

Supporting Information

A regioselective synthesis of 3,4-diaryl-1*H*-pyrazoles through a 1,3-dipolar cycloaddition of tosylhydrazones and nitroalkenes.

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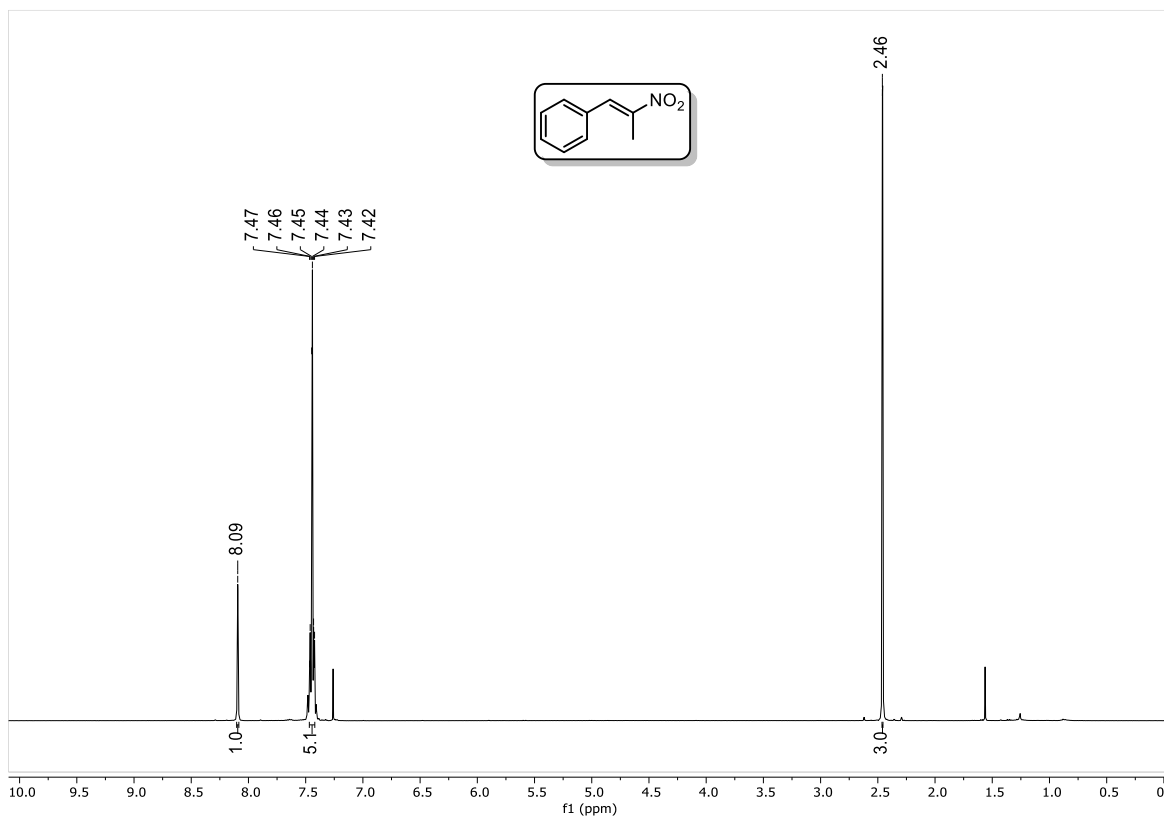
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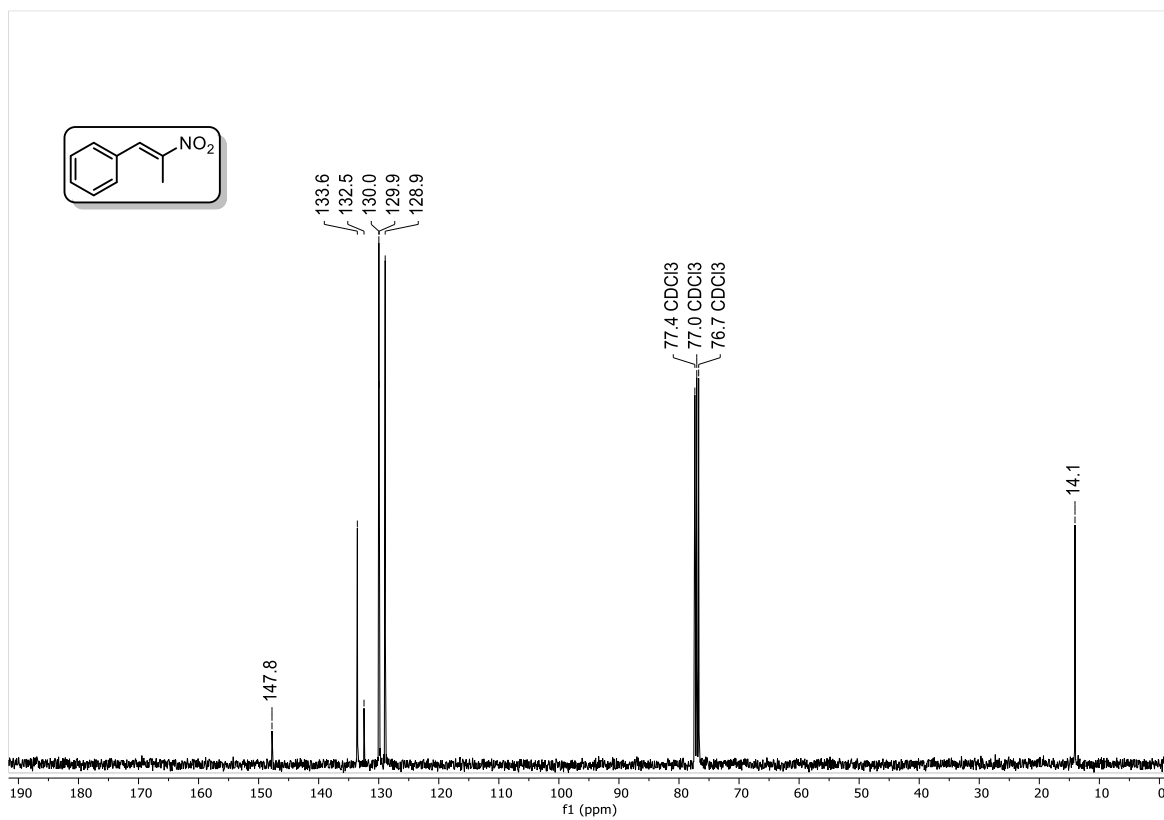
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^1H -NMR and ^{13}C -NMR and HRMS spectra of the
nitroalkenes.



Spectrum 1. ¹H-NMR of 6a.



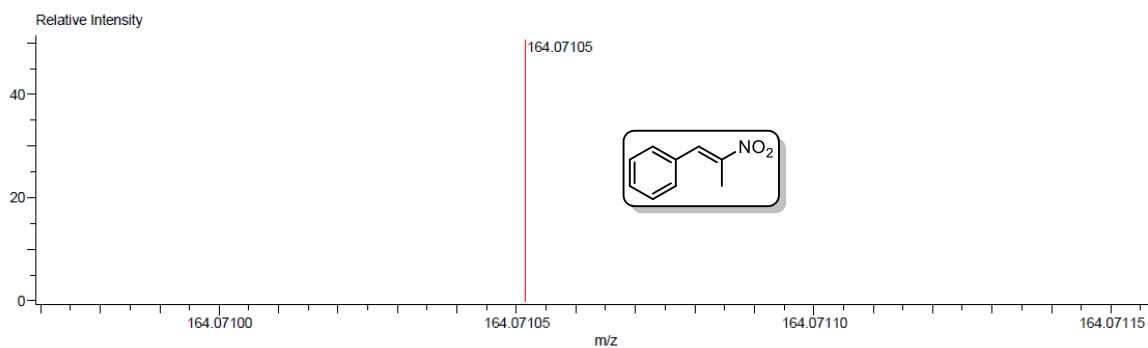
Spectrum 2. ¹³C-NMR of 6a.

Description:
Ionization Mode:ESI+
History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:CaI_PEG_600
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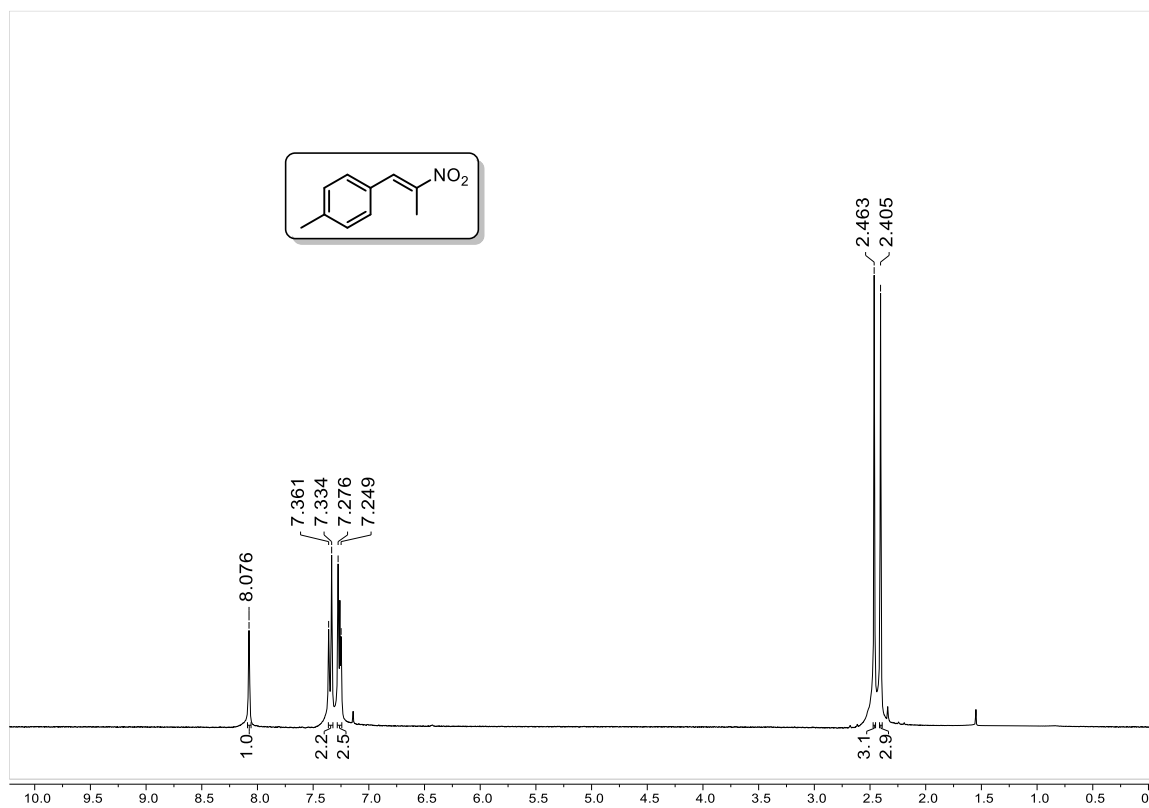
Charge number:1
Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)
Element:¹²C:0 .. 9, ¹H:0 .. 50, ³⁵Cl:0 .. 1, ¹⁴N:0 .. 1, ¹⁶O:0 .. 2, ³²S:0 .. 0

Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

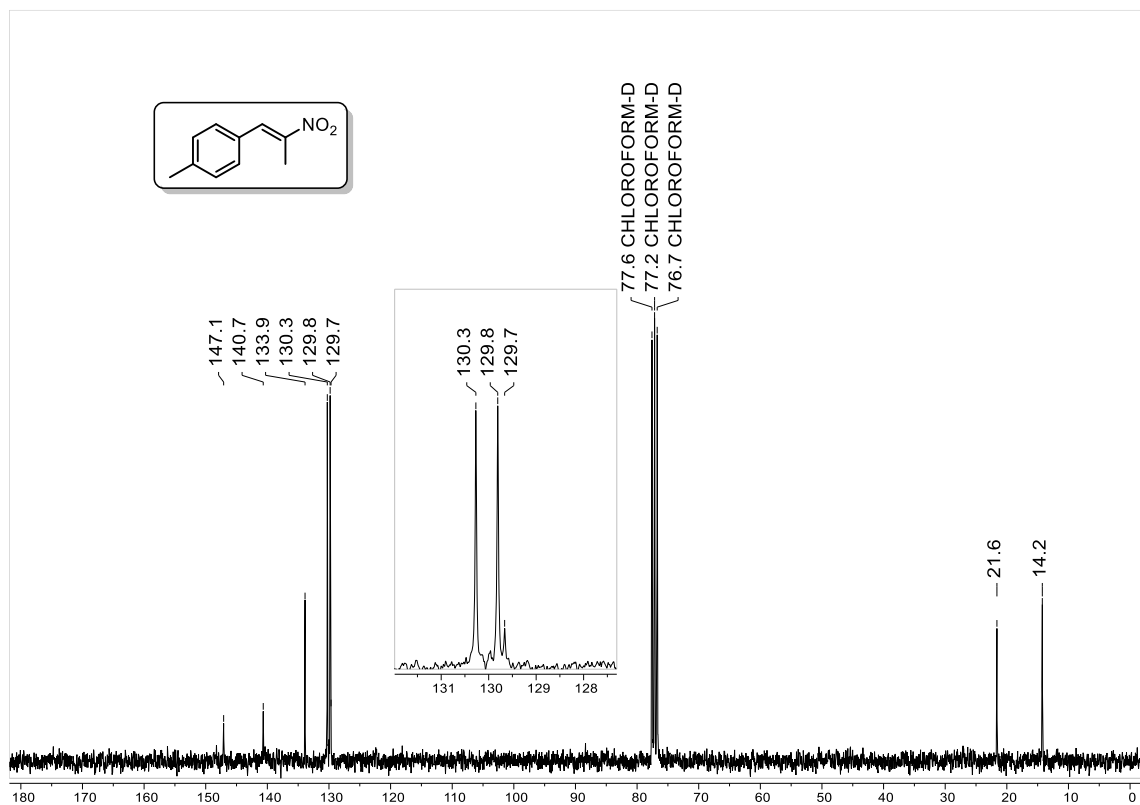


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
164.07105	11180.34	164.07115	-0.10	-0.62	¹² C ₉ ¹ H ₁₀ ¹⁴ N ₁ ¹⁶ O ₂	5.5

Spectrum 3. HRMS of 6a.



Spectrum 4. ¹H-NMR of 6b.



Spectrum 5. ¹³C-NMR of 6b.

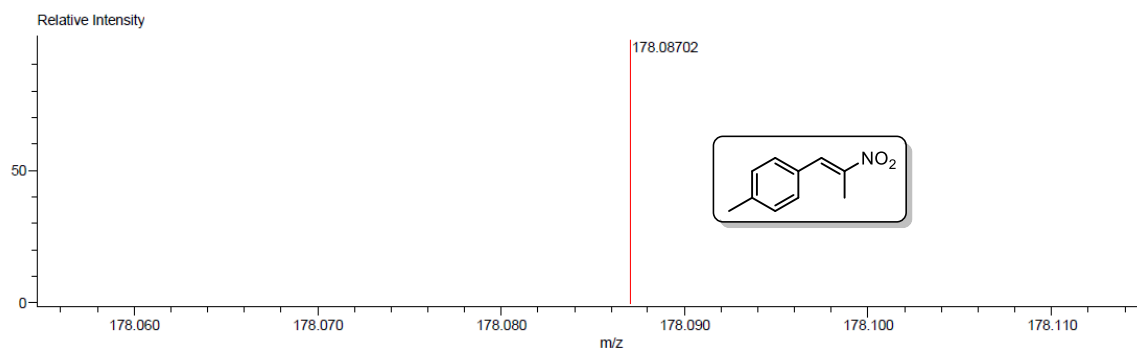
Description: Ionization Mode: ESI+
 History: Determine m/z [Peak Detect [Centroid, 30, Area]; Correct Base[]; Smooth [5]]; Correct Base [5.0%]; Average (MS [...])

Mass Calibration data: Cal_PEG_600
 Created: 6/1/2023 10:08:27 AM
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Charge number: 1
 Element: ¹²C: 0 .. 22, ¹H: 0 .. 28, ³⁵Cl: 0 .. 0, ¹⁹F: 0 .. 0, ¹⁴N: 0 .. 1, ¹⁶O: 0 .. 2, ³²S: 0 .. 0

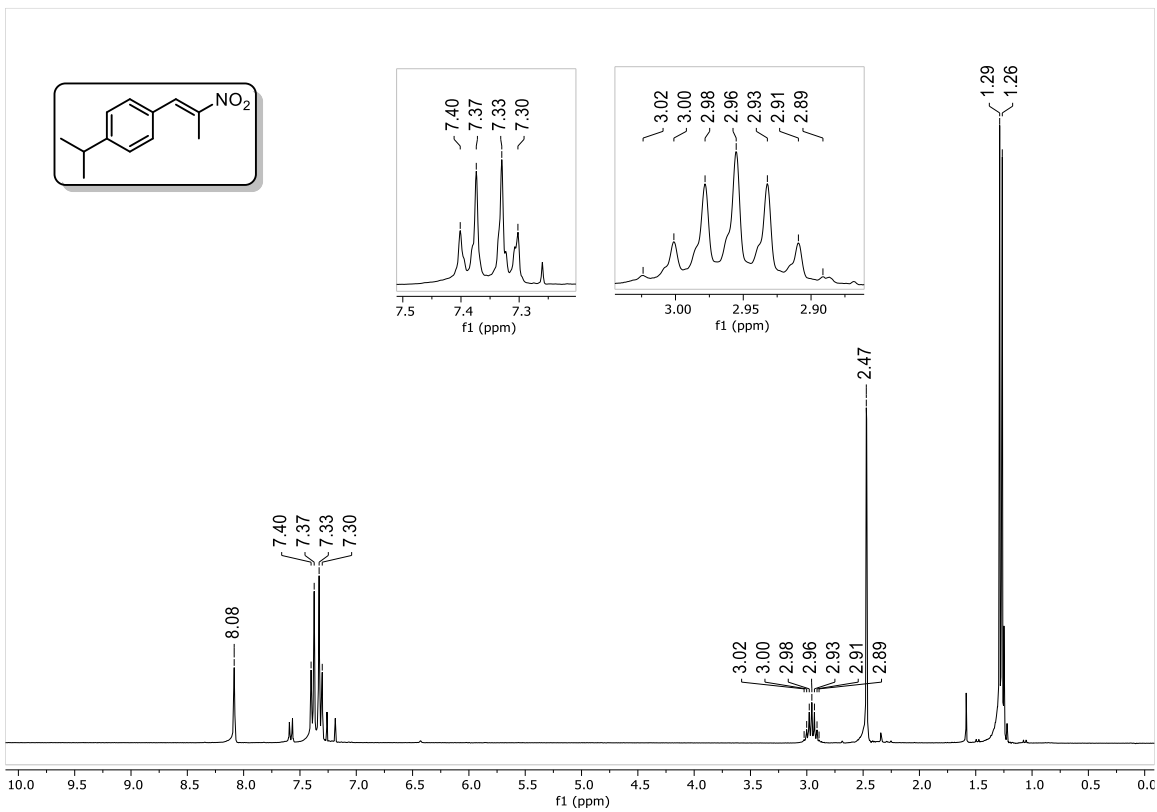
Tolerance: 5.00 (ppm), 5.00 .. 15.00 (mmu)

Unsaturation Number: -1.0 .. 60.0 (Fraction: Both)

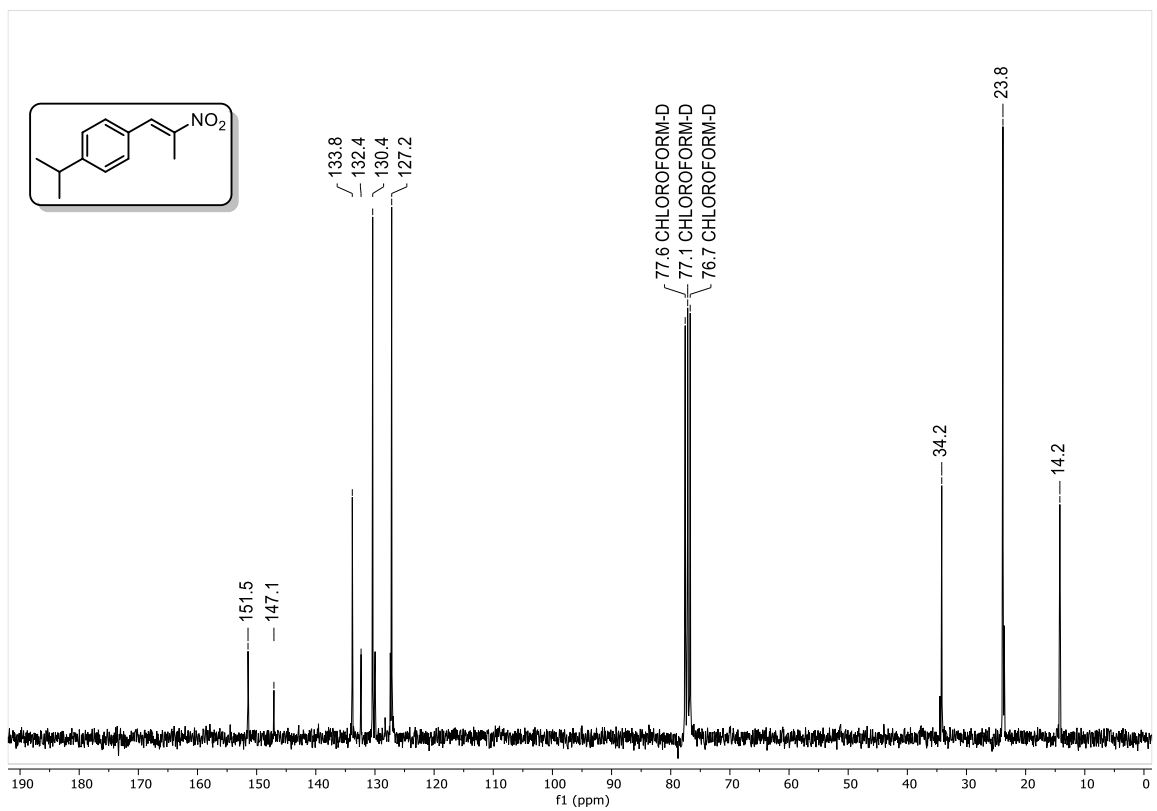


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
178.08702	20008.16	178.08680	0.22	1.23	¹² C ₁₀ ¹ H ₁₂ ¹⁴ N ₁ ¹⁶ O ₂	5.5

Spectrum 6. HRMS of 6b.



Spectrum 7. ¹H-NMR of 6c.



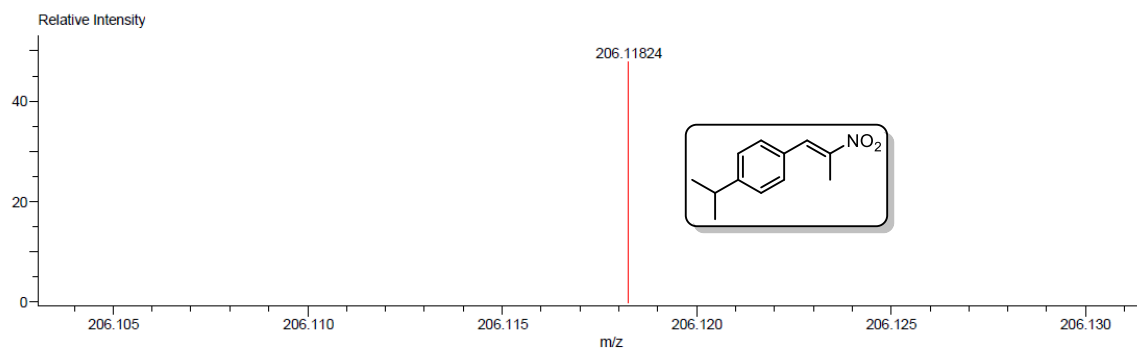
Spectrum 8. ¹³C-NMR of 6c.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:Cal_PEG_600
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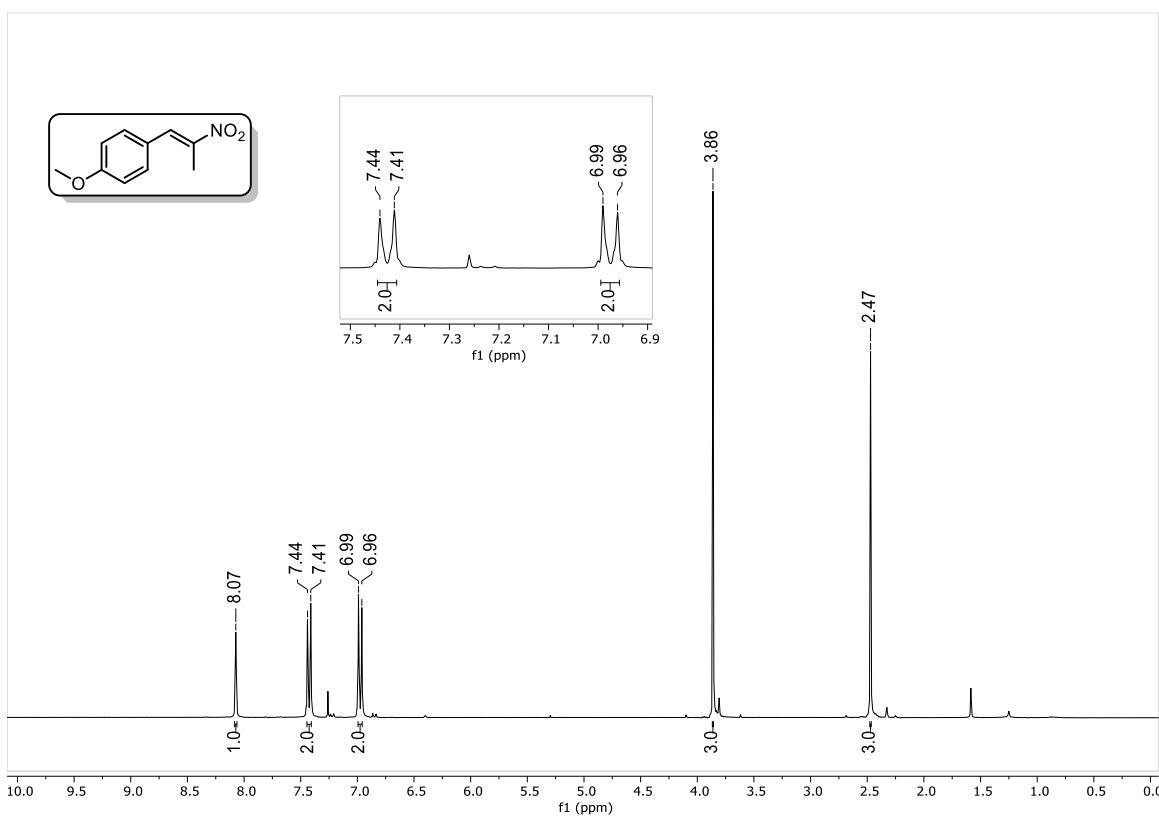
Charge number:1
 Tolerance:5.00(ppm), 5.00 .. 15.00(mmu)
 Element:¹²C:0 .. 22, ¹H:0 .. 28, ³⁵Cl:0 .. 0, ¹⁹F:0 .. 0, ¹⁴N:0 .. 1, ¹⁶O:0 .. 2, ³²S:0 .. 0

Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

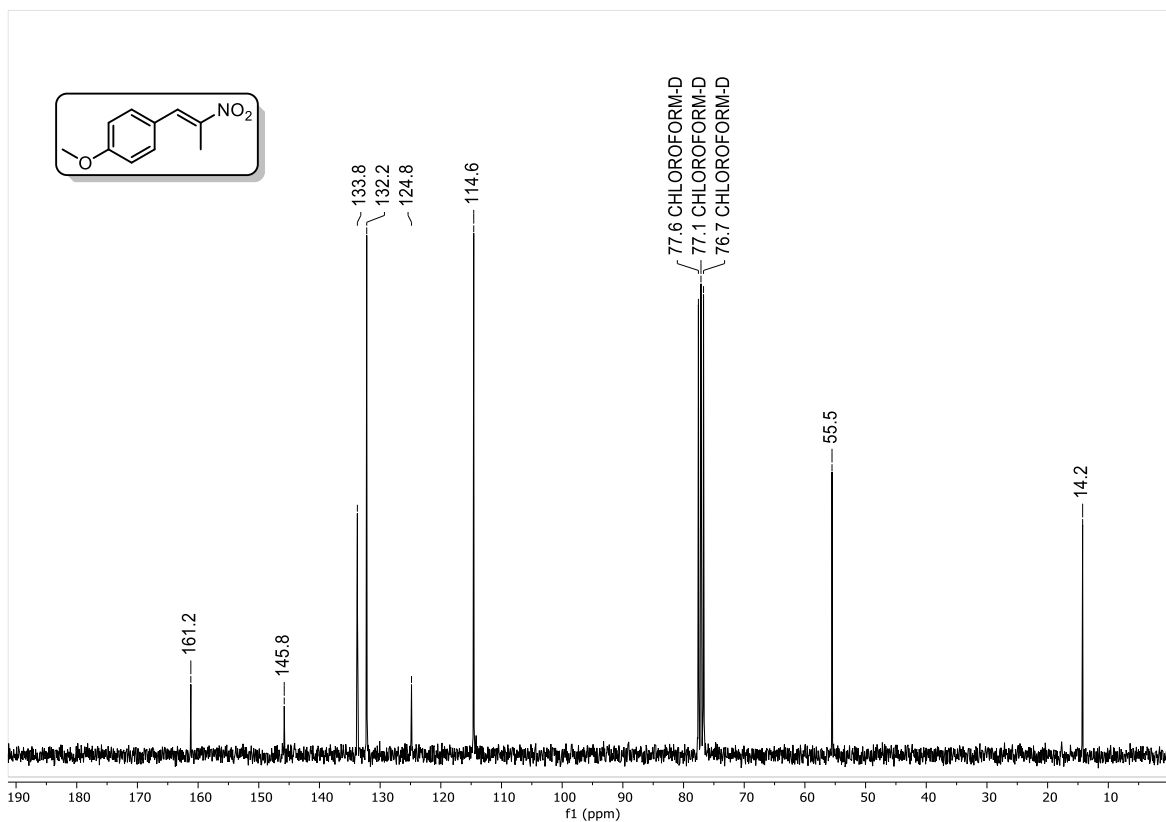


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
206.11824	3006.77	206.11810	0.14	0.68	¹² C ₁₂ ¹ H ₁₆ ¹⁴ N ₁ ¹⁶ O ₂	5.5

Spectrum 9. HRMS of 6c.



Spectrum 10. ¹H-NMR of 6d.



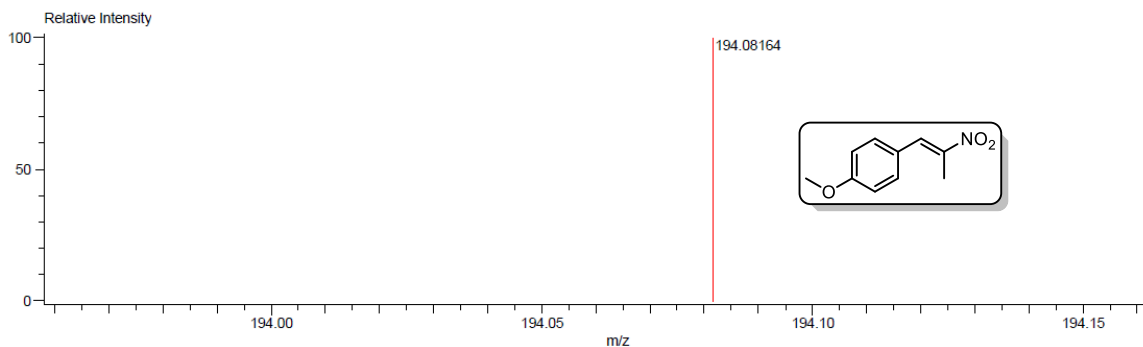
Spectrum 11. ¹³C-NMR of 6d.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
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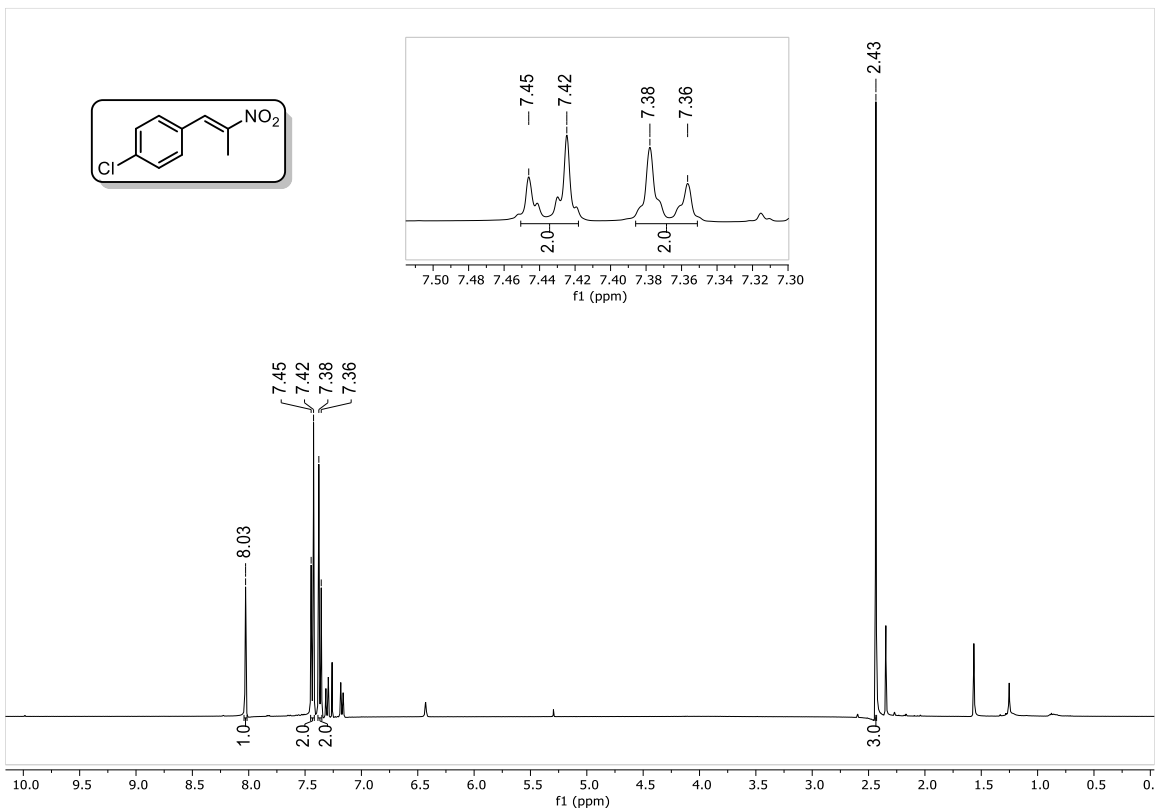
Charge number:1
 Element:¹²C:0 .. 10, ¹H:0 .. 28, ³⁵Cl:0 .. 0, ¹⁹F:0 .. 0, ¹⁴N:0 .. 1, ¹⁶O:0 .. 3, ³²S:0 .. 0
 Tolerance:5.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

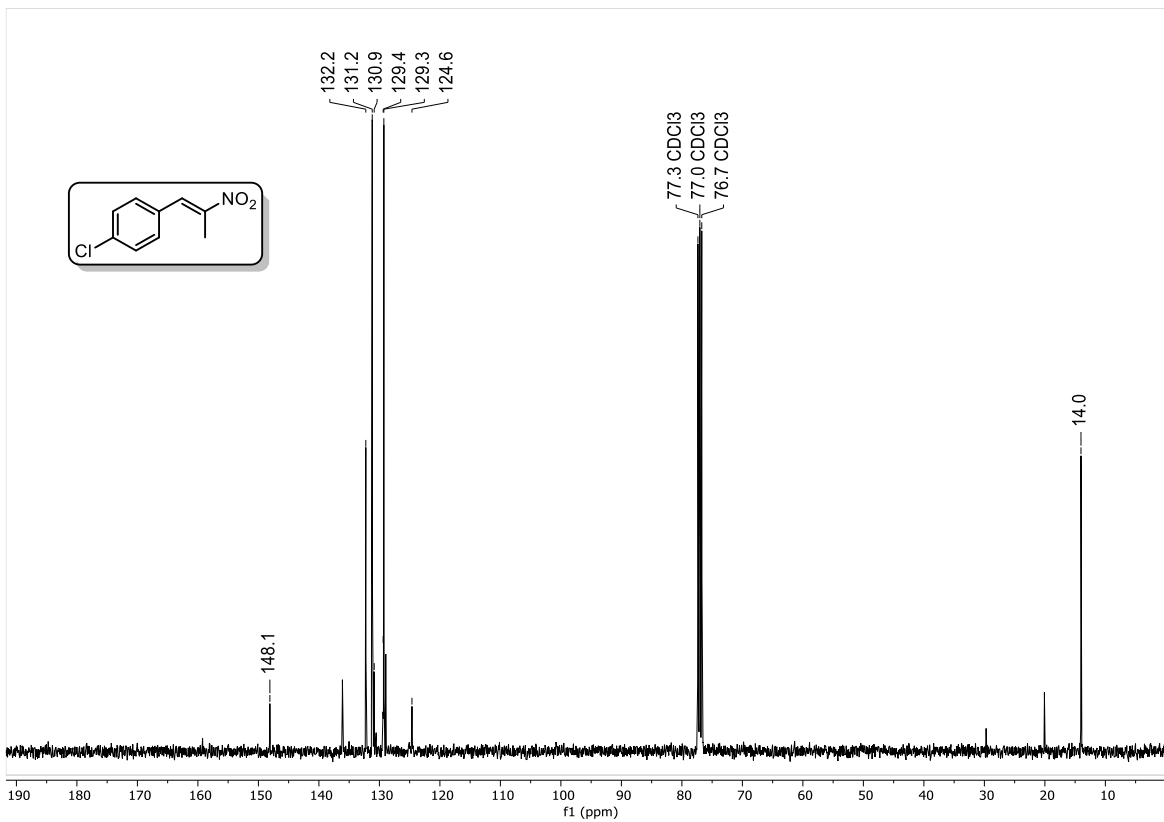


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
194.08164	6888.90	194.08172	-0.07	-0.38	¹² C ₁₀ ¹ H ₁₂ ¹⁴ N ₁ ¹⁶ O ₃	5.5

Spectrum 12. HRMS of 6d.



Spectrum 13. $^1\text{H-NMR}$ of 6e.



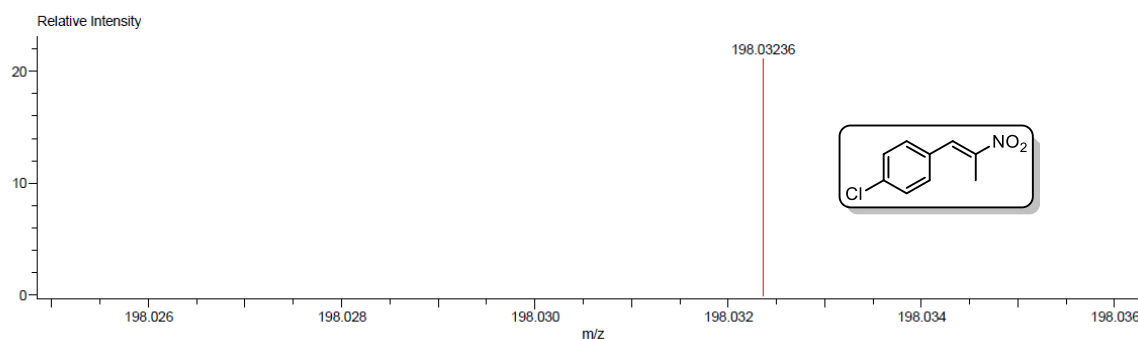
Spectrum 14. $^{13}\text{C-NMR}$ of 6e.

Description:
 Ionization Mode: ESI+
 History: Determine m/z [Peak Detect [Centroid, 30, Area]; Correct Base[]; Smooth [5]]; Correct Base [5.0%]; Average (MS[...

Mass Calibration data: Cal_PEG_600
 Created: 6/2/2023 9:58:59 AM
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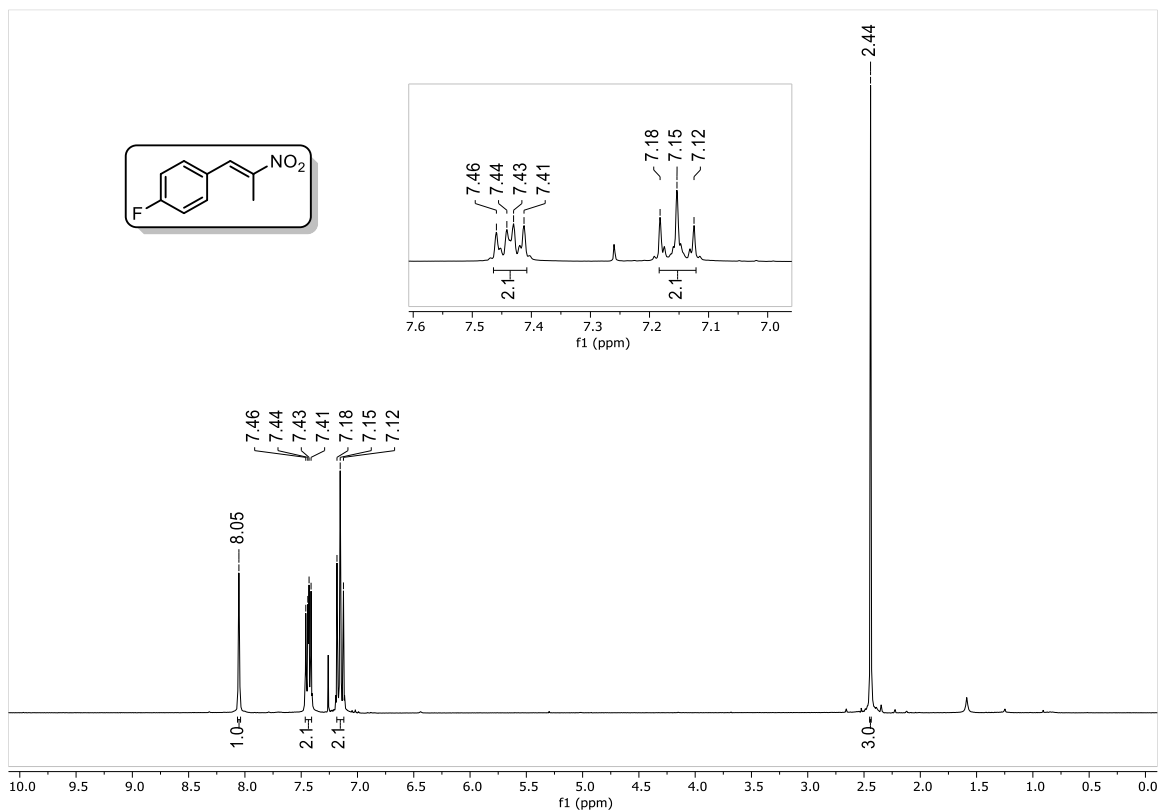
Charge number: 1
 Element: ^{12}C : 0 .. 9, ^1H : 0 .. 50, ^{35}Cl : 0 .. 1, ^{14}N : 0 .. 1, ^{16}O : 0 .. 2, ^{32}S : 0 .. 0
 Tolerance: 50.00 (ppm), 5.00 .. 15.00 (mmu)

Unsaturation Number: -1.0 .. 60.0 (Fraction: Both)

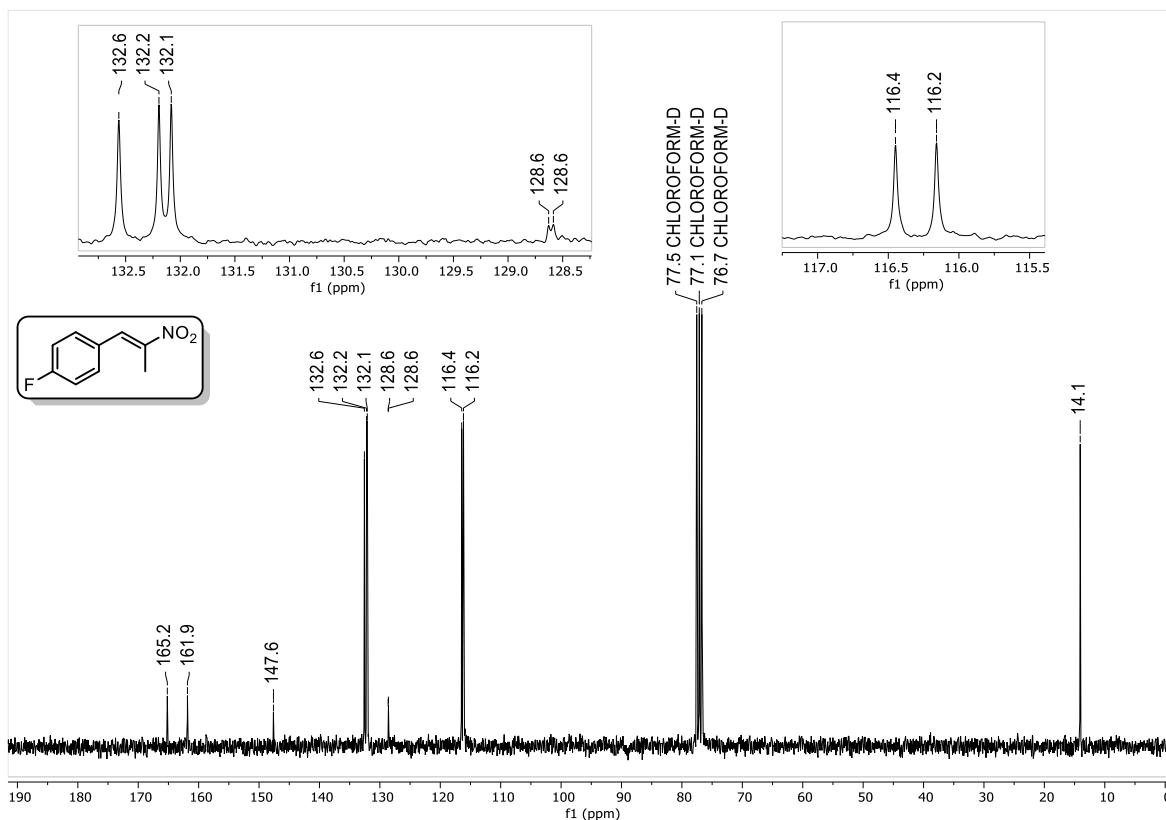


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
198.03236	1682.79	198.03218	0.18	0.91	$^{12}\text{C}_9\text{H}_9\text{Cl}_1\text{N}_1\text{O}_2$	5.5

Spectrum 15. HRMS of 6e.



Spectrum 16. ^1H -NMR of 6f.



Spectrum 17. ^{13}C -NMR of **6f**.

