

**Iodine promoted dual oxidative C (sp³) -H amination of 2-methyl-3-aryl quinazolin-4(3H)-ones:
A facile route to 1,4-diarylimidazo [1,5-a]quinazoline-5(4H)-ones**

Kavitha Donthiboina,^{a,b} Namballa Hari Krishna,^{a,b} Siddiq Pasha Shaik,^b Jagadeesh Babu Nanubolu,^c
Nagula Shankaraiah,^{*a} Ahmed Kamal^{*a,b}

^a*Department of Medicinal Chemistry, National Institute of Pharmaceutical Education and Research (NIPER), Hyderabad - 500 037, India.*

^b*Department of Medicinal Chemistry and Biotechnology, CSIR-Indian Institute of Chemical Technology, Hyderabad-500007, India.*

^c*Laboratory of X-ray Crystallography, CSIR-Indian Institute of Chemical Technology, Hyderabad-500007, India..*

Corresponding authors: Dr. Ahmed Kamal, E-mail: ahmedkamal@iict.res.in; Dr. NagulaShankaraiah, shankar@niperhyd.ac.in; shankarnbs@gmail.com

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X-ray crystallographic information and data

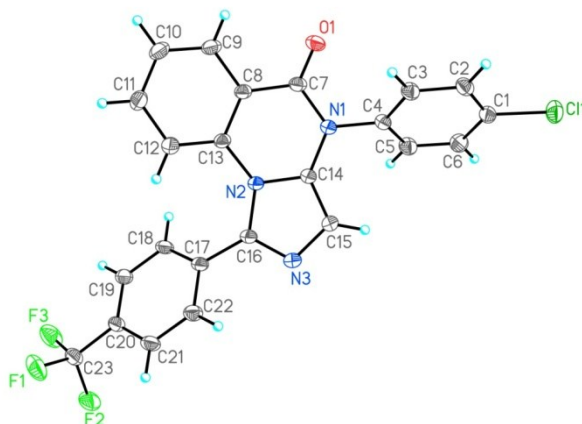


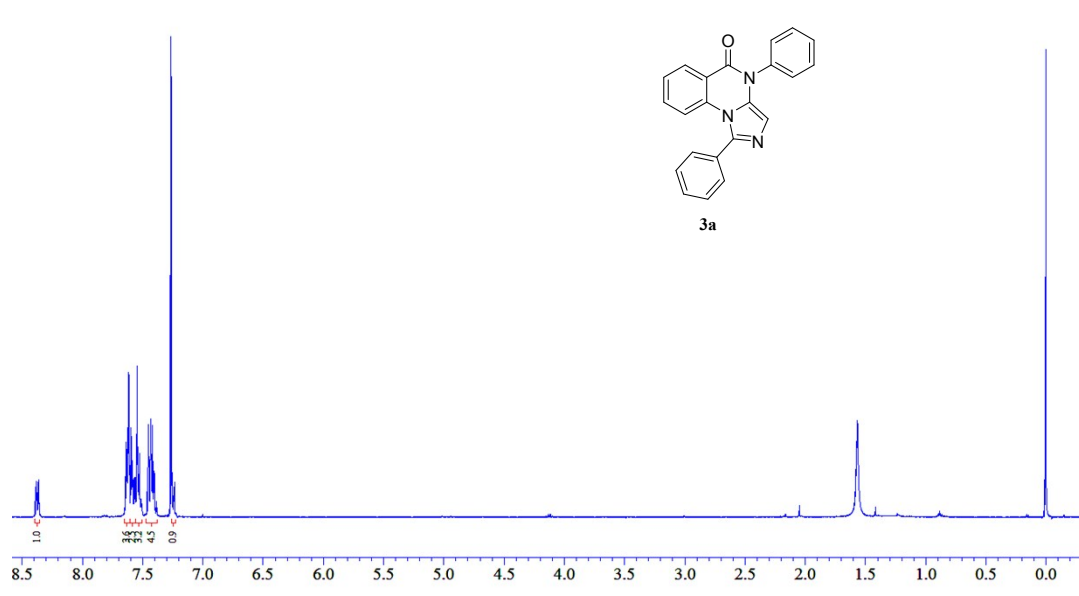
Figure caption: The molecular structure of KA129 with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radius.

Crystal data for KA129: $C_{23}H_{13}ClN_3O$, $M = 439.81$, crystal size $0.43 \times 0.32 \times 0.22 \text{ mm}^3$, triclinic, space group $P\bar{1}$ (No. 2), $a = 9.584(5)$, $b = 9.866(4)$, $c = 11.366(6) \text{ \AA}$, $\alpha = 80.384(9)$, $\beta = 84.059(12)$, $\gamma = 66.028(10)^\circ$, $V = 967.5(8) \text{ \AA}^3$, $Z = 2$, $D_c = 1.510 \text{ g/cm}^3$, $F_{000} = 448$, PHOTON 100 area detector, MoK α radiation, $\lambda = 0.71073 \text{ \AA}$, $T = 293(2) \text{ K}$, $2\theta_{\text{max}} = 61^\circ$, 24696 reflections collected, 5909 unique ($R_{\text{int}} = 0.019$), Final $GooF = 1.05$, $R_I = 0.0424$, $wR2 = 0.1245$, R indices based on 4261 reflections with $I > 2\sigma(I)$ (refinement on F^2), 280 parameters, $\mu = 0.247 \text{ mm}^{-1}$, Min. and Max. Resd.Dens. = -0.35 , $0.71 \text{ e/\AA}^3$. CCDC **1579886** 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/>

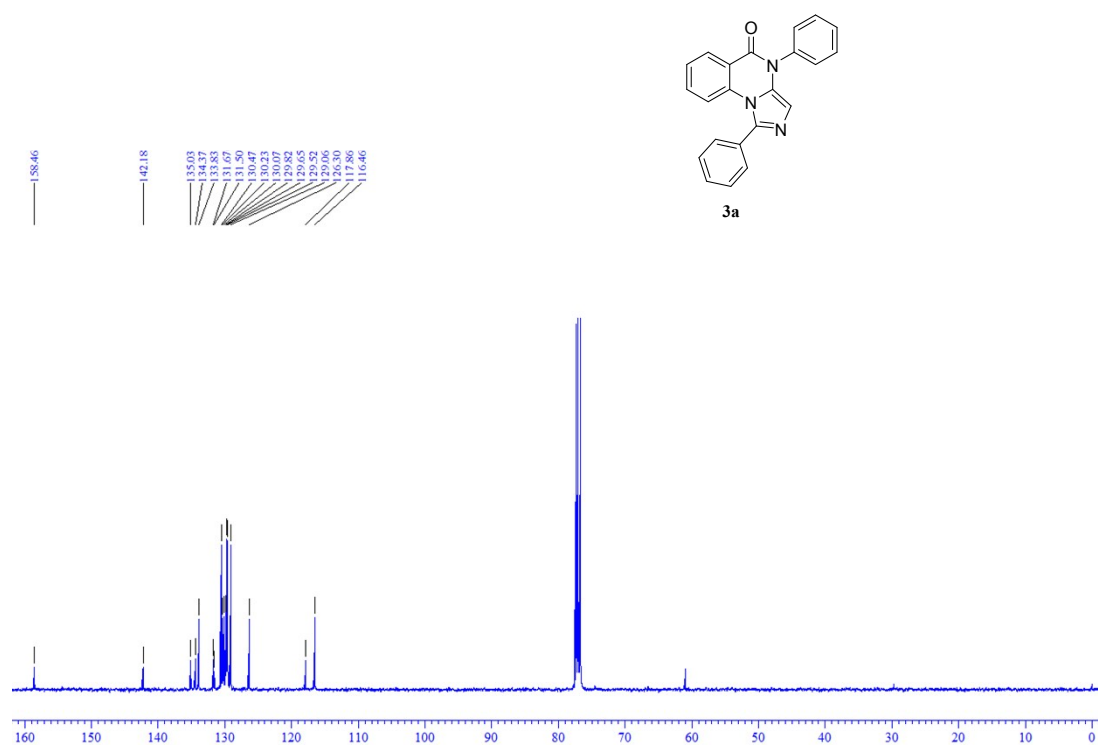
Data collection and structure solution of KA129: Single crystal X-ray data were collected at room temperature on a Bruker D8 QUEST equipped with a four circle kappa diffractometer and Photon 100 detector. An I μ smicrofocus Mo source ($\lambda=0.71073 \text{ \AA}$) supplied the multi-mirror monochromated incident beam. A combination of Phi and Omega scans were used to collect the necessary data. Unit cell dimensions were determined using 9954 reflections. 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 $0.93\text{--}0.97 \text{ \AA}$, and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}$ for methyl atoms.

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.

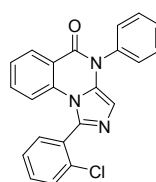
Copies of spectra



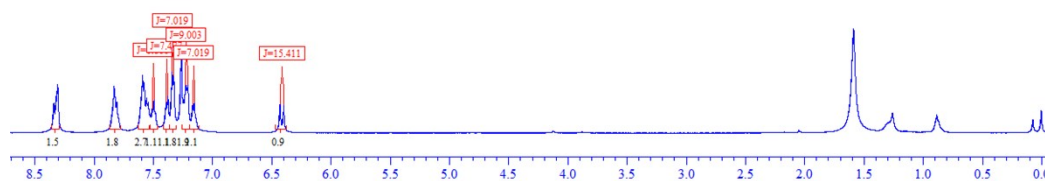
¹H NMR (400 MHz, CDCl₃) spectrum of compound 3a



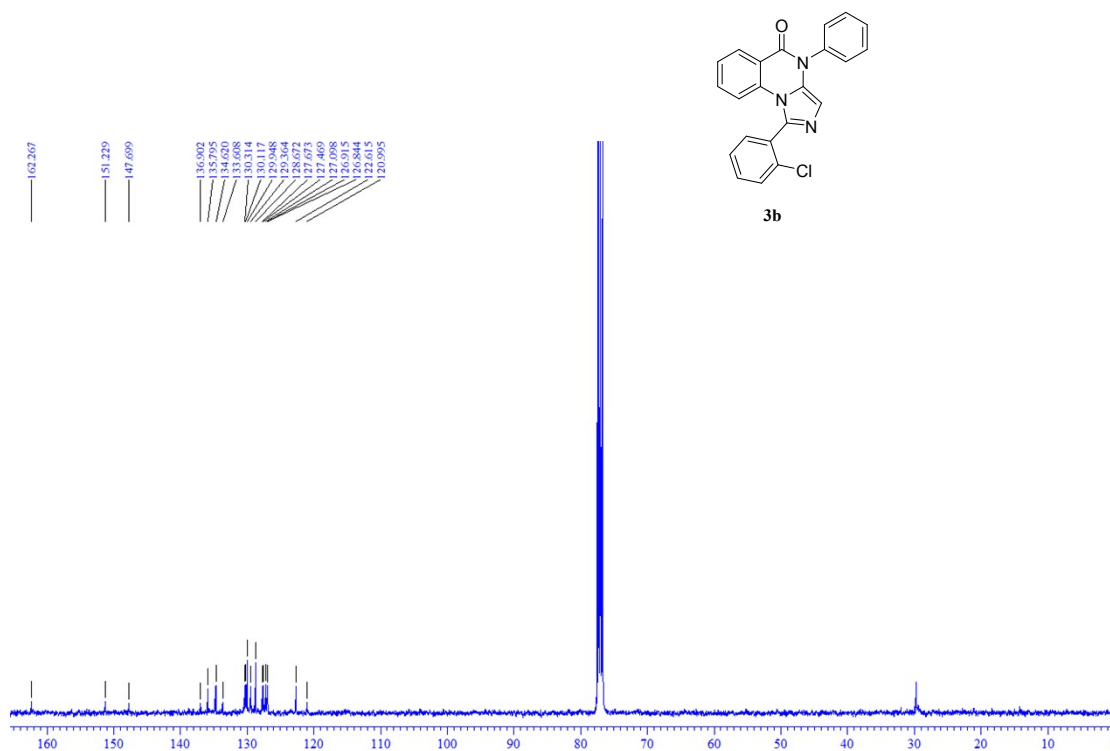
¹³C NMR (100 MHz, CDCl₃) spectrum of compound 3a



