

Supporting Information

Copper-Catalyzed Domino Addition, Hydroamination and Cyclization: A Multicomponent Approach to Spiro Oxazolidinone Derivatives

Gal Reddy Potuganti, ^{†‡} Divakar Reddy Indukuri, ^{†‡} Jagadeesh Babu Nanubolu[†] Manjula Alla.^{*†‡}

[†] Fluoro-Agrochemicals Division, CSIR-Indian Institute of Chemical Technology, Tarnaka, Hyderabad, 500007, India.

[†] Centre for X-ray Crystallography, CSIR-Indian Institute of Chemical Technology, Tarnaka, Hyderabad 500 007, India.

[‡] Academy of Scientific and Innovative Research, CSIR- Indian Institute of Chemical Technology, Tarnaka, Hyderabad 500007, India.

Email: manjula@iict.res.in

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1. Crystal structure analysis of compound 4e (#CCDC 1863694)

Table S1 Crystal data and structure refinement for compound 4e.

Identification code	KA409MF
Empirical formula	C ₂₄ H ₁₇ ClN ₂ O ₃
Formula weight	416.84
Temperature/K	293(2)
Crystal system	monoclinic
Space group	P2 ₁ /c
a/Å	6.16338(8)
b/Å	33.2578(5)
c/Å	9.94383(14)
α /°	90
β /°	100.2051(8)
γ /°	90
Volume/Å ³	2006.05(5)
Z	4
$\rho_{\text{calc}}/\text{cm}^3$	1.380
μ/mm^{-1}	0.220
F(000)	864.0
Crystal size/mm ³	0.300 × 0.210 × 0.150
Radiation	MoK α (λ = 0.71073)
2 Θ range for data collection/°	4.338 to 55
Index ranges	-8 ≤ h ≤ 8, -43 ≤ k ≤ 43, -12 ≤ l ≤ 12
Reflections collected	19219
Independent reflections	4589 [R_{int} = 0.0370, R_{sigma} = 0.0323]
Data/restraints/parameters	4589/6/270
Goodness-of-fit on F ²	1.128
Final R indexes [$I \geq 2\sigma(I)$]	R_1 = 0.0506, wR_2 = 0.1467
Final R indexes [all data]	R_1 = 0.0697, wR_2 = 0.1603

Largest diff. peak/hole / e Å⁻³ 0.26/-0.43

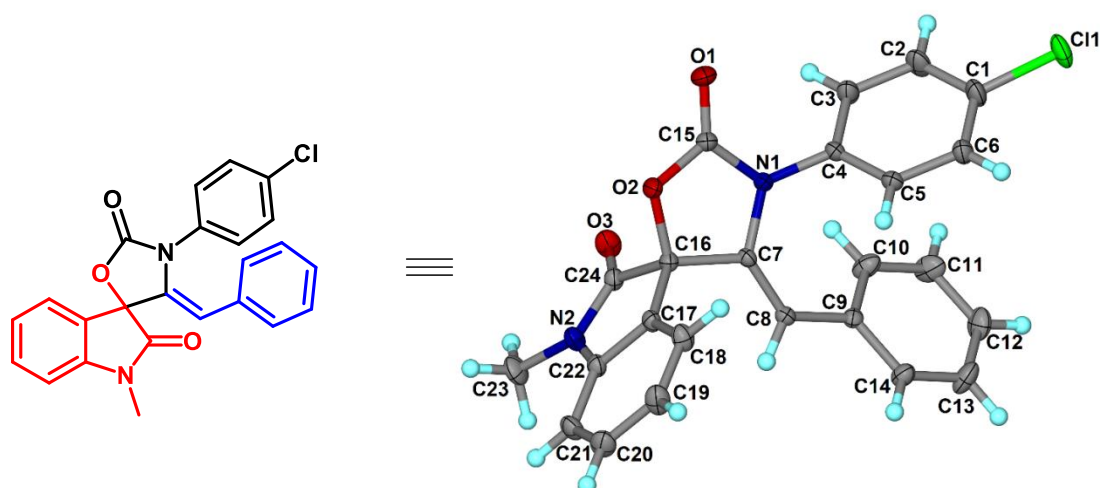


Figure S1. X-ray crystal structure of compound 4e

Figure S1: X-ray crystal structure of compound 4e with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radius. Only major component of the disordered atoms is shown for clarity.

3. Crystal structure analysis of compound 6f (#CCDC 1863695)

Table S2 Crystal data and structure refinement for compound 6f.

Identification code	KA471MF
Empirical formula	C ₂₁ H ₂₀ ClNO ₂
Formula weight	353.83
Temperature/K	293(2)
Crystal system	monoclinic
Space group	C2/c
a/Å	27.583(2)
b/Å	13.6001(10)
c/Å	10.2013(10)
α/°	90
β/°	104.589(4)
γ/°	90

Volume/Å ³	3703.5(5)
Z	8
$\rho_{\text{calc}}/\text{cm}^3$	1.269
μ/mm^{-1}	0.220
F(000)	1488.0
Crystal size/mm ³	0.400 × 0.360 × 0.320
Radiation	MoK α (λ = 0.71073)
2 Θ range for data collection/°	5.016 to 49.98
Index ranges	-32 ≤ h ≤ 32, -16 ≤ k ≤ 16, -12 ≤ l ≤ 11
Reflections collected	21907
Independent reflections	3261 [R_{int} = 0.0611, R_{sigma} = 0.0393]
Data/restraints/parameters	3261/240/290
Goodness-of-fit on F ²	1.066
Final R indexes [$I \geq 2\sigma(I)$]	R_1 = 0.0550, wR_2 = 0.1412
Final R indexes [all data]	R_1 = 0.0921, wR_2 = 0.1641
Largest diff. peak/hole / e Å ⁻³	0.15/-0.28

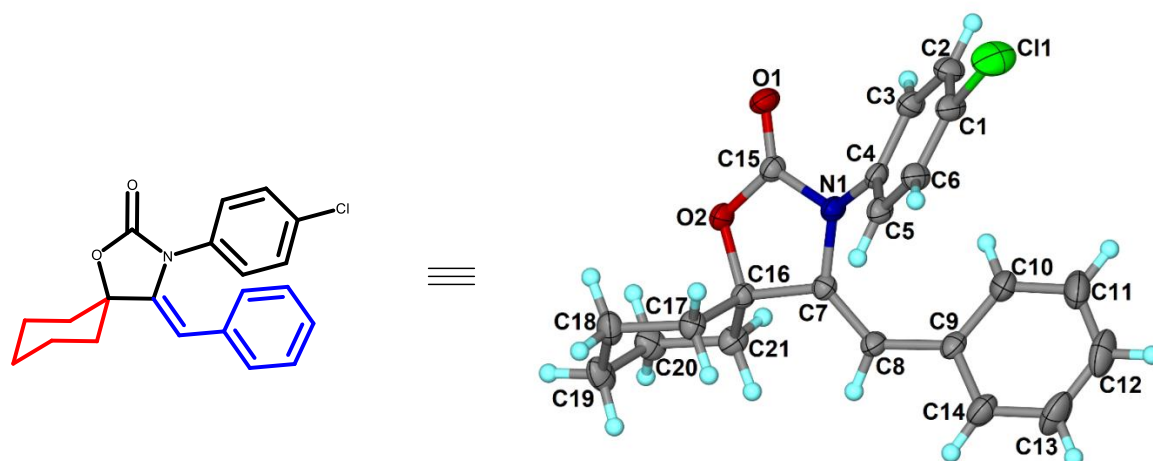


Figure S2. X-ray crystal structure of compound 6f

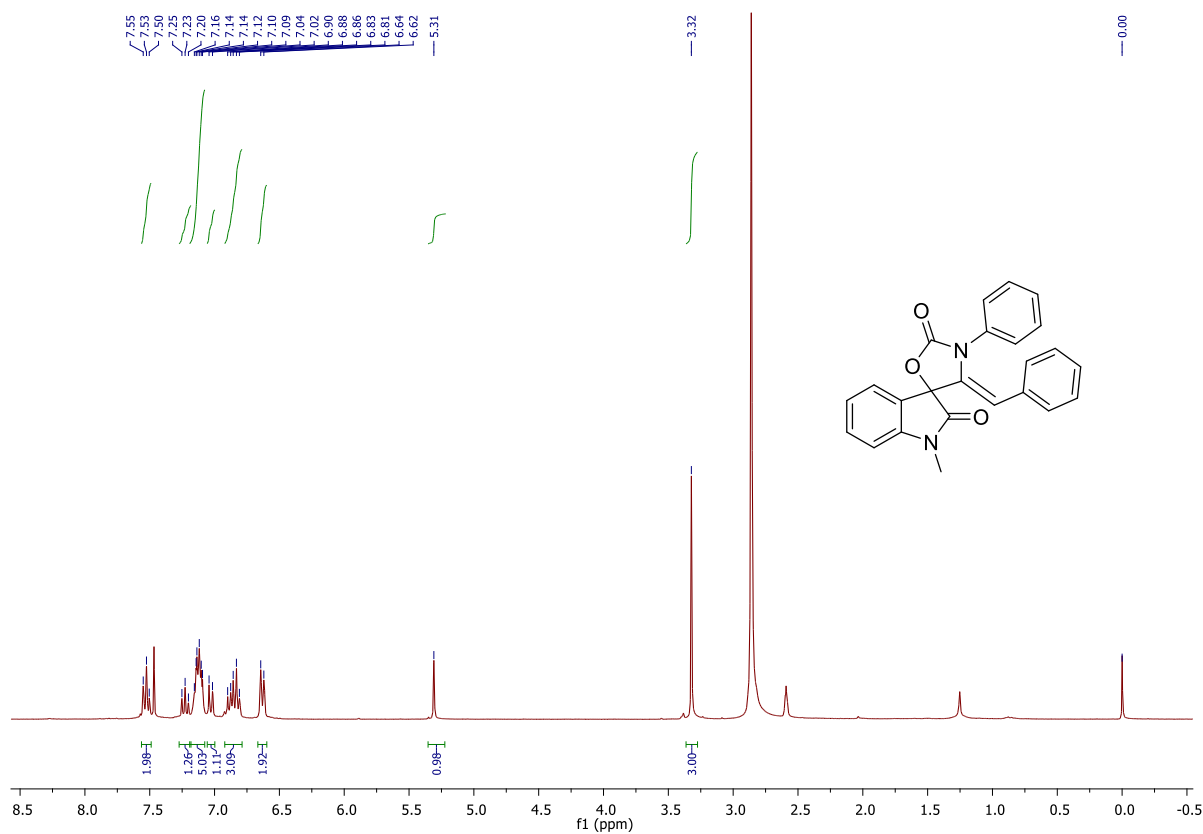
Figures S2: X-ray crystal structure of compound 6f with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radius. Only major component of the disorder atoms is shown for clarity.

#CCDC 1863694 & 1863695 contains the supplementary crystallographic data for this paper. These data can be obtained free of charge from The Cambridge Crystallographic Data Centre via <https://www.ccdc.cam.ac.uk/structures/>

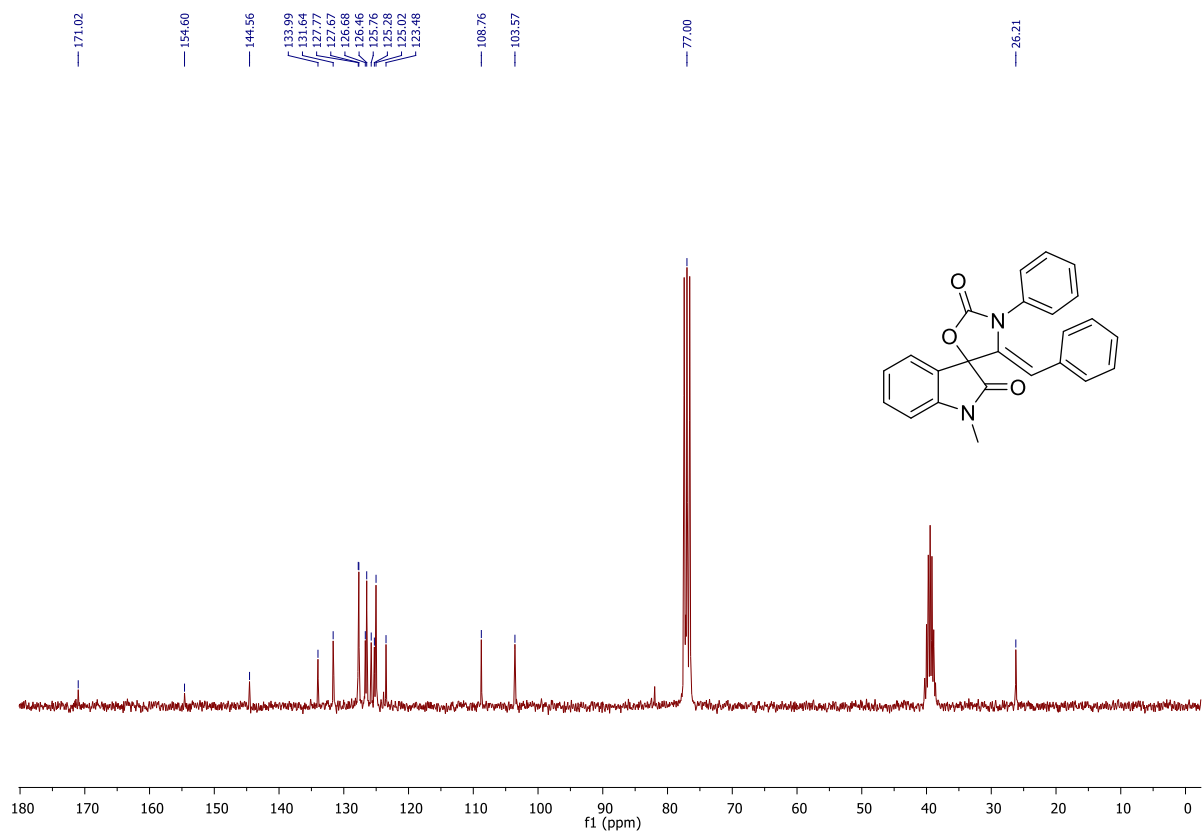
Data collection and structure solution of 4e (KA409) and 6f (KA471): Single crystal X-ray data for two compounds were collected at room temperature on a Bruker D8 QUEST equipped with a four-circle kappa diffractometer and Photon 100 detector. An I μ s microfocus Mo source ($\lambda=0.71073\text{\AA}$) supplied the multi-mirror monochromated incident beam. A combination of Phi and Omega scans were used to collect the necessary data. Unit cell dimensions were determined using 7490 reflections for KA409 & 5607 reflection for KA417 data sets. Integration and scaling of intensity data were accomplished using SAINT program.¹ The structures were solved by Direct Methods using SHELXS97² and refinement was carried out by full-matrix least-squares technique using SHELXL-2014/7.²⁻³ Anisotropic displacement parameters were included for all non-hydrogen atoms. All H atoms were positioned geometrically and treated as riding on their parent C atoms with C-H distances of 0.93–0.97 \AA , and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}$ for methyl atoms. In KA409, the chlorine atom on the phenyl ring was disordered over two sites, with the site occupancy factor of 0.57(2) for Cl1 atom (major component) & 0.43(2) for Cl1D atom (minor component). In KA416, the chlorophenyl ring was disordered over two sites, with the site occupancy factor of 0.52(2) for Cl1/C1/C2/C3/C4/C5/C6 atoms (major component) & 0.48(2) for Cl1D/C1D/C2D/C3D/C4D/C5D/C6D atoms (minor component). The anisotropic displacement parameters of the disordered carbon atoms were restrained to be similar (SIMU instruction) and the direction of motion along the axis between these atoms was also restrained (DELU instruction).³ The C-C bond distances of disordered ethyl groups were restrained to their expected values with DFIX instruction and performed the final cycle of refinement. The phenyl ring atoms were also treated as split models joining isopropyl group major and minor components respectively.

1. SMART & SAINT. Software Reference manuals. Versions 6.28a & 5.625, Bruker Analytical X-ray Systems Inc., Madison, Wisconsin, U.S.A., 2001.
2. Sheldrick, G. M. SHELXS97 and SHELXL Version 2014/7, <http://shelx.uni-ac.gwdg.de/SHELX/index.php>
3. Muller, P, Herbst-Imer, R, Spek, A. L, Schneider, T. R, and Sawaya, M. R. Crystal Structure Refinement: A Crystallographer's Guide to SHELXL. Muller, P. Ed. 2006 Oxford University Press: Oxford, New York, pp. 57–91

