

Pharmacophore-based virtual screening, synthesis, biological evaluation, and molecular docking study of novel pyrrolizines bearing urea/thiourea moieties with potential anticancer and CDK inhibitory activities

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Chemical structures of the compound library

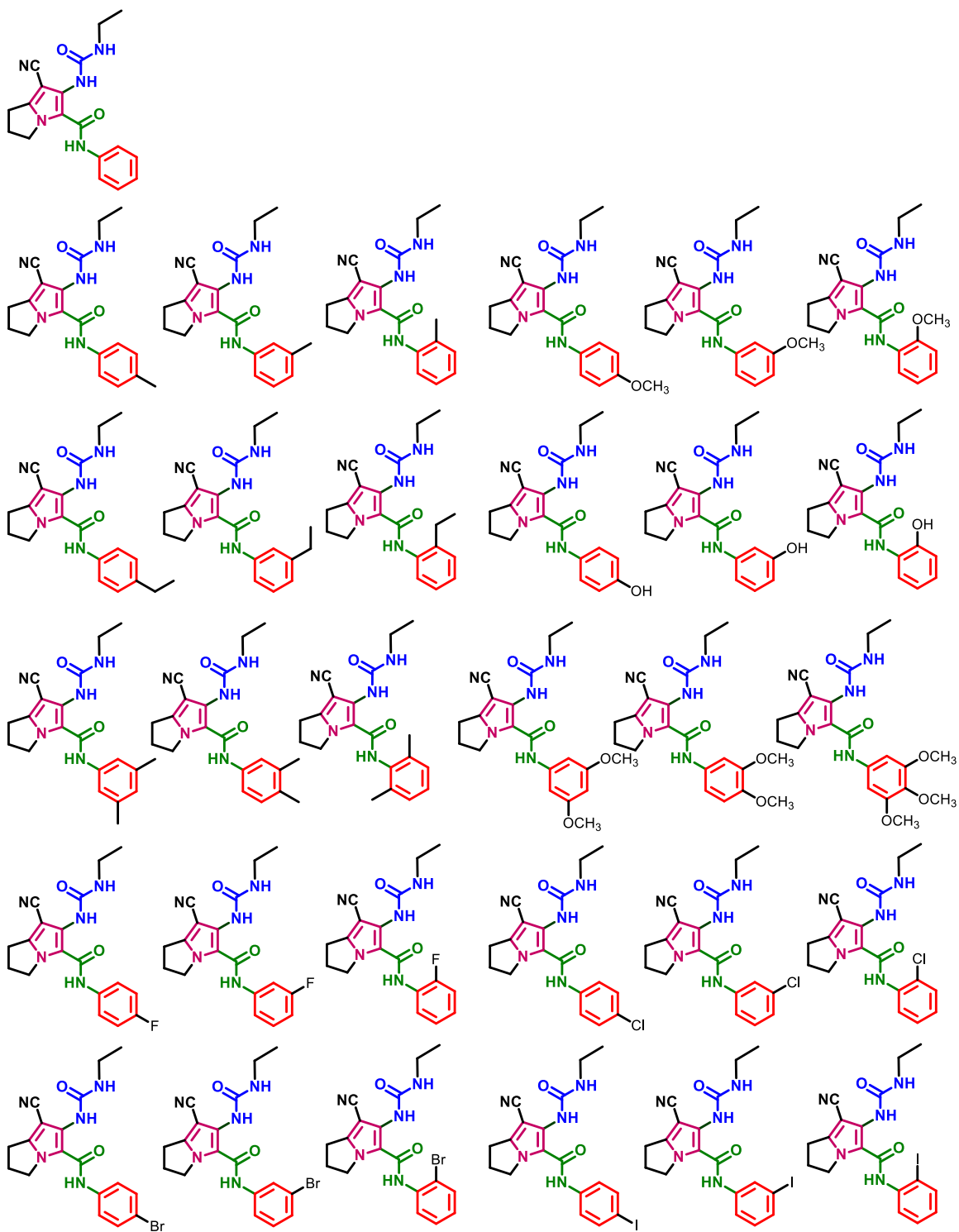


Figure S1. Chemical structure of compound 1-31 (Etiso)

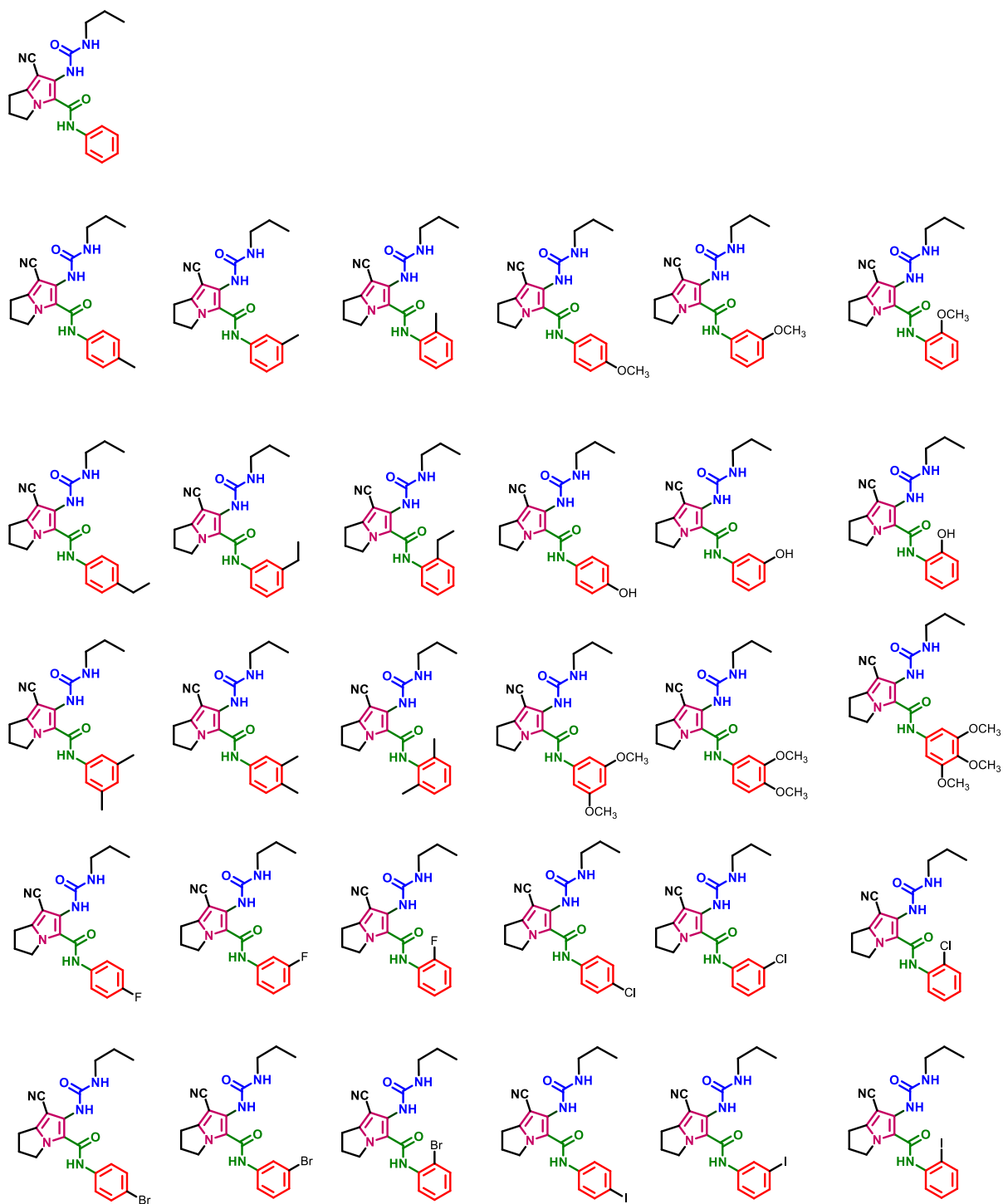


Figure S2. Chemical structure of compound 32-62 (Propyliso)

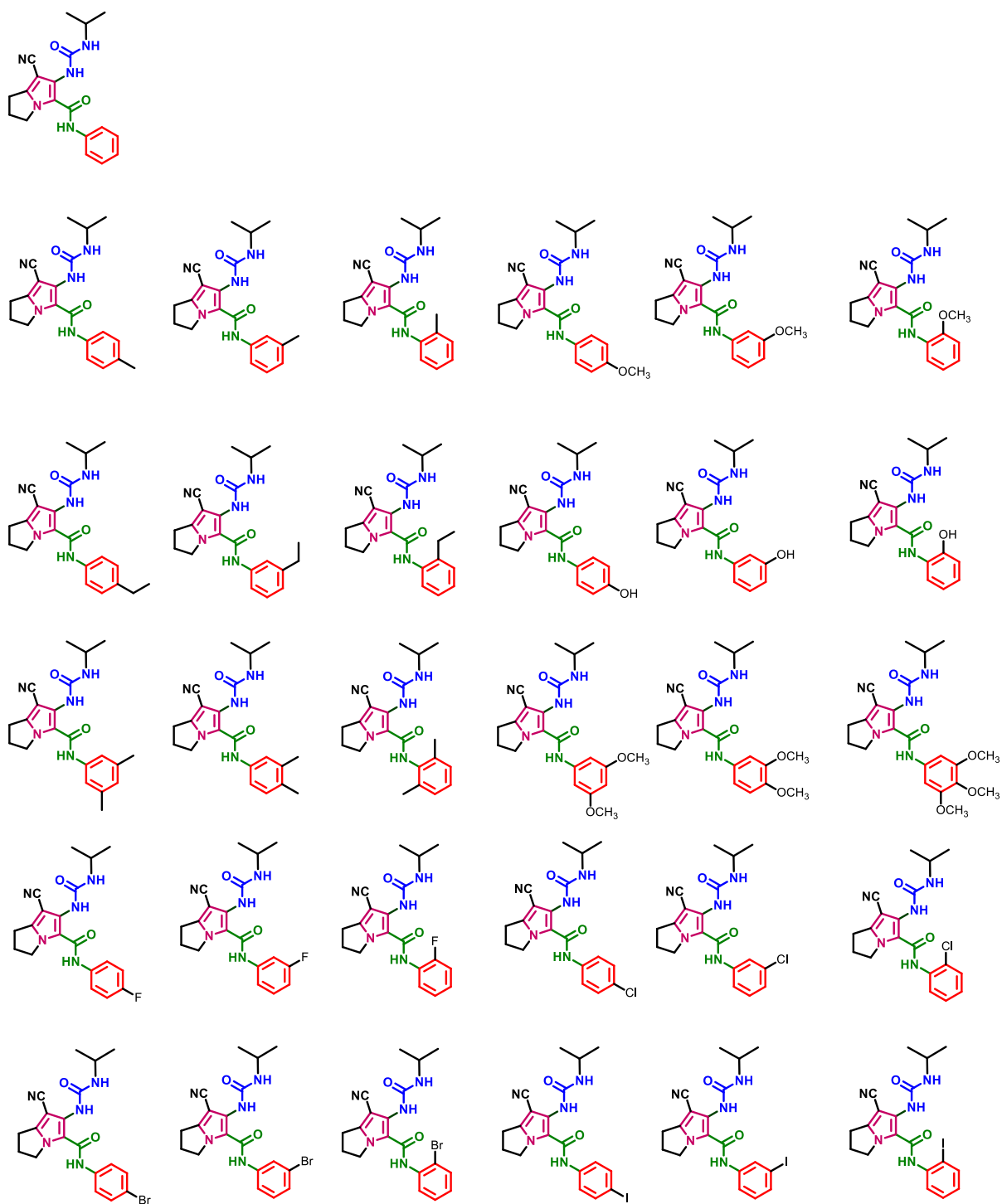


Figure S3. Chemical structure of compound 63-93 (Isopropyliso)

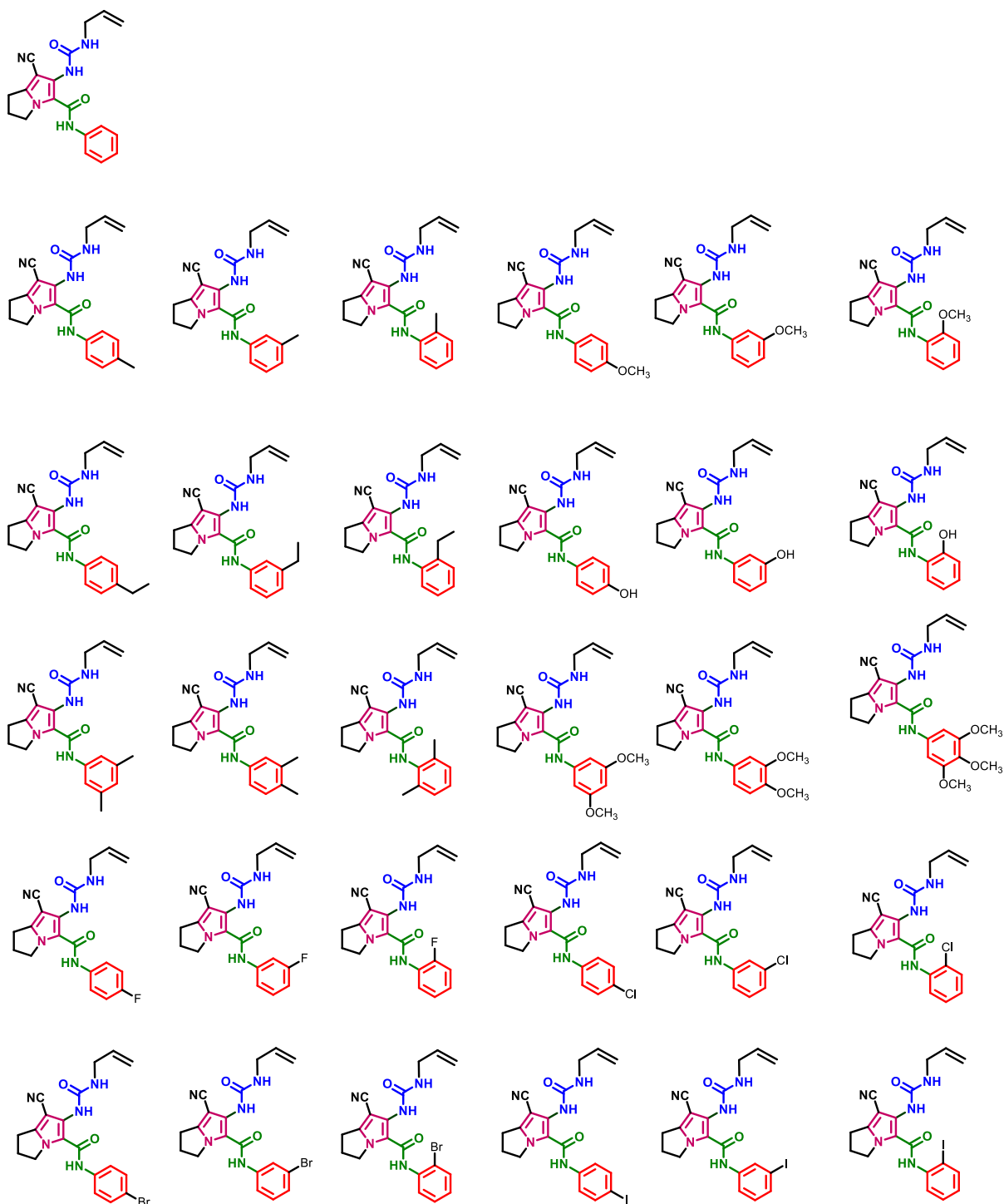


Figure S4. Chemical structure of compound 94-124 (allyliso)

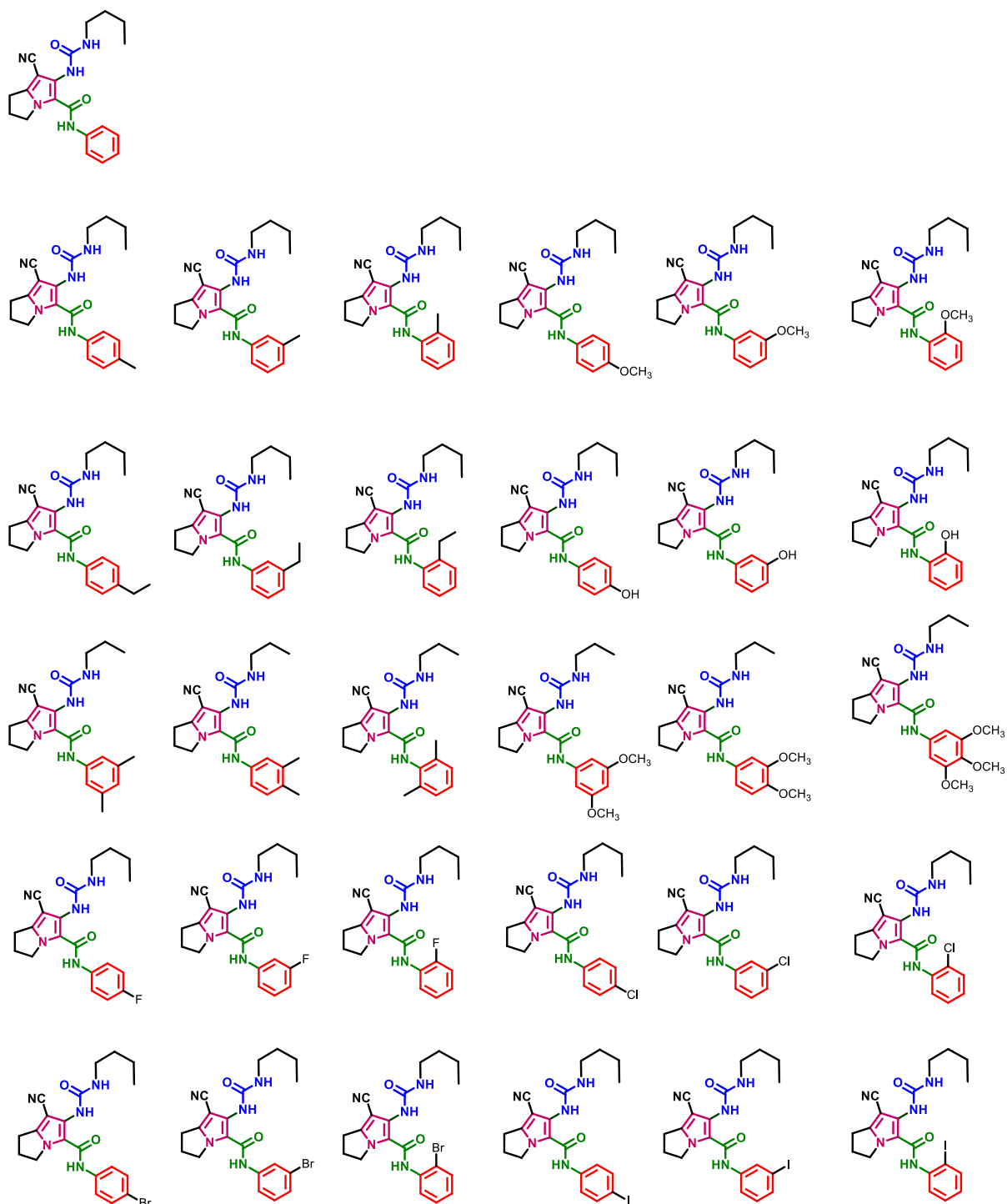


Figure S5. Chemical structure of compound 125-155 (butyliso)

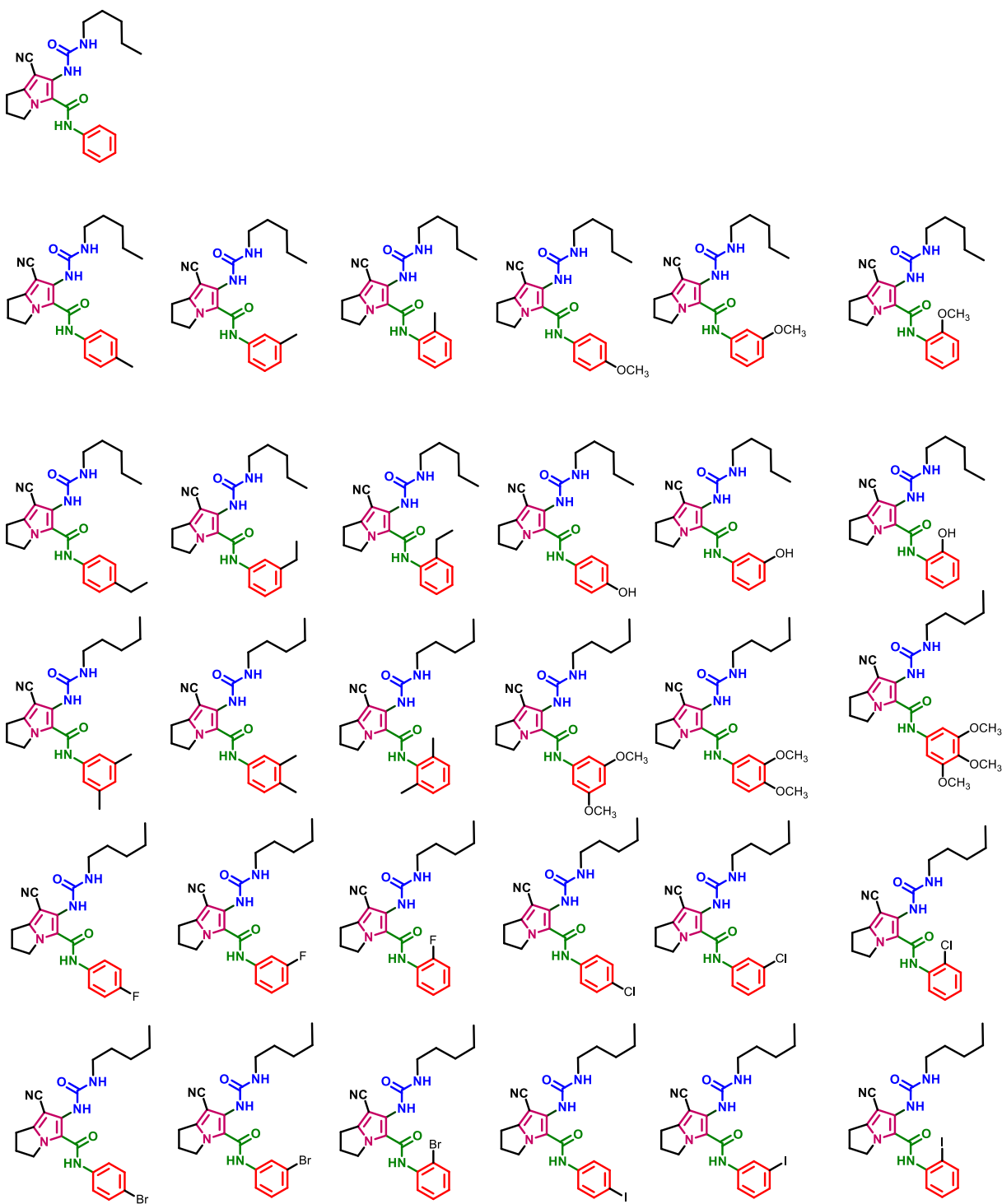


Figure S6. Chemical structure of compound 156-186 (pentyliso)

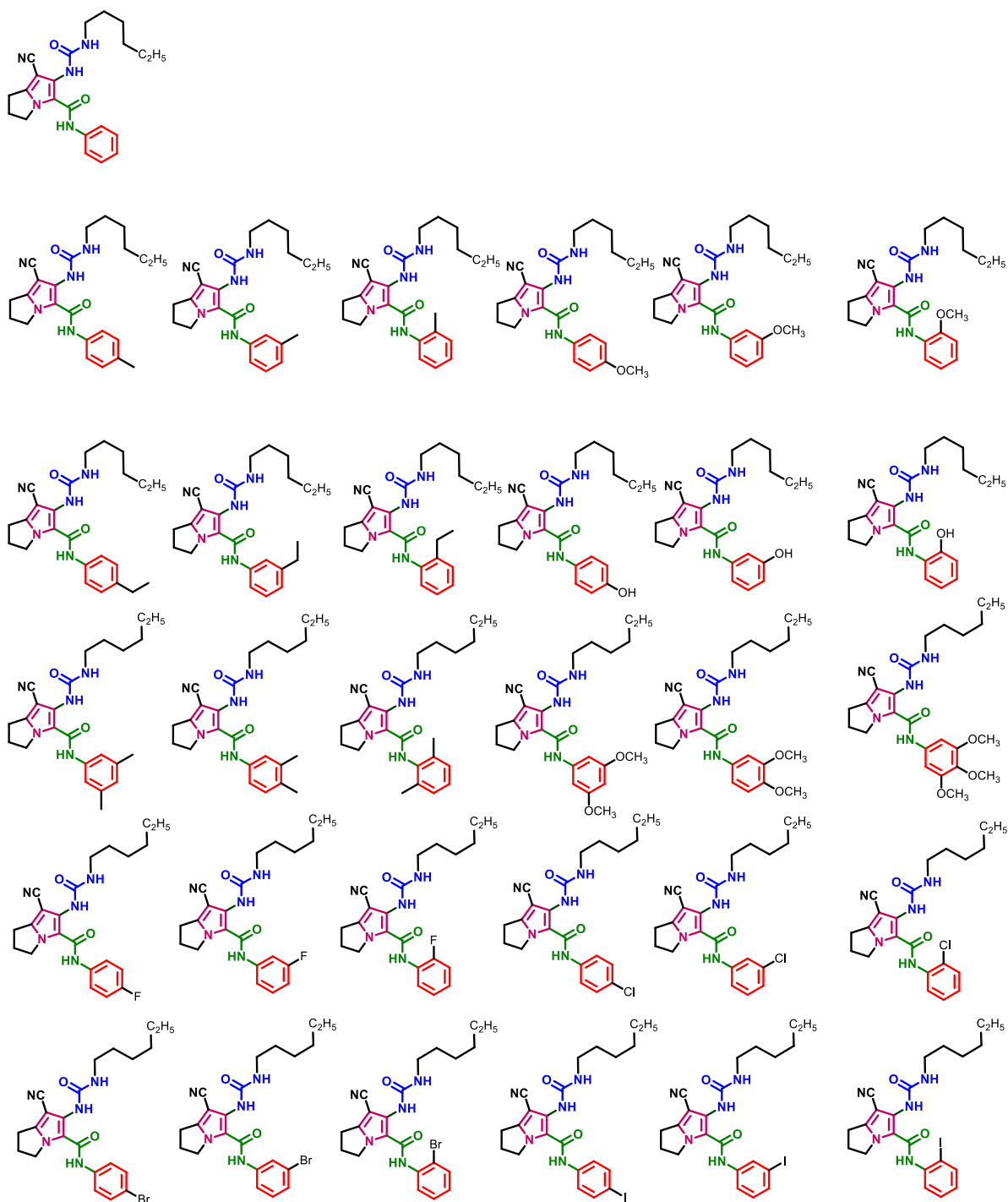


Figure S7. Chemical structure of compound 187-217 (hexyliso)

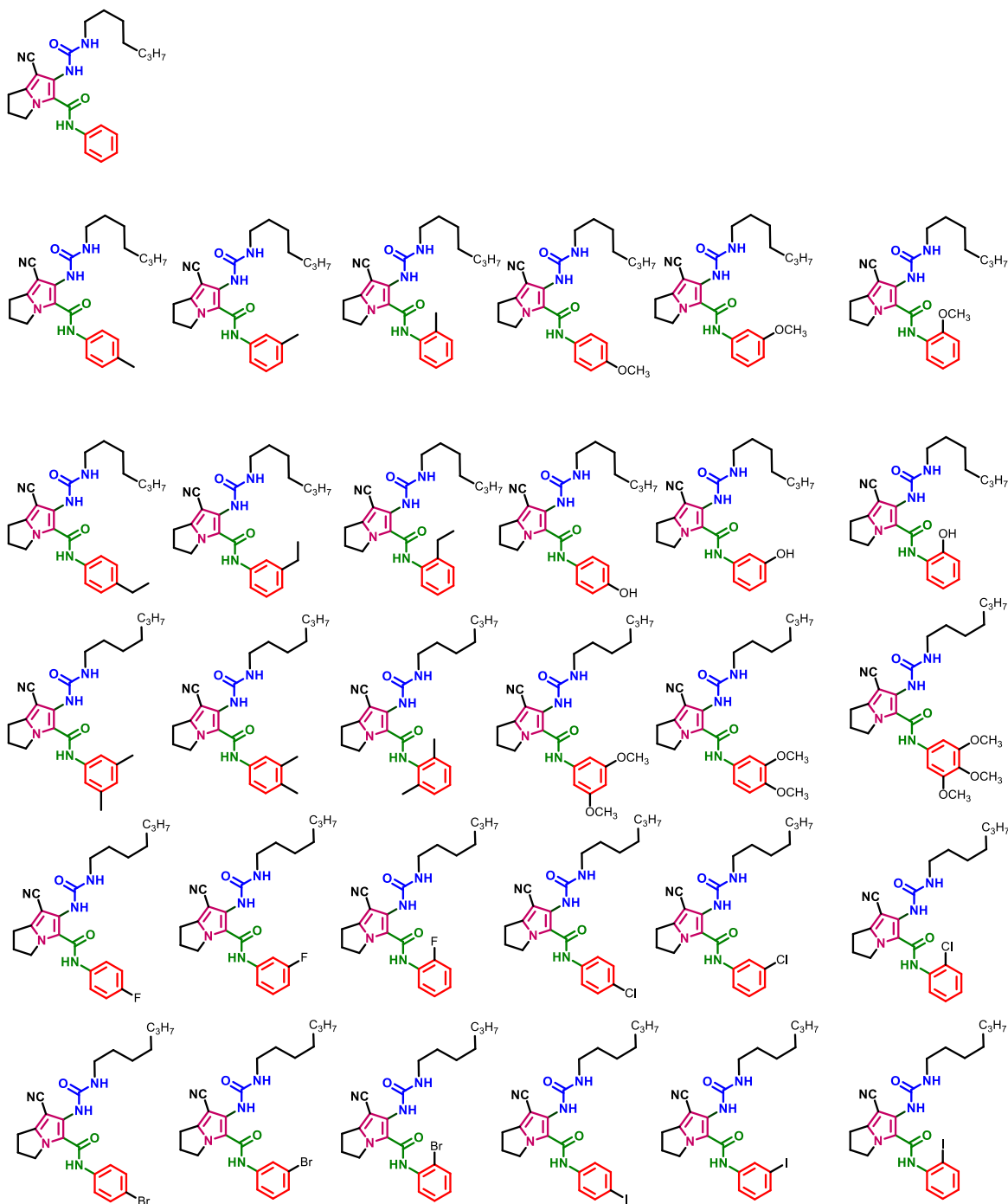


Figure S8. Chemical structure of compound 218-248 (heptyliso)

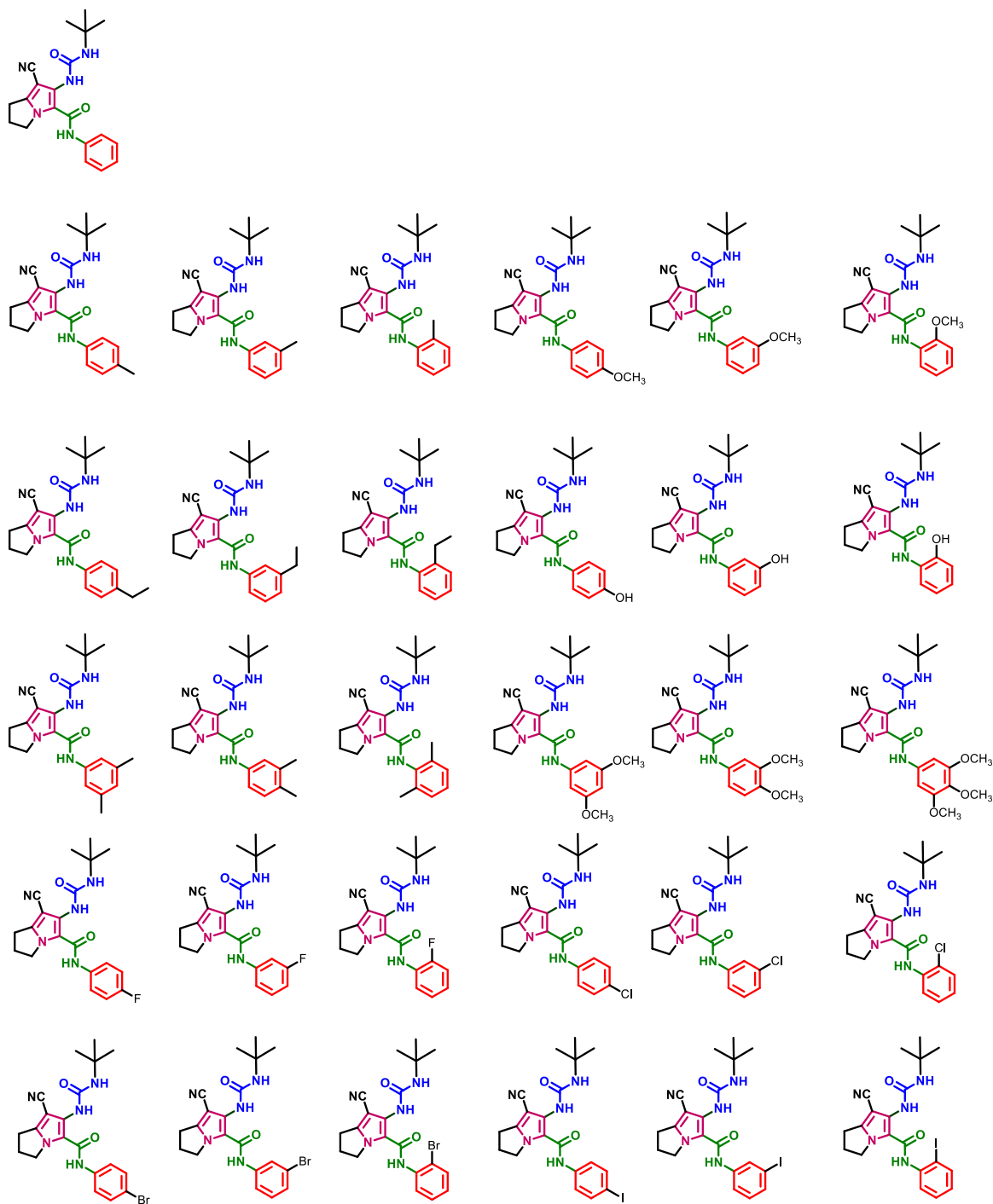


Figure S9. Chemical structure of compound 249-279 (tertbutyl)

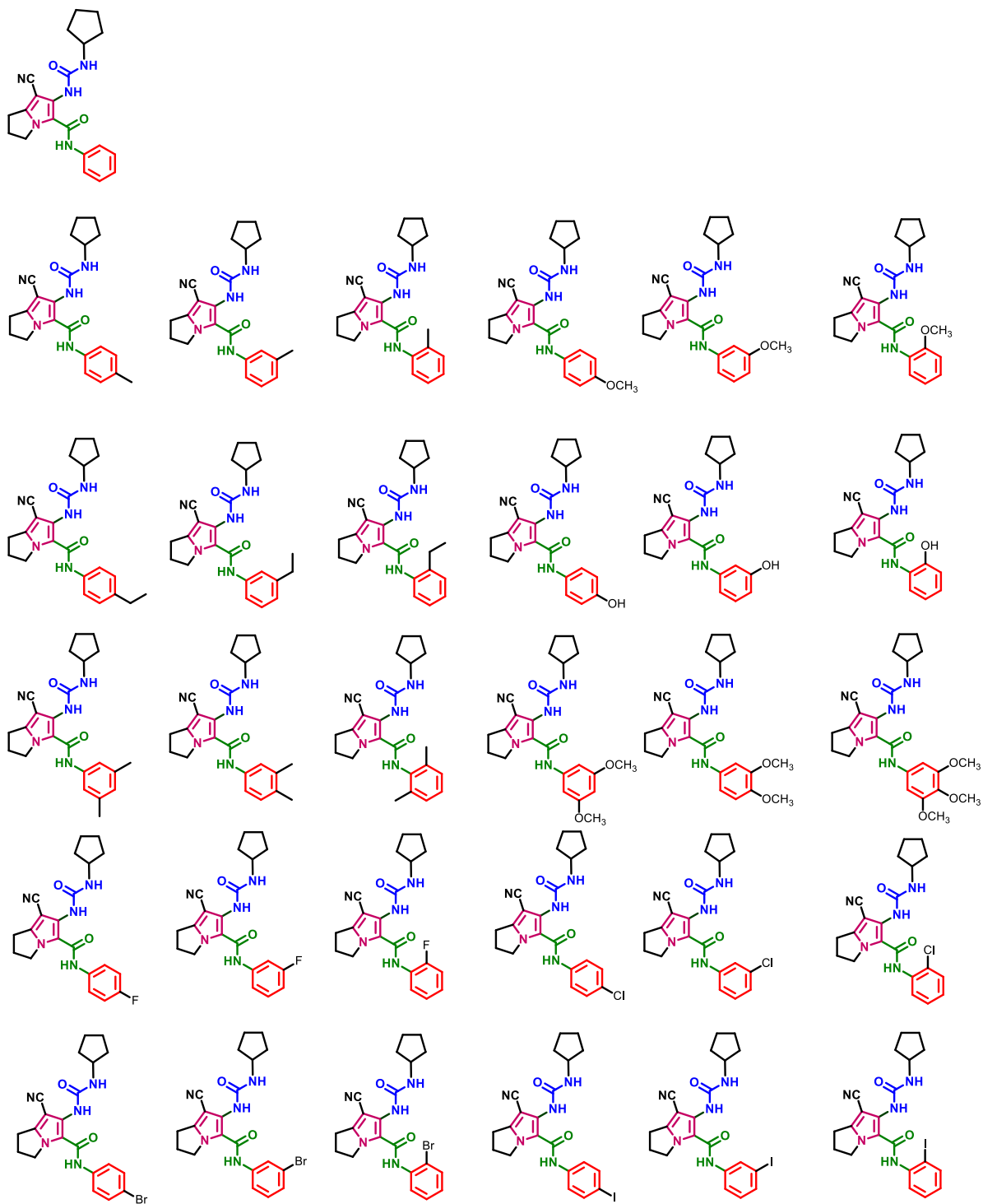


Figure S10. Chemical structure of compound 280-310 (cyclopentyl)