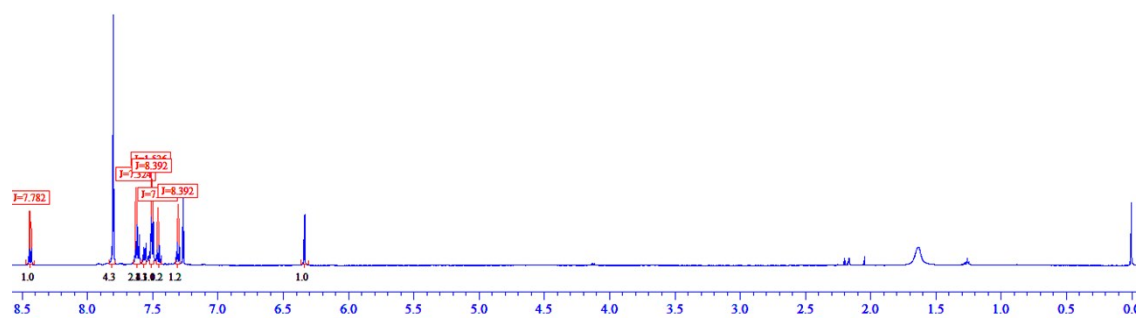
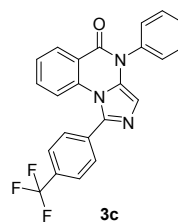
3b



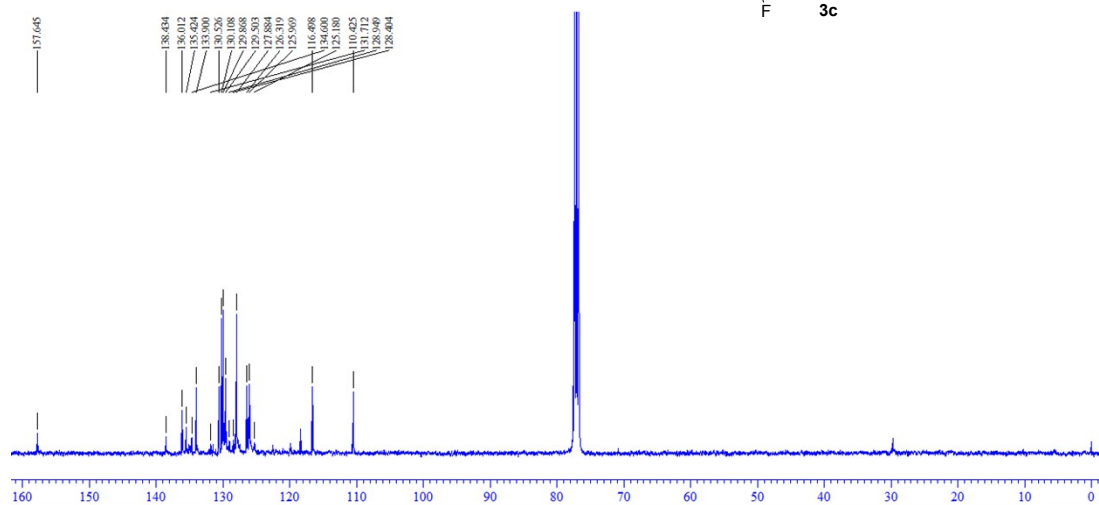
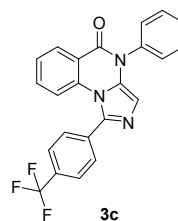
^1H NMR (400 MHz, CDCl_3) spectrum of compound **3b**



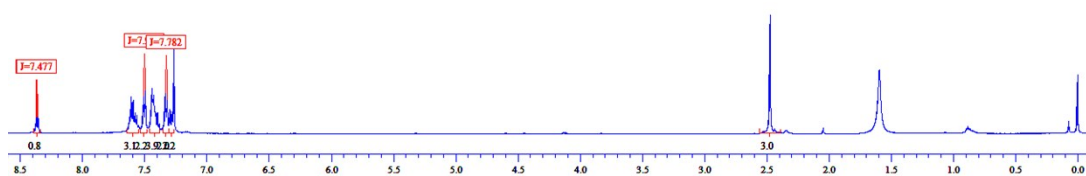
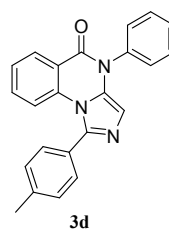
^{13}C NMR (100 MHz, CDCl_3) spectrum of compound **3b**



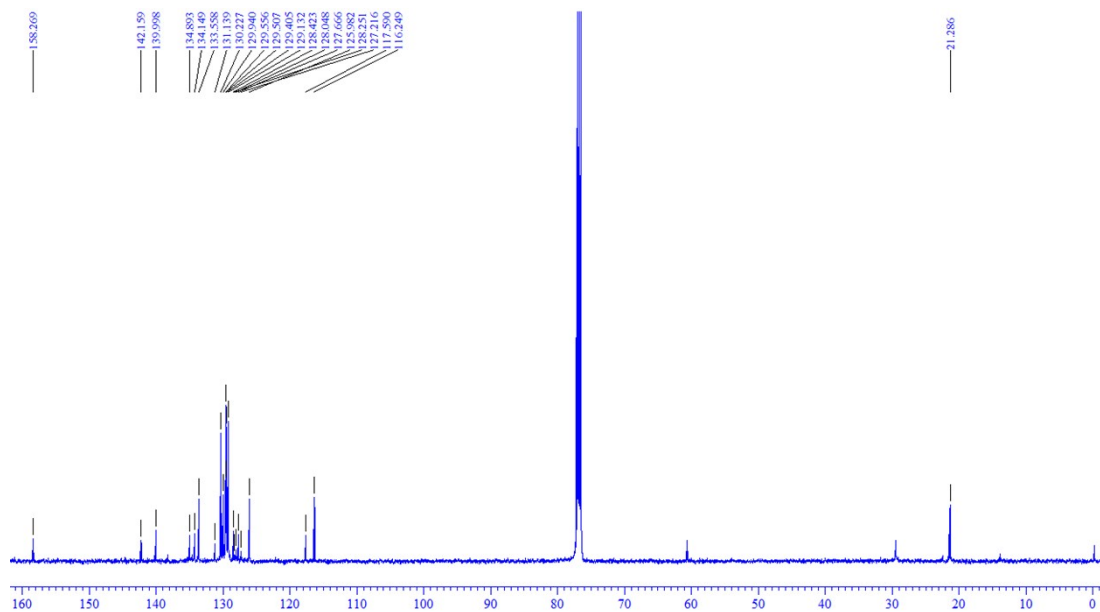
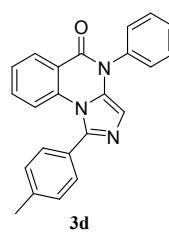
^1H NMR (500 MHz, CDCl_3) spectrum of compound **3c**



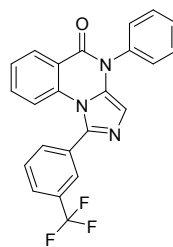
^{13}C NMR (125 MHz, CDCl_3) spectrum of compound **3c**