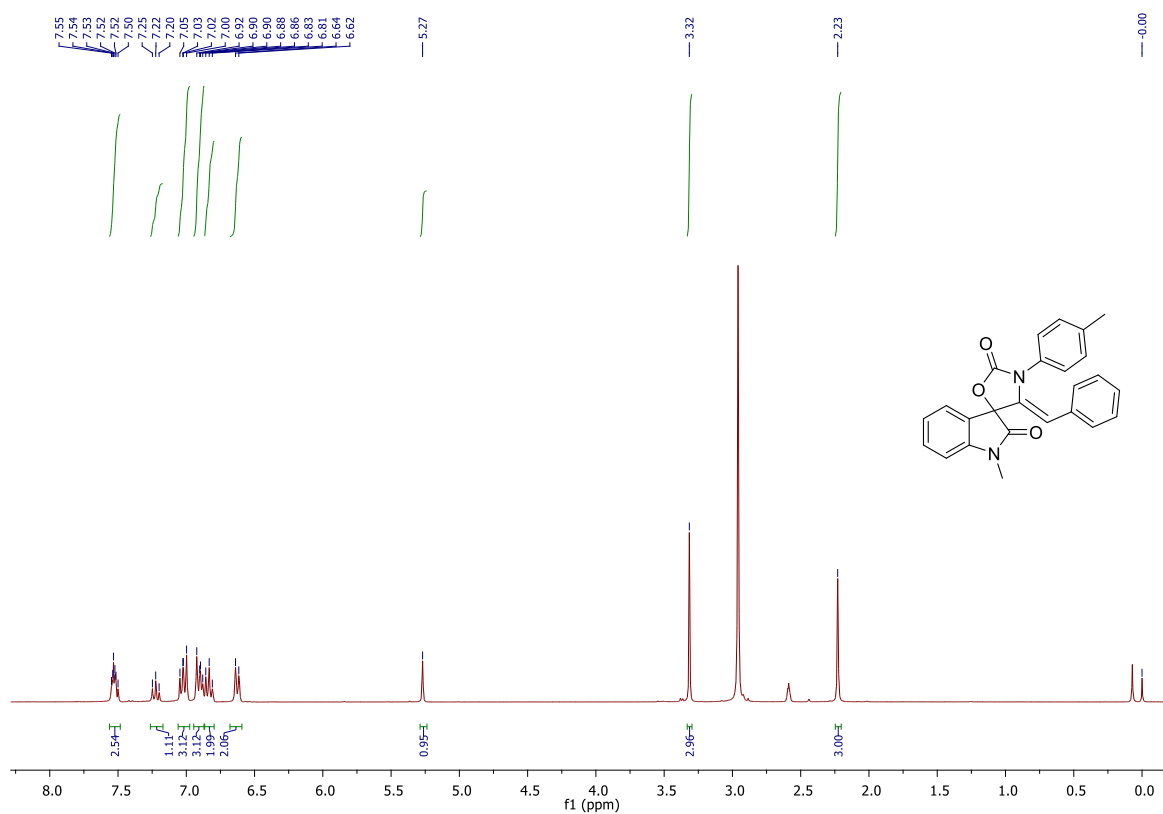
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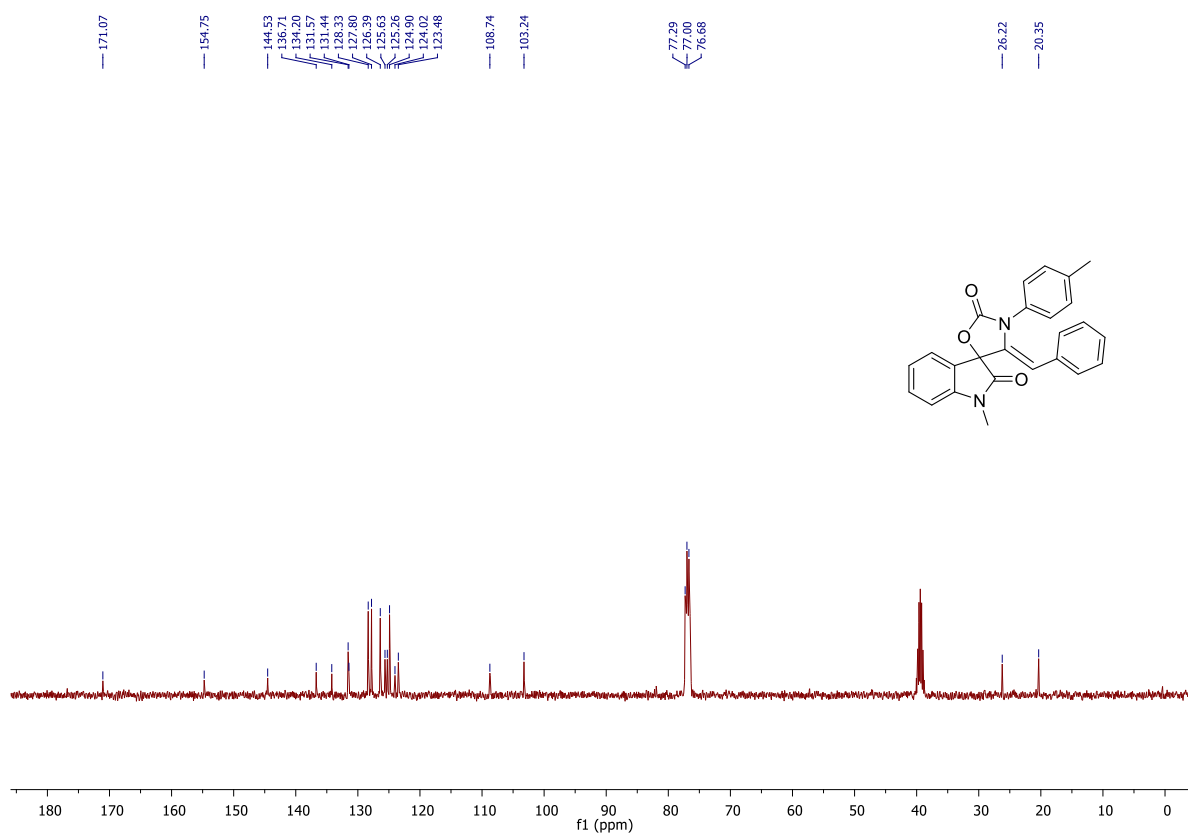
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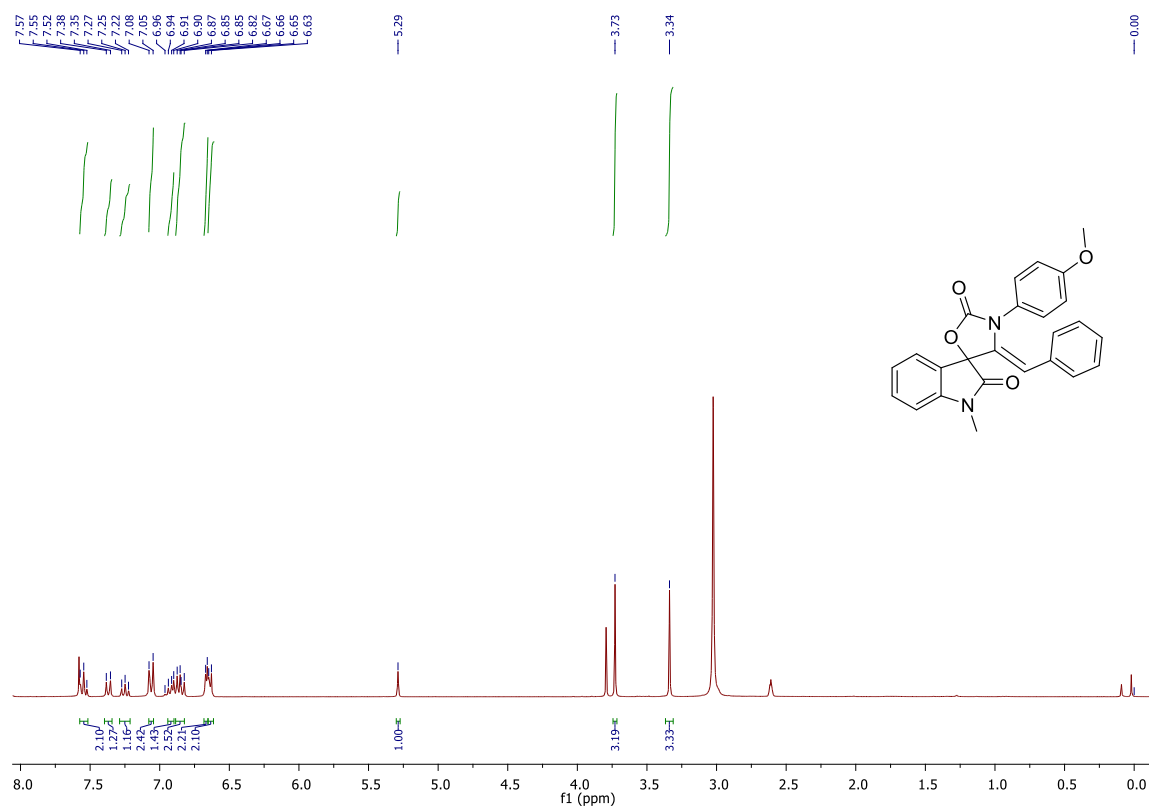
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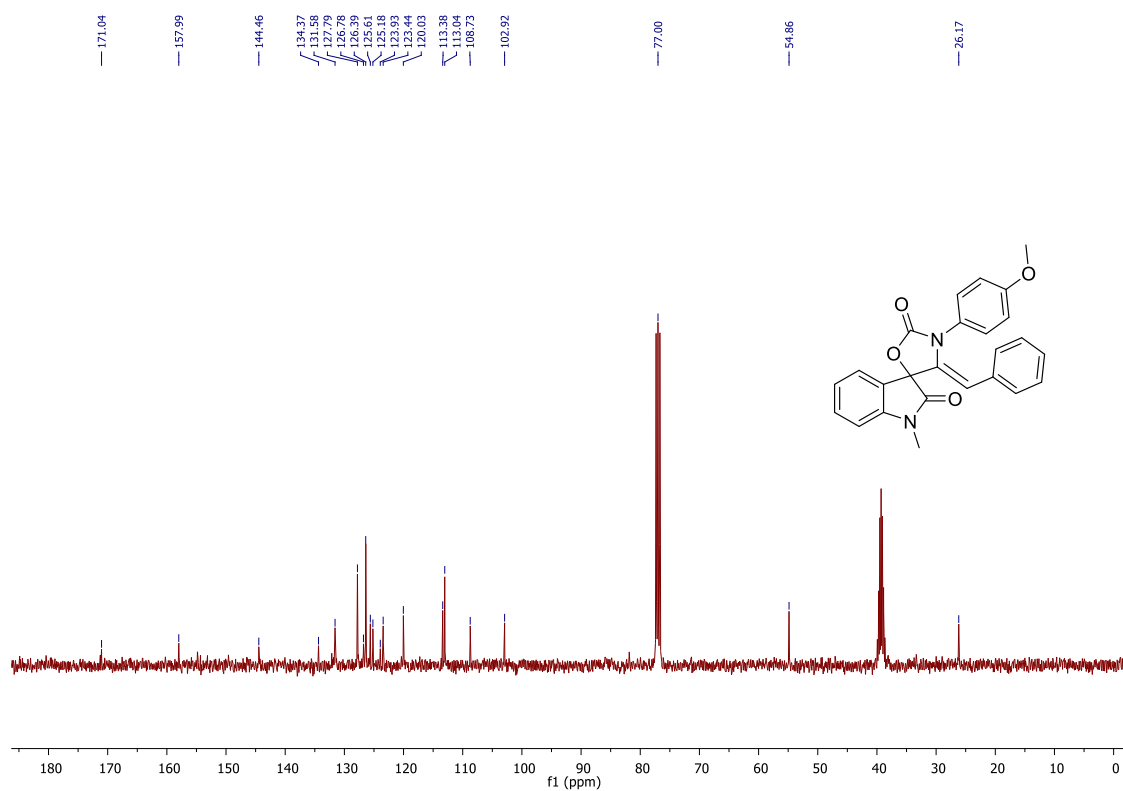
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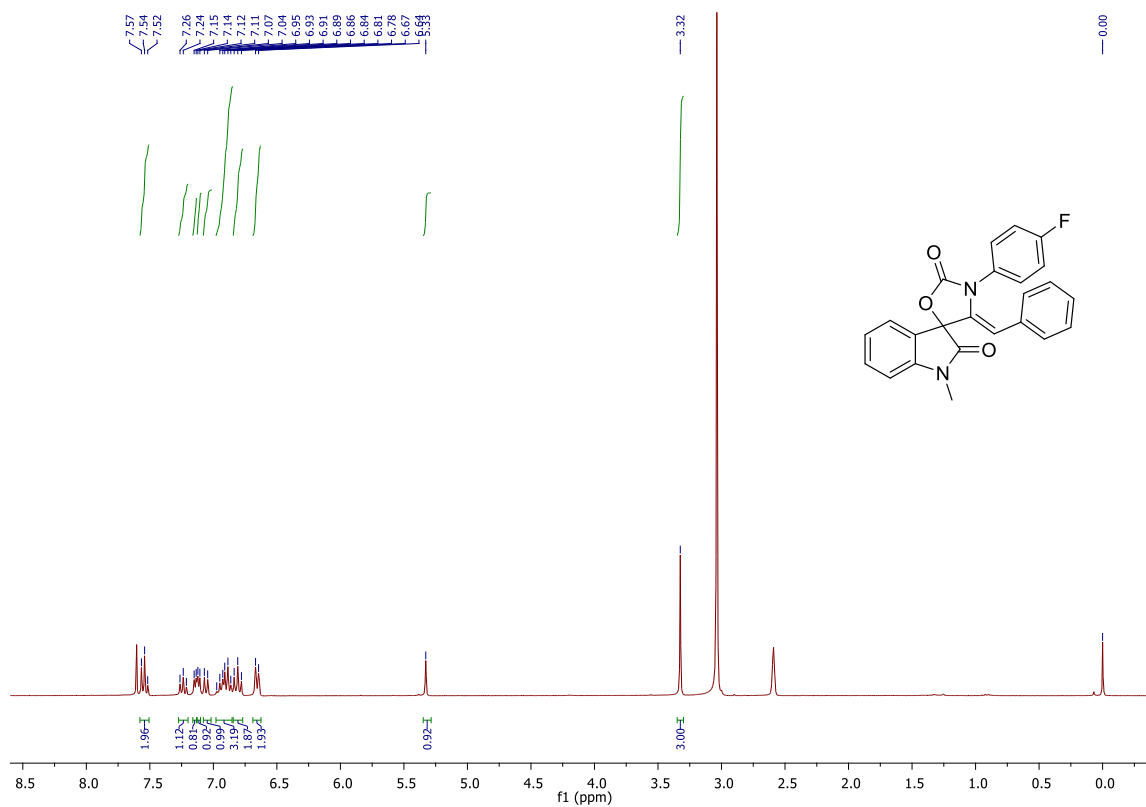
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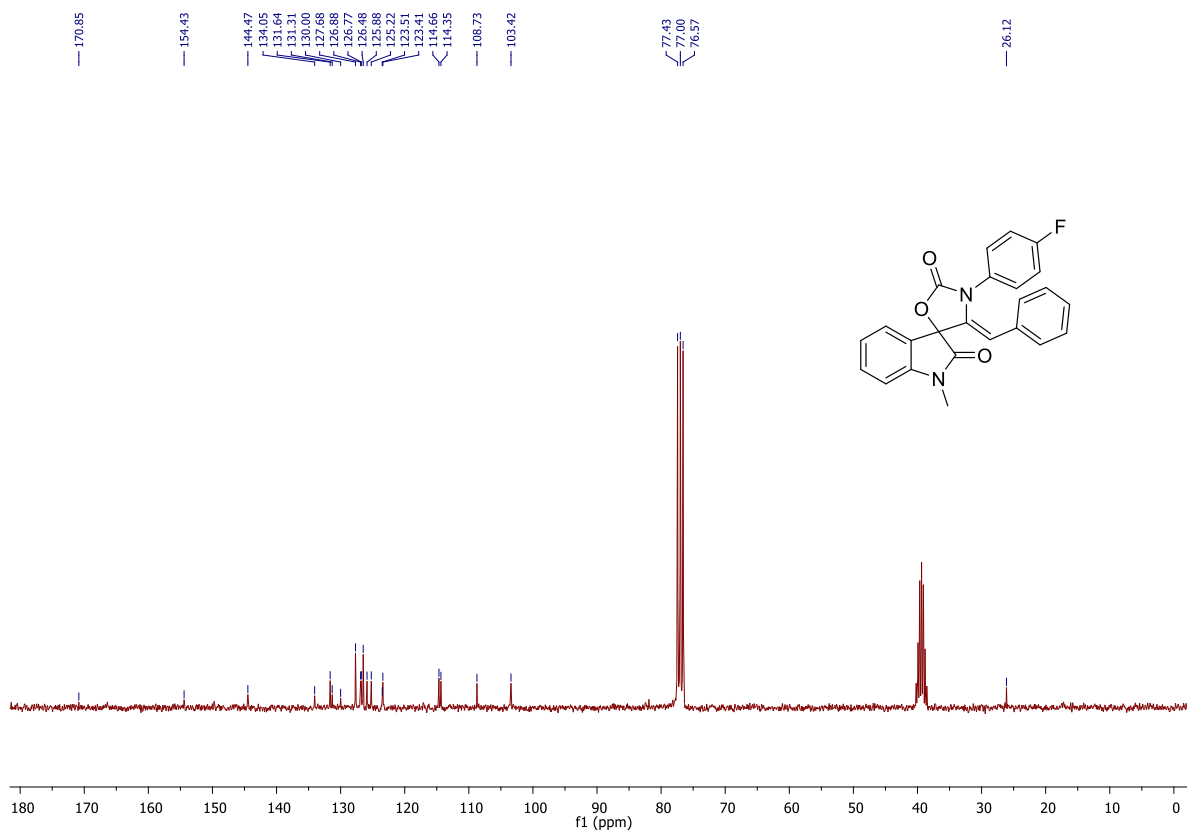
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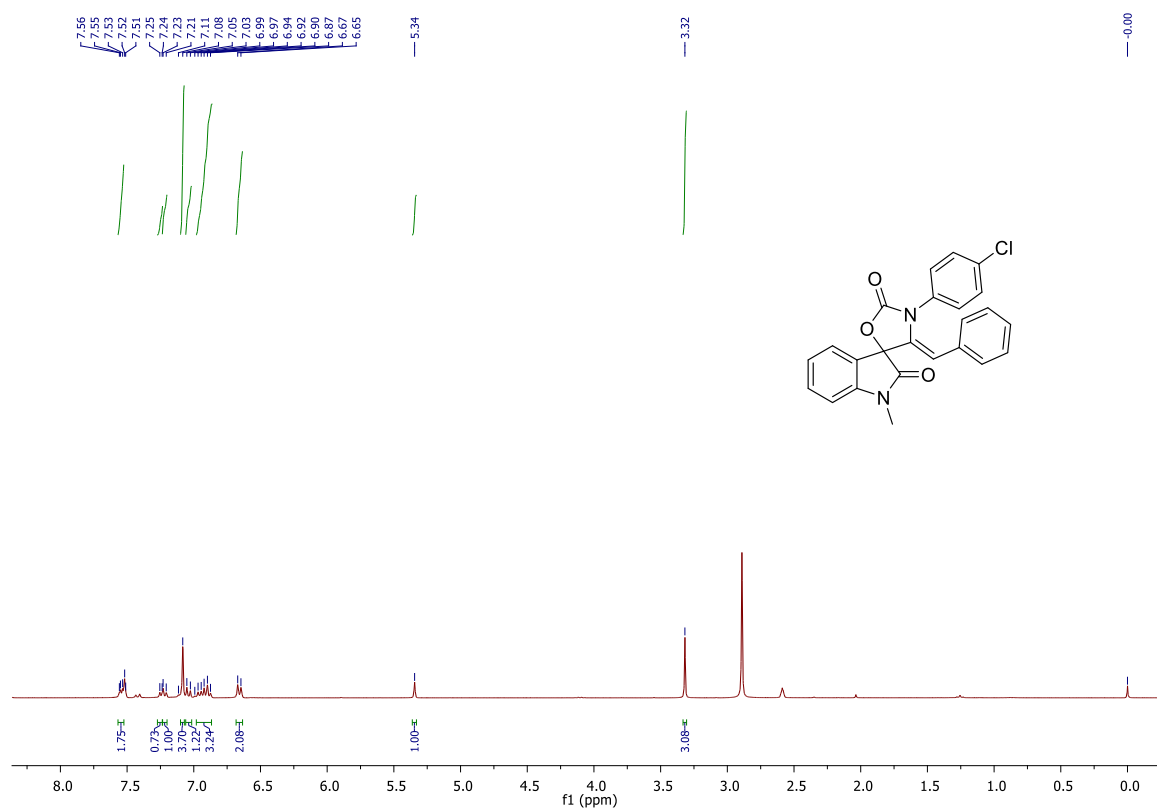
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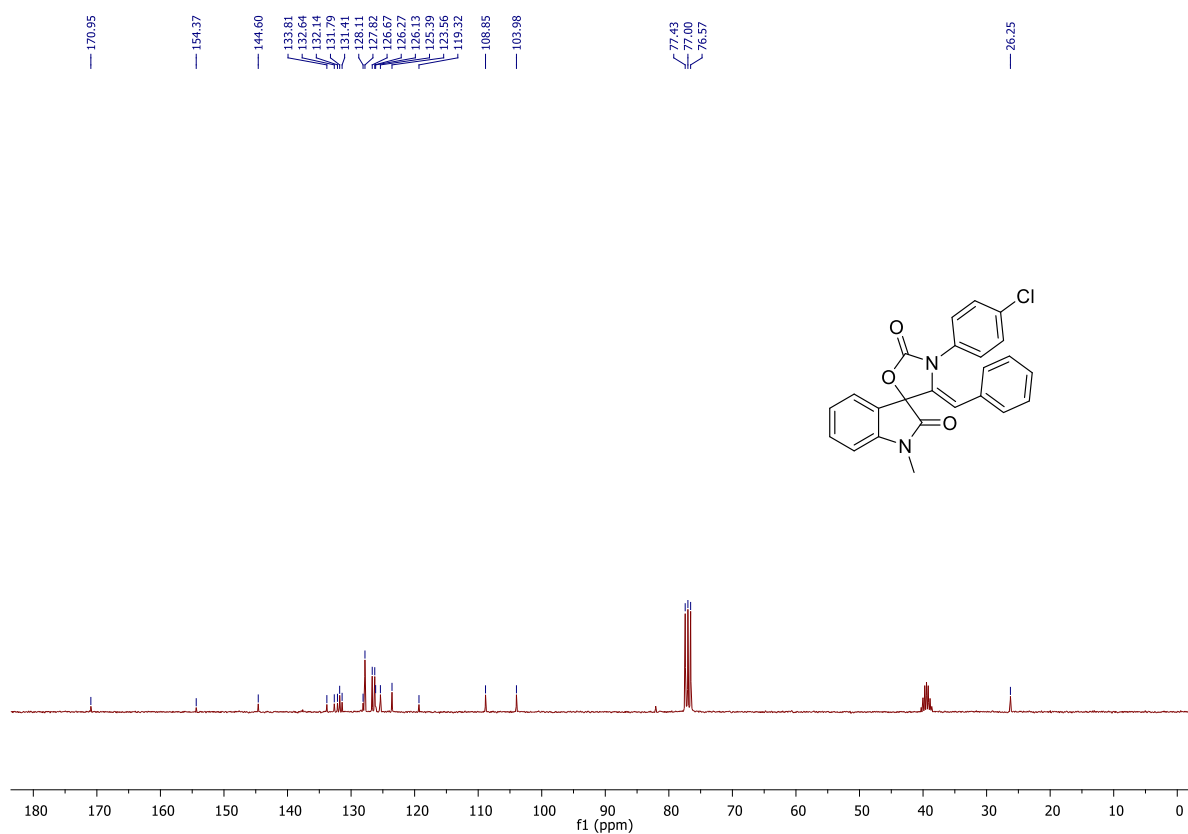