Description:
Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600

Created:6/1/2023 10:01:51 AM

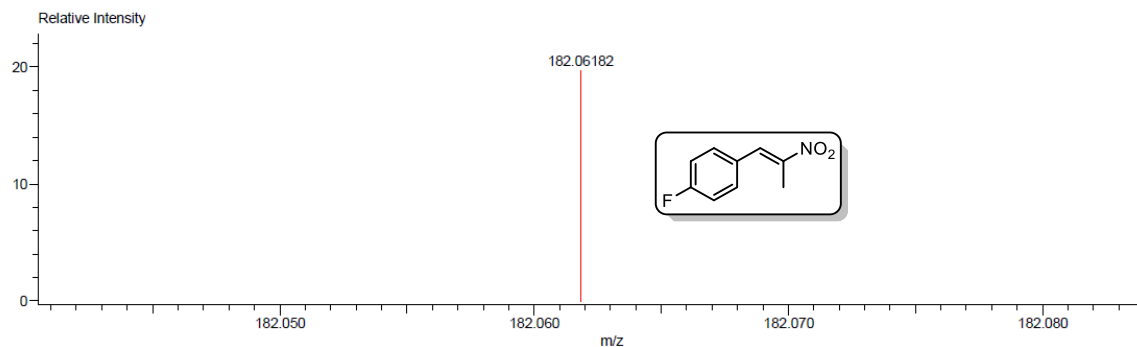
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Charge number:1

Tolerance:5.00(ppm), 5.00 .. 15.00(mmu)

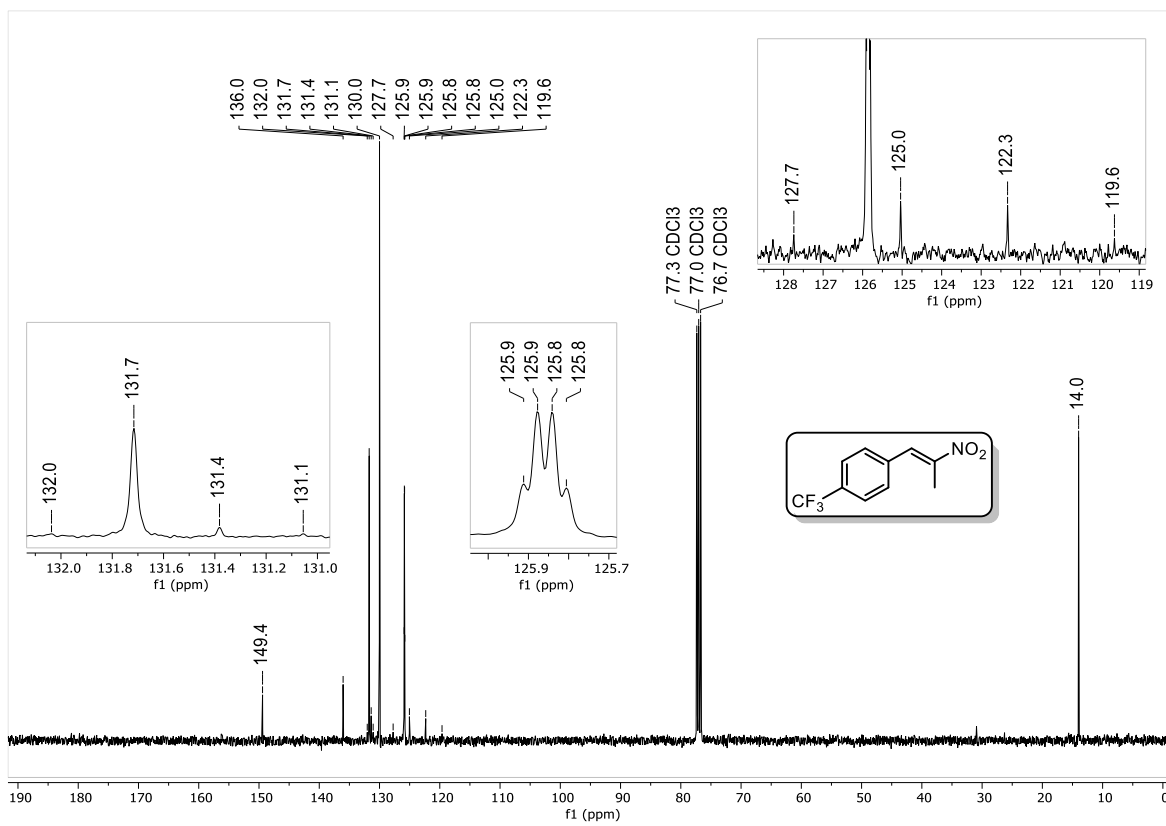
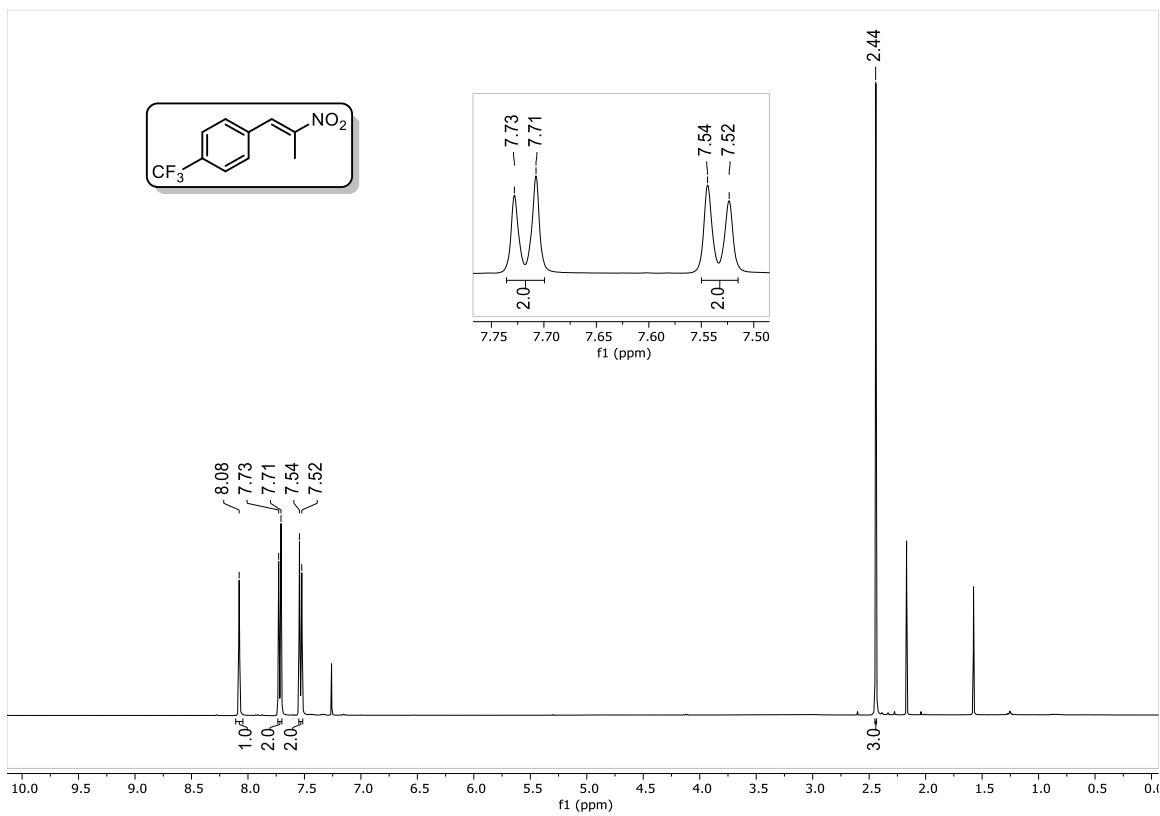
Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

Element: ^{12}C :0 .. 22, ^1H :0 .. 28, ^{35}Cl :0 .. 0, ^{19}F :1 .. 1, ^{14}N :0 .. 1, ^{16}O :0 .. 2, ^{32}S :0 .. 0



Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
182.06182	58399.13	182.06173	0.09	0.49	$^{12}\text{C}_9\text{H}_6\text{F}_1\text{N}_1\text{O}_2$	5.5

Spectrum 18. HRMS of **6f**.



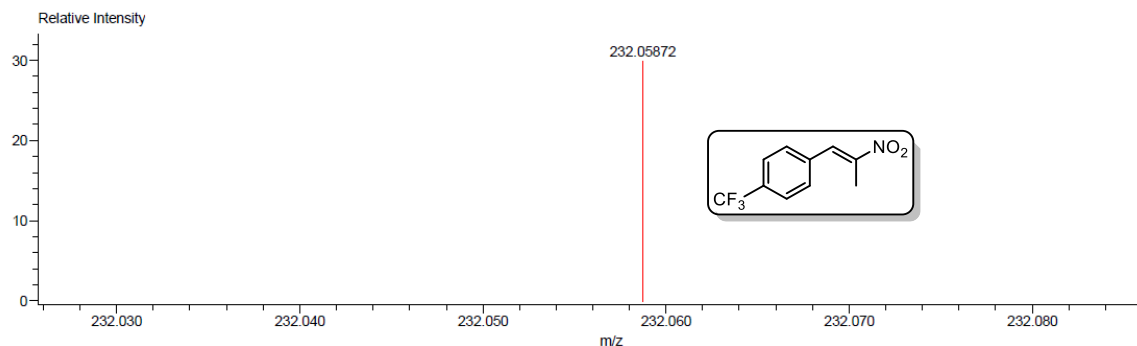
Spectrum 20. ¹³C-NMR of 6g.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:Cal_PEG_600
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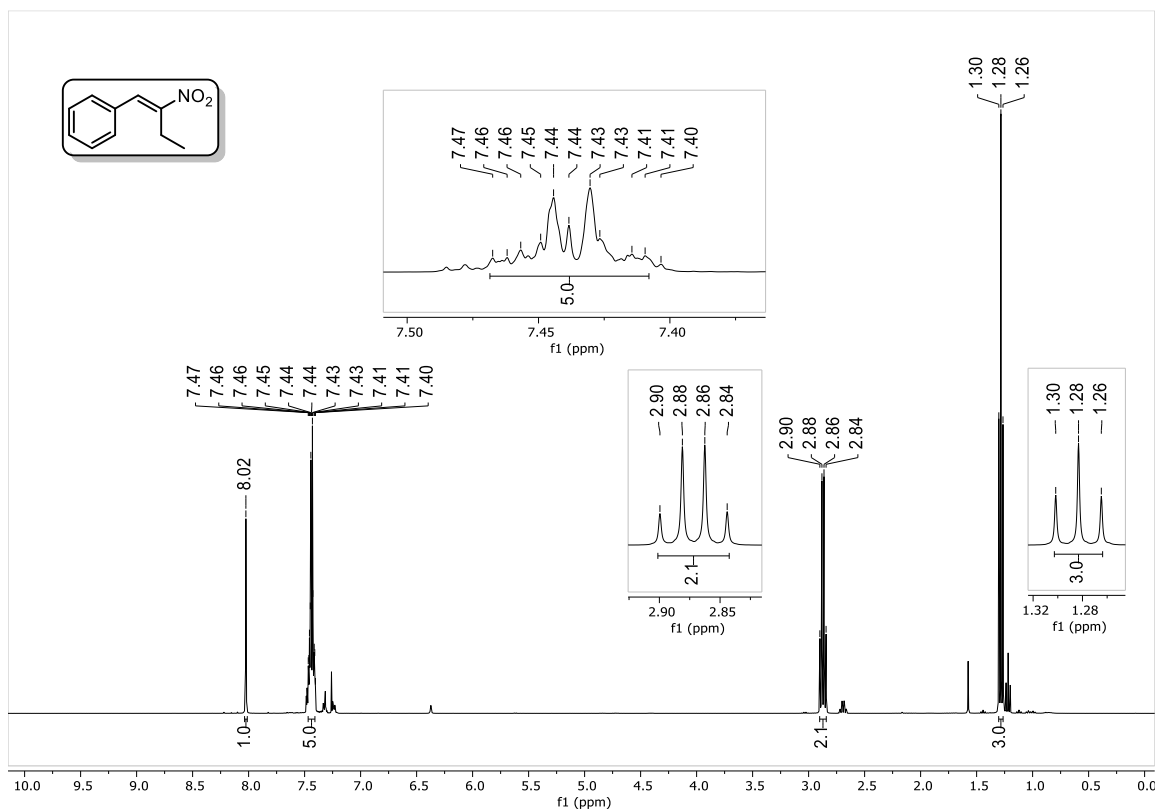
Charge number:1
 Element:¹²C:0 .. 10, ¹H:0 .. 28, ³⁵Cl:0 .. 0, ¹⁹F:0 .. 3, ¹⁴N:0 .. 1, ¹⁶O:0 .. 2, ³²S:0 .. 0
 Tolerance:5.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

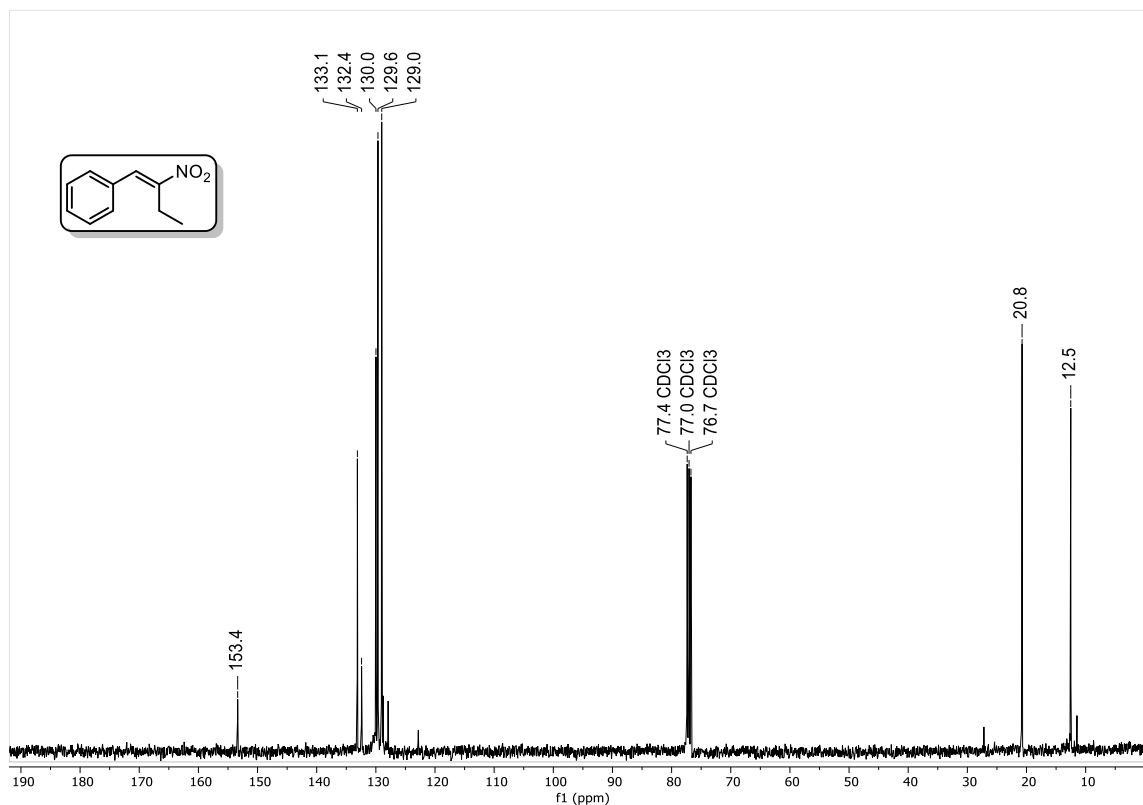


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
232.05872	23637.22	232.05854	0.18	0.79	¹² C ₁₀ ¹ H ₆ ¹⁹ F ₃ ¹⁴ N ₁ ¹⁶ O ₂	5.5

Spectrum 21. HRMS of 6g.



Spectrum 22. ¹H-NMR of 6h.



Spectrum 23. ^{13}C -NMR of 6h.

Description:
Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:Cal_PEG_600

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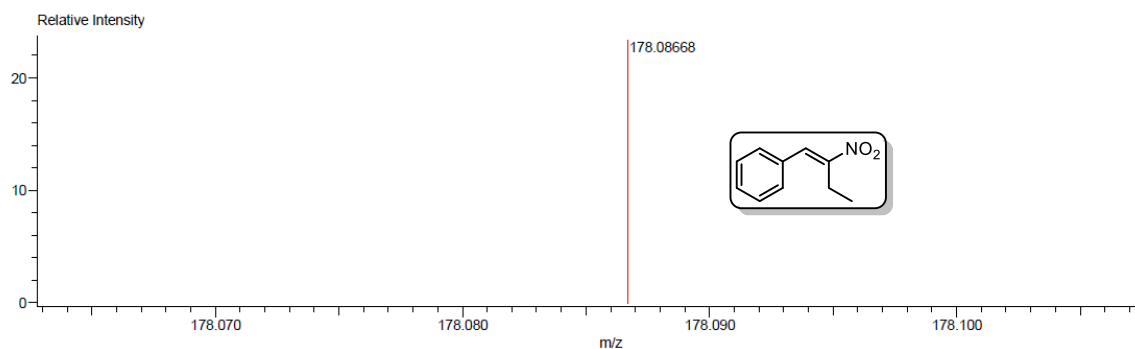
Created by:

Charge number:1

Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)

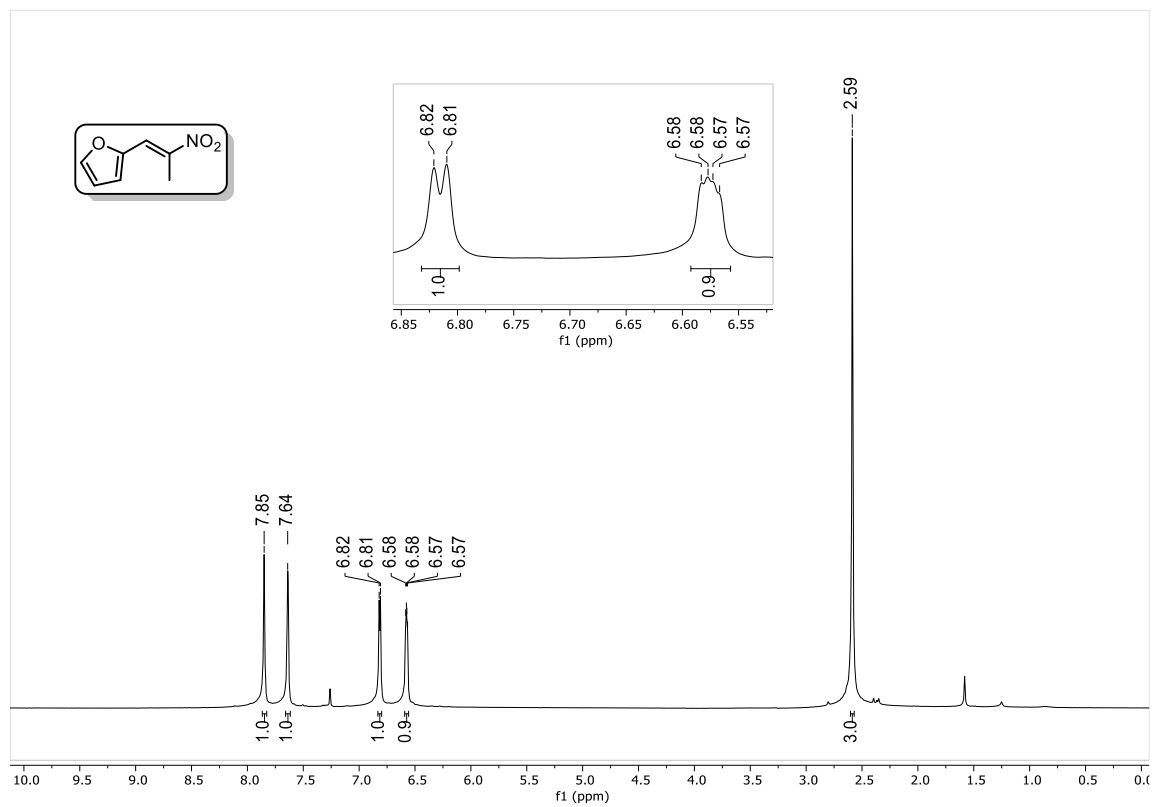
Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

Element: ^{12}C :0 .. 12, ^1H :0 .. 50, ^{35}Cl :0 .. 0, ^{14}N :0 .. 3, ^{16}O :0 .. 2, ^{32}S :0 .. 0

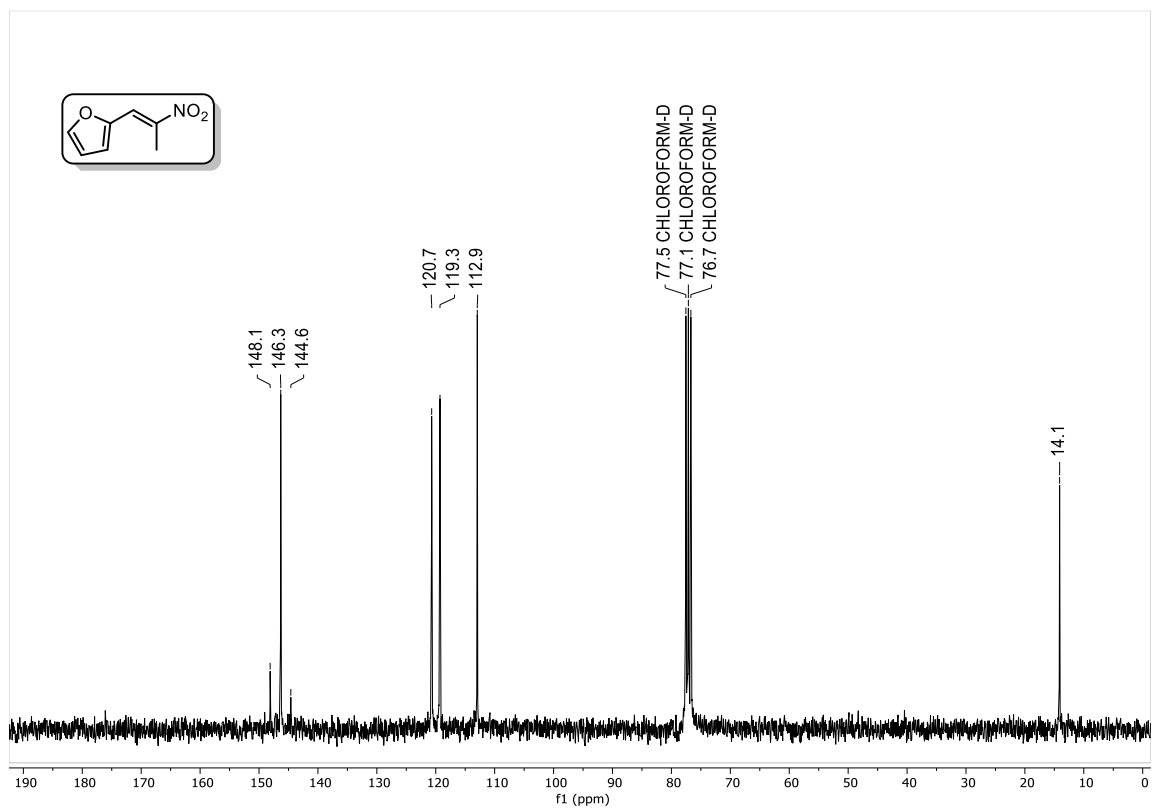


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
178.08668	1529.45	178.08680	-0.13	-0.71	$^{12}\text{C}_{10}^{1}\text{H}_{12}^{14}\text{N}_1^{16}\text{O}_2$	5.5

Spectrum 24. HRMS of 6h.



Spectrum 25. ¹H-NMR of 6j.



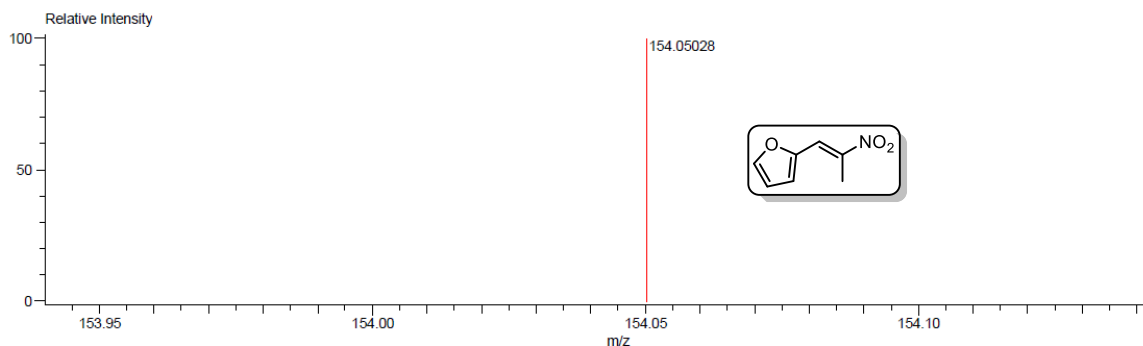
Spectrum 26. ¹³C-NMR of 6j.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
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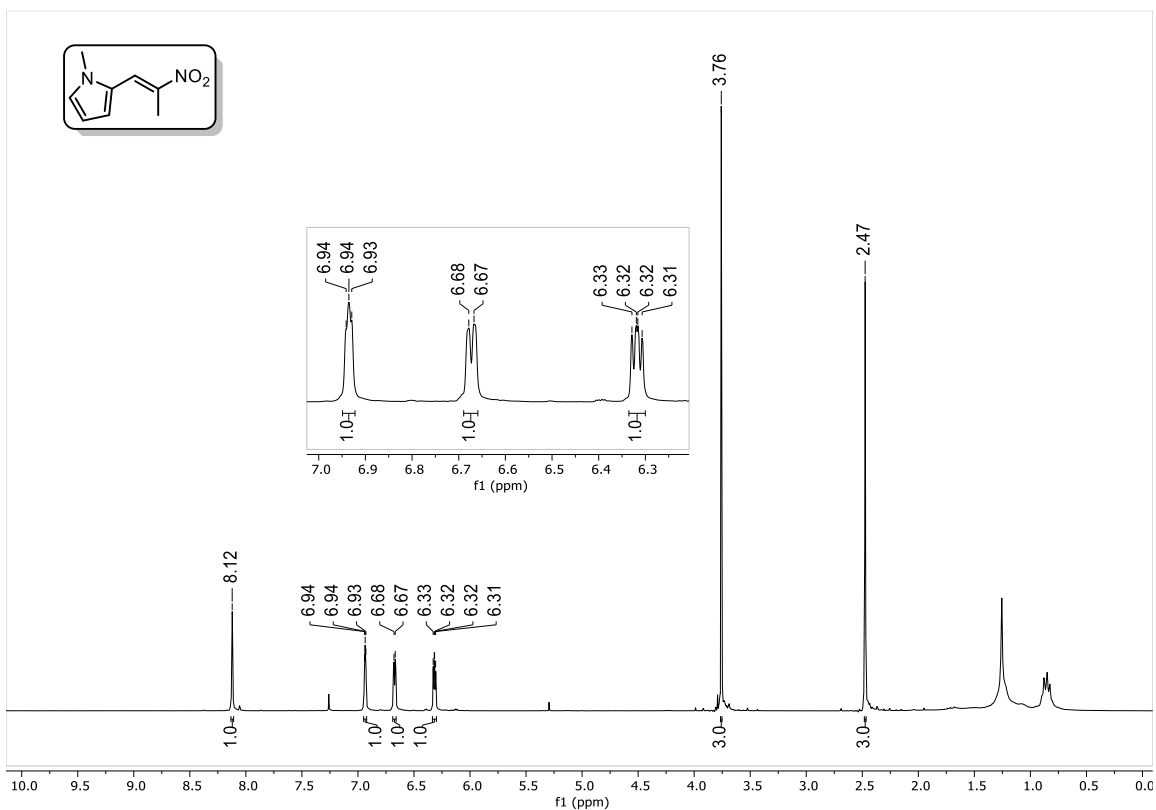
Charge number:1
 Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)
 Element:¹²C:0 .. 7, ¹H:0 .. 50, ³⁵Cl:0 .. 0, ¹⁴N:0 .. 1, ¹⁶O:0 .. 3, ³²S:0 .. 0

Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

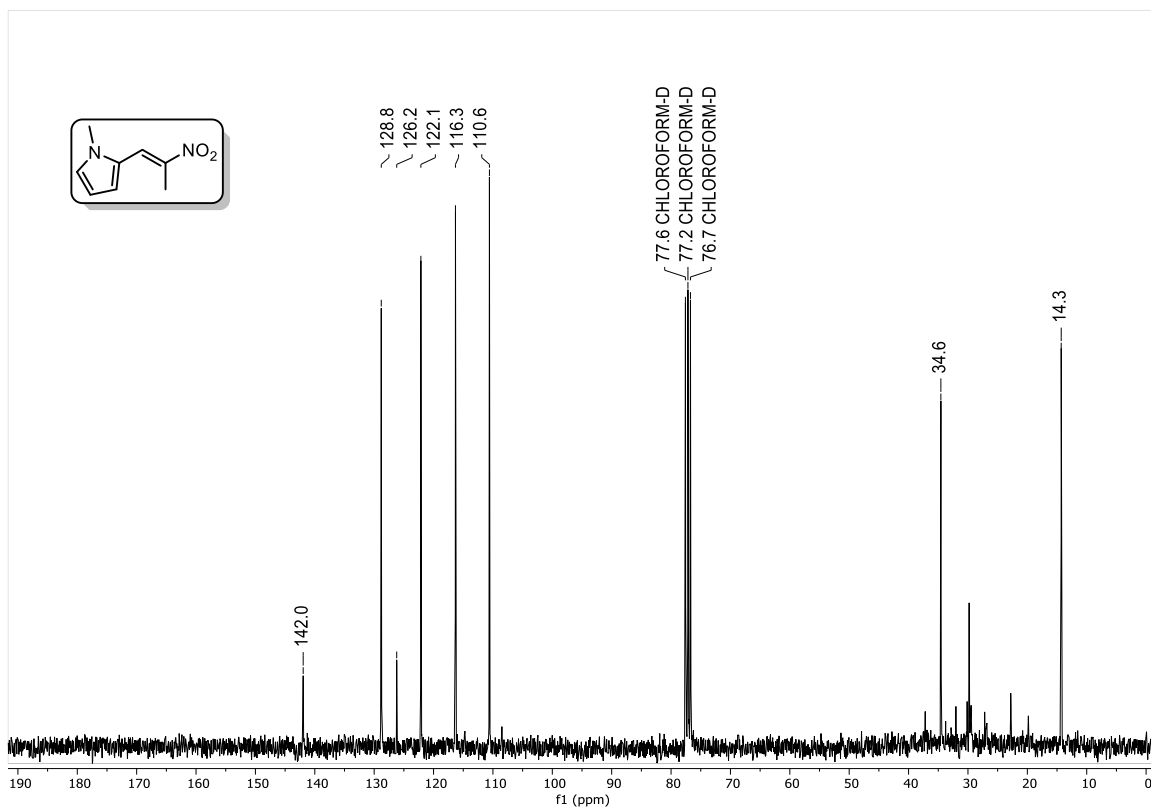


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
154.05028	92551.73	154.05042	-0.13	-0.87	¹² C ₇ ¹ H ₈ ¹⁴ N ₁ ¹⁶ O ₃	4.5

Spectrum 27. HRMS of 6j.



Spectrum 28. ¹H-NMR of 6k.



Spectrum 29. ^{13}C -NMR of 6k.

Description:
Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:Cal_PEG_600

Created:6/2/2023 10:11:27 AM

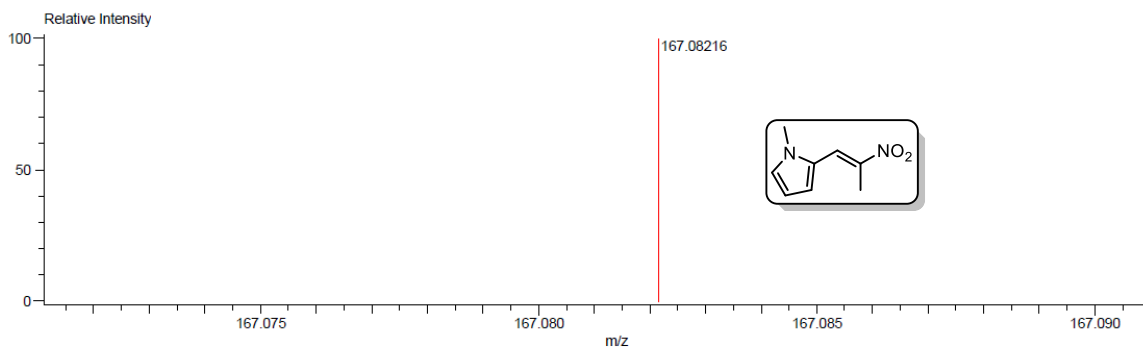
Created by:

Charge number:1

Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)

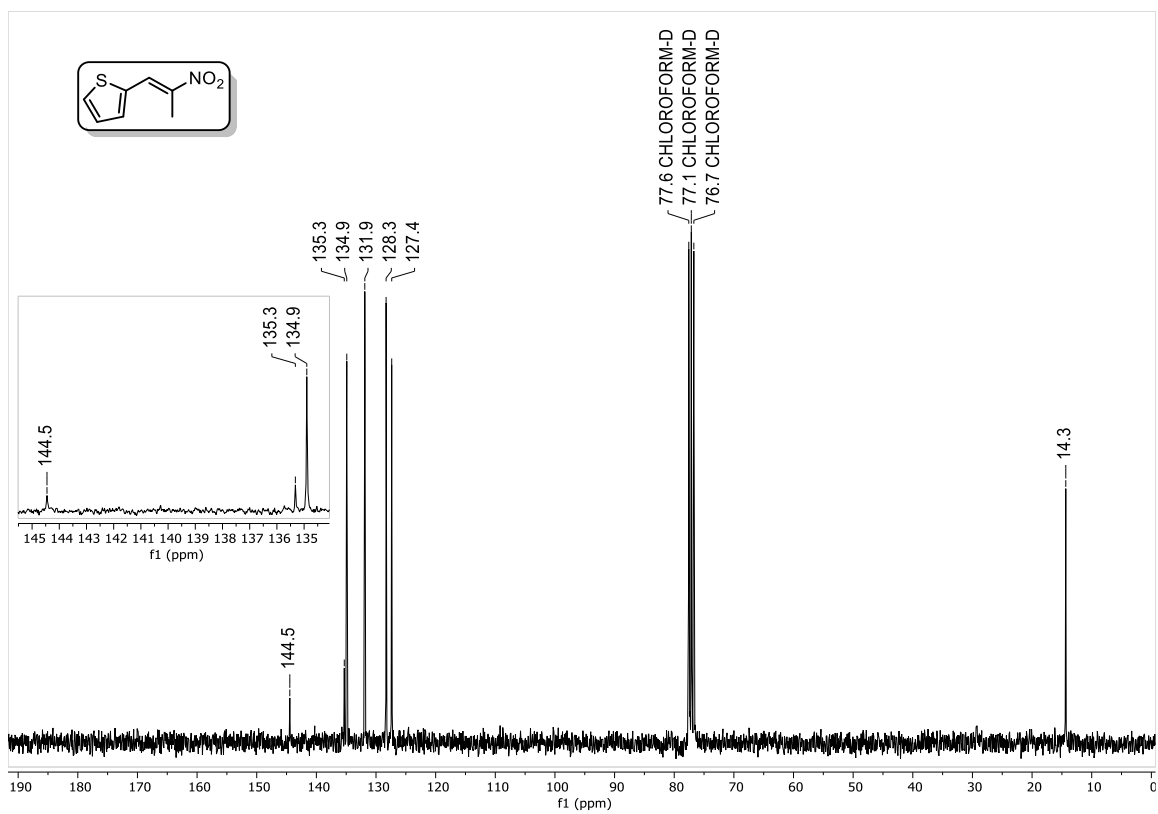
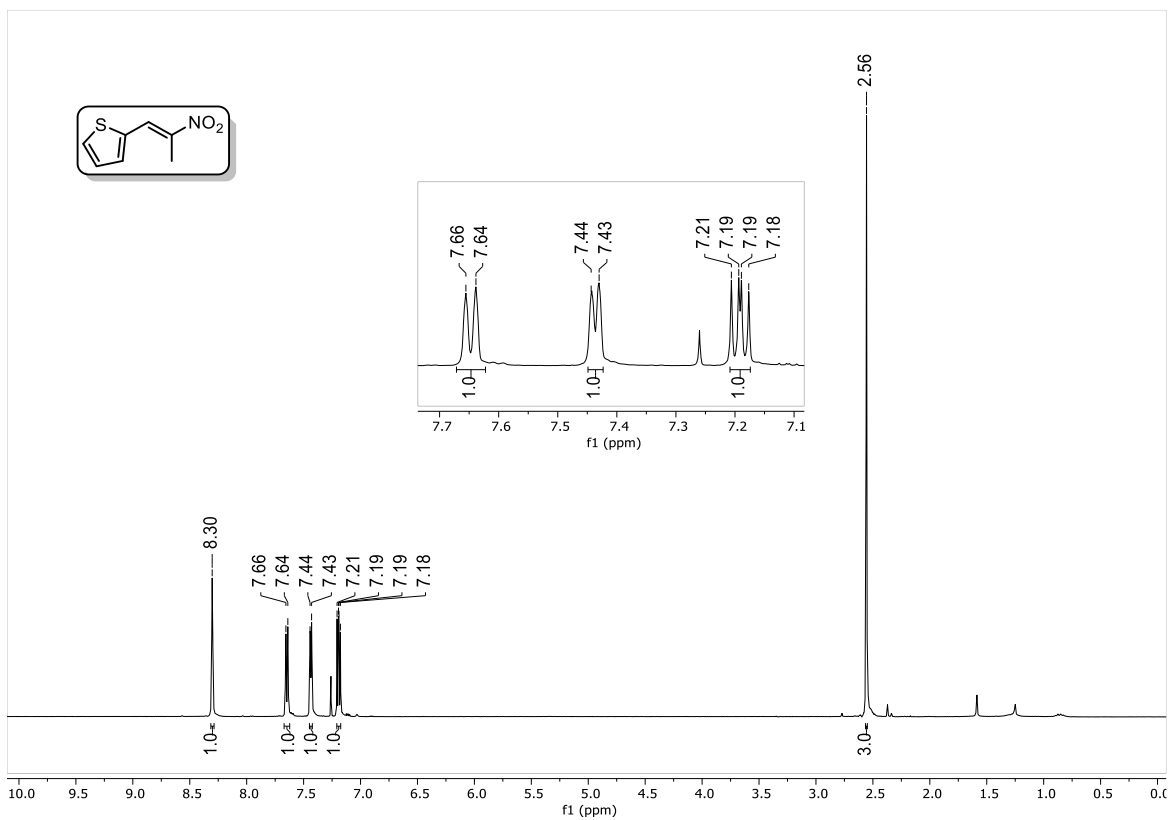
Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

Element: ^{12}C :0 .. 8, ^1H :0 .. 50, ^{35}Cl :0 .. 0, ^{14}N :0 .. 2, ^{16}O :0 .. 2, ^{32}S :0 .. 0



Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
167.08216	6406.48	167.08205	0.11	0.66	$^{12}\text{C}_8^{1}\text{H}_{11}^{14}\text{N}_2^{16}\text{O}_2$	4.5

Spectrum 30. HRMS of 6k.

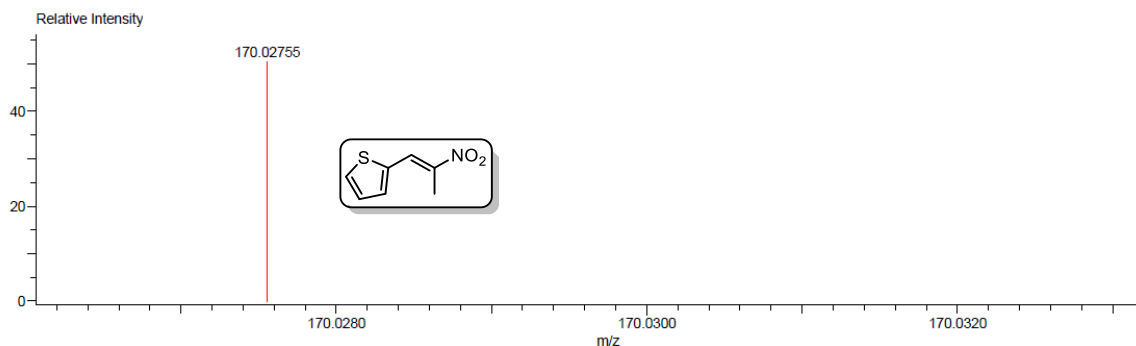


Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:Cal_PEG_600
 Created:6/2/2023 9:53:05 AM
 Created by:

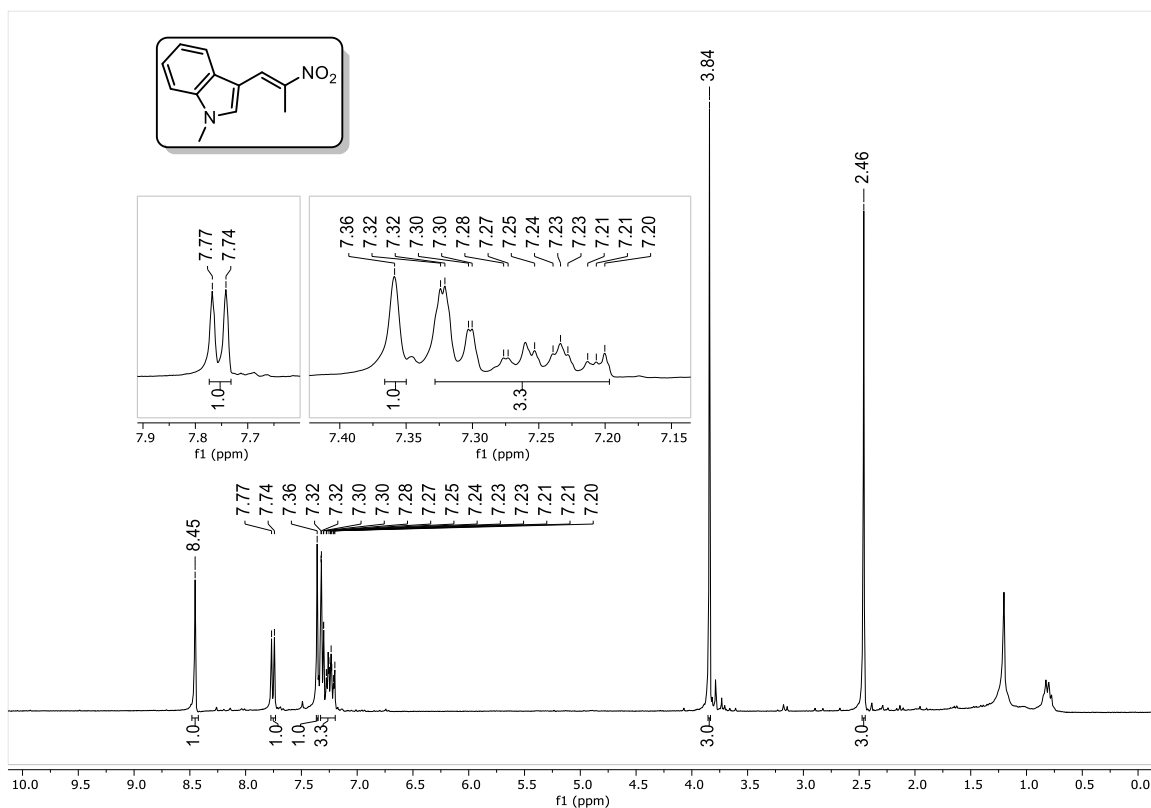
Charge number:1
 Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)
 Element:¹²C:0 .. 8, ¹H:0 .. 50, ³⁵Cl:0 .. 0, ¹⁴N:0 .. 1, ¹⁶O:0 .. 2, ³²S:0 .. 1

Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

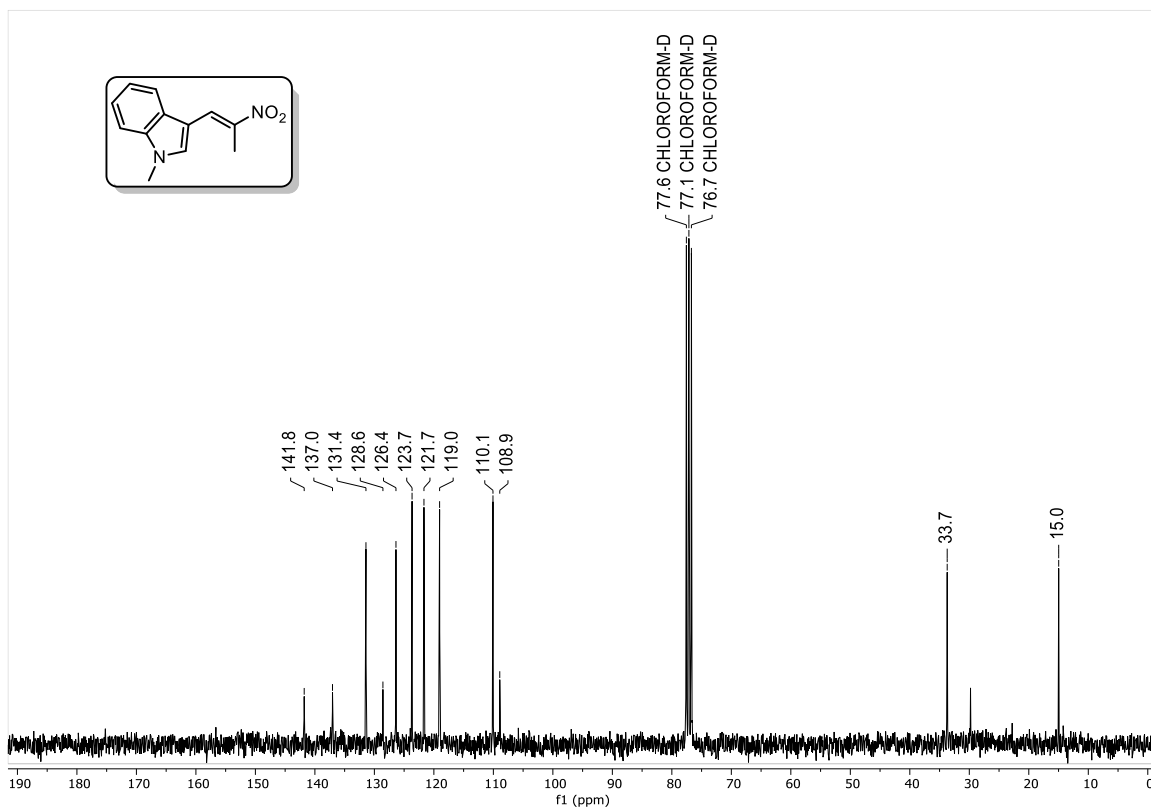


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
170.02755	3602.55	170.02757	-0.02	-0.12	¹² C ₇ ¹ H ₈ ¹⁴ N ₁ ¹⁶ O ₂ ³² S ₁	5.5

Spectrum 33. HRMS of 6I.



Spectrum 34. ¹H-NMR of 6m.



Spectrum 35. ¹³C-NMR of 6m.

Description:
Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area],Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

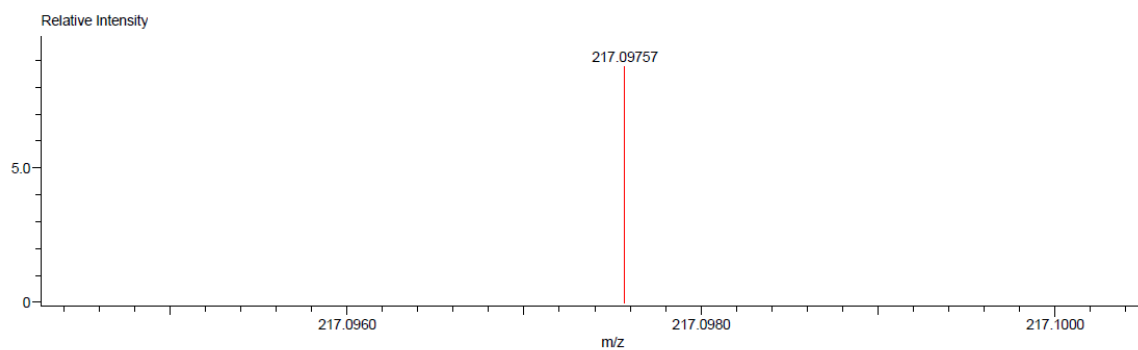
Mass Calibration data:Cal_PEG_600
Created:6/6/2023 10:43:33 AM
Created by:AccuTOF

Charge number:1

Tolerance:100.00(ppm), 5.00 .. 15.00(mmu)

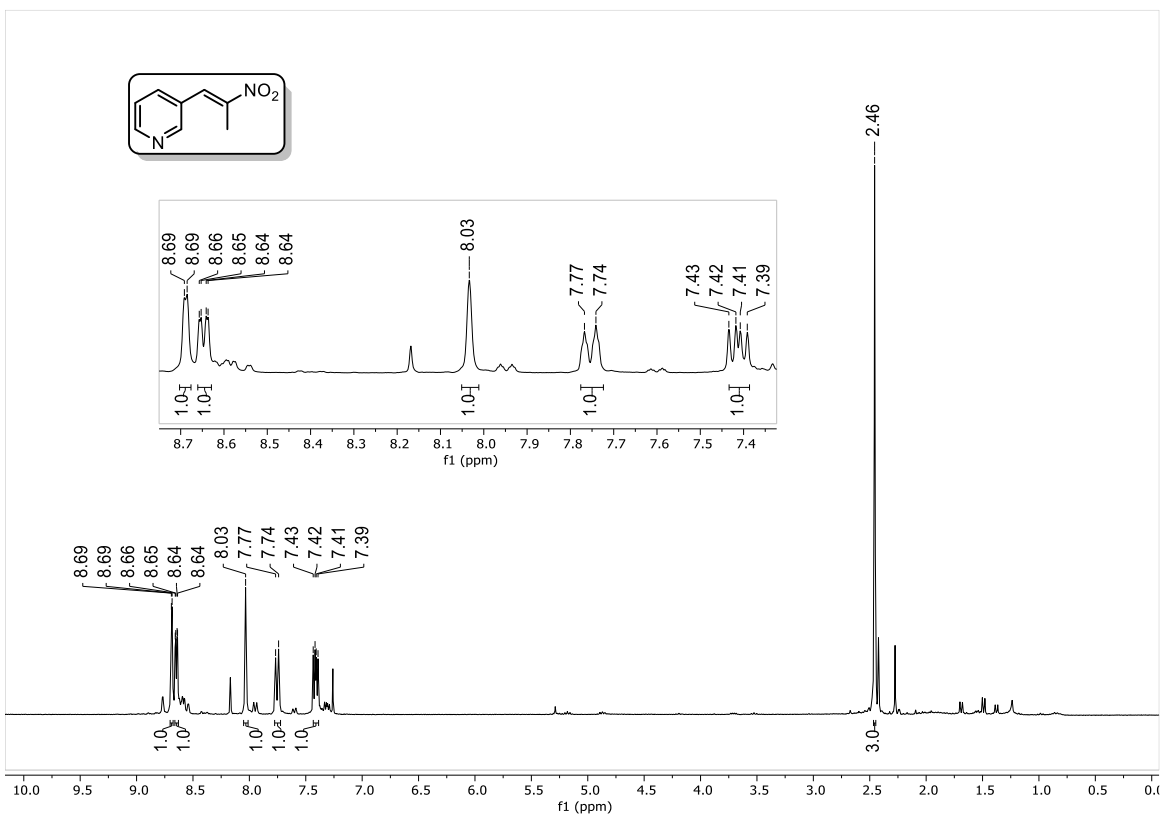
Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)

Element:¹²C:12 .. 12, ¹H:13 .. 30, ¹⁴N:0 .. 2, ¹⁶O:2 .. 2

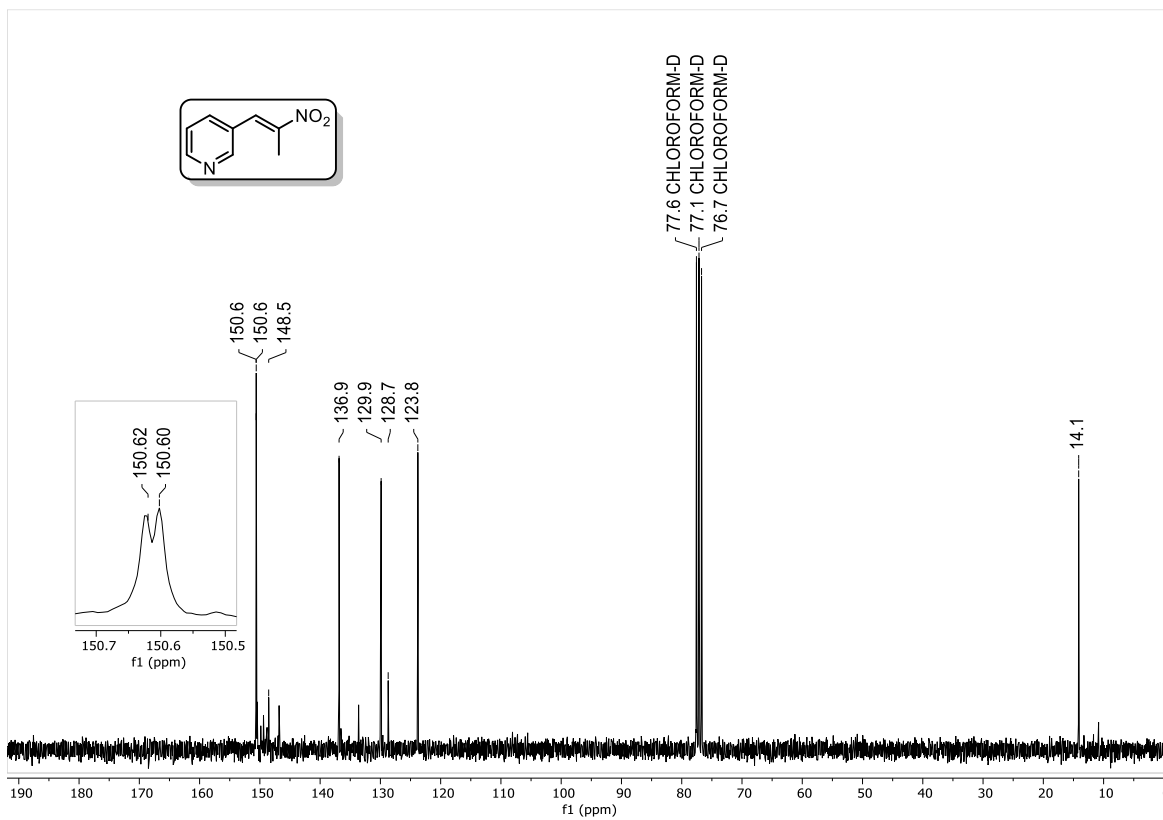


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
217.09757	8308.68	217.09770	-0.13	-0.61	¹² C ₁₂ ¹ H ₁₃ ¹⁴ N ₂ ¹⁶ O ₂	7.5

Spectrum 36. HRMS of 6m.



