

Supplementary Material

Inhibitory effects of a novel chrysin-derivative, CPD 6, on acute and chronic skin inflammation

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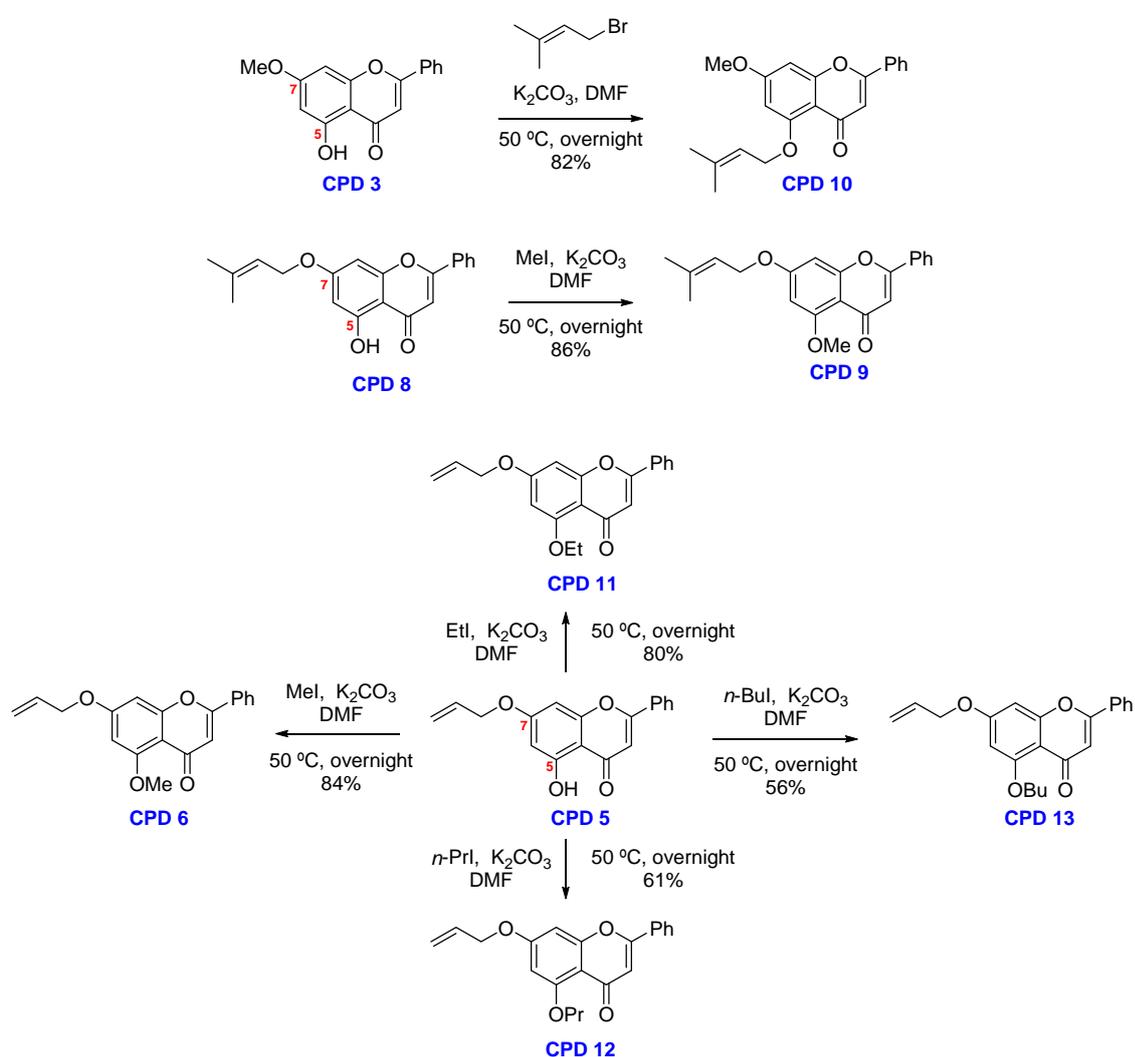
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Synthesis of CPD 1-14

CPD 1 (chrysin) was purchased from Sigma-Aldrich (MO, USA). CPDs 2-8 and CPD14 were synthesized by according to the literature procedure [1-7]. CPD 9 and CPD 10 were prepared from CPD 8 and CPD 3 by methylation and prenylation, respectively (Scheme 1). CPDs 6 and 11-13 were prepared from CPD 5 by alkylation (methylation, ethylation, *n*-propylation, and *n*-butylation).

Scheme 1. Preparation of CPDs 6, 9-13



CPD-6: m.p 128.1 °C; ¹H NMR (600 MHz, CDCl₃) δ 7.89-7.85 (m, 2 H), 7.52-7.48 (m, 3 H), 6.69 (d, *J* = 0.6 Hz, 1 H), 6.58 (d, *J* = 2.4 Hz, 1 H), 6.43 (d, *J* = 2.4 Hz, 1 H), 6.09 (dddd, *J* = 17.4, 10.8, 5.4, 5.4 Hz, 1 H), 5.48 (dddd, *J* = 17.4, 1.8, 1.8, 1.8 Hz, 1 H), 5.37 (dddd, *J* = 10.8, 1.2, 1.2, 1.2 Hz, 1 H), 4.65 (ddd, *J* = 13.2, 7.8, 4.8 Hz, 2 H), 3.97 (s, 3 H); ¹³C NMR (150 MHz, CDCl₃) δ 177.6, 163.0, 161.0, 160.7, 159.8, 132.1, 131.2, 128.9, 126.0, 118.6, 109.4, 109.1, 96.6, 93.7, 69.3, 56.5.

CPD-9: m.p 121.2 °C; ¹H NMR (600 MHz, CDCl₃) δ 7.90-7.85 (m, 2 H), 7.54-7.46 (m, 3 H), 6.68 (s, 1 H), 6.59 (d, *J* = 2.4 Hz, 1 H), 6.40 (d, *J* = 2.4 Hz, 1 H), 5.53 (dddd, *J* = 7.2, 7.2, 1.2, 1.2 Hz, 1 H), 4.61 (d, *J* = 7.2 Hz, 2 H), 3.95 (s, 3 H), 1.84 (s, 3 H), 1.80 (s, 3 H); ¹³C NMR (150 MHz, CDCl₃) δ 177.6, 163.3, 160.9, 160.6, 159.9, 139.6, 131.6, 131.1, 128.9, 126.0, 118.5, 109.3, 109.1, 96.7, 93.5, 65.3, 56.4, 25.8, 18.3.

CPD-10: m.p 138.3 °C; ¹H NMR (600 MHz, CDCl₃) δ 7.91-7.85 (m, 2 H), 7.54-7.45 (m, 3 H), 6.65 (s, 1 H), 6.56 (dd, *J* = 2.4, 0.6 Hz, 1 H), 6.38 (d, *J* = 2.4 Hz, 1 H), 5.58 (dddd, *J* = 6.6, 6.6, 1.2, 1.2 Hz, 2 H), 4.69 (d, *J* = 6.6 Hz, 2 H), 3.90 (s, 3 H), 1.79 (s, 3 H), 1.76 (s, 3 H); ¹³C NMR (150 MHz, CDCl₃) δ 177.4, 163.8, 160.5, 160.2, 159.9, 137.4, 131.7, 131.1, 128.9, 125.9, 119.5, 109.7, 109.1, 97.6, 92.9, 66.6, 55.7, 25.8, 18.4.