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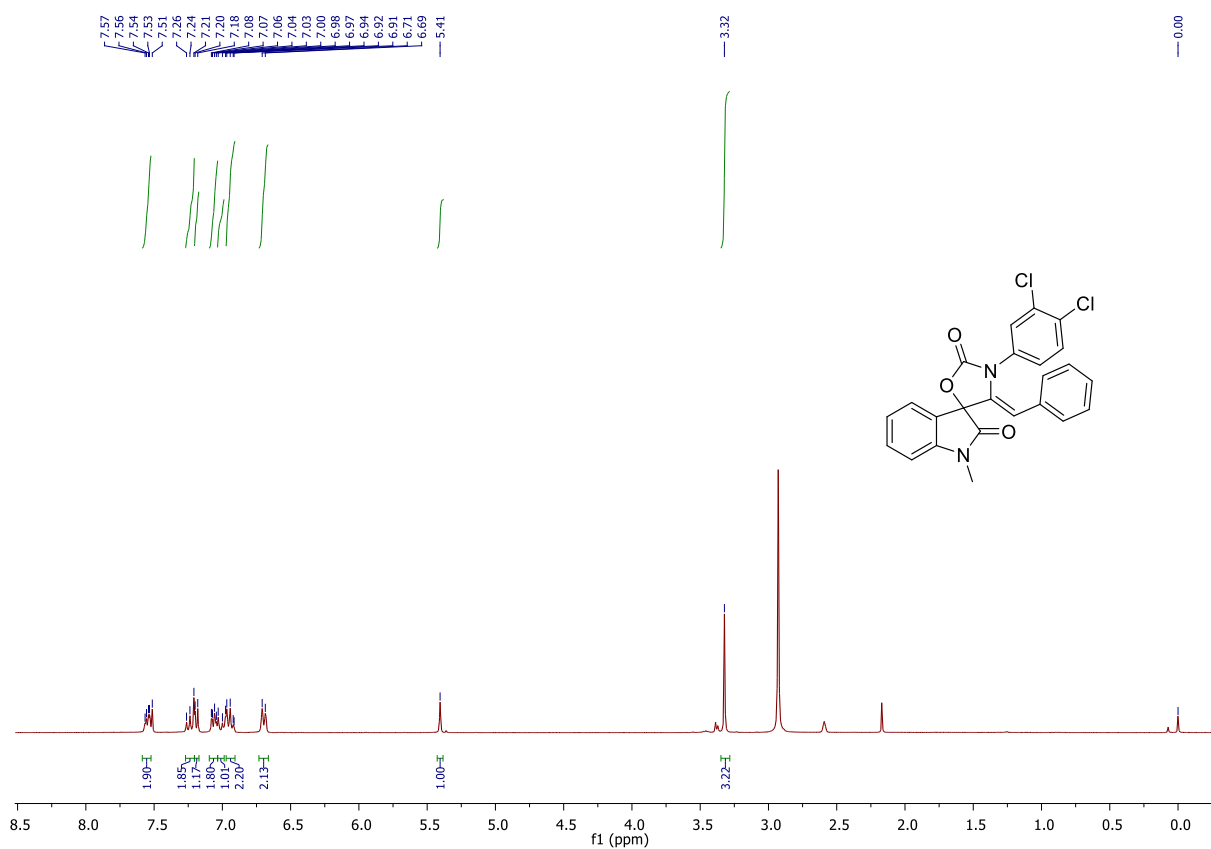
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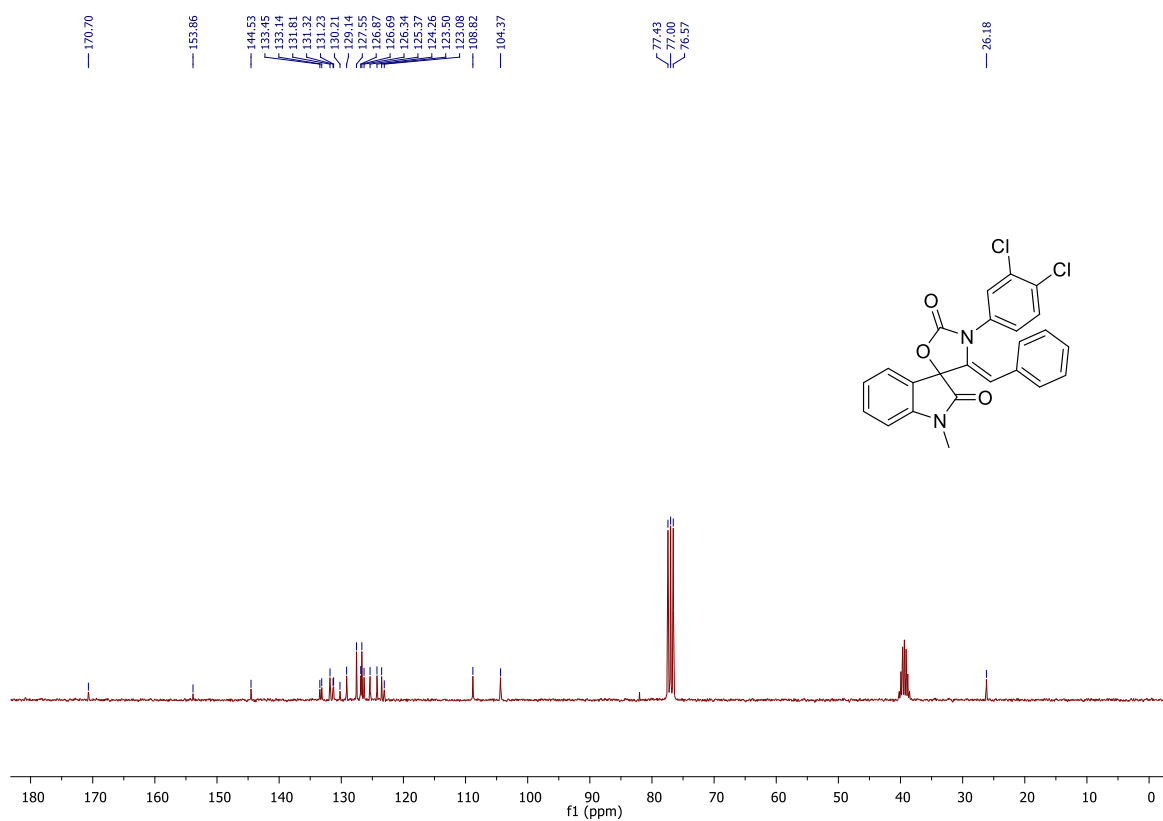
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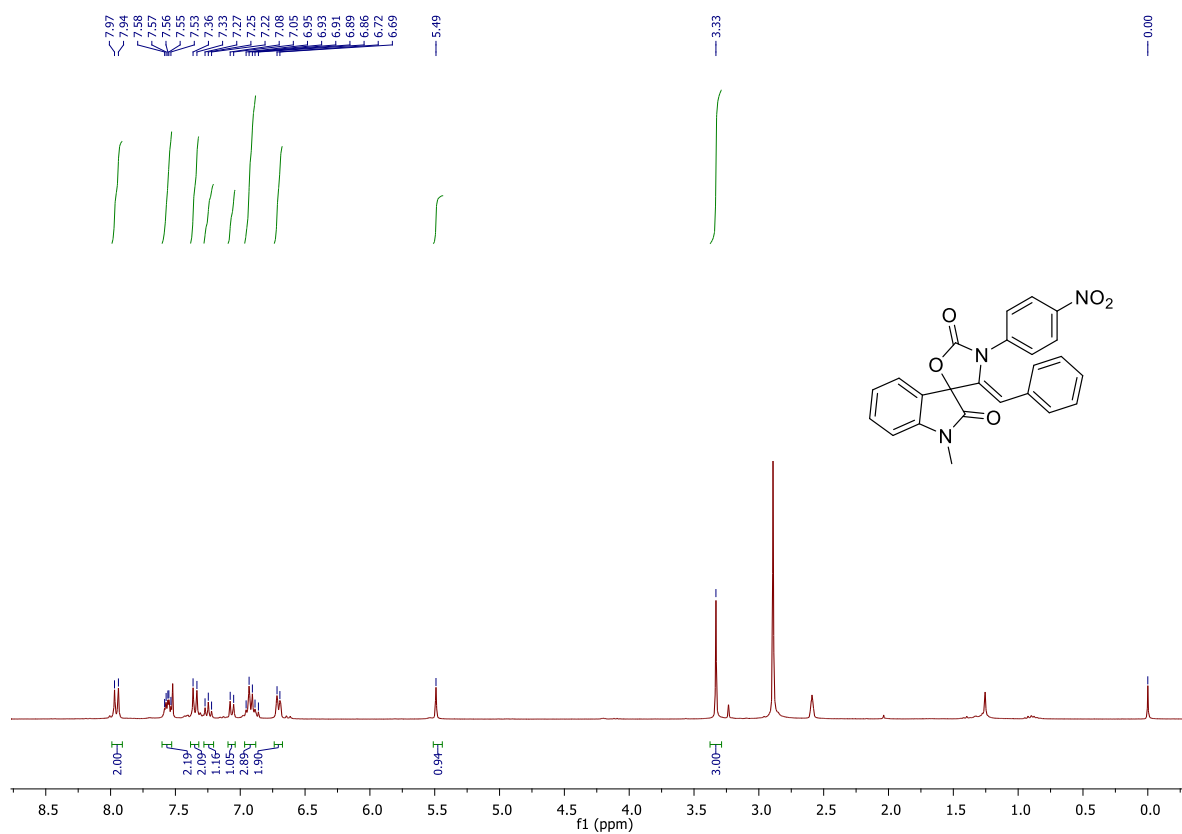
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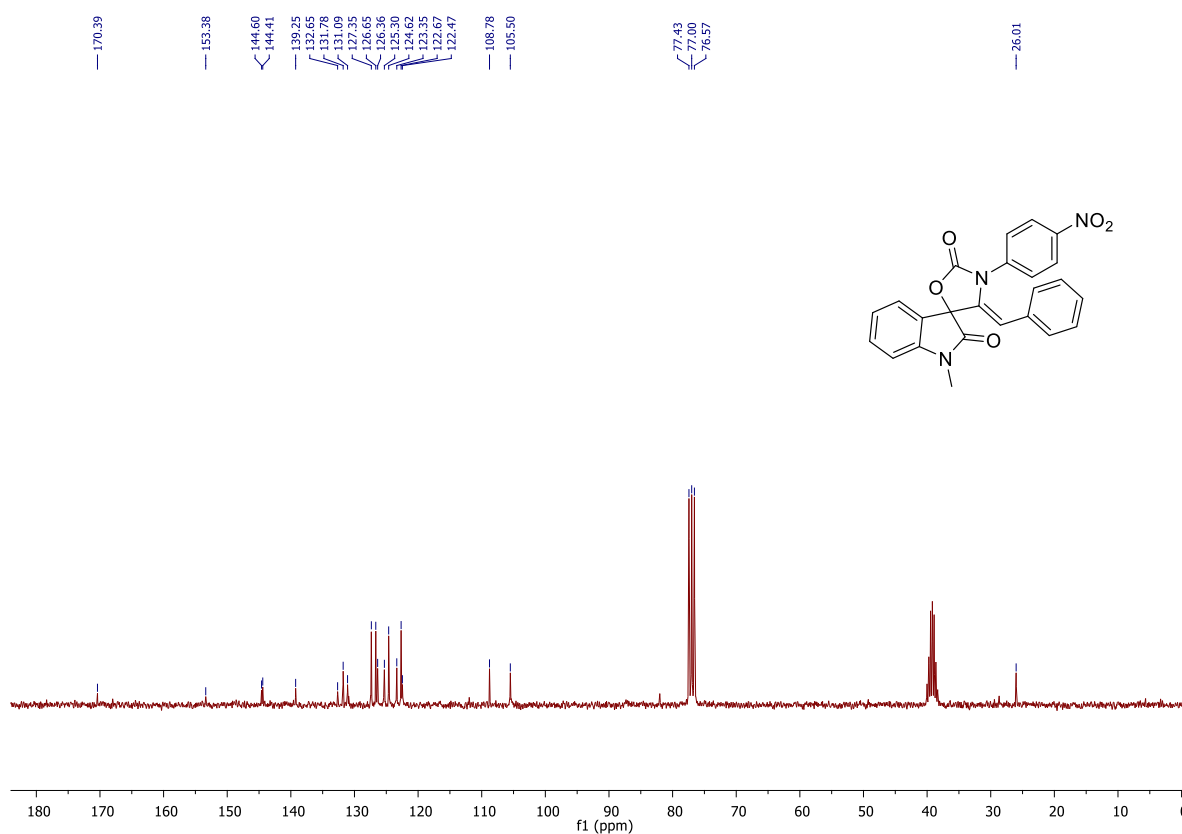
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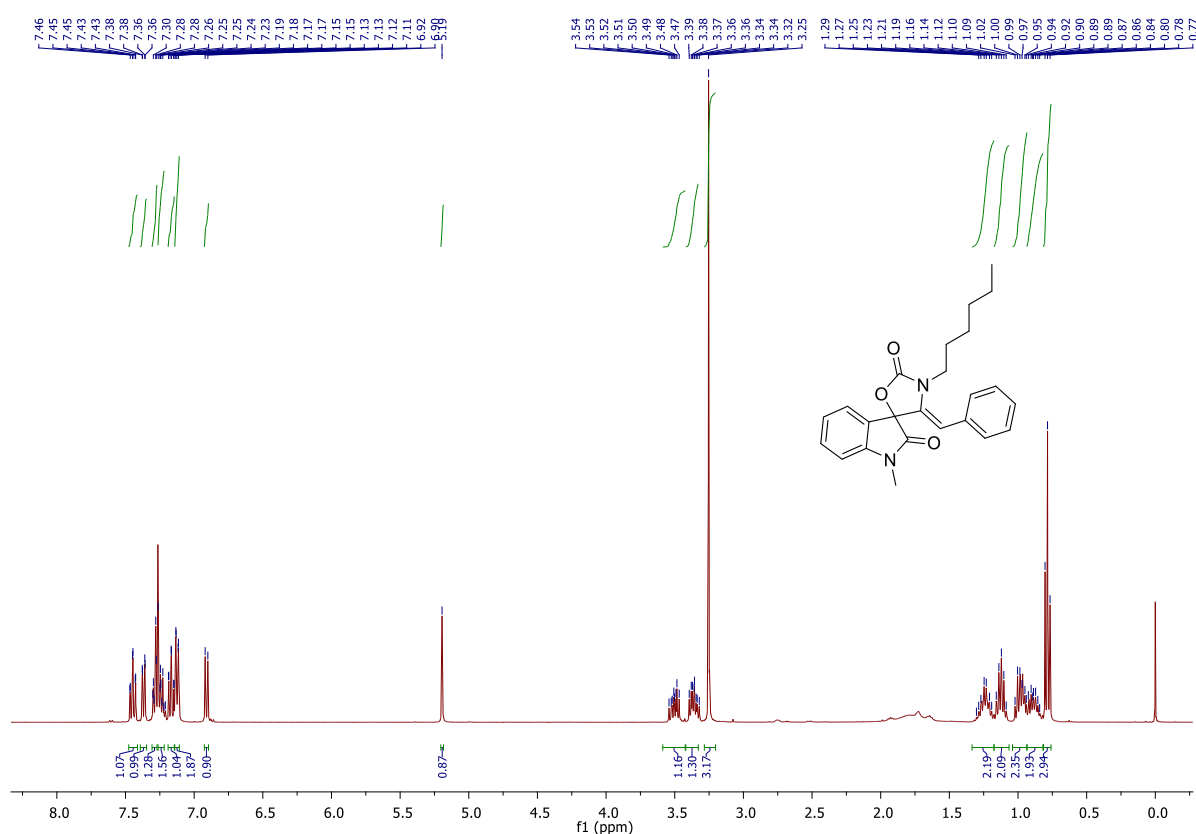
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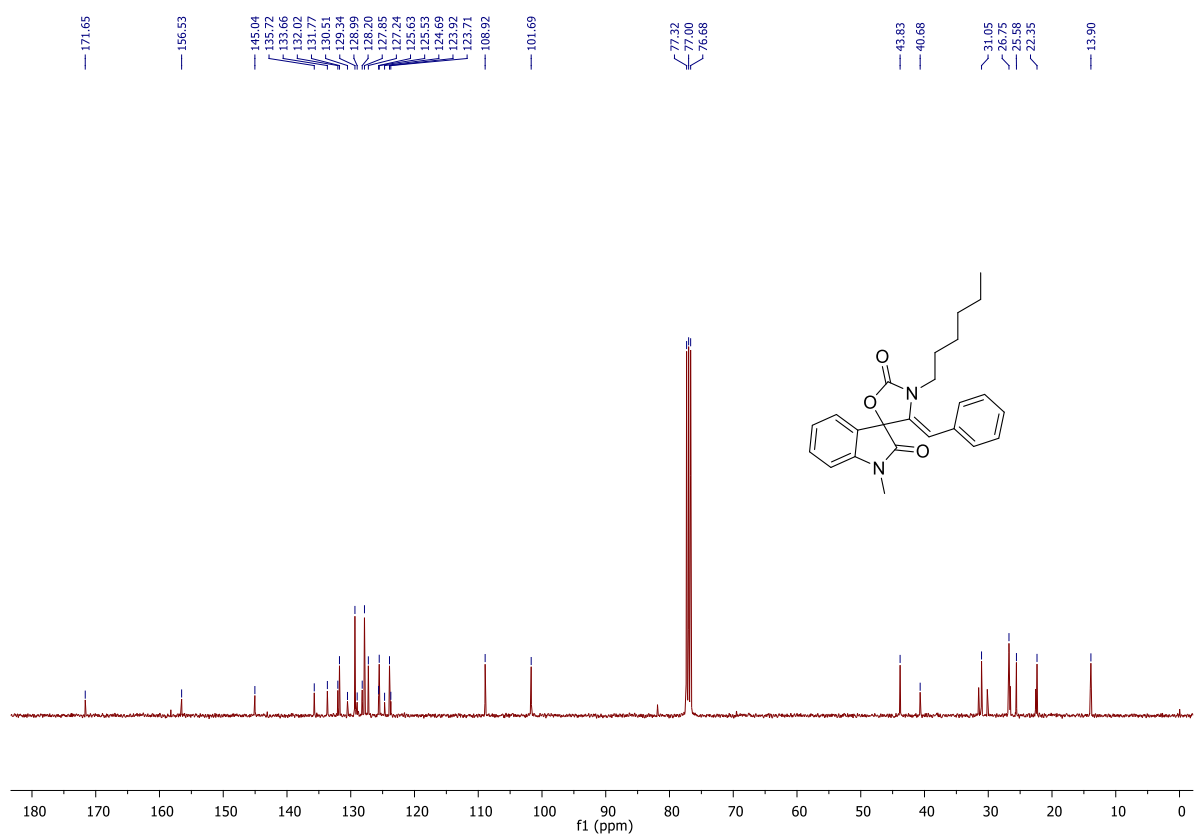
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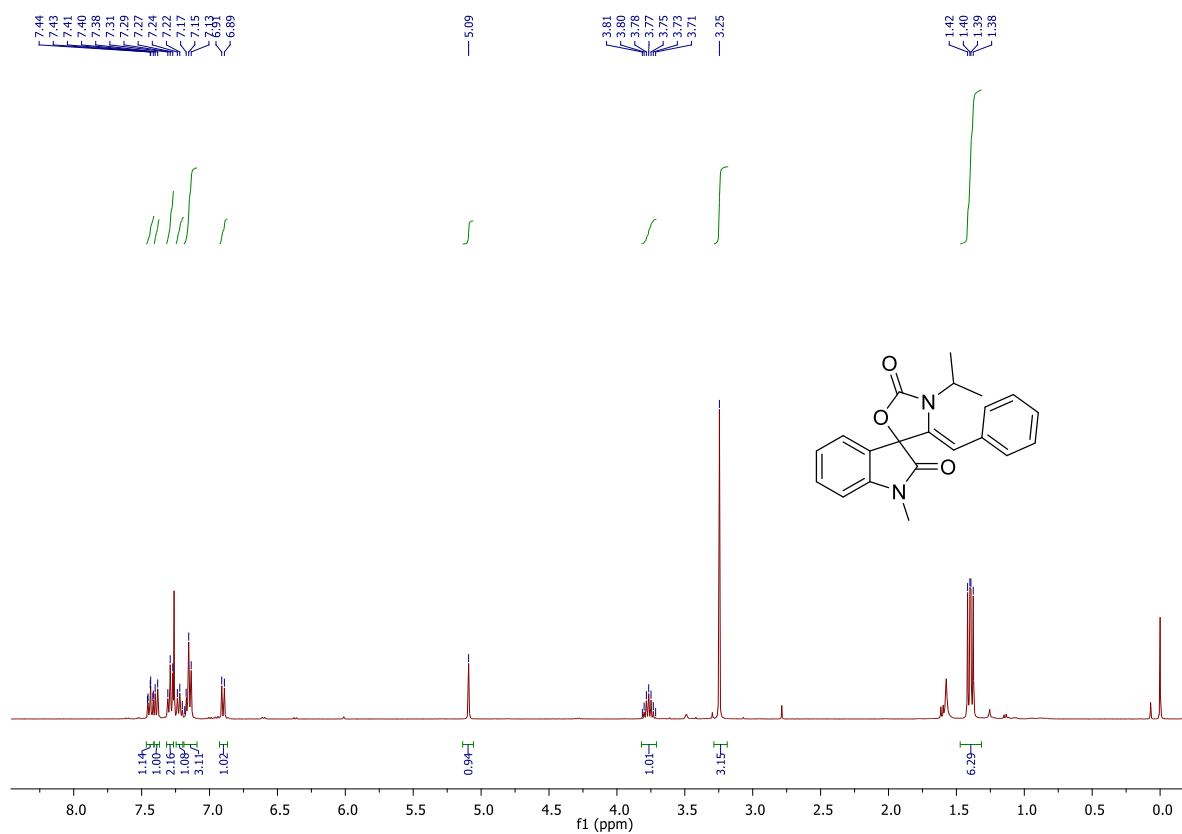
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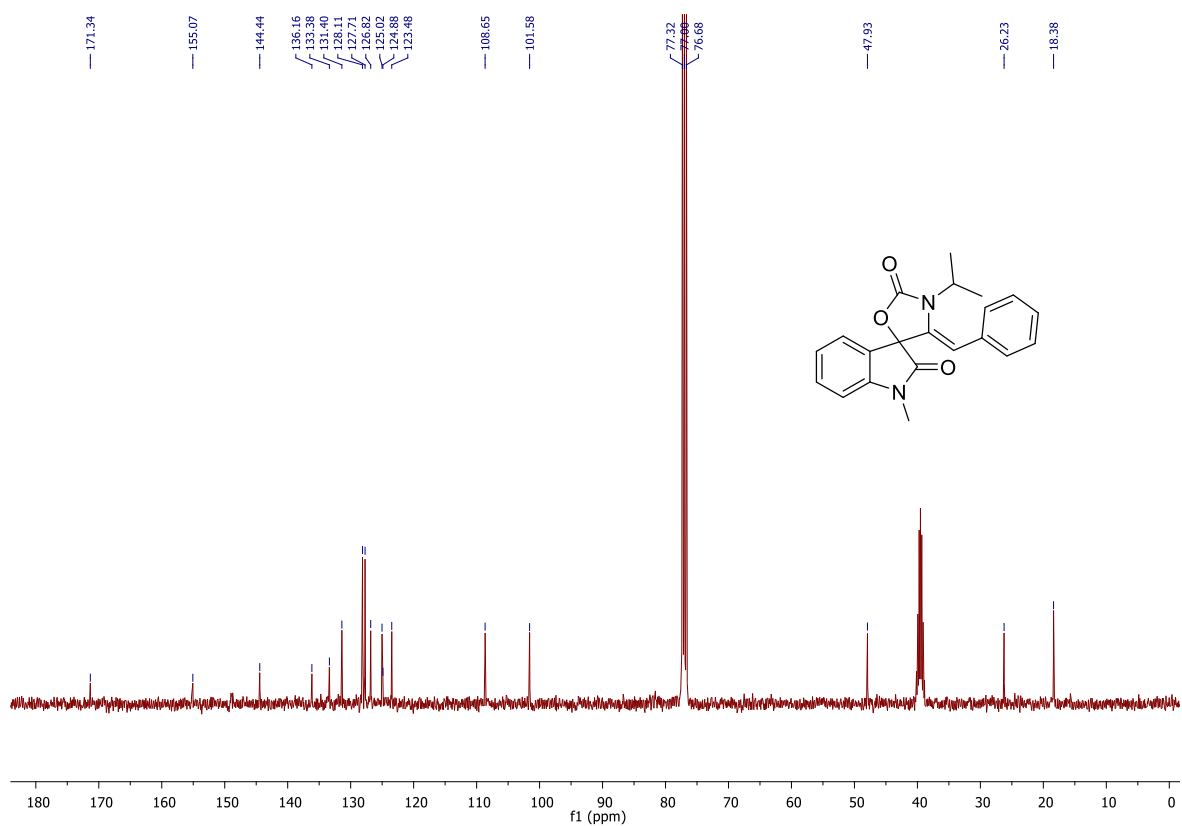