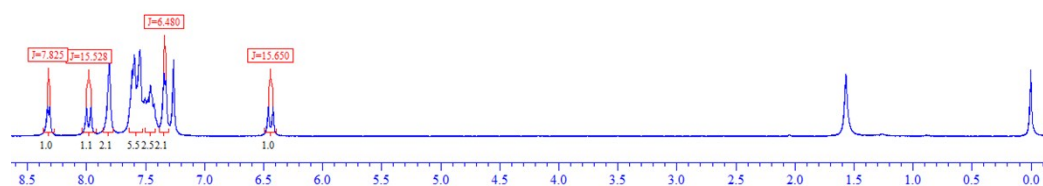
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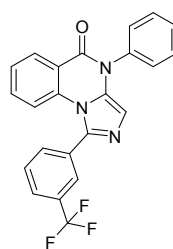
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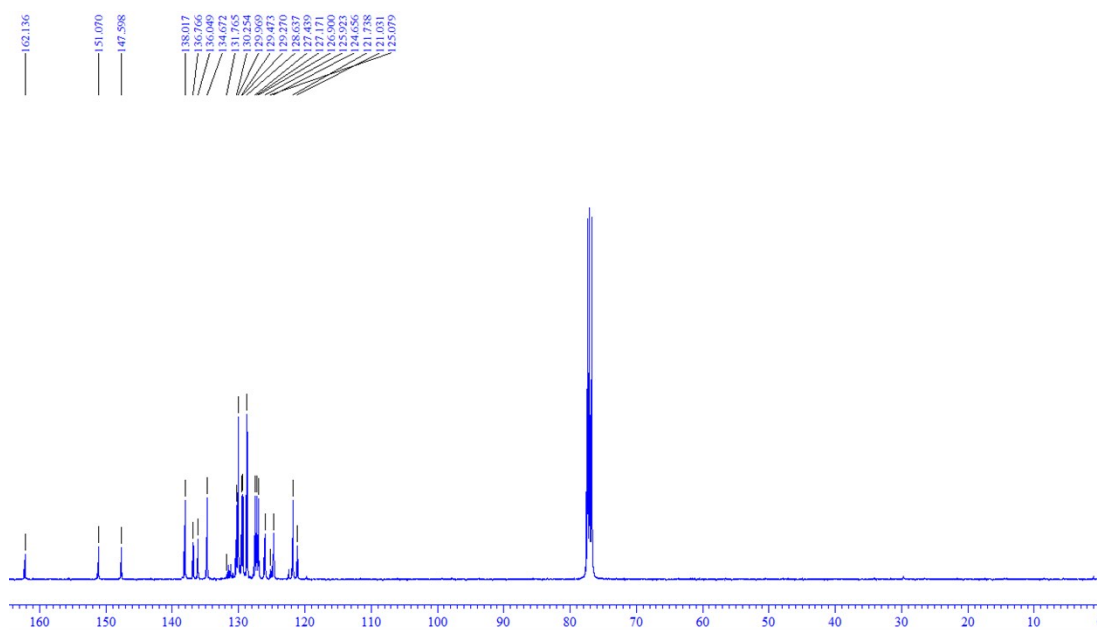
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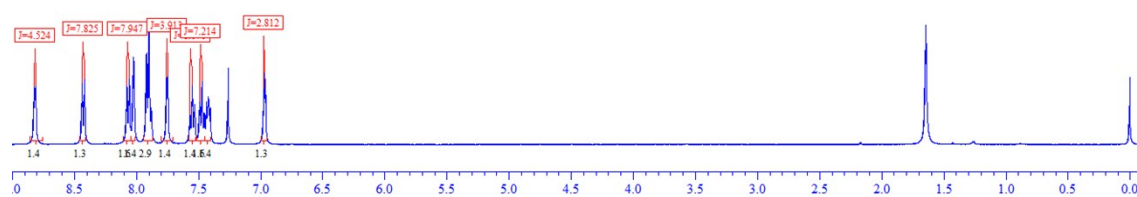
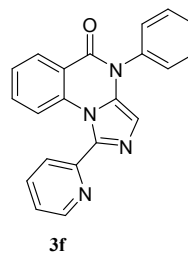
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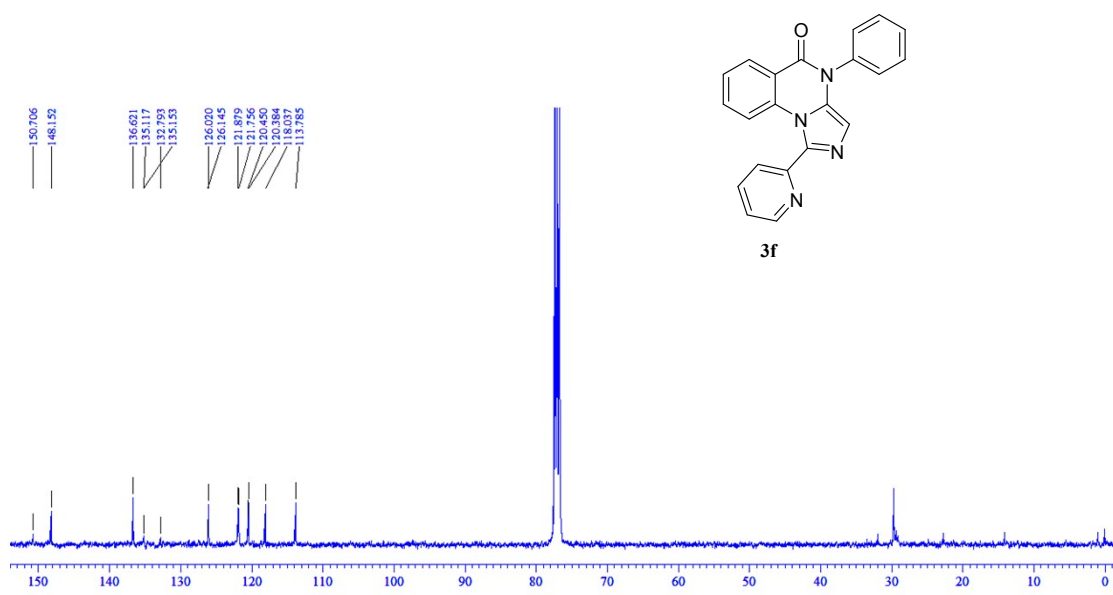
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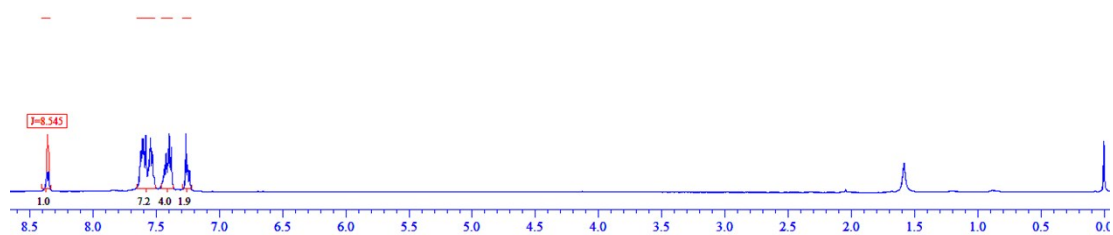
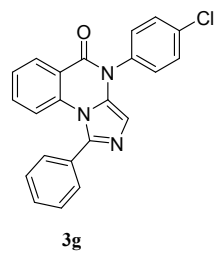
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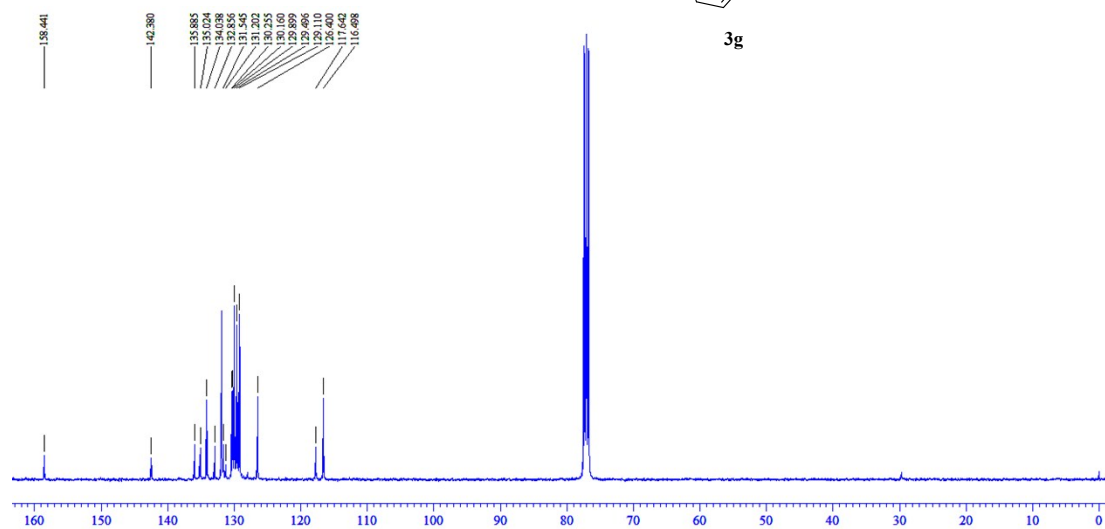
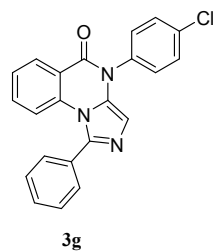
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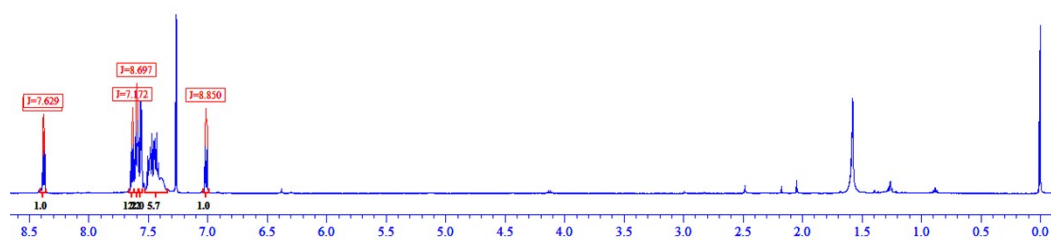
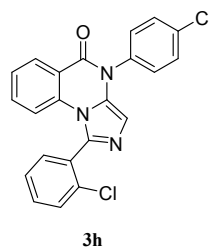
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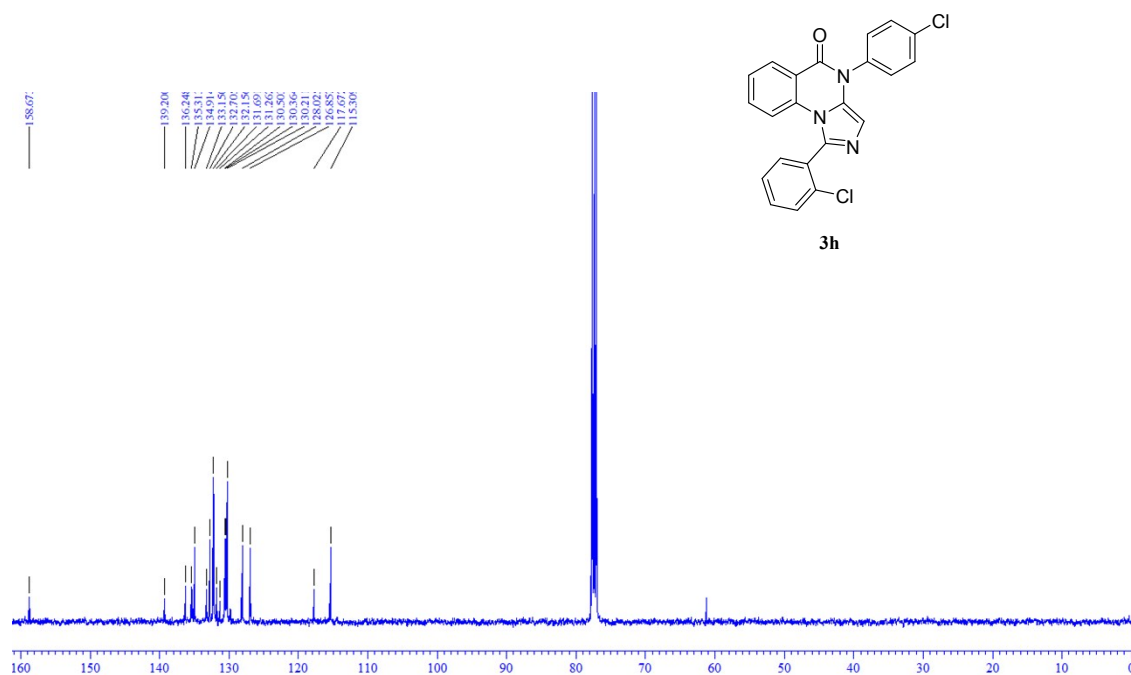
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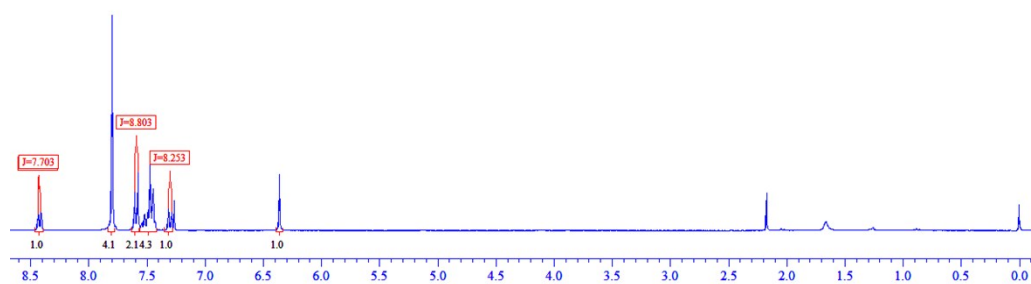
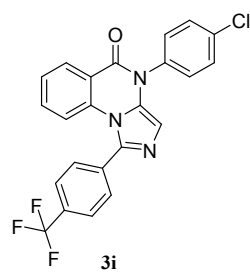
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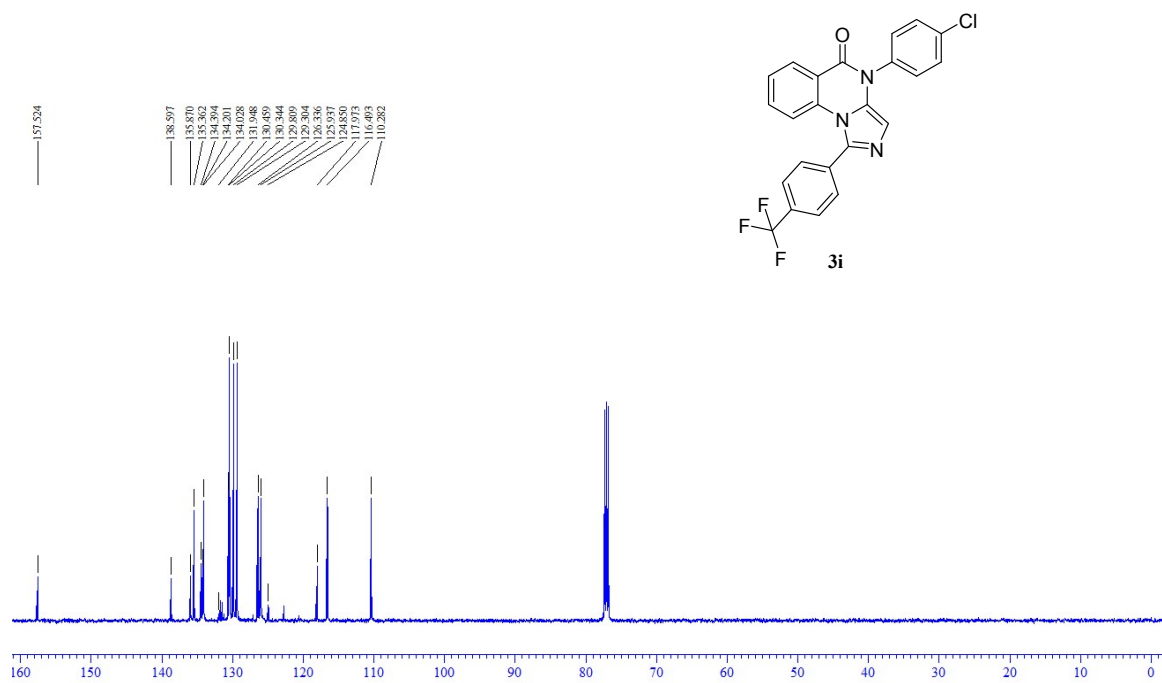
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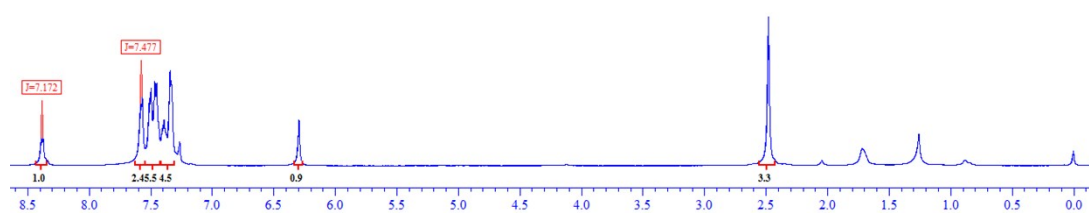
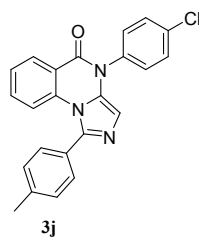
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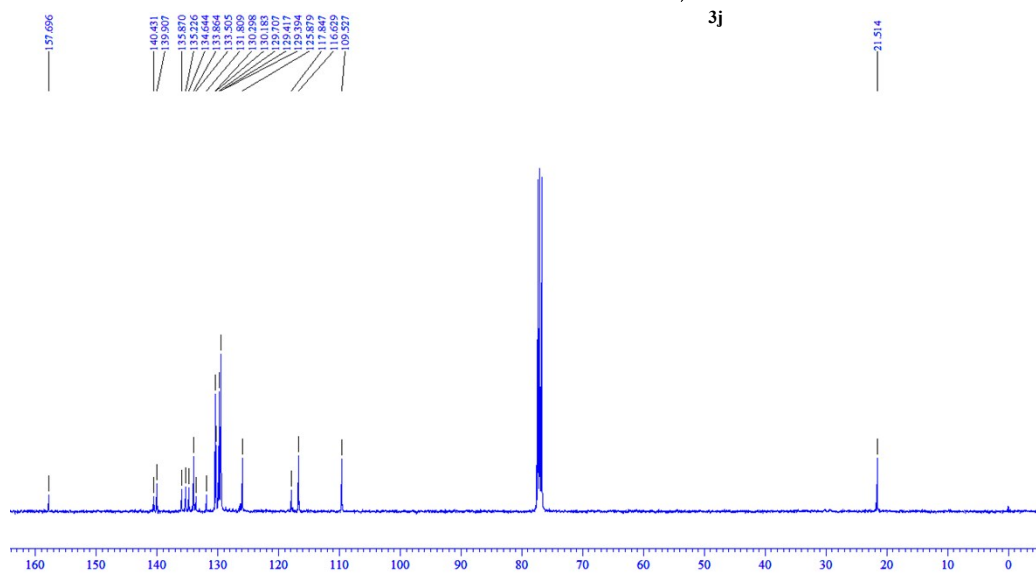
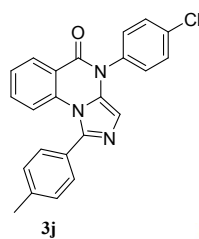
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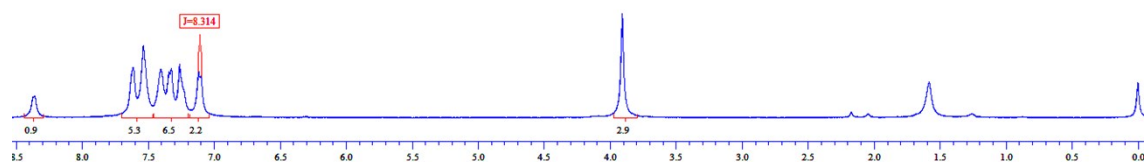
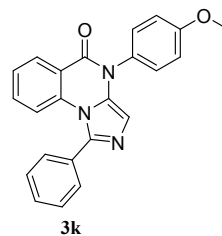
