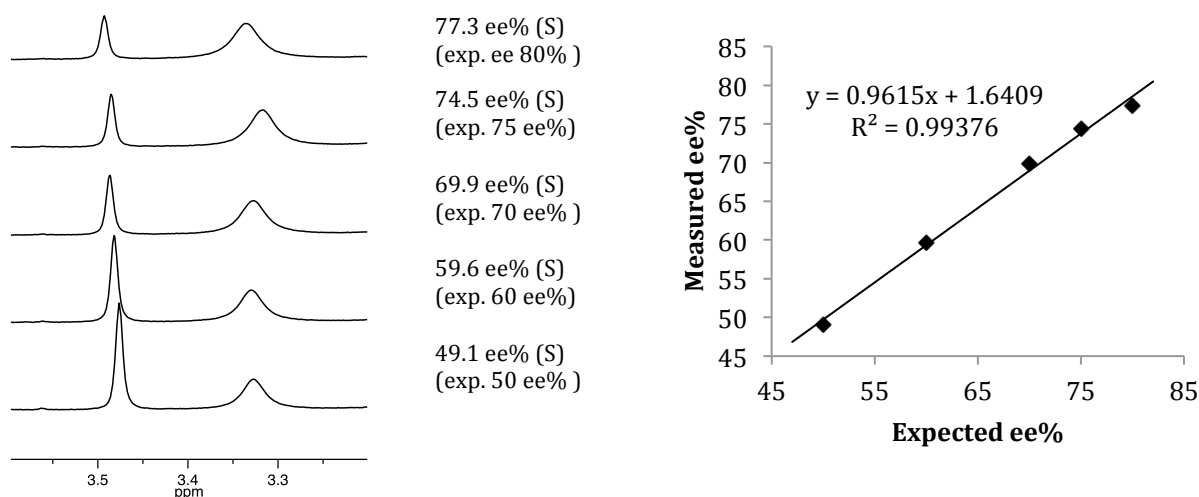


Figure 15. Results from ee% determination (^1H NMR).

4.2 Enantiomeric excess measurements of 8 with 6b

Due to solubility problems of **6b** ee% measurements were performed in opposite manner compared to **6a**. Solutions of racemic **8** and *S*-**8** were prepared (46.6 mM) as well as solution containing the host (2.0 mM). To NMR tube containing the host (0.5 mL, 2.0 mM, 1.0 eq) was added different portions of solutions containing racemic **8** and *S*-**8** (46.6 mM) to yield total 60.0 μ L (2.8 eq) of guest. (Higher equivalent rate was chosen to ease the calculation of *S*:*R* ratio and addition of analyte.)

Table 4. ee% determined from ^1H NMR spectra

<i>S</i> (μ L)	<i>R/S</i> (μ L)	Expected <i>S/R</i>	Measured <i>S/R</i>	Area <i>S</i>	Area <i>R</i>	Expected ee% (<i>S</i>)	Measured ee% (<i>S</i>)
0.0	60.0	1	0.982591241	10594.285	10781.986	50	49.56095944
10.0	50.0	1.4	1.445427826	13368.496	9248.816	58	59.10735988
20.0	40.0	2	1.976496042	15224.866	7702.958	67	66.40344936
30.0	30.0	3	3.020116878	16154.578	5348.991	75	75.12510133
25.0	35.0	3.8	3.705853512	16159.793	4360.613	79	78.74986976

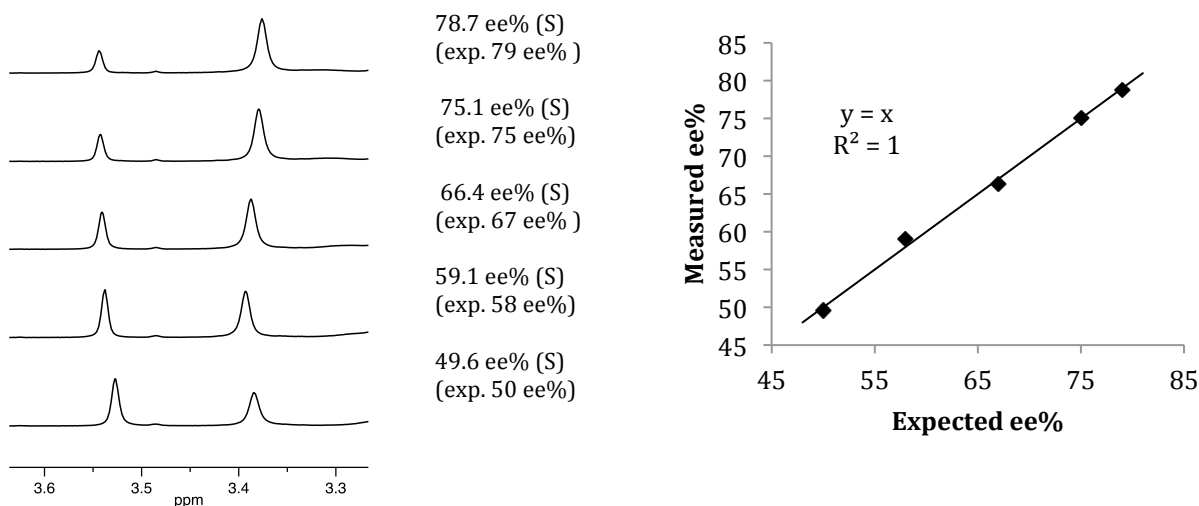


Figure 16. Results from ee% determination (¹H NMR).

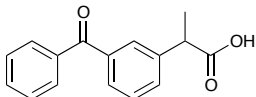
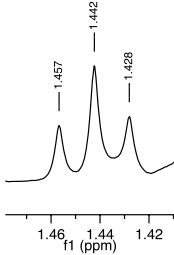
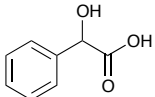
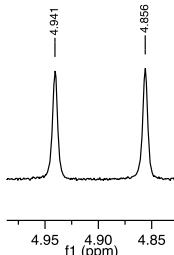
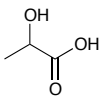
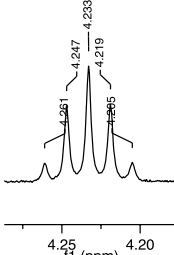
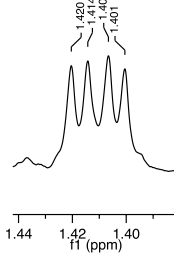
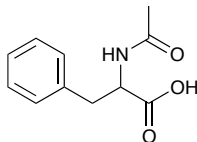
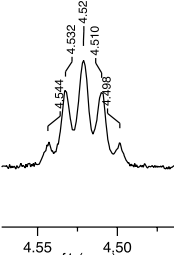
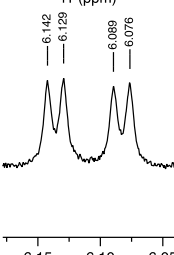
5 SEPARATION OF CARBOXYLIC ACIDS

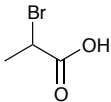
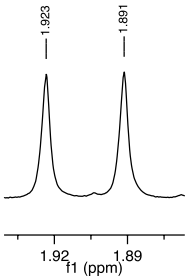
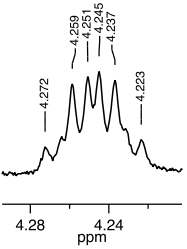
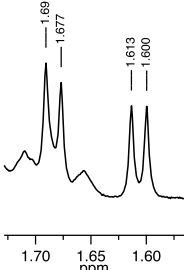
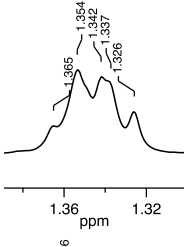
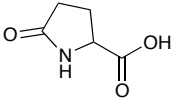
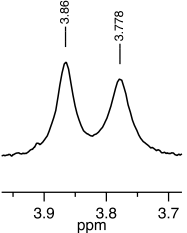
5.1 Separation of carboxylic acids with 6a

In chiral recognition ability determination study of racemic carboxylic acids, solutions of different racemic carboxylic acids were prepared (2.0 mM). Experiment was performed by adding 11.0 μL of solution containing host (46.6 mM, 0.5 eq) to NMR tube containing 0.5 mL of guest (2.0 mM, 1.0 eq) and recorded spectra.

Table 5. Determination of the chemical shift difference between enantiomers ($\Delta\delta$) of five racemic carboxylic acids in the presence of **6a**, using ¹H NMR (500 MHz) in CDCl₃ at 27 °C.

No.	Racemic carboxylic acid	Actual multiplet	Spectra (if split is detected)	$\Delta\delta$ (ppm)	
				0.002	
9a		H	q		0.0052
		CH ₂	d		0.0062

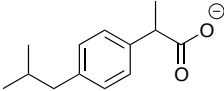
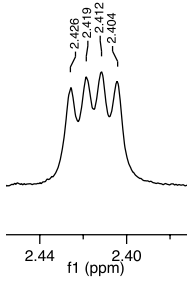
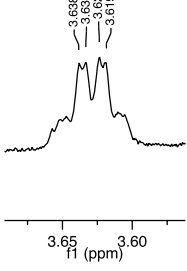
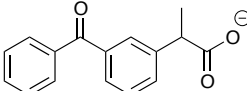
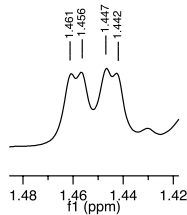
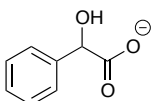
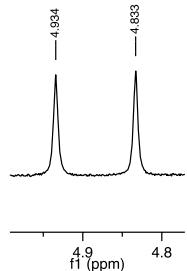
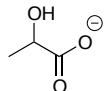
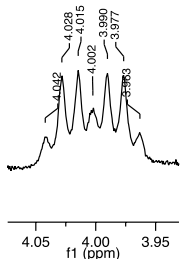
10a		Me	d		0.014
		H	q	nd	nd
11a		H	s		0.084
12a		H	q		0.014
		Me	d		0.0062
13a		H	q		0.01
		NH	d		0.052

14a		Me	s		0.033
		H	q		0.014
		Me	d		0.077
		H	t		0.016
15a		NH	br. s		0.088

5.2 Separation of carboxylic acids with 6b

In chiral recognition ability determination study of racemic carboxylic acids, solutions of different $[\text{Bu}_4\text{N}]^+$ salts of carboxylic acids were prepared (46.6 mM). Experiment was performed by adding 22.5 μL of solution containing guest to NMR tube containing 0.5 mL of host (2.0 mM, 1.0 eq) and recorded spectra.

Table 6. Determination of the chemical shift difference between enantiomers ($\Delta\delta$) of five racemic carboxylic acid $[\text{Bu}_4\text{N}]^+$ salts in the presence of **6**, using ^1H NMR (500 MHz) in CDCl_3 at 27 $^\circ\text{C}$.

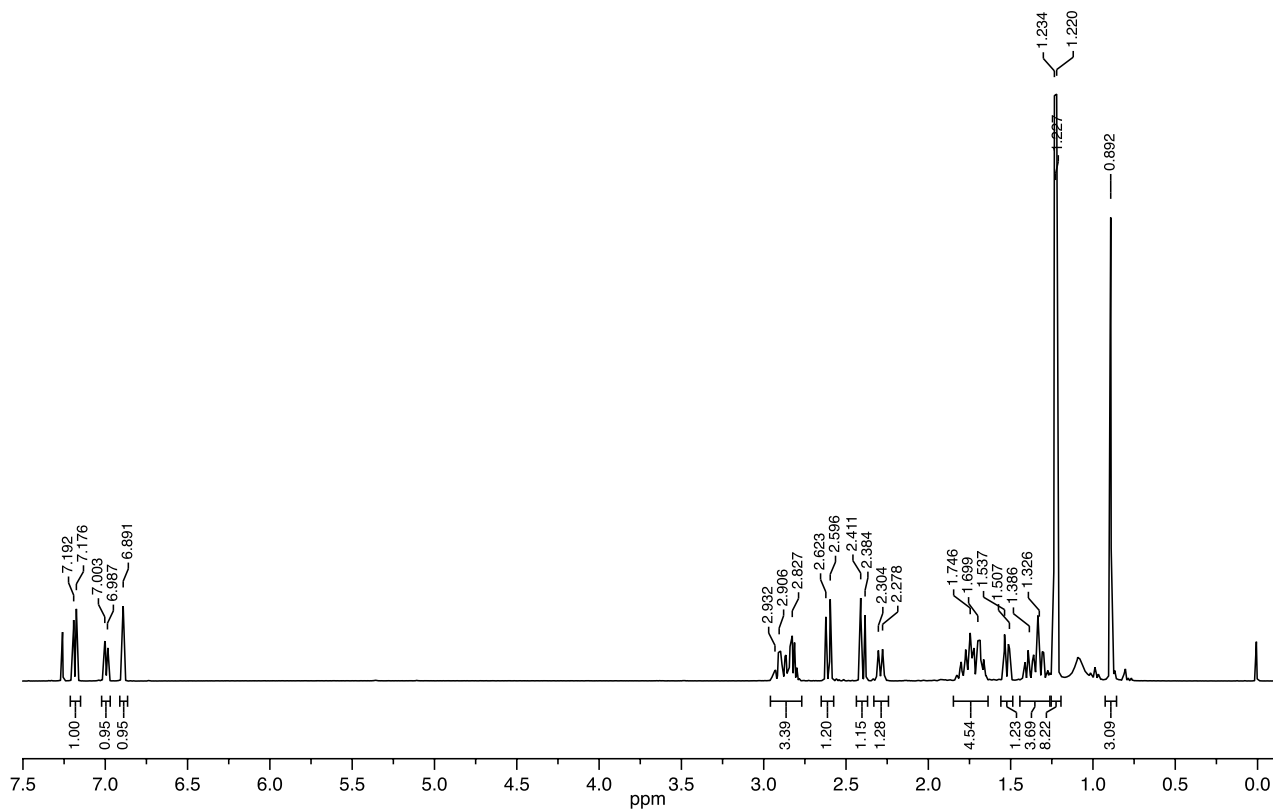
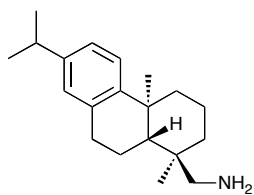
No.	[Bu ₄ N] ⁺ salt of racemic carboxylic acid	Actual multiplet	Spectra	Δδ (ppm)
9b		Me		0.0099
		H		0.0053
		CH ₂	-	-[a]
10b		Me		0.01
		H	nd	nd
11b		H		0.10
12b		H		0.038
		Me	-	-[a]
13b		H	nd	nd