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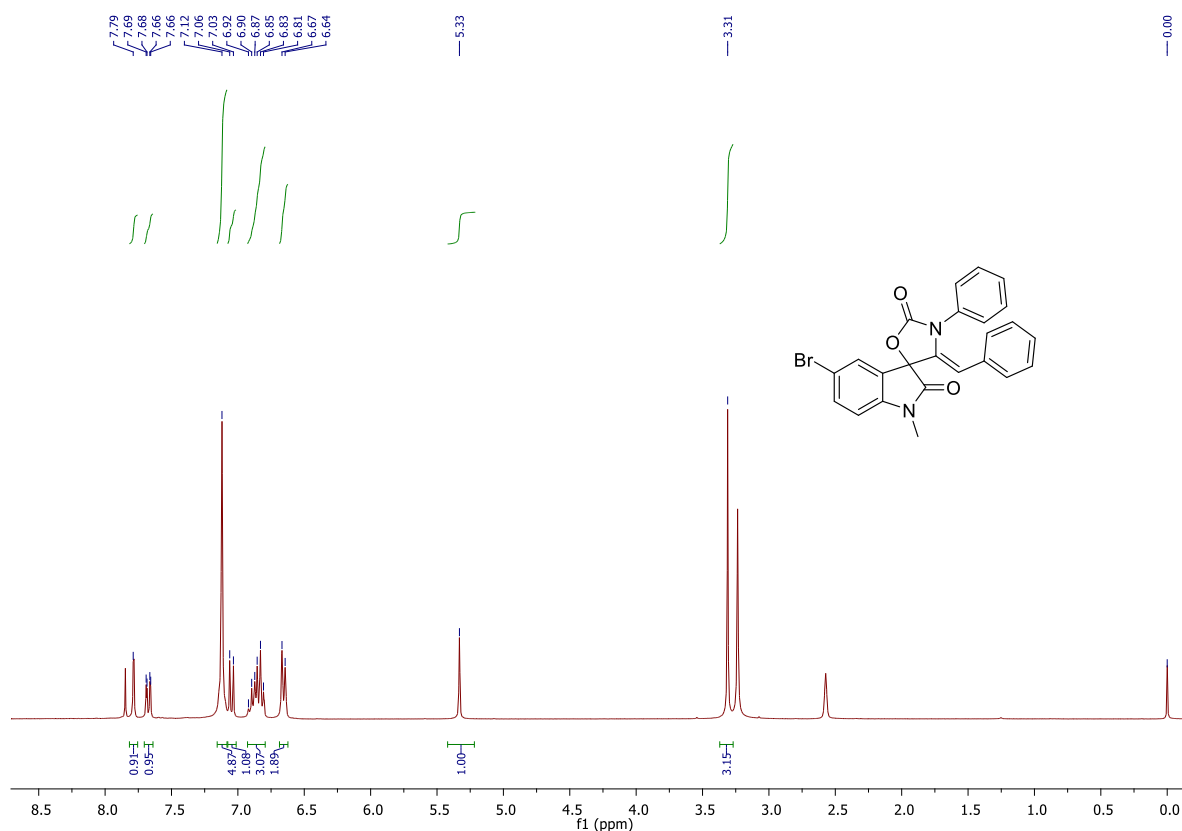
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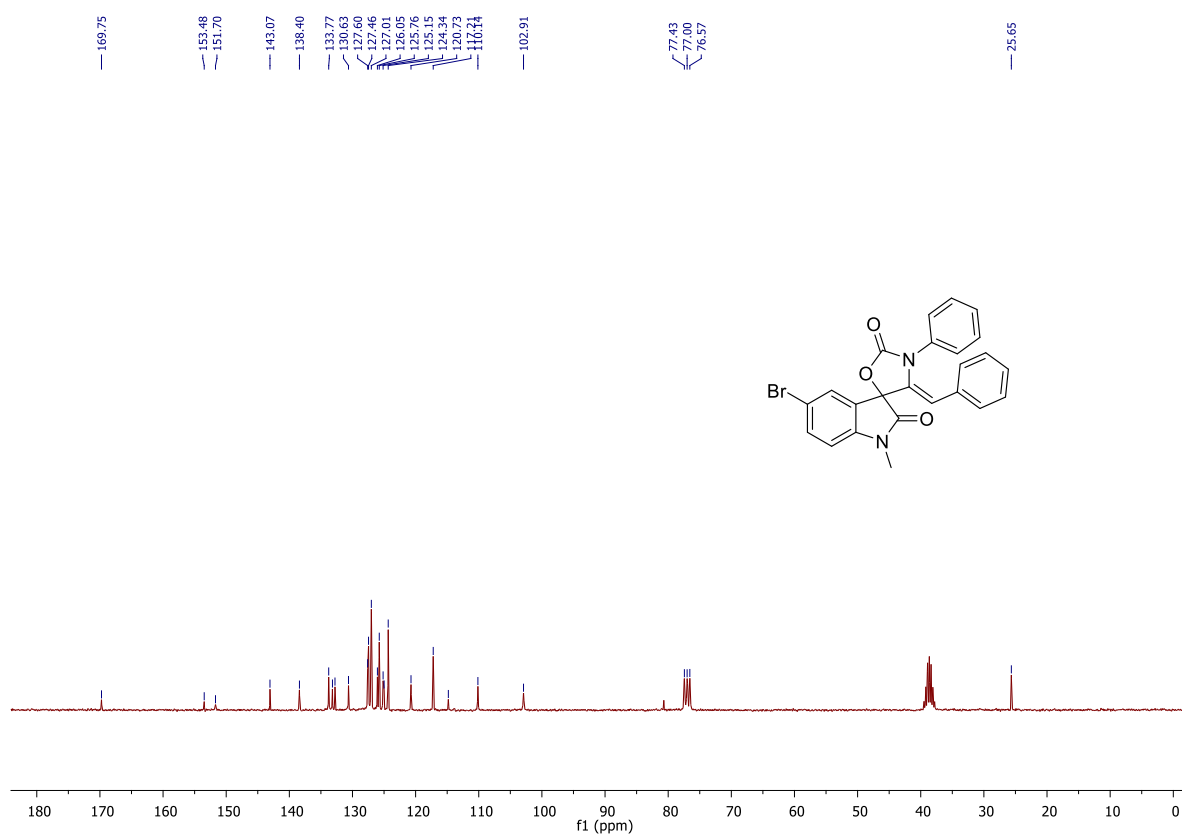
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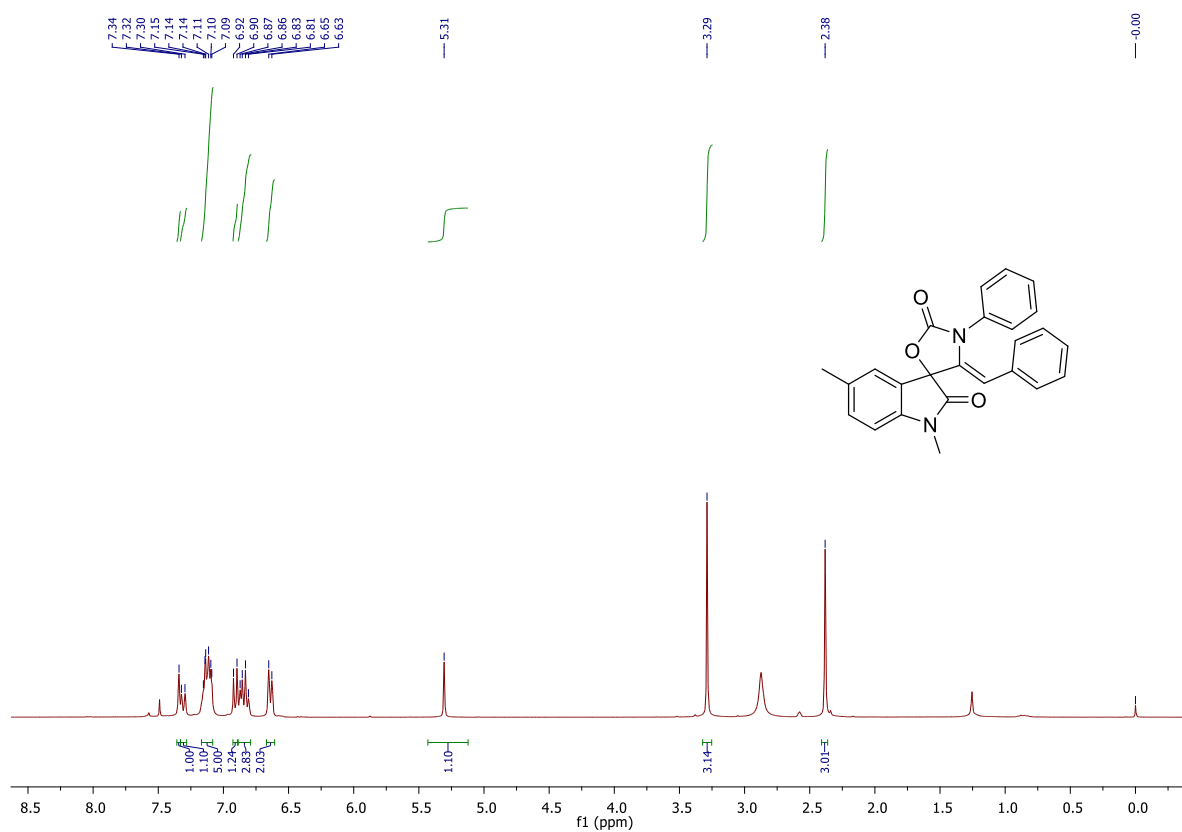
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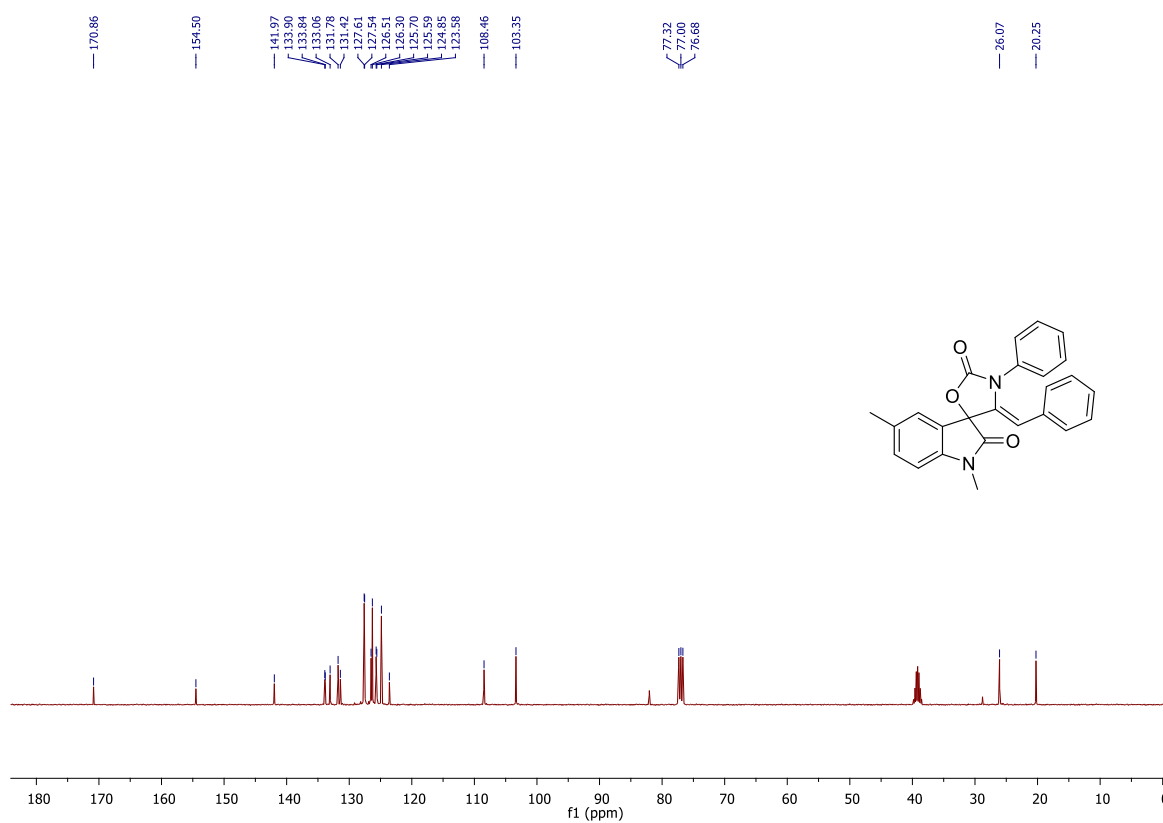
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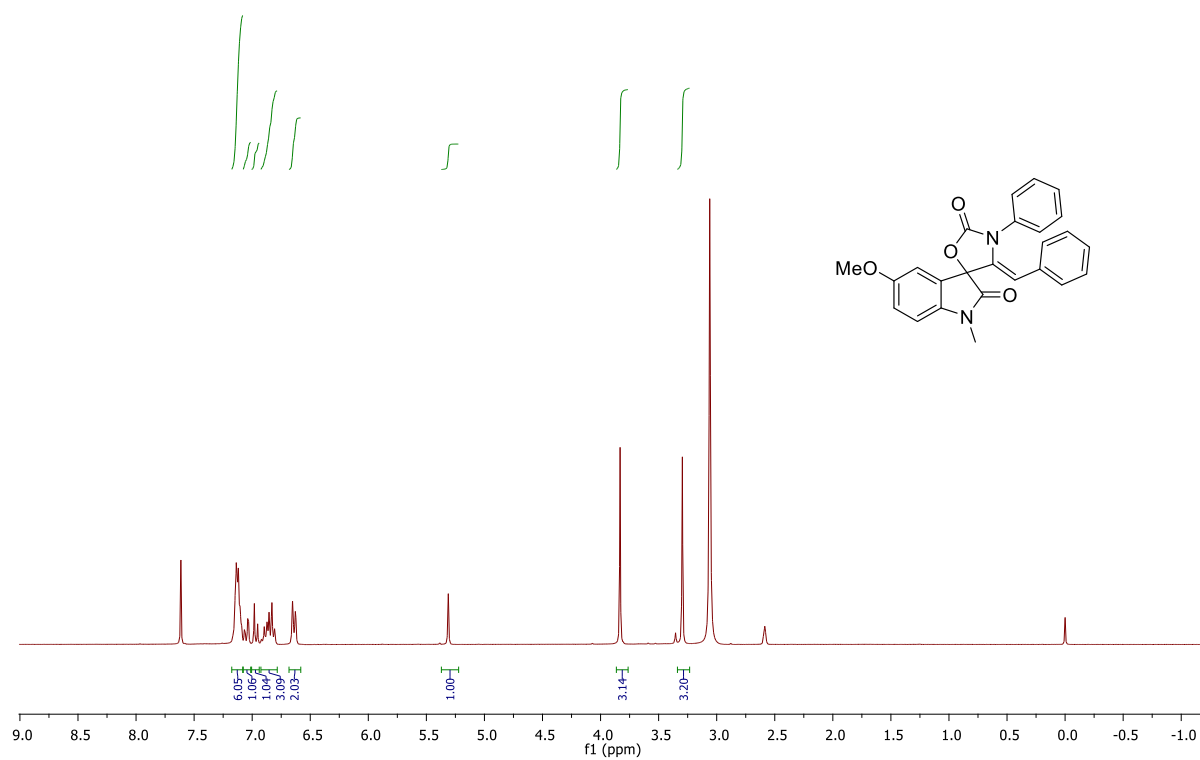
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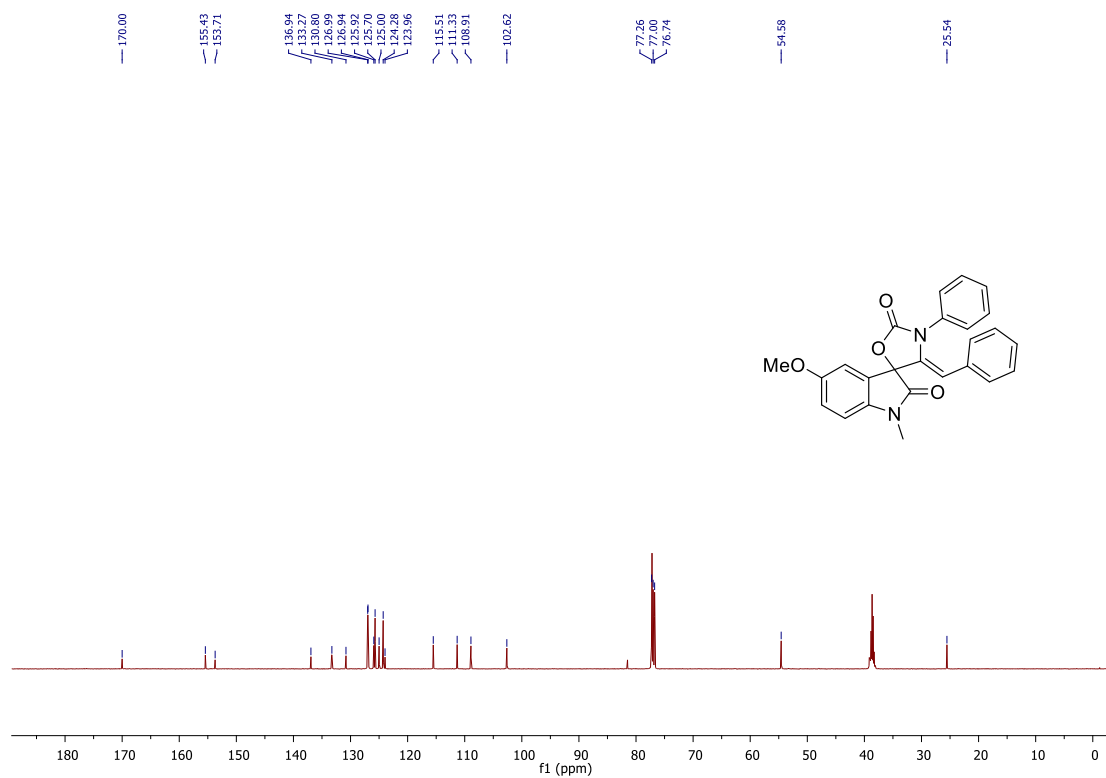
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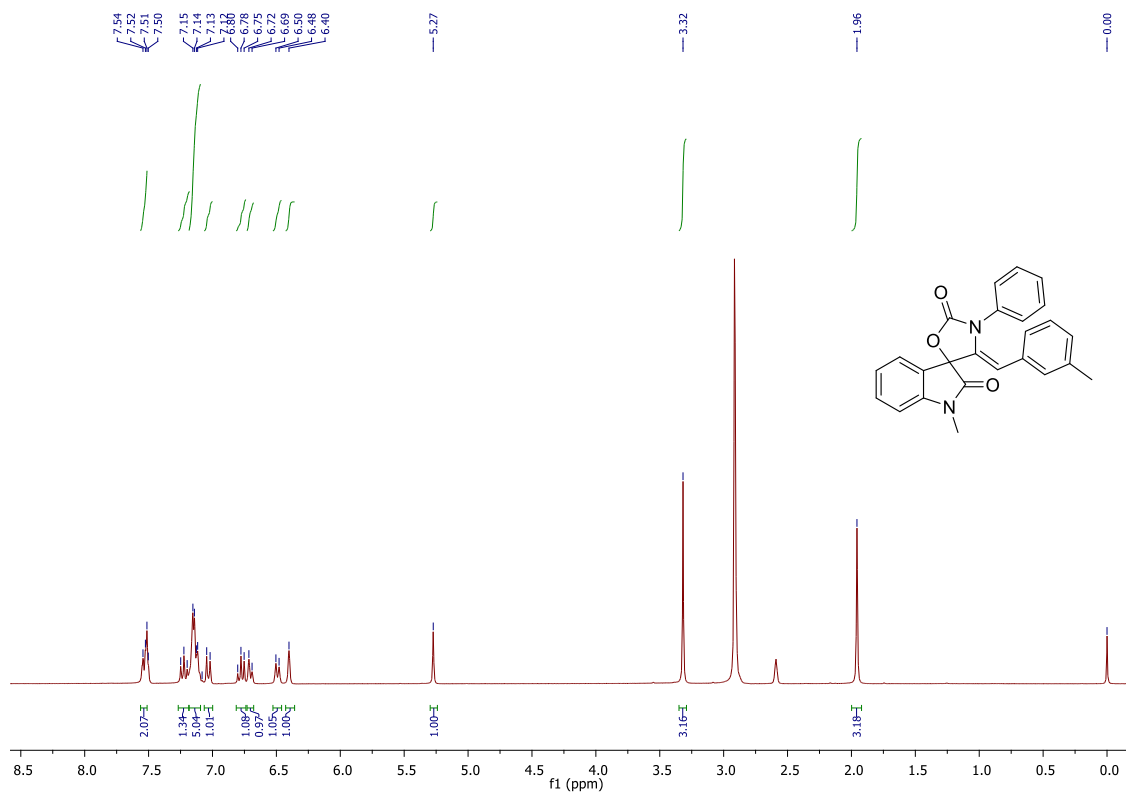