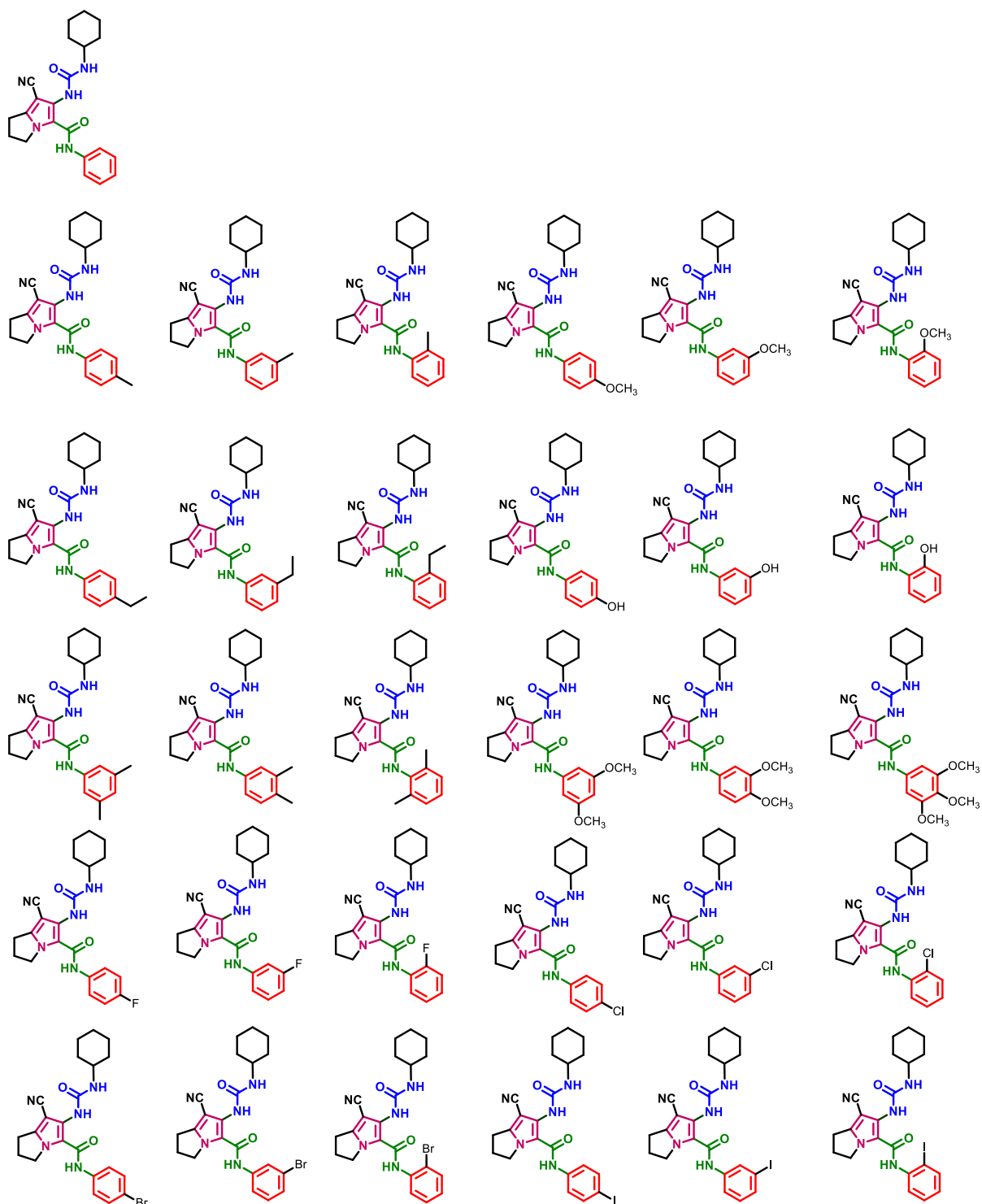


Figure S11. Chemical structure of compound 311-341 (cyclohexyl)

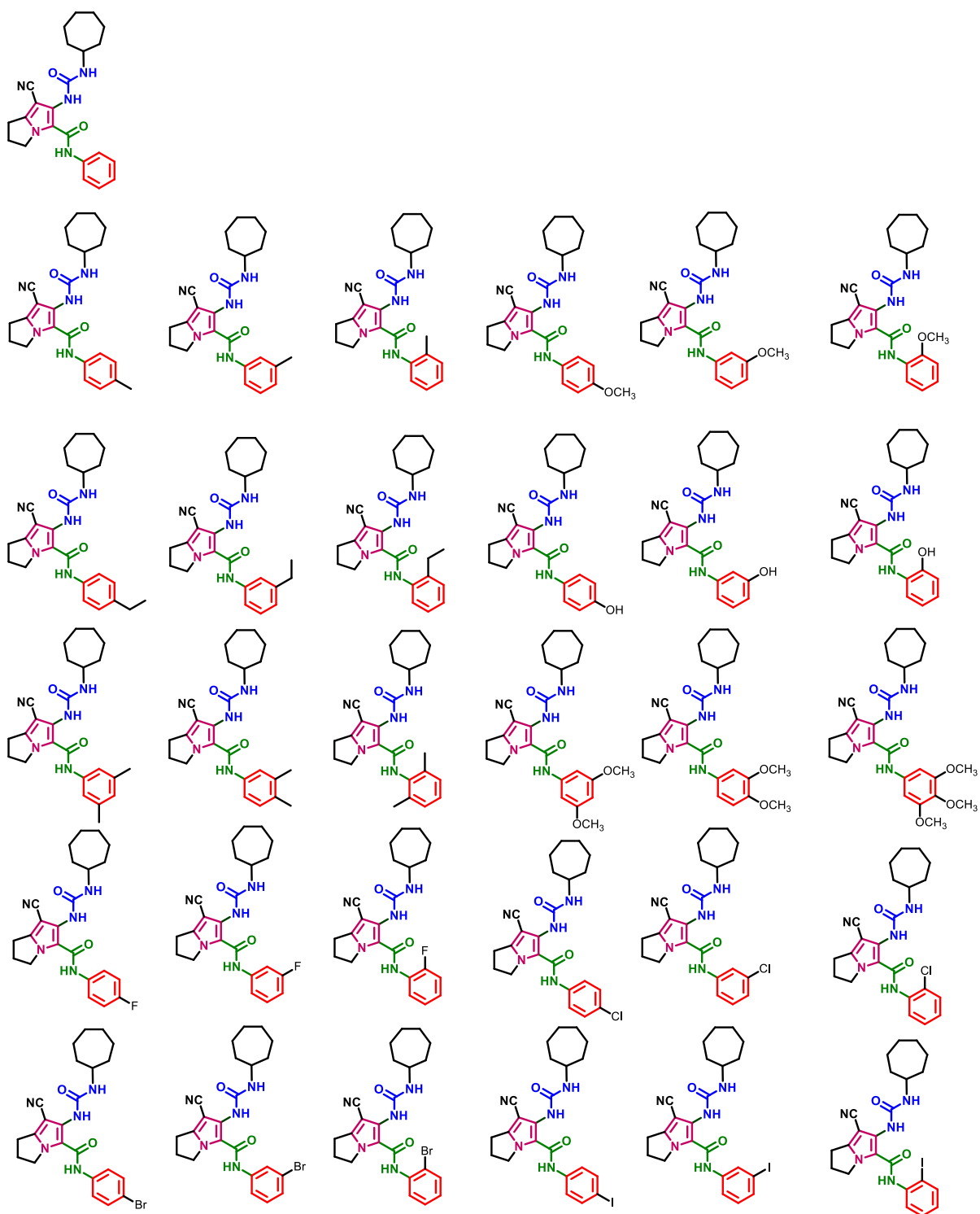


Figure S12. Chemical structure of compound 342-372 (cycloheptyl)

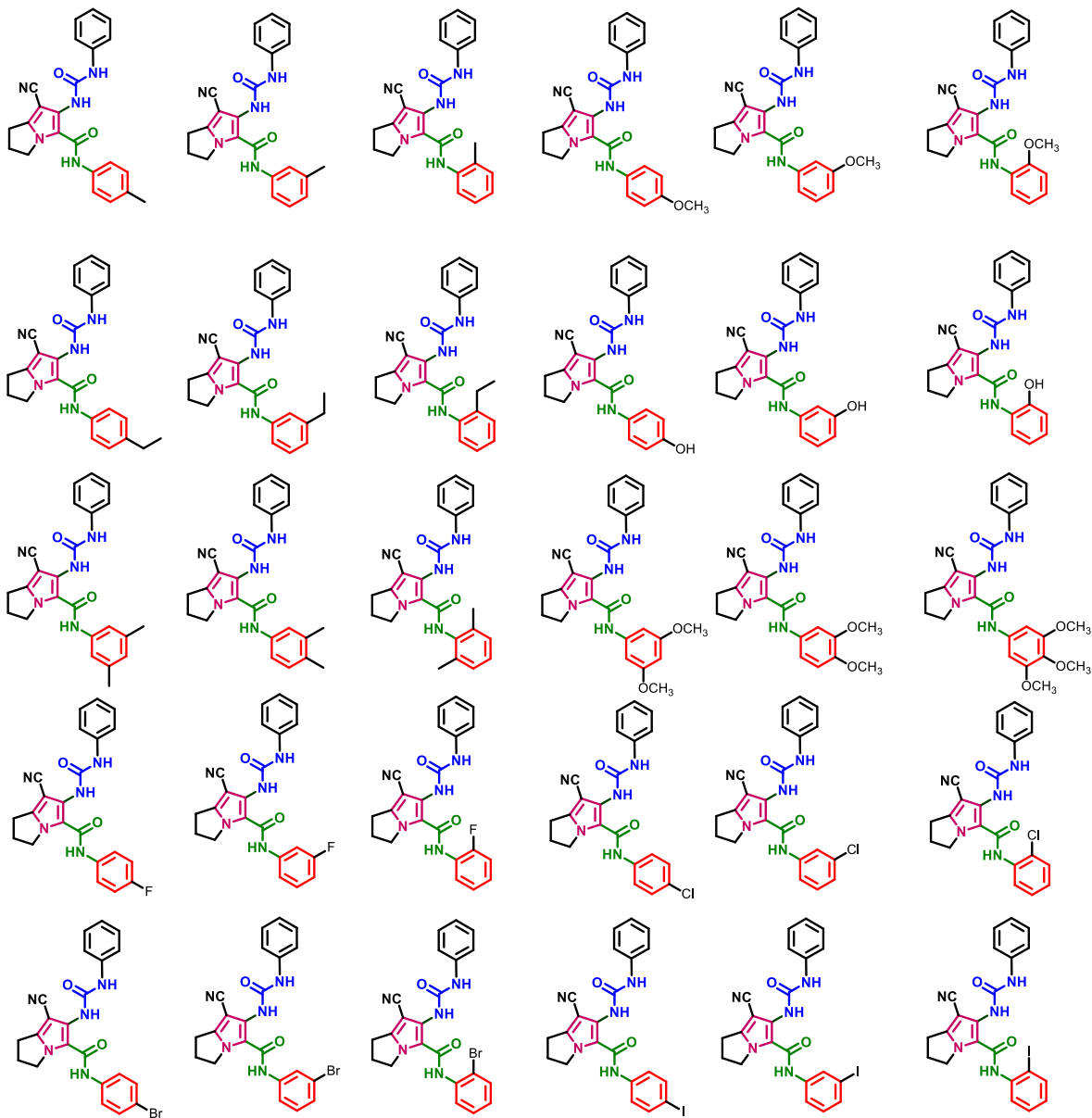
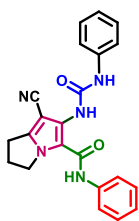


Figure S13. Chemical structure of compound 373-403 (phenyliso)

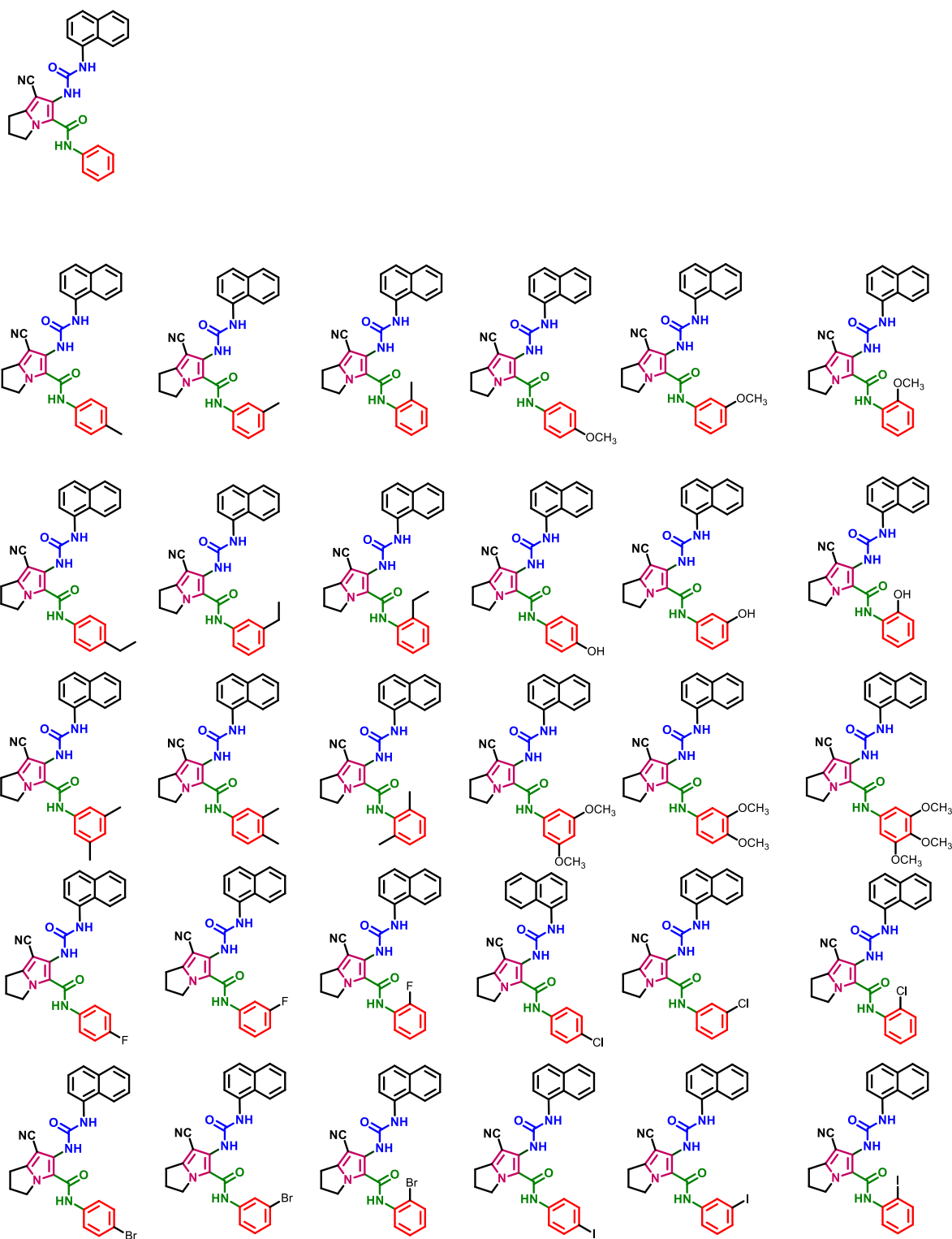


Figure S14. Chemical structure of compound 404-434 (1naphthyliso)

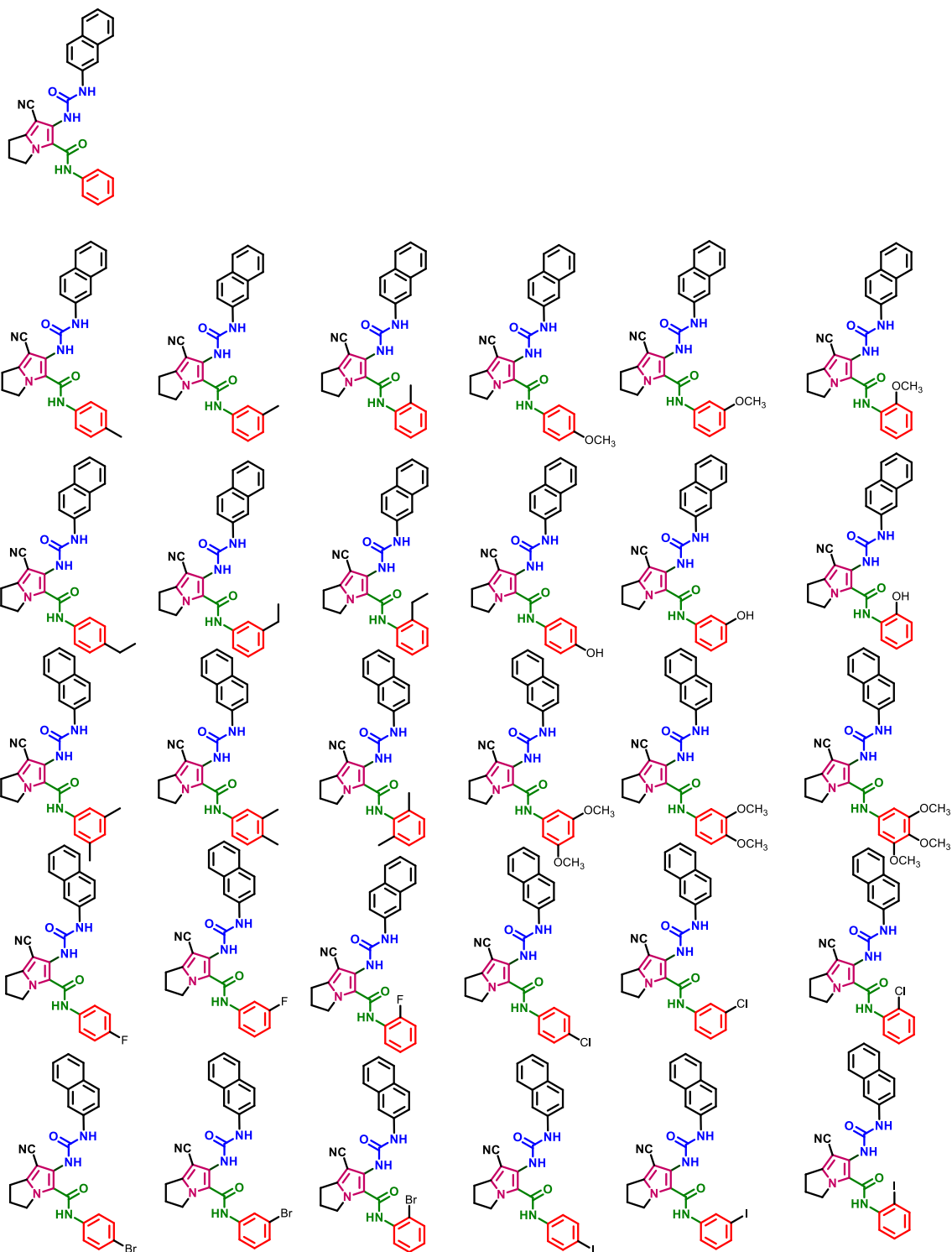


Figure S15. Chemical structure of compound 435-465 (2naphthyliso)

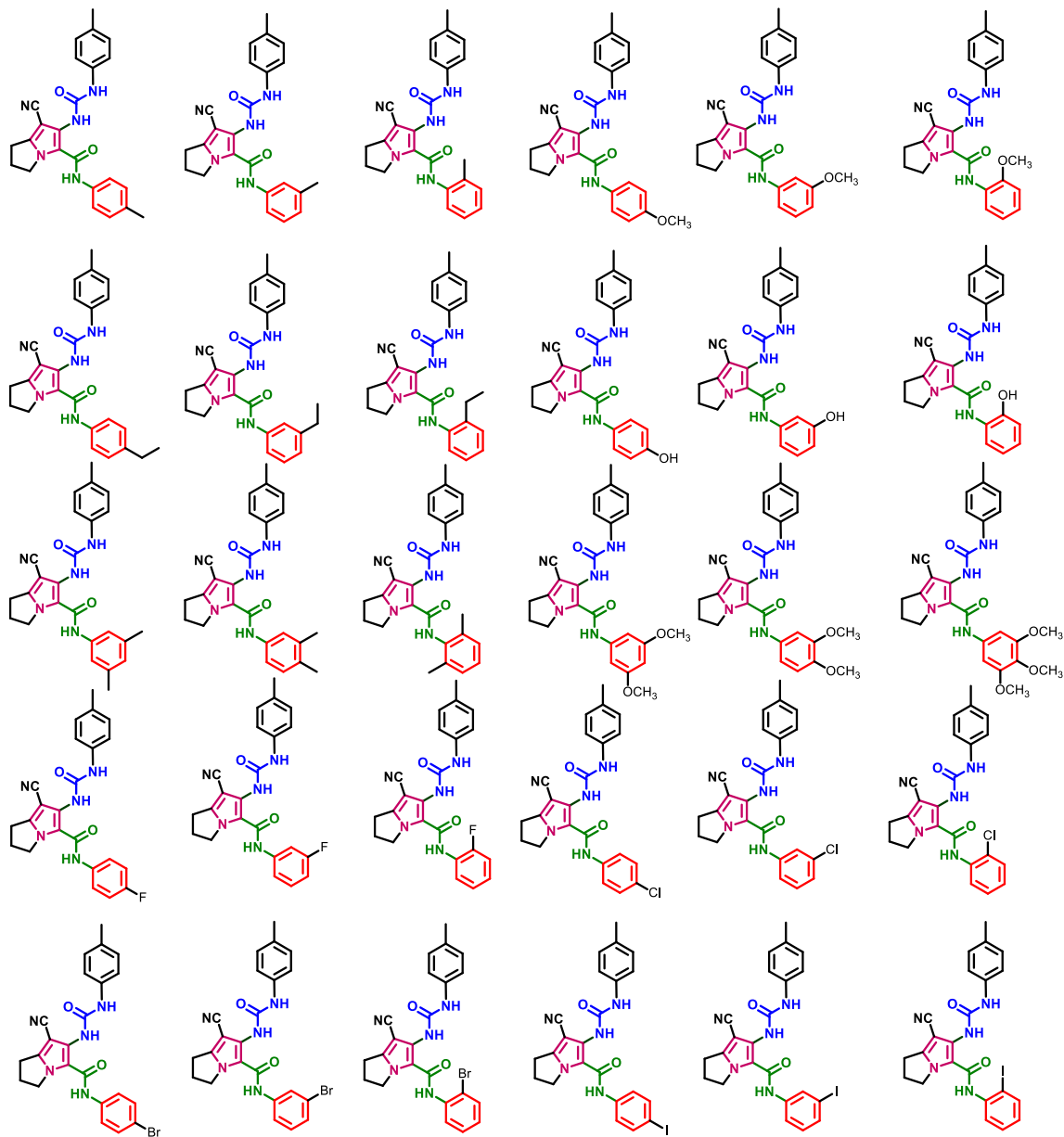
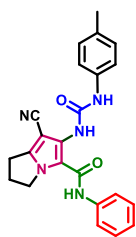


Figure S16. Chemical structure of compound 466-496 (4methylphenyliso)

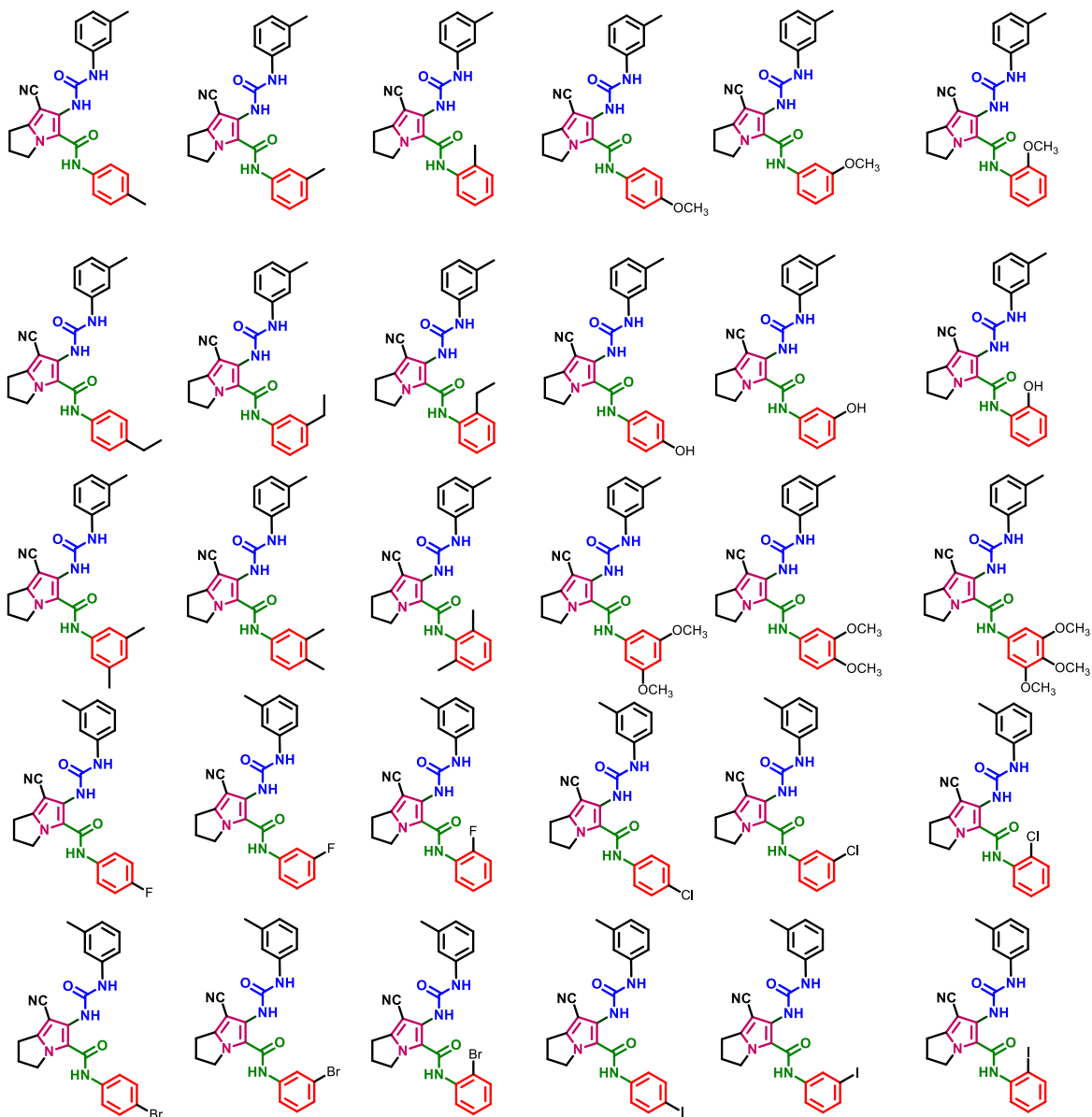
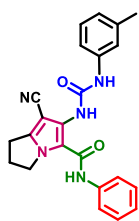


Figure S17. Chemical structure of compound 497-527 (3methylphenyliso)

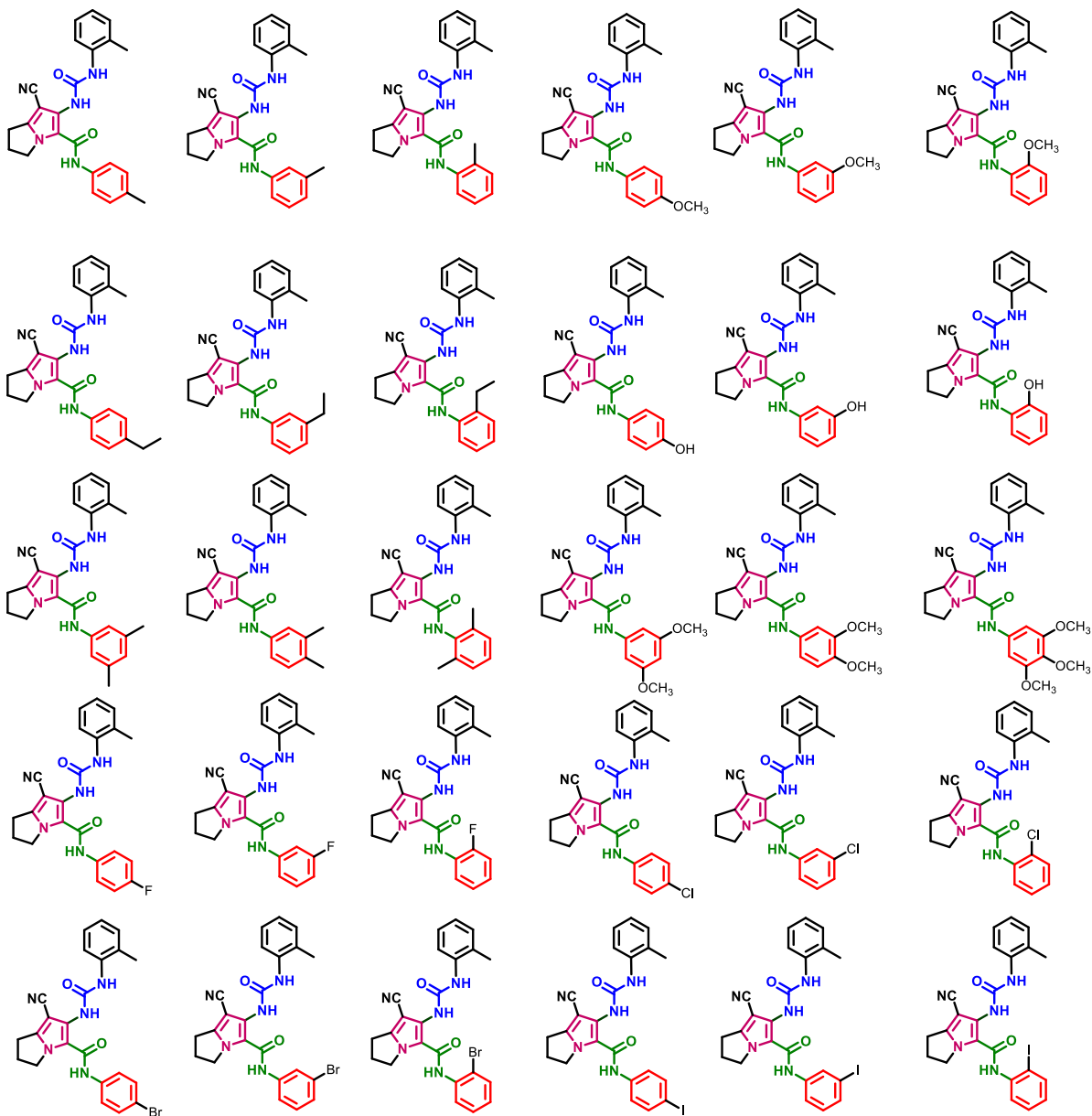
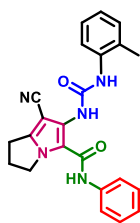


Figure S18. Chemical structure of compound 528-558 (2methylphenyliso)

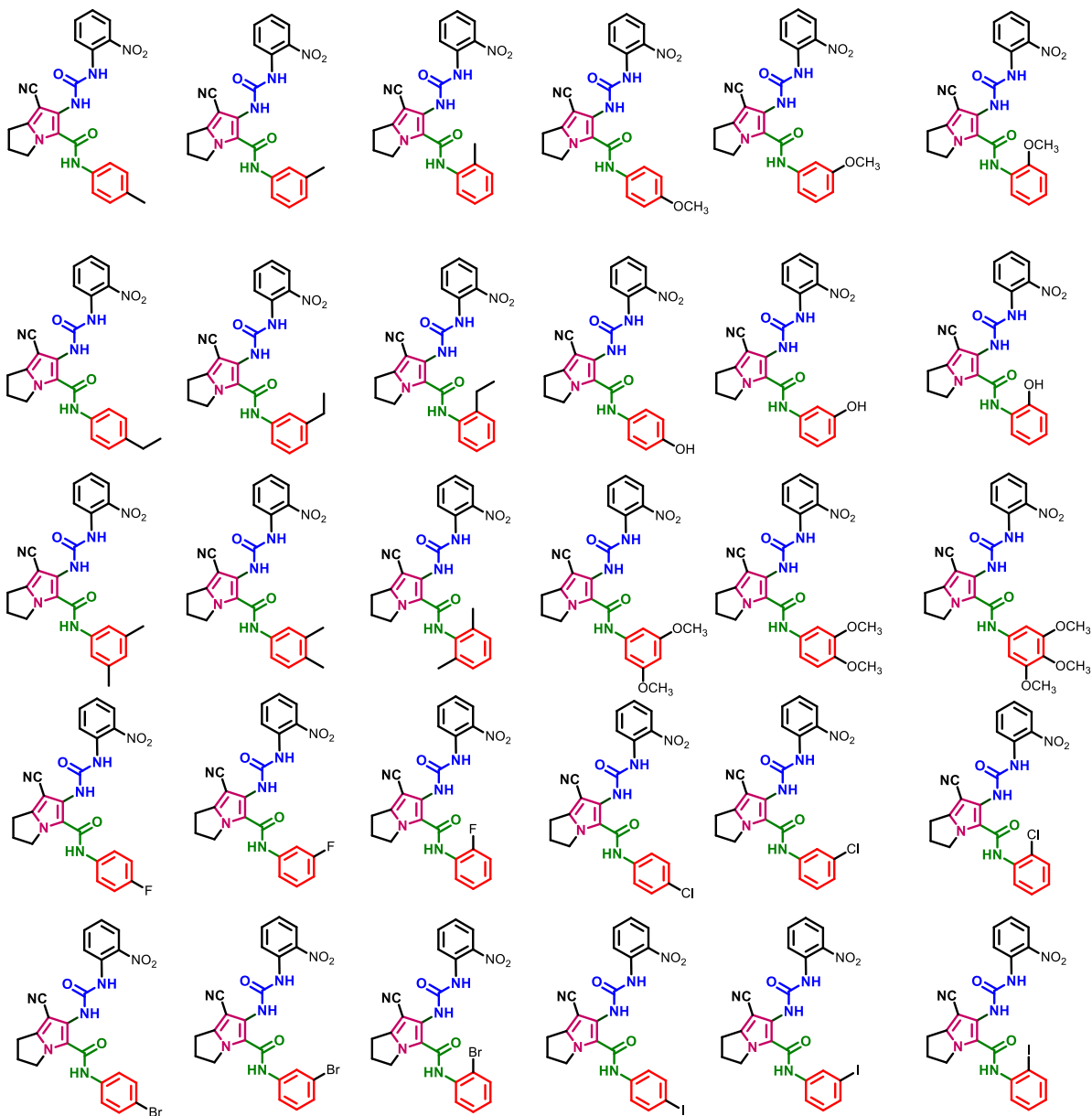
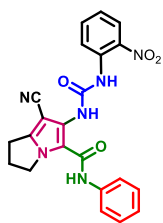


Figure S19. Chemical structure of compound 559-589 (2nitrophenyliso)

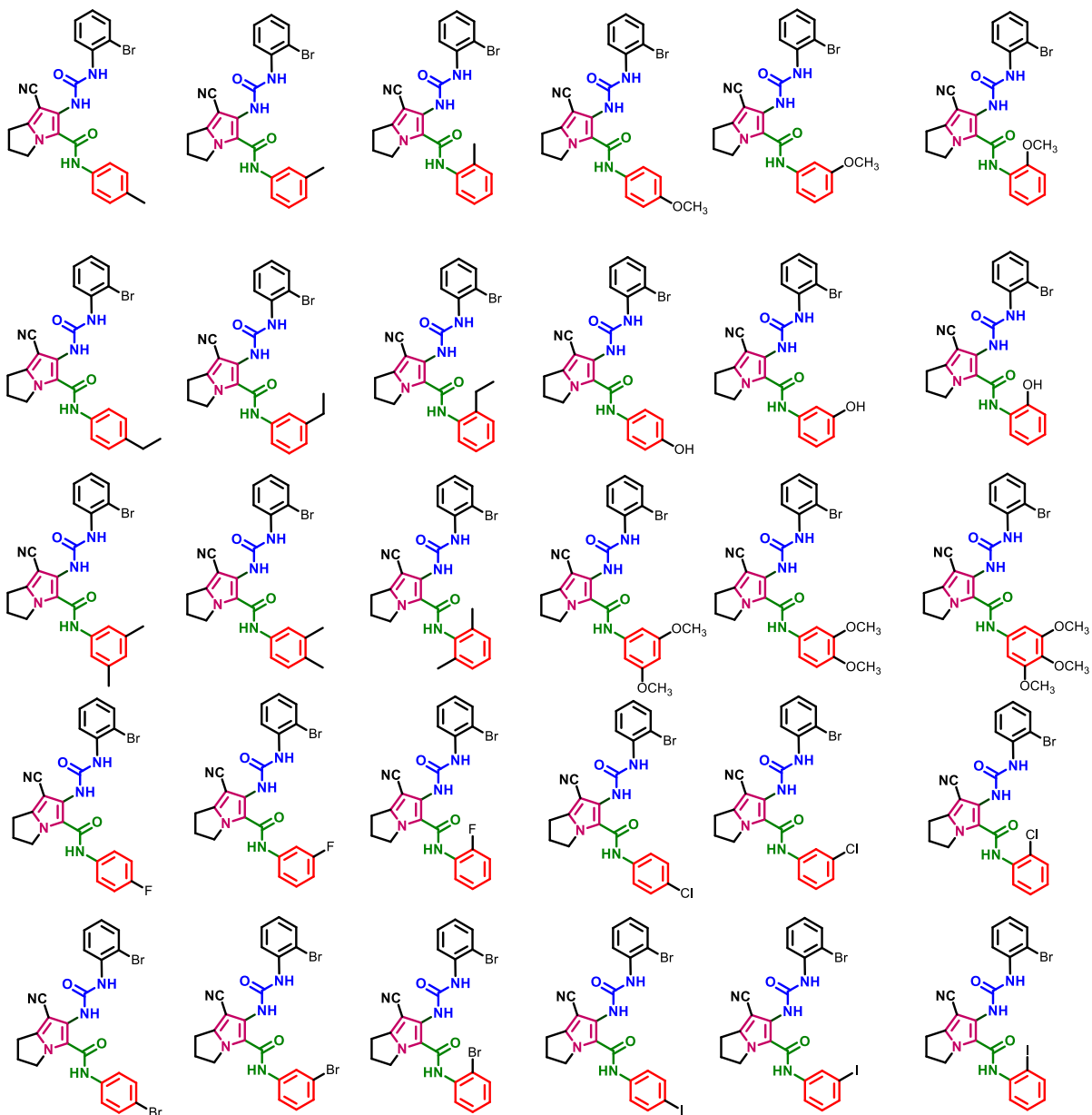
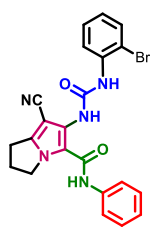


