

Supplementary Material

A modular approach for the installation of functionalized phosphonates to heterocycles

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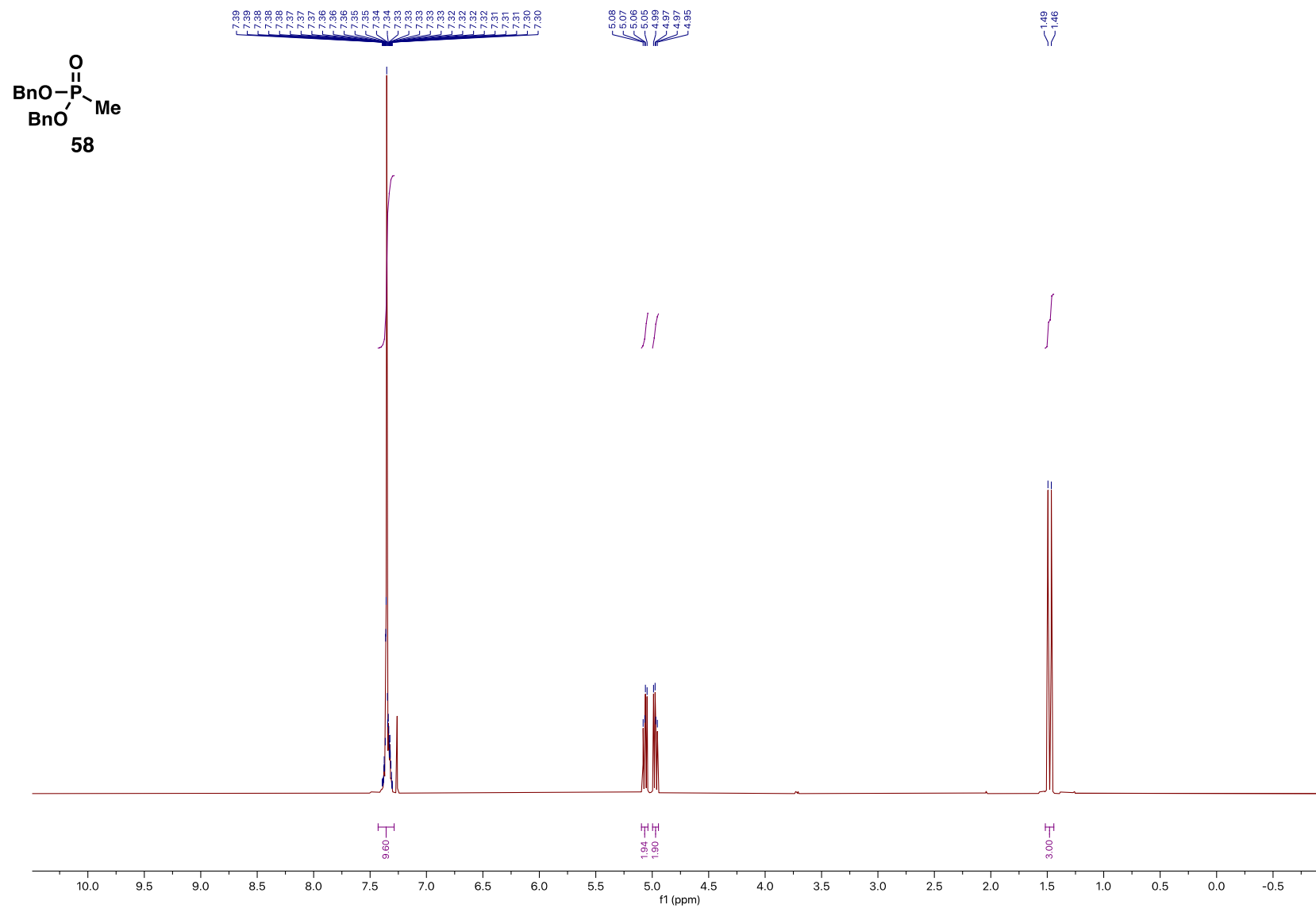
^b Department of Oncologic Sciences, College of Medicine, University of South Florida,
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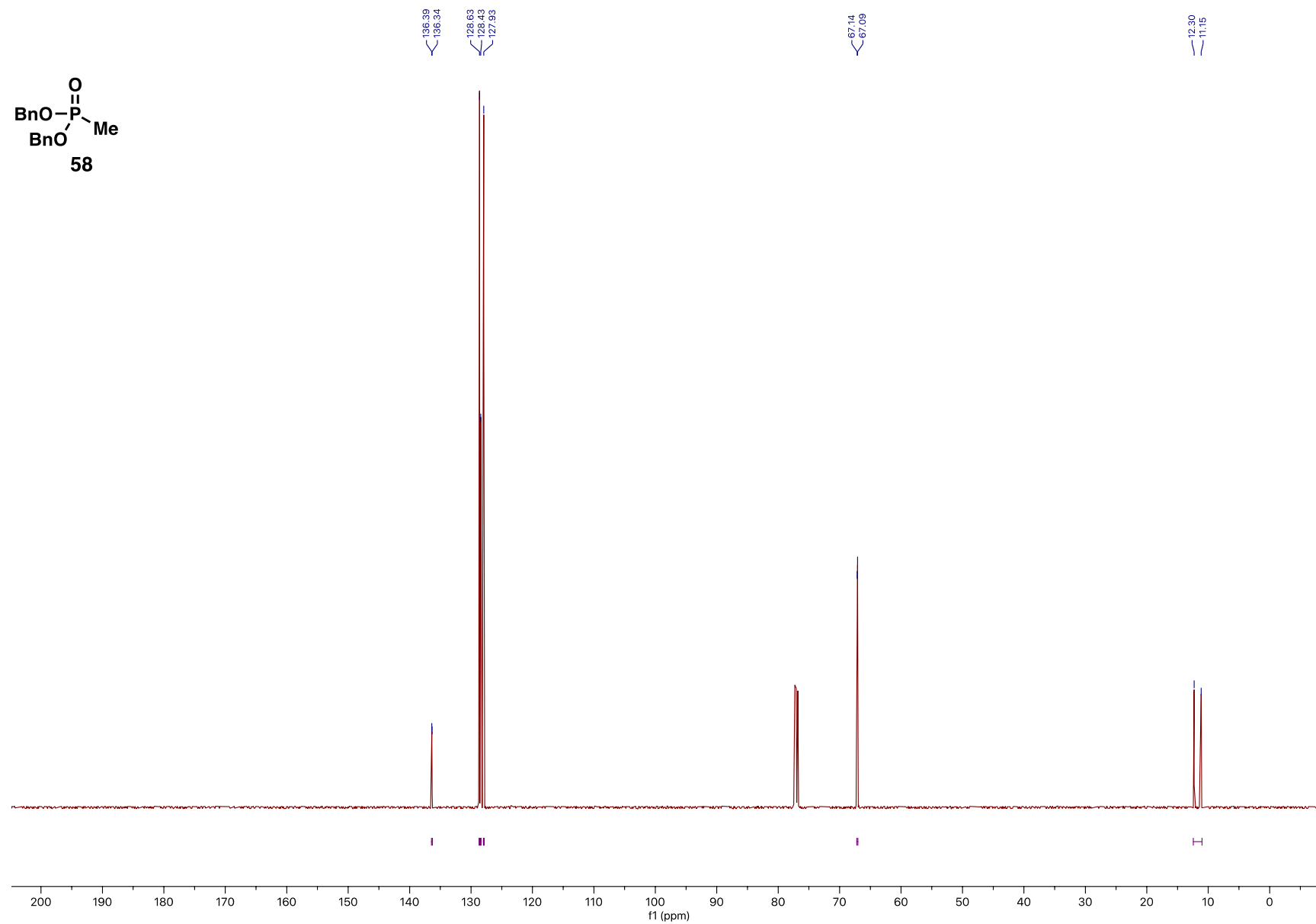
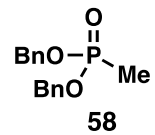
^c Department of Chemistry, University of South Florida, Tampa, Florida 33620, USA
Email: justin.lopchuk@moffitt.org

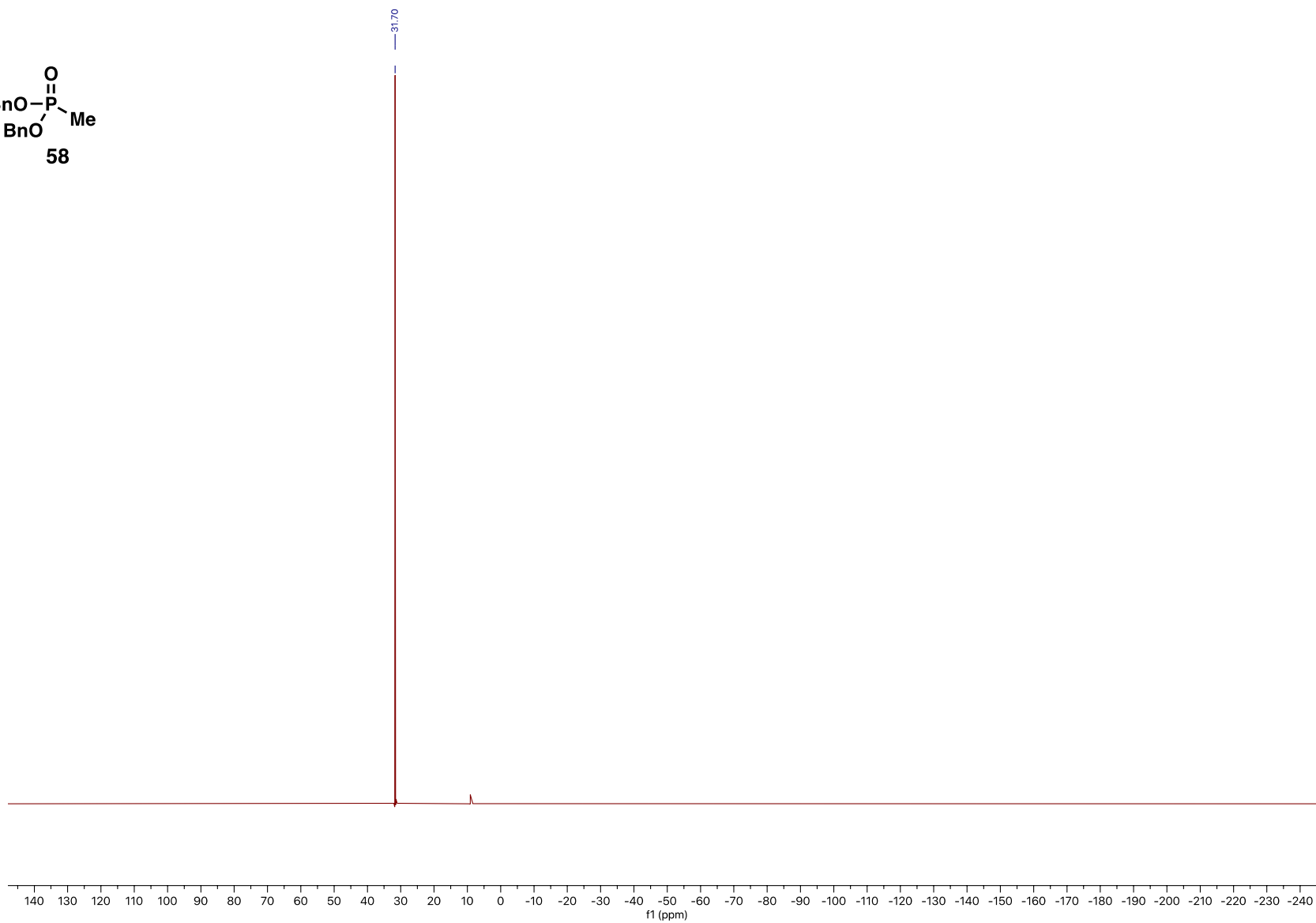
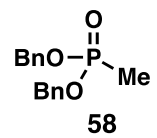
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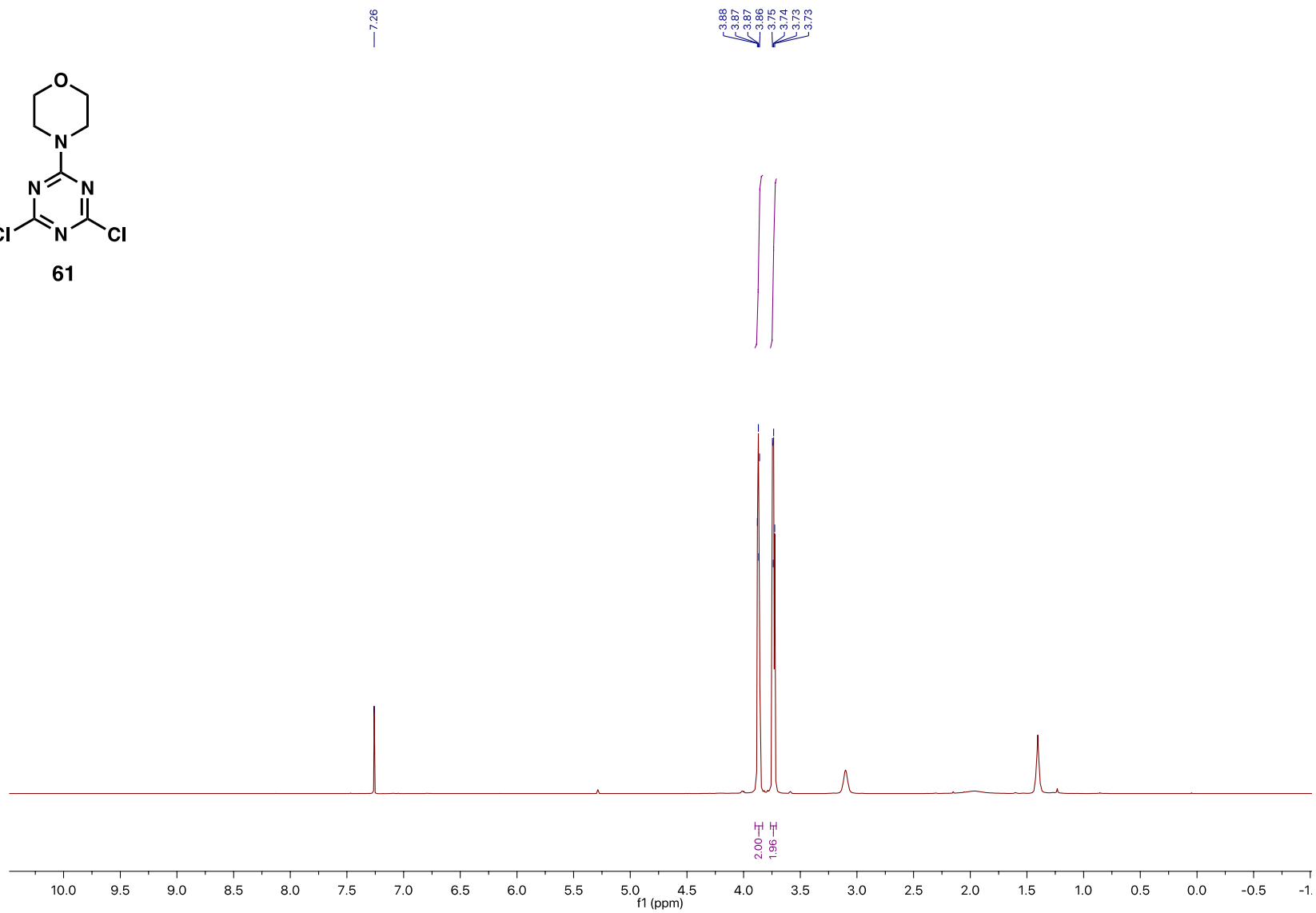
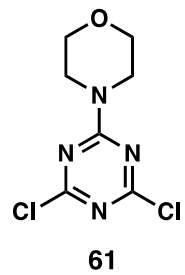
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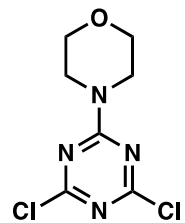
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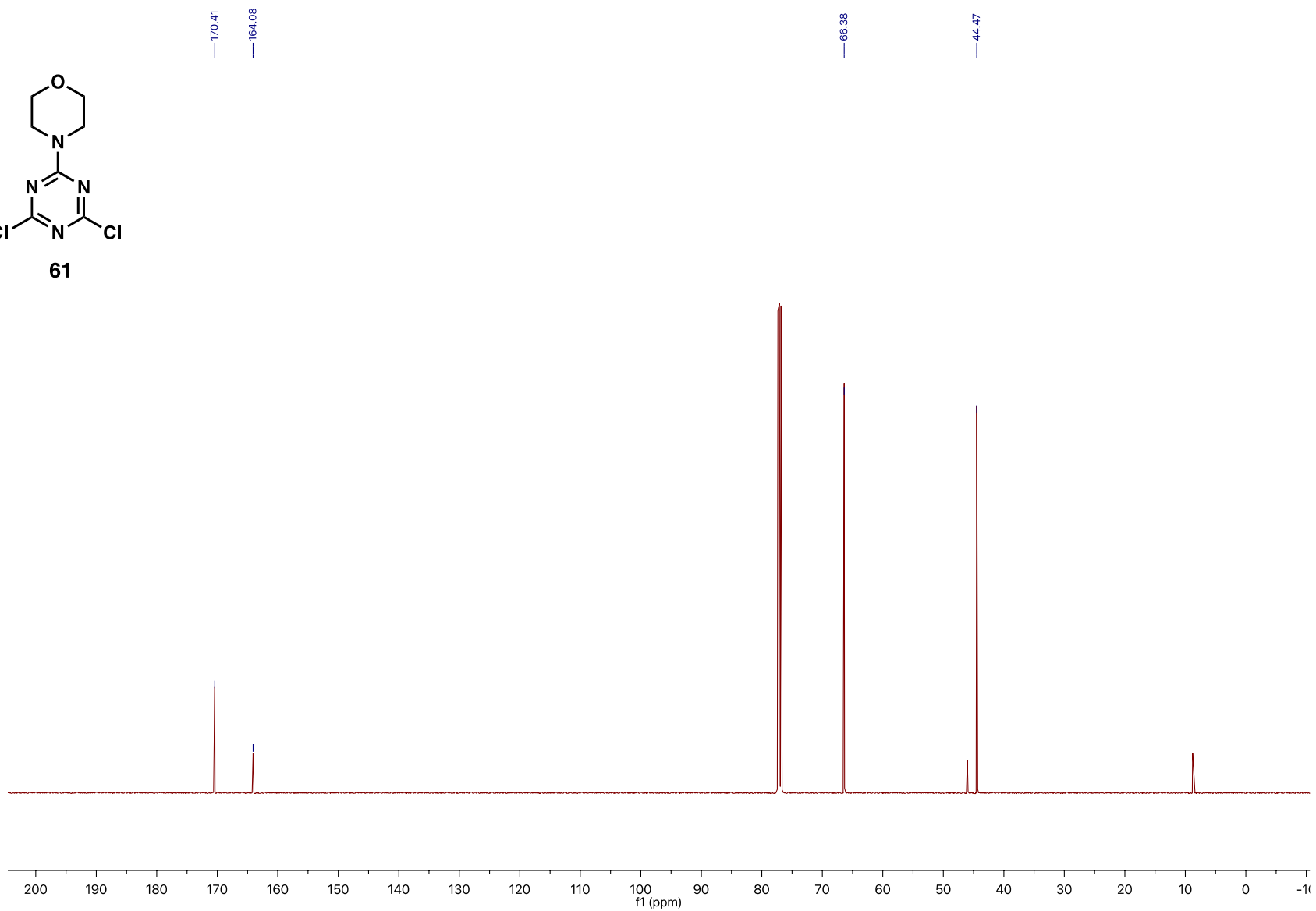


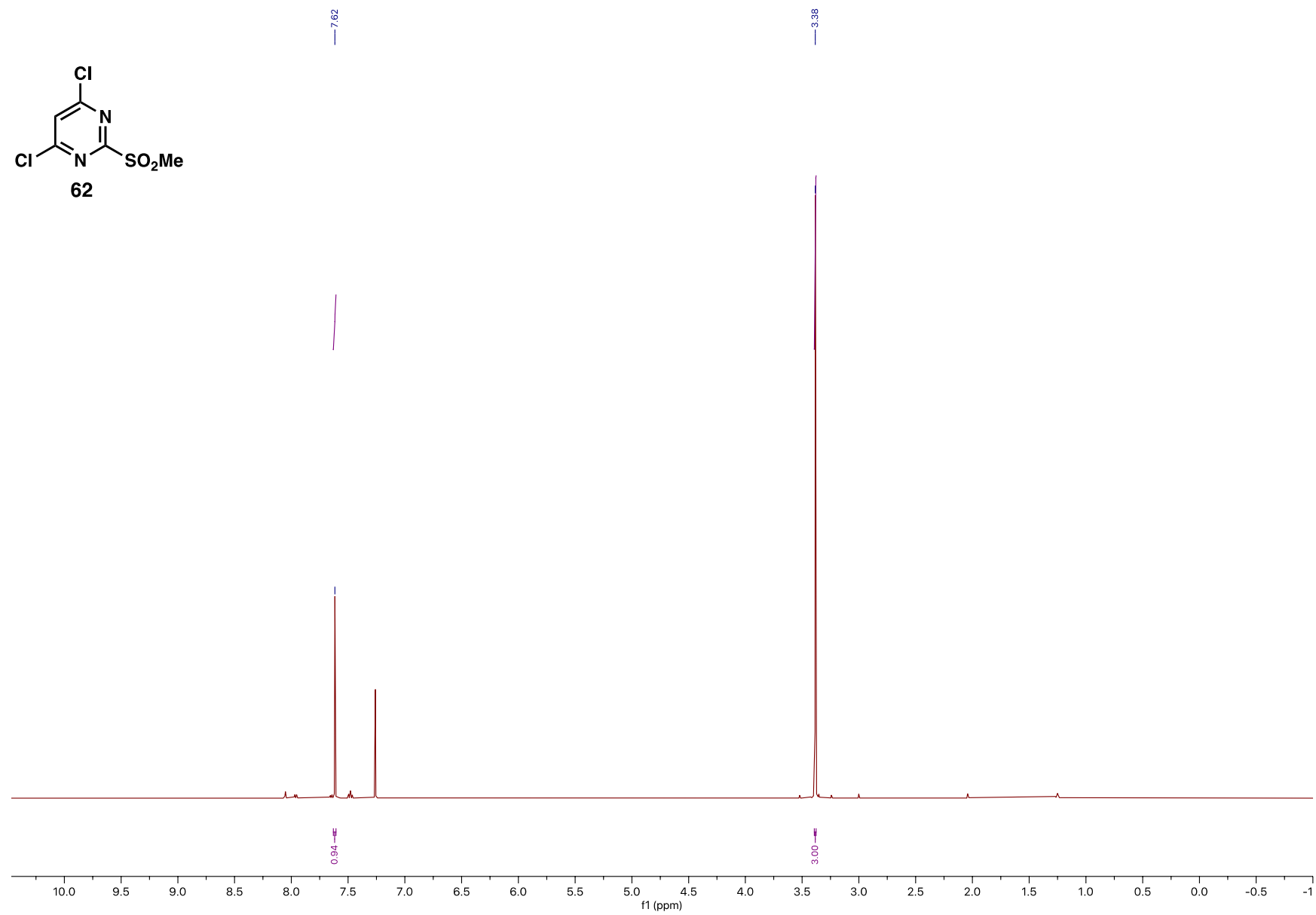
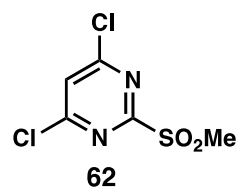


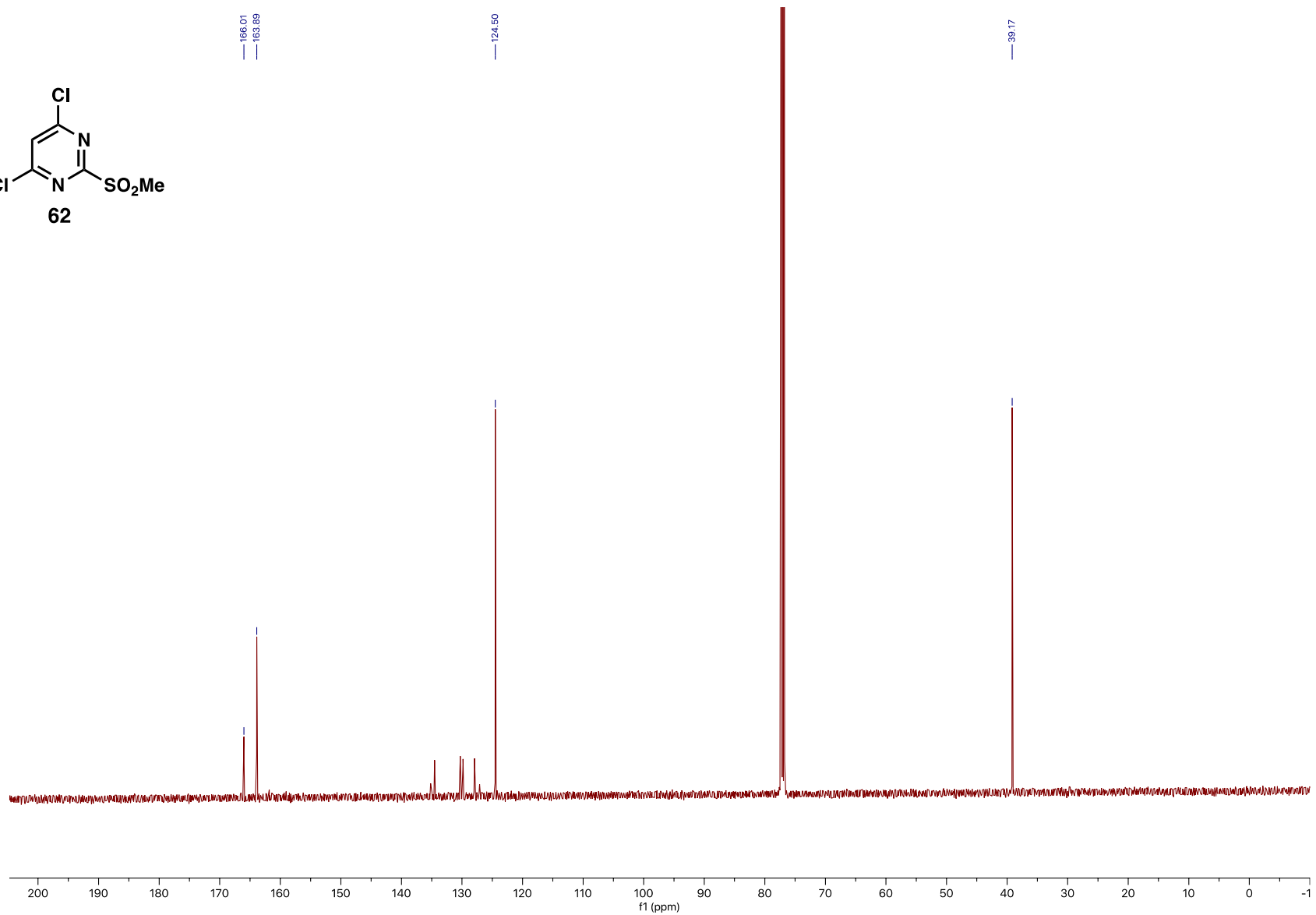
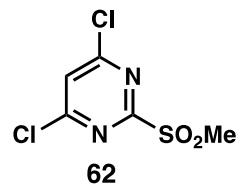


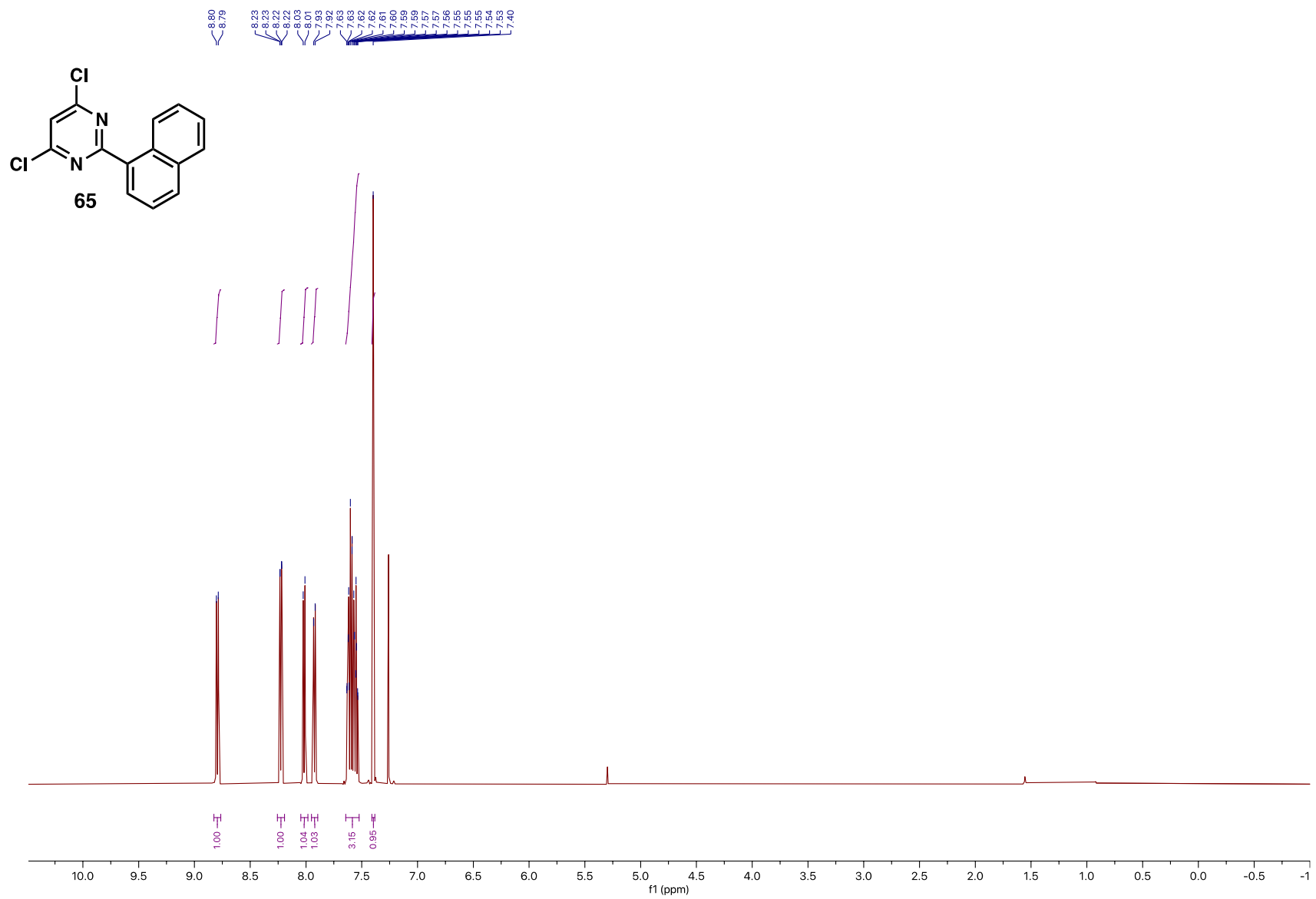


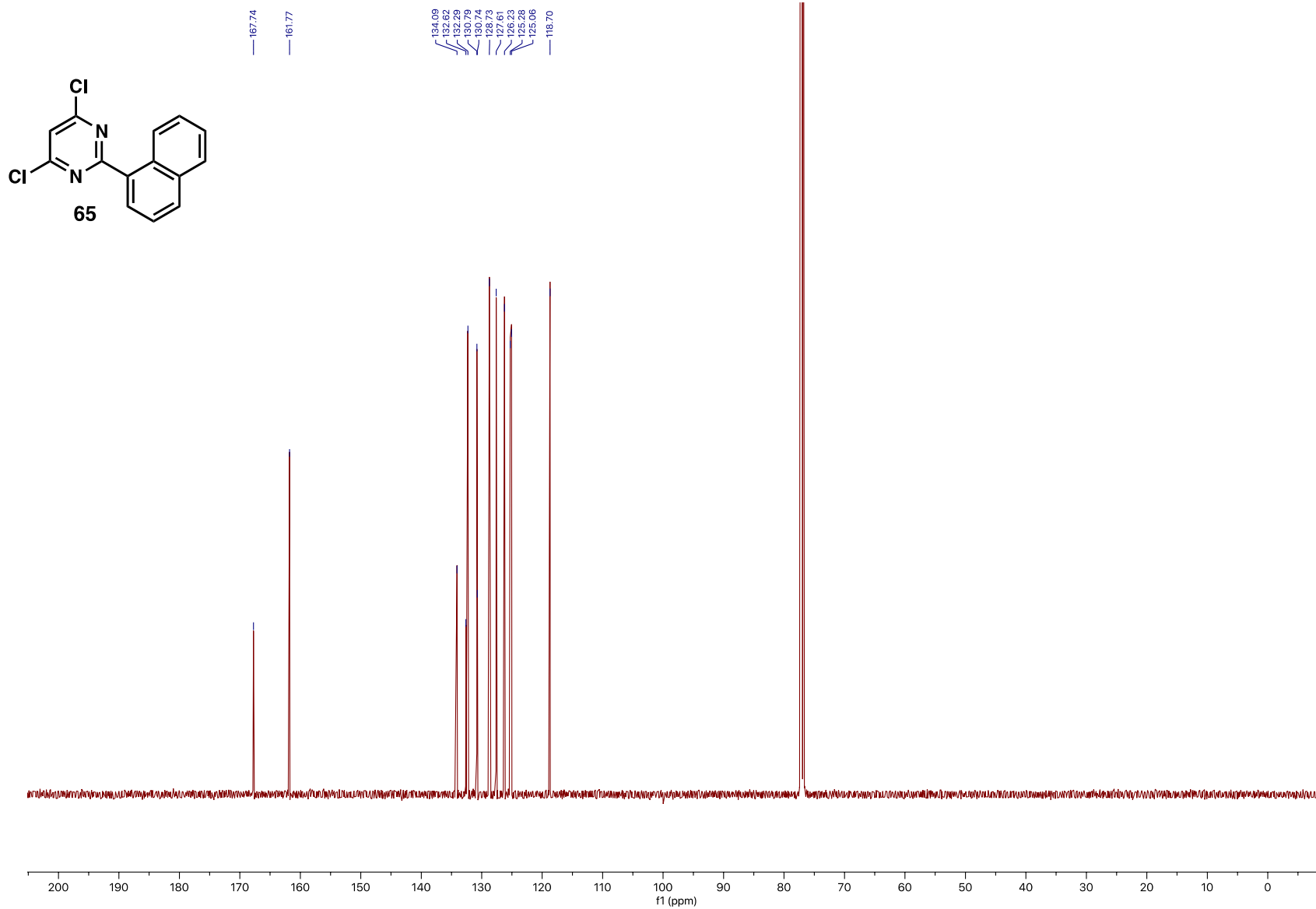
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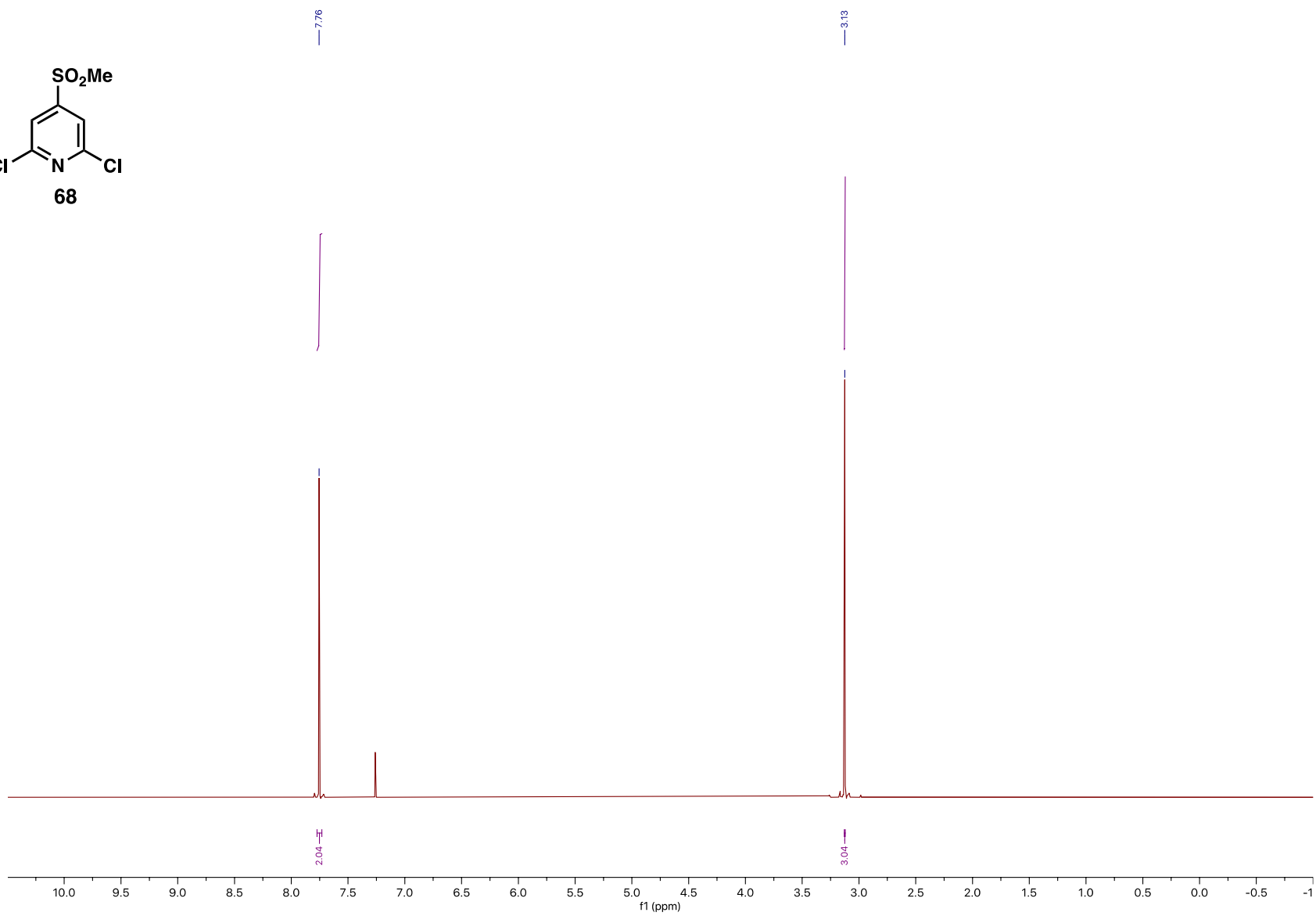
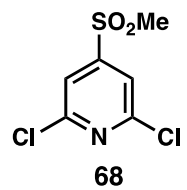


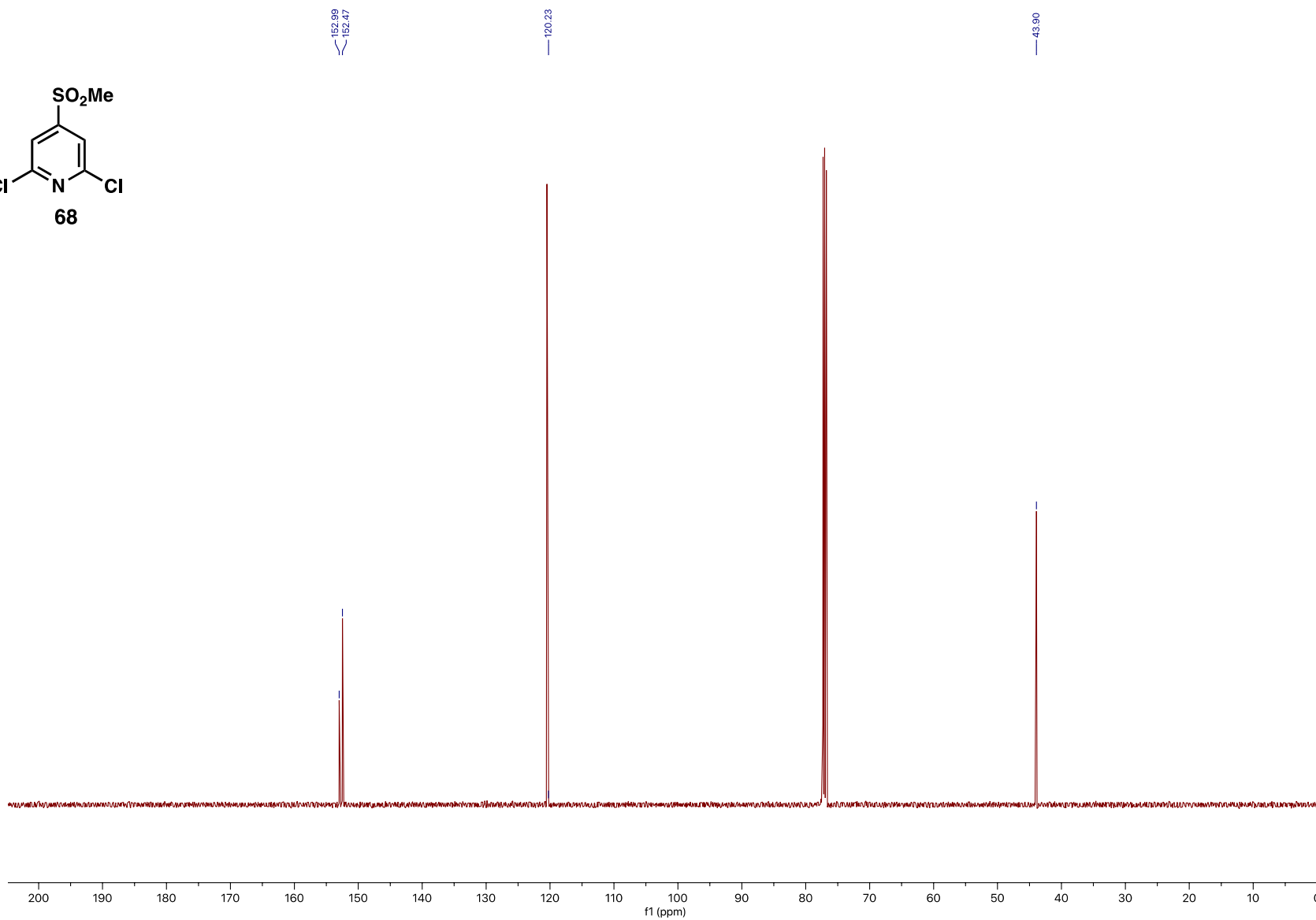
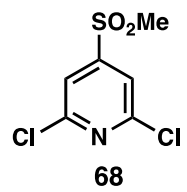


