

Supporting Information  
for DOI: 10.1055/s-0039-1690179

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# Synthesis of Aza-polyquinanes via Fischer indolization and Ring-Rearrangement Metathesis as Key Steps

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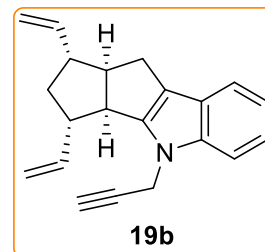
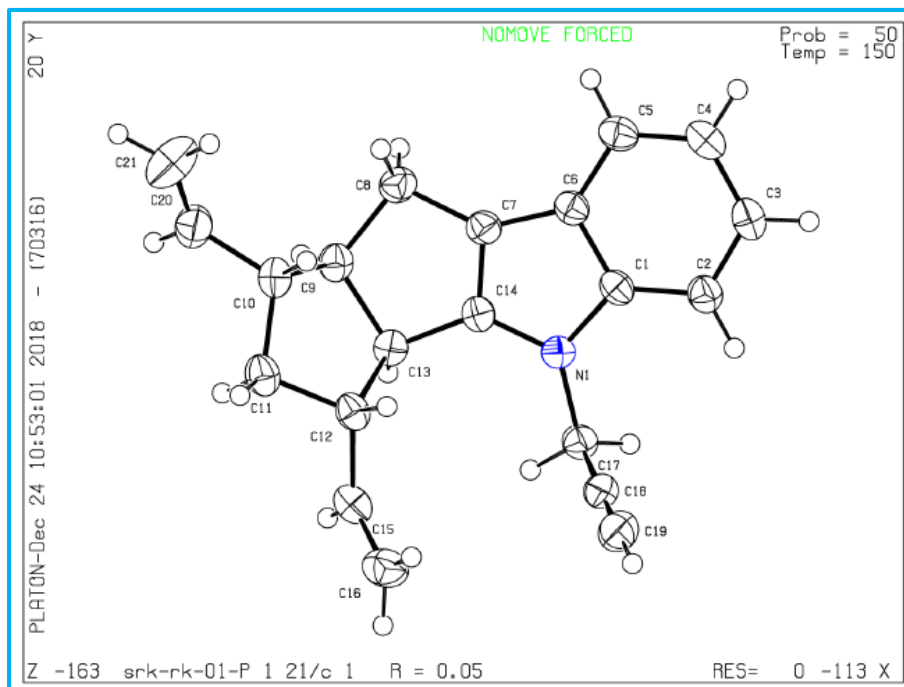
## Supporting Information

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**1. X-ray data and refinement parameters for target compounds  
(1R,3S,3aS,9aR)-4-(prop-2-yn-1-yl)-1,3-divinyl-2,3,3a,4,9,9a-hexahydro-1H-pentaleno[1,2-b]indole (19b)**

CCDC Number = 1886978



**Table S1.** X-ray crystallographic data and refinement parameters for **19b** (CCDC 1886978)

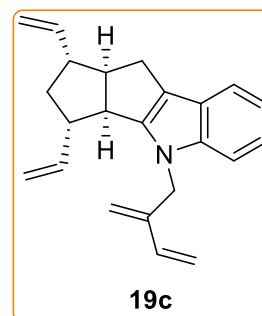
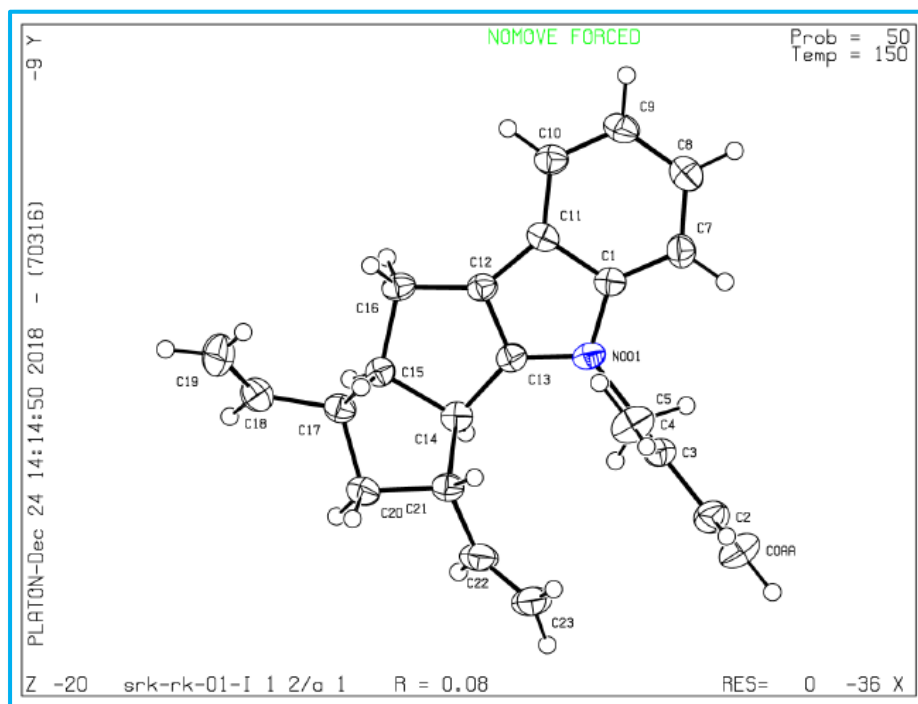
Identification code	SRK-RK-01-50-2_mo
Empirical formula	C <sub>21</sub> H <sub>21</sub> N
Formula weight	287.39
Temperature	150 K
Crystal system	Monoclinic
Space group	P2 <sub>1</sub> /c
Unit cell dimensions	a = 16.9888 (12) Å    α = 90° b = 4.9020 (4) Å    β = 90.857° (7) c = 19.1064 (15) Å    γ = 90°
Volume	1591.0 (2) Å <sup>3</sup>
Z	4
Density (calculated)	1.200 g/cm <sup>3</sup>
Absorption coefficient (μ)	0.069 mm <sup>-1</sup>
Absorption correction	Multi-scan
Max. and Min. transmission	0.997-0.992

F (000)	616.0
Crystal size	0.172 x 0.099 x 0.038 mm <sup>3</sup>
Index ranges	-20 ≤ h ≤ 19, -5 ≤ k ≤ 4, -19 ≤ l ≤ 22
Theta range for data collection	2.398 to 24.994°
Reflections collected	0.0544 (1930)
Diffraction radiation wavelength	0.71073
Independent reflections	2792 [ R <sub>(int)</sub> = 0.0734]
Completeness to θ = 24.994°	99.8%
Refinement method	Full-matrix least-squares on F <sup>2</sup>
Data/restraints/parameters	2792/0/199
Goodness-of-fit on F <sup>2</sup>	1.059
Final R indices [I ≥ 2σ (I)]	R1 = 0.0544, wR2 = 0.1180
R indices (all data)	R1 = 0.0851, wR2 = 0.1429
Largest diff. peak and hole	0.24/-0.24 e Å <sup>-3</sup>

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**(1R,3S,3aS,9aR)-4-(2-methylenebut-3-en-1-yl)-1,3-divinyl-2,3,3a,4,9,9a-hexahydro-1H-pentaleno[1,2-b]indole (19c)**

CCDC Number = 1887151



**Table S2.** X-ray crystallographic data and refinement parameters for **19c** (CCDC 1887151)

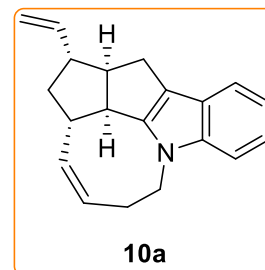
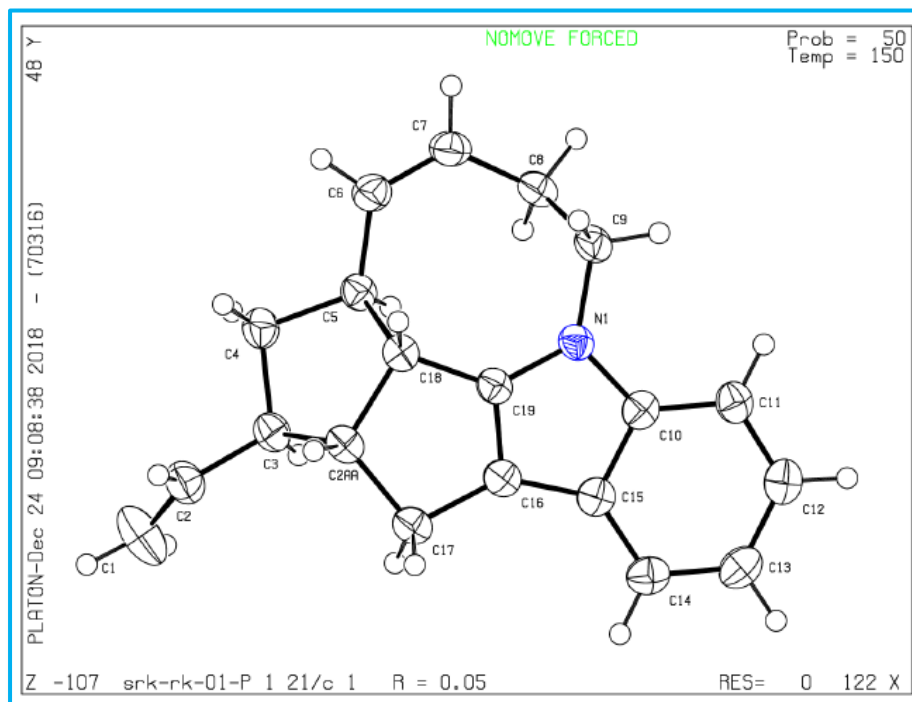
Identification code	SRK-RK-01-87_cu
Empirical formula	C <sub>23</sub> H <sub>25</sub> N
Formula weight	315.44
Temperature	150 K
Crystal system	Monoclinic
Space group	I2/a
Unit cell dimensions	a = 22.8136 (15) Å    α = 90° b = 8641 (4) Å    β = 109.643° (7) c = 24.7742 (13) Å    γ = 90°
Volume	3653.8 (4) Å <sup>3</sup>
Z	8
Density (calculated)	1.147 g/cm <sup>3</sup>
Absorption coefficient (μ)	0.494 mm <sup>-1</sup>
Absorption correction	Multi-scan
Max. and Min. transmission	0.980-0.900
F (000)	1360.0

Crystal size	0.213 x 0.198 x 0.041 mm <sup>3</sup>
Index ranges	-26 ≤ h ≤ 24, -8 ≤ k ≤ 4, -28 ≤ l ≤ 29
Theta range for data collection	3.789 to 64.998°
Reflections collected	0.0833 (1837)
Diffraction radiation wavelength	1.54184
Independent reflections	3029 [ R <sub>(int)</sub> = 0.1256]
Completeness to θ = 64.998°	97.6%
Refinement method	Full-matrix least-squares on F <sup>2</sup>
Data/restraints/parameters	3029/0/217
Goodness-of-fit on F <sup>2</sup>	1.074
Final R indices [I ≥ 2σ (I)]	R1 = 0.0833, wR2 = 0.2020
R indices (all data)	R1 = 0.1370, wR2 = 0.2750
Largest diff. peak and hole	0.55/-0.58 e Å <sup>-3</sup>

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## Compound 10a

CCDC Number = 1886977



**Table S3.** X-ray crystallographic data and refinement parameters for **10a** (CCDC 1886977)

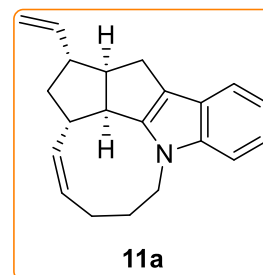
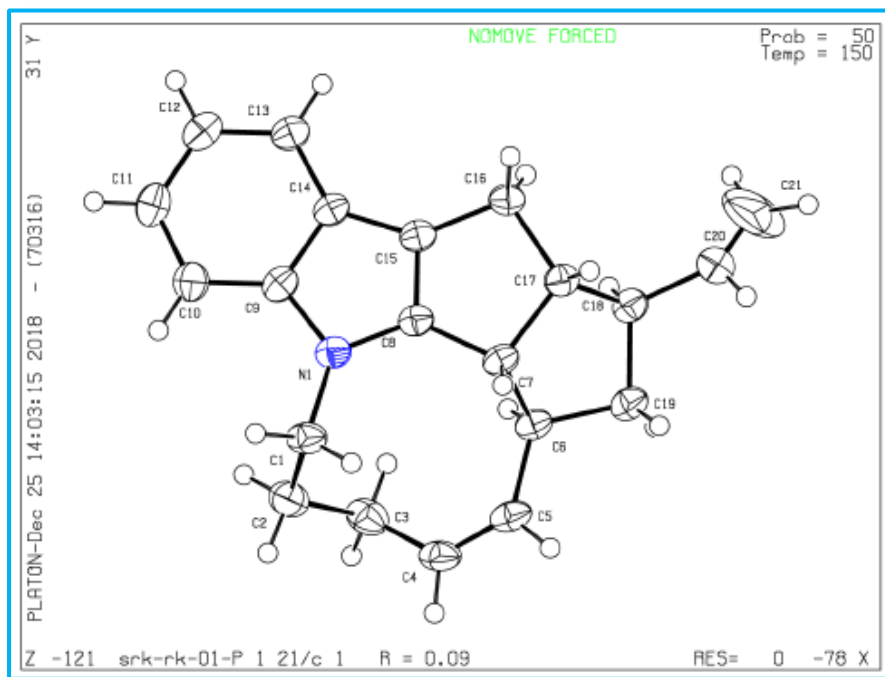
Identification code	SRK-RK-01-81-2_mo	
Empirical formula	C <sub>20</sub> H <sub>21</sub> N	
Formula weight	275.38	
Temperature	150 K	
Crystal system	Monoclinic	
Space group	P2 <sub>1</sub> /c	
Unit cell dimensions	a = 5.3134(3) Å	α = 90°
	b = 12.2320(6) Å	β = 96.223° (5)
	c = 23.2718(11) Å	γ = 90°
Volume	1503.60(13) Å <sup>3</sup>	
Z	4	
Density (calculated)	1.217 g/cm <sup>3</sup>	
Absorption coefficient (μ)	0.070 mm <sup>-1</sup>	
Absorption correction	Multi-scan	
Max. and Min. transmission	0.994-0.992	
F (000)	592.0	

Crystal size	0.235 x 0.097 x 0.08 mm <sup>3</sup>
Index ranges	-6 ≤ h ≤ 6, -14 ≤ k ≤ 14, -27 ≤ l ≤ 27
Theta range for data collection	2.423 to 25.000°
Reflections collected	0.0499 (1990)
Diffraction radiation wavelength	0.71073
Independent reflections	2549 [ R <sub>(int)</sub> = 0.0753]
Completeness to θ = 24.99°	96.3%
Refinement method	Full-matrix least-squares on F <sup>2</sup>
Data/restraints/parameters	2549/0/190
Goodness-of-fit on F <sup>2</sup>	1.057
Final R indices [I ≥ 2σ (I)]	R1 = 0.0499, wR2 = 0.1211
R indices (all data)	R1 = 0.0685, wR2 = 0.1397
Largest diff. peak and hole	0.25/-0.33 e Å <sup>-3</sup>

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## Compound 11a

CCDC Number = 1887288



**Table S4.** X-ray crystallographic data and refinement parameters for **11a** (CCDC 1887288)

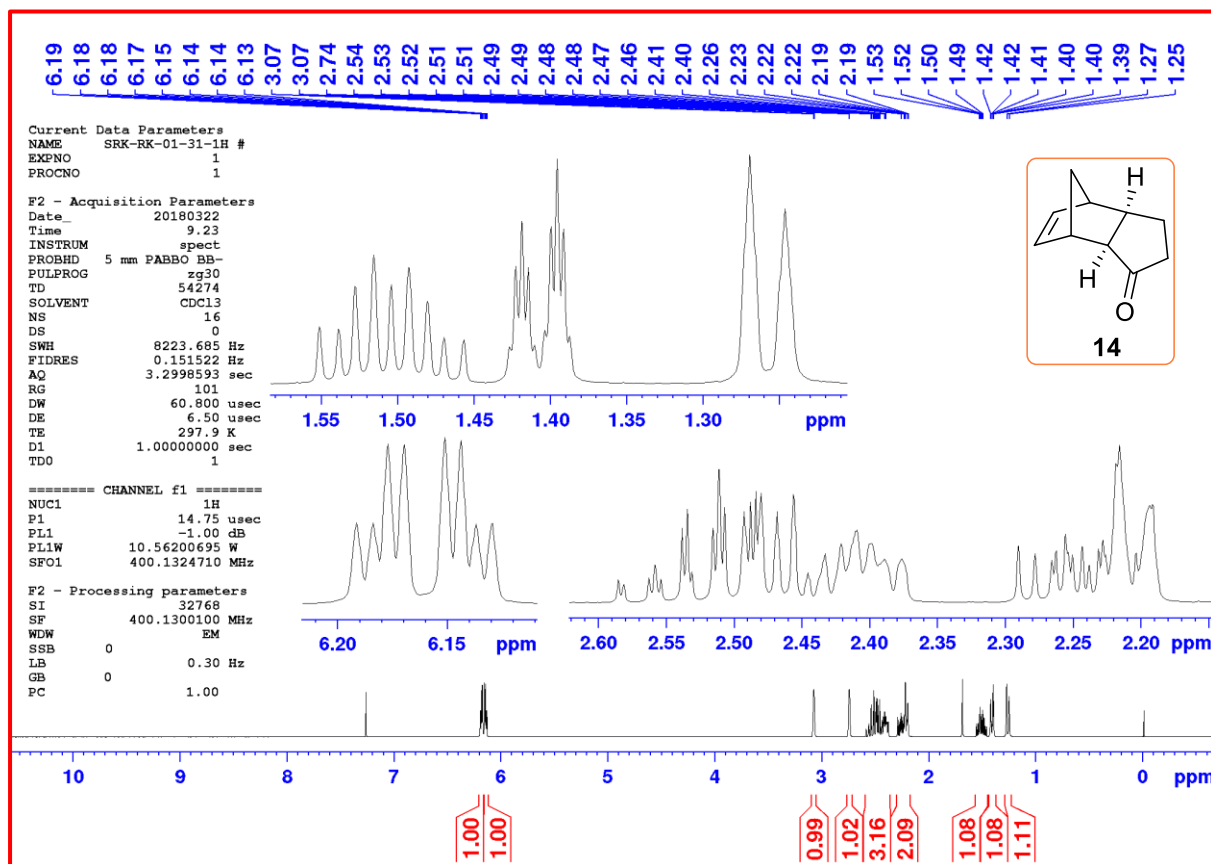
Identification code	SRK-RK-01-82_mo
Empirical formula	C <sub>21</sub> H <sub>23</sub> N
Formula weight	289.40
Temperature	150 K
Crystal system	Monoclinic
Space group	P2 <sub>1</sub> /c
Unit cell dimensions	a = 14.3351 (8) Å     α = 90° b = 5.0684 (3) Å     β = 98.988° (5) c = 22.3710 (14) Å     γ = 90°
Volume	1605.43 (17) Å <sup>3</sup>
Z	4
Density (calculated)	1.197 g/cm <sup>3</sup>
Absorption coefficient (μ)	0.069 mm <sup>-1</sup>
Absorption correction	Multi-scan
Max. and Min. transmission	0.998-0.993
F (000)	624.0
Crystal size	0.302 x 0.087 x 0.035 mm <sup>3</sup>