Spectrum 37. $^1\text{H-NMR}$ of 6n.



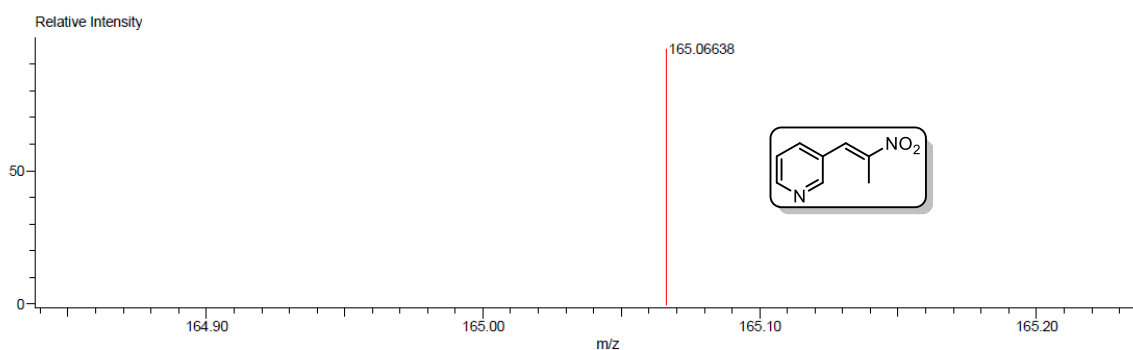
Spectrum 38. $^{13}\text{C-NMR}$ of 6n.

Description:
Ionization Mode:ESI+
History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:Cal_PEG_600
Created:6/2/2023 10:20:25 AM
Created by:

Charge number:1
Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)
Element:¹²C:0 .. 12, ¹H:0 .. 50, ³⁵Cl:0 .. 0, ¹⁴N:0 .. 3, ¹⁶O:0 .. 2, ³²S:0 .. 0

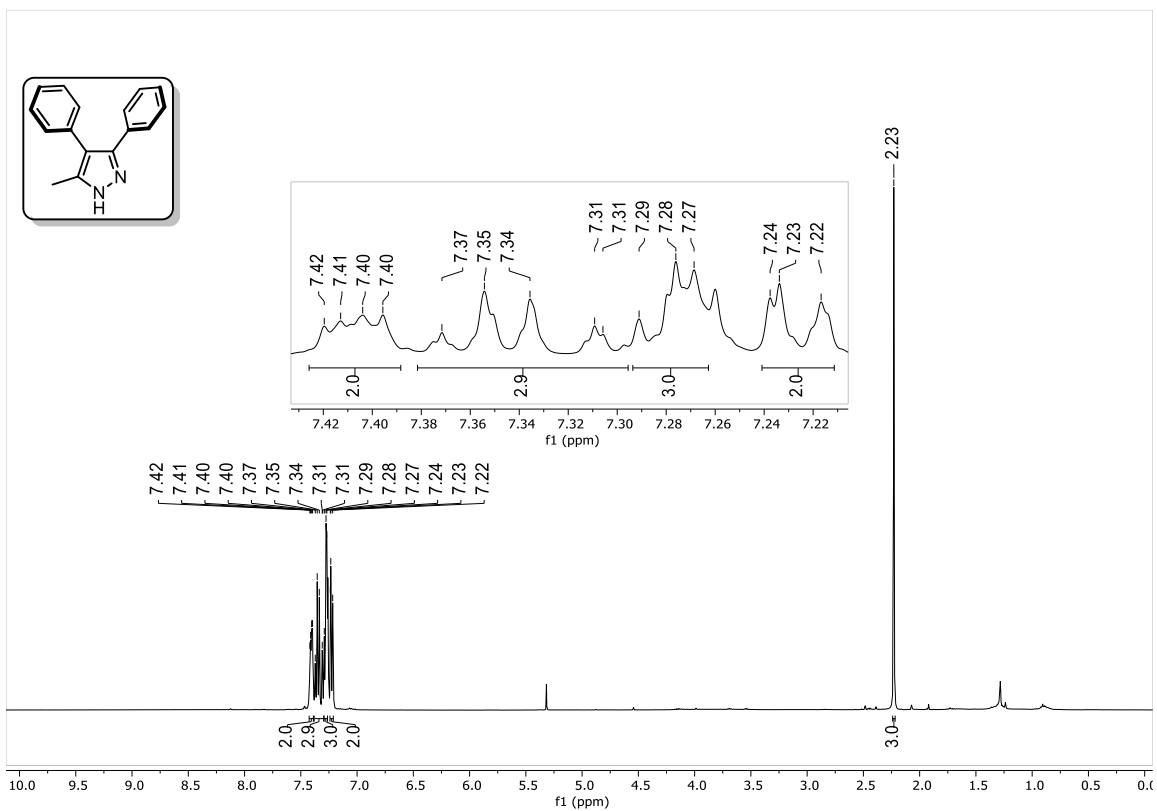
Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)



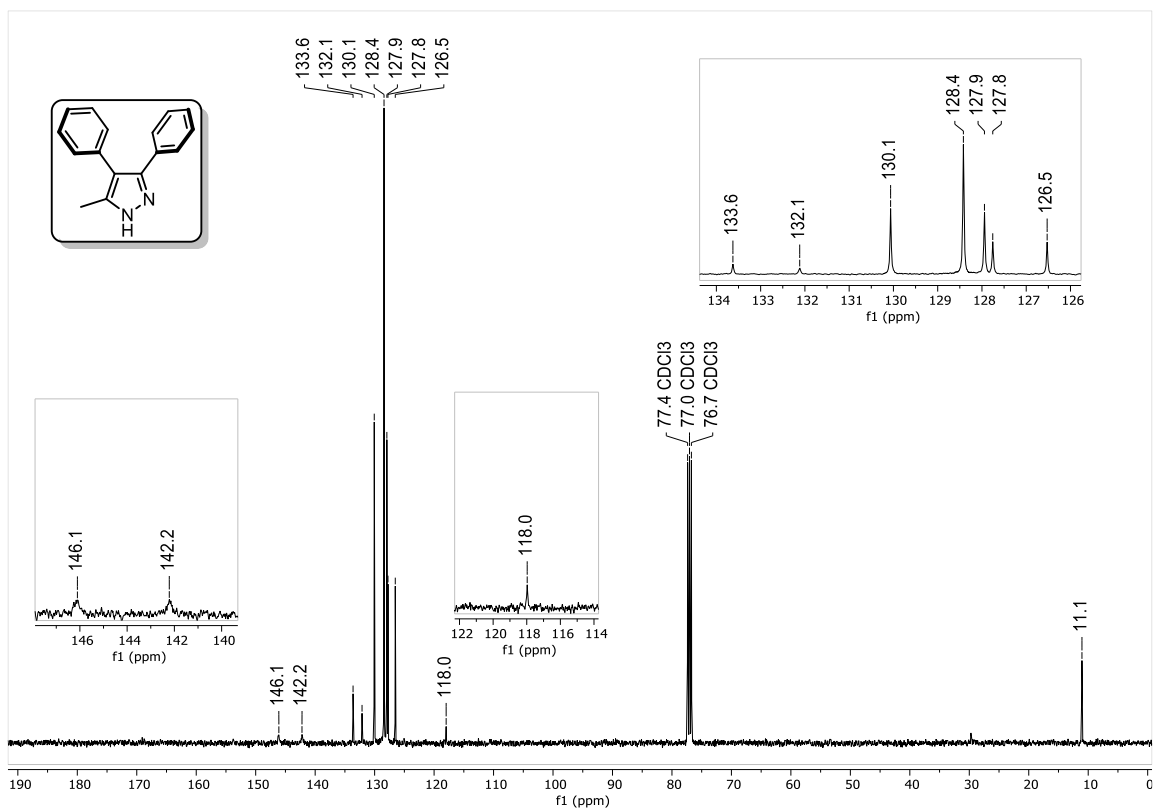
Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
165.06638	3005.60	165.06640	-0.02	-0.11	¹² C ₈ ¹ H ₈ ¹⁴ N ₂ ¹⁶ O ₂	5.5

Spectrum 39. HRMS of 6n.

^1H -NMR, ^{13}C -NMR, and HRMS spectra of the
pyrazoles.



Spectrum 40. ¹H-NMR of 8a.



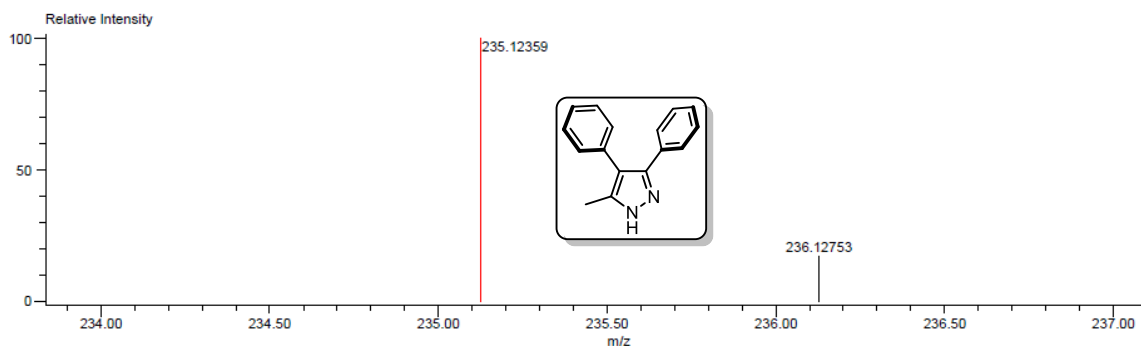
Spectrum 41. ¹³C-NMR of 8a.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:CaI_PEG_600
 Created:11/28/2022 2:08:11 PM
 Created by:AccuTOF

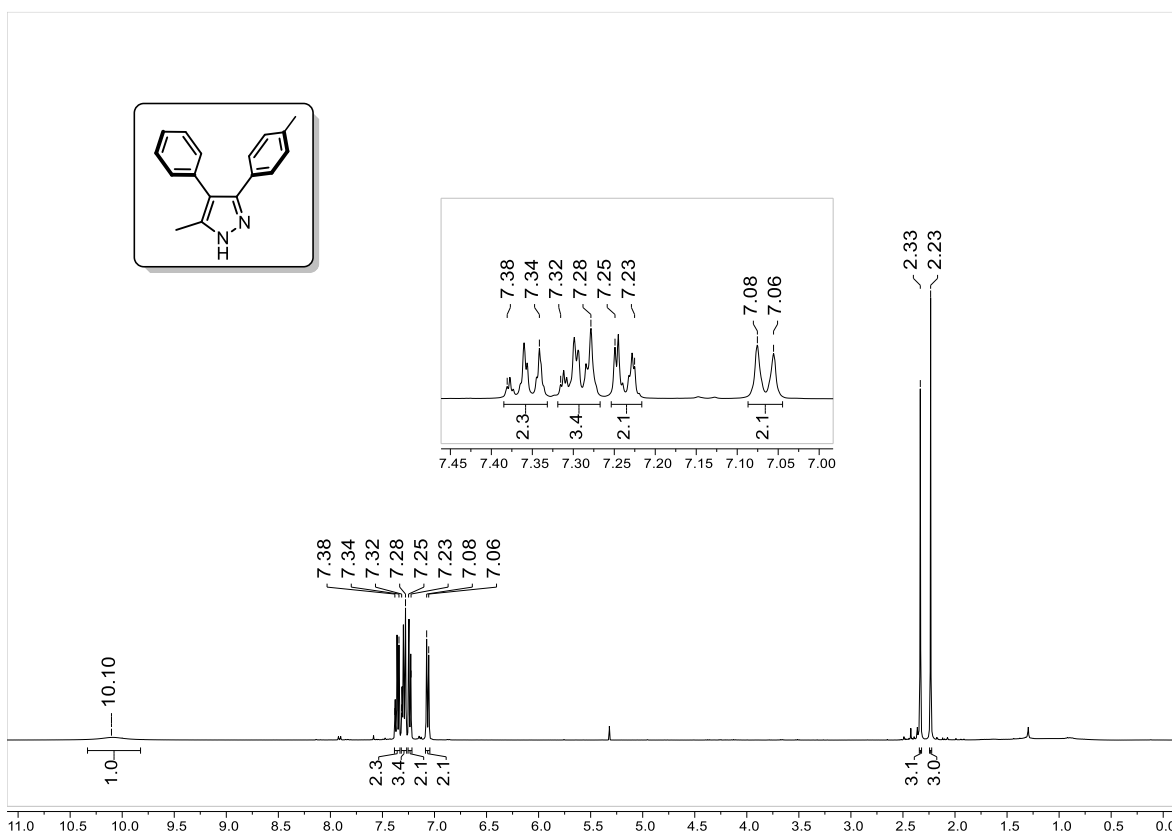
Charge number:1
 Element:¹²C:0 .. 16, ¹H:0 .. 50, ⁷⁹Br:0 .. 0, ¹⁴N:2 .. 2, ¹⁶O:0 .. 0
 Tolerance:200.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 30.0 (Fraction:Both)

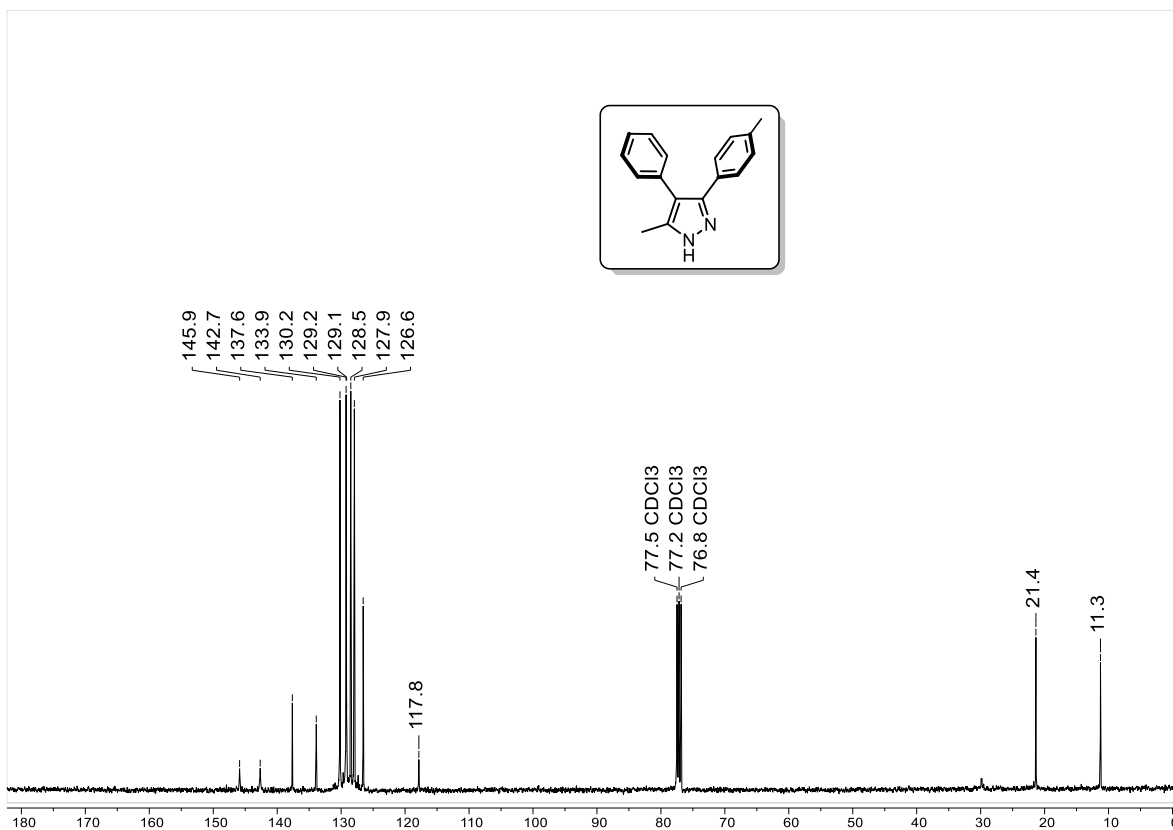


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
235.12359	176868.61	235.12352	0.06	0.27	¹² C ₁₆ ¹ H ₁₅ ¹⁴ N ₂	10.5

Spectrum 42. HRMS of 8a.



Spectrum 43. ¹H-NMR of 8b.



Spectrum 44. ¹³C-NMR of 8b.

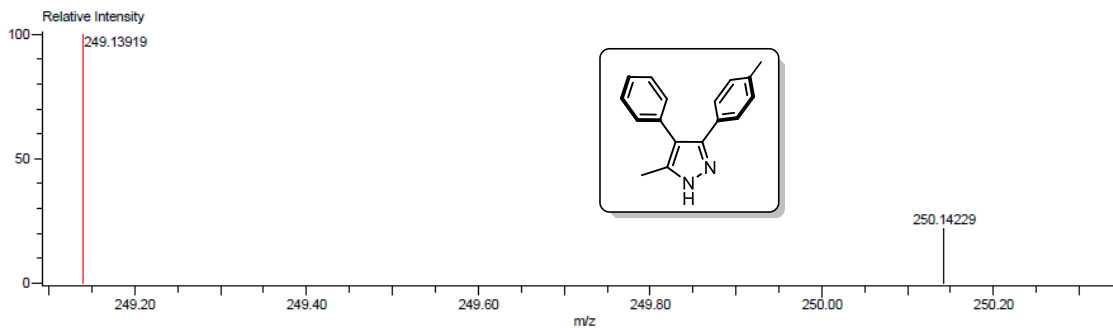
Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area],Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
 Created:2/23/2023 12:46:57 PM
 Created by:AccuTOF

Charge number:1
 Element:¹²C:0 .. 20, ¹H:0 .. 18, ¹⁴N:0 .. 3, ¹⁶O:0 .. 1

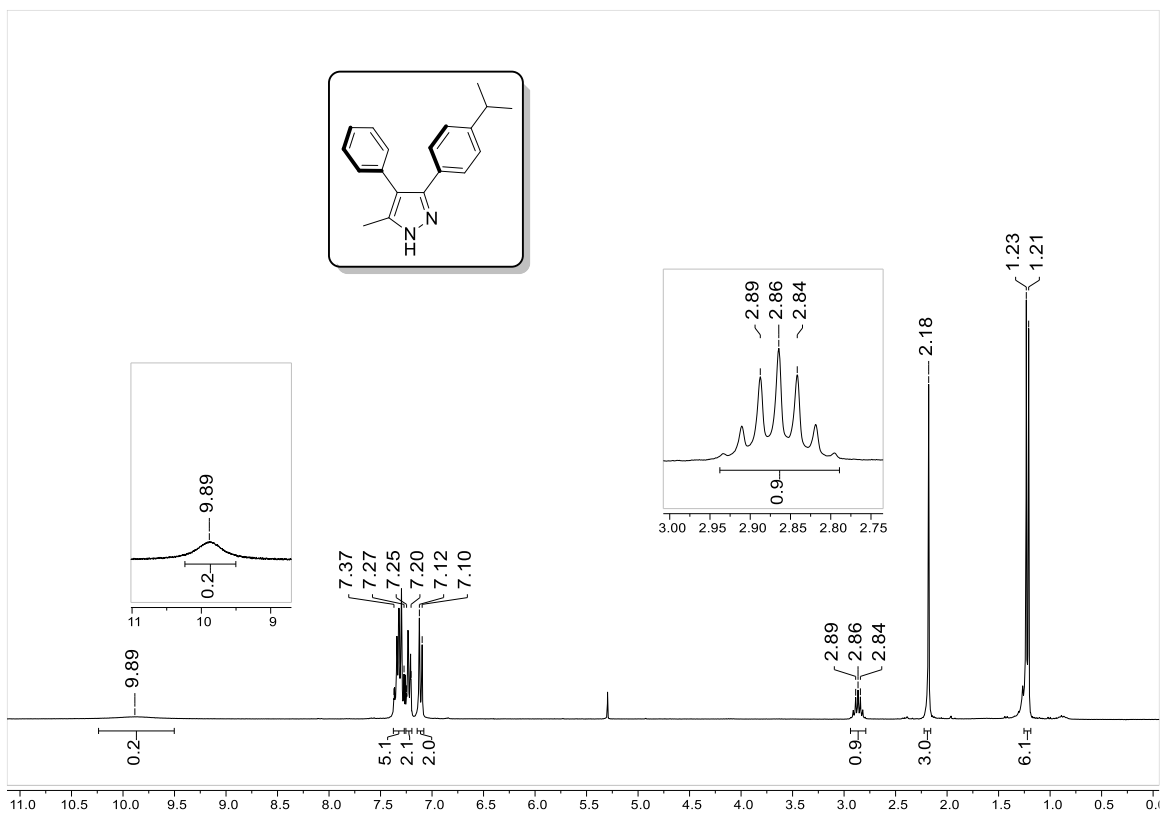
Tolerance:4.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 40.0 (Fraction:Both)

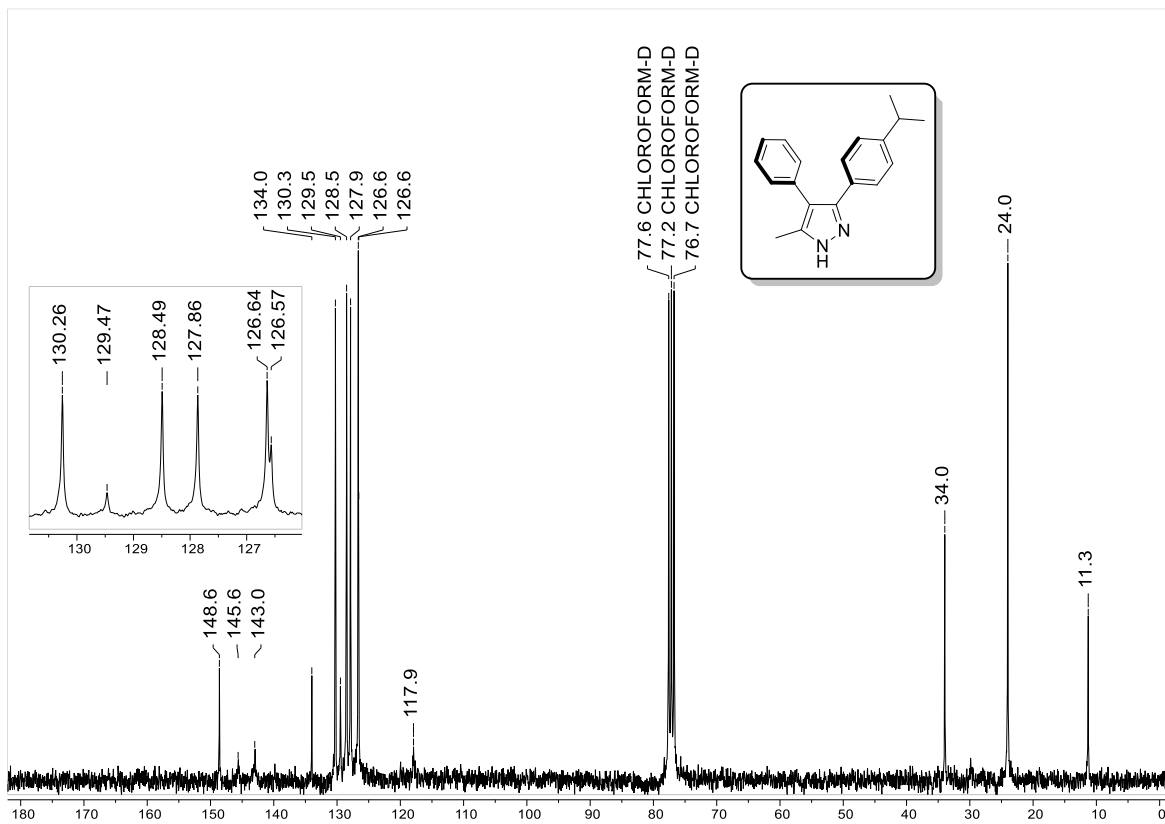


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
249.13919	5395.11	249.13917	0.02	0.07	¹² C ₁₇ ¹ H ₁₇ ¹⁴ N ₂	10.5

Spectrum 45. HRMS of 8b.



Spectrum 46. ¹H-NMR of 8c.



Spectrum 47. ¹³C-NMR of 8c.

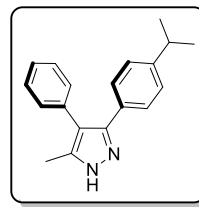
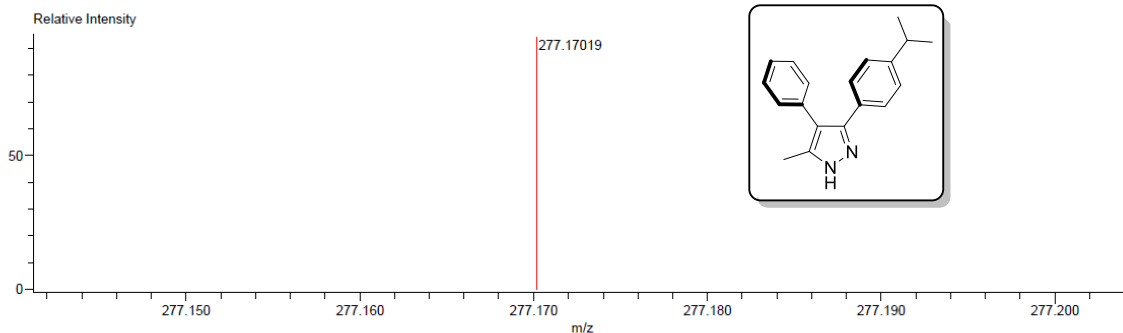
Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
 Created:2/3/2023 12:32:06 PM
 Created by:AccuTOF

Charge number:1
 Element:¹²C:0 .. 19, ¹H:0 .. 50, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0

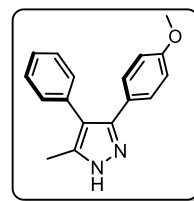
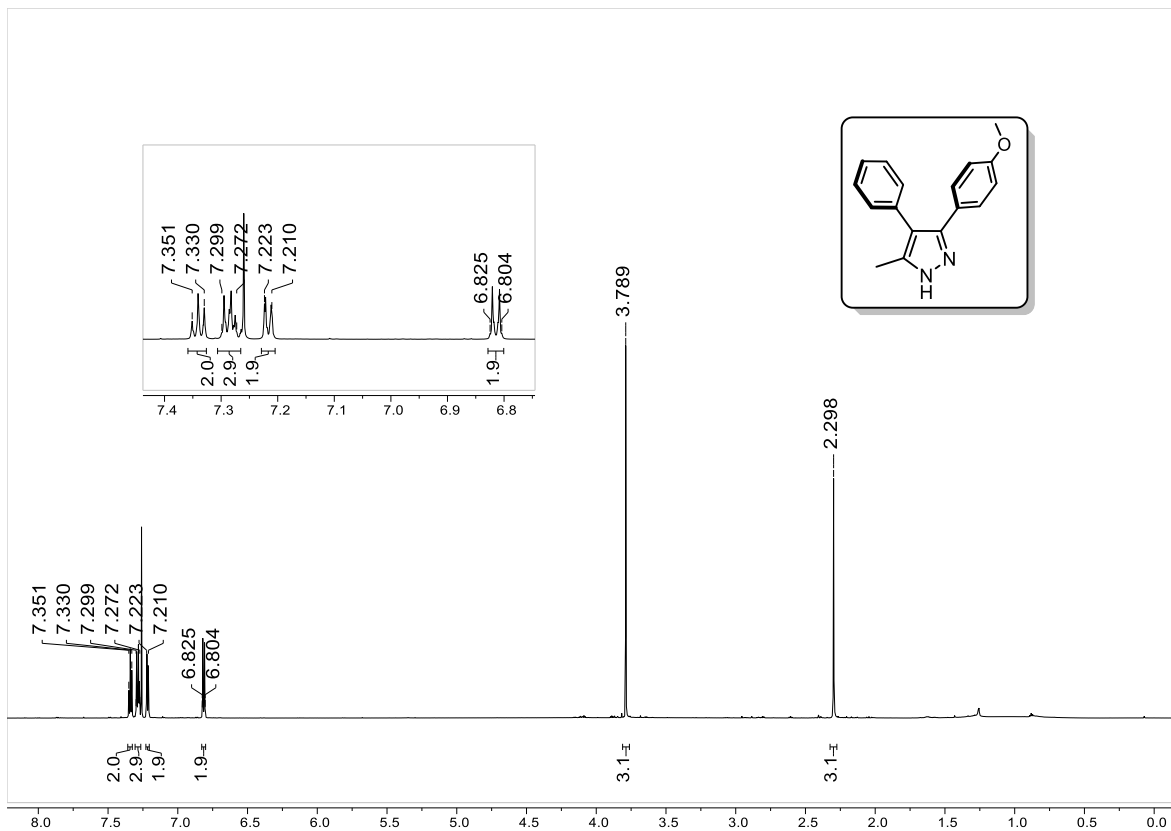
Tolerance:100.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 50.0 (Fraction:.5)

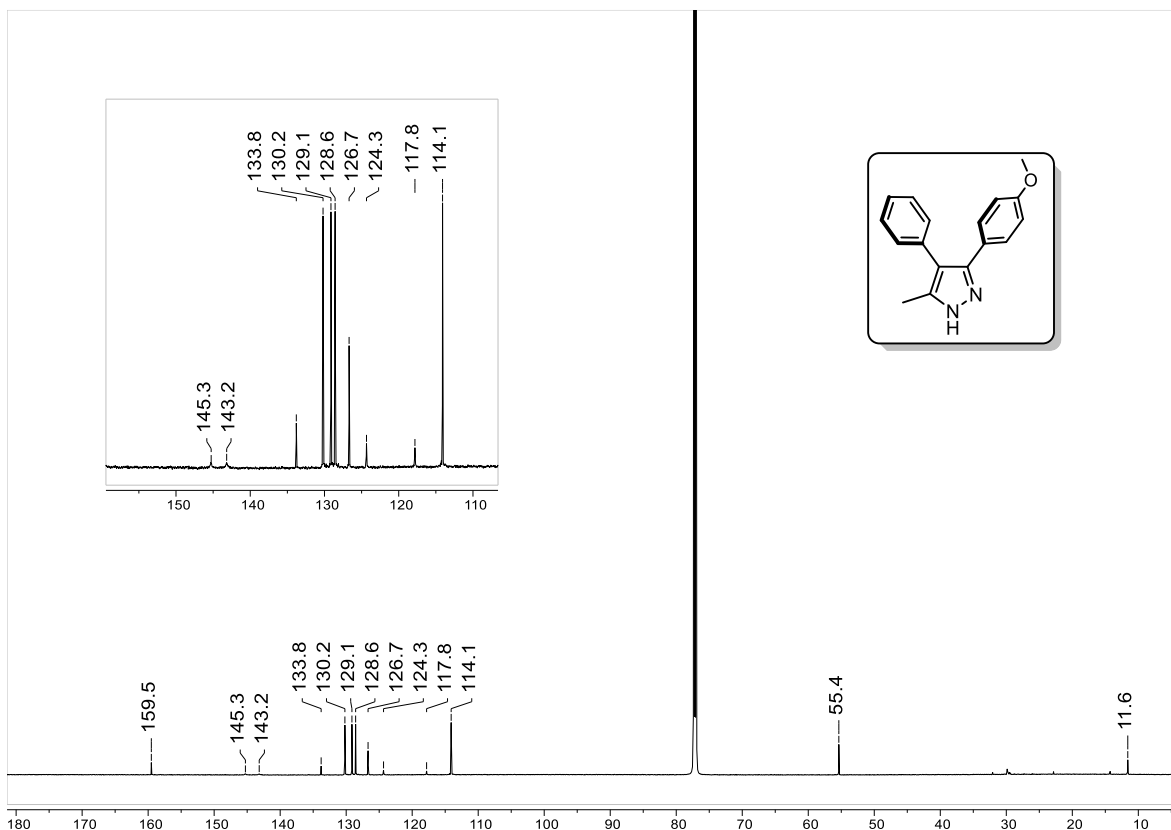


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
277.17019	3715.29	277.17047	-0.28	-1.01	¹² C ₁₉ ¹ H ₂₁ ¹⁴ N ₂	10.5

Spectrum 48. HRMS of 8c.



Spectrum 49. ¹H-NMR of 8d.



Spectrum 50. ^{13}C -NMR of **8d**.

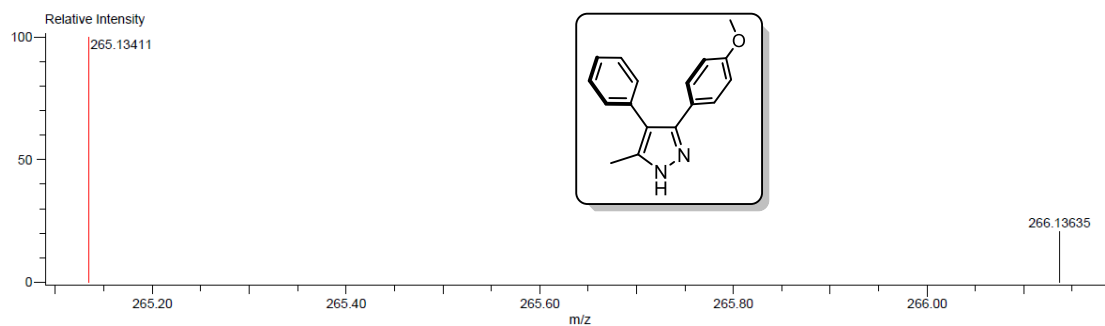
Description:
 Ionization Mode: ESI+
 History: Determine m/z [Peak Detect [Centroid, 30, Area], Correct Base[]; Smooth [5]]; Correct Base [5.0%]; Average [MS[...

Mass Calibration data: Cal_PEG_600
 Created: 2/17/2023 11:40:57 AM
 Created by: AccuTOF

Charge number: 1
 Element: ^{12}C : 0 .. 17, ^1H : 0 .. 21, ^{14}N : 0 .. 2, ^{16}O : 0 .. 1

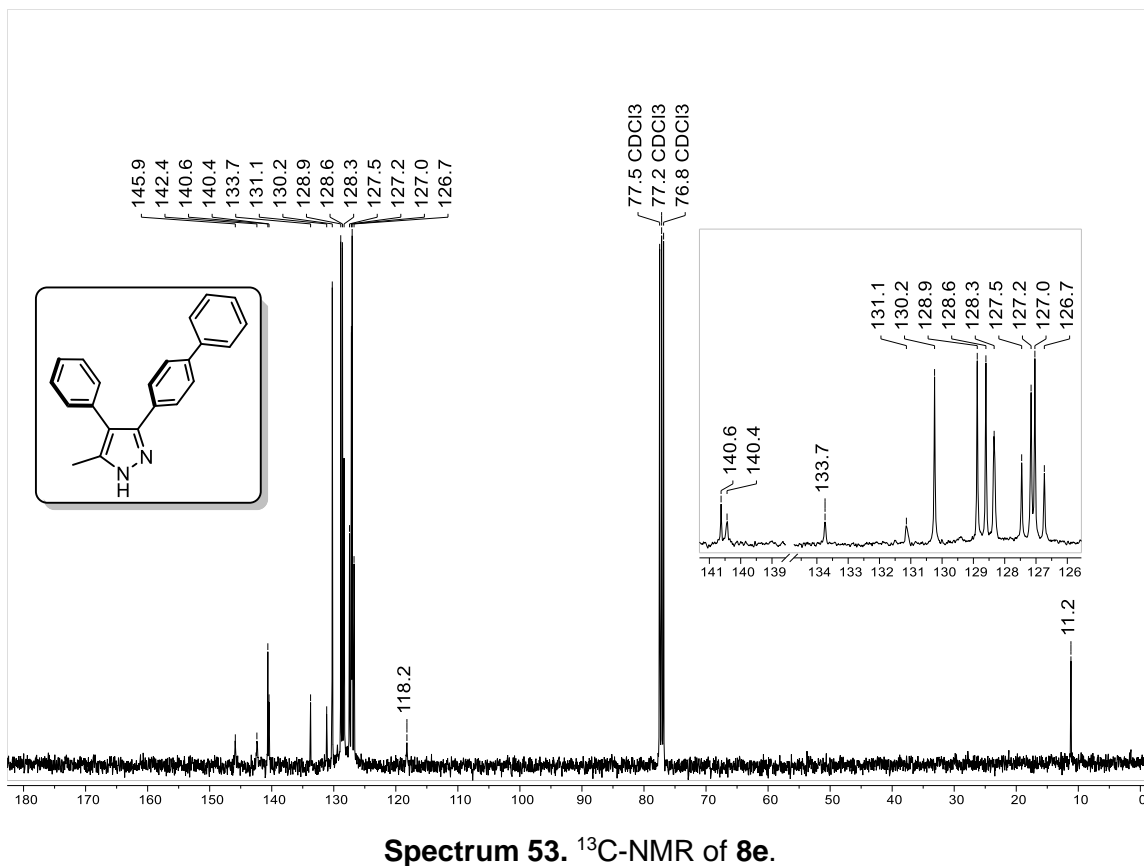
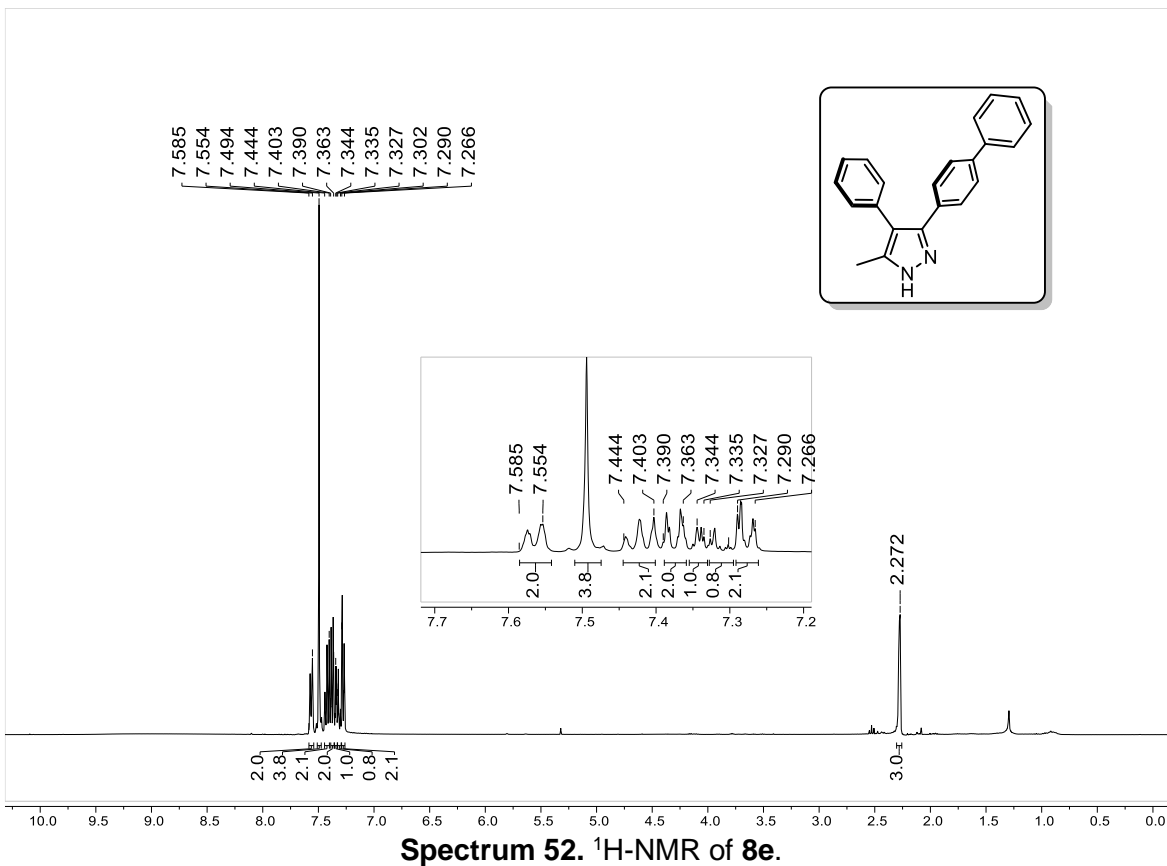
Tolerance: 5.00 (ppm), 5.00 .. 15.00 (mmu)

Unsaturation Number: 0.0 .. 50.0 (Fraction: Both)



Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
265.13411	13136.52	265.13409	0.02	0.08	$^{12}\text{C}_{17}\text{H}_{17}\text{N}_2\text{O}_1$	10.5

Spectrum 51. HRMS of **8d**.



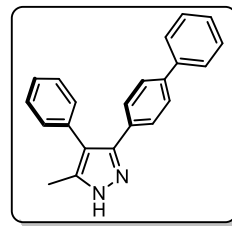
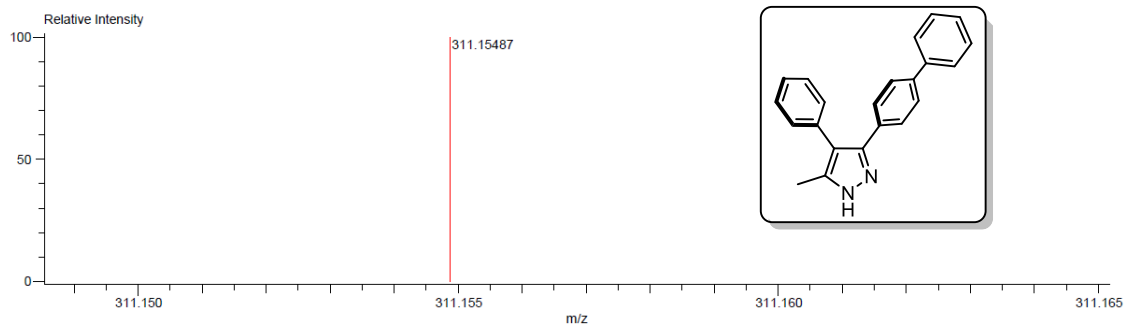
Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5];Correct Base[5.0%];Average(MS[...]

Charge number:1
 Element:¹²C:0 .. 22, ¹H:0 .. 50, ¹⁹F:0 .. 0, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0

Mass Calibration data:Cal_PEG_600
 Created:2/2/2023 10:25:58 AM
 Created by:AccuTOF

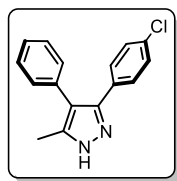
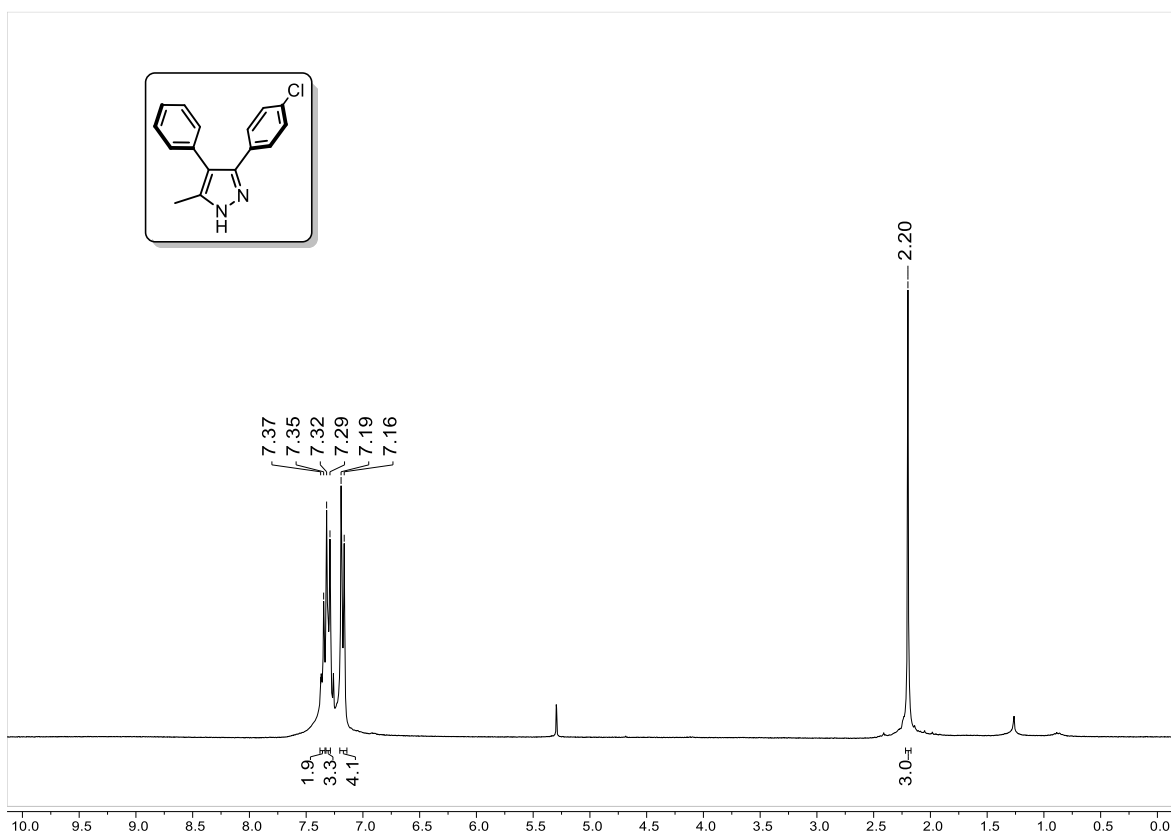
Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 50.0 (Fraction:.5)

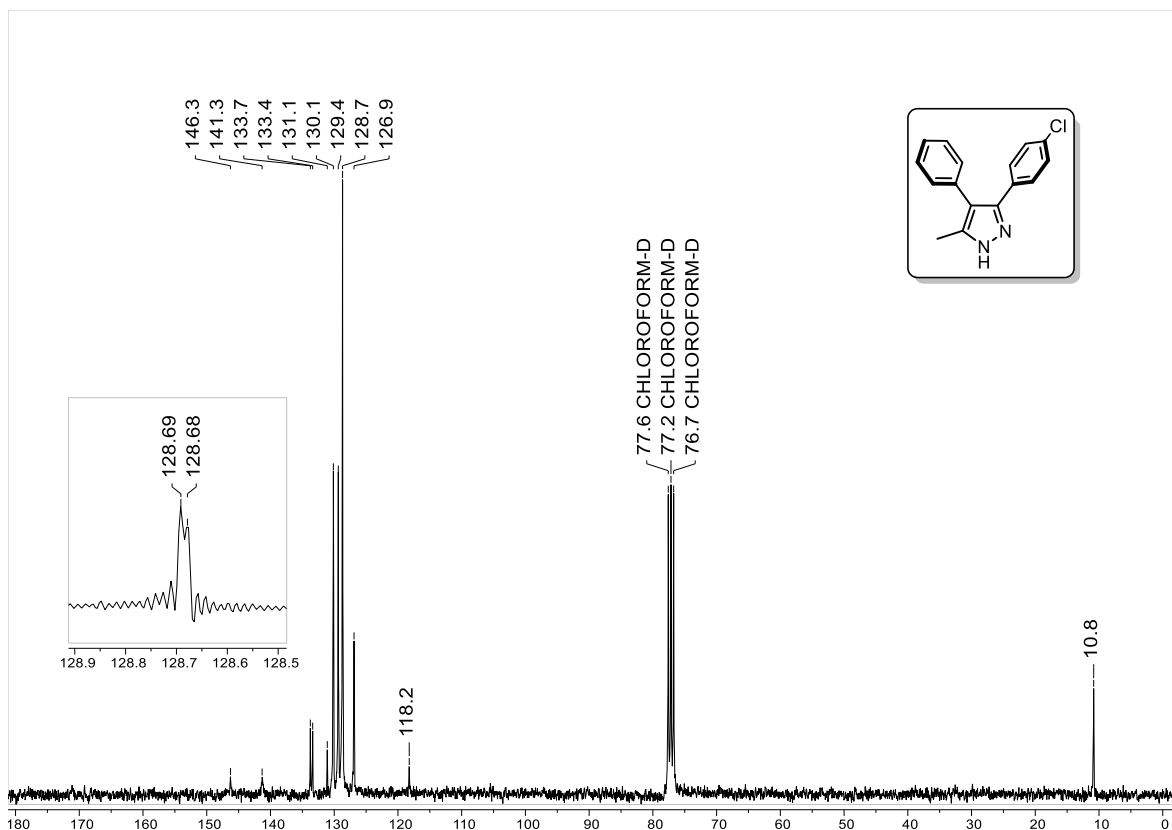


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
311.15487	265293.61	311.15482	0.05	0.15	¹² C ₂₂ ¹ H ₁₉ ¹⁴ N ₂	14.5

Spectrum 54. HRMS of 8e.