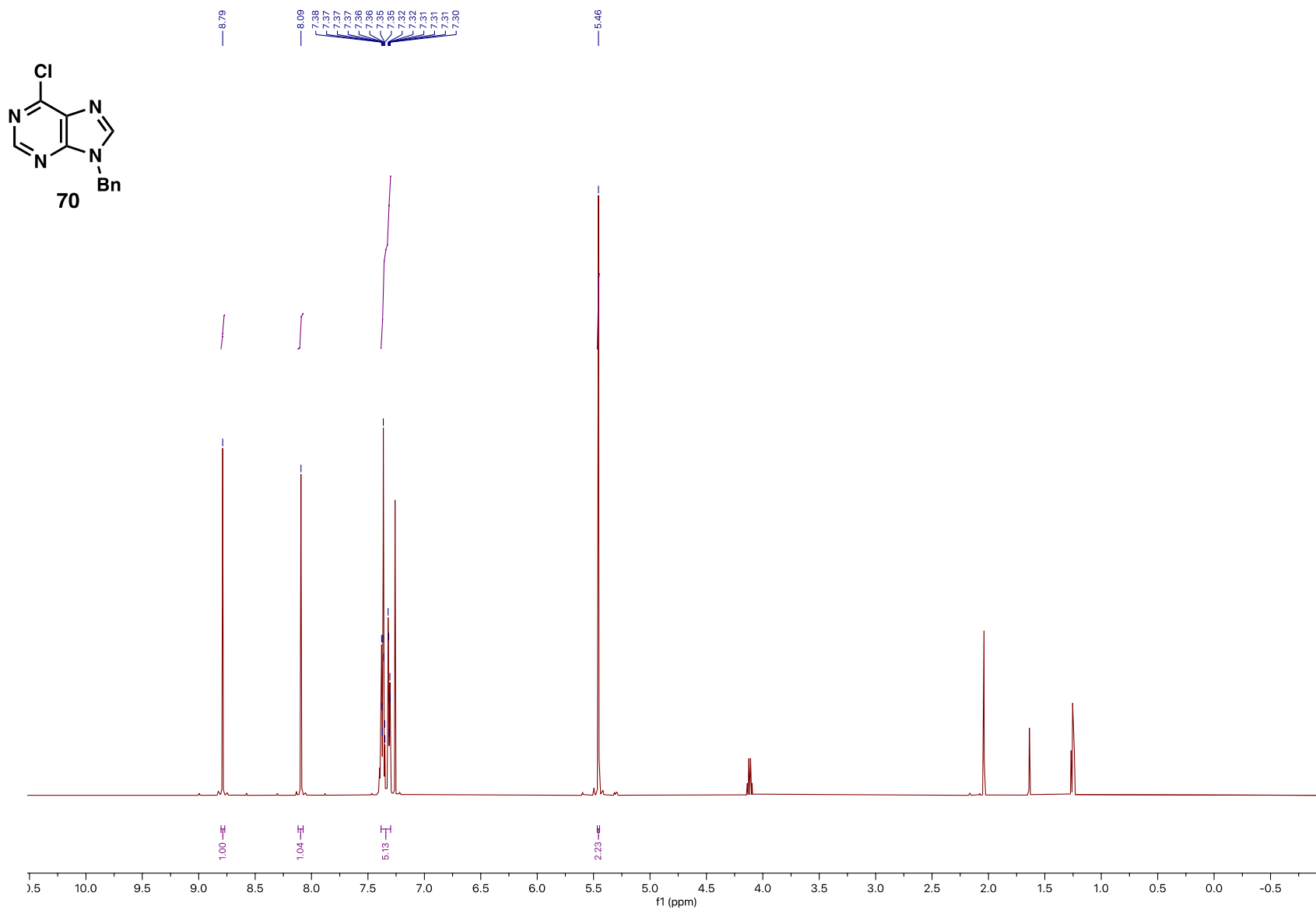




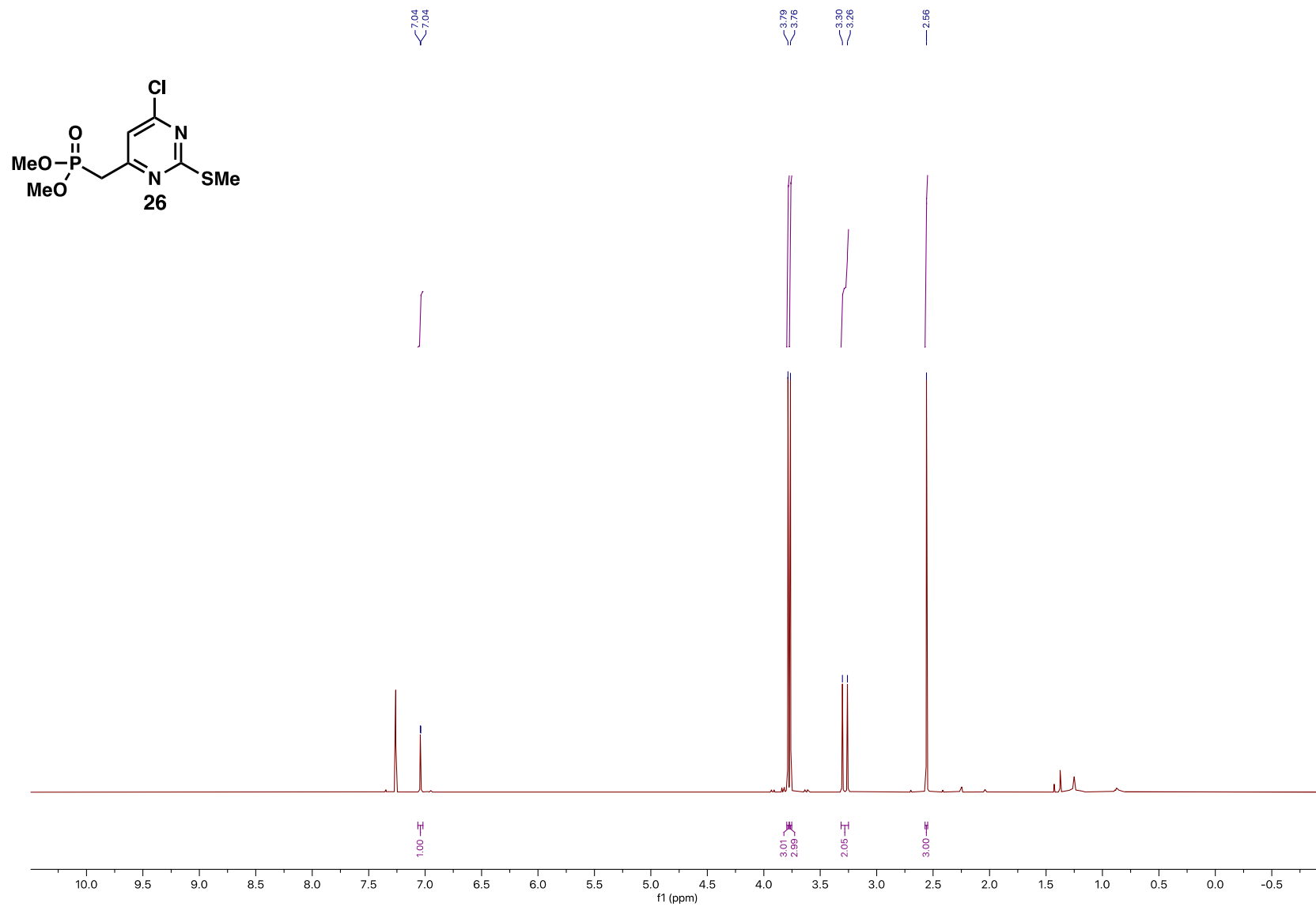


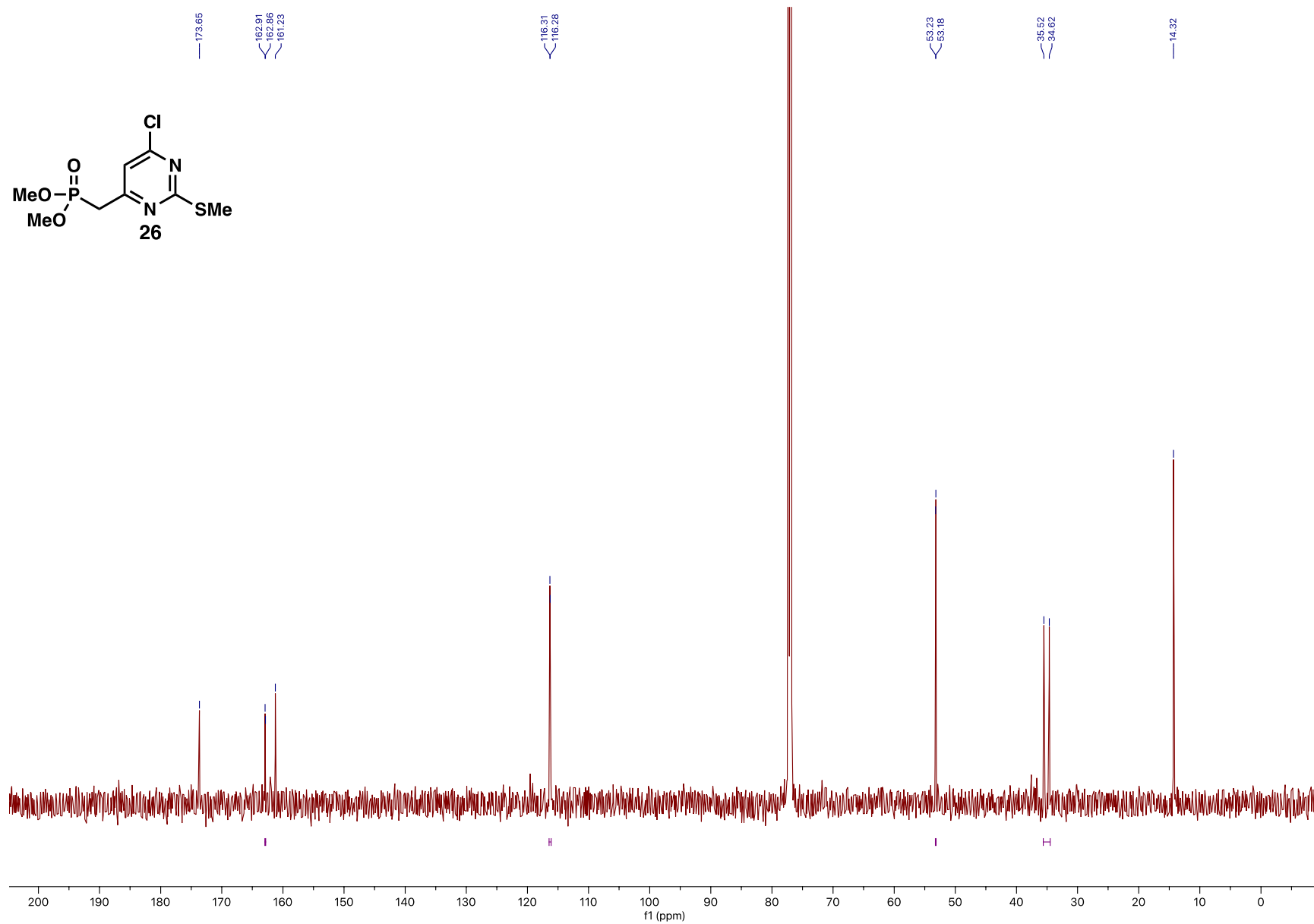
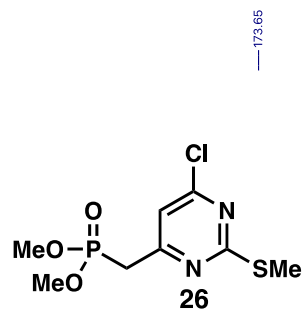


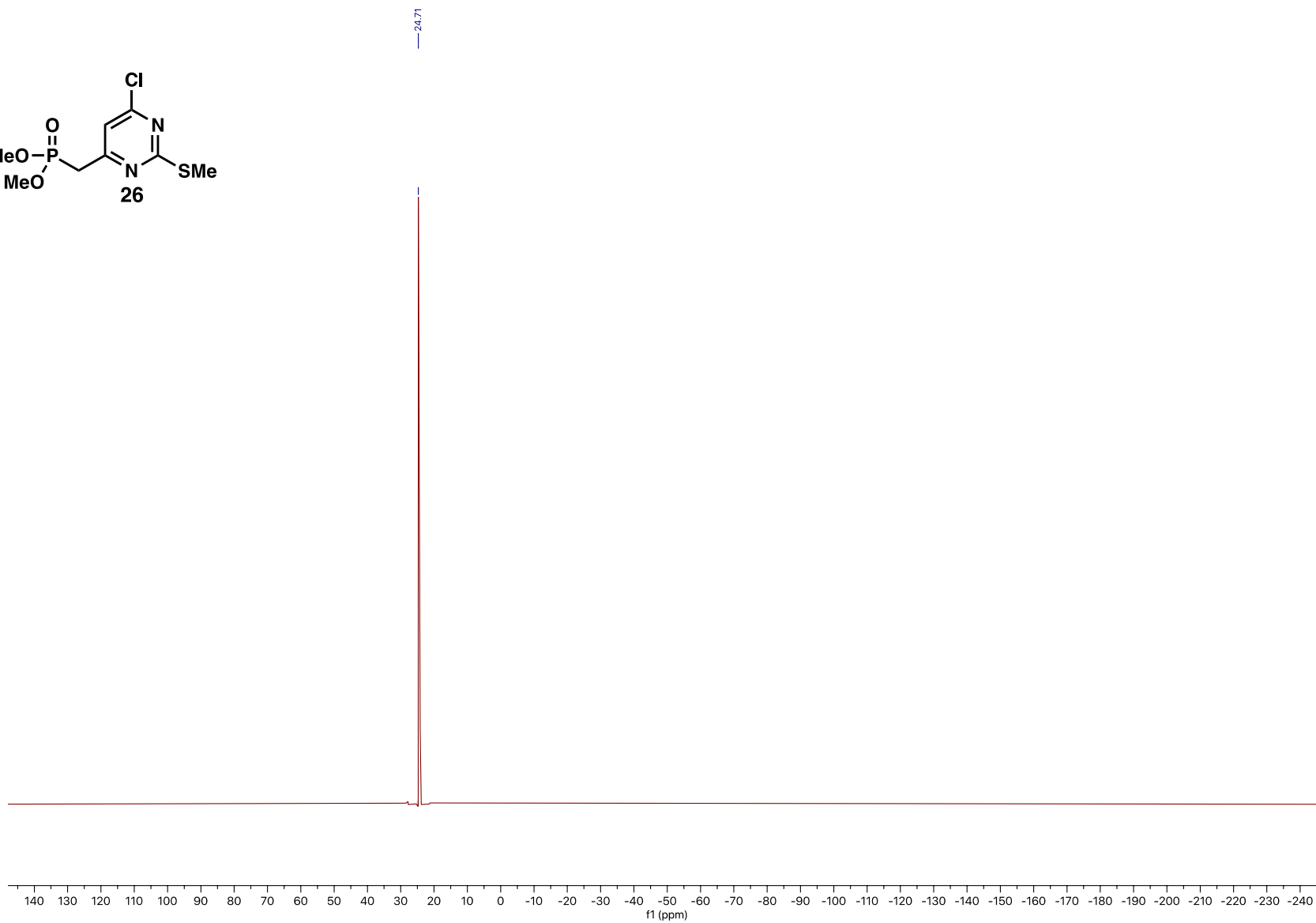
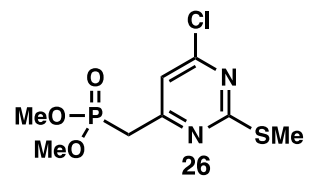


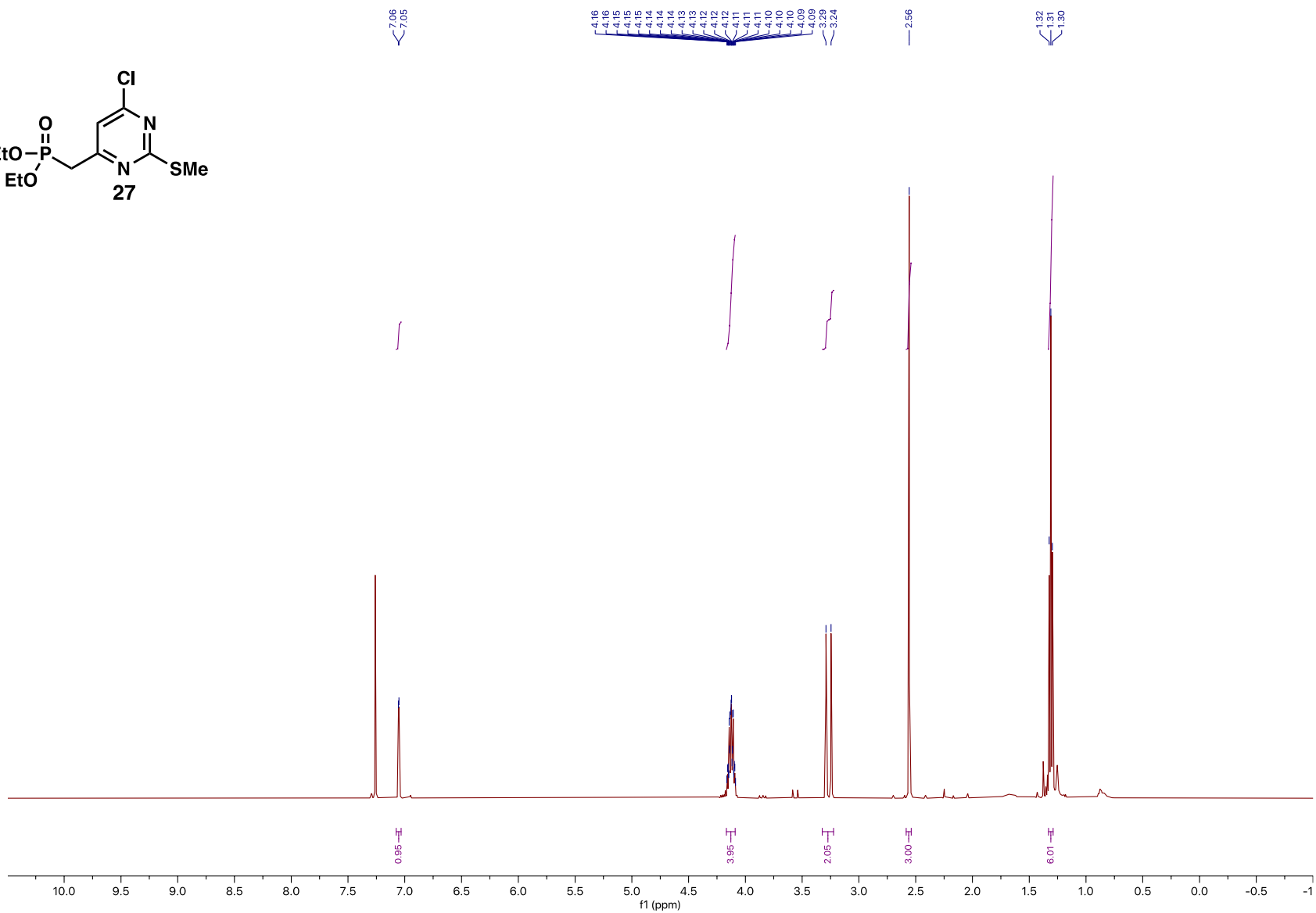
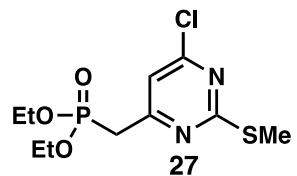


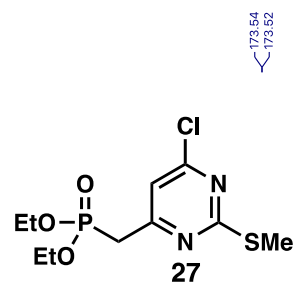
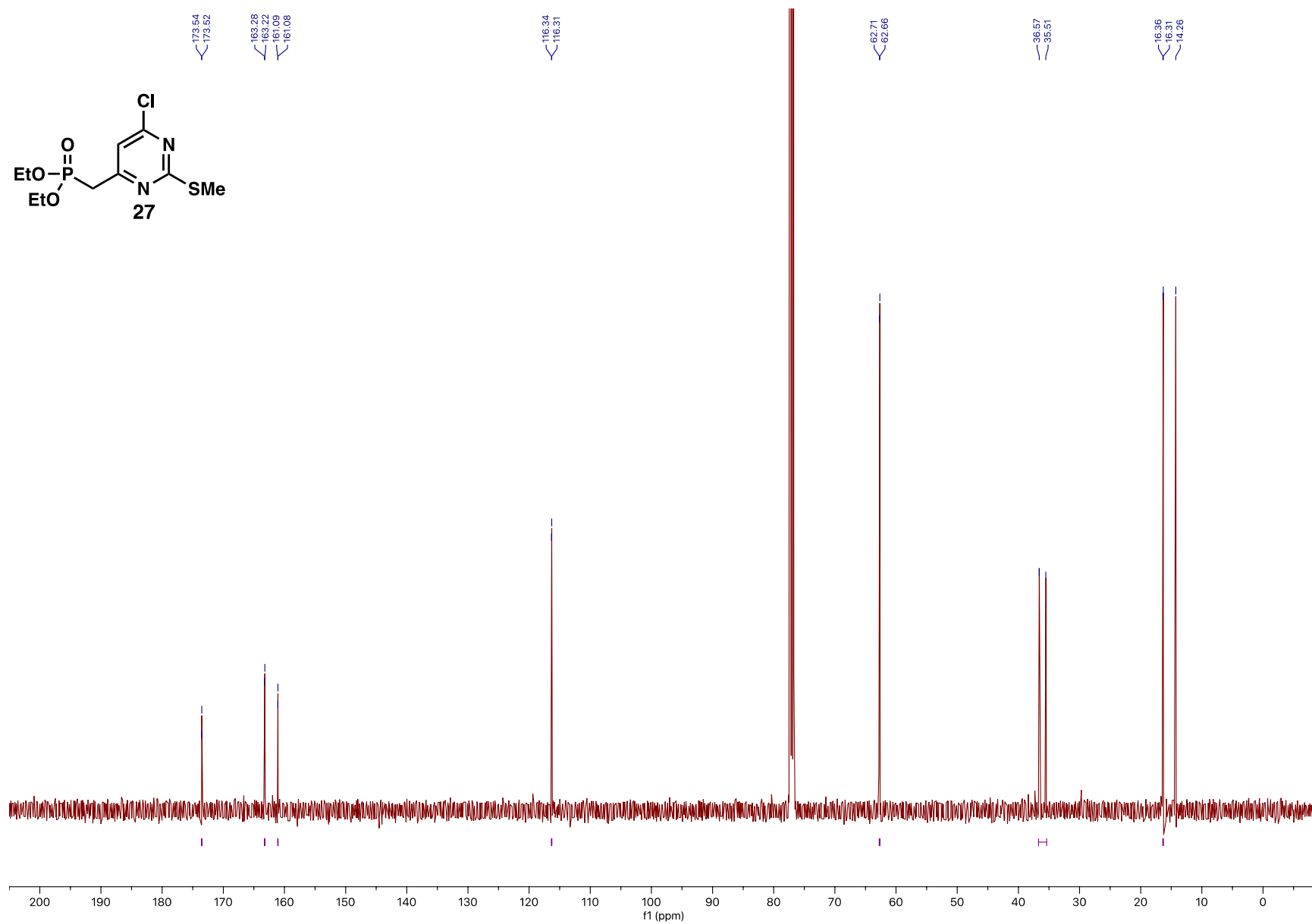
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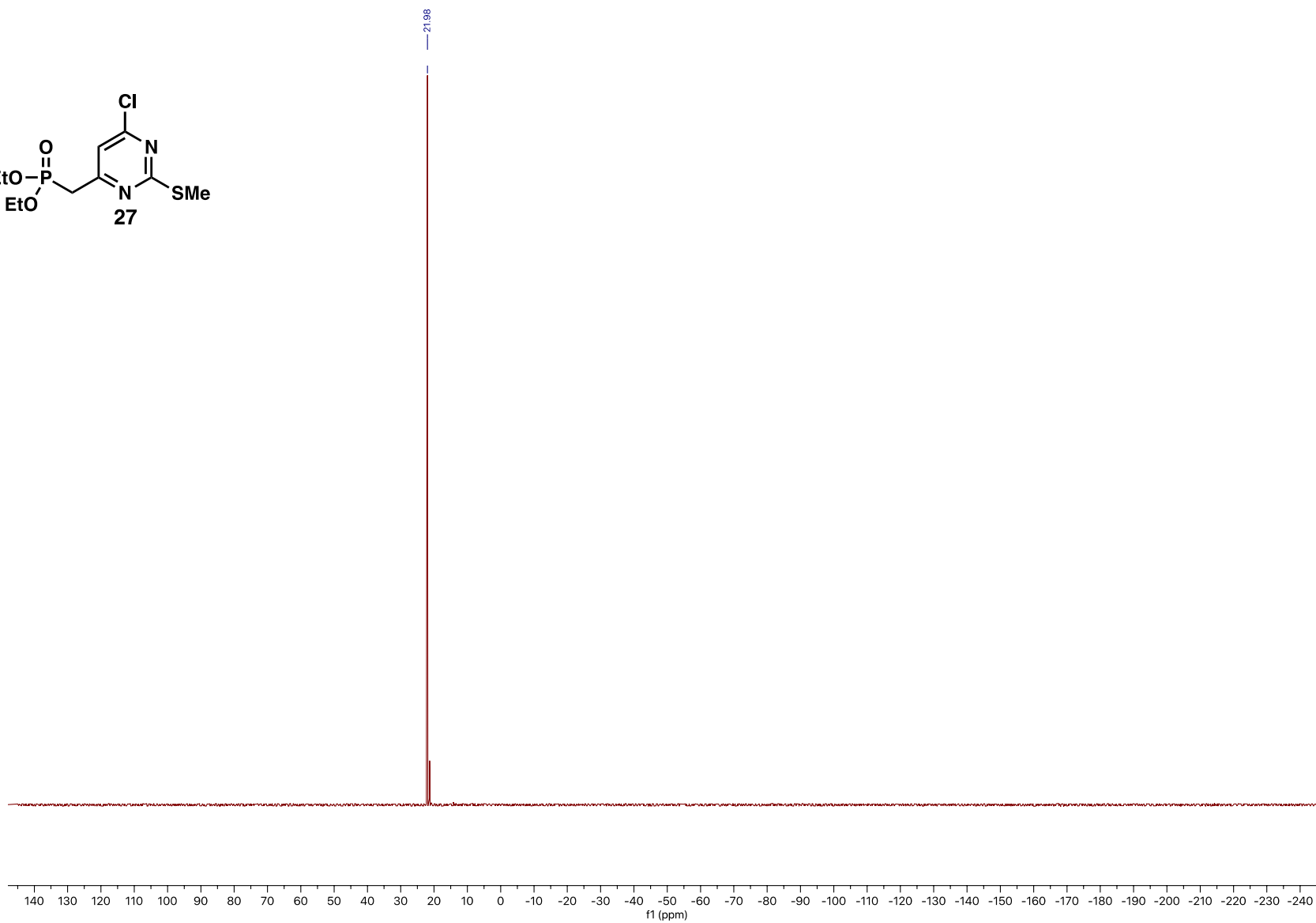
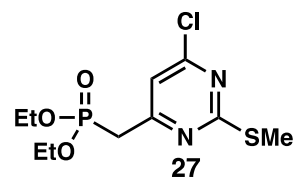


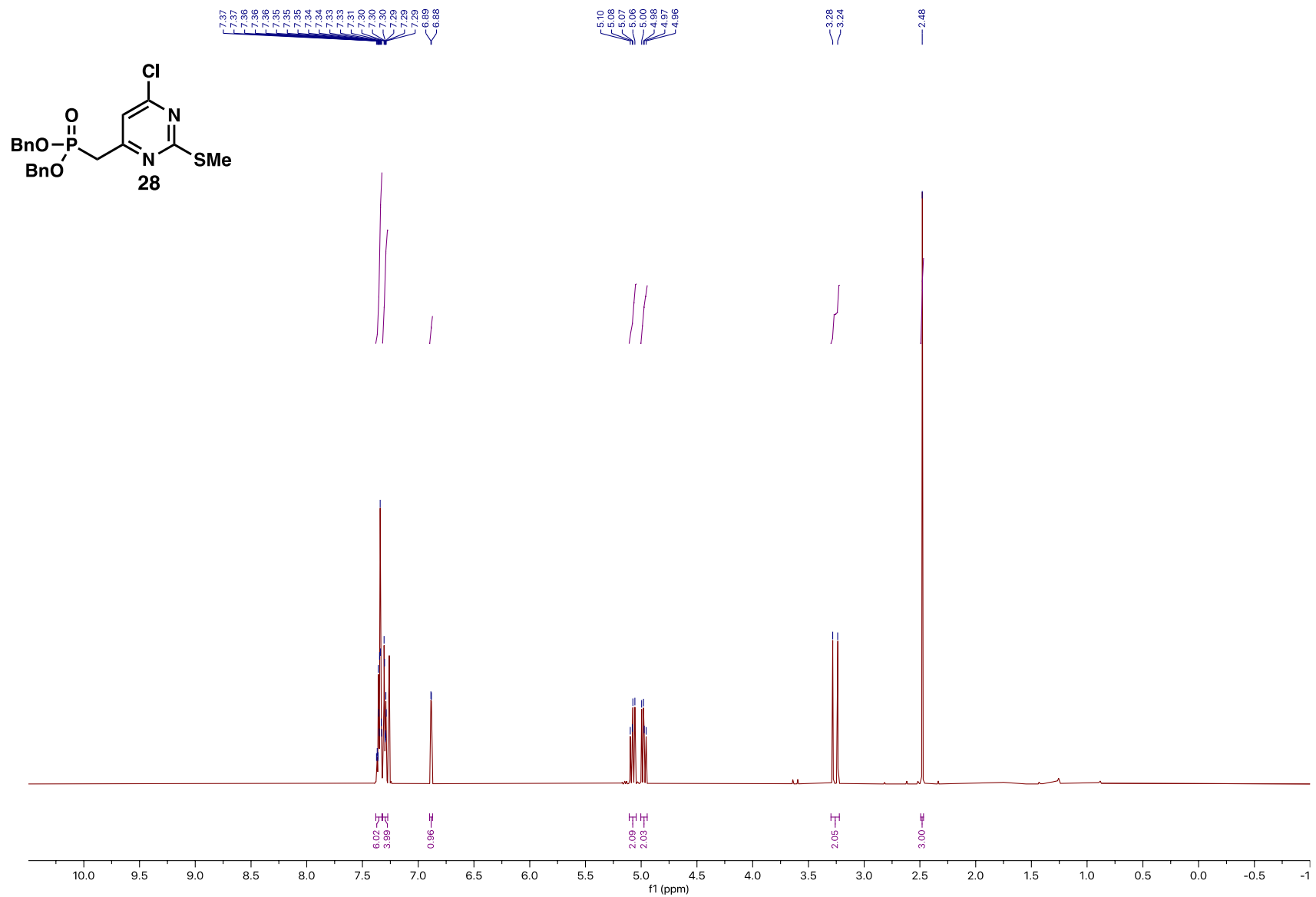


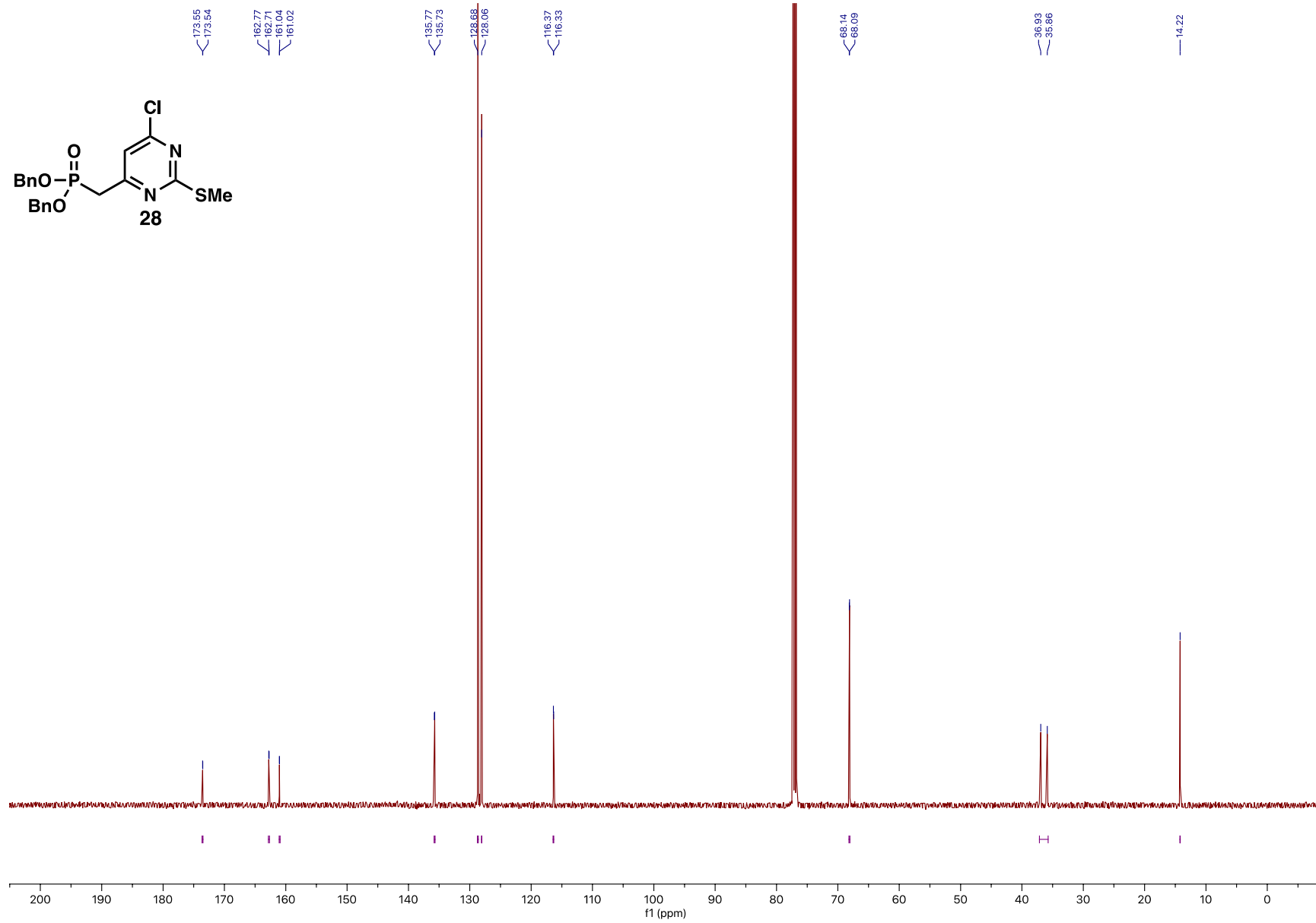


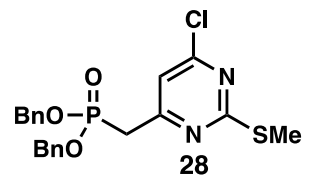


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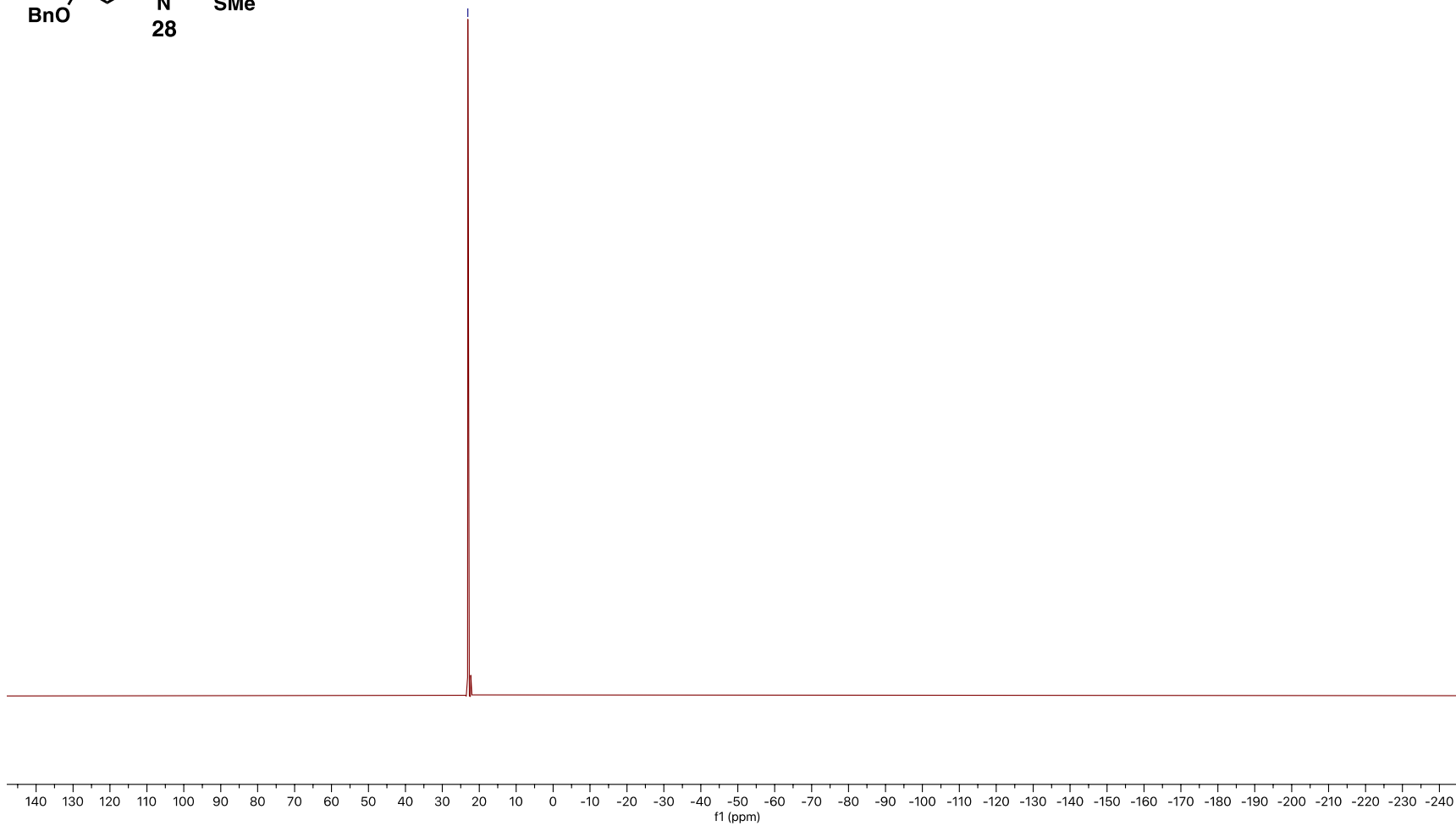