Index ranges	-17 ≤ h ≤ 17, -6 ≤ k ≤ 5, -26 ≤ l ≤ 26
Theta range for data collection	1.843 to 24.998°
Reflections collected	0.0857 (2063)
Diffraction radiation wavelength	0.71073
Independent reflections	2804 [ R <sub>(int)</sub> = 0.0778]
Completeness to θ = 24.998°	100%
Refinement method	Full-matrix least-squares on F <sup>2</sup>
Data/restraints/parameters	2804/0/199
Goodness-of-fit on F <sup>2</sup>	1.075
Final R indices [I ≥ 2σ(I)]	R1 = 0.0857, wR2 = 0.2324
R indices (all data)	R1 = 0.1153, wR2 = 0.2734
Largest diff. peak and hole	1.48/-1.11 e Å <sup>-3</sup>

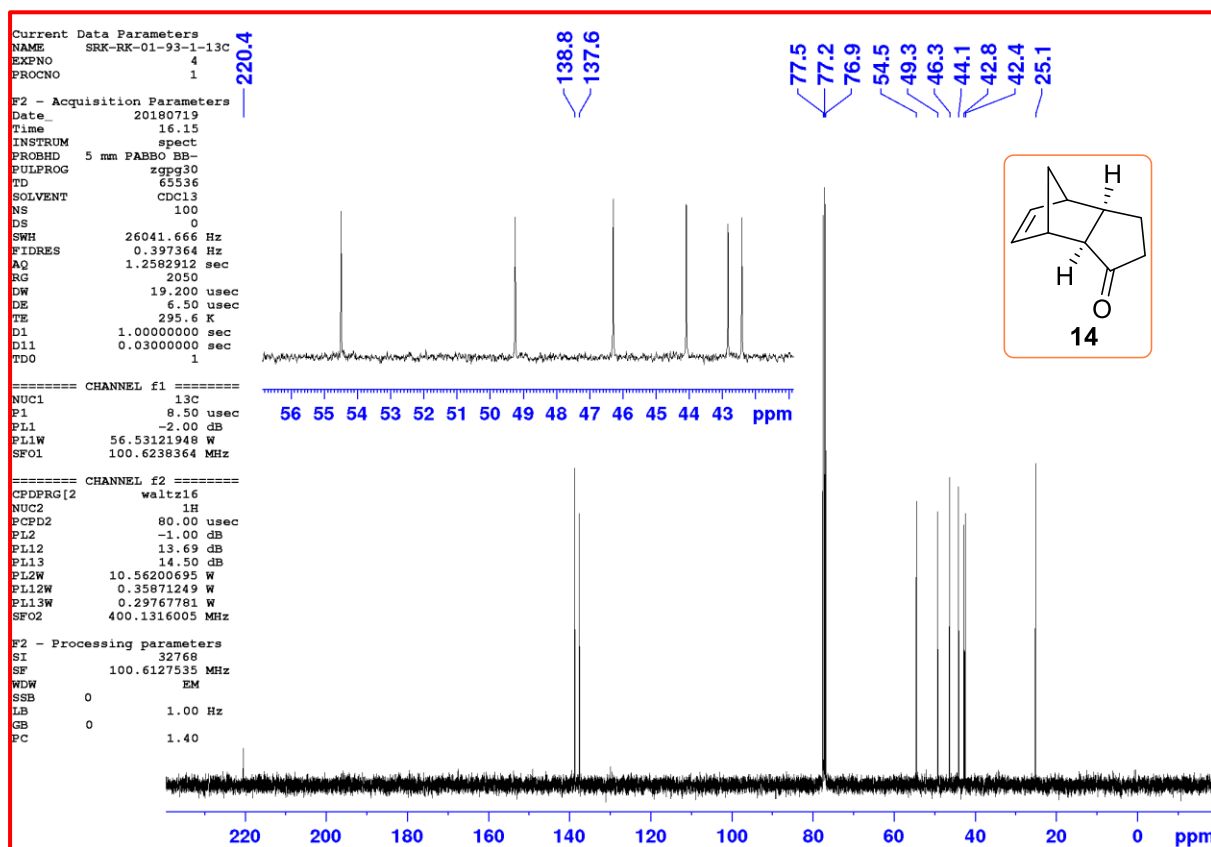
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## 2. Plots of $^1\text{H}$ NMR, $^{13}\text{C}$ NMR and DEPT-135 spectral data all new compounds

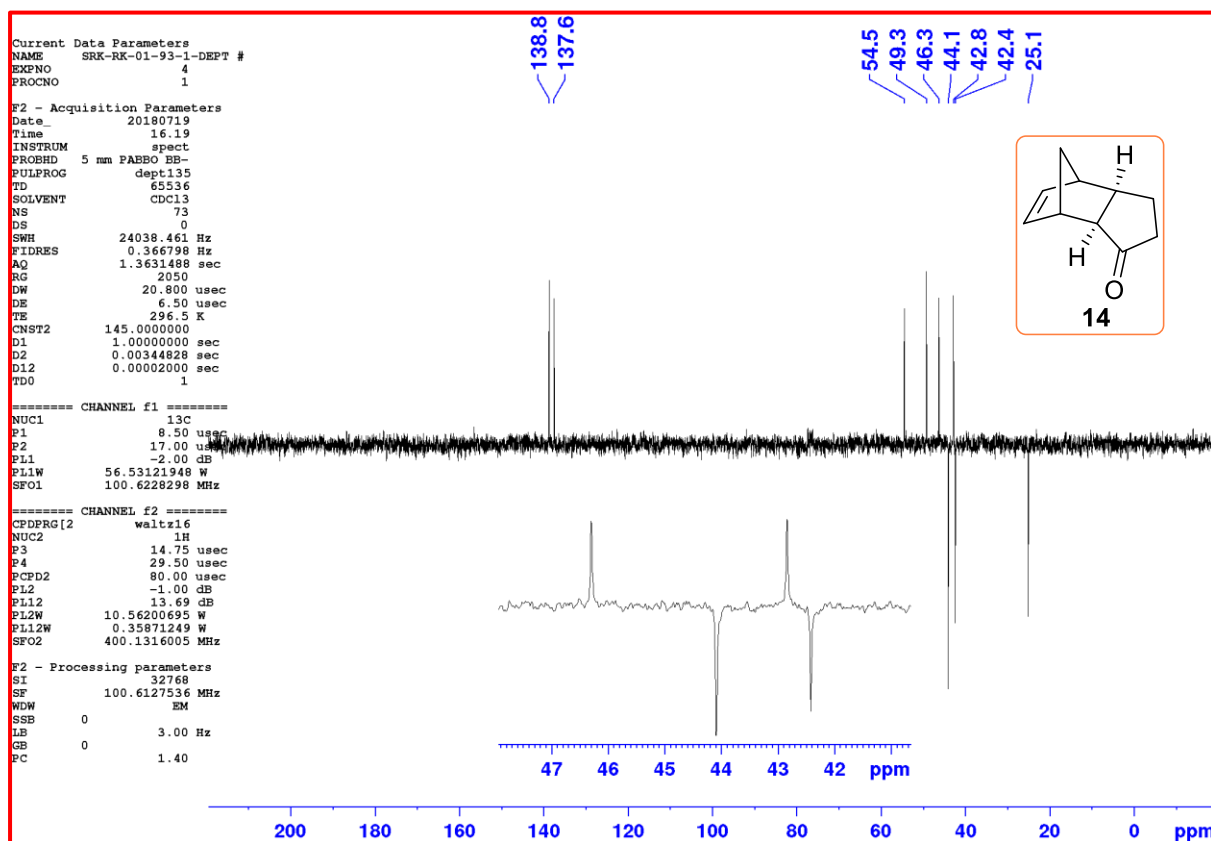
$^1\text{H}$  NMR (400 MHz,  $\text{CDCl}_3$ ) of **14**



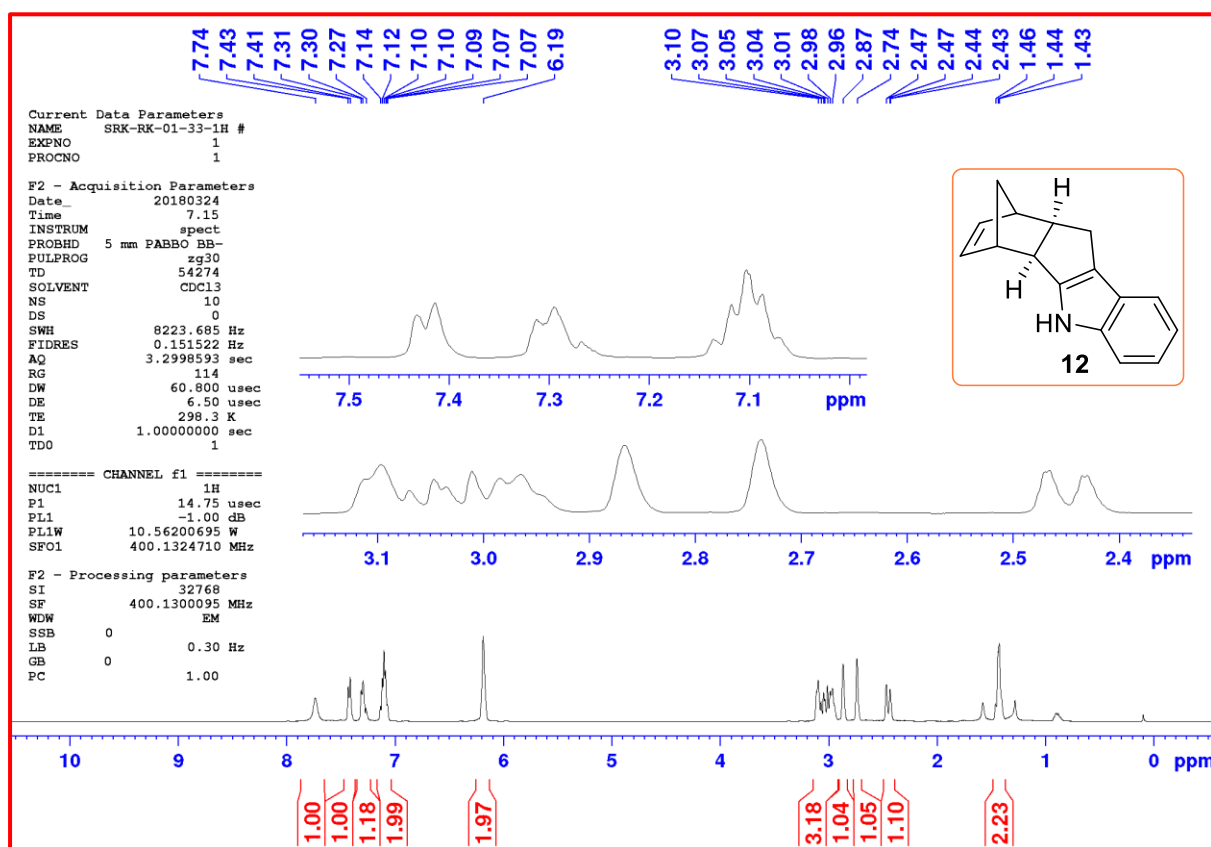
$^{13}\text{C}$  NMR (100 MHz,  $\text{CDCl}_3$ ) of **14**



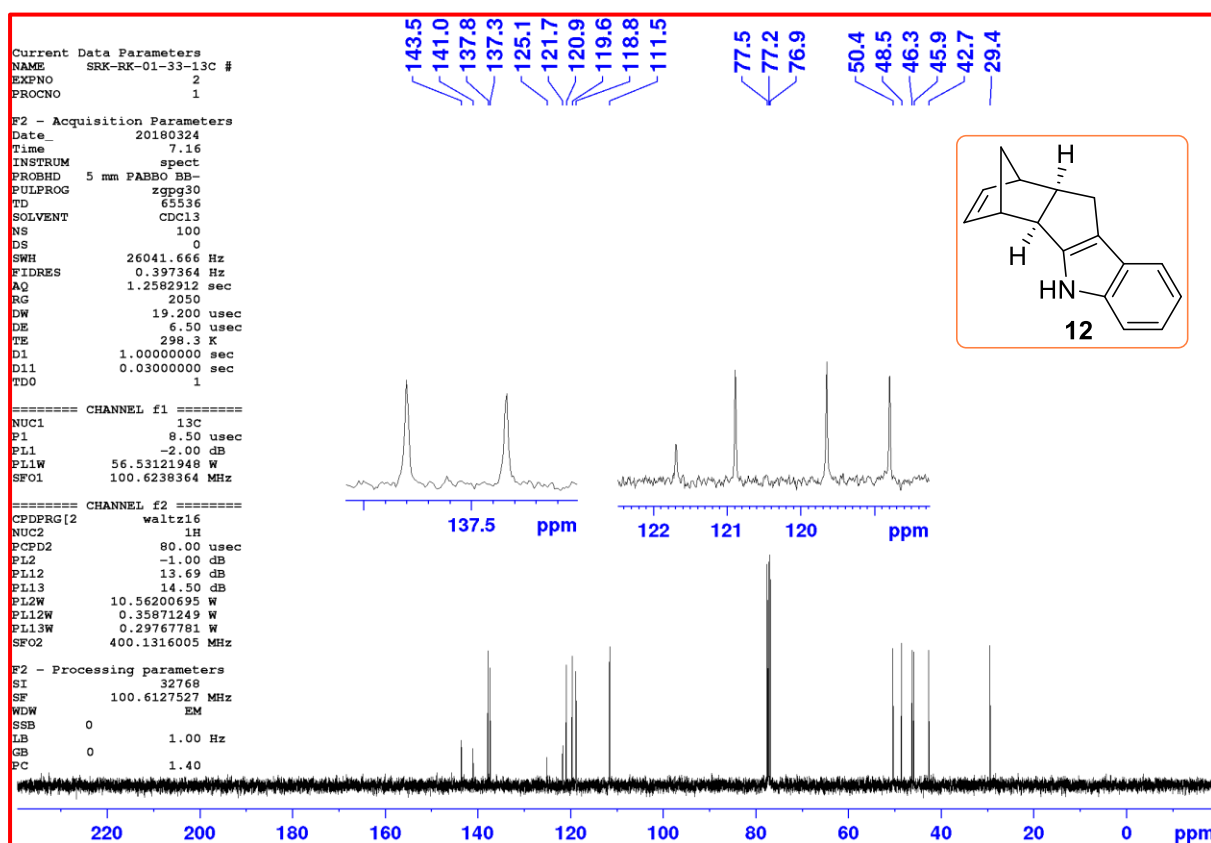
DEPT-135 (100 MHz,  $\text{CDCl}_3$ ) of **14**



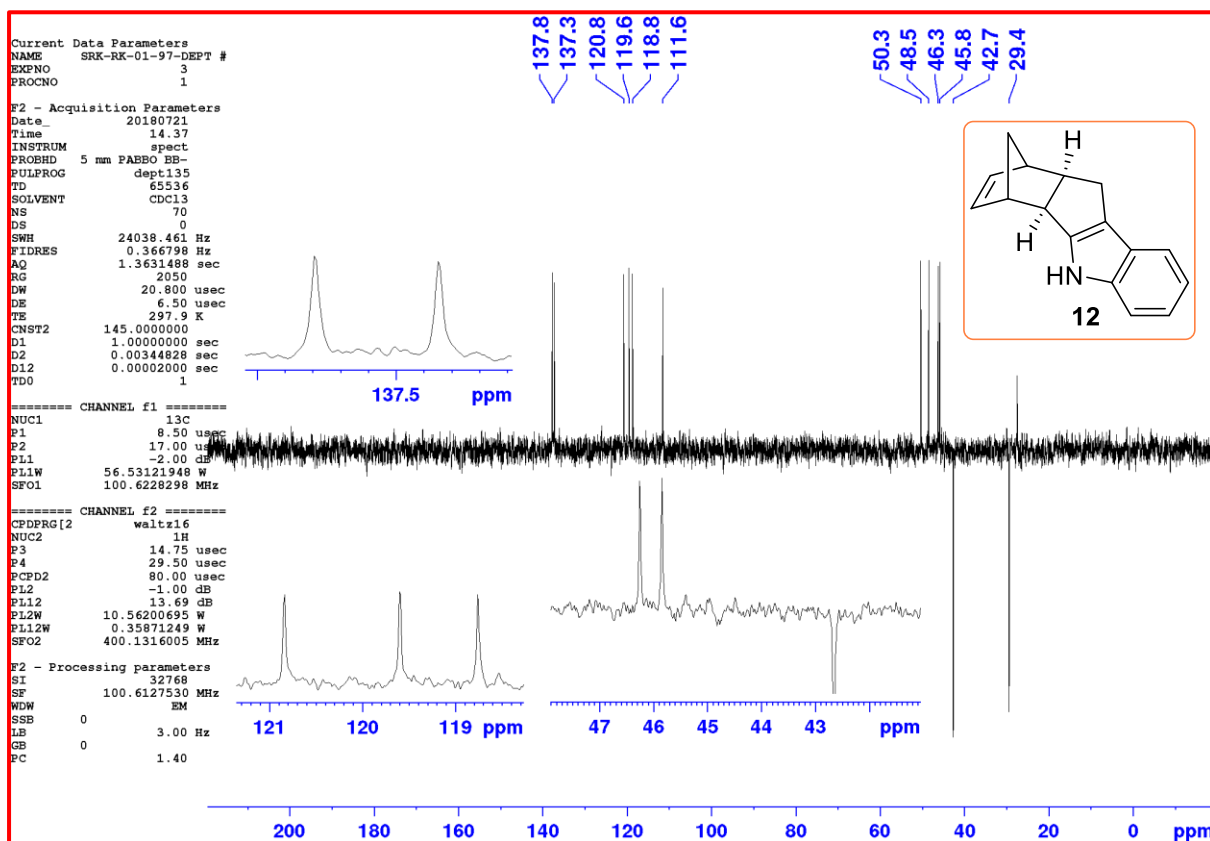
# <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of 12



