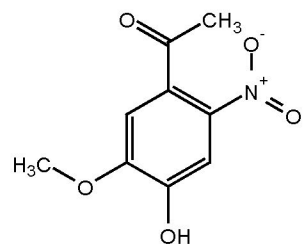
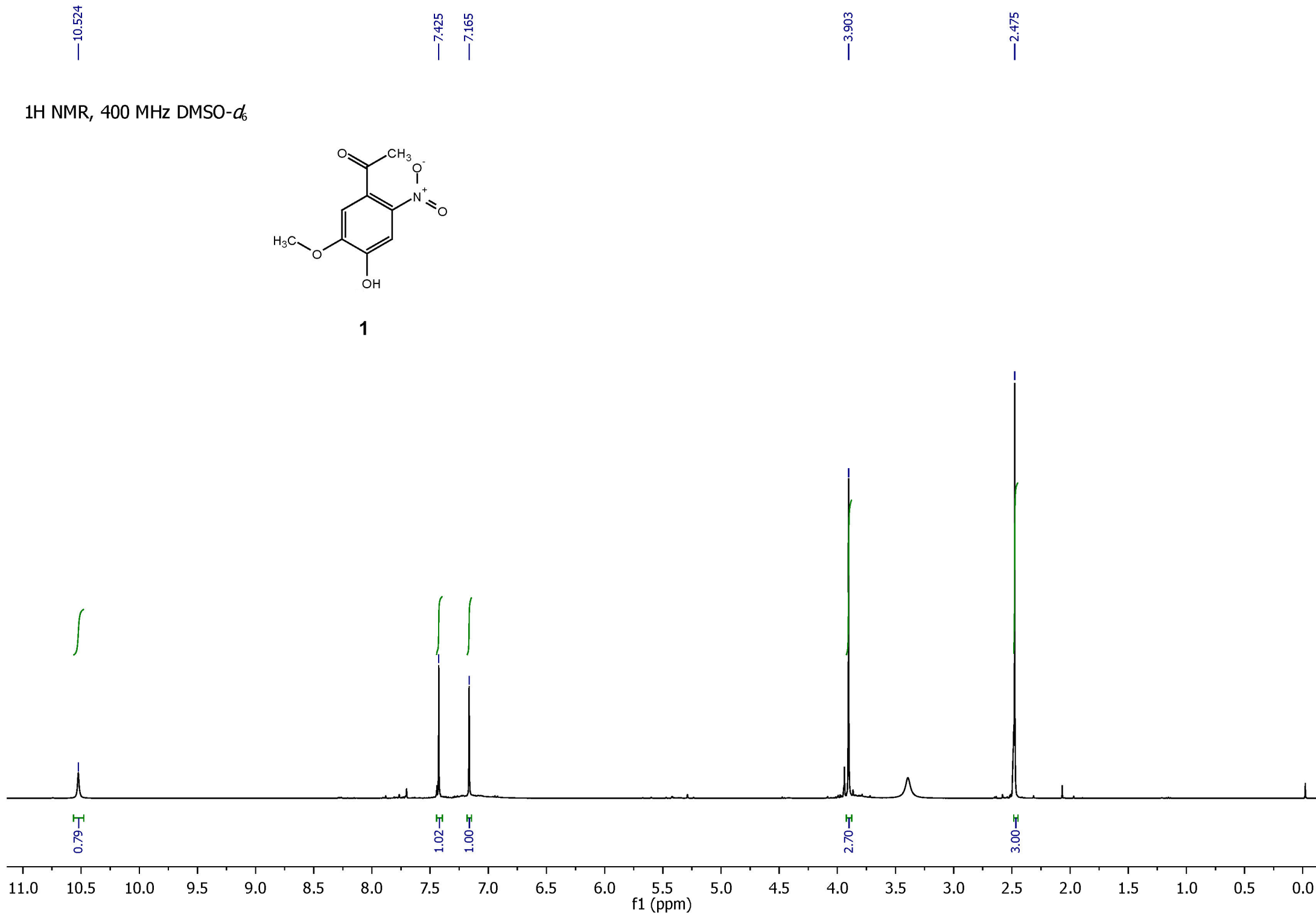