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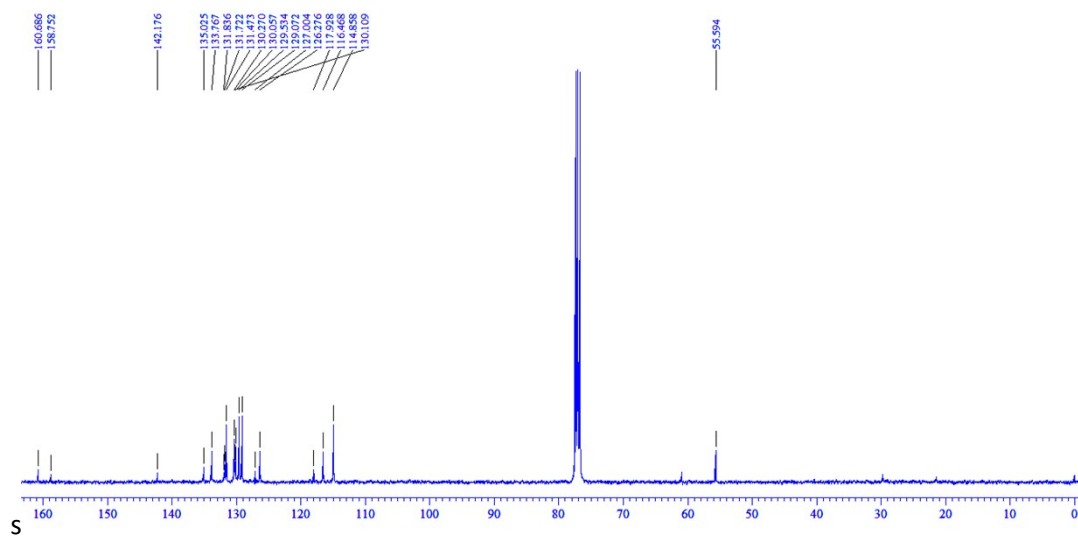
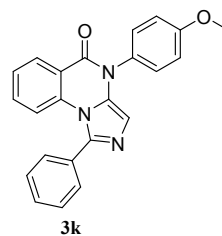
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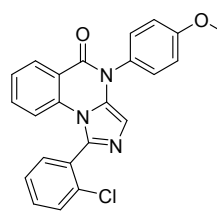
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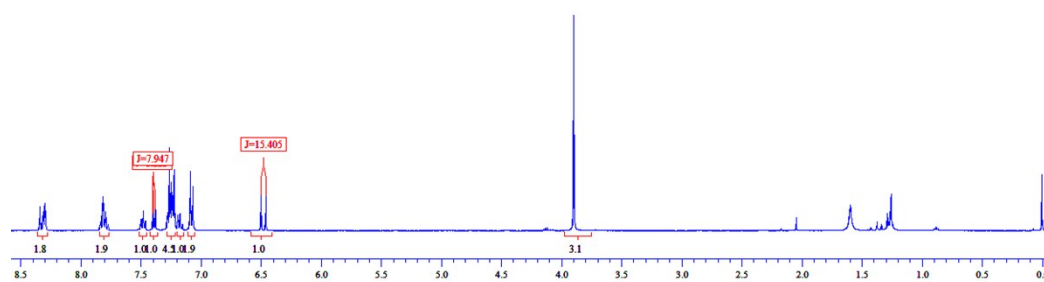
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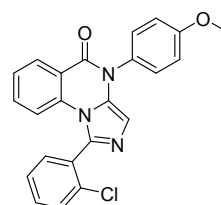
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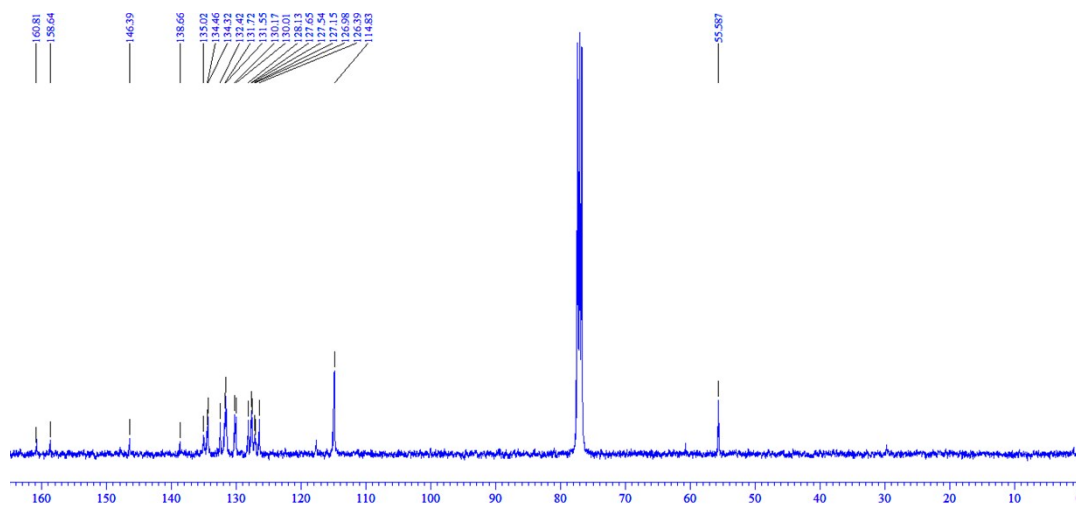
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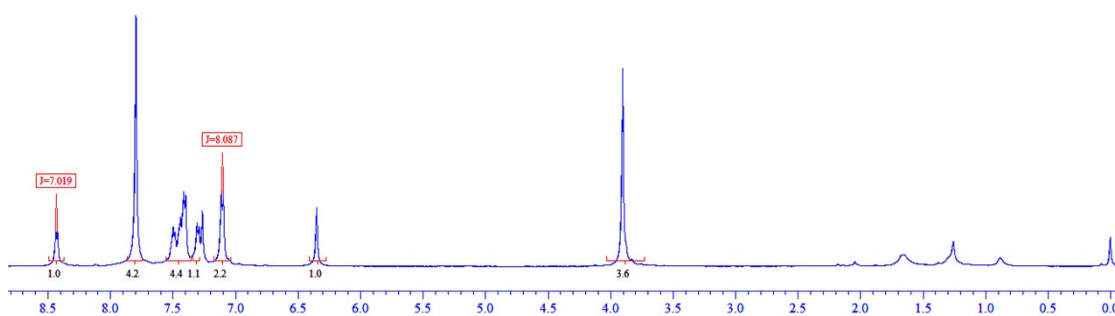
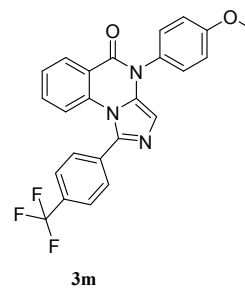
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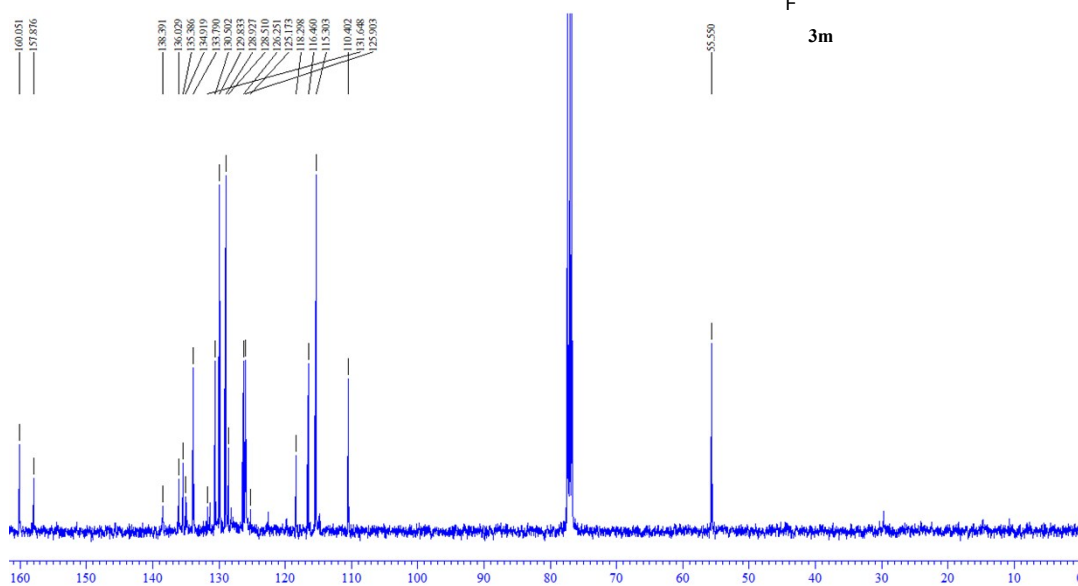
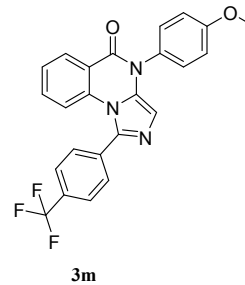
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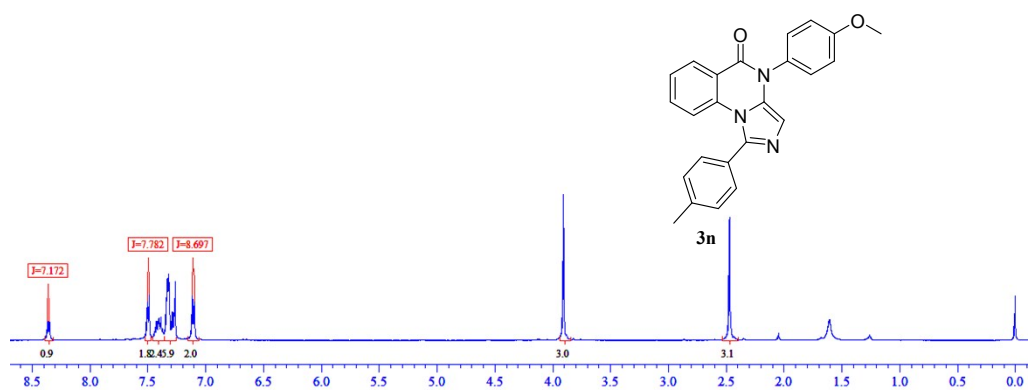
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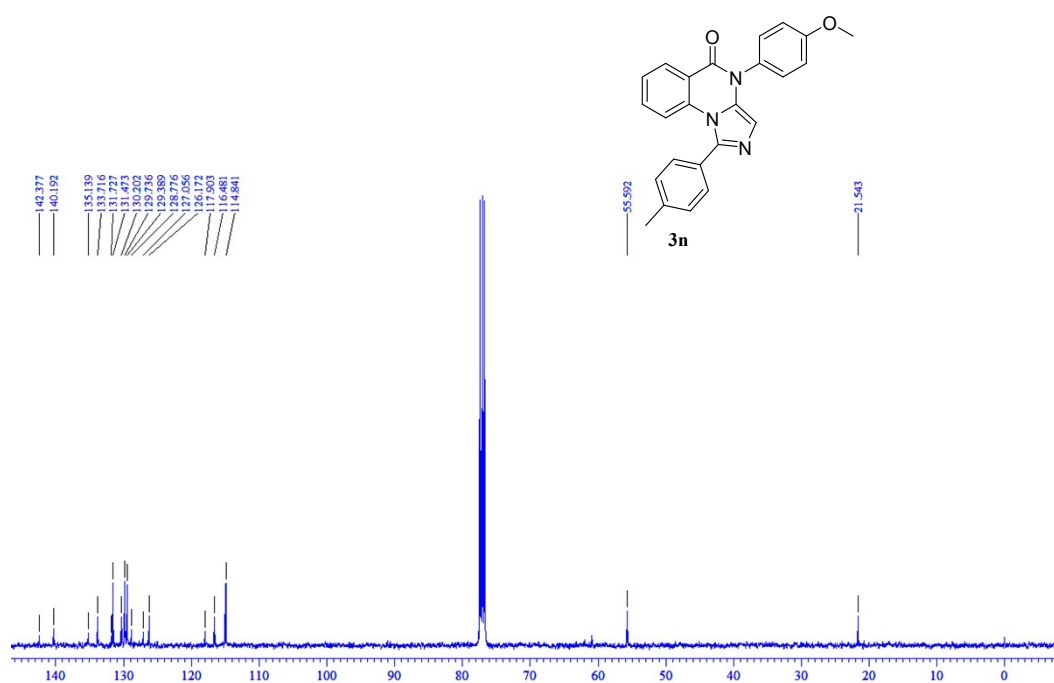
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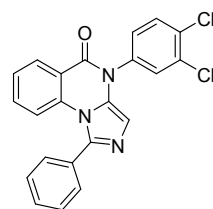
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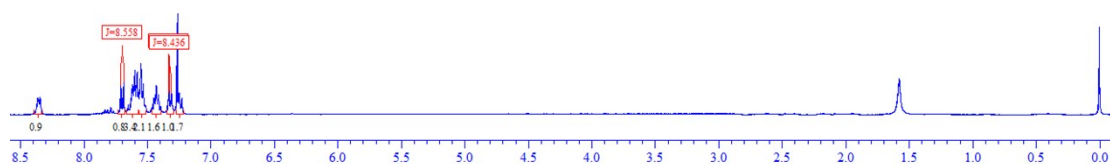
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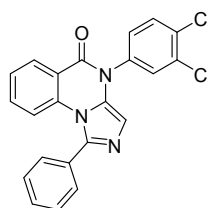
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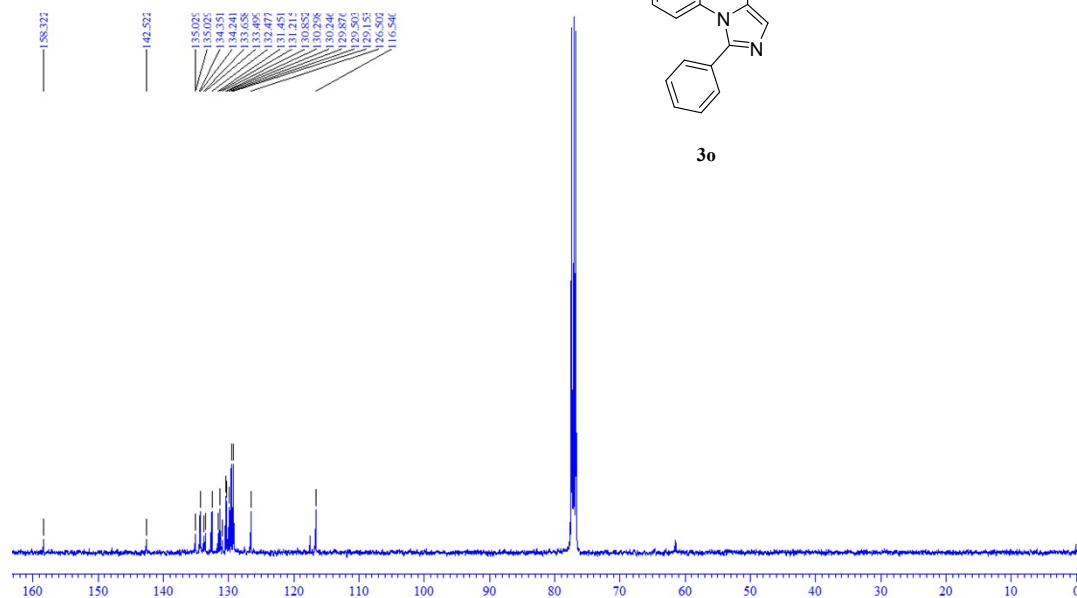
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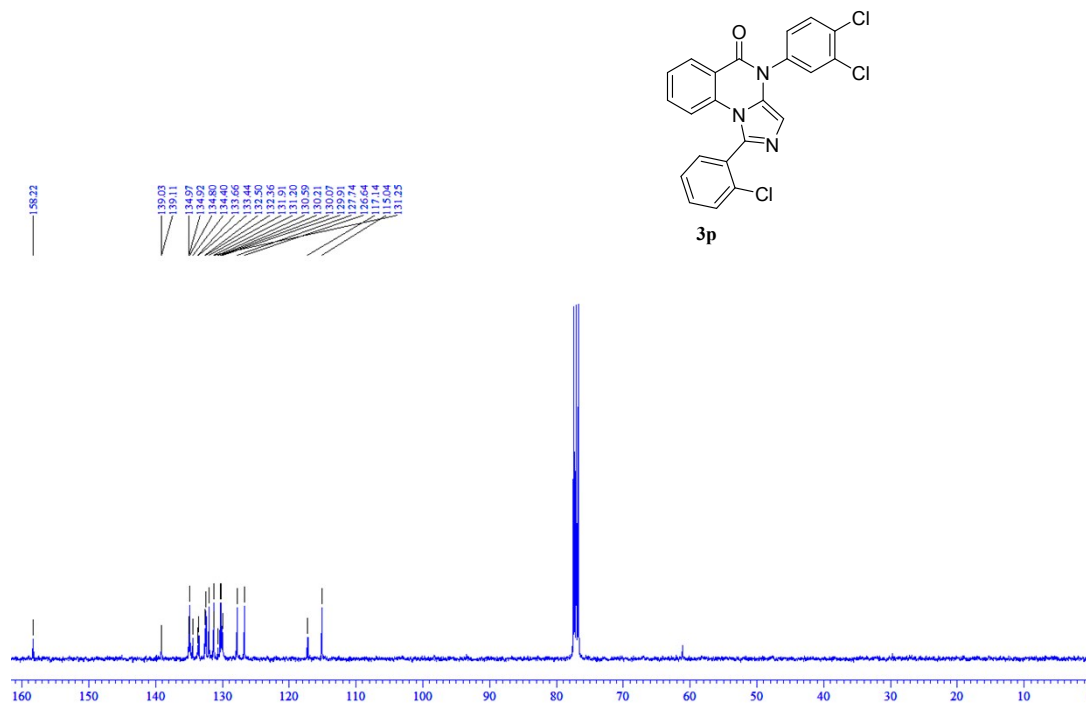
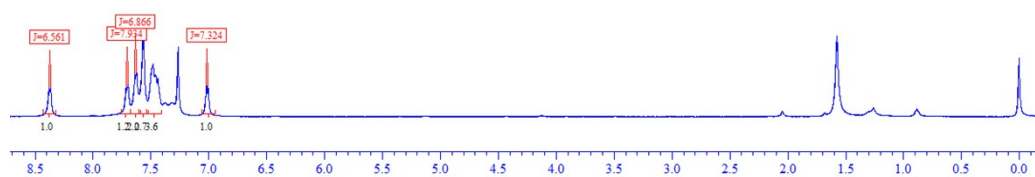
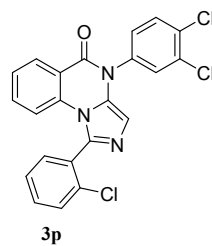
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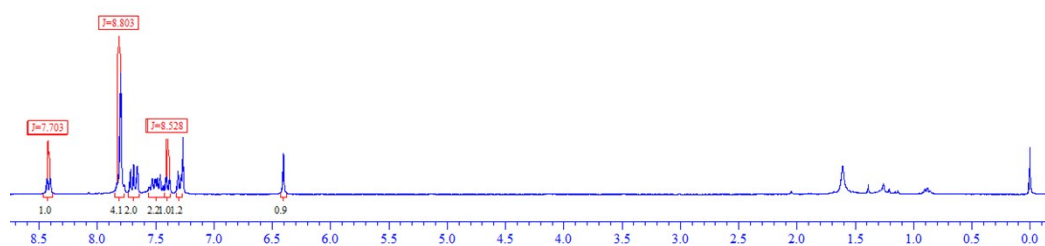
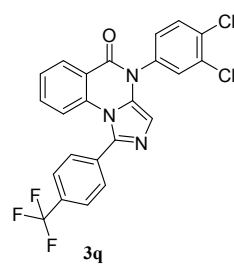