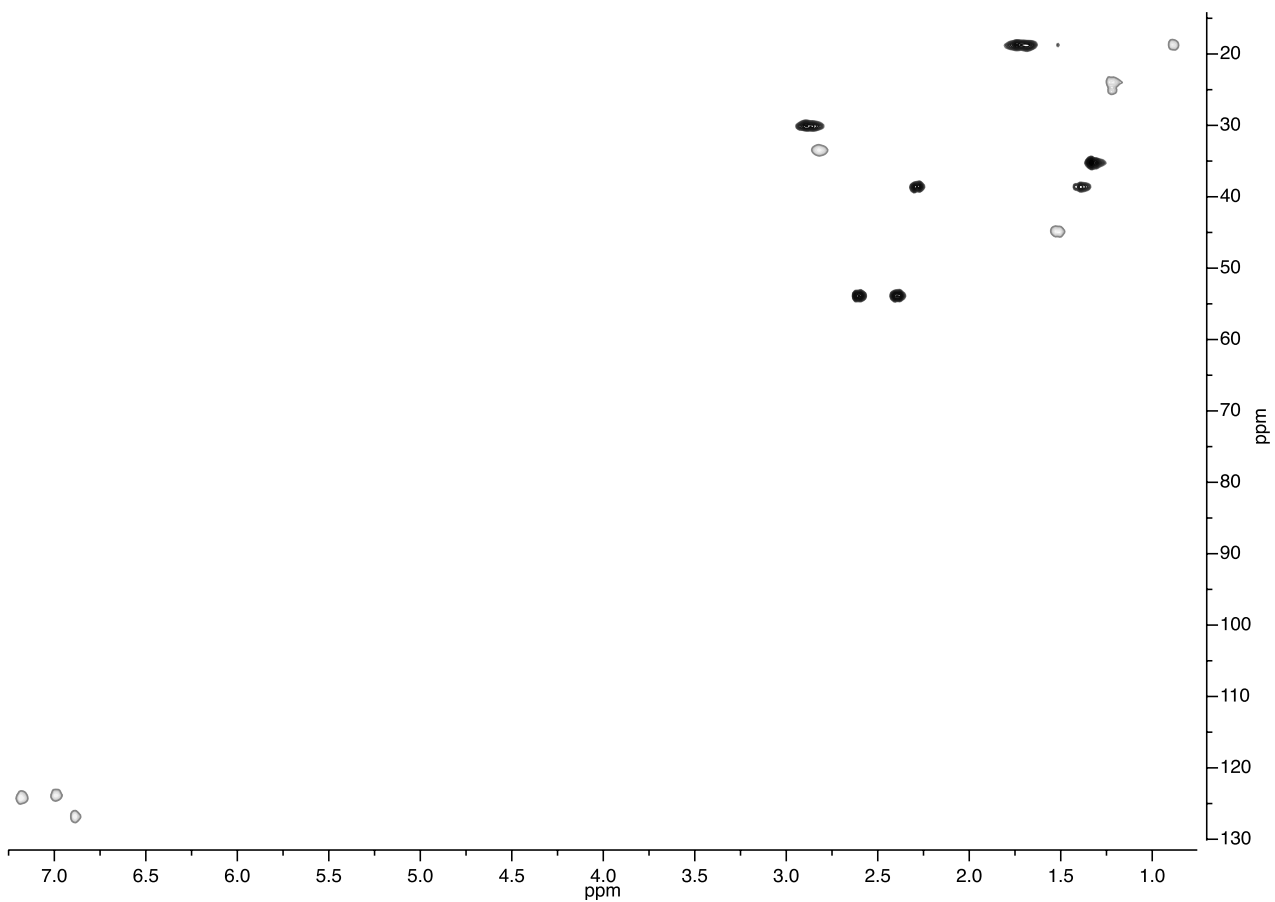
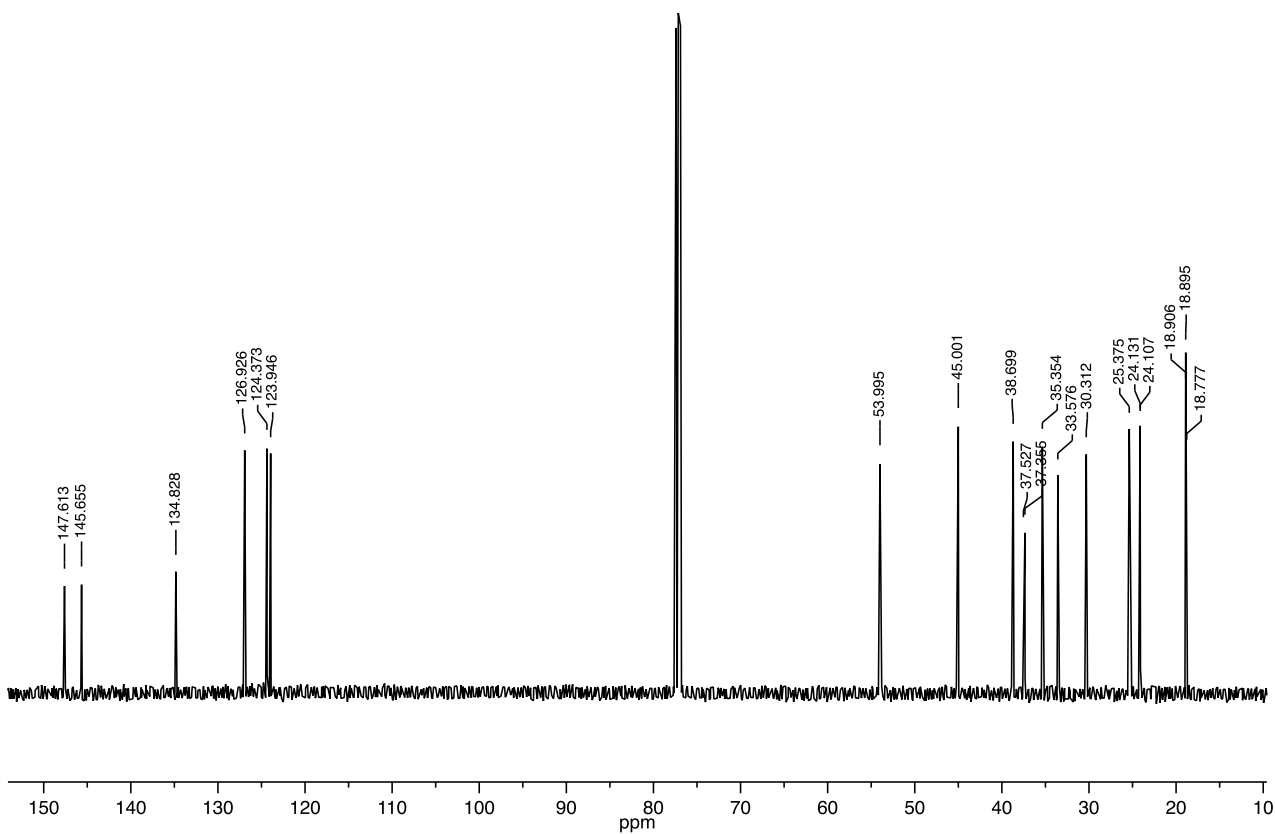
		NH	d		0.051
		Me	s		0.033
14a		H	q		0.016
		Me	d	-	-[a]
15a		H	t		0.017
		NH	br. s		0.072

[a] Peaks overlapped, nd (resolution not detected)

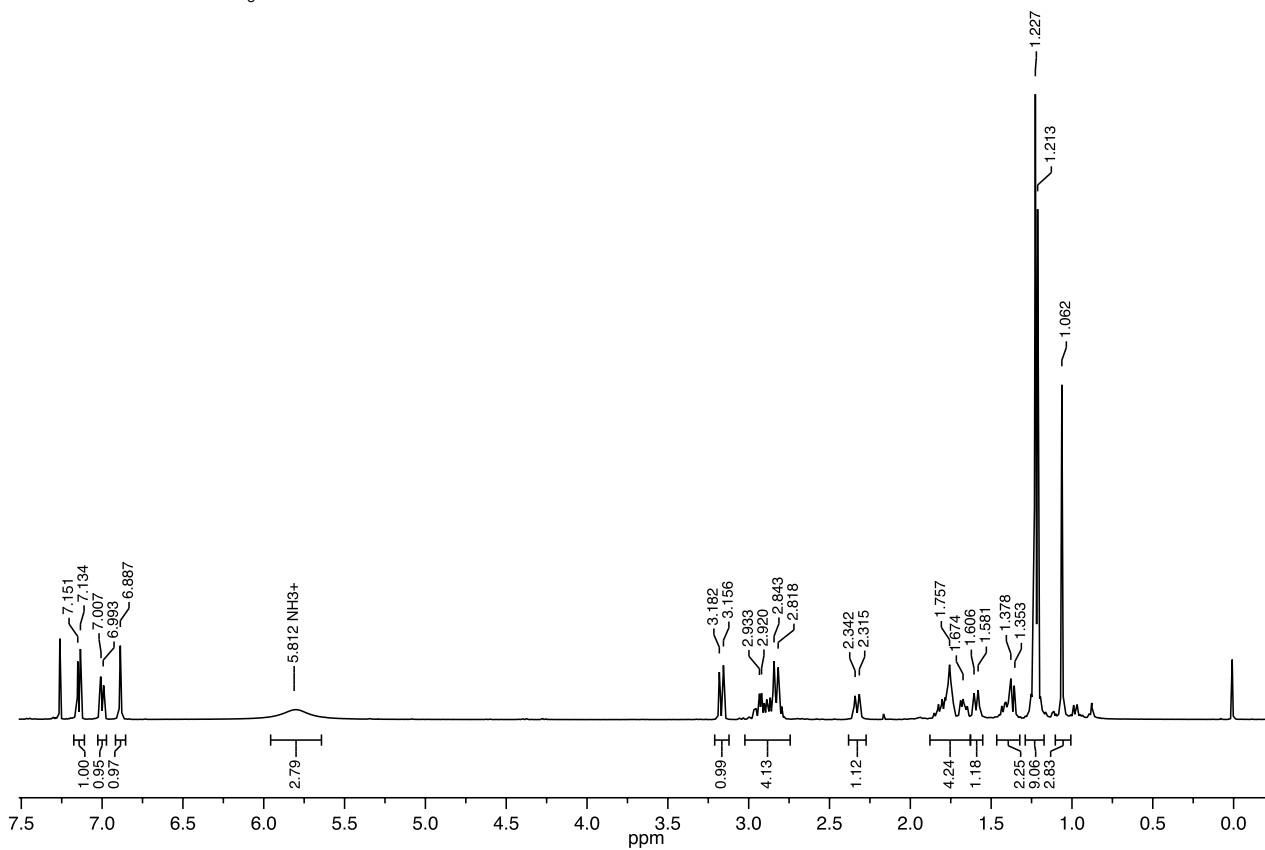
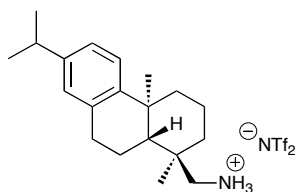
6 NMR SPECTRA OF SYNTHESISED CSAS

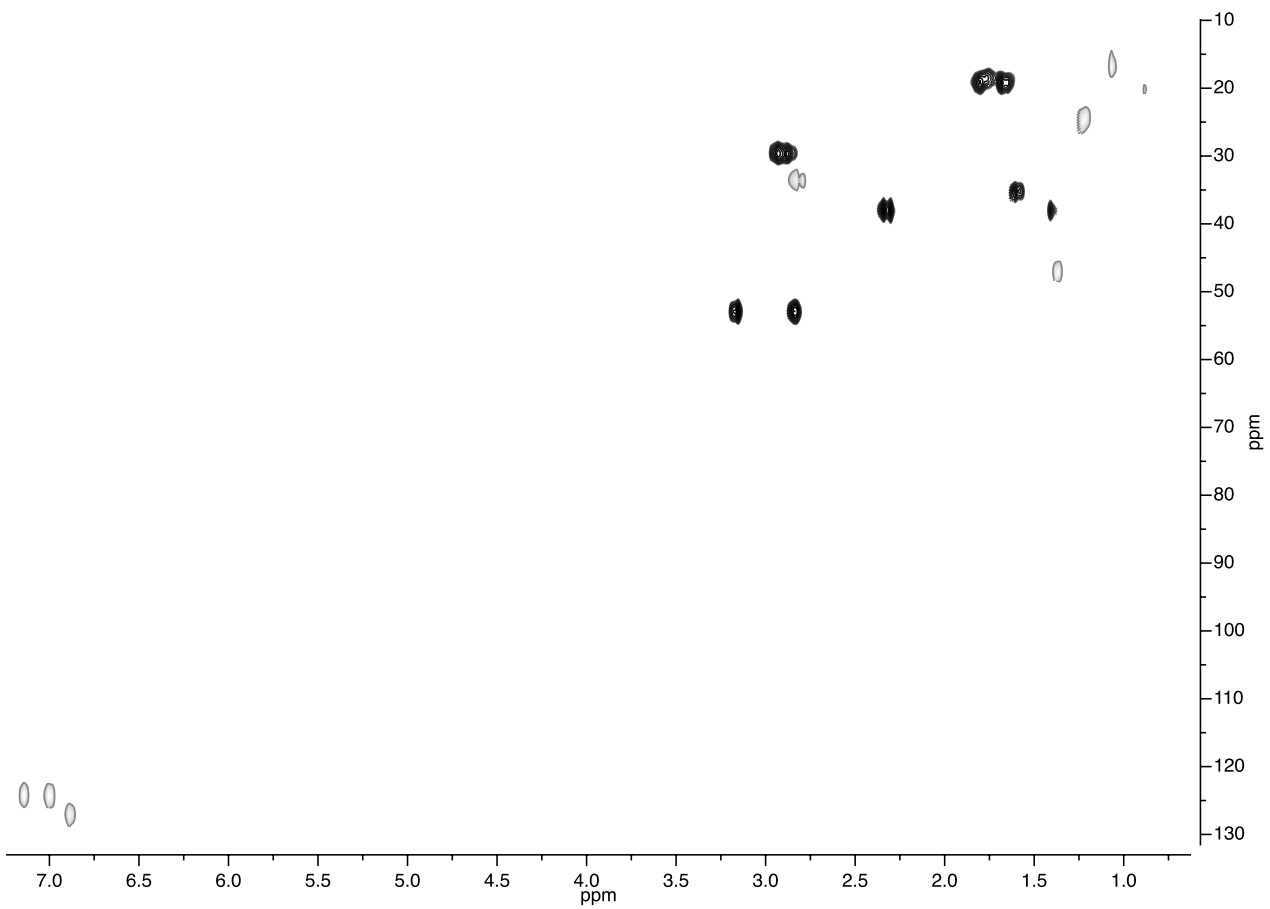
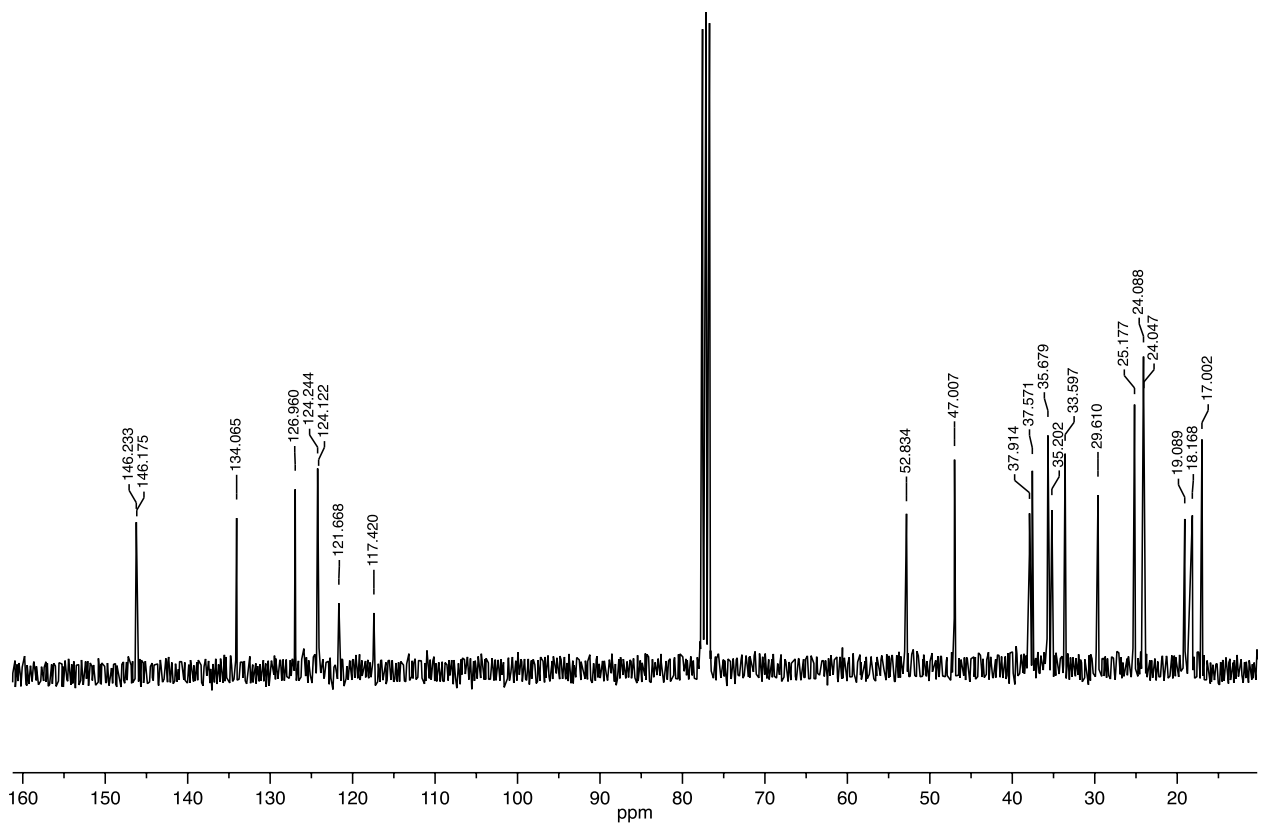
1H, 13C and HSQC spectra of compound 1a