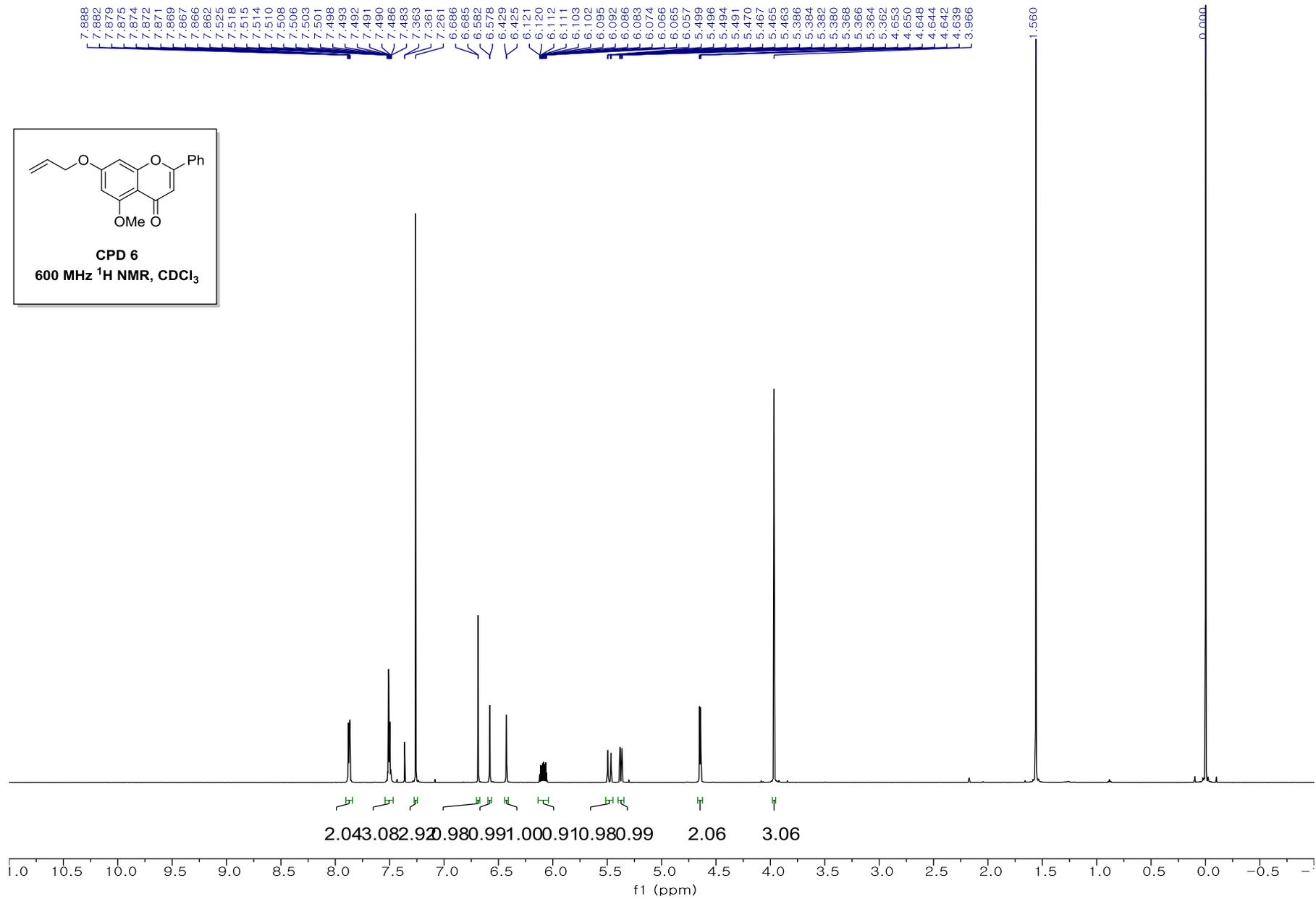
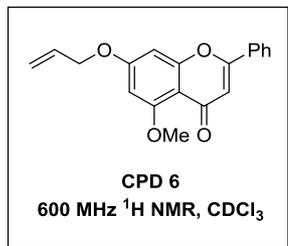
CPD-11: m.p 146.3 °C; ¹H NMR (600 MHz, CDCl₃) δ 7.89-7.86 (m, 2 H), 7.52-7.47 (m, 3 H), 6.65 (s, 1 H), 6.56 (d, *J* = 2.4 Hz, 1 H), 6.41 (d, *J* = 2.4 Hz, 1 H), 6.08 (dddd, *J* = 17.4, 10.8, 5.4, 5.4 Hz, 1 H), 5.47 (dddd, *J* = 17.4, 1.8, 1.8, 1.8 Hz, 1 H), 5.37 (dddd, *J* = 10.8, 1.2, 1.2, 1.2 Hz, 1 H), 4.63 (ddd, *J* = 5.4, 1.8, 1.8 Hz, 2 H), 4.16 (ddd, *J* = 7.2, 7.2, 7.2 Hz, 2 H), 1.56 (dd, *J* = 7.2, 7.2 Hz, 3 H); ¹³C NMR (150 MHz, CDCl₃) δ 177.5, 162.9, 160.5, 160.3, 159.8, 132.2, 131.7, 131.1, 128.9, 126.0, 118.5, 109.6, 109.1, 97.5, 93.6, 69.2, 65.0, 14.5.

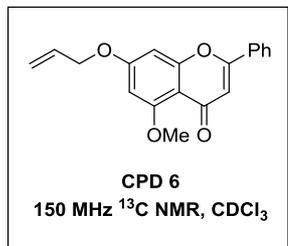
CPD-12: mp 112.6 °C; ¹H NMR (600 MHz, CDCl₃) δ 7.89-7.85 (m, 2 H), 7.53-7.46 (m, 3 H), 6.63 (s, 1 H), 6.55 (d, *J* = 2.4 Hz, 1 H), 6.40 (d, *J* = 2.4 Hz, 1 H), 6.08 (dddd, *J* = 17.4, 10.8, 5.4, 5.4 Hz, 1 H), 5.47 (dddd, *J* = 17.4, 1.8, 1.8, 1.8 Hz, 1 H), 5.36 (dddd, *J* = 10.8, 1.2, 1.2, 1.2 Hz, 1 H), 4.63 (ddd, *J* = 5.4, 1.8, 1.8 Hz, 2 H), 4.03 (dd, *J* = 6.6, 6.6 Hz, 2 H), 1.96 (dddd, *J* = 7.2, 7.2, 7.2, 7.2, 7.2 Hz, 2 H), 1.13 (dd, *J* = 7.2, 7.2 Hz, 3 H); ¹³C NMR (150 MHz, CDCl₃) δ 177.4, 162.9, 160.5, 160.5, 159.8, 132.2, 131.7, 131.1, 128.9, 126.0, 118.5, 109.7, 109.2, 97.4, 93.5, 70.9, 69.2, 22.3, 10.5.

CPD-13: mp 117.9 °C; ¹H NMR (600 MHz, CDCl₃) δ 7.91-7.84 (m, 2 H), 7.54-7.46 (m, 3 H), 6.62 (s, 1 H), 6.55 (d, *J* = 2.4 Hz, 1 H), 6.40 (d, *J* = 2.4 Hz, 1 H), 6.08 (dddd, *J* = 17.4, 10.2, 6.0, 6.0 Hz, 1 H), 5.47 (dddd, *J* = 17.4, 1.8, 1.8, 1.8 Hz, 1 H), 5.36 (dddd, *J* = 10.8, 1.2, 1.2, 1.2 Hz, 1 H), 4.63 (ddd, *J* = 5.4, 1.8, 1.8 Hz, 2 H), 4.07 (dd, *J* = 6.6, 6.6 Hz, 2 H), 1.97-1.87 (m, 2 H), 1.64-1.56 (m, 2 H), 1.00 (dd, *J* = 7.2, 7.2 Hz, 3 H); ¹³C NMR (150 MHz, CDCl₃) δ 177.4, 162.9, 160.6, 160.5, 159.8, 132.2, 131.7, 131.1, 128.9, 126.0, 118.5, 109.7, 109.2, 97.4, 93.5, 69.2, 30.9, 19.2, 13.8.