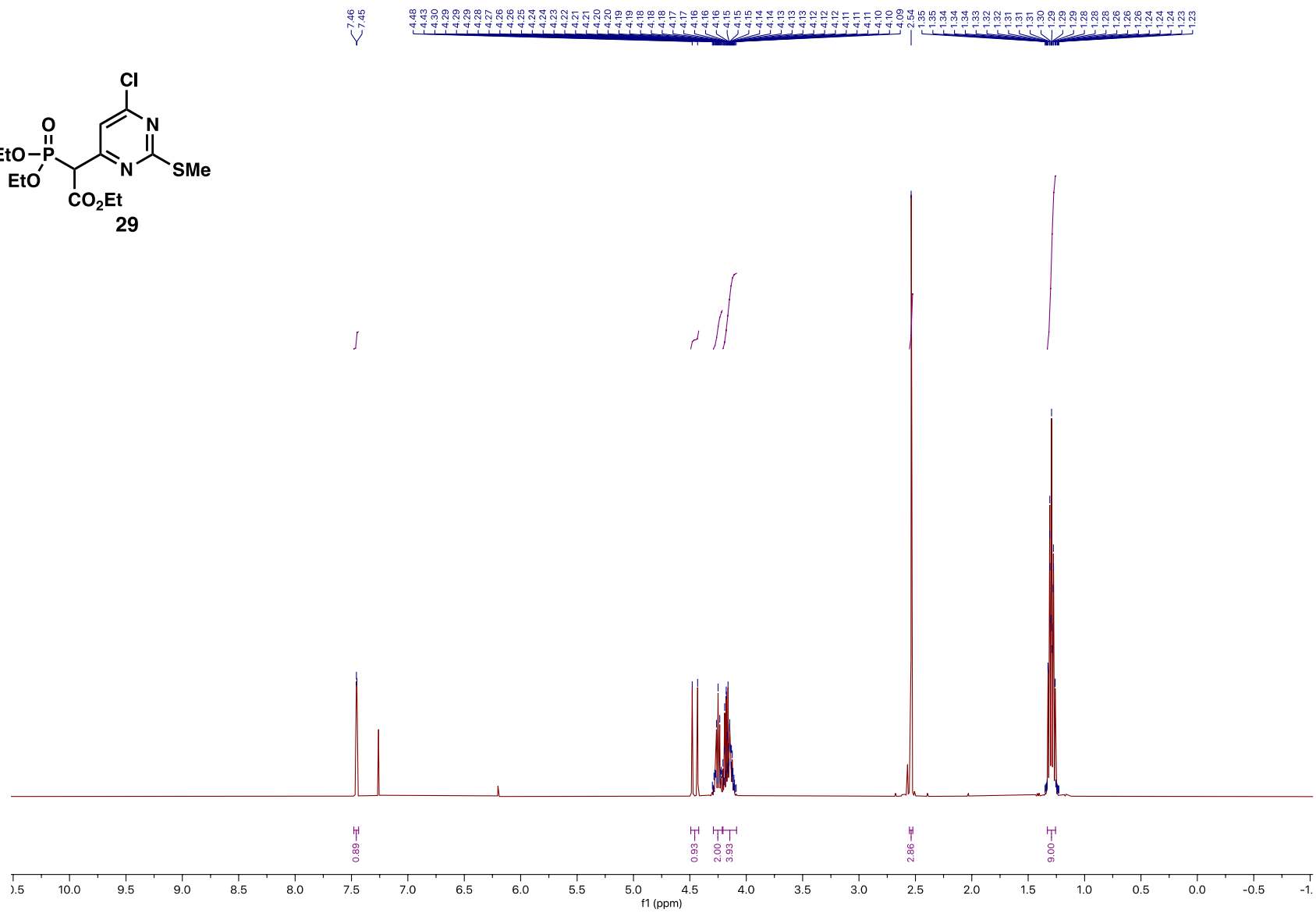
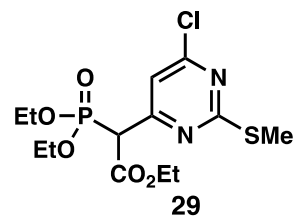


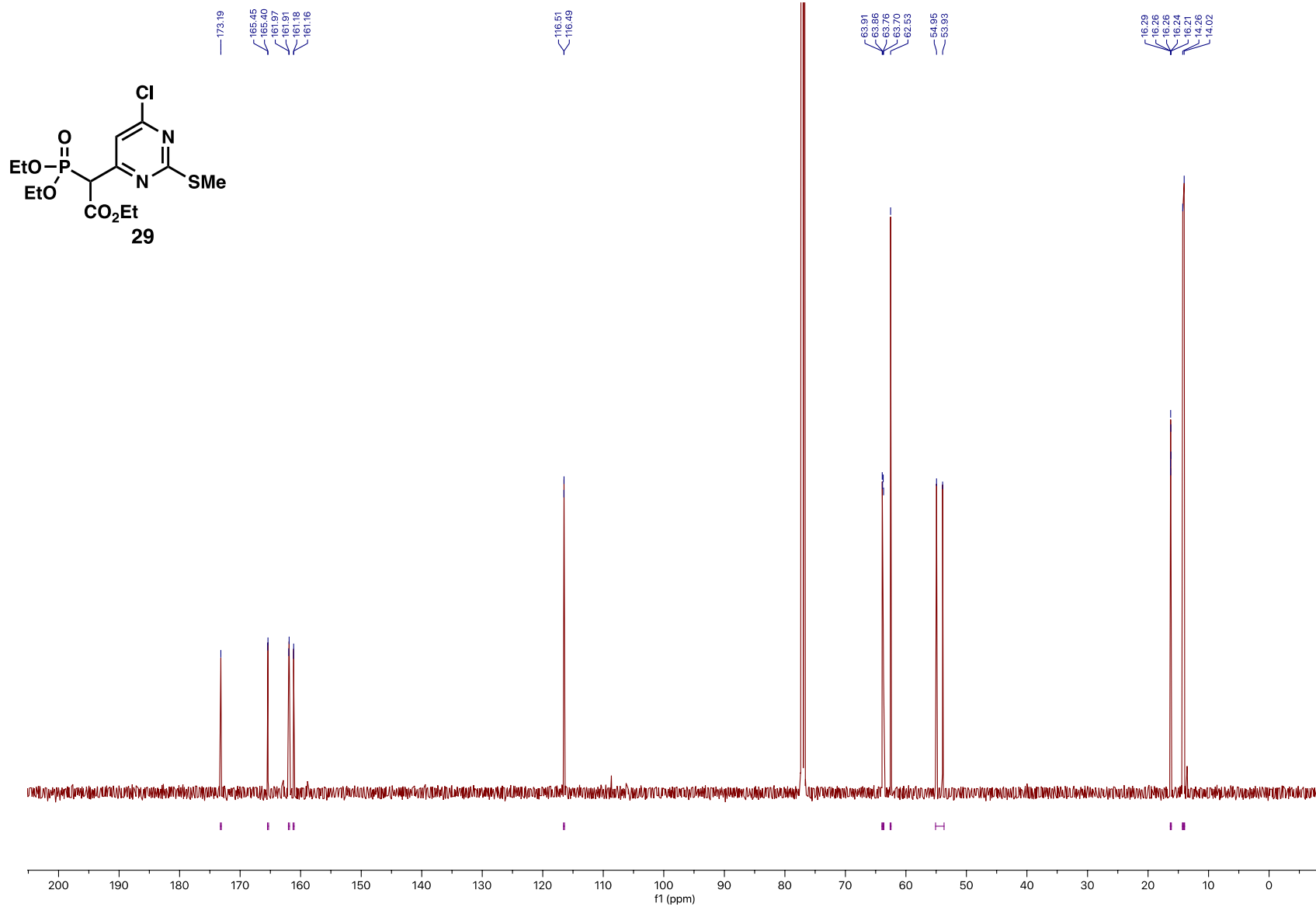


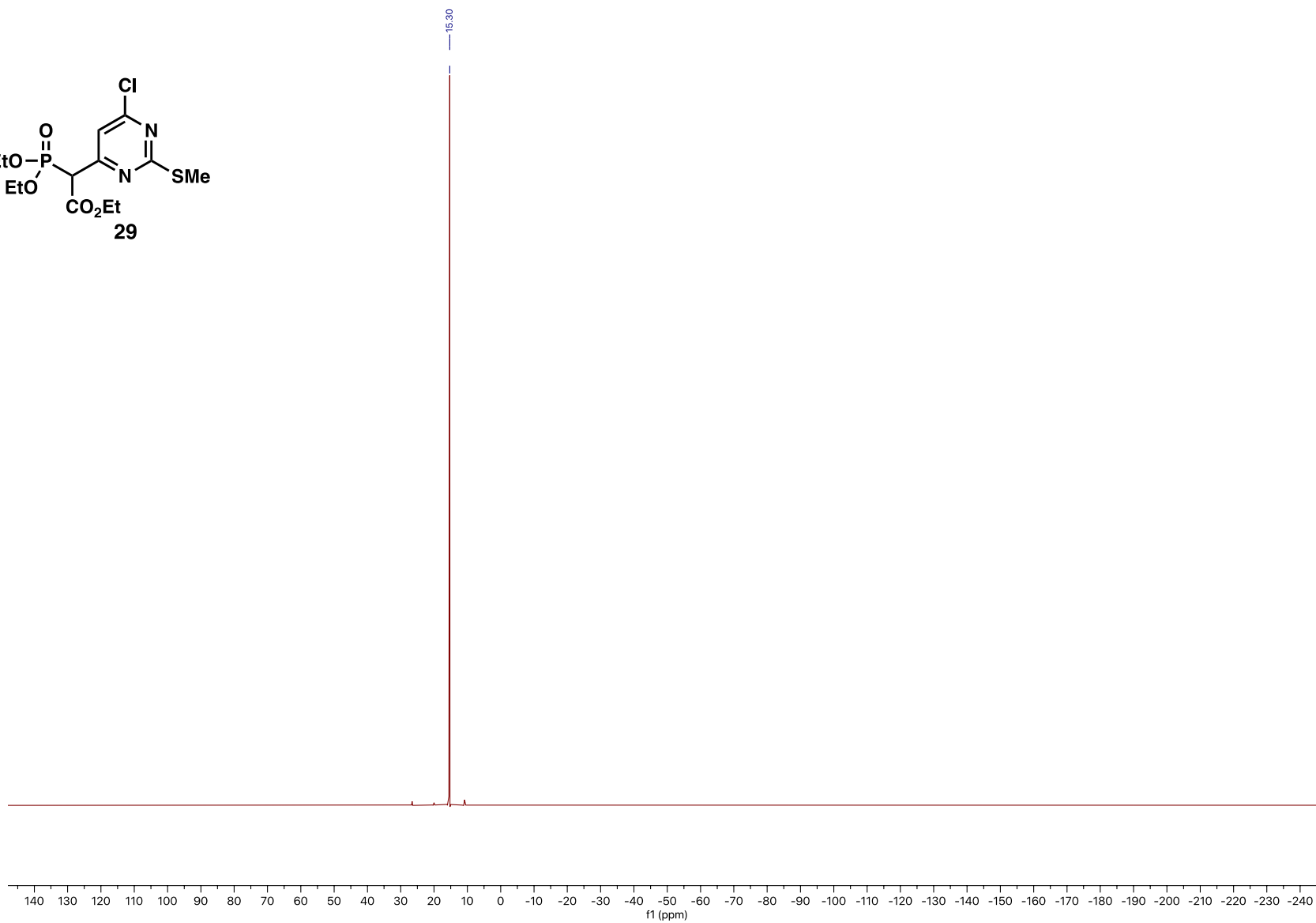
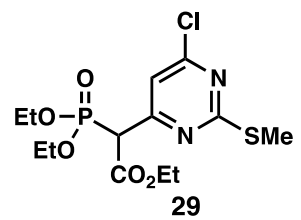


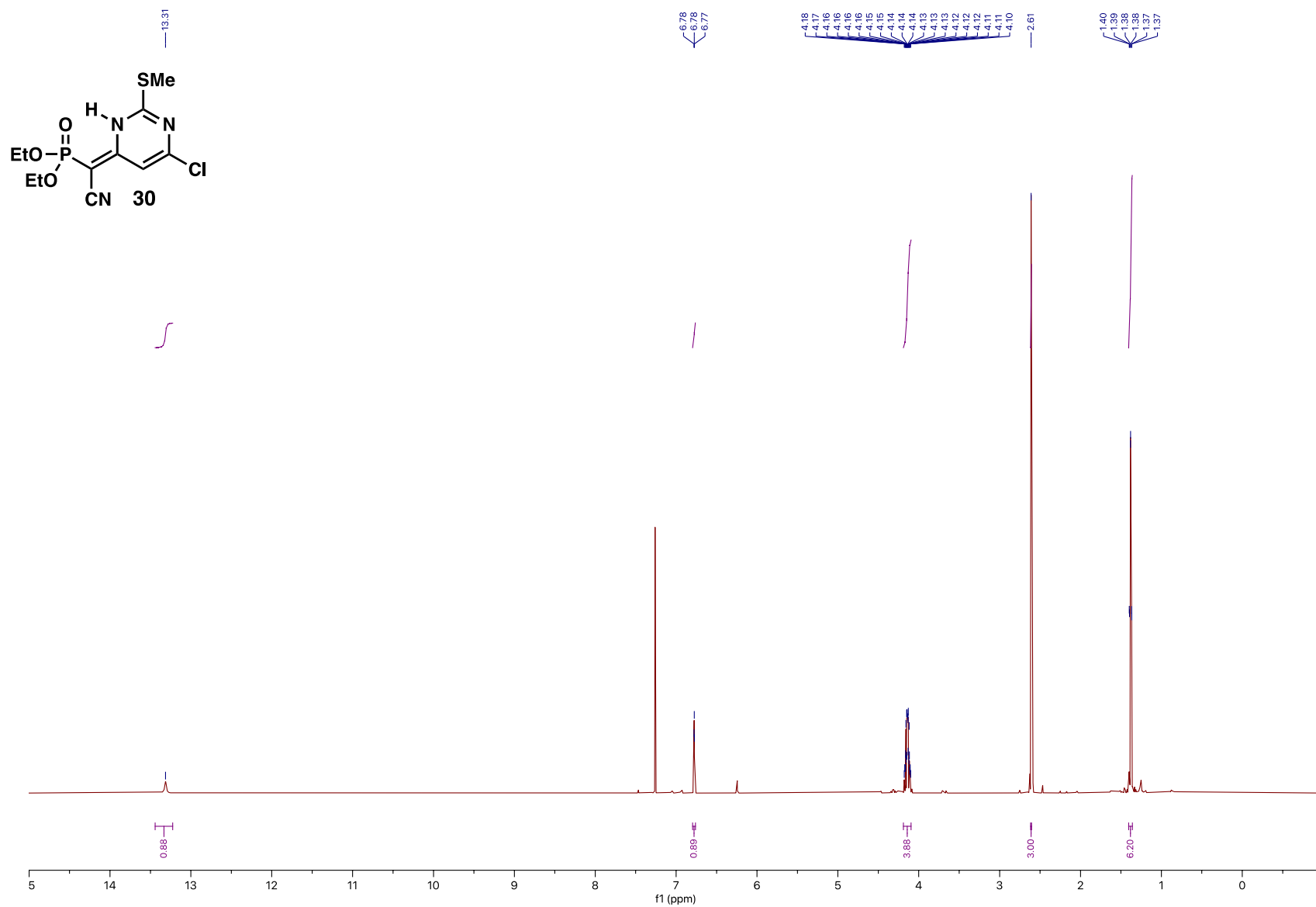
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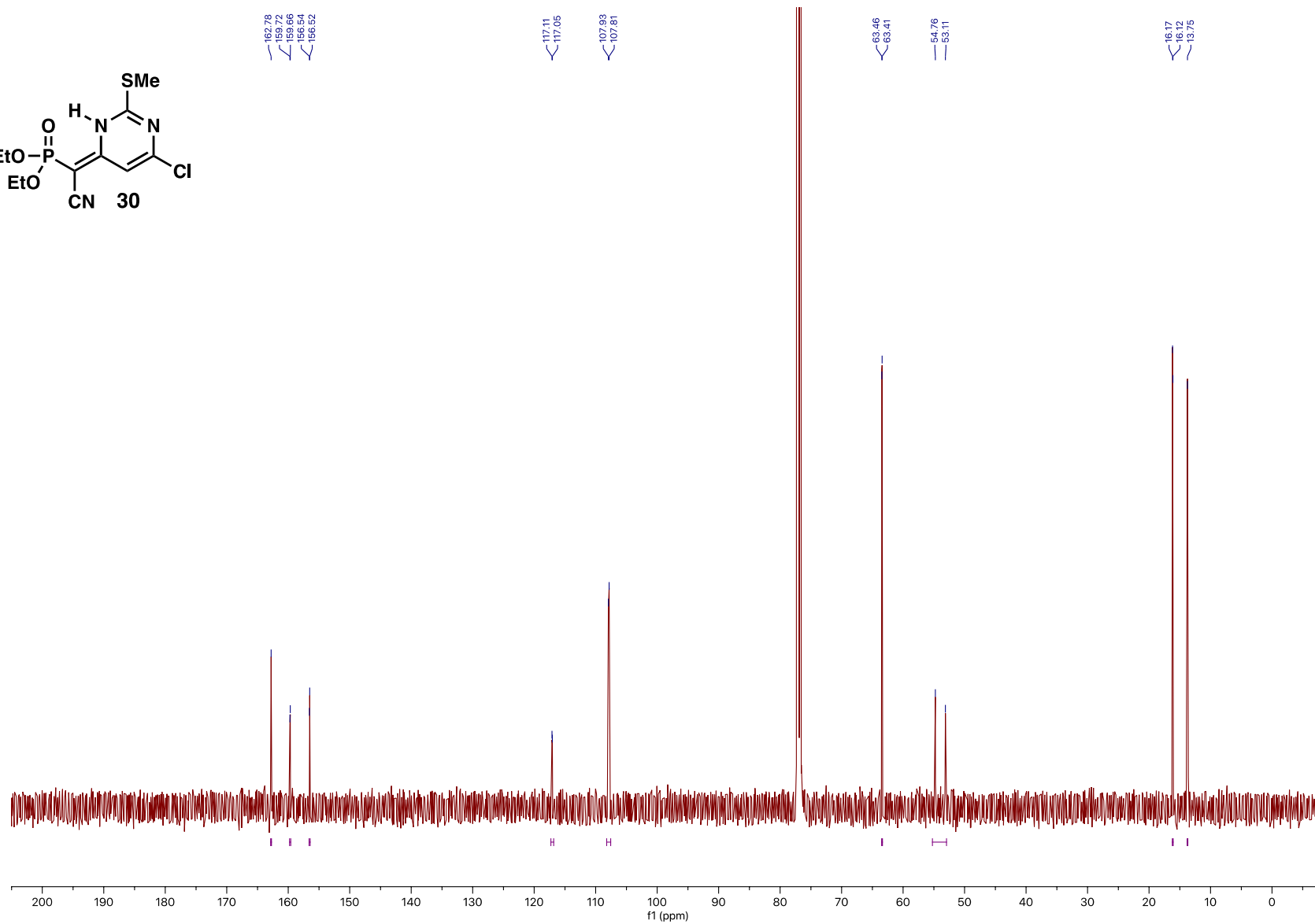
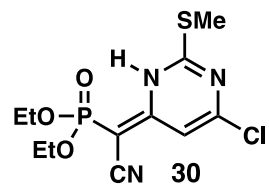


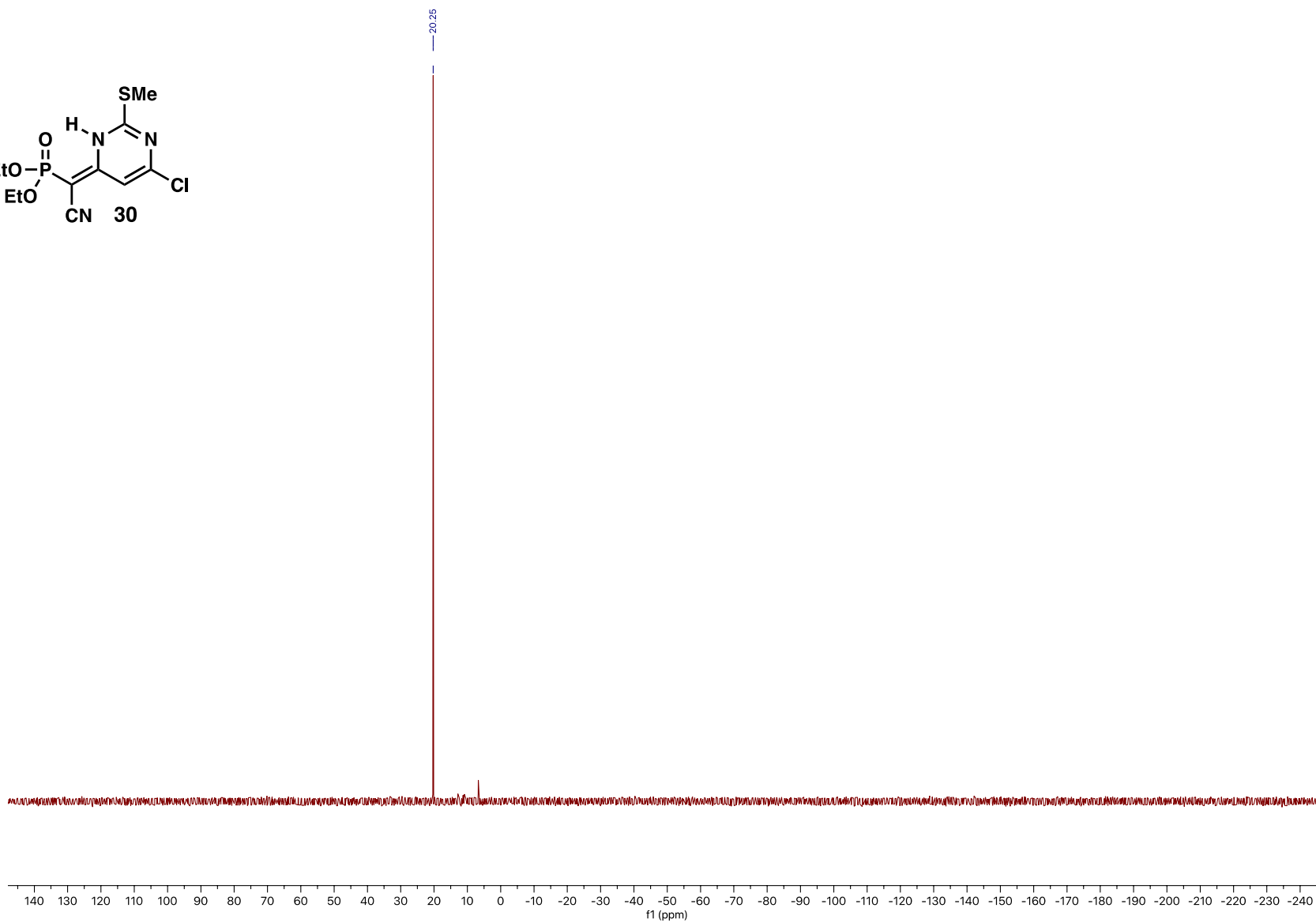
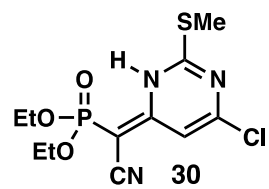


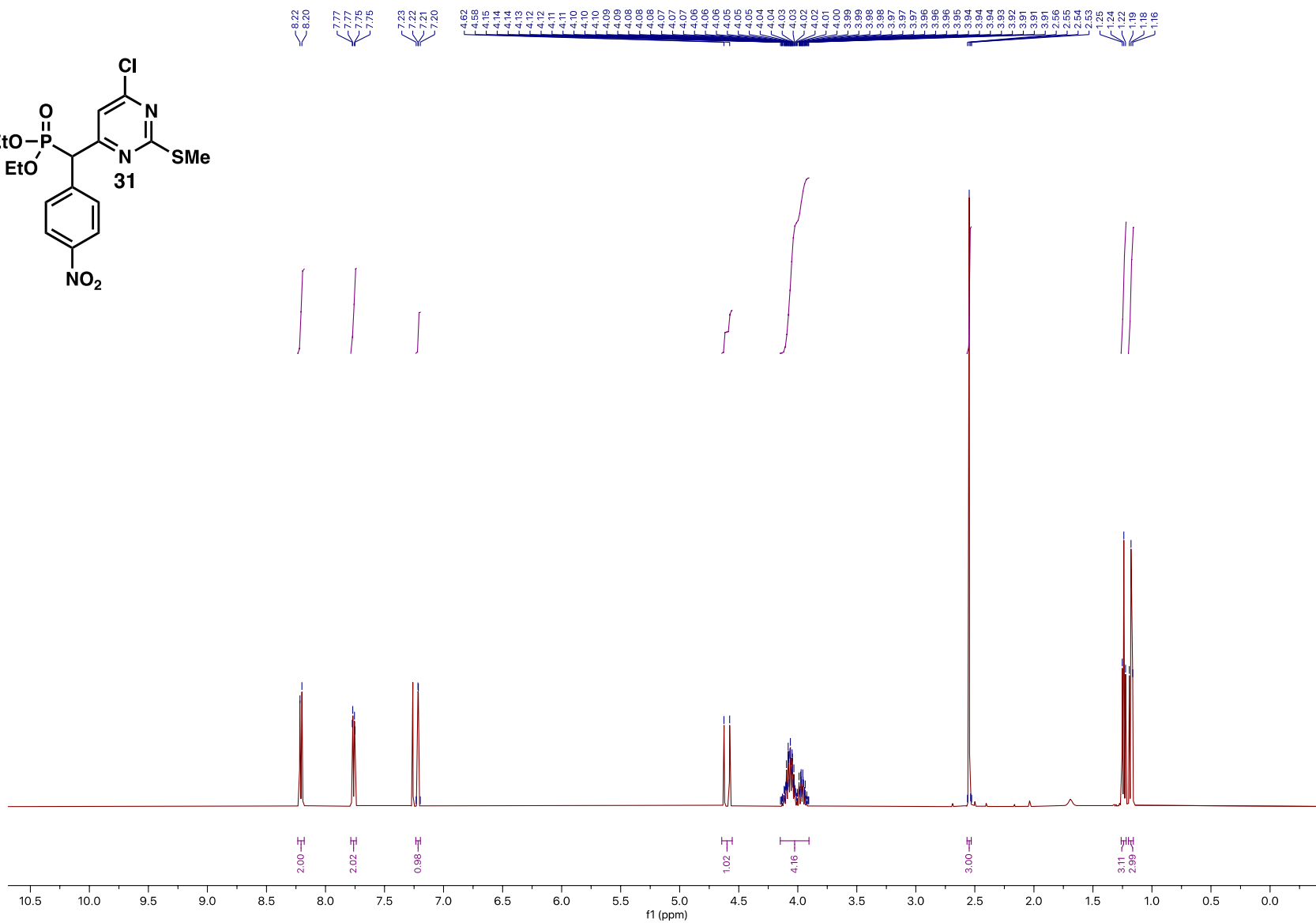
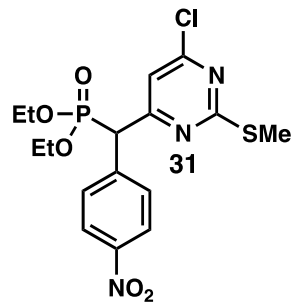


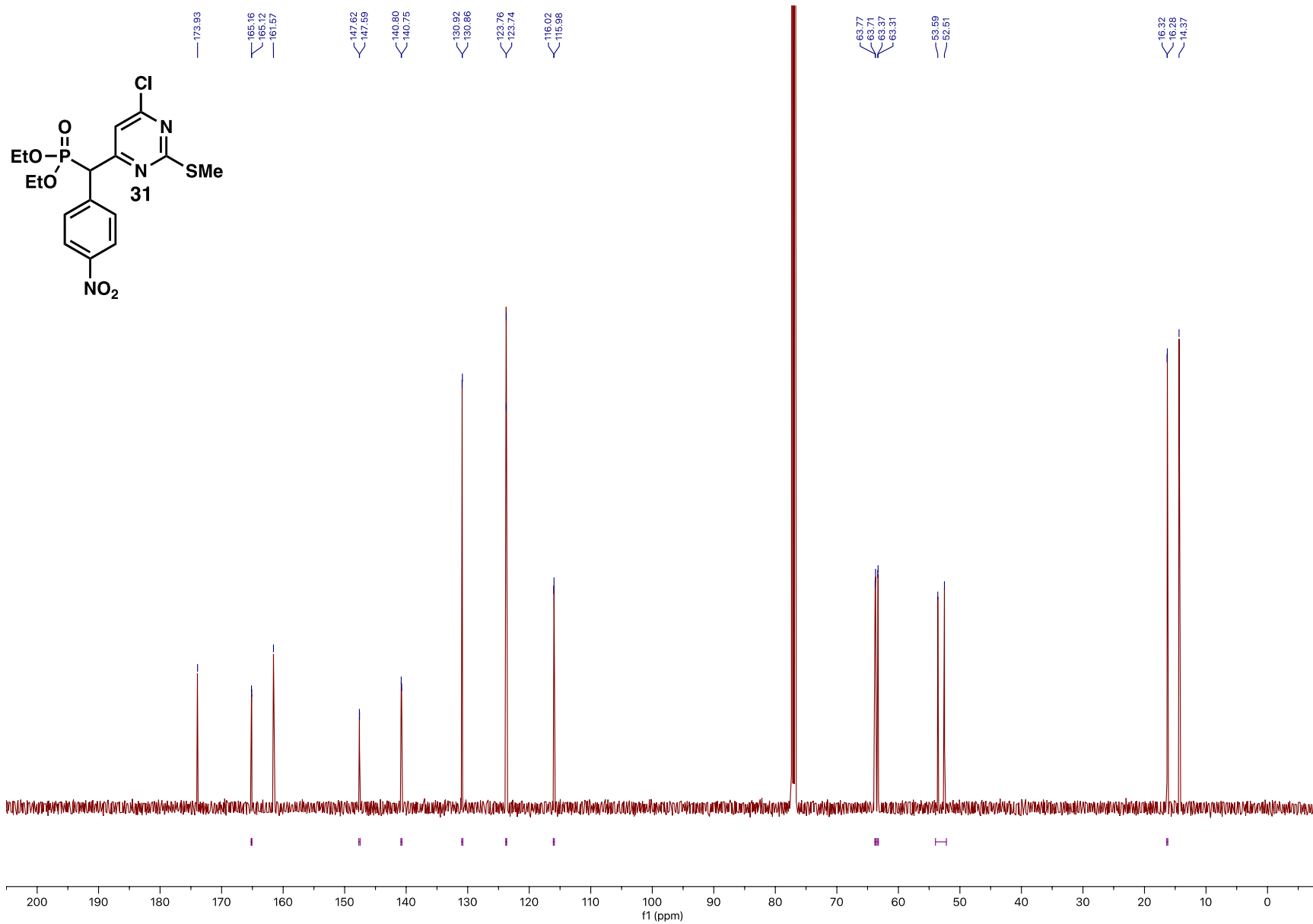


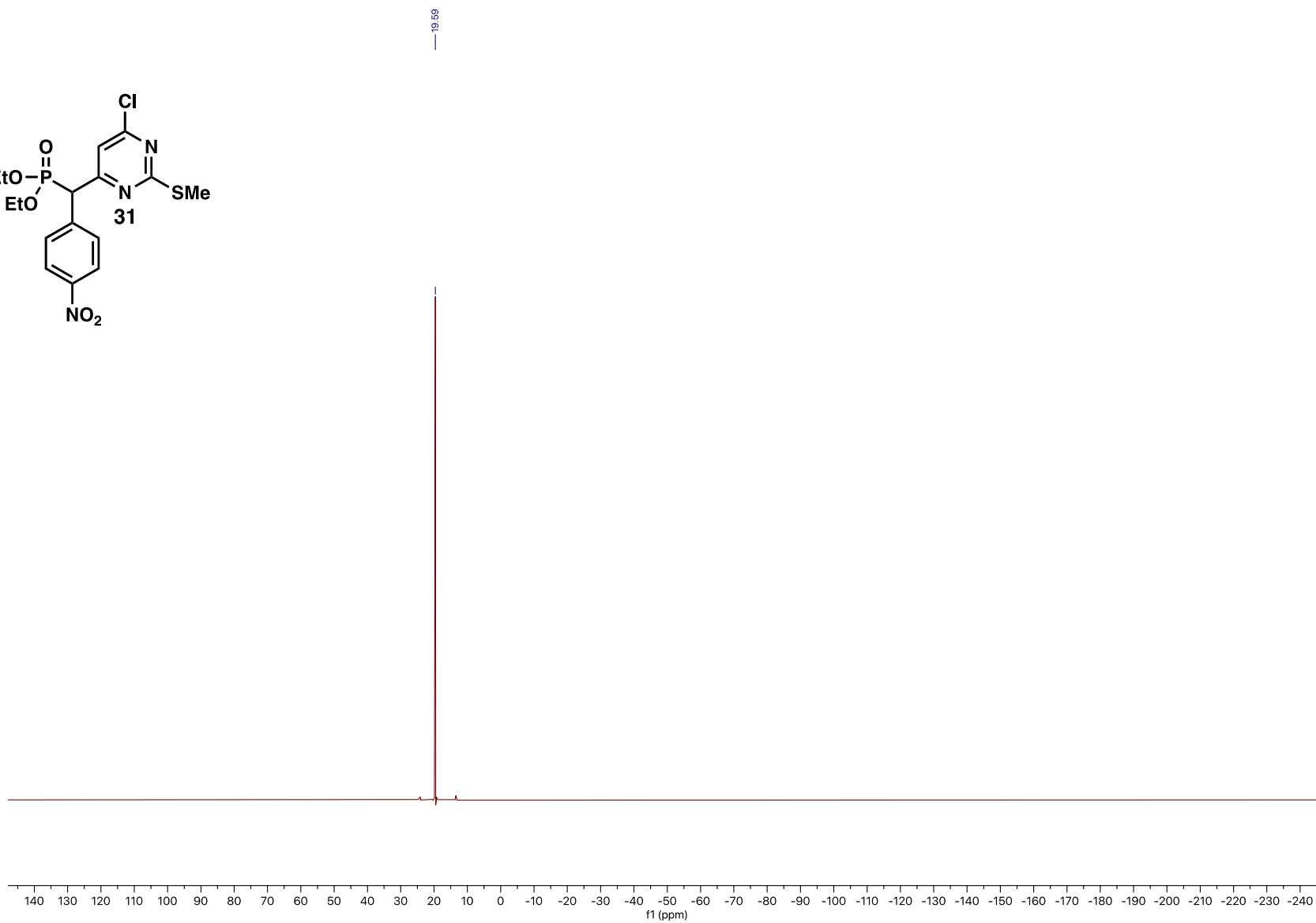
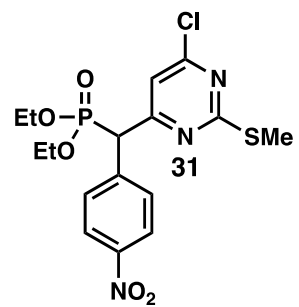


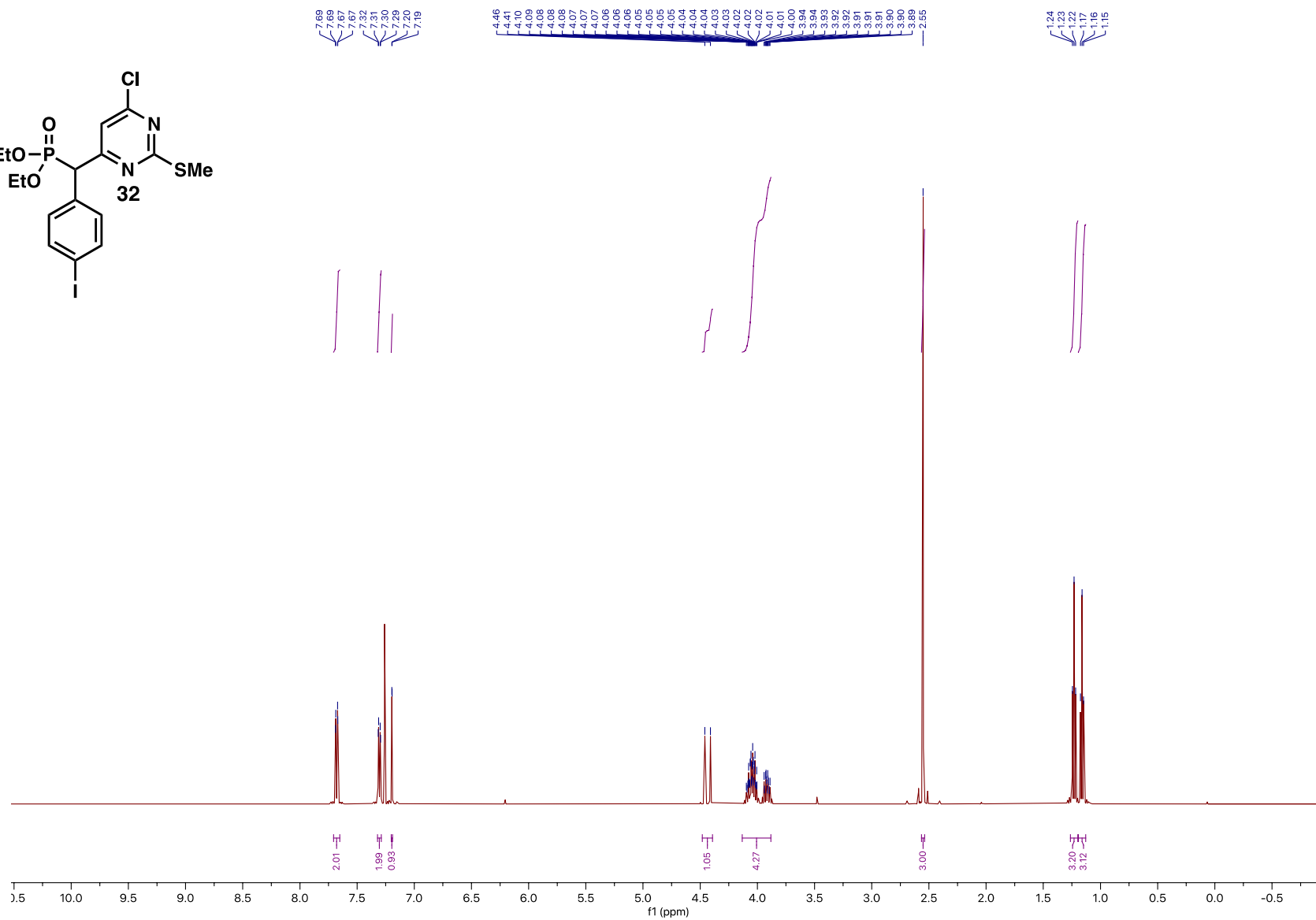
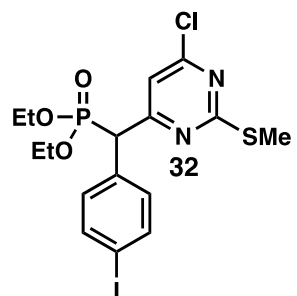