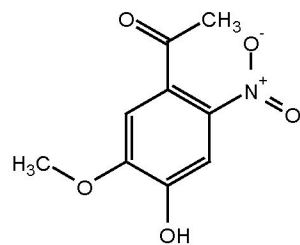
¹H NMR, 400 MHz DMSO-*d*₆



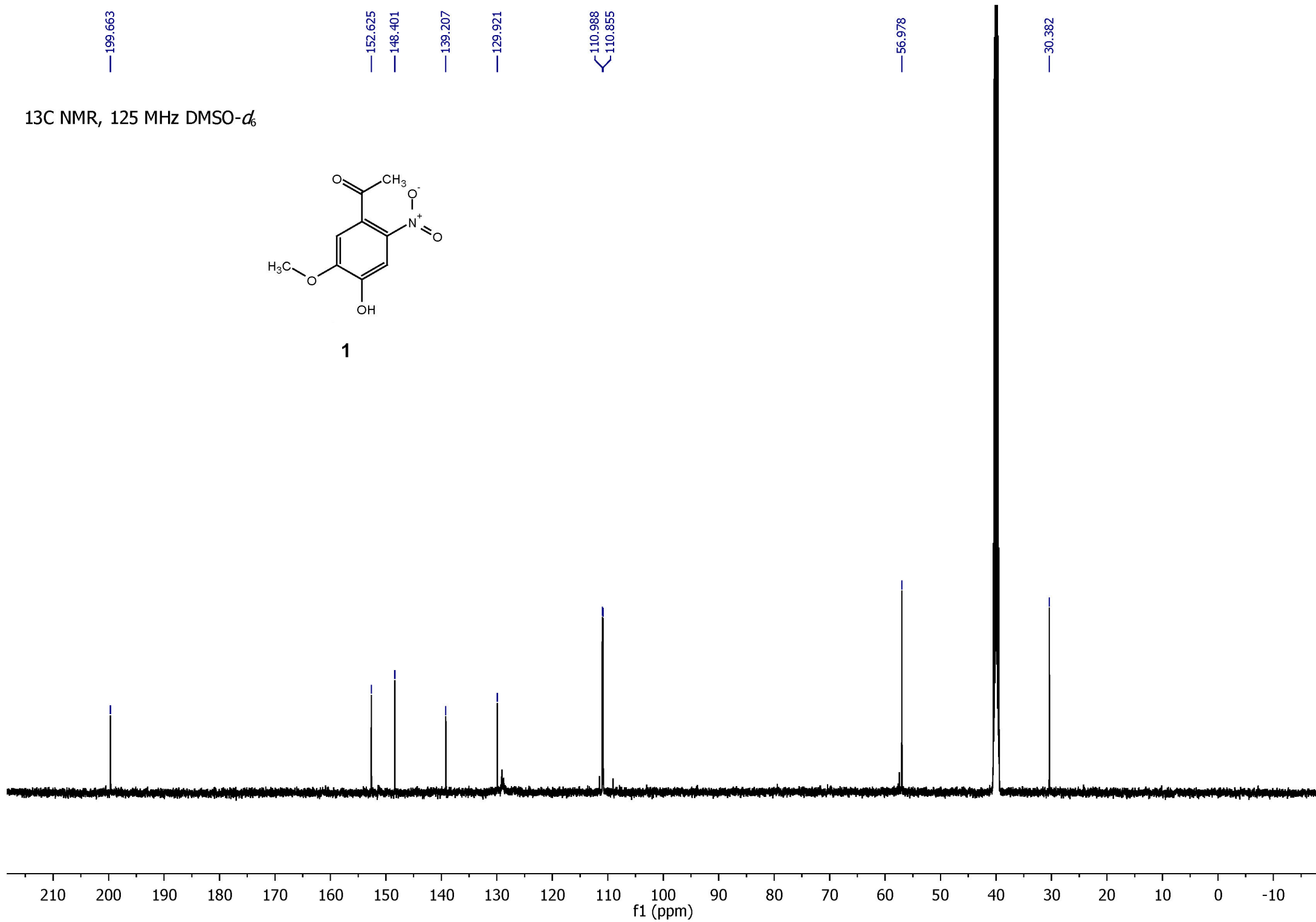
1



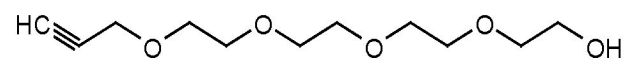
¹³C NMR, 125 MHz DMSO-*d*₆



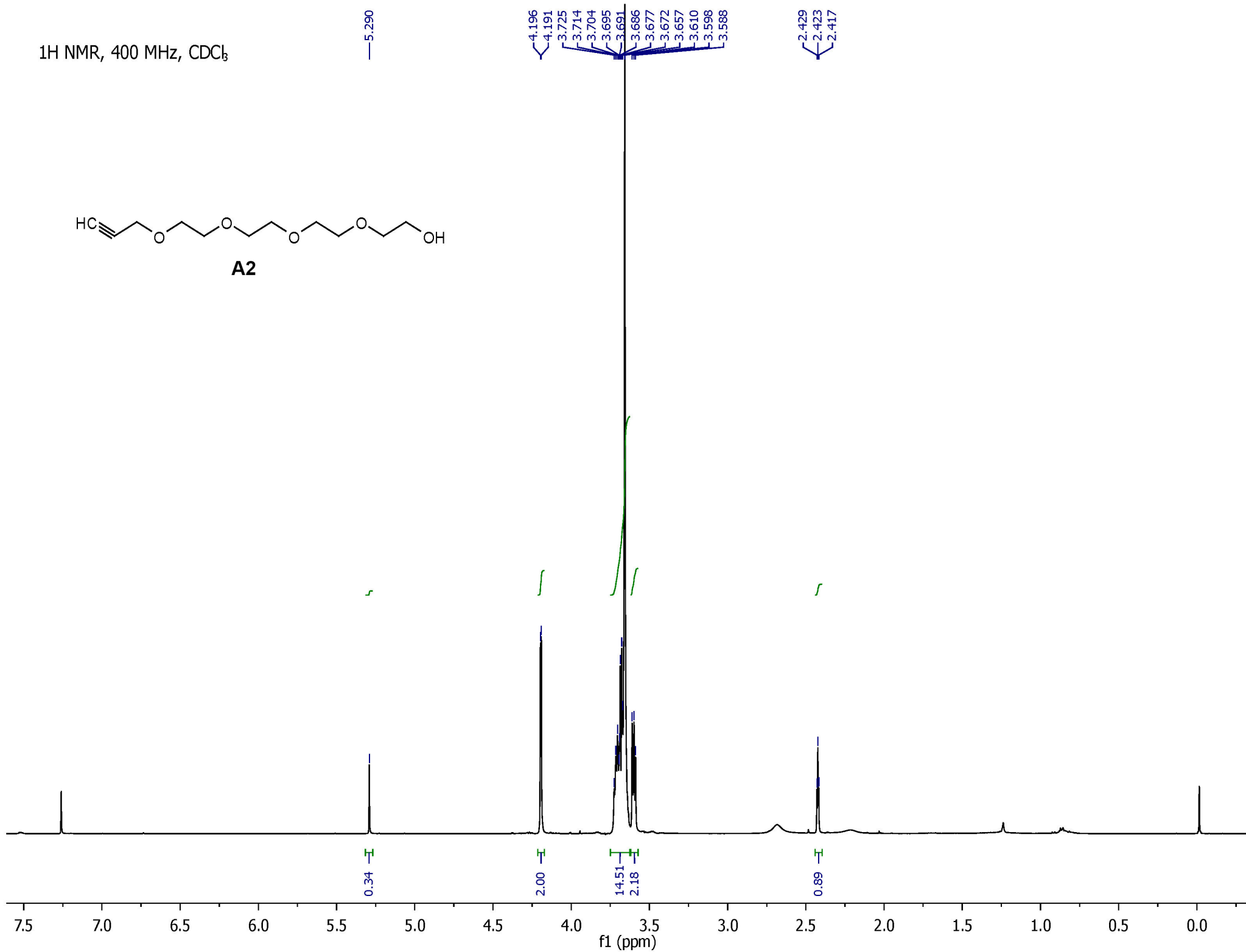
1



¹H NMR, 400 MHz, CDCl₃

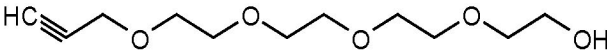


A2

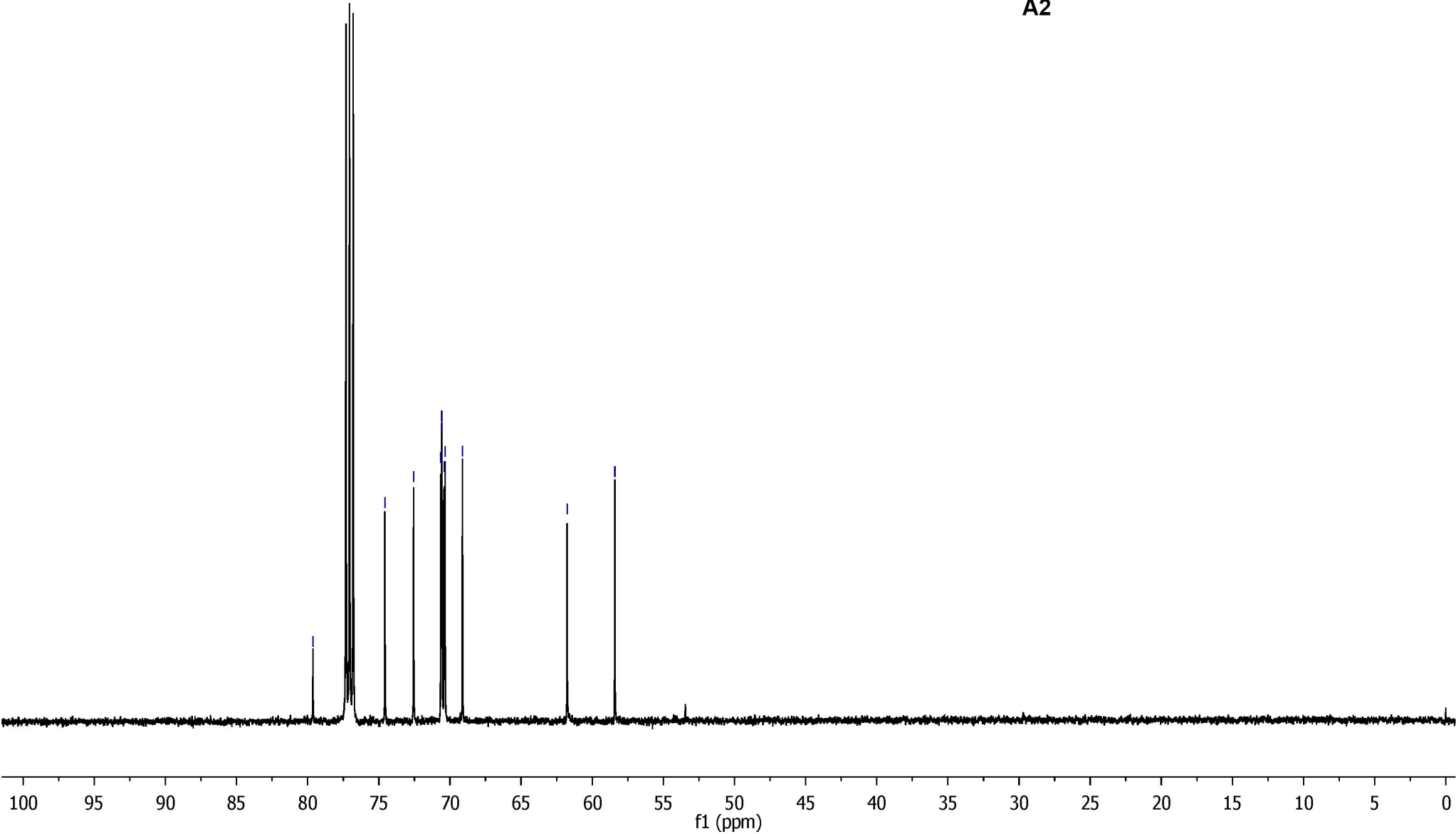


¹³C NMR, 125 MHz, CDCl₃

79.624
74.568
72.549
70.636
70.583
70.554
70.406
70.339
69.118
61.763
58.409



A2



7.338
7.317