Figure S20. Chemical structure of compound 590-620 (2bromophenyliso)

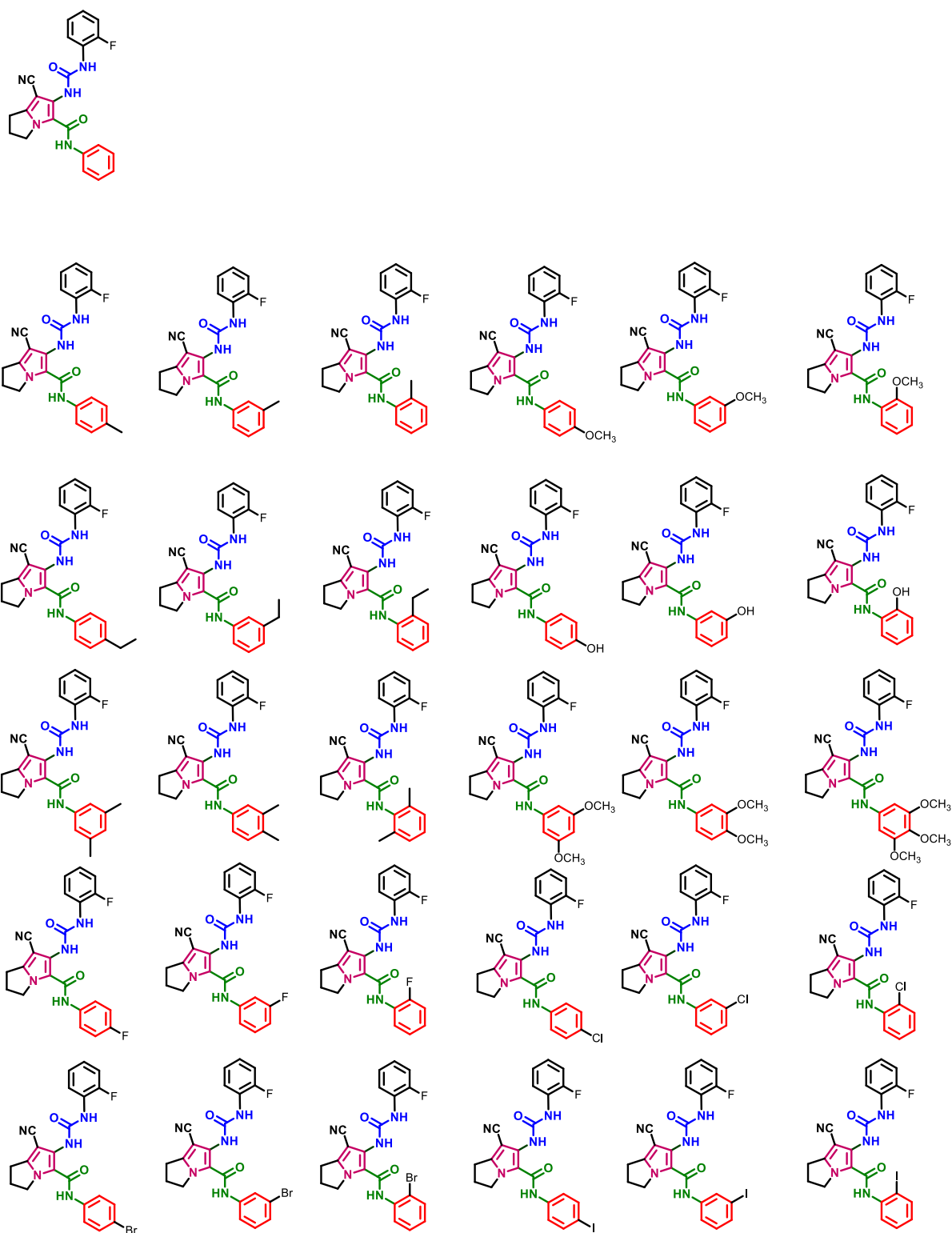


Figure B21. Chemical structure of compound 621-651 (2fluorophenyliso)

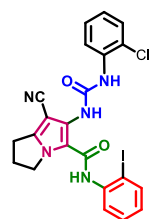
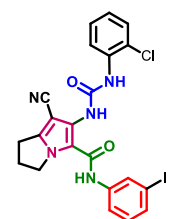
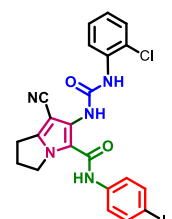
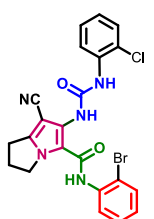
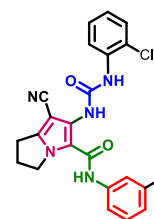
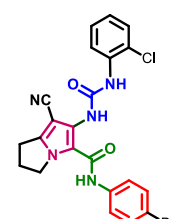
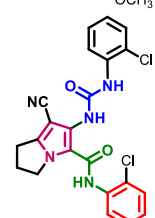
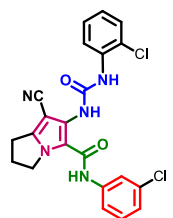
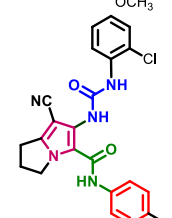
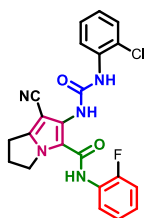
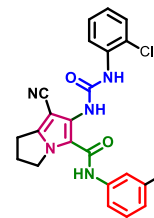
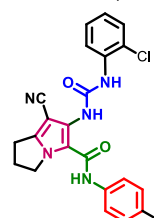
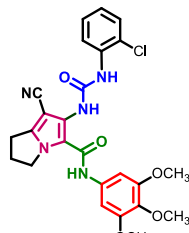
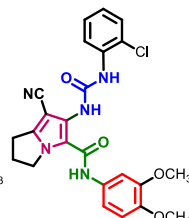
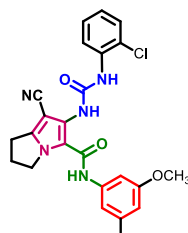
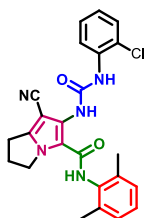
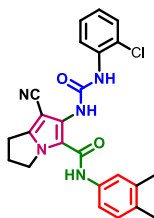
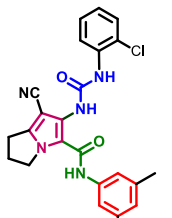
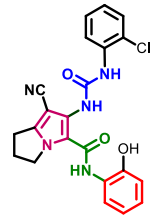
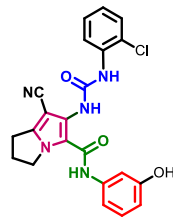
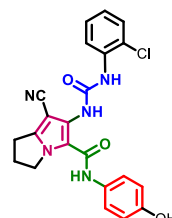
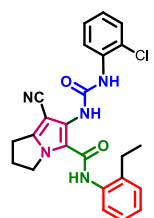
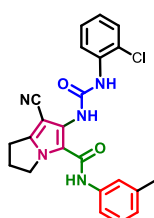
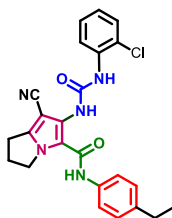
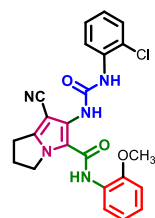
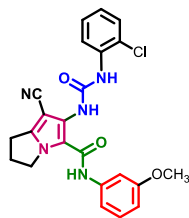
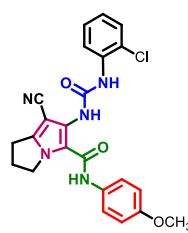
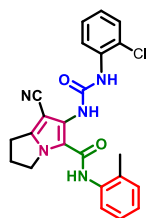
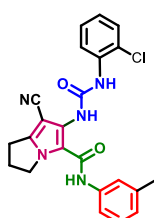
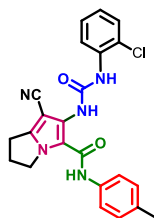
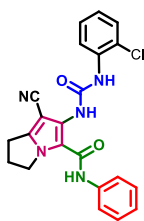


Figure S22. Chemical structure of compound 652-682 (2chlorophenyliso)

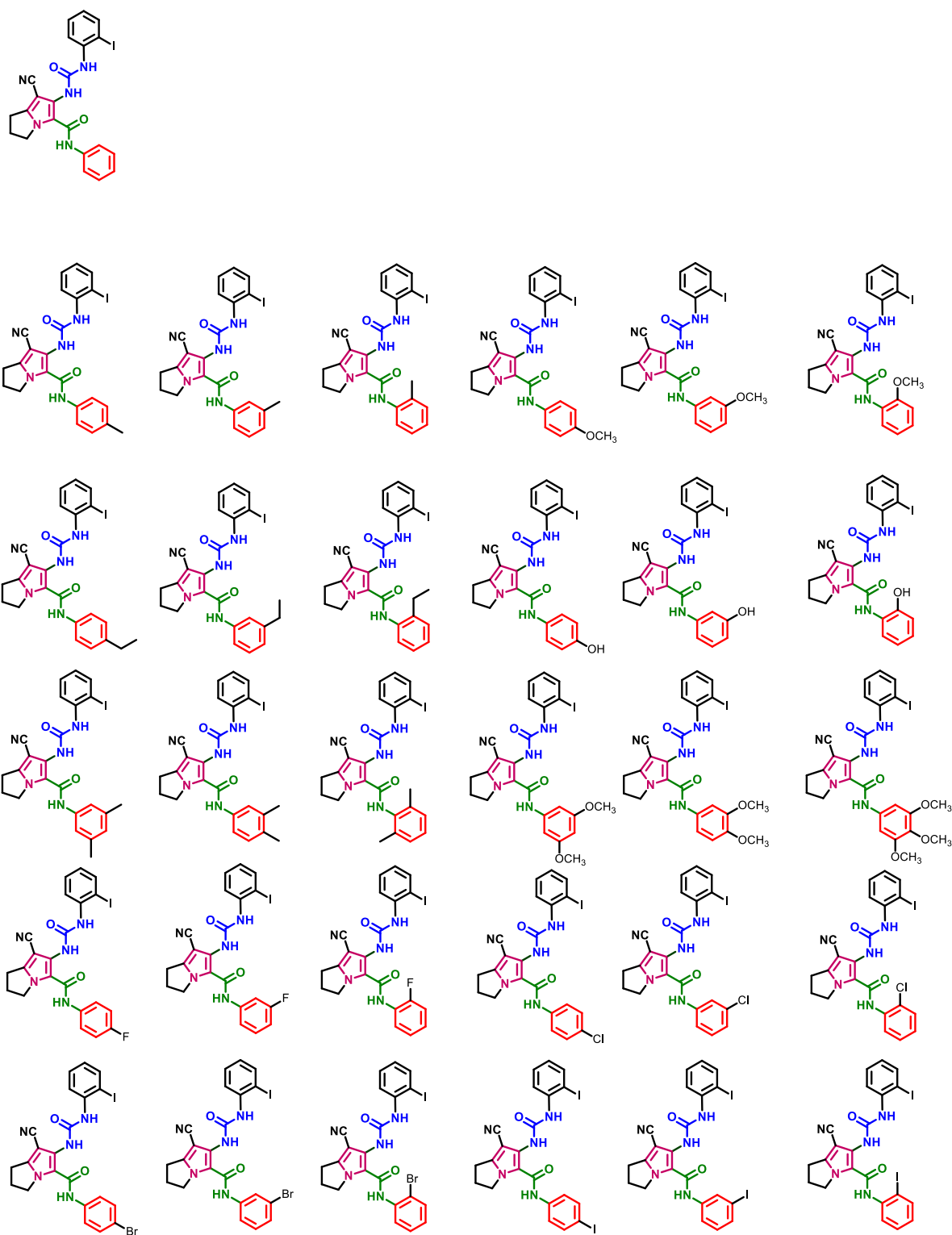


Figure S23. Chemical structure of compound 683-713 (2iodophenyliso)

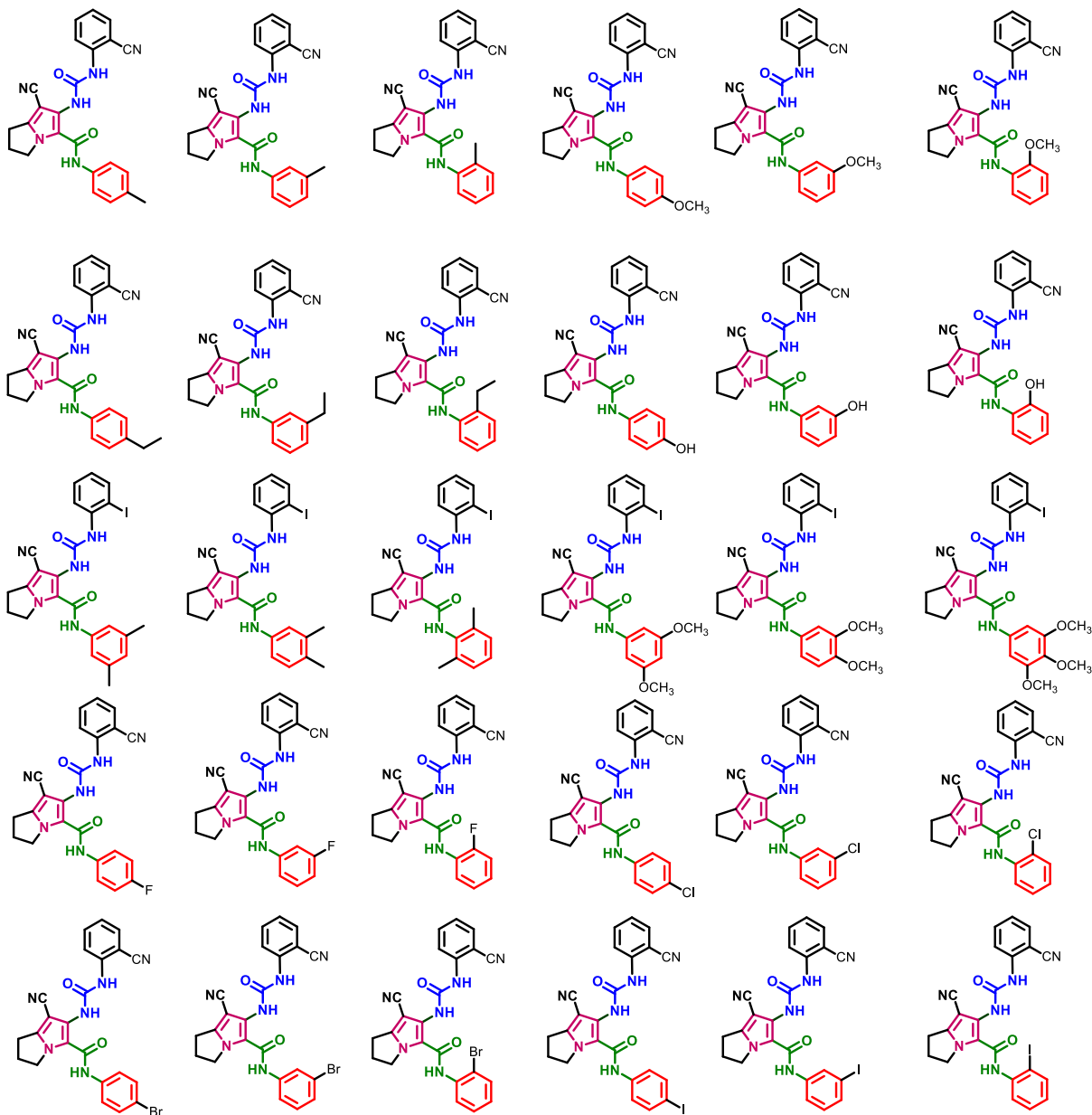
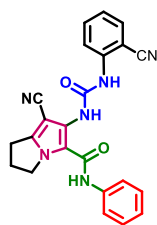


Figure S24. Chemical structure of compound 714-744 (2cyanophenyliso)

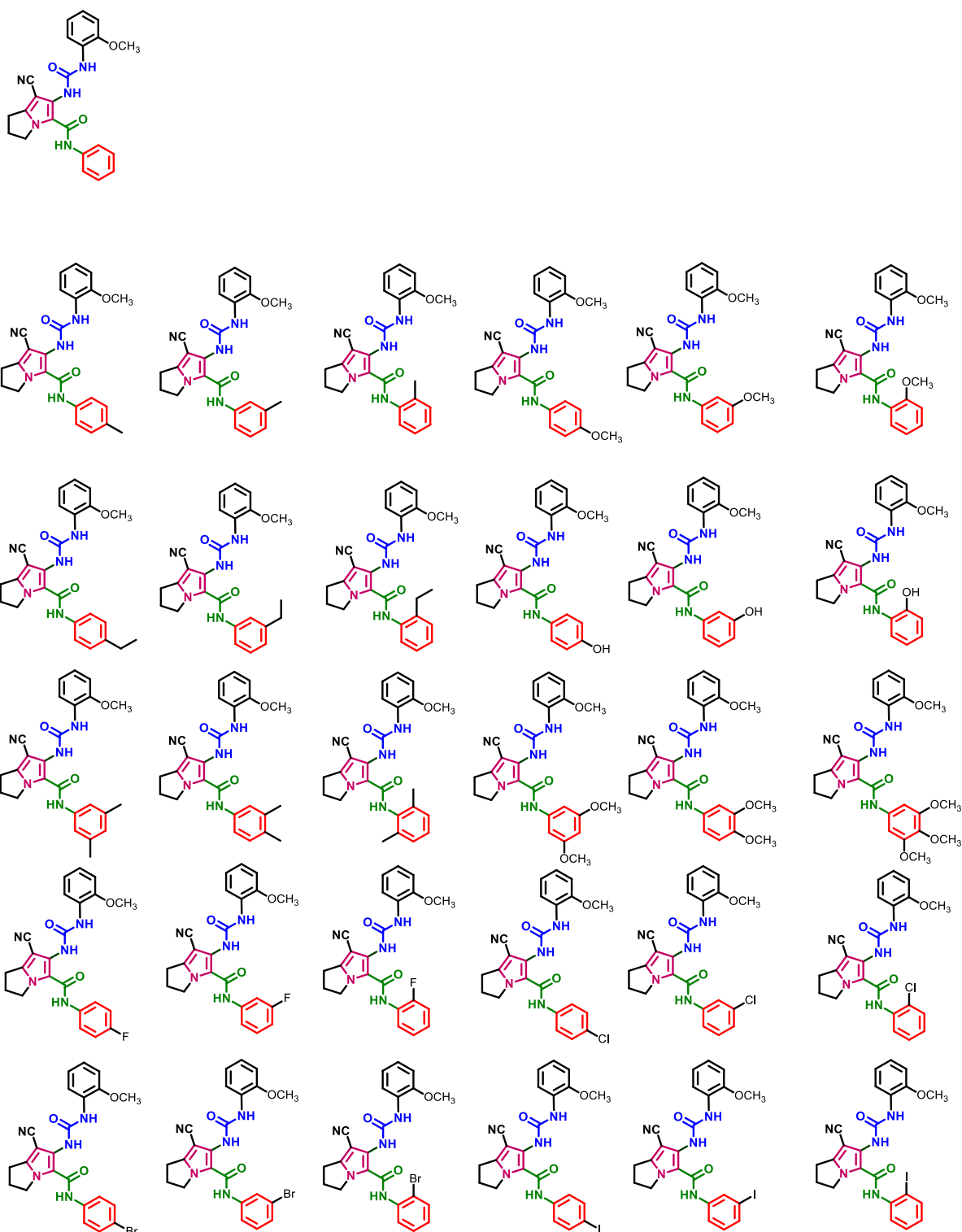


Figure S25. Chemical structure of compound 745-775 (2methoxyphenyliso)

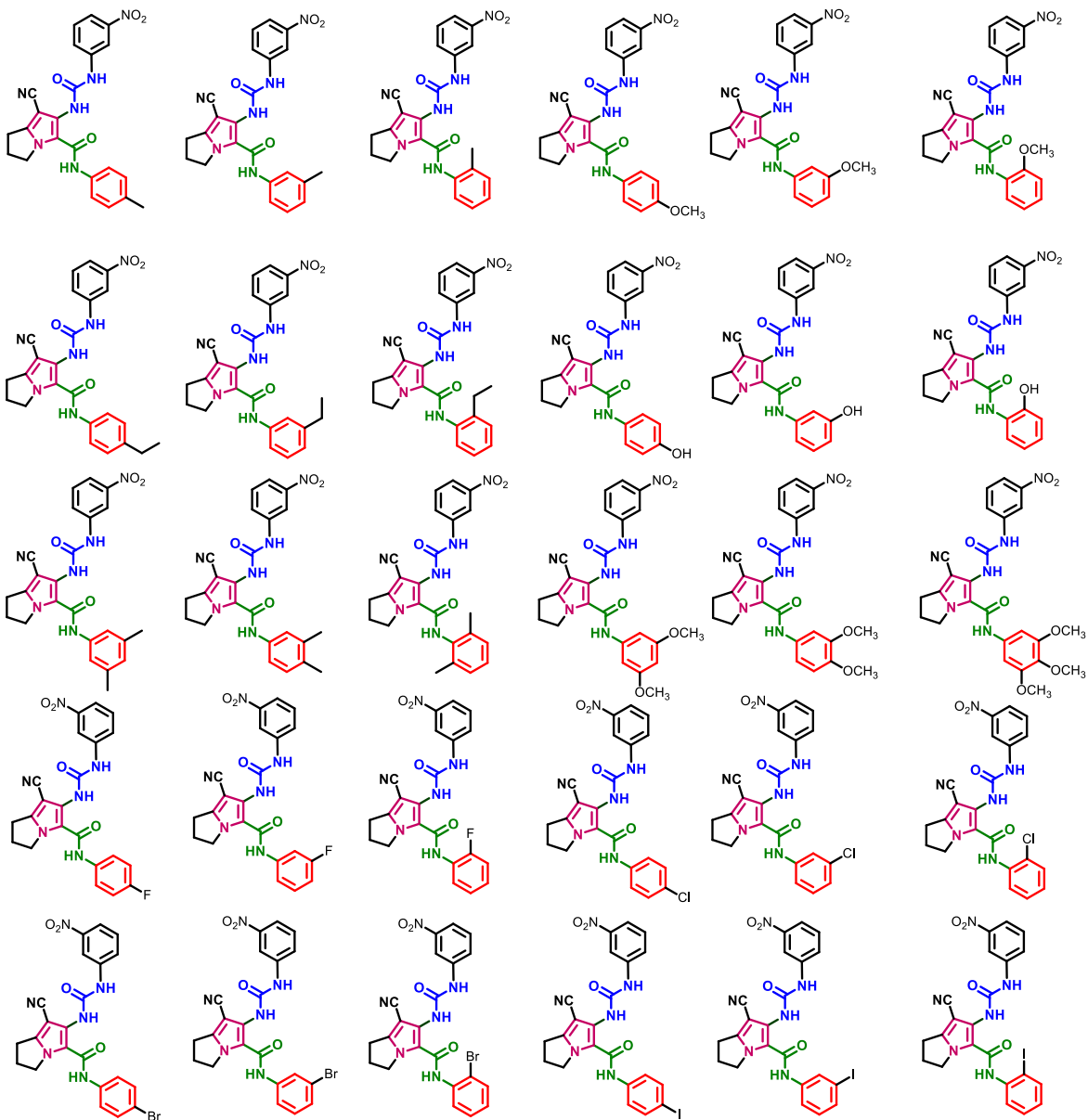
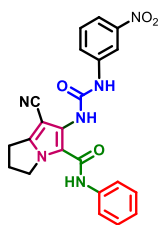


Figure S26. Chemical structure of compound 776-806 (3nitrophenyliso)

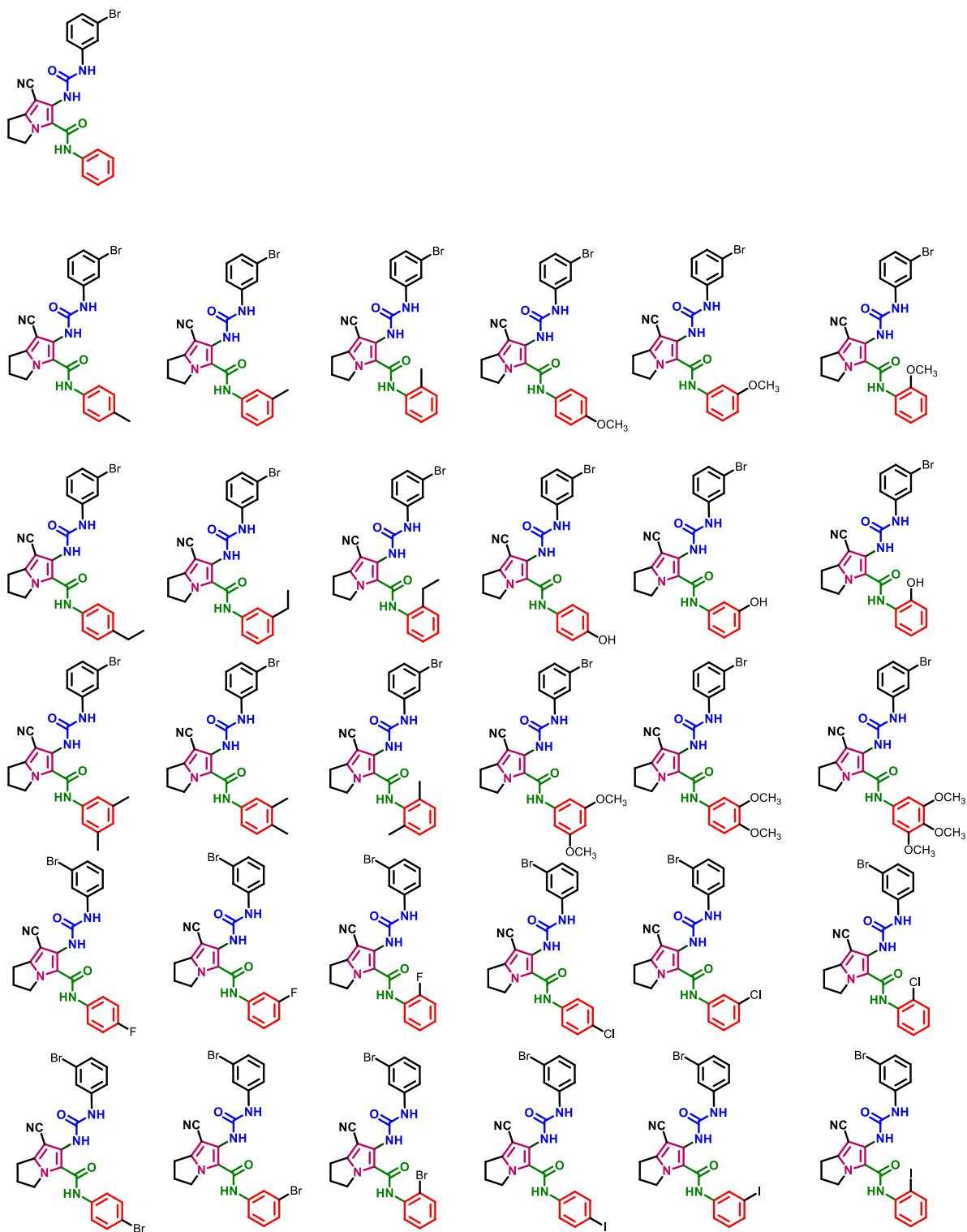


Figure S27. Chemical structure of compound 807-837 (3bromophenyliso)

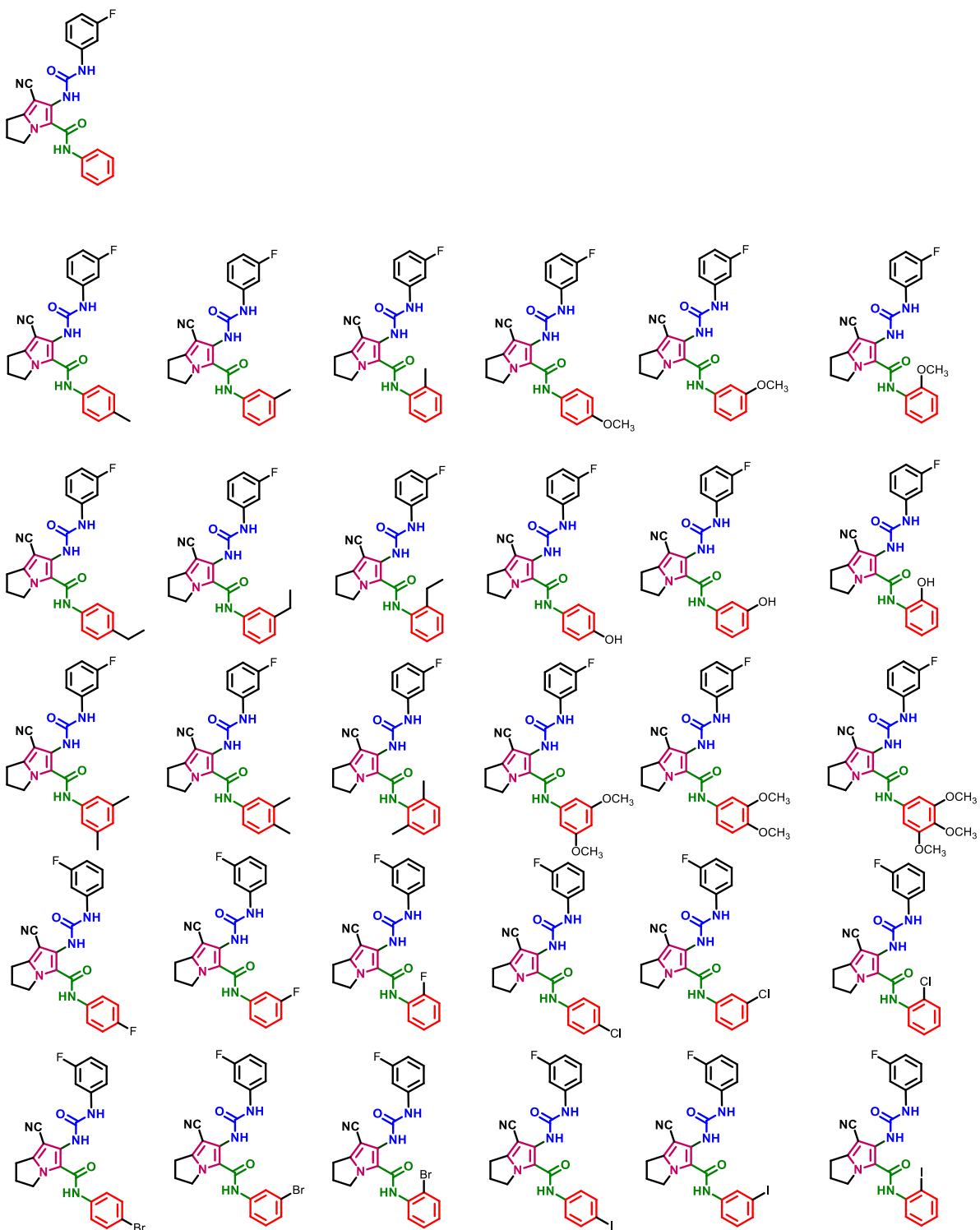


Figure S28. Chemical structure of compound 838-868 (3fluorophenyliso)

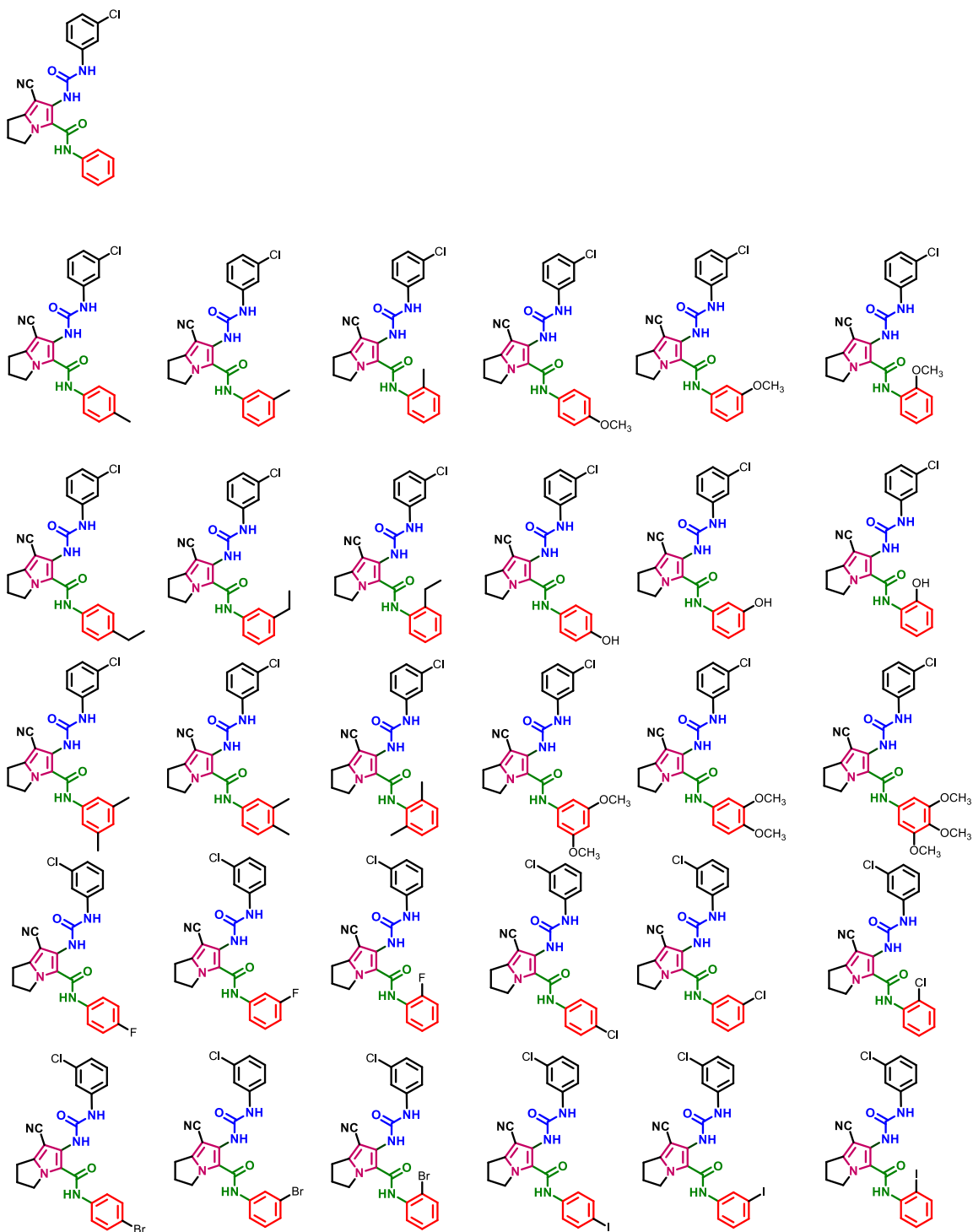


Figure S29. Chemical structure of compound 869-899 (3chlorophenyliso)