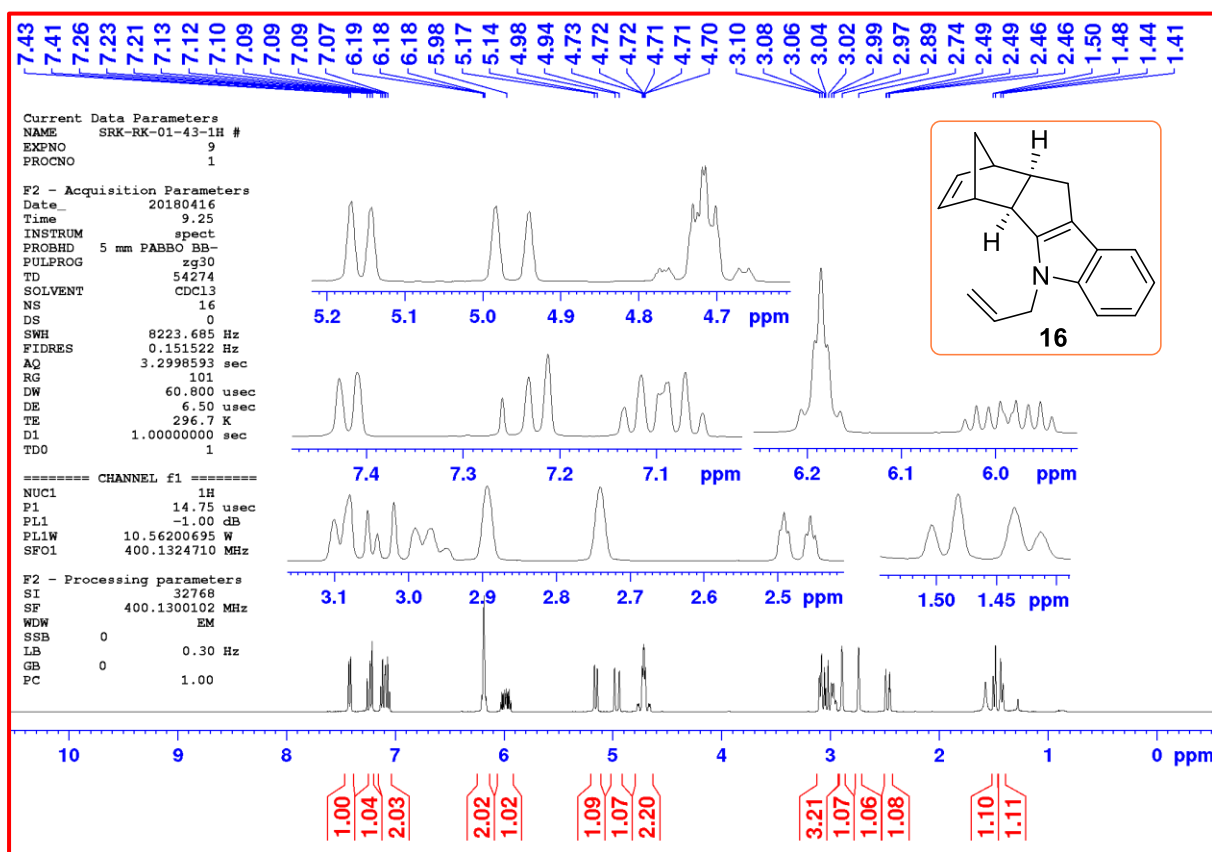
# <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) of 12



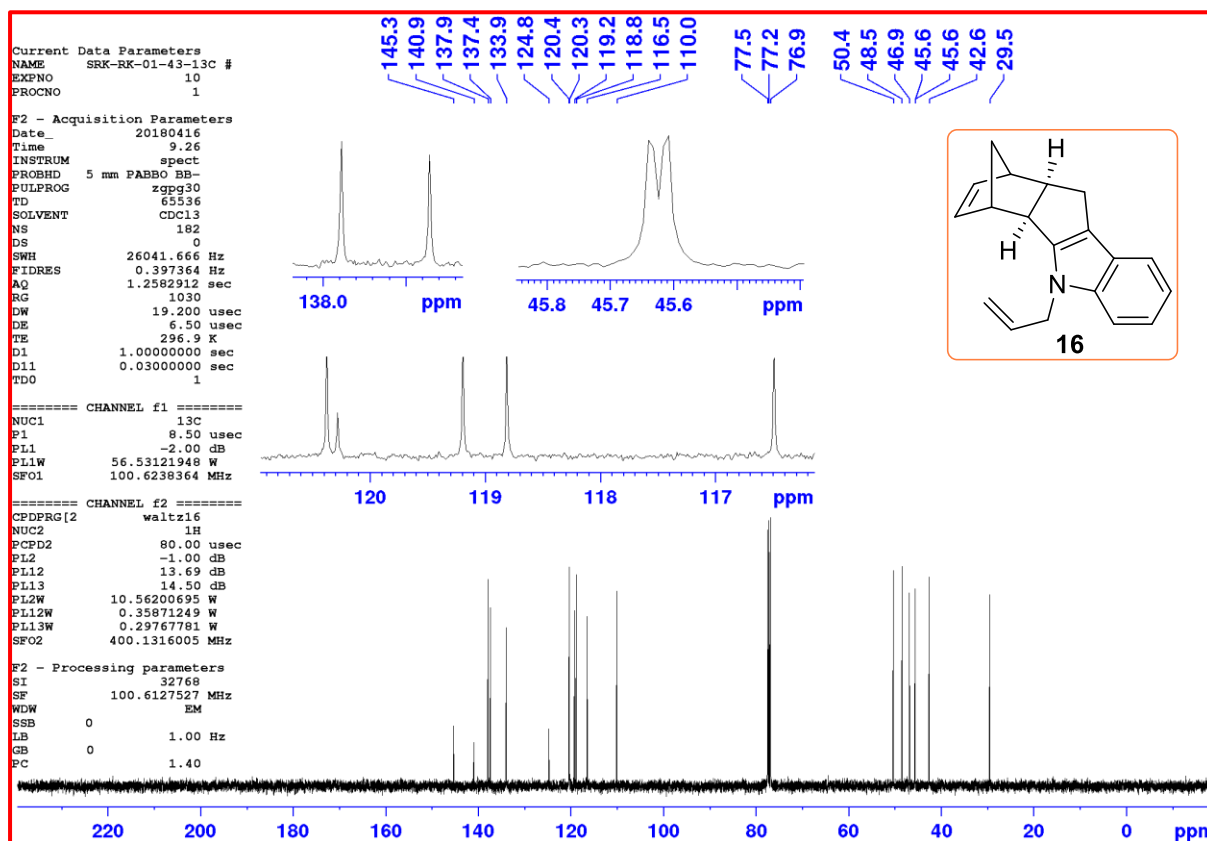
DEPT-135 (100 MHz, CDCl<sub>3</sub>) of 12



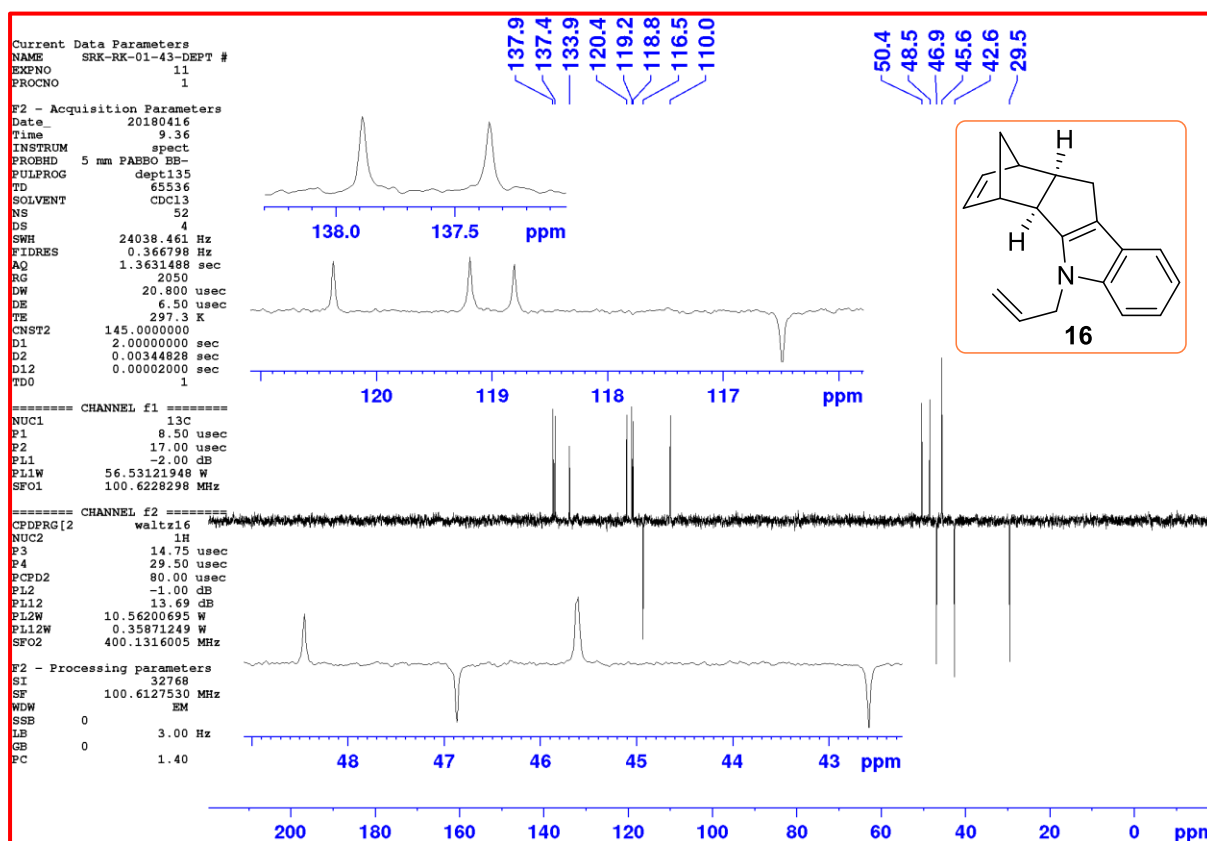
<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of 16



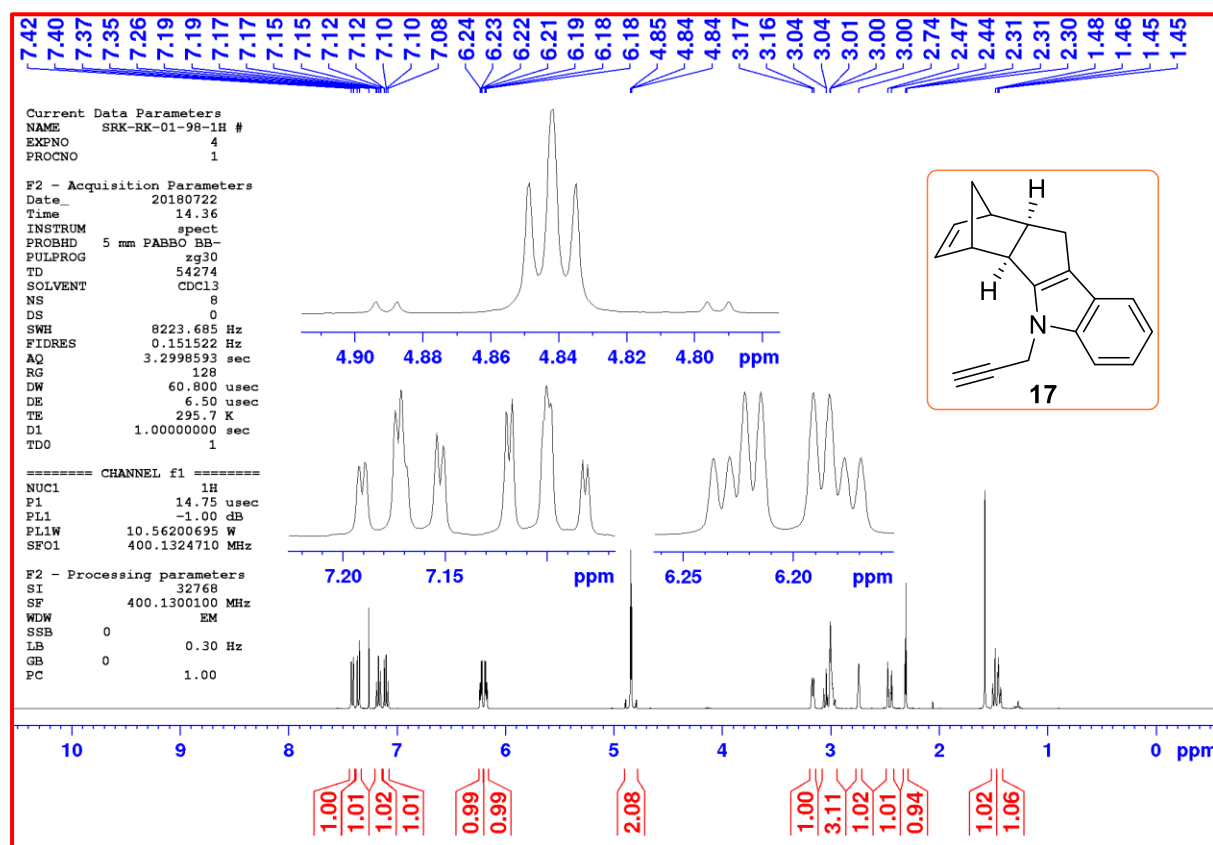
<sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) of **16**



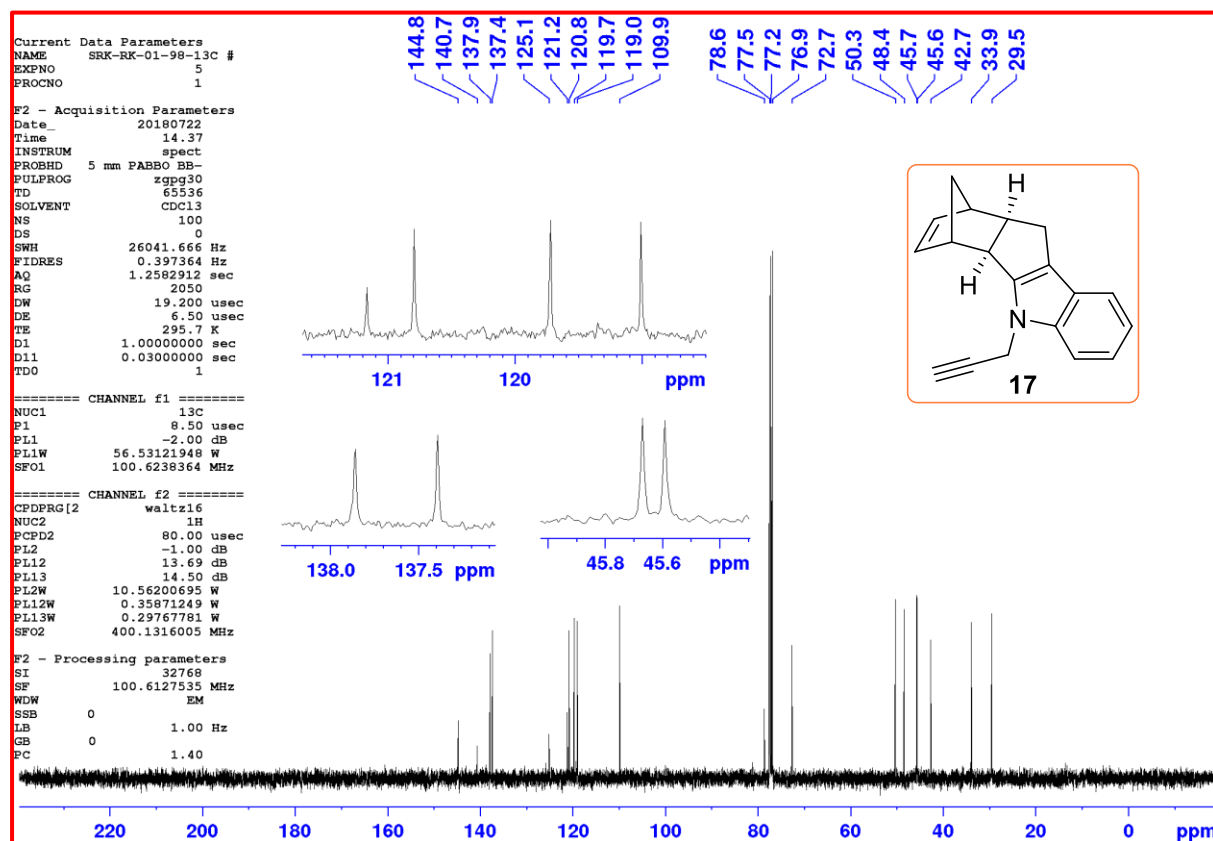
DEPT-135 (100 MHz, CDCl<sub>3</sub>) of **16**



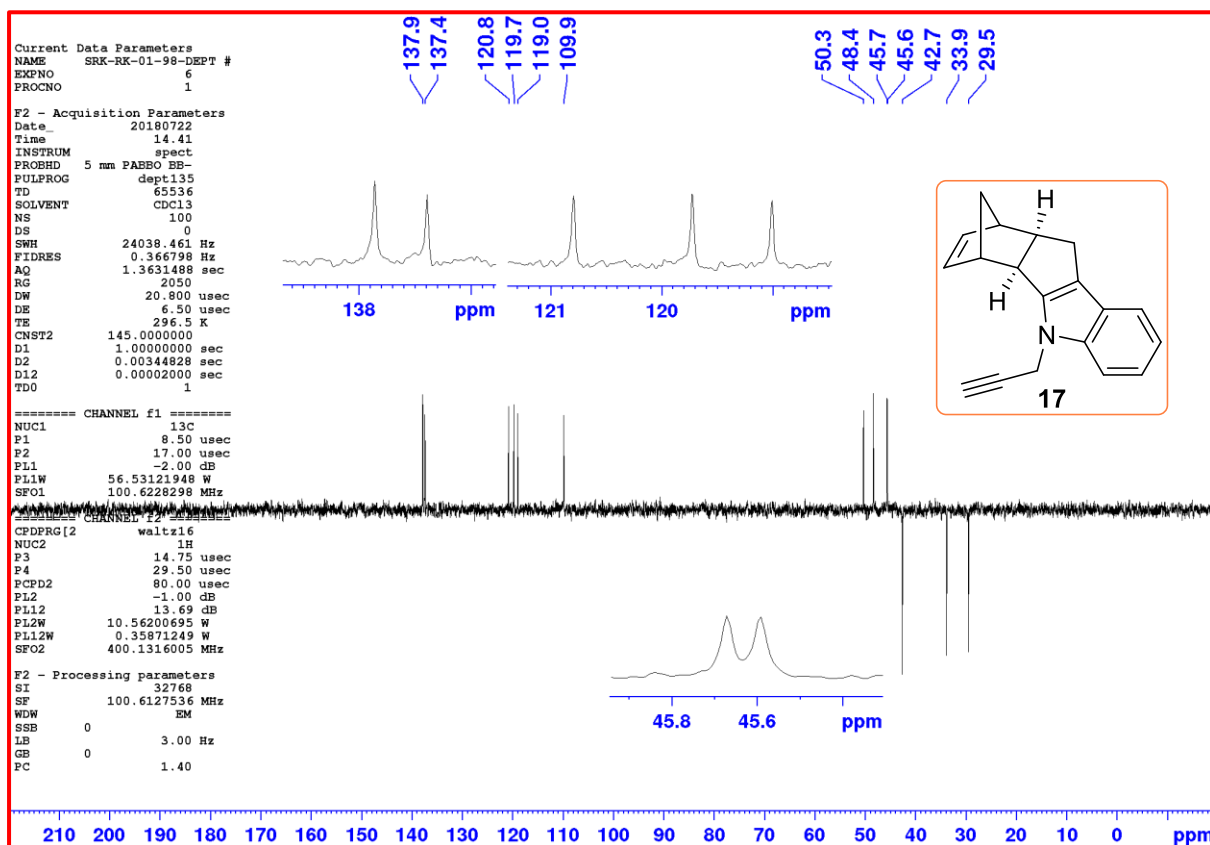
# <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of 17