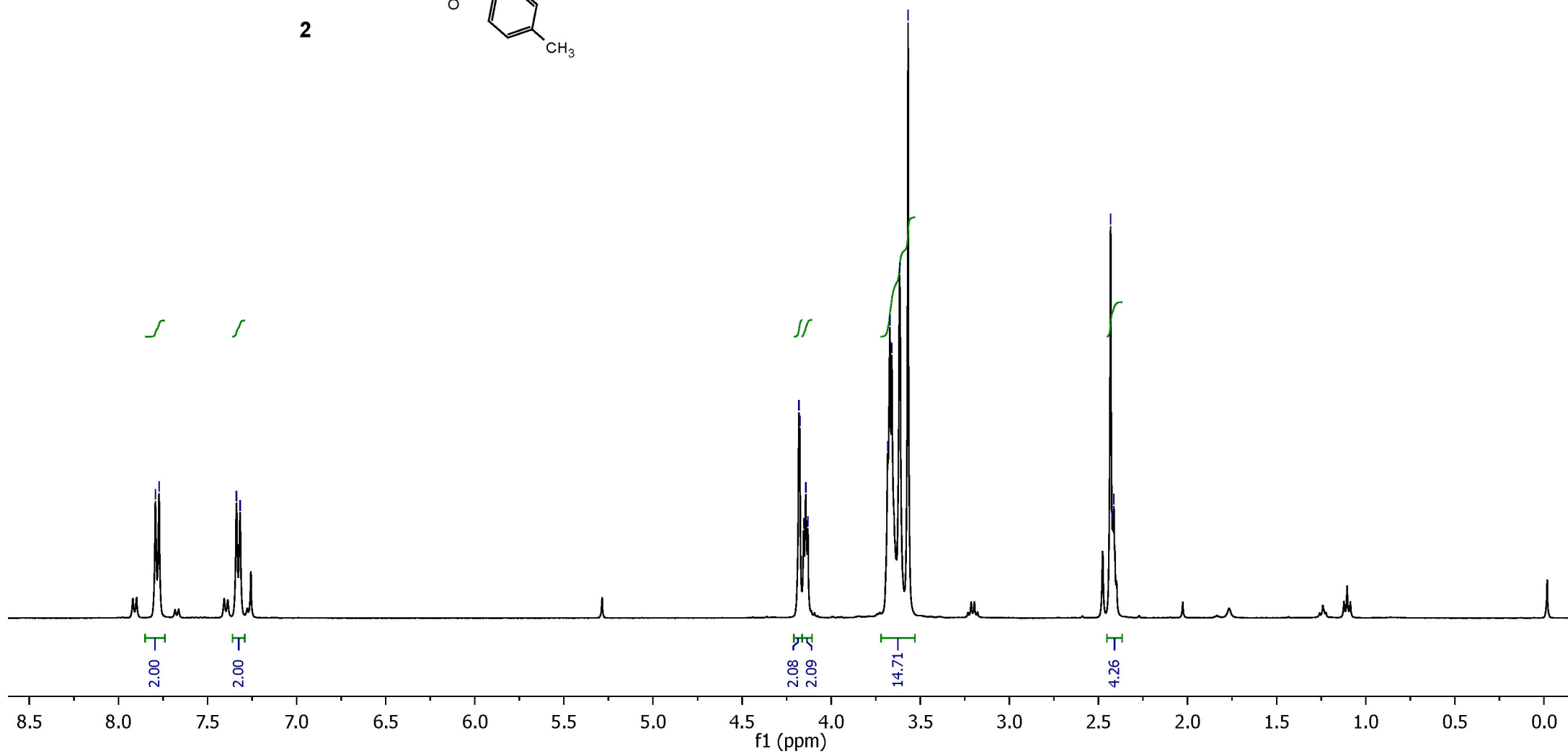
Figure 1 displays two line graphs showing the evolution of the average number of nodes per cluster (N_c) over time (t). The x-axis represents time t from 0 to 100, and the y-axis represents N_c from 0 to 5. The left graph (blue lines) shows N_c increasing from approximately 1.5 to 4.1. The right graph (red lines) shows N_c increasing from approximately 1.5 to 3.6. Both models show a sharp increase in N_c around $t=20$, followed by a plateau.