Spectrum 55. ¹H-NMR of 8f.



Spectrum 56. ¹³C-NMR of 8f.

Description:

Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600

Created:2/3/2023 12:43:22 PM

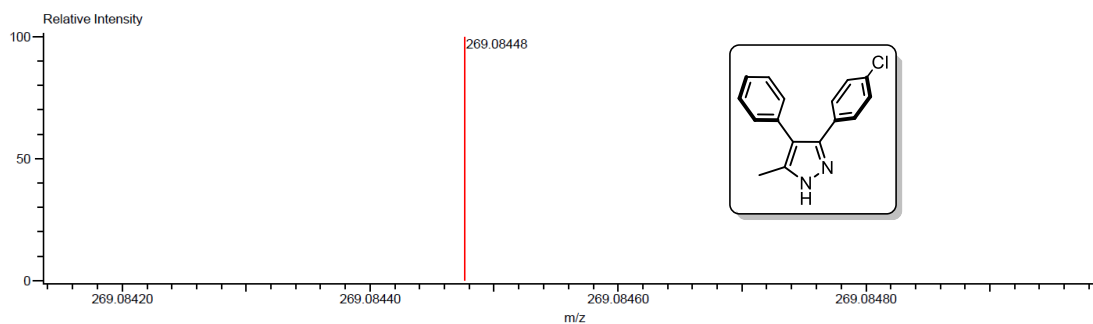
Created by:AccuTOF

Charge number:1

Tolerance:100.00(ppm), 5.00 .. 15.00(mmu)

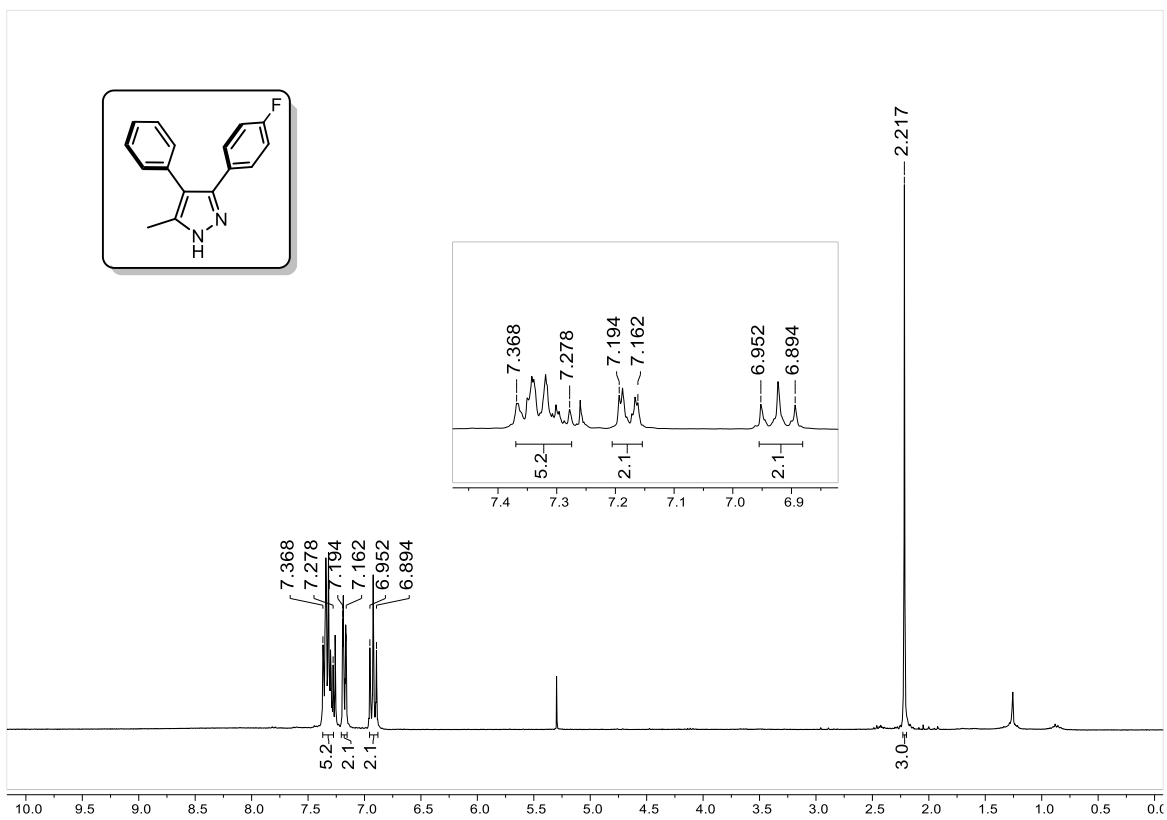
Unsaturation Number:0.0 .. 50.0 (Fraction:.5)

Element:¹²C:0 .. 19, ¹H:0 .. 50, ³⁵Cl:0 .. 1, ¹⁹F:0 .. 0, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0

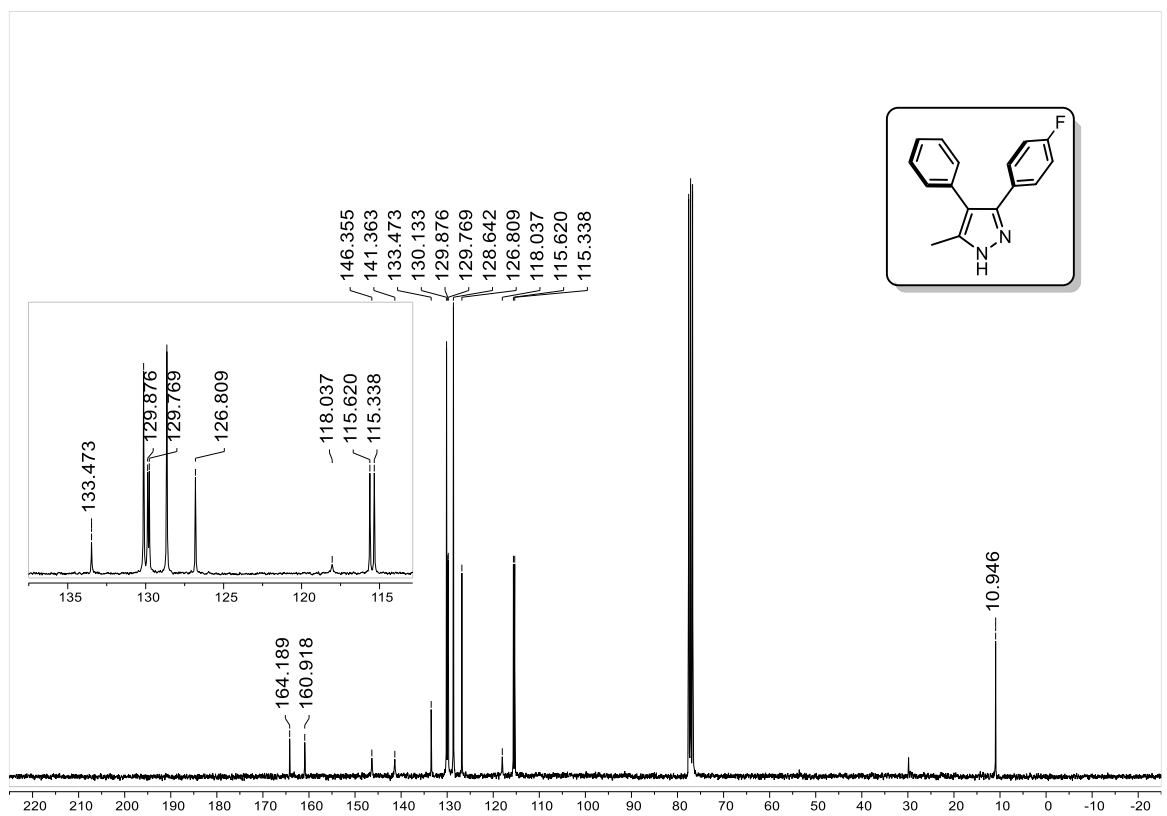


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
269.08448	878215.03	269.08455	-0.07	-0.28	¹² C ₁₆ ¹ H ₁₄ ³⁵ Cl ₁ ¹⁴ N ₂	10.5

Spectrum 57. HRMS of 8f.



Spectrum 58. ¹H-NMR of 8g.



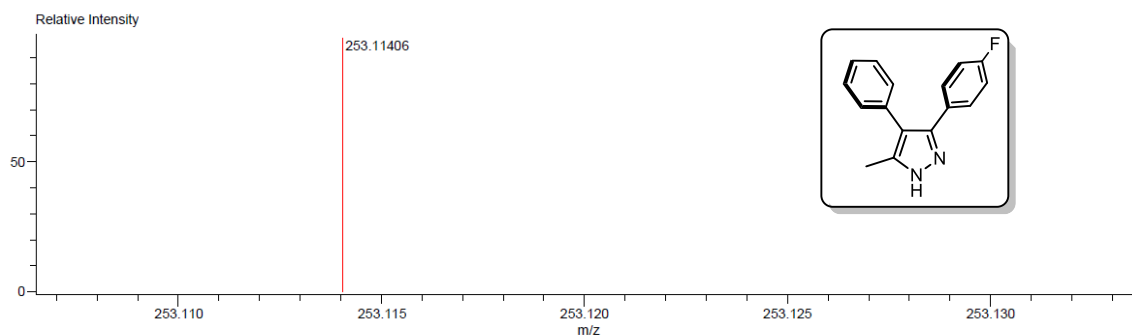
Spectrum 59. ¹³C-NMR of 8g.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area],Correct Base[],Smooth[5]],Correct Base[5.0%],Average(MS[...

Mass Calibration data:Cal_PEG_600
 Created:2/15/2023 11:15:26 AM
 Created by:

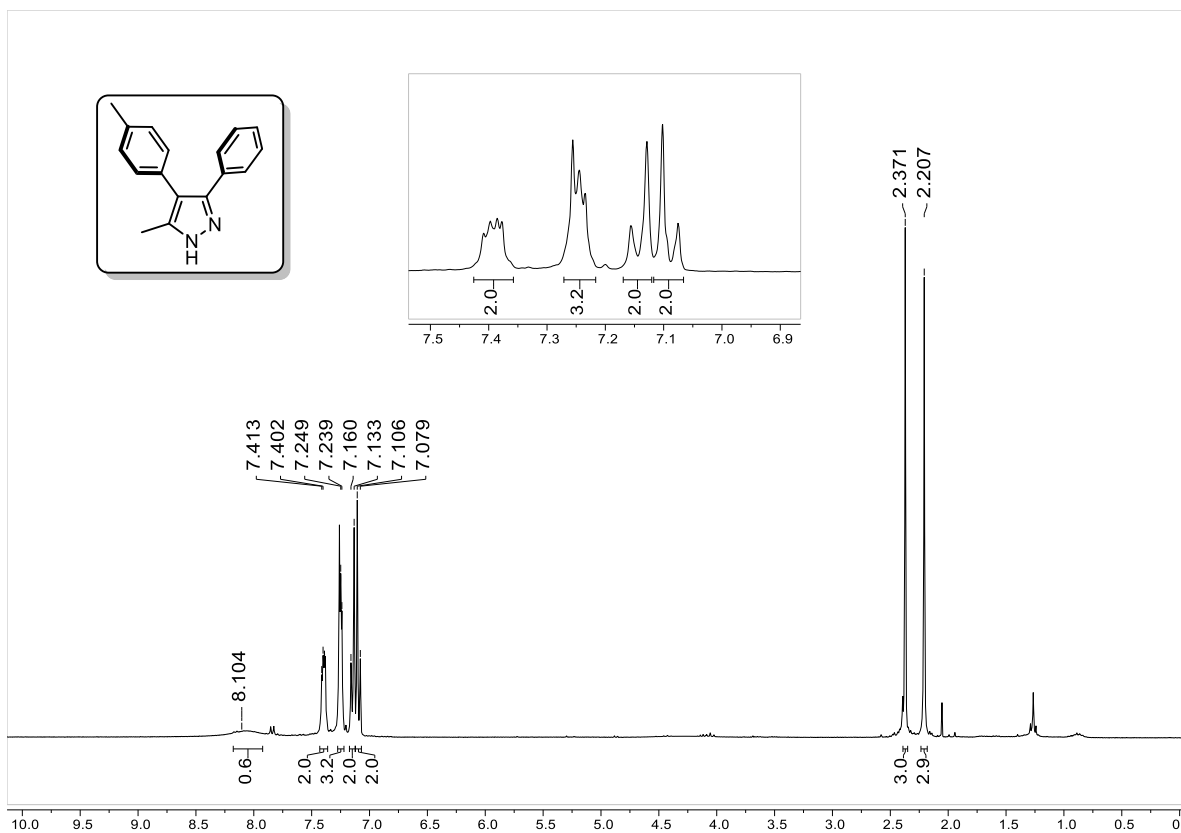
Charge number:1
 Element:¹²C:0 .. 16, ¹H:0 .. 22, ³⁵Cl:0 .. 0, ¹⁹F:0 .. 1, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0
 Tolerance:100.00(mmu)

Unsaturation Number:0.0 .. 20.0 (Fraction:Both)

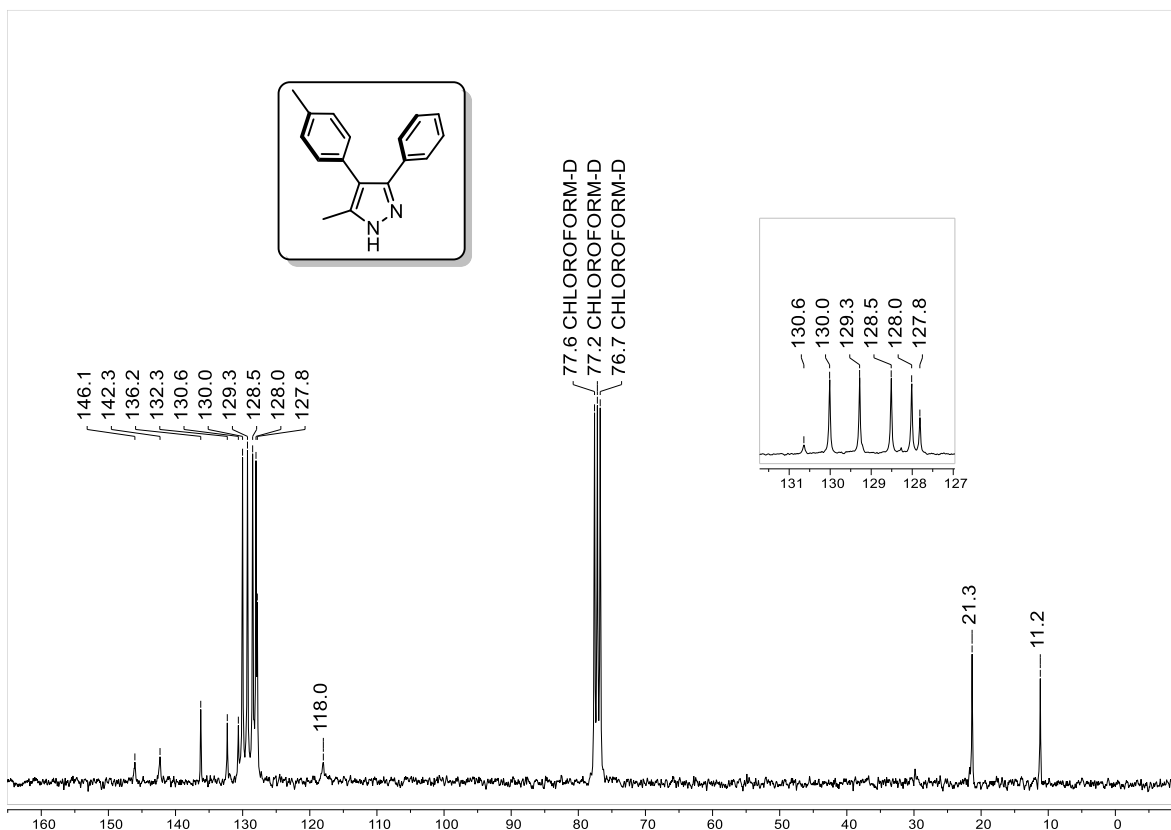


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
253.11406	762947.17	253.11410	-0.04	-0.17	¹² C ₁₆ ¹ H ₁₄ ¹⁹ F ₁ ¹⁴ N ₂	10.5

Spectrum 60. HRMS of 8g.



Spectrum 61. ¹H-NMR of 8h.



Spectrum 62. ^{13}C -NMR of **8h**.

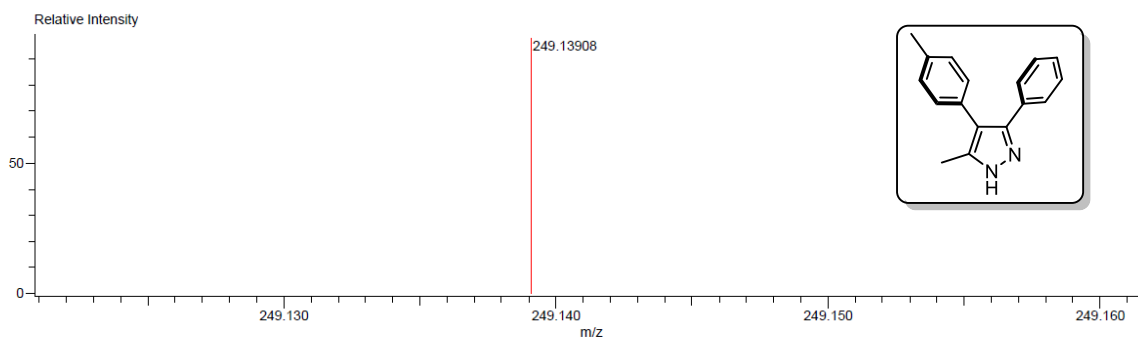
Description:
 Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
 Created:1/24/2023 2:20:24 PM
 Created by:AccuTOF

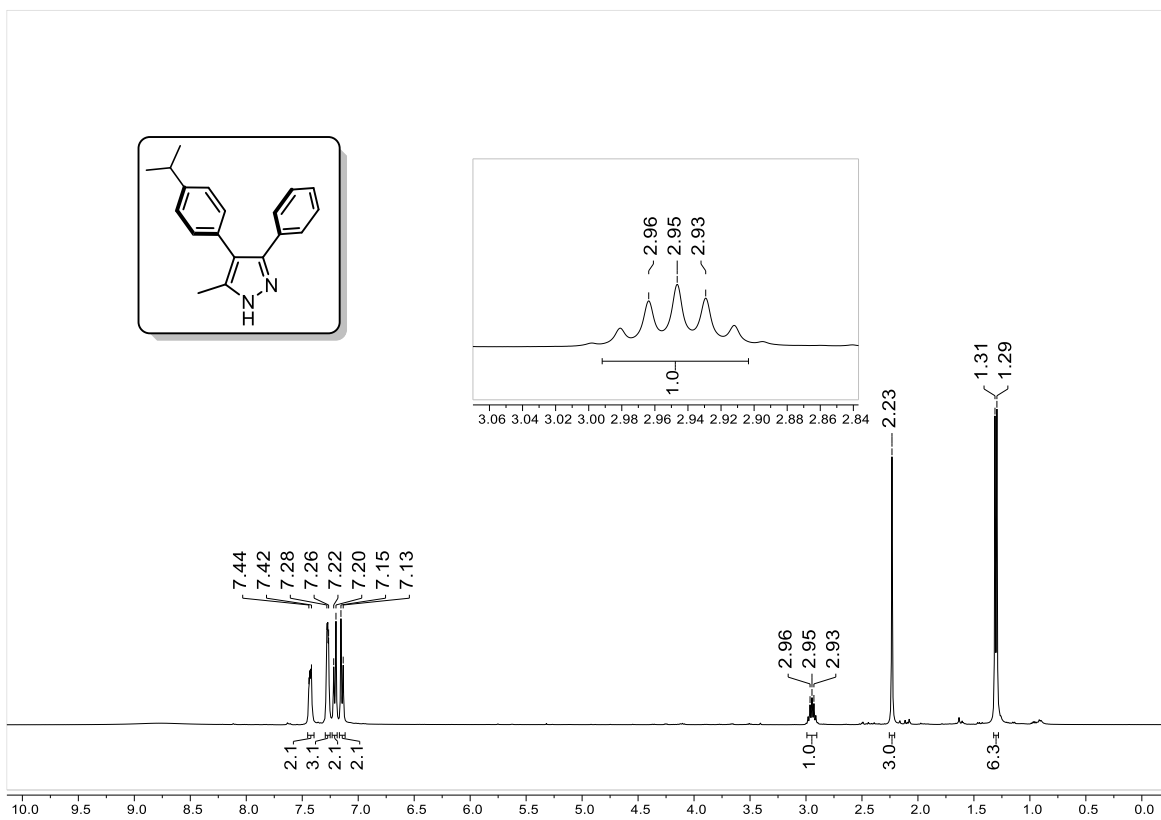
Charge number:1
 Tolerance:5.00(ppm), 5.00 .. 15.00(mmu)
 Element: ^{12}C :0 .. 17, ^1H :0 .. 20, ^{35}Cl :0 .. 0, ^{14}N :0 .. 2, ^{16}O :0 .. 0

Unsaturation Number:0.0 .. 45.0 (Fraction:Both)

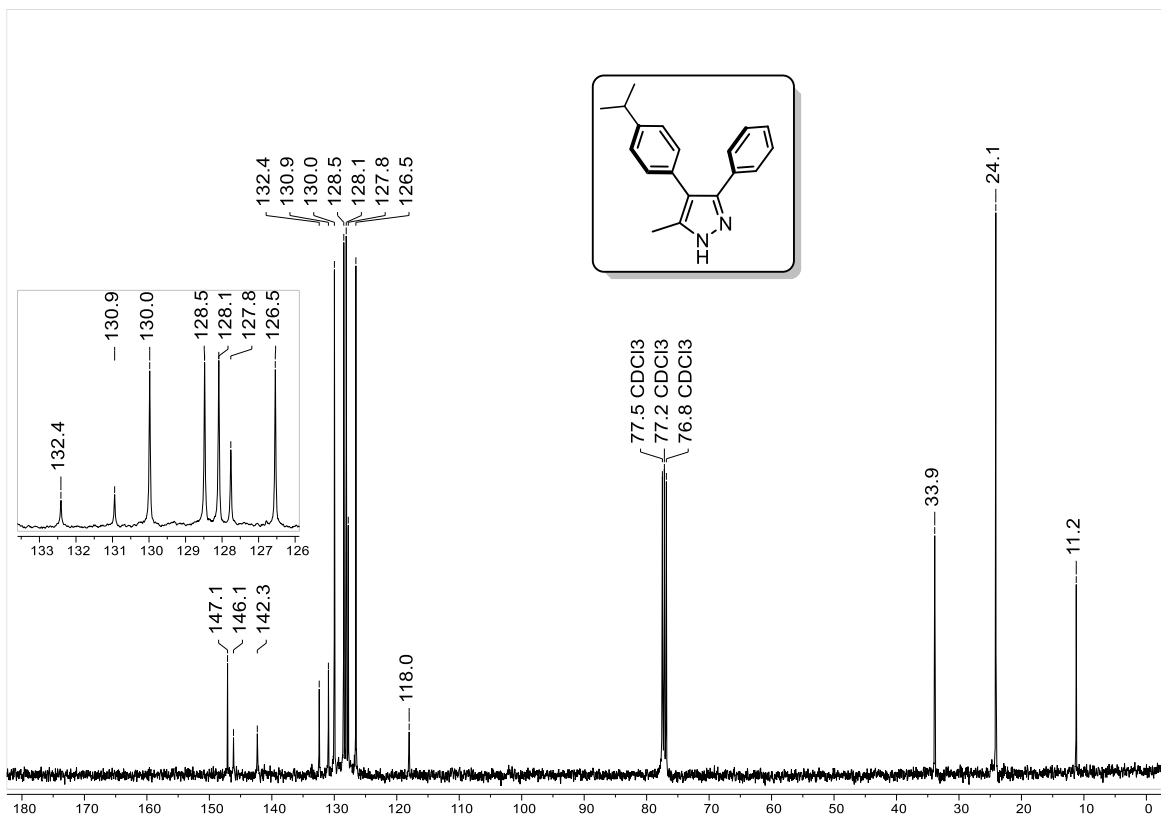


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
249.13908	3737.98	249.13917	-0.09	-0.37	$^{12}\text{C}_{17}\text{H}_{17}\text{N}_2$	10.5

Spectrum 63. HRMS of **8h**.



Spectrum 64. ¹H-NMR of 8i.

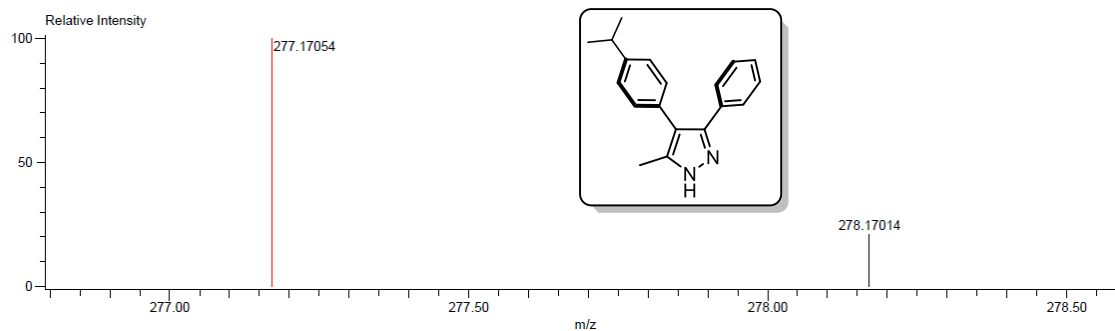


Spectrum 65. ¹³C-NMR of 8i.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5];Correct Base[5.0%];Average[MS[...

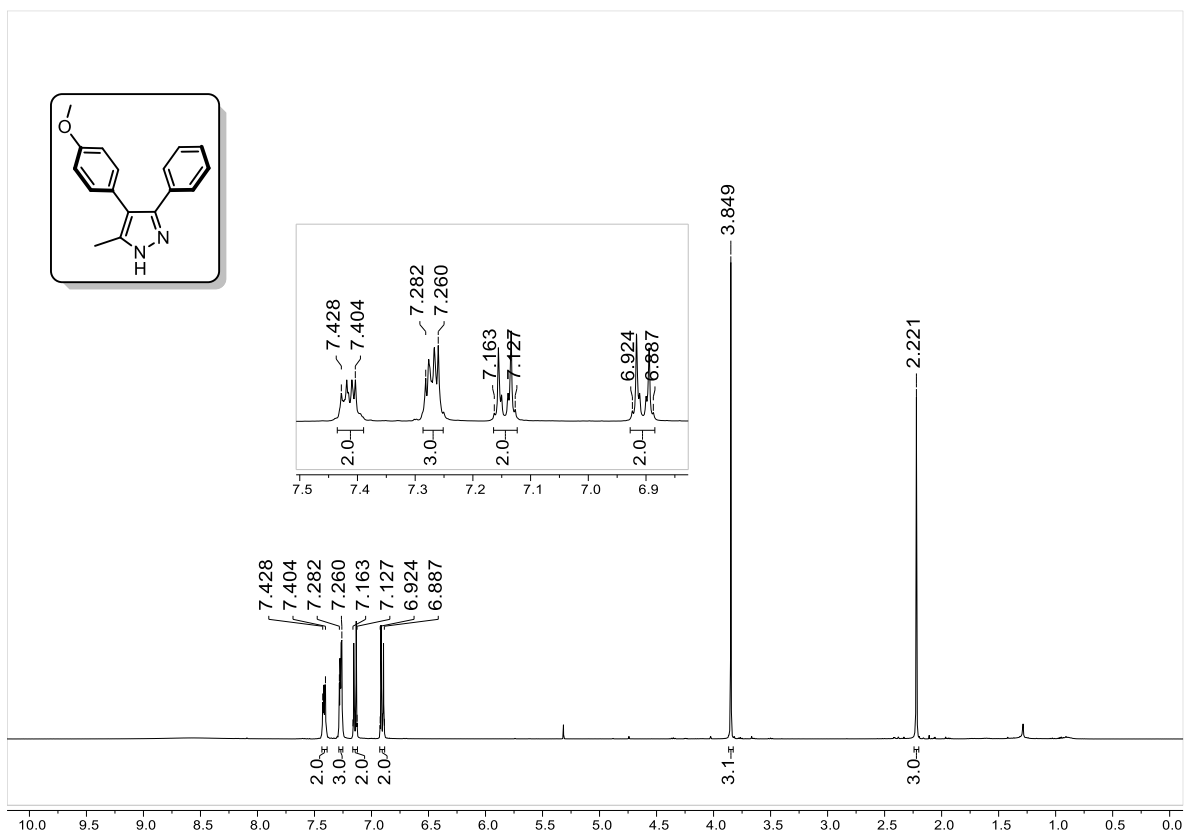
Mass Calibration data:Cal_PEG_600
 Created:2/17/2023 11:37:13 AM
 Created by:AccuTOF

Charge number:1 Tolerance:5.00(ppm), 5.00 .. 15.00(mmu) Unsaturation Number:0.0 .. 50.0 (Fraction:Both)
 Element:¹²C:0 .. 19, ¹H:0 .. 21, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0

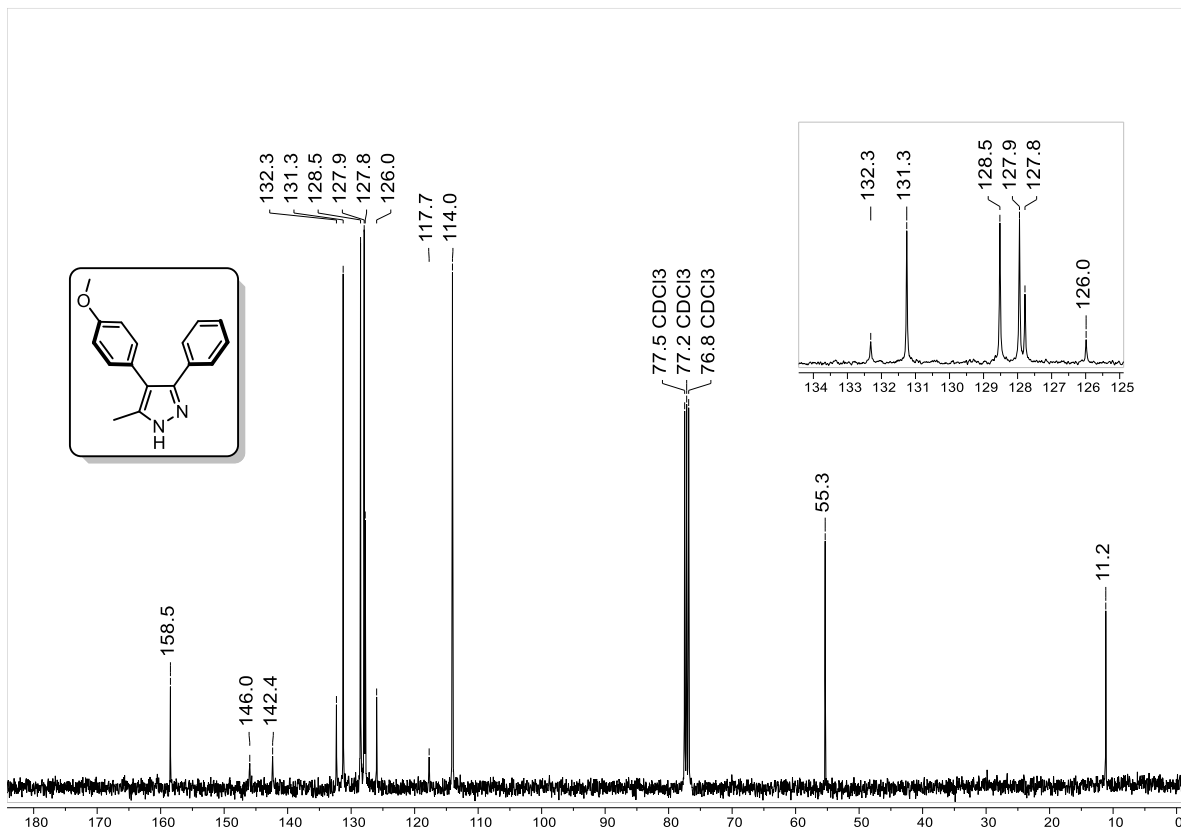


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
277.17054	8995.62	277.17047	0.07	0.24	¹² C ₁₉ ¹ H ₂₁ ¹⁴ N ₂	10.5

Spectrum 66. HRMS of 8i.



Spectrum 67. ¹H-NMR of 8j.



Spectrum 68. ¹³C-NMR of 8j.

Description:

Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area],Correct Base[],Smooth[5]],Correct Base[5.0%];Average[MS[...

Mass Calibration data:CaI_PEG_600

Created:2/23/2023 12:48:51 PM

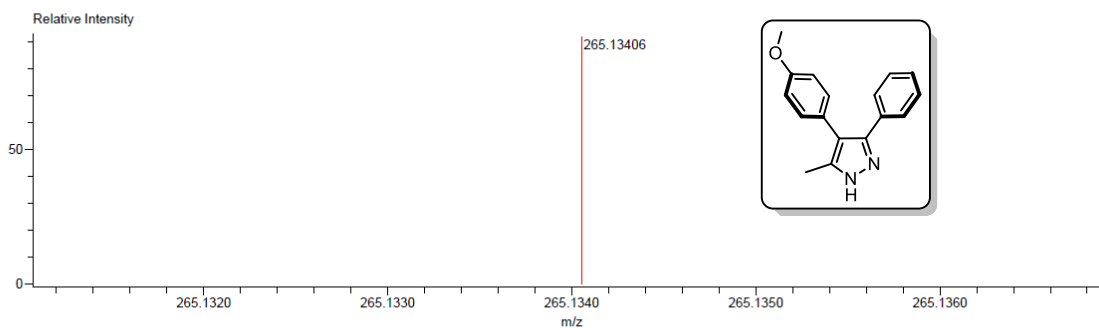
Created by:AccuTOF

Charge number:1

Tolerance:4.00(ppm), 5.00 .. 15.00(mmu)

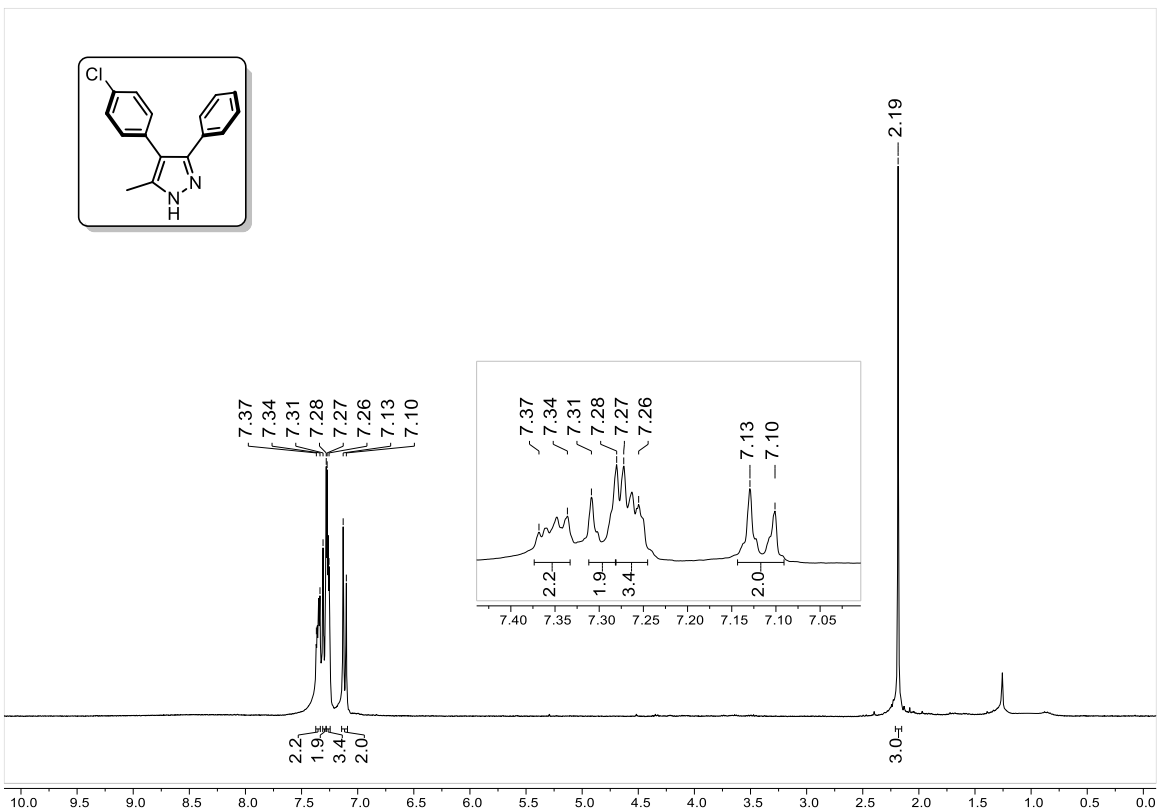
Unsaturation Number:0.0 .. 40.0 (Fraction:Both)

Element:¹²C:0 .. 17, ¹H:0 .. 18, ¹⁴N:0 .. 2, ¹⁶O:0 .. 1

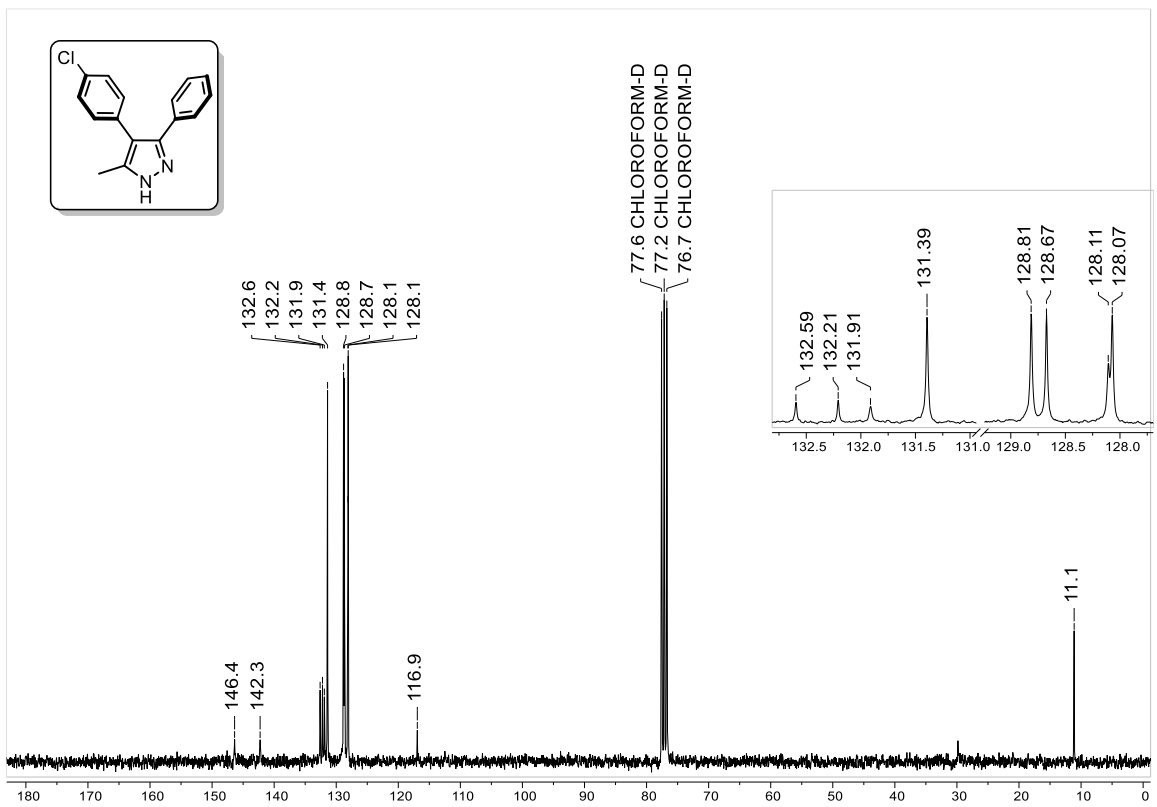


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
265.13406	3779.12	265.13409	-0.03	-0.12	¹² C ₁₇ ¹ H ₁₇ ¹⁴ N ₂ ¹⁶ O ₁	10.5

Spectrum 69. HRMS of 8j.



Spectrum 70. $^1\text{H-NMR}$ of 8k.



Spectrum 71. $^{13}\text{C-NMR}$ of 8k.

Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area],Correct Base[],Smooth[5]],Correct Base[5.0%];Average[MS[...

Created:1/25/2023 10:56:47 AM

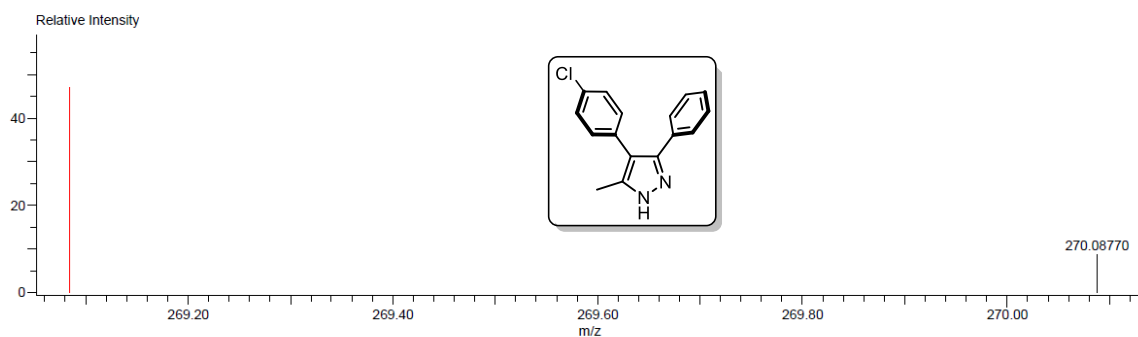
Created by:AccuTOF

Charge number:1

Tolerance:3.00(ppm), 5.00 .. 15.00(mmu)

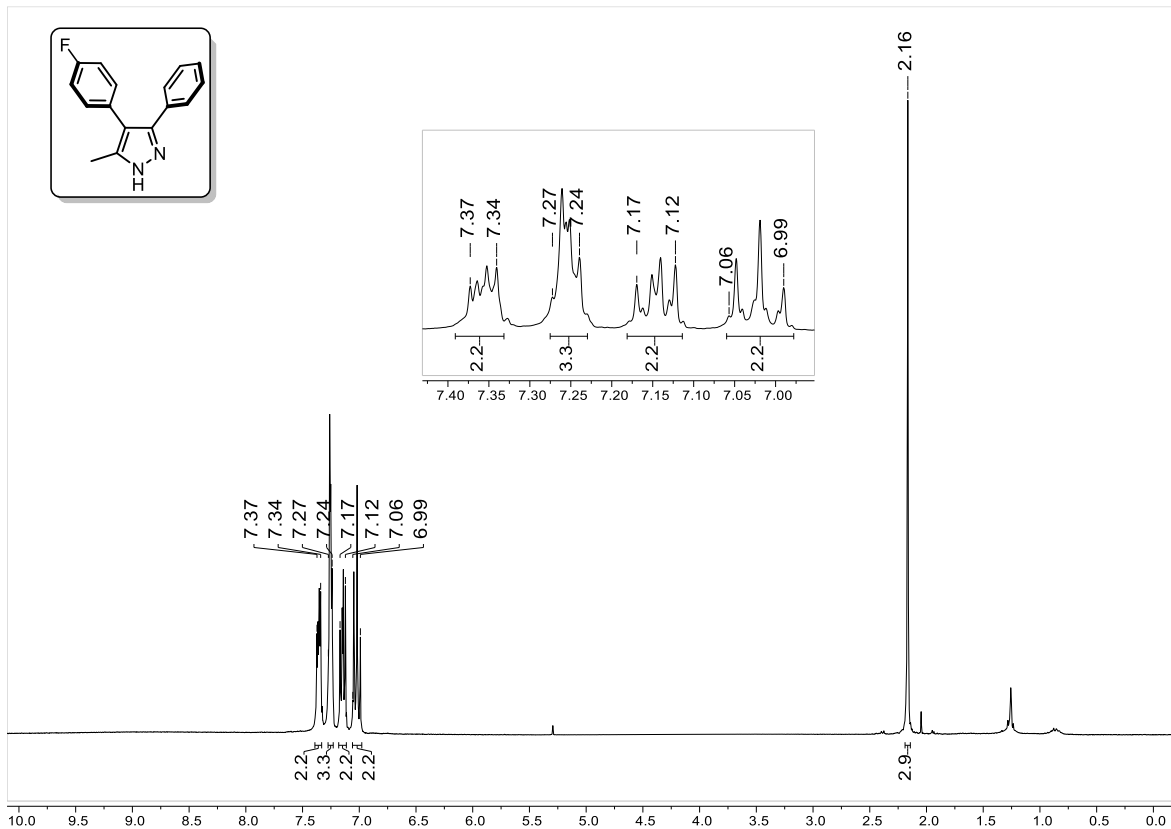
Unsaturation Number:0.0 .. 45.0 (Fraction:Both)

Element:¹²C:0 .. 16, ¹H:0 .. 20, ³⁵Cl:0 .. 1, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0

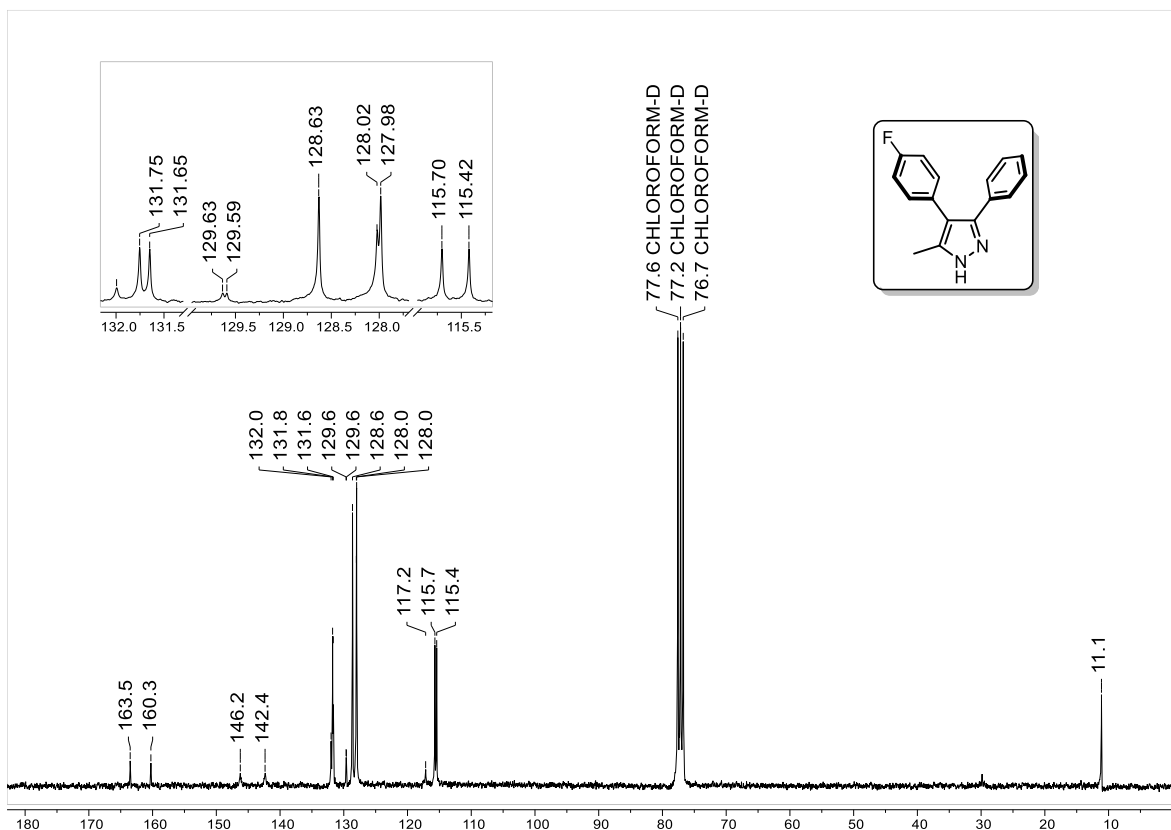


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
269.08460	6551.40	269.08455	0.05	0.18	¹² C ₁₆ ¹ H ₁₄ ³⁵ Cl ₁ ¹⁴ N ₂	10.5

Spectrum 72. HRMS of 8k.



Spectrum 73. ¹H-NMR of 8l.



Spectrum 74. ¹³C-NMR of 8I.