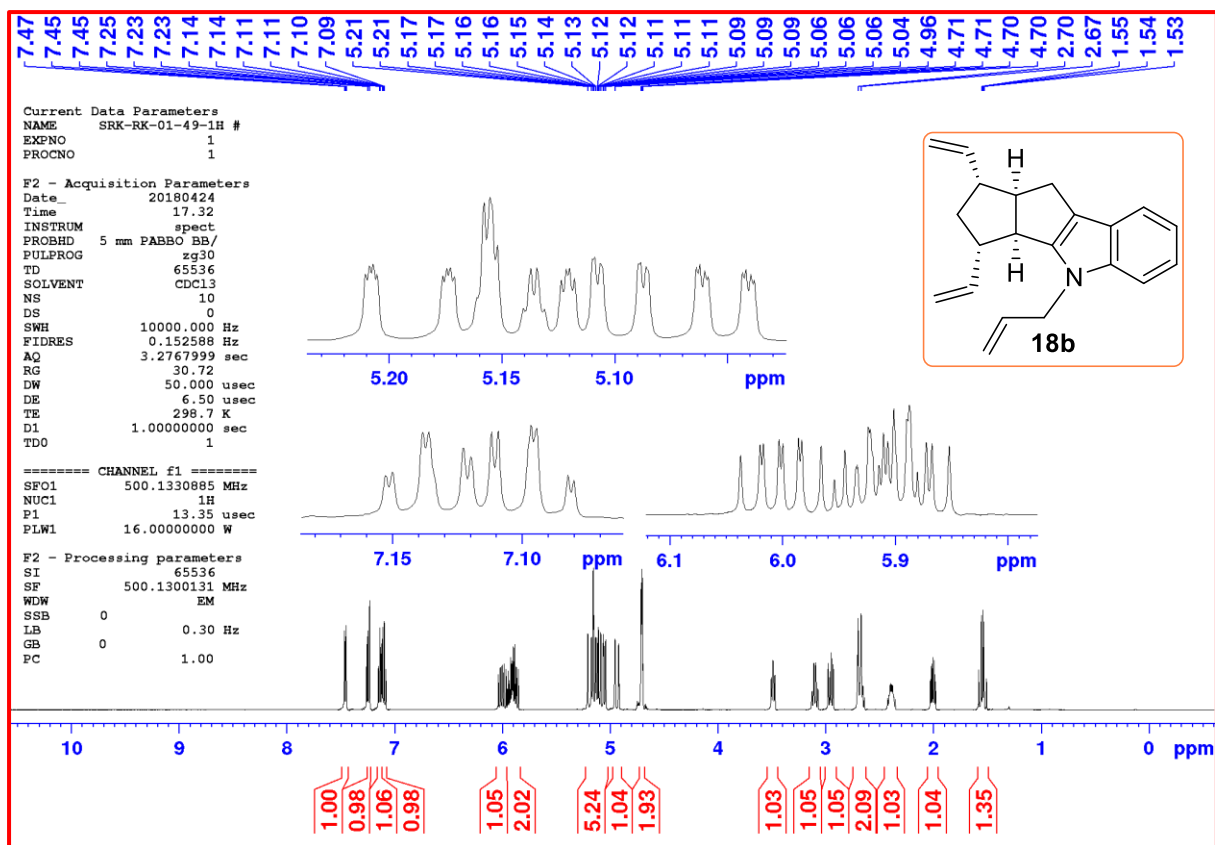
# <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) of 17



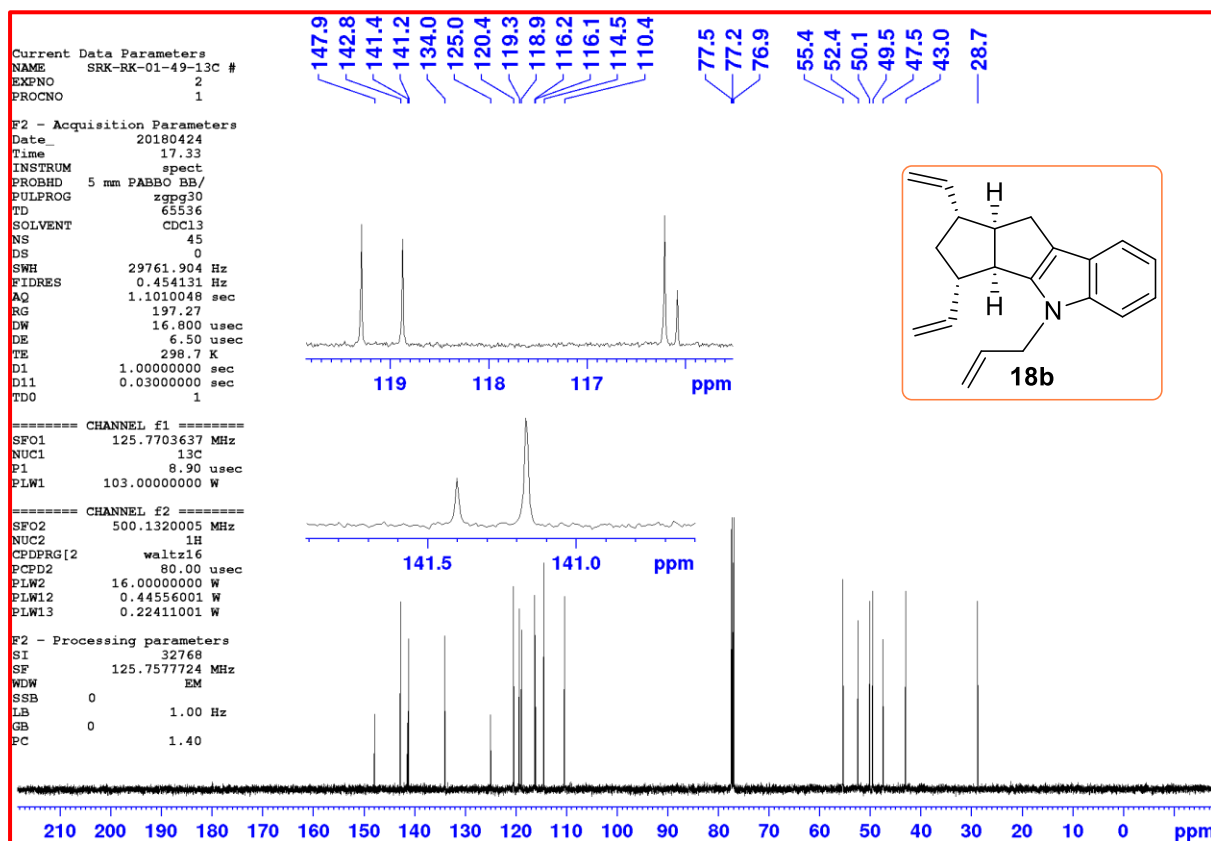
DEPT-135 (100 MHz, CDCl<sub>3</sub>) of **17**



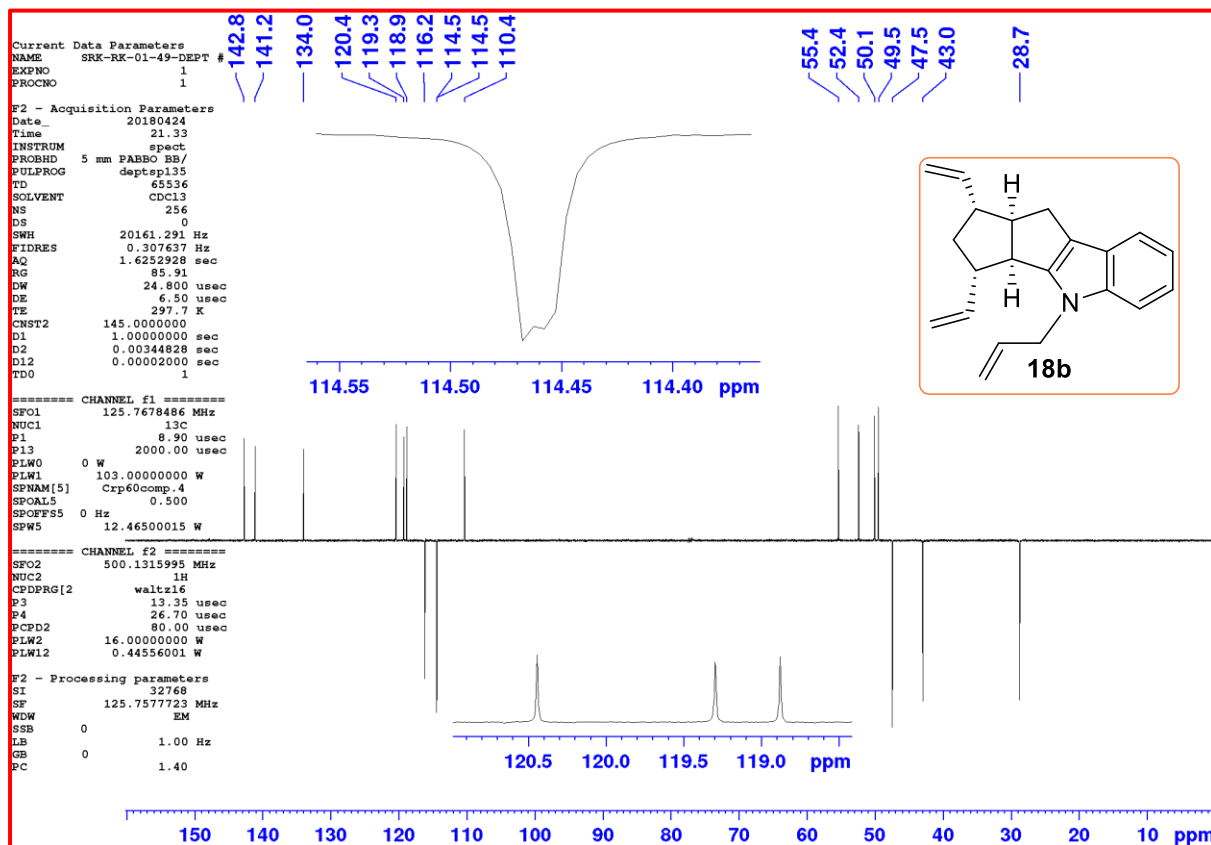
<sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>) of **18b**



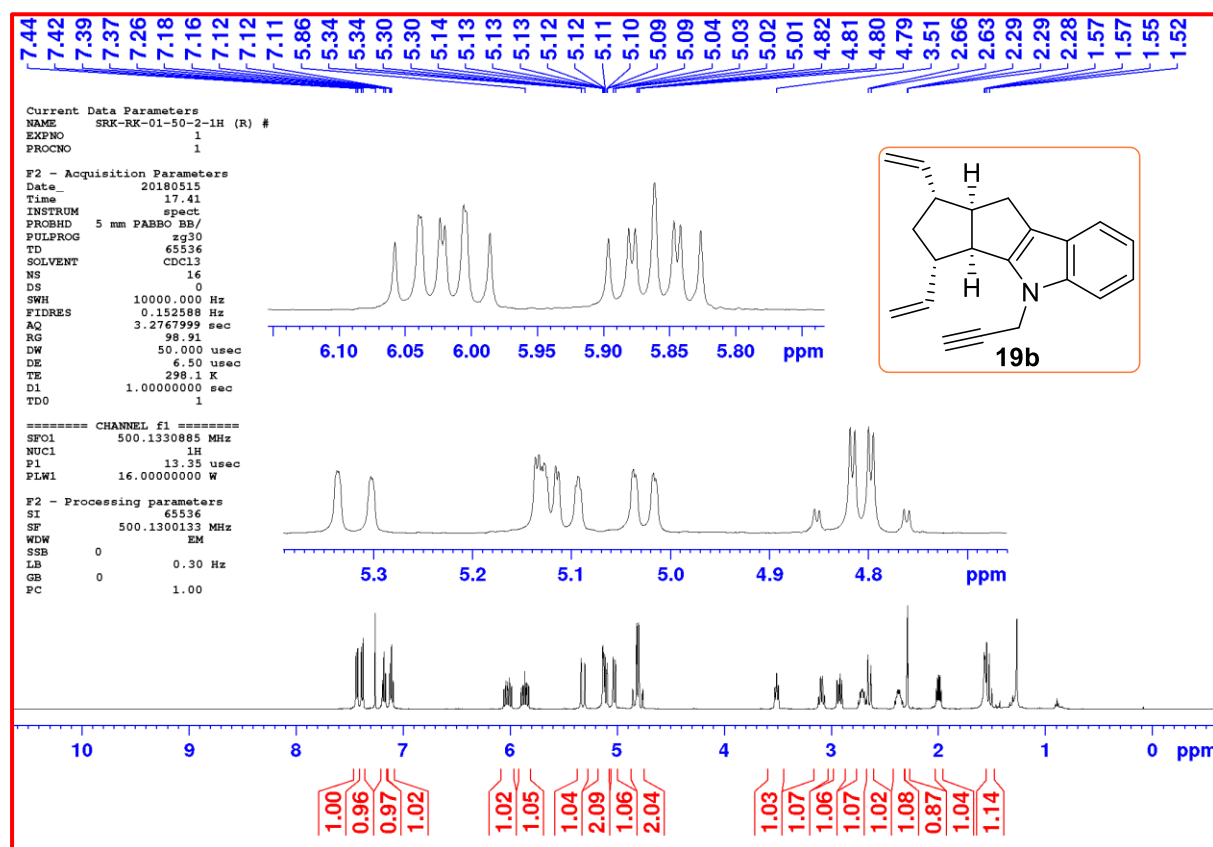
<sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>) of **18b**



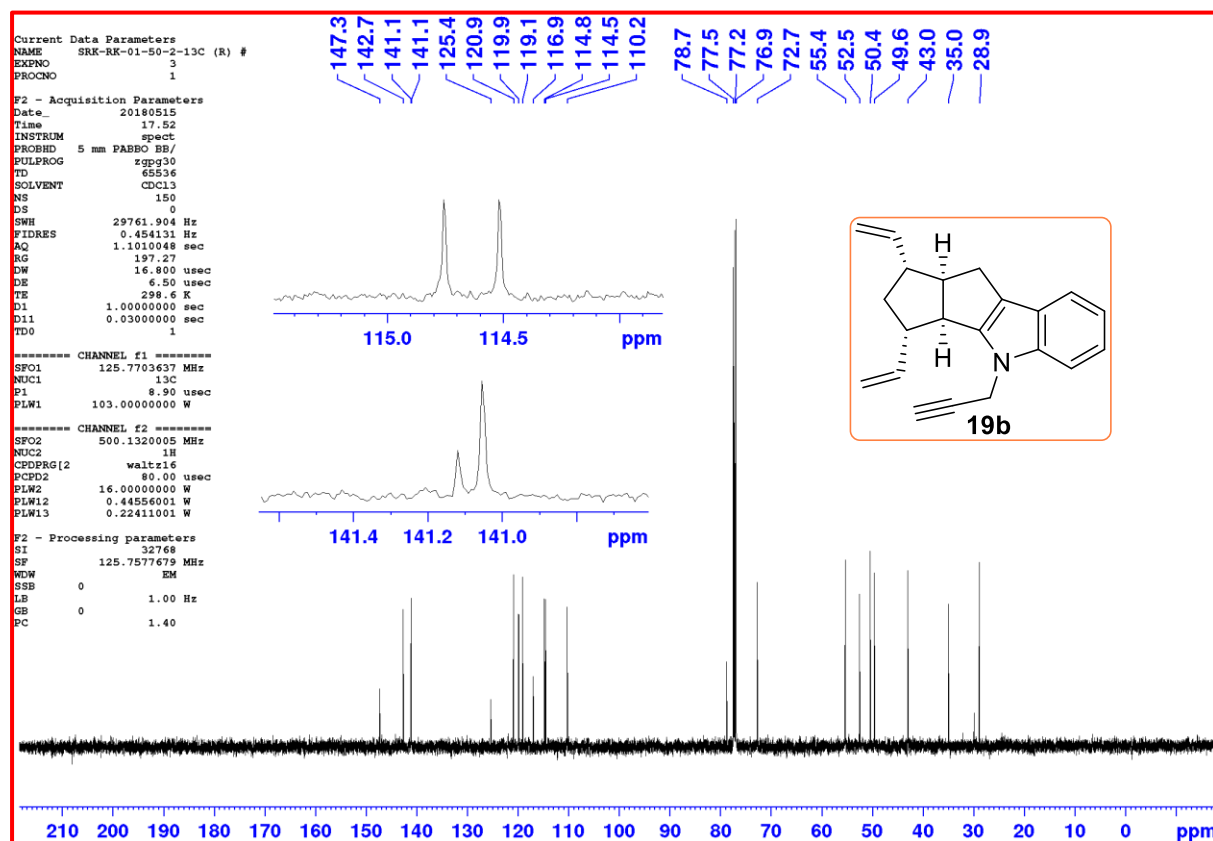
DEPT-135 (125 MHz, CDCl<sub>3</sub>) of **18b**