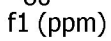
Time (t)	Left Model (N_c)	Right Model (N_c)
0	1.5	1.5
20	4.182	3.683
40	4.176	3.671
60	4.155	3.660
80	4.143	3.616
100	4.131	3.569

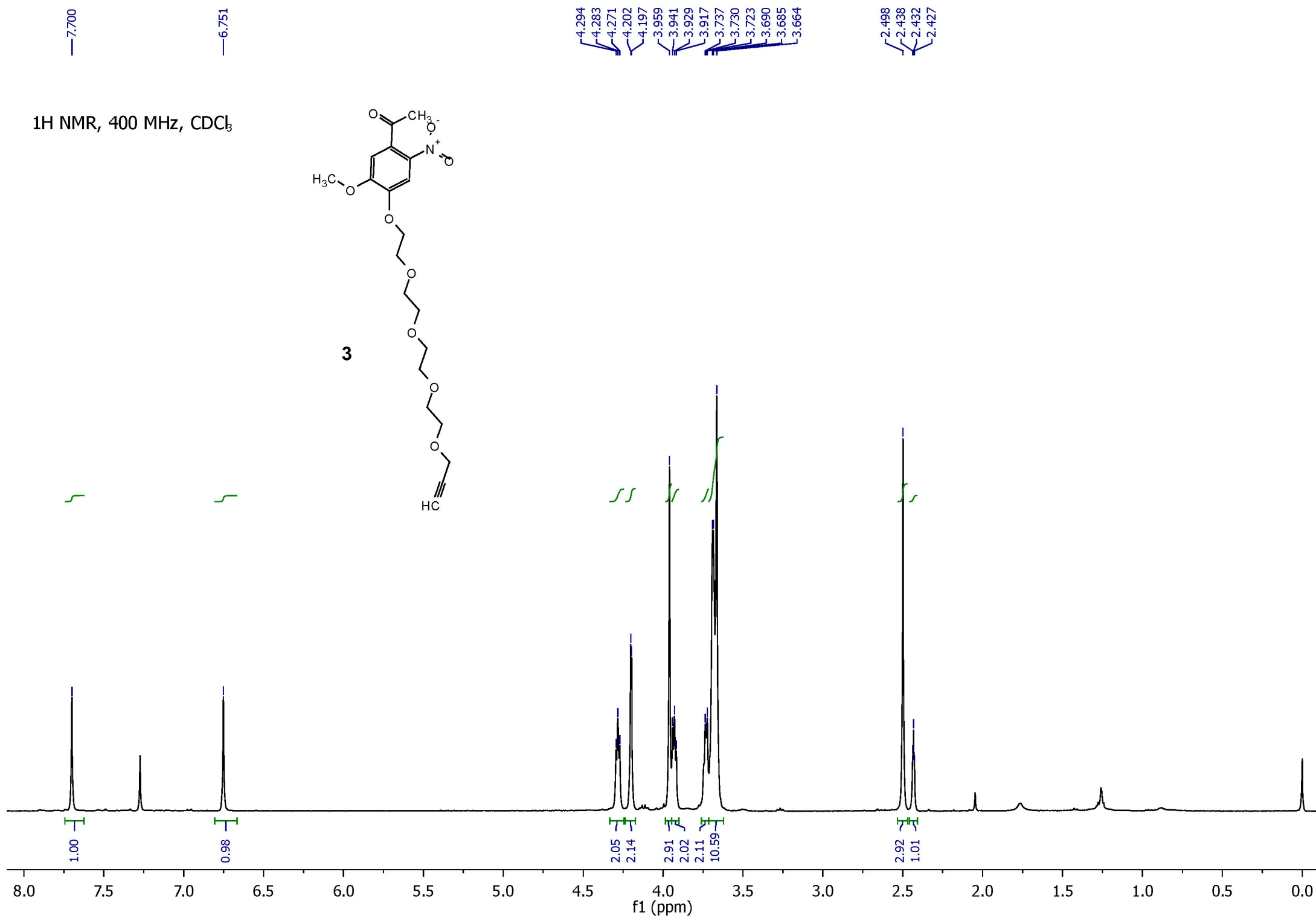
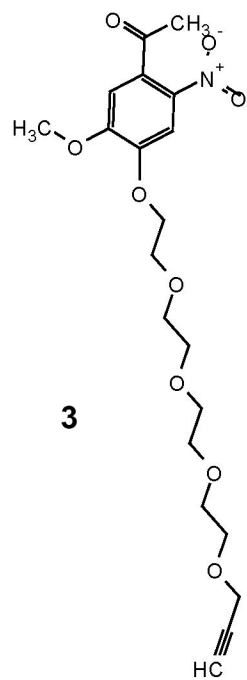
2.432
2.419
2.414

C#CCOCCOCCOCCOCCOS(=O)(=O)c1ccc(C)cc1

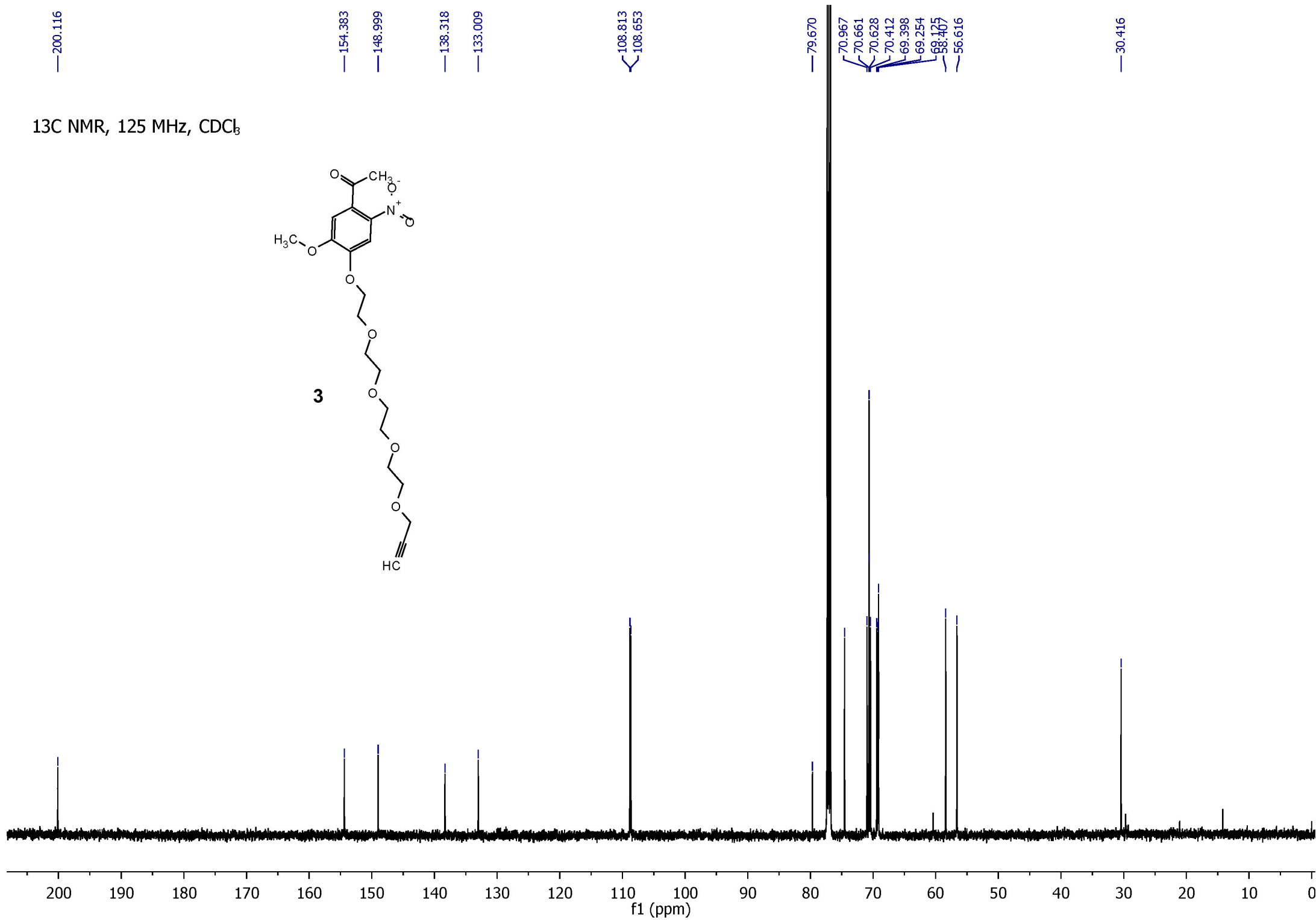
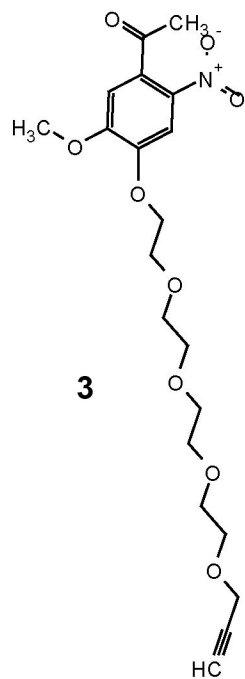
2