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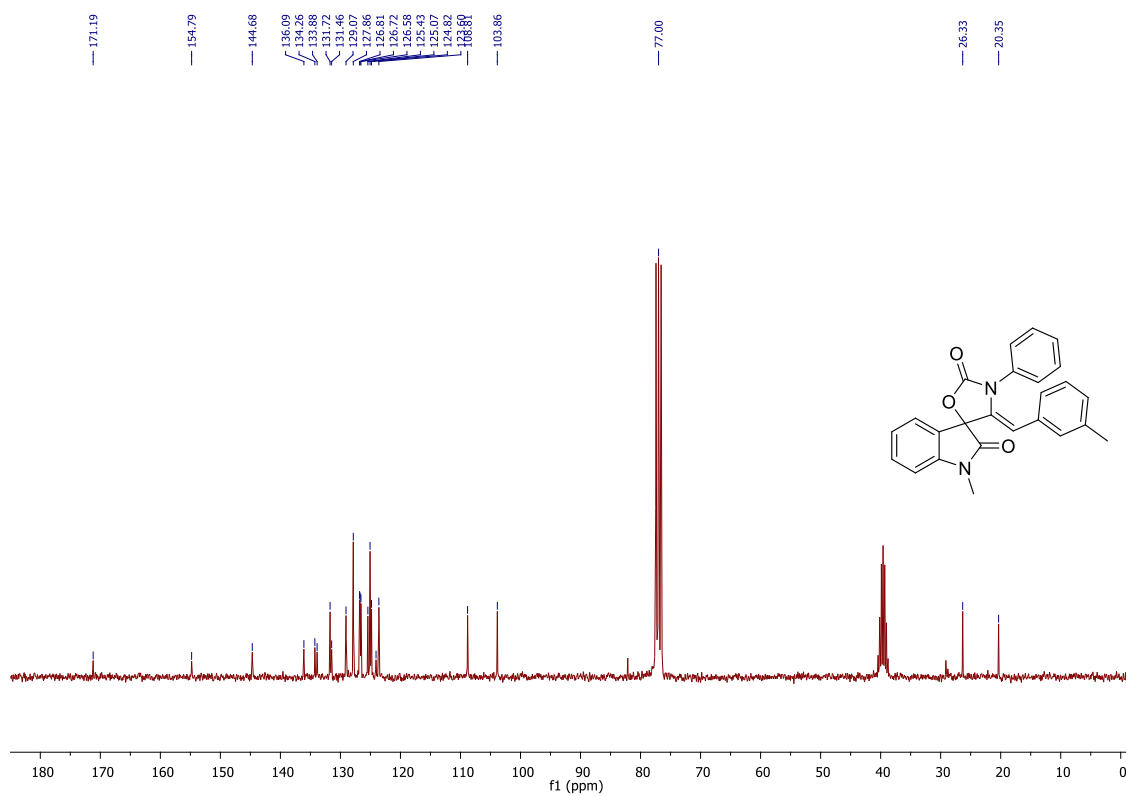
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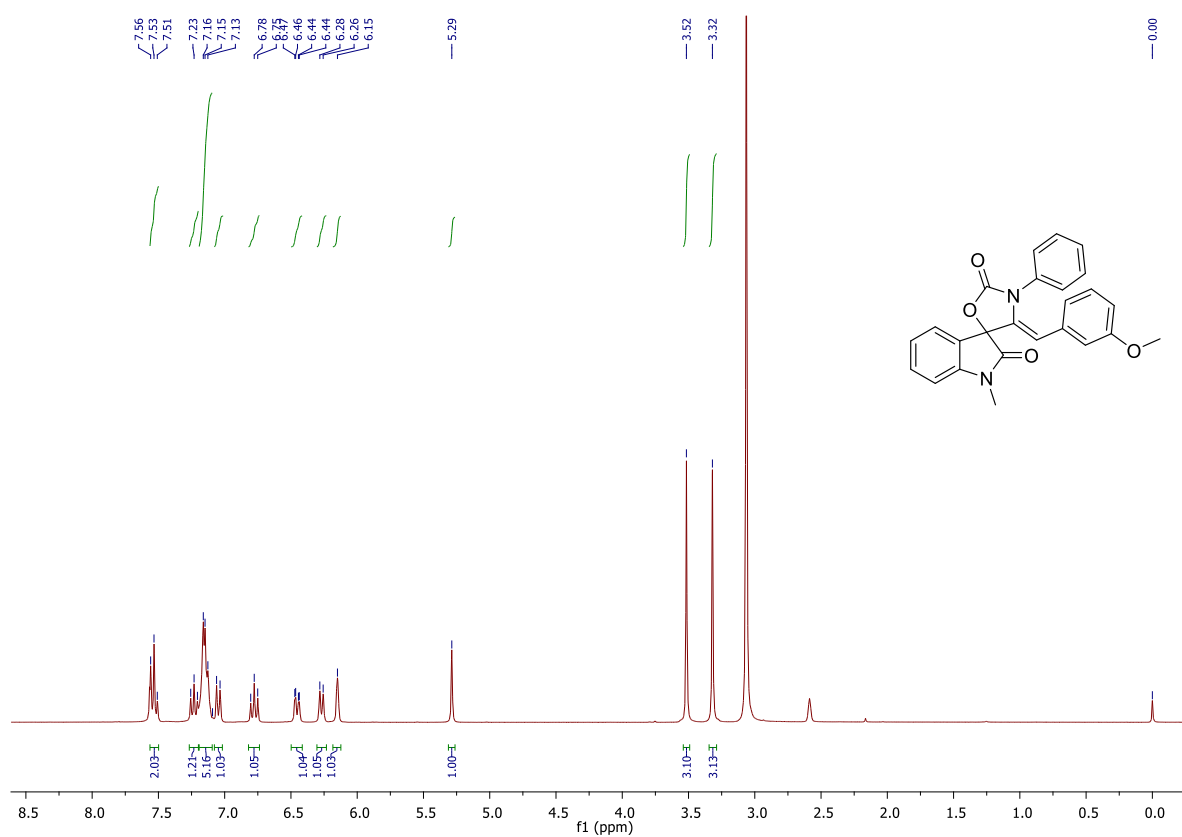
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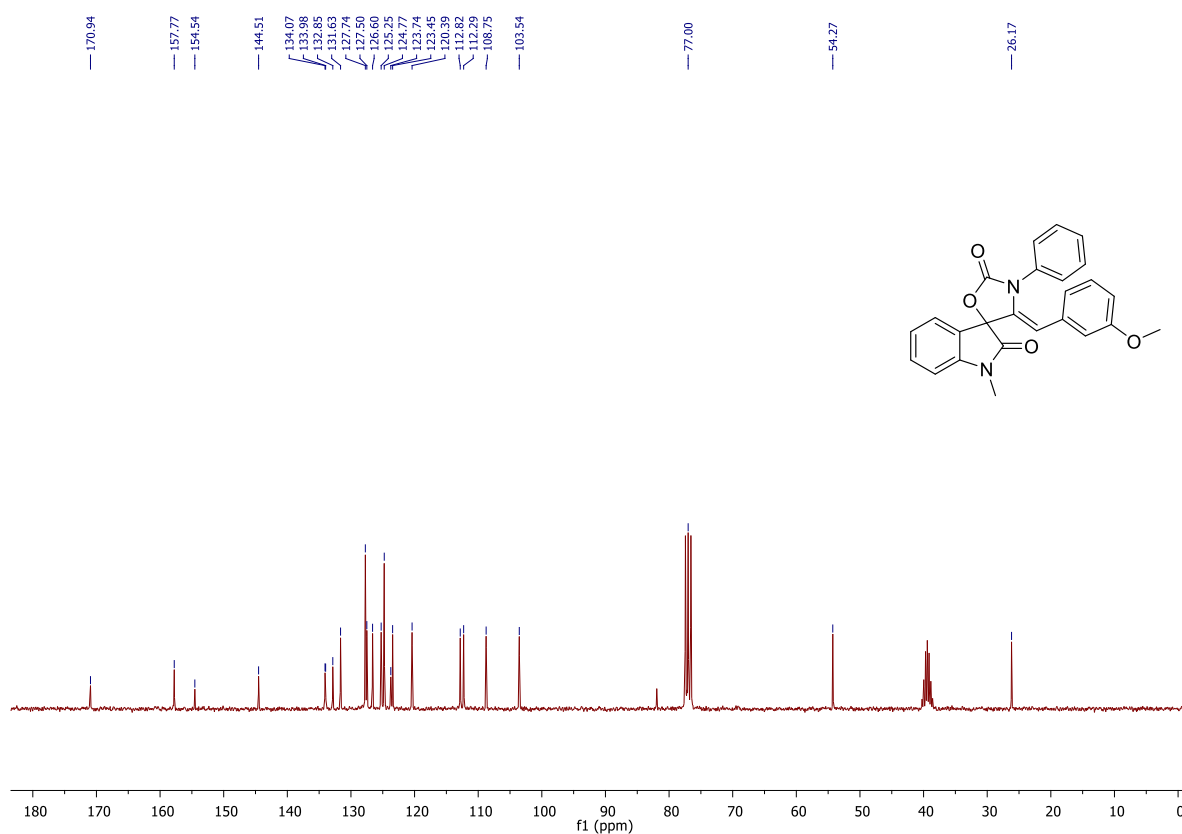
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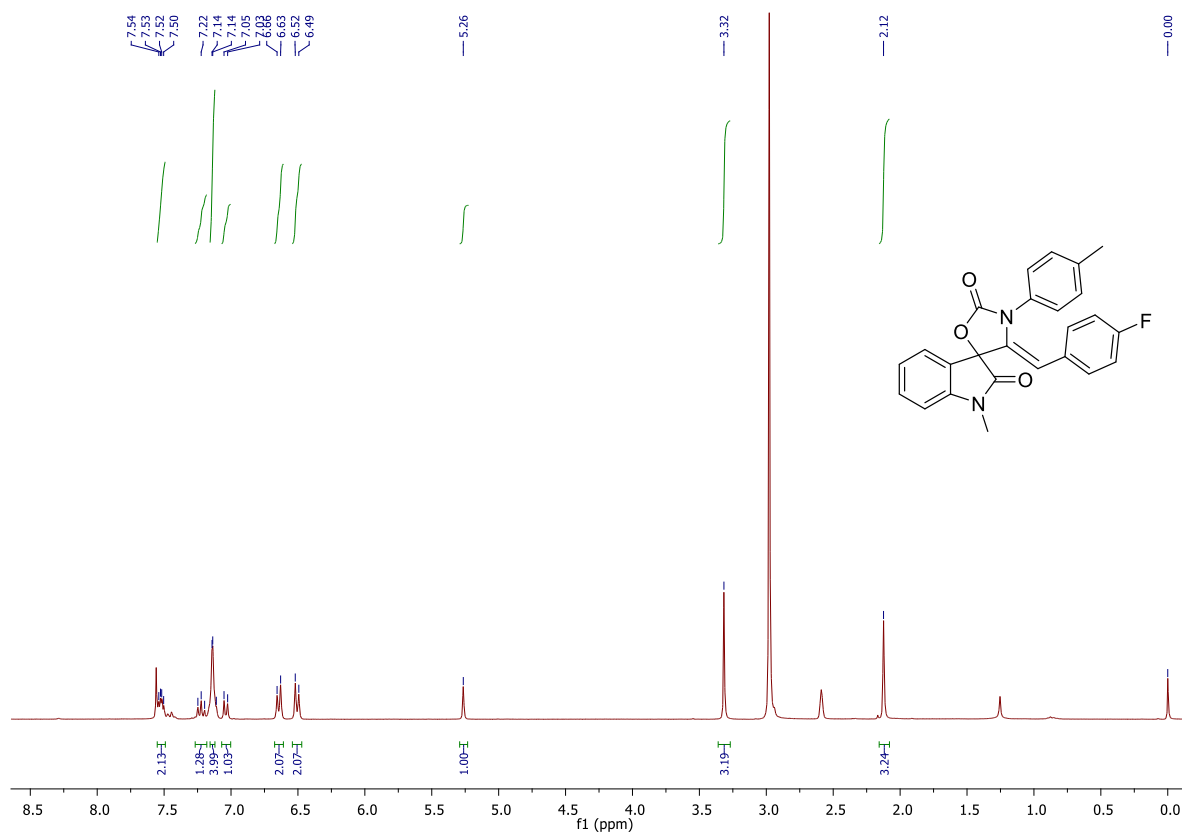
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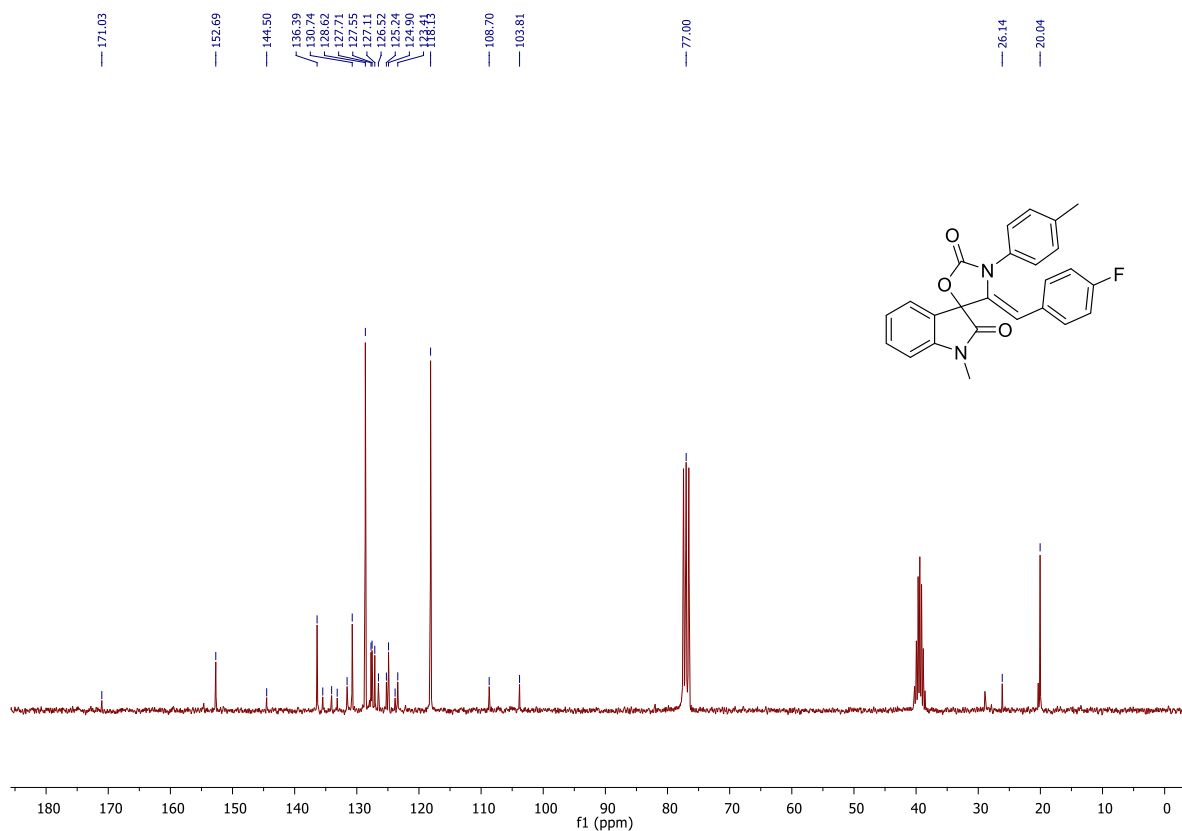
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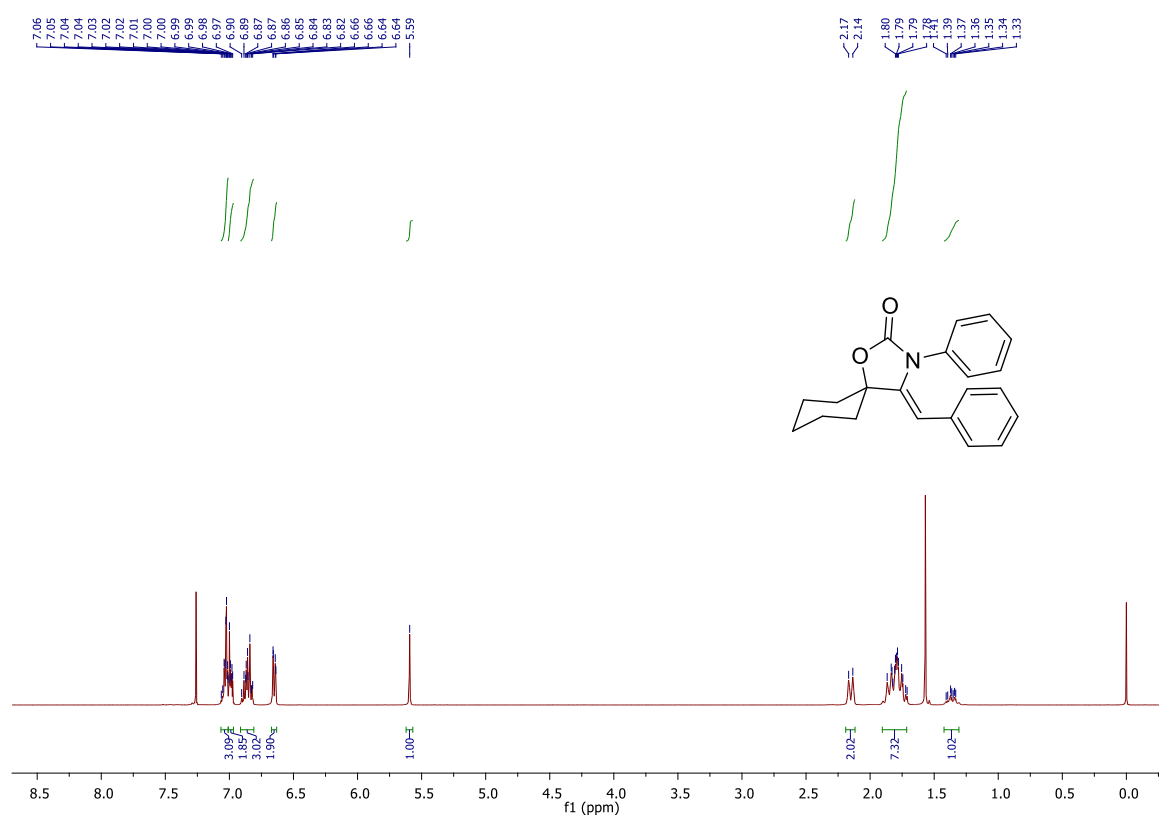
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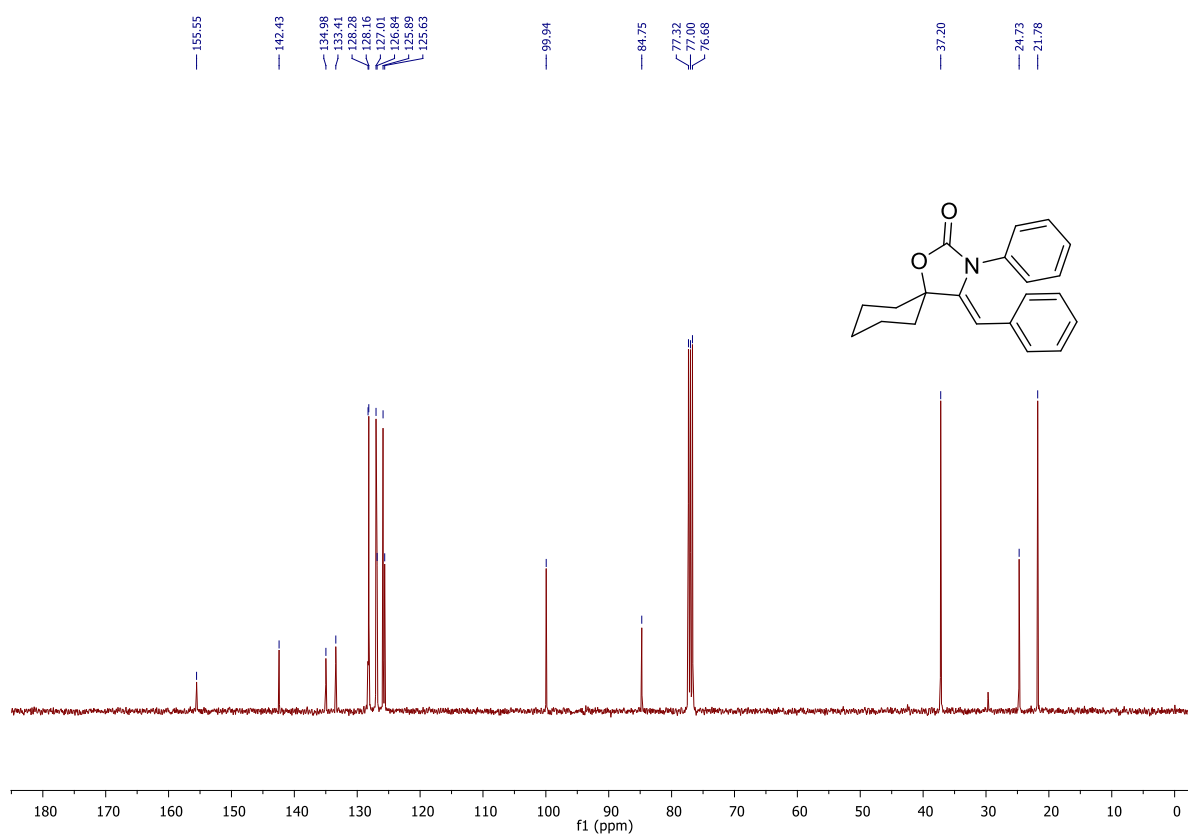
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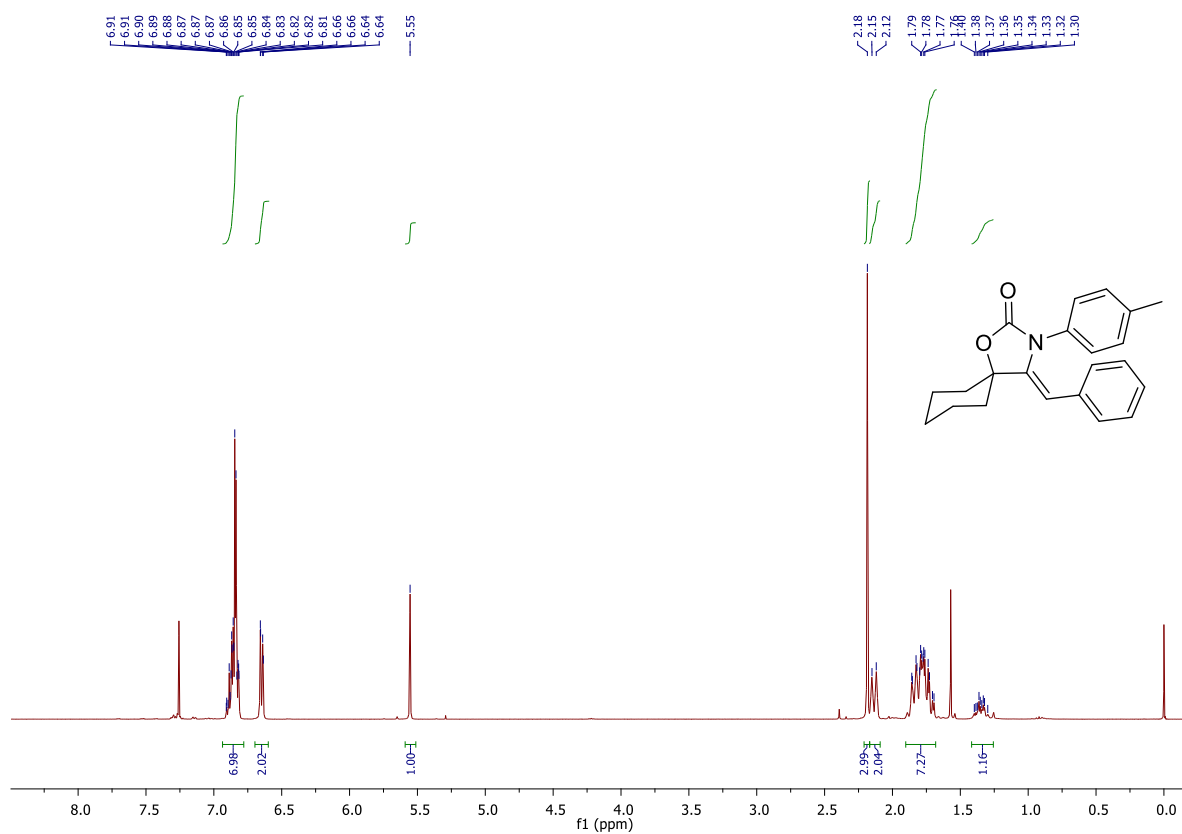