Description:

Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600

Created:2/3/2023 12:41:09 PM

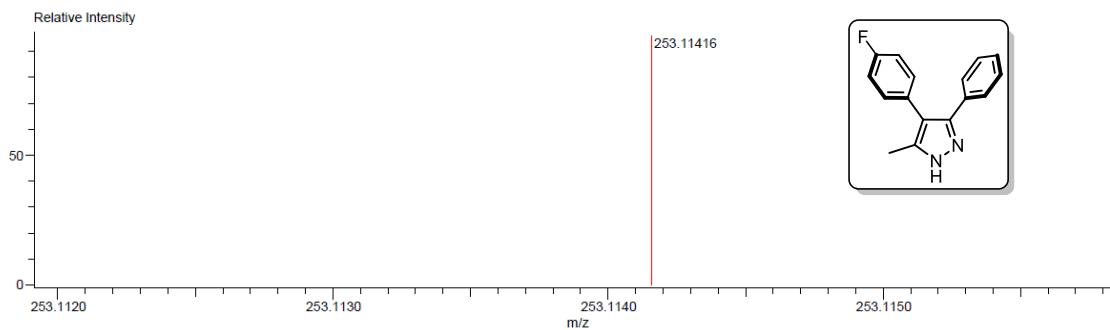
Created by:AccuTOF

Charge number:1

Tolerance:100.00(ppm), 5.00 .. 15.00(mmu)

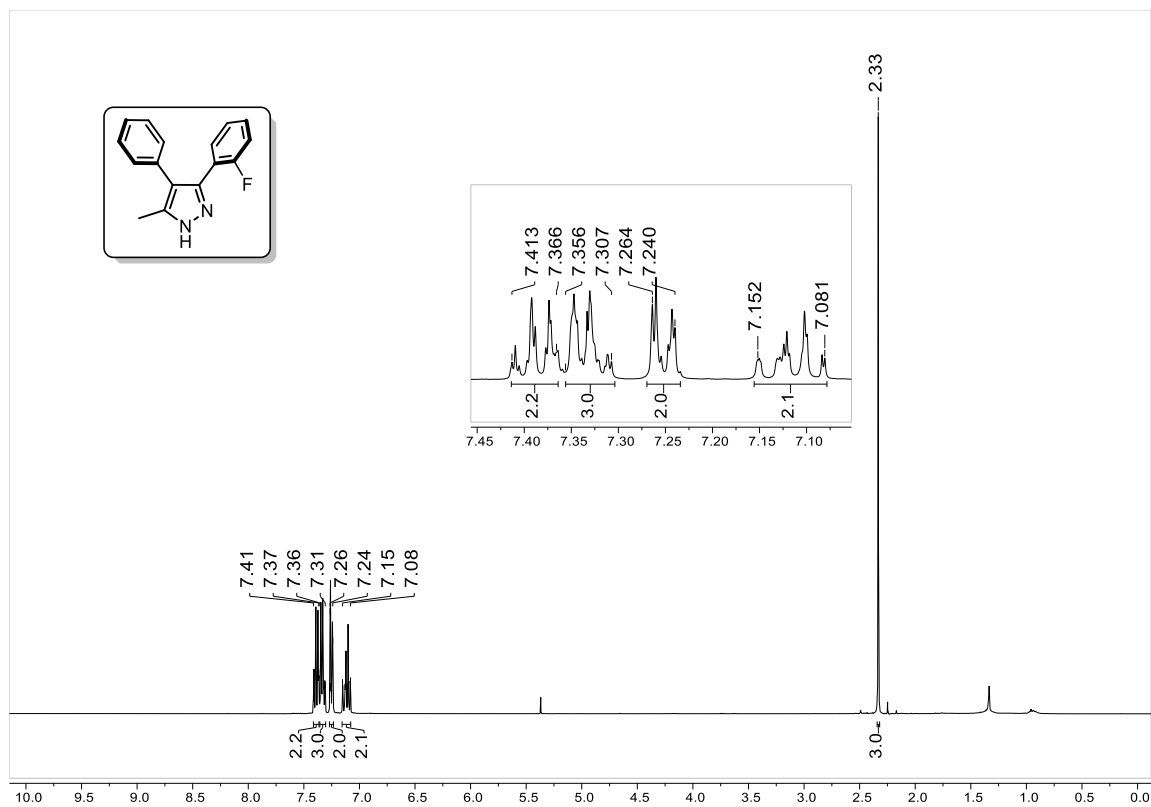
Unsaturation Number:0.0 .. 50.0 (Fraction:.5)

Element:¹²C:0 .. 19, ¹H:0 .. 50, ¹⁹F:0 .. 1, ¹⁴N:0 .. 2, ¹⁸O:0 .. 0

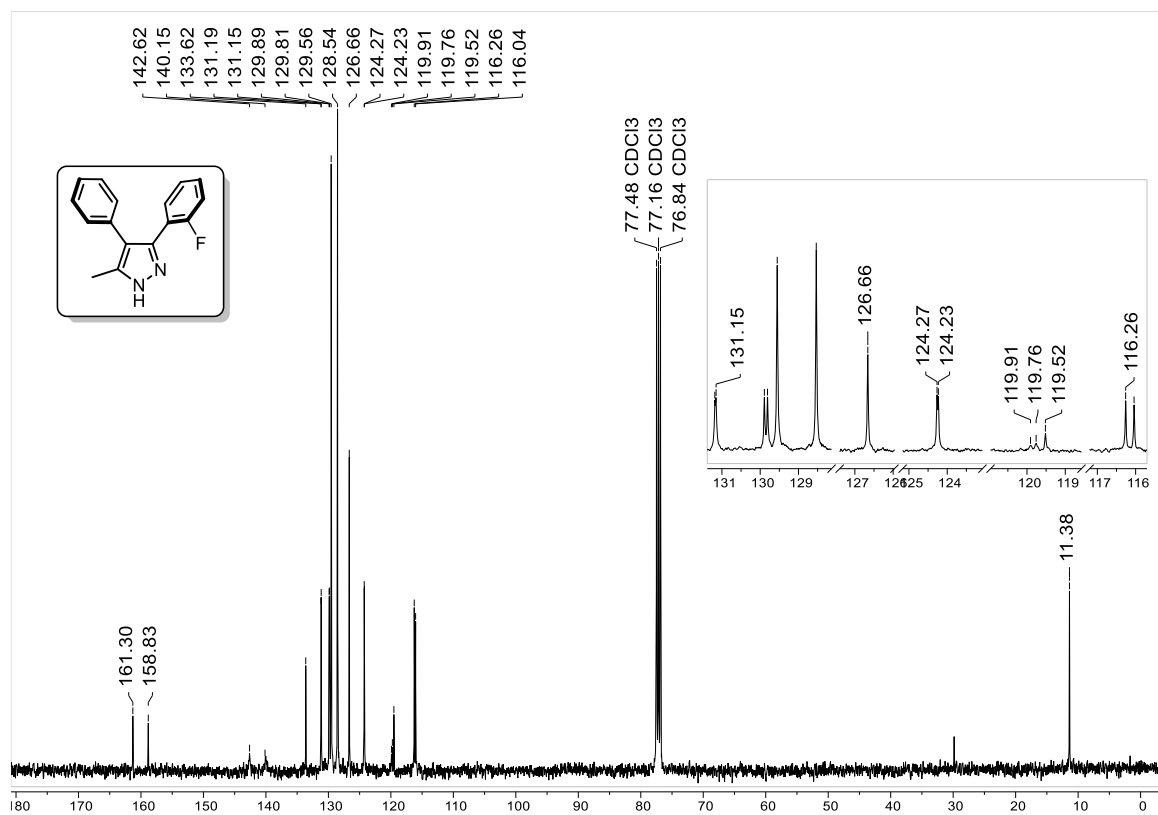


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
253.11416	844768.97	253.11410	0.06	0.23	¹² C ₁₆ ¹ H ₁₄ ¹⁹ F ₁ ¹⁴ N ₂	10.5

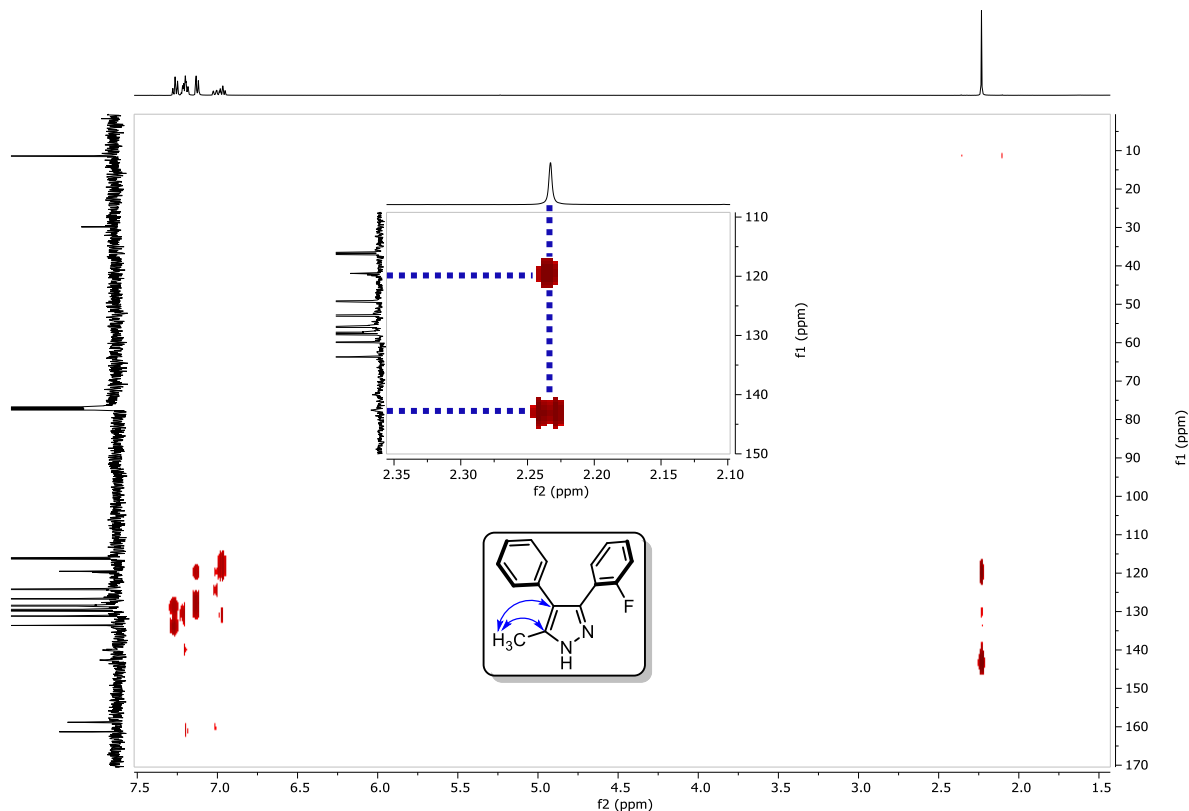
Spectrum 75. HRMS of 8I.



Spectrum 76. ¹H-NMR of 8m.



Spectrum 77. ¹³C-NMR of 8m.



Spectrum 78. HMBC of 8m.

Description:
Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:CaI_PEG_600

Created:3/10/2023 9:36:56 AM

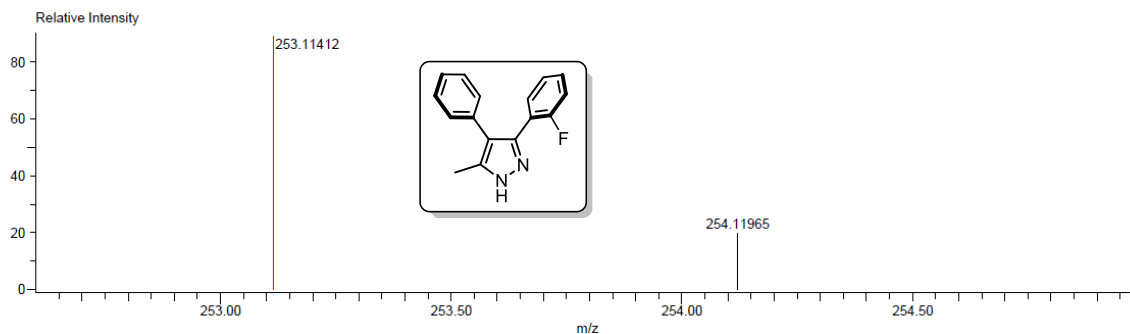
Created by:AccuTOF

Charge number:1

Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)

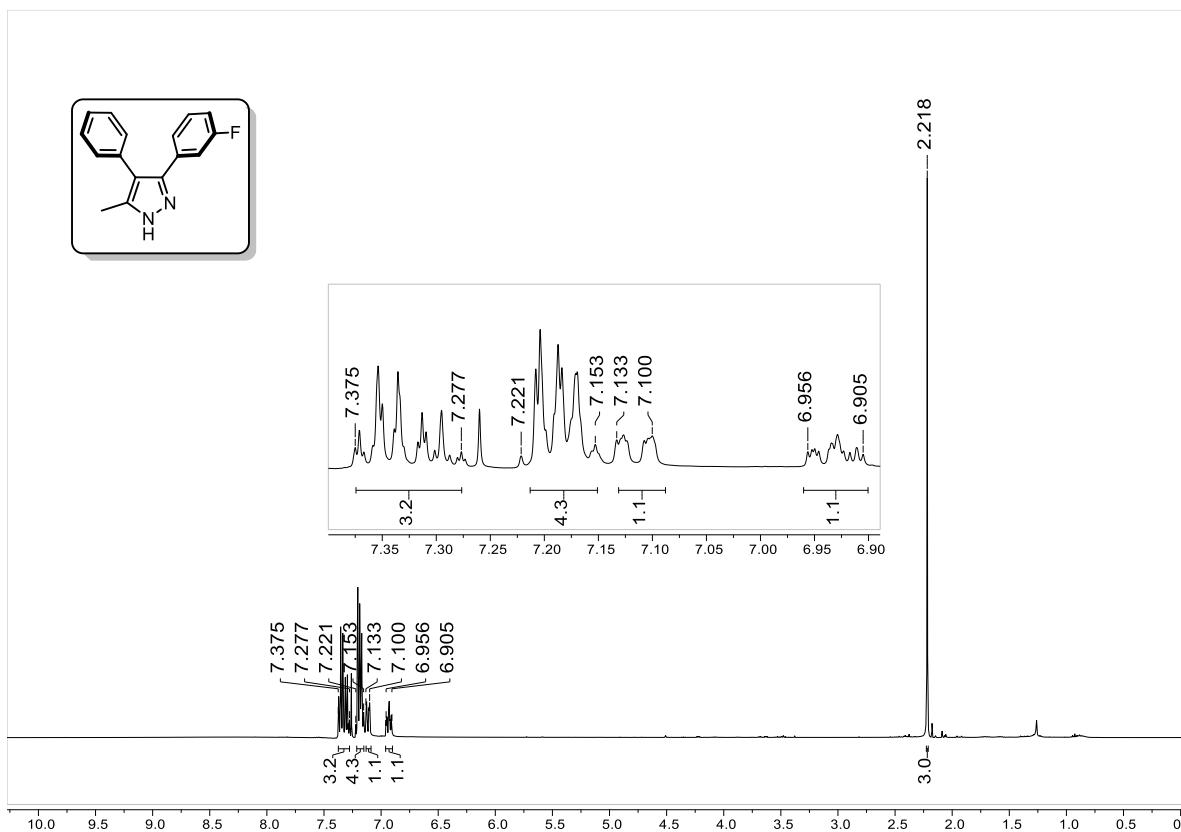
Unsaturation Number:0.0 .. 50.0 (Fraction:Both)

Element:¹²C:0 .. 16, ¹H:0 .. 50, ³⁵Cl:0 .. 0, ¹⁹F:0 .. 1, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0, ³²S:0 .. 0

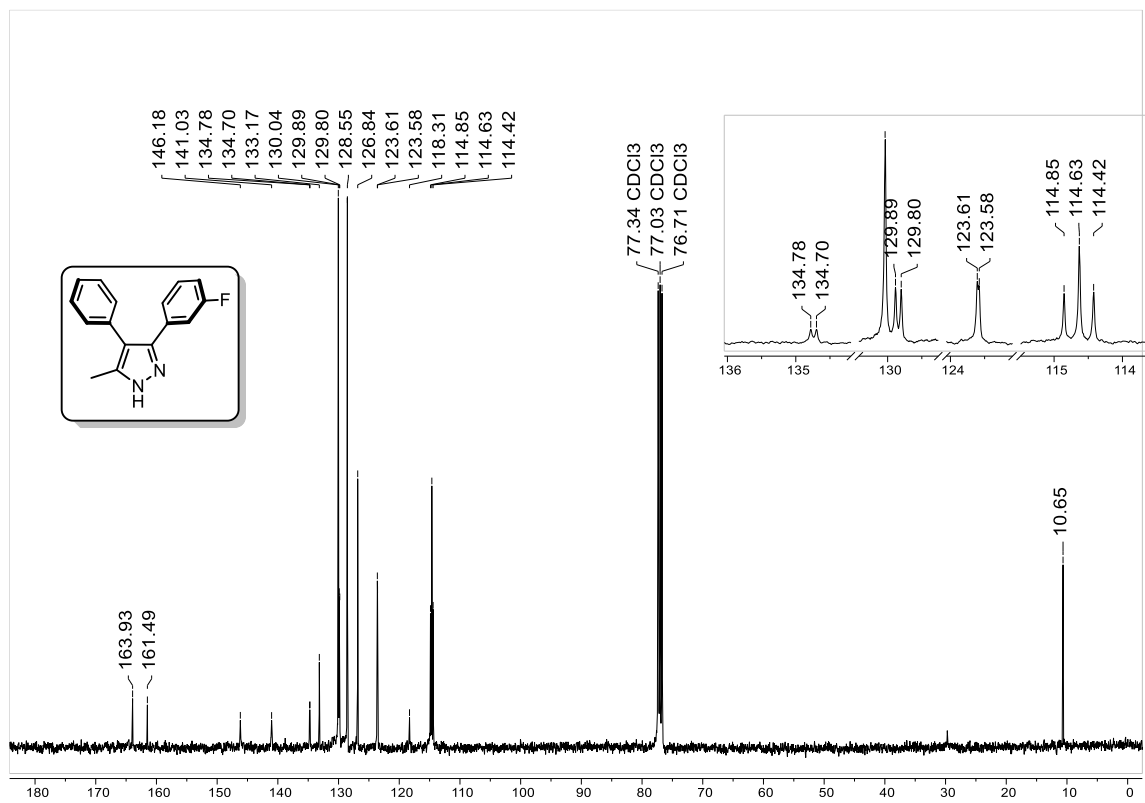


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
253.11412	14151.66	253.11410	0.02	0.06	¹² C ₁₆ ¹ H ₁₄ ¹⁹ F ₁ ¹⁴ N ₂	10.5

Spectrum 79. HRMS of 8m.



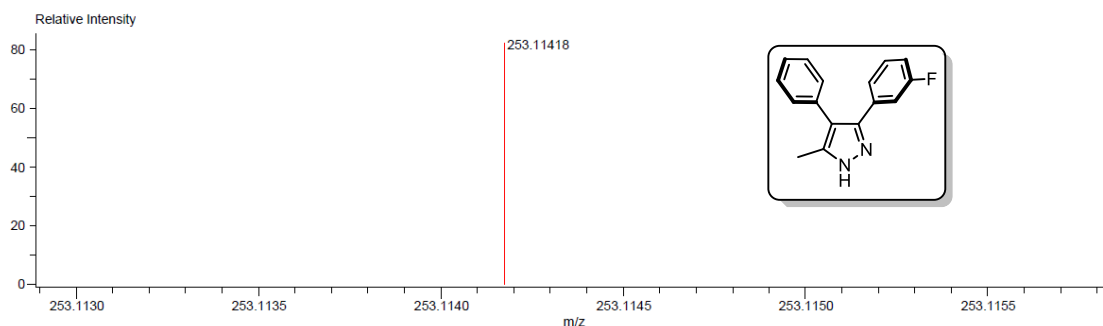
Spectrum 80. ¹H-NMR of 8n.



Spectrum 81. ¹³C-NMR of 8n.

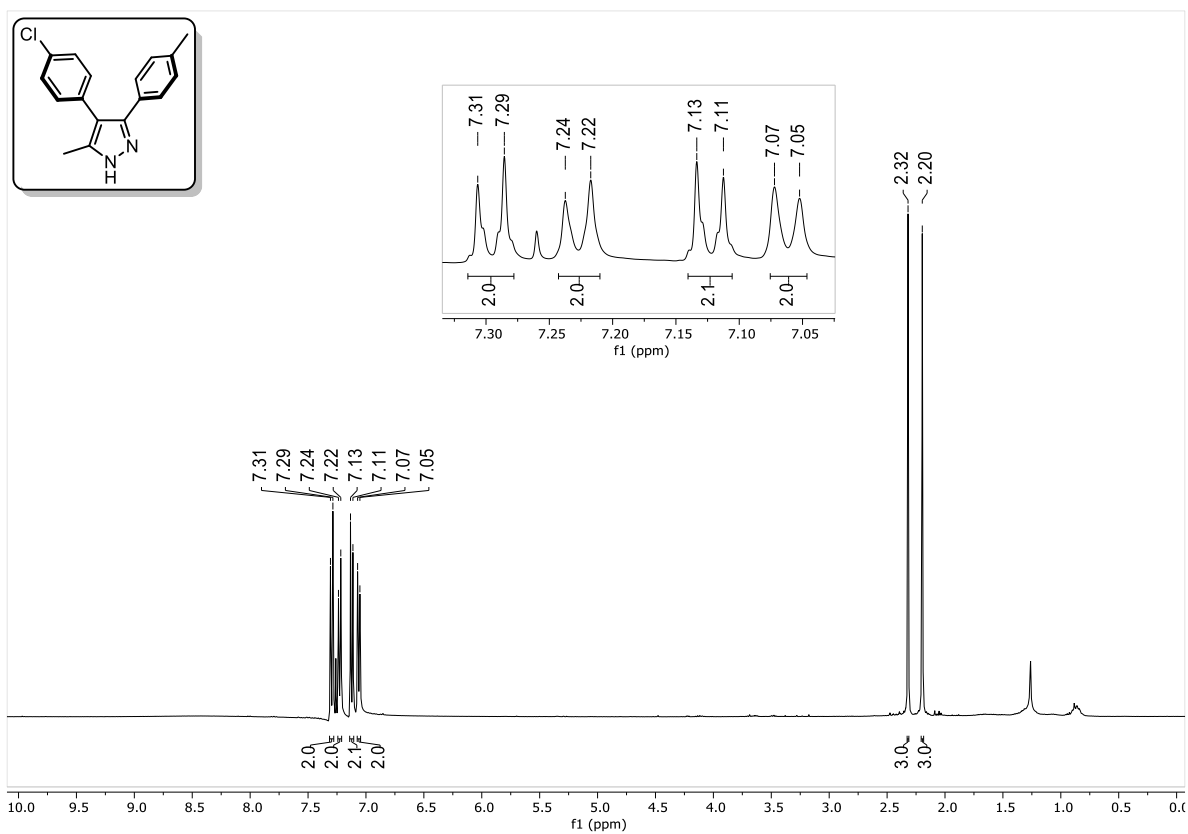
Description:
 Ionization Mode: ESI+
 History: Determine m/z [Peak Detect [Centroid, 30, Area], Correct Base [], Smooth [5]], Correct Base [5.0%], Average (MS...
 Charge number: 1
 Element: ¹²C: 0 .. 16, ¹H: 0 .. 20, ³⁵Cl: 0 .. 1, ¹⁹F: 1 .. 1, ¹⁴N: 2 .. 2

Mass Calibration data: Cal_PEG_600
 Created: 2/27/2023 2:14:34 PM
 Created by: AccuTOF
 Tolerance: 200.00 (ppm), 5.00 .. 15.00 (mmu)
 Unsaturation Number: 0.0 .. 30.0 (Fraction: Both)

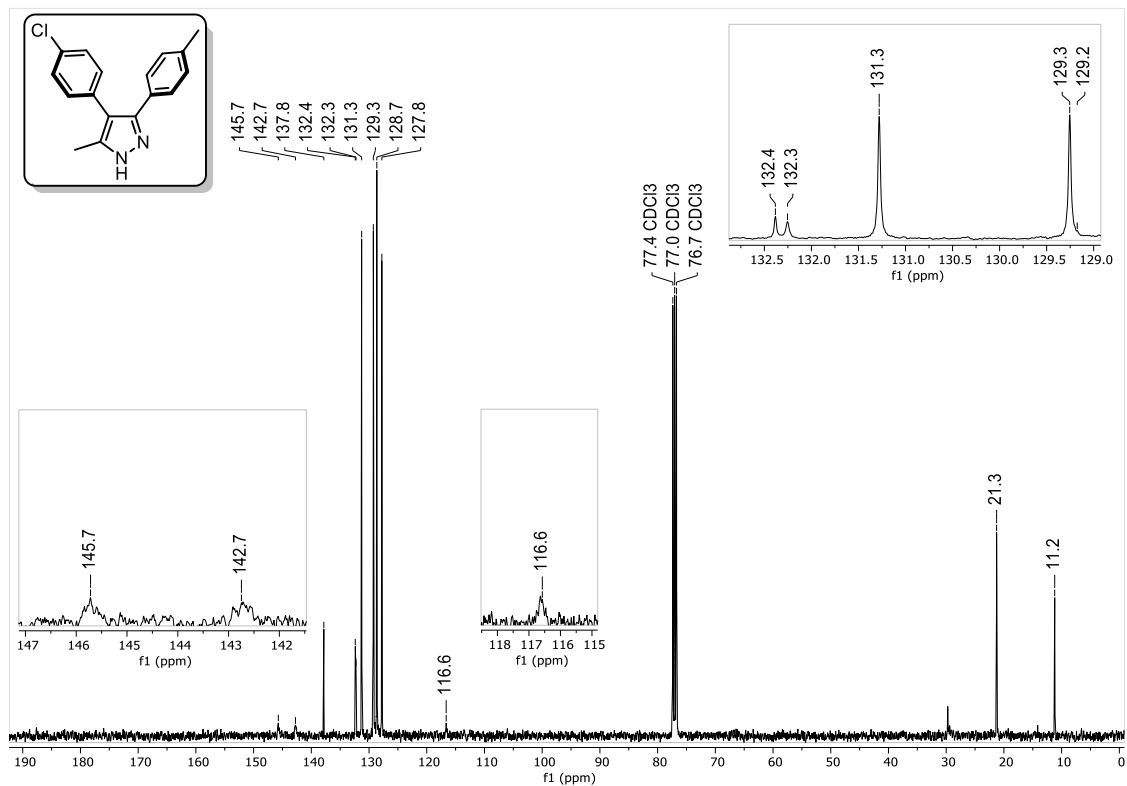


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
253.11418	62940.07	253.11410	0.07	0.29	¹² C ₁₆ ¹ H ₁₄ ¹⁹ F ₁ ¹⁴ N ₂	10.5

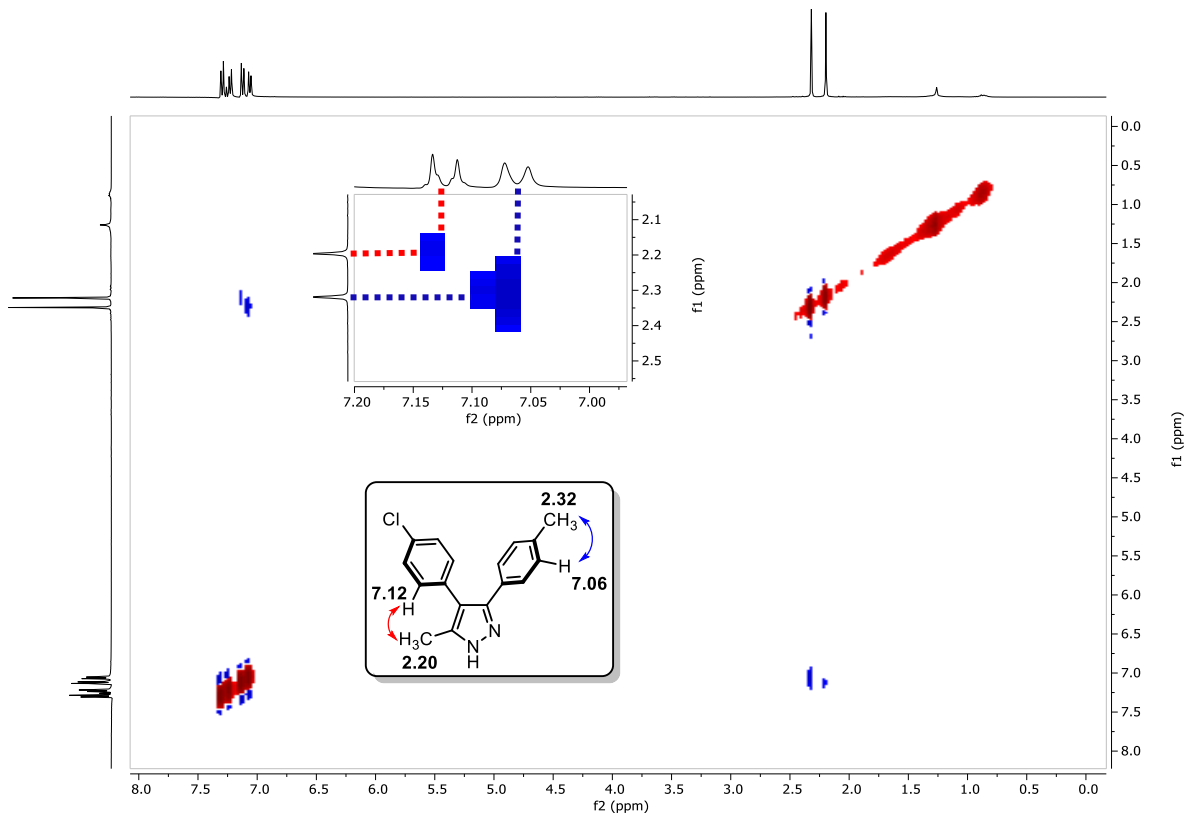
Spectrum 82. HRMS of 8n.



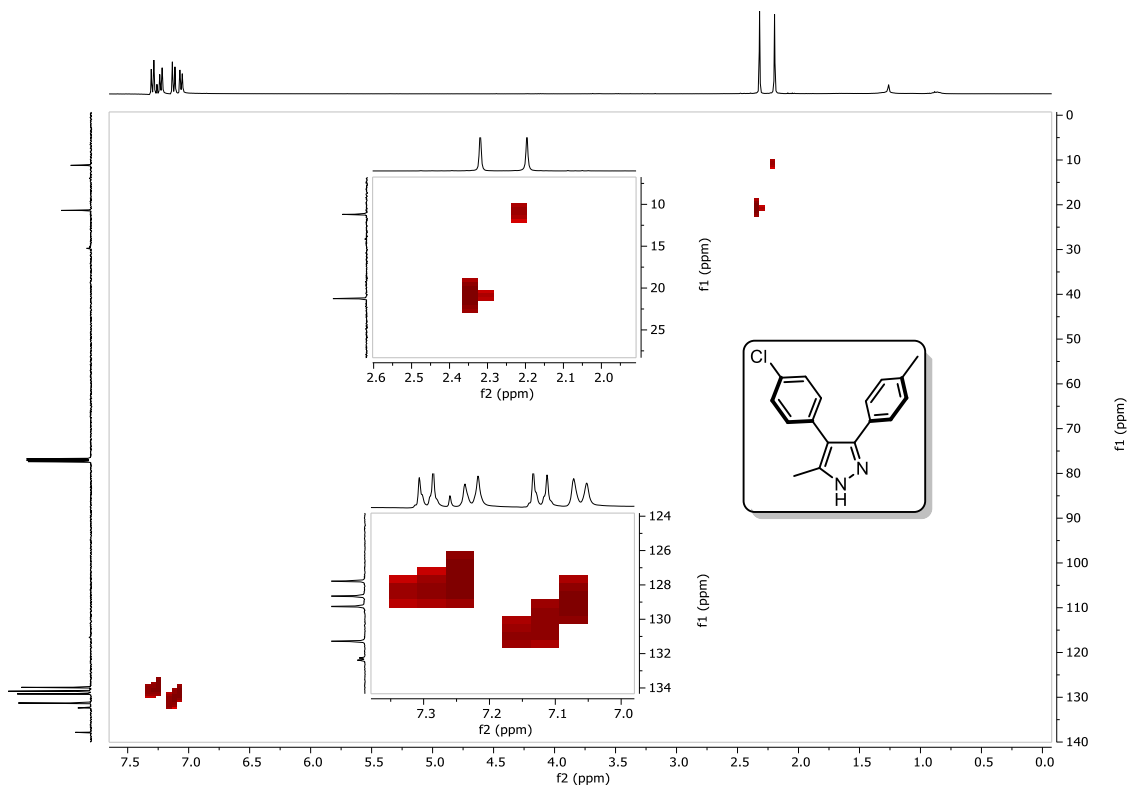
Spectrum 83. ¹H-NMR of 8o.



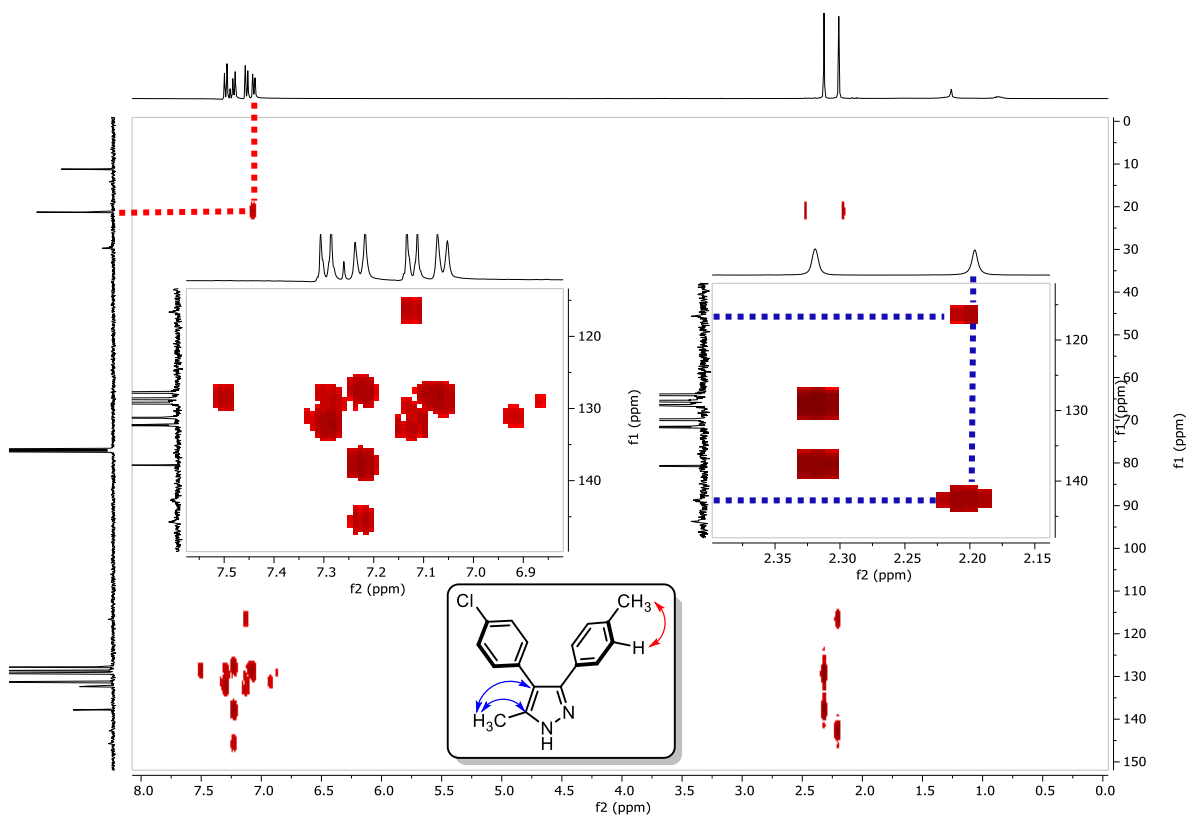
Spectrum 84 ^{13}C -NMR of 8o.



Spectrum 85. 2D-NOESY of 8o.



Spectrum 86. HSQC of 8o.



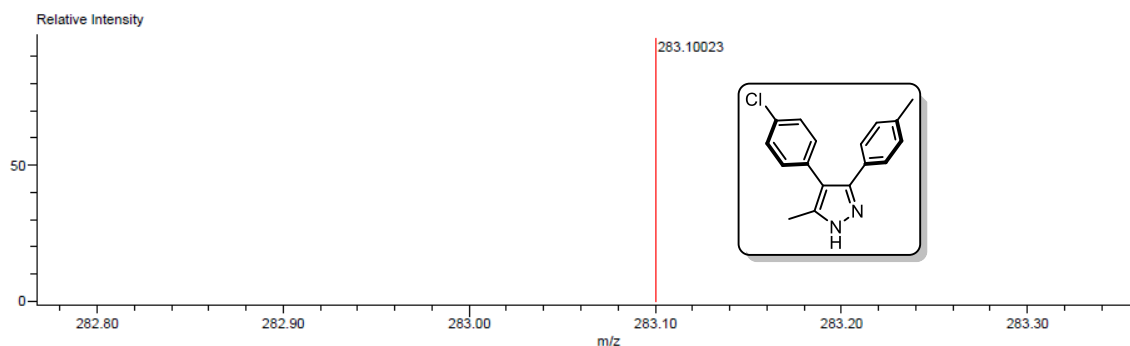
Spectrum 87. HMBC of 8o.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
 Created:1/25/2023 10:59:37 AM
 Created by:AccuTOF

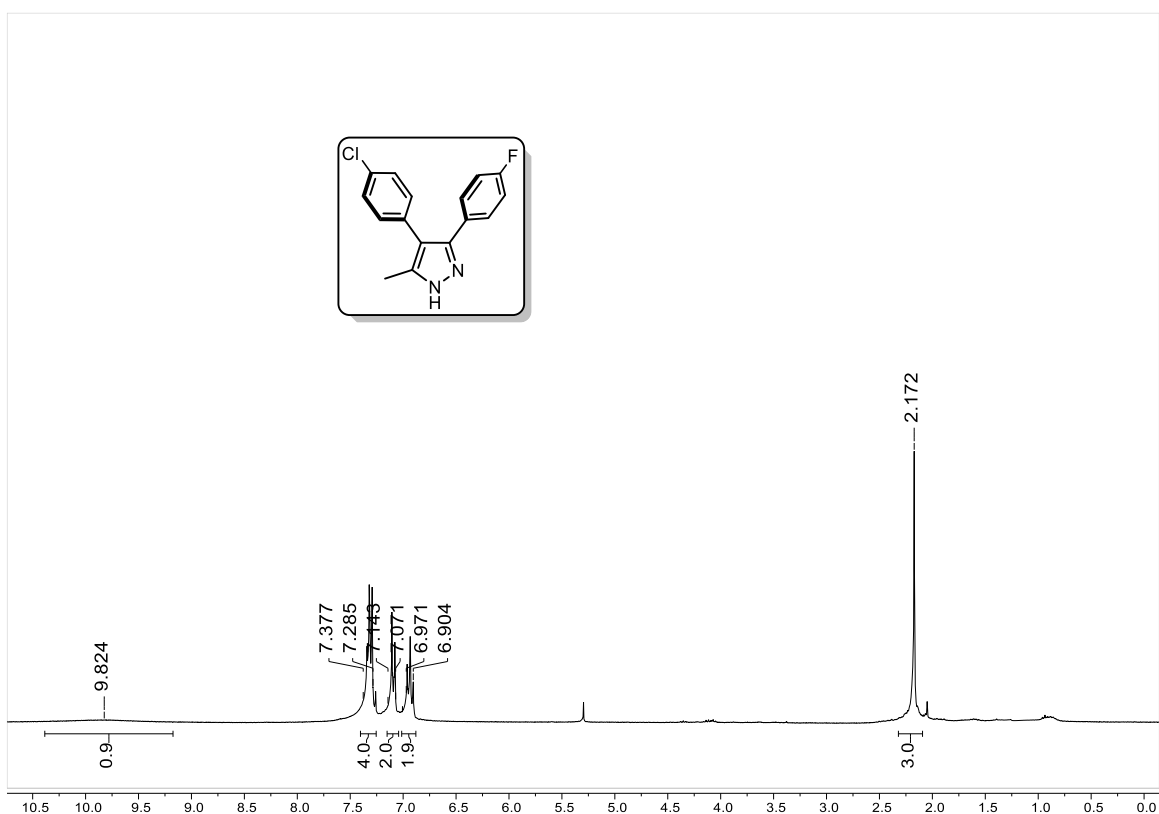
Charge number:1
 Element:¹²C:0 .. 17, ¹H:0 .. 20, ³⁵Cl:0 .. 1, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0
 Tolerance:3.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 45.0 (Fraction:Both)

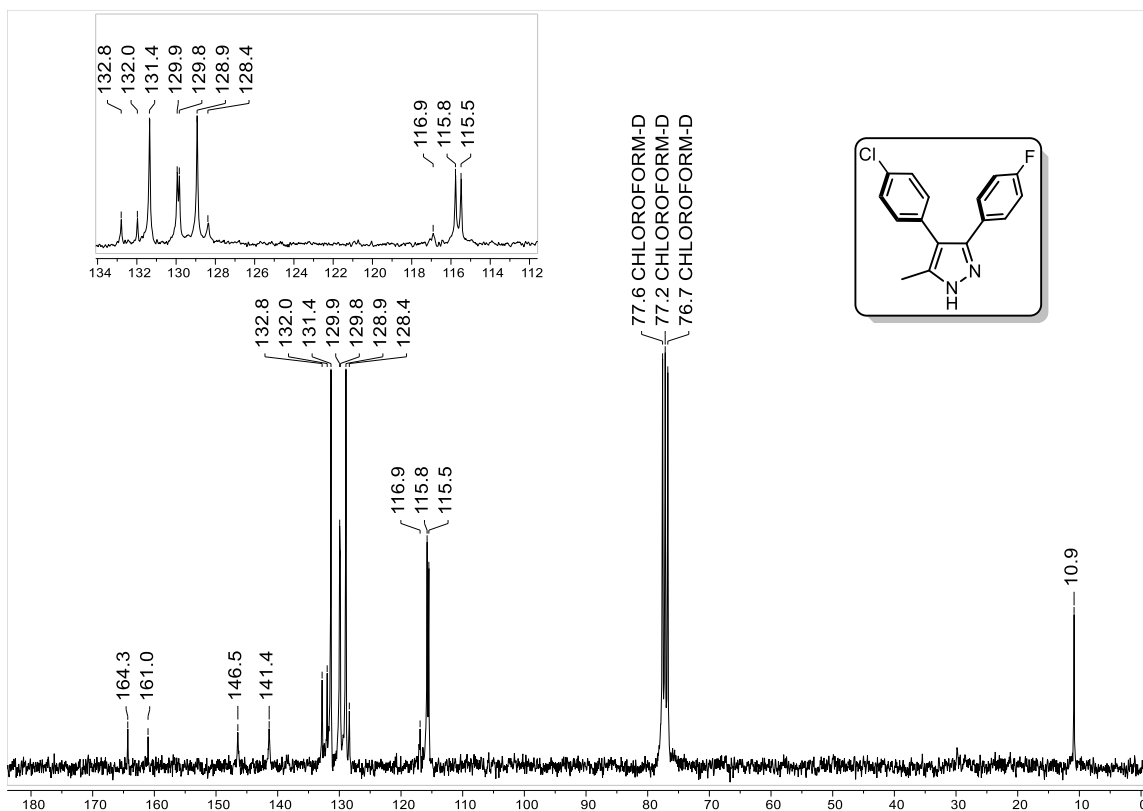


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
283.10023	1926911.20	283.10020	0.03	0.11	¹² C ₁₇ ¹ H ₁₆ ³⁵ Cl ₁ ¹⁴ N ₂	10.5

Spectrum 88. HRMS of 8o.



Spectrum 89. ¹H-NMR of 8p.



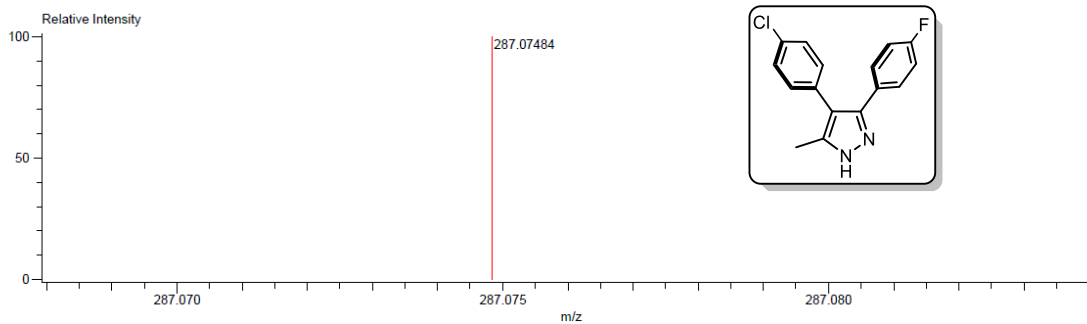
Spectrum 90. ¹³C-NMR of 8p.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:Cal_PEG_600
 Created:2/15/2023 11:00:07 AM
 Created by:

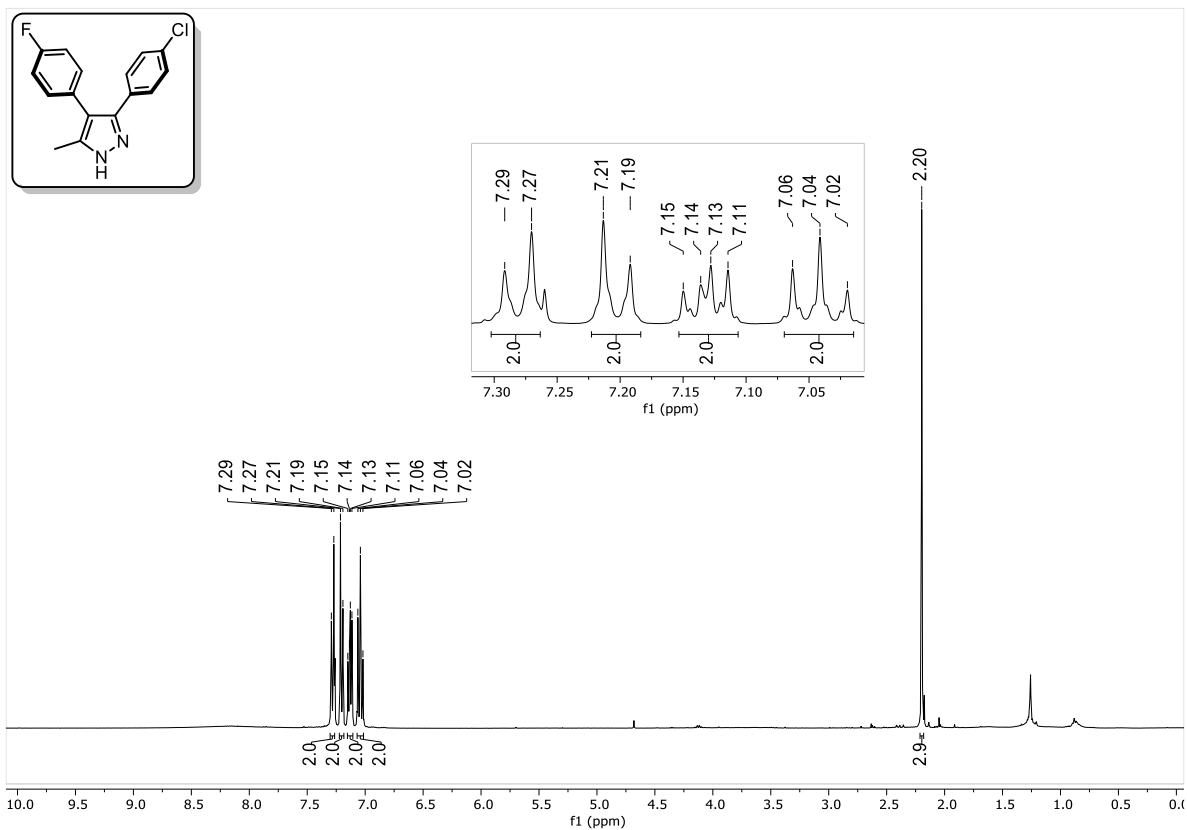
Charge number:1
 Element:¹²C:0 .. 16, ¹H:0 .. 22, ³⁵Cl:0 .. 1, ¹⁹F:0 .. 1, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0
 Tolerance:100.00(mmu)

Unsaturation Number:0.0 .. 20.0 (Fraction:Both)

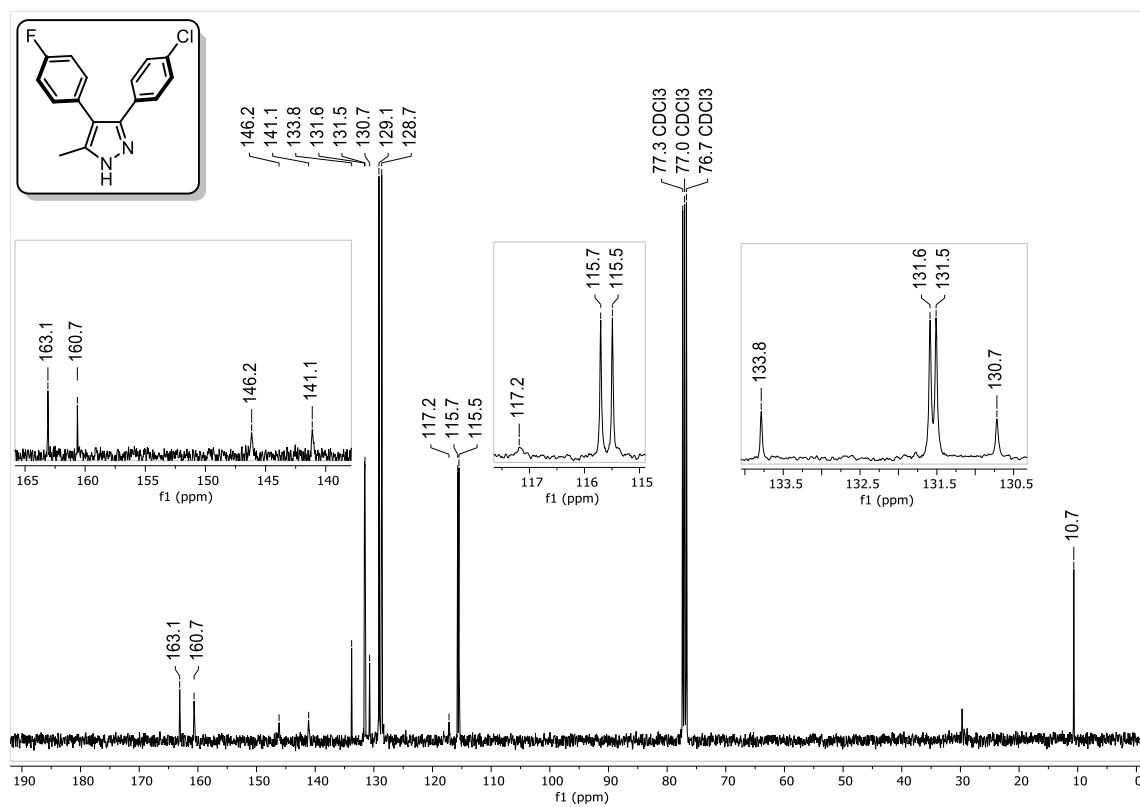


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
287.07484	14179.37	287.07513	-0.29	-1.00	¹² C ₁₈ ¹ H ₁₃ ³⁵ Cl ₁ ¹⁹ F ₁ ¹⁴ N ₂	10.5

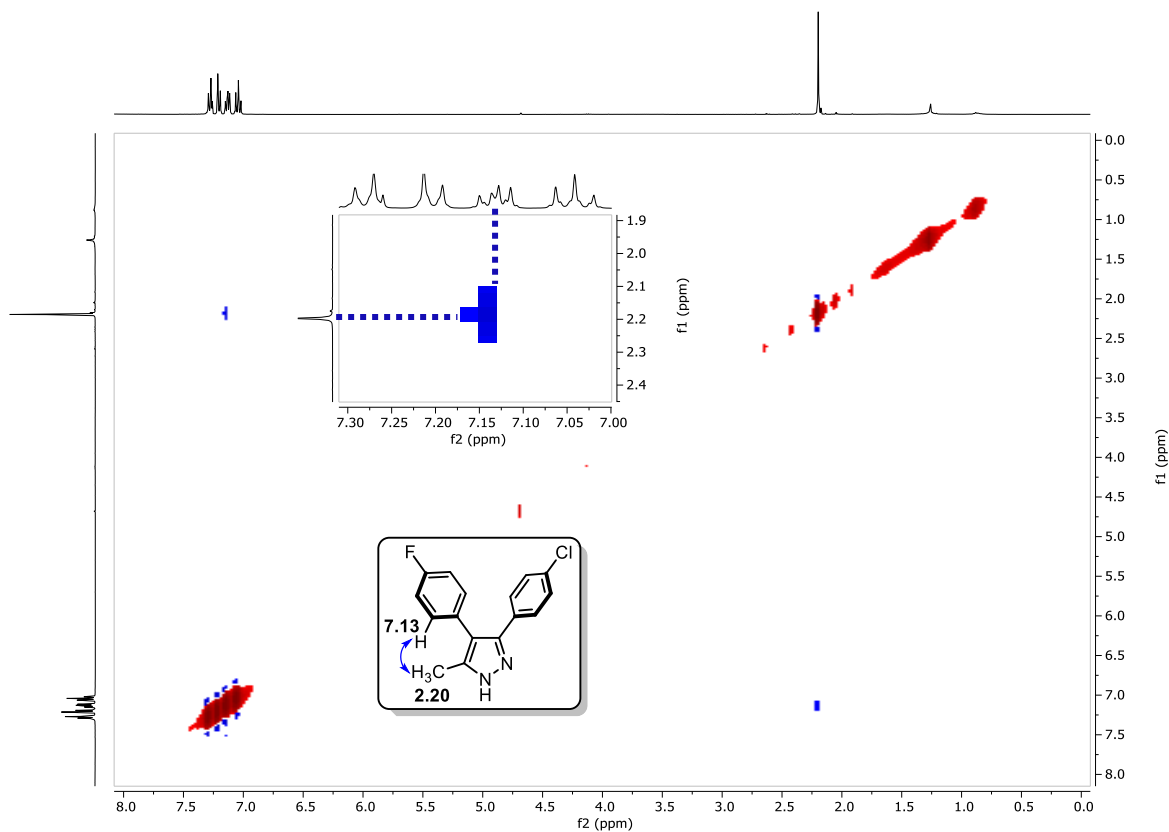
Spectrum 91. HRMS of 8p.