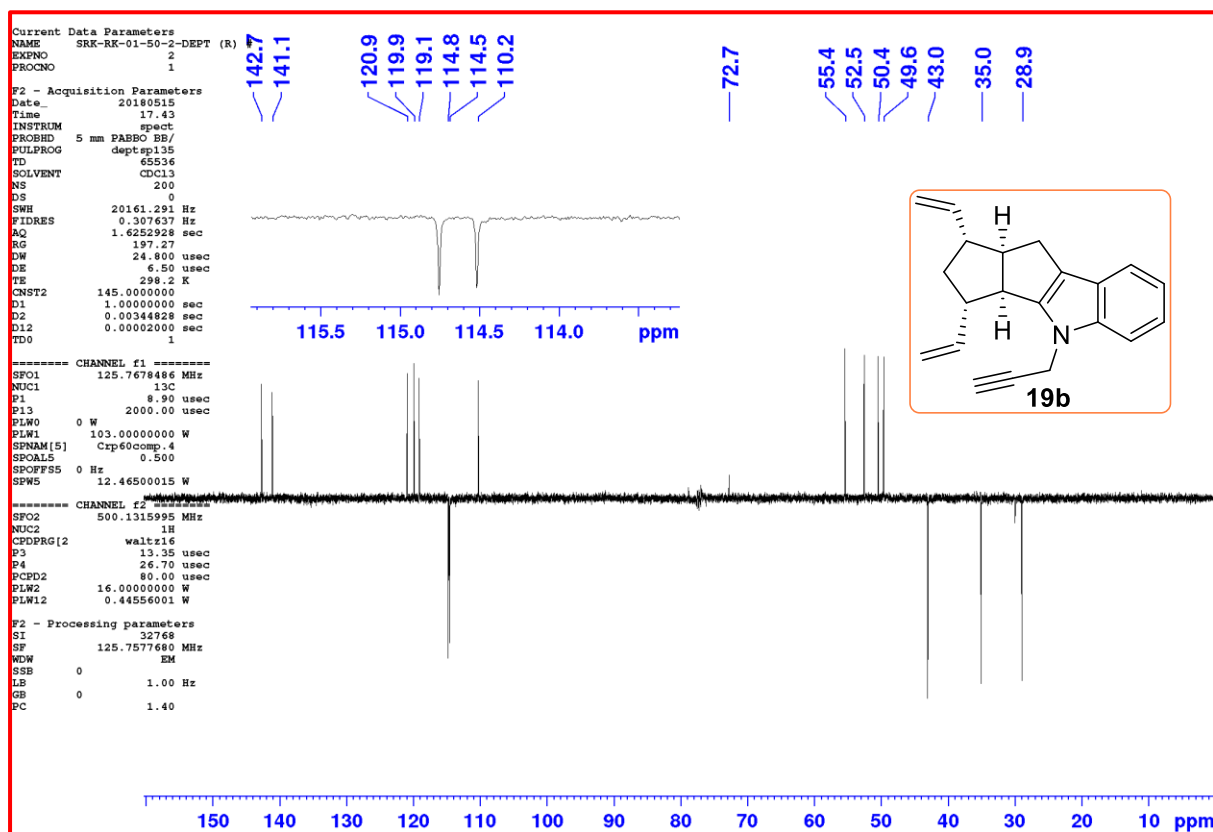
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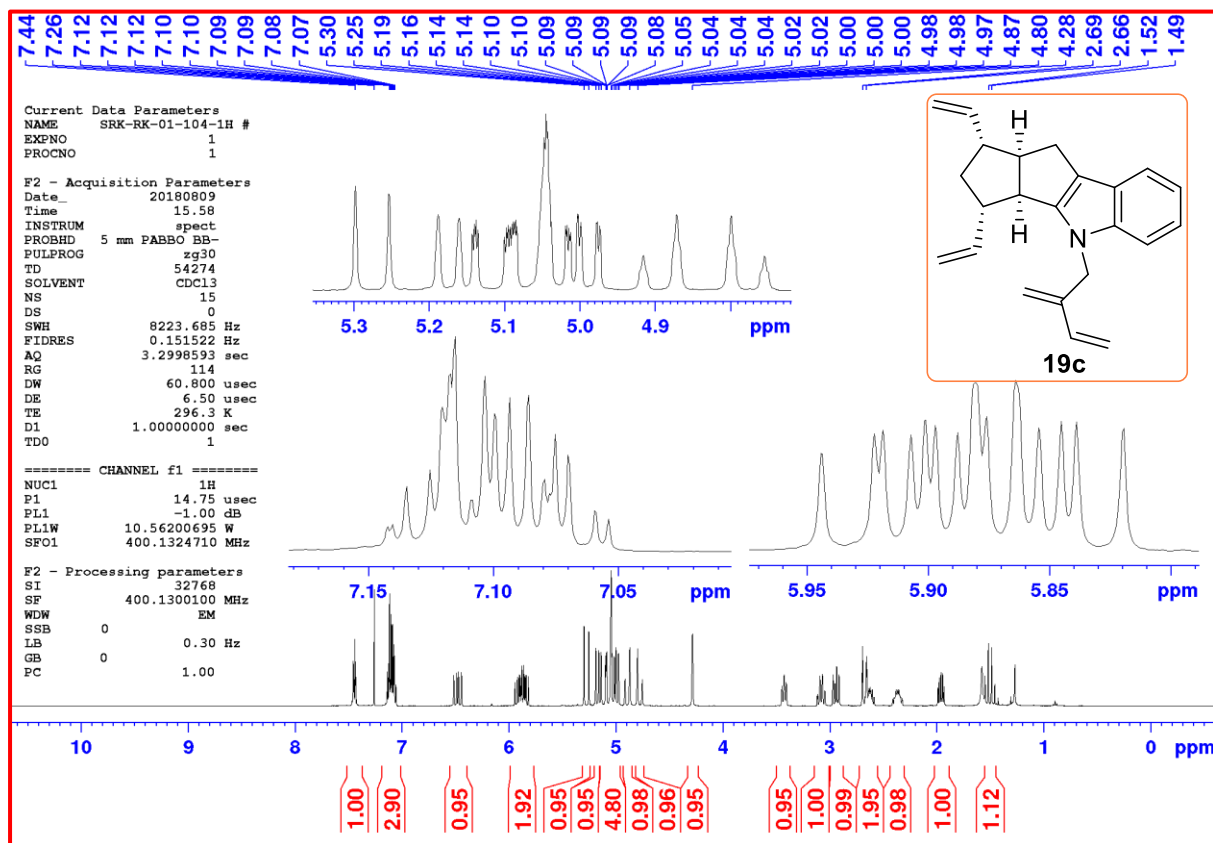
<sup>13</sup>C NMR (125 MHz, CDCl<sub>3</sub>) of **19b**



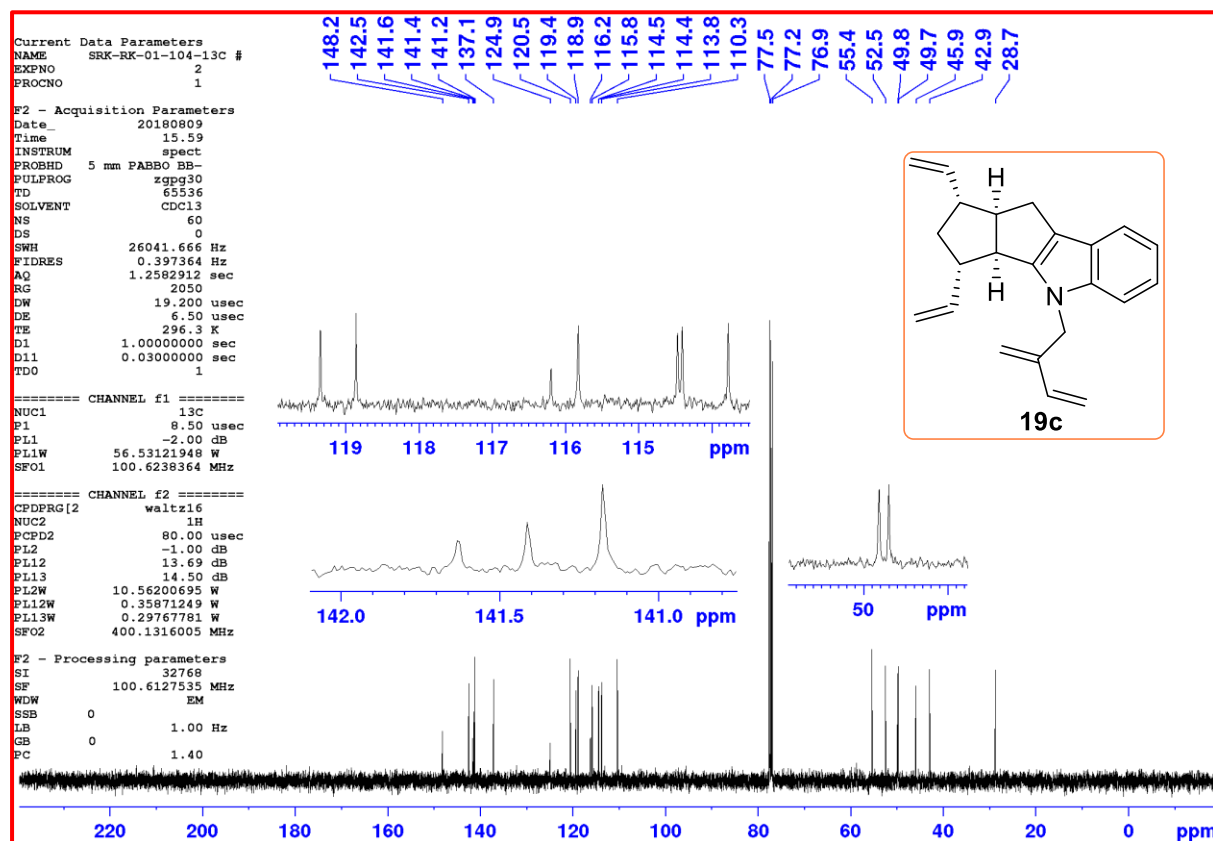
DEPT-135 (125 MHz, CDCl<sub>3</sub>) of **19b**



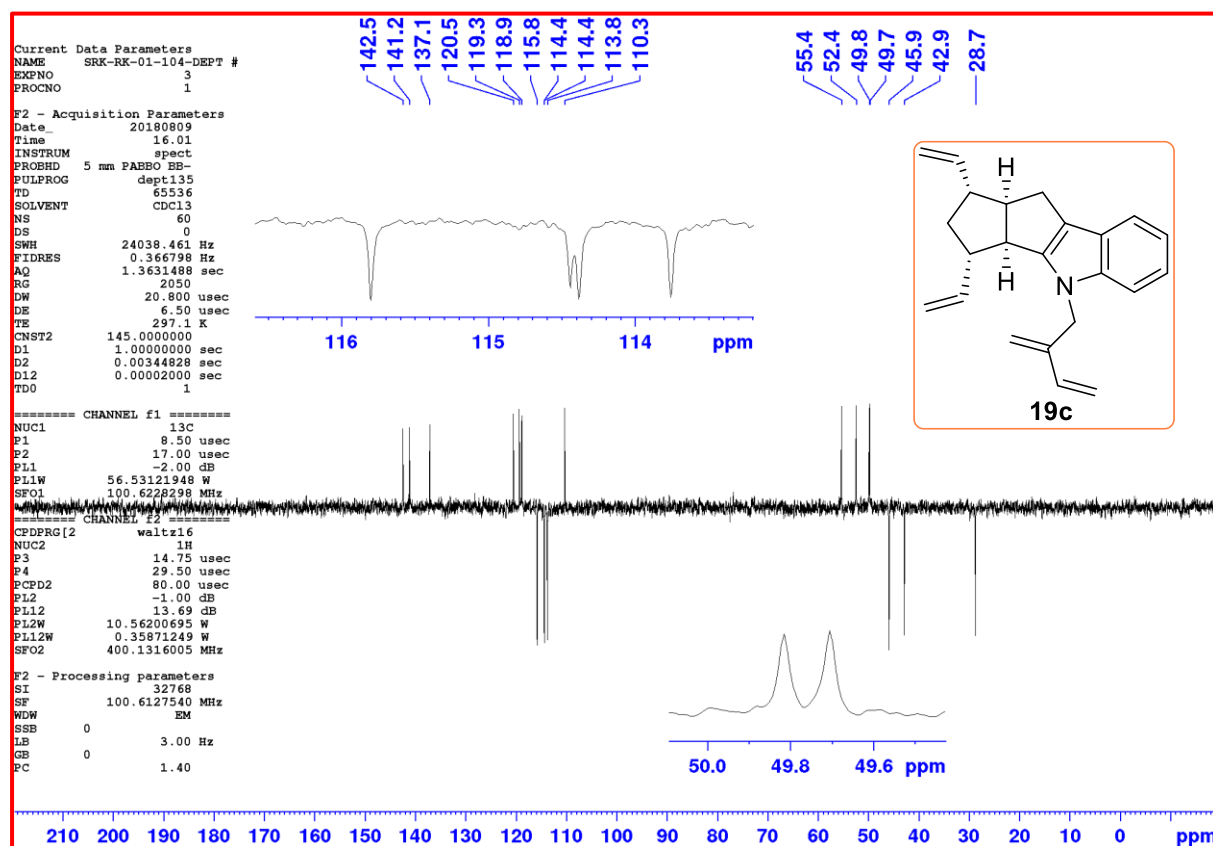
<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of **19c**



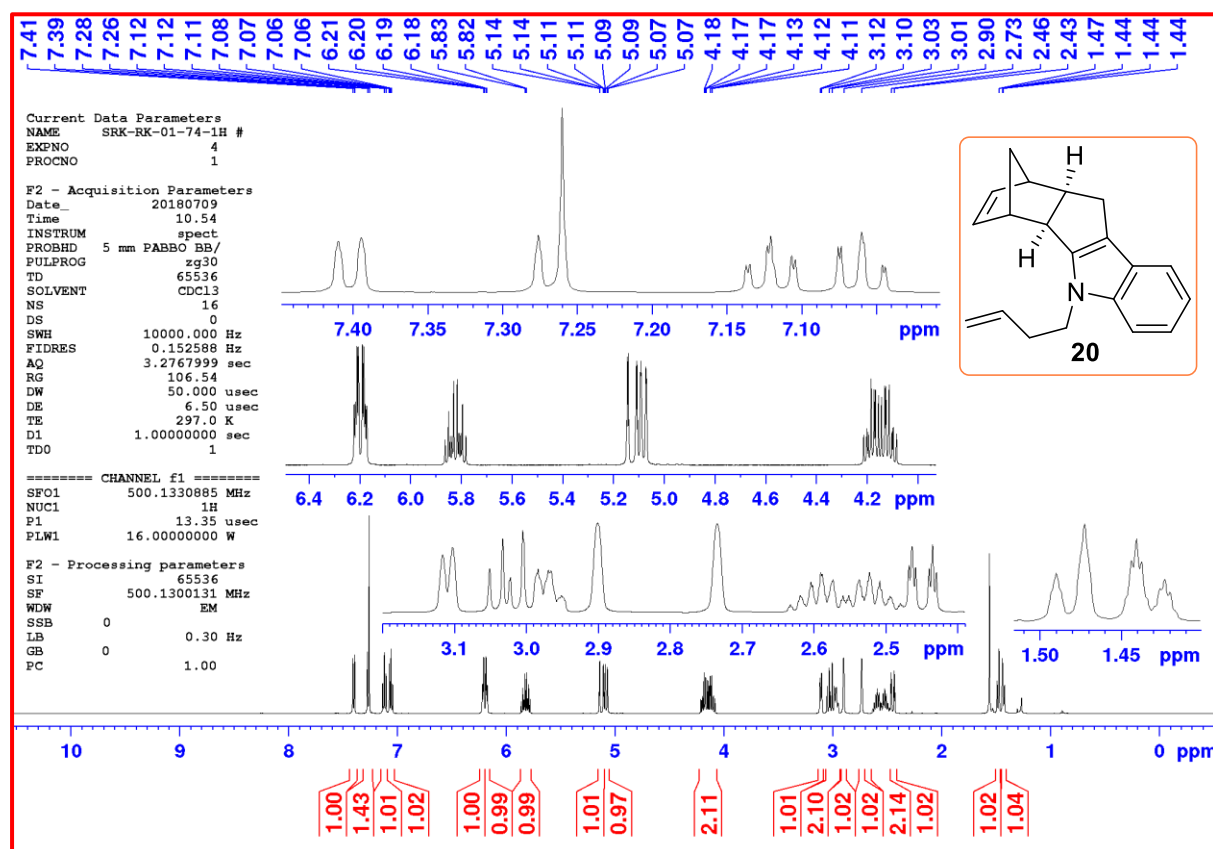
<sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) of **19c**



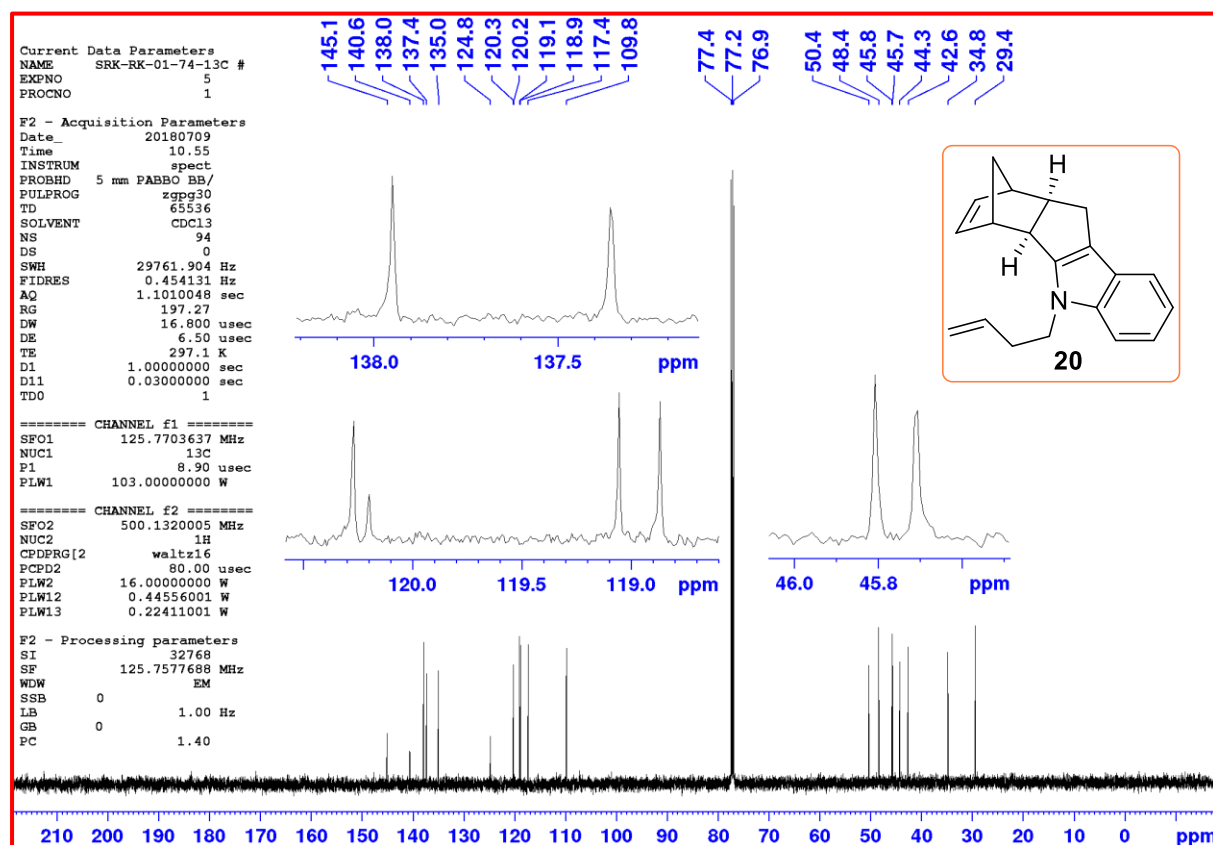
DEPT-135 (100 MHz, CDCl<sub>3</sub>) of **19c**



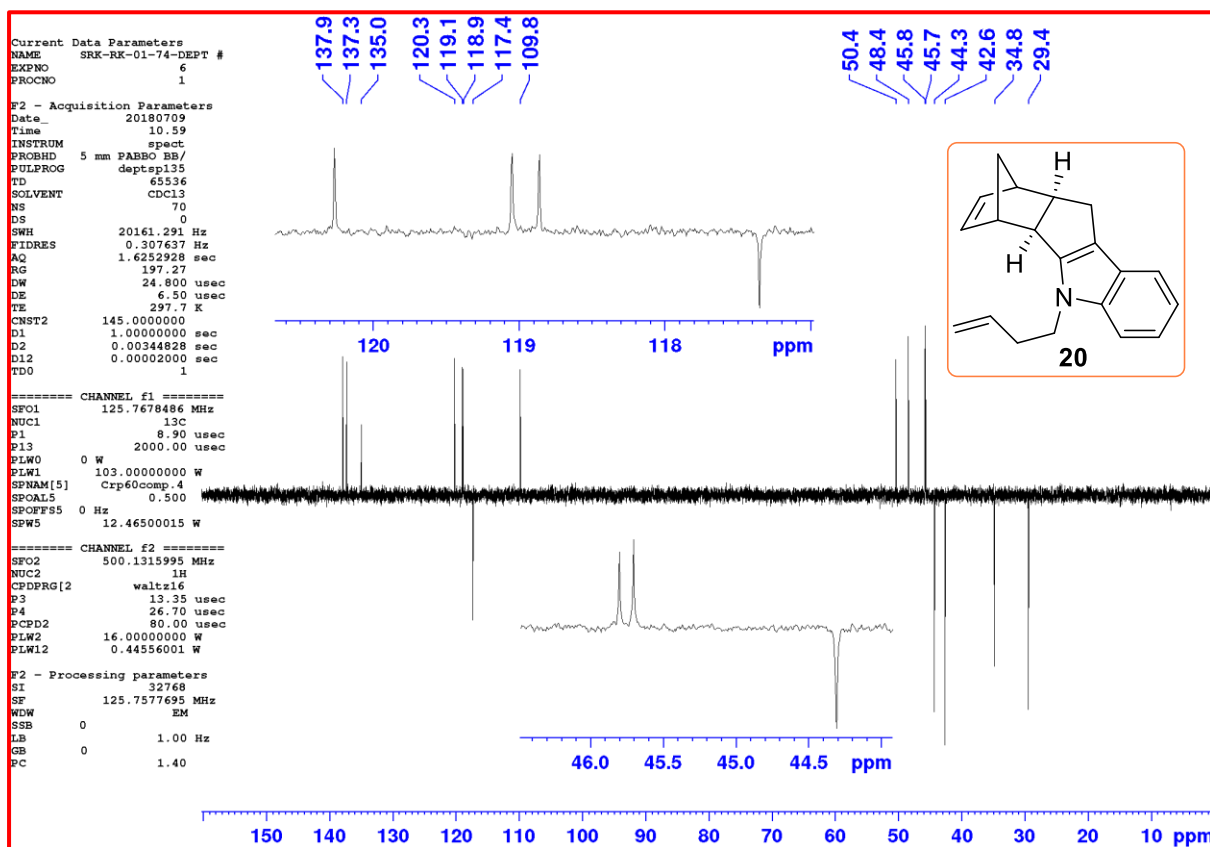
# <sup>1</sup>H NMR (500 MHz, CDCl<sub>3</sub>) of **20**