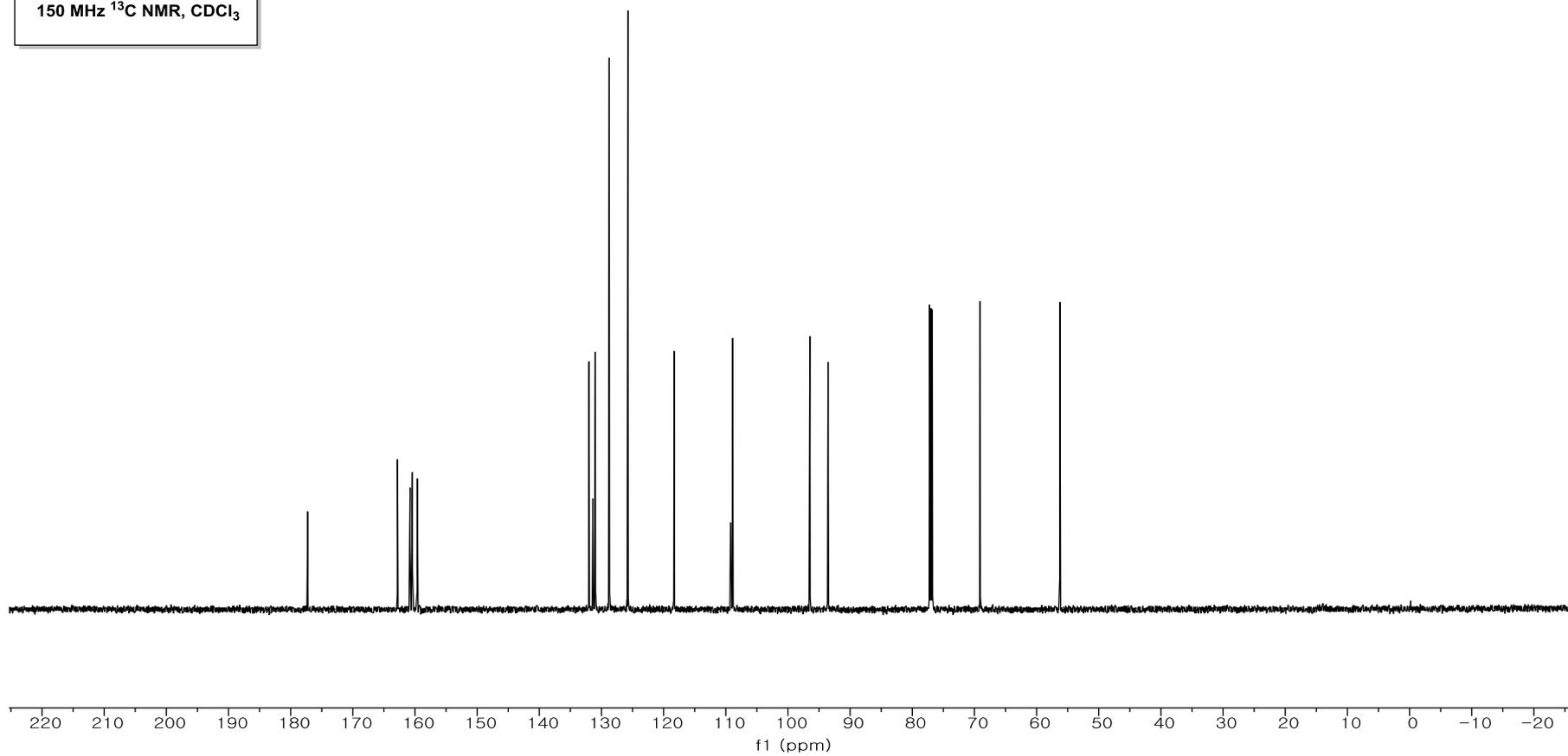
References

1. For **CPD 2**: Daskiewicz, J.B.; Depeint, F.; Viorner, L.; Bayet, C.; Comte-Sarrazin, G.; Comte, G.; Gee, J.M.; Johnson, I.T.; Ndjoko, K.; Hostettmann, K.; Barron, D. Effects of Flavonoids on Cell Proliferation and Caspase Activation in a Human Colonic Cell Line HT29: An SAR Study. *J. Med. Chem.* **2005**, *48*, 2790–2804.
2. For **CPD 3**: Moreira, J.; Ribeiro, D.; Silva, P.M. A.; Nazareth, N.; Monteiro, M.; Palmeira, A.; Saraiva, L.; Pinto, M.; Bousbaa, H.; Cidade, H. New Alkoxy Flavone Derivatives Targeting Caspases: Synthesis and Antitumor Activity Evaluation. *Molecules* **2018**, *24*, 129.
3. For **CPD 4**: Khanapur, M.; Pinna, N.K.; Badiger, J. Synthesis and anti-inflammatory in vitro, in silico, and in vivo studies of flavone analogues. *Med. Chem. Res.* **2015**, *24*, 2656–2669.
4. For **CPD 5**: Kim, H.; Lim, D.; Shin, I.; Lee, D. Gram-scale synthesis of anti-pancreatic flavonoids (\pm)-8-[1-(4'-hydroxy-3'-methoxyphenyl)prop-2-en-1-yl]-chrysin and -galangin. *Tetrahedron* **2014**, *70*, 4738–4744.
5. For **CPD 6**: Aneja, R.; Khanna, R. N.; Seshadri, T. R. 6-Methoxyfuroflavone, a new component of the seeds of *Pongamia glabra*. *J. Chem. Soc.* **1963**, 163–168.
6. For **CPD 7**: Khanna, R. N.; Seshadri, T. R. Synthesis of isomers of kanjone. *Indian. J. Chem.*, **1963**, *1*, 385–387.