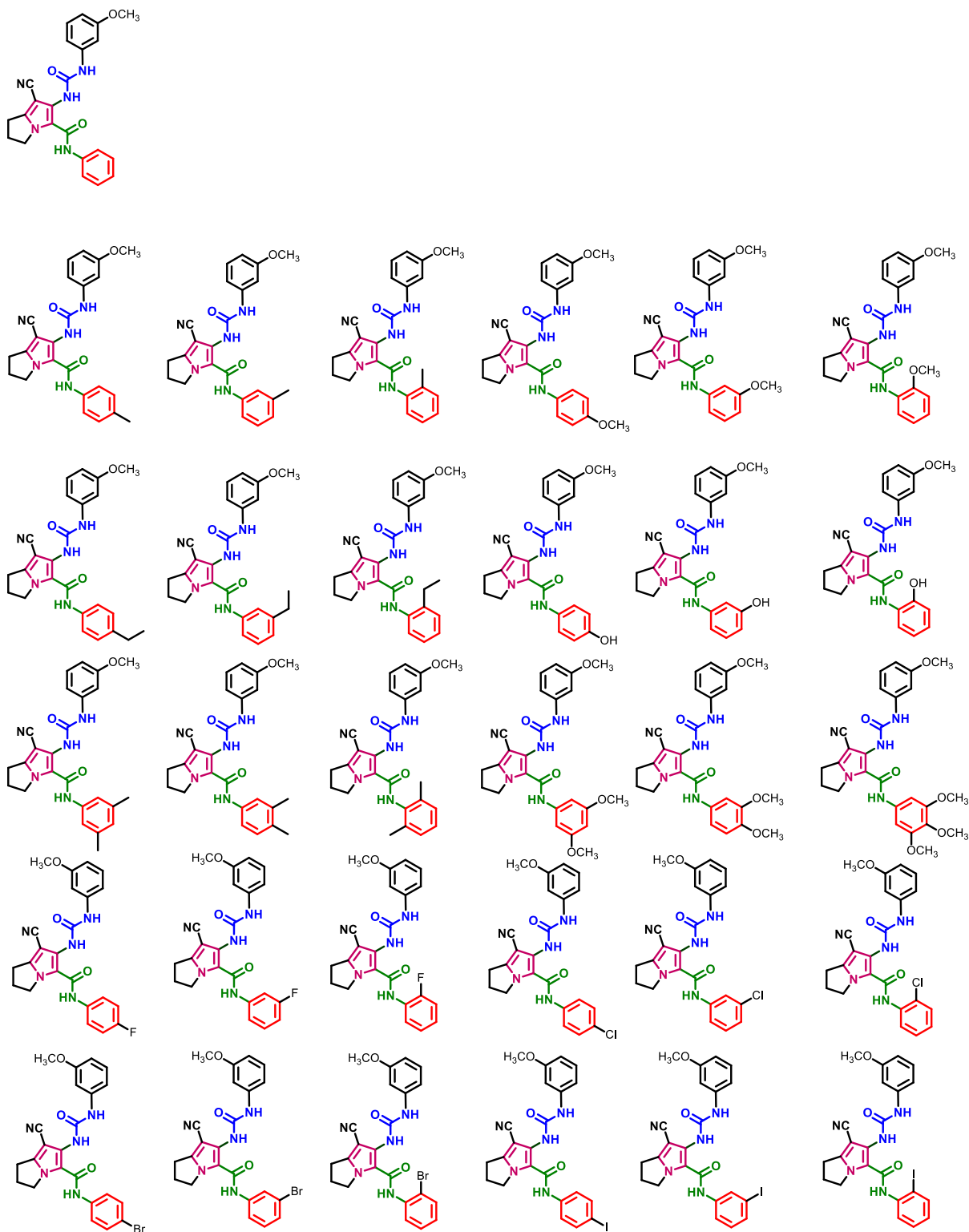


Figure S30. Chemical structure of compound 900-930 (3methoxyphenyliso)

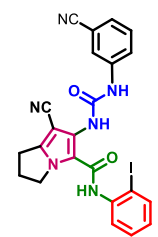
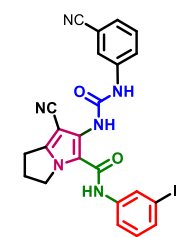
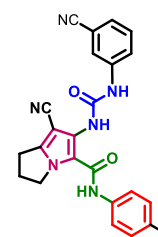
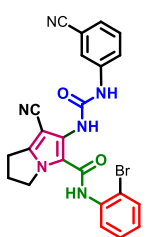
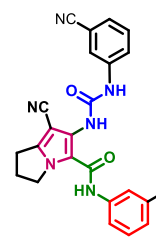
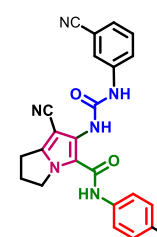
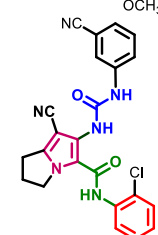
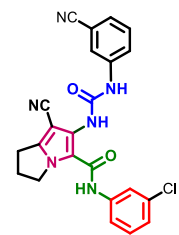
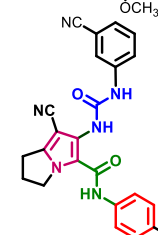
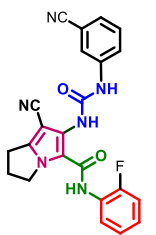
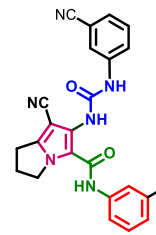
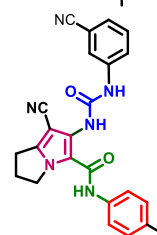
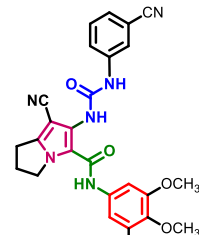
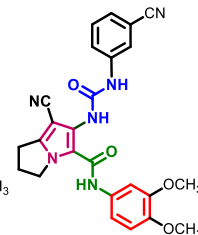
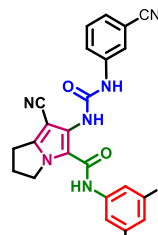
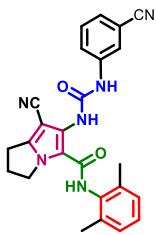
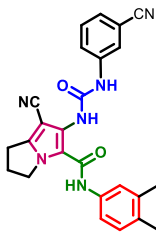
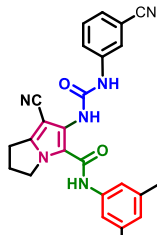
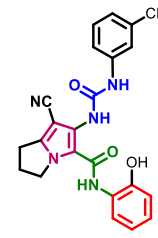
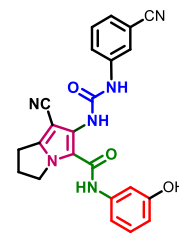
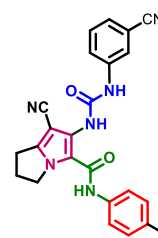
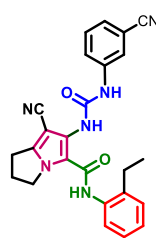
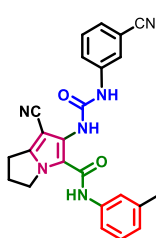
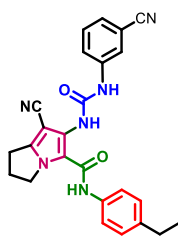
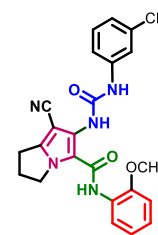
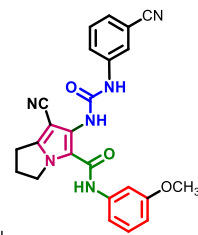
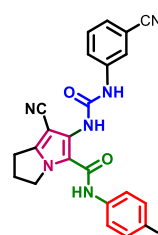
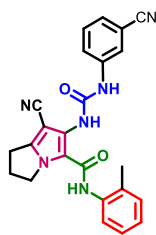
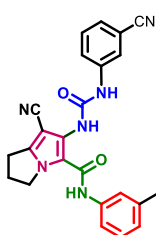
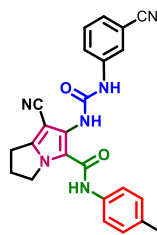
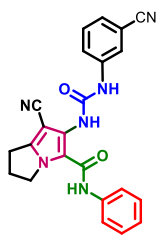


Figure S31. Chemical structure of compound 931-961 (3cyanophenyliso)

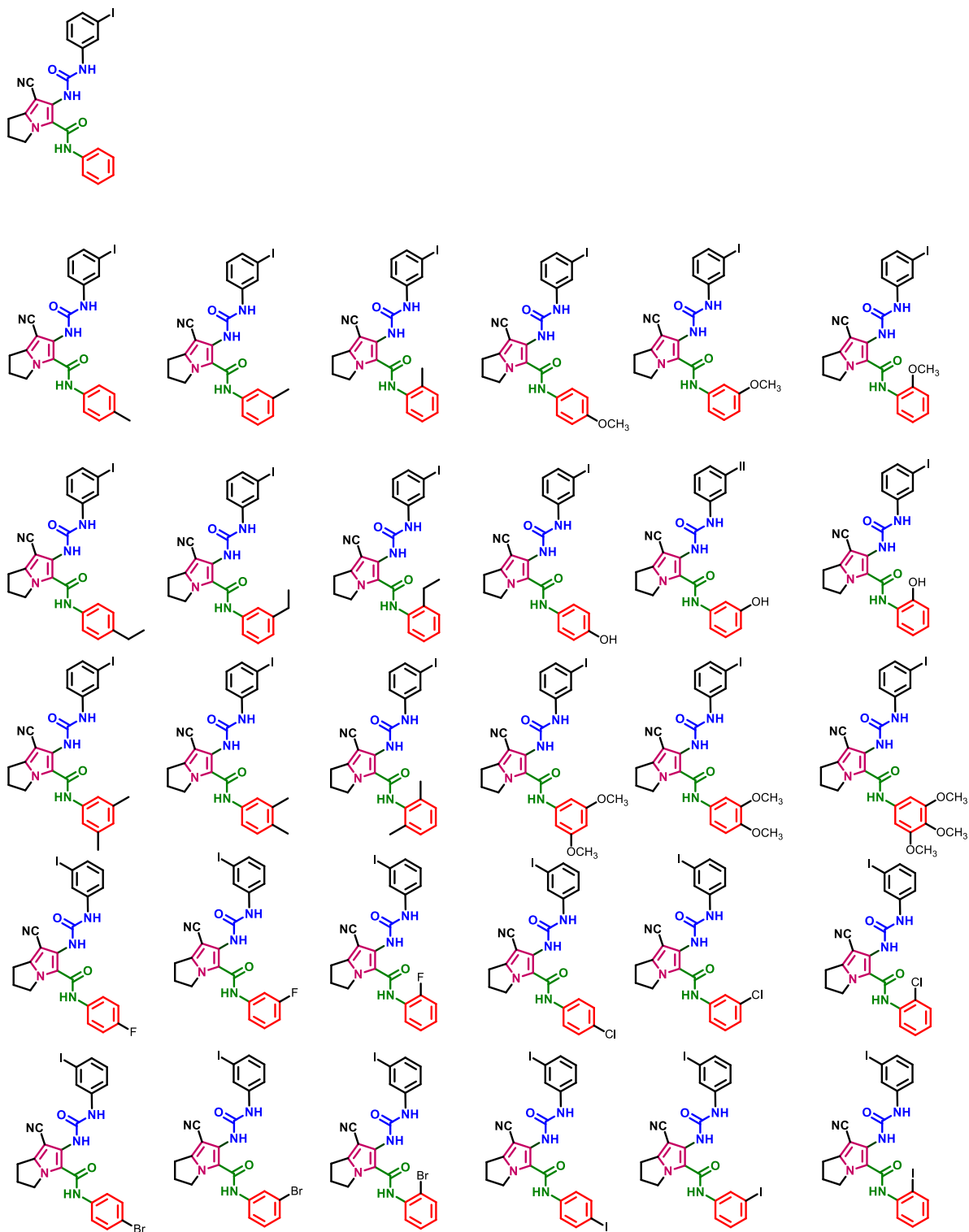


Figure S32. Chemical structure of compound 962-992 (3iodophenyliso)

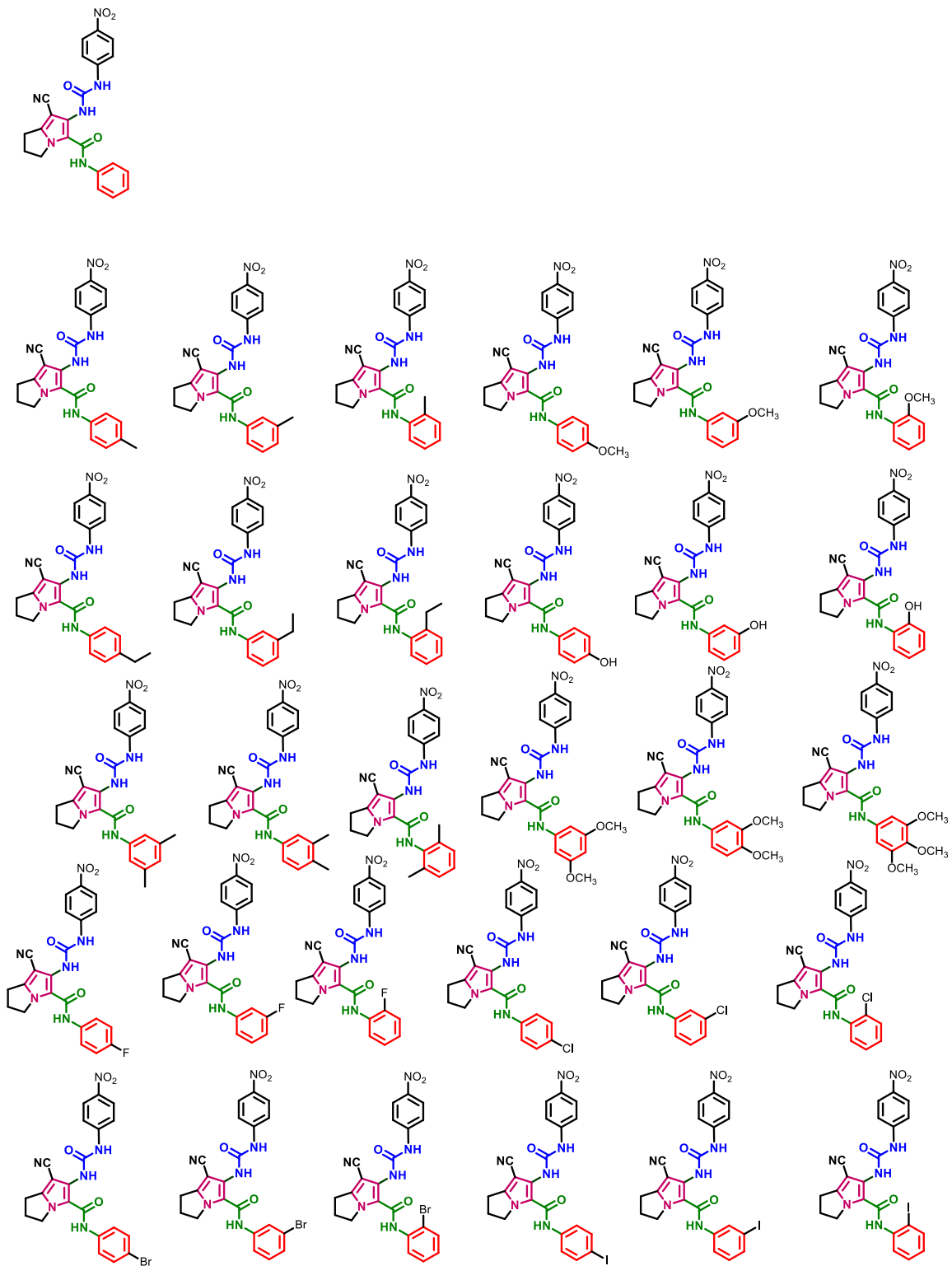


Figure S33. Chemical structure of compound 993-1023 (4nitrophenyliso)

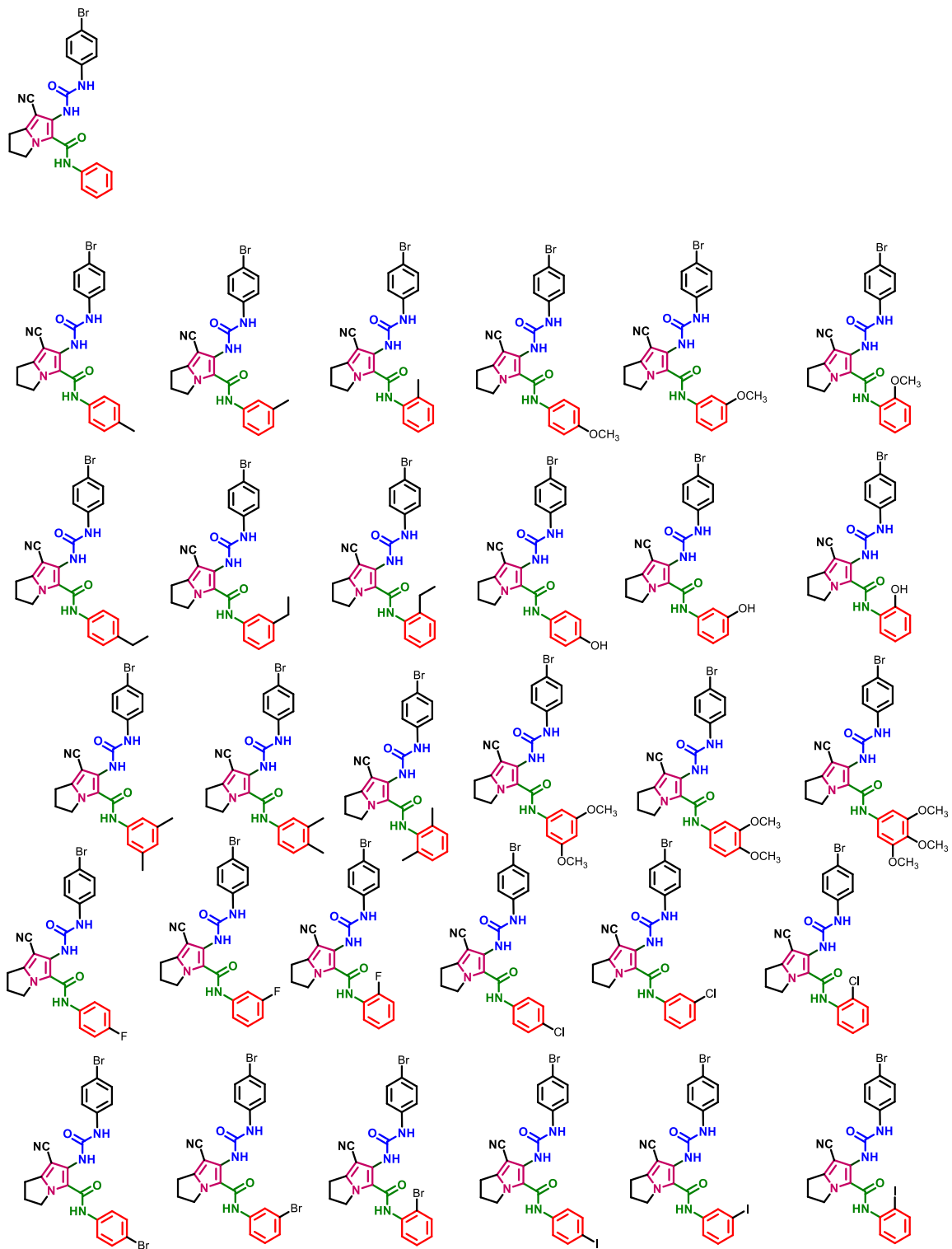


Figure S34. Chemical structure of compound 1024-1054 (4bromophenyliso)

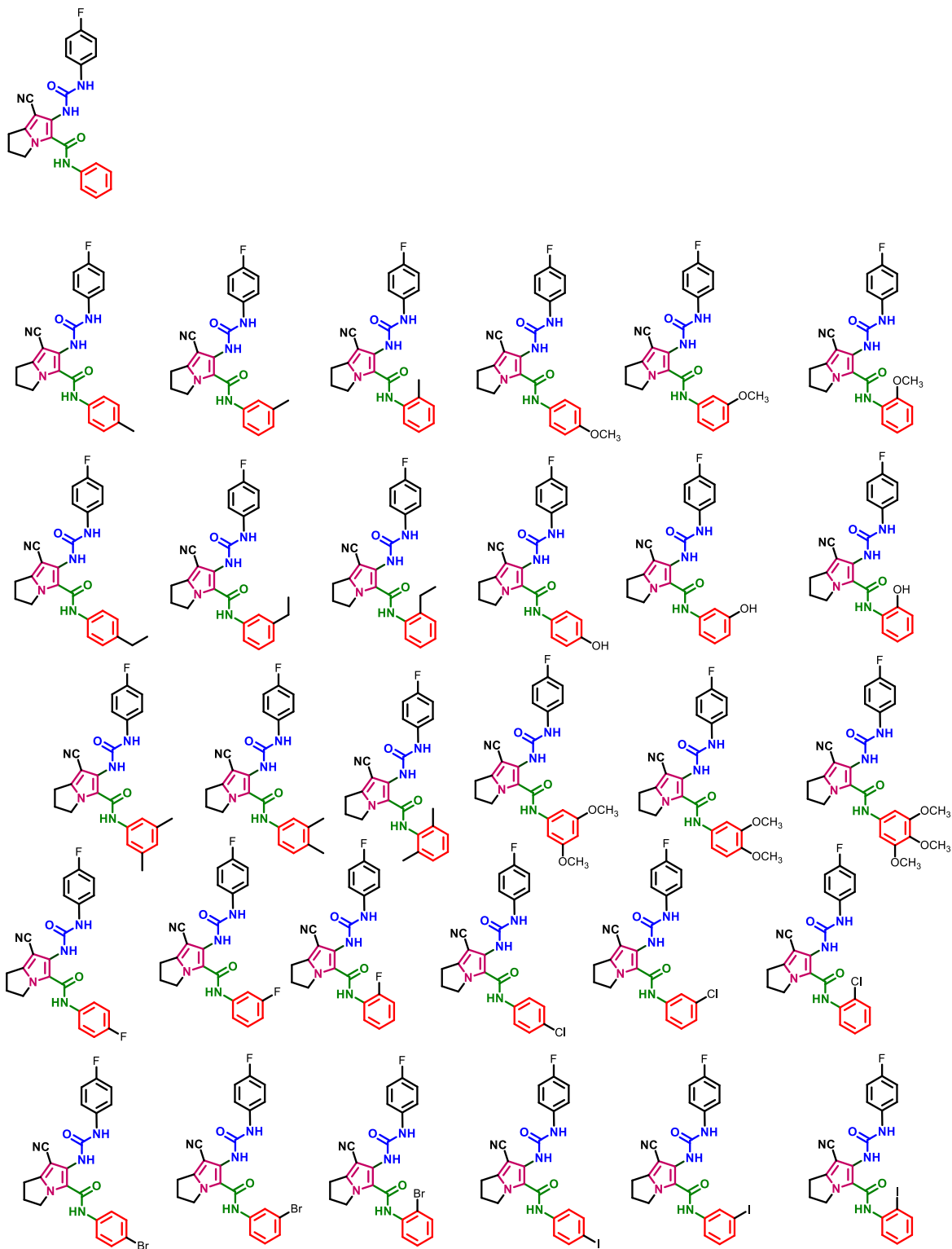


Figure S35. Chemical structure of compound 1055-1085 (4fluorophenyliso)

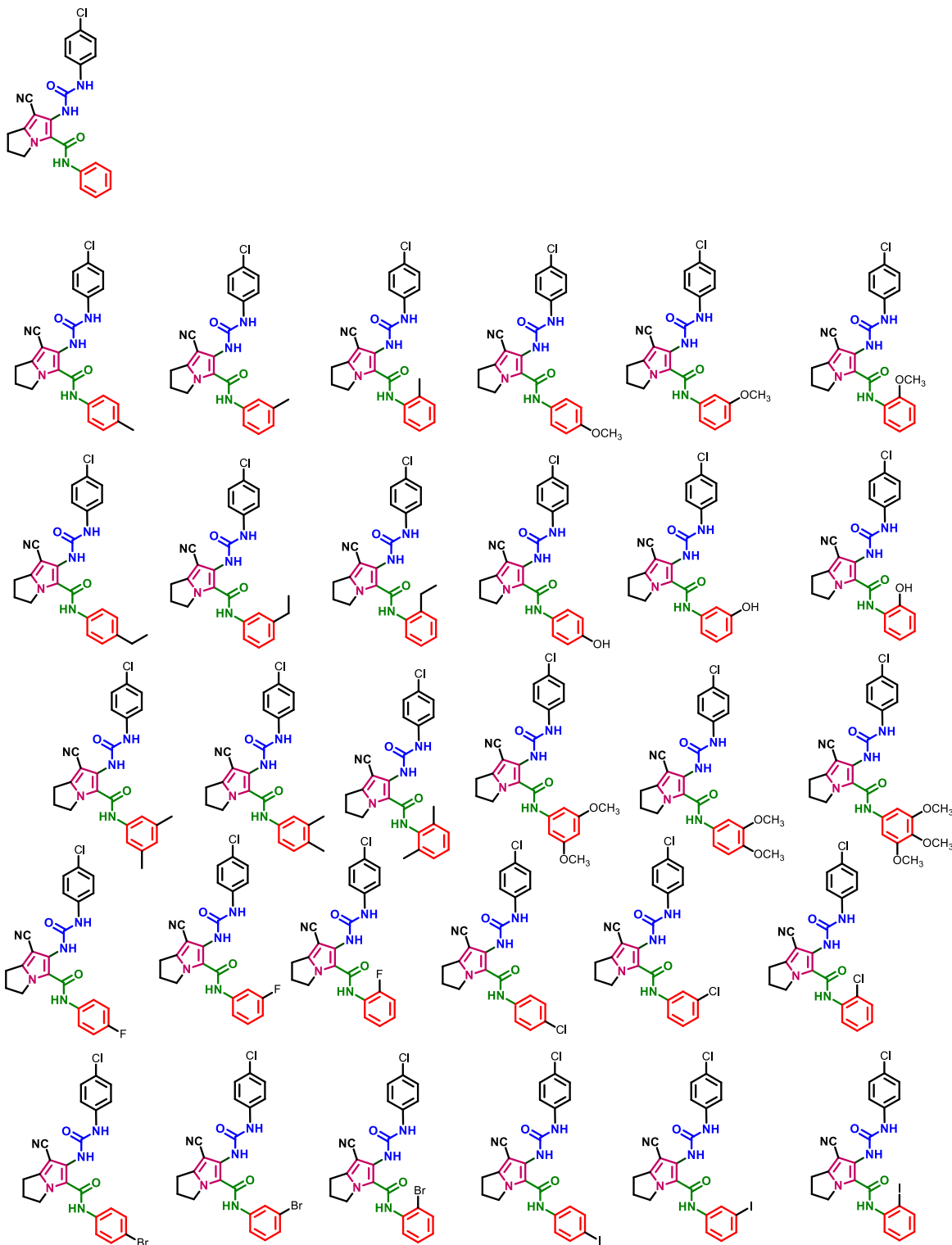


Figure S36. Chemical structure of compound 1086-1116 (4chlorophenyliso)

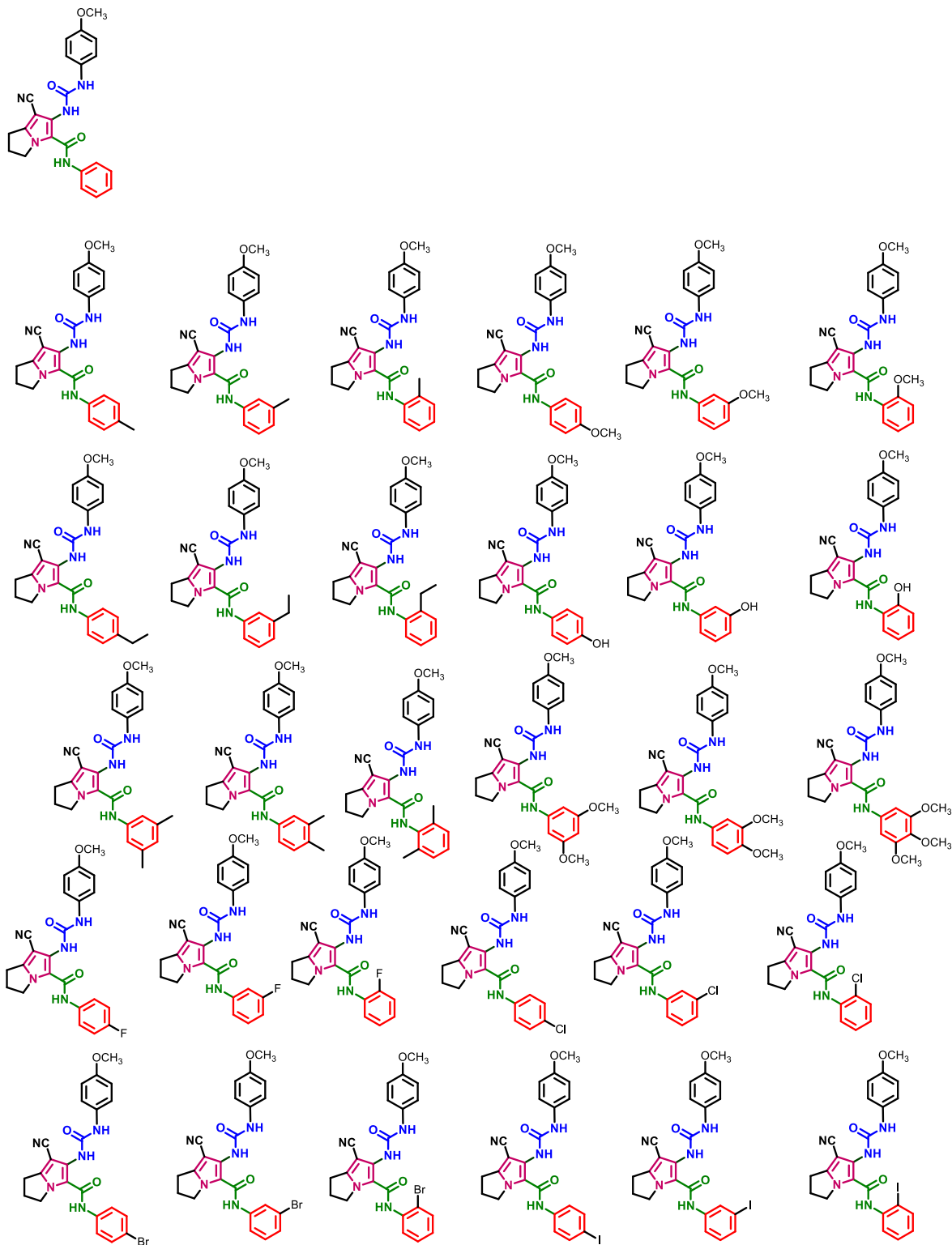


Figure S37. Chemical structure of compound 1117-1147 (4methoxyphenyliso)

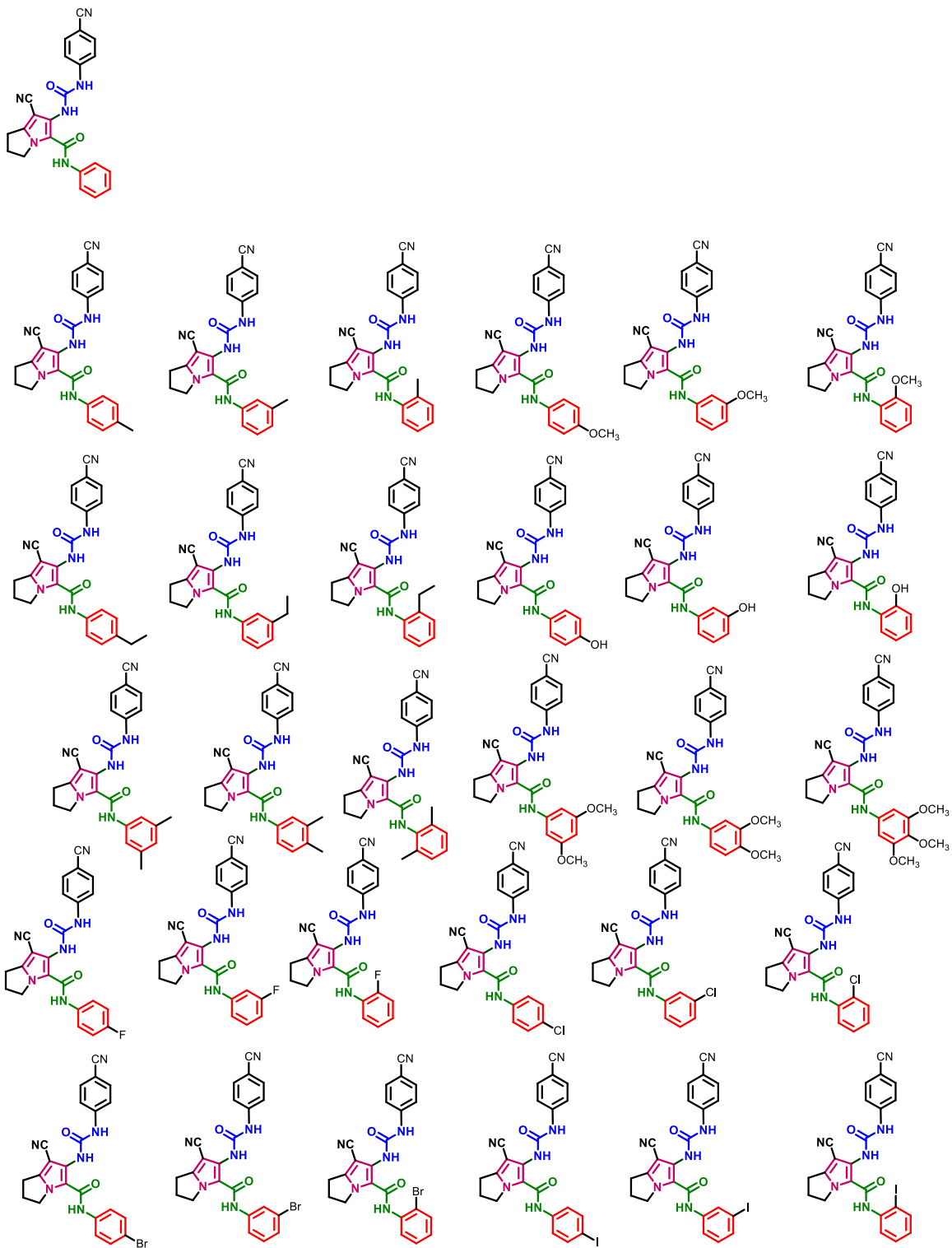


Figure S38. Chemical structure of compound 1148-1178 (4cyanophenyliso)

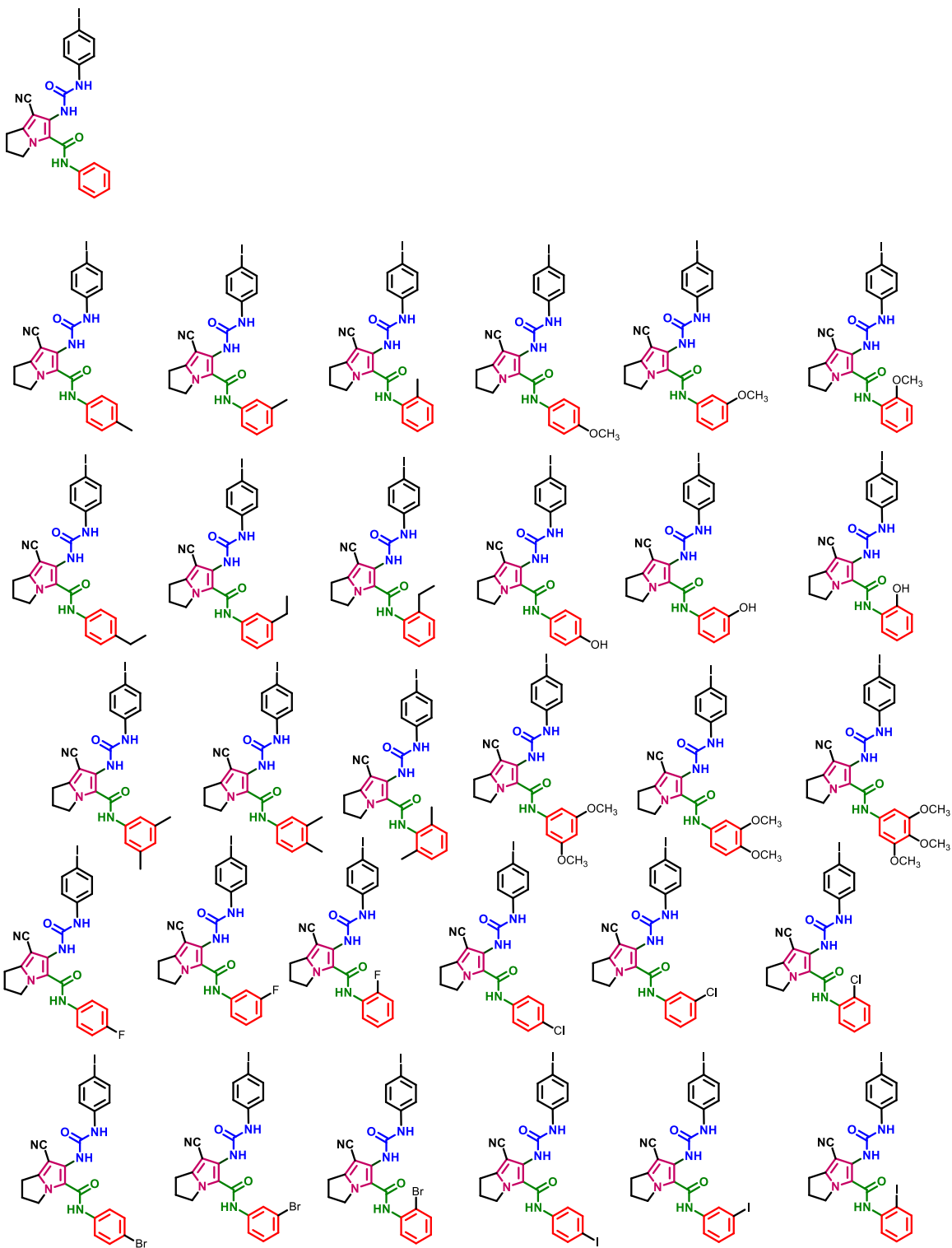


Figure S39. Chemical structure of compound 1179-1209 (4iodophenyliso)