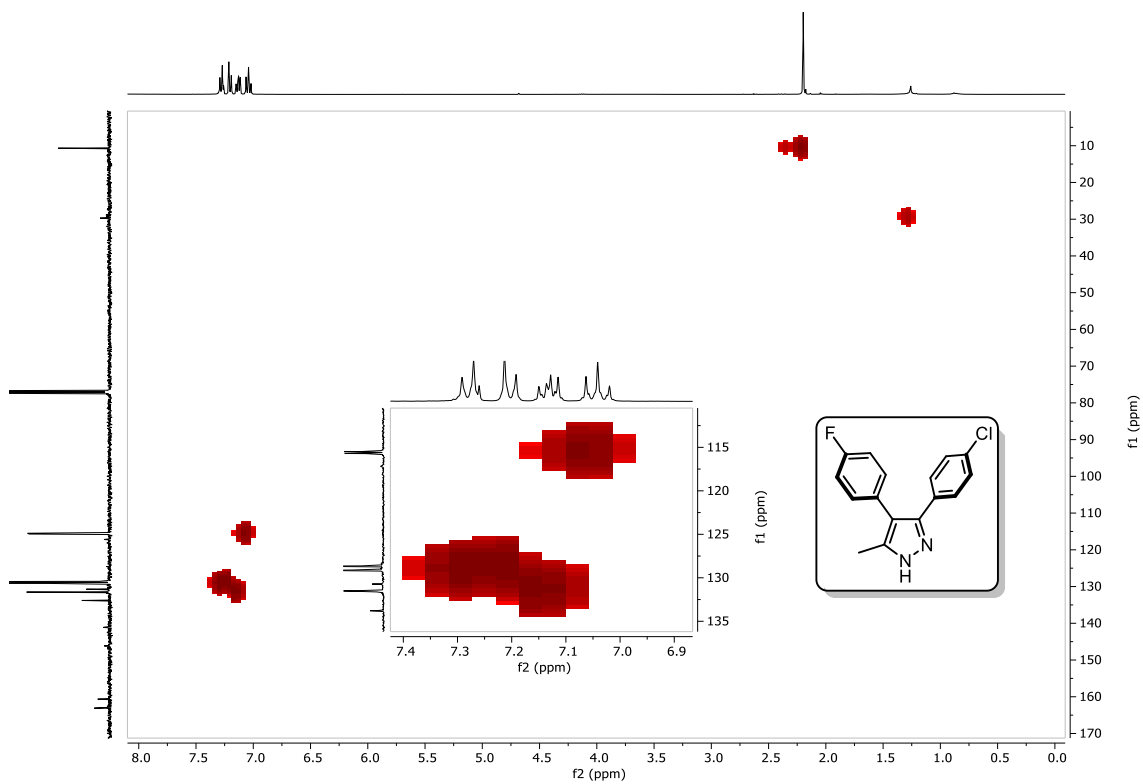
Spectrum 92. ¹H-NMR of 8q.



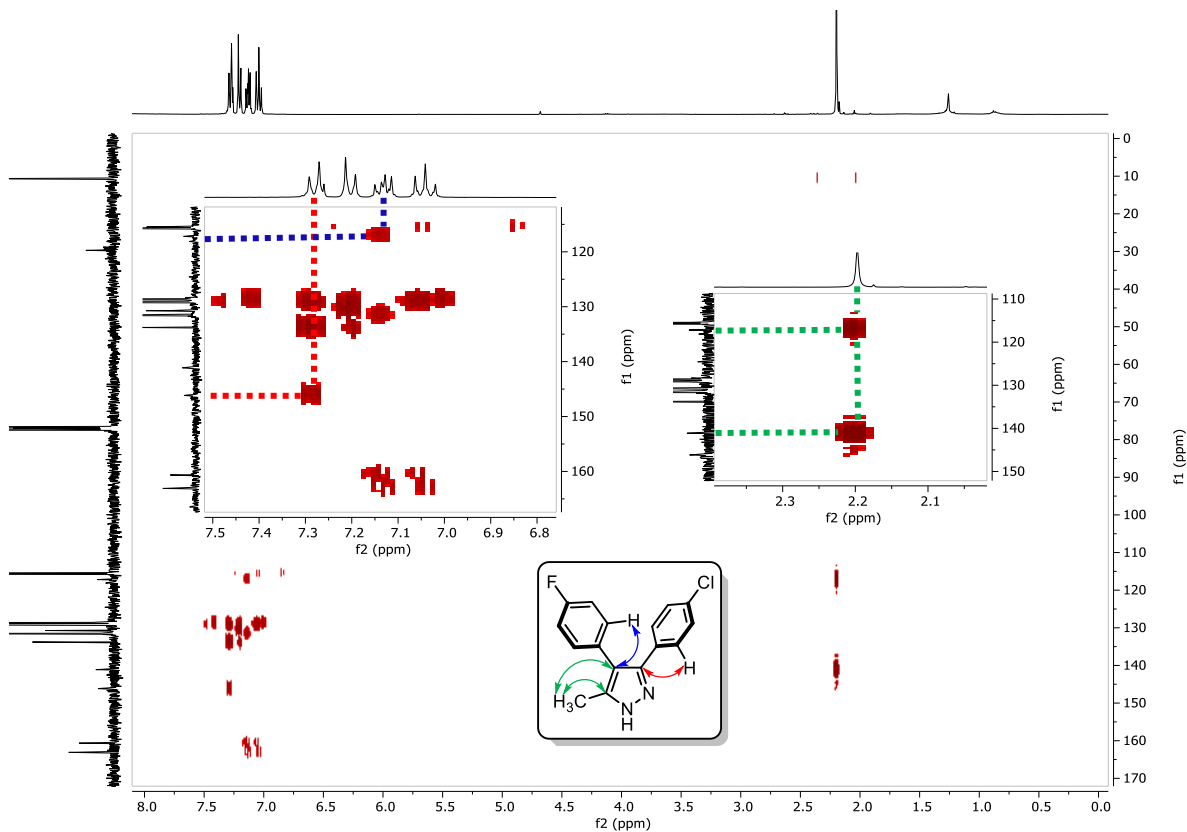
Spectrum 93. ¹³C-NMR of 8q.



Spectrum 94. 2D-NOESY of 8q.



Spectrum 95. HSQC of 8q.

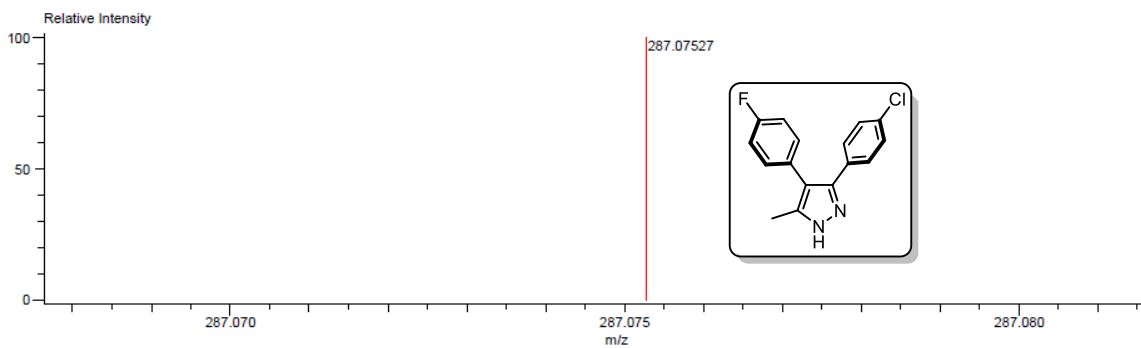


Spectrum 96. HMBC of 8q.

Description: Ionization Mode: ESI+
 History: Determine m/z [Peak Detect [Centroid, 30, Area]; Correct Base[]; Smooth [5]]; Correct Base [5.0%]; Average (MS[...])

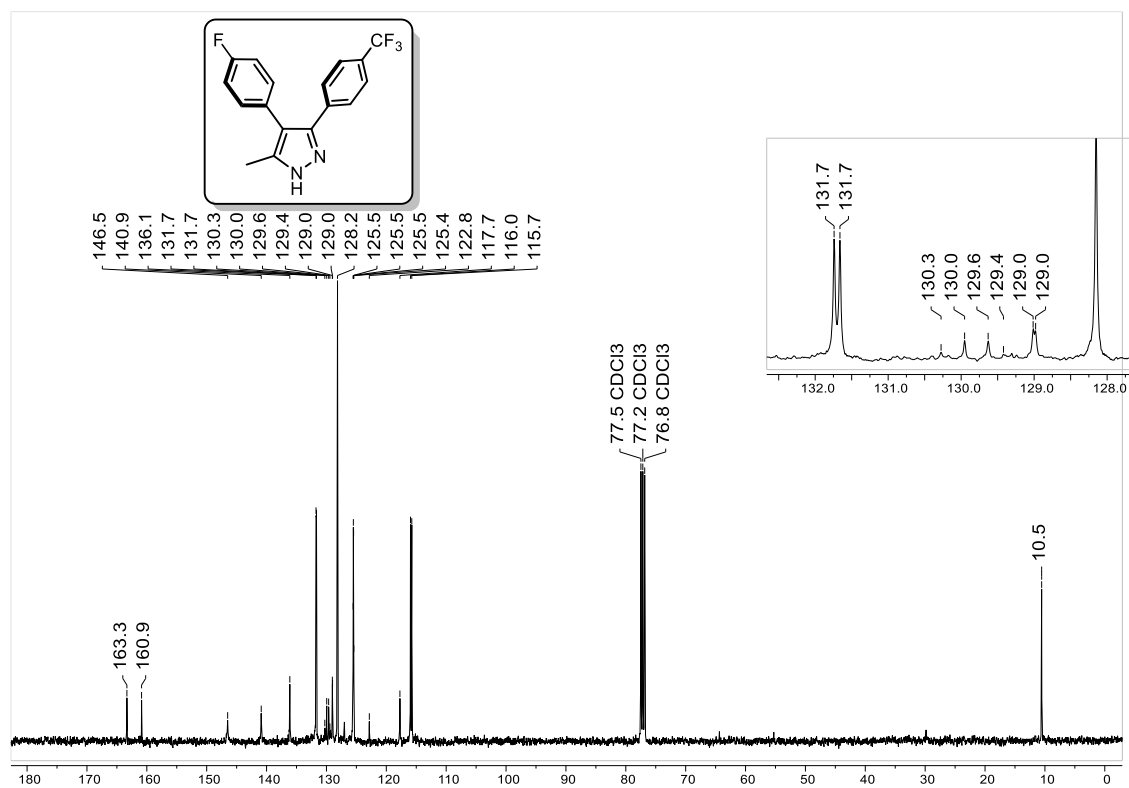
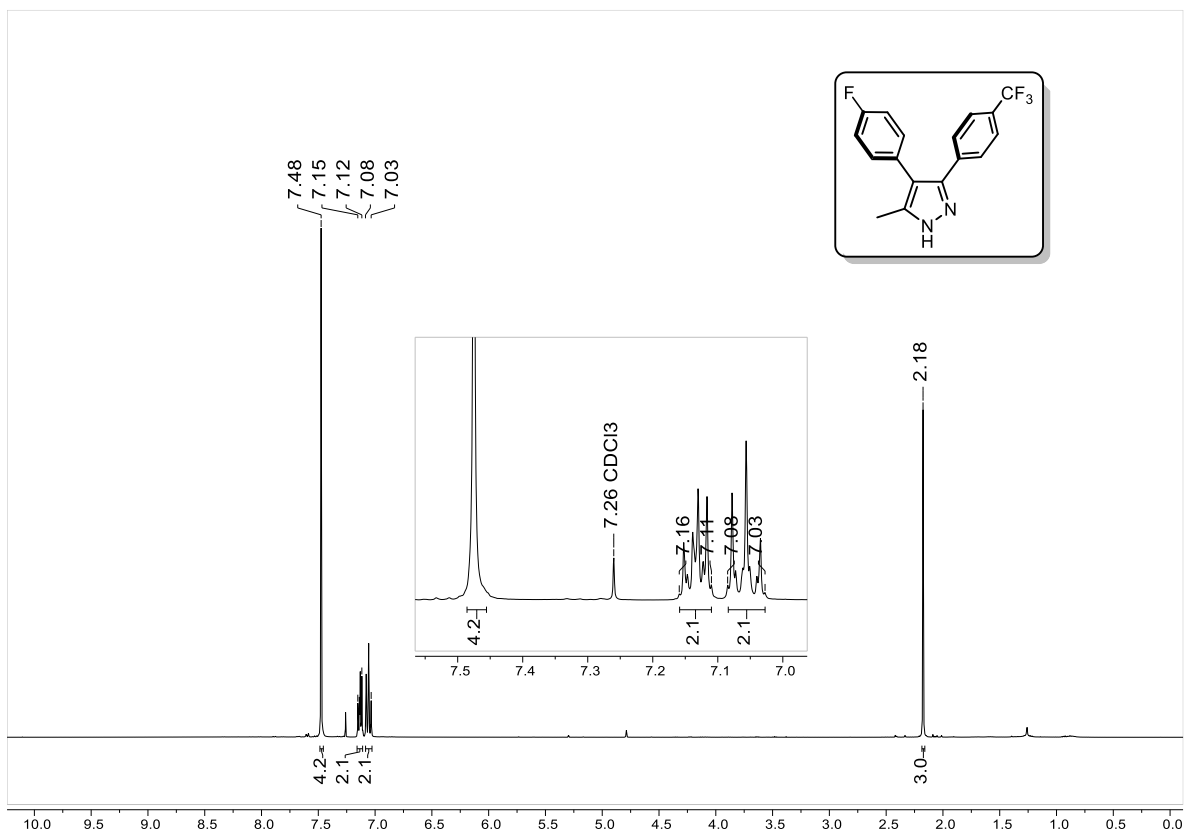
Mass Calibration data: Cal_PEG_600
 Created: 1/25/2023 11:02:36 AM
 Created by: AccuTOF

Charge number: 1 Tolerance: 3.00 (ppm), 5.00 .. 15.00 (mmu) Unsaturation Number: 0.0 .. 45.0 (Fraction: Both)
 Element: ¹²C: 0 .. 16, ¹H: 0 .. 20, ³⁵Cl: 0 .. 1, ¹⁹F: 1 .. 1, ¹⁴N: 0 .. 2, ¹⁶O: 0 .. 0



Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
287.07527	17703.09	287.07513	0.14	0.49	¹² C ₁₆ ¹ H ₁₃ ³⁵ Cl ₁ ¹⁹ F ₁ ¹⁴ N ₂	10.5

Spectrum 97. HRMS of 8q.

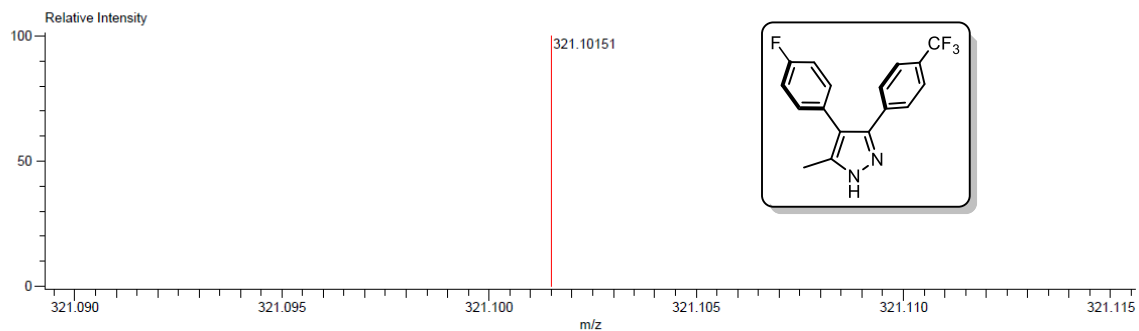


Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
 Created:2/2/2023 10:33:12 AM
 Created by:AccuTOF

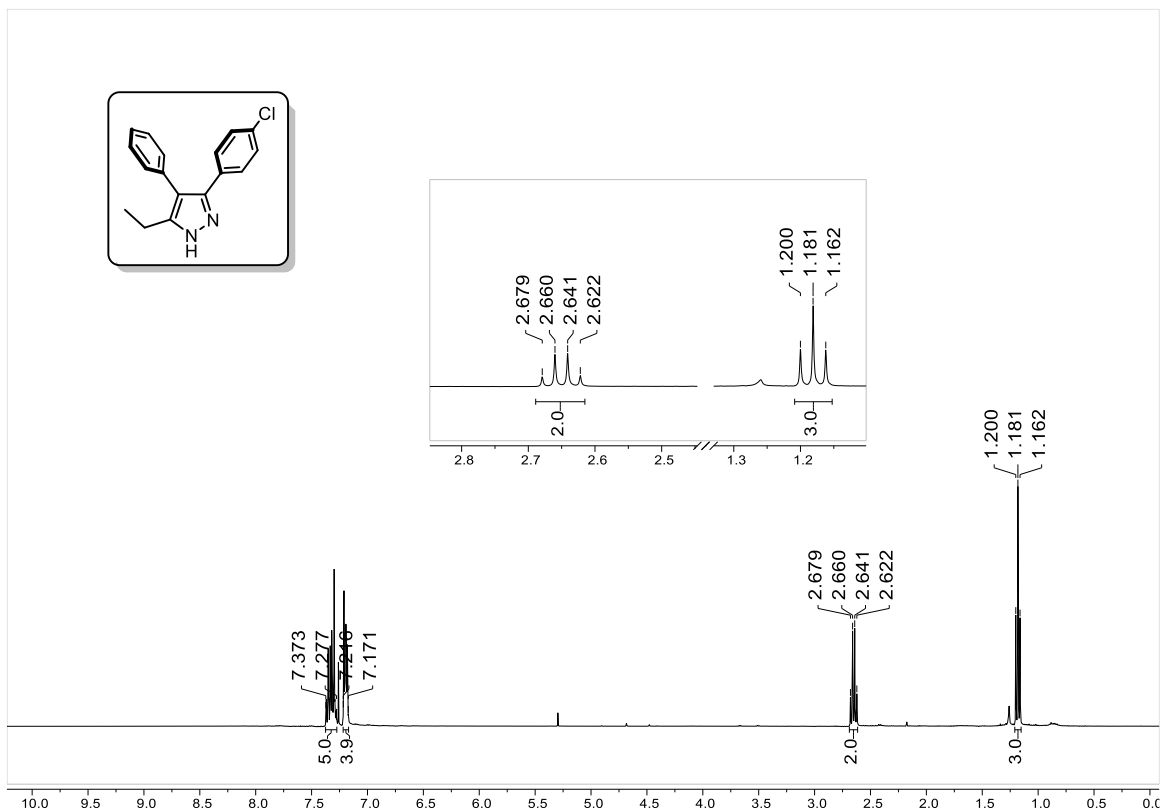
Charge number:1
 Element:¹²C:0 .. 17, ¹H:0 .. 50, ¹⁹F:0 .. 4, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0
 Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 50.0 (Fraction:.5)

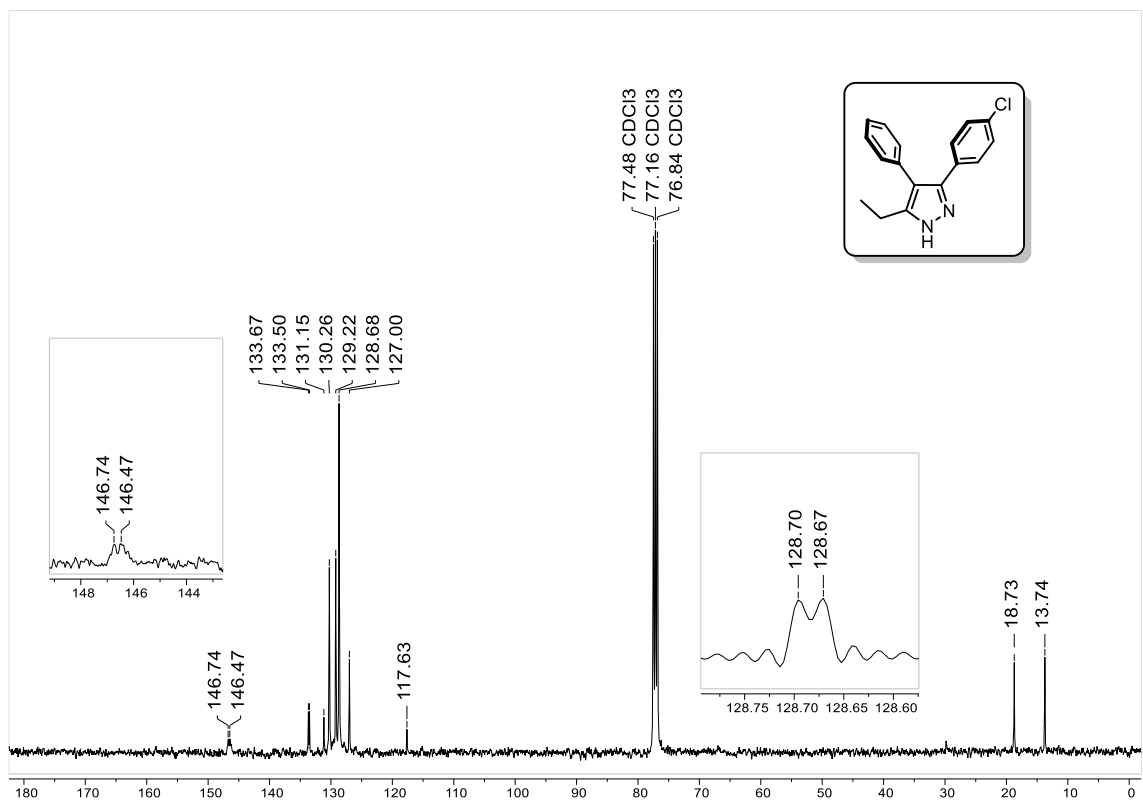


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
321.10151	4047.61	321.10149	0.02	0.07	¹² C ₁₇ ¹ H ₁₃ ¹⁹ F ₄ ¹⁴ N ₂	10.5

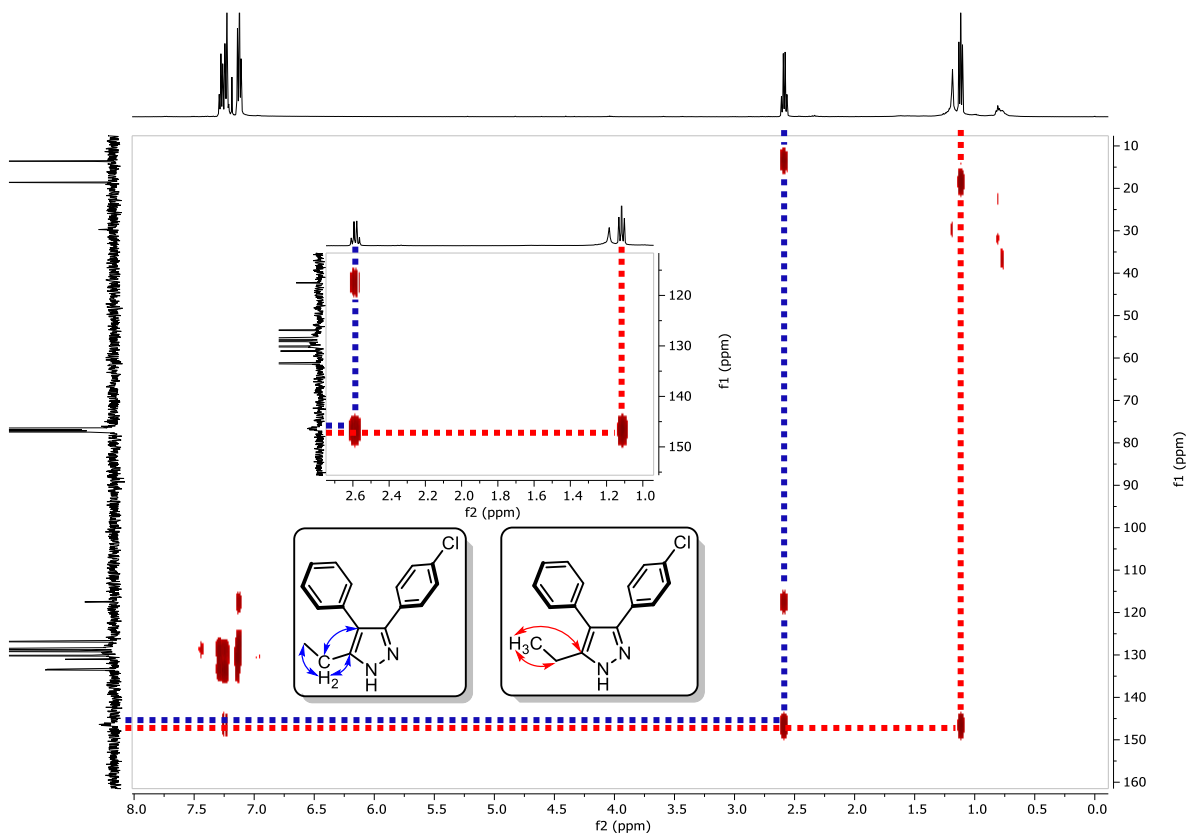
Spectrum 100. HRMS of 8r.



Spectrum 101. ¹H-NMR of 8s.



Spectrum 102. ¹³C-NMR of 8s.



Spectrum 103. HMBC of 8s.

Ionization Mode: ESI+

History: Determine m/z [Peak Detect [Centroid, 30, Area], Correct Base [], Smooth [5]], Correct Base [5.0%], Average [MS]...

Created: 3/10/2023 9:26:52 AM

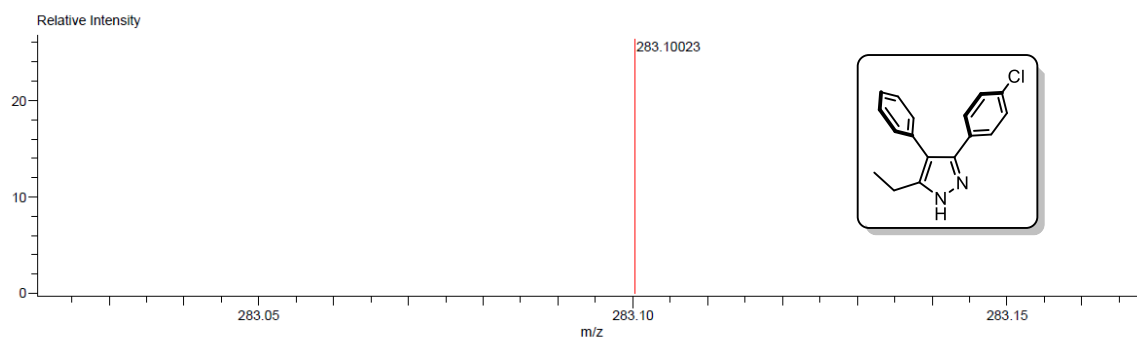
Created by: AccuTOF

Charge number: 1

Tolerance: 50.00 (ppm), 5.00 .. 15.00 (mmu)

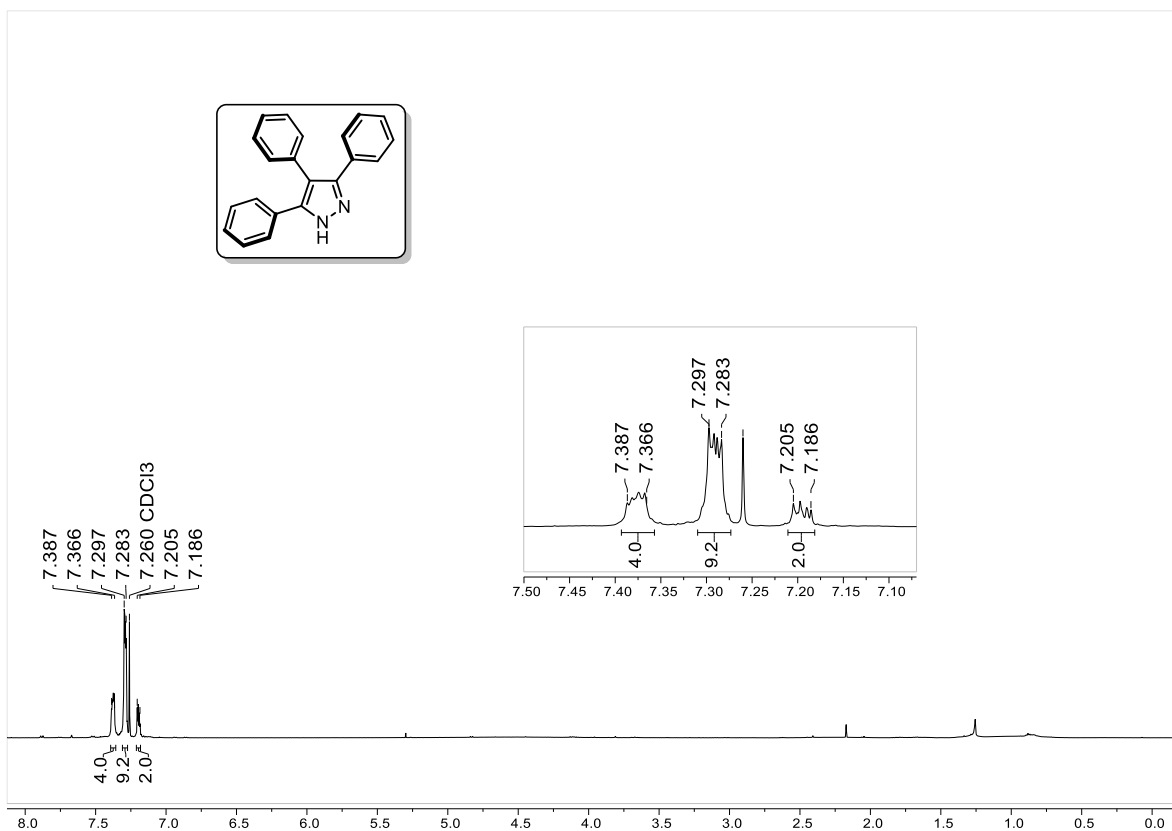
Unsaturation Number: 0.0 .. 50.0 (Fraction: Both)

Element: ¹²C: 0 .. 17, ¹H: 0 .. 50, ³⁵Cl: 0 .. 1, ¹⁹F: 0 .. 0, ¹⁴N: 0 .. 2, ¹⁶O: 0 .. 0, ³²S: 0 .. 0

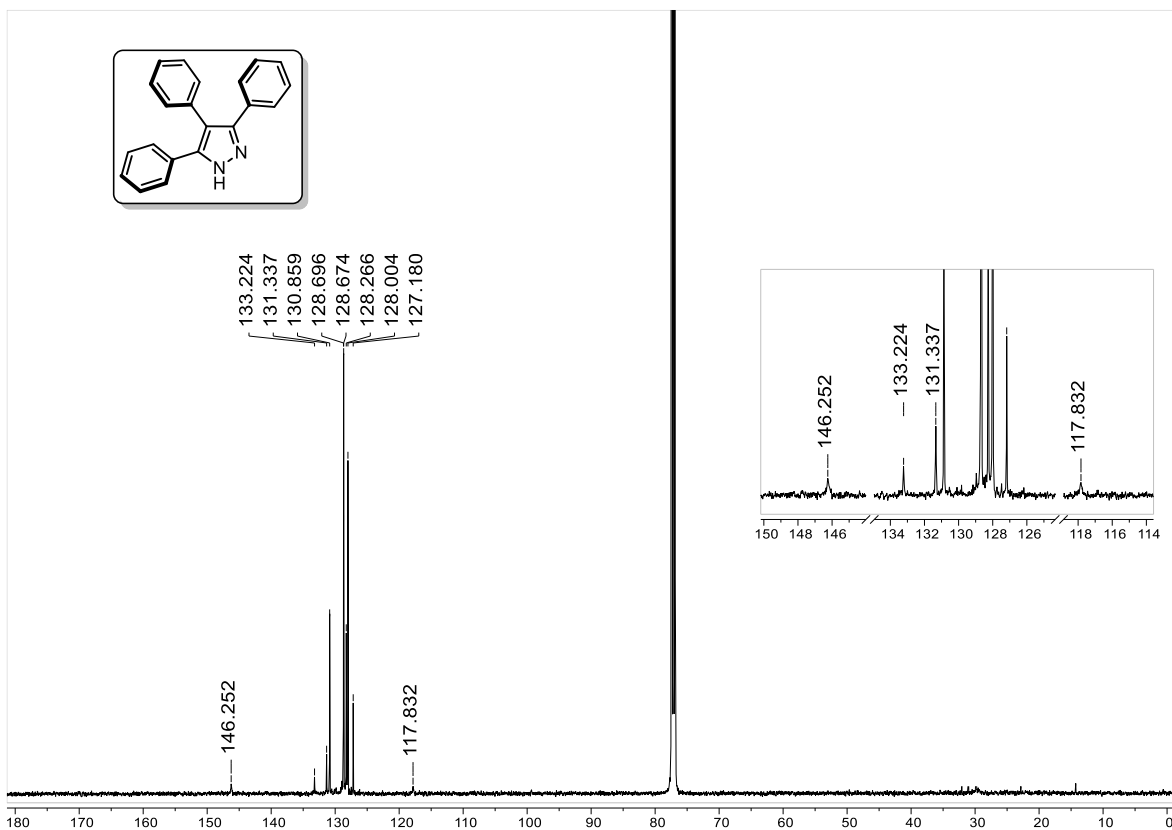


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
283.10023	377996.21	283.10020	0.03	0.12	¹² C ₁₇ ¹ H ₁₈ ³⁵ Cl ₁ ¹⁴ N ₂	10.5

Spectrum 104. HRMS of 8s.



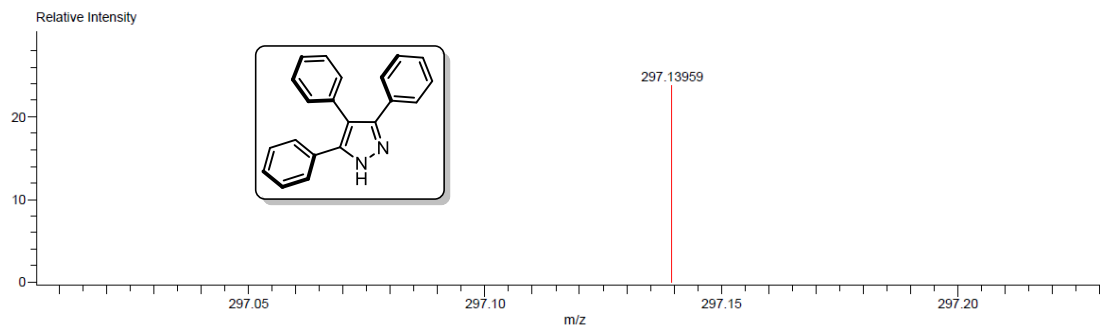
Spectrum 105. ¹H-NMR of 8t.



Spectrum 106. ^{13}C -NMR of 8t.

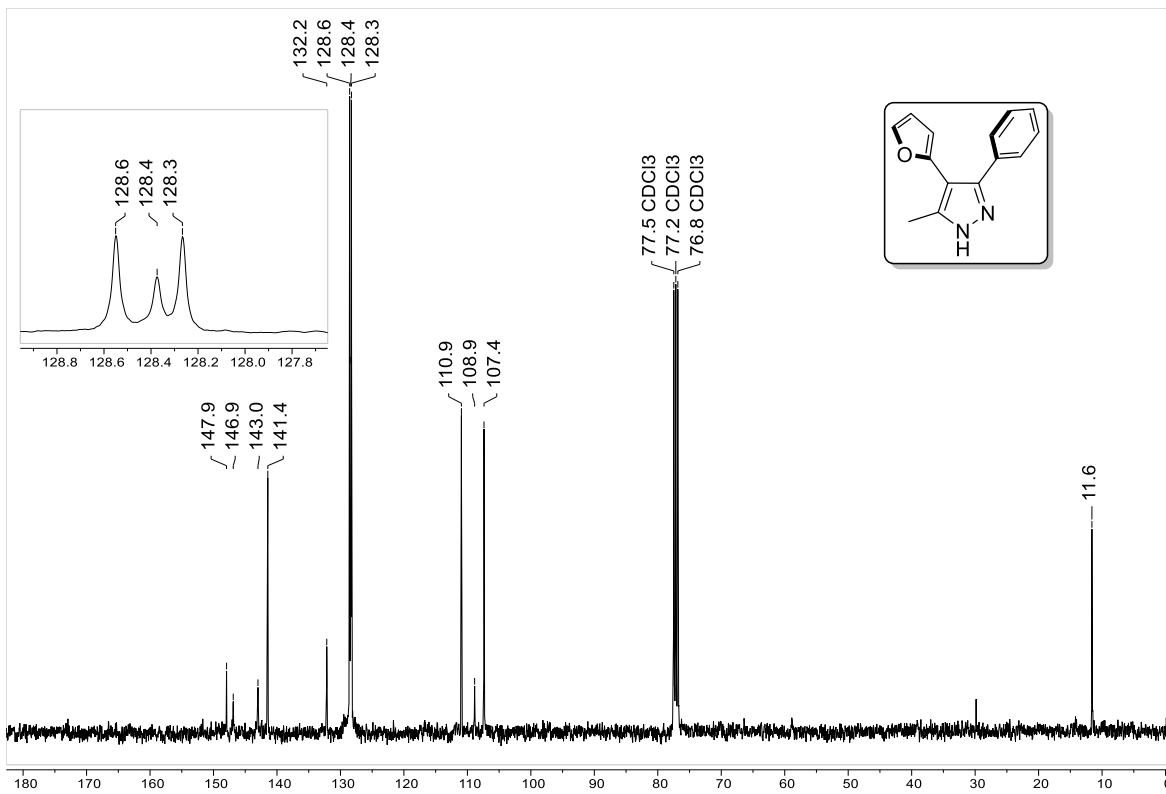
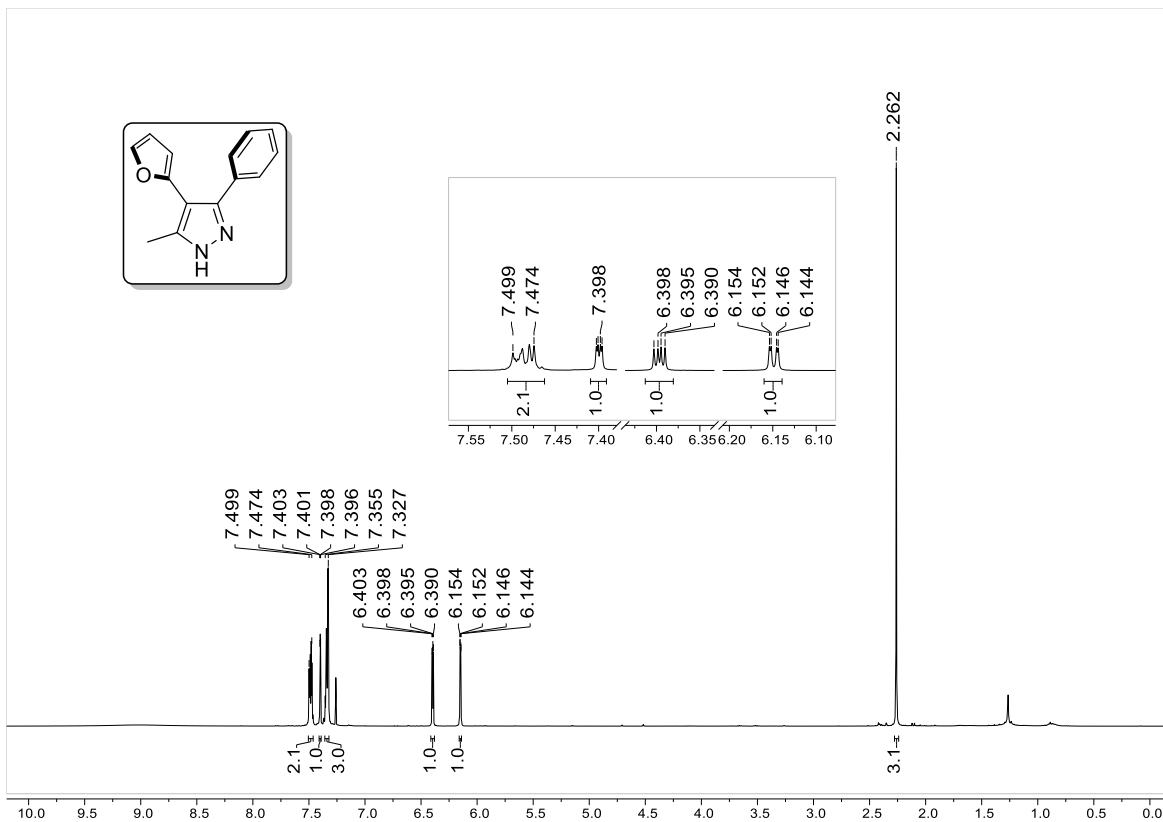
Description: Mass Calibration data: Cal_PEG_600
 Ionization Mode: ESI+ Created: 3/17/2023 11:01:00 AM
 History: Determine m/z [Peak Detect [Centroid, 30, Area]; Correct Base[]; Smooth [5]]; Correct Base [5.0%]; Average [MS...]
 Created by: AccuTOF

Charge number: 1 Tolerance: 5.00 (mmu) Unsaturation Number: 0.0 .. 50.0 (Fraction: Both)
 Element: ^{12}C : 0 .. 21, ^1H : 0 .. 18, ^{14}N : 0 .. 2, ^{32}S : 0 .. 0



Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
297.13959	22230.20	297.13917	0.42	1.42	$^{12}\text{C}_{21}\text{H}_{17}\text{N}_2$	14.5

Spectrum 107. HRMS of 8t.



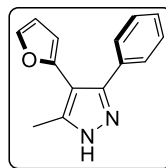
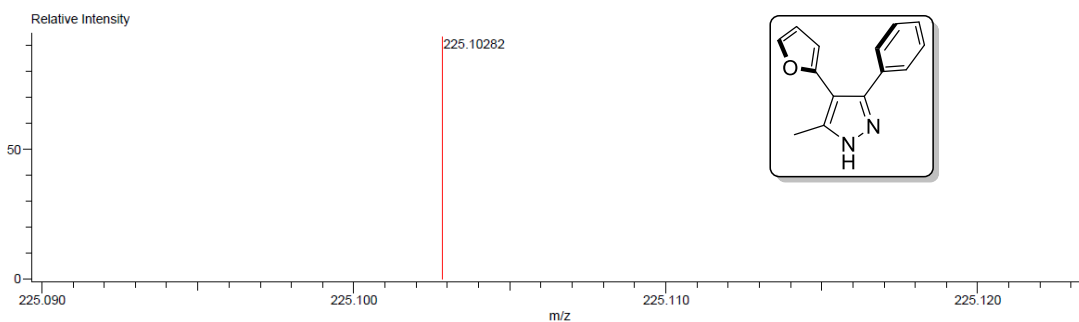
Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[;Smooth[5];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
 Created:2/23/2023 12:57:25 PM
 Created by:AccuTOF

Charge number:1
 Element:¹²C:0 .. 17, ¹H:0 .. 18, ¹⁴N:0 .. 2, ¹⁶O:0 .. 1

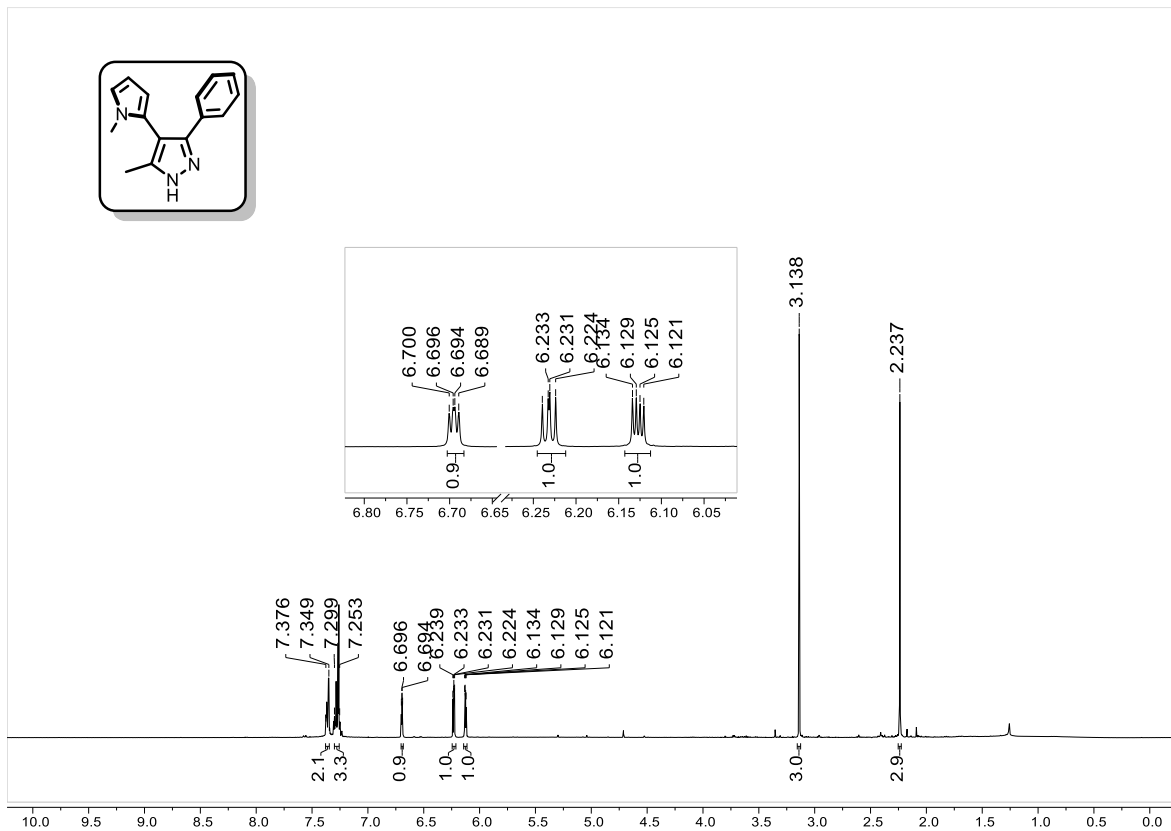
Tolerance:4.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 40.0 (Fraction:Both)

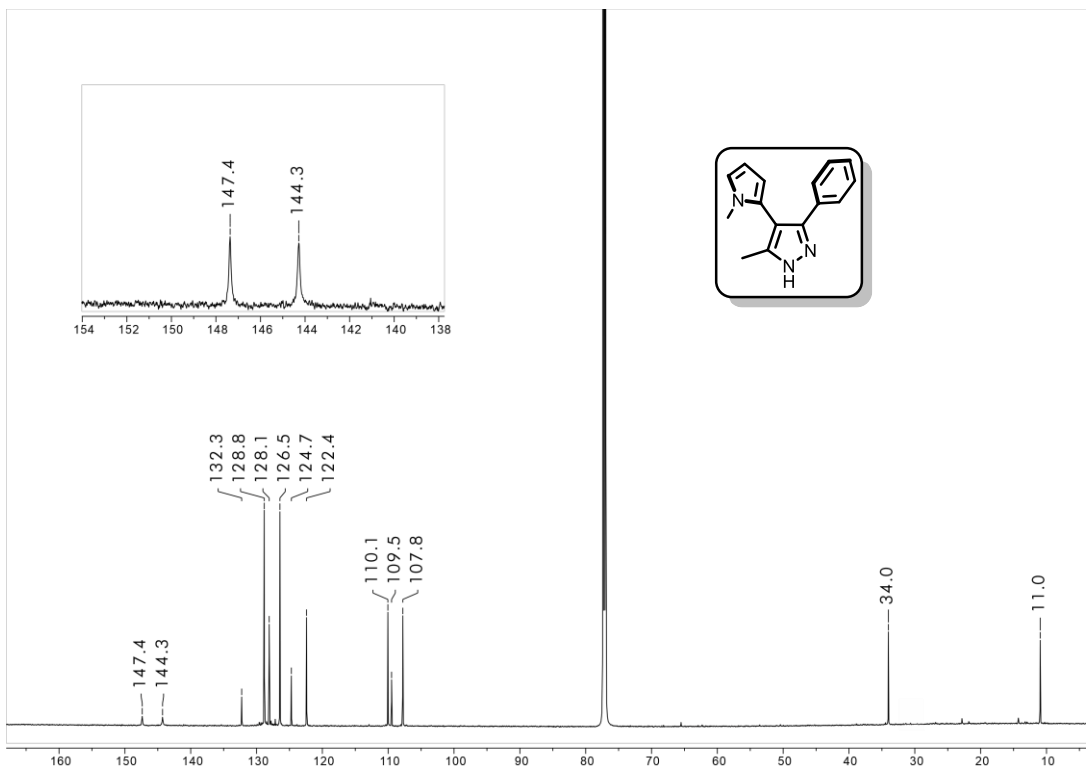


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
225.10282	5968.48	225.10279	0.03	0.14	¹² C ₁₄ ¹ H ₁₃ ¹⁴ N ₂ ¹⁶ O ₁	9.5

Spectrum 110. HRMS of 8u.