7. For **CPD 8**: a) Hwang, S. H.; Kim, H. Y.; Zuo, G.; Wang, Z.; Lee, J.-Y.; Lim, S. S. Anti-glycation, Carbonyl Trapping and Anti-inflammatory Activities of Chrysin Derivatives. *Molecules* **2018**, *23*, 1752. b) Wang, S.H.; Chen, C.H.; Lo, C.Y.; Feng, J.Z.; Lin, H.J.; Chang, P.Y.; Yang, L. L.; Chen, L.G.; Liu, Y. W.; Kuo, C. D.; Wu, J.Y. Synthesis and biological evaluation of novel 7-O-lipophilic substituted baicalein derivatives as potential anticancer agents. *Med. Chem. Comm.* **2015**, *6*, 1864–1873.





177.281
162.813
160.783
160.413
159.633
131.998
131.384
130.983
128.745
125.750
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109.239
108.883
96.474
93.554
77.212
77.006
76.788
69.079
56.253



State Parameter Editor

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 Ion Source Gas 2 (GS2): 0.0
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 Declustering Potential (DP): 70.0
 Entrance Potential (EP): 10.0

 Q1 Resolution: Unit
 Ion Energy 1 (IE1): 1.0

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Mass Spectrometer Method Properties

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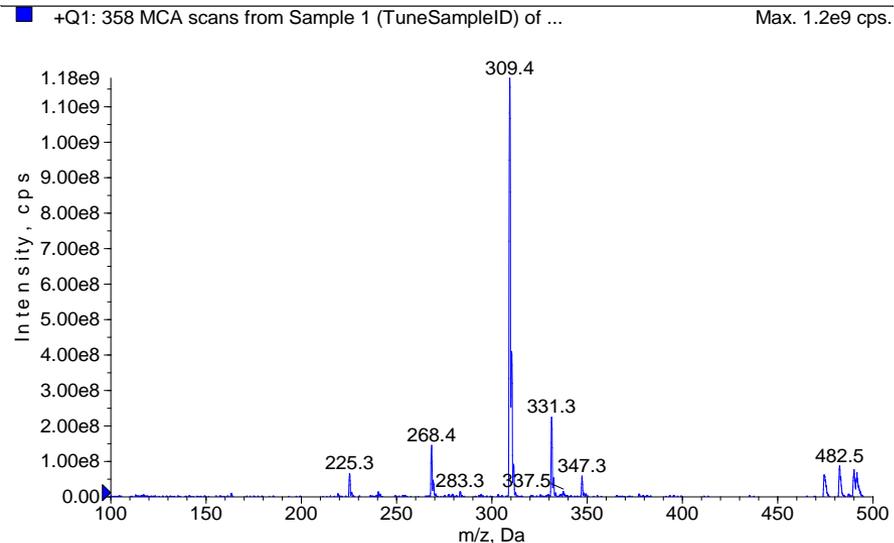
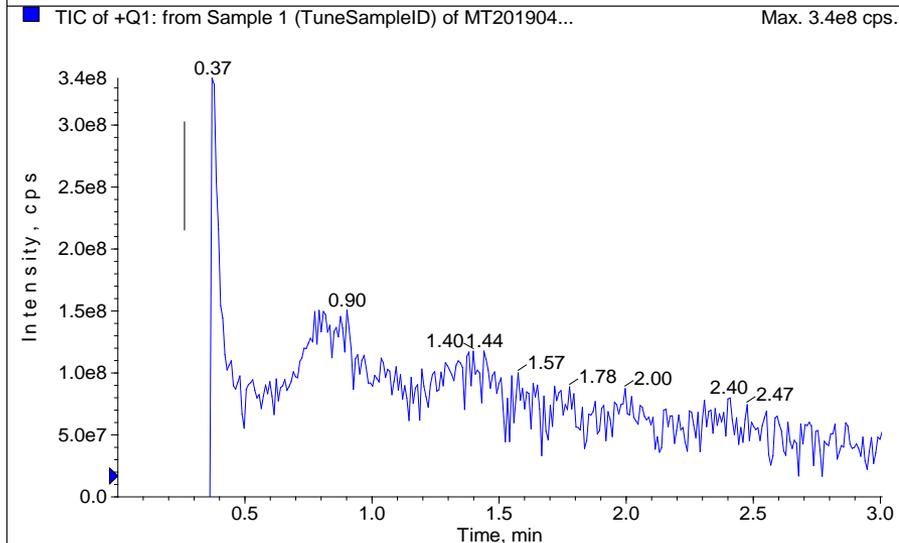
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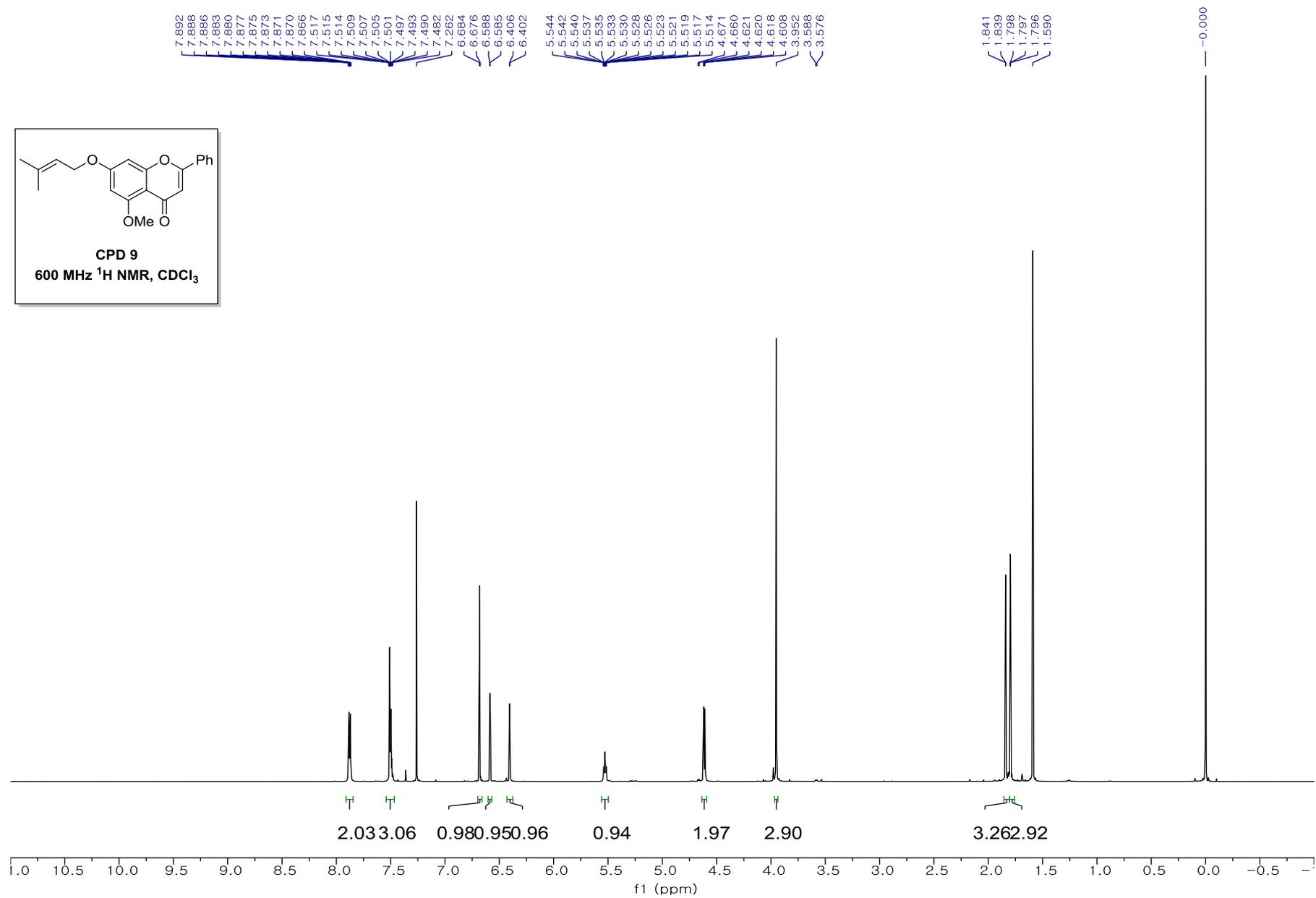
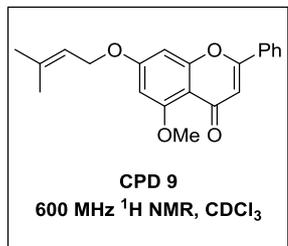
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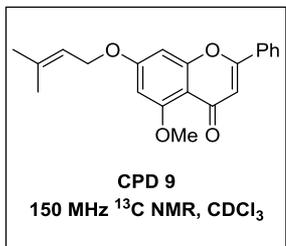
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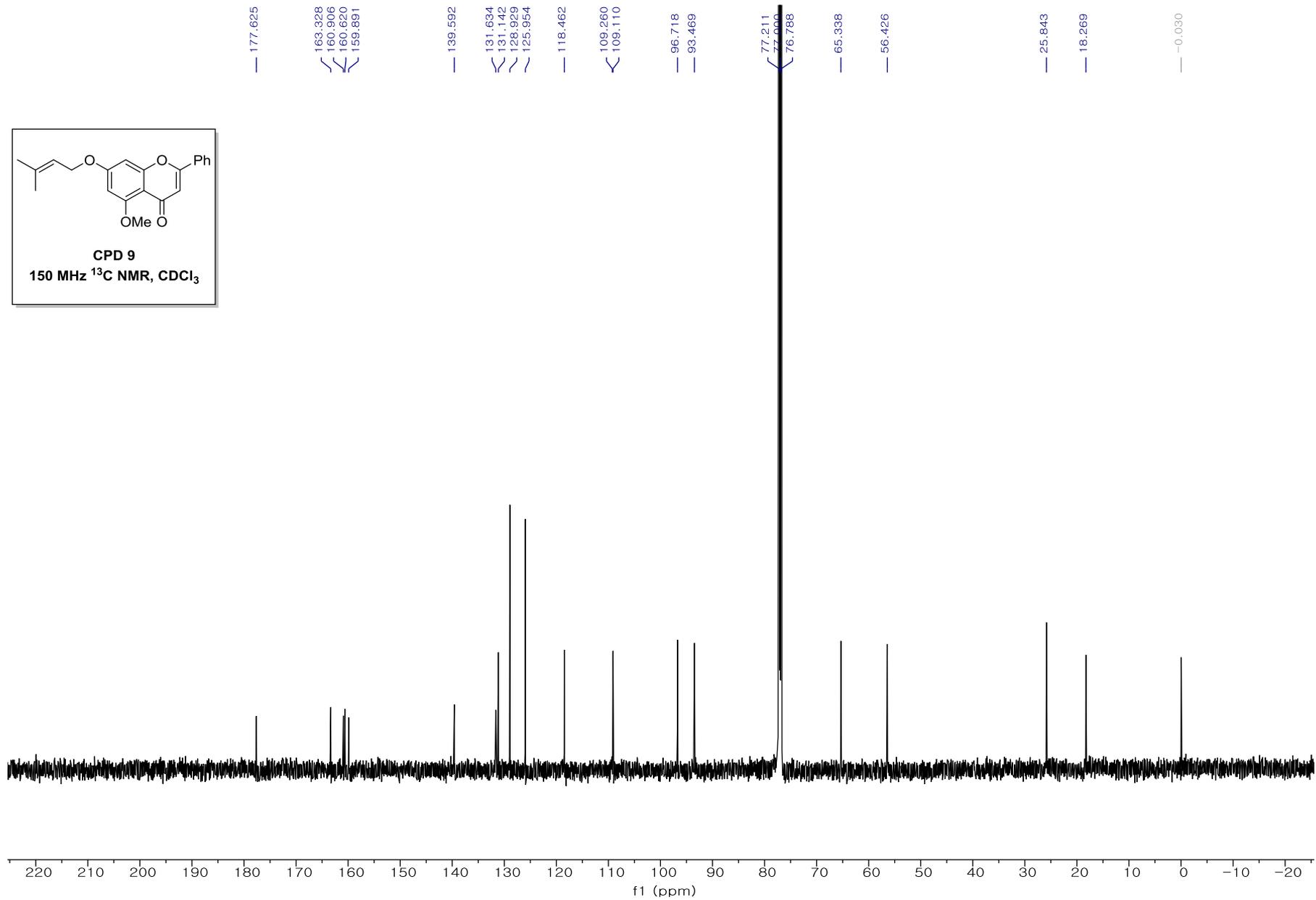
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 DP: 70.00
 EP: 10.00