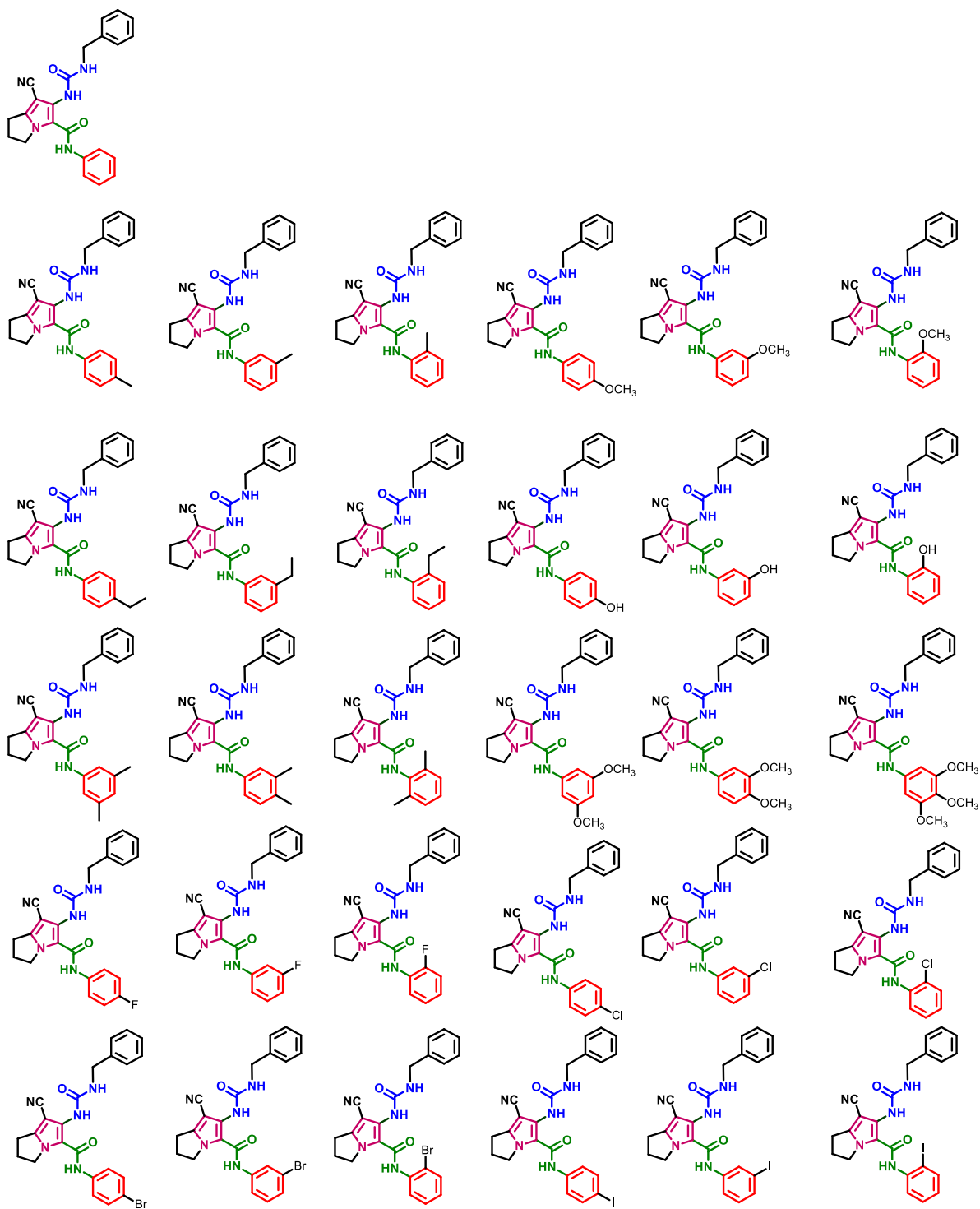


Figure S40. Chemical structure of compound 1210-1240 (benzyliso)

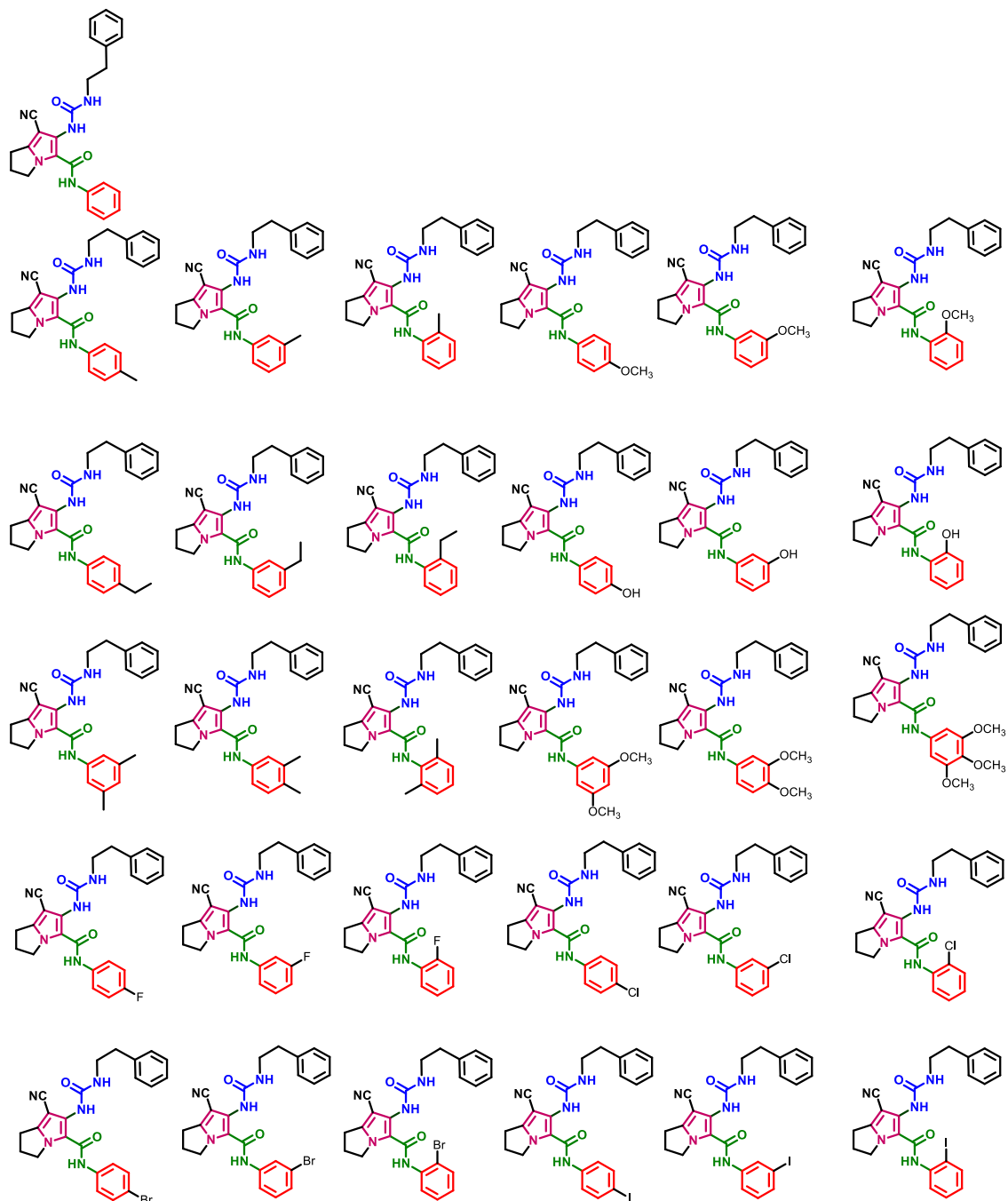


Figure S41. Chemical structure of compound 1241-1271 (phenylethyl)

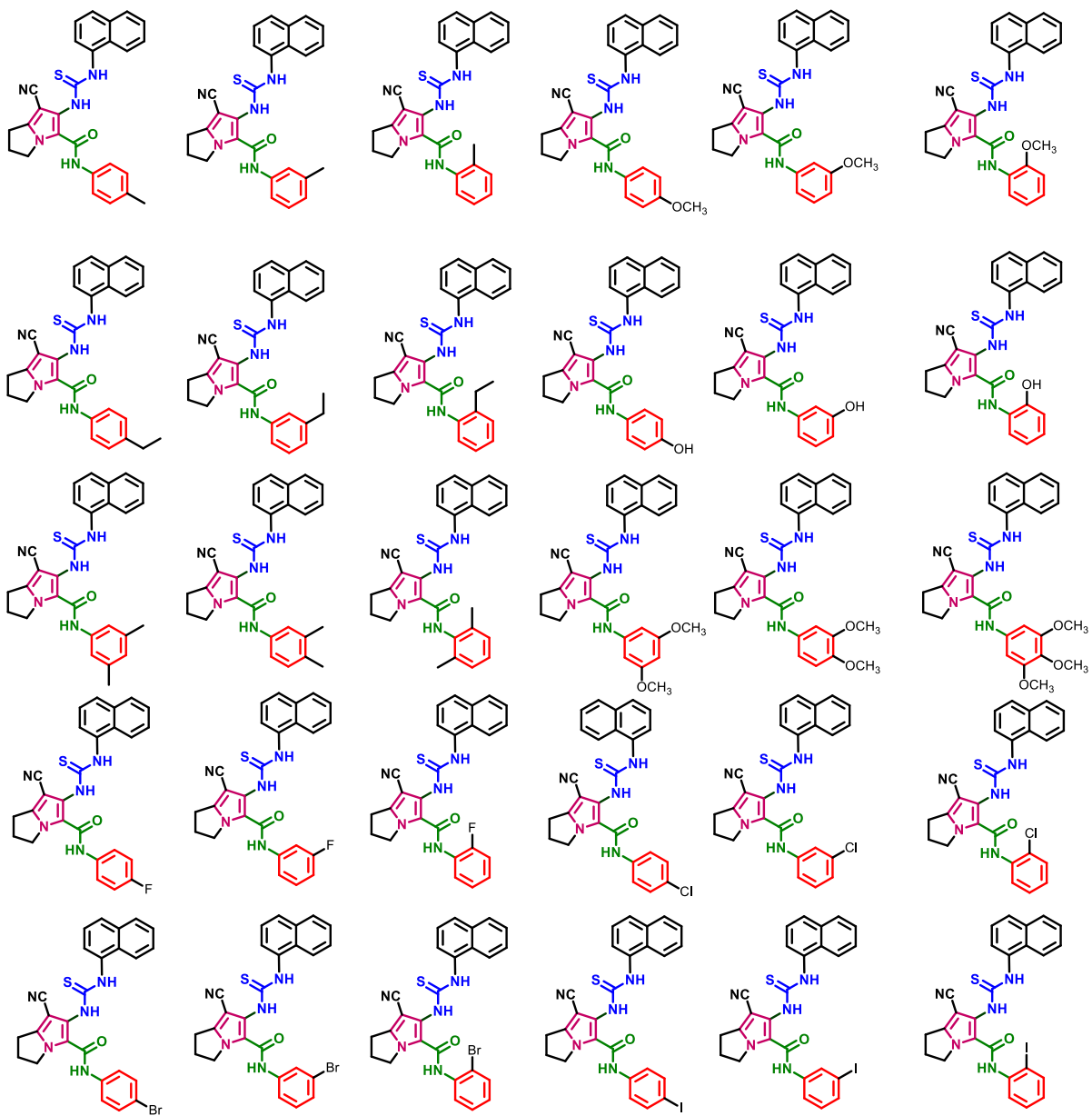
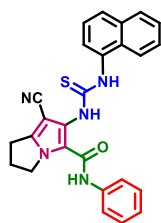
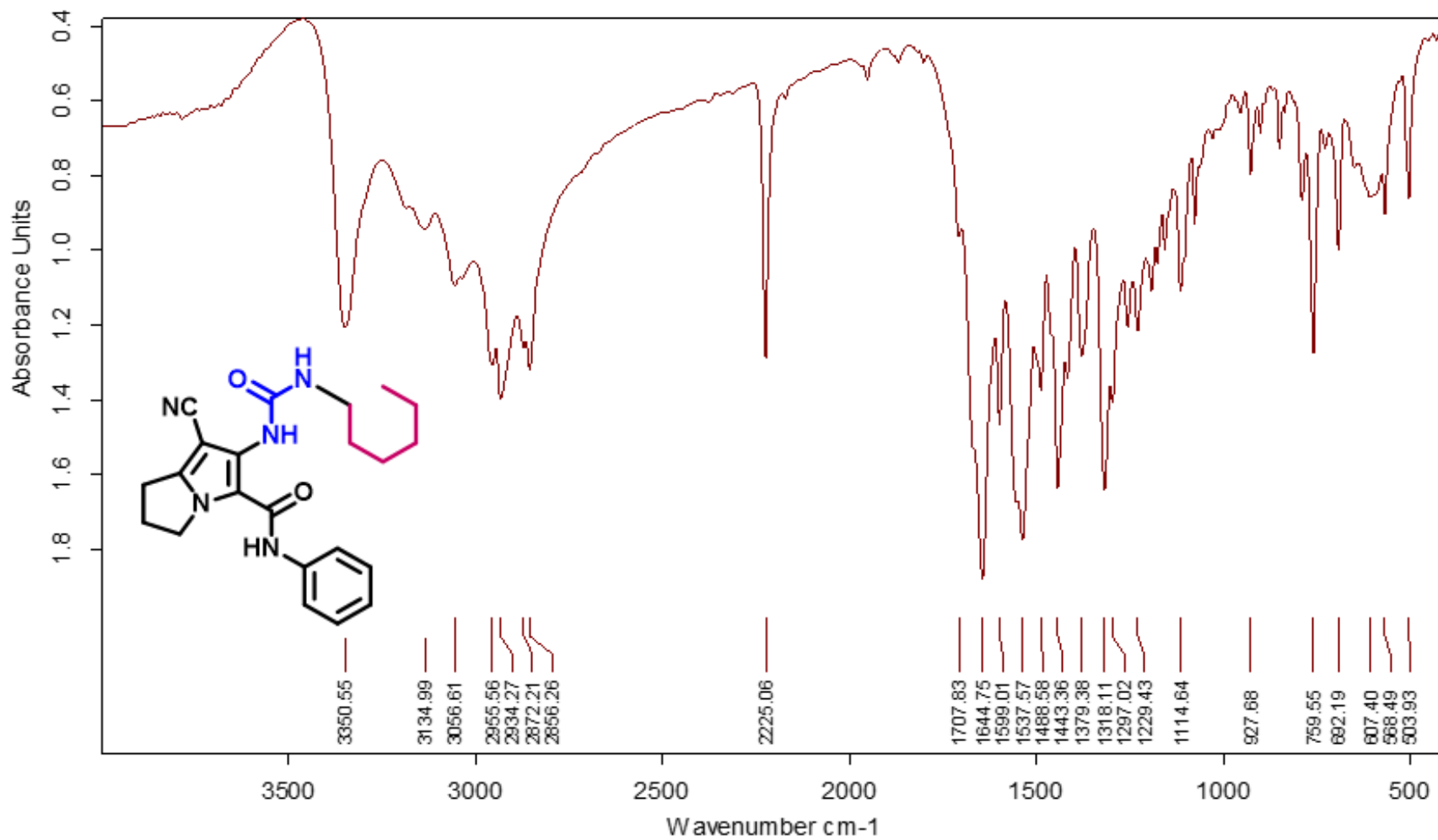


Figure S42. Chemical structure of compound 1272-1302 (phenylethyl)

IR Spectra

Infrared spectra (IR) were done using BRUKER TENSOR 37 spectrophotometer and absorption were expressed in wave number (cm^{-1}) using KBr Disk.

Figure S43. IR spectrum of compound 16a



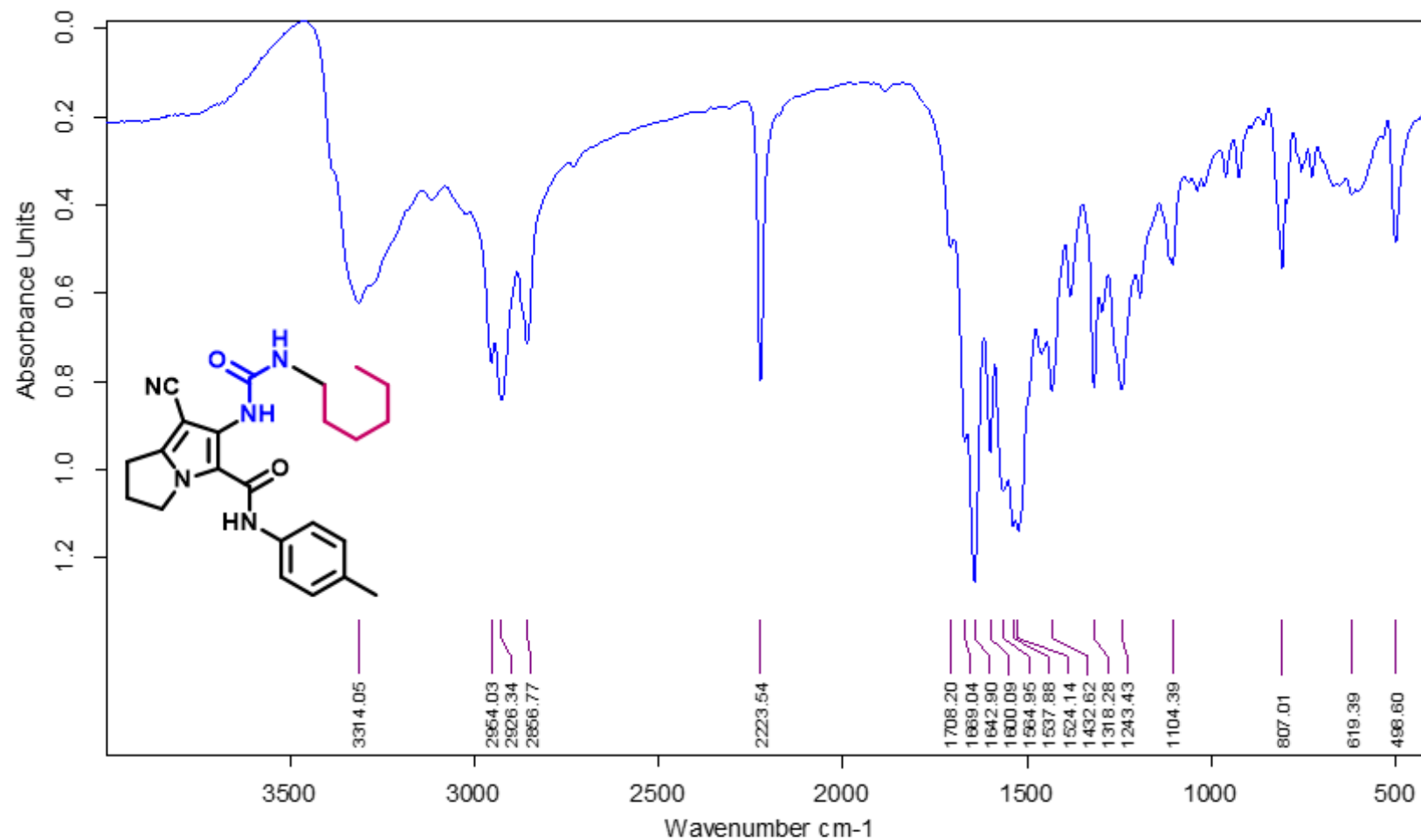
C:\Program Files\OPUS_65MEAS\Protein.247

Protein

AquaSpec

31/01/2018

Figure S44. IR spectrum of compound 16b



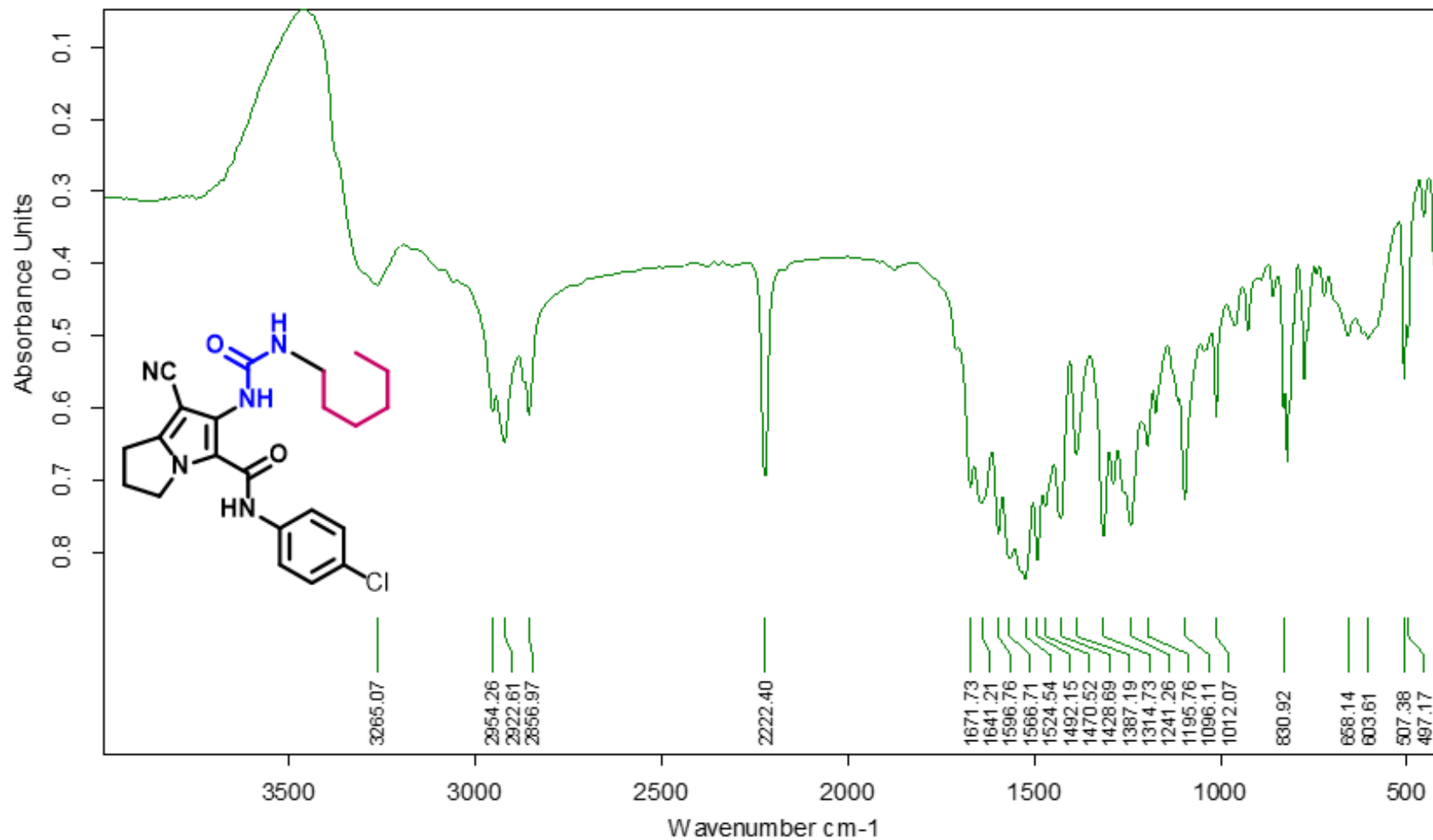
C:\Program Files\OPUS_85\MEAS\Protein.248

Protein

AquaSpec

31/01/2018

Figure S45. IR spectrum of compound 16c



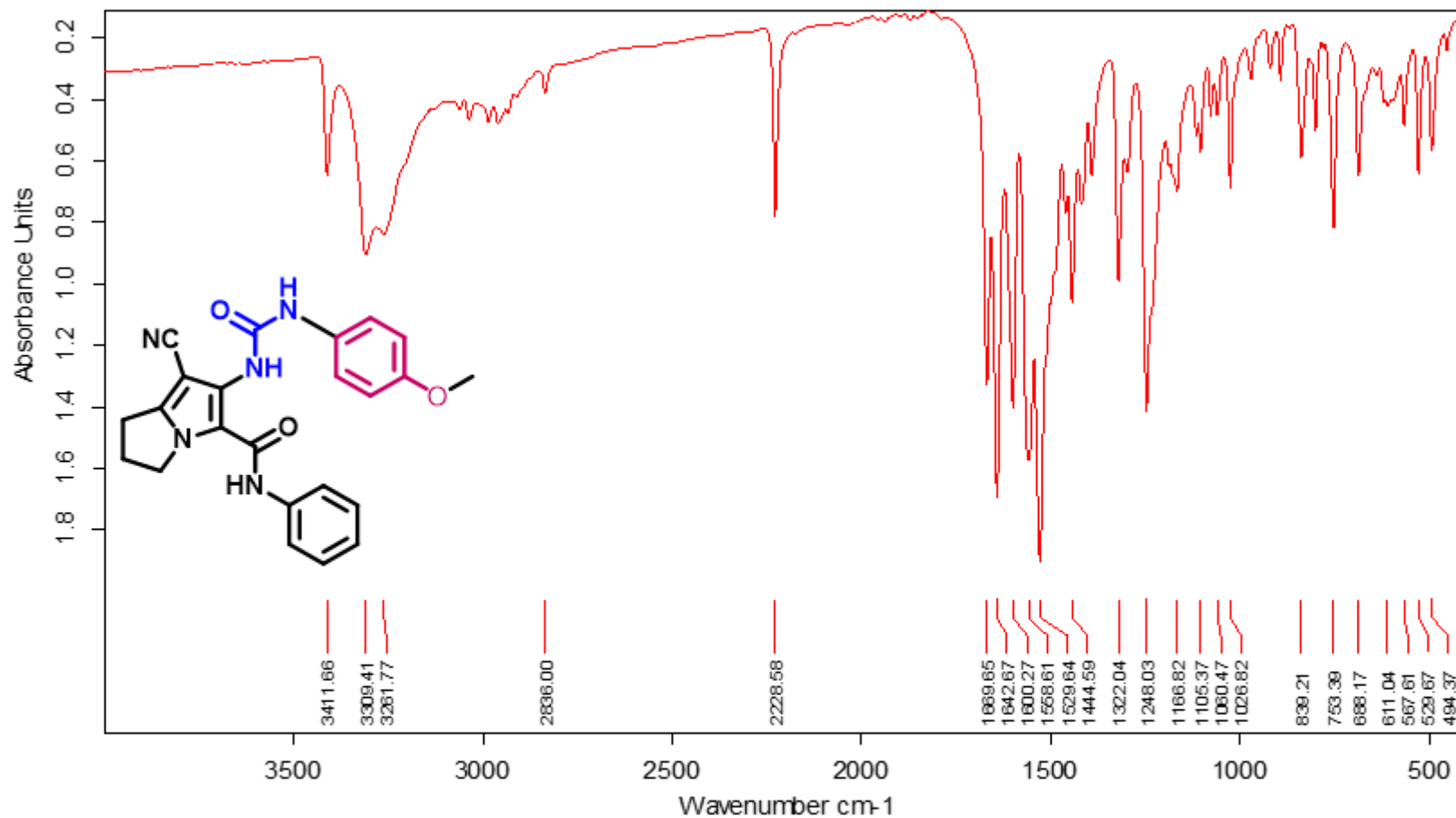
C:\Program Files\OPUS_65MEAS\Protein.249

Protein

AquaSpec

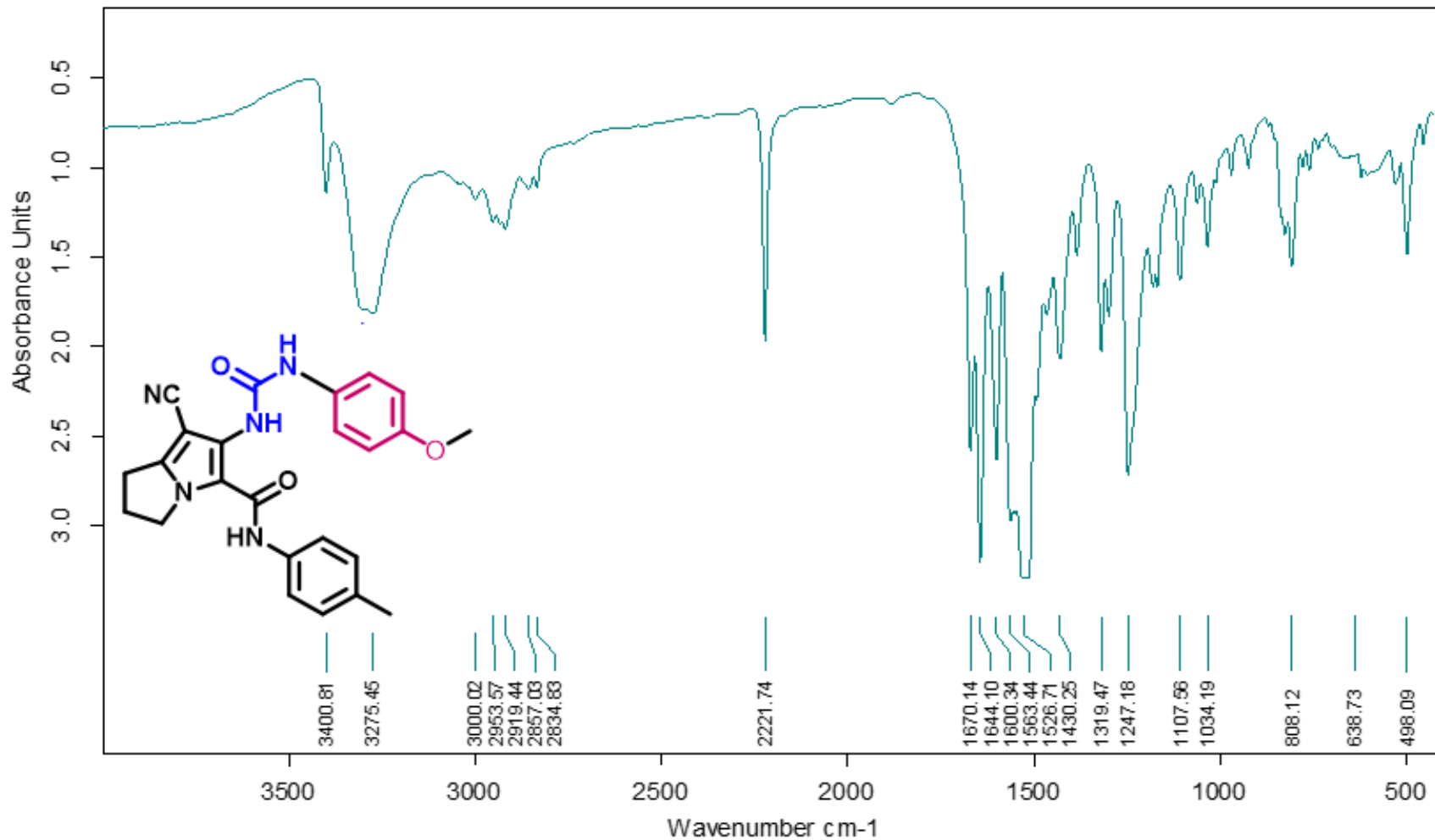
31/01/2018

Figure S46. IR spectrum of compound 17a



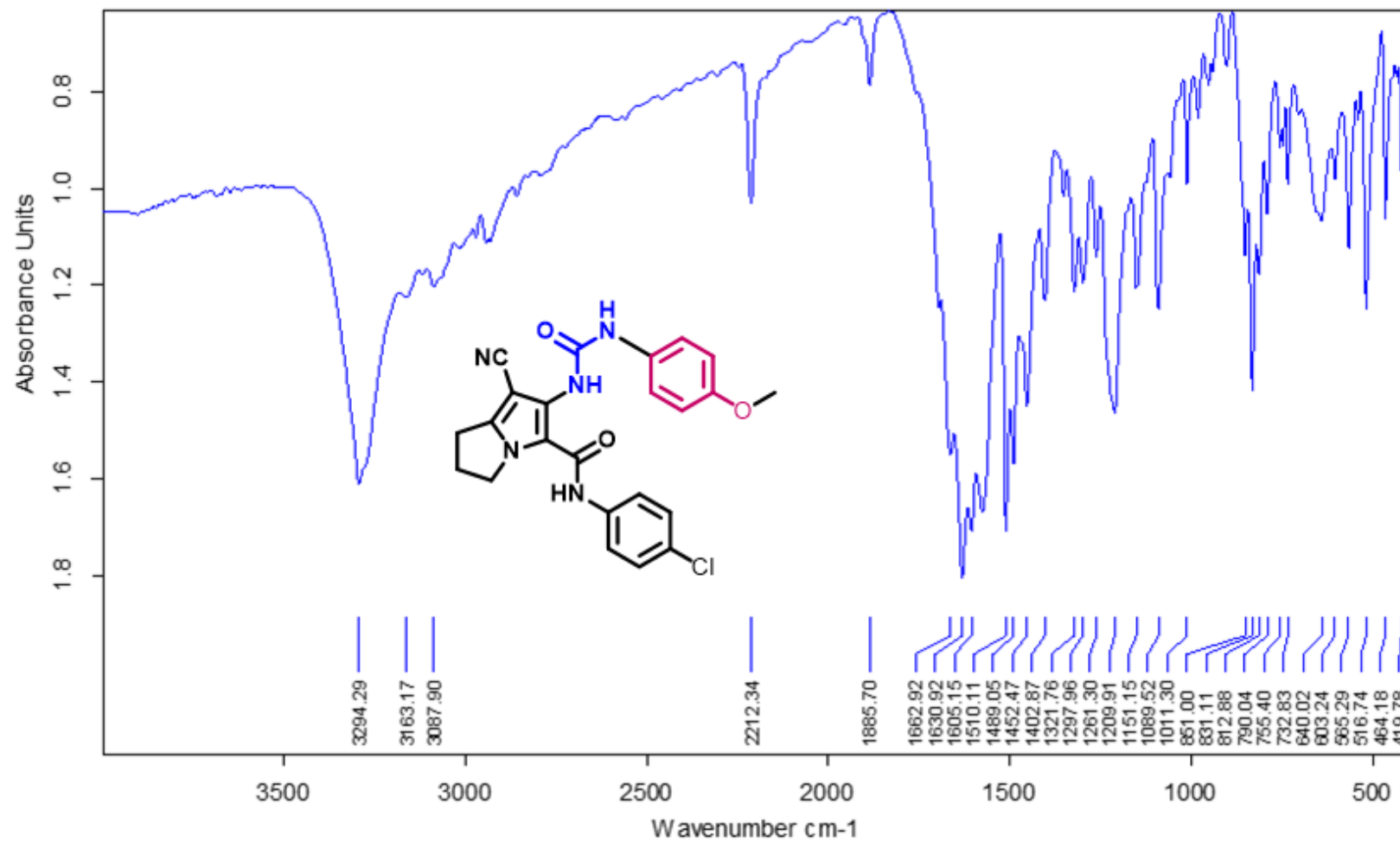
C:\Program Files\OPUS_65\MEAS\Protein.206	Protein	AquaSpec	26/10/2017
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Figure S47. IR spectrum of compound 17b



C:\Program Files\OPUS_65\MEAS\Protein.250	Protein	AquaSpec	31/01/2018
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Figure S48. IR spectrum of compound 17c



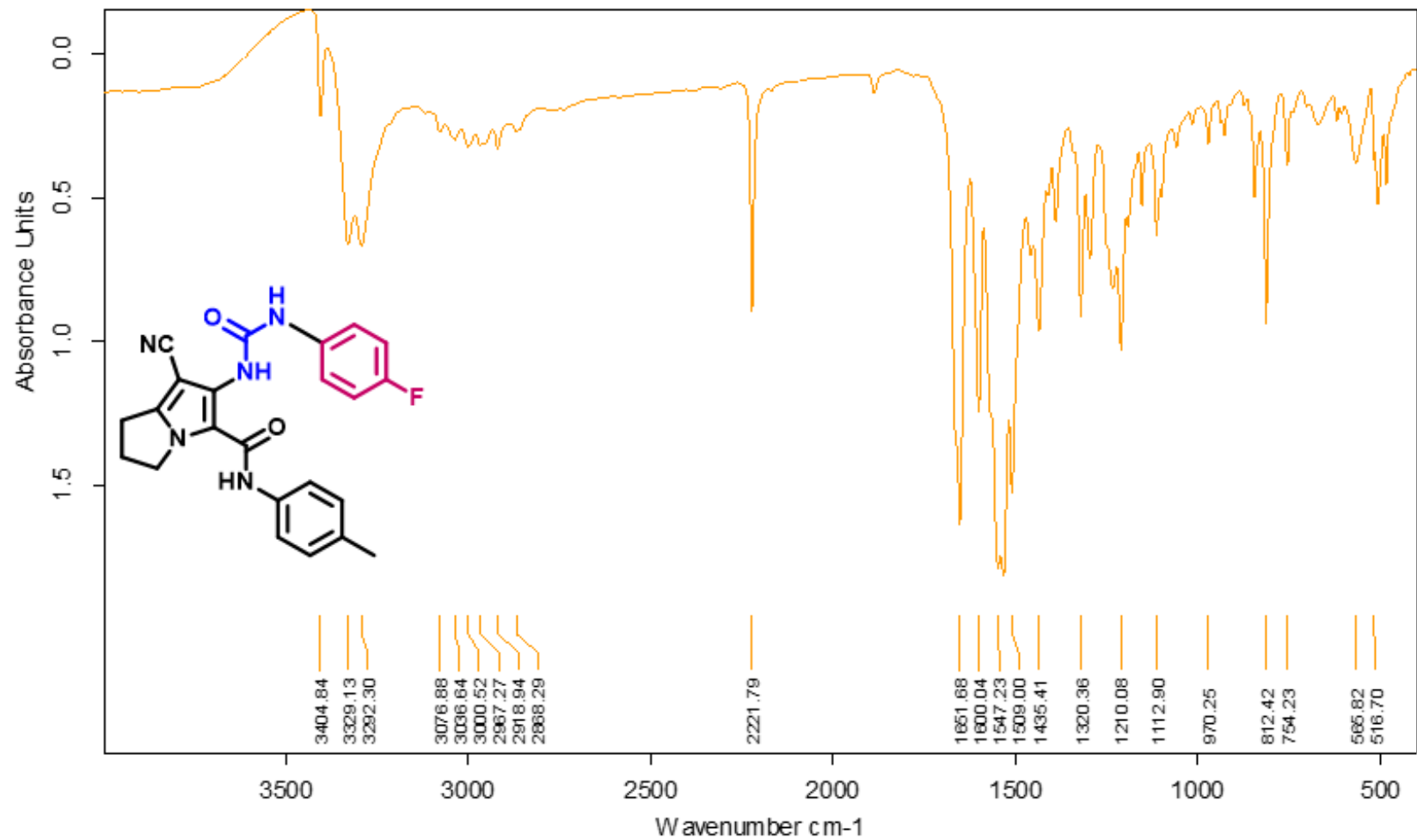
C:\Program Files\OPUS_65\MEAS\Protein.220