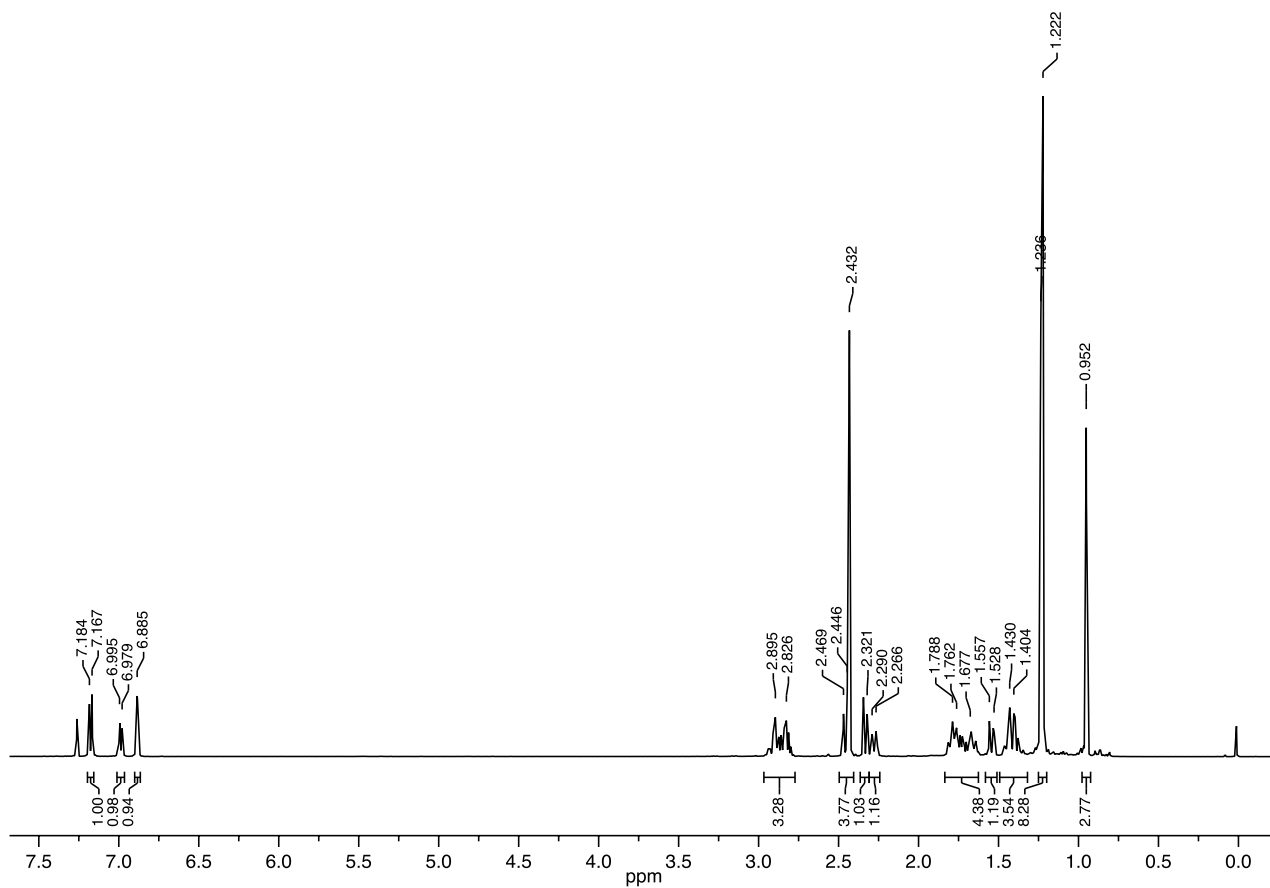
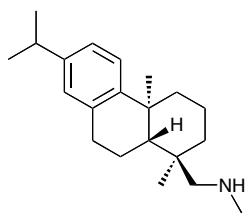


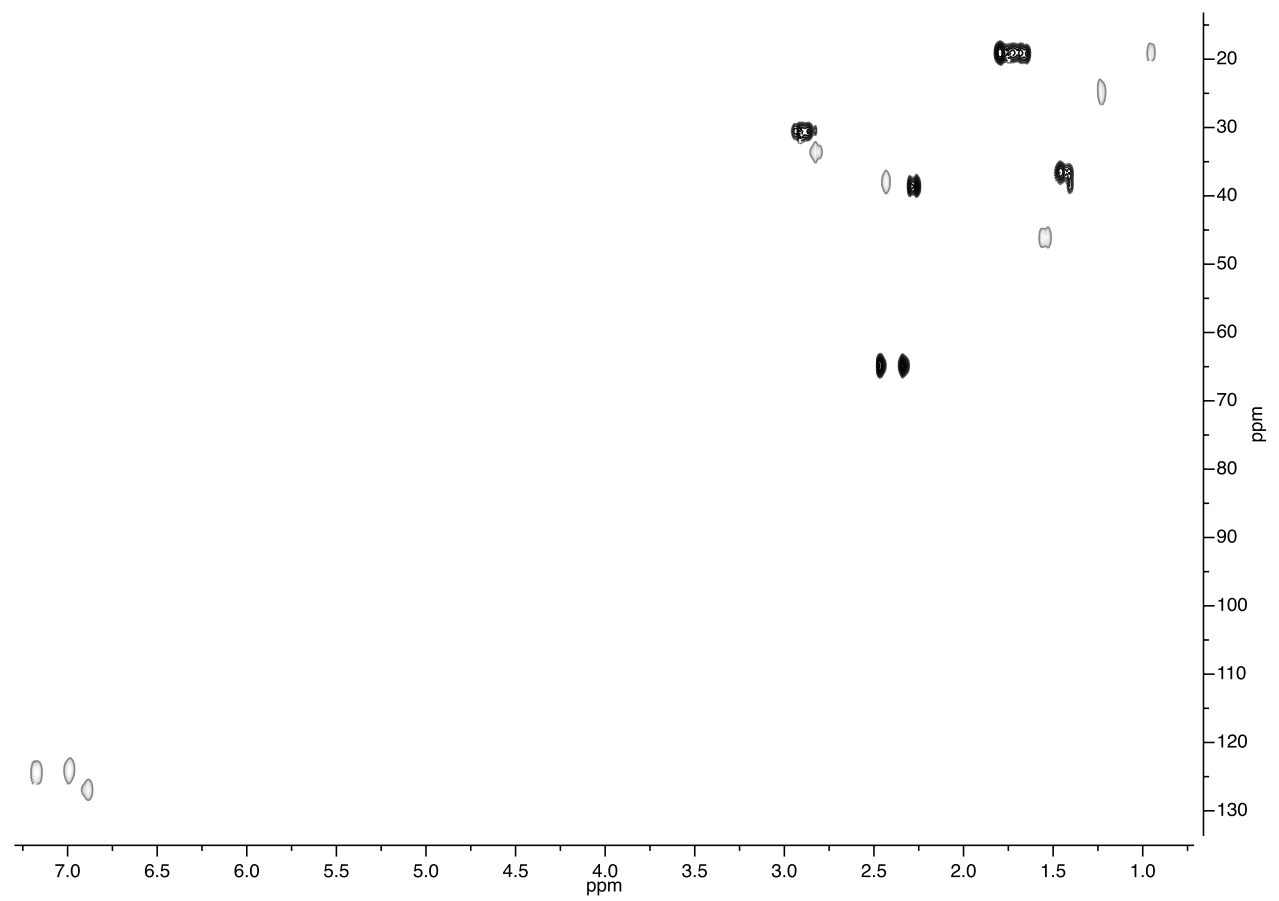
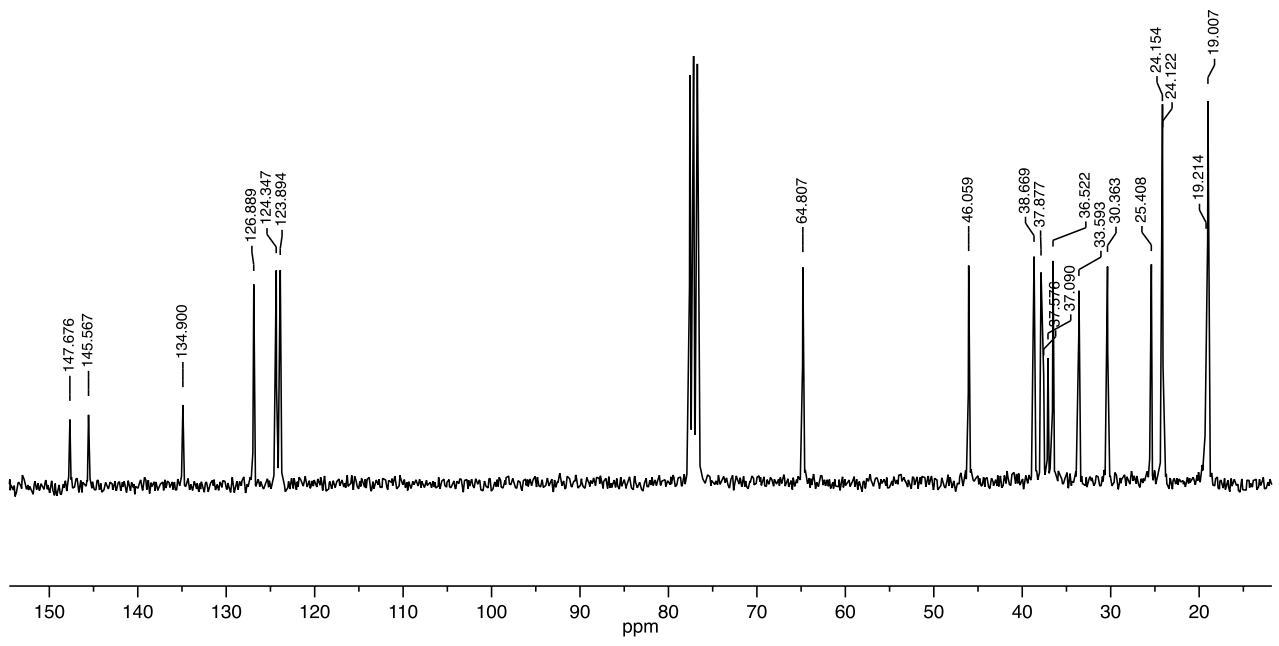
1H, 13C and HSQC spectra of compound 1b



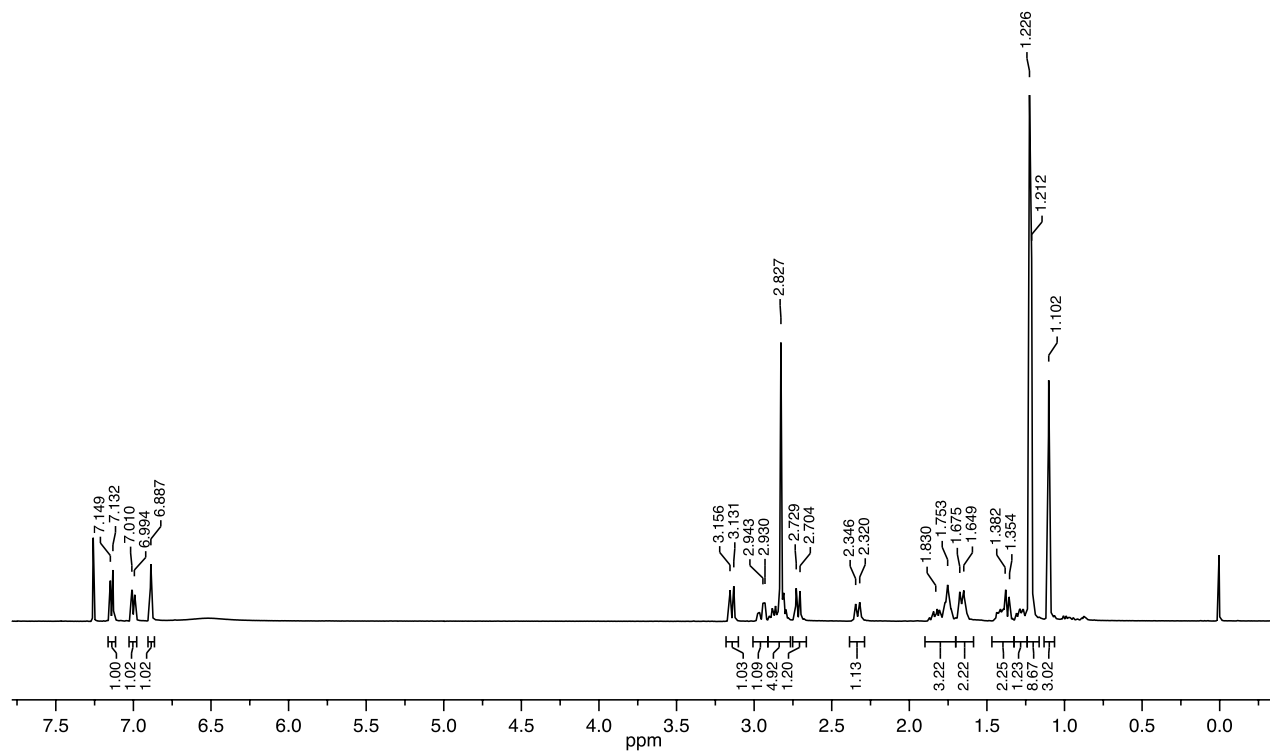
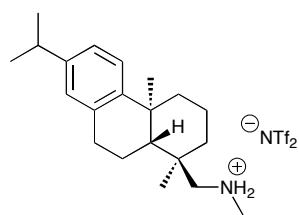