- 177.625
- 163.328
- 160.906
- 160.620
- 159.891
- 139.592
- 131.654
- 131.142
- 128.929
- 125.954
- 118.462
- 109.260
- 109.110
- 96.718
- 93.469
- 77.211
- 77.000
- 76.788
- 65.338
- 56.426
- 25.843
- 18.269
- 0.030



State Parameter Editor

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 Ion Source Gas 2 (GS2): 0.0
 Interface Heater (ihe): On

 Declustering Potential (DP): 70.0
 Entrance Potential (EP): 10.0

 Q1 Resolution: Unit
 Ion Energy 1 (IE1): 1.0

 Deflector (DF): -200.0
 CEM (CEM): 2500.0

Mass Spectrometer Method Properties

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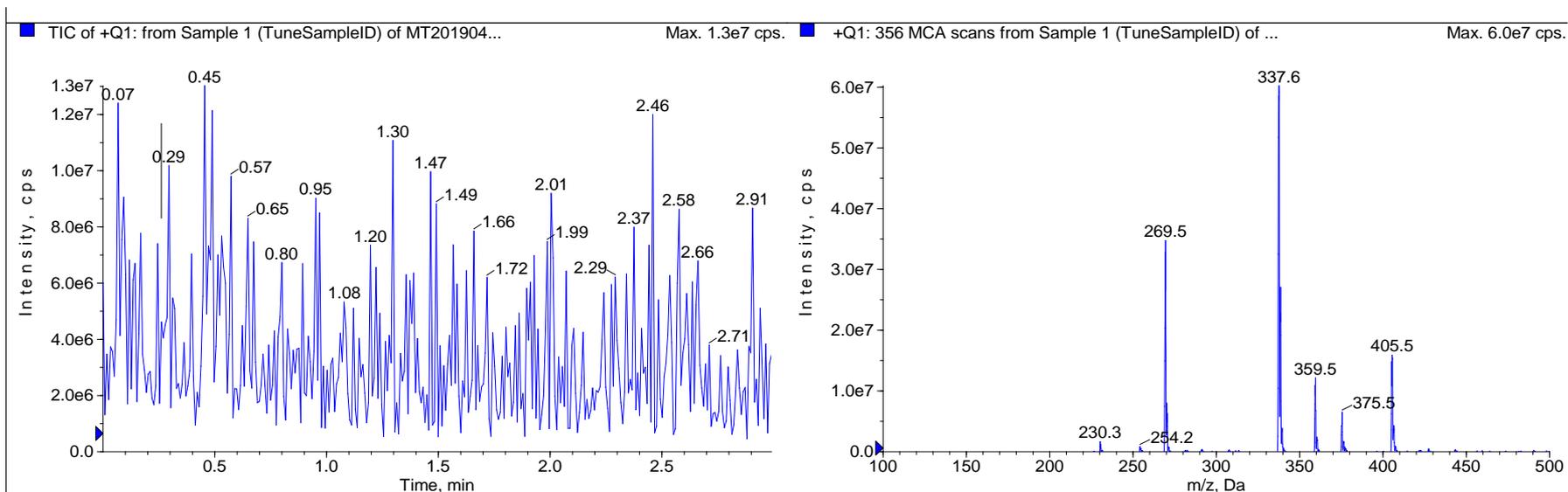
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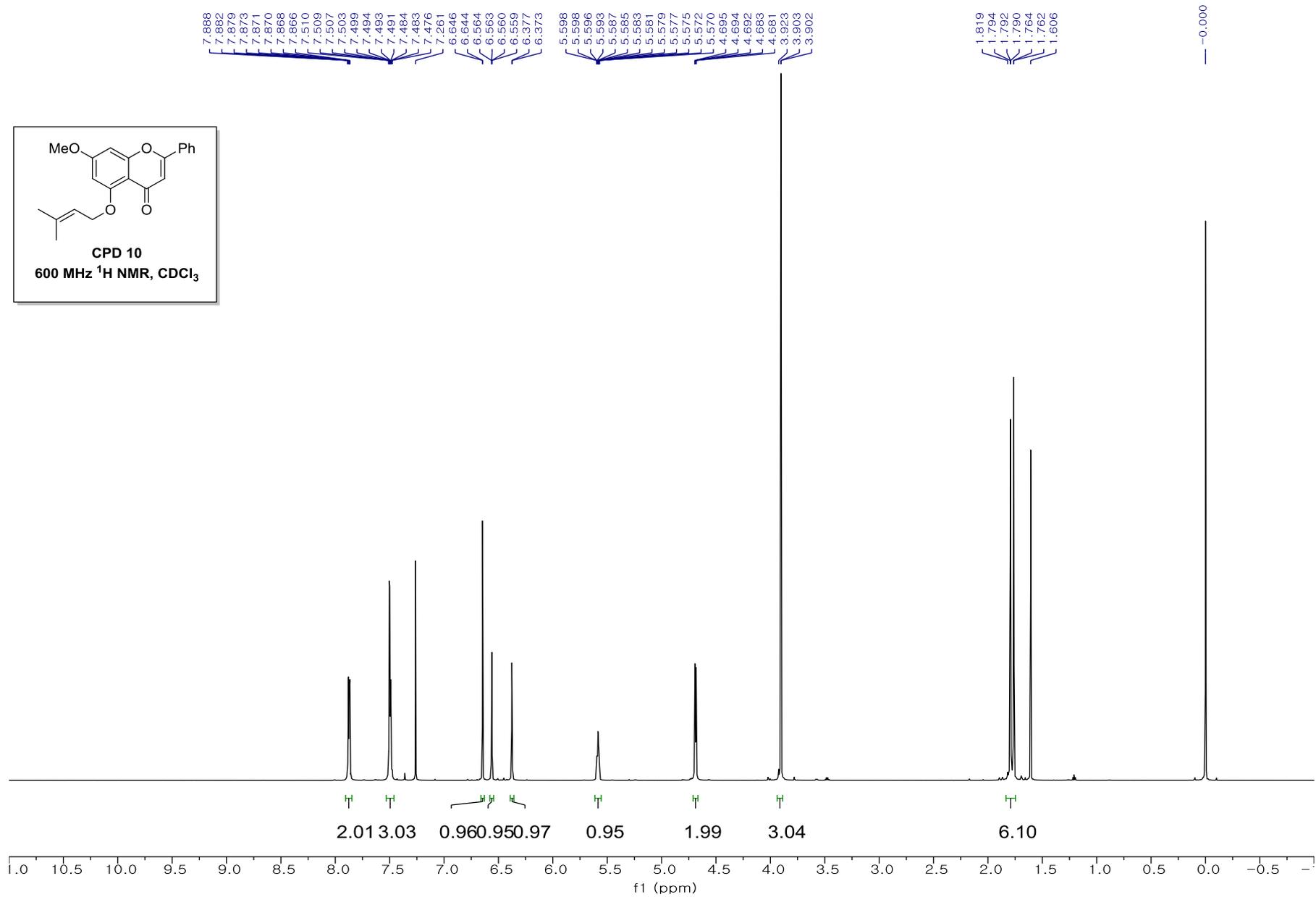
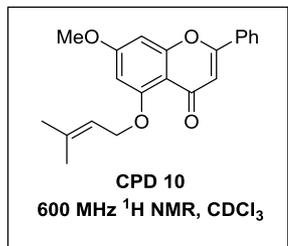
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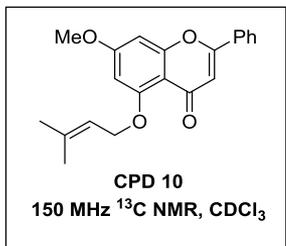
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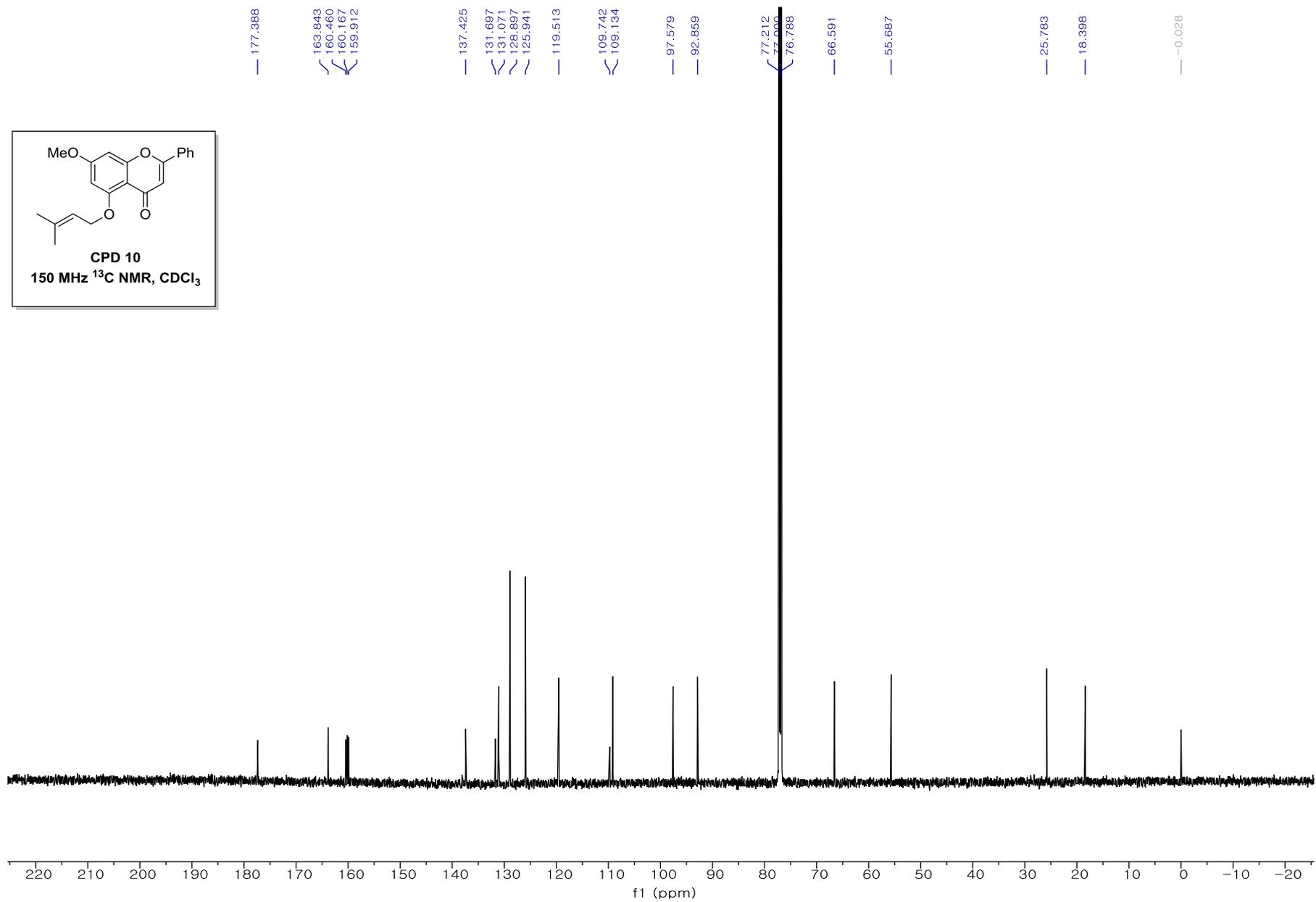




S11



- 177.388
- 163.843
- 160.460
- 160.167
- 159.912
- 137.425
- 131.697
- 131.071
- 128.897
- 125.941
- 119.513
- 109.742
- 109.134
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- 92.859
- 77.212
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State Parameter Editor

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 Interface Heater (ihe): On

 Declustering Potential (DP): 70.0
 Entrance Potential (EP): 10.0

 Q1 Resolution: Unit
 Ion Energy 1 (IE1): 1.0

 Deflector (DF): -200.0
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Mass Spectrometer Method Properties