Protein

AquaSpec

26/10/2017

Figure S50. IR spectrum of compound 18b



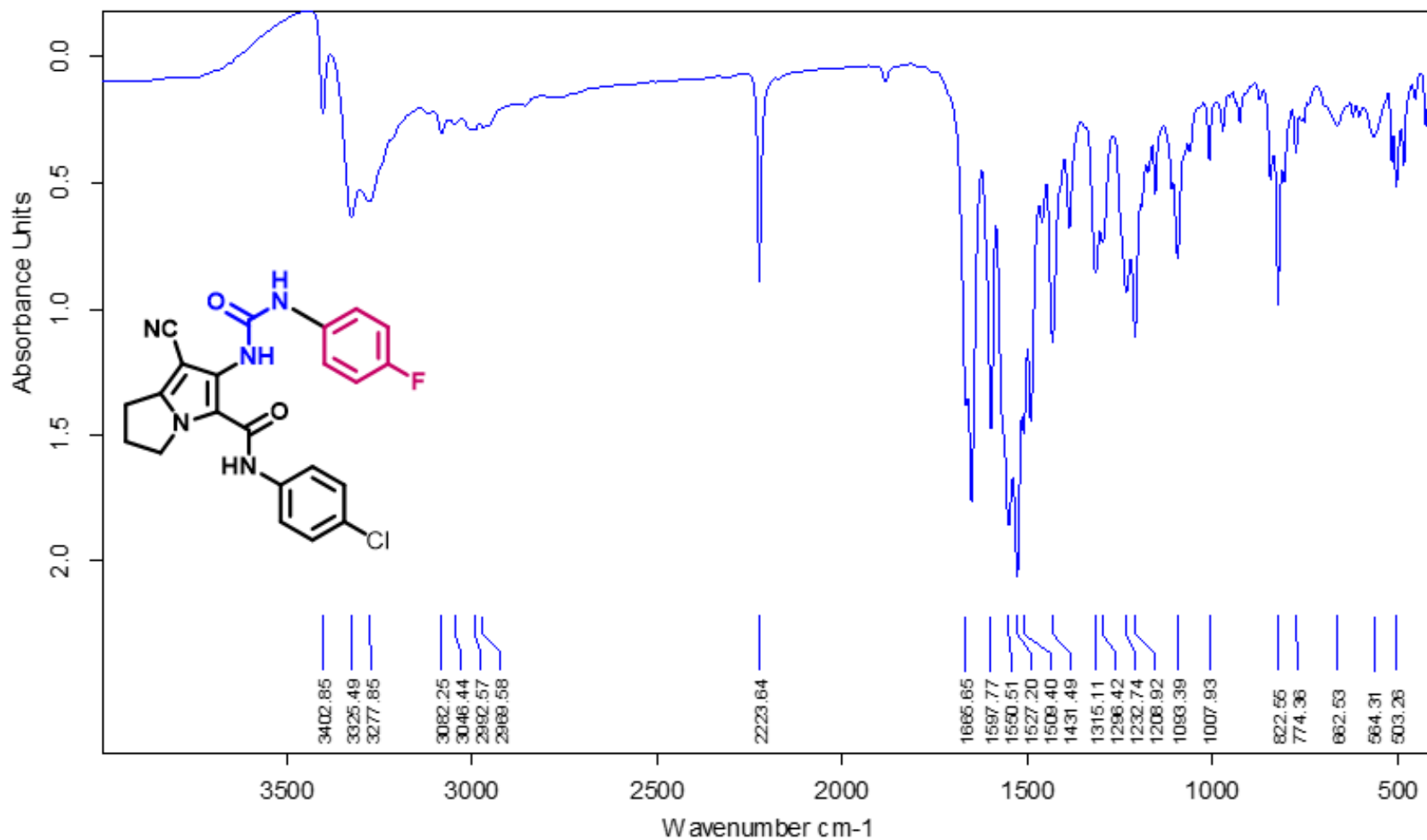
C:\Program Files\OPUS_65\MEAS\Protein.252

Protein

AquaSpec

31/01/2018

Figure S51. IR spectrum of compound 18c



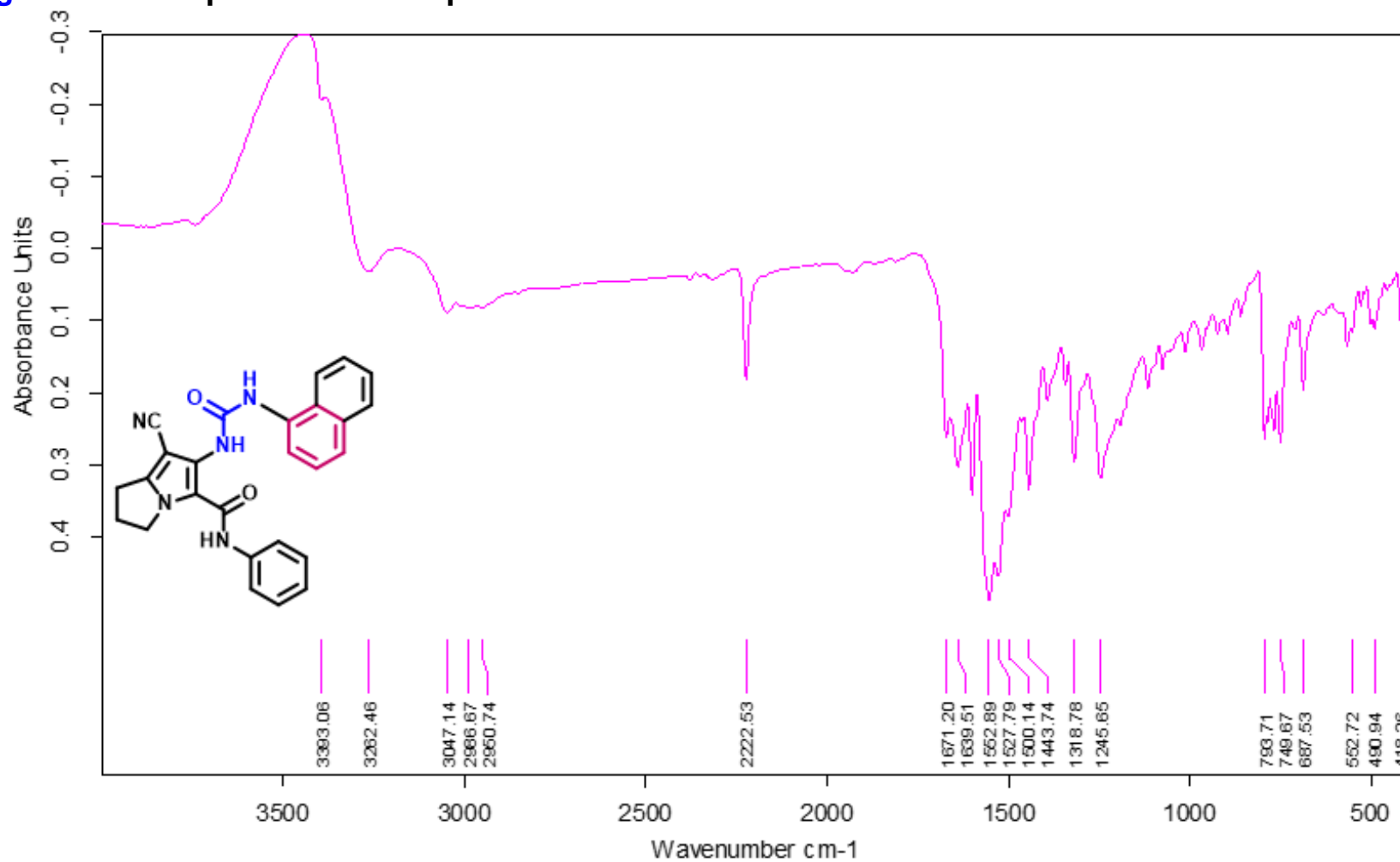
C:\Program Files\OPUS_65\MEAS\Protein.253

Protein

AquaSpec

31/01/2018

Figure S52. IR spectrum of compound 19a



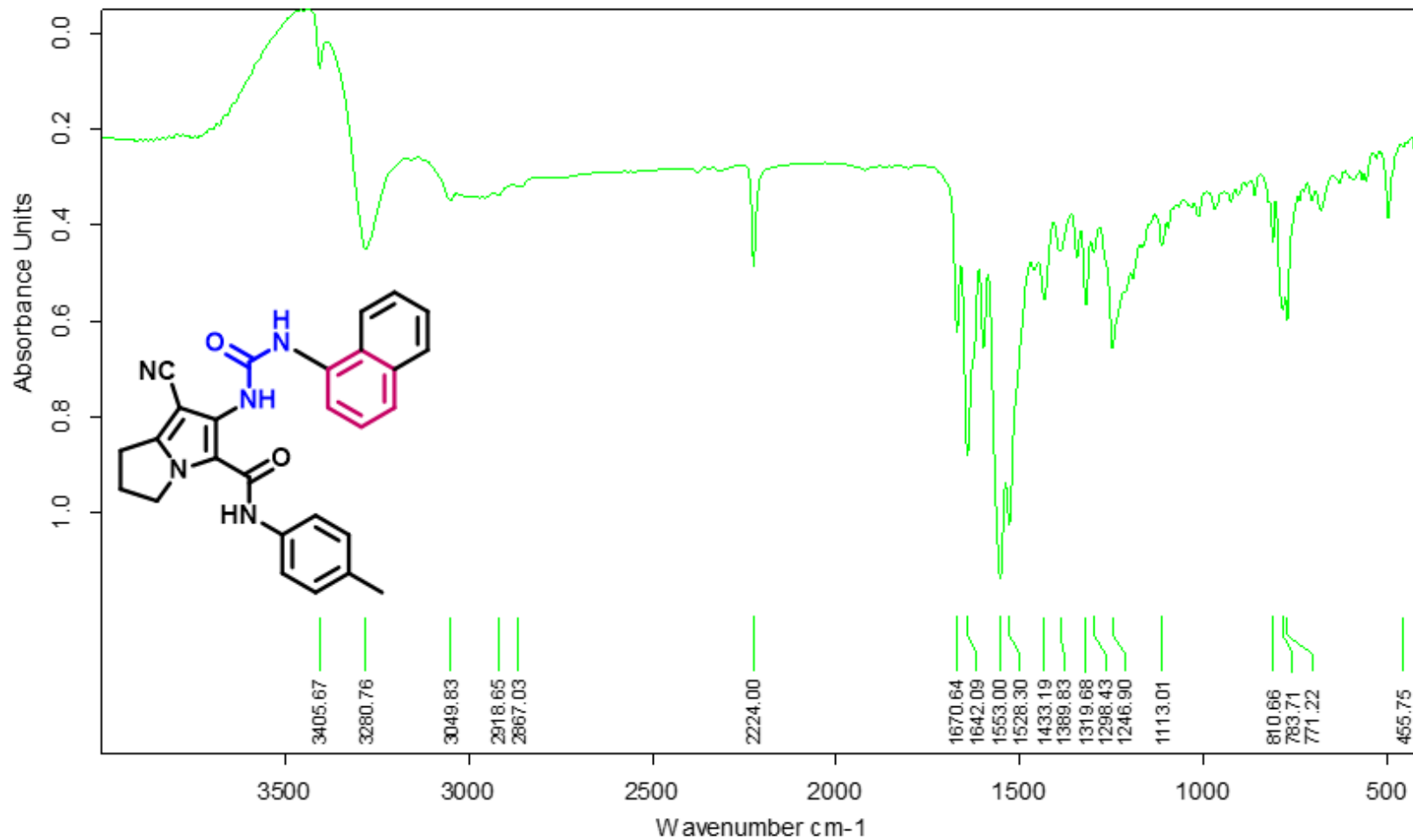
C:\Program Files\OPUS_65\MEAS\Protein.255

Protein

AquaSpec

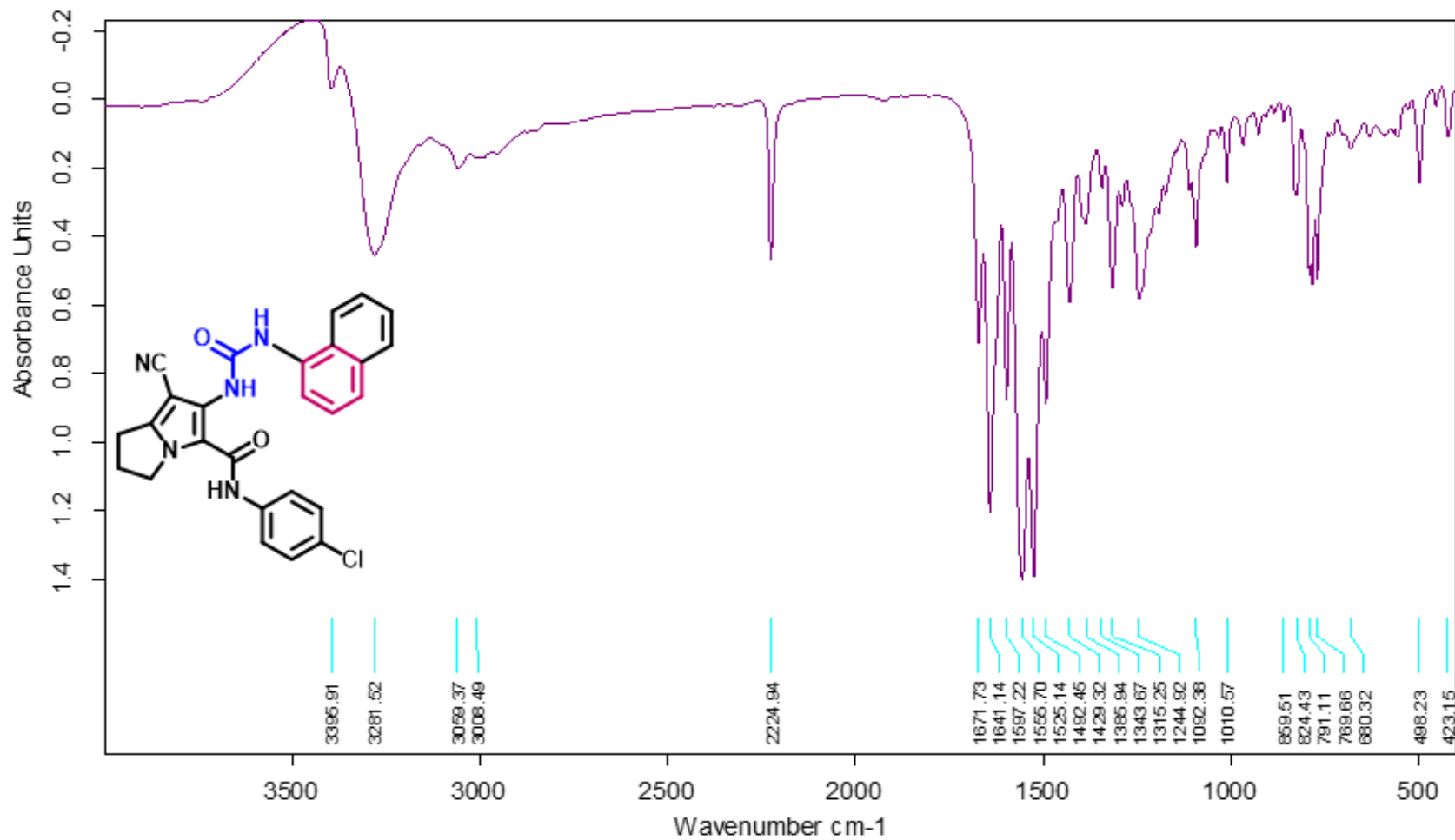
31/01/2018

Figure S53. IR spectrum of compound 19b



C:\Program Files\OPUS_65\MEAS\Protein.256	Protein	AquaSpec	31/01/2018
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Figure S54. IR spectrum of compound 19c



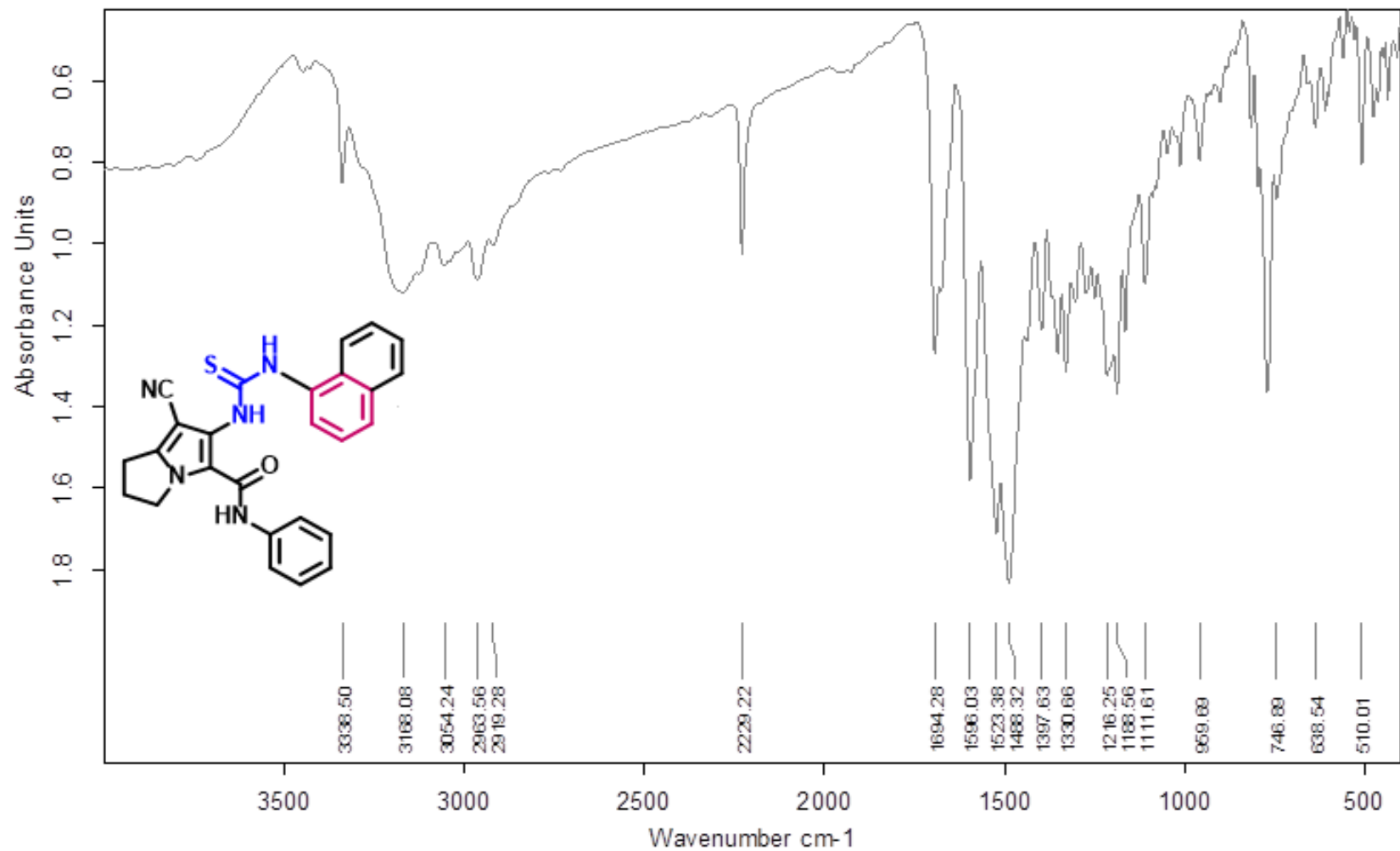
C:\Program Files\OPUS_85\MEAS\Protein.257

Protein

AquaSpec

31/01/2018

Figure S55. IR spectrum of compound 20a



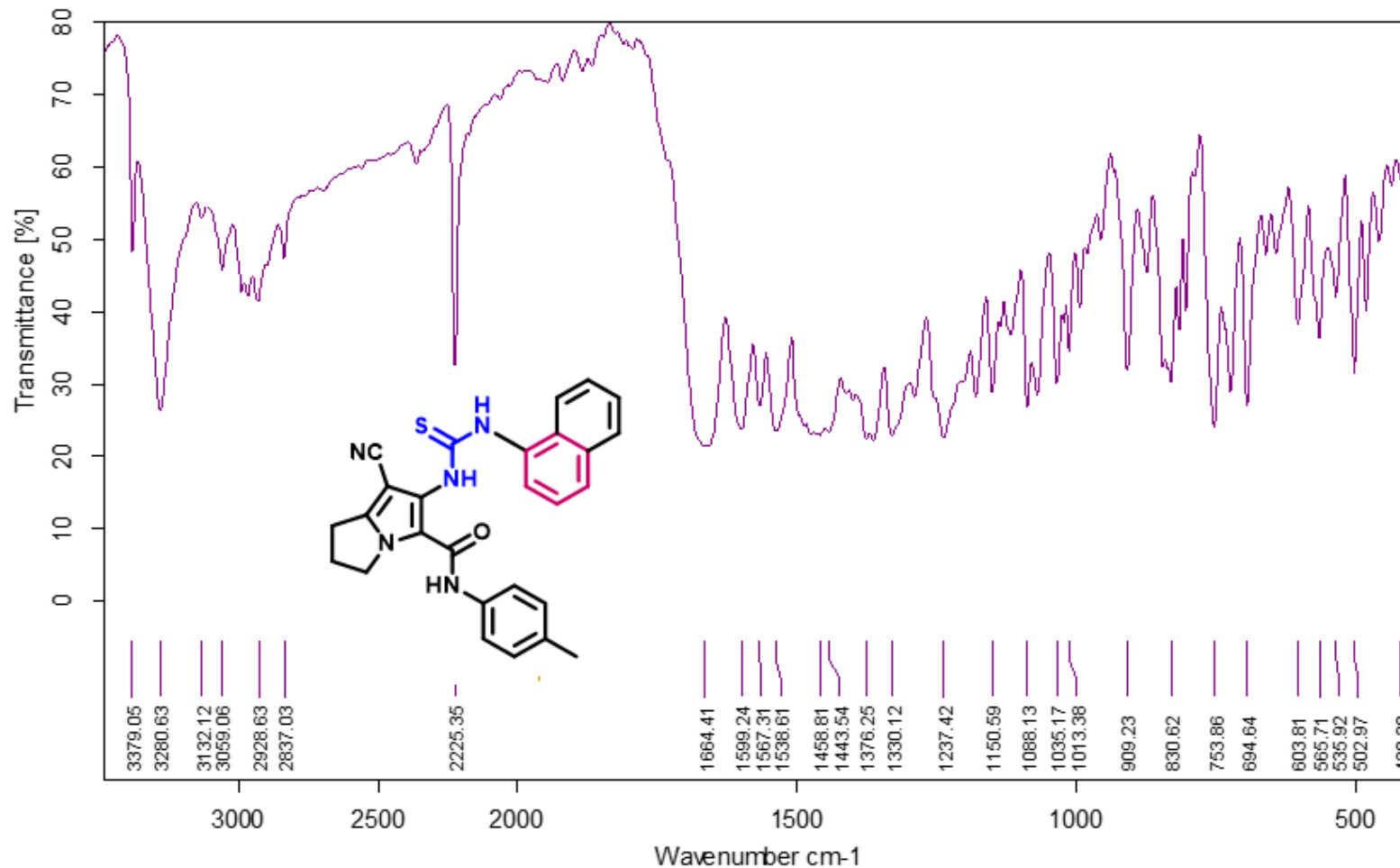
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Protein

AquaSpec

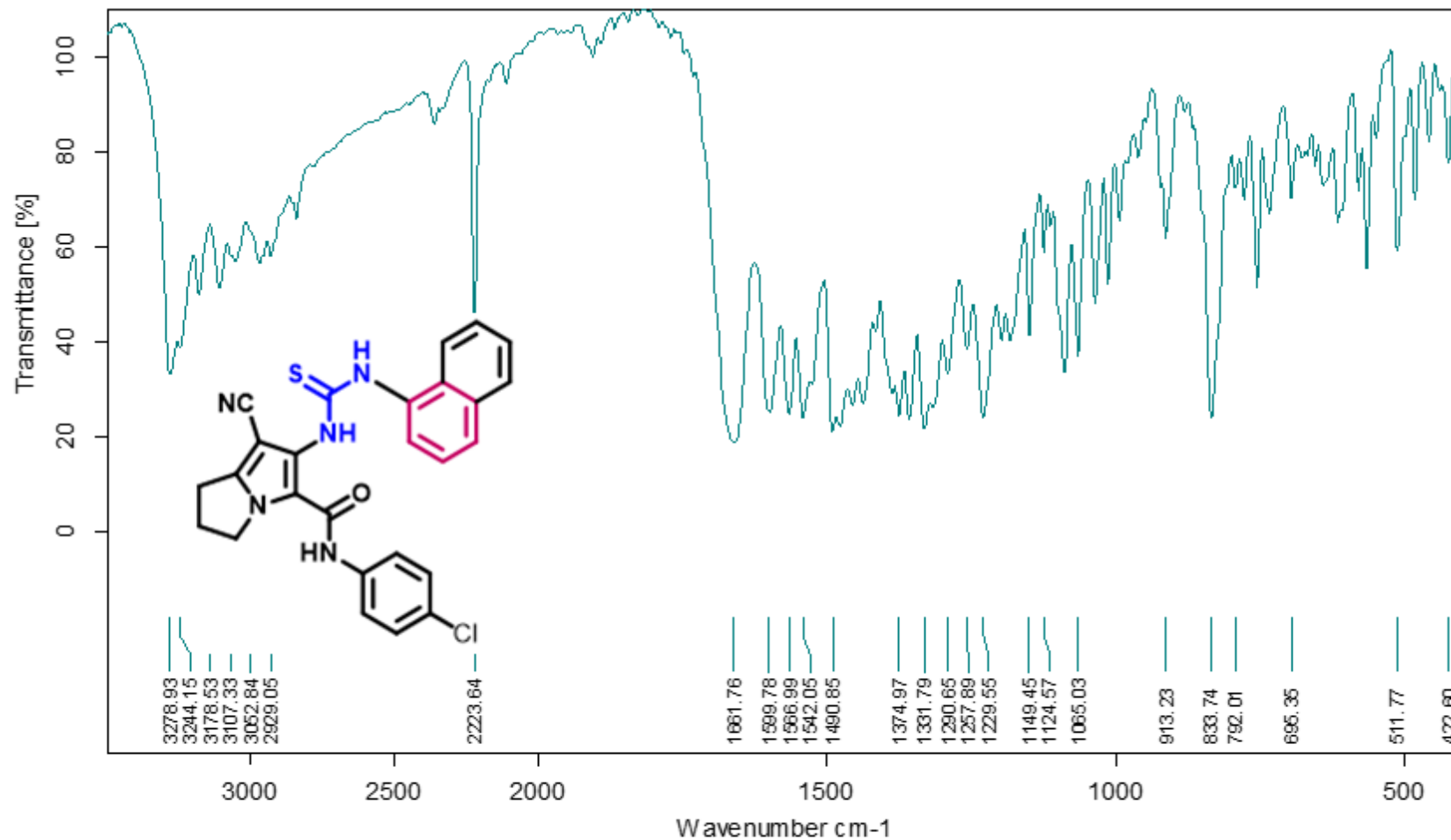
31/01/2018

Figure S56. IR spectrum of compound 20b



C:\Program Files\OPUS_65\MEAS\MLT.128 MLT Instrument type and / or accessory

Figure S57. IR spectrum of compound 20c



C:\Program Files\OPUS_65\MEAS\MLT.130

MLT

Instrument type and / or accessory

^1H -NMR, ^{13}C -NMR and DEPT C^{135} Spectra

^1H -NMR spectra were recorded on a BRUKER AVANCE III spectrometer (at the faculty of pharmacy, Umm Al-Qura University) at 500 MHz in the specified solvent, chemical shifts were reported on the δ (ppm) scale and were related to that of the solvent and J values are given in Hz. ^{13}C NMR and DEPT C^{135} spectra were obtained on a BRUKER AVANCE III at 125 MHz (at the faculty of pharmacy, Umm Al-Qura University).

Figure S59. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound 16a (zoom on aliphatic Hs)

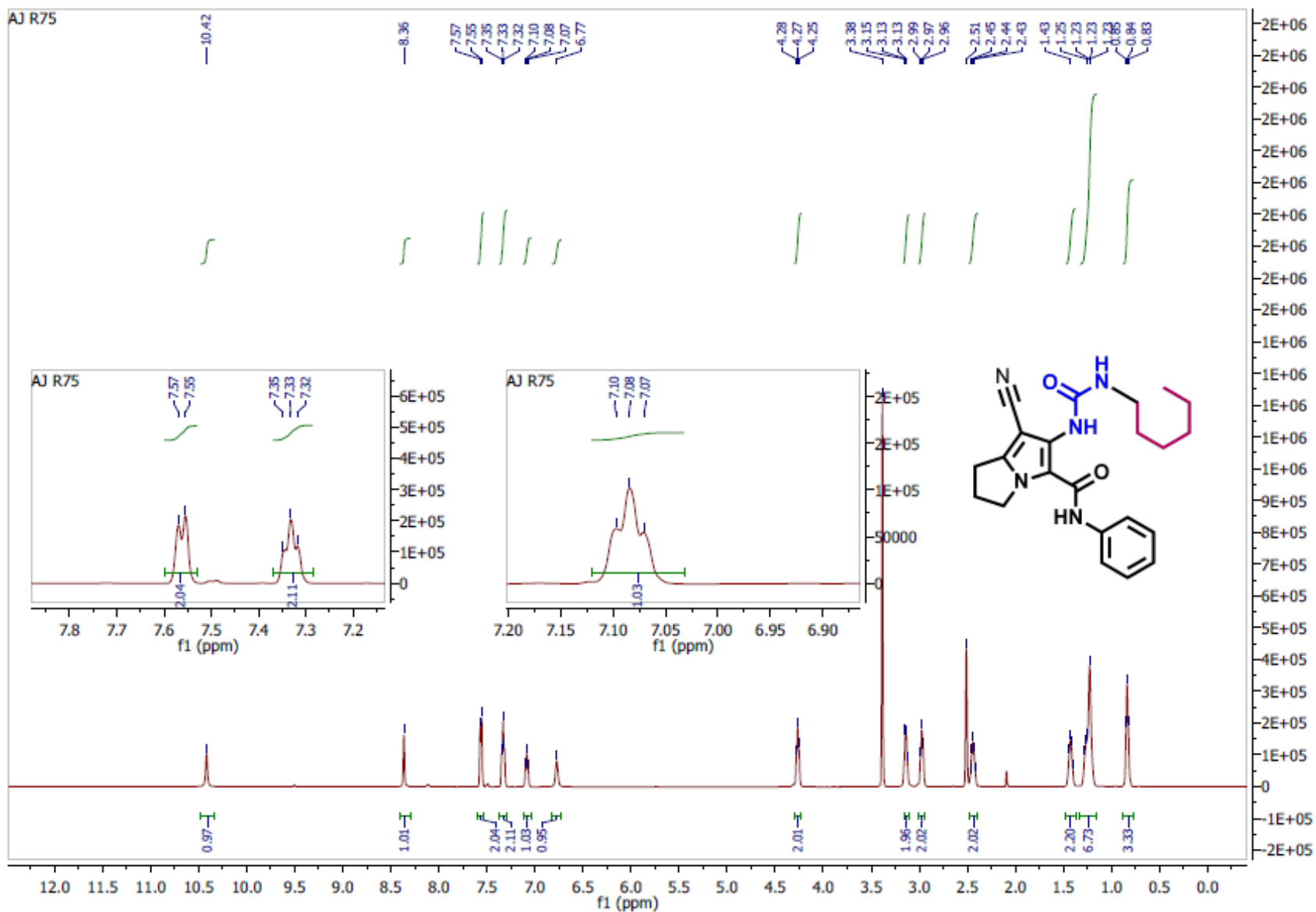


Figure S60. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **16a** (zoom on aliphatic Hs)

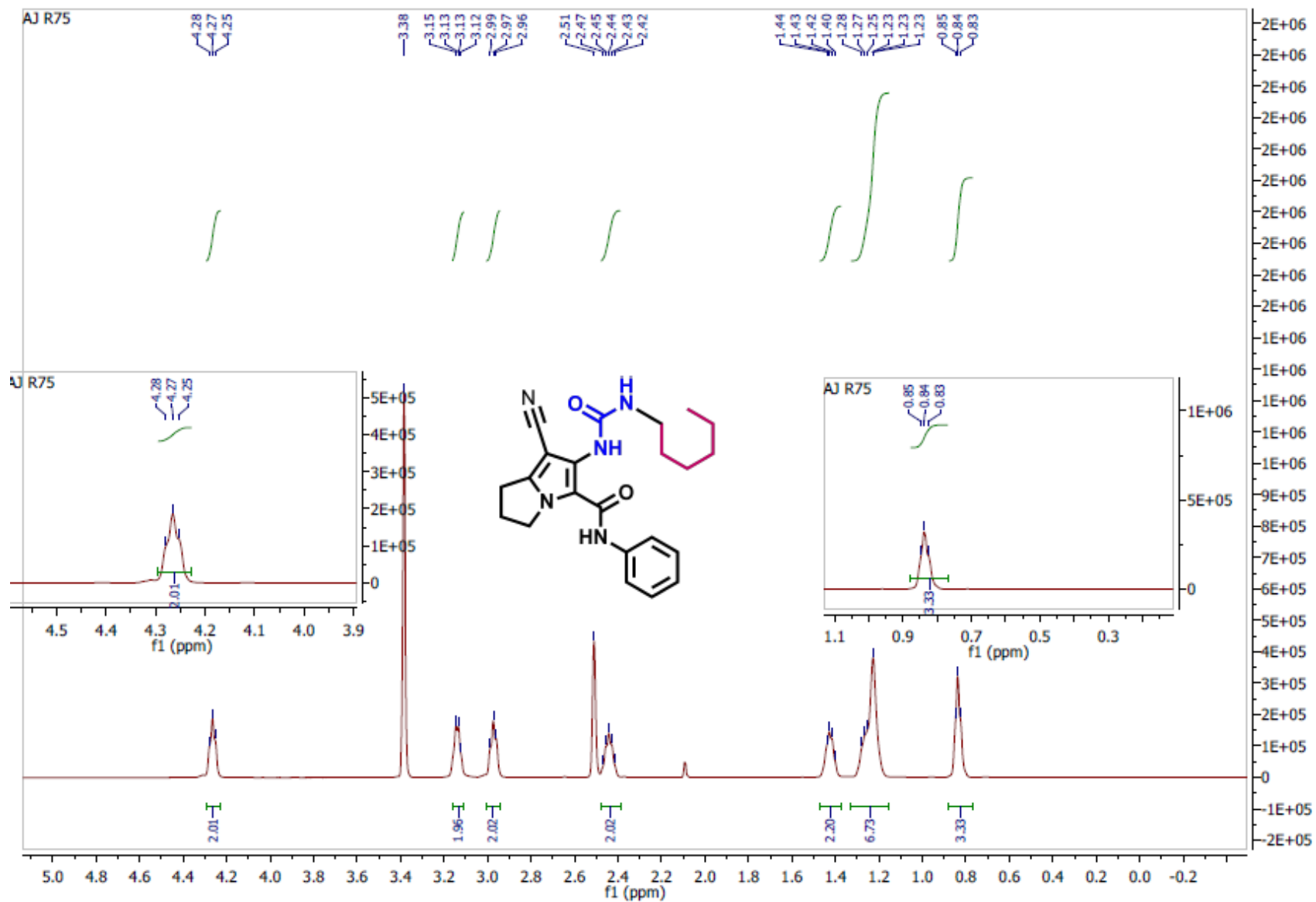


Figure S61. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **16a** (zoom on aliphatic Hs)