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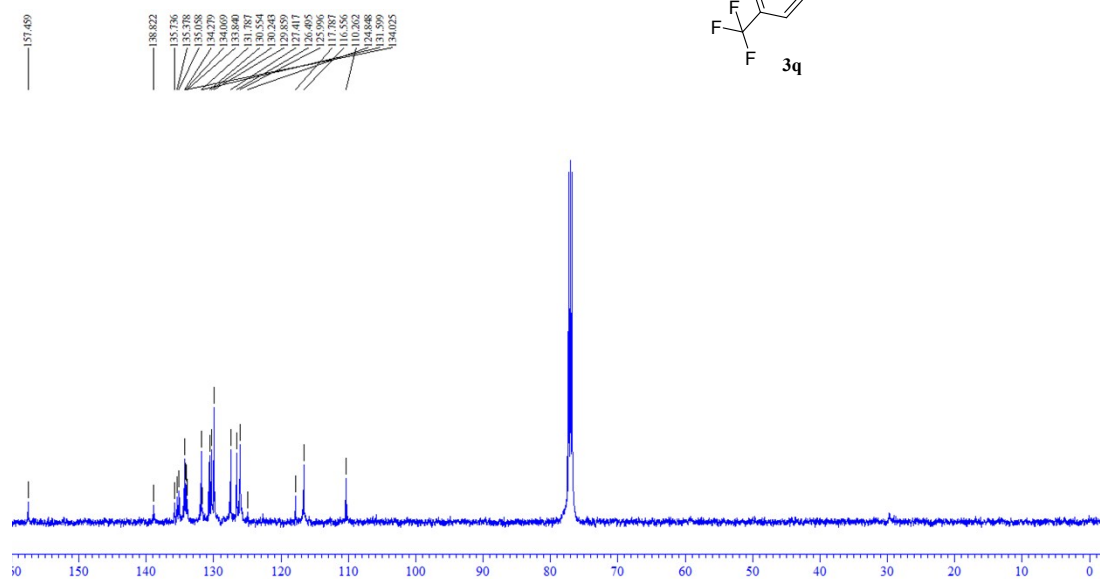
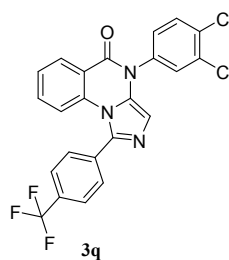