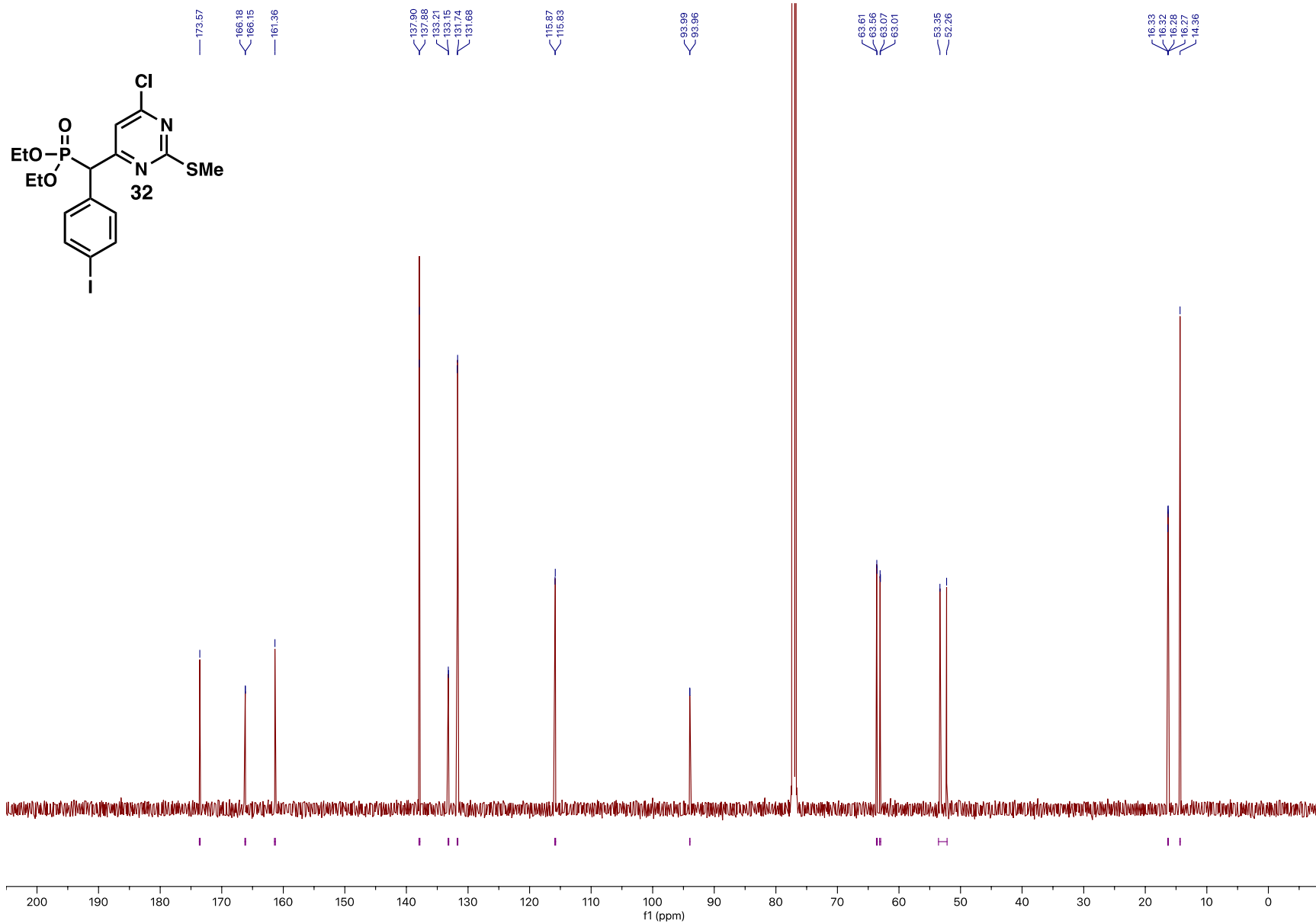


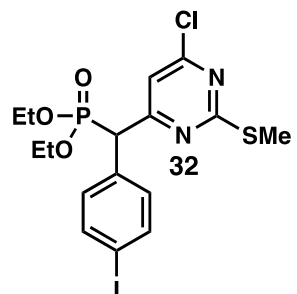




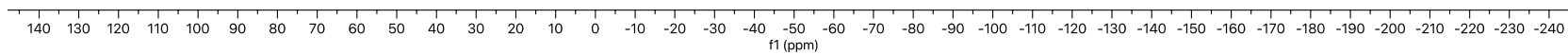


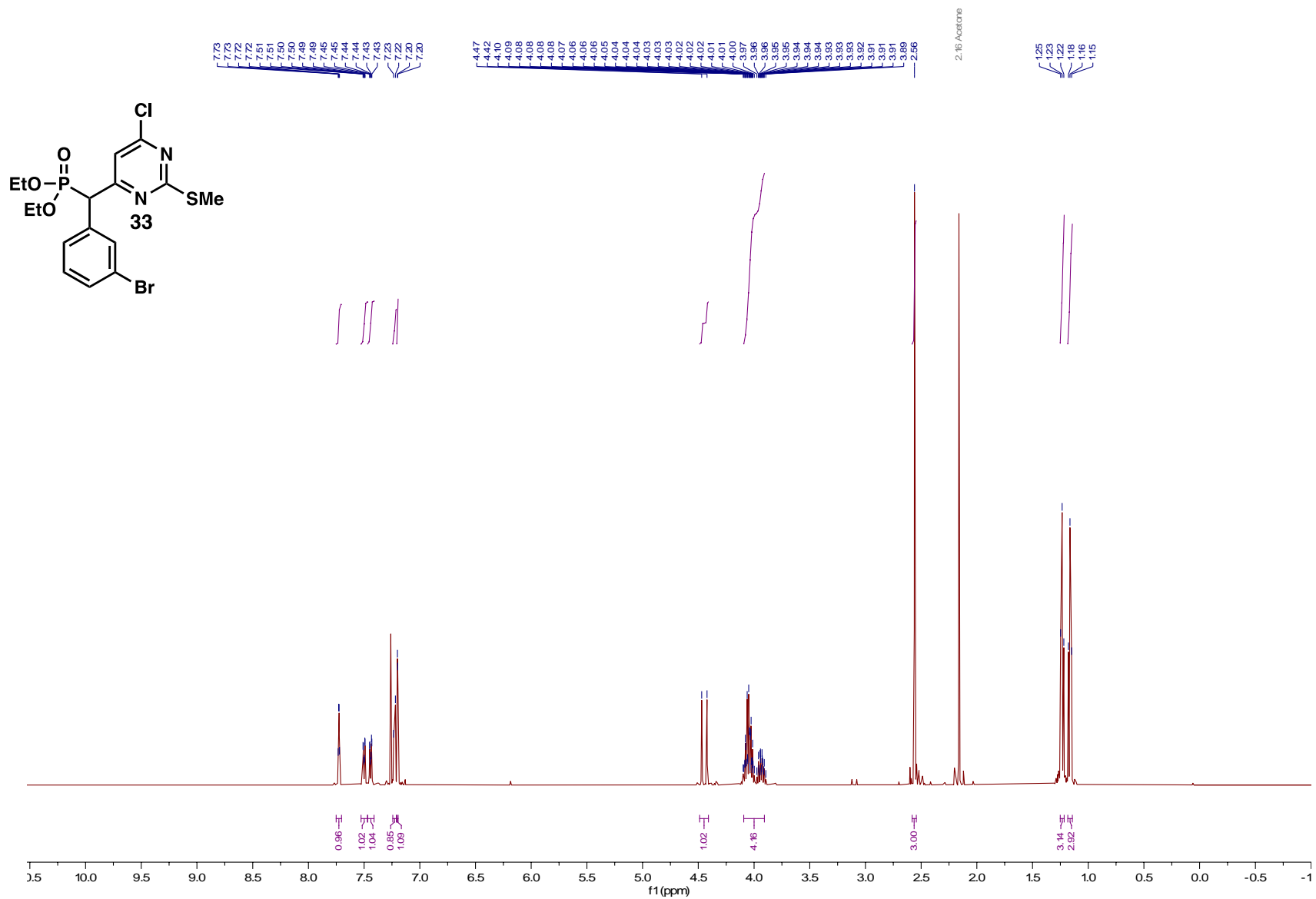


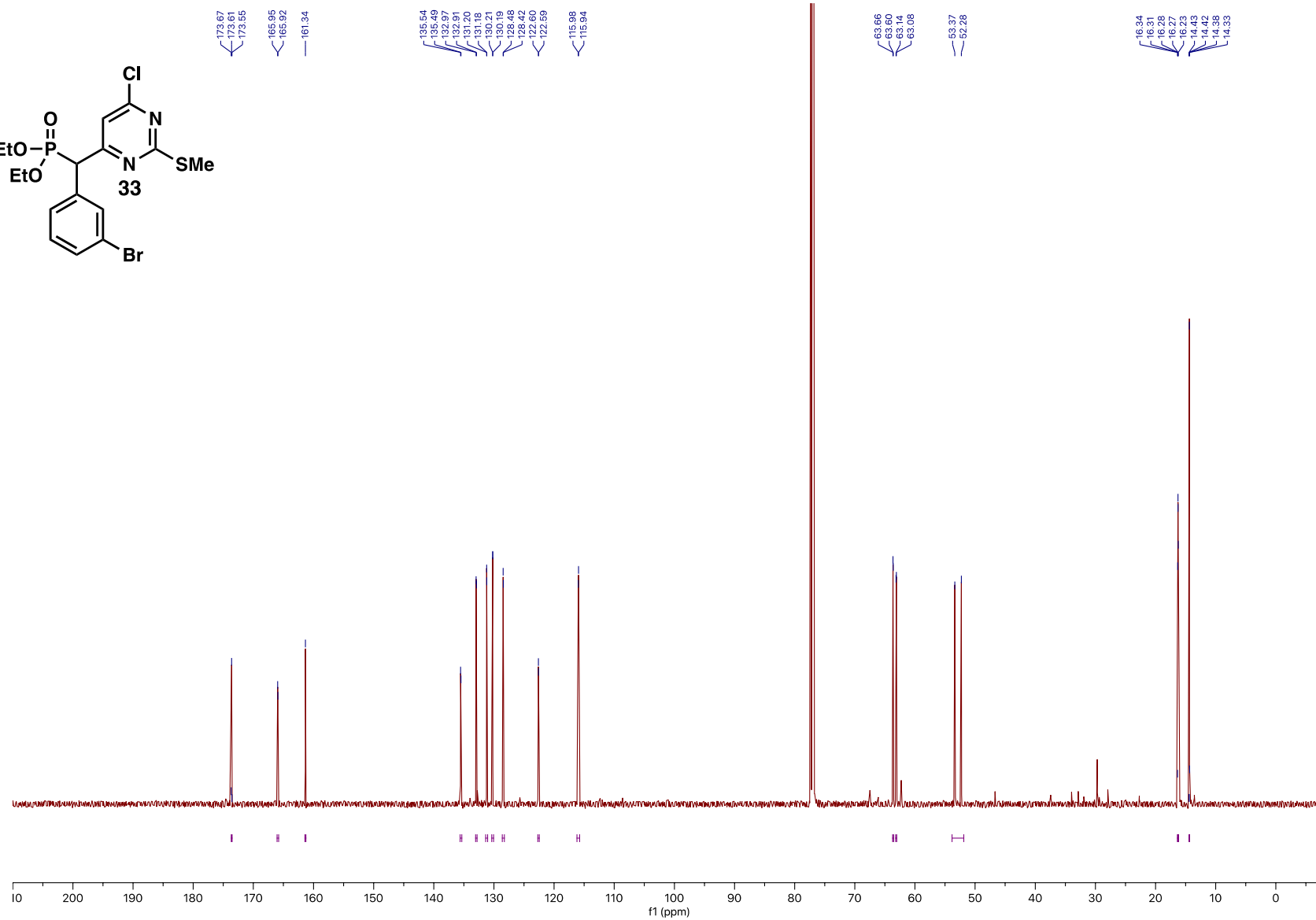
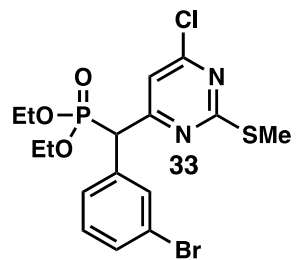


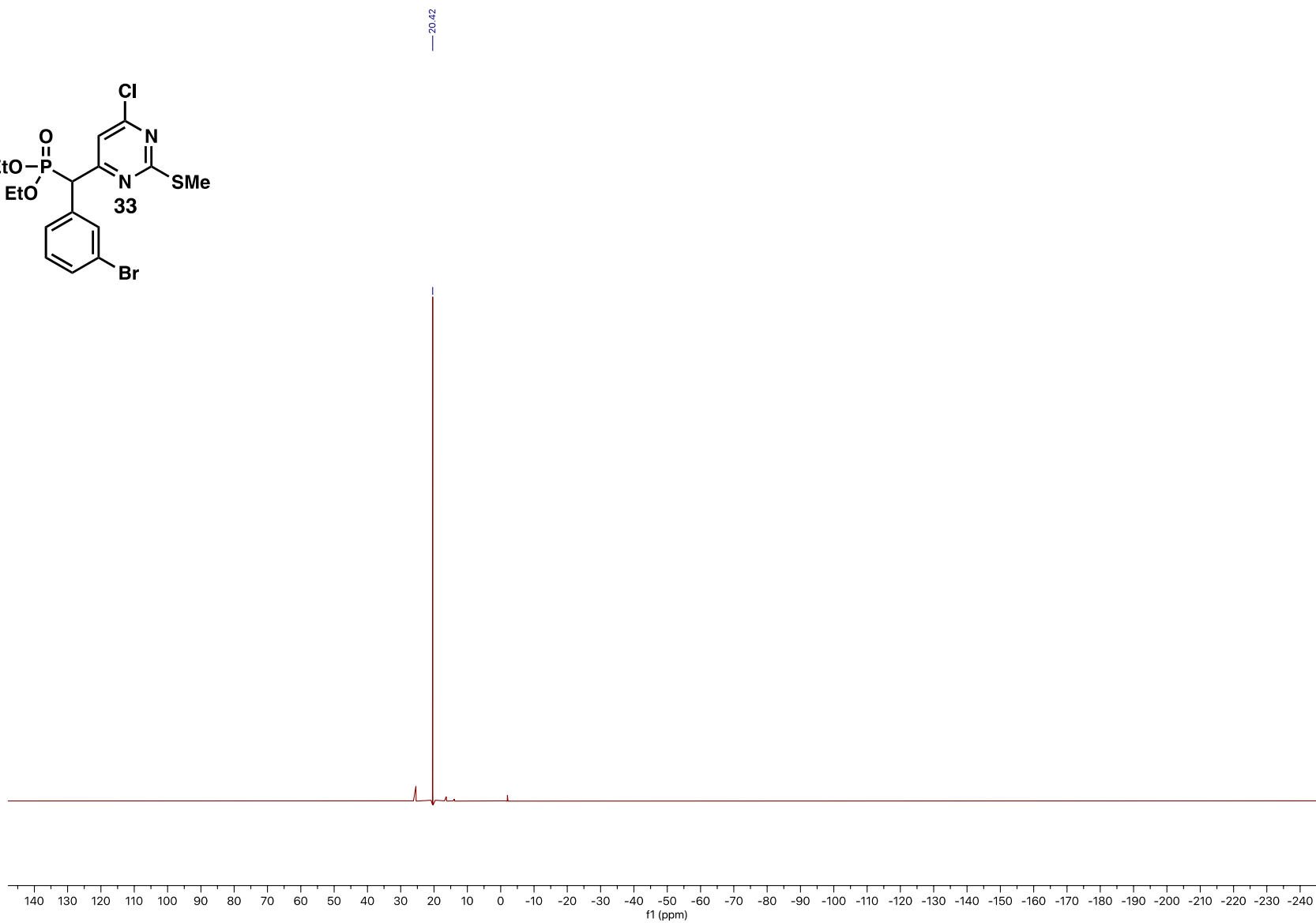
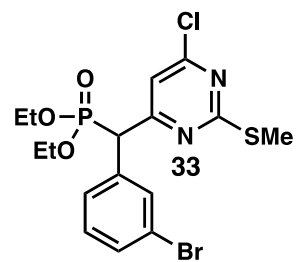


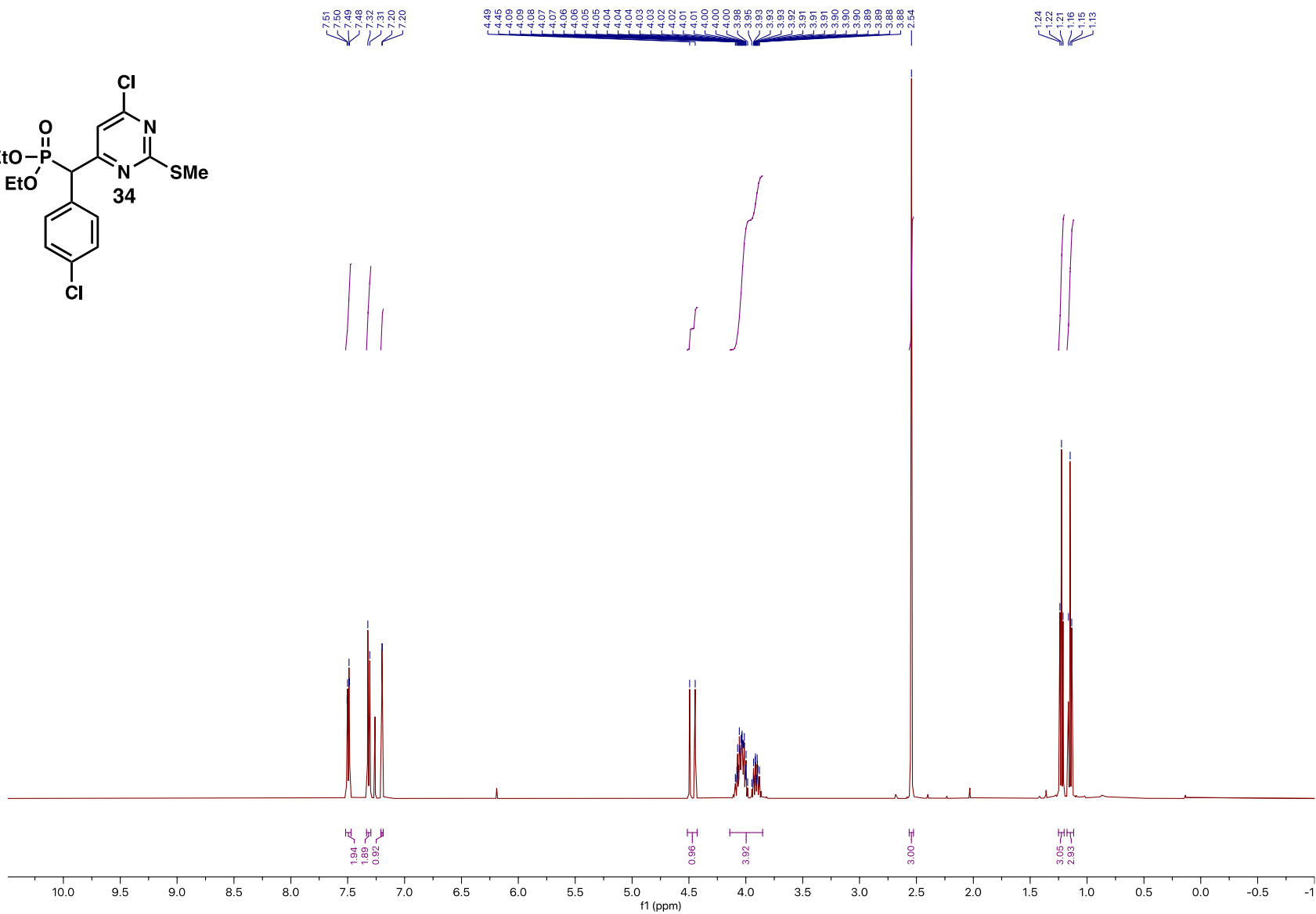
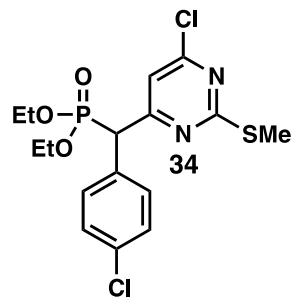
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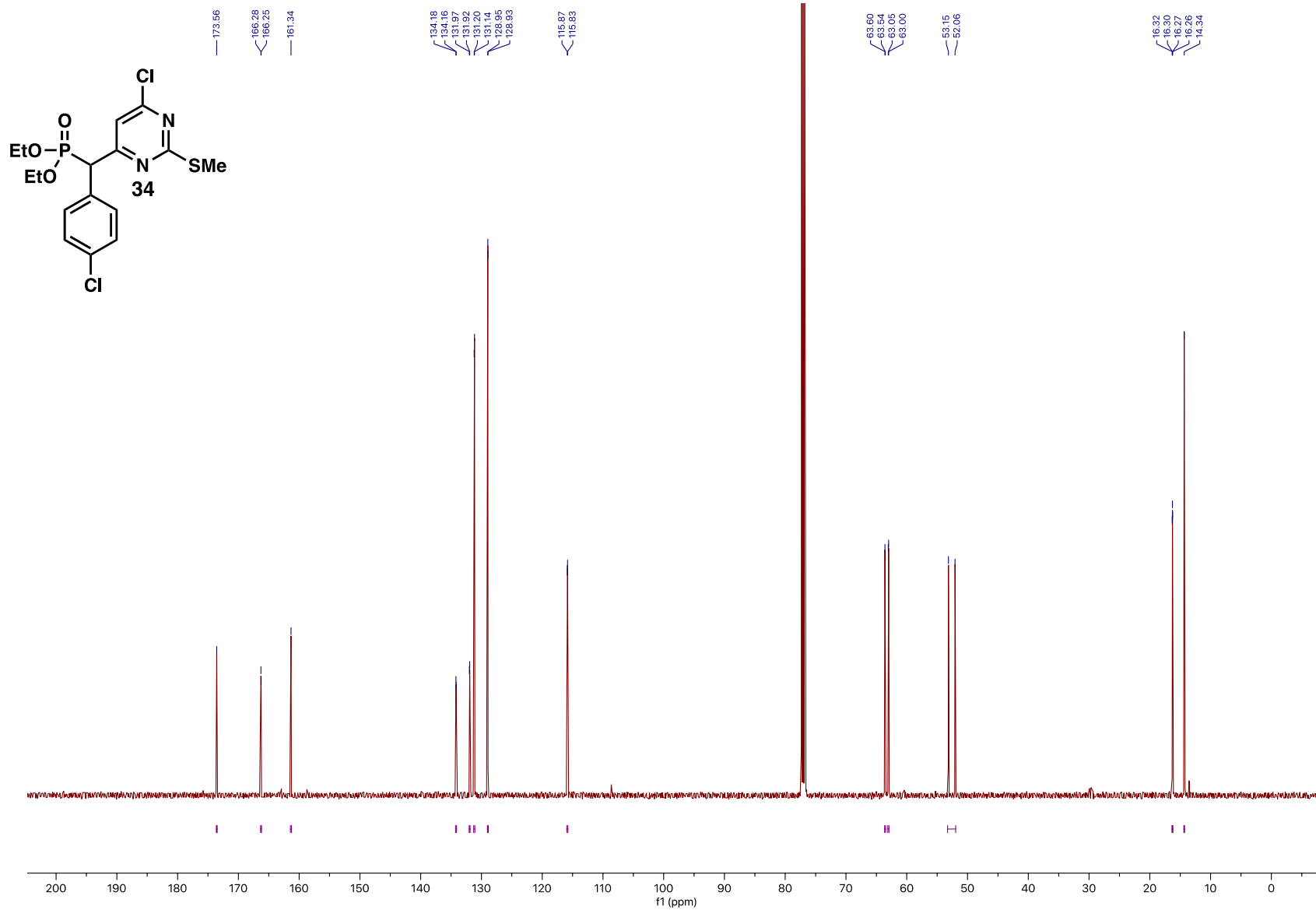


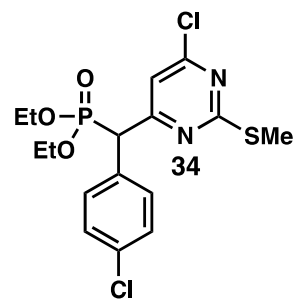




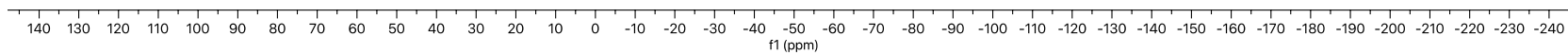


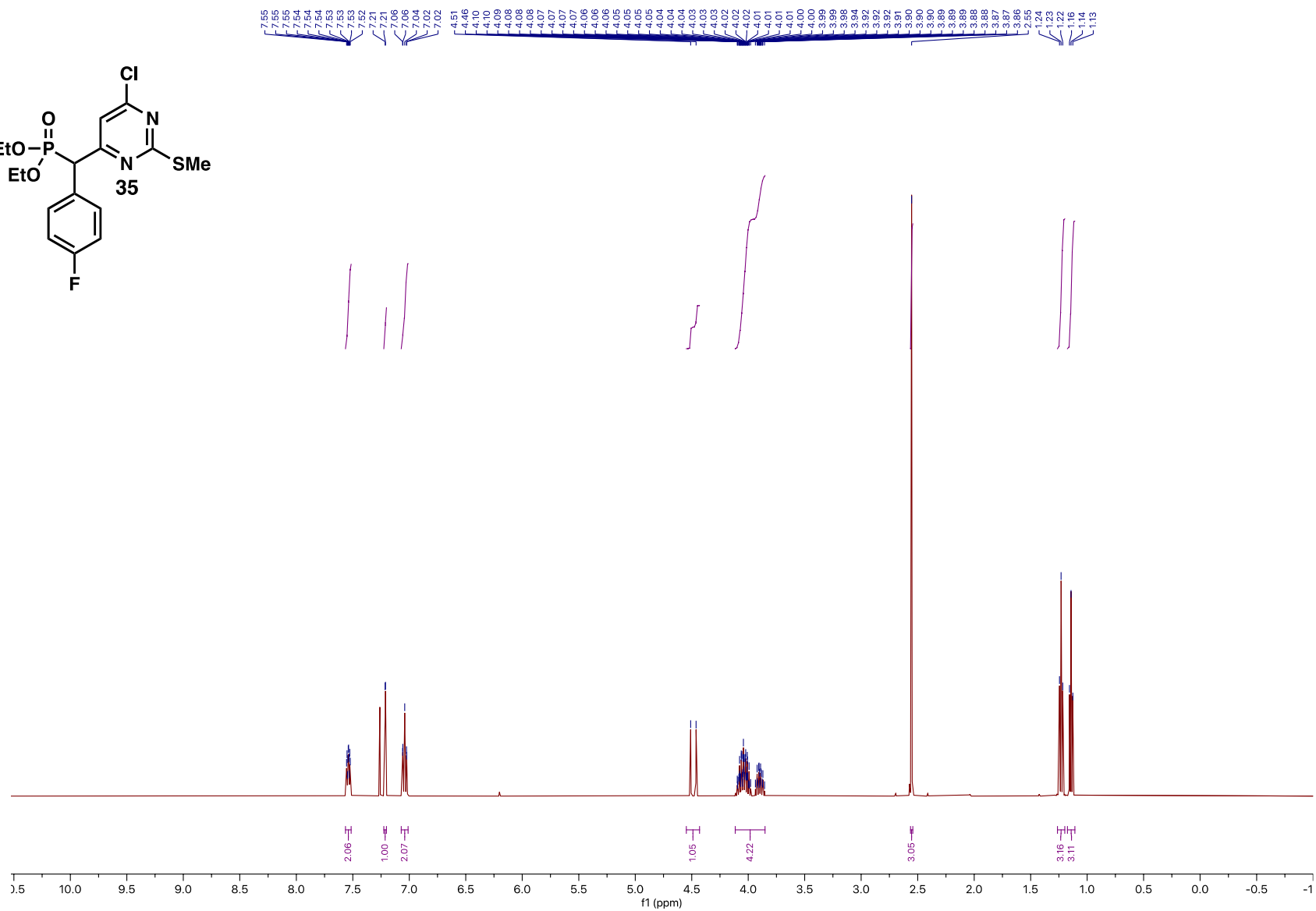
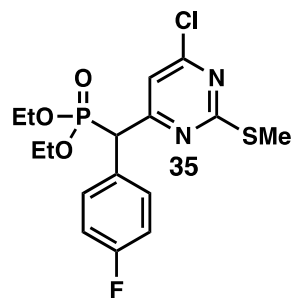


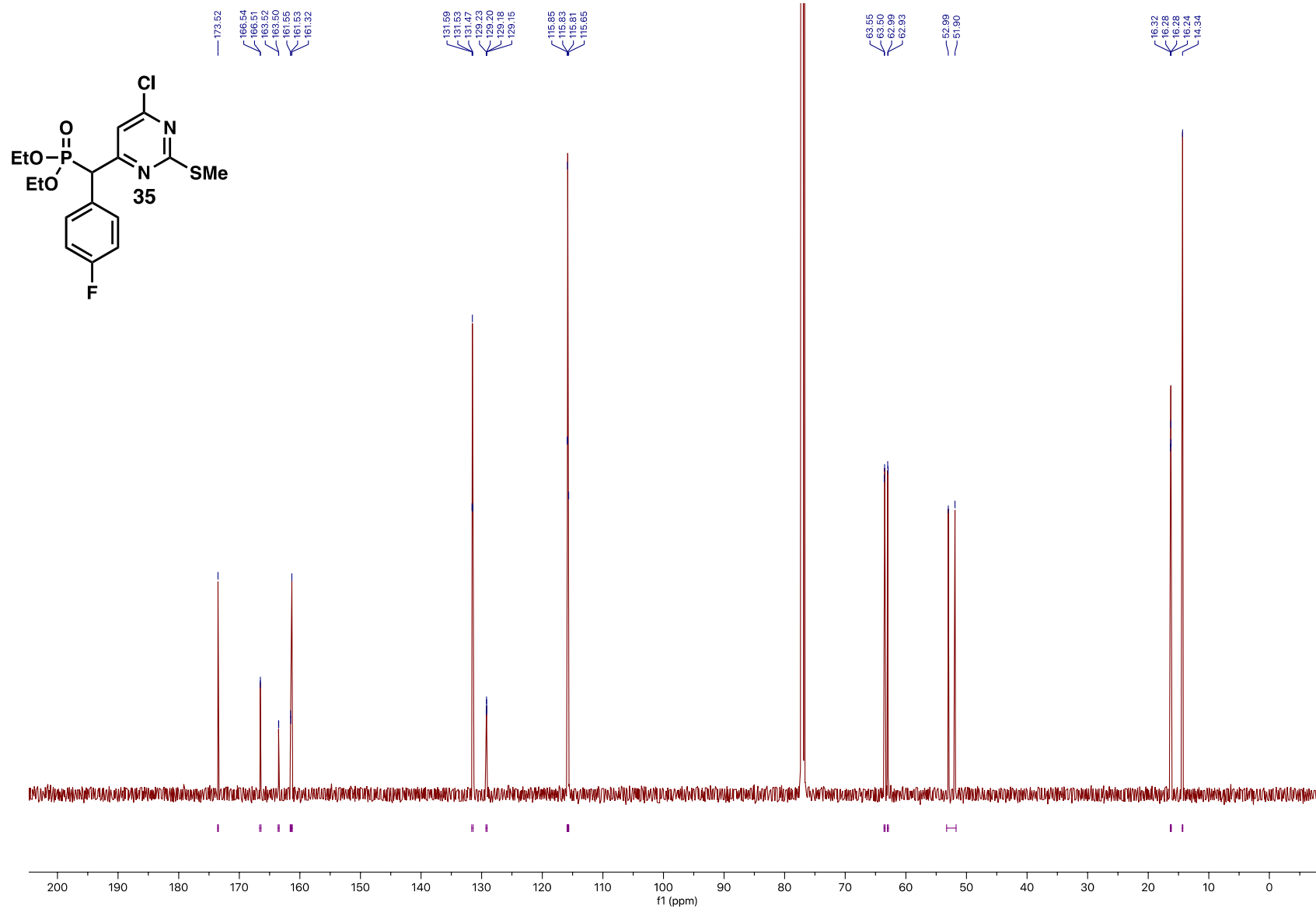


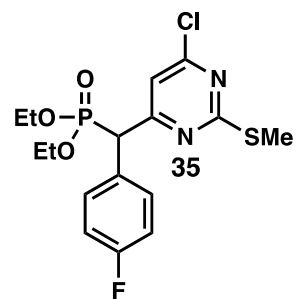
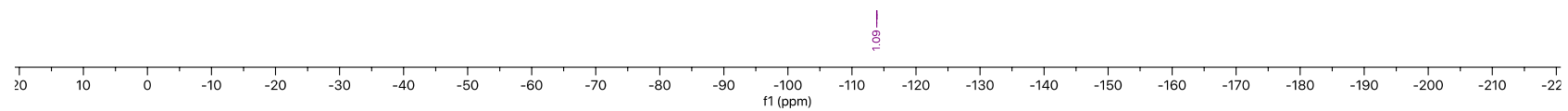


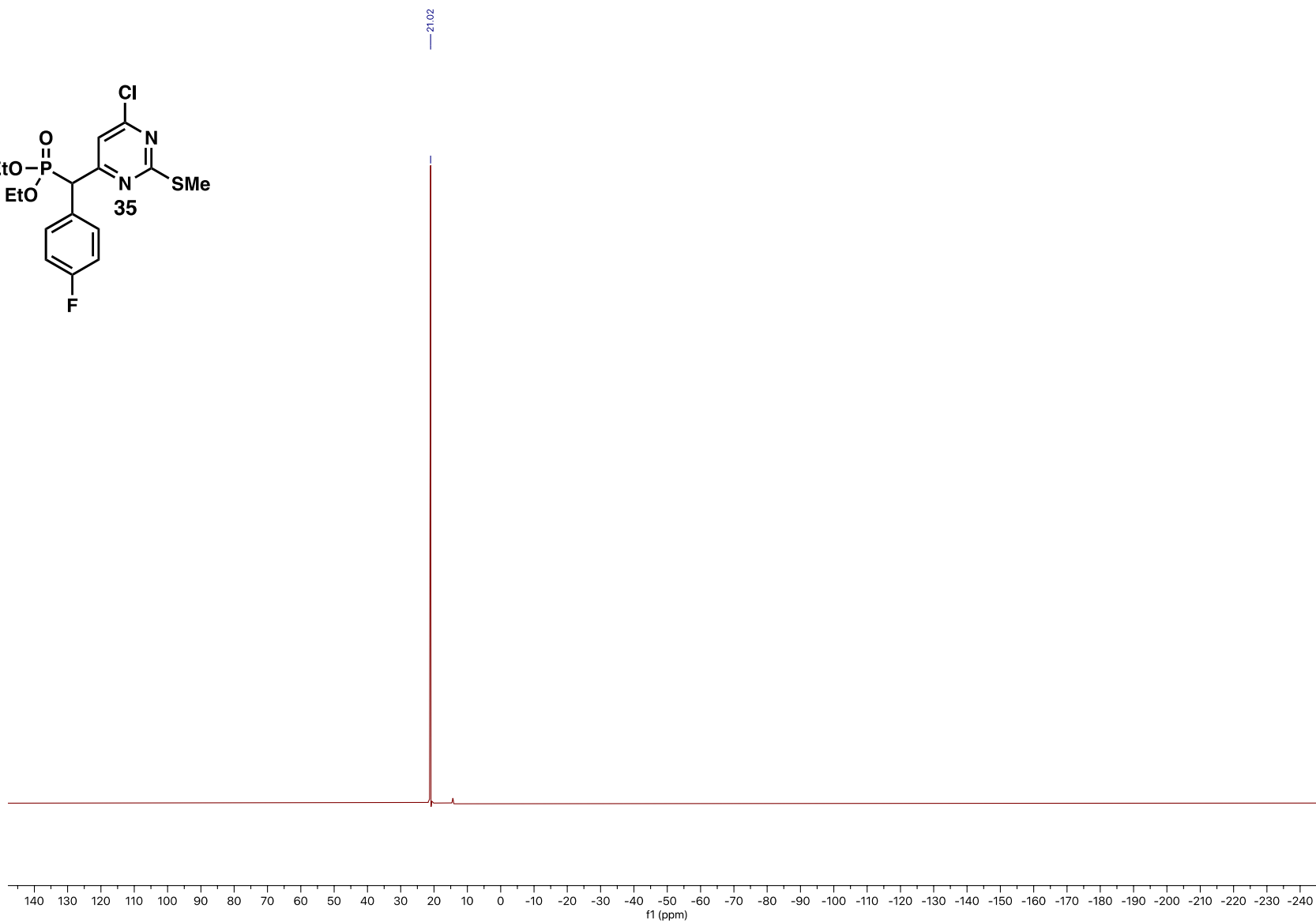
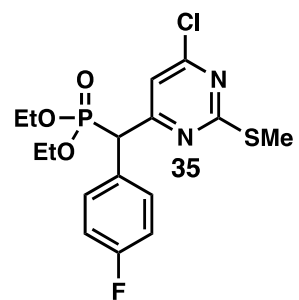
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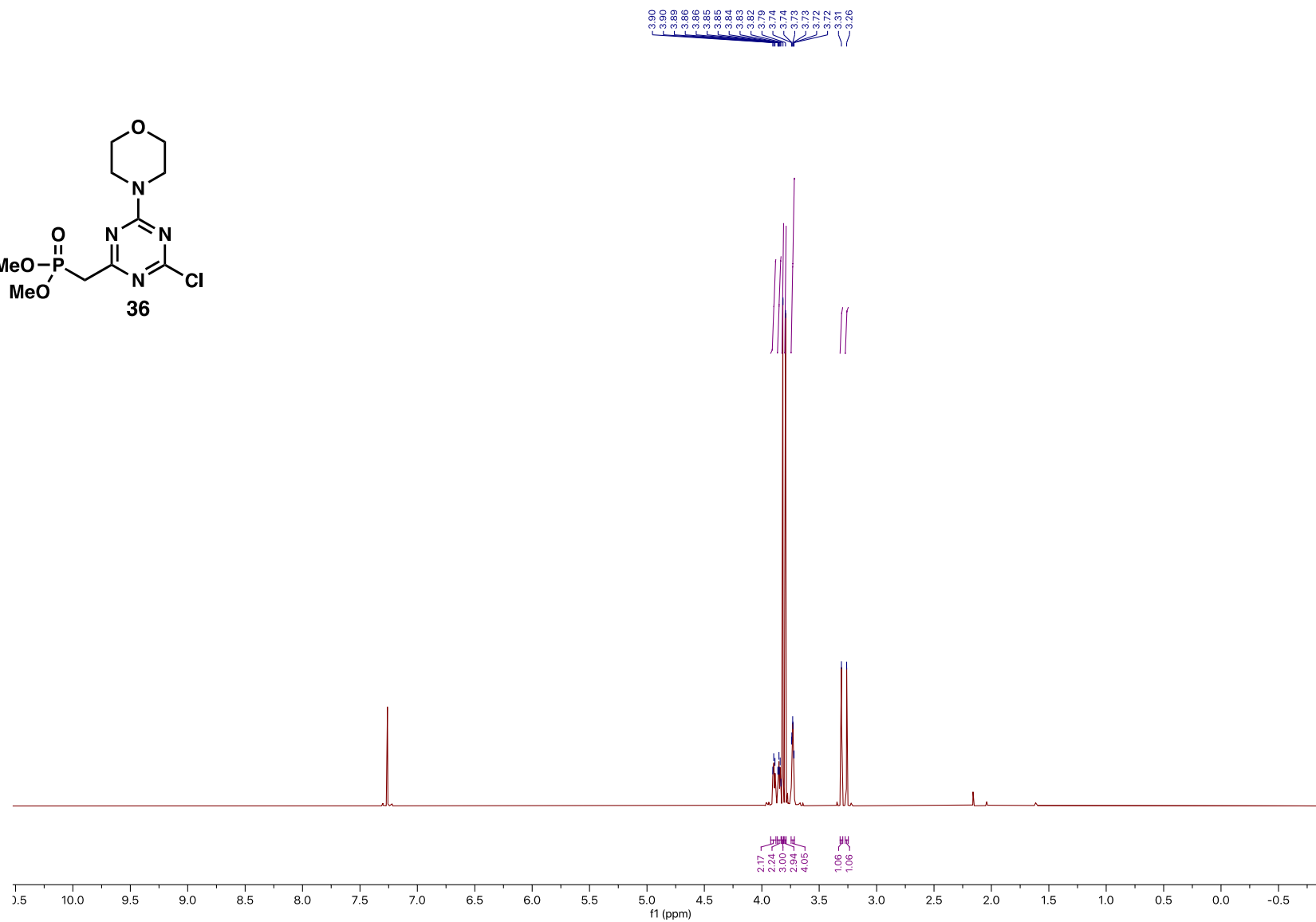
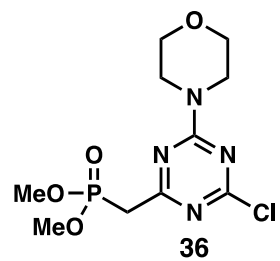


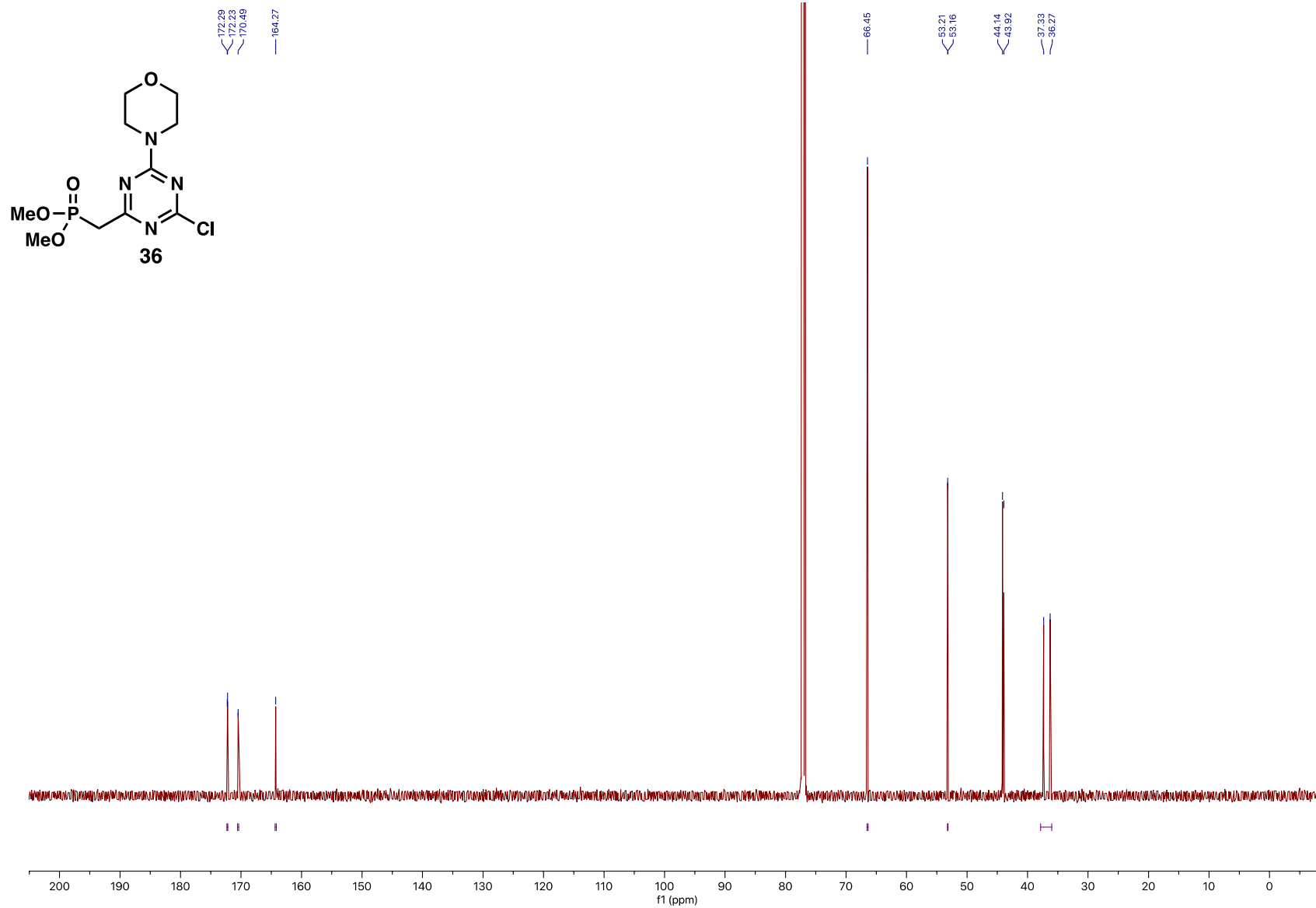


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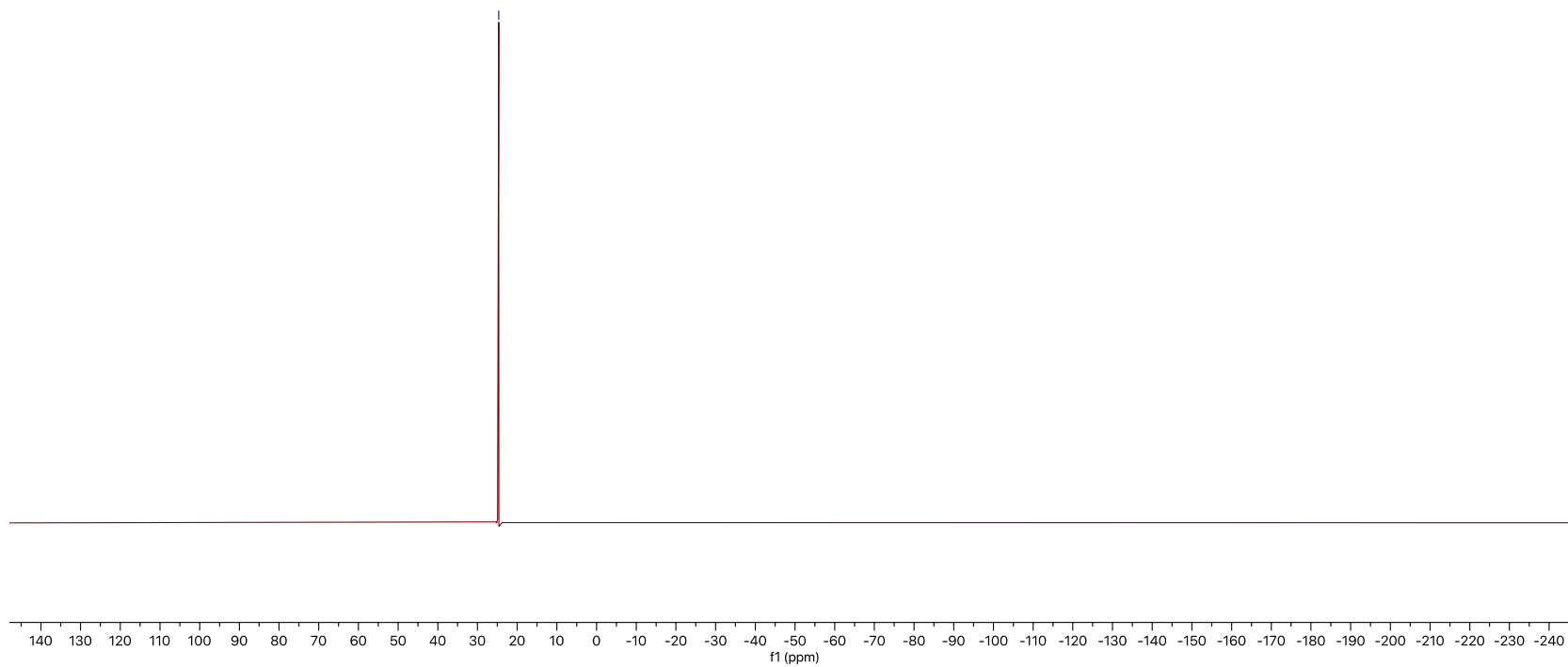
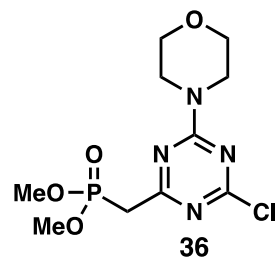