Spectrum 111. ¹H-NMR of 8v.



Spectrum 112. ^{13}C -NMR of **8v**.

Description:

Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base];Smooth[5];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600

Created:3/13/2023 2:53:20 PM

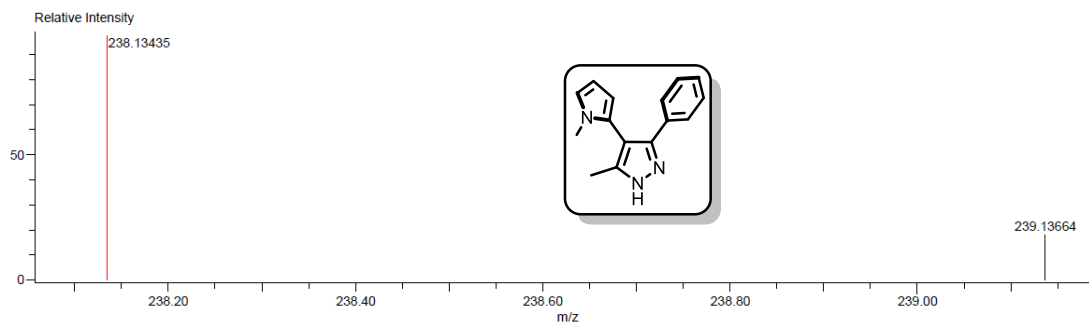
Created by:AccuTOF

Charge number:1

Tolerance:100.00(mmu)

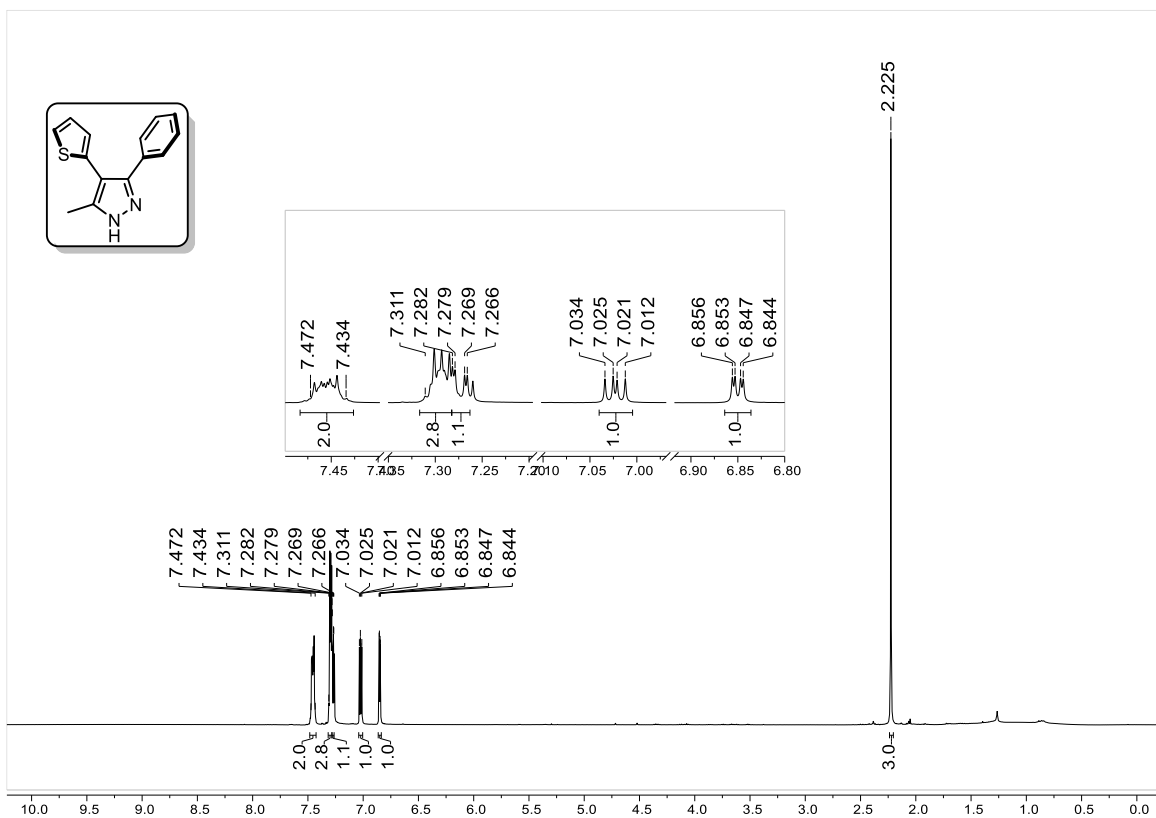
Unsaturation Number:0.0 .. 40.0 (Fraction:Both)

Element: ^{12}C :0 .. 15, ^1H :0 .. 16, ^{14}N :3 .. 3, ^{16}O :0 .. 0, ^{32}S :0 .. 0

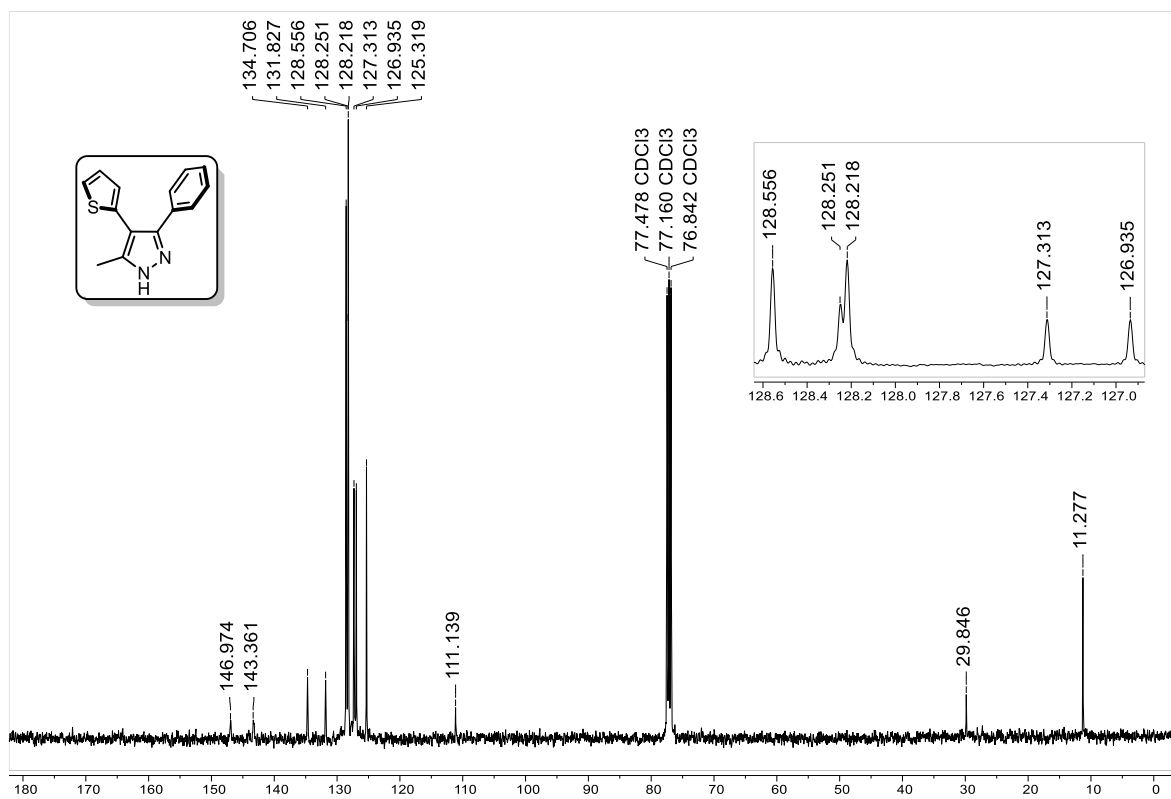


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
238.13435	21076.71	238.13442	-0.07	-0.29	$^{12}\text{C}_{15}\text{H}_{16}\text{N}_3$	9.5

Spectrum 113. HRMS of **8v**.



Spectrum 114. ¹H-NMR of 8w.



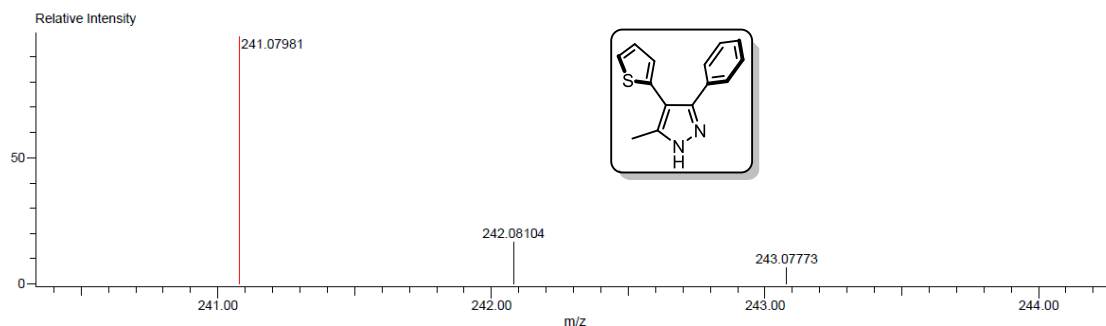
Spectrum 115. ¹³C-NMR of 8w.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:CaI_PEG_600
 Created:3/10/2023 9:16:20 AM
 Created by:AccuTOF

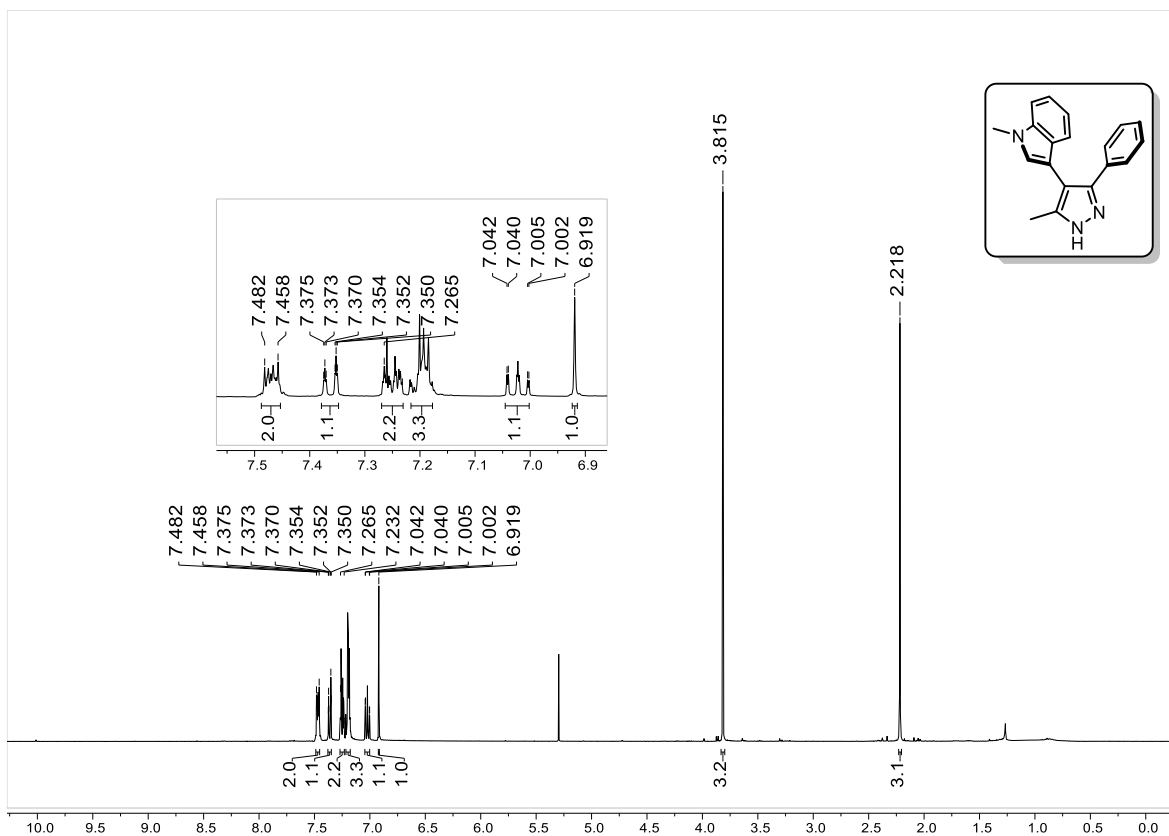
Charge number:1
 Element:¹²C:0 .. 14, ¹H:0 .. 50, ³⁵Cl:0 .. 0, ¹⁹F:0 .. 0, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0, ³²S:0 .. 1
 Tolerance:50.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 50.0 (Fraction:Both)

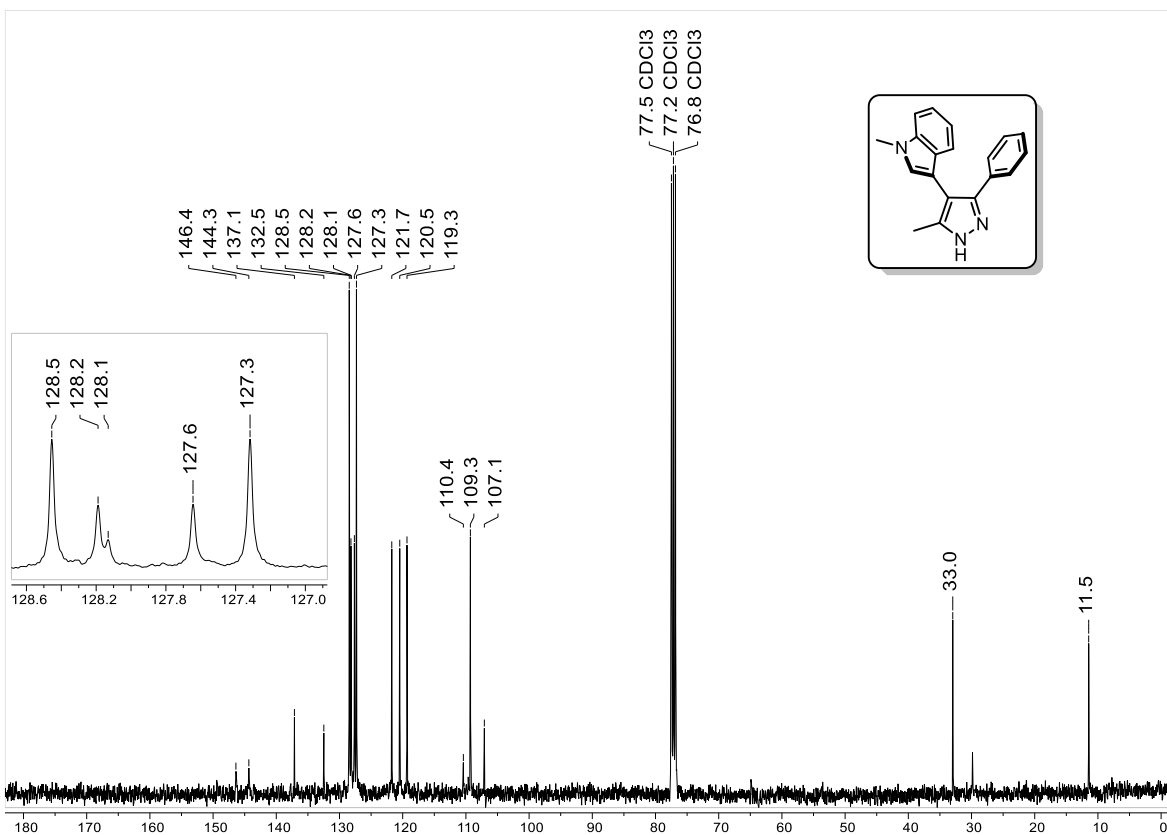


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
241.07981	20510.44	241.07994	-0.13	-0.55	¹² C ₁₄ ¹ H ₁₃ ¹⁴ N ₂ ³² S ₁	9.5

Spectrum 116. HRMS of 8w.



Spectrum 117. ¹H-NMR of 8x.



Spectrum 118. ^{13}C -NMR of **8x**.

Description:

Ionization Mode:ESI+

History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:CaI_PEG_600

Created:3/1/2023 10:40:03 AM

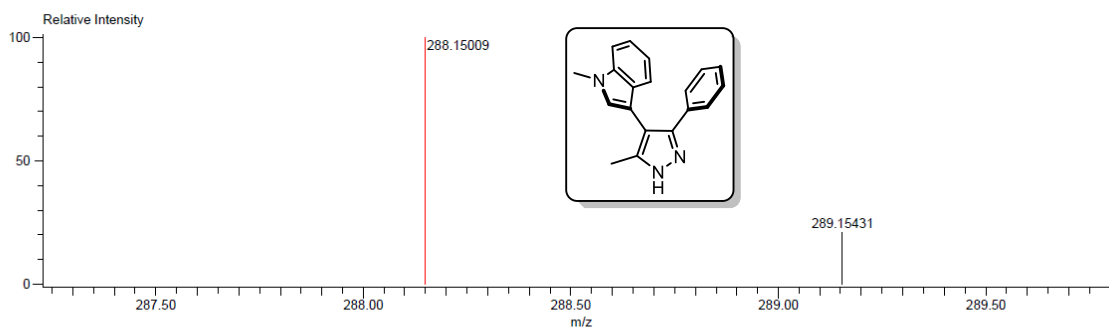
Created by:AccuTOF

Charge number:1

Tolerance:3.00(mmu)

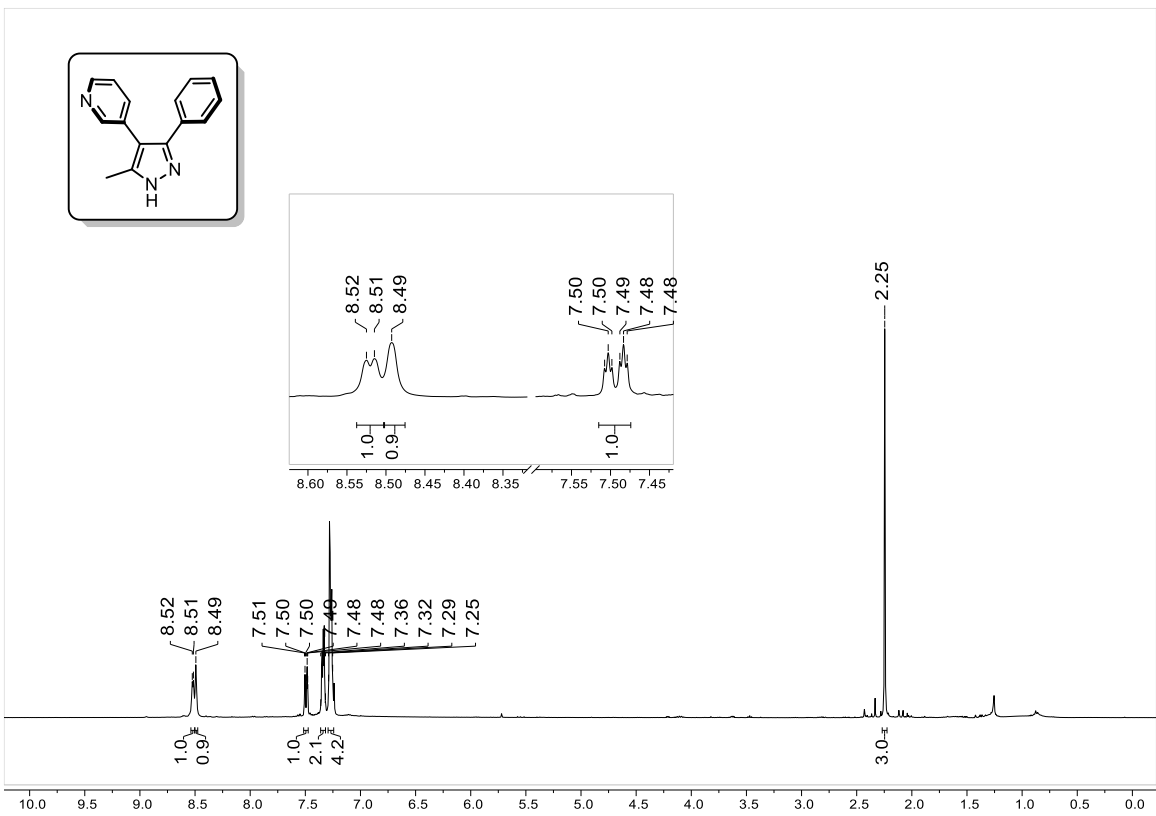
Unsaturation Number:0.0 .. 30.0 (Fraction:Both)

Element: ^{12}C :0 .. 19, ^1H :0 .. 20, ^{79}Br :0 .. 0, ^{81}Br :0 .. 0, ^{14}N :0 .. 3, ^{15}N :0 .. 0, ^{16}O :0 .. 0, ^{32}S :0 .. 0

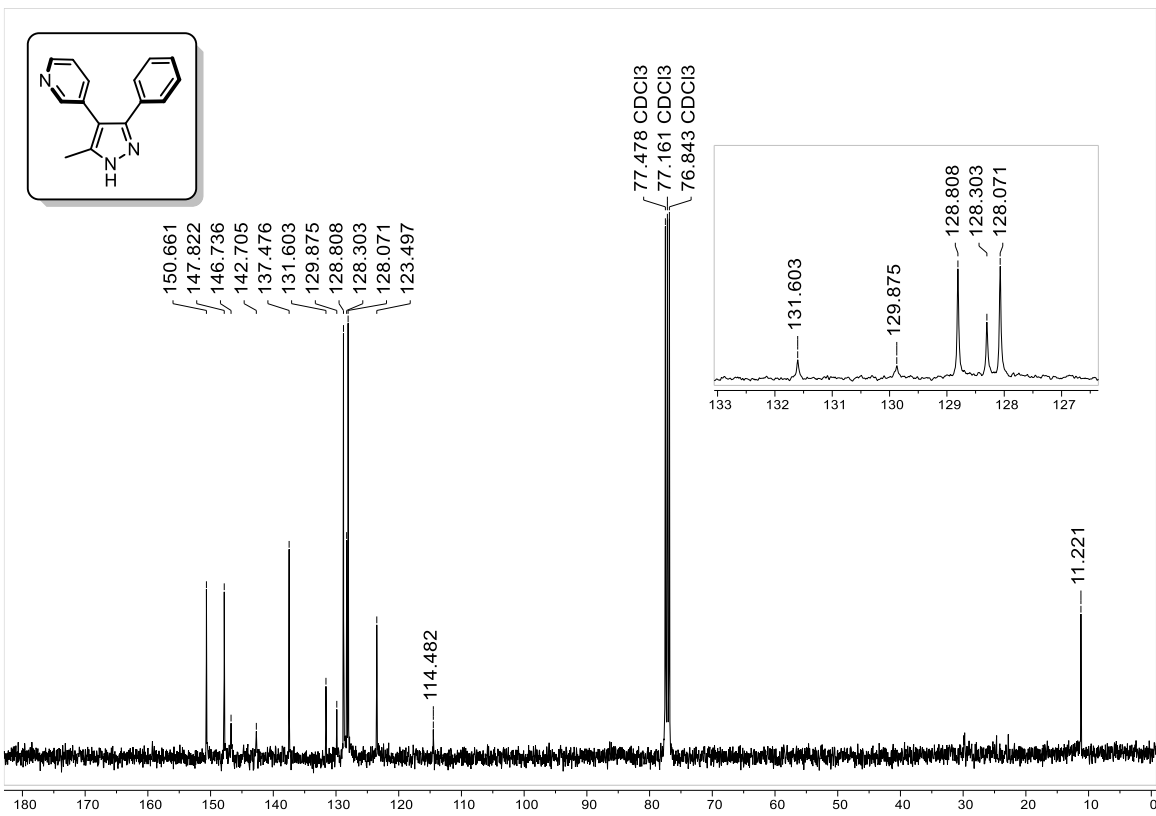


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
288.15009	208716.91	288.15007	0.01	0.05	$^{12}\text{C}_{19}\text{H}_{18}\text{N}_3$	12.5

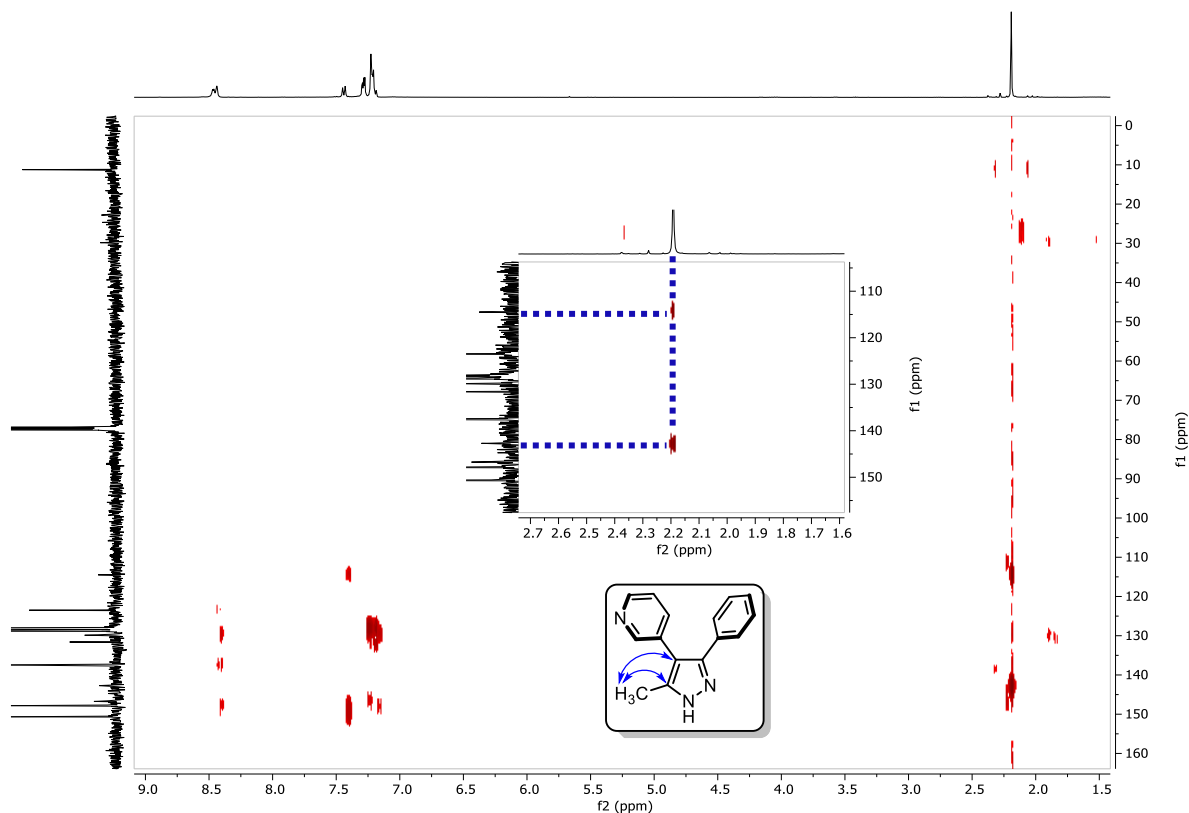
Spectrum 119. HRMS of **8x**.



Spectrum 120. ¹H-NMR of 8y.



Spectrum 121. ¹³C-NMR of 8y.

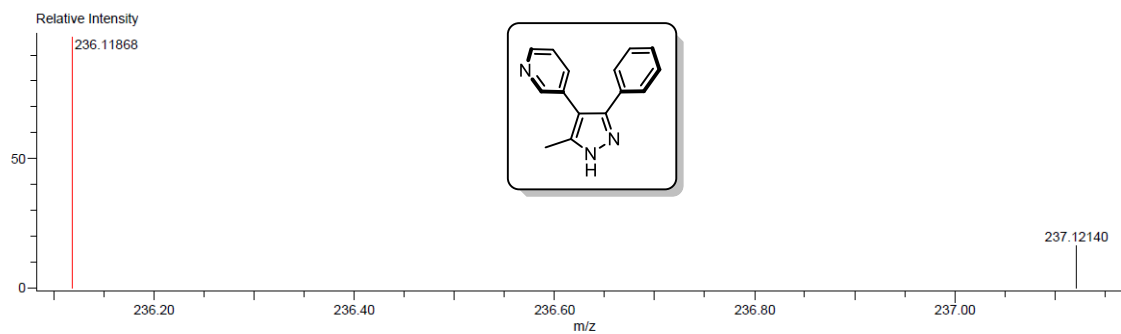


Spectrum 122. ^{13}C -NMR of **8y**.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...]
 Charge number:1
 Element: ^{12}C :0 .. 15, ^1H :0 .. 22, ^{79}Br :0 .. 0, ^{14}N :0 .. 3, ^{16}O :0 .. 0

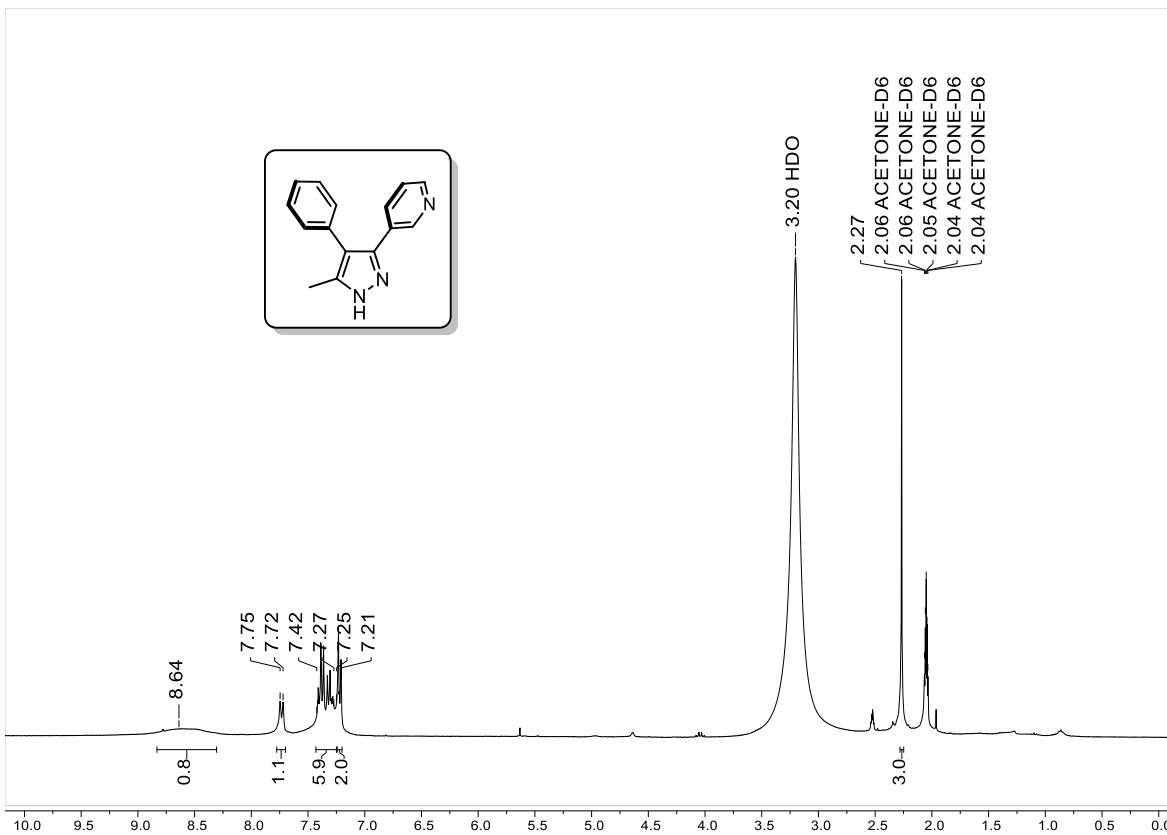
Mass Calibration data:Cal_PEG_600
 Created:2/21/2023 10:03:16 AM
 Created by:AccuTOF

Unsaturat. Number:0.0 .. 30.0 (Fraction:5)

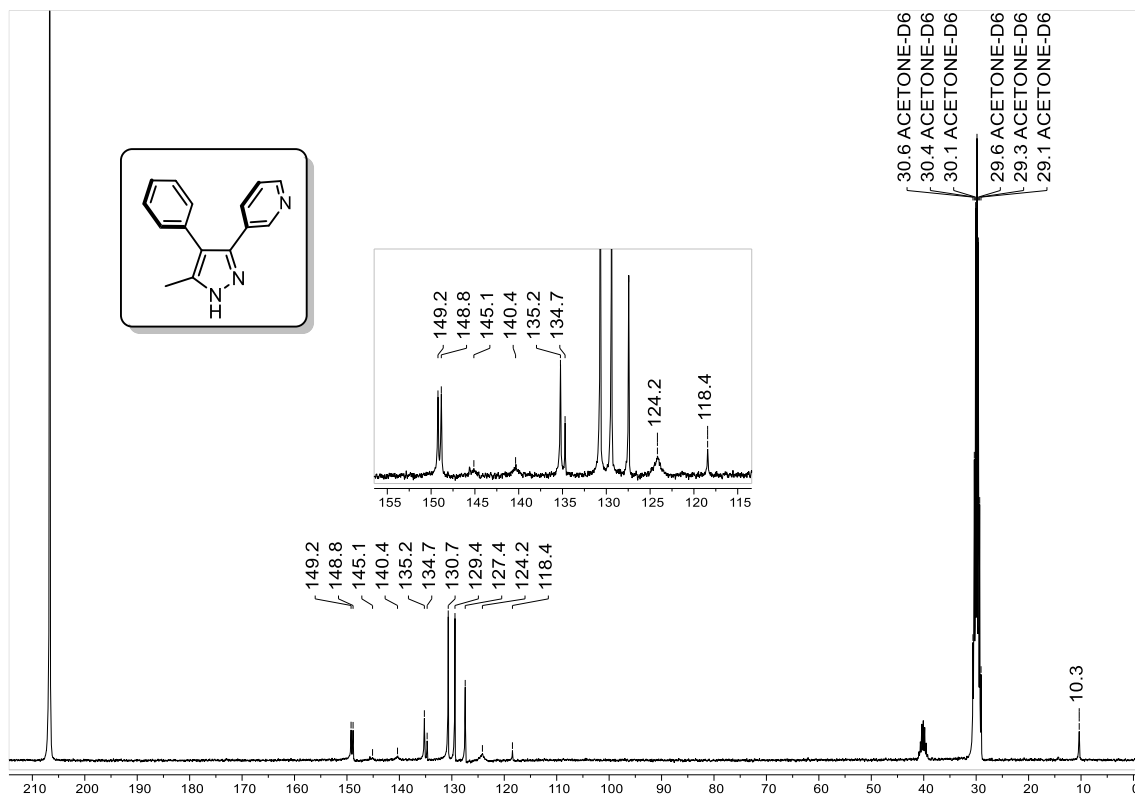


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturat. Number
236.11868	41968.84	236.11877	-0.10	-0.41	$^{12}\text{C}_{15}\text{H}_{14}\text{N}_3$	10.5

Spectrum 123. HRMS of **8y**.



Spectrum 124. $^1\text{H-NMR}$ of 8z.



Spectrum 125. $^{13}\text{C-NMR}$ of 8z.

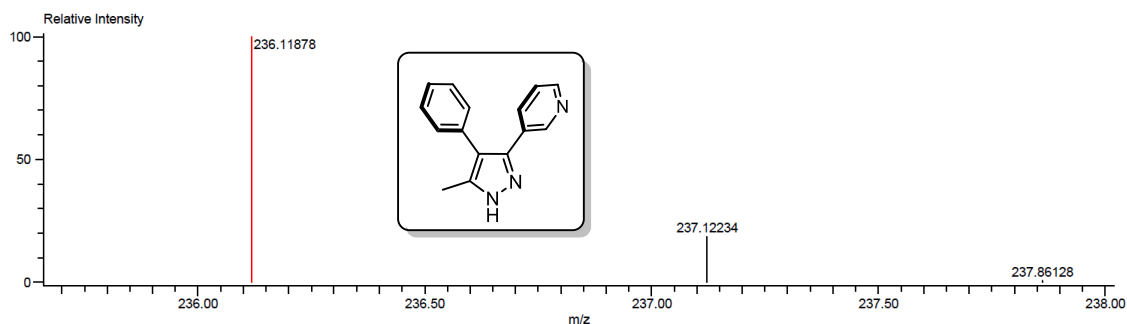
Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average(MS[...]

Charge number:1
 Element:¹²C:0 .. 15, ¹H:0 .. 22, ³⁵Cl:0 .. 0, ¹⁴N:3 .. 3, ¹⁶O:0 .. 0

Mass Calibration data:Cal_PEG_600
 Created:1/27/2023 10:29:43 AM
 Created by:AccuTOF

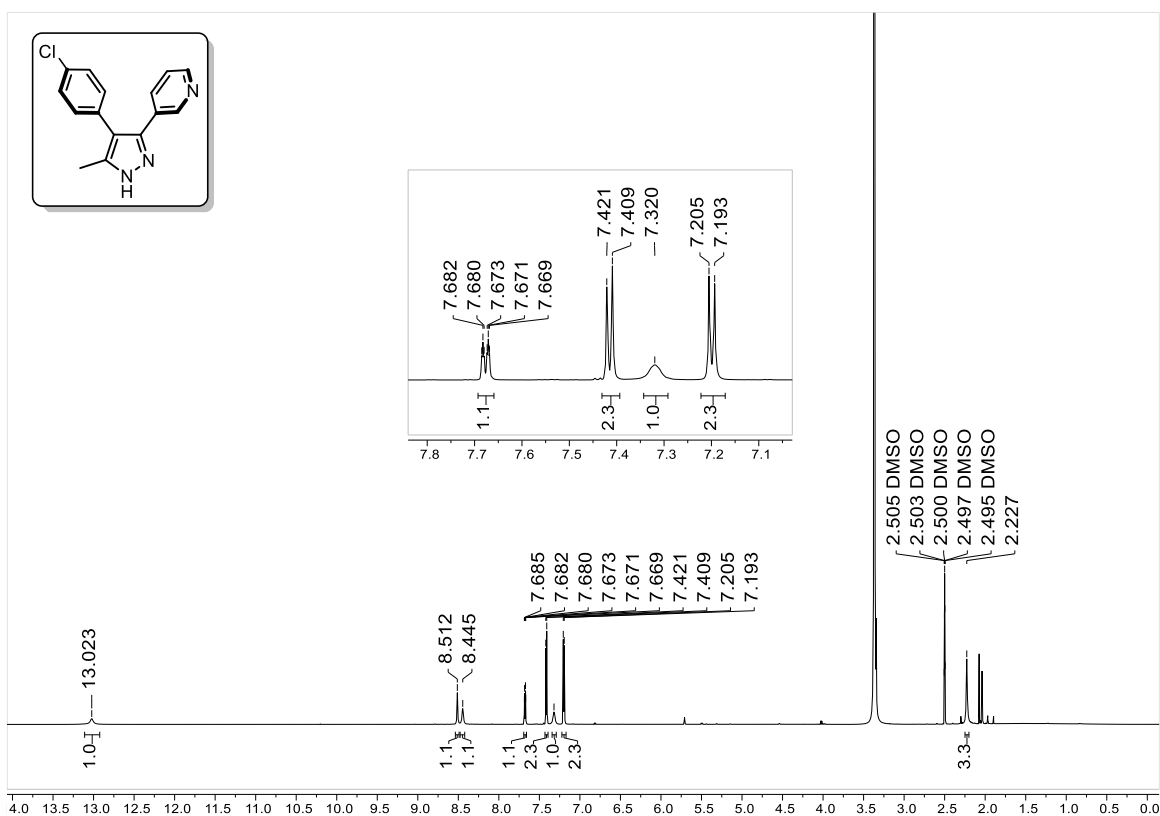
Tolerance:5.00(ppm), 5.00 .. 15.00(mmu)

Unsaturation Number:0.0 .. 45.0 (Fraction:Both)

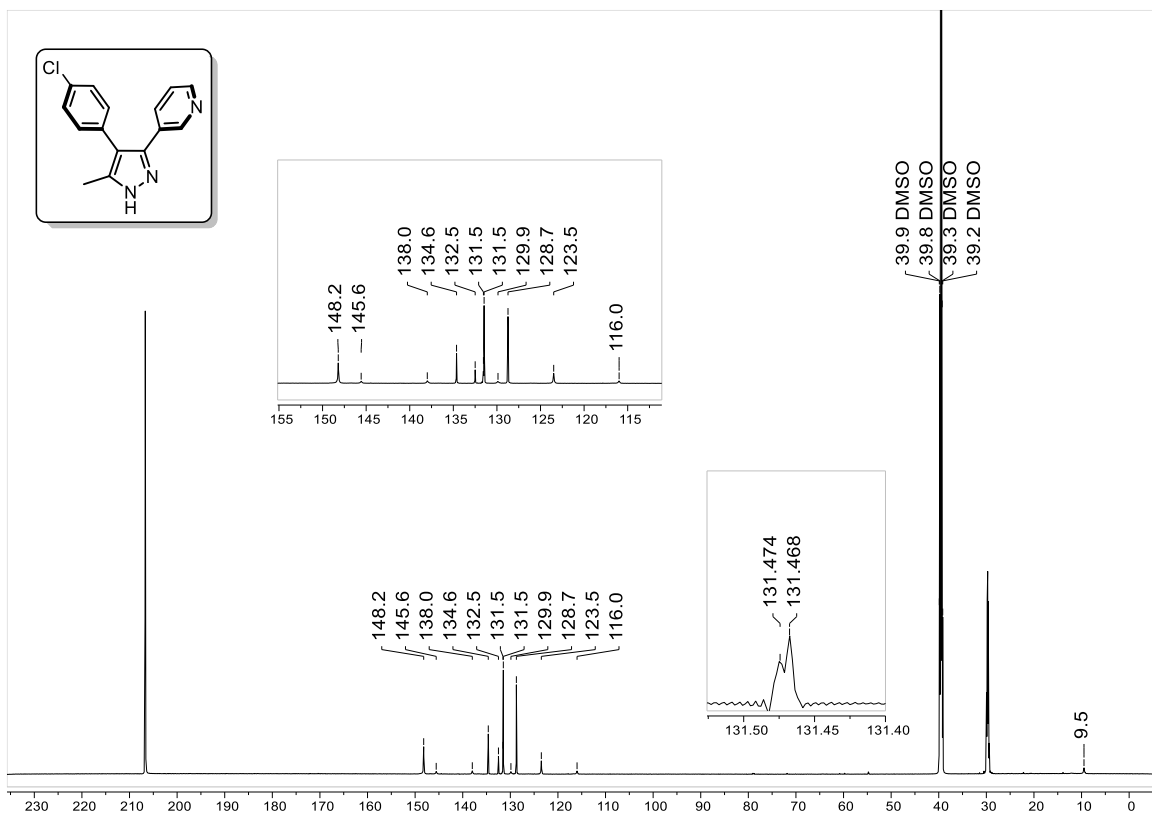


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
236.11878	11340.26	236.11877	0.01	0.04	¹² C ₁₅ ¹ H ₁₄ ¹⁴ N ₃	10.5

Spectrum 126. HRMS of 8z.



Spectrum 127. ¹H-NMR of 8aa.



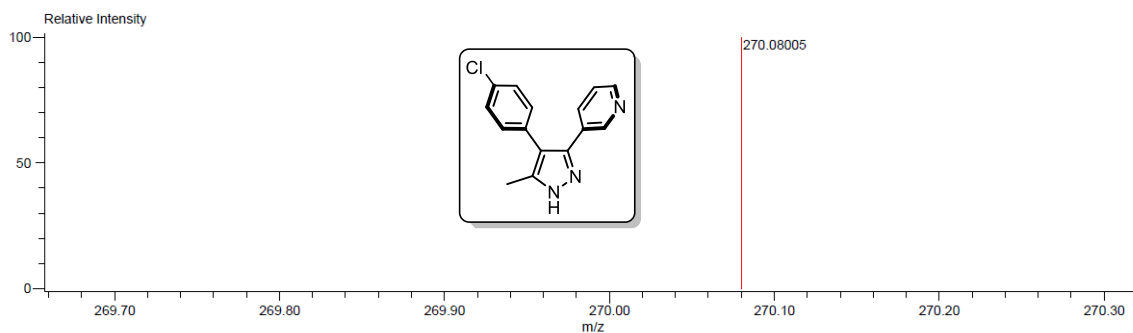
Spectrum 128. ¹³C-NMR of 8aa.

Description: Ionization Mode: ESI+
 History: Determine m/z [Peak Detect [Centroid, 30, Area], Correct Base[]; Smooth [5]]; Correct Base [5.0%]; Average [MS] ...