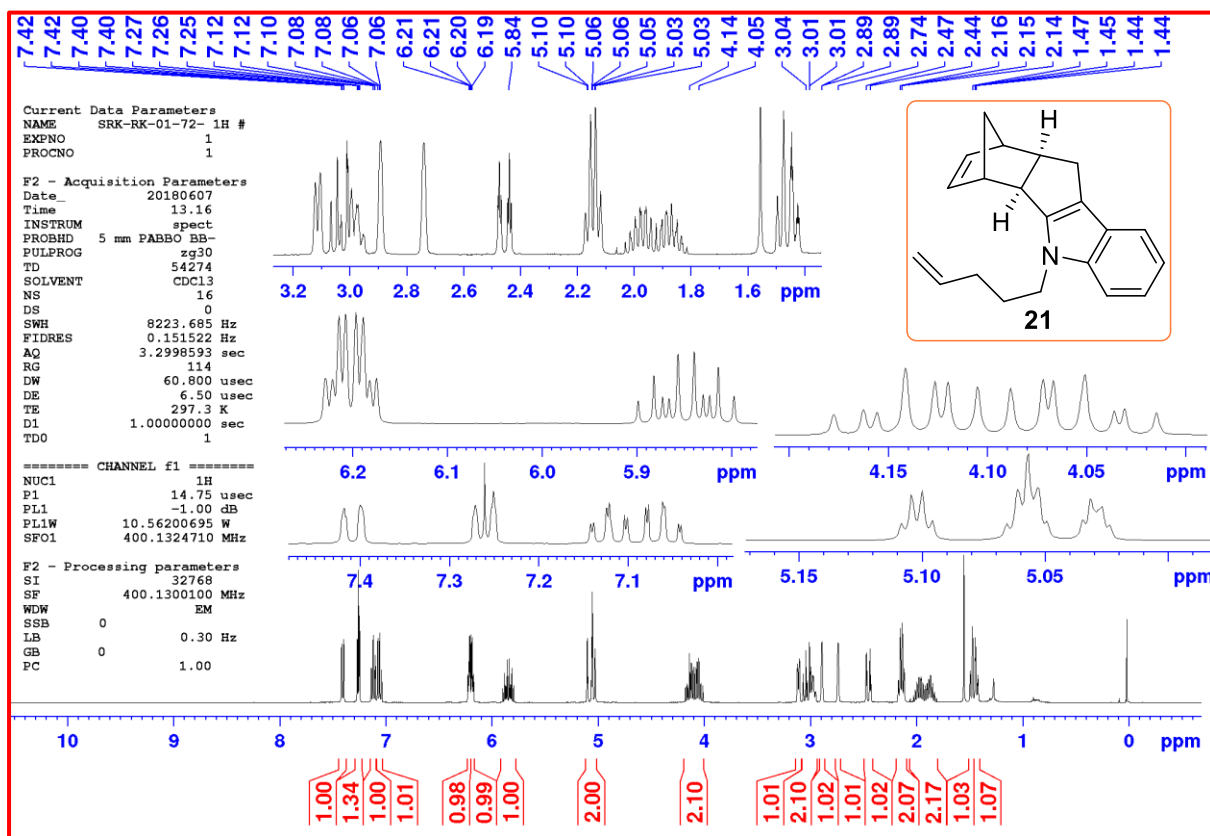
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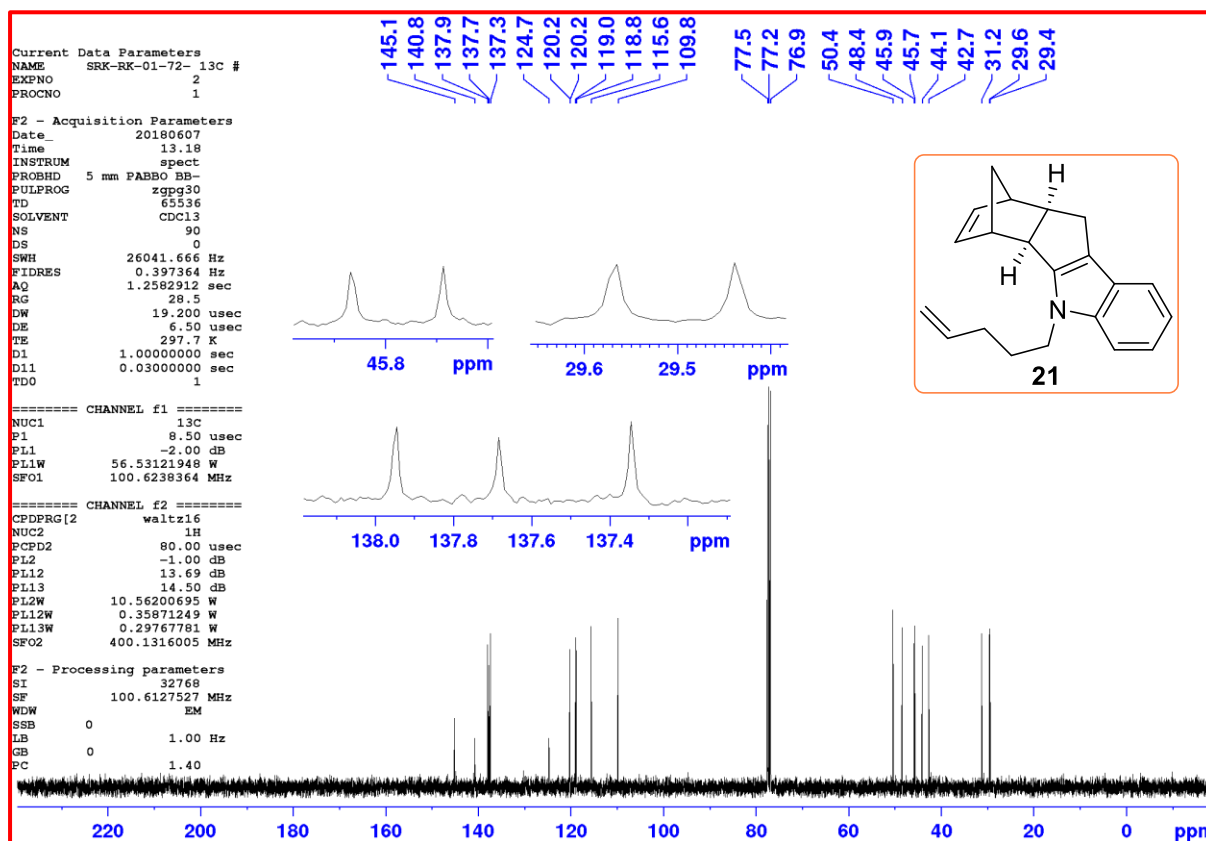
DEPT-135 (125 MHz, CDCl<sub>3</sub>) of **20**



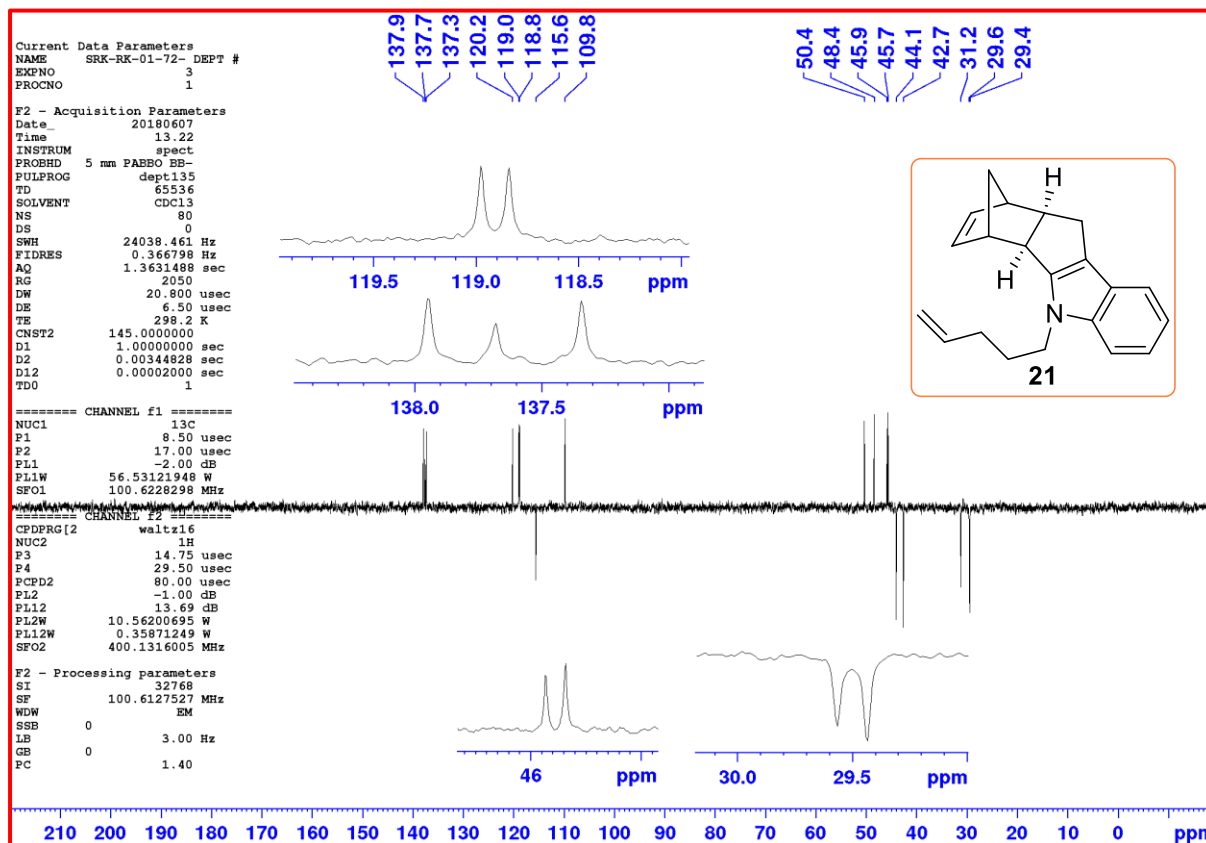
<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of **21**



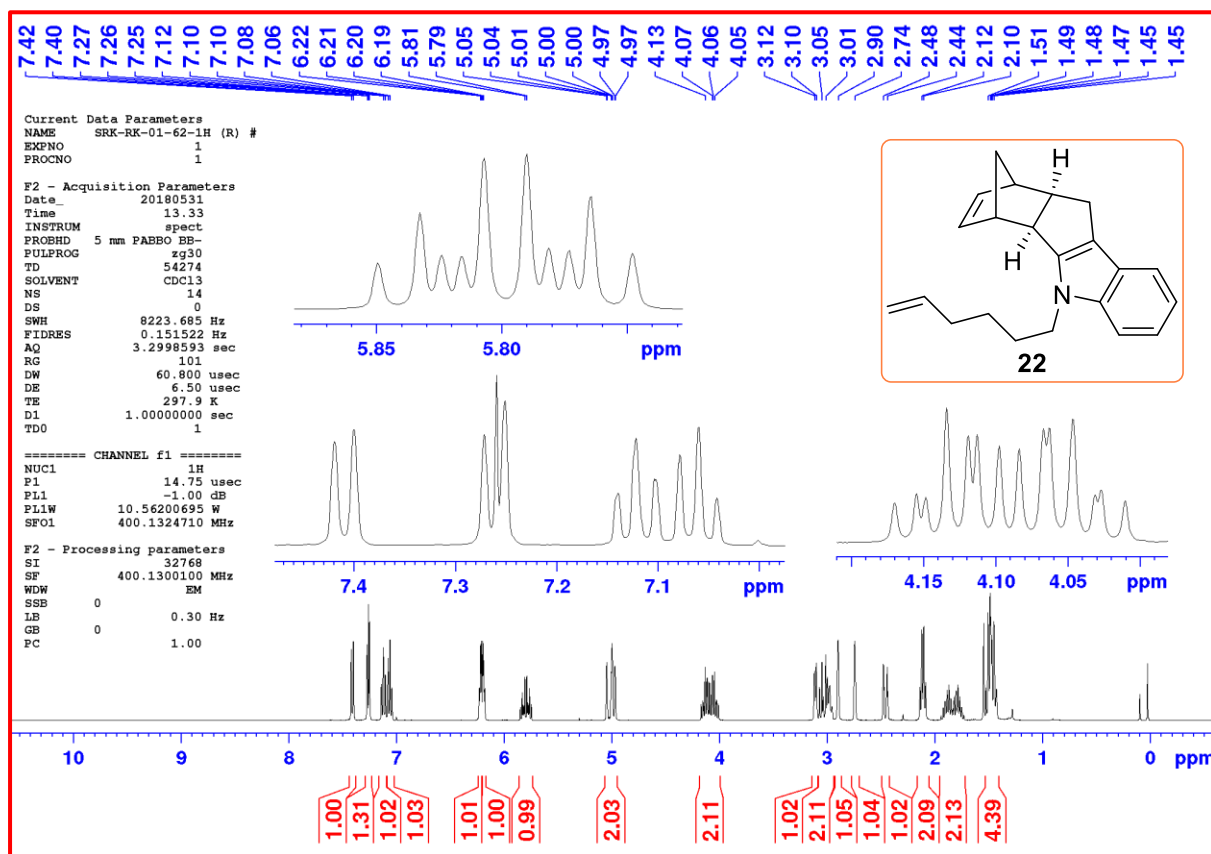
<sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) of **21**



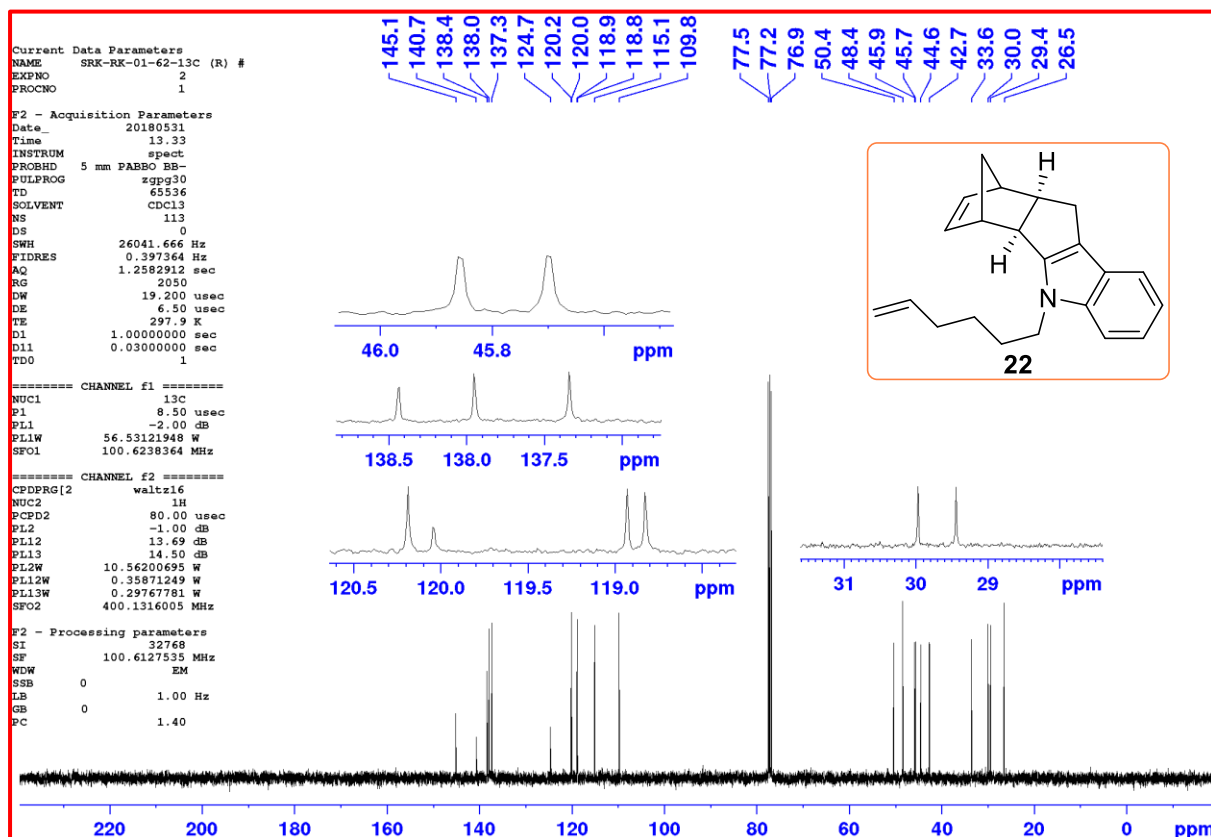
DEPT-135 (100 MHz, CDCl<sub>3</sub>) of **21**



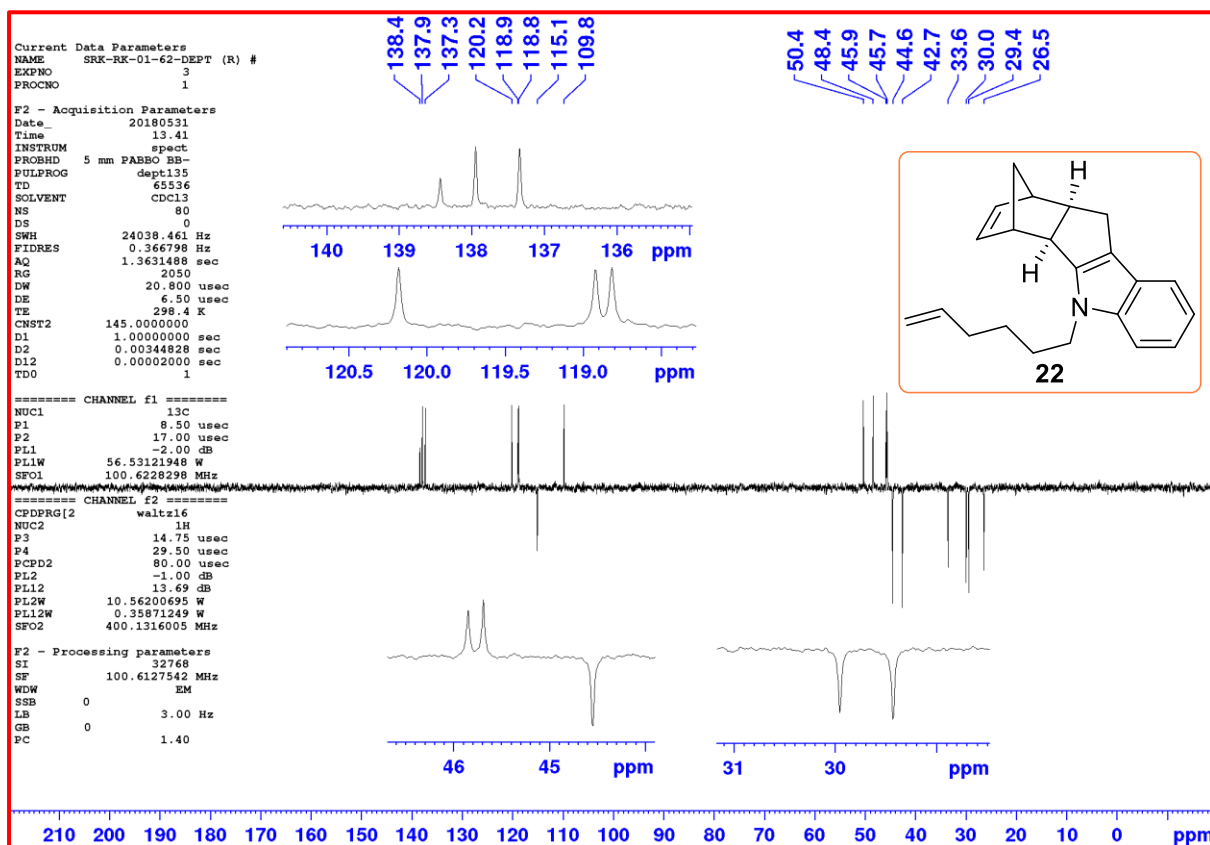
# <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of 22