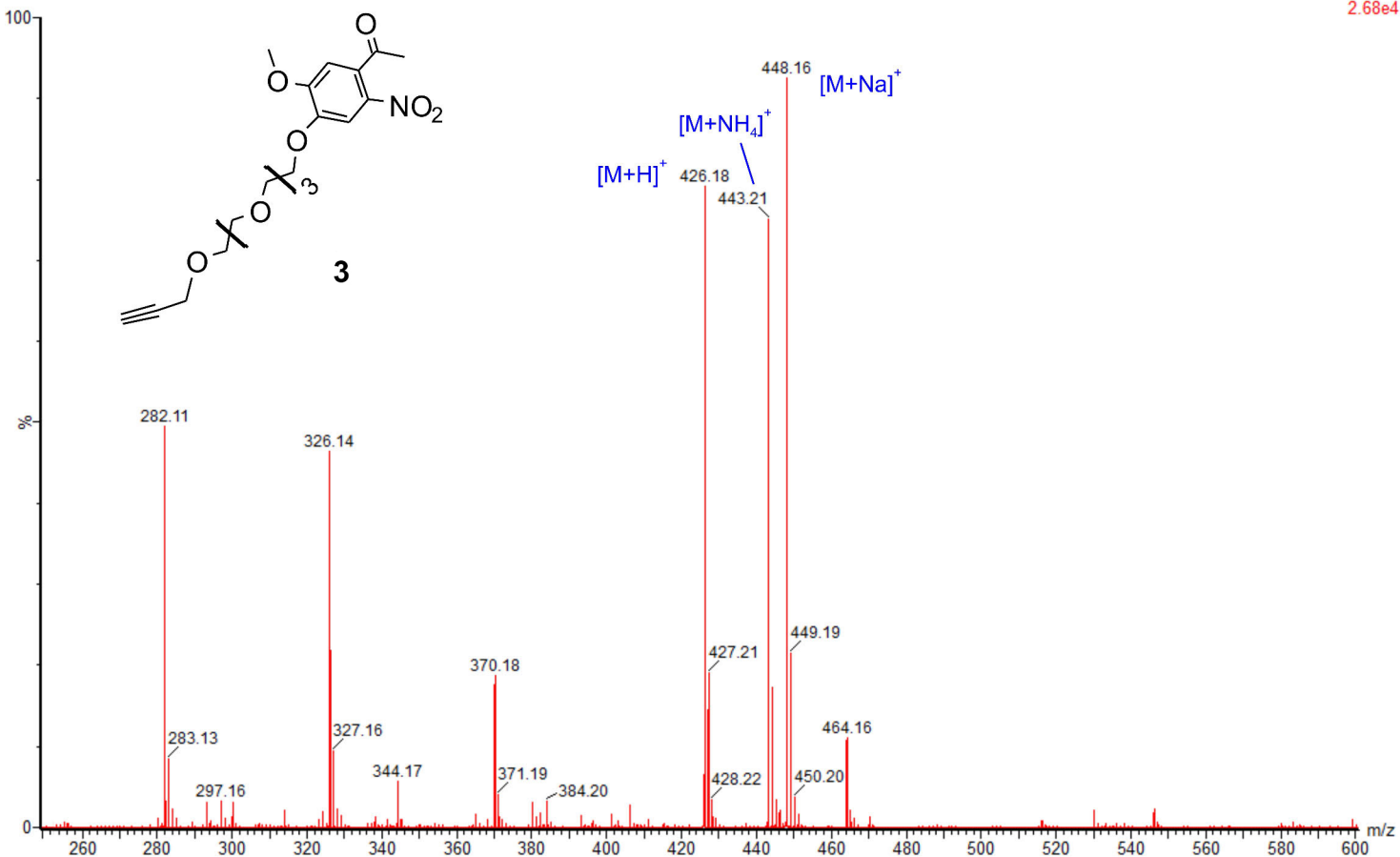
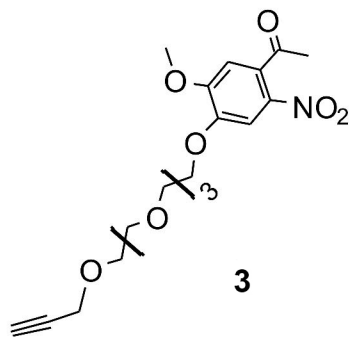