Mass Calibration data: Cal_PEG_600
 Created: 2/2/2023 11:17:31 AM
 Created by: AccuTOF

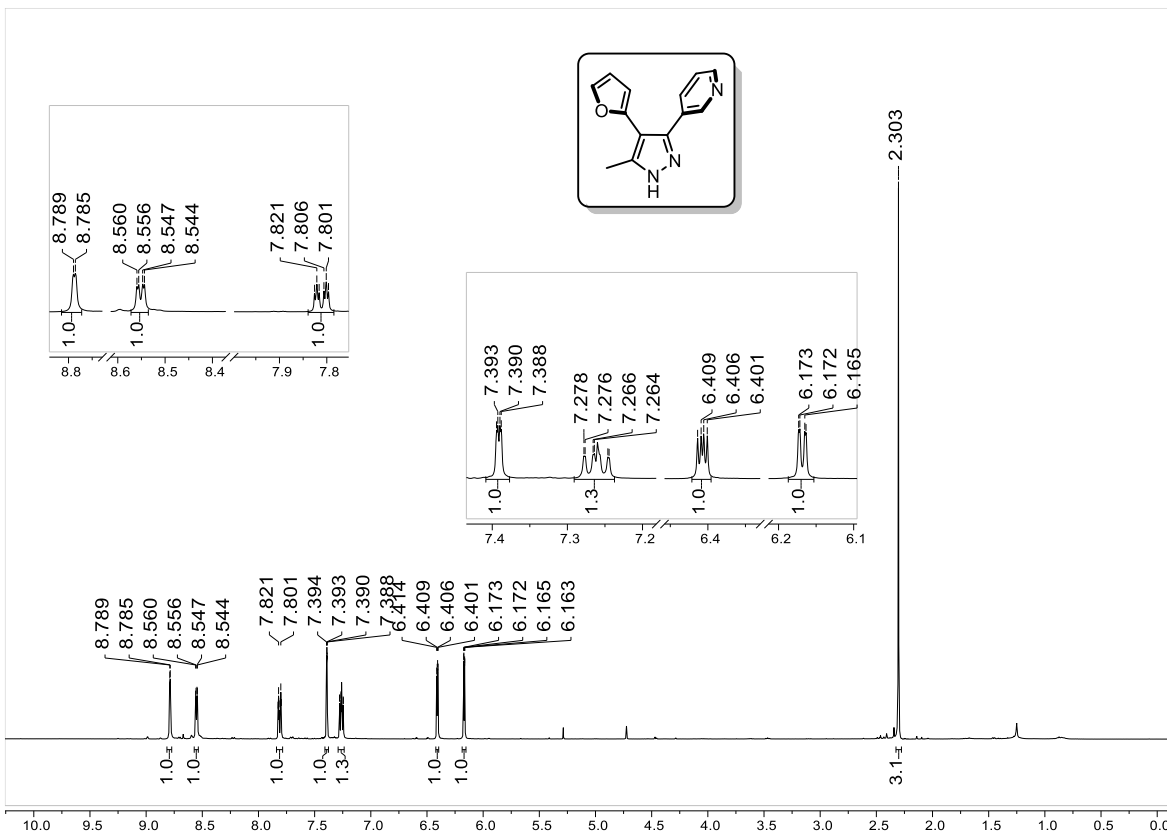
Charge number: 1
 Element: ¹²C: 0 .. 15, ¹H: 0 .. 50, ³⁵Cl: 0 .. 1, ¹⁹F: 0 .. 0, ¹⁴N: 0 .. 3, ¹⁸O: 0 .. 0

Tolerance: 50.00 (ppm), 5.00 .. 15.00 (mmu)
 Unsaturation Number: 0.0 .. 50.0 (Fraction: 5)

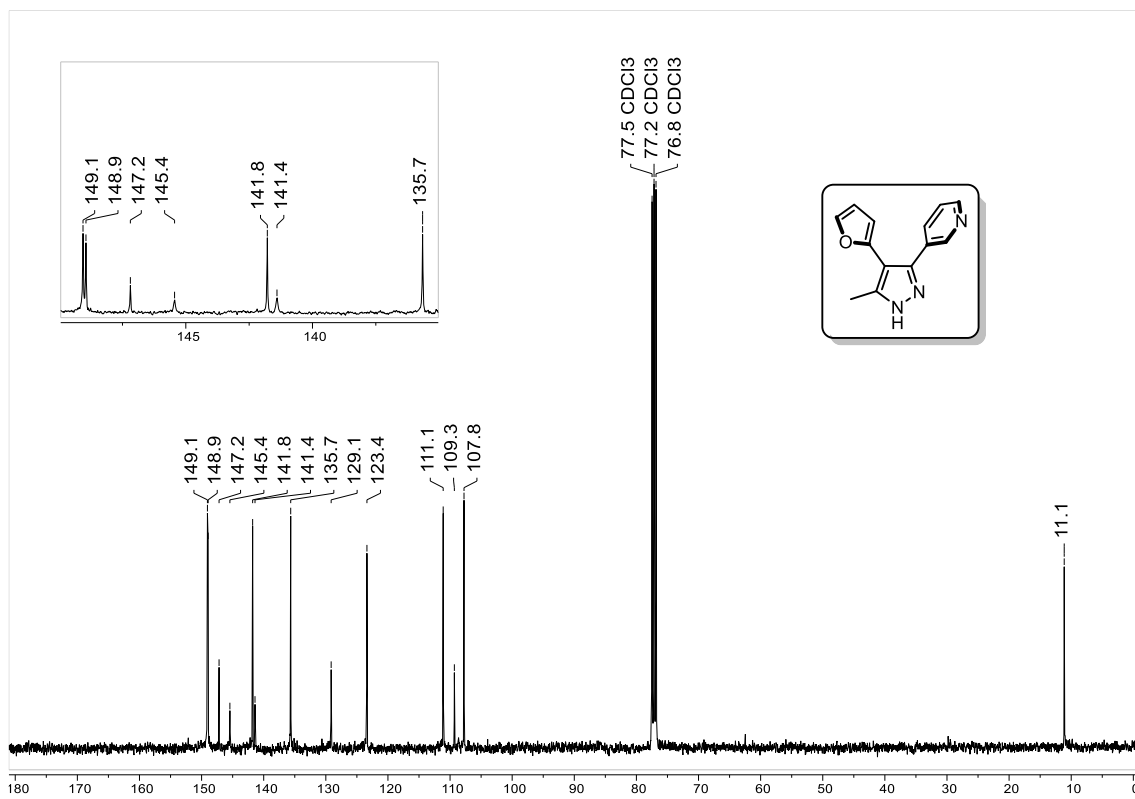


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
270.08005	4805.12	270.07980	0.25	0.92	¹² C ₁₅ ¹ H ₁₃ ³⁵ Cl ₁ ¹⁴ N ₃	10.5

Spectrum 129. HRMS of 8aa.



Spectrum 130. ¹H-NMR of 8ab.



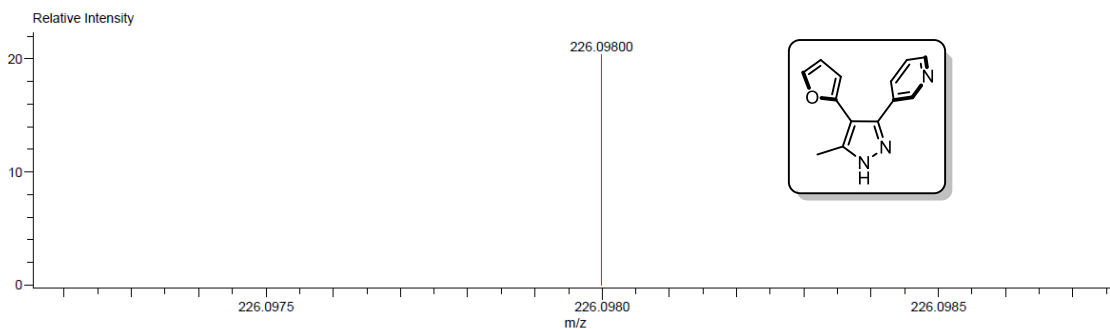
Spectrum 131. ¹³C-NMR of 8ab.

Description:
 Ionization Mode: ESI+
 History: Determine m/z [Peak Detect [Centroid, 30, Area], Correct Base [], Smooth [5]], Correct Base [5.0%], Average (MS[...

Mass Calibration data: Cal_PEG_600
 Created: 2/27/2023 2:22:47 PM
 Created by: AccuTOF

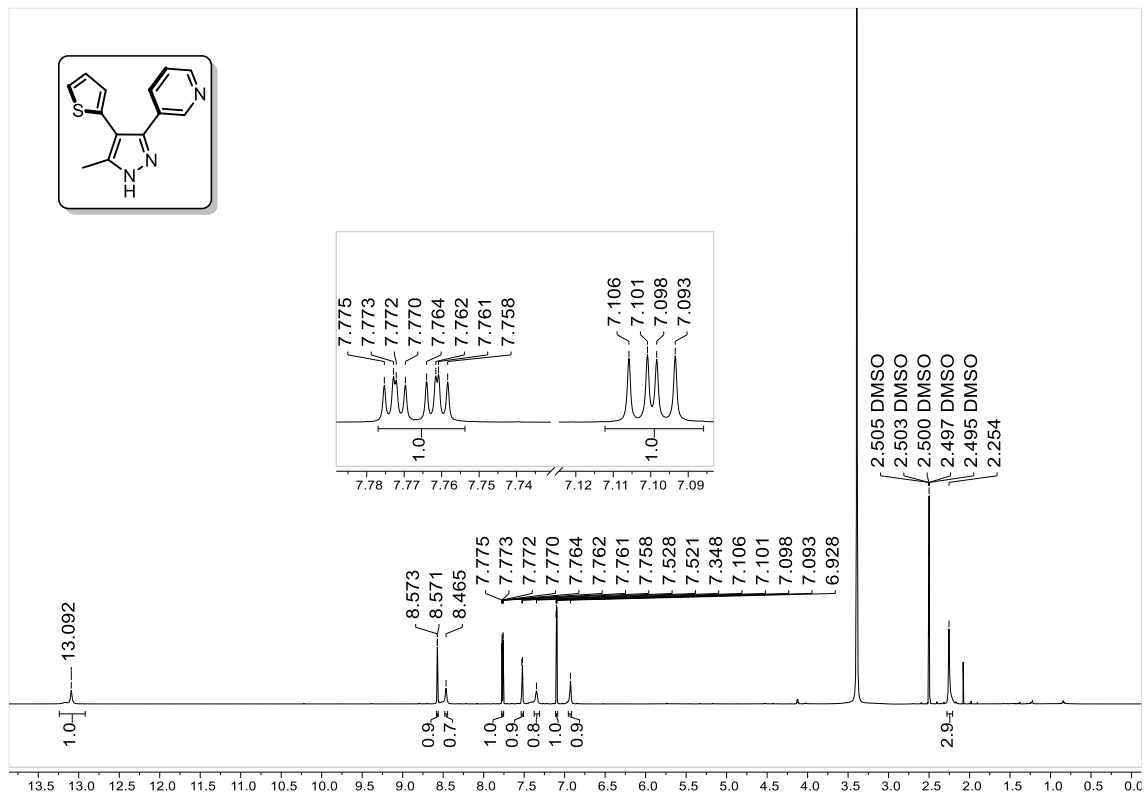
Charge number: 1
 Element: ^{12}C : 0 .. 13, ^1H : 0 .. 20, ^{35}Cl : 0 .. 0, ^{19}F : 0 .. 0, ^{14}N : 2 .. 3, ^{16}O : 0 .. 1

Unsaturat. Number: 0.0 .. 30.0 (Fraction: Both)

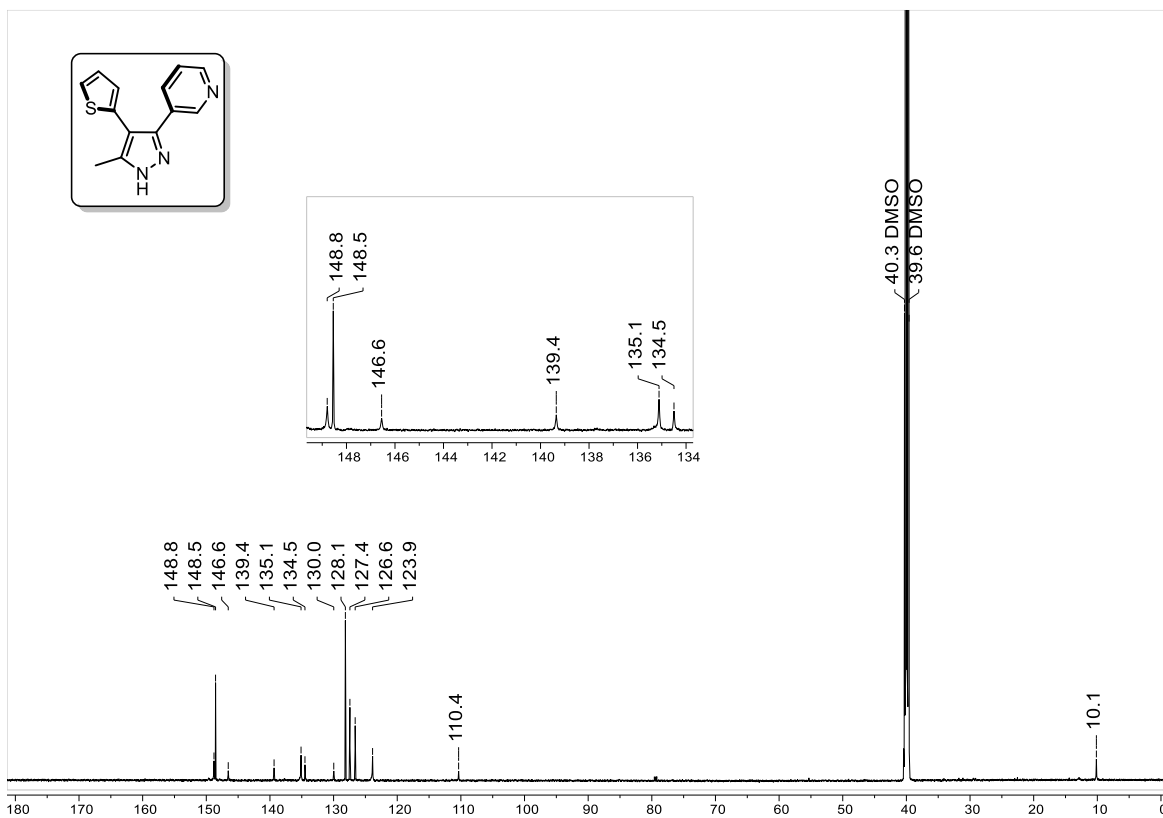


Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturat. Number
226.09800	18117.10	226.09804	-0.04	-0.17	$^{12}\text{C}_{13}\text{H}_{12}\text{N}_3\text{O}_1$	9.5

Spectrum 132. HRMS of 8ab.



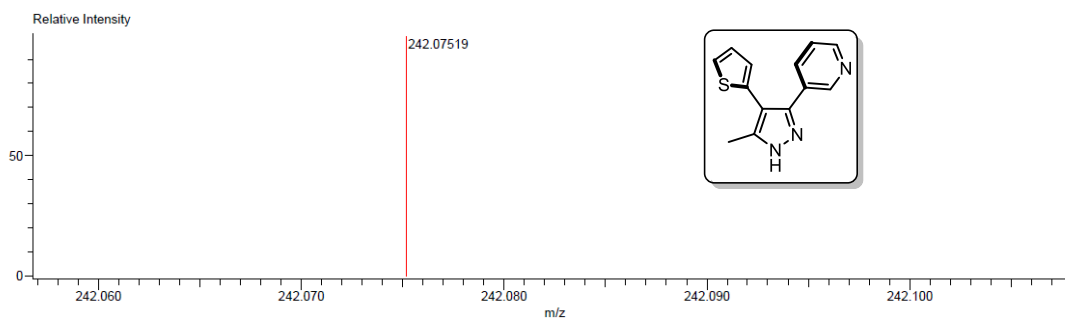
Spectrum 133. ^1H -NMR of 8ac.



Spectrum 134. ¹³C-NMR of 8ac.

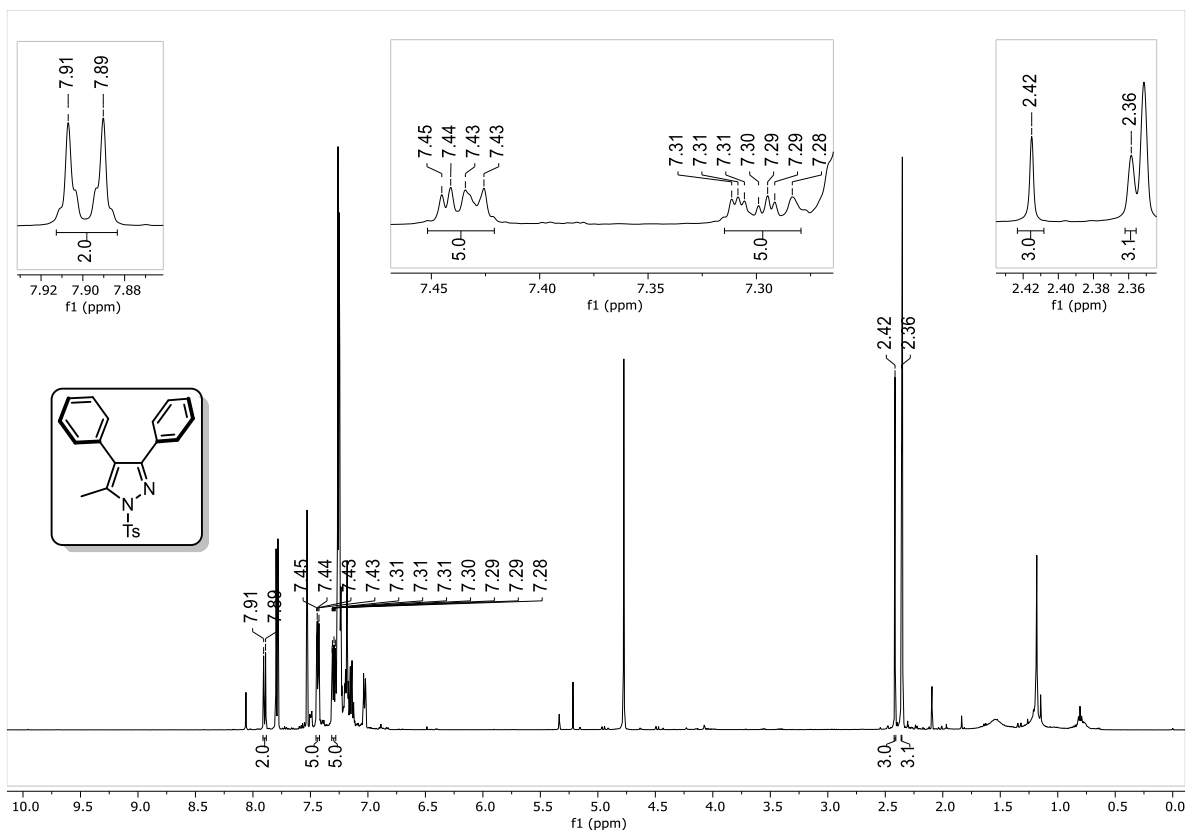
Description: Ionization Mode: ESI+
 History: Determine m/z [Peak Detect [Centroid, 30, Area], Correct Base [], Smooth [5]], Correct Base [5.0%], Average (MS...
 Mass Calibration data: Cal_PEG_600
 Created: 3/17/2023 11:26:55 AM
 Created by: AccuTOF

Charge number: 1
 Element: ¹²C: 0 .. 13, ¹H: 0 .. 18, ¹⁴N: 0 .. 3, ³²S: 1 .. 1
 Tolerance: 5.00 (mmu)
 Unsaturation Number: 0.0 .. 50.0 (Fraction: Both)



Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
242.07519	11321.62	242.07519	0.00	-0.01	¹² C ₁₃ ¹ H ₁₂ ¹⁴ N ₃ ³² S ₁	10.5

Spectrum 135. HRMS of 8ac.



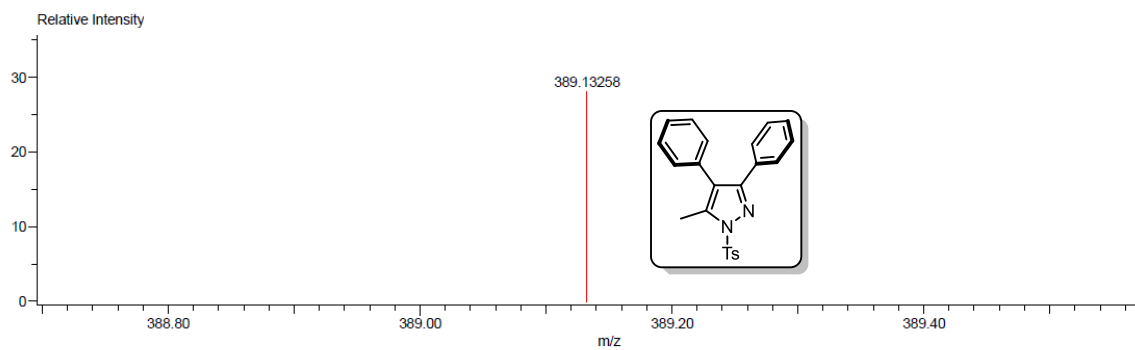
Spectrum 136. ¹H-NMR of 9.

Description:
 Ionization Mode:ESI+
 History:Determine m/z[Peak Detect[Centroid,30,Area];Correct Base[];Smooth[5]];Correct Base[5.0%];Average[MS[...

Mass Calibration data:Cal_PEG_600
 Created:6/14/2023 9:50:50 AM
 Created by:AccuTOF

Charge number:1
 Element:¹²C:0 .. 23, ¹H:0 .. 25, ¹⁴N:0 .. 2, ¹⁶O:2 .. 2, ³²S:0 .. 1
 Tolerance:4.00(mmu)

Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)



Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
389.13258	246534.04	389.13237	0.20	0.52	¹² C ₂₃ ¹ H ₂₁ ¹⁴ N ₂ ¹⁶ O ₂ ³² S ₁	15.5

Spectrum 137. HRMS of 9.

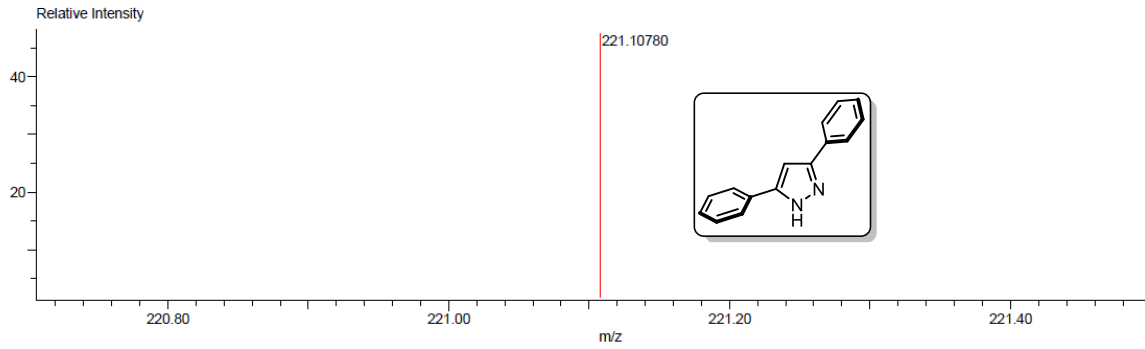
Description:
Ionization Mode:ESI+
History:Determine m/z[Peak Detect[Centroid,30,Area],Correct Base[],Smooth[5]];Correct Base[5.0%];Average(MS[...

Mass Calibration data:Cal_PEG_600
Created:6/14/2023 3:22:25 PM
Created by:AccuTOF

Charge number:1
Element:¹²C:0 .. 15, ¹H:0 .. 13, ¹⁴N:0 .. 2, ¹⁶O:0 .. 0

Tolerance:3.00(mmu)

Unsaturation Number:-1.0 .. 60.0 (Fraction:Both)



Mass	Intensity	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	Possible Formula	Unsaturation Number
221.10780	11449.31	221.10787	-0.07	-0.34	¹² C ₁₅ ¹ H ₁₃ ¹⁴ N ₂	10.5

Spectrum 138. HRMS of 11.

Comparative TLC of the methodologies.

Although Tang's and our conditions are similar, TLC outcomes reveal a different pattern. In our methodology, the 3,4-diaryl-1*H*-pyrazole ($R_f = 0.06$; Hex-AcOEt 8:2) is more polar than the 3,5-isomer ($R_f = 0.48$), which was obtained only traces (< 5%). It is noteworthy to mention that the yield of 3,4-regioisomer decreases when our protocol is carried out at 80 °C, while 3,5-regioisomer increases (Figure S1).

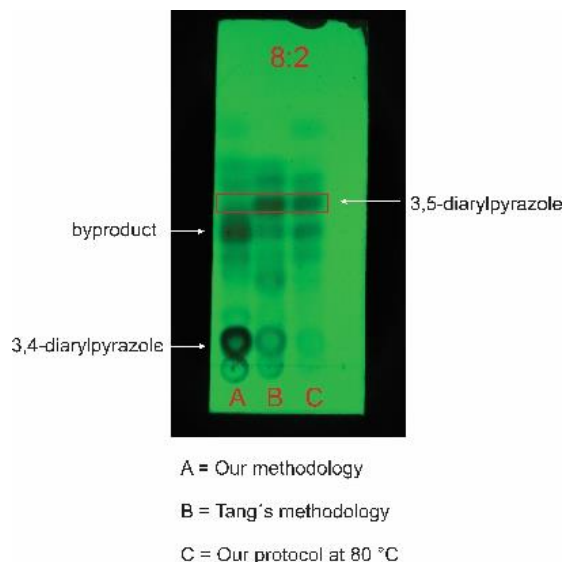


Figure S1. Comparative TLC of the methodologies (Eluent: Hex-AcOEt 8:2).

To bring a proof about the mechanism of 3,4-aza-heterocyclic synthesis, we tested our conditions without base (K_2CO_3). The result indicated the importance of the base for converting the *N*-tosylhydrazones to the corresponding diazo compounds; thus, the reaction proceeds *via* a [3+2] cycloaddition. Moreover, we identified that the most abundant byproduct as the tosylated pyrazole.

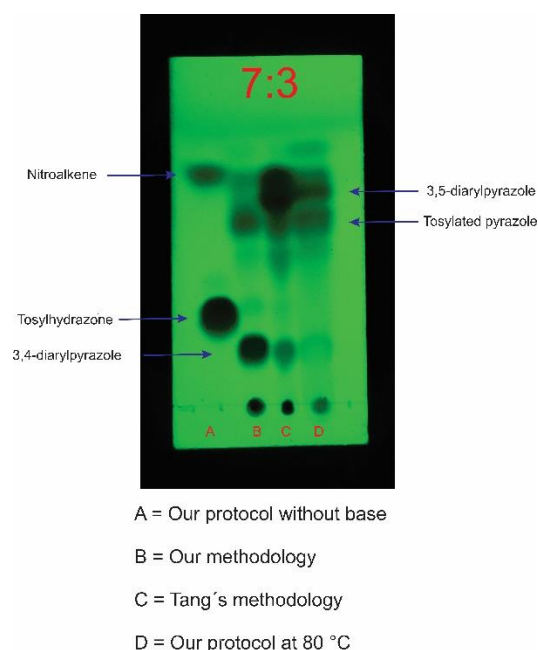


Figure S2. Comparative TLC of the methodologies (Eluent: Hex-AcOEt 7:3).

X-Ray Data.

Table S1. X-ray data.

Identification code	089HVE23	
Empirical formula	C ₁₅ H ₁₂ Cl N ₃	
Formula weight	269.73	
Temperature	100(2) K	
Wavelength	0.71073 Å	
Crystal system	Monoclinic	
Space group	P2 ₁ /c	
Unit cell dimensions	a = 14.1324(5) Å	α = 90°.
	b = 8.3919(3) Å	β = 103.5870(10)°.
	c = 11.1269(4) Å	γ = 90°.
Volume	1282.69(8) Å ³	
Z	4	
Density (calculated)	1.397 Mg/m ³	
Absorption coefficient	0.286 mm ⁻¹	
F(000)	560	
Crystal size	0.360 x 0.261 x 0.208 mm ³	
Theta range for data collection	2.844 to 27.948°.	
Index ranges	-18<=h<=18, -11<=k<=11, -14<=l<=14	
Reflections collected	18891	
Independent reflections	3087 [R(int) = 0.0187]	
Completeness to theta = 25.242°	99.9 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.7456 and 0.7186	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	3087 / 1 / 176	
Goodness-of-fit on F ²	1.070	
Final R indices [I>2sigma(I)]	R1 = 0.0304, wR2 = 0.0845	
R indices (all data)	R1 = 0.0323, wR2 = 0.0861	
Largest diff. peak and hole	0.407 and -0.222 e.Å ⁻³	

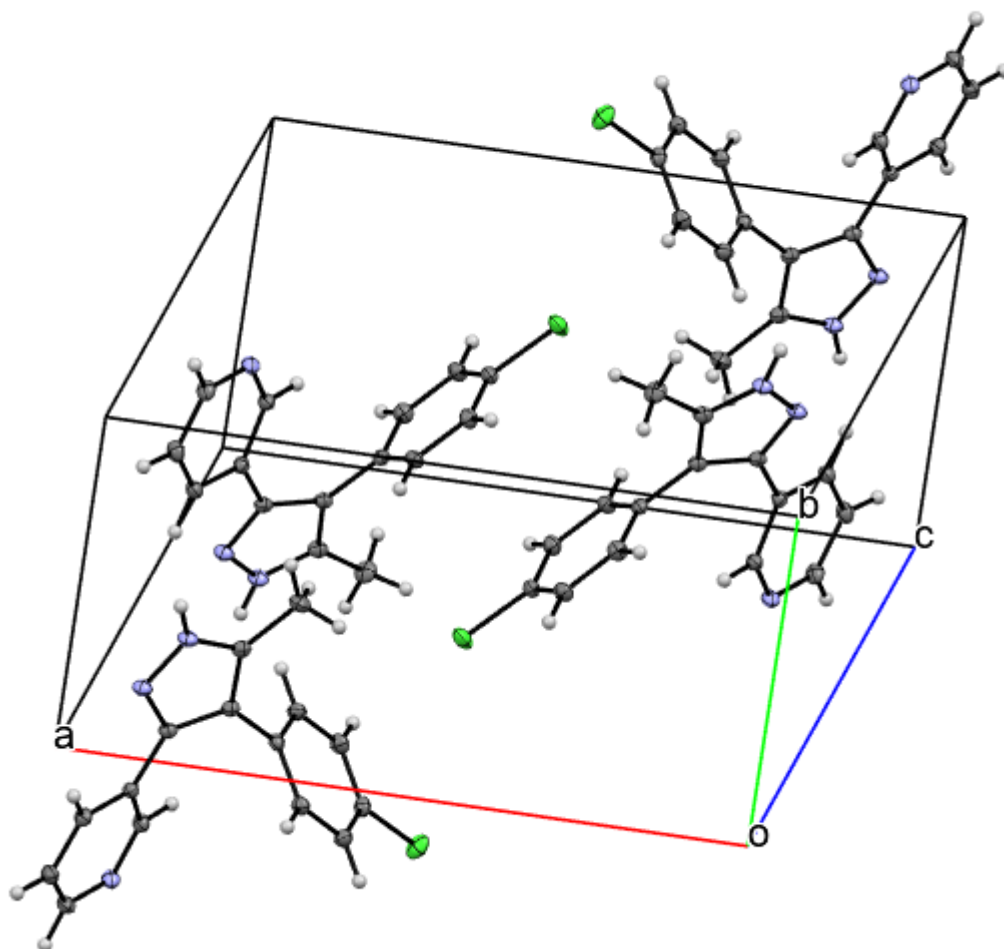


Figure S3. X-Ray of **8aa**.

Coordinates of optimized geometries.

Nitroalkene (6a).

@<TRIPOS>MOLECULE

21 21

SMALL

NO_CHARGES

@<TRIPOS>ATOM

1	C1	-3.7717	0.1353	0.0155	C
2	C2	-3.2296	-1.1239	0.2943	C
3	C3	-1.8505	-1.3157	0.2381	C
4	C4	-0.9791	-0.2471	-0.0677	C
5	C5	-1.5427	1.0120	-0.3651	C
6	C6	-2.9247	1.1958	-0.3245	C
7	H7	-4.8468	0.2864	0.0482	H
8	H8	-3.8805	-1.9565	0.5448	H
9	H9	-1.4323	-2.2979	0.4415	H
10	H10	-0.9097	1.8387	-0.6648	H
11	H11	-3.3425	2.1686	-0.5673	H
12	C12	0.4528	-0.5442	-0.0989	C
13	C13	1.5067	0.2822	0.0863	C
14	H14	0.6963	-1.5879	-0.2748	H
15	N15	2.8346	-0.3380	-0.0696	N
16	O16	2.9358	-1.5514	-0.3012	O
17	O17	3.8190	0.4075	0.0335	O
18	C18	1.5375	1.7302	0.4586	C
19	H19	1.7567	2.3724	-0.4024	H
20	H20	0.5749	2.0279	0.8762	H
21	H21	2.3106	1.9171	1.2075	H

@<TRIPOS>BOND

1 1 2 Ar
2 1 6 Ar
3 1 7 1
4 2 3 Ar
5 2 8 1
6 3 4 Ar
7 3 9 1
8 4 5 Ar
9 4 12 1
10 5 6 Ar
11 5 10 1
12 6 11 1
13 12 13 2
14 12 14 1
15 13 15 1
16 13 18 1
17 15 16 Ar
18 15 17 Ar
19 18 19 1
20 18 20 1
21 18 21 1

Diazo compound.

@<TRIPOS>MOLECULE

15 15

SMALL

NO_CHARGES

@<TRIPOS>ATOM

1	C1	-0.1288	-0.3729	0.0000	C
2	C2	0.9303	-1.3039	-0.0000	C
3	C3	0.1822	1.0026	0.0000	C
4	C4	2.2563	-0.8713	-0.0000	C
5	H5	0.7062	-2.3674	-0.0000	H
6	C6	-1.4979	-0.8781	0.0000	C
7	C7	1.5116	1.4258	0.0000	C
8	H8	-0.6138	1.7427	0.0000	H
9	C9	2.5572	0.4954	-0.0000	C

10 H10 3.0570 -1.6061 -0.0000 H
11 H11 -1.7300 -1.9370 0.0001 H
12 H12 1.7308 2.4902 0.0000 H
13 H13 3.5903 0.8304 -0.0000 H
14 N14 -2.5235 -0.0826 0.0000 N
15 N15 -3.4202 0.6342 -0.0000 N

@<TRIPOS>BOND

1 1 2 Ar
2 1 3 Ar
3 1 6 1
4 2 4 Ar
5 2 5 1
6 3 7 Ar
7 3 8 1
8 4 9 Ar
9 4 10 1
10 6 11 1
11 6 14 2
12 7 9 Ar
13 7 12 1
14 9 13 1
15 14 15 2

TS-1.

@<TRIPOS>MOLECULE

36 38

SMALL

NO_CHARGES

@<TRIPOS>ATOM

1 C1 3.3870 -2.9826 -0.2050 C
2 C2 2.6110 -2.7105 -1.3351 C
3 C3 1.7508 -1.6096 -1.3472 C
4 C4 1.6600 -0.7477 -0.2389 C
5 C5 2.4339 -1.0442 0.8981 C
6 C6 3.2900 -2.1471 0.9129 C
7 H7 4.0542 -3.8397 -0.1910 H
8 H8 2.6690 -3.3563 -2.2069 H
9 H9 2.3594 -0.4236 1.7859 H
10 H10 3.8774 -2.3570 1.8025 H
11 C11 0.7212 0.4088 -0.3366 C
12 C12 1.1231 1.7488 -0.1544 C
13 H13 -0.0274 0.2867 -1.1130 H
14 N14 0.3317 2.7315 -0.8427 N
15 O15 -0.8003 2.4248 -1.2798 O
16 O16 0.8047 3.8781 -0.9774 O
17 C17 2.4492 2.2454 0.3440 C
18 H18 2.3387 3.1620 0.9276 H
19 H19 2.9307 1.4941 0.9681 H
20 H20 3.1240 2.4702 -0.4922 H
21 C21 -1.9681 -0.5788 0.6024 C
22 C22 -2.1461 -1.9724 0.5604 C
23 C23 -2.9578 0.2590 0.0579 C
24 C24 -3.2925 -2.5196 -0.0193 C
25 H25 -1.3865 -2.6243 0.9834 H
26 C26 -0.7074 -0.0513 1.1693 C
27 C27 -4.1078 -0.2928 -0.5048 C
28 H28 -2.8236 1.3362 0.0685 H
29 C29 -4.2781 -1.6821 -0.5494 C
30 H30 -3.4185 -3.5981 -0.0472 H
31 H31 -0.1504 -0.7555 1.7869 H
32 H32 -4.8701 0.3627 -0.9159 H
33 H33 -5.1738 -2.1070 -0.9932 H
34 N34 -0.7189 1.1786 1.7702 N
35 N35 -0.2550 2.2317 1.7577 N
36 H36 1.1469 -1.4079 -2.2280 H

@<TRIPOS>BOND

1 1 2 Ar
2 1 6 Ar

3 1 7 1
4 2 3 Ar
5 2 8 1
6 3 4 Ar
7 3 36 1
8 4 5 Ar
9 4 11 1
10 5 6 Ar
11 5 9 1
12 6 10 1
13 11 12 Ar
14 11 13 1
15 11 26 Wk
16 12 14 1
17 12 17 1
18 12 35 Wk
19 14 15 Ar
20 14 16 Ar
21 17 18 1
22 17 19 1
23 17 20 1
24 21 22 Ar
25 21 23 Ar
26 21 26 1
27 22 24 Ar
28 22 25 1
29 23 27 Ar
30 23 28 1
31 24 29 Ar
32 24 30 1
33 26 31 1
34 26 34 Ar
35 27 29 Ar
36 27 32 1
37 29 33 1
38 34 35 2

TS-1'

@<TRIPOS>MOLECULE

36 38

SMALL

NO_CHARGES

@<TRIPOS>ATOM

1 C1	0.1519	0.5779	0.7236	C
2 C2	-1.2716	-0.3034	-0.9733	C
3 C3	2.5856	-0.3659	0.3146	C
4 C4	3.5608	-0.5269	1.3189	C
5 C5	3.0194	-0.2448	-1.0178	C
6 C6	4.9223	-0.5244	1.0106	C
7 H7	3.2458	-0.6475	2.3524	H
8 C8	1.1598	-0.4199	0.7509	C
9 C9	4.3815	-0.2567	-1.3273	C
10 H10	2.2951	-0.1461	-1.8161	H
11 C11	5.3389	-0.3877	-0.3171	C
12 H12	5.6544	-0.6364	1.8057	H
13 H13	1.0480	-1.0452	1.6344	H
14 H14	4.6932	-0.1644	-2.3642	H
15 H15	6.3972	-0.3923	-0.5627	H
16 N16	0.4749	-1.8574	-0.5281	N
17 N17	-0.4972	-1.4090	-1.0049	N
18 H18	-1.0192	0.3954	-1.7702	H
19 C19	-2.6711	-0.3864	-0.5591	C
20 C20	-3.5182	0.6981	-0.8595	C
21 C21	-3.1829	-1.4741	0.1765	C
22 C22	-4.8429	0.7002	-0.4227	C
23 H23	-3.1324	1.5362	-1.4337	H
24 C24	-4.5107	-1.4706	0.5985	C

25 H25 -2.5485 -2.3264 0.4034 H
26 C26 -5.3441 -0.3835 0.3056 C
27 H27 -5.4848 1.5441 -0.6582 H
28 H28 -4.8987 -2.3183 1.1559 H
29 H29 -6.3782 -0.3859 0.6374 H
30 N30 0.3632 1.8068 -0.0033 N
31 O31 1.0202 1.7917 -1.0602 O
32 O32 -0.1471 2.8481 0.4505 O
33 C33 -0.7998 0.6926 1.8880 C
34 H34 -1.7339 1.1863 1.6159 H
35 H35 -1.0329 -0.3089 2.2599 H
36 H36 -0.3492 1.2613 2.7106 H

@<TRIPOS>BOND

1 1 8 Ar
2 1 30 1
3 1 33 1
4 1 2 Wk
5 2 17 Ar
6 2 18 1
7 2 19 1
8 3 4 Ar
9 3 5 Ar
10 3 8 1
11 4 6 Ar
12 4 7 1
13 5 9 Ar
14 5 10 1
15 6 11 Ar
16 6 12 1
17 8 13 1
18 8 16 Wk
19 9 11 Ar
20 9 14 1
21 11 15 1
22 16 17 2
23 19 20 Ar
24 19 21 Ar
25 20 22 Ar
26 20 23 1
27 21 24 Ar
28 21 25 1
29 22 26 Ar
30 22 27 1
31 24 26 Ar
32 24 28 1
33 26 29 1
34 30 31 Ar
35 30 32 Ar
36 33 34 1
37 33 35 1
38 33 36 1

PS-1 (8a).

@<TRIPOS>MOLECULE

32 34

SMALL

NO_CHARGES

@<TRIPOS>ATOM

1 C1 2.7777 -2.8764 0.2604 C
2 C2 3.1791 -1.9363 -0.6932 C
3 C3 2.5221 -0.7059 -0.7893 C
4 C4 1.4543 -0.3865 0.0698 C
5 C5 1.0645 -1.3416 1.0281 C
6 C6 1.7162 -2.5731 1.1200 C
7 H7 3.2864 -3.8335 0.3346 H
8 H8 4.0004 -2.1611 -1.3685 H
9 H9 0.2500 -1.1131 1.7090 H
10 H10 1.3982 -3.2938 1.8687 H

11 C11	0.7764	0.9263	-0.0133	C
12 C12	1.4020	2.1755	0.0048	C
13 C13	2.8402	2.5786	0.0689	C
14 H14	3.2680	2.7152	-0.9315	H
15 H15	2.9547	3.5225	0.6119	H
16 H16	3.4311	1.8153	0.5818	H
17 C17	-1.7877	0.3093	-0.1043	C
18 C18	-2.9657	0.6265	0.5966	C
19 C19	-1.7571	-0.8717	-0.8667	C
20 C20	-4.0792	-0.2143	0.5382	C
21 H21	-3.0009	1.5340	1.1916	H
22 C22	-0.6249	1.2229	-0.0508	C
23 C23	-2.8709	-1.7129	-0.9222	C
24 H24	-0.8650	-1.1275	-1.4289	H
25 C25	-4.0357	-1.3898	-0.2188	C
26 H26	-4.9787	0.0461	1.0896	H
27 H27	-2.8290	-2.6191	-1.5204	H
28 H28	-4.9005	-2.0462	-0.2613	H
29 N29	-0.8328	2.5471	-0.0535	N
30 N30	0.4020	3.0915	-0.0162	N
31 H31	2.8311	0.0082	-1.5474	H
32 H32	0.5014	4.0987	-0.0094	H

@<TRIPOS>BOND

1 1 2 Ar
2 1 6 Ar
3 1 7 1
4 2 3 Ar
5 2 8 1
6 3 4 Ar
7 3 31 1
8 4 5 Ar
9 4 11 1
10 5 6 Ar
11 5 9 1
12 6 10 1
13 11 12 Ar
14 11 22 1
15 12 13 1
16 12 30 Ar
17 13 14 1
18 13 15 1
19 13 16 1
20 17 18 Ar
21 17 19 Ar
22 17 22 1
23 18 20 Ar
24 18 21 1
25 19 23 Ar
26 19 24 1
27 20 25 Ar
28 20 26 1
29 22 29 Ar
30 23 25 Ar
31 23 27 1
32 25 28 1
33 29 30 Ar
34 30 32 1

PS-1'

@<TRIPOS>MOLECULE

32 34
SMALL
NO_CHARGES

@<TRIPOS>ATOM

1 C1	0.0092	0.4407	-0.0319	C
2 C2	1.1186	-0.4539	0.0026	C
3 C3	-2.5600	-0.1225	-0.0204	C
4 C4	-3.0969	0.9930	-0.6902	C

5 C5 -3.4429 -0.9869 0.6574 C
6 C6 -4.4720 1.2361 -0.6787 C
7 H7 -2.4427 1.6620 -1.2380 H
8 C8 -1.1140 -0.3946 -0.0293 C
9 C9 -4.8173 -0.7444 0.6603 C
10 H10 -3.0529 -1.8426 1.2019 H
11 C11 -5.3385 0.3697 -0.0055 C
12 H12 -4.8664 2.1013 -1.2044 H
13 H13 -5.4799 -1.4214 1.1924 H
14 H14 -6.4078 0.5610 0.0013 H
15 N15 -0.6308 -1.6698 0.0028 N
16 N16 0.7121 -1.7346 0.0119 N
17 C17 2.5691 -0.1550 0.0192 C
18 C18 3.1125 0.7847 0.9129 C
19 C19 3.4393 -0.8383 -0.8491 C
20 C20 4.4859 1.0420 0.9292 C
21 H21 2.4618 1.2994 1.6135 H
22 C22 4.8126 -0.5843 -0.8285 C
23 H23 3.0320 -1.5671 -1.5434 H
24 C24 5.3409 0.3602 0.0576 C
25 H25 4.8882 1.7681 1.6304 H
26 H26 5.4690 -1.1209 -1.5083 H
27 H27 6.4087 0.5604 0.0714 H
28 C28 0.0320 1.9416 -0.0969 C
29 H29 1.0127 2.3347 0.1771 H
30 H30 -0.1889 2.3081 -1.1071 H
31 H31 -0.7063 2.3863 0.5792 H
32 H32 -1.1676 -2.5270 -0.0203 H

@<TRIPOS>BOND

1 1 2 Ar
2 1 8 Ar
3 1 28 1
4 2 16 Ar
5 2 17 1
6 3 4 Ar
7 3 5 Ar
8 3 8 1
9 4 6 Ar
10 4 7 1
11 5 9 Ar
12 5 10 1
13 6 11 Ar
14 6 12 1
15 8 15 Ar
16 9 11 Ar
17 9 13 1
18 11 14 1
19 15 16 1
20 15 32 1
21 17 18 Ar
22 17 19 Ar
23 18 20 Ar
24 18 21 1
25 19 22 Ar
26 19 23 1
27 20 24 Ar
28 20 25 1
29 22 24 Ar
30 22 26 1
31 24 27 1
32 28 29 1
33 28 30 1
34 28 31 1

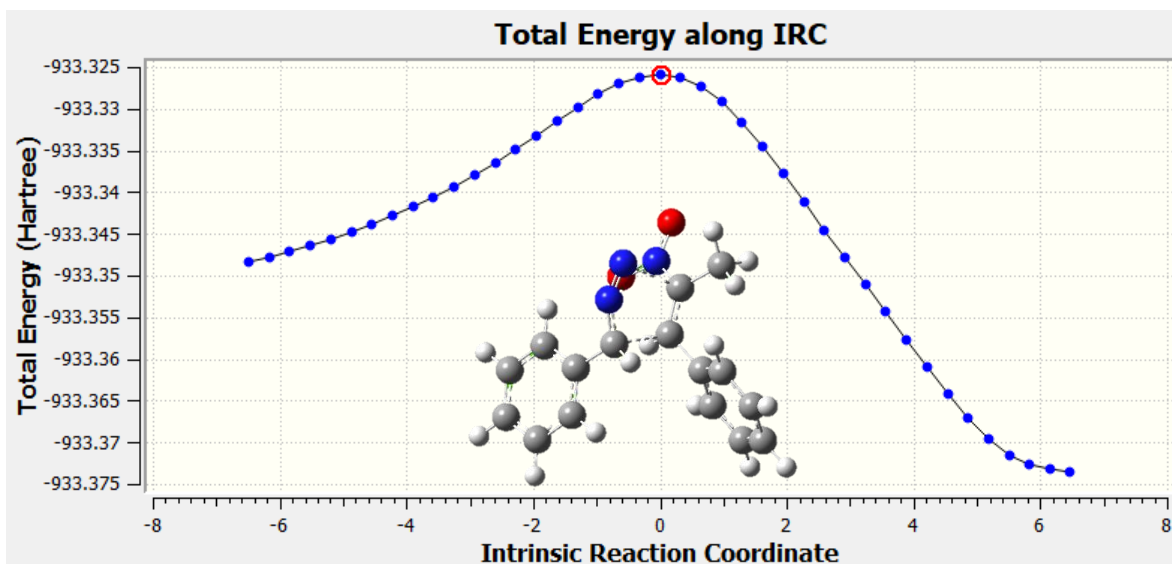
Table S2. Calculated energy values.

Molecules	G	H	E-opt
Diazo Compound	-379.751428	-379.710897	-379.833602
Nitroalkene (6a)	-553.394375	-553.345723	-553.521571
TS-1	-933.093703	-933.026443	-933.325866
TS-1'	-933.082648	-933.015308	-933.314513
PS-1 (8a)	-727.488152	-727.430168	-727.706292
PS-1'	-727.486576	-727.428357	-727.704591

Values shown at Hartree at 298.15 K

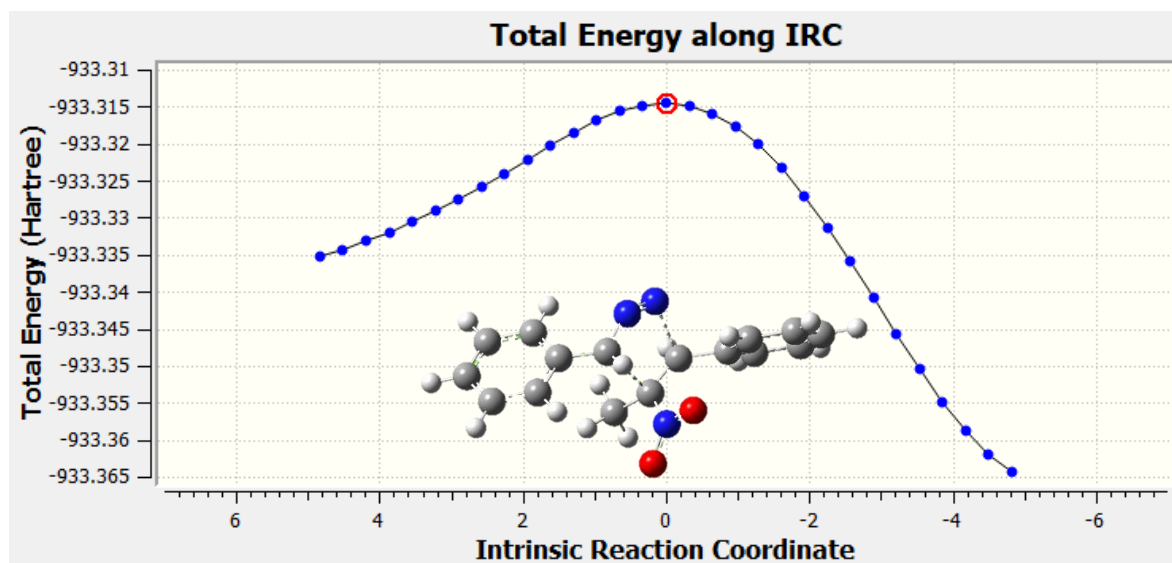
Intrinsic Reaction Coordinate (IRC) of Transition States.

IRC TS-1.



Negative frequency magnitude: -409.46

IRC TS-1'.



Negative frequency magnitude: -405.71

Results of Cytotoxicity screening.

Table S3. Cytotoxicity activity of pyrazoles in different cell lines.

Compound	% growth inhibition by cell line ^a			
	PC-3 ^b	HCT-15 ^c	MCF-7 ^d	COS-7 ^e
8a	12.29	0.13	4.35	NC ^f
8b	13.71	5.66	11.78	8.50
8c	18.22	10.92	9.90	16.20
8d	NC	0.14	20.68	12.37
8e	24.21	29.69	30.23	32.80
8f	25.08	NC	6.09	2.55
8g	11.94	1.86	NC	1.17
8h	11.72	8.22	45.56	28.46
8i	36.34	27.27	26.38	18.50
8j	4.04	8.08	35.47	31.39
8k	28.59	12.91	15.52	30.07
8l	7.06	NC	21.93	15.80
8m	14.41	2.50	2.90	4.40
8n	6.57	NC	NC	NC
8o	27.07	21.88	36.78	42.52
8p	8.98	7.29	34.90	34.33
8q	11.54	9.97	35.82	23.37
8r	4.84	5.12	27.68	12.92
8s	23.37	10.74	NC	16.83
8t	5.29	16.03	1.45	NC
8u	8.83	3.61	NC	NC
8v	18.91	21.14	20.14	7.42
8w	5.88	NC	9.81	4.85
8x	17.74	1.14	22.28	6.94
8y	6.82	0.40	NC	NC
8aa	10.68	2.73	5.90	7.26
8ab	6.42	NC	0.32	3.19
8ac	15.69	10.09	10.63	10.61

a) Concentration: 25 μ M; Vehicle: DMSO. b) Prostate cancer. c) Colon cancer. d) Breast cancer. e) Monkey's kidney cell line (not cancerous). f) Non-cytotoxic.