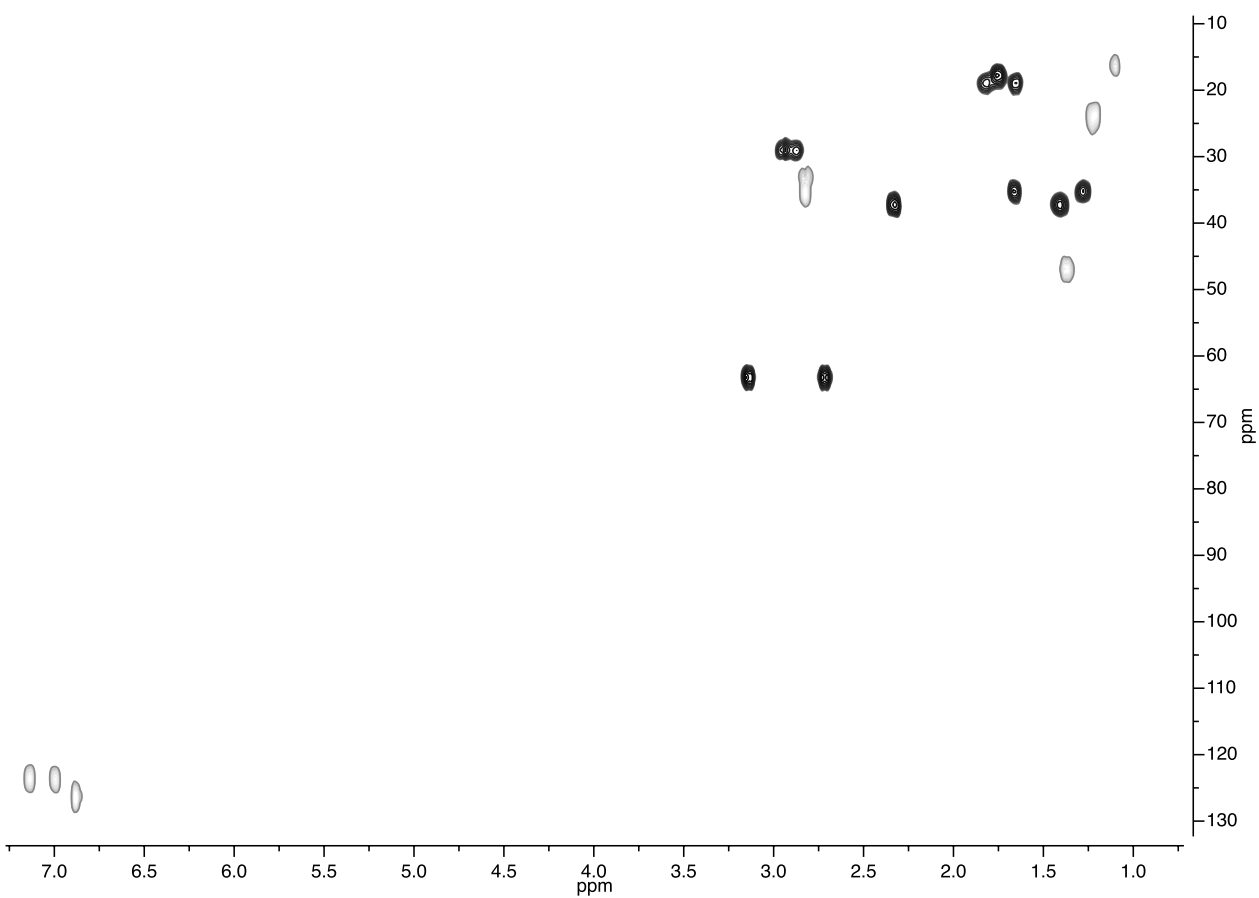
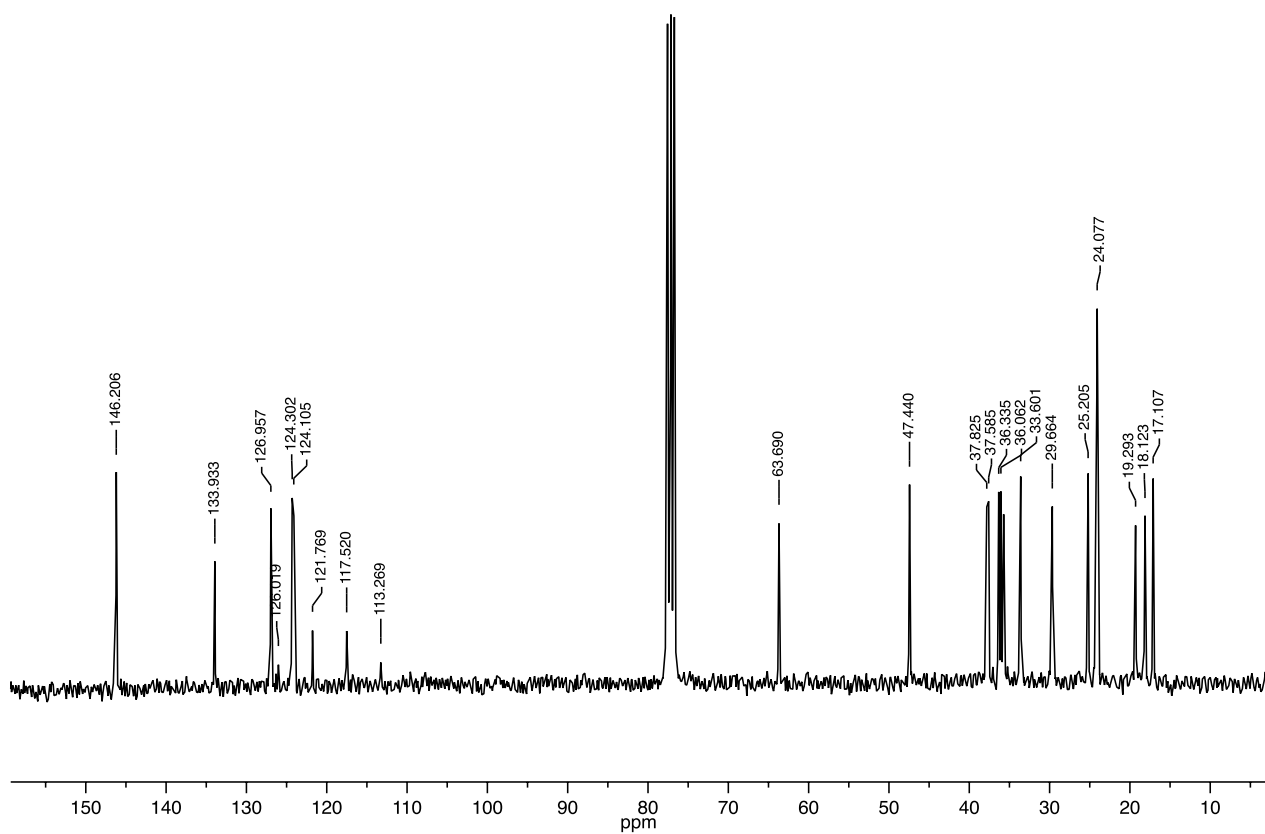
1H, 13C and HSQC spectra of compound 2a



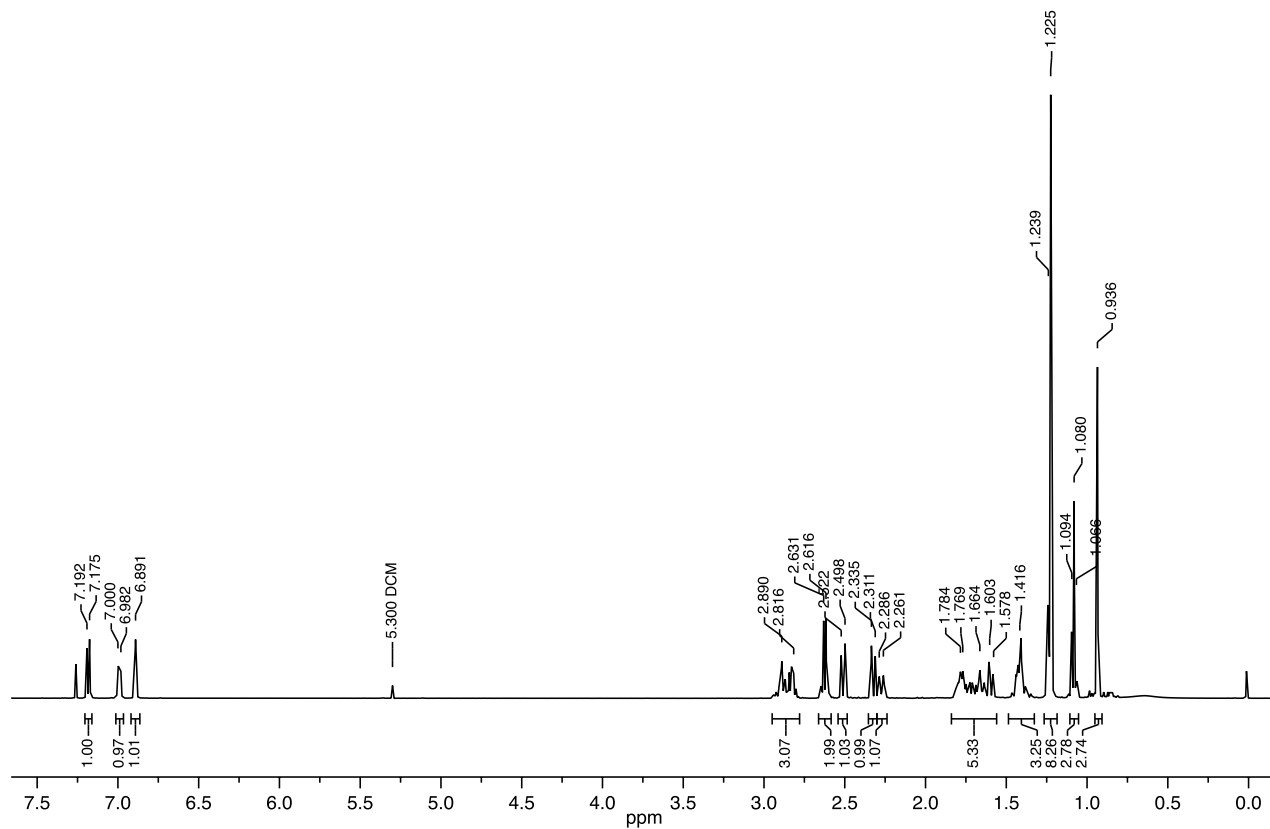
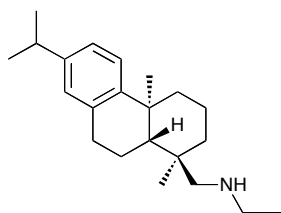


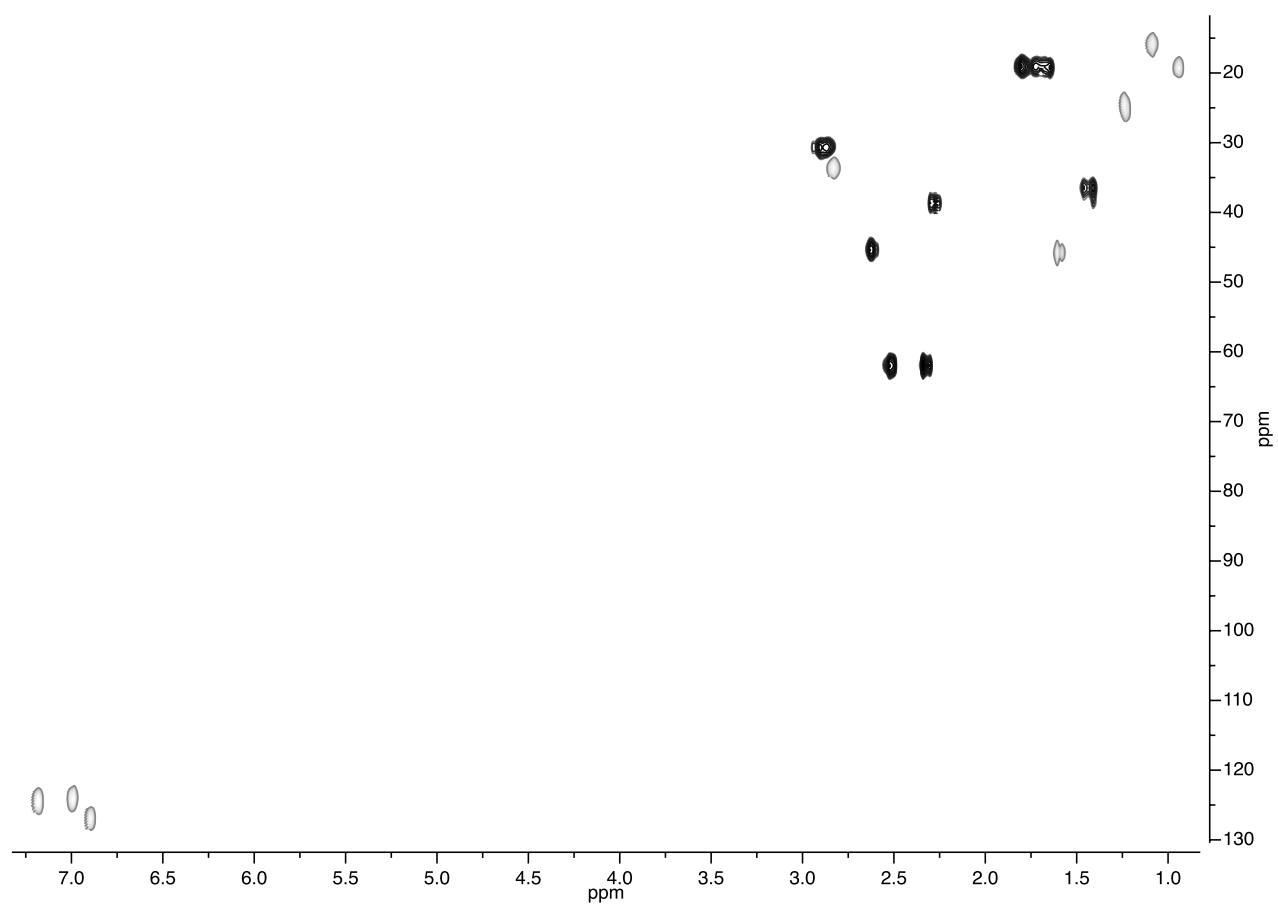
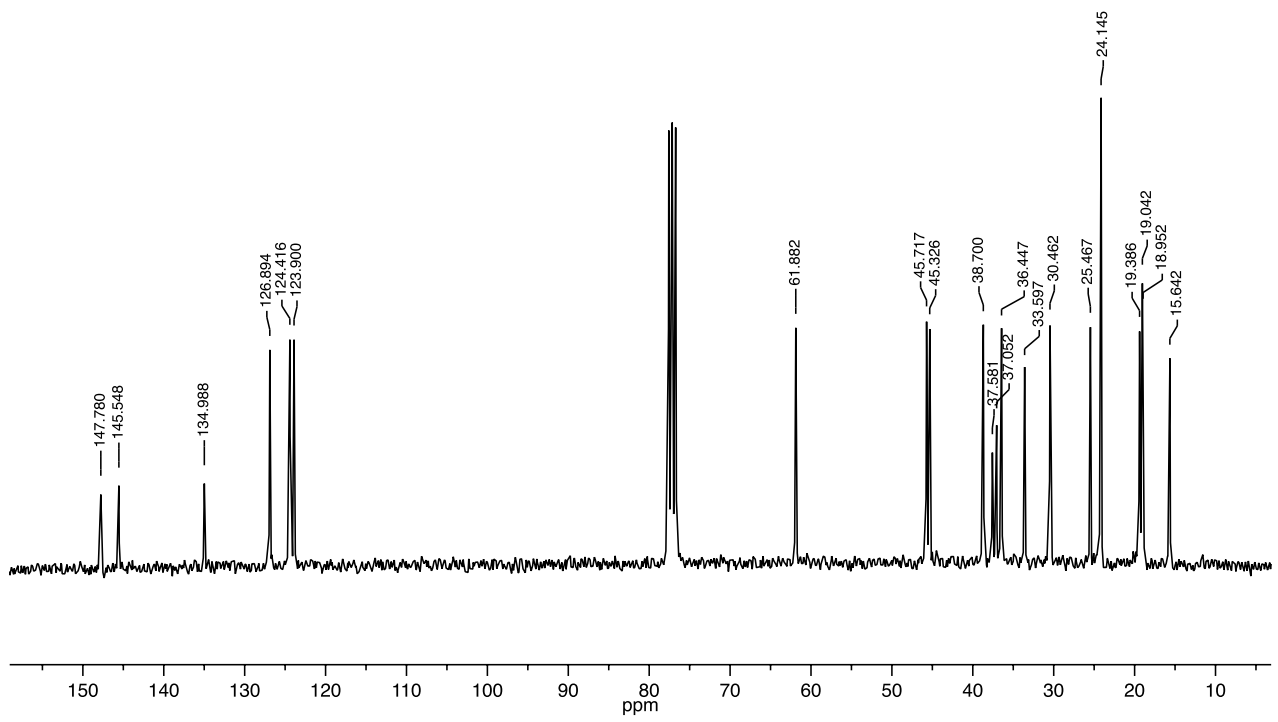
1H, 13C and HSQC spectra of compound 2b