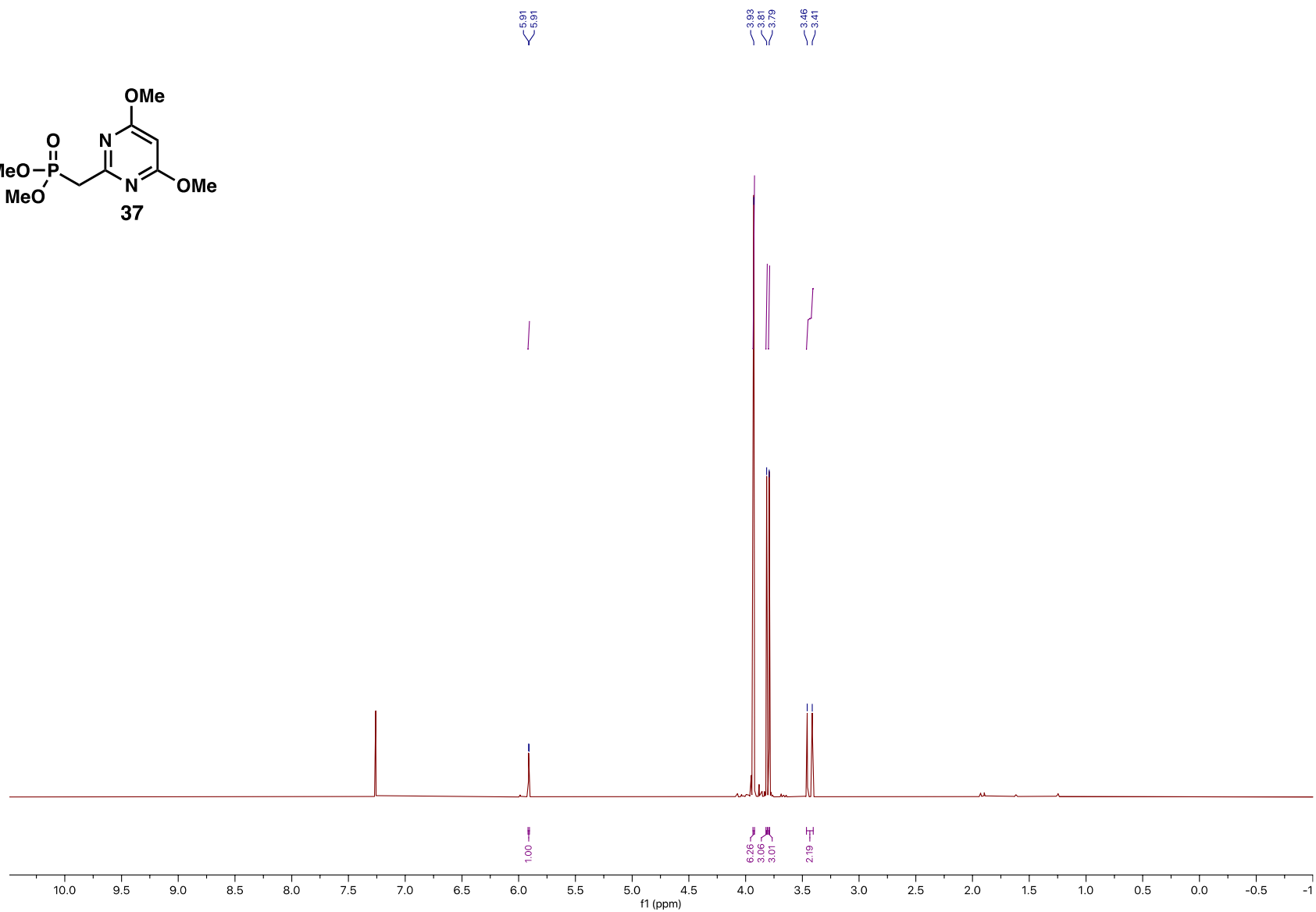
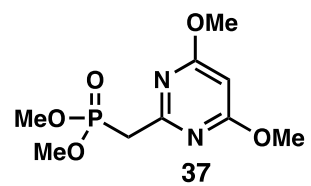
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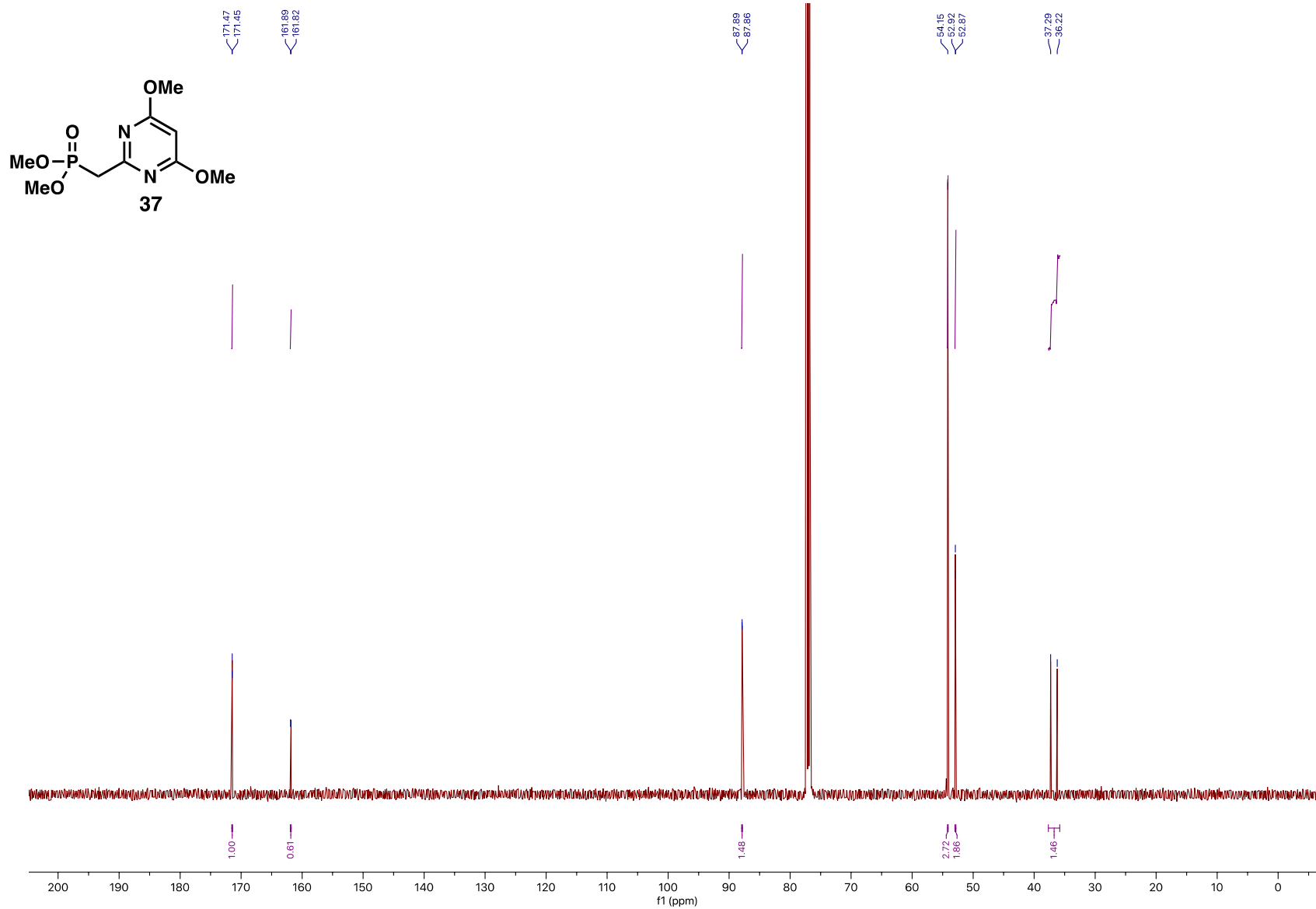


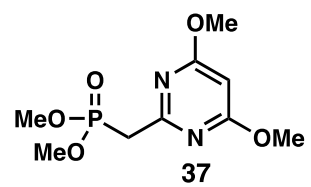


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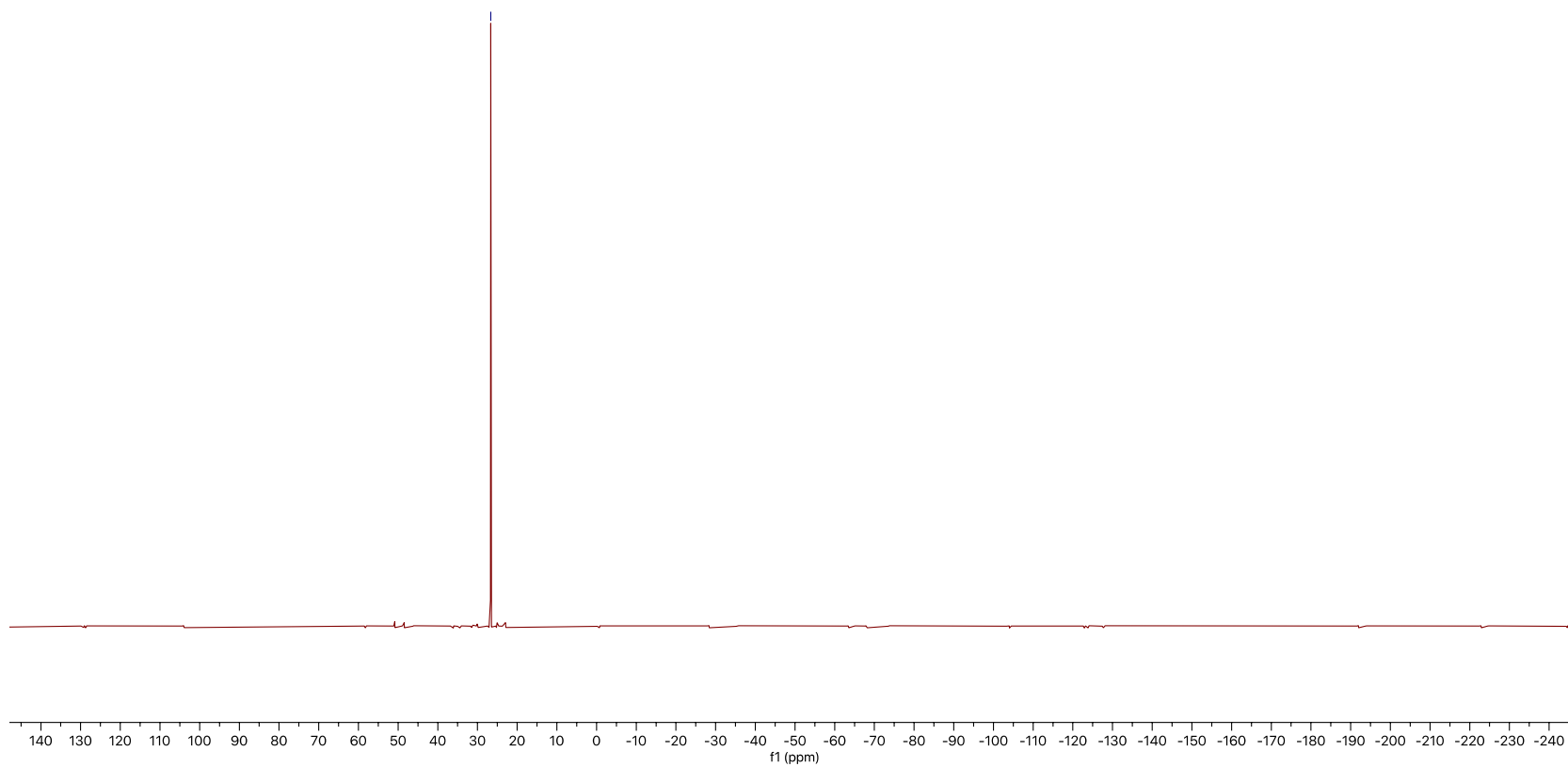


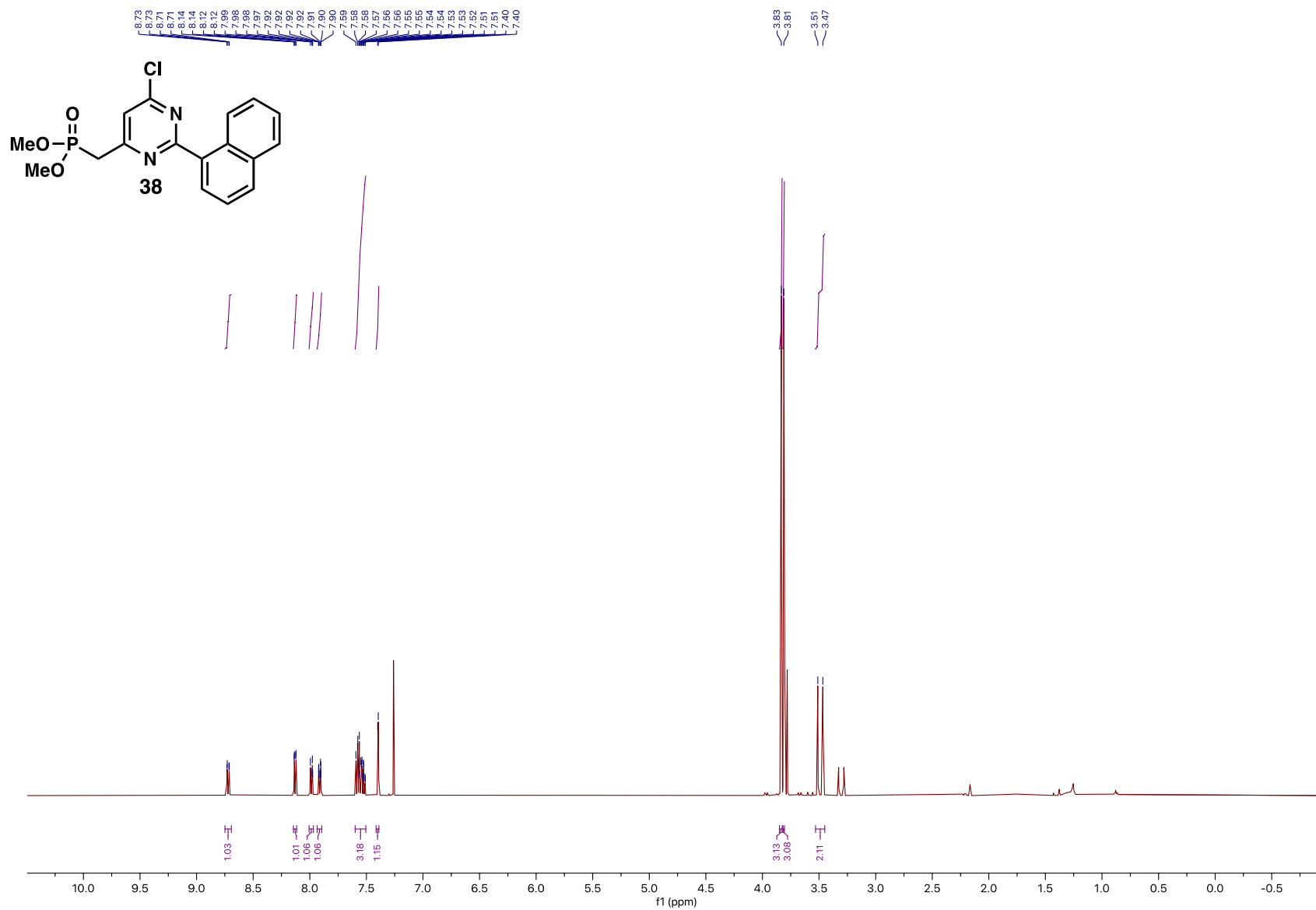


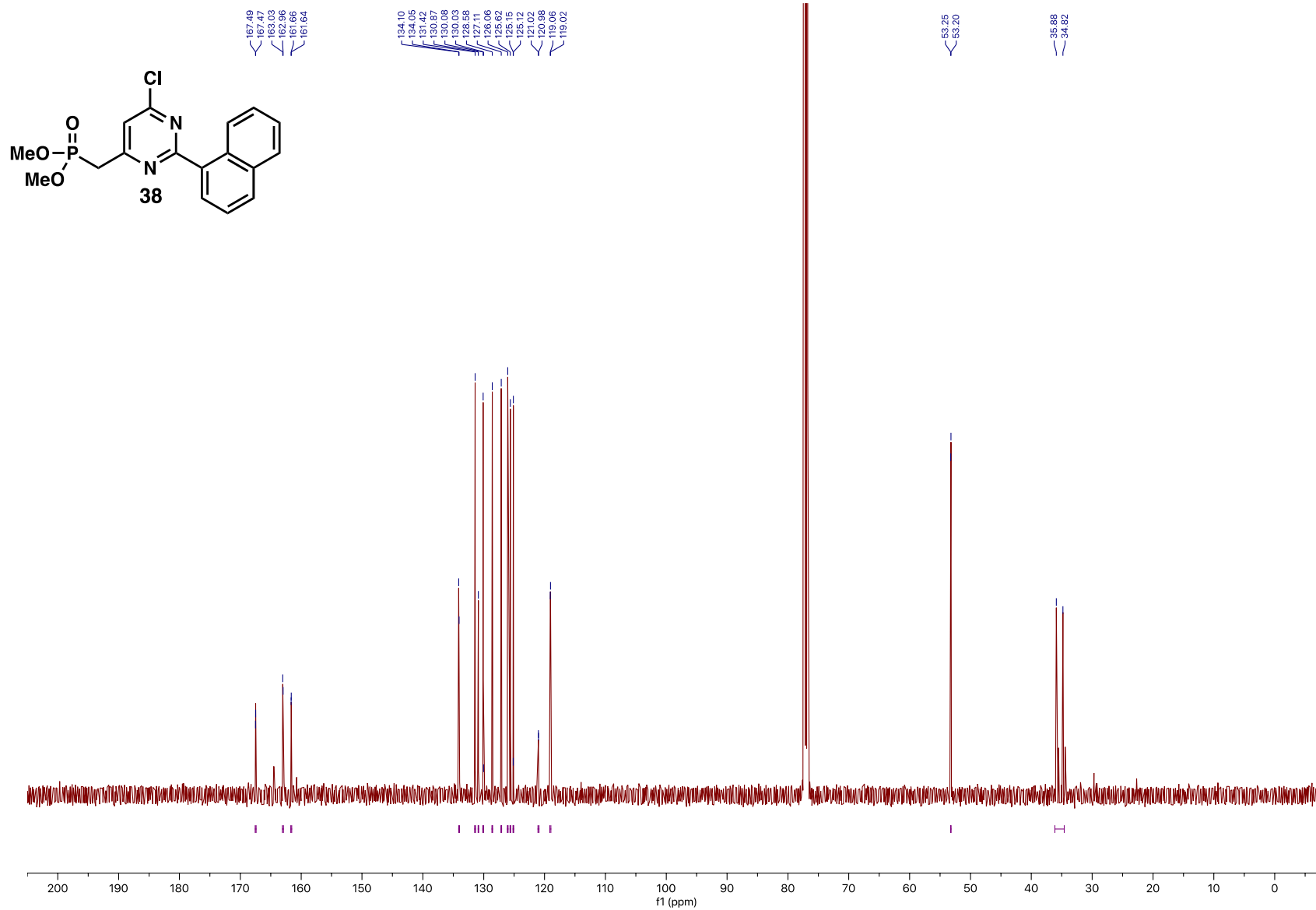


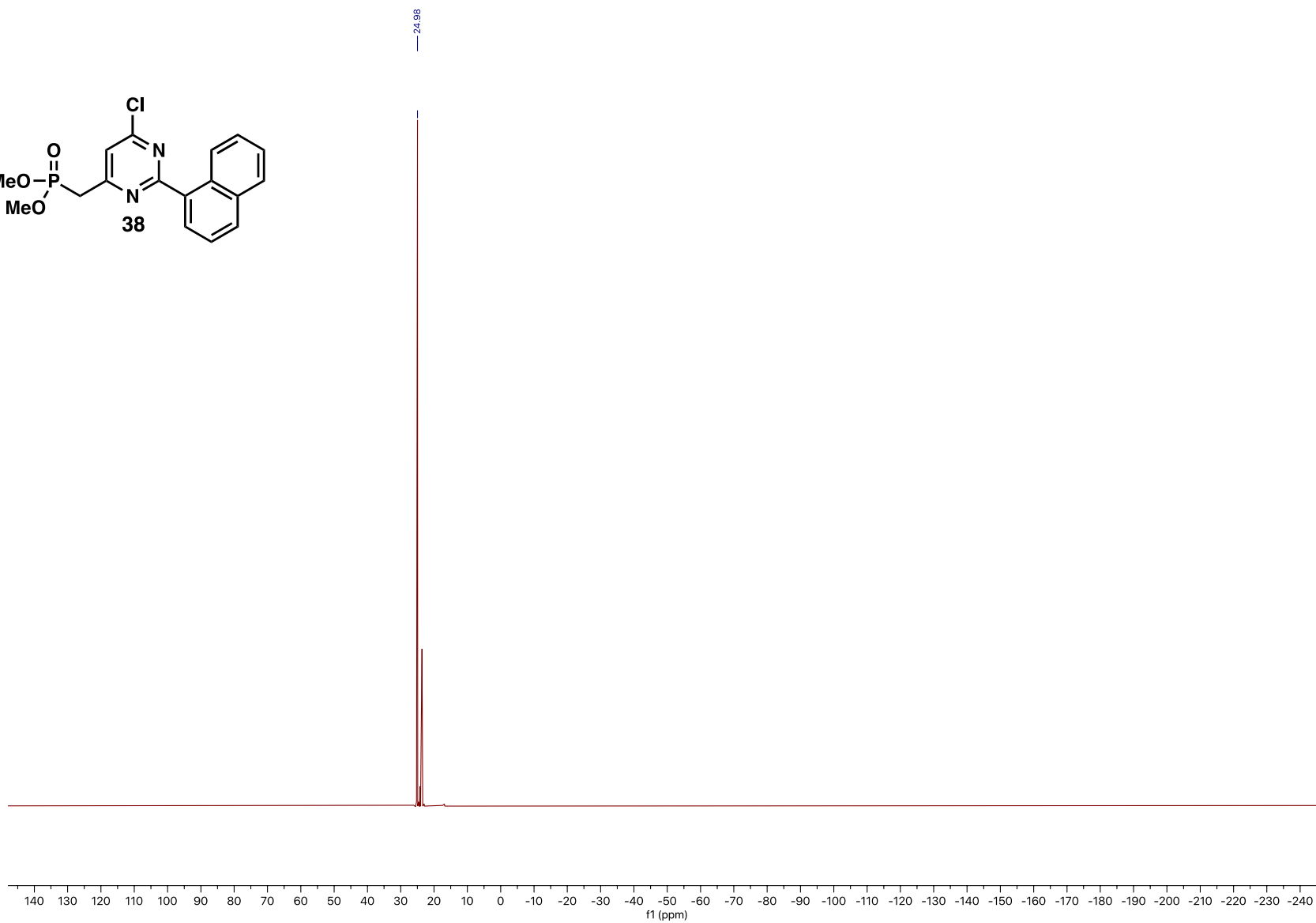
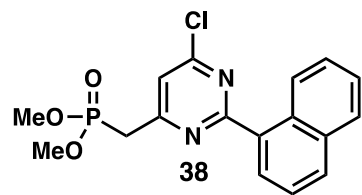


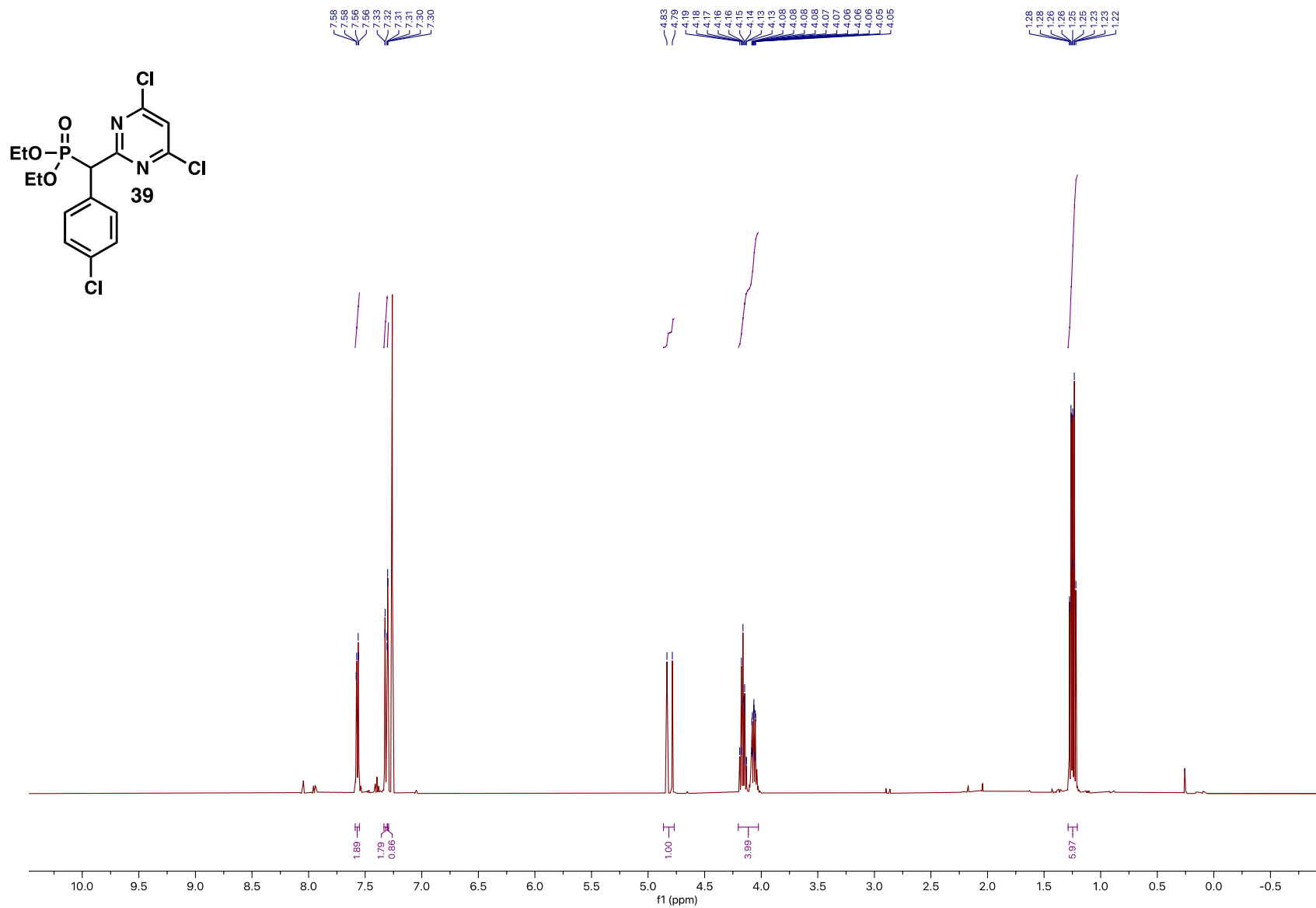
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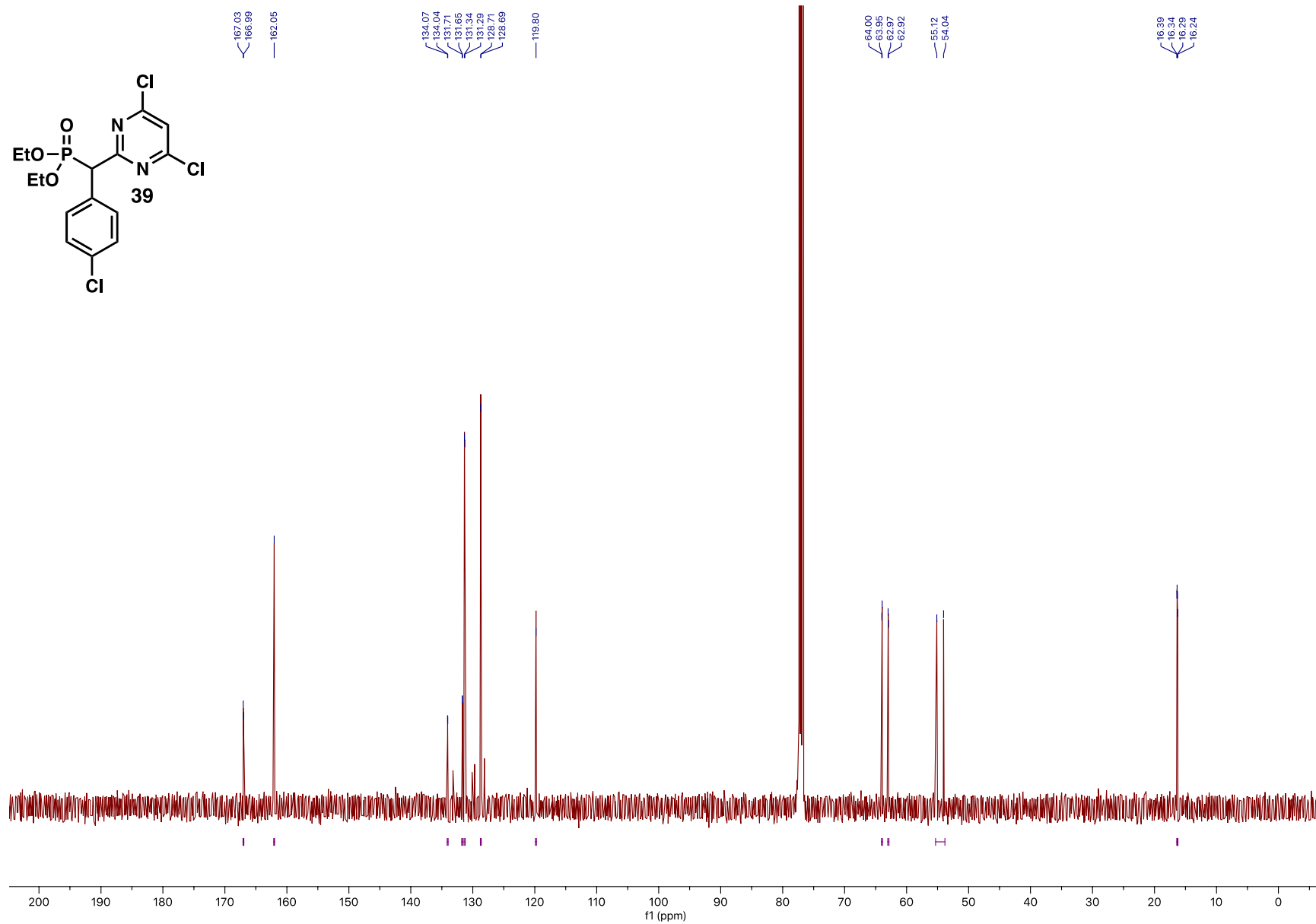
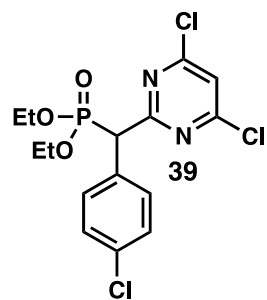


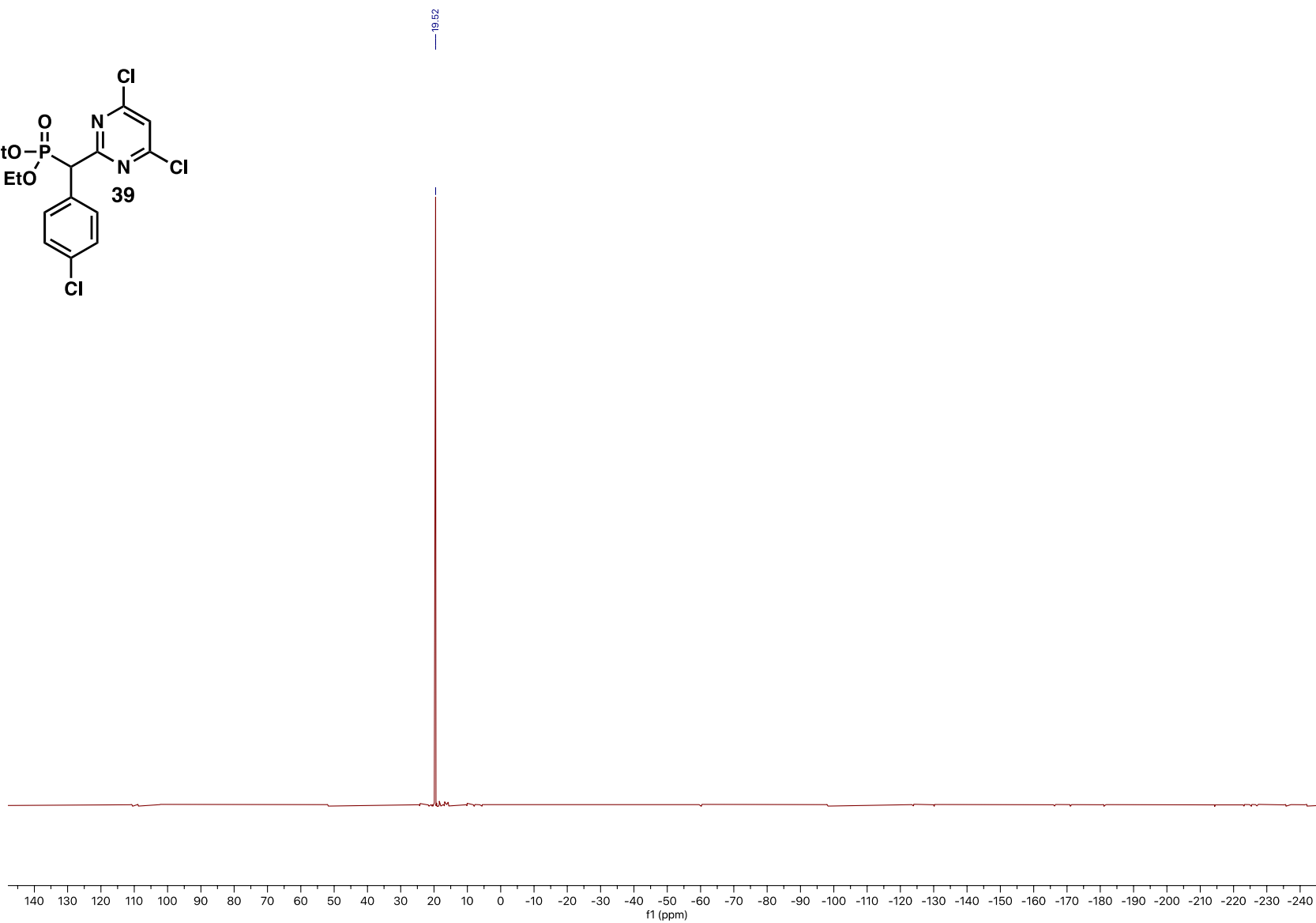
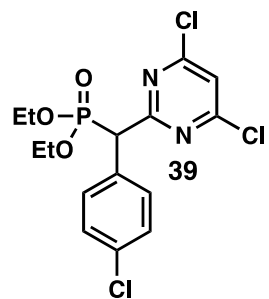


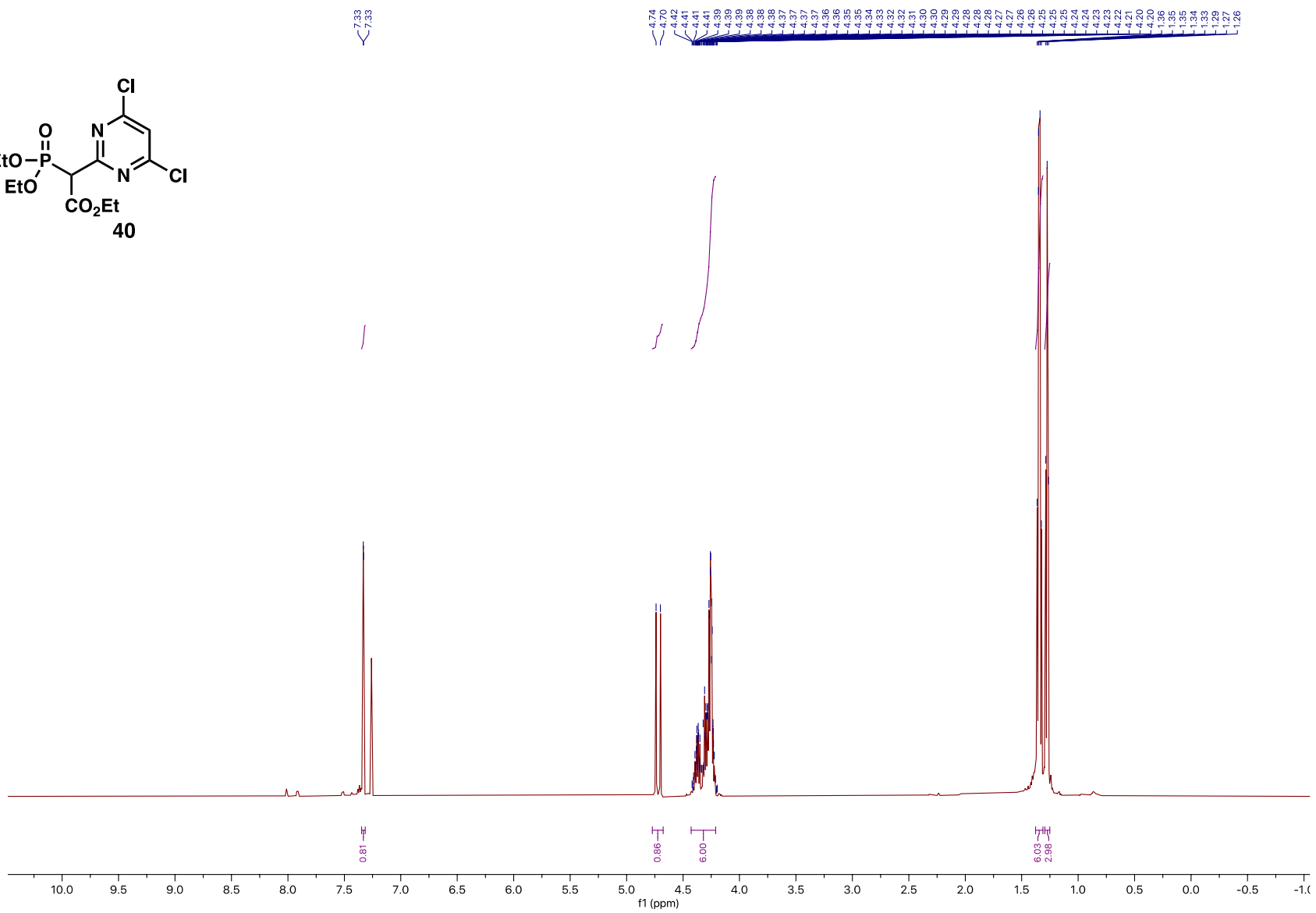
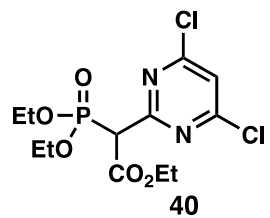