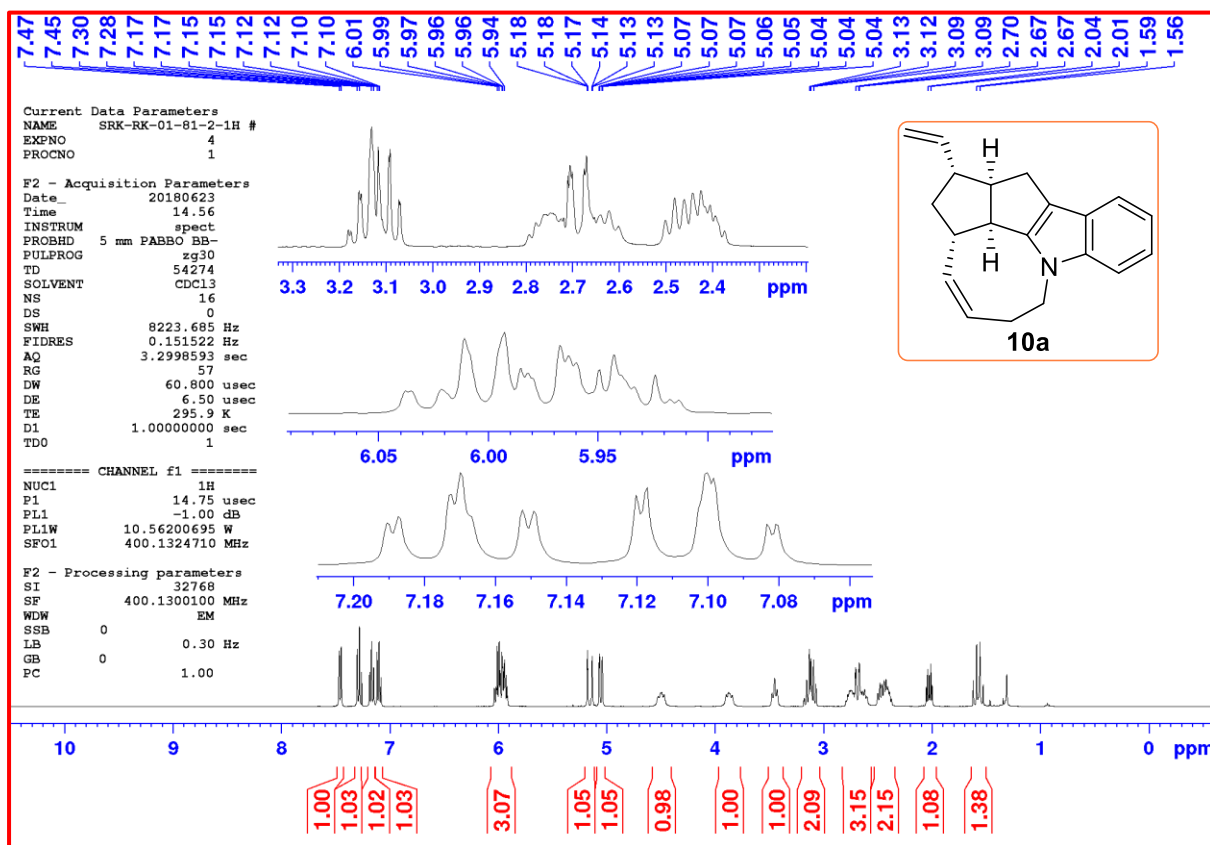
# <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) of 22



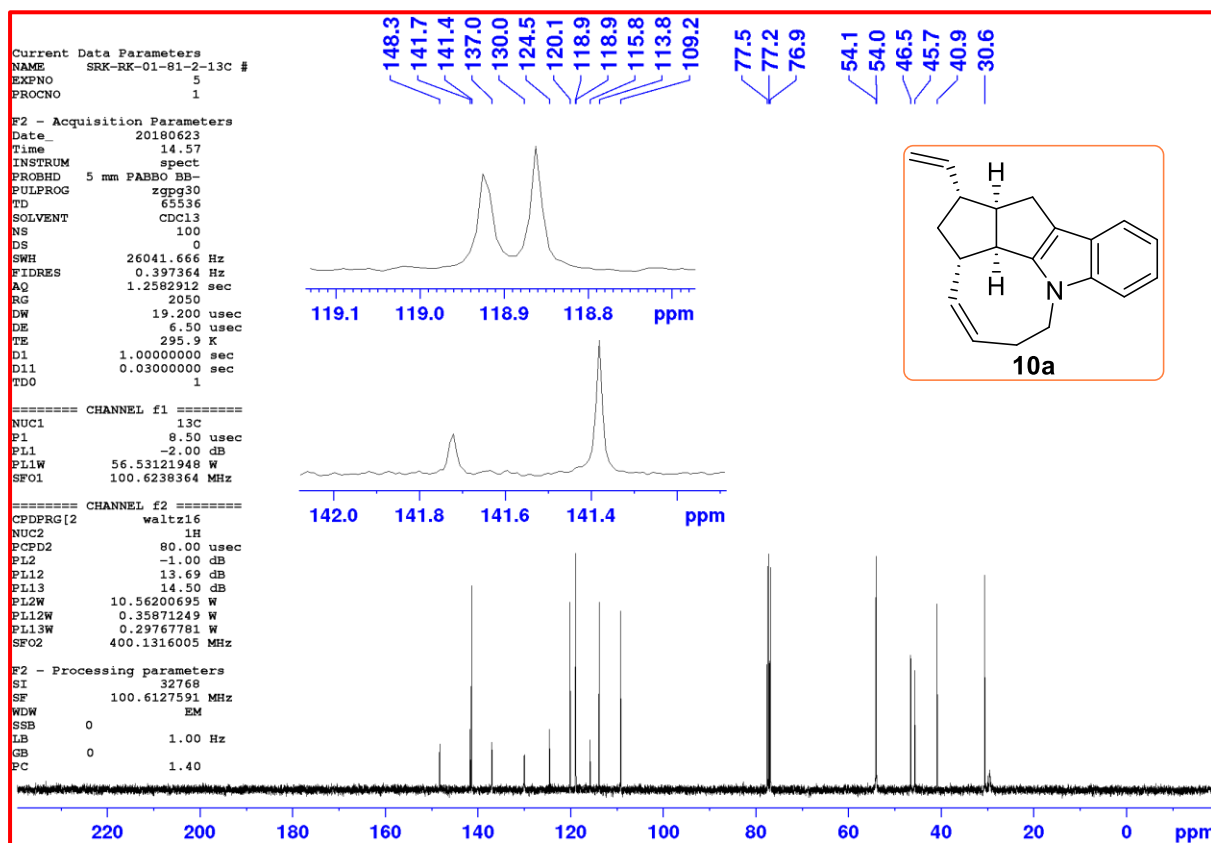
DEPT-135 (100 MHz, CDCl<sub>3</sub>) of **22**



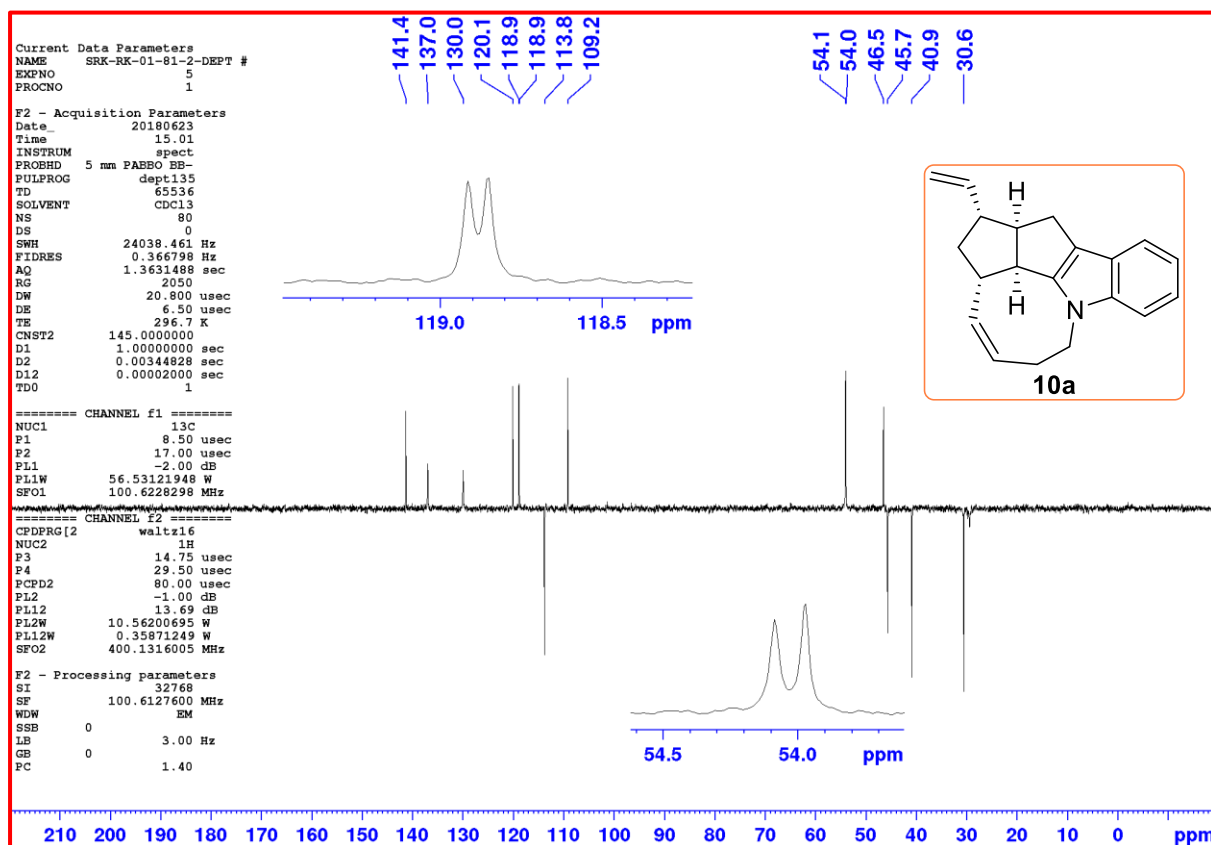
<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of **10a**



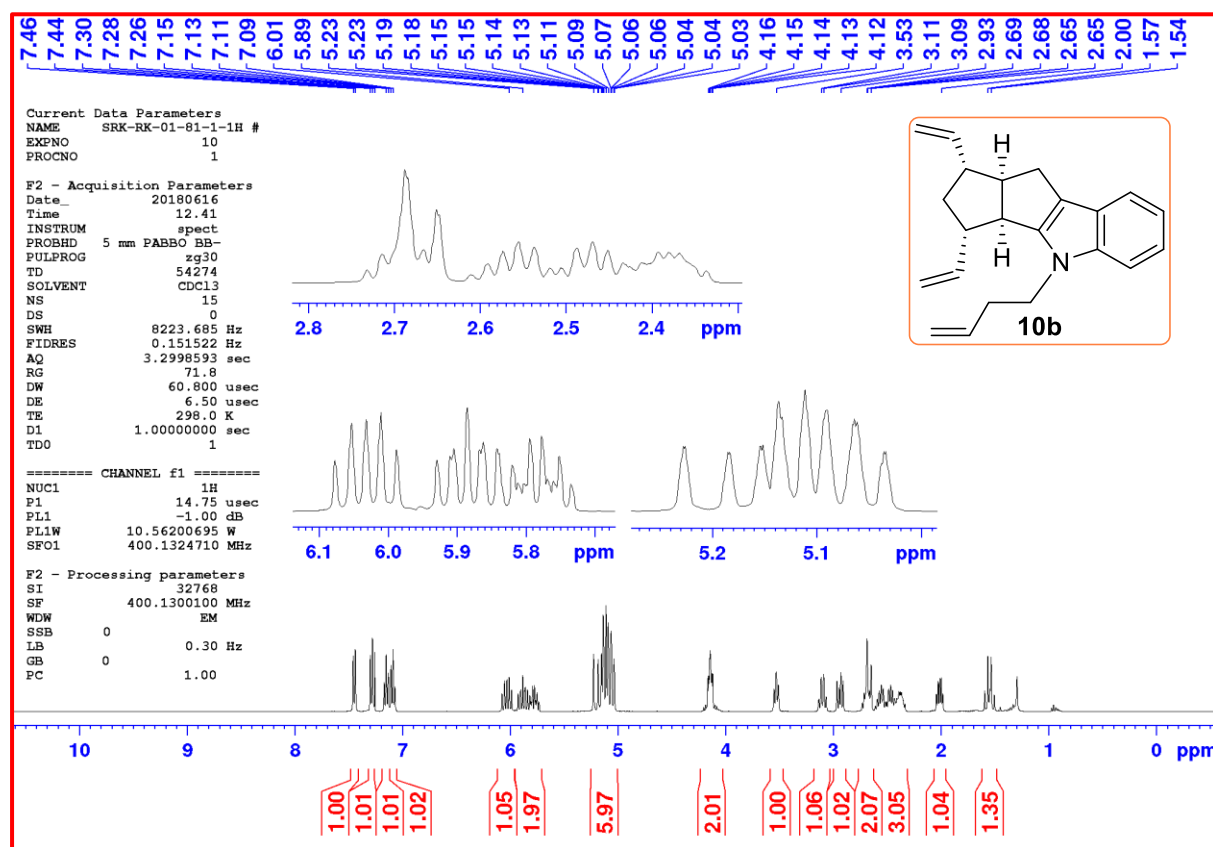
$^{13}\text{C}$  NMR (100 MHz,  $\text{CDCl}_3$ ) of **10a**



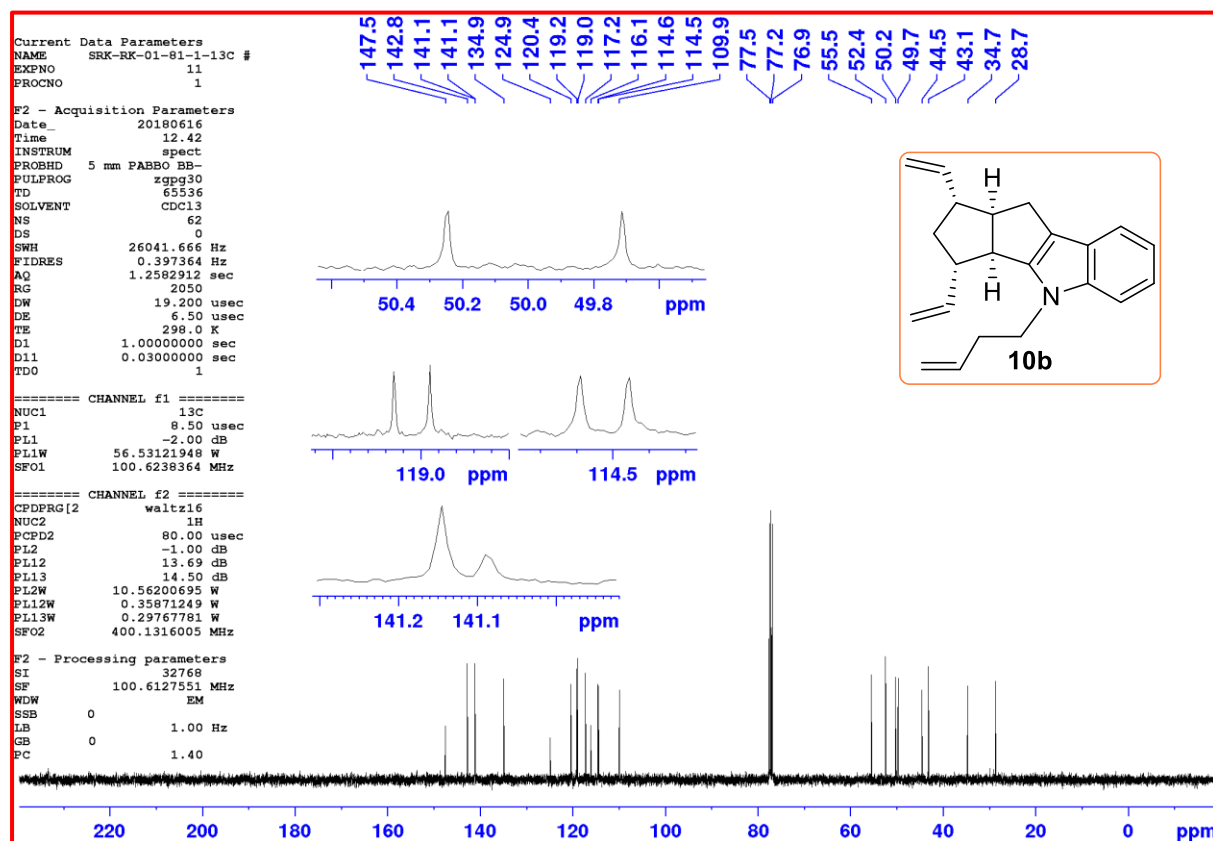
DEPT-135 (100 MHz,  $\text{CDCl}_3$ ) of **10a**