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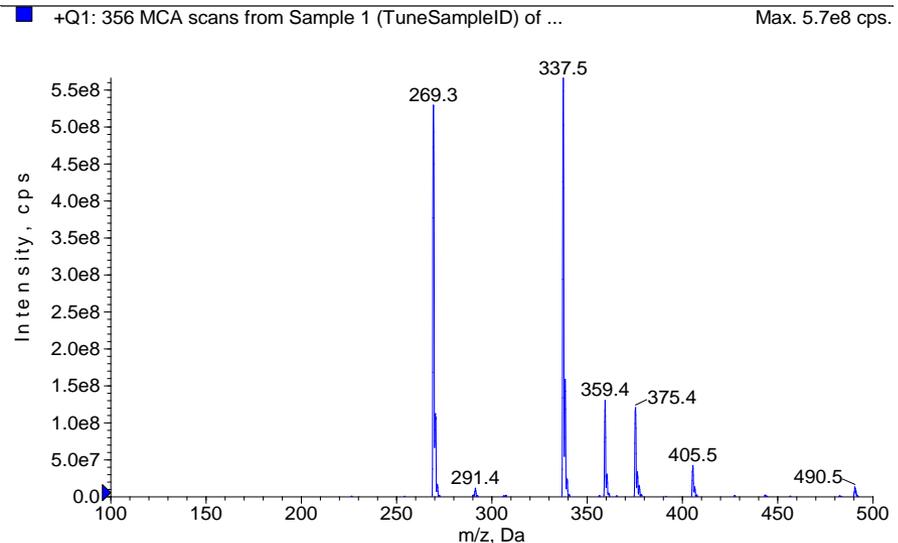
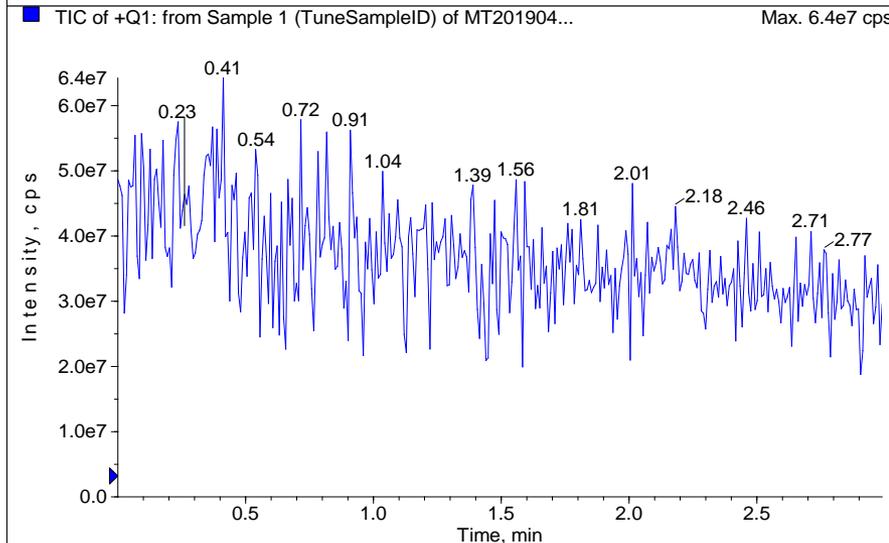
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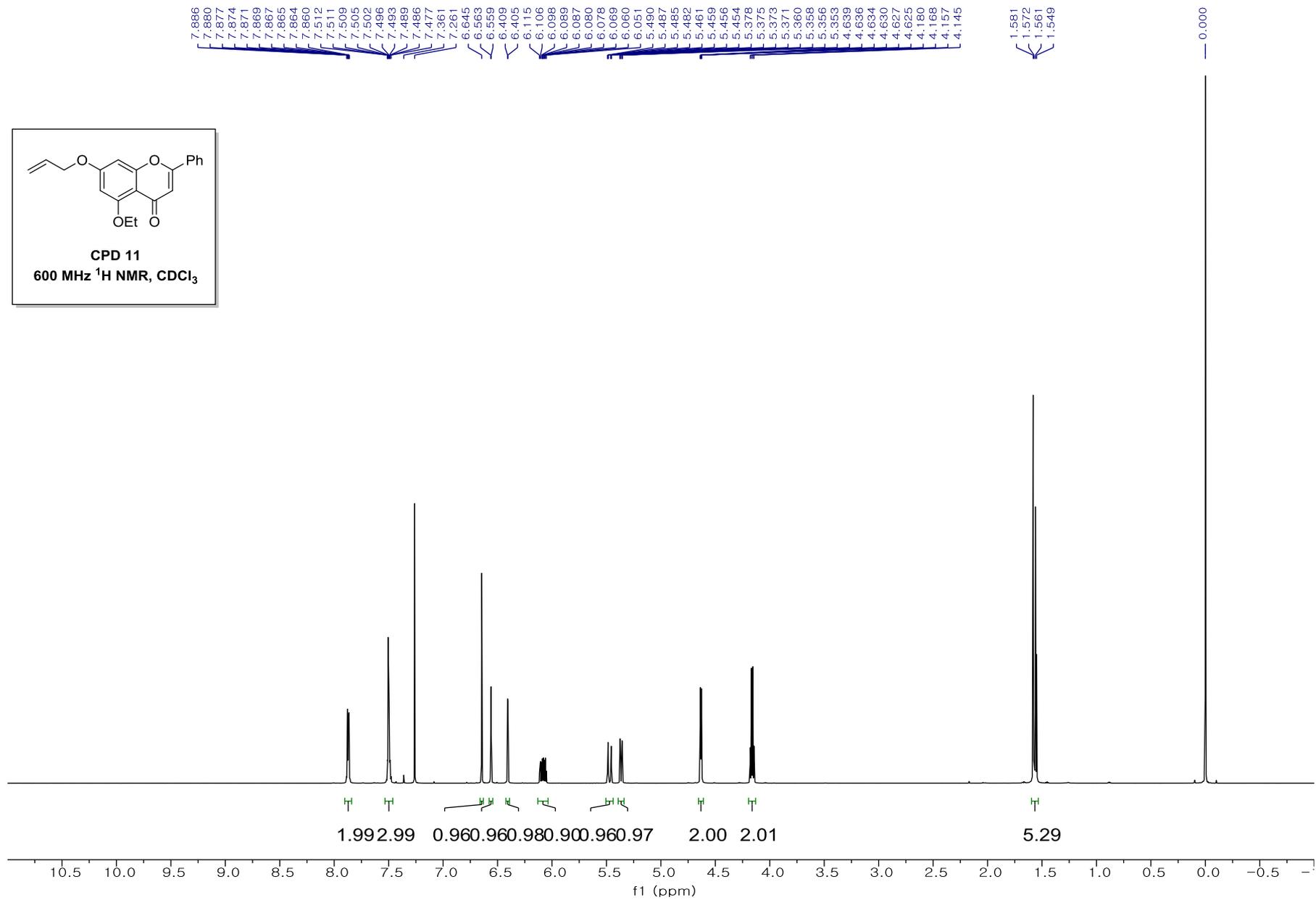
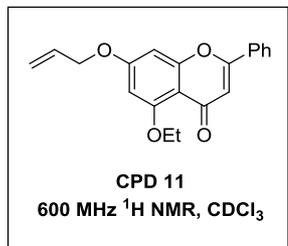
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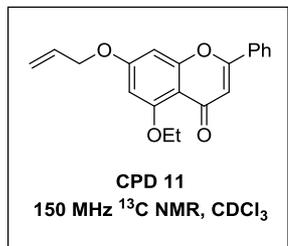
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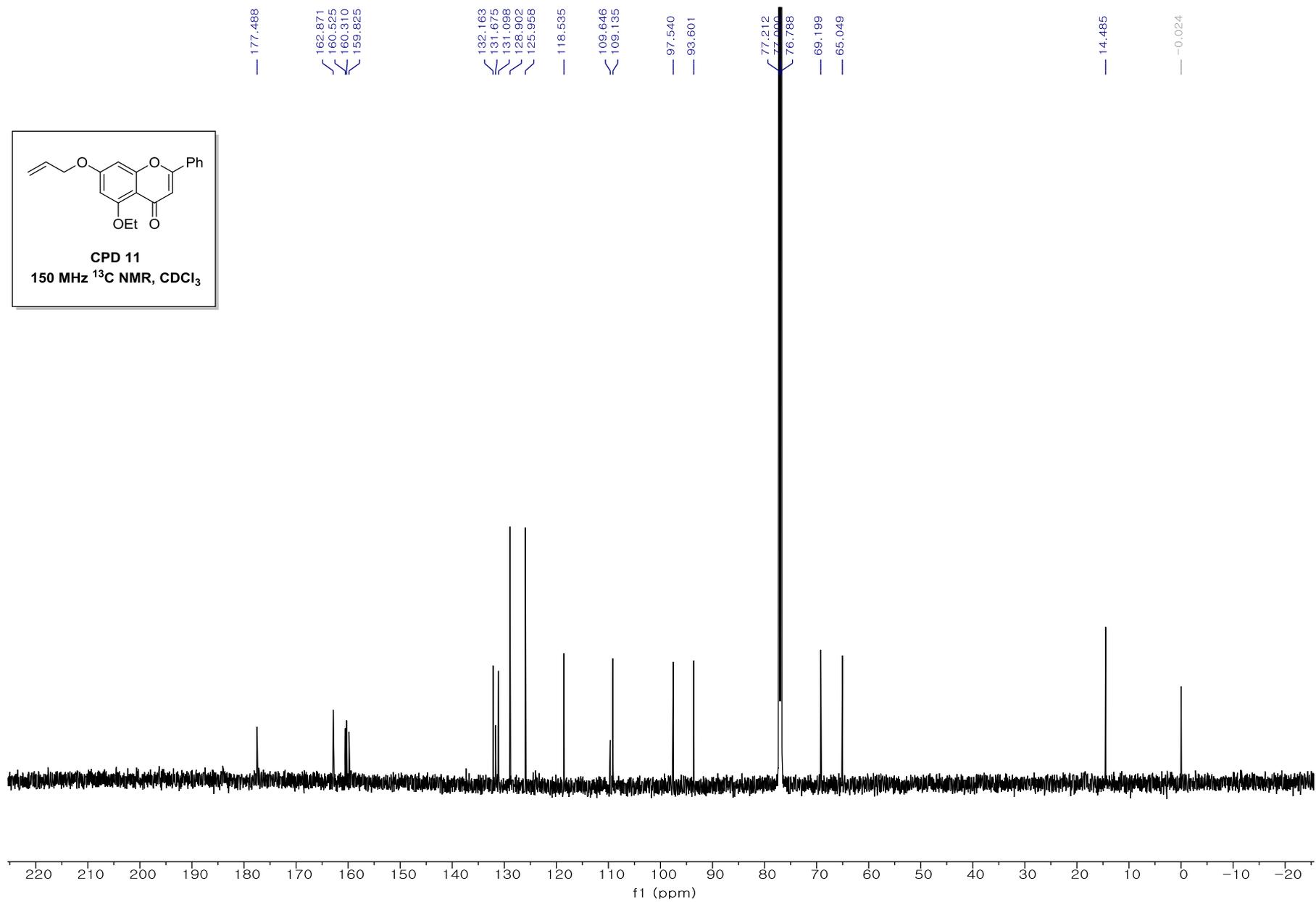
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- 162.871
- 160.525
- 160.310
- 159.825
- 132.163
- 131.675
- 131.098
- 128.902
- 125.958
- 118.535
- 109.646
- 109.135
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- 93.601
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- 14.485
- 0.024



State Parameter Editor

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 Ion Source Gas 2 (GS2): 0.0
 Interface Heater (ihe): On

 Declustering Potential (DP): 70.0
 Entrance Potential (EP): 10.0

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 Ion Energy 1 (IE1): 1.0

 Deflector (DF): -200.0
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Mass Spectrometer Method Properties