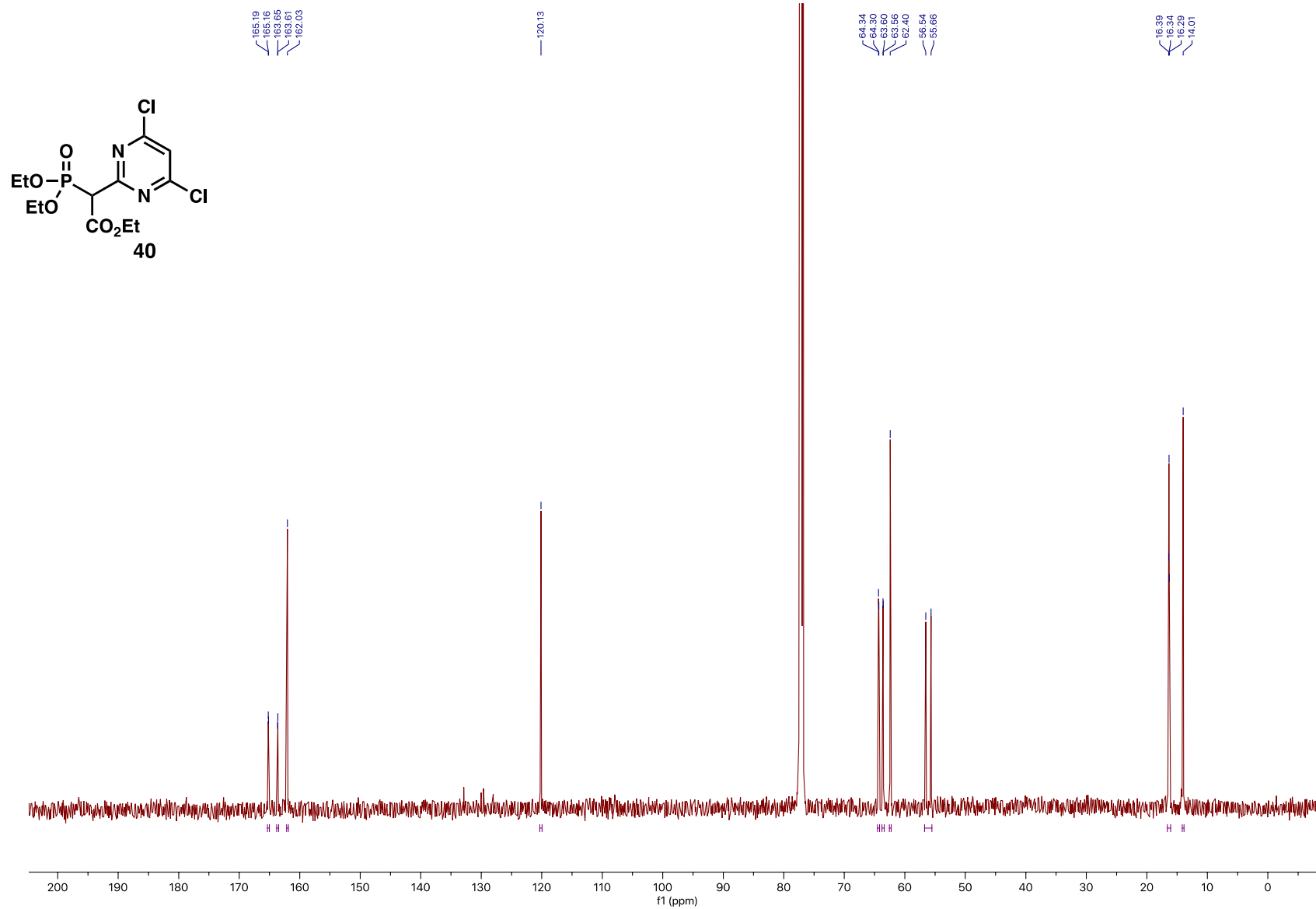


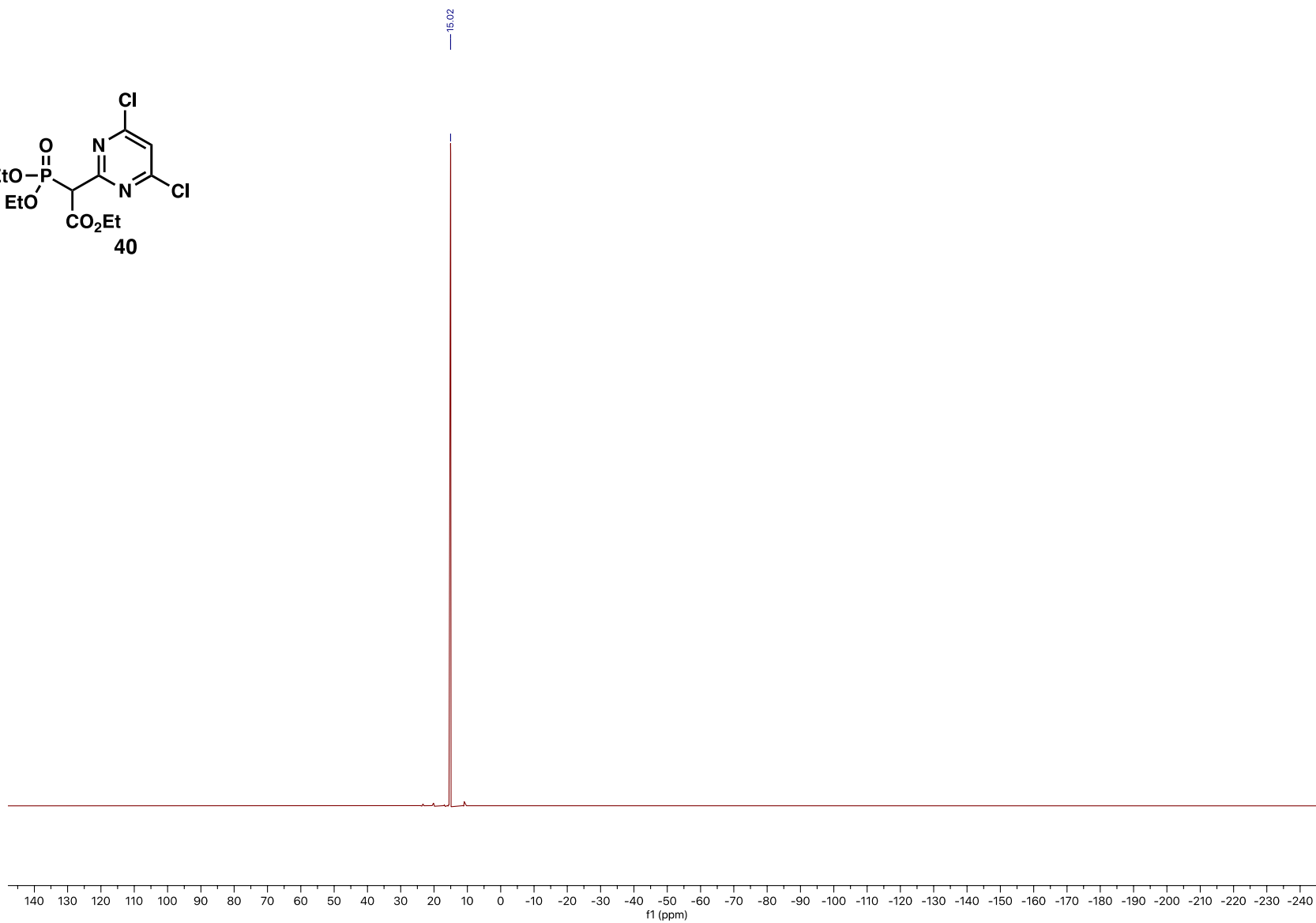
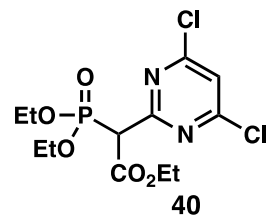


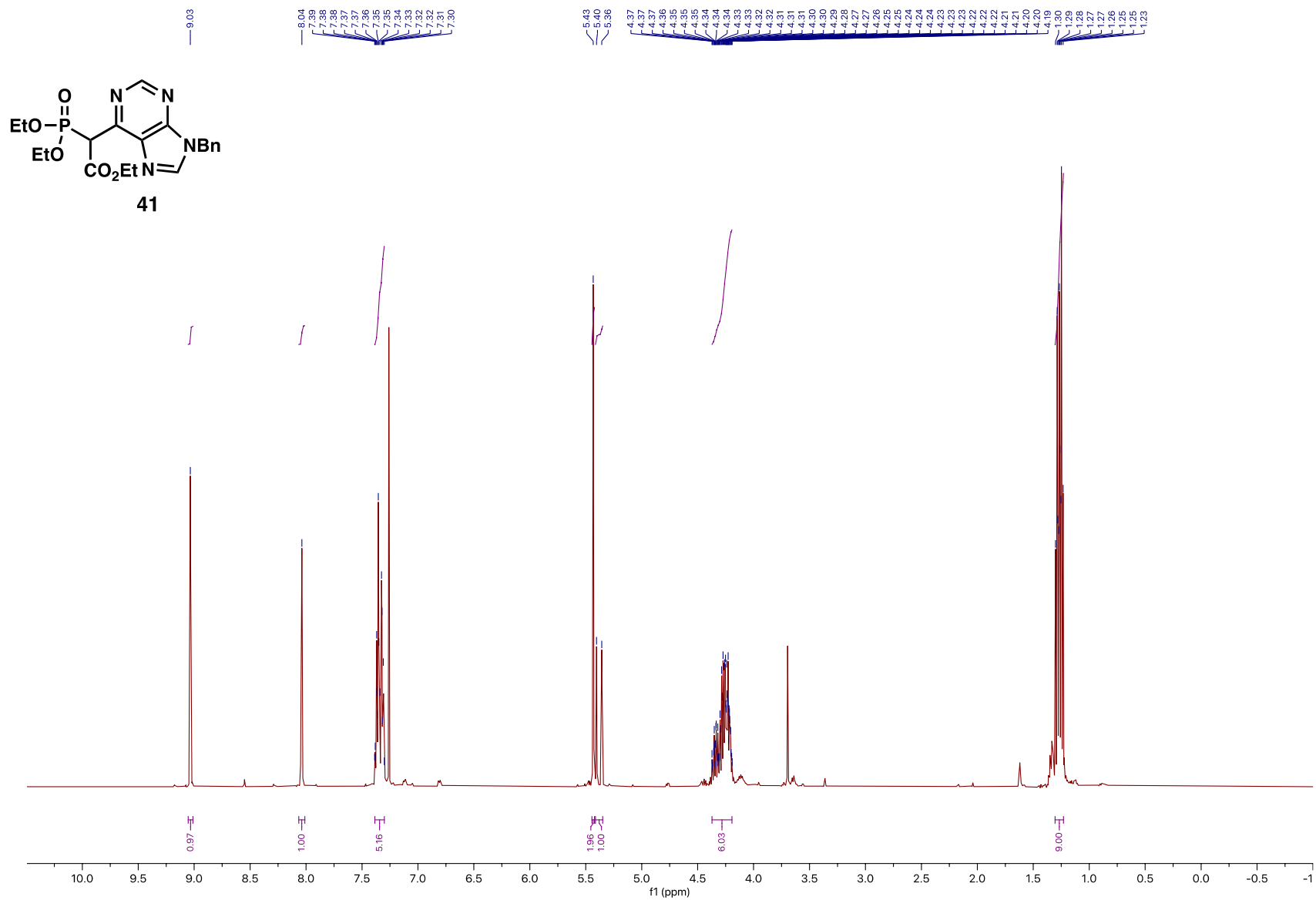


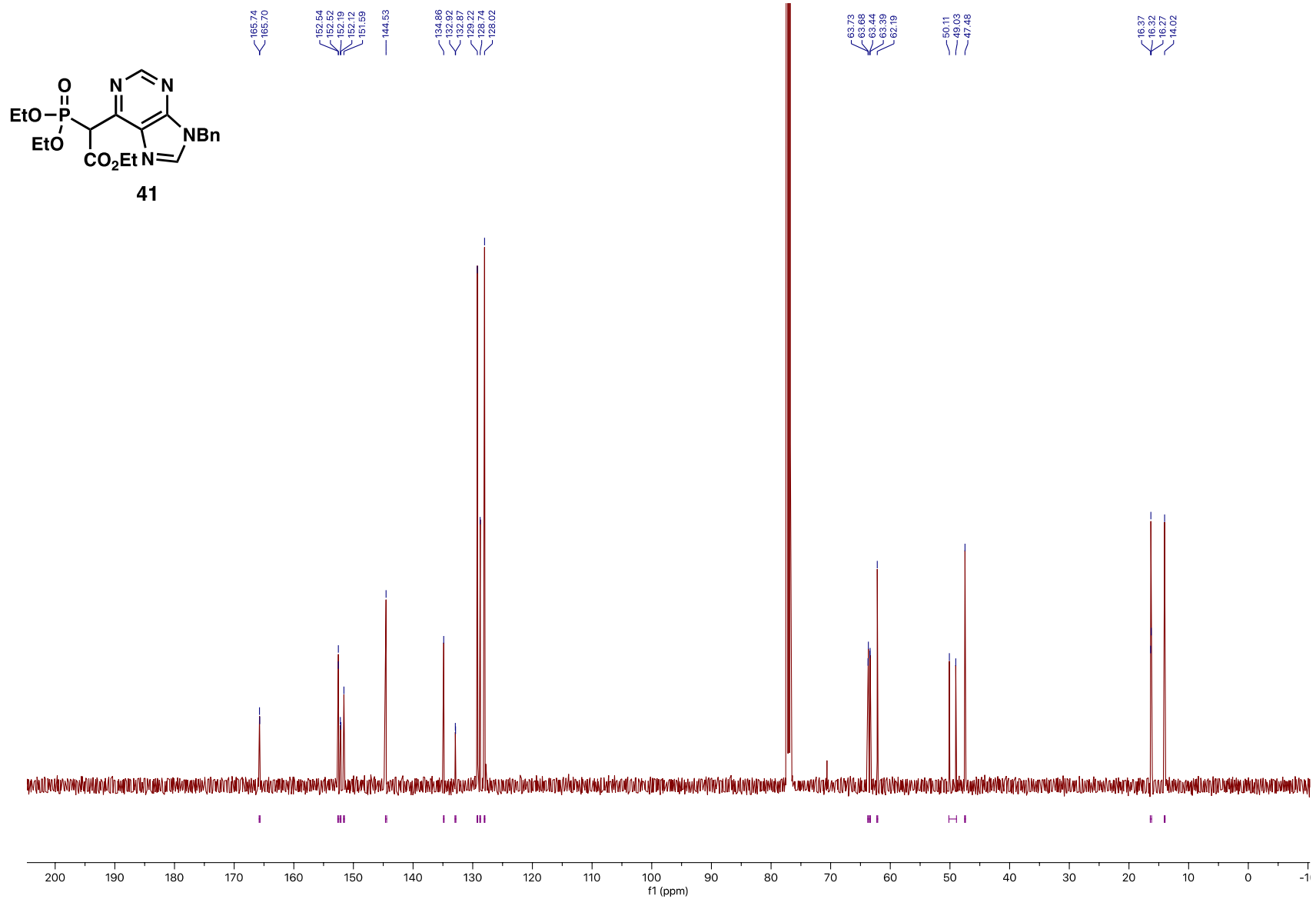


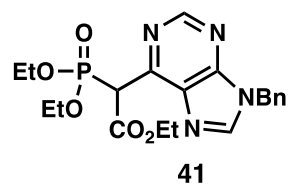




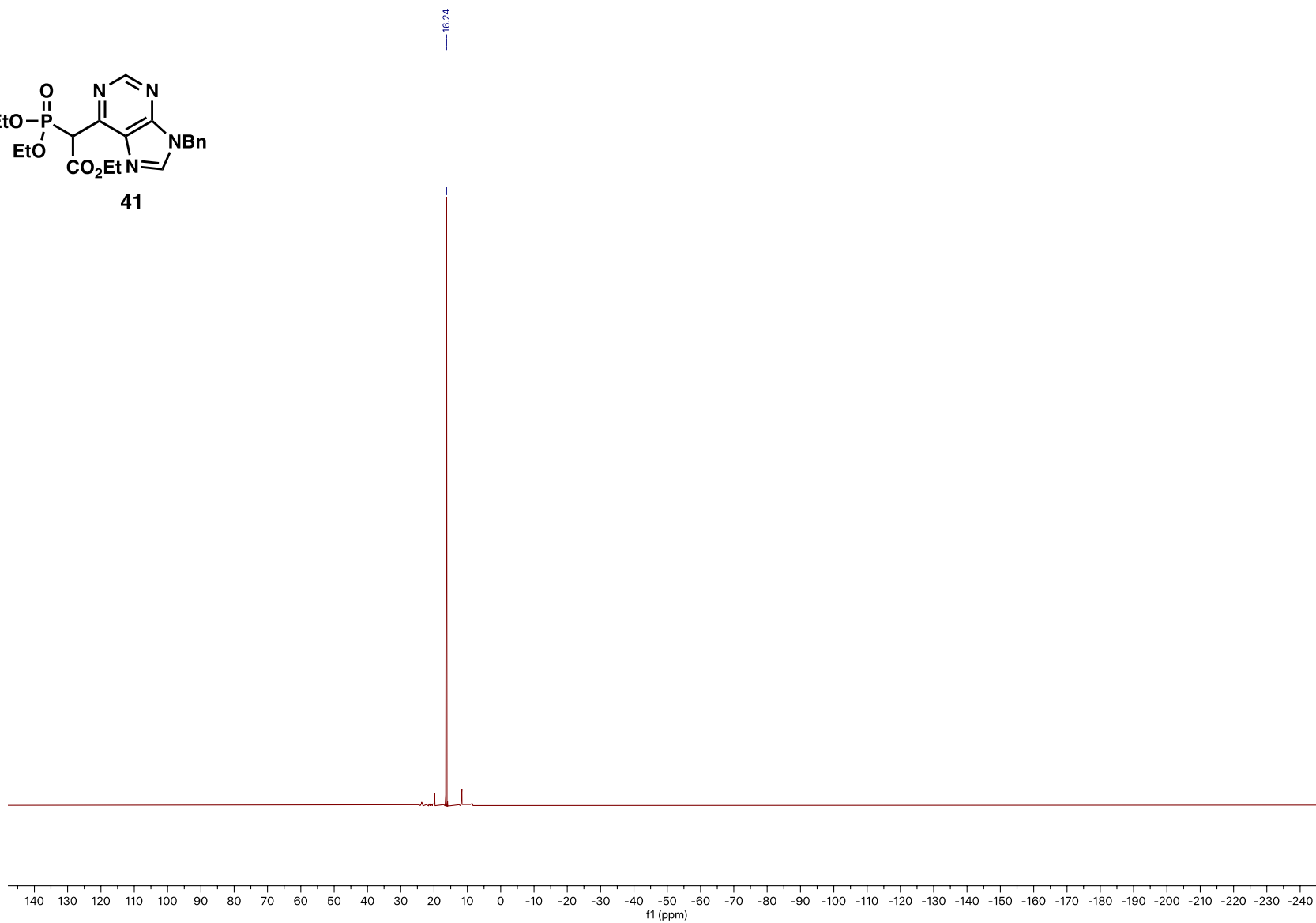


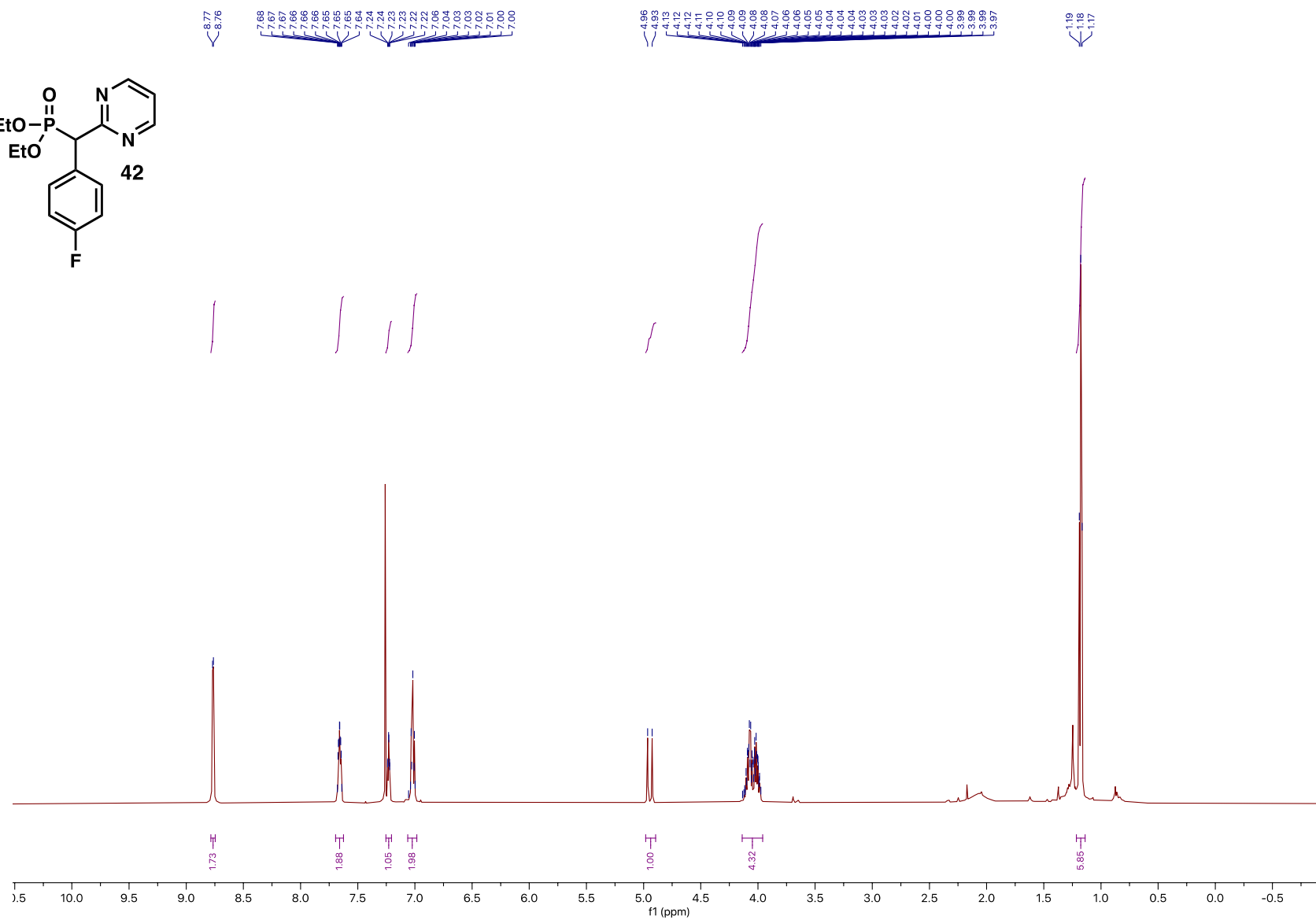
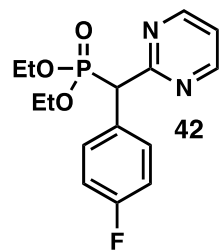


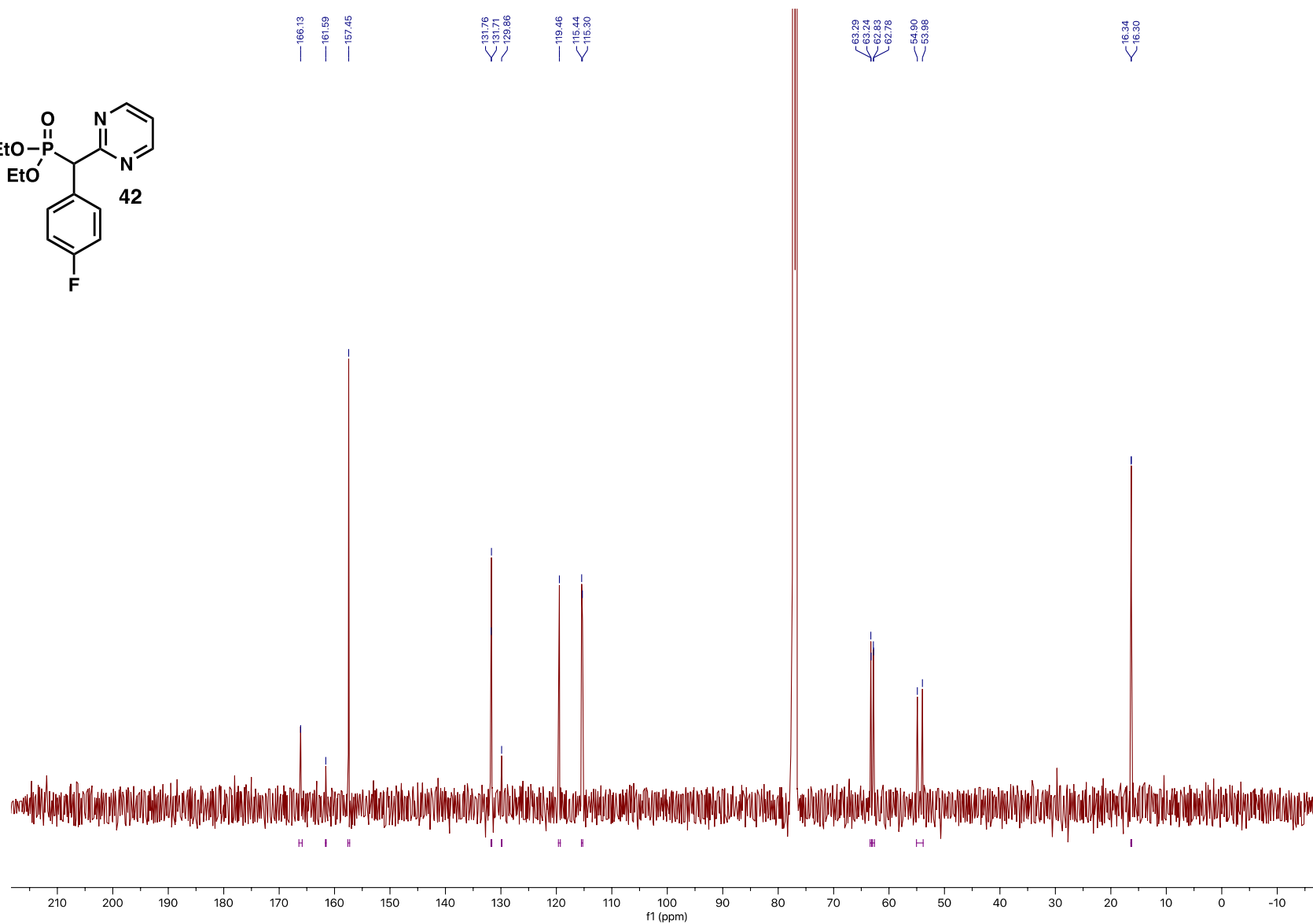
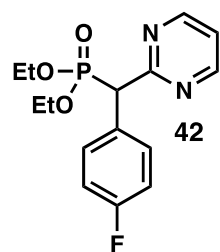


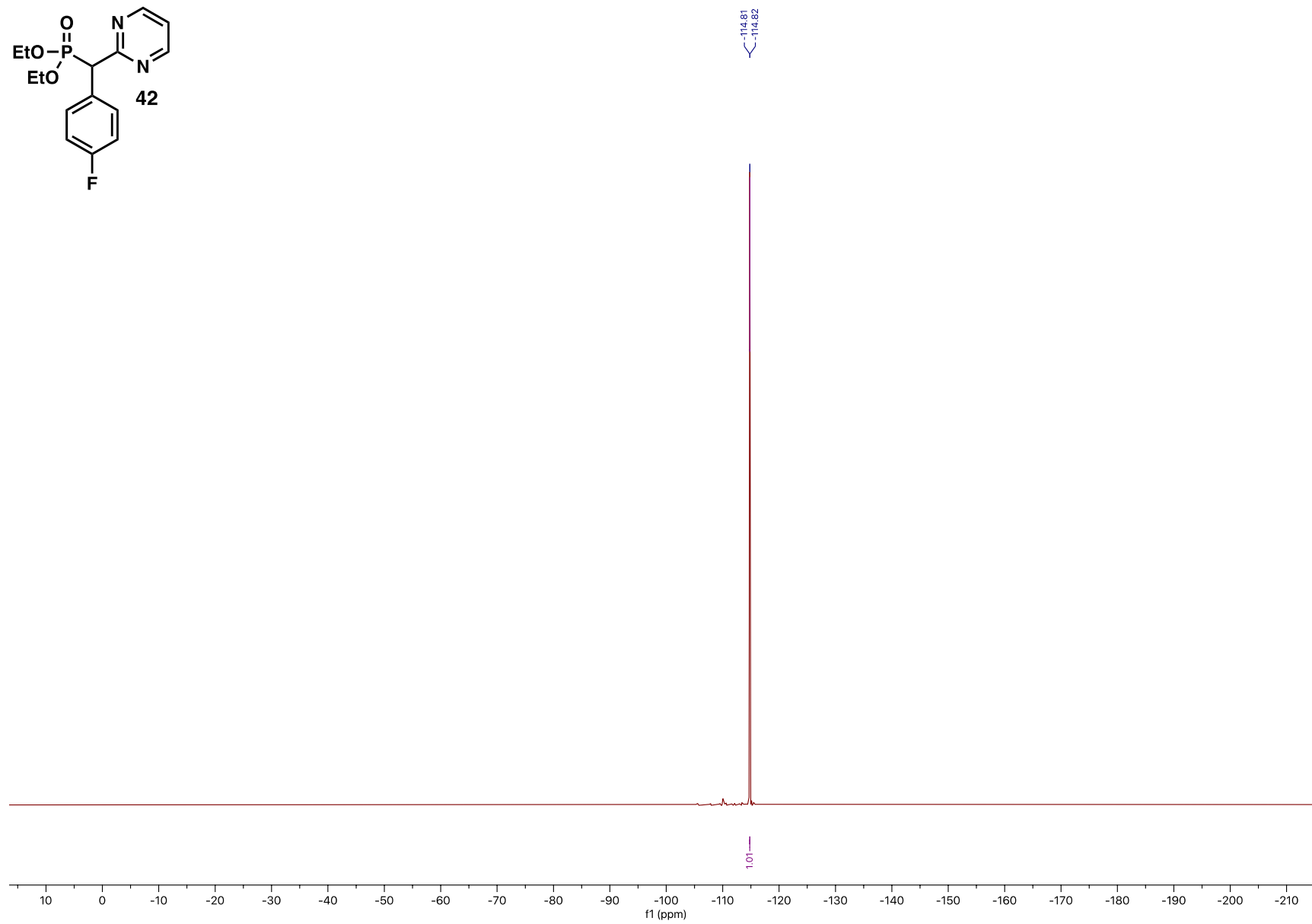
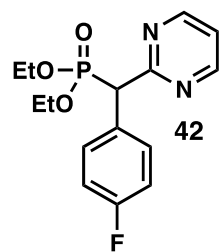


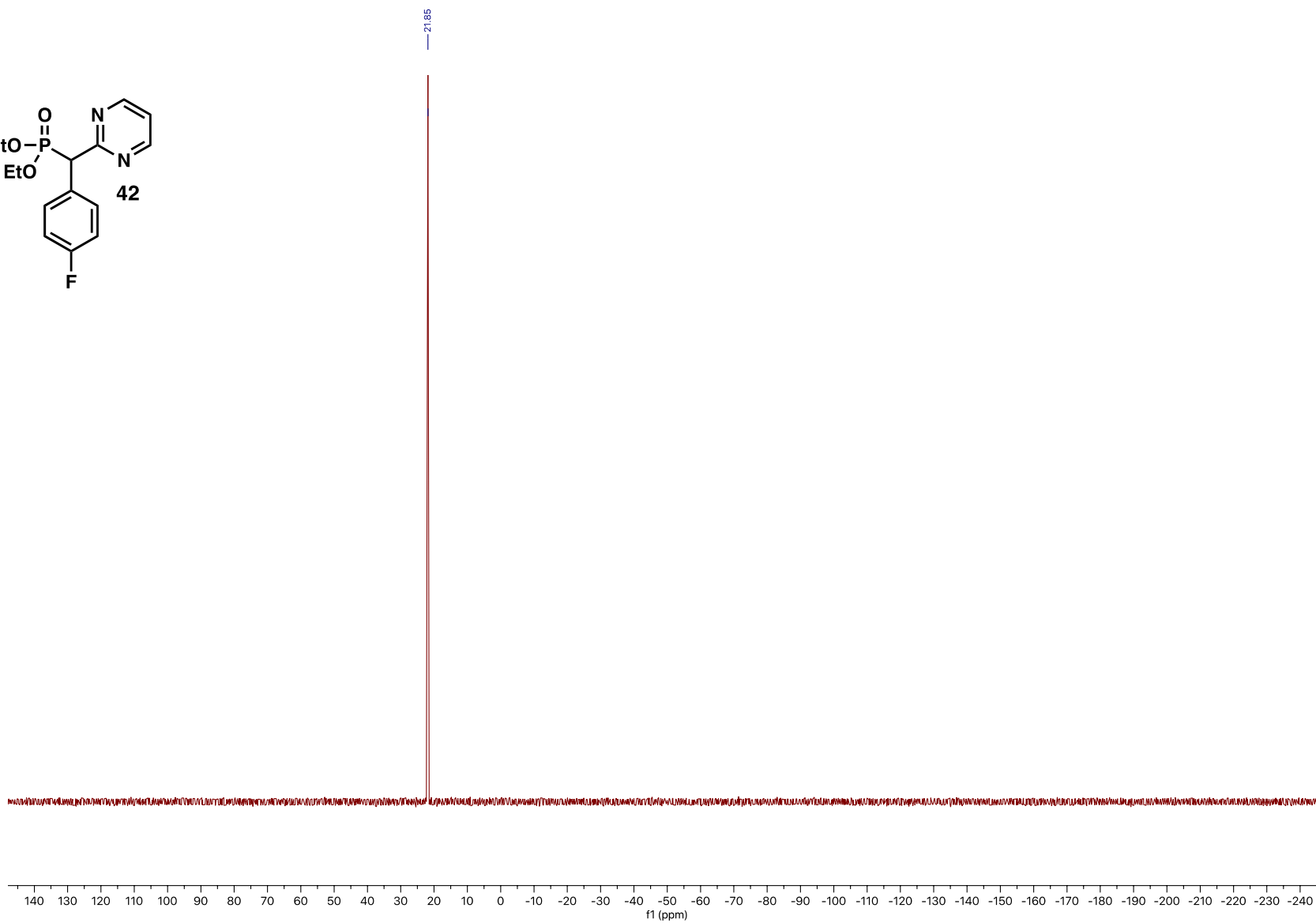
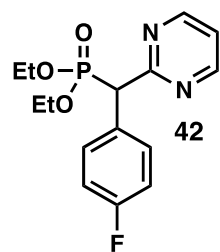
41

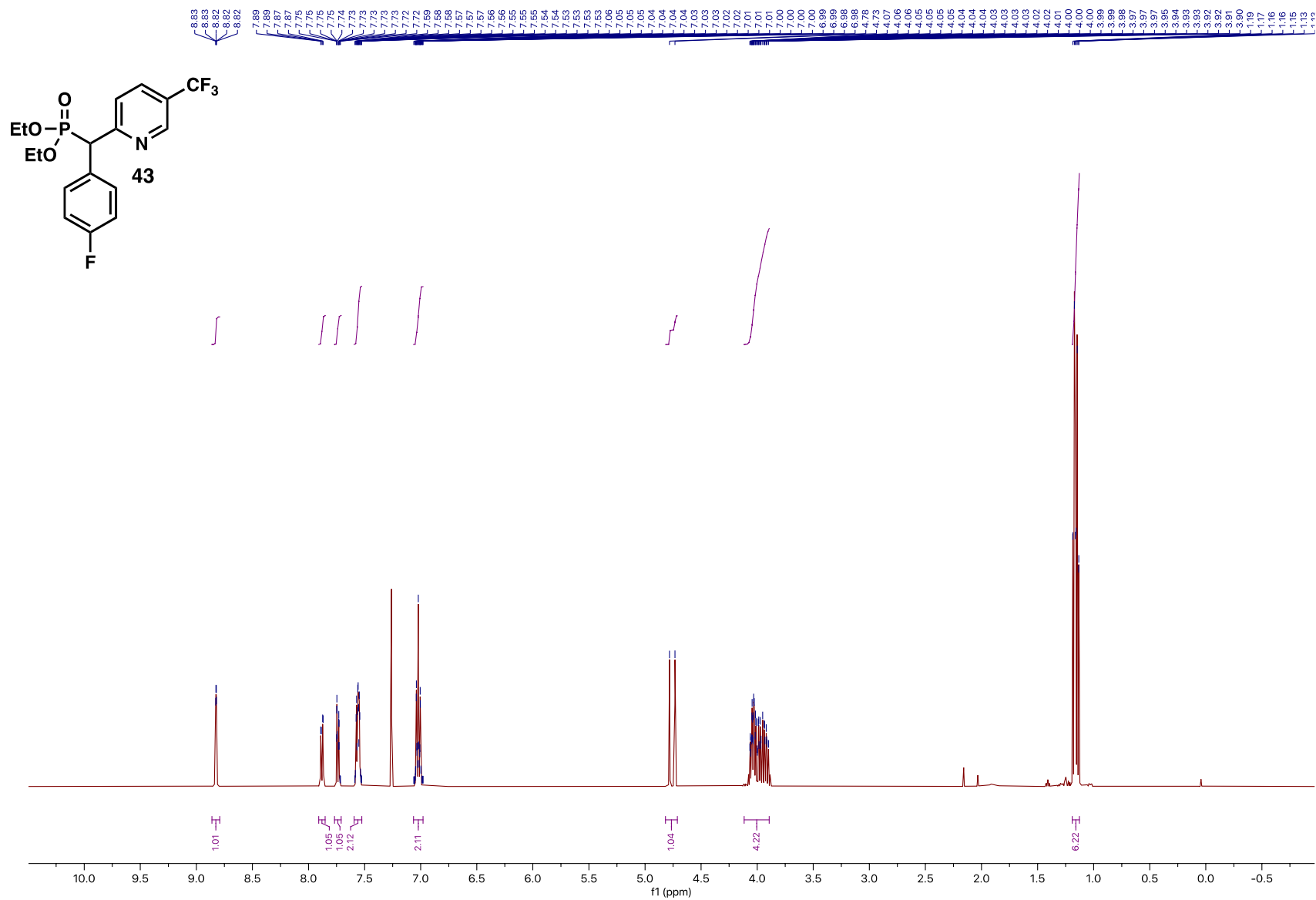


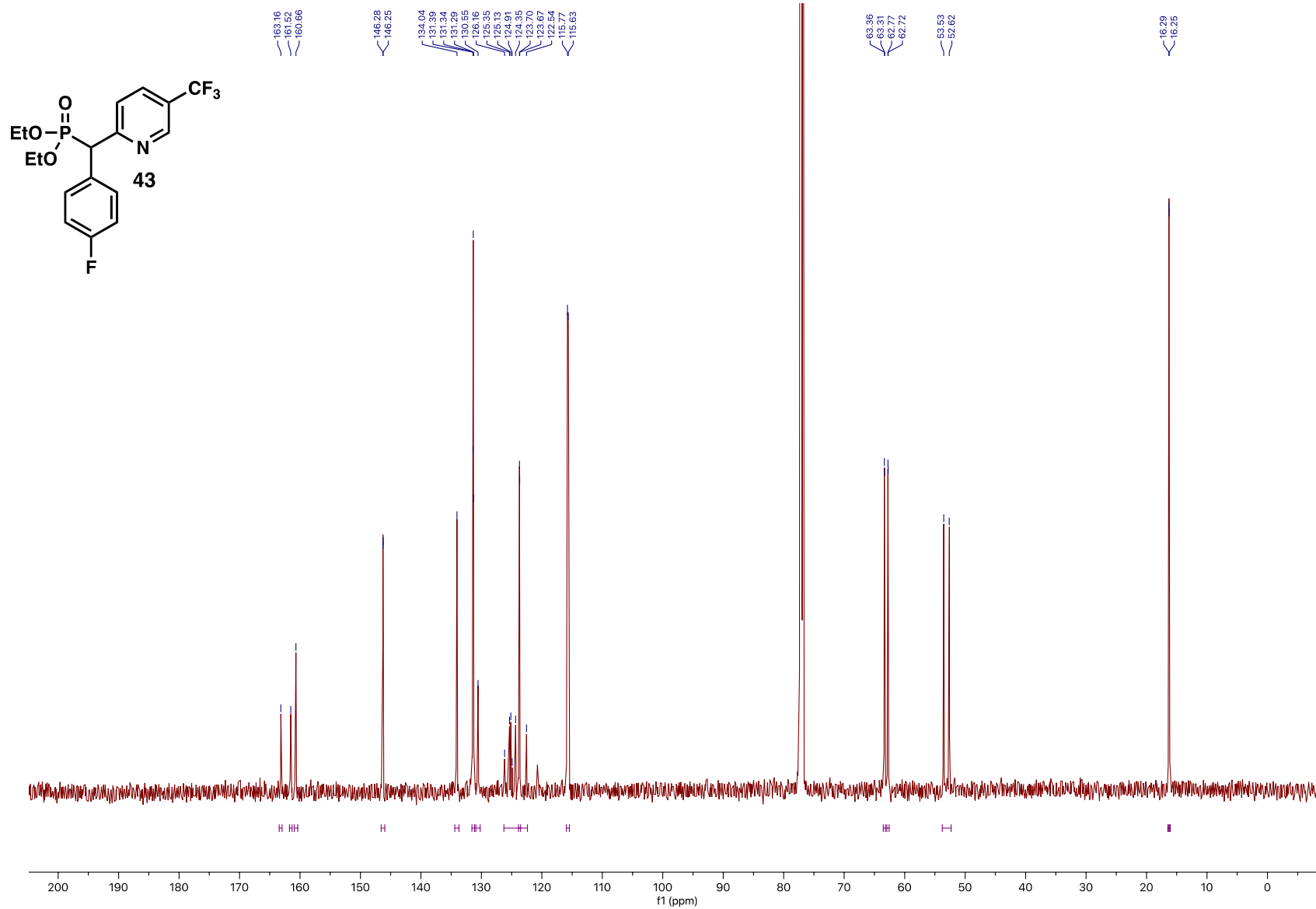


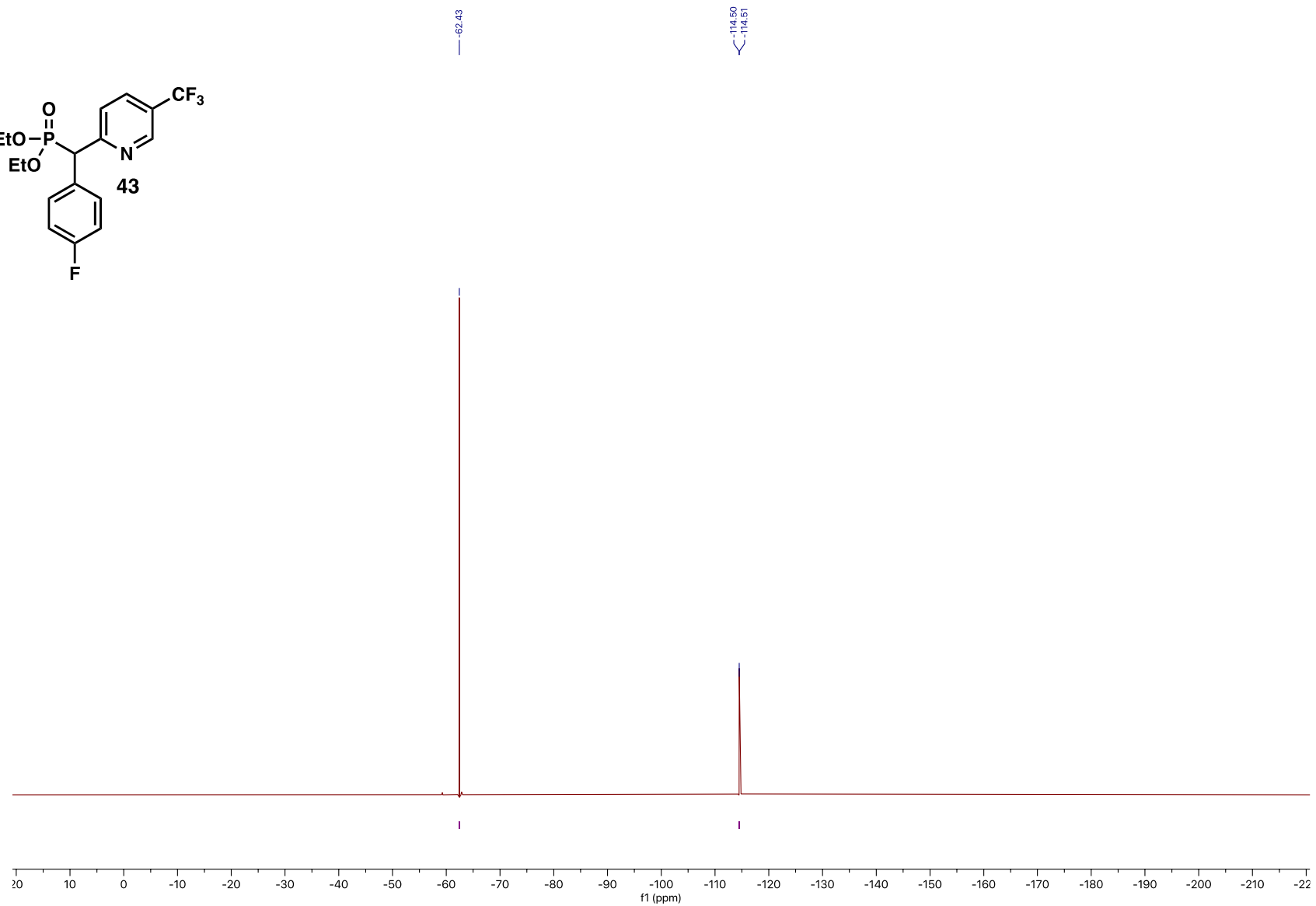
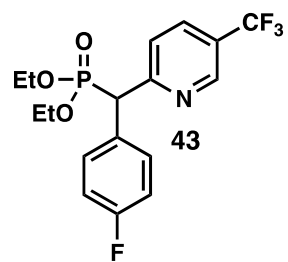