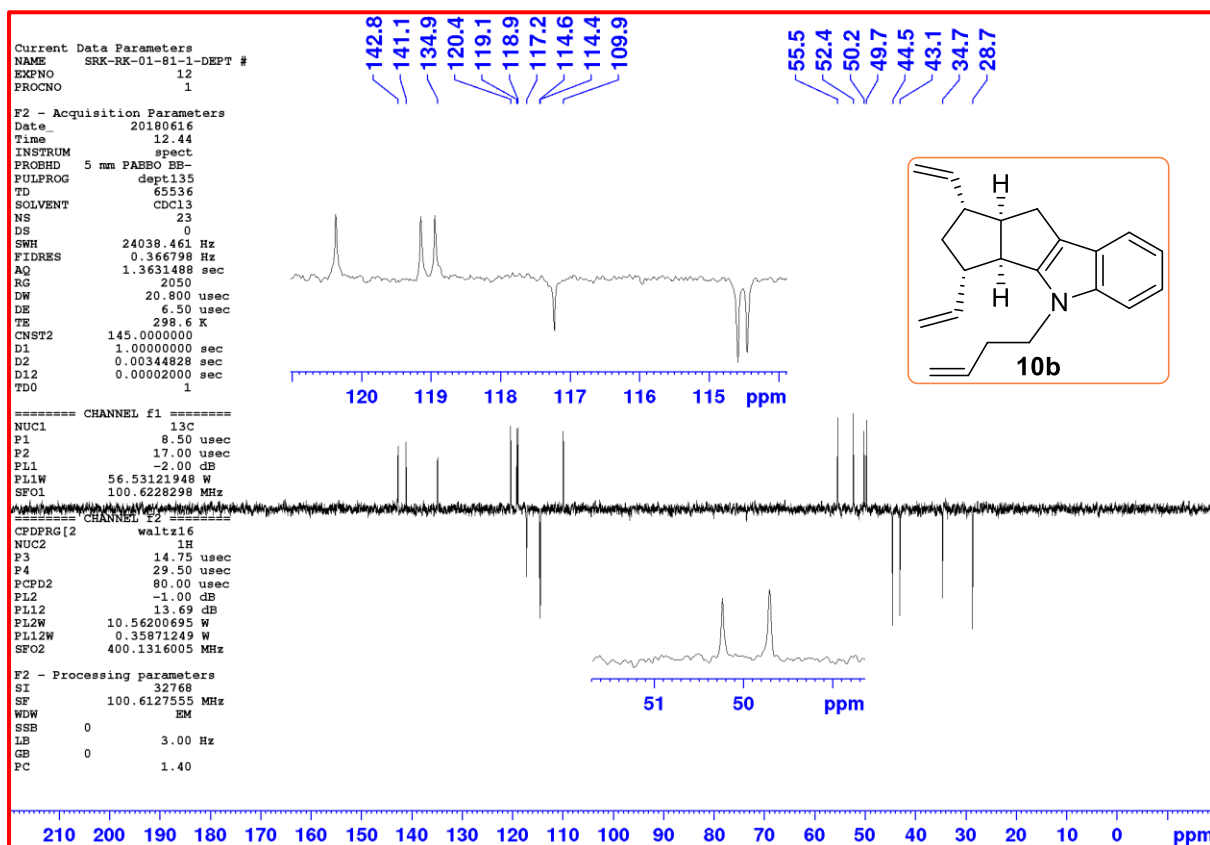
<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of **10b**



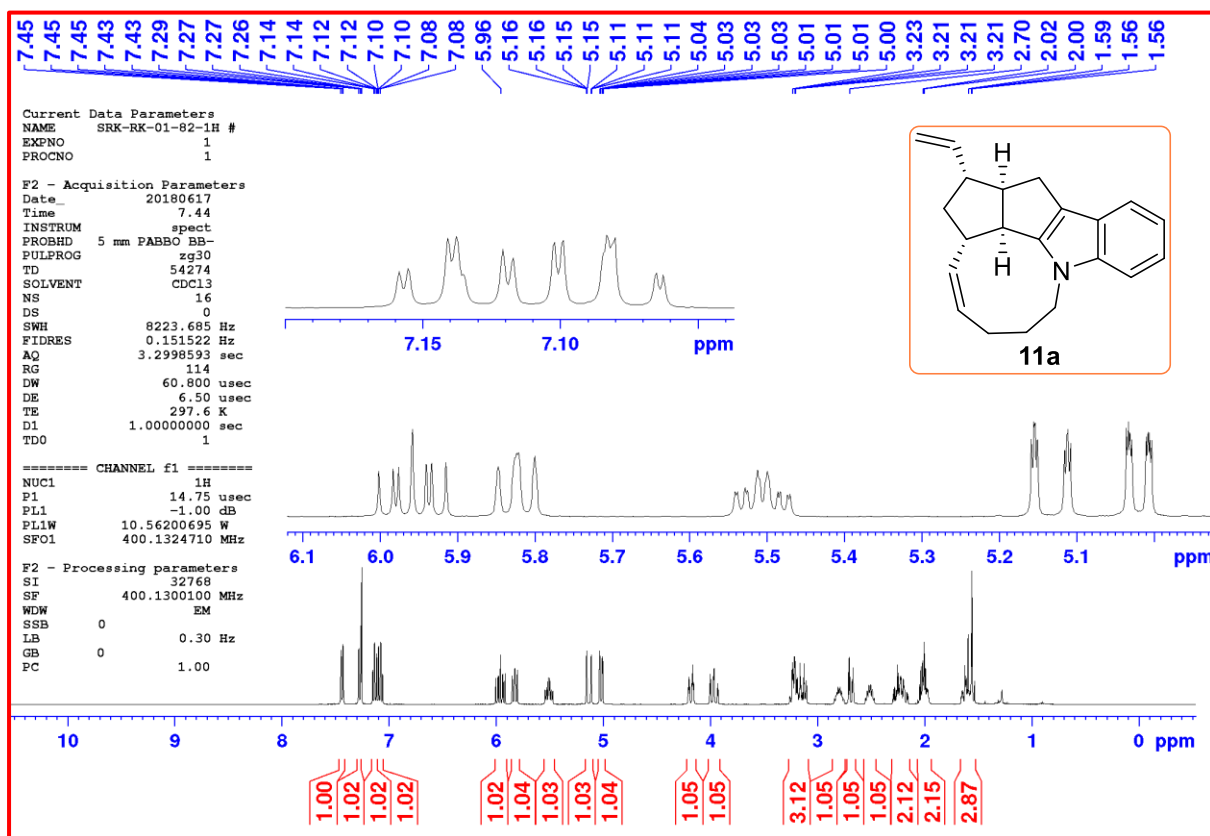
<sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) of **10b**



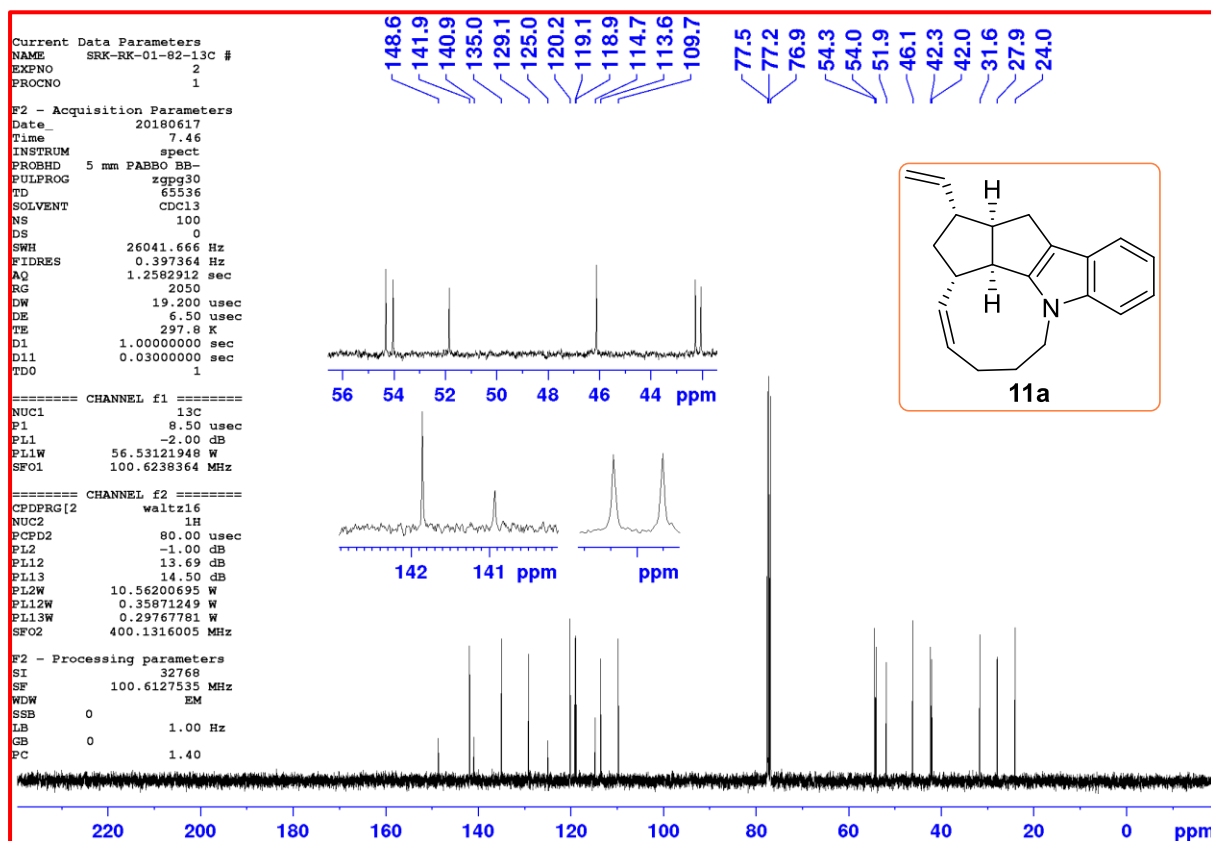
DEPT-135 (100 MHz, CDCl<sub>3</sub>) of **10b**



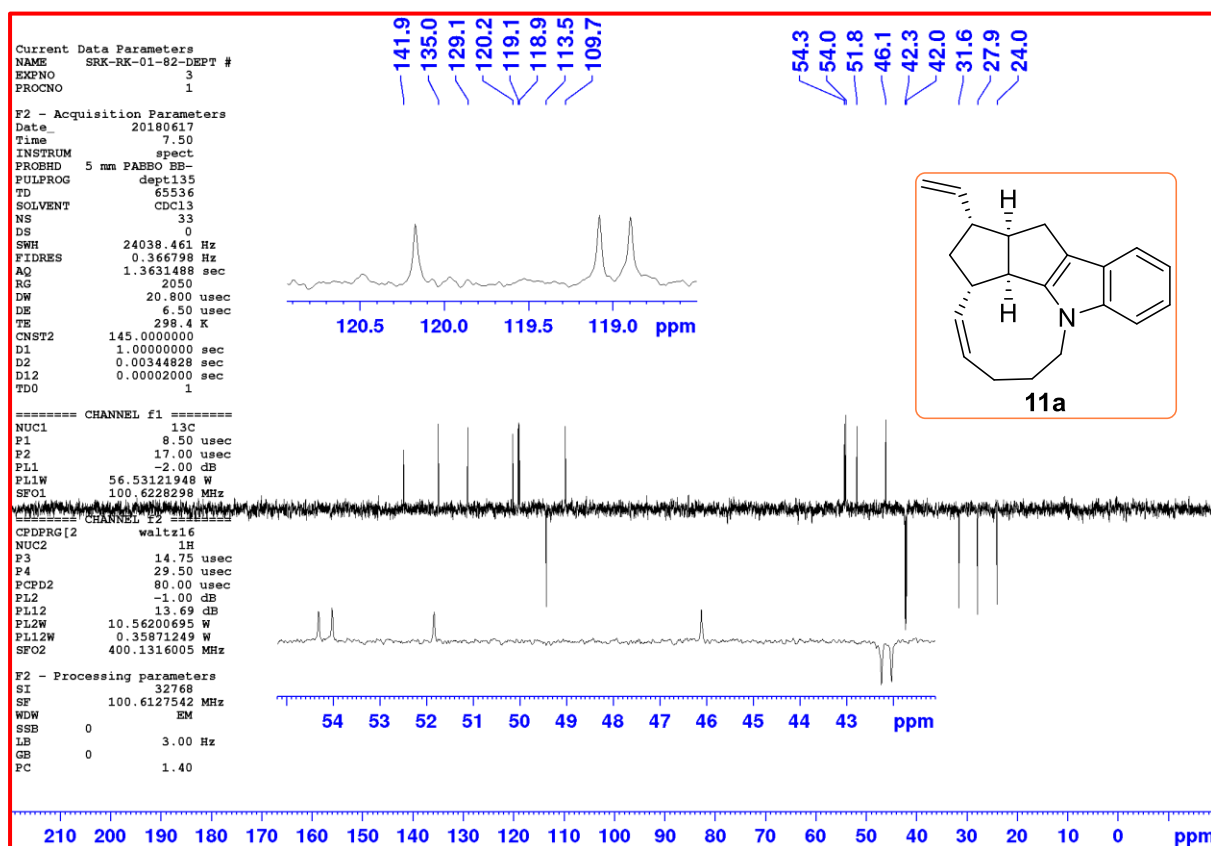
<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of **11a**



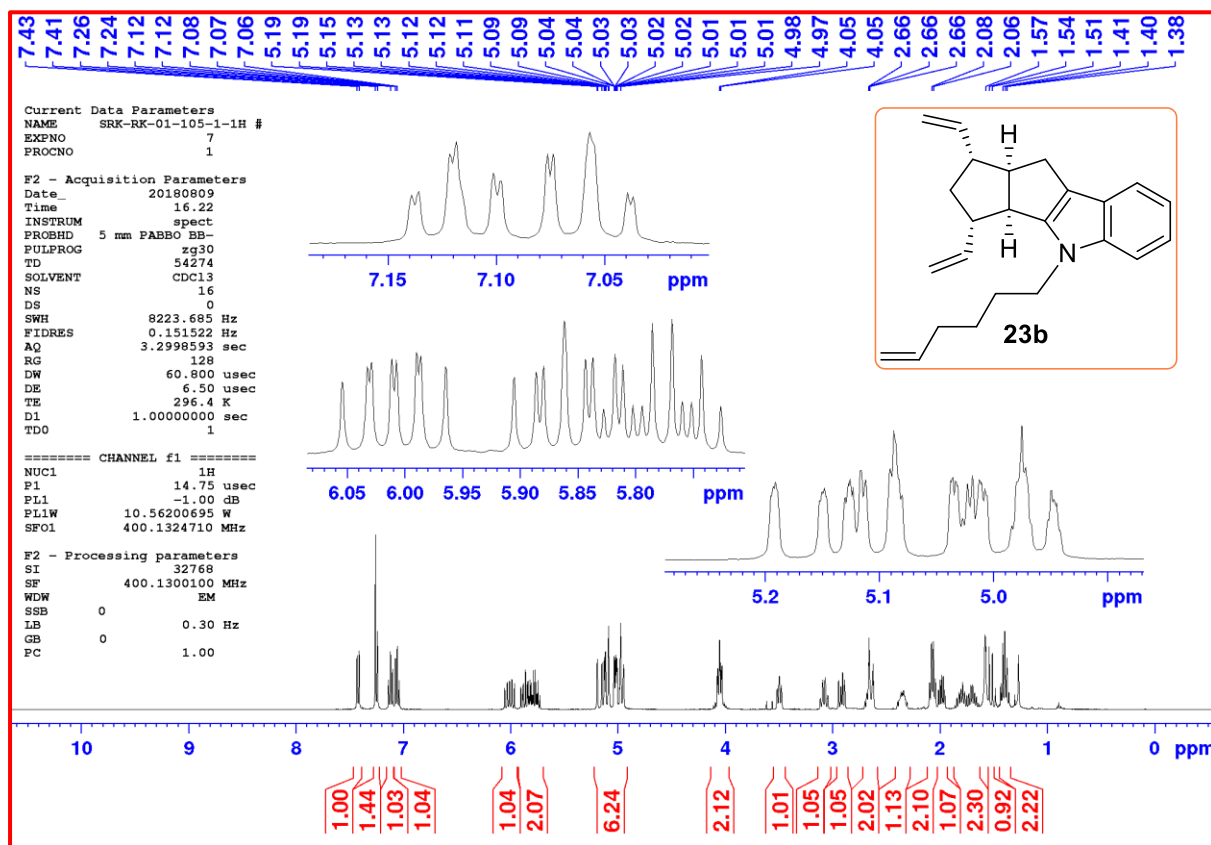
### $^{13}\text{C}$ NMR (100 MHz, $\text{CDCl}_3$ ) of **11a**



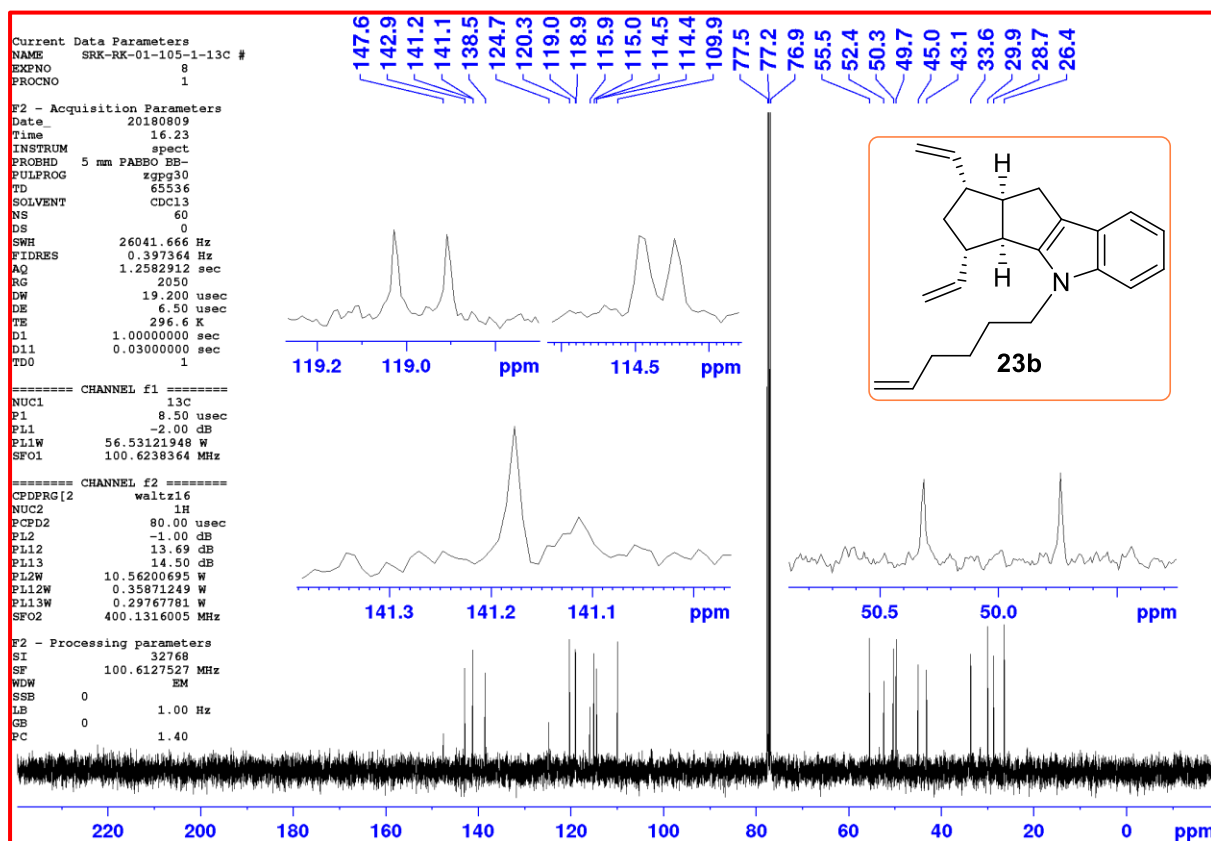
### DEPT-135 (100 MHz, $\text{CDCl}_3$ ) of **11a**



<sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) of **23b**



<sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) of **23b**



DEPT-135 (100 MHz, CDCl<sub>3</sub>) of **23b**