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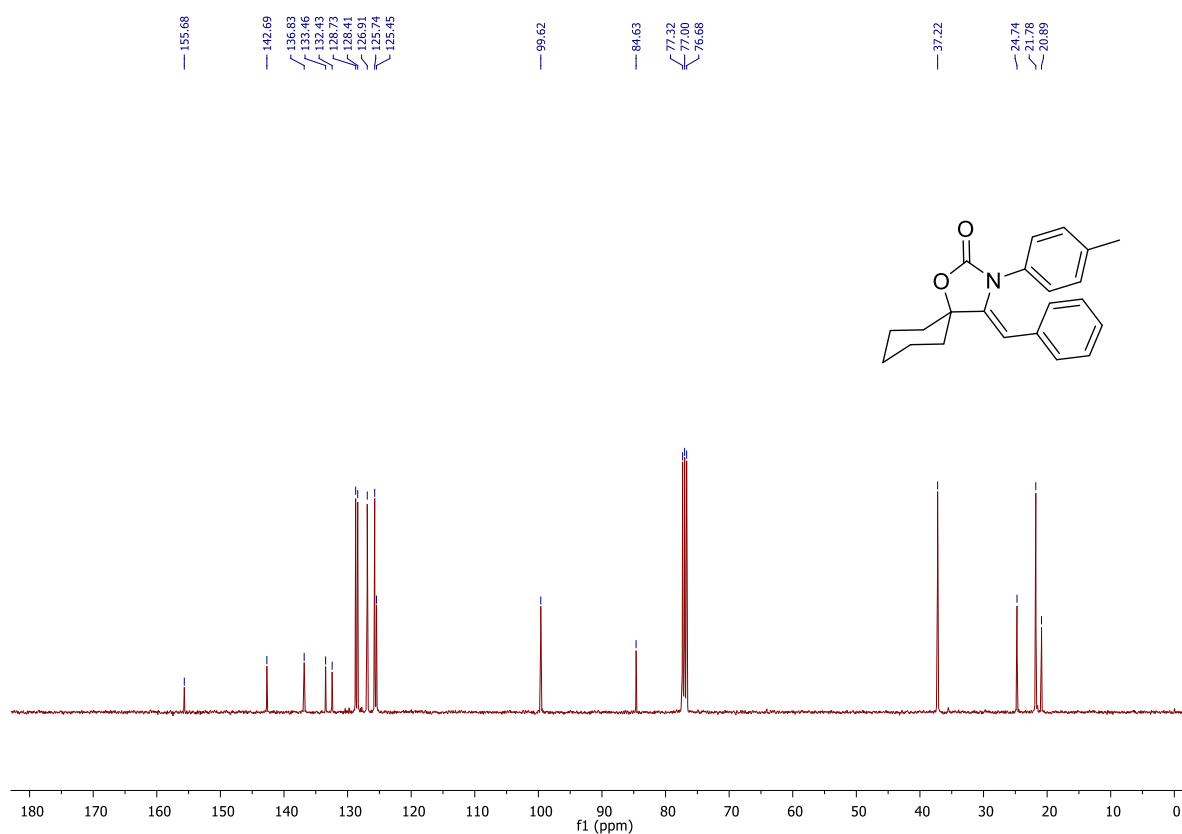
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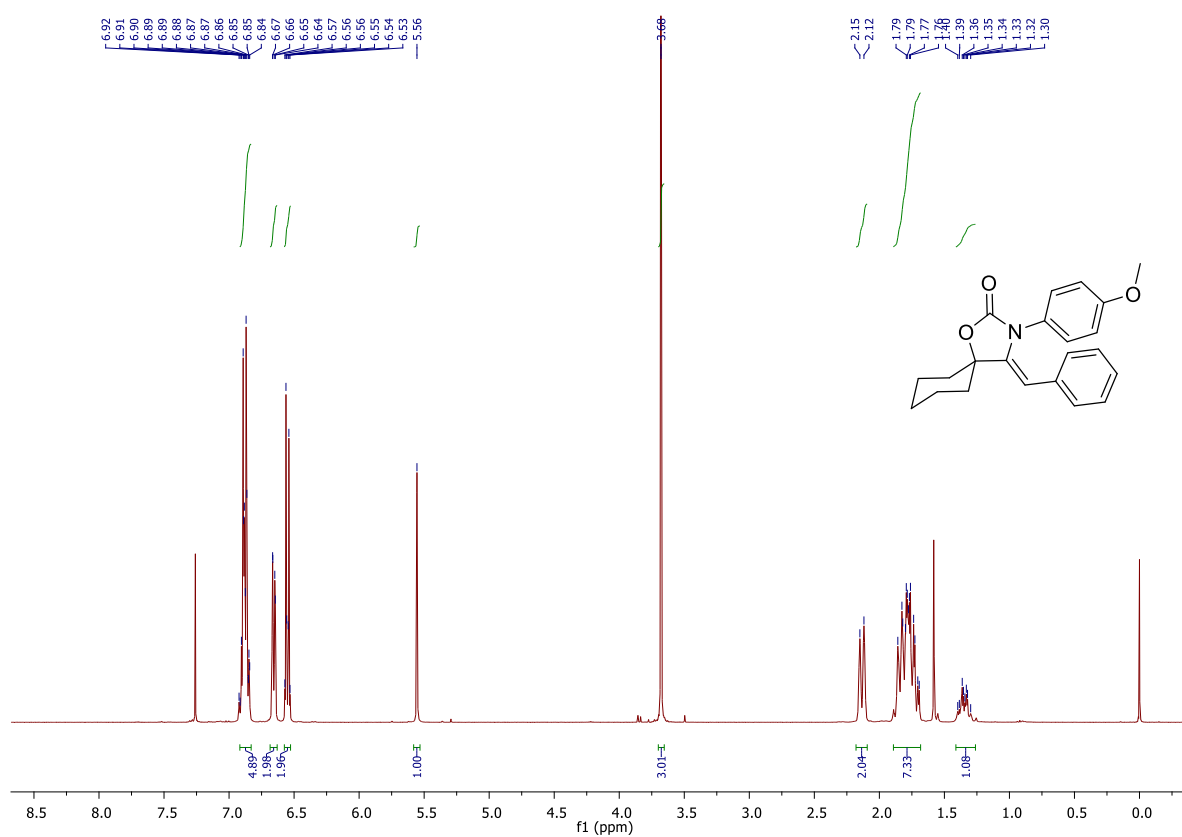
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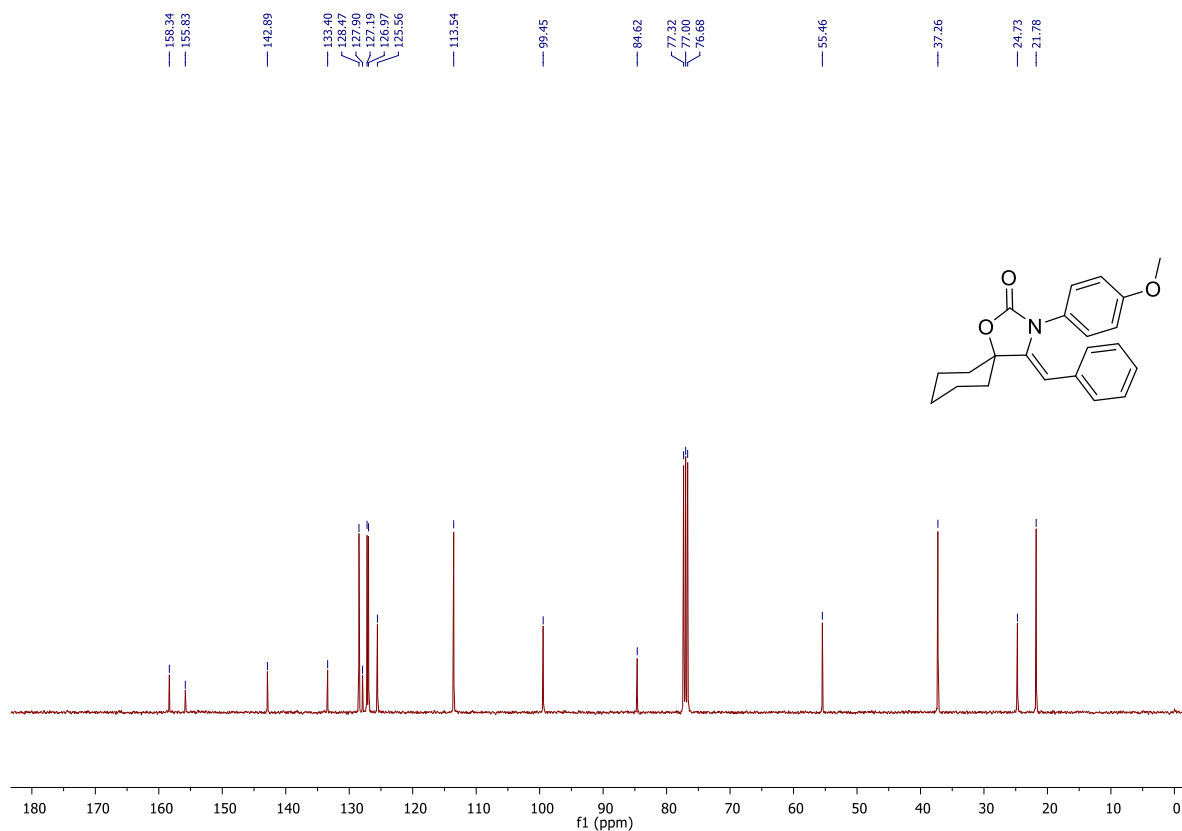
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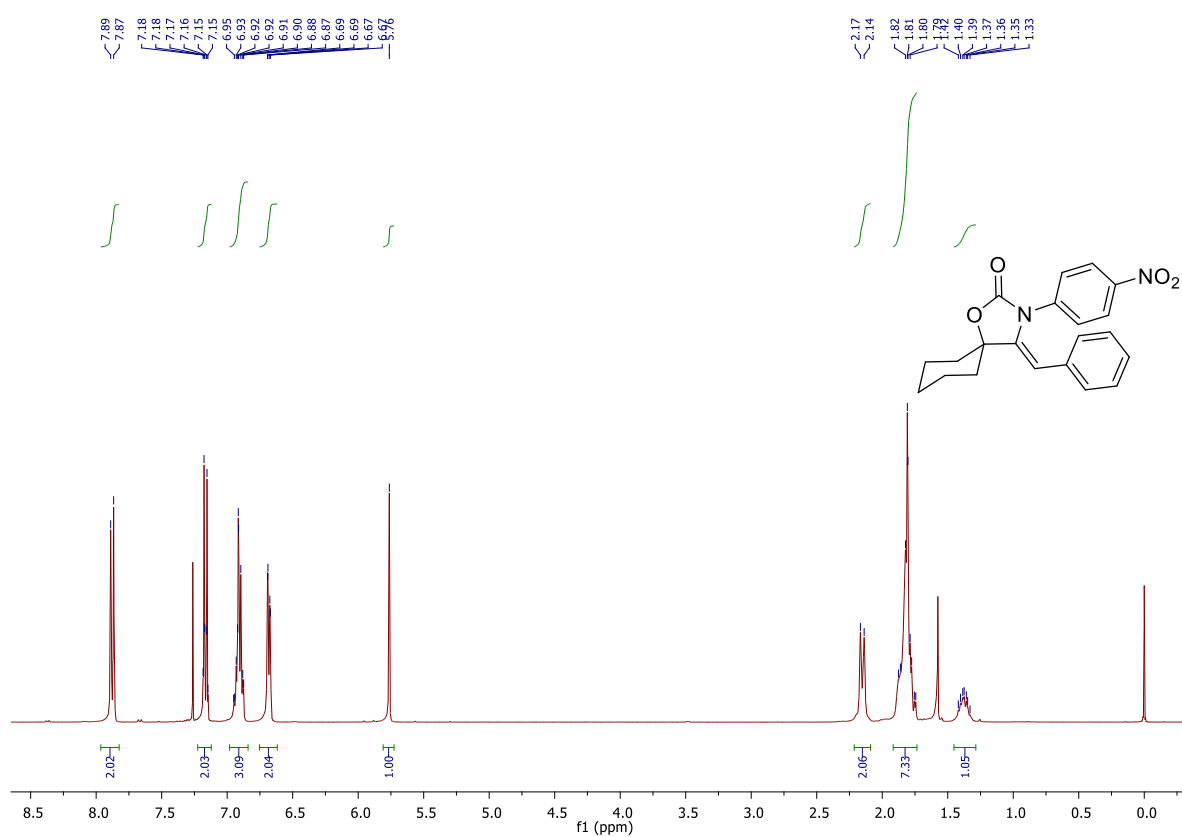
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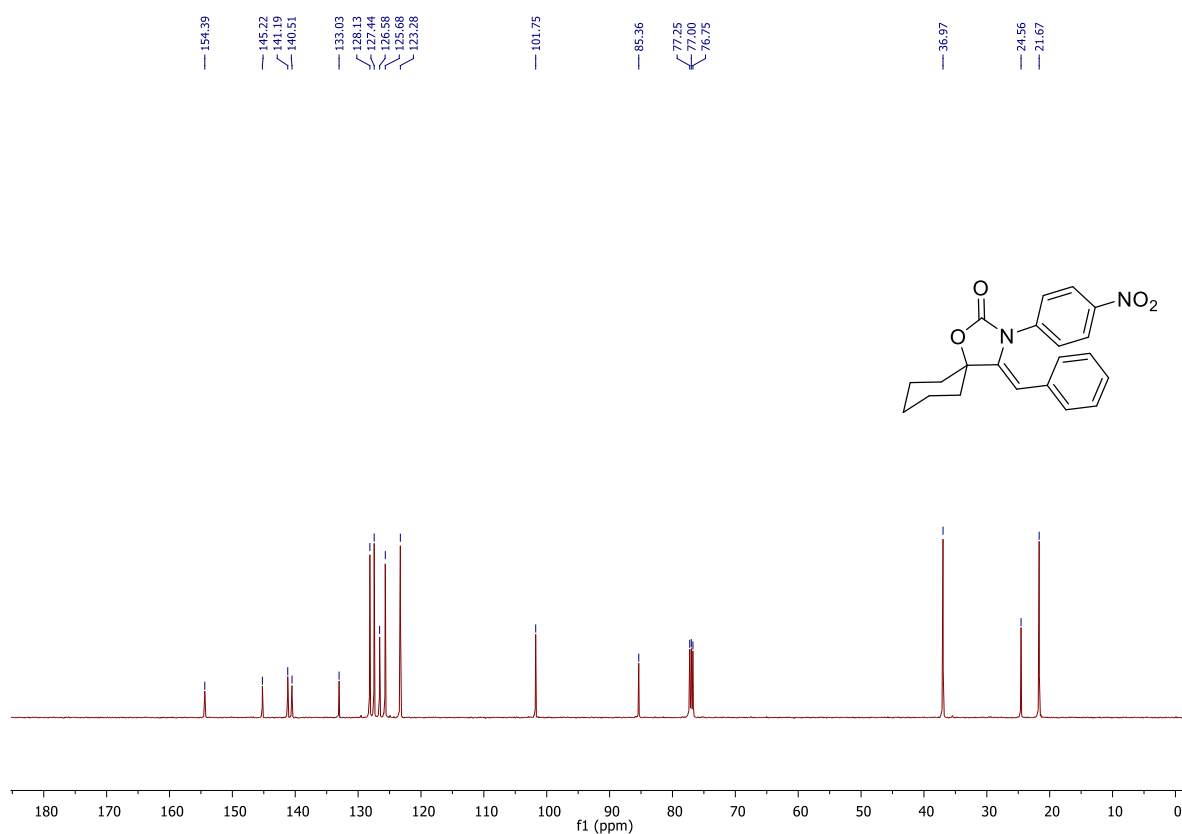
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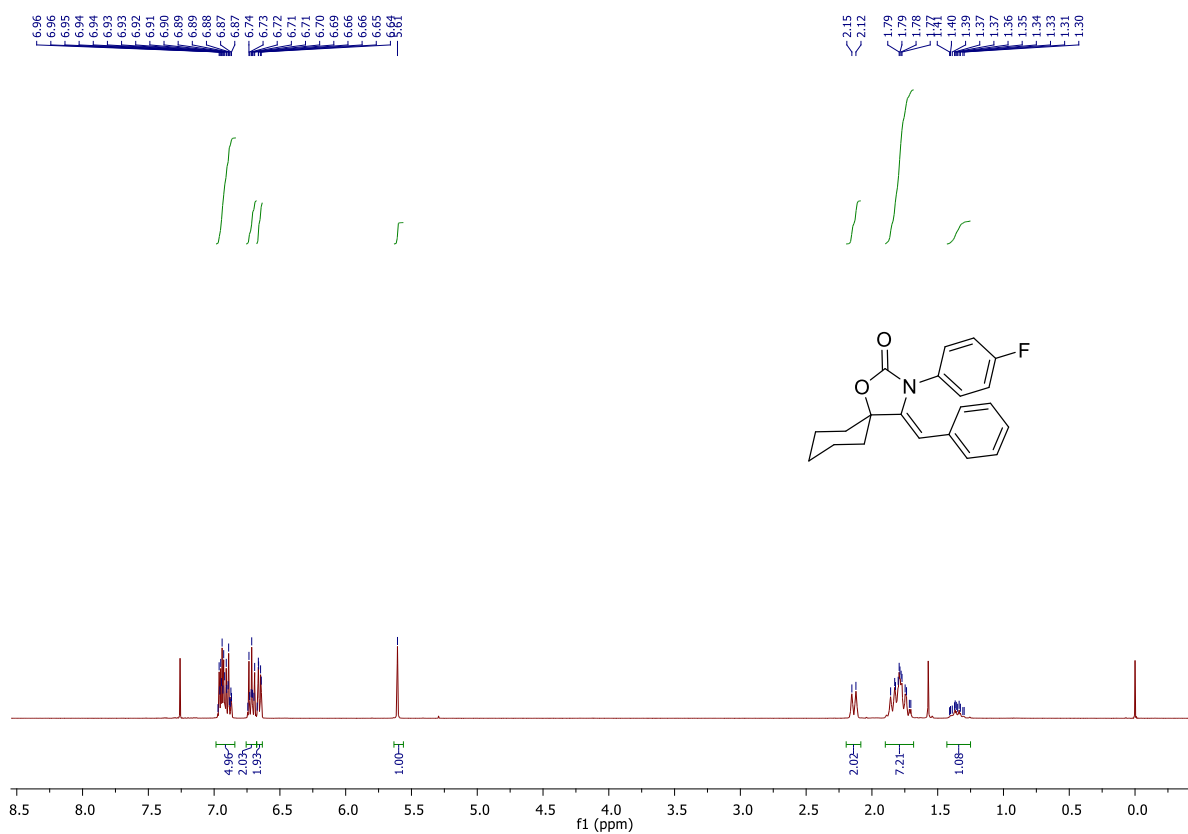
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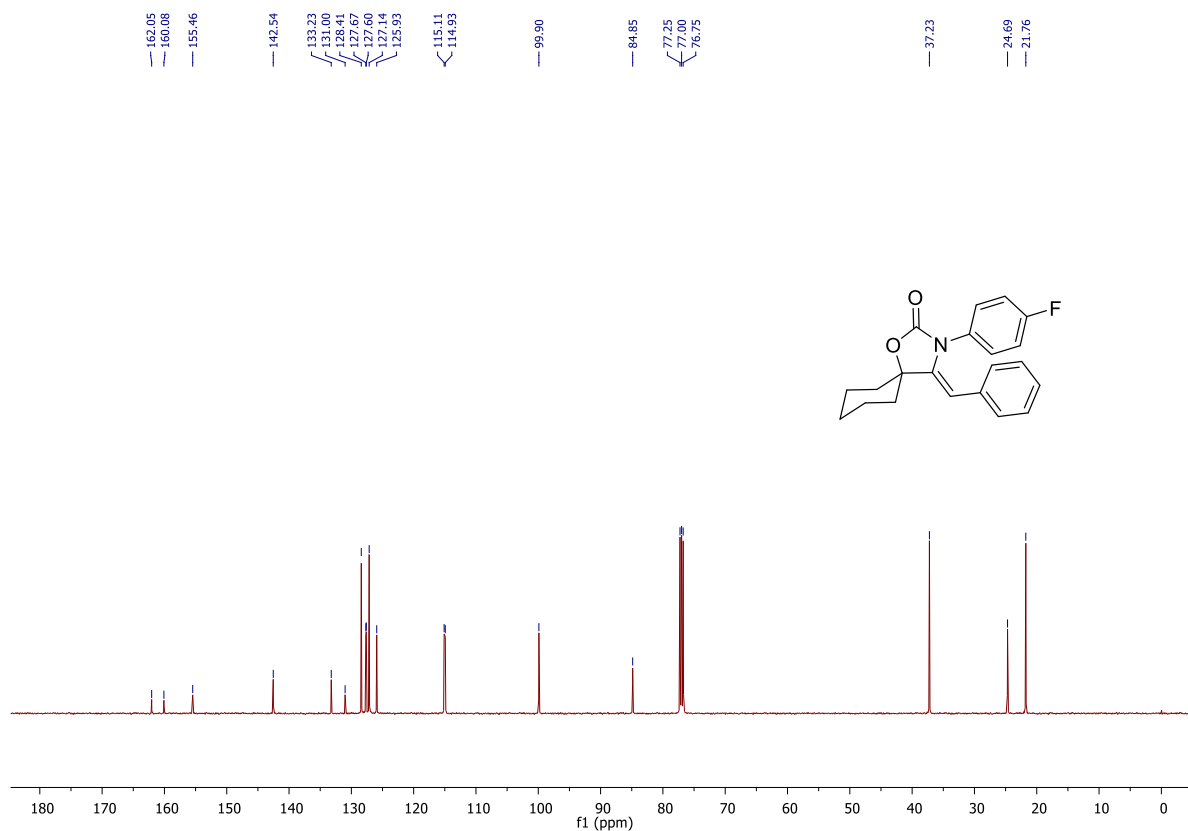