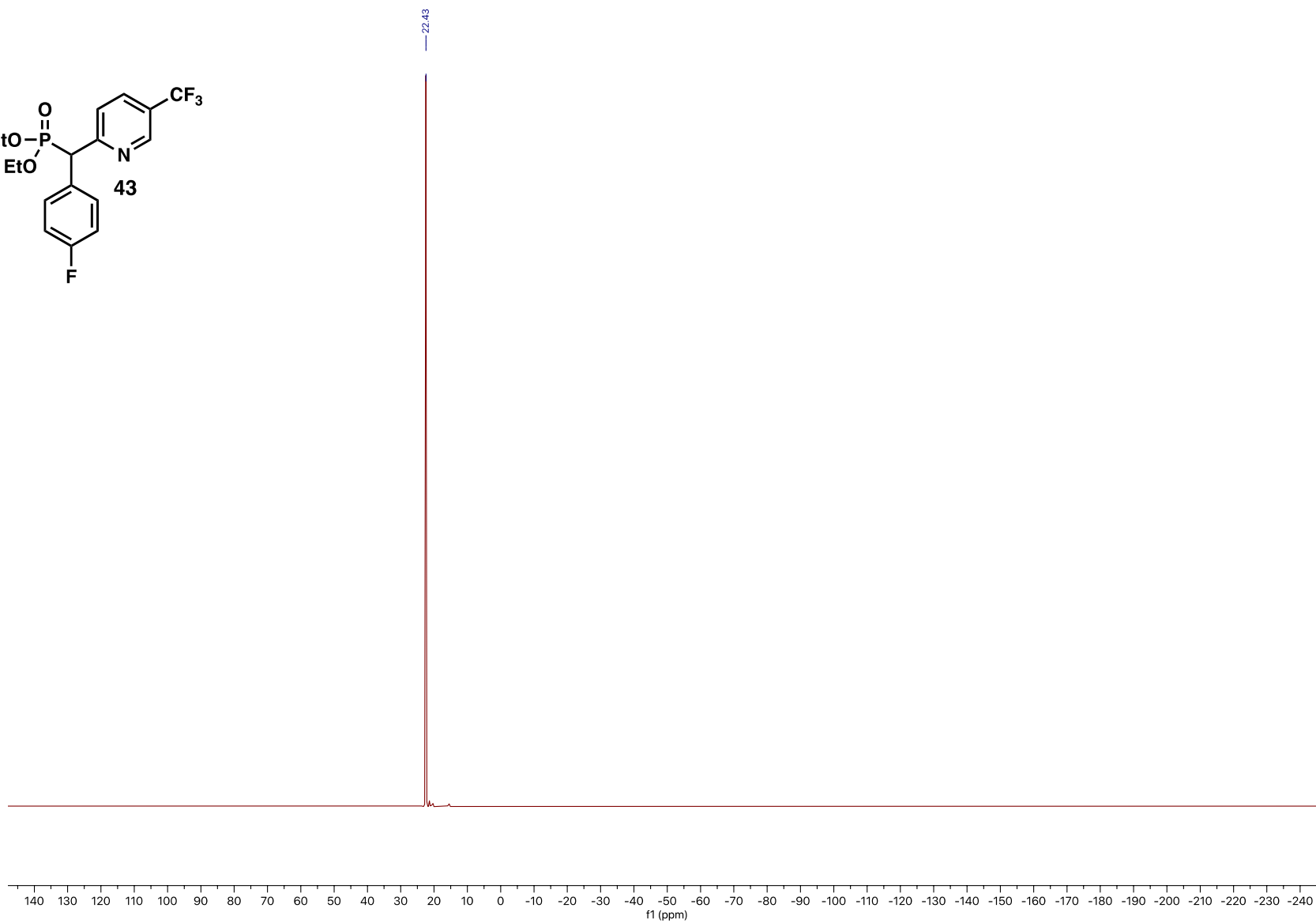
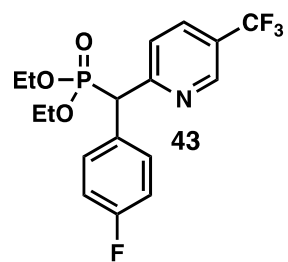


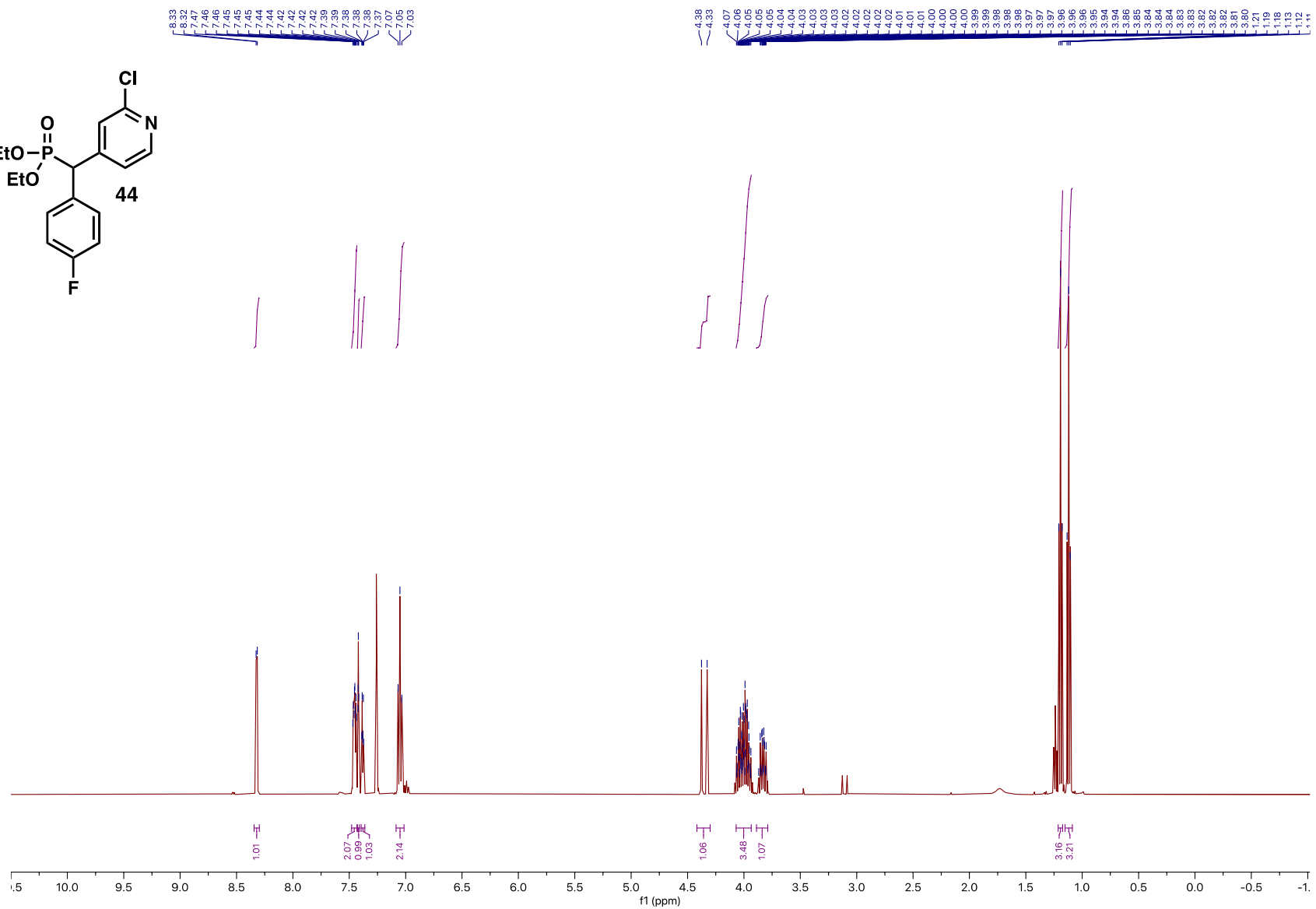
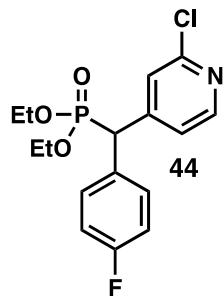


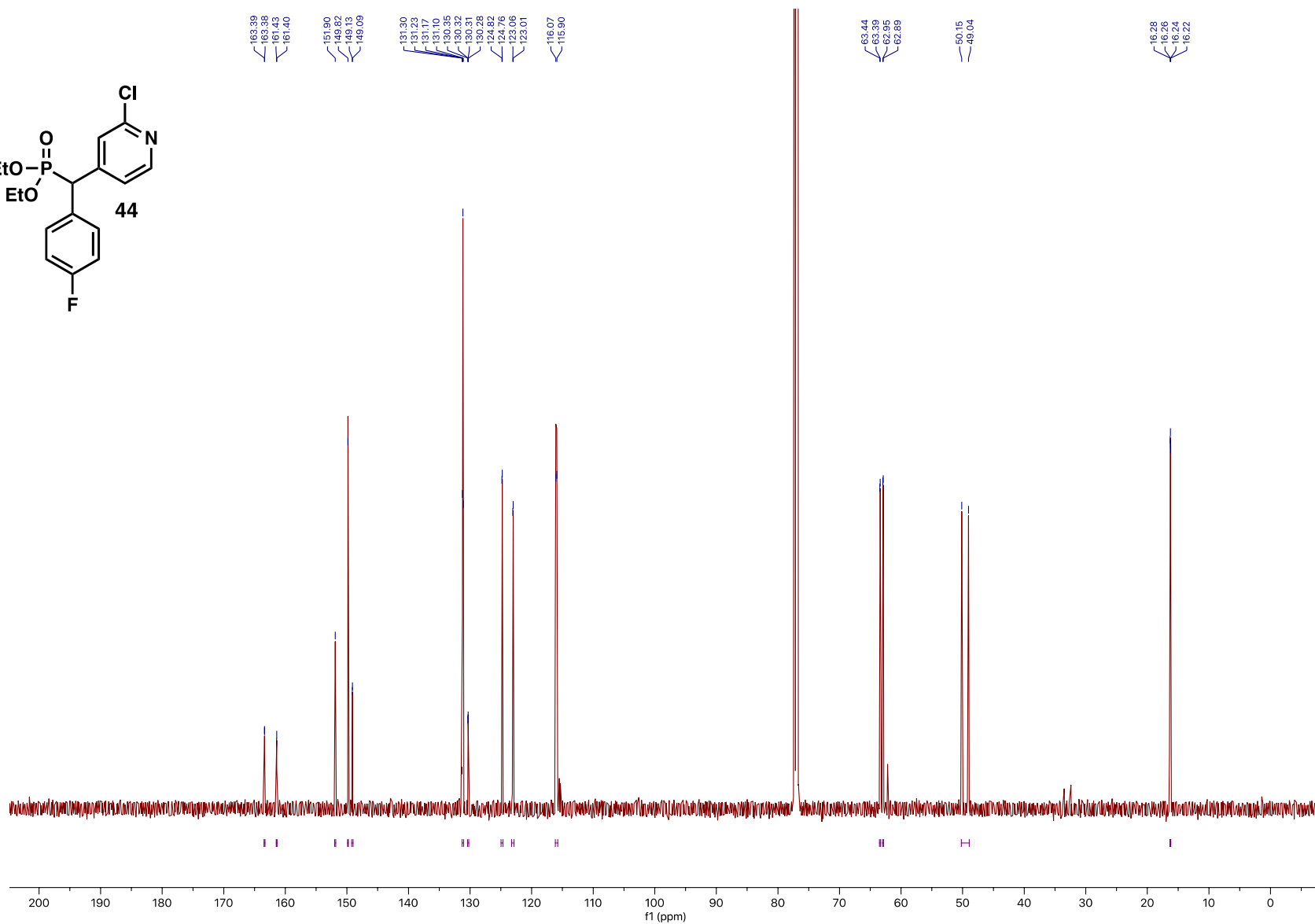
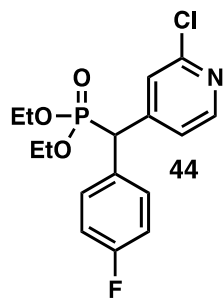


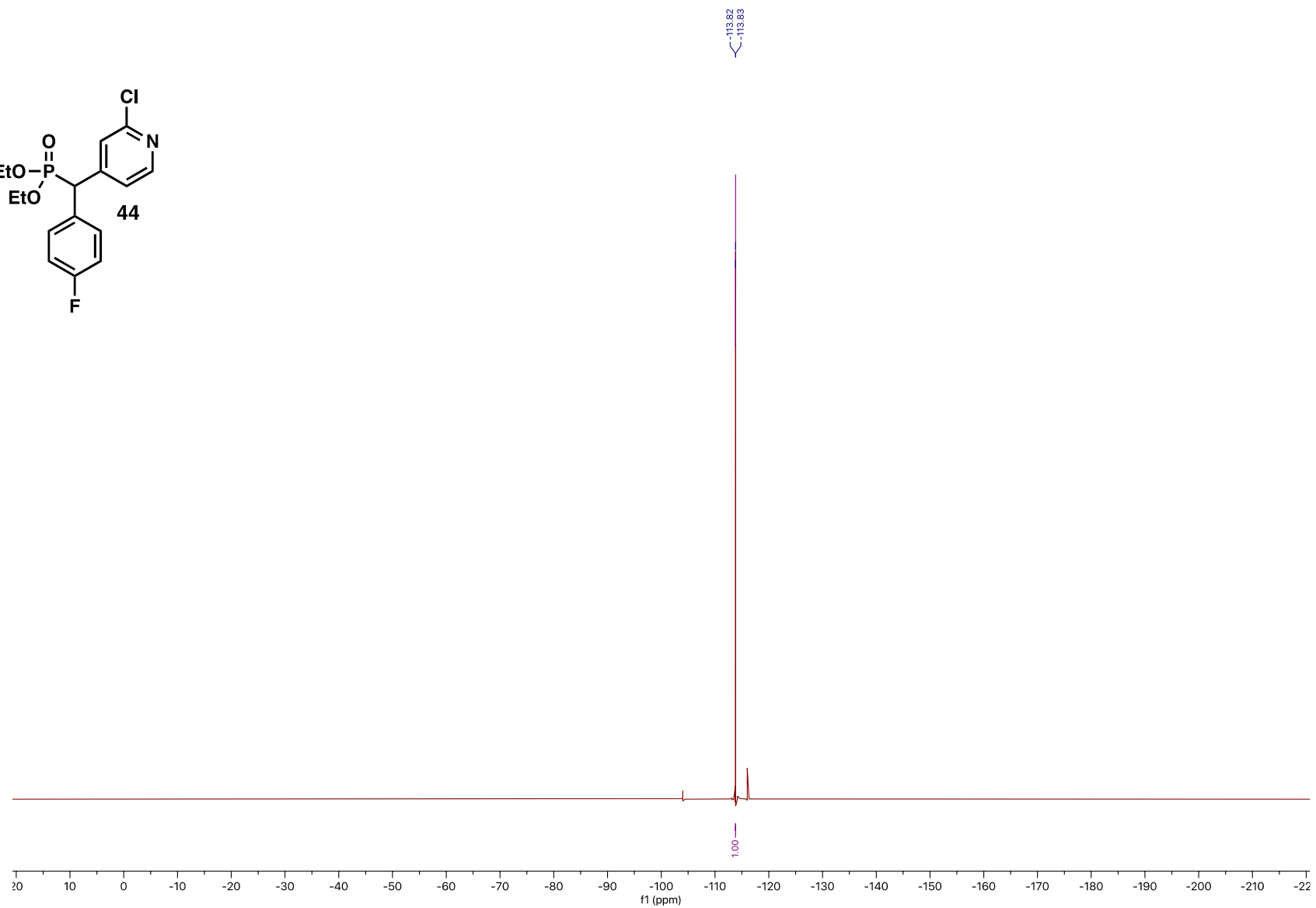
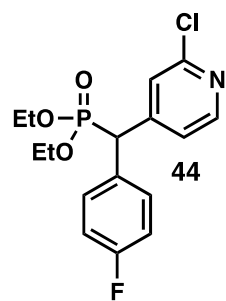


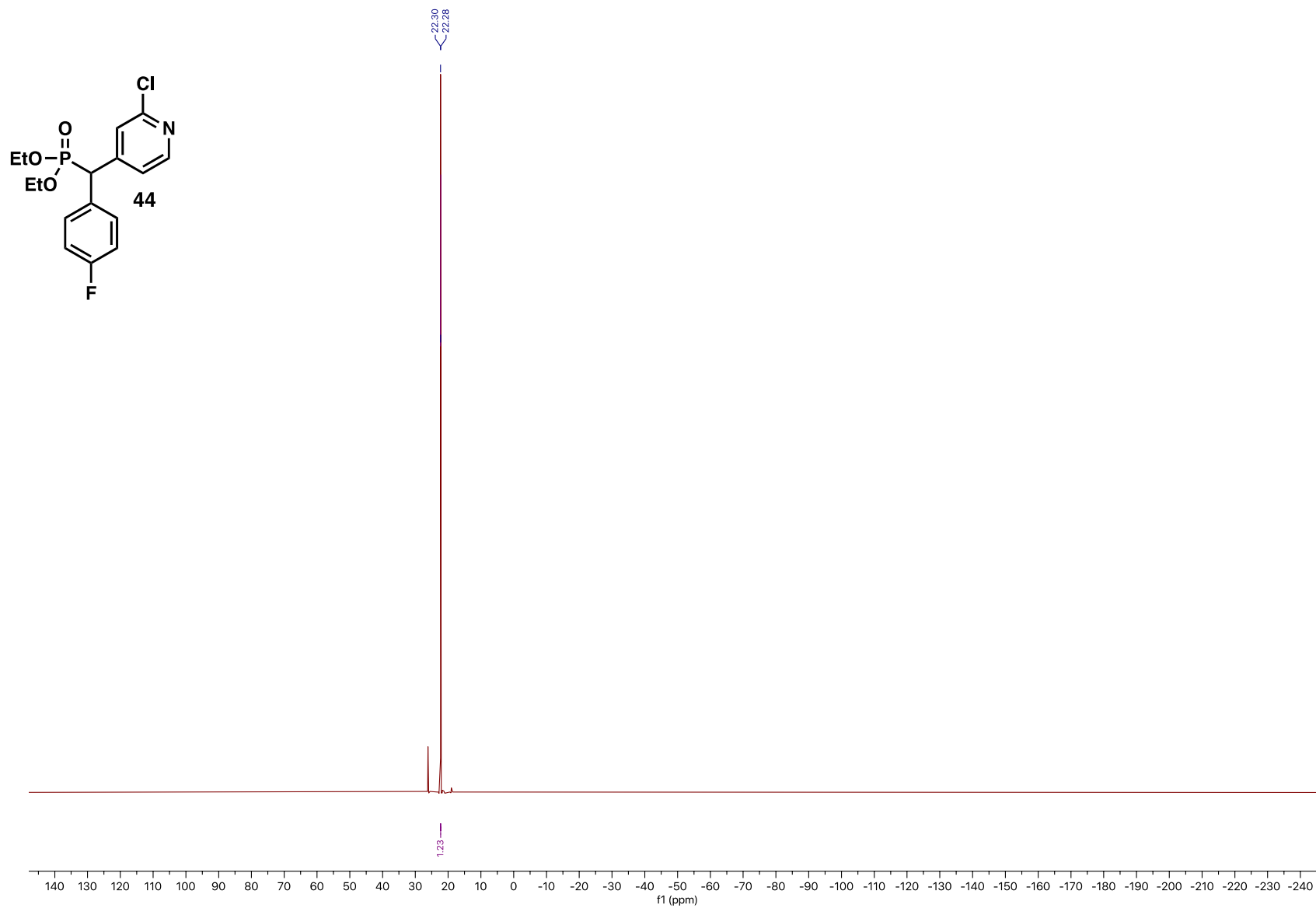


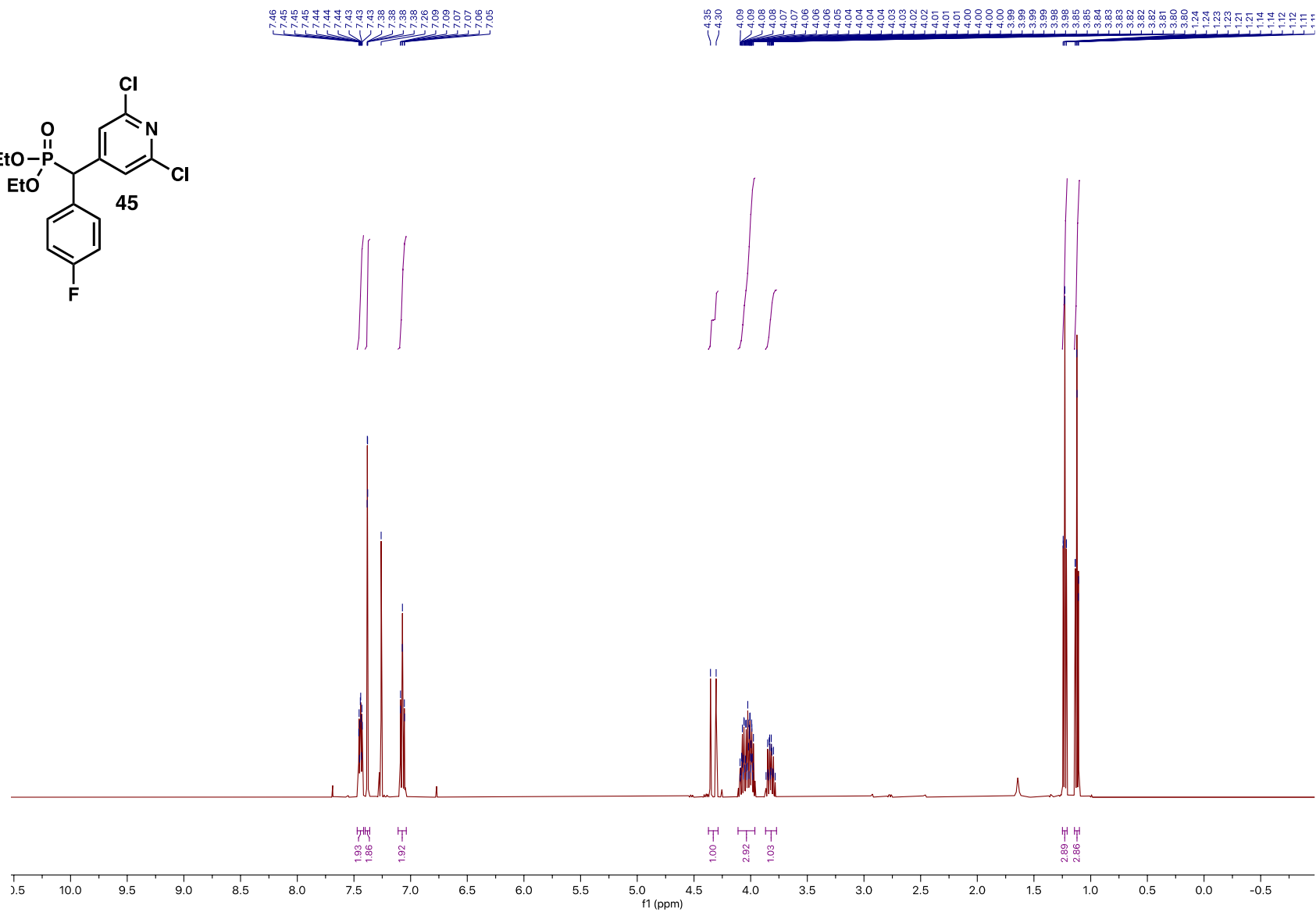
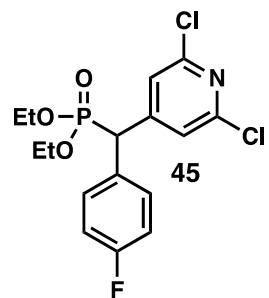


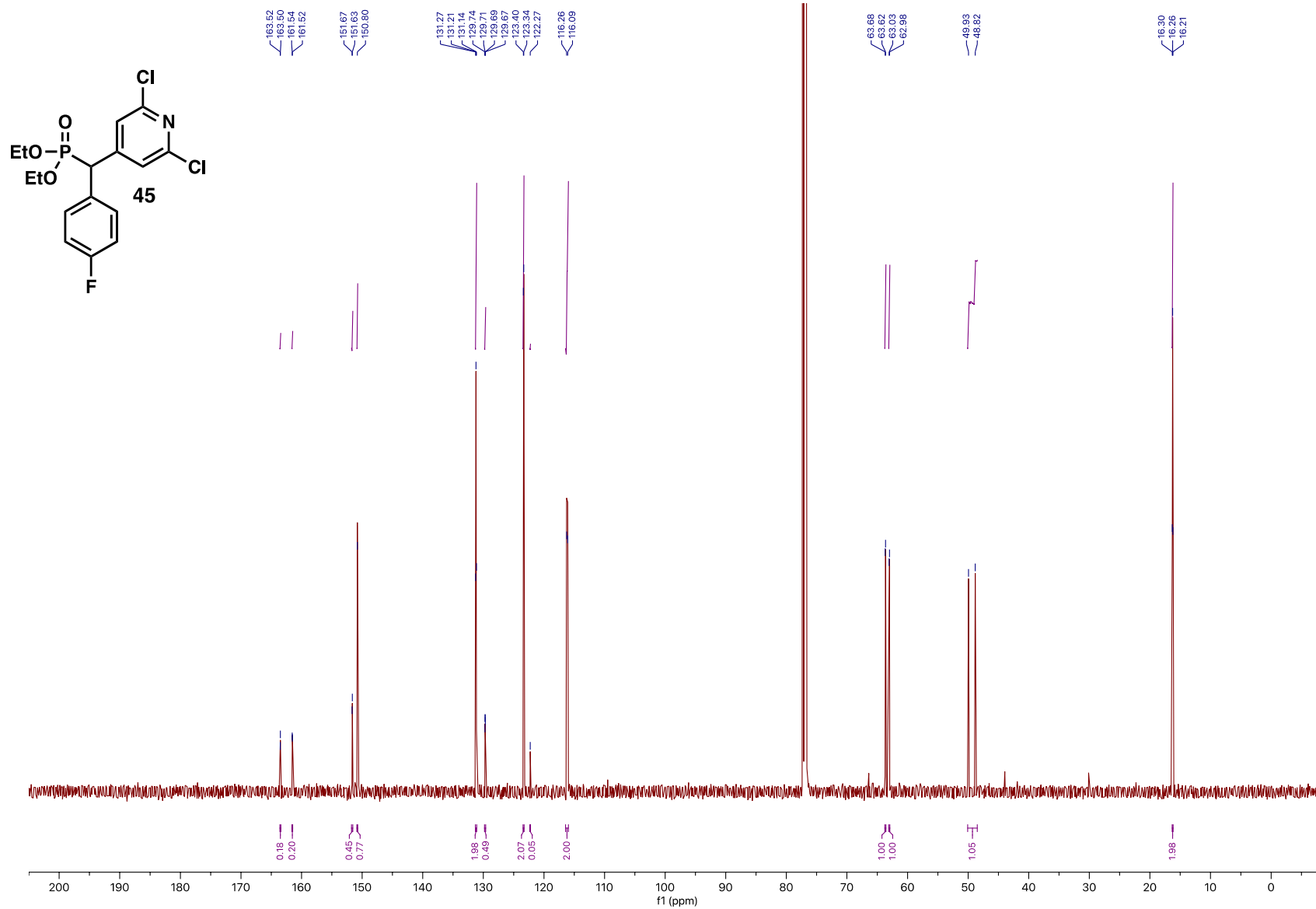


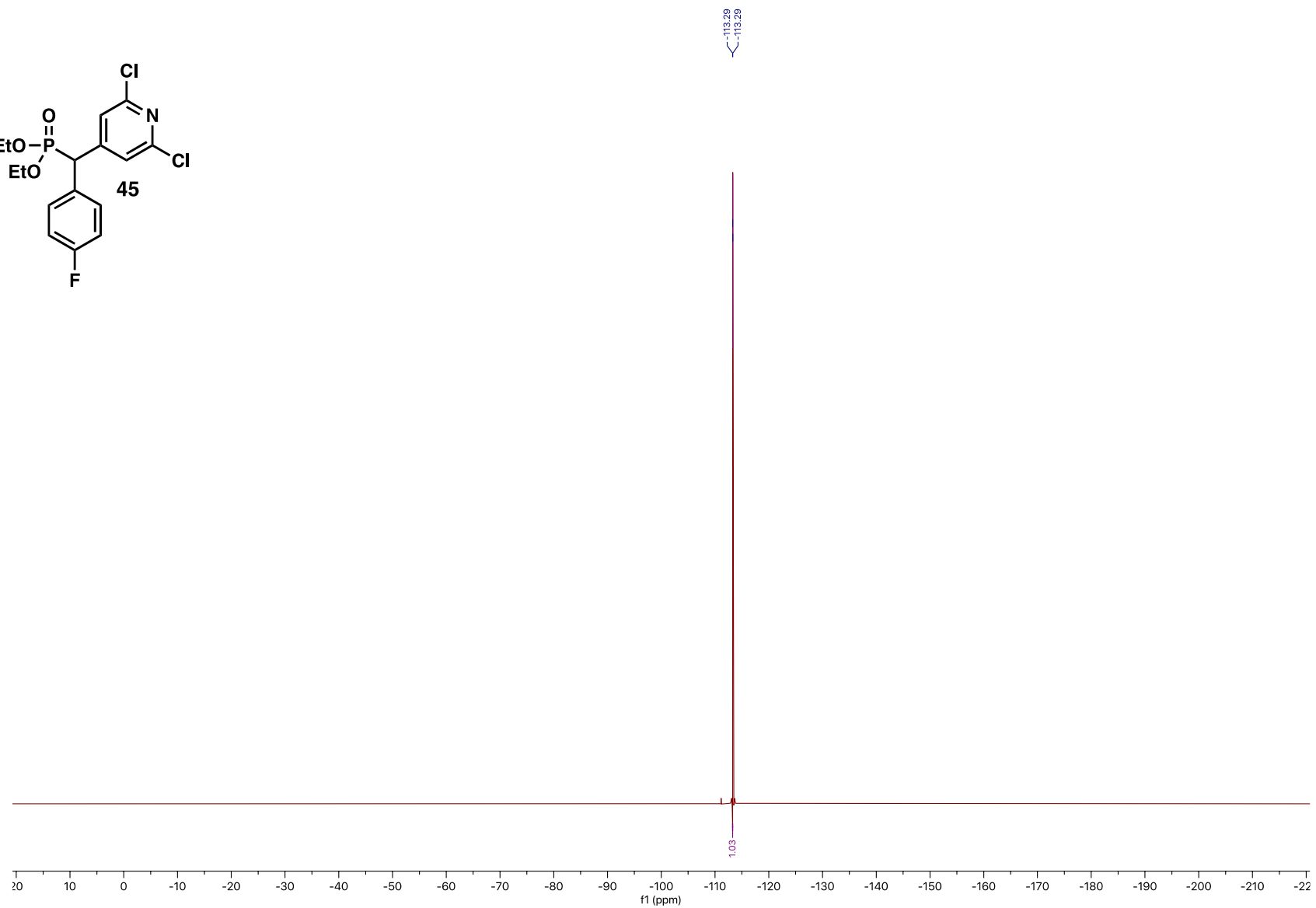
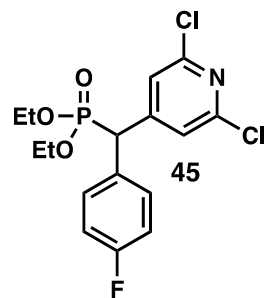


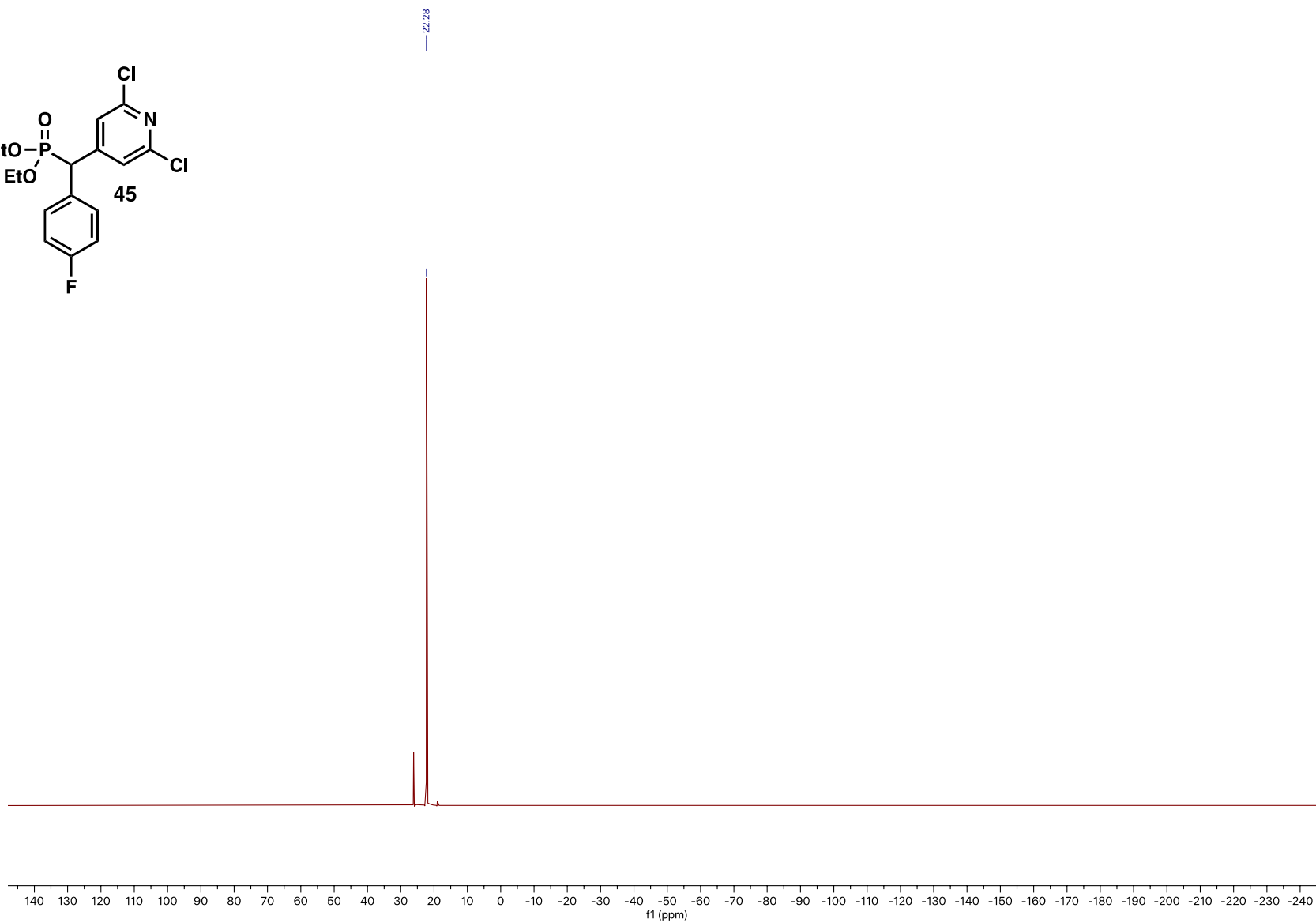
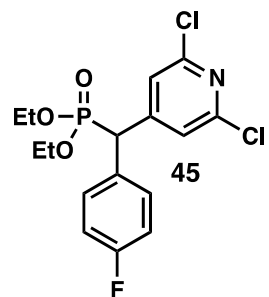