^1H NMR, 400 MHz, CDCl_3 

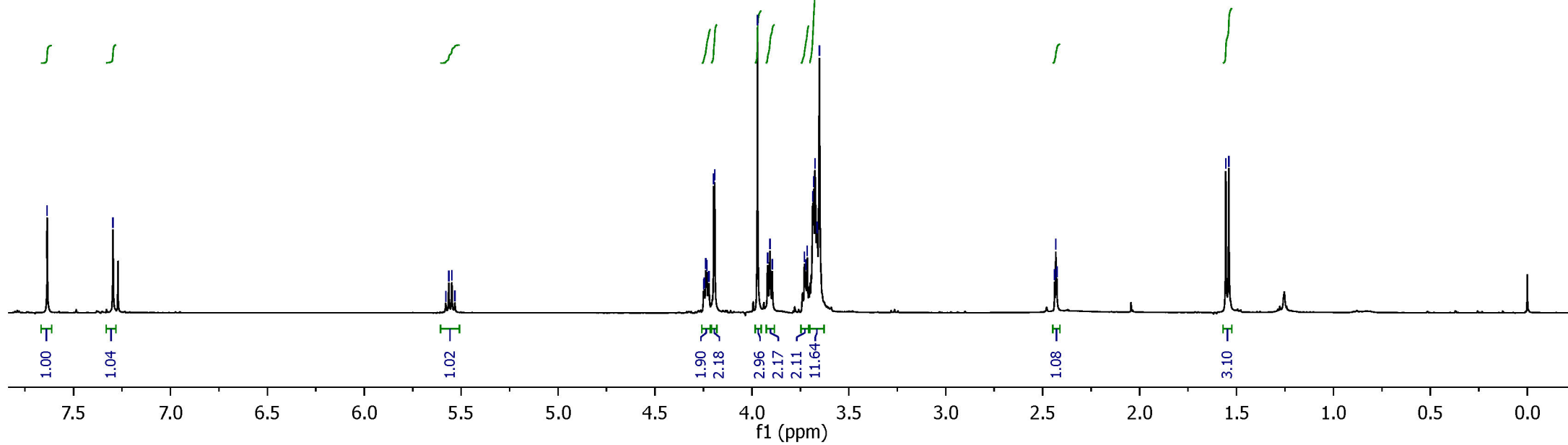
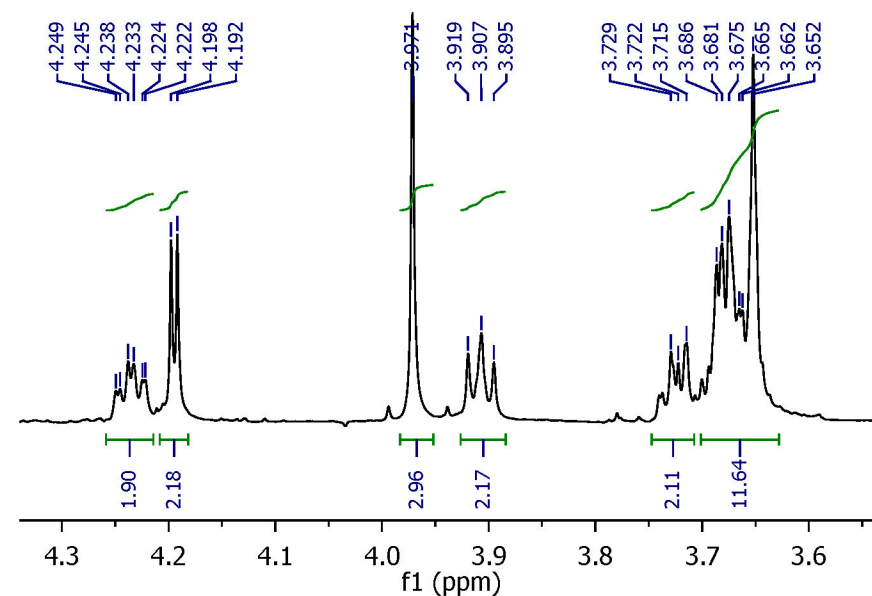
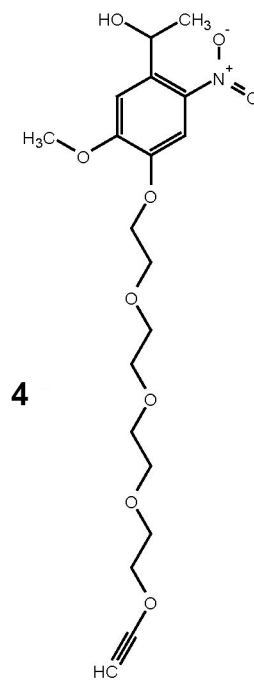
¹³C NMR, 125 MHz, CDCl₃



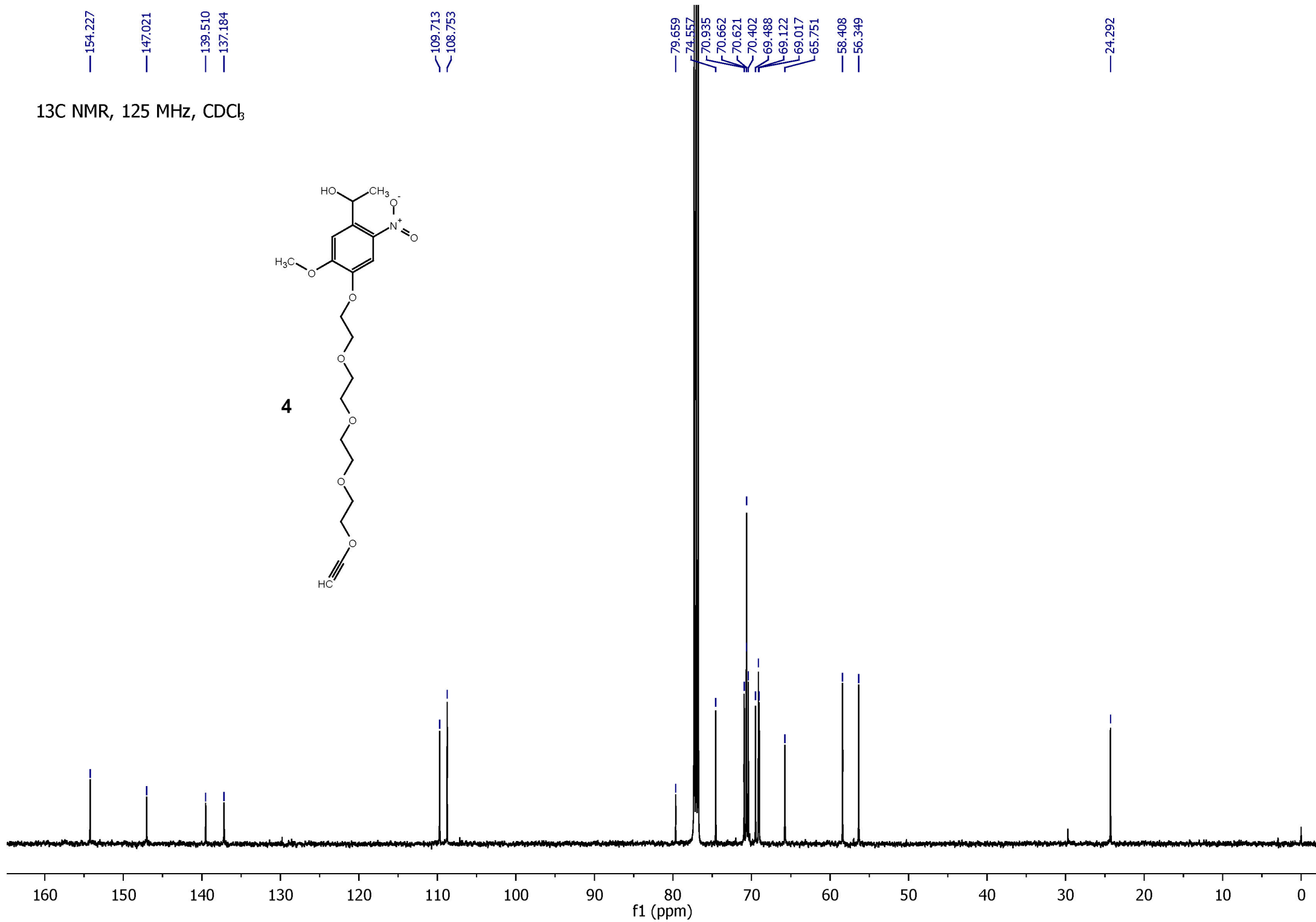
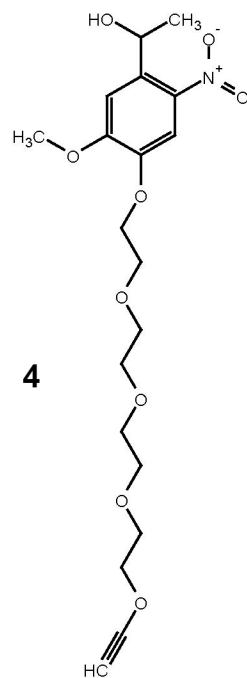