^{13}C NMR (100 MHz, CDCl_3) spectrum of compound **3o**

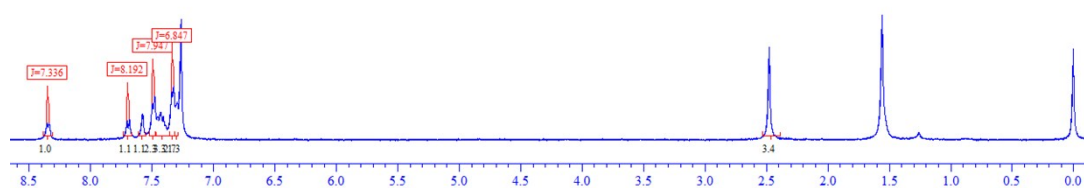
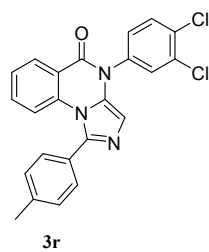




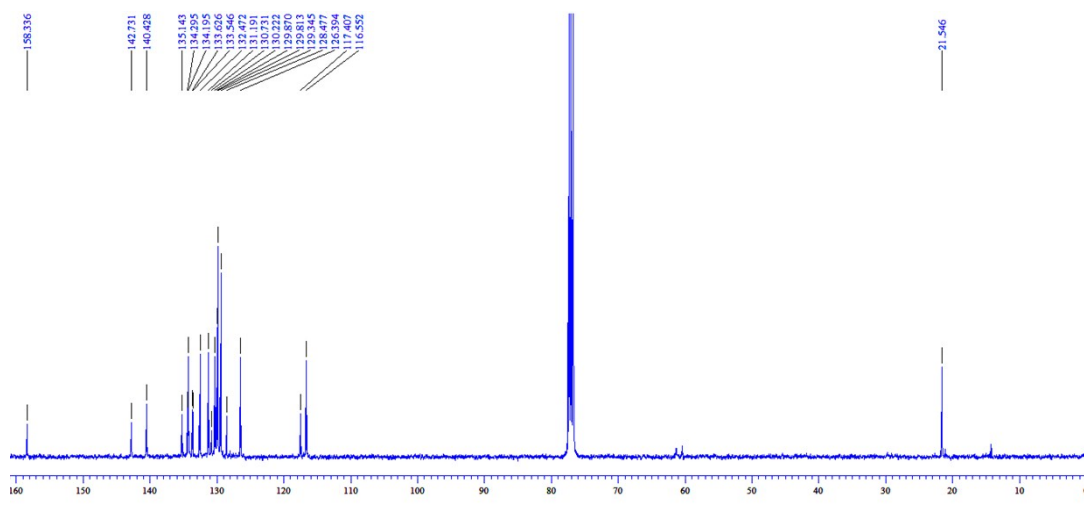
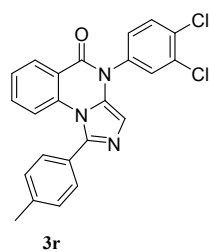
^1H NMR (500 MHz, CDCl_3) spectrum of compound **3q**



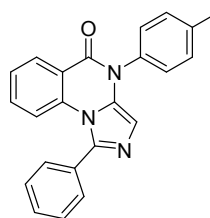
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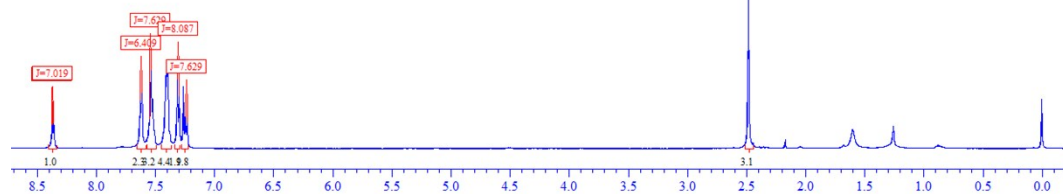
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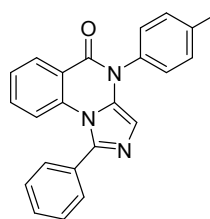
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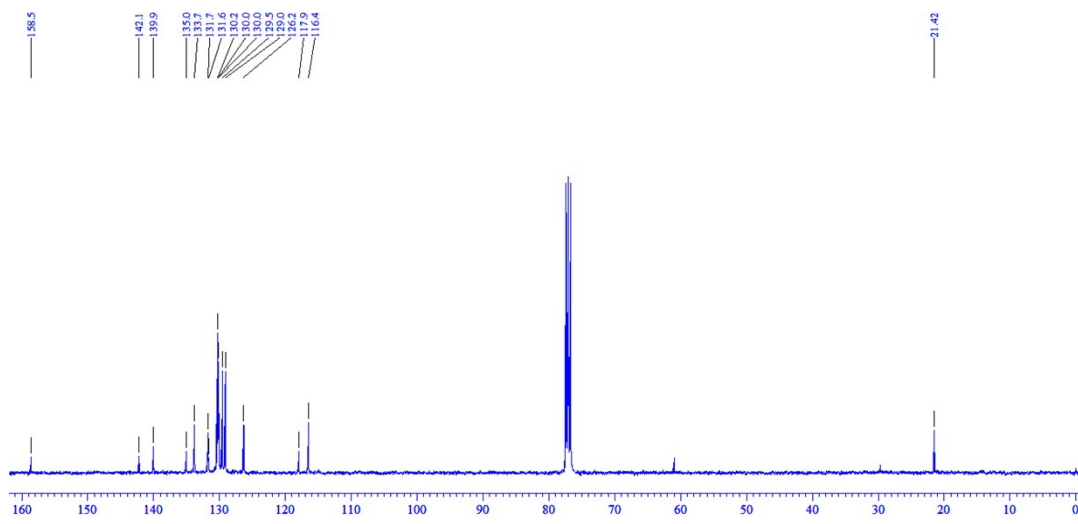
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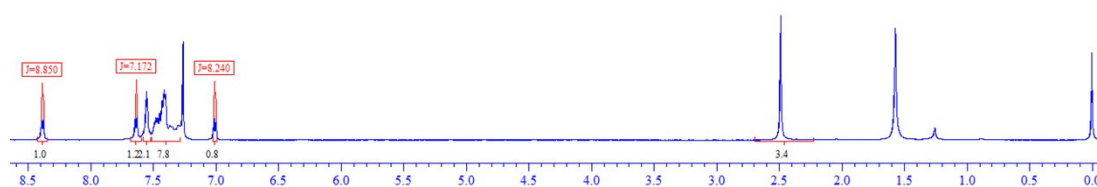
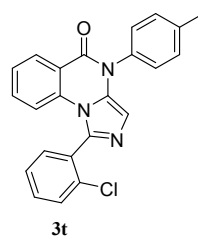
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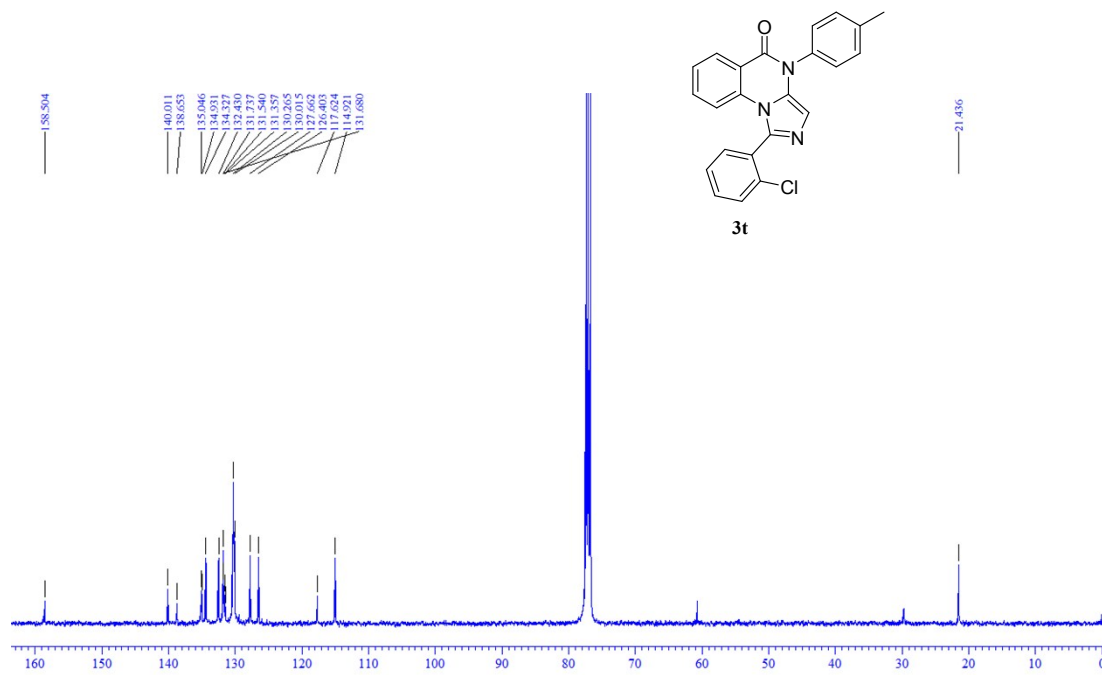
3s