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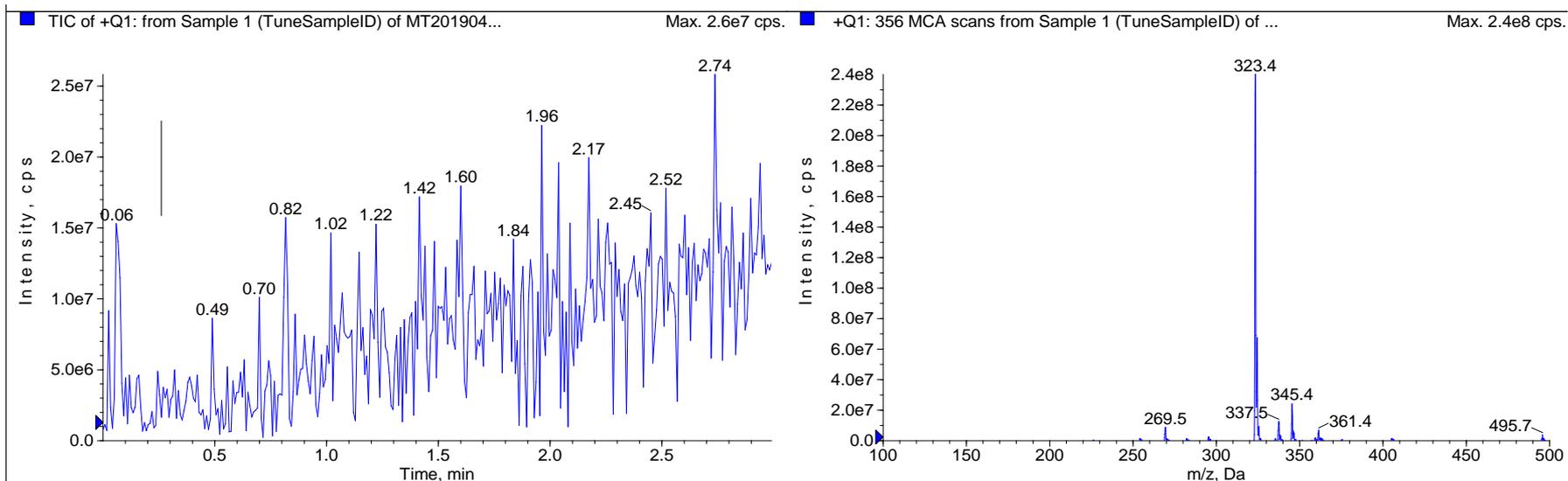
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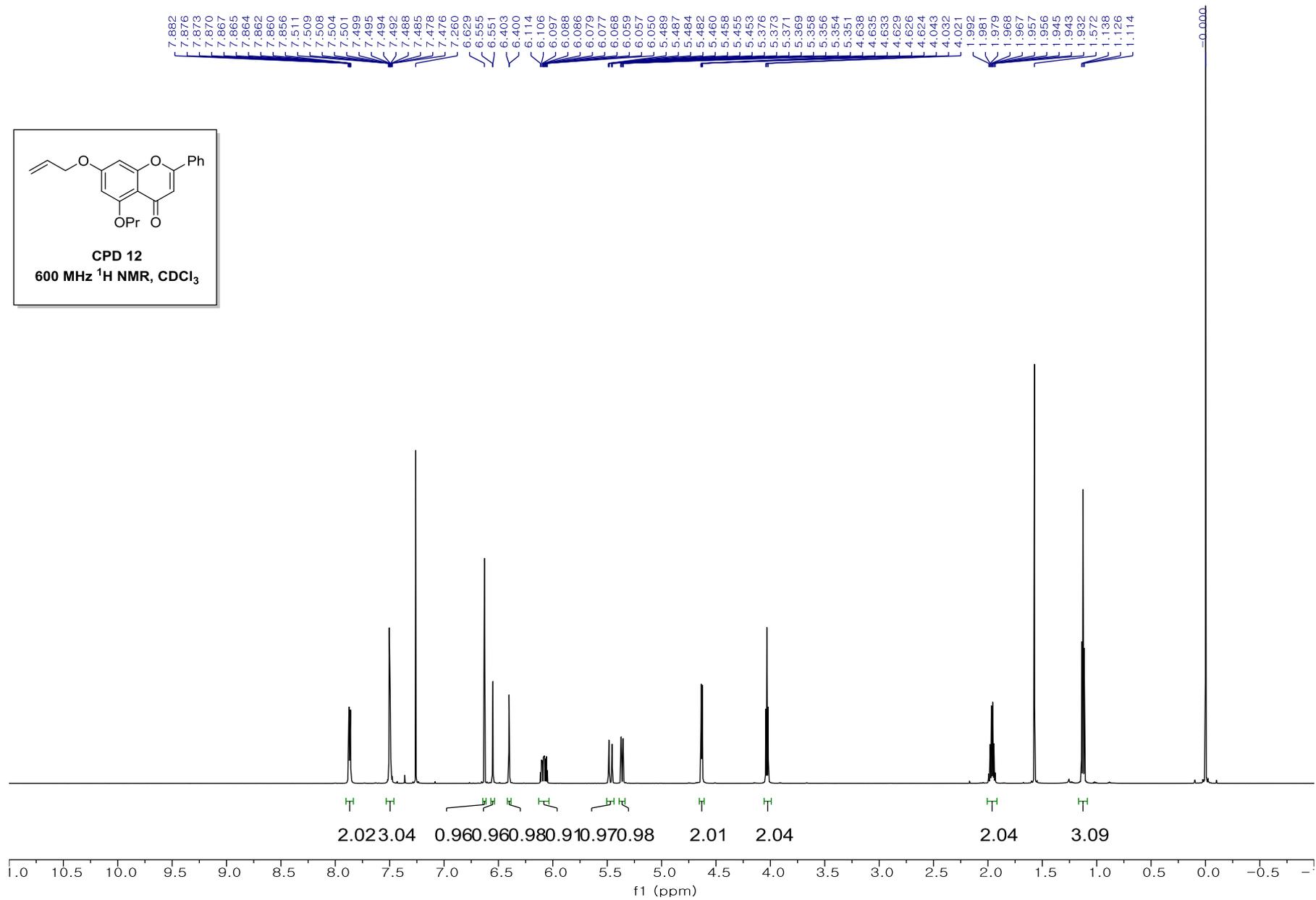
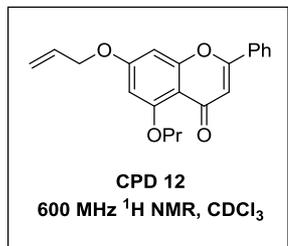
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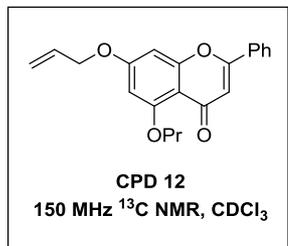
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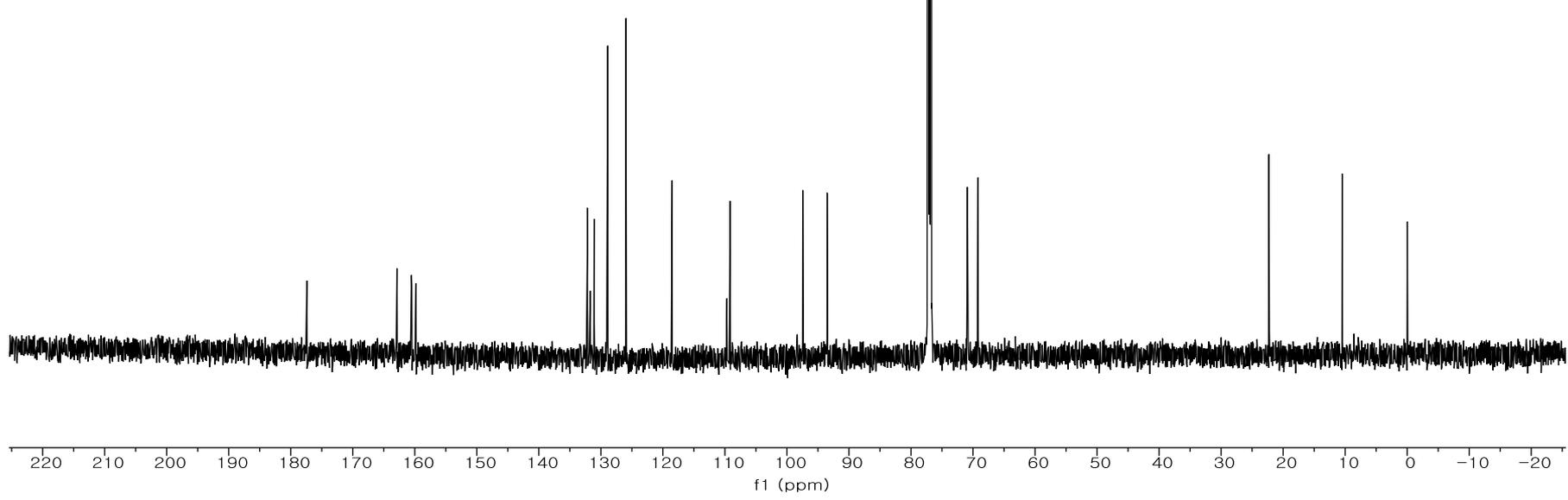
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- 177.403
- 162.867
- 160.540
- 160.485
- 159.804
- 132.187
- 131.718
- 131.073
- 128.901
- 125.958
- 118.508
- 109.681
- 109.171
- 97.447
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- 77.006
- 76.788
- 70.904
- 69.184
- 22.293
- 10.469
- 0.027



State Parameter Editor

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 Ion Source Gas 1 (GS1): 12.0
 Ion Source Gas 2 (GS2): 0.0
 Interface Heater (ihe): On

 Declustering Potential (DP): 70.0
 Entrance Potential (EP): 10.0

 Q1 Resolution: Unit
 Ion Energy 1 (IE1): 1.0

 Deflector (DF): -200.0
 CEM (CEM): 2500.0

Mass Spectrometer Method Properties

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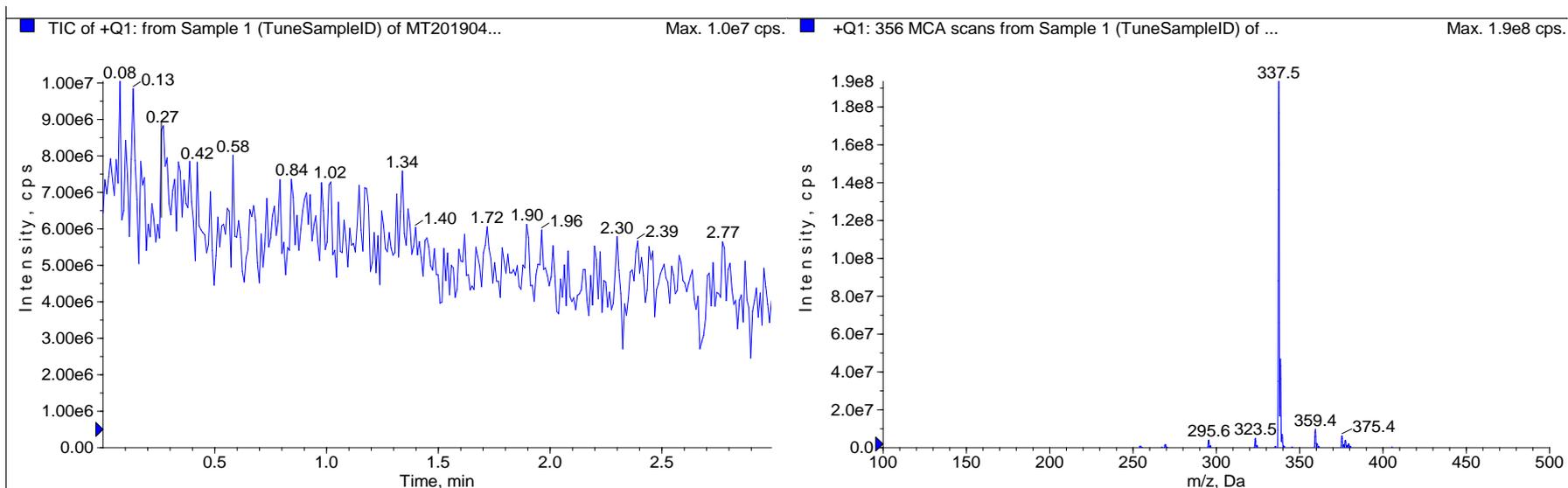
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 Relative Start Time: 0.00 msec
 Experiments in Period: 1