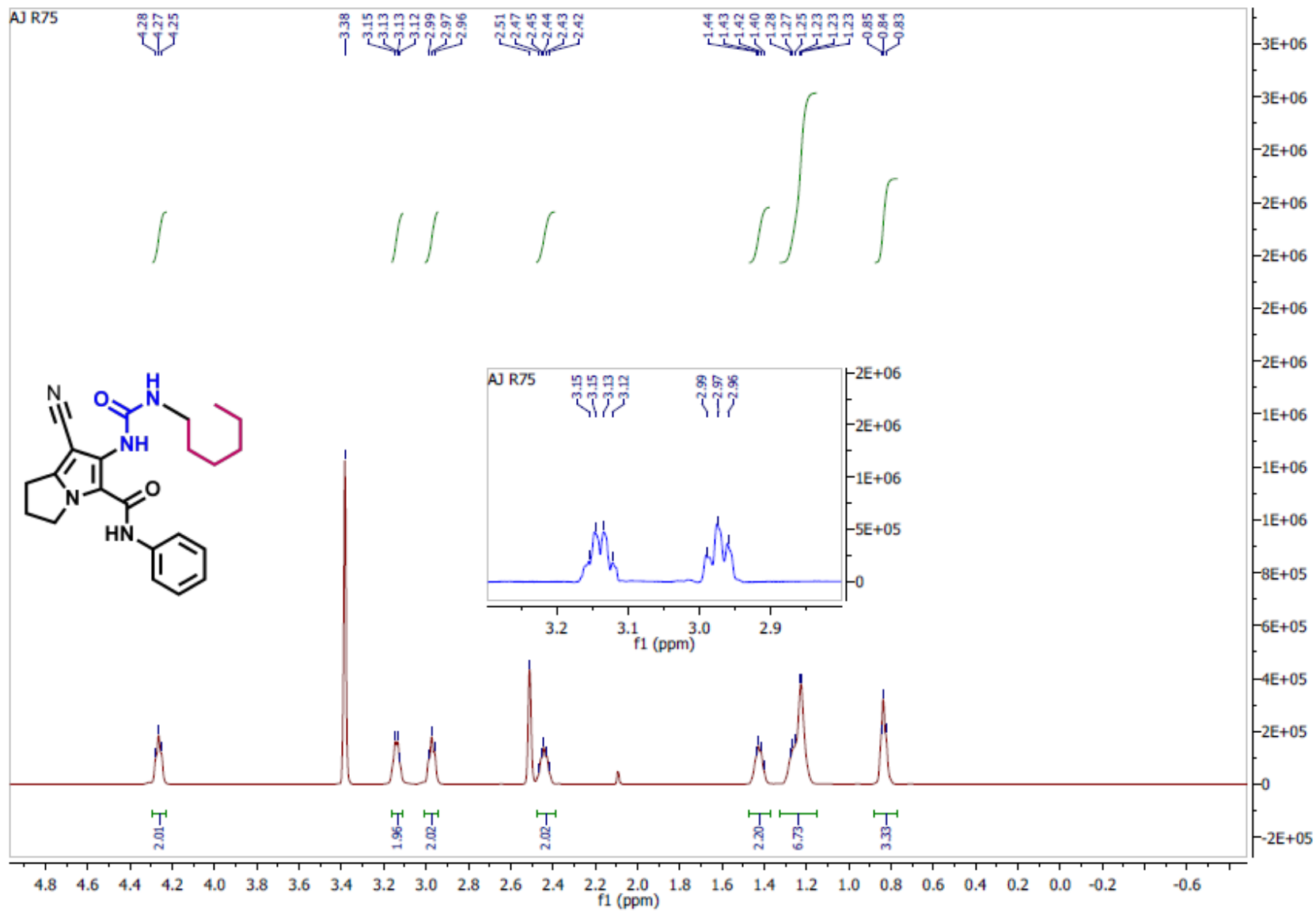


Figure S62. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **16a** (zoom on NHs & aromatic Hs)

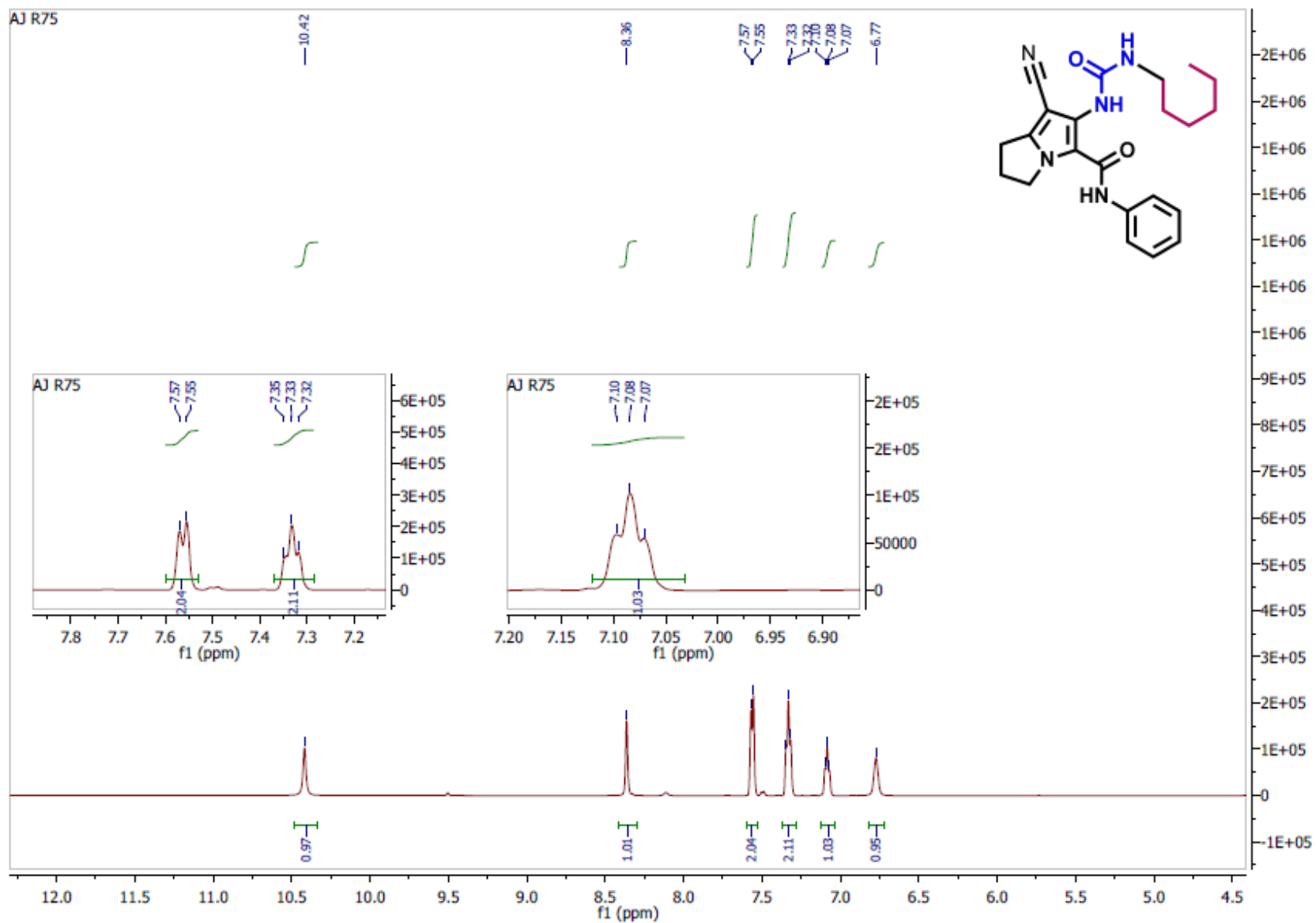


Figure S63. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **16a**

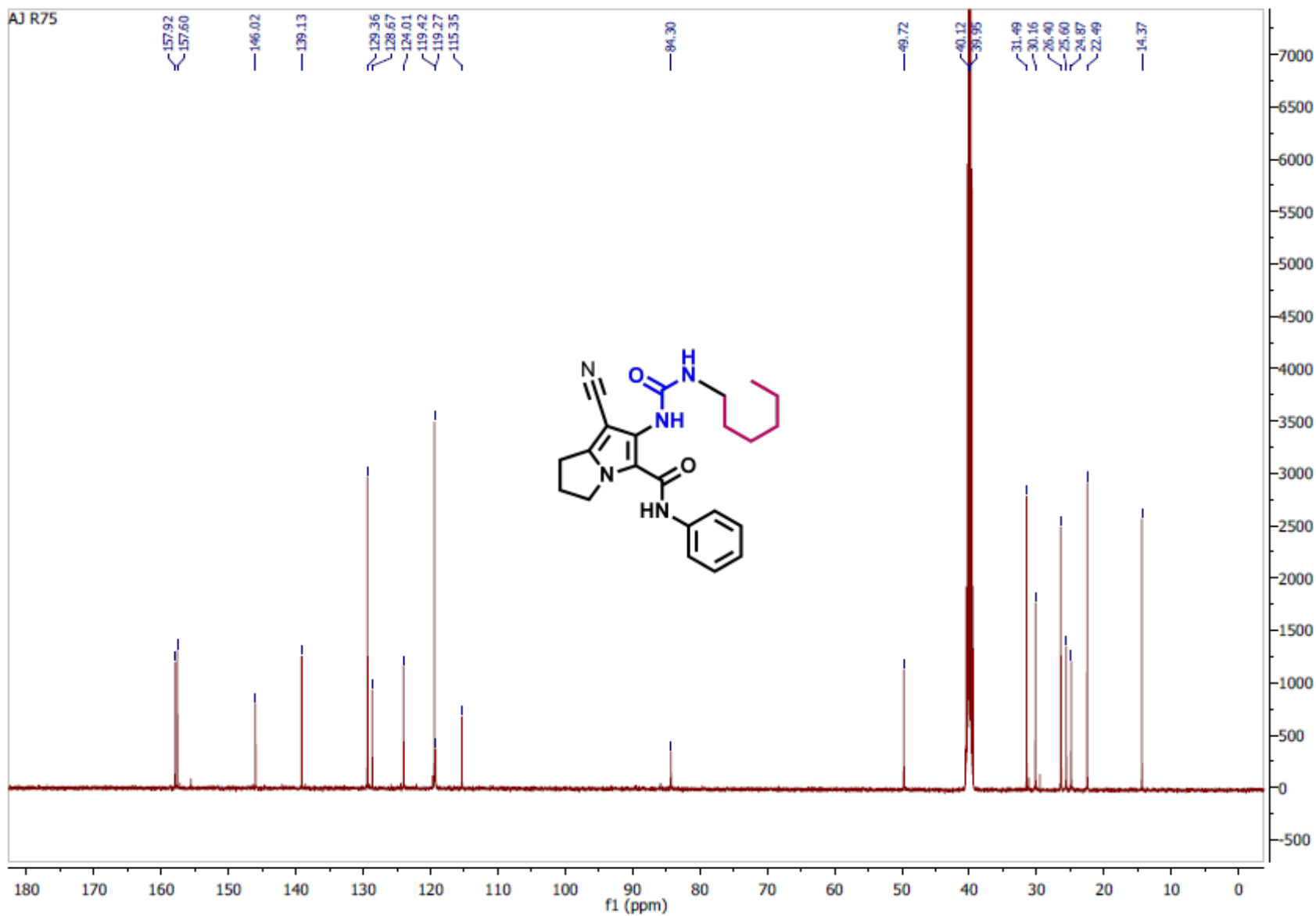


Figure S64. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **16a** (zoom on aliphatic Cs)

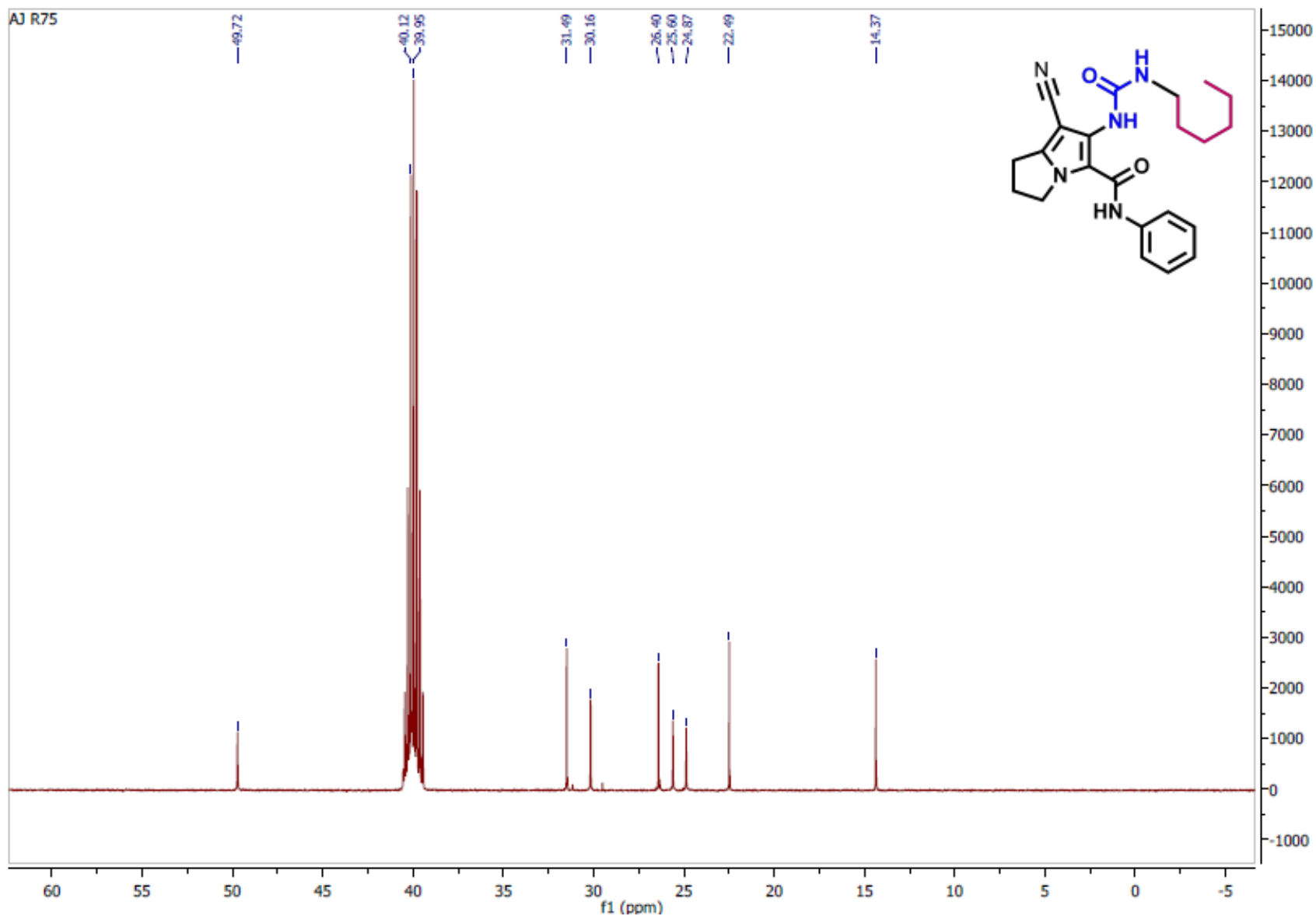


Figure S65. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **16a** (zoom on aromatic Cs)

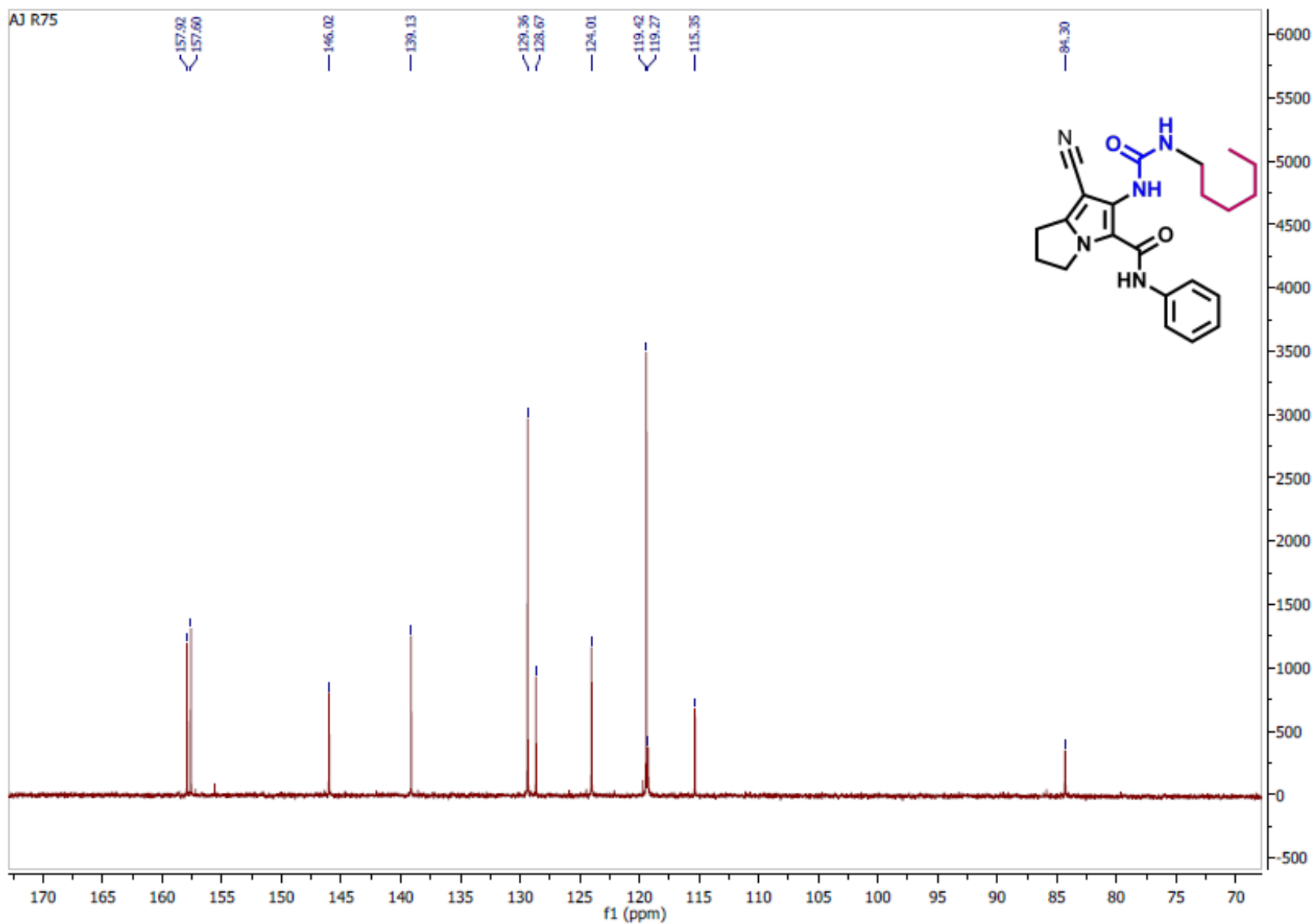


Figure S66. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound **16a**.

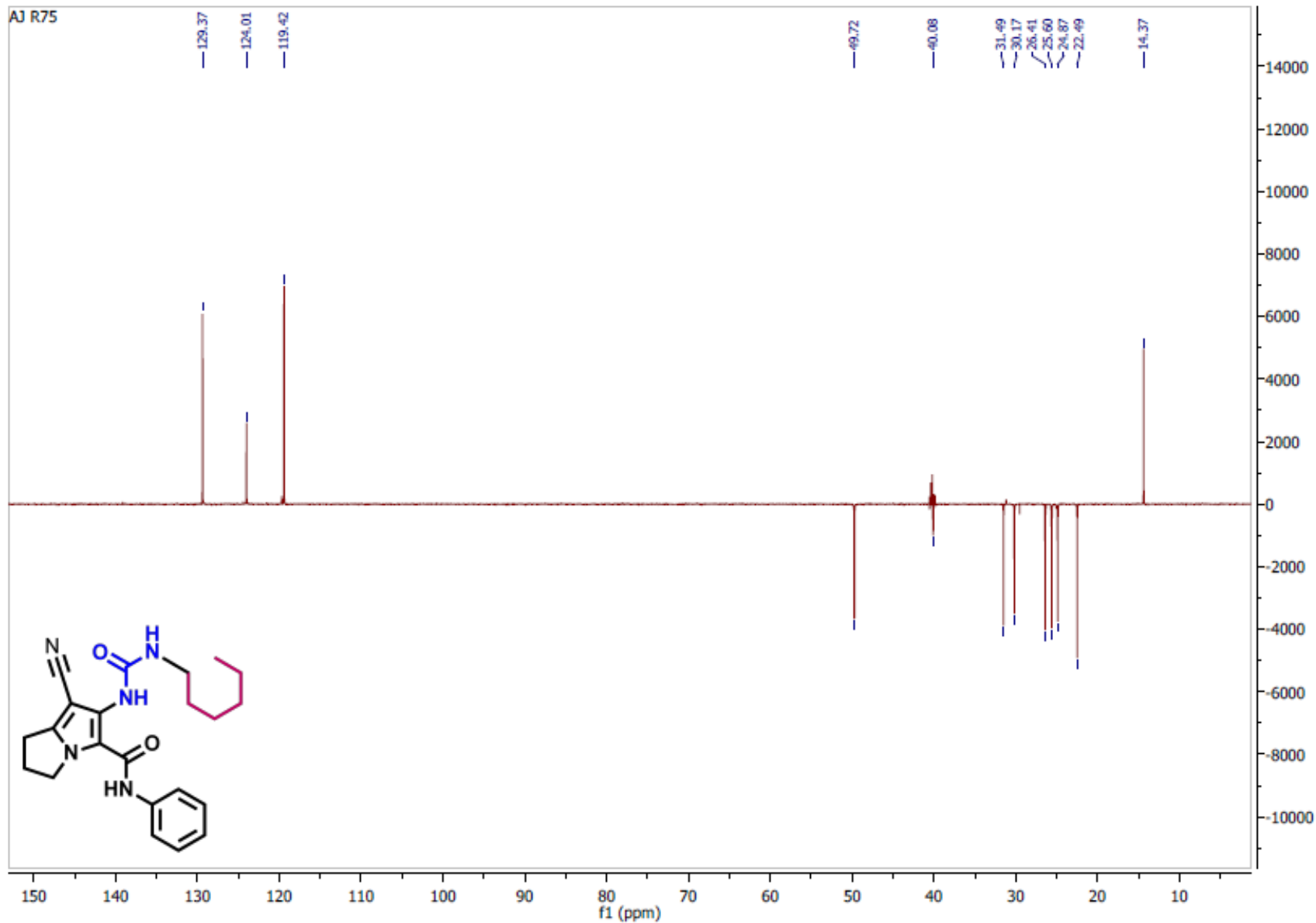


Figure S67. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound 16b.

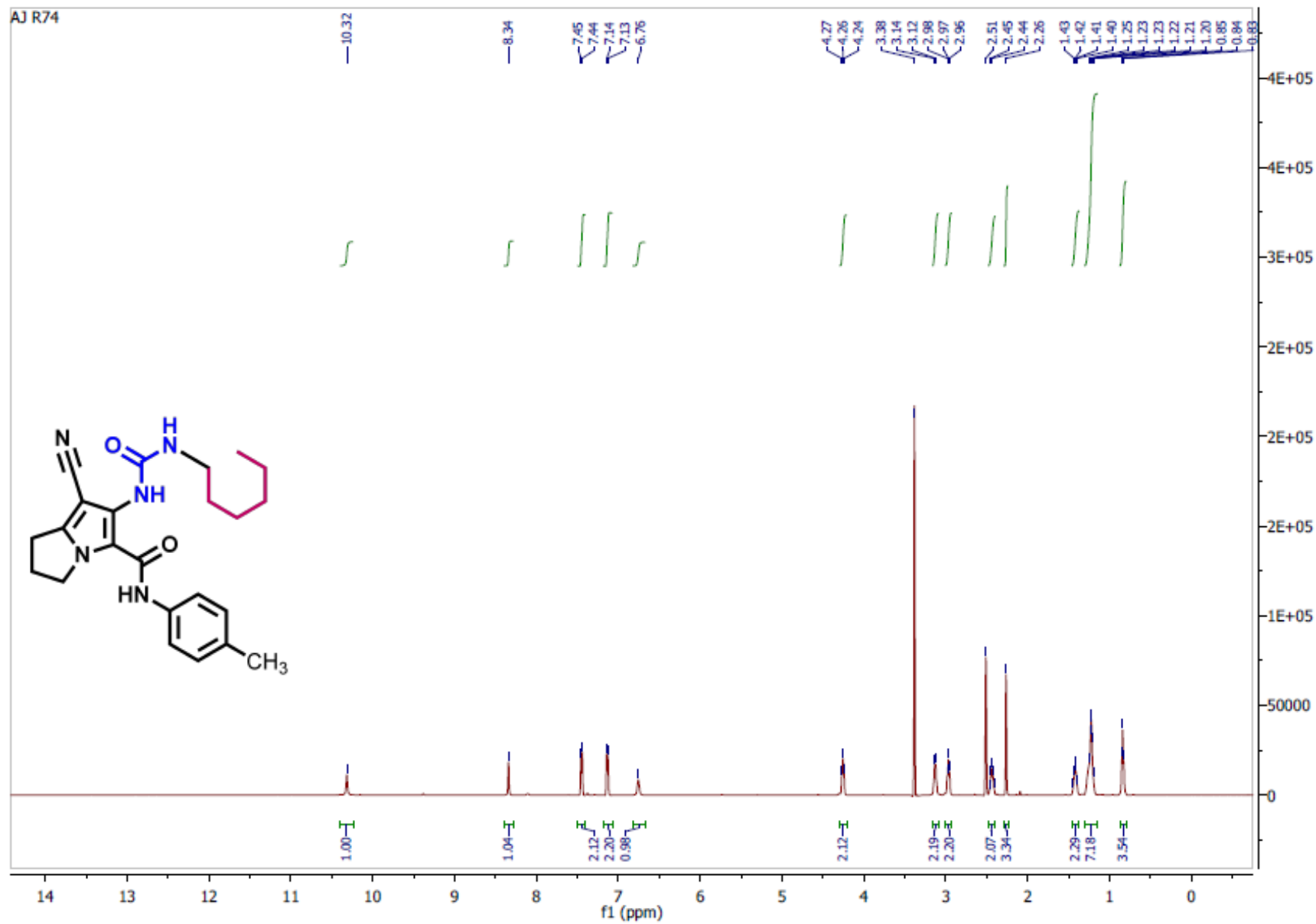


Figure S68. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound **16b** (zoom on aliphatic Hs).

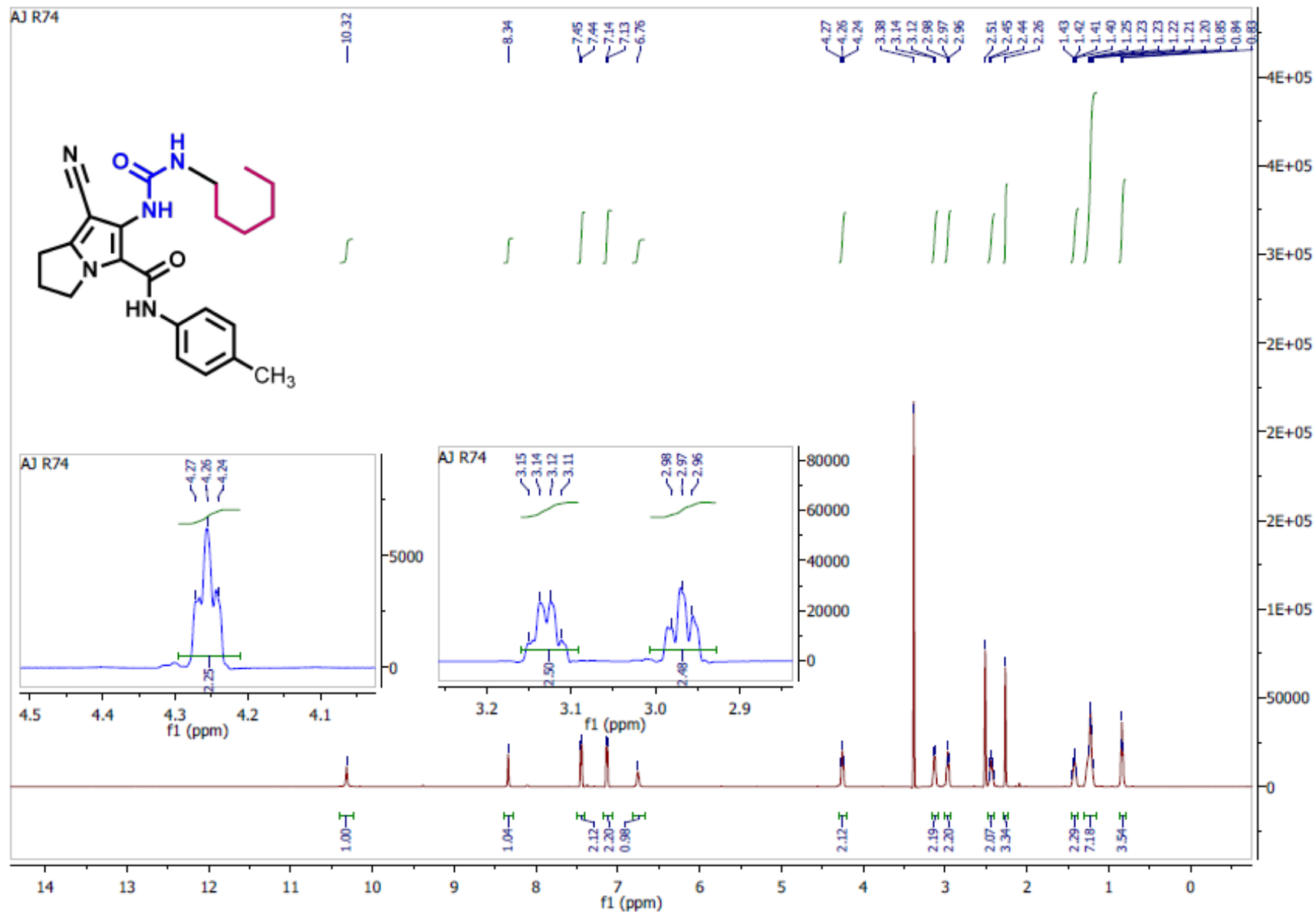


Figure S69. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound **16b** (zoom on aliphatic Hs).

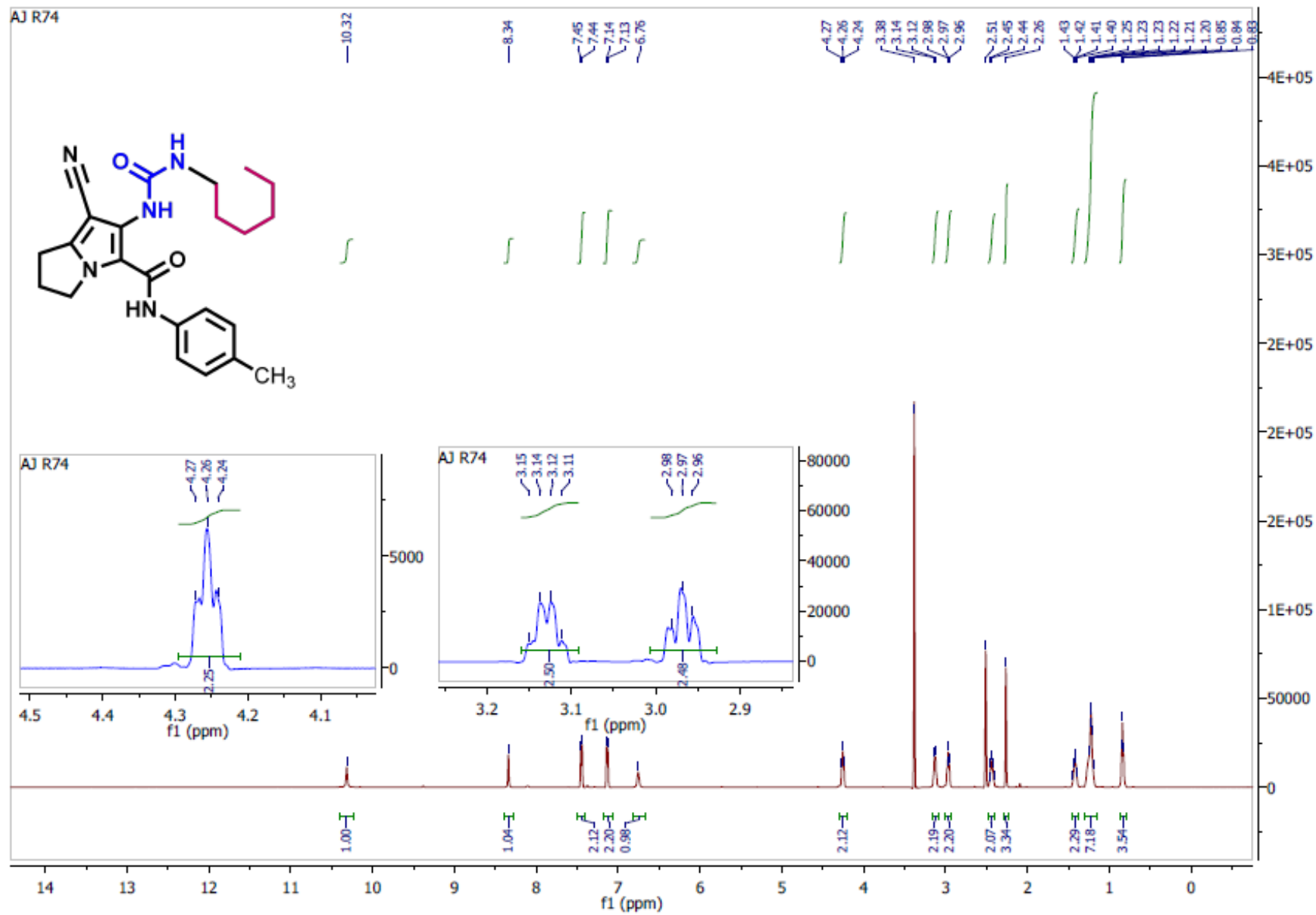


Figure S70. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **16b** (zoom on aromatic Hs).

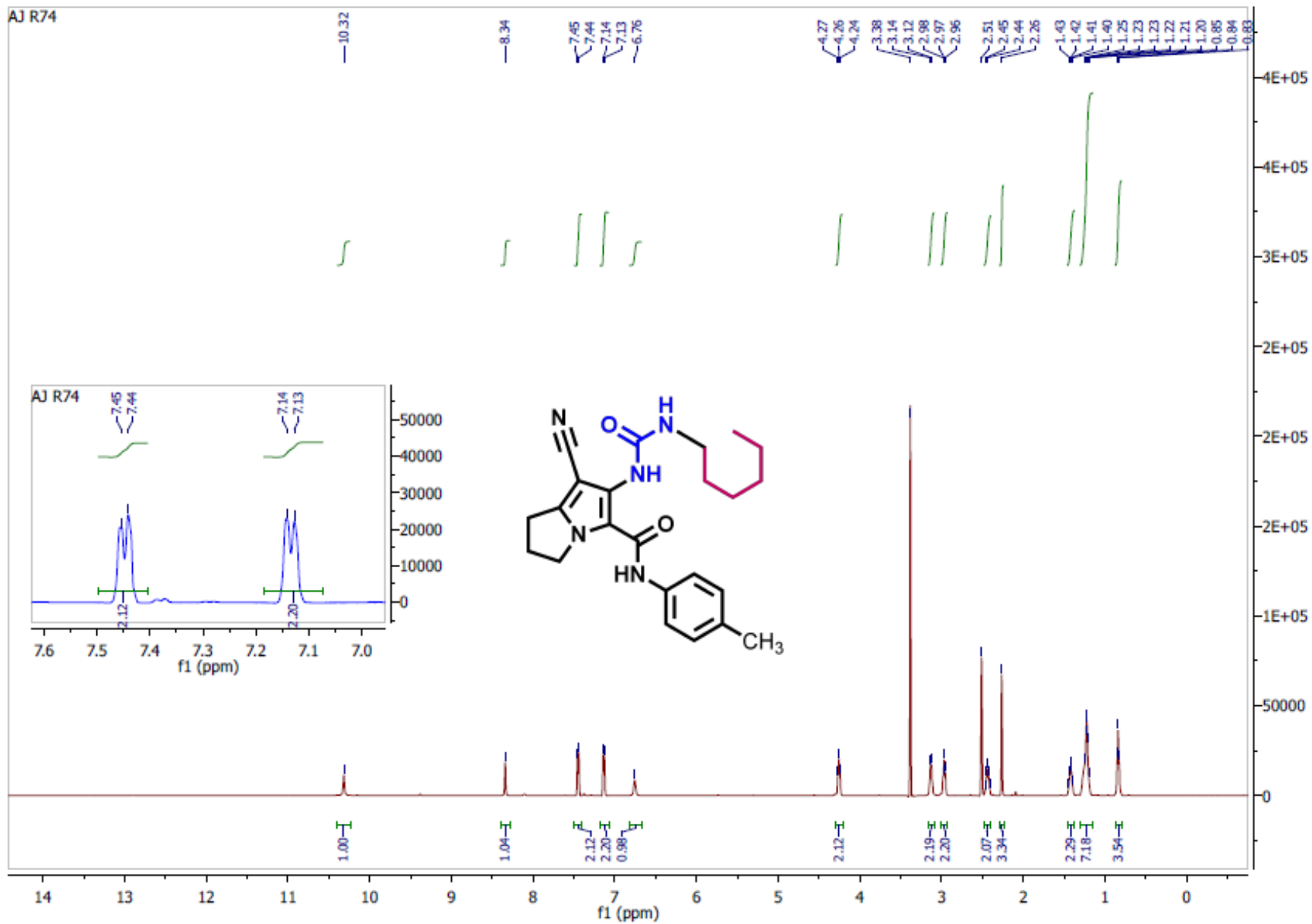


Figure S71. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **16b**.