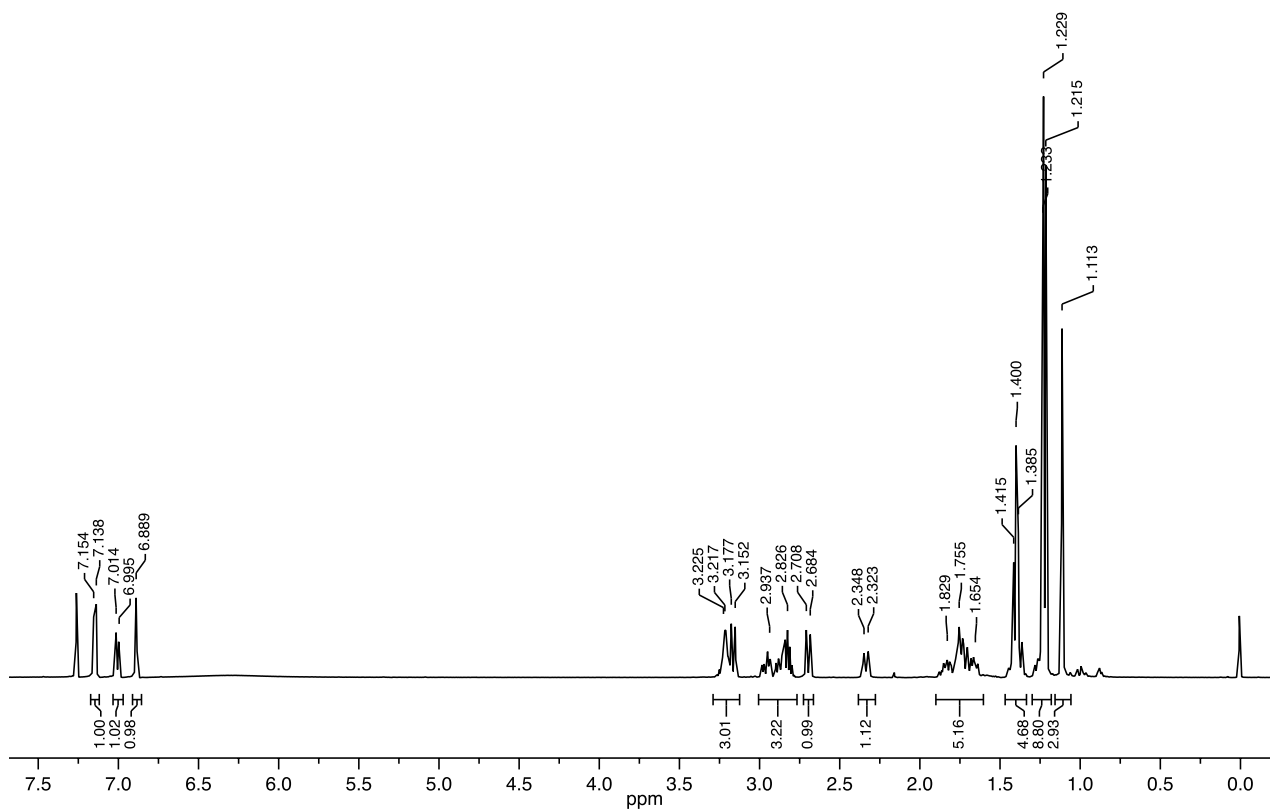
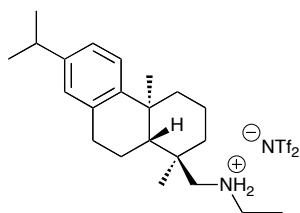


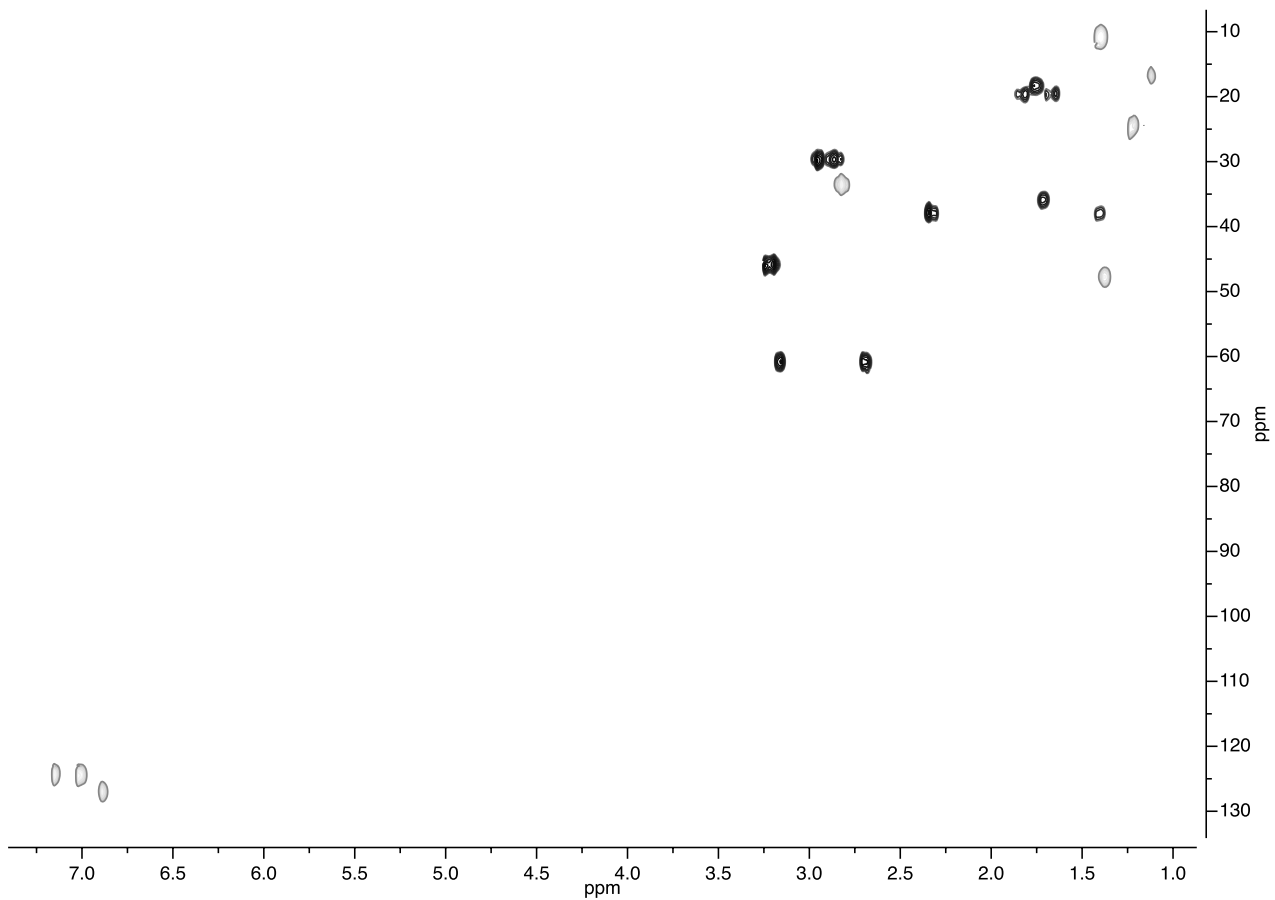
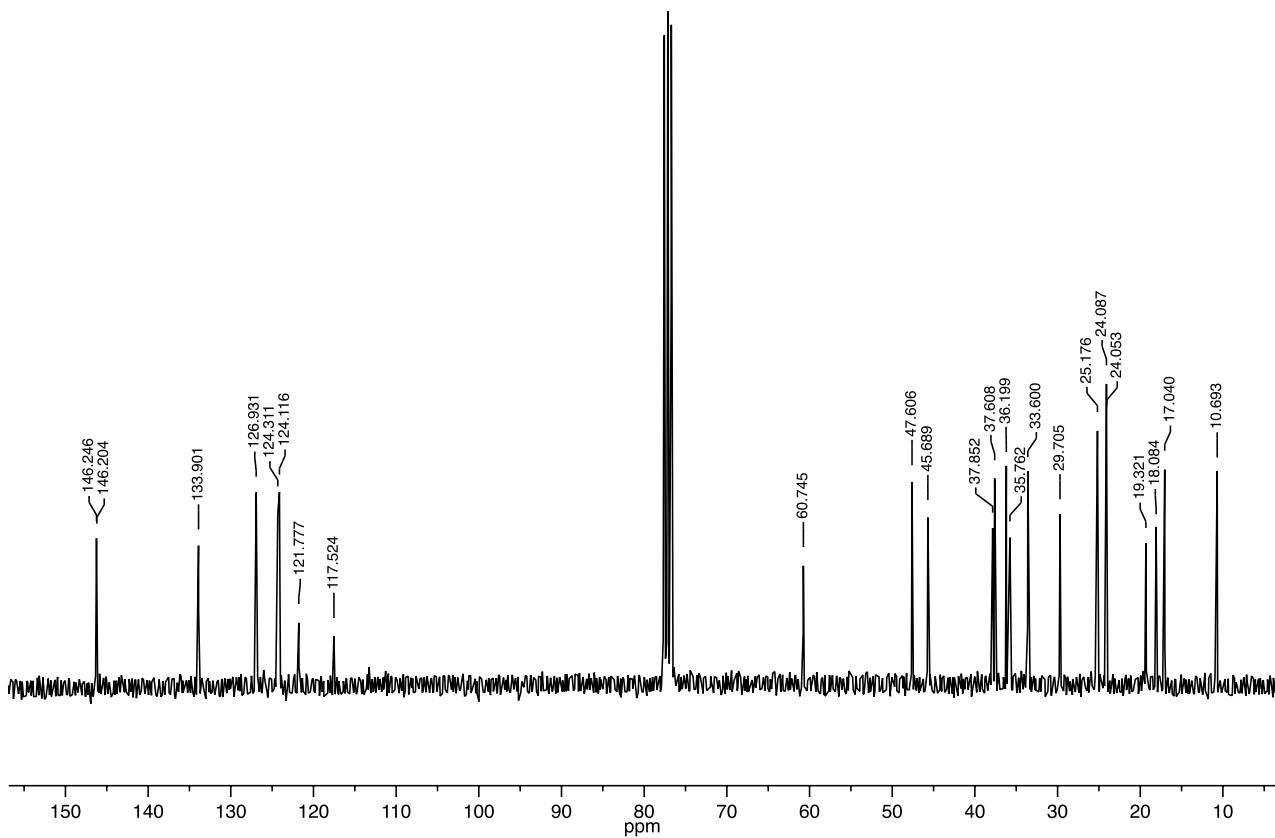
1H, 13C and HSQC spectra of compound 3a



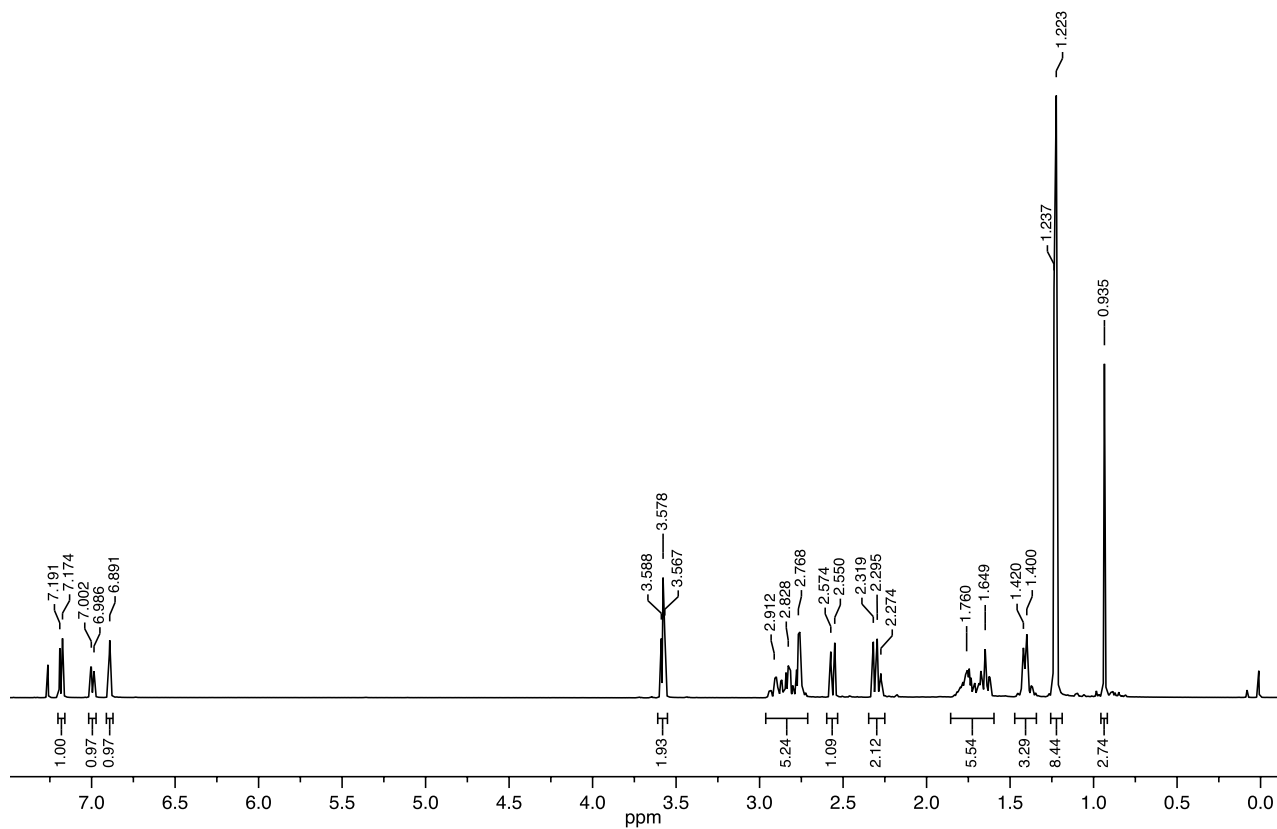
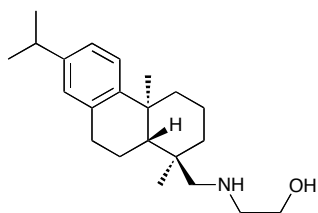