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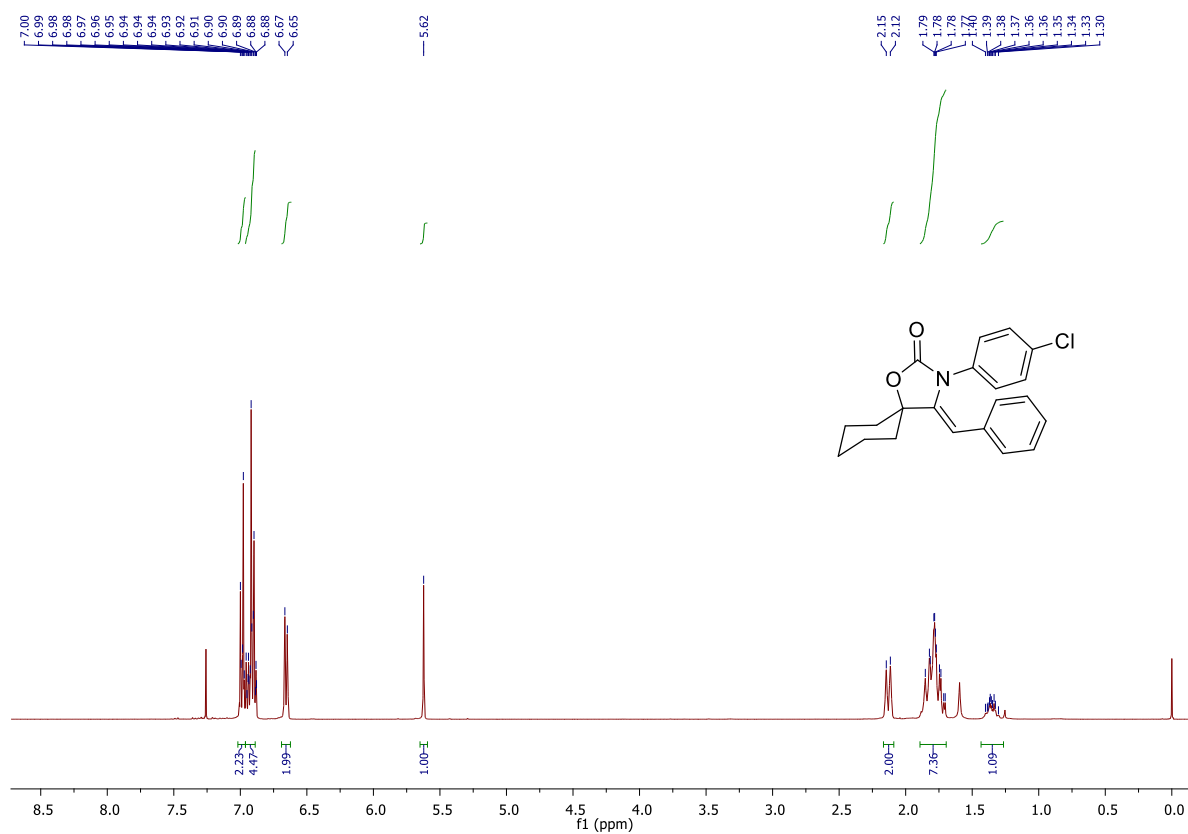
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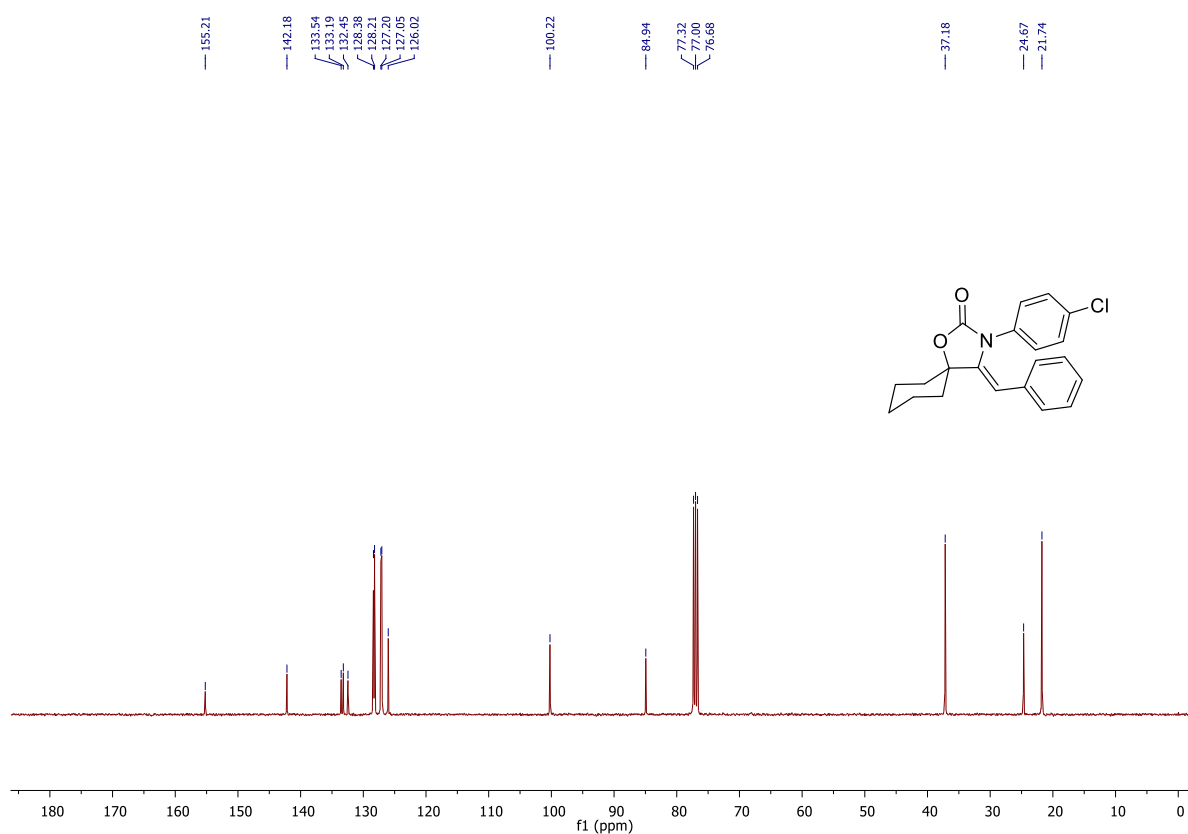
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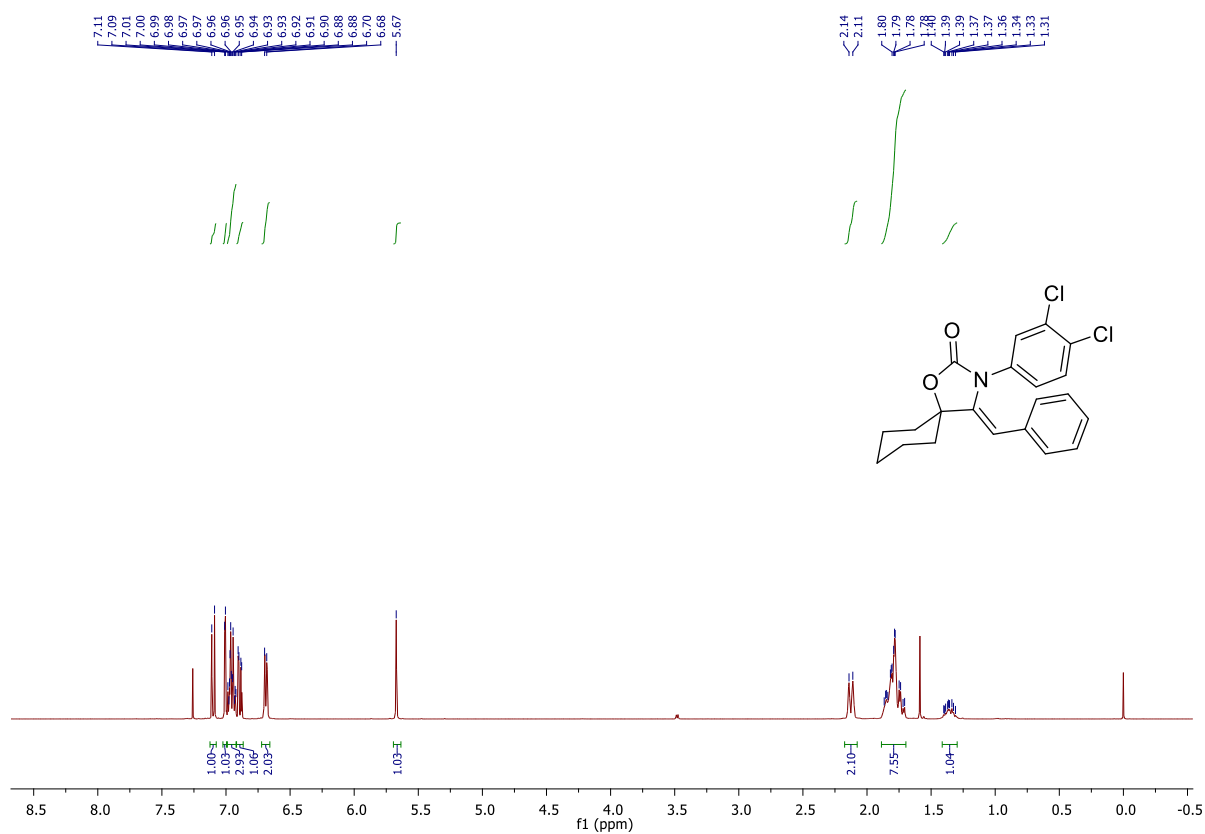
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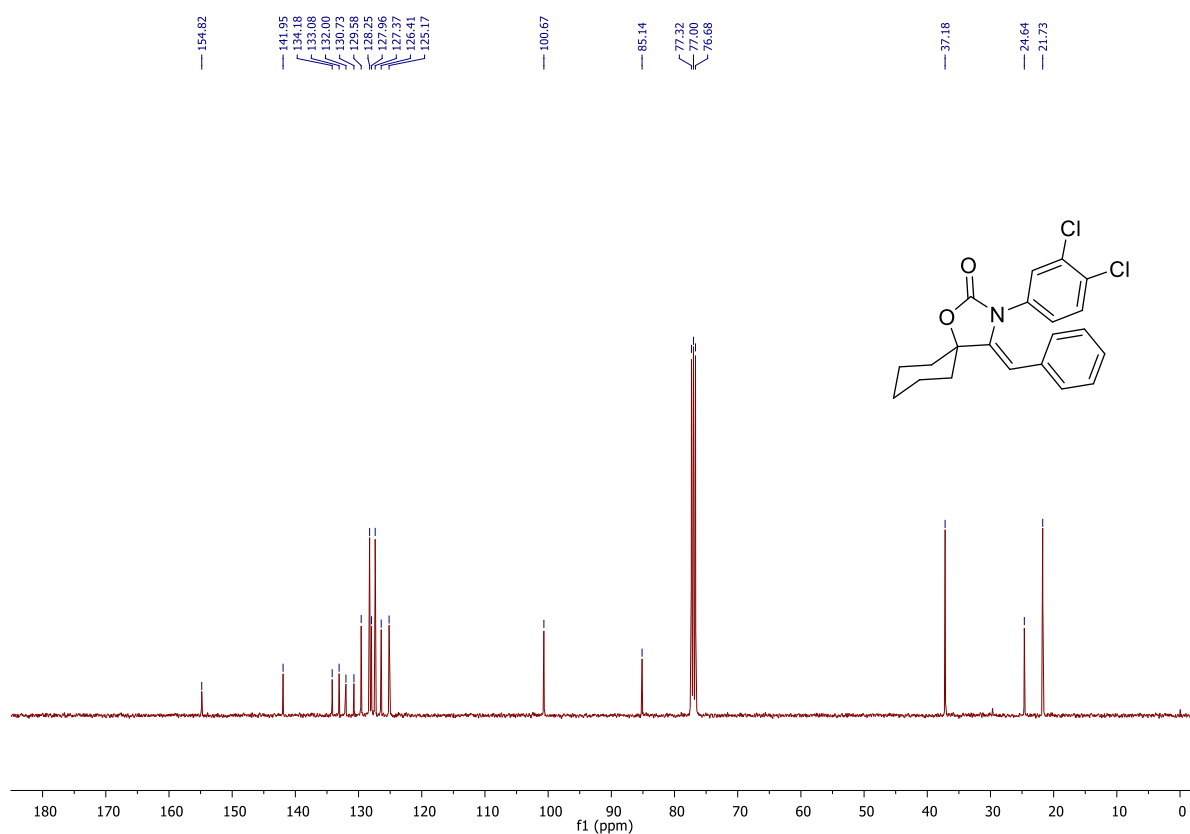
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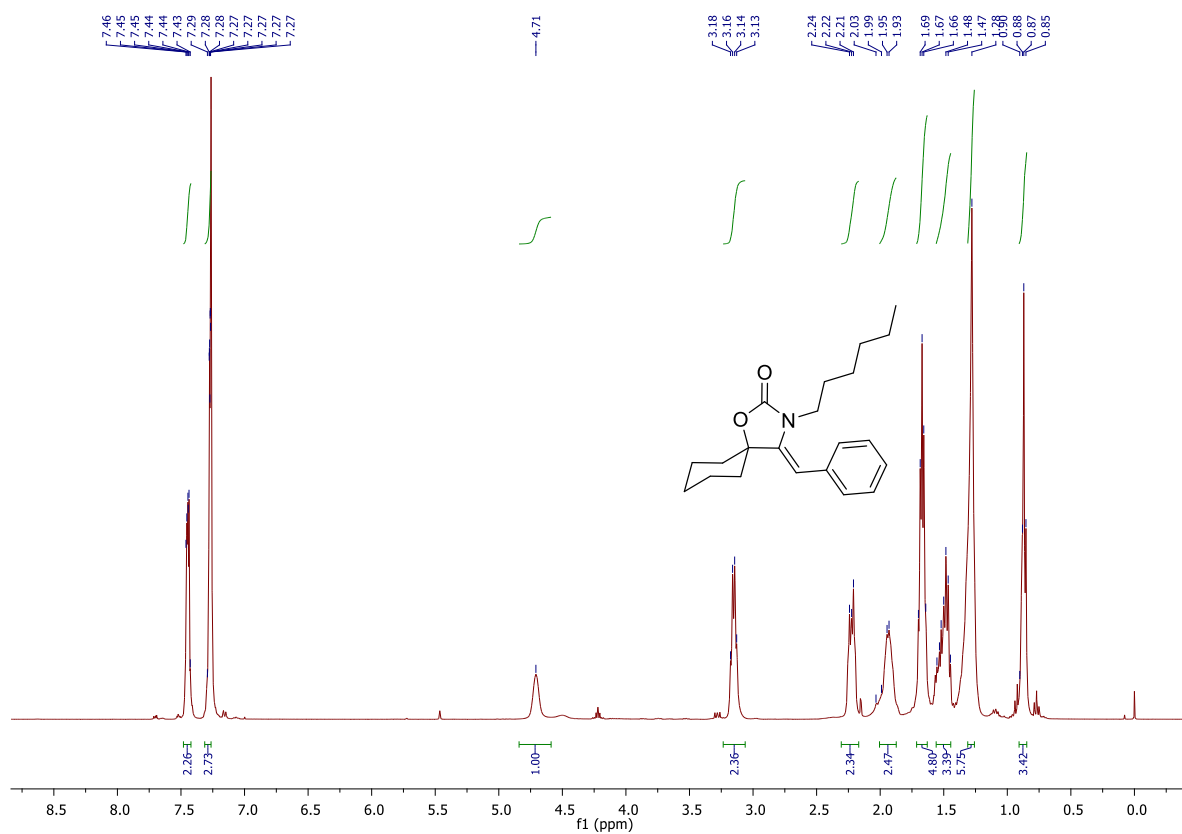
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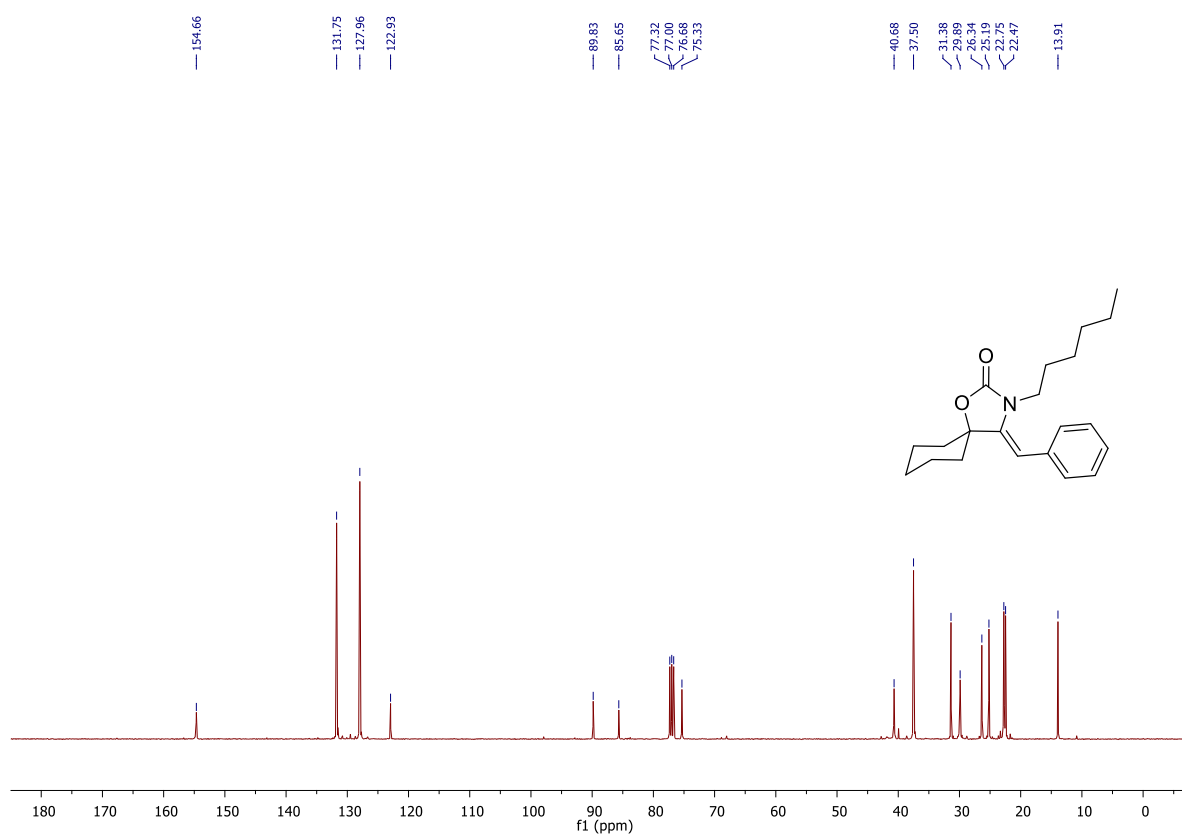
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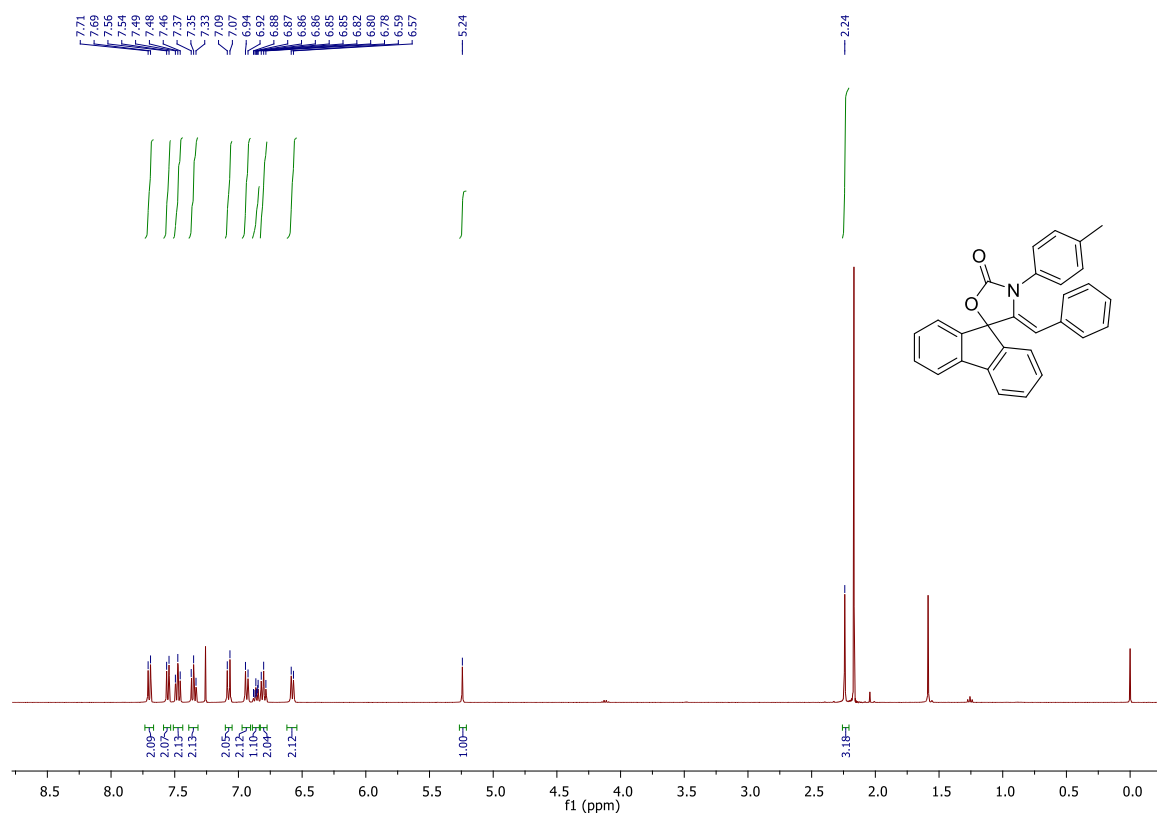
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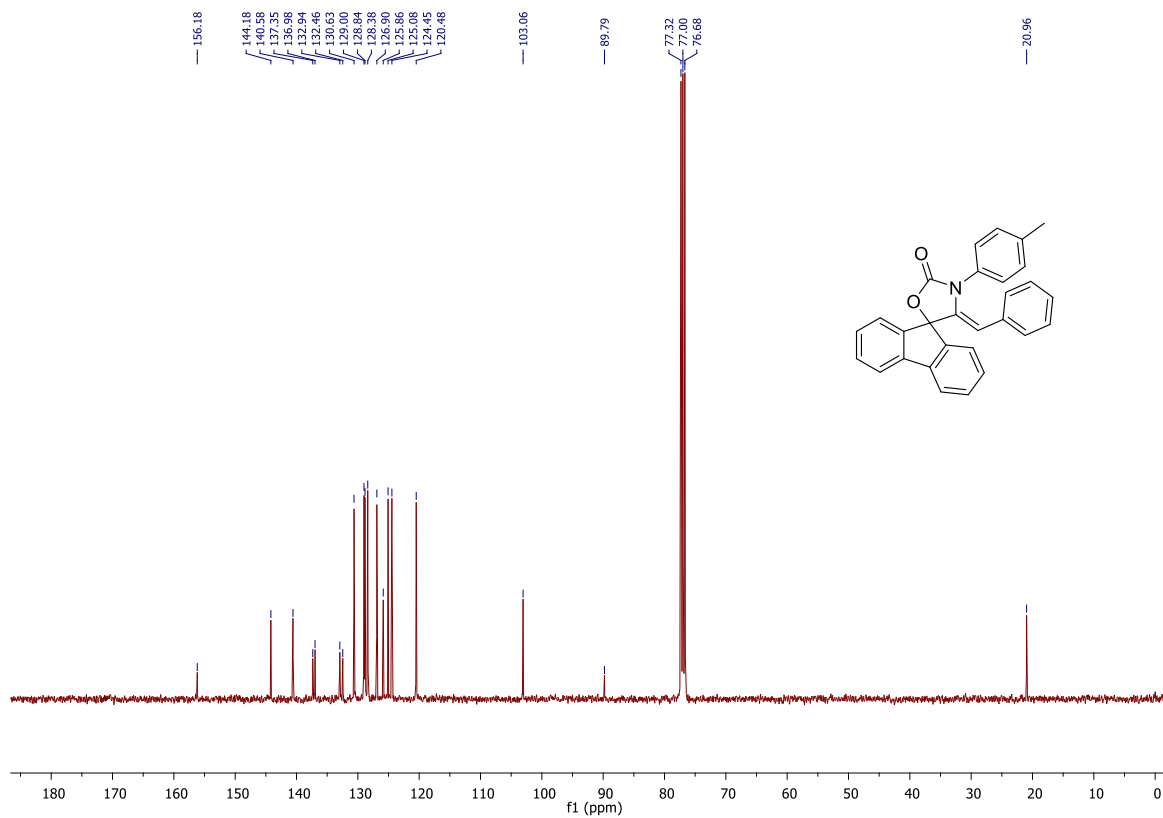