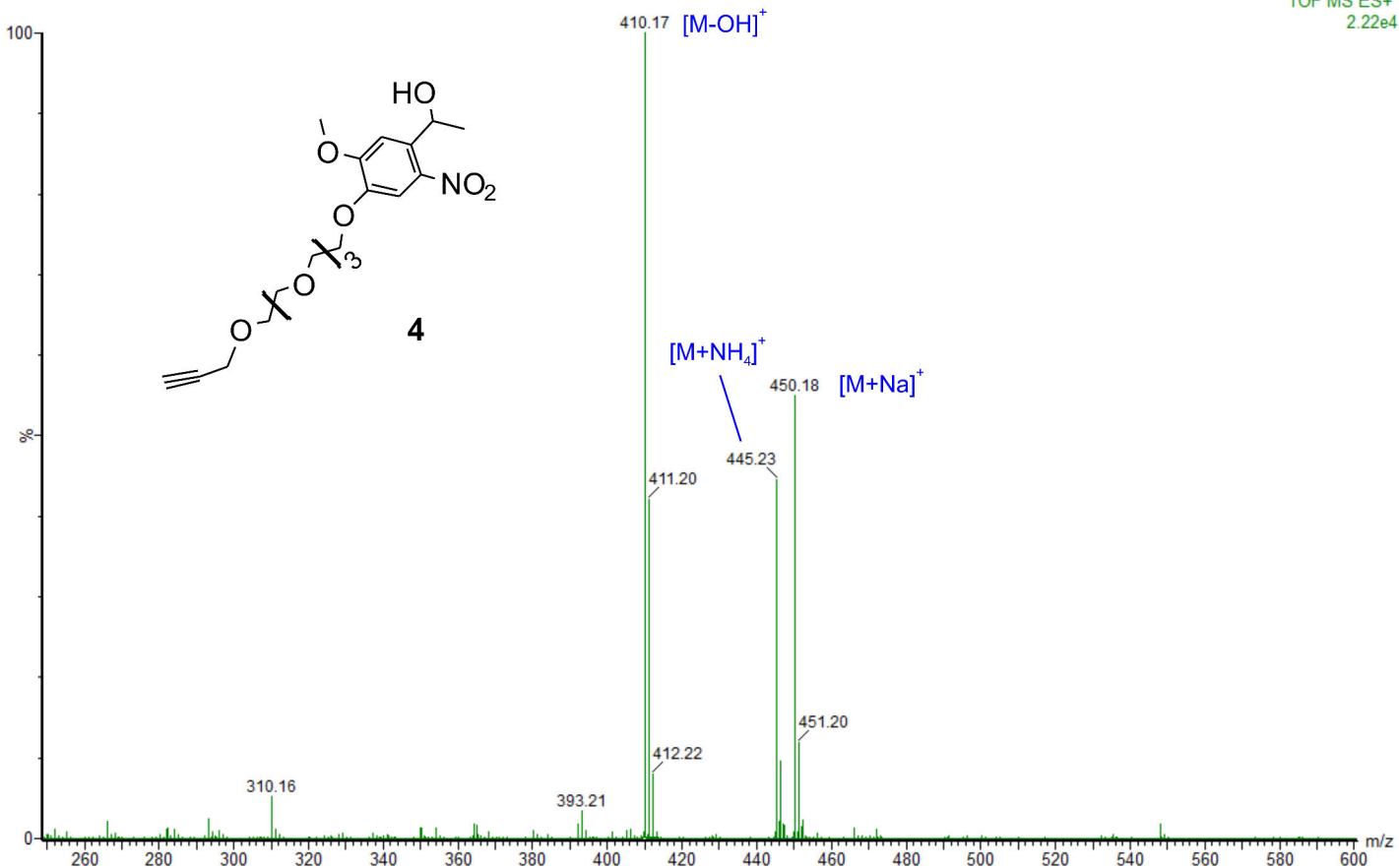
—7.636

—7.296

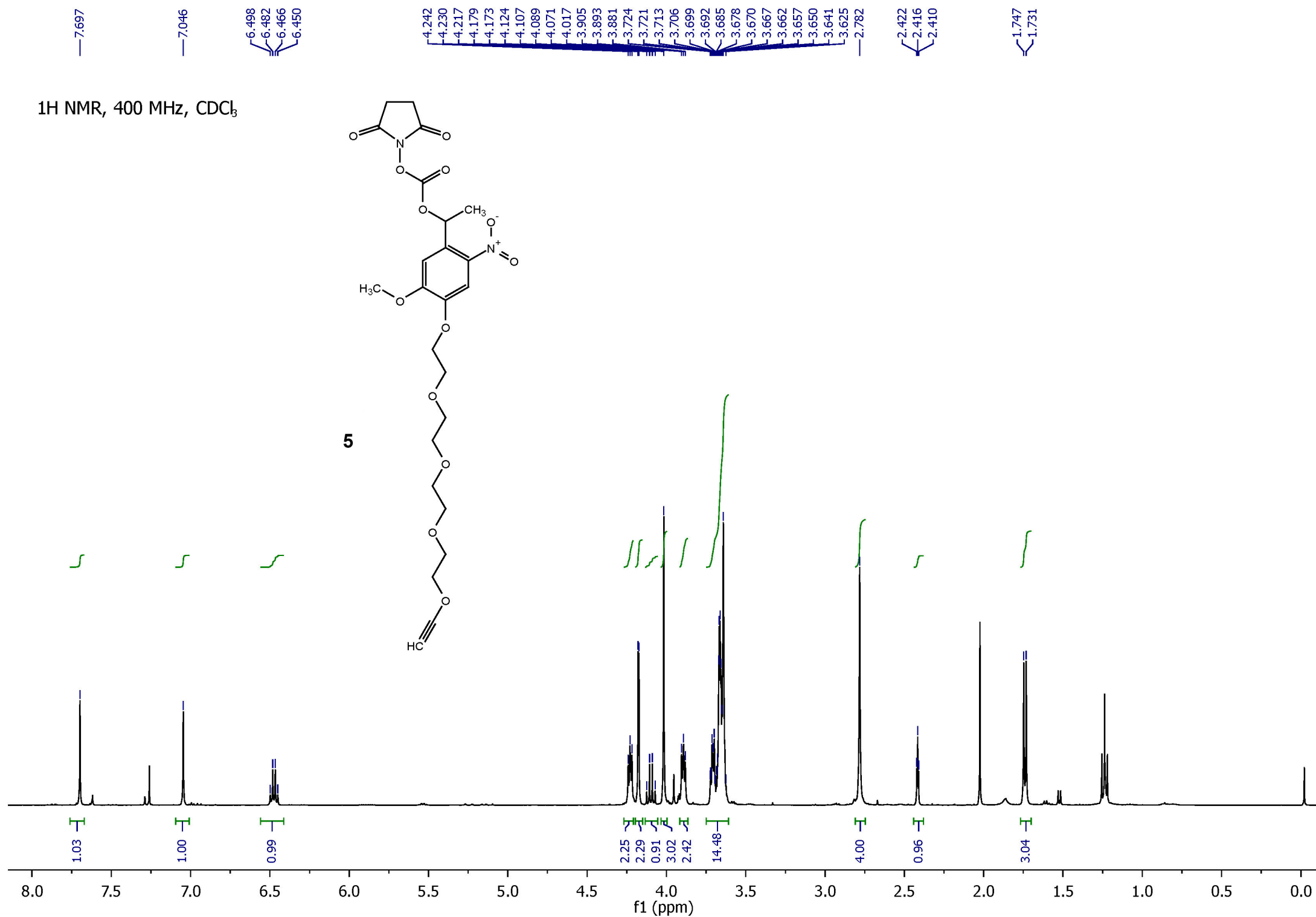
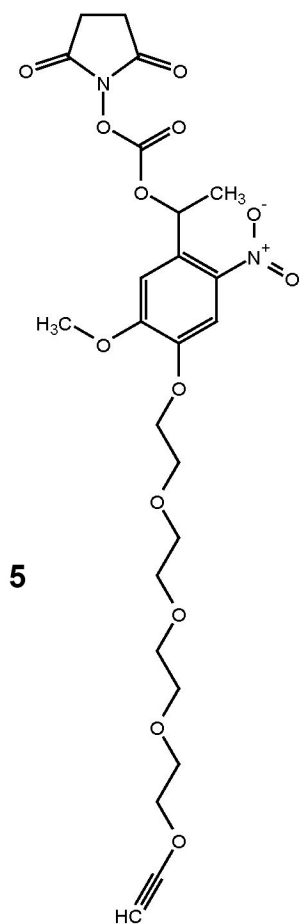
5.580
5.564
5.548
5.5334.249
4.245
4.238
4.233
4.224
4.222
4.198
4.192
3.971
3.919
3.907
3.895
3.729
3.722
3.715
3.686
3.681
3.675
3.665
3.662
3.6522.438
2.432
2.4261.555
1.5401H NMR, 400 MHz, CDCl₃