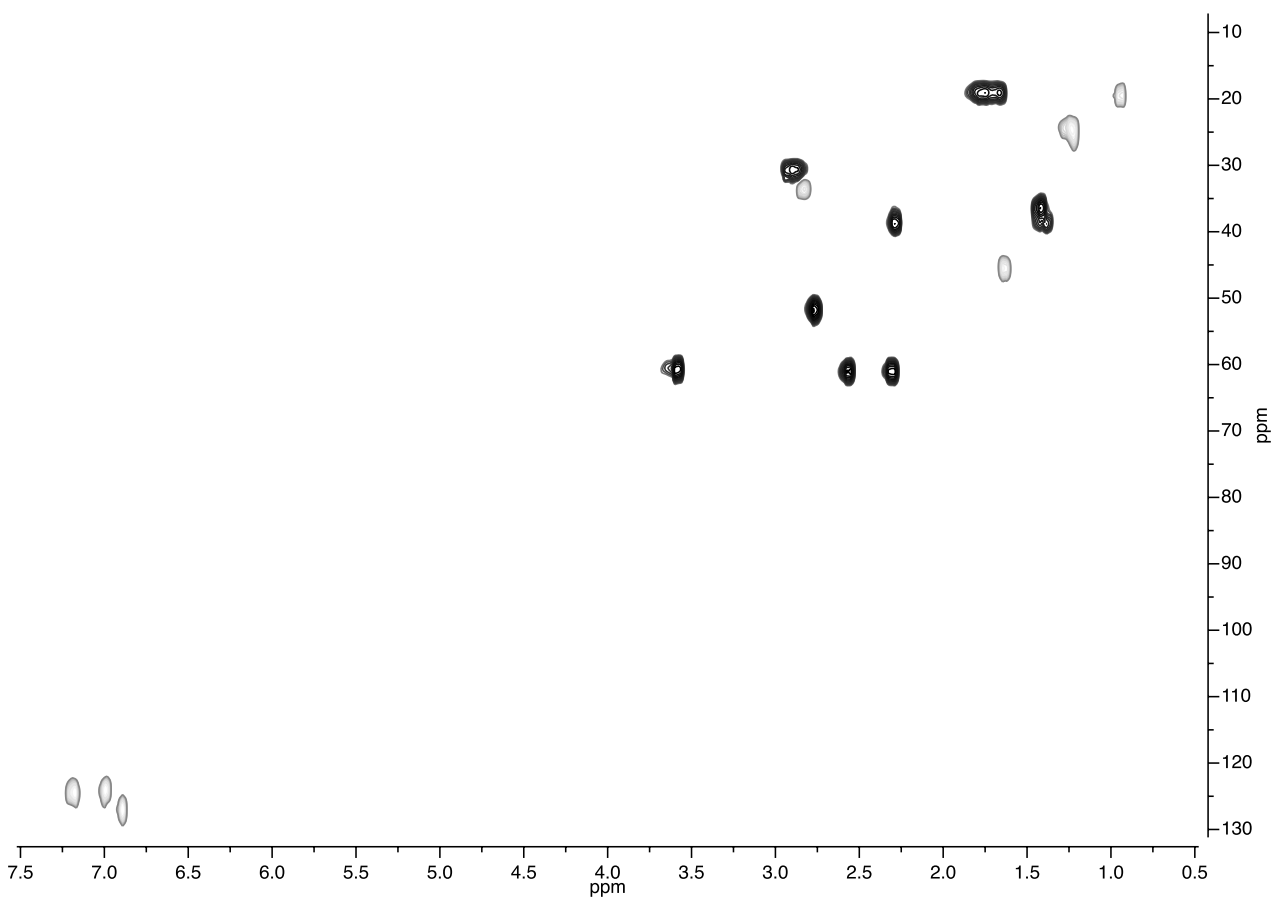
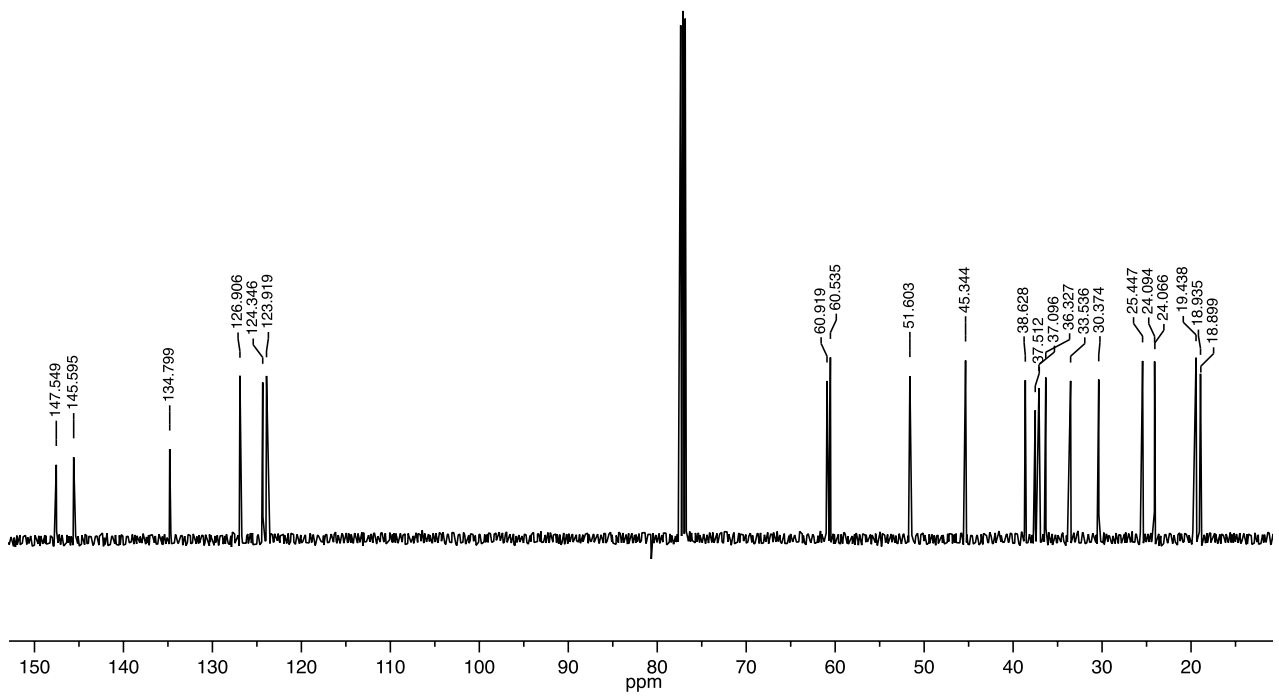
1H, 13C and HSQC spectra of compound 3b



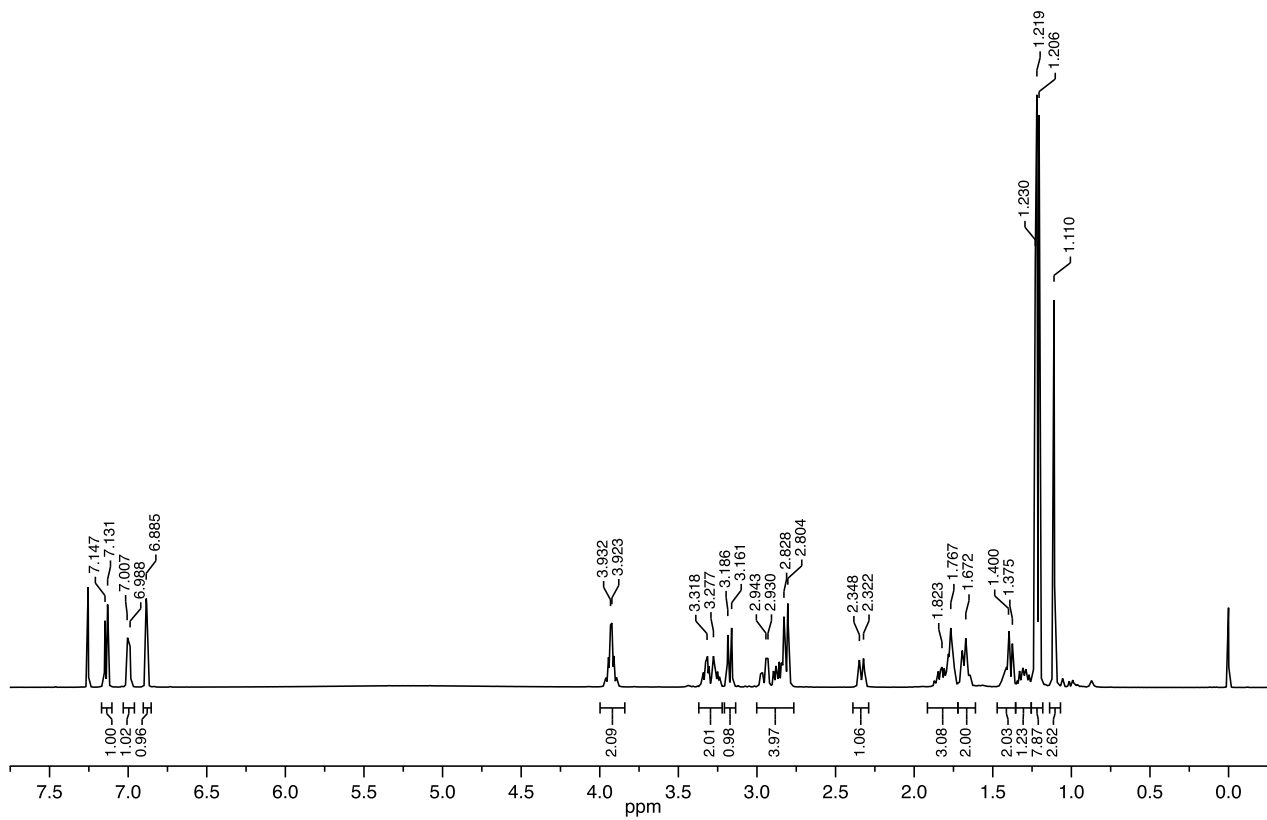
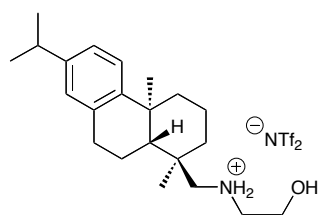


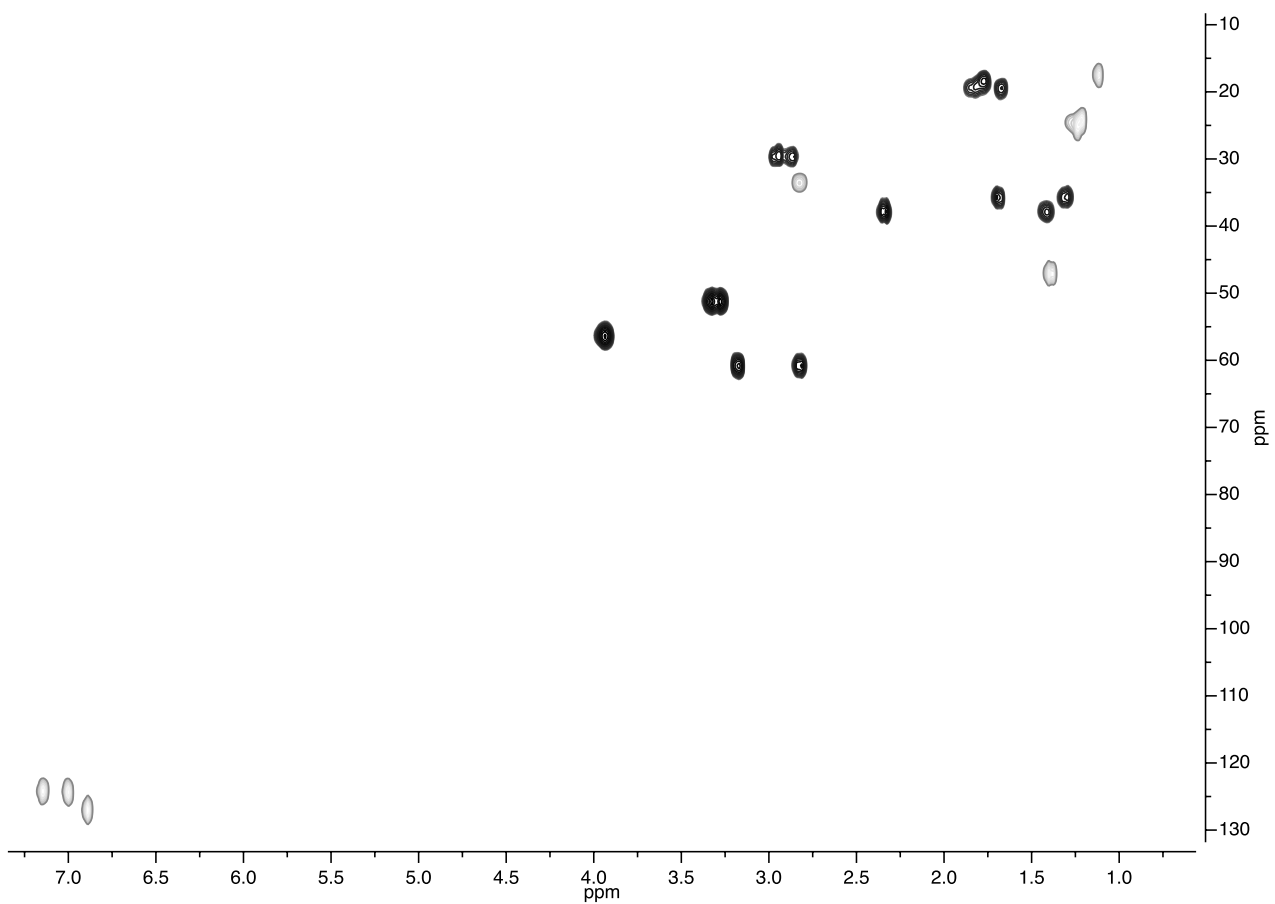
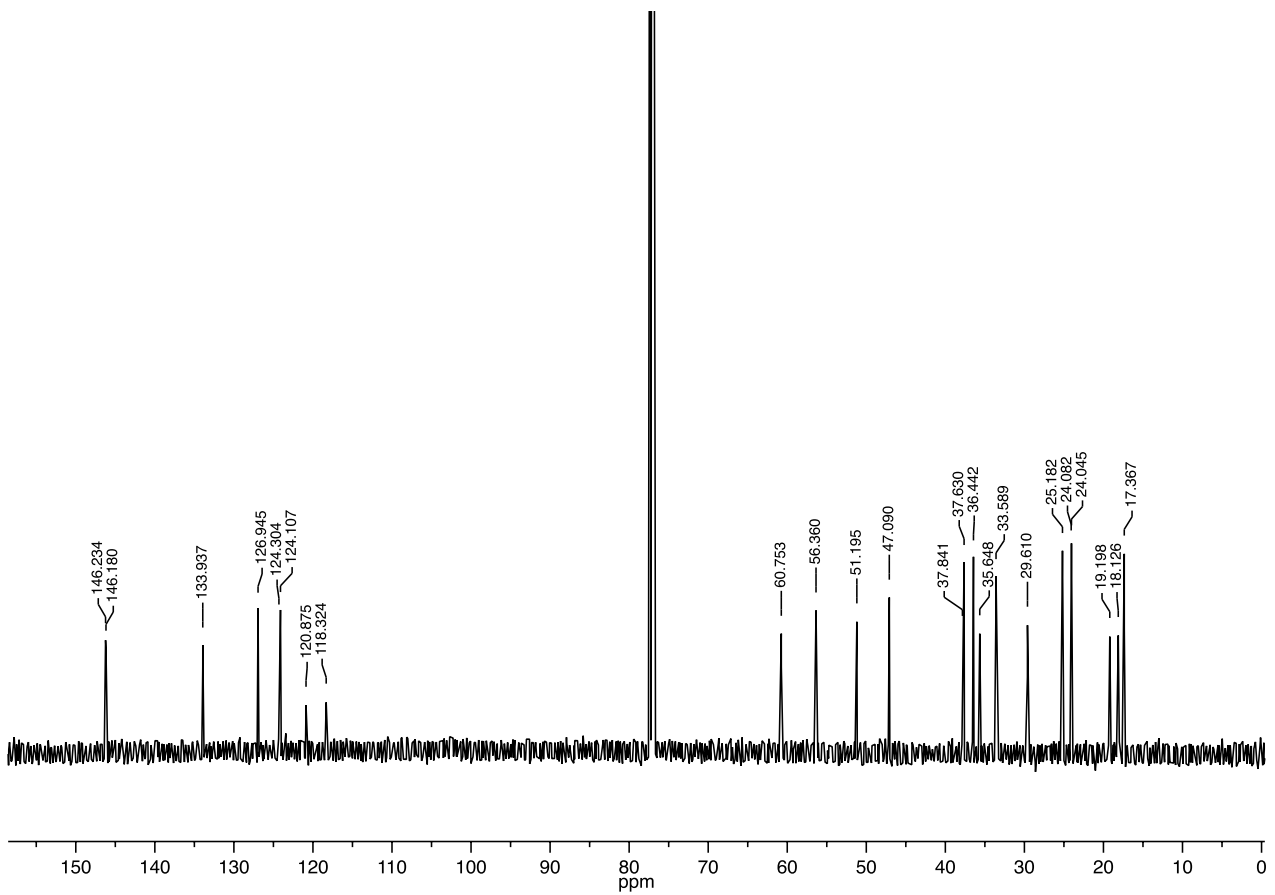
¹H, ¹³C and HSQC spectra of compound 4a