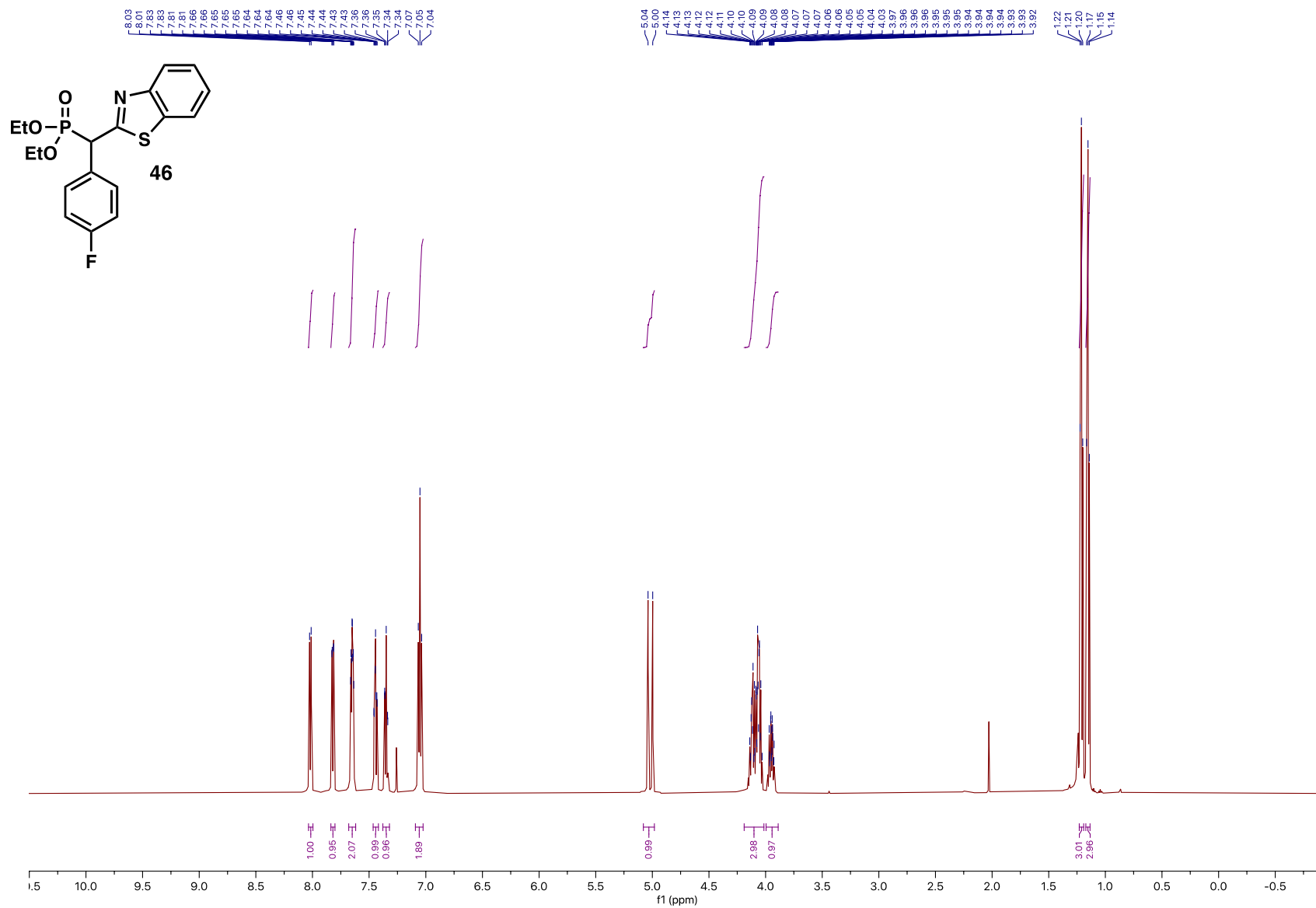


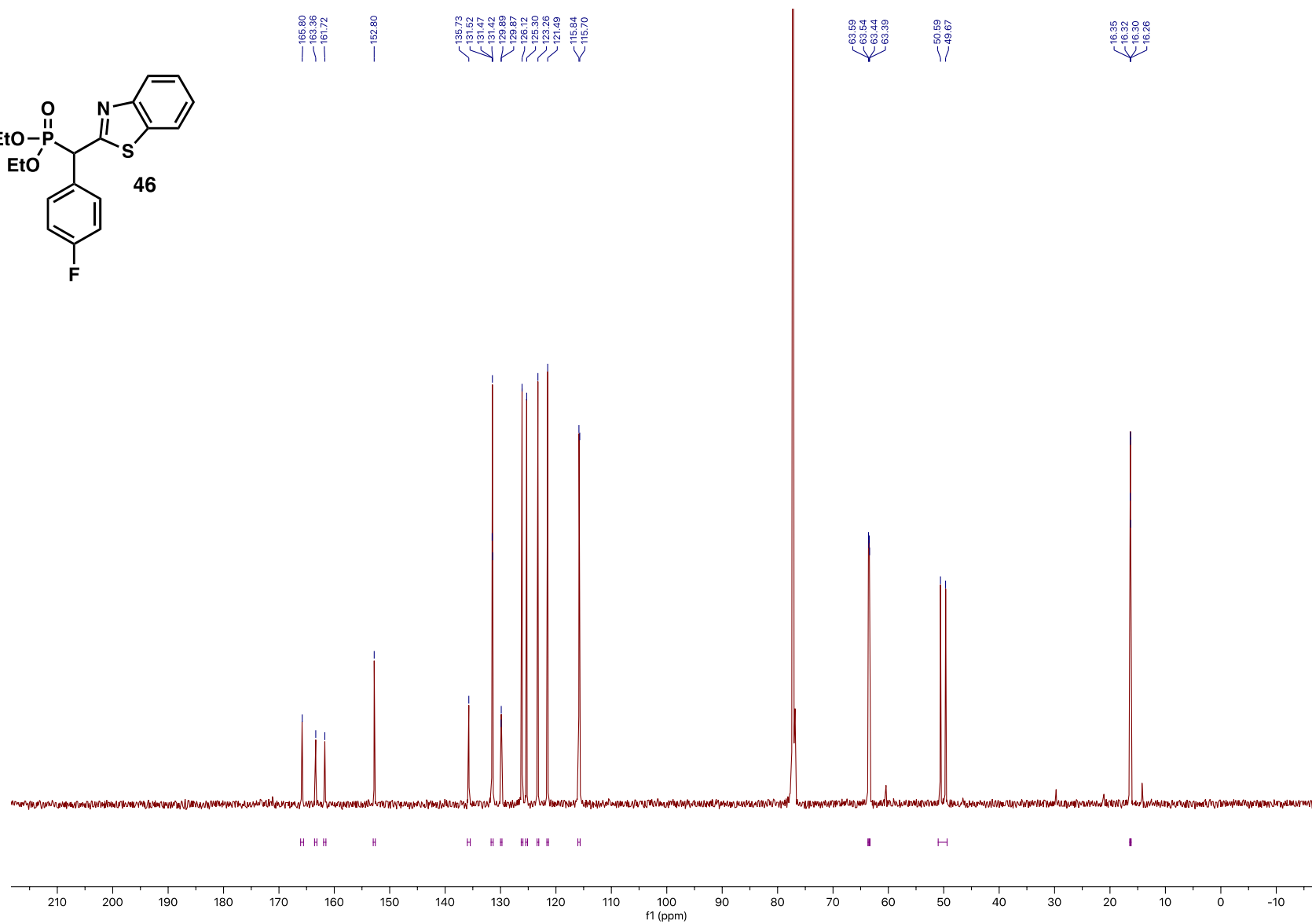
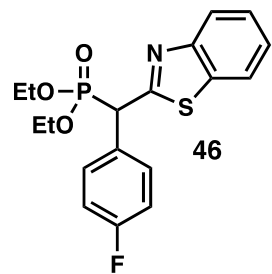


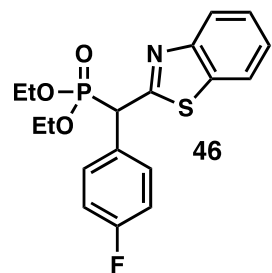




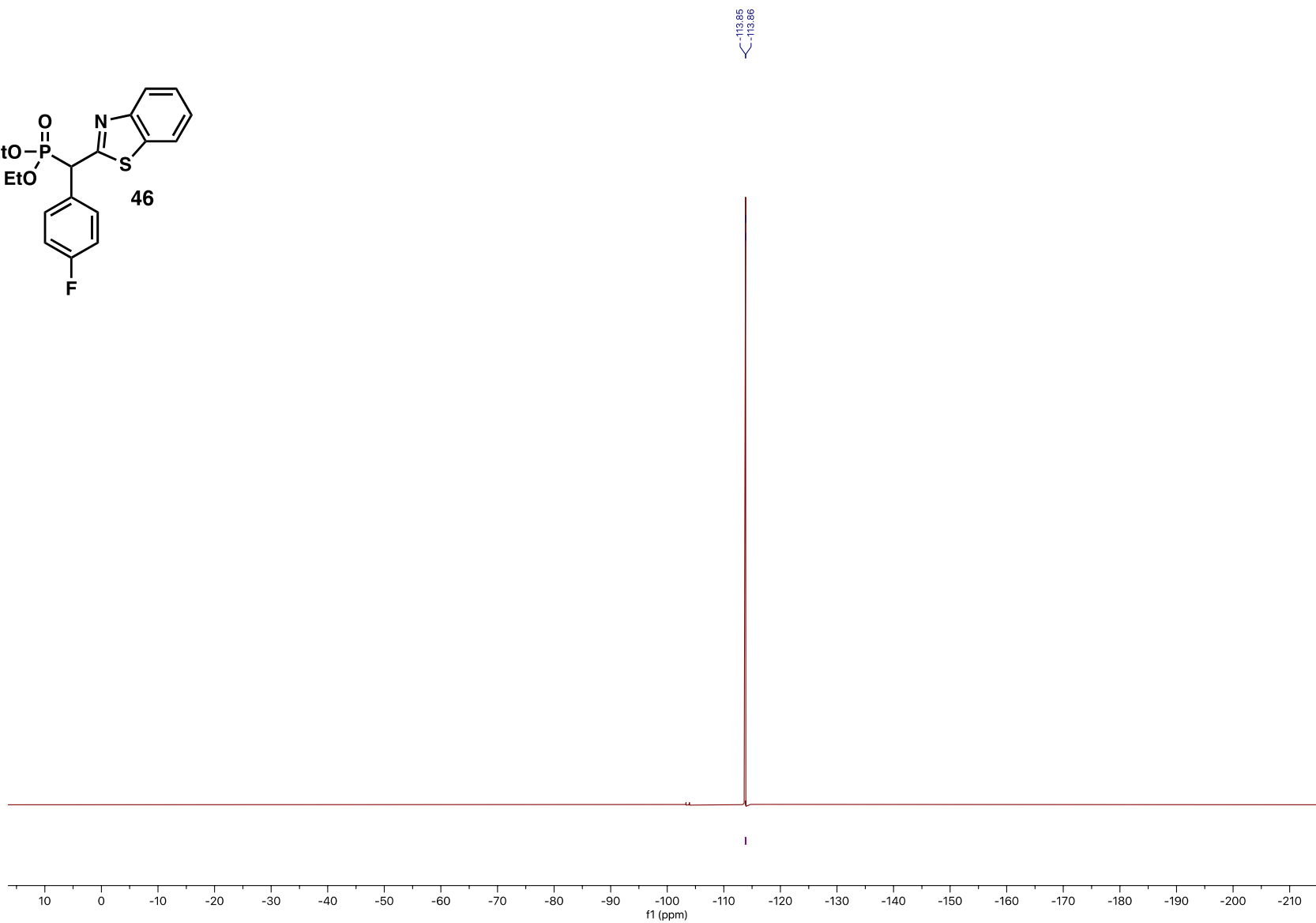


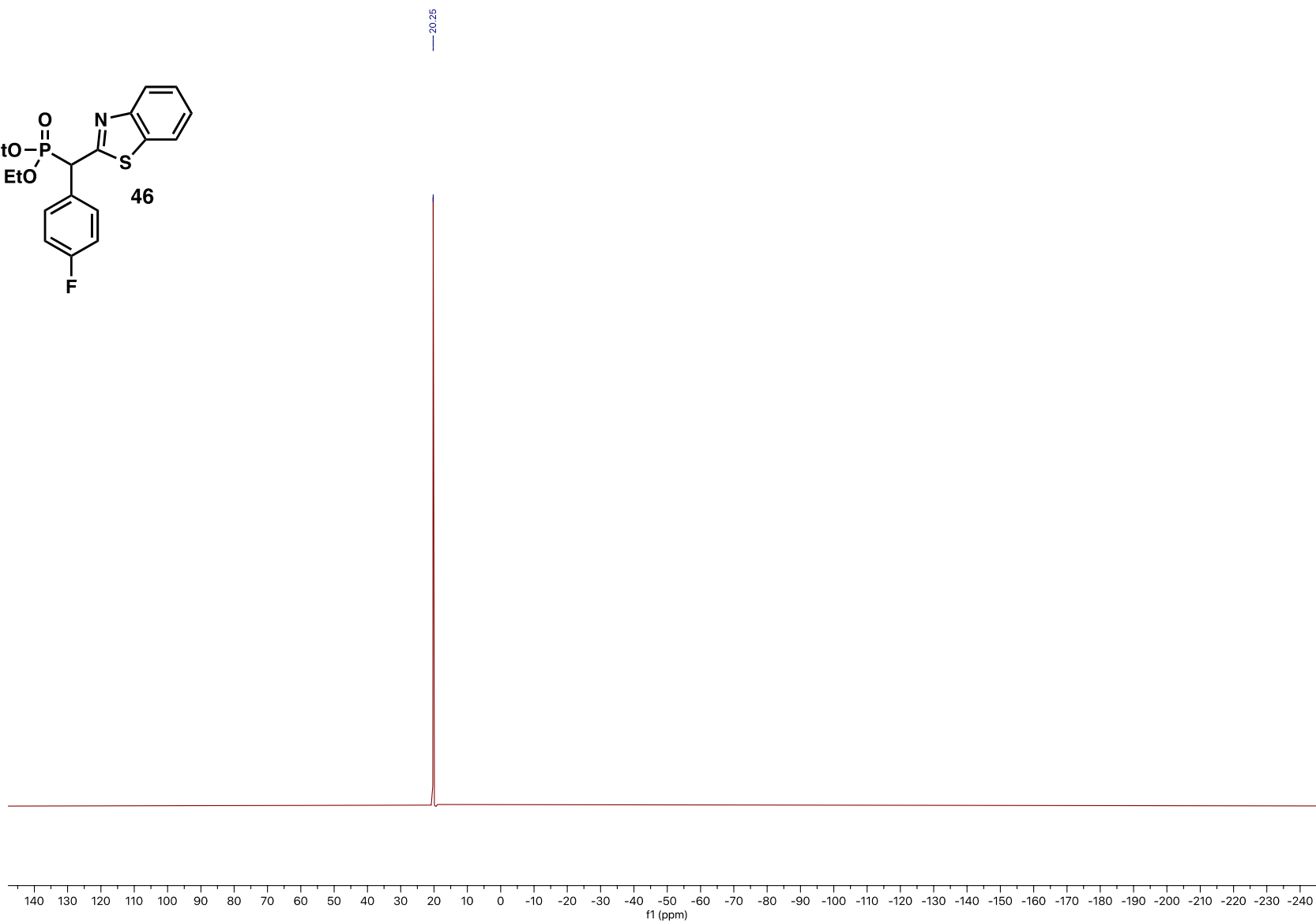
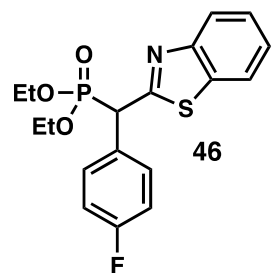


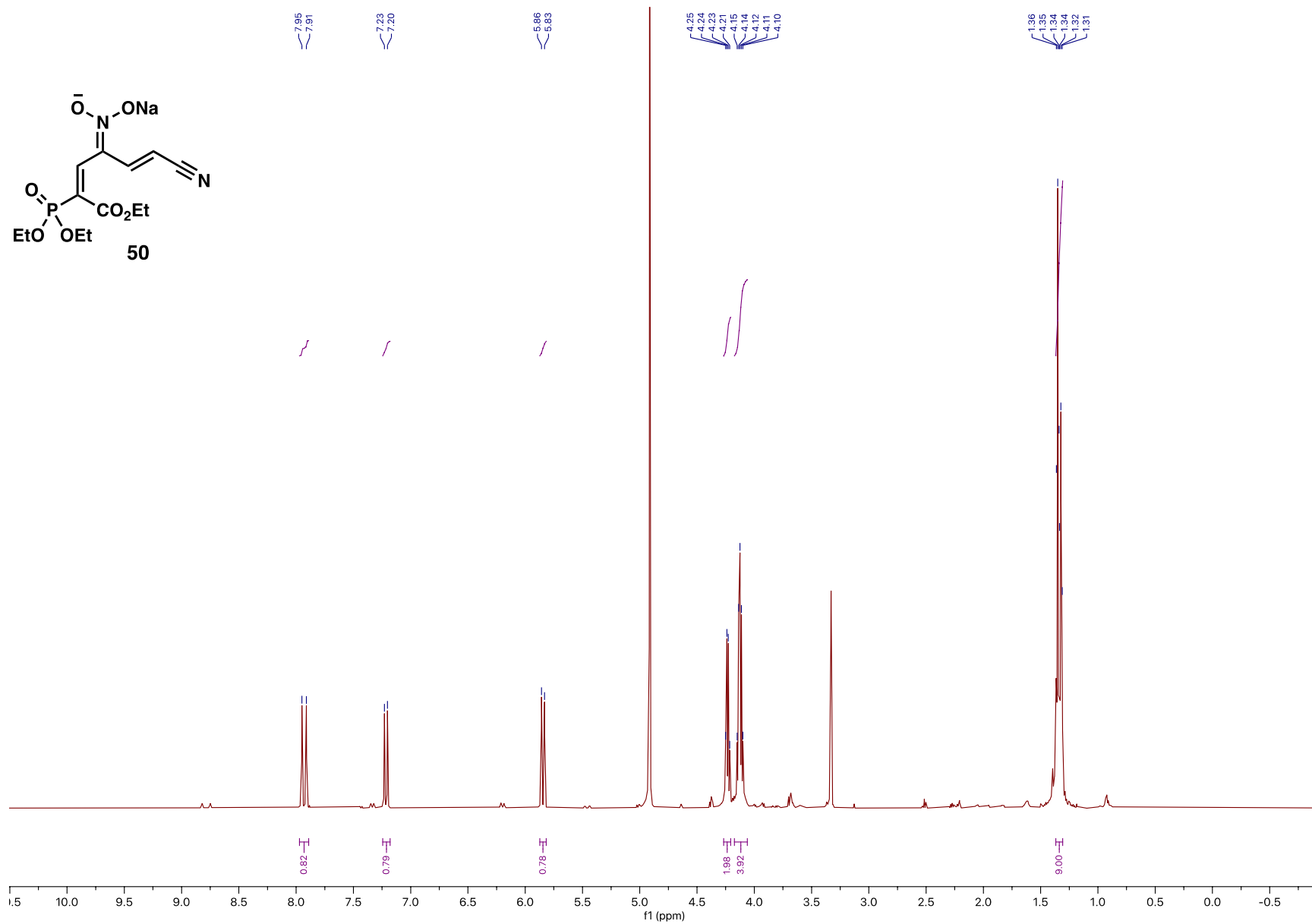


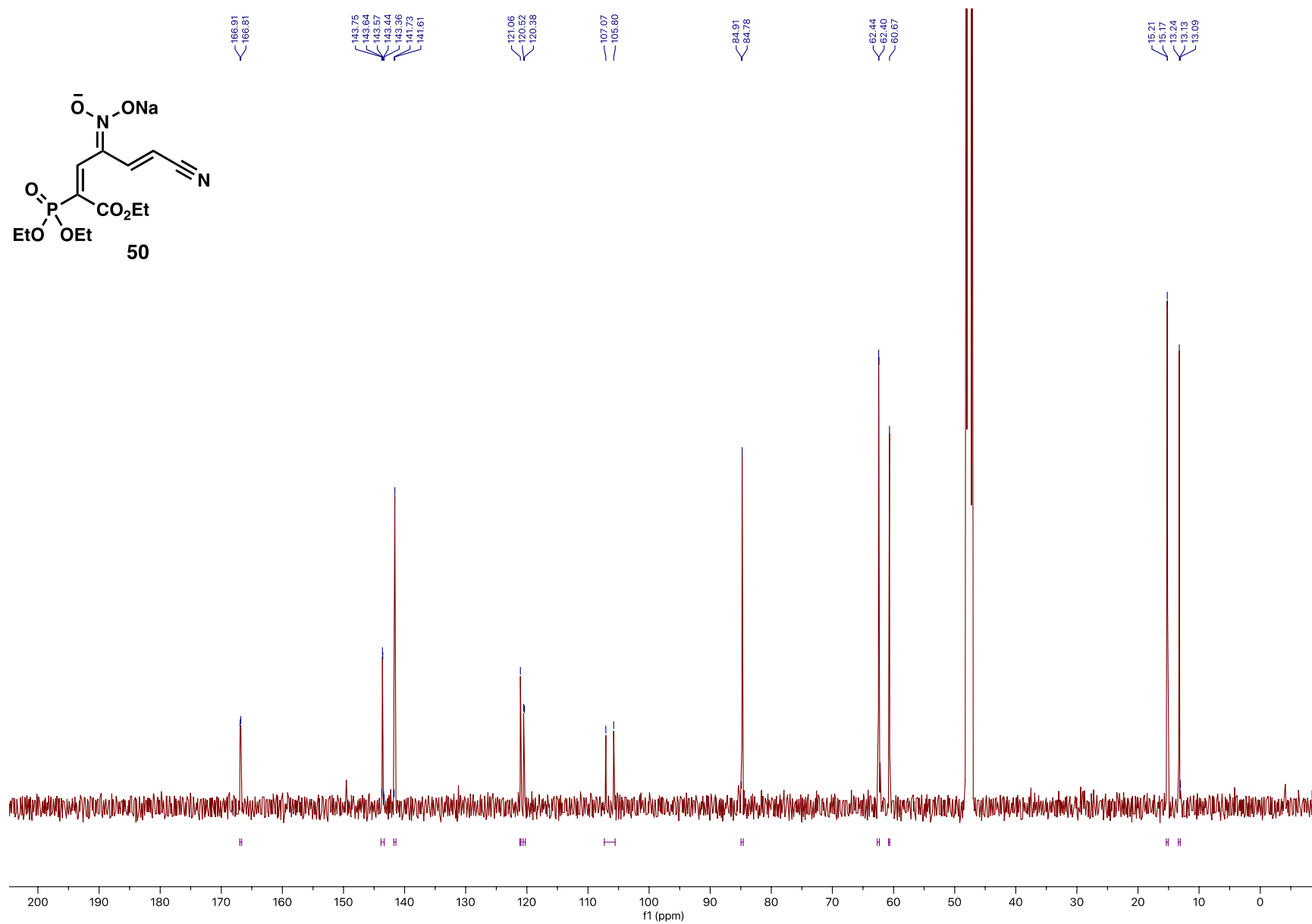
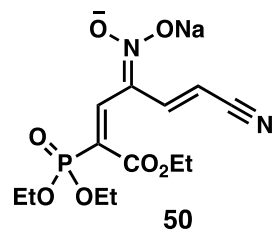


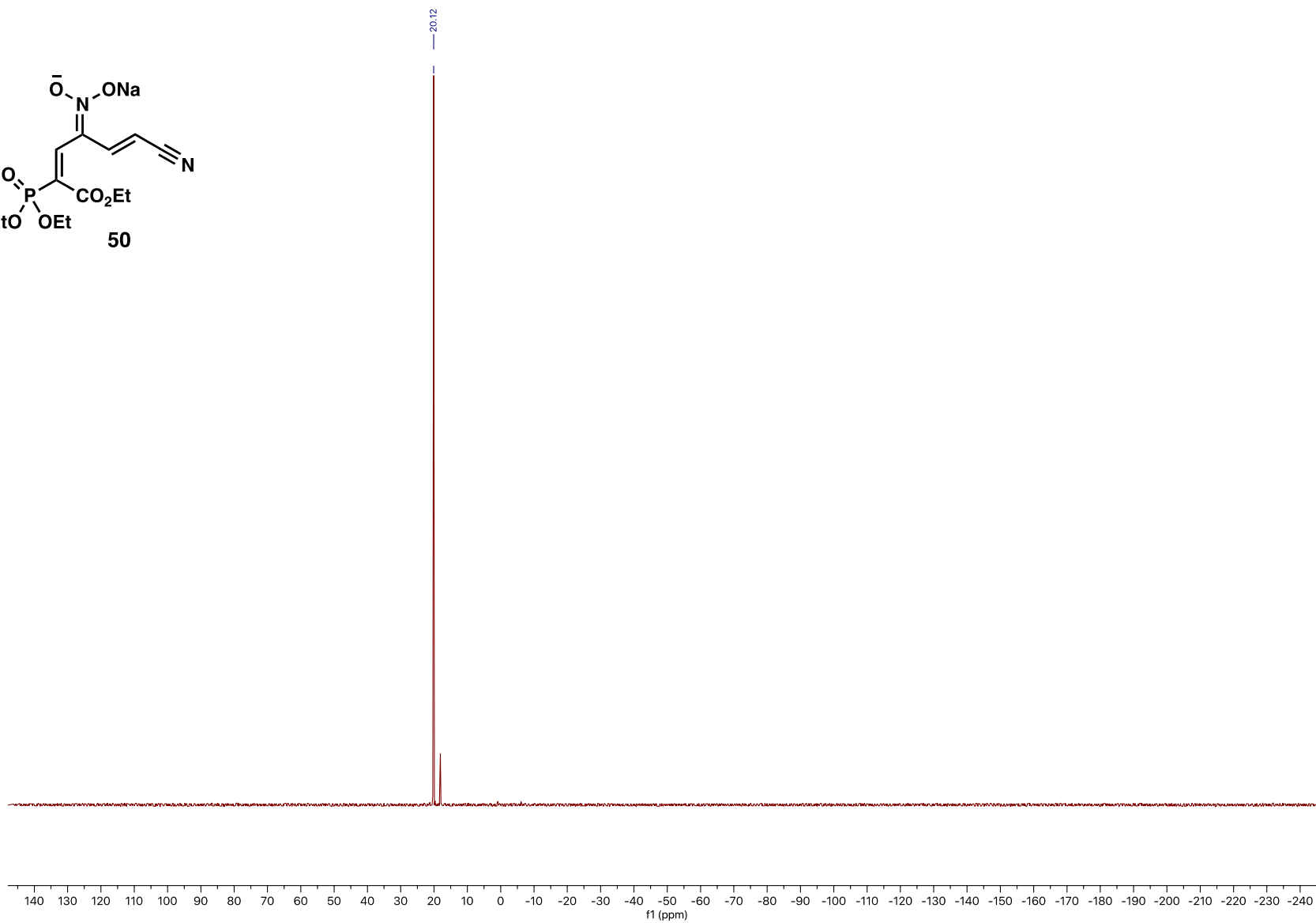
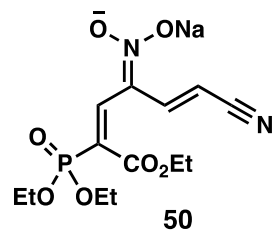
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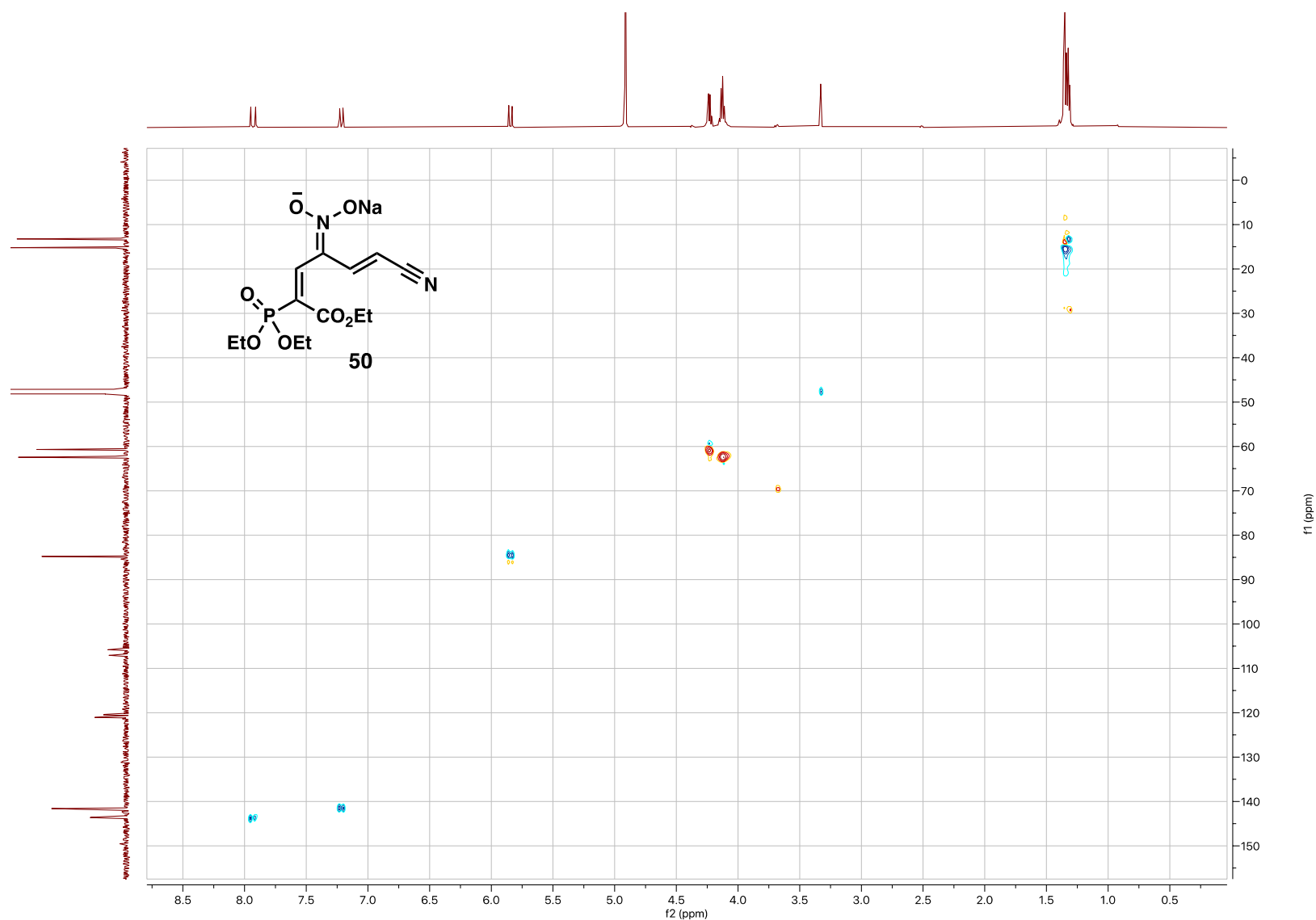








HSQC of 50:



Single-Crystal X-Ray Crystallography of Compound 30

X-ray diffraction data were measured on Bruker D8 Venture PHOTON II CPAD diffractometer equipped with a Cu K α INCOATEC ImuS micro-focus source ($\lambda = 1.54178 \text{ \AA}$). Indexing was performed using *APEX3* [1] (Difference Vectors method). Data integration and reduction were performed using SaintPlus [2]. Absorption correction was performed by multi-scan method implemented in SADABS [3]. Space groups were determined using XPREP

implemented in APEX3 [1]. Structure was solved using SHELXT [4] and refined using SHELXL-2018 [5] (full-matrix least-squares on F^2) through OLEX2 interface program [6]. Disordered -OEt group was refined with restraints / constraint. Crystal data and refinement conditions are shown in Table 1.

[1] Bruker (2019). *APEX3* Bruker AXS Inc., Madison, Wisconsin, USA.

[2] Bruker (2019) SAINT V8.35A. Data Reduction Software.

[3] Sheldrick, G. M. (1996). *SADABS. Program for Empirical Absorption Correction*. University of Gottingen, Germany.

[4] XT, G.M. Sheldrick, *Acta Cryst.* (2015). A71, 3-8

[5] XL, Sheldrick, G. M. (2008). *Acta Cryst.* A64, 112-122.

[6] Dolomanov, O.V.; Bourhis, L.J.; Gildea, R.J.; Howard, J.A.K.; Puschmann, H., OLEX2: A complete structure solution, refinement and analysis program (2009). *J. Appl. Cryst.*, 42, 339-341

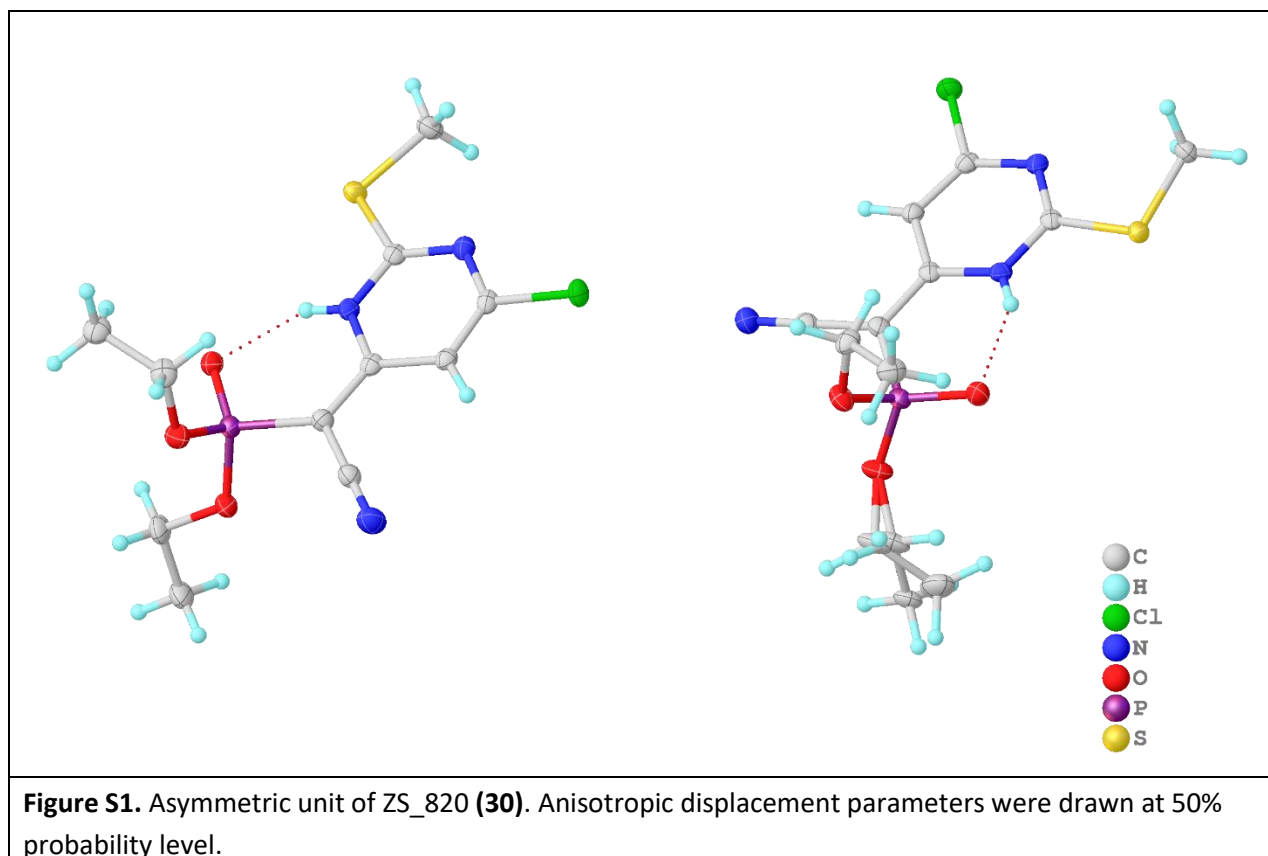


Table S1. Crystal data and structure refinement for ZS_820 (**30**).

Identification code	ZS_820
Empirical formula	C ₁₁ H ₁₅ ClN ₃ O ₃ PS
Formula weight	335.74
Temperature/K	100.0
Crystal system	triclinic
Space group	P-1
a/Å	10.5147(4)
b/Å	12.3363(4)
c/Å	12.9255(5)
α/°	63.702(2)
β/°	83.475(2)
γ/°	83.211(2)
Volume/Å ³	1489.00(10)
Z	4
ρ _{calc} /cm ³	1.498
μ/mm ⁻¹	4.706
F(000)	696.0
Crystal size/mm ³	0.67 × 0.08 × 0.01
Radiation	CuKα (λ = 1.54178)
2θ range for data collection/°	7.648 to 158.31
Index ranges	-13 ≤ h ≤ 13, -15 ≤ k ≤ 15, -16 ≤ l ≤ 16
Reflections collected	27051
Independent reflections	6068 [R _{int} = 0.0640, R _{sigma} = 0.0516]
Data/restraints/parameters	6068/33/389
Goodness-of-fit on F ²	1.038
Final R indexes [I ≥ 2σ (I)]	R ₁ = 0.0404, wR ₂ = 0.1016
Final R indexes [all data]	R ₁ = 0.0492, wR ₂ = 0.1086
Largest diff. peak/hole / e Å ⁻³	0.51/-0.39

Single-Crystal X-Ray Crystallography for Compound 50

X-ray diffraction data were measured on Bruker D8 Venture PHOTON II CPAD diffractometer equipped with a Cu K α INCOATEC ImuS micro-focus source ($\lambda = 1.54178 \text{ \AA}$). Indexing was performed using APEX3 [1] (Difference Vectors method). Data integration and reduction were performed using SaintPlus [2]. Absorption correction was performed by multi-scan method implemented in TWINABS [3]. Space group was determined using XPREP implemented in APEX3 [1]. Structure was solved using SHELXT [4] and refined using SHELXL-2018/3 [5] (full-matrix least-squares on F²) through OLEX2 interface program [6]. Ellipsoid plot was drawn with Platon [7]. Crystal was a twin. Data were integrated with Bruker-Saint using two orientation matrices determined in APEX3/RLATT based on two manually selected reciprocal lattices. Twin law from Saint: -1.00005 0.00008 0.02110 / -0.00011 -1.00040 0.08615 / 0.00404 -0.01146 1.00045. Reflections were scaled and merged with TWINABS. Detwinned data were used for refinement. Data and refinement conditions are shown in Table 1.

[1] Bruker (2019). APEX3. Bruker AXS LLC, Madison, Wisconsin, USA.

[2] Bruker (2019) SAINT. Bruker AXS LLC, Madison, Wisconsin, USA.

[3] Krause, L., Herbst-Irmer, R., Sheldrick, G. M., Stalke, D. (2015). "Comparison of silver and molybdenum microfocus X-ray sources for single-crystal structure determination" J. Appl. Cryst. 48, 3-10.

[4] Sheldrick, G. M. (2015). "SHELXT - Integrated space-group and crystal-structure determination", Acta Cryst. A71, 3-8.

[5] Sheldrick, G. M. (2015) "Crystal structure refinement with SHELXL", Acta Cryst., C71, 3-8

[6] Dolomanov, O.V.; Bourhis, L.J.; Gildea, R.J.; Howard, J.A.K.; Puschmann, H., OLEX2: A complete structure solution, refinement and analysis program (2009). J. Appl. Cryst., 42, 339-341

[7] Spek, A. L. (2009). "Structure validation in chemical crystallography", Acta Cryst. D65, 148-155.

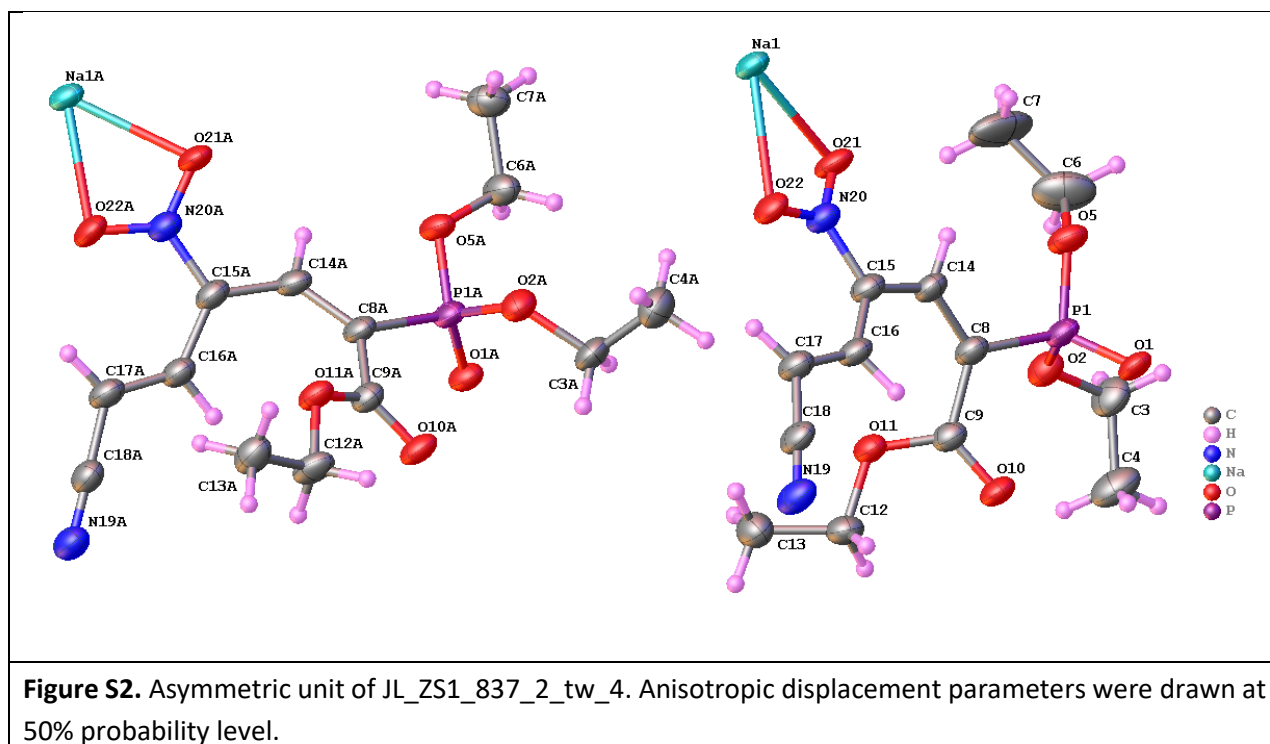


Table S2. Crystal data and structure refinement for JL_ZS1_837_2_tw_4 (50).

Identification code	JL_ZS1_837_2_tw_4
Empirical formula	C ₁₃ H ₁₈ N ₂ NaO ₇ P
Formula weight	368.25
Temperature/K	100.0
Crystal system	triclinic
Space group	P-1
a/Å	9.3192(3)
b/Å	11.1085(4)
c/Å	18.3315(6)
α/°	85.750(2)
β/°	88.791(2)
γ/°	66.538(2)
Volume/Å ³	1735.96(10)
Z	4
ρ _{calc} /cm ³	1.409
μ/mm ⁻¹	1.999
F(000)	768.0
Crystal size/mm ³	0.33 × 0.13 × 0.03
Radiation	CuKα (λ = 1.54178)
2θ range for data collection/°	4.834 to 133.788
Index ranges	-11 ≤ h ≤ 11, -13 ≤ k ≤ 13, 0 ≤ l ≤ 21
Reflections collected	6147
Independent reflections	6147 [R _{int} = 0.0946, R _{sigma} = 0.0622]
Data/restraints/parameters	6147/3/439
Goodness-of-fit on F ²	1.044
Final R indexes [I ≥ 2σ (I)]	R ₁ = 0.0752, wR ₂ = 0.2218
Final R indexes [all data]	R ₁ = 0.0887, wR ₂ = 0.2288