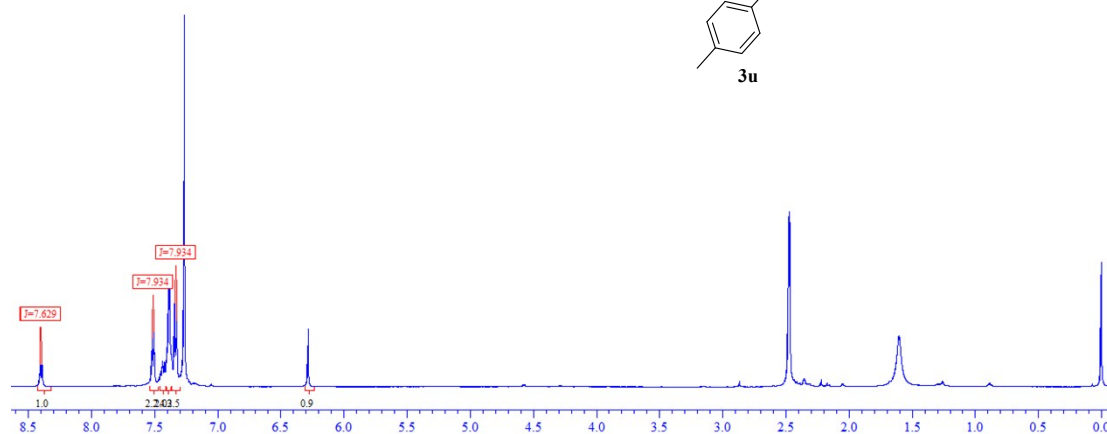
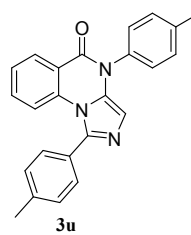
^{13}C NMR (100 MHz, CDCl_3) spectrum of compound 3s



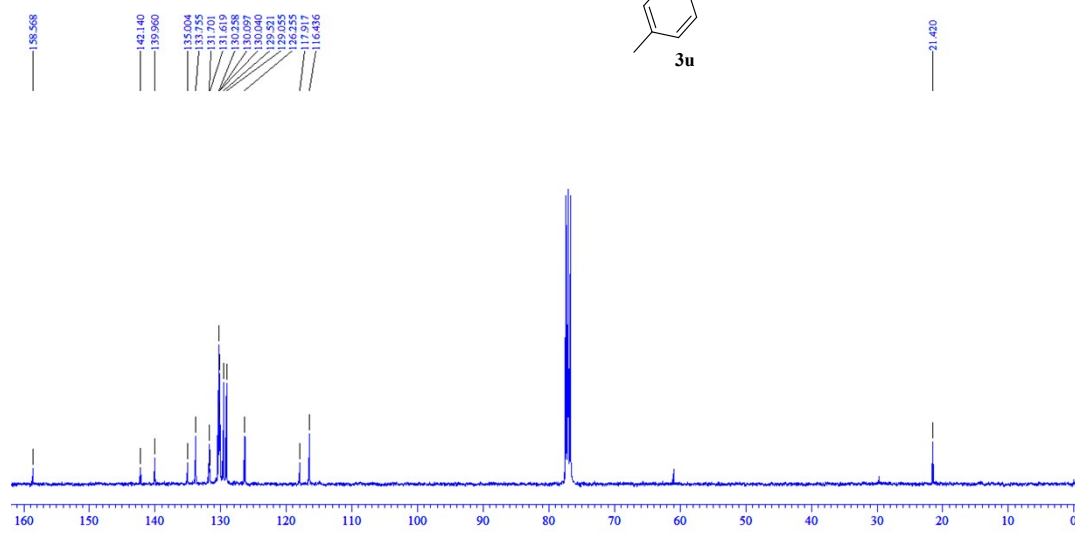
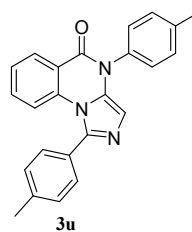
^1H NMR (500 MHz, CDCl_3) spectrum of compound **3t**



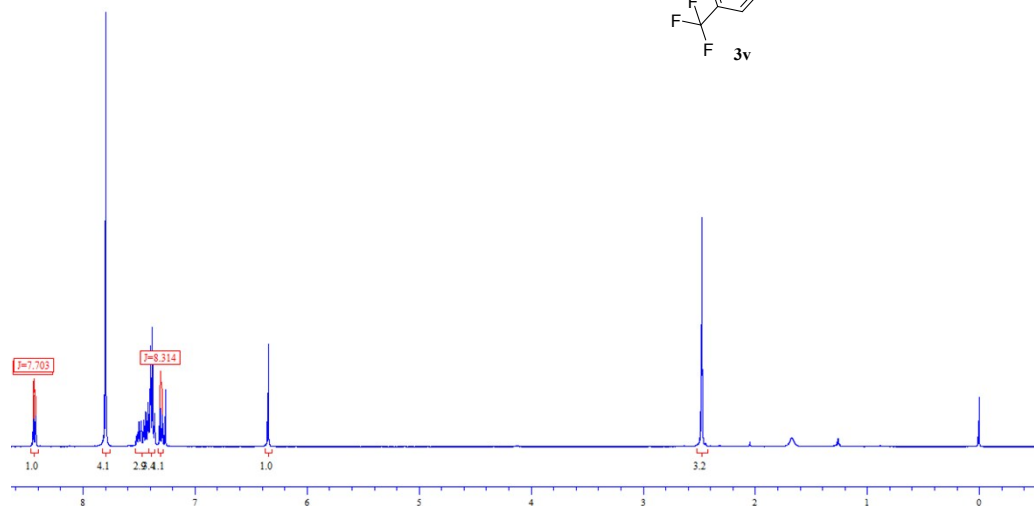
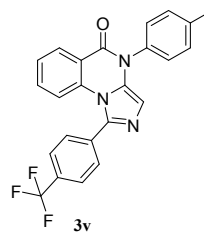
^{13}C NMR (100 MHz, CDCl_3) spectrum of compound **3t**



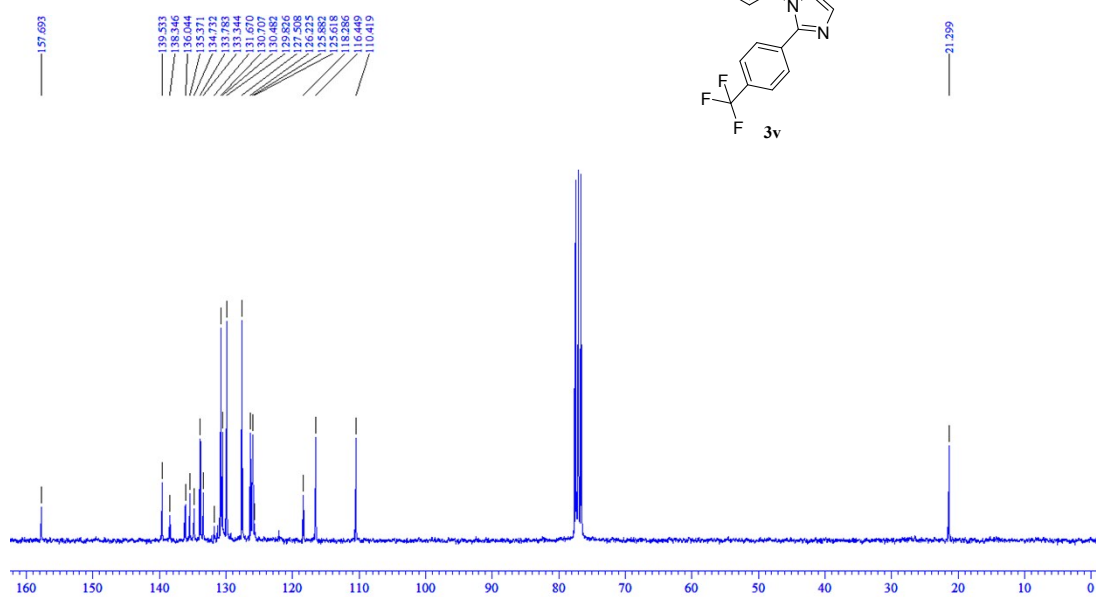
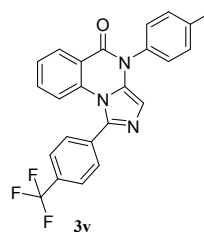
^1H NMR (500 MHz, CDCl_3) spectrum of compound **3u**



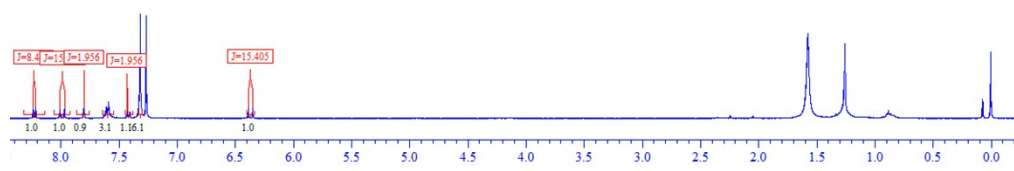
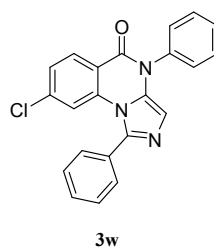
^{13}C NMR (100 MHz, CDCl_3) spectrum of compound **3u**



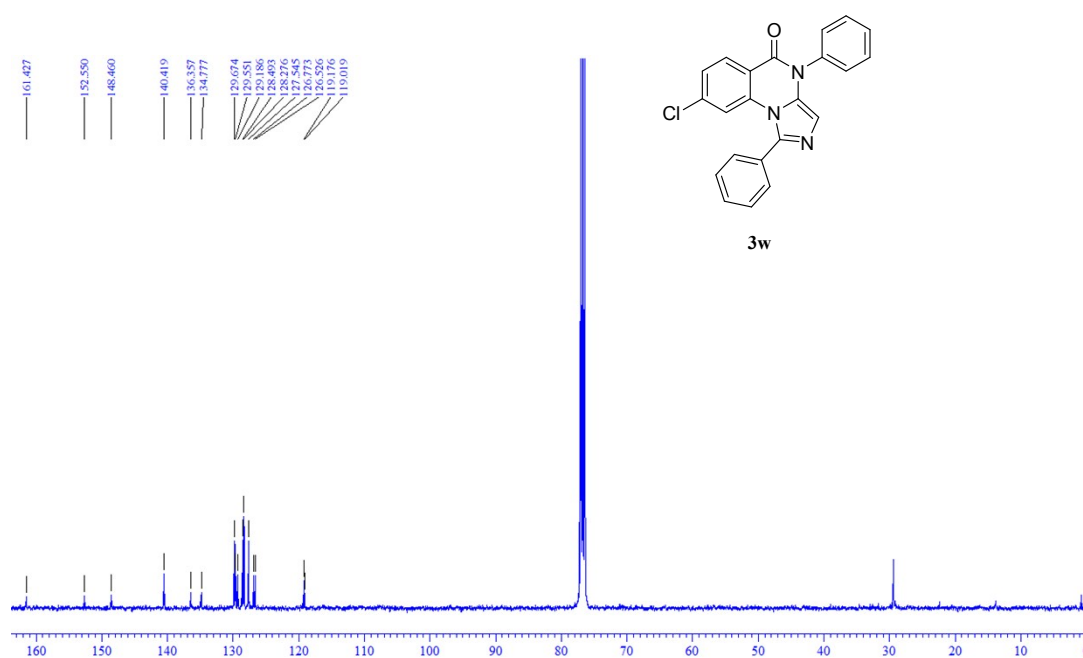
^1H NMR (500 MHz, CDCl_3) spectrum of compound 3v