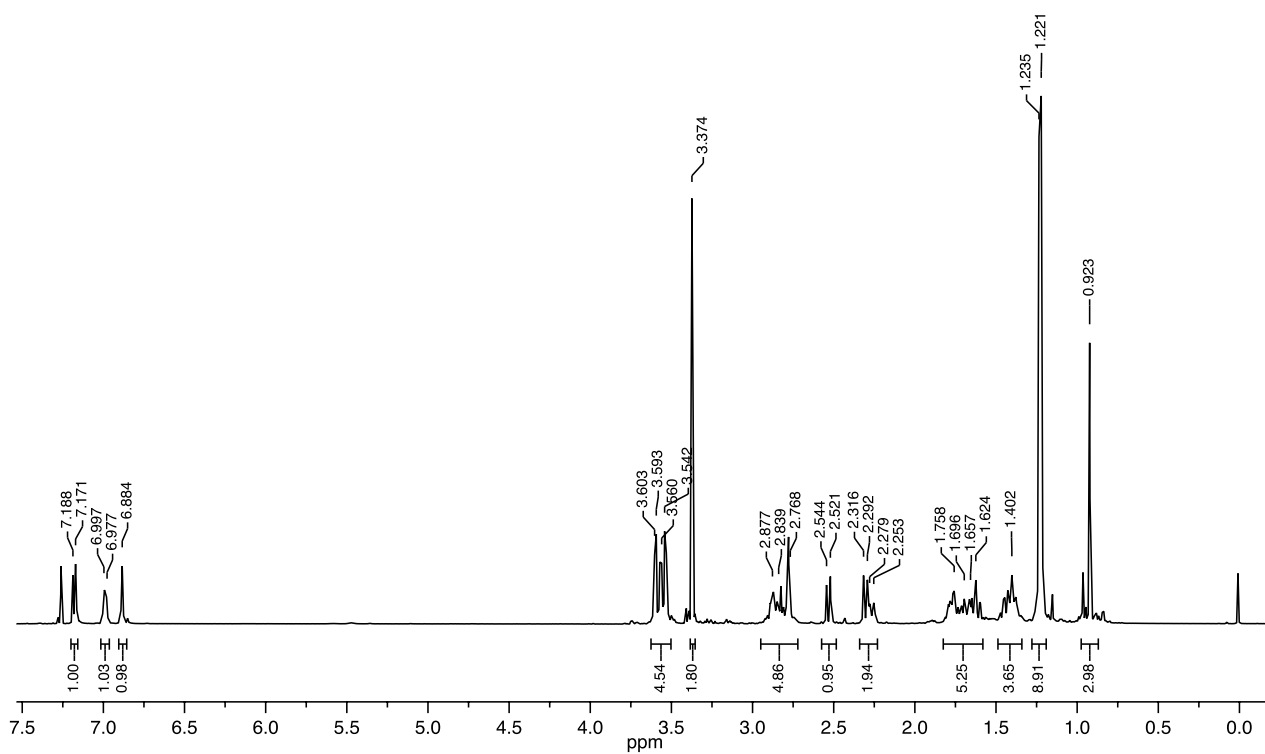
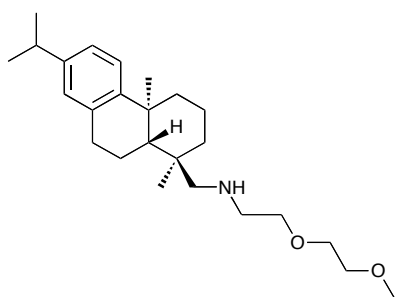


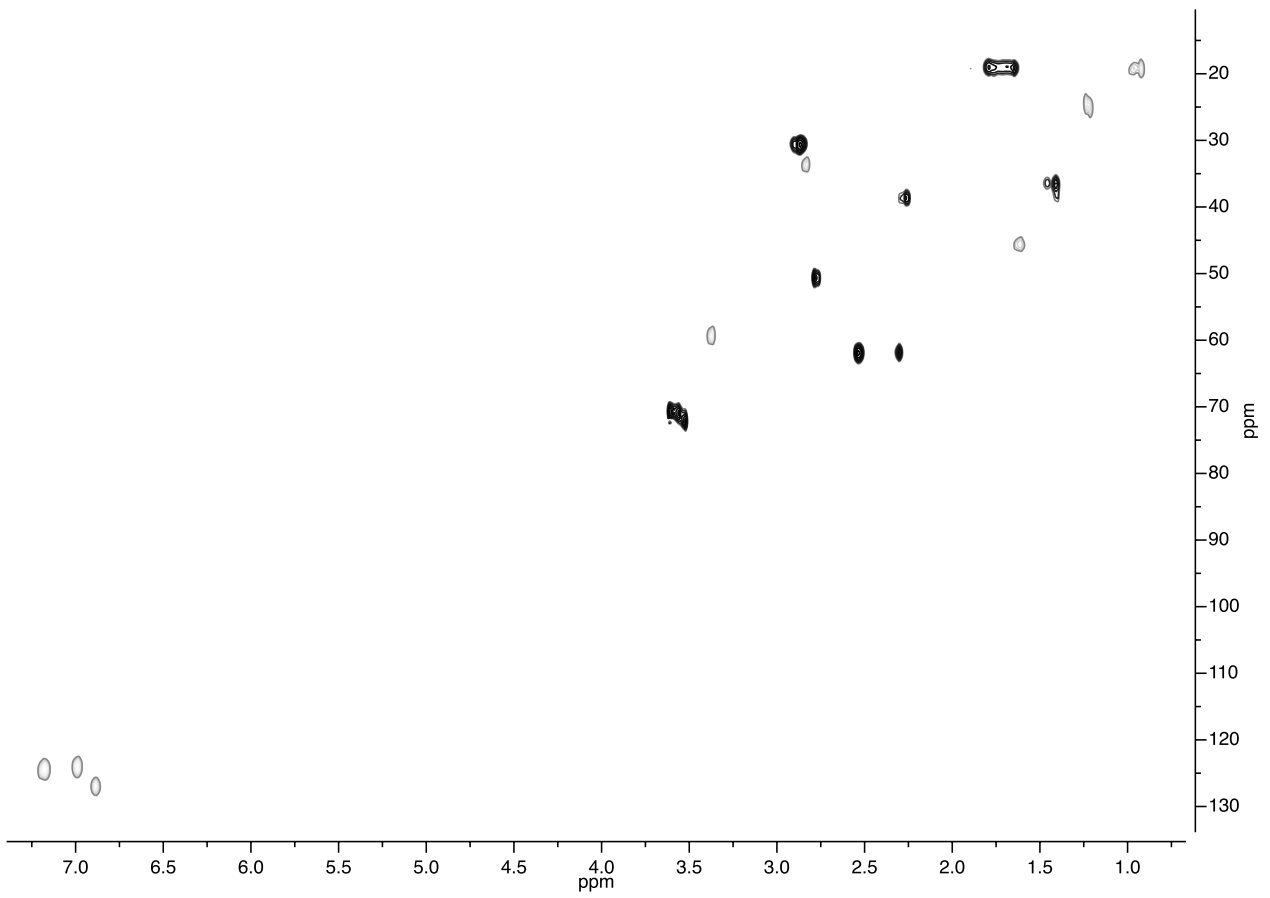
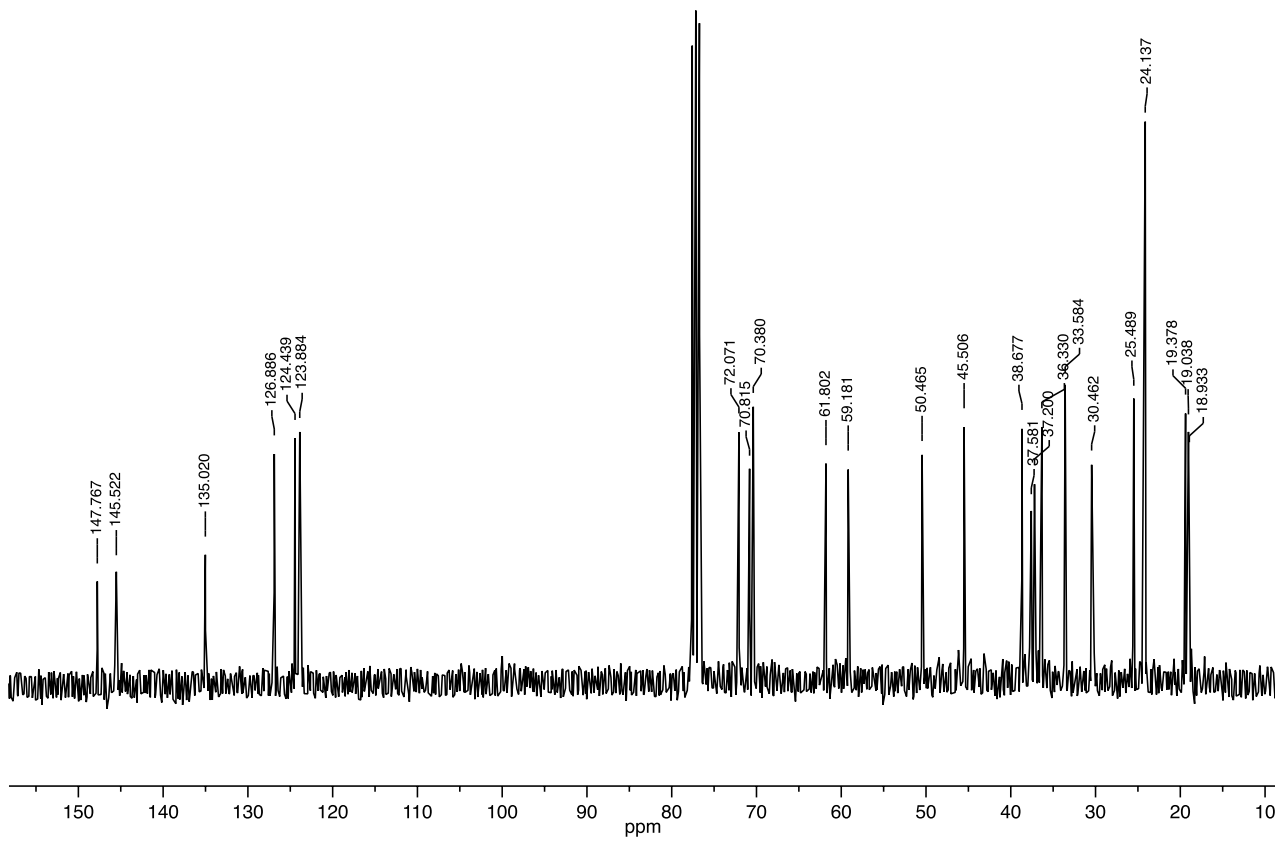
1H, 13C and HSQC spectra of compound 4b



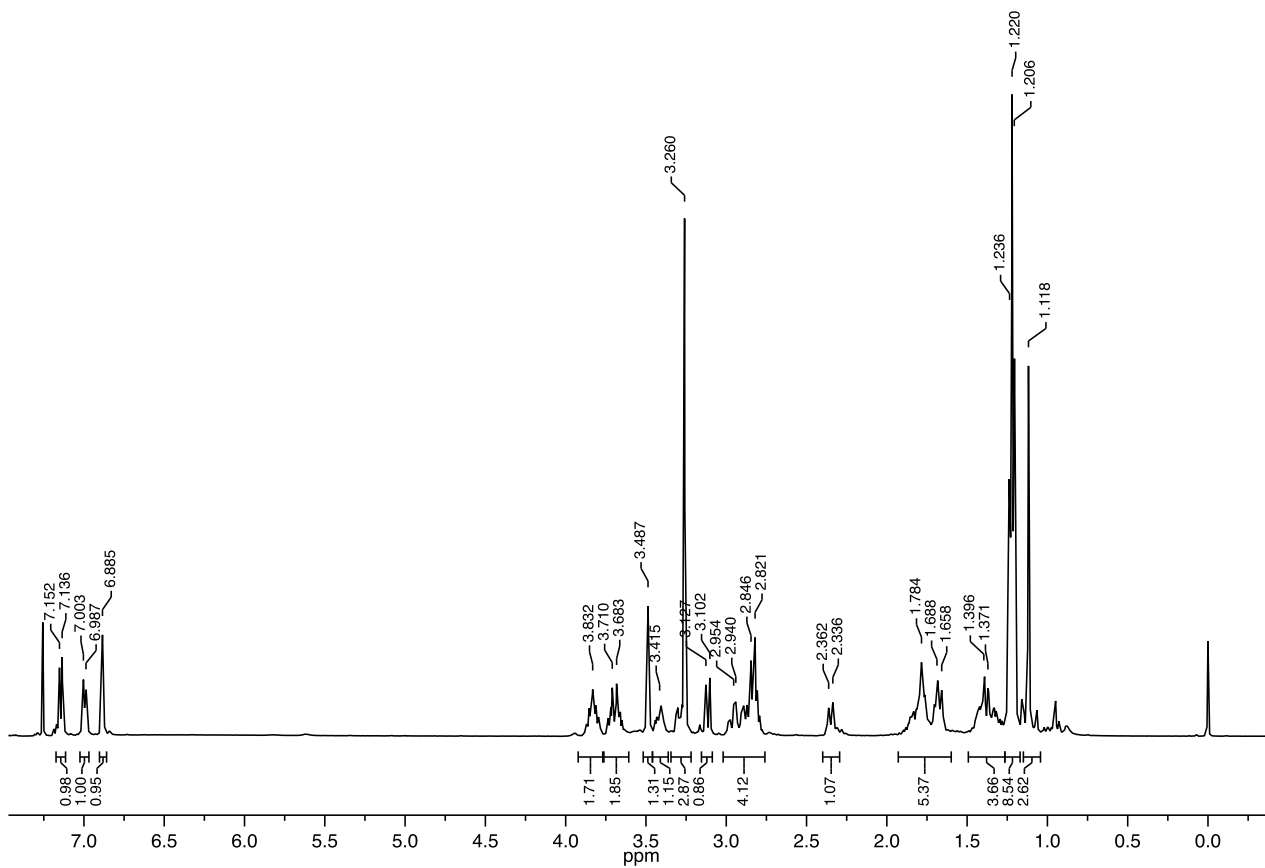
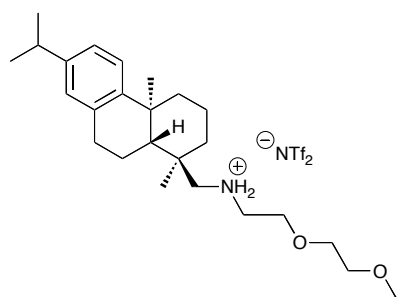