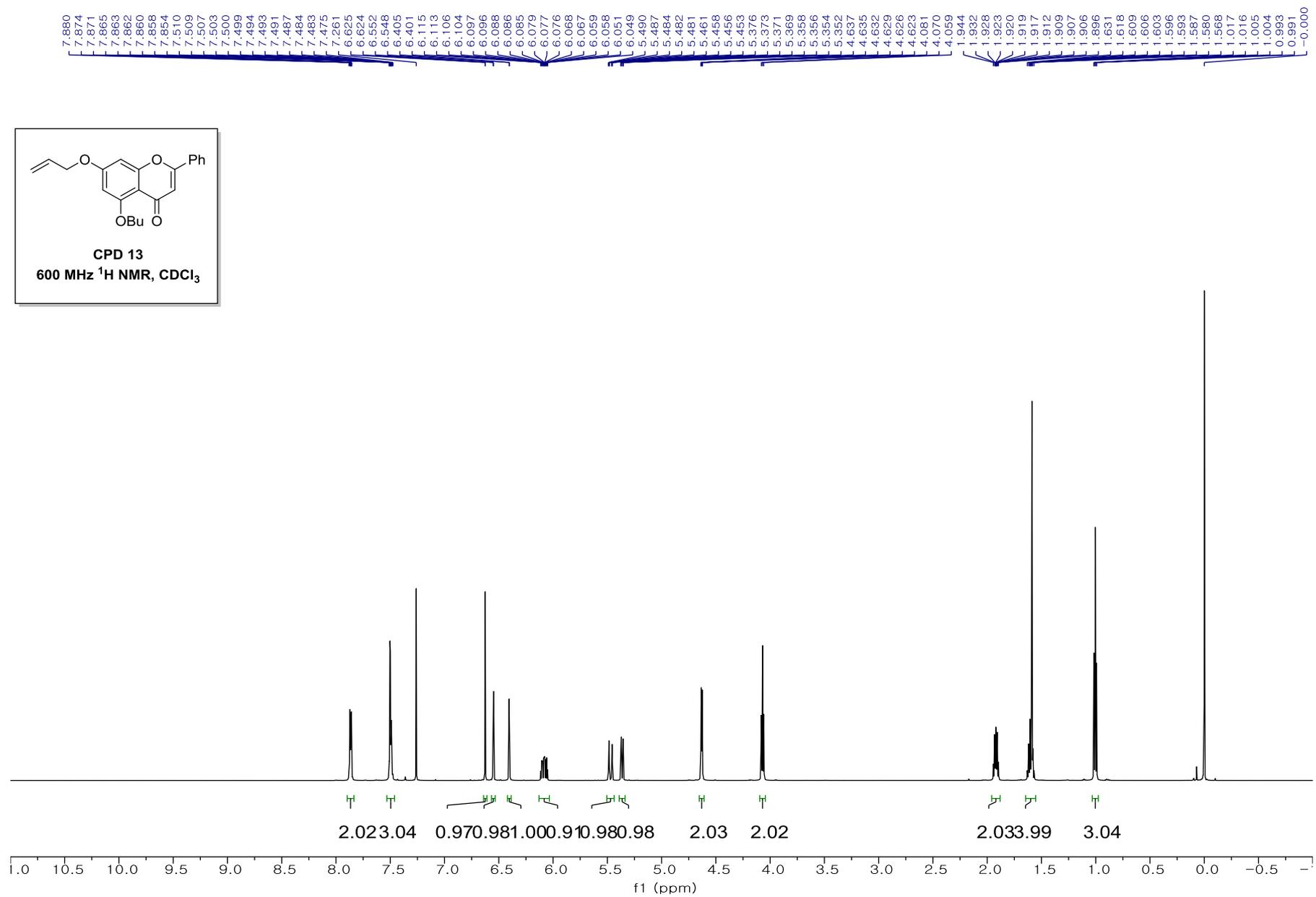
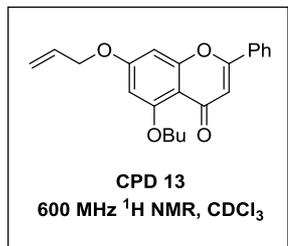
 Period 1 Experiment 1:

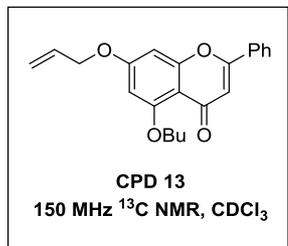
 Scan Type: Q1 MS (Q1)
 Polarity: Positive
 Scan Mode: Profile
 Ion Source: Turbo Spray
 Resolution Q1: Unit
 Intensity Thres.: 0.00 cps
 Settling Time: 0.0000 msec
 MR Pause: 5.0070 msec
 MCA: Yes
 Center/Width: No
 Step Size: 0.10 Da

Start (Da)	Stop (Da)	Time (sec)	Param	Start	Stop
100.00	500.00	0.50	CEP	7.84	19.33

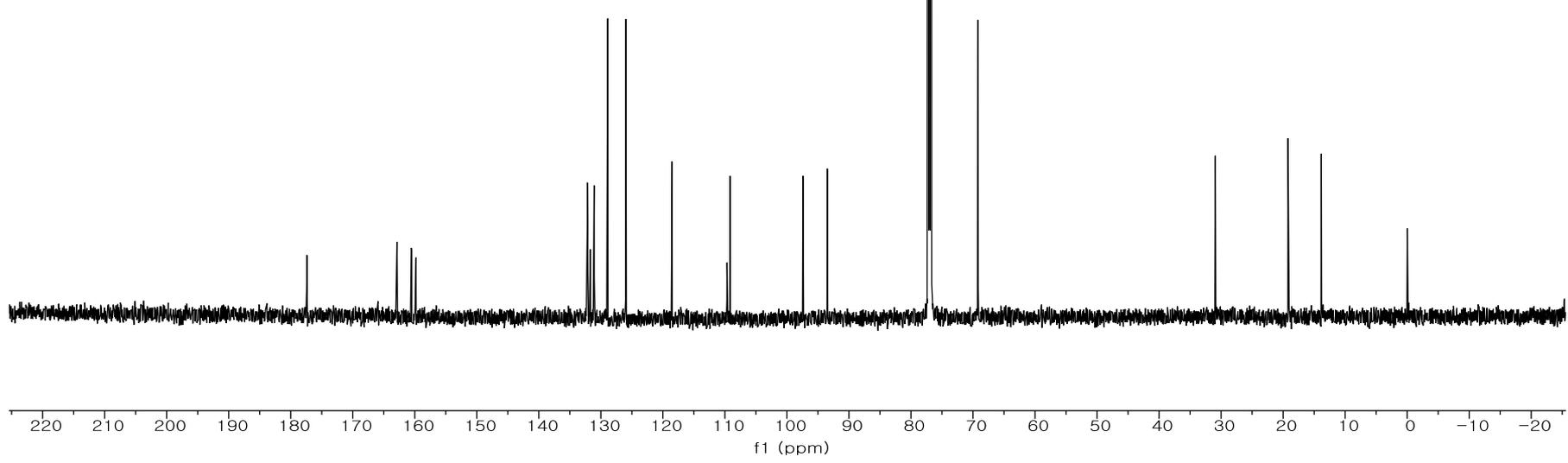
 Parameter Table(Period 1 Experiment 1):
 CUR: 10.00
 TEM: 0.00
 GS1: 12.00
 GS2: 0.00
 ihe: ON
 IS: 5500.00
 DP: 70.00
 EP: 10.00







- 177.394
- 162.865
- 160.557
- 160.486
- 159.803
- 132.185
- 131.716
- 131.070
- 128.896
- 125.956
- 118.510
- 109.664
- 109.163
- 97.419
- 93.490
- 77.211
- 77.000
- 76.788
- 69.181
- 30.937
- 19.202
- 13.849
- 0.030



State Parameter Editor

Ion Source: Turbo Spray
 Ion Source Temperature Reached
 Curtain Gas (CUR): 10.0
 Ion Spray Voltage (IS): 5500.0
 Temperature (TEM): 0.0
 Ion Source Gas 1 (GS1): 12.0
 Ion Source Gas 2 (GS2): 0.0
 Interface Heater (ihe): On

 Declustering Potential (DP): 70.0
 Entrance Potential (EP): 10.0

 Q1 Resolution: Unit
 Ion Energy 1 (IE1): 1.0

 Deflector (DF): -200.0
 CEM (CEM): 2500.0

Mass Spectrometer Method Properties

Period 1:

 Scans in Period: 356
 Relative Start Time: 0.00 msec
 Experiments in Period: 1

 Period 1 Experiment 1:

 Scan Type: Q1 MS (Q1)
 Polarity: Positive
 Scan Mode: Profile
 Ion Source: Turbo Spray
 Resolution Q1: Unit
 Intensity Thres.: 0.00 cps
 Settling Time: 0.0000 msec
 MR Pause: 5.0070 msec
 MCA: Yes
 Center/Width: No
 Step Size: 0.10 Da

Start (Da)	Stop (Da)	Time (sec)	Param	Start	Stop
100.00	500.00	0.50	CEP	7.84	19.33

 Parameter Table(Period 1 Experiment 1):
 CUR: 10.00
 TEM: 0.00
 GS1: 12.00
 GS2: 0.00
 ihe: ON
 IS: 5500.00
 DP: 70.00
 EP: 10.00