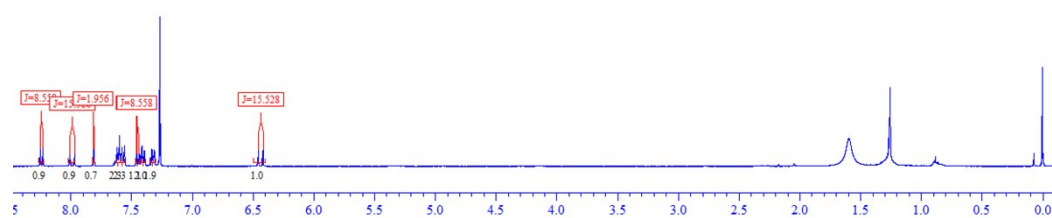
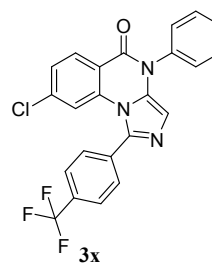
^{13}C NMR (100 MHz, CDCl_3) spectrum of compound 3v



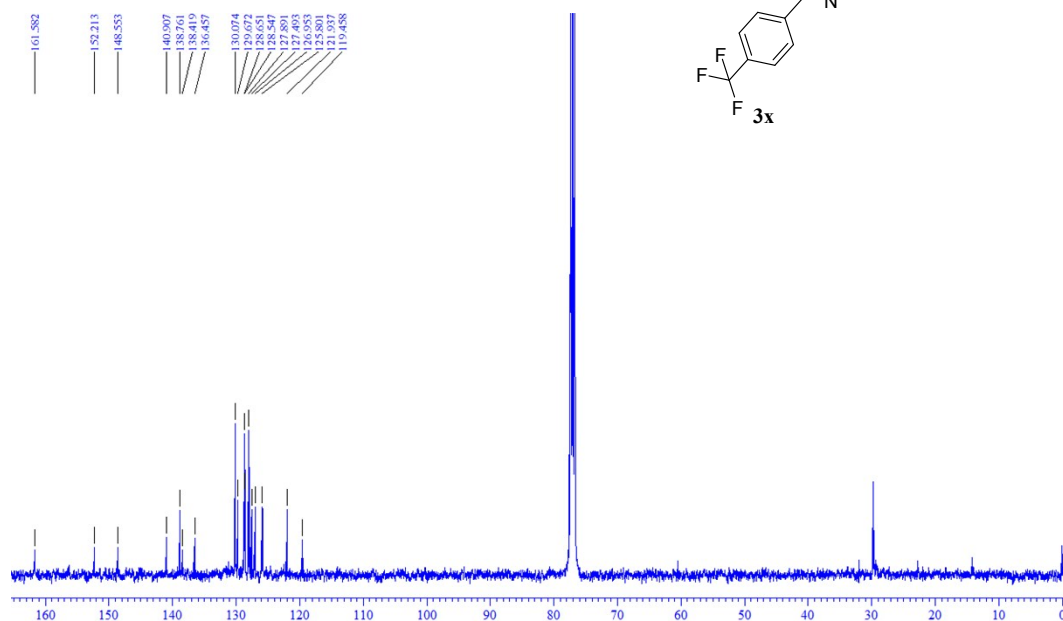
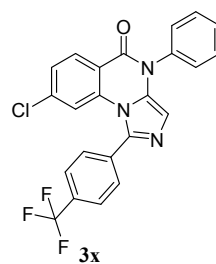
^1H NMR (500 MHz, CDCl_3) spectrum of compound **3w**



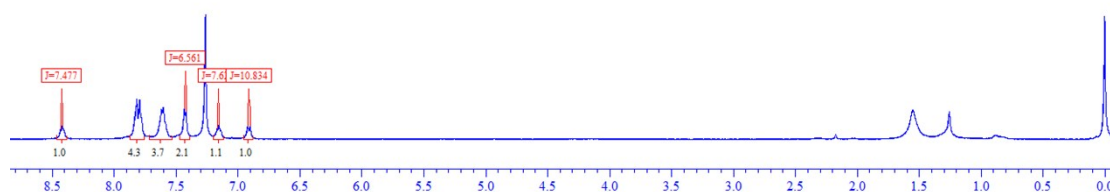
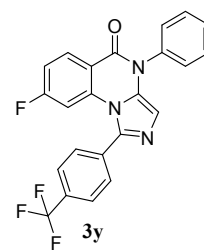
^{13}C NMR (100 MHz, CDCl_3) spectrum of compound **3w**



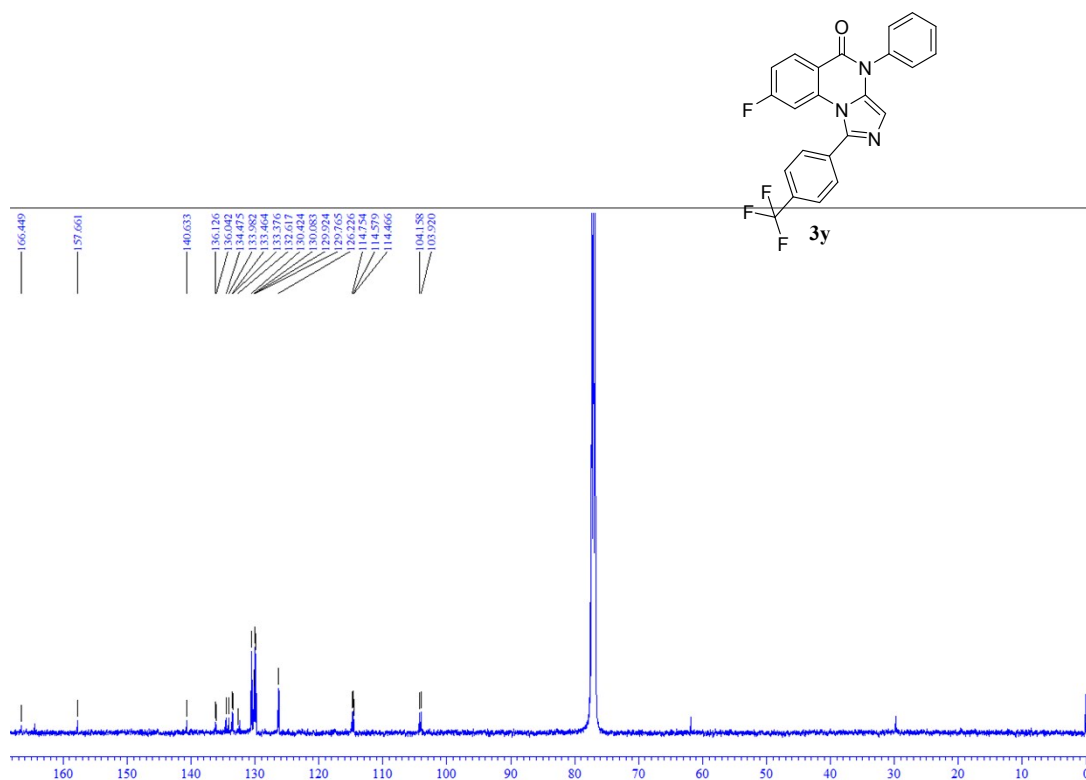
¹H NMR (500 MHz, CDCl₃) spectrum of compound 3x



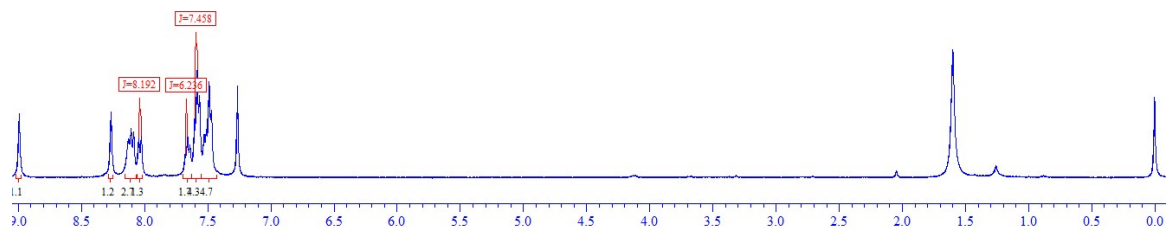
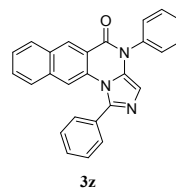
¹³C NMR (100 MHz, CDCl₃) spectrum of compound 3x



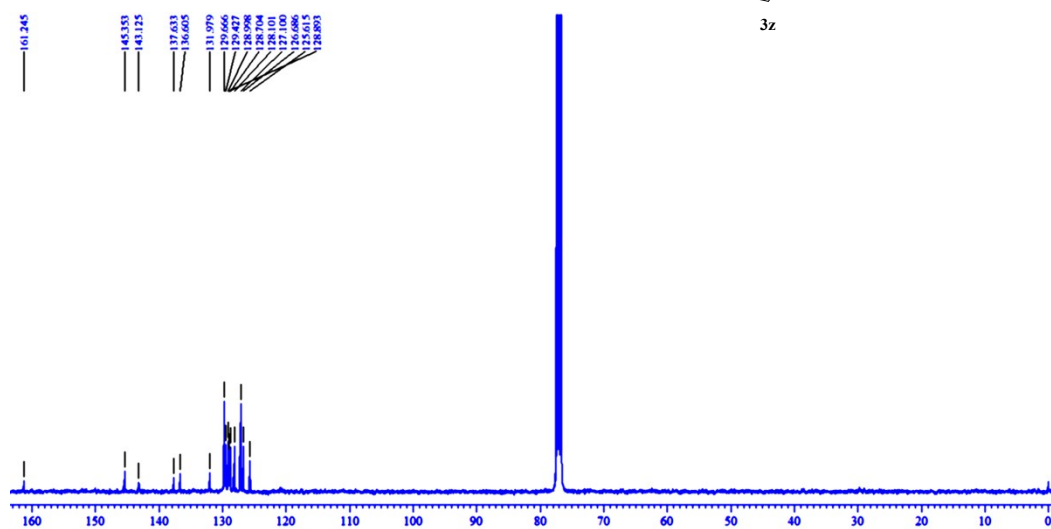
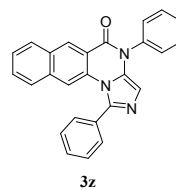
^1H NMR (500 MHz, CDCl_3) spectrum of compound **3y**



^{13}C NMR (100 MHz, CDCl_3) spectrum of compound **3y**



^1H NMR (500 MHz, CDCl_3) spectrum of compound 3z



^{13}C NMR (100 MHz, CDCl_3) spectrum of compound 3z