¹³C NMR, 125 MHz, CDCl₃

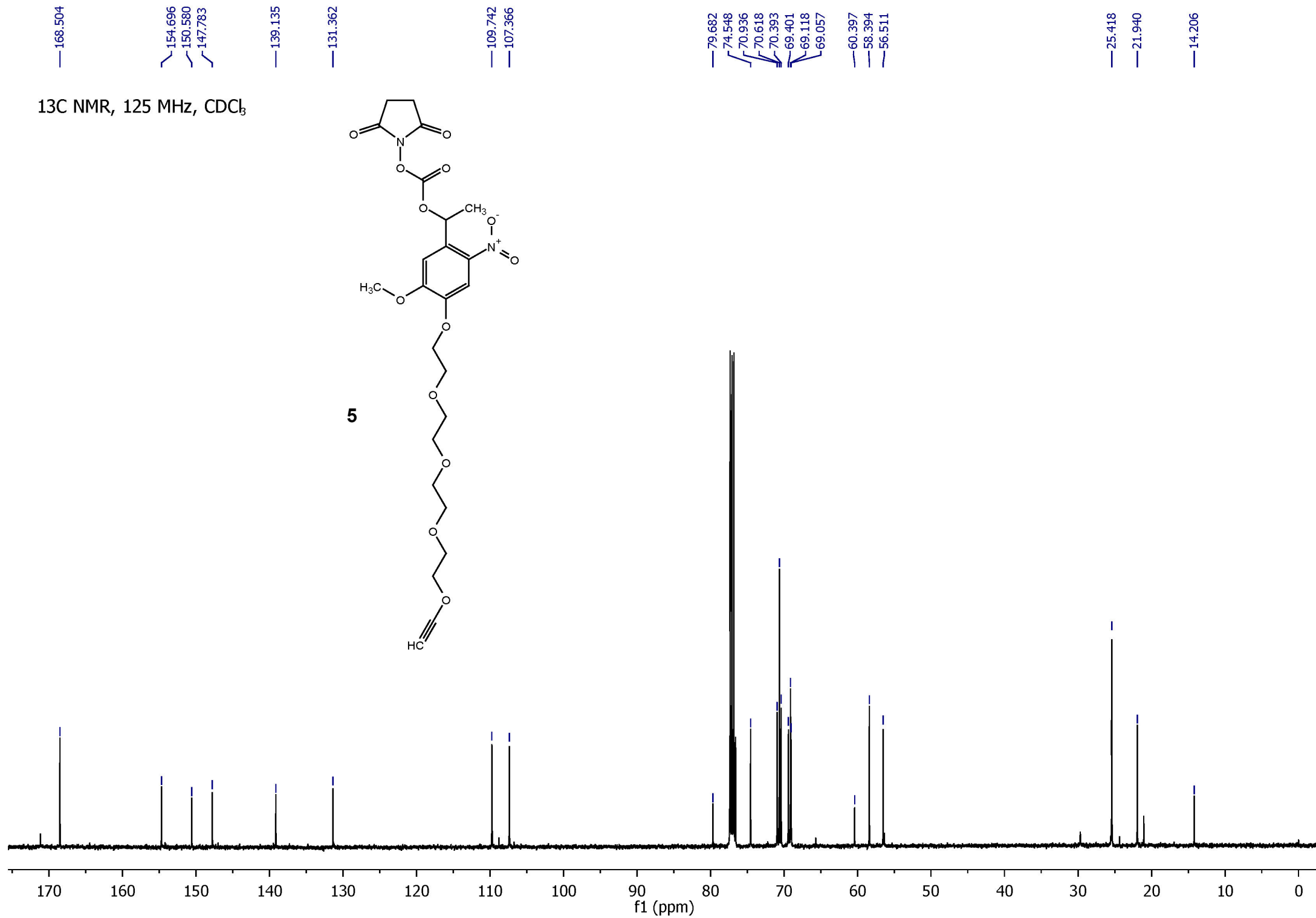
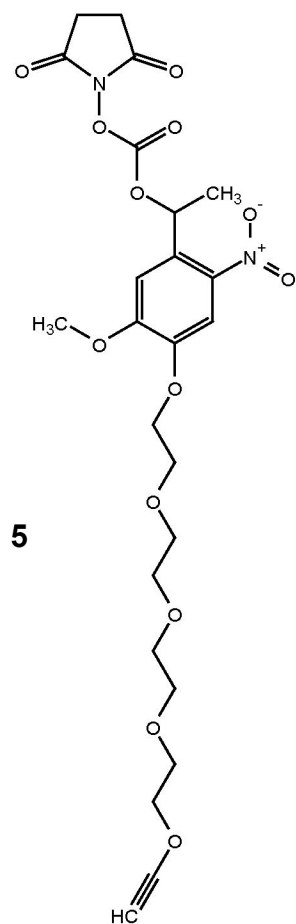


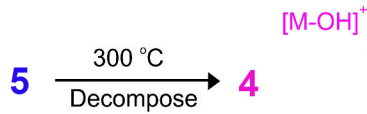


¹H NMR, 400 MHz, CDCl₃



¹³C NMR, 125 MHz, CDCl₃





[M-OH]⁺