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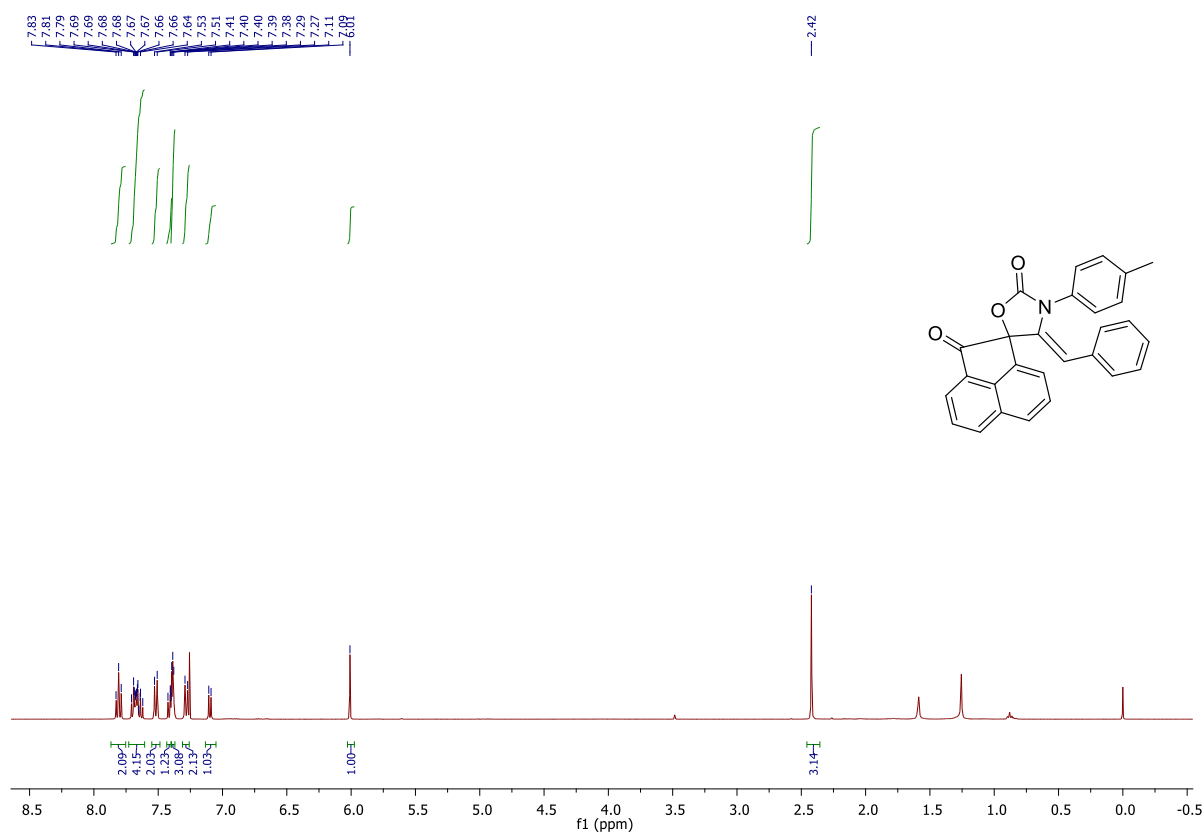
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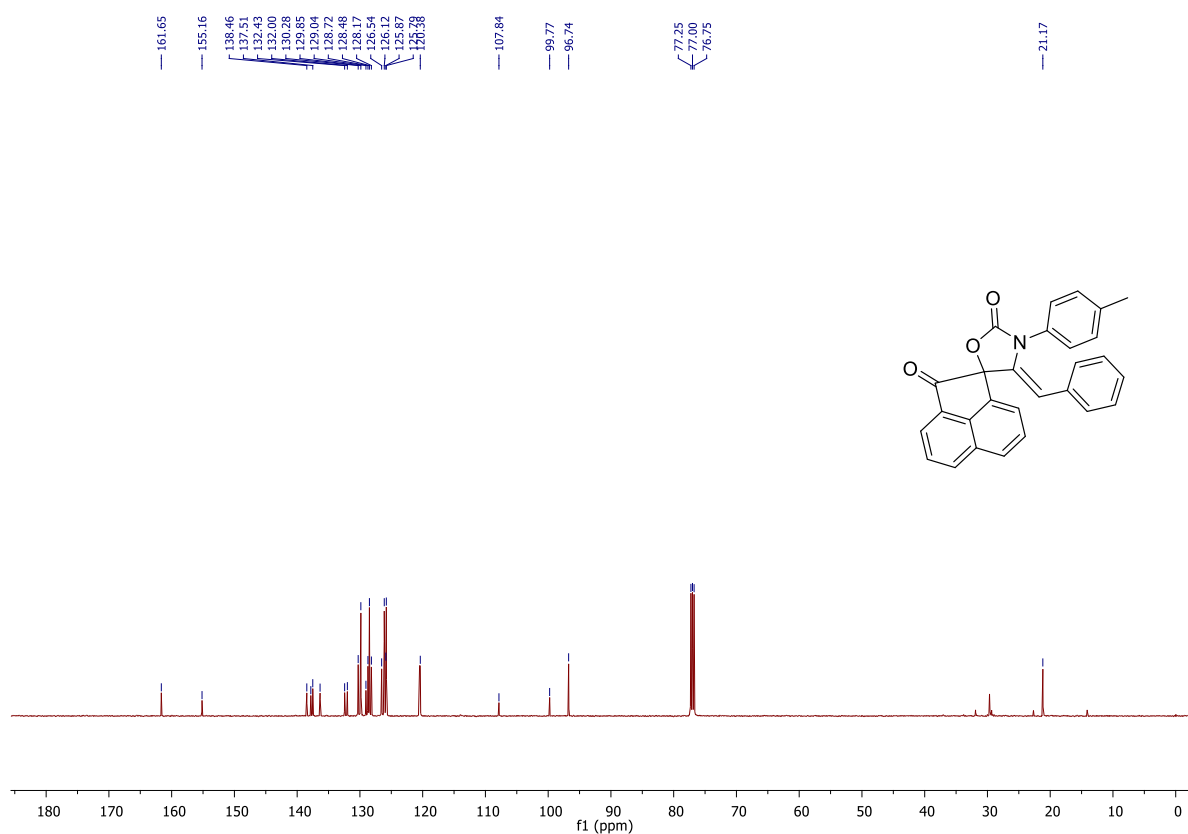
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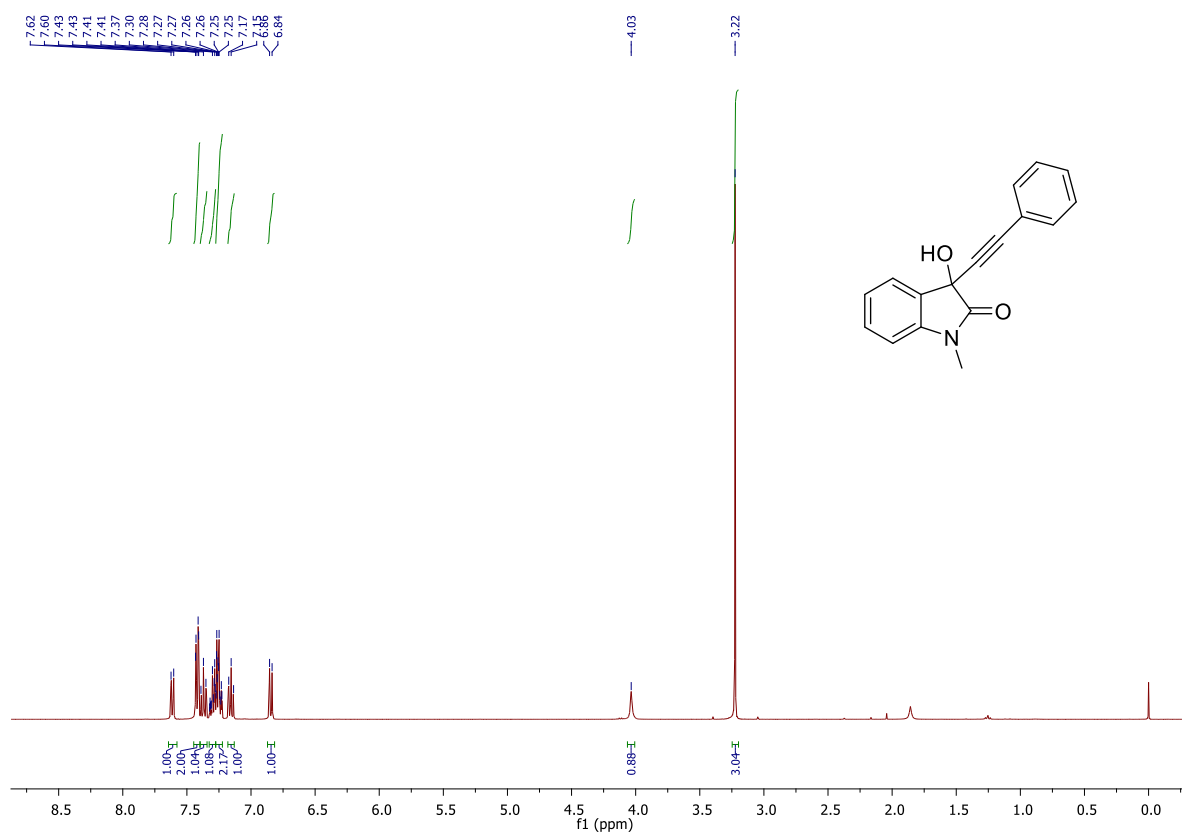
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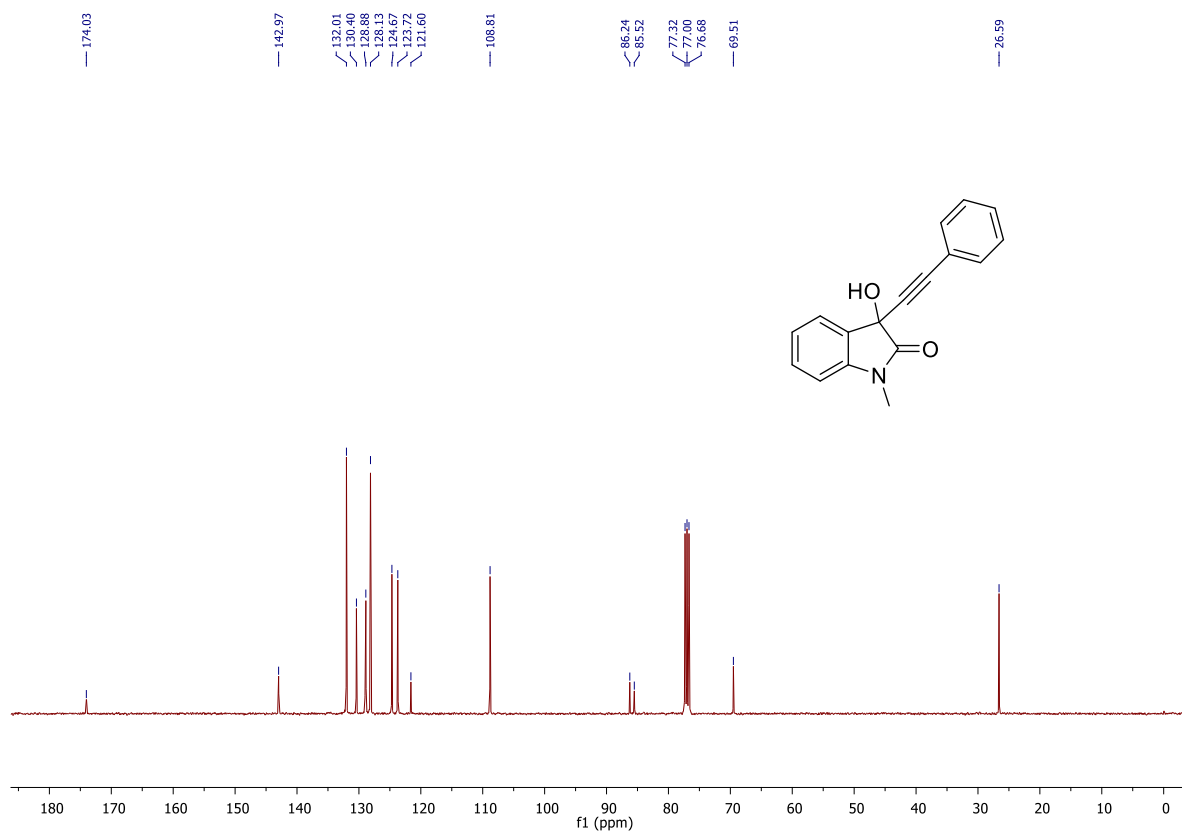
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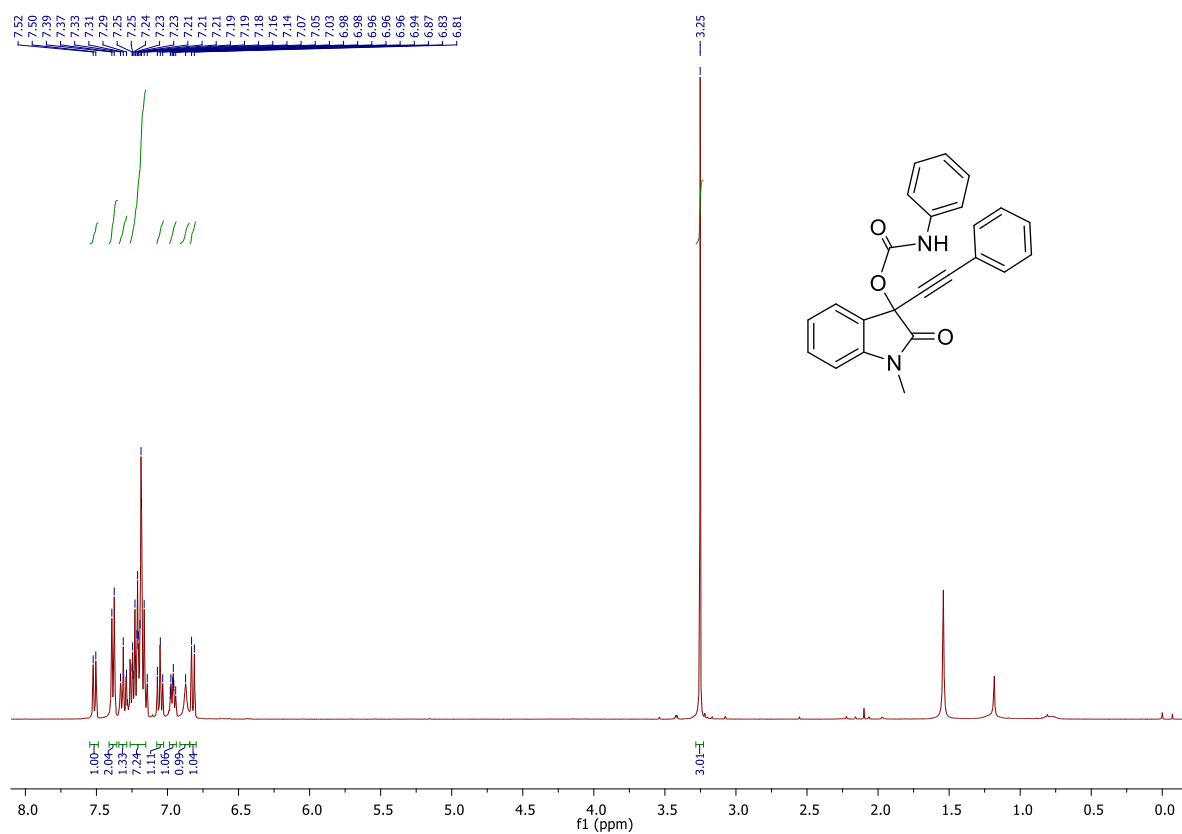
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¹³C NMR (100 MHz, CDCl₃) spectrum of compound 11



^1H NMR (400 MHz, CDCl_3) spectrum of compound 12



^{13}C NMR (100 MHz, CDCl_3) spectrum of compound 12