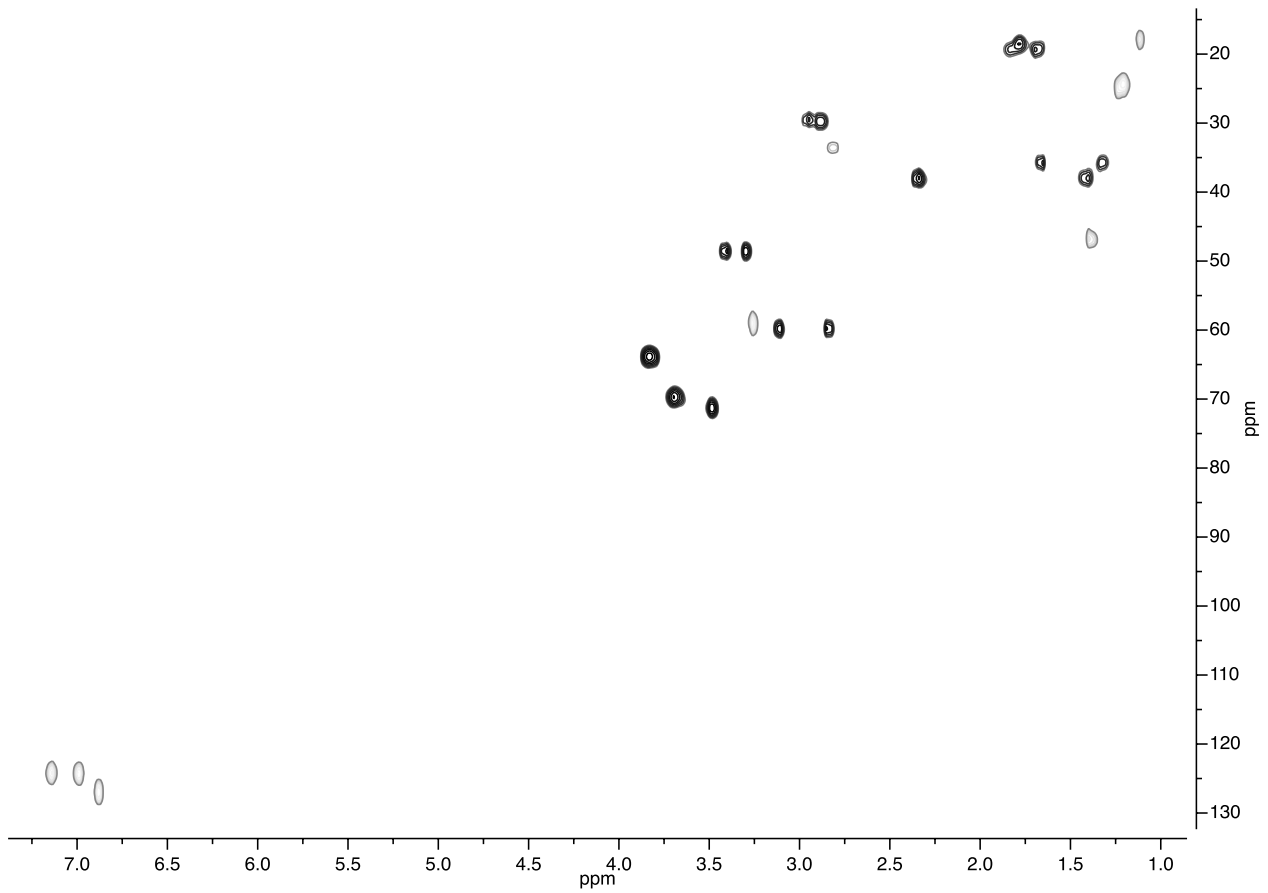
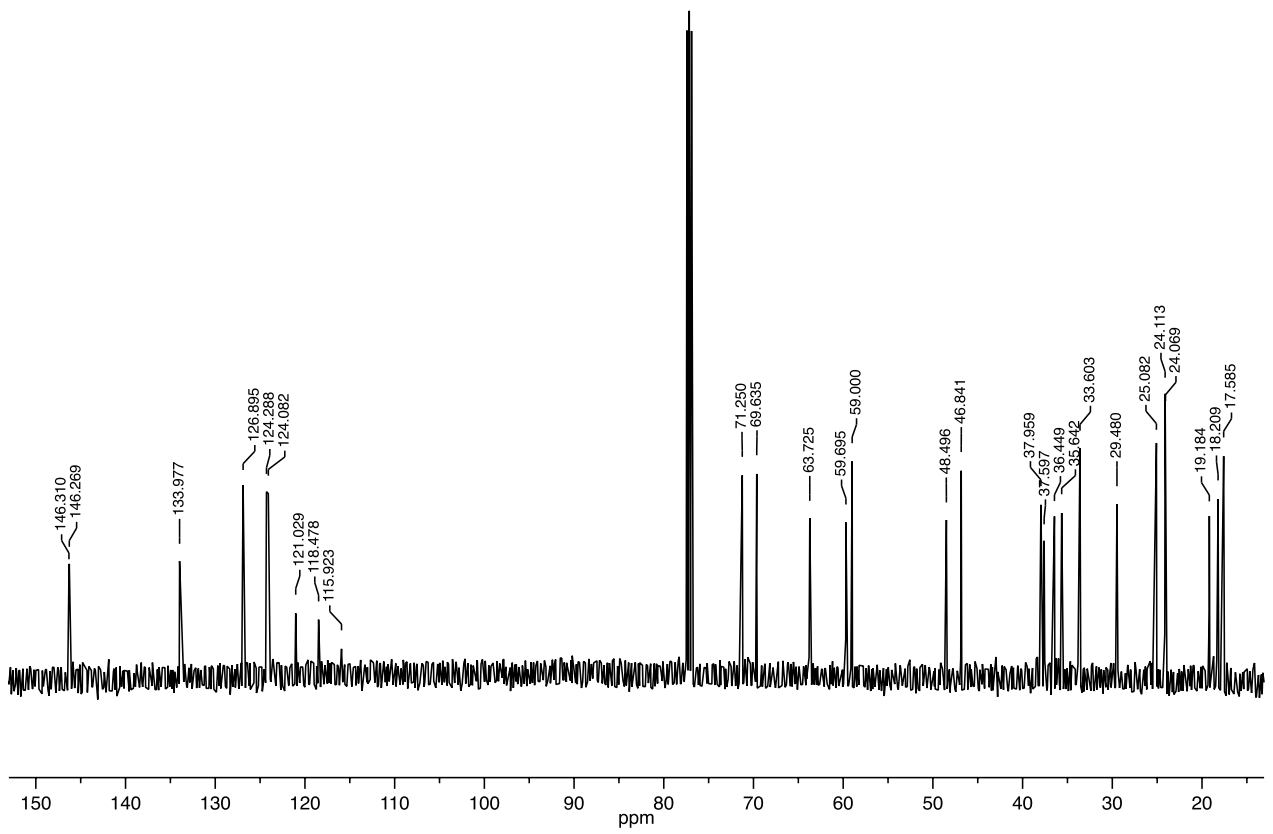
1H, 13C and HSQC spectra of compound 5a



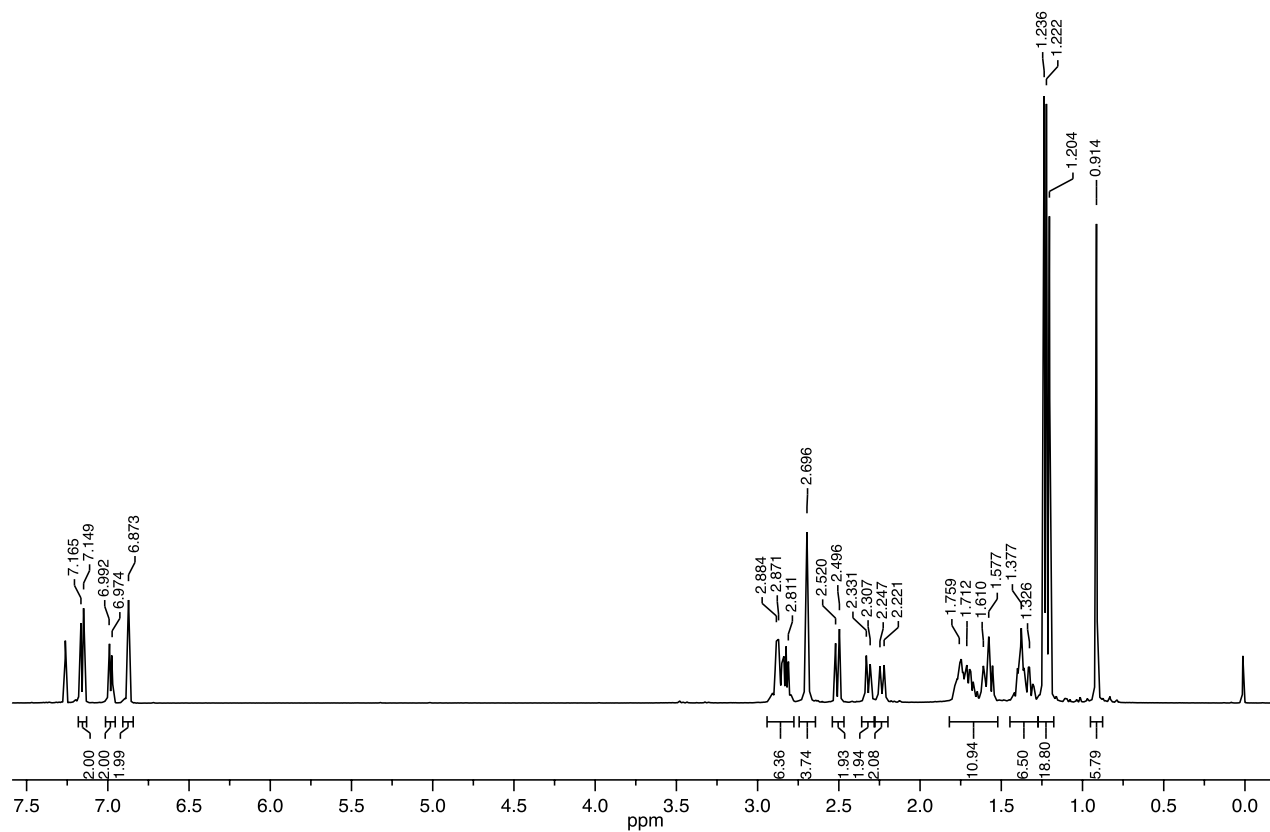
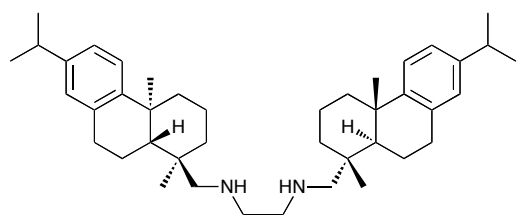


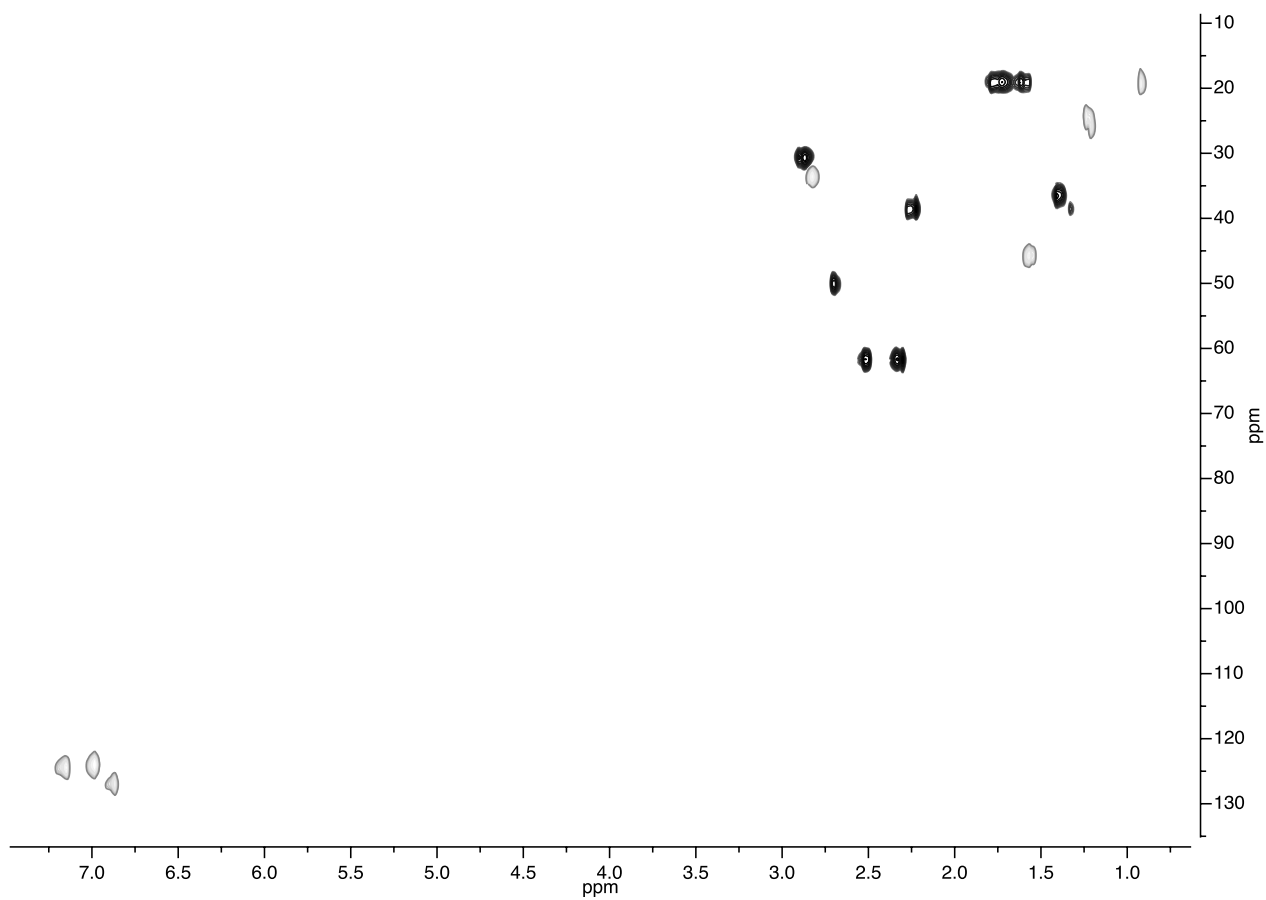
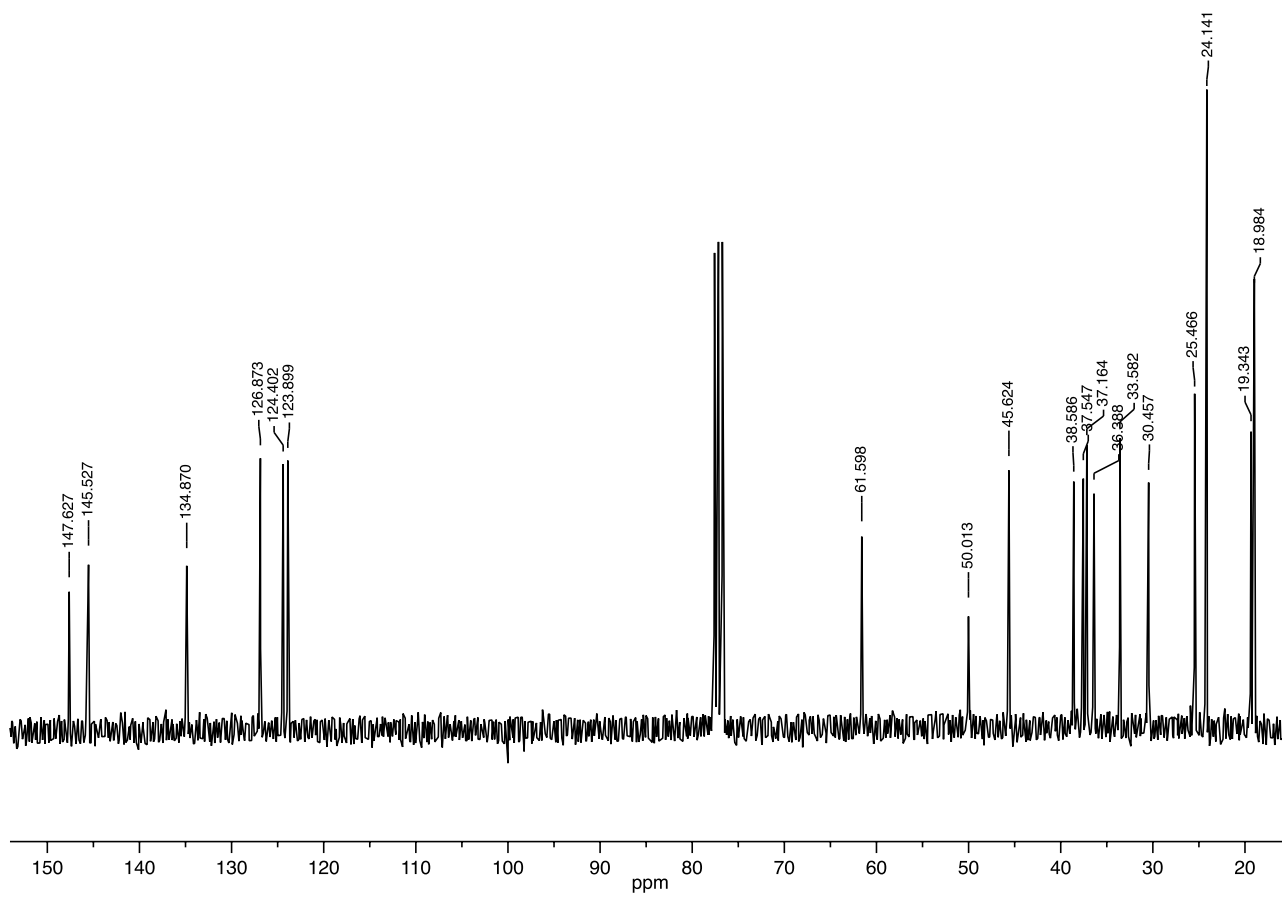
1H, 13C and HSQC spectra of compound 5b





¹H, ¹³C and HSQC spectra of compound 6a





1H, 13C and HSQC spectra of compound 6b