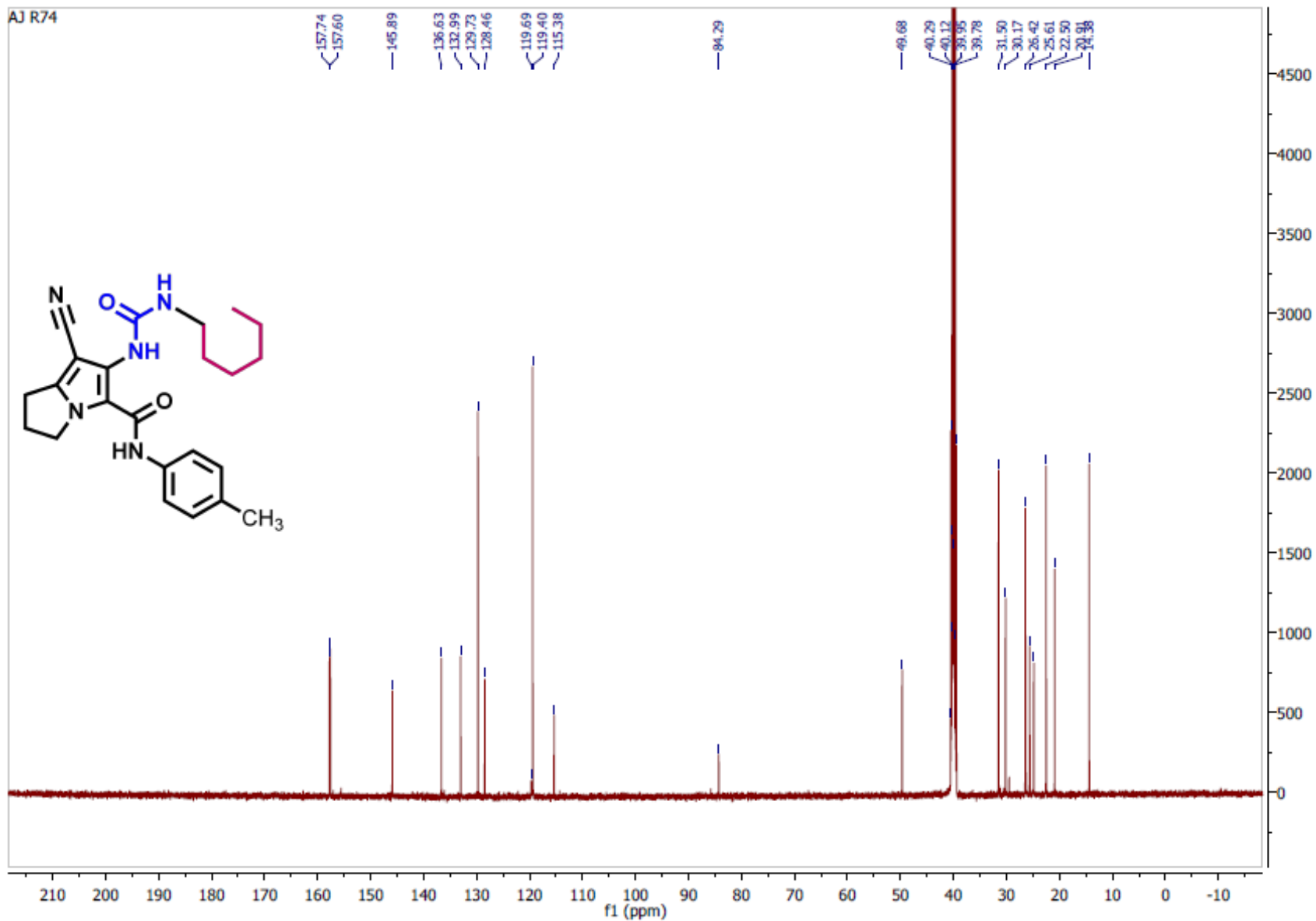


Figure S72. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **16b** (zoom on aliphatic Cs).

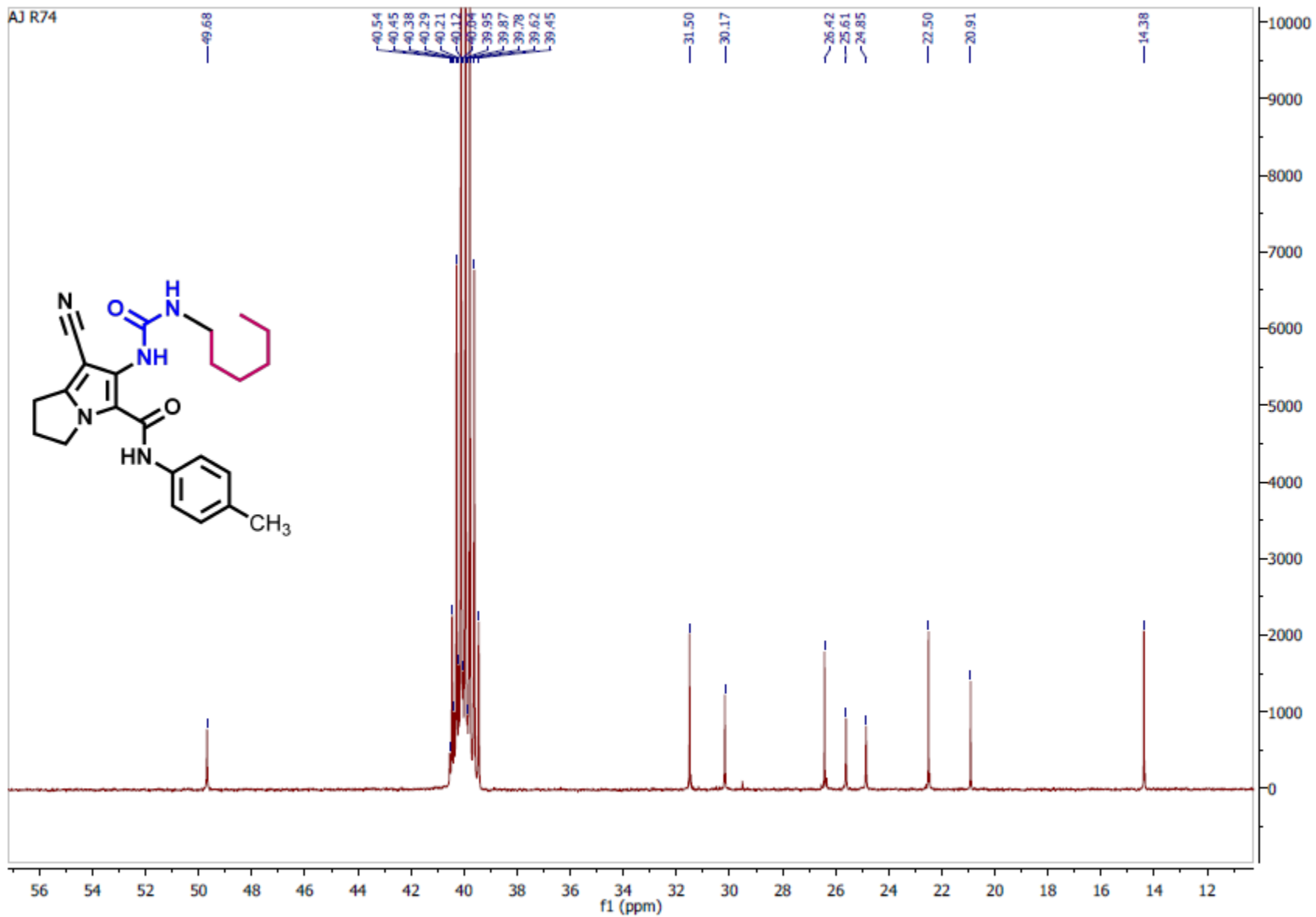


Figure S73. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **16b** (zoom on aromatic Cs).

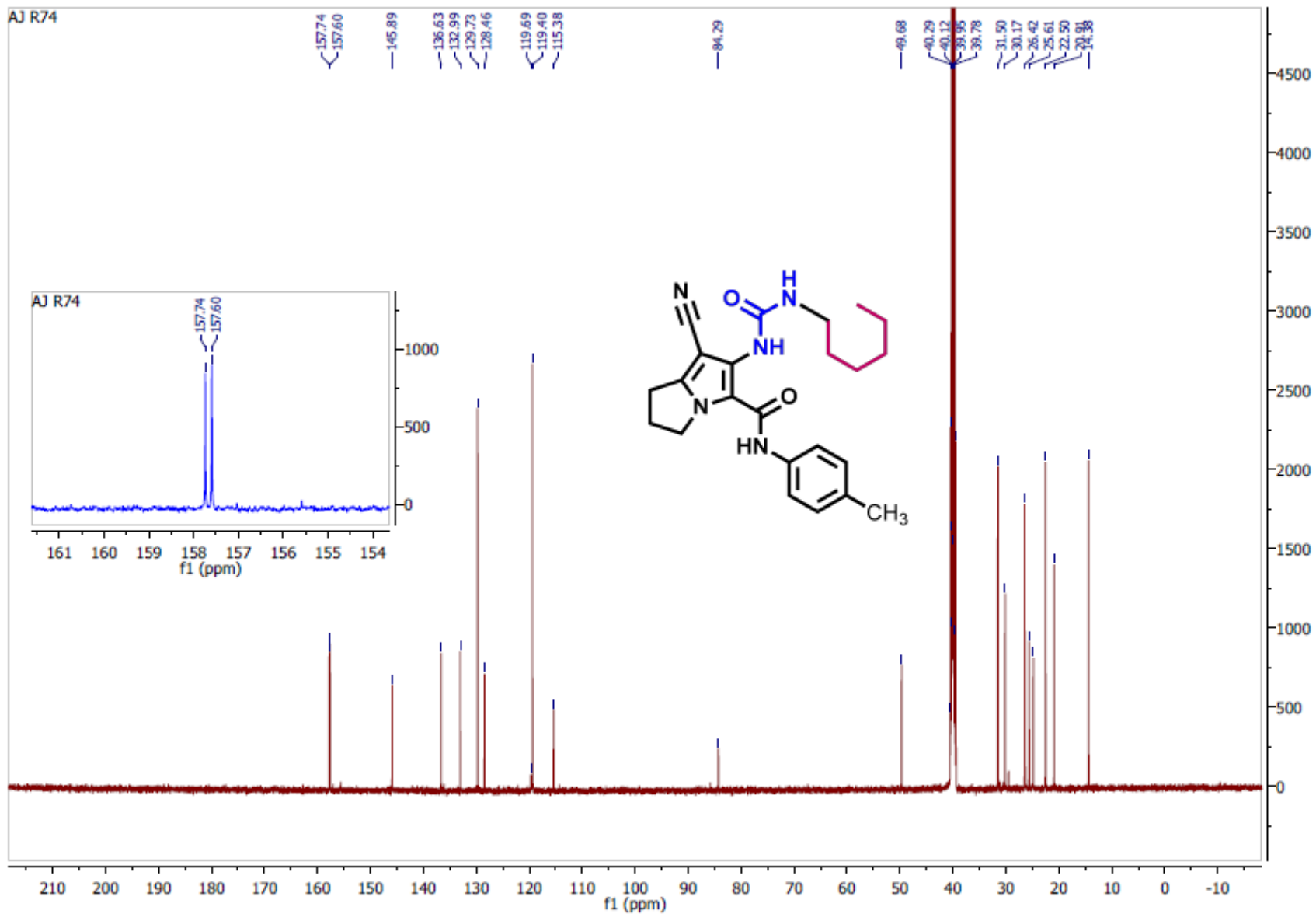


Figure S74. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound **16b**.

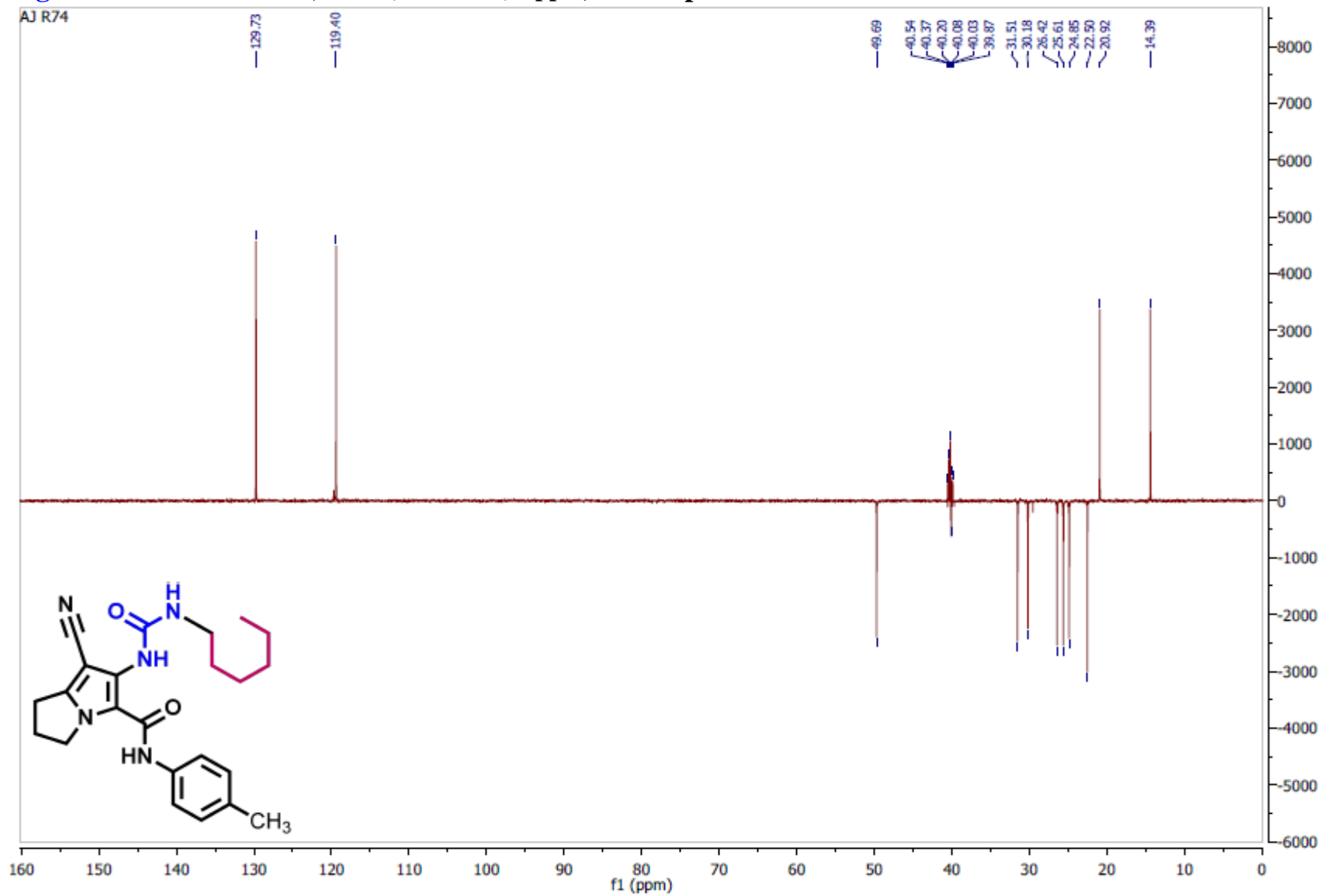


Figure S75. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound **16b** (zoom on aliphatic Cs).

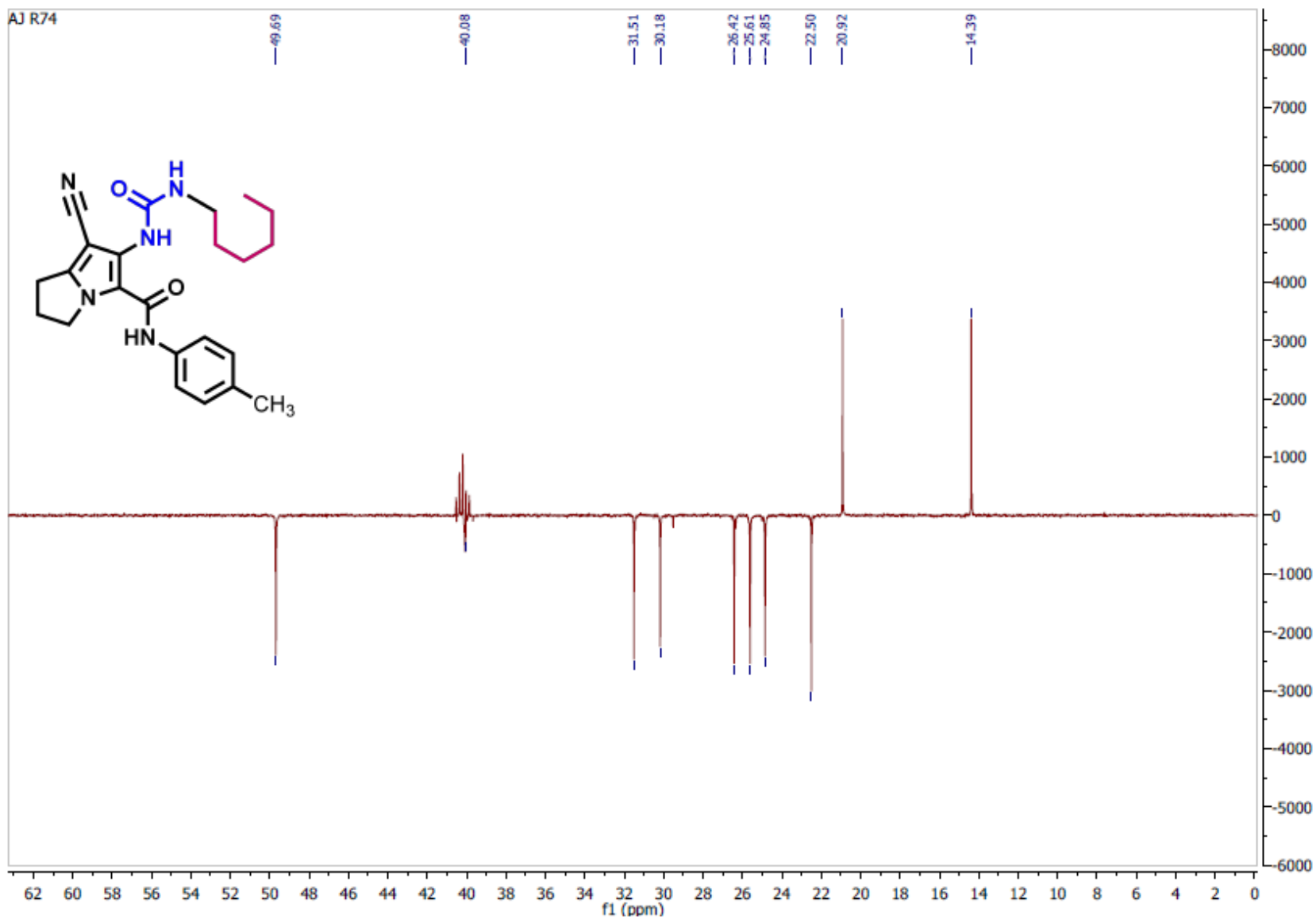


Figure S76. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound **16b** (zoom on aliphatic Cs).

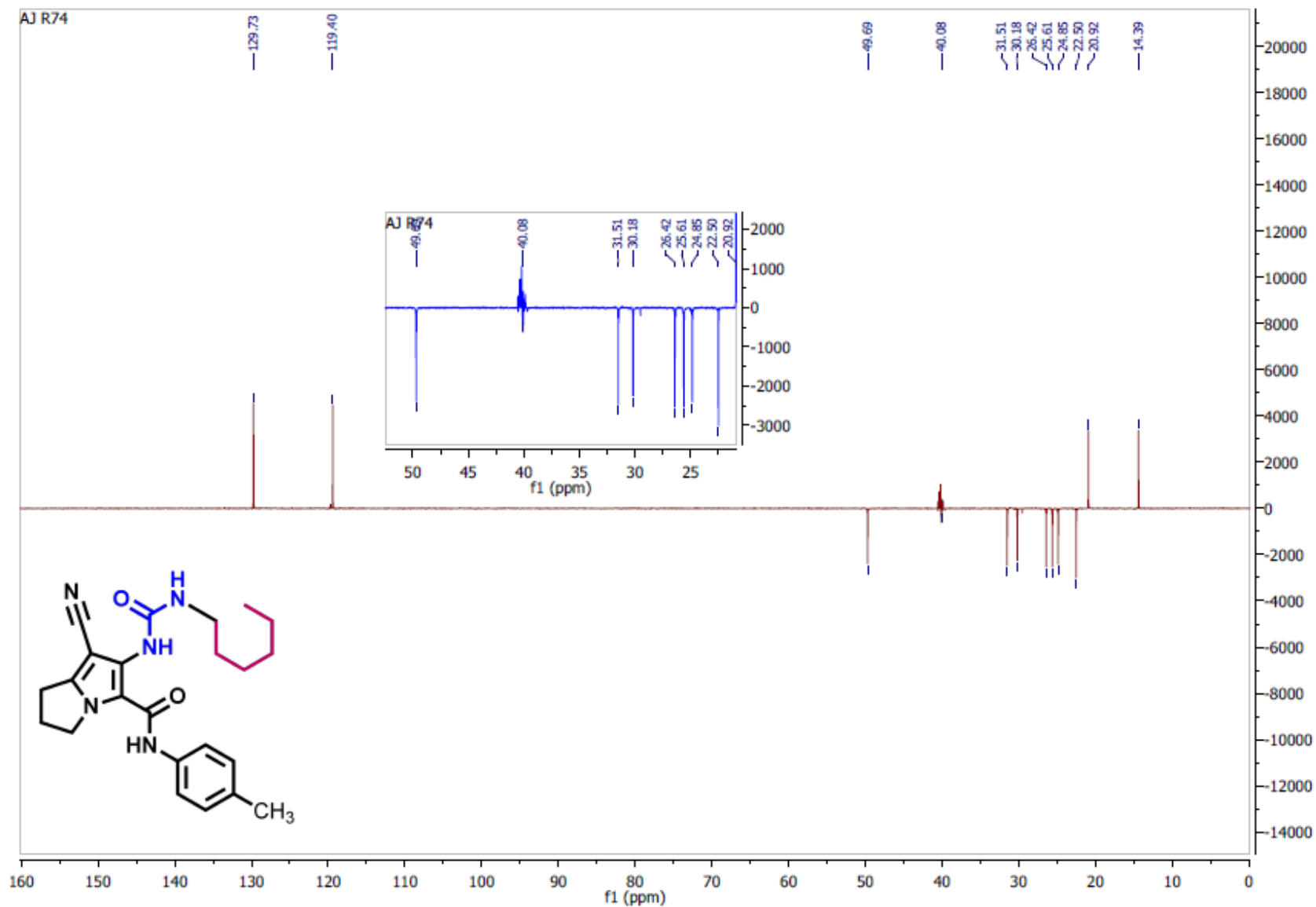


Figure S77. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound 16c.

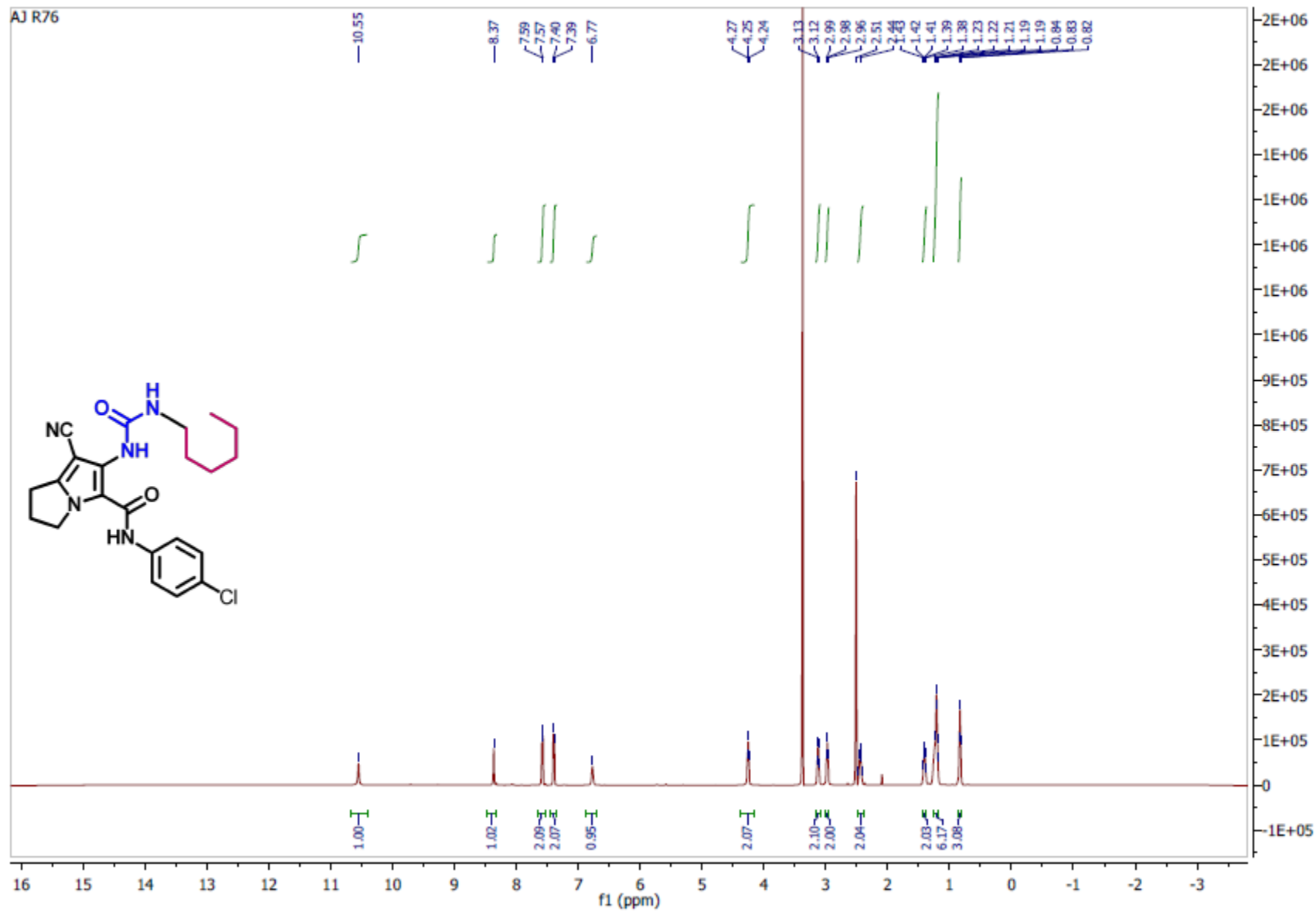


Figure S80. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound **16c** (zoom on aliphatic/aromatic Hs).

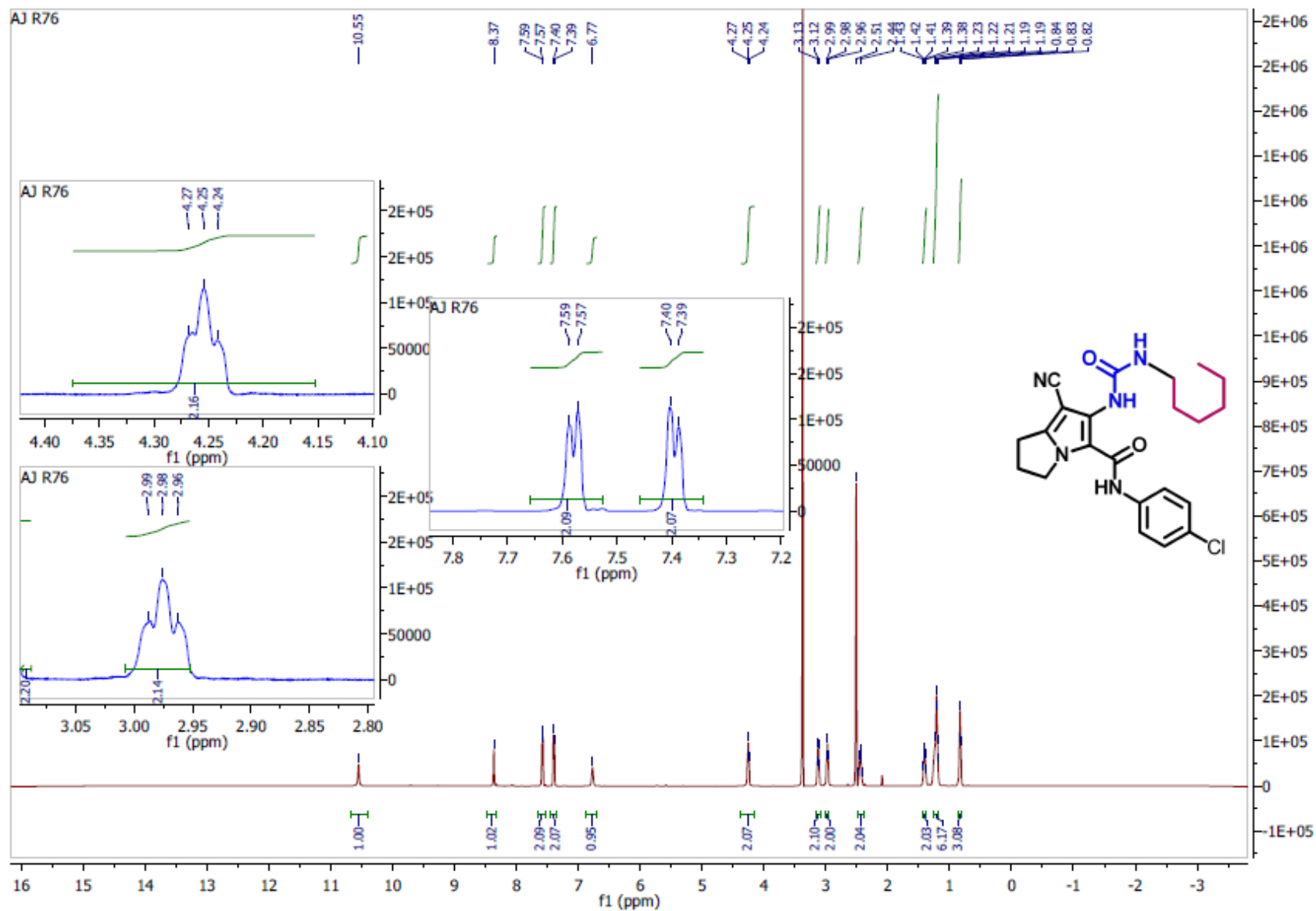


Figure S81. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **16c**.

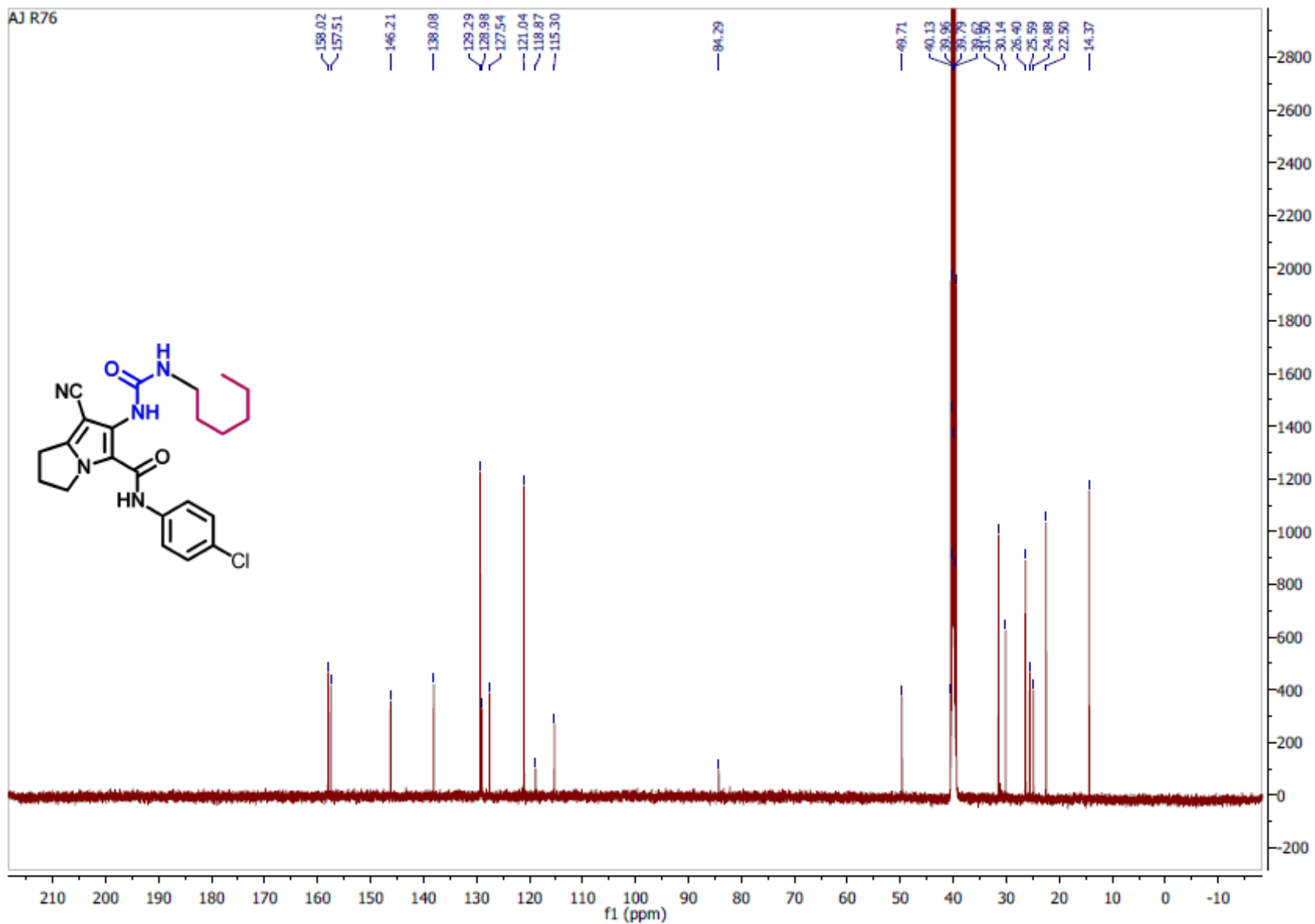


Figure S82. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **16c** (zoom on aliphatic/aromatic Cs).

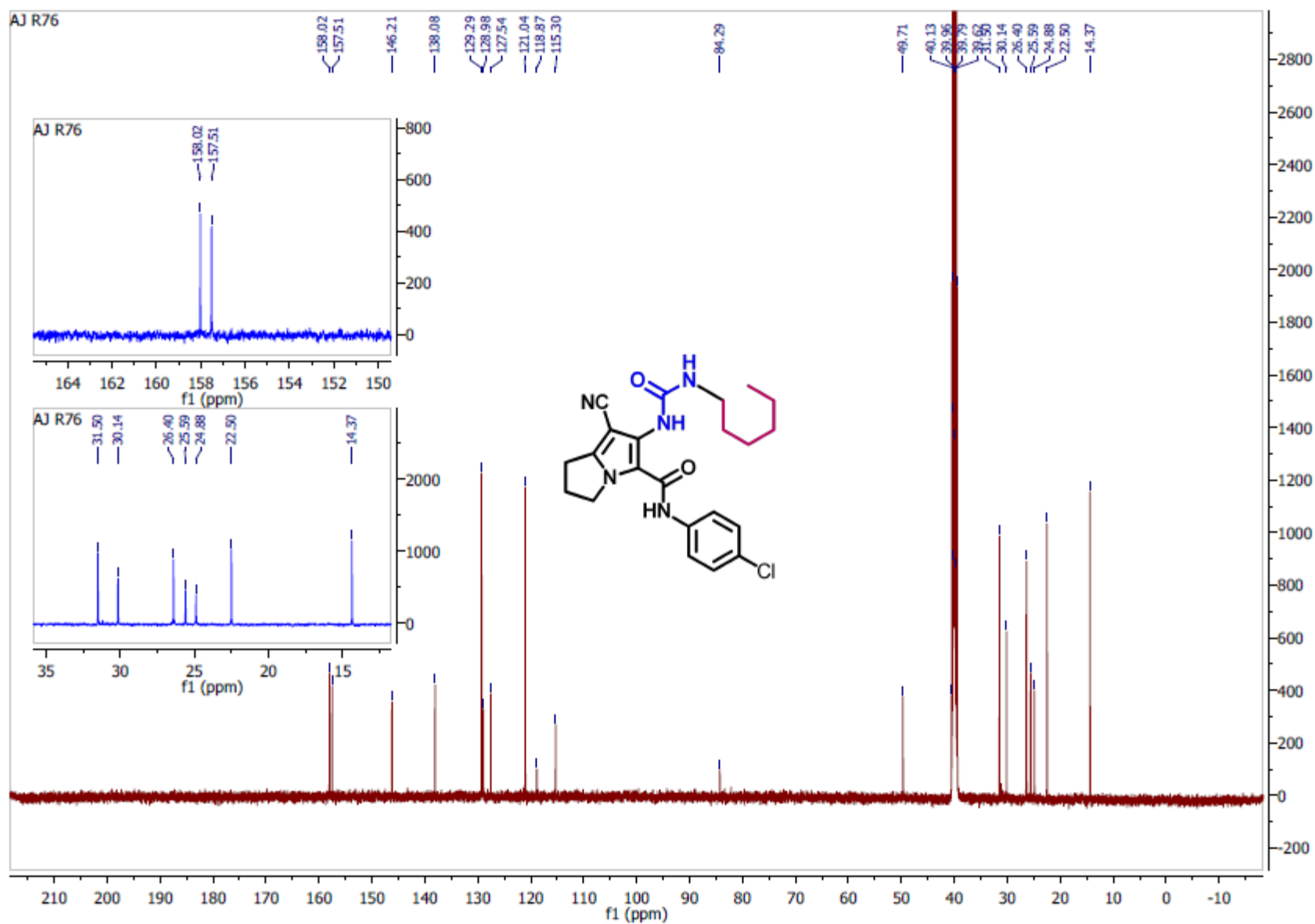


Figure S83. DEPT C^{135} (DMSO, 125 MHz, δ ppm) of compound **16c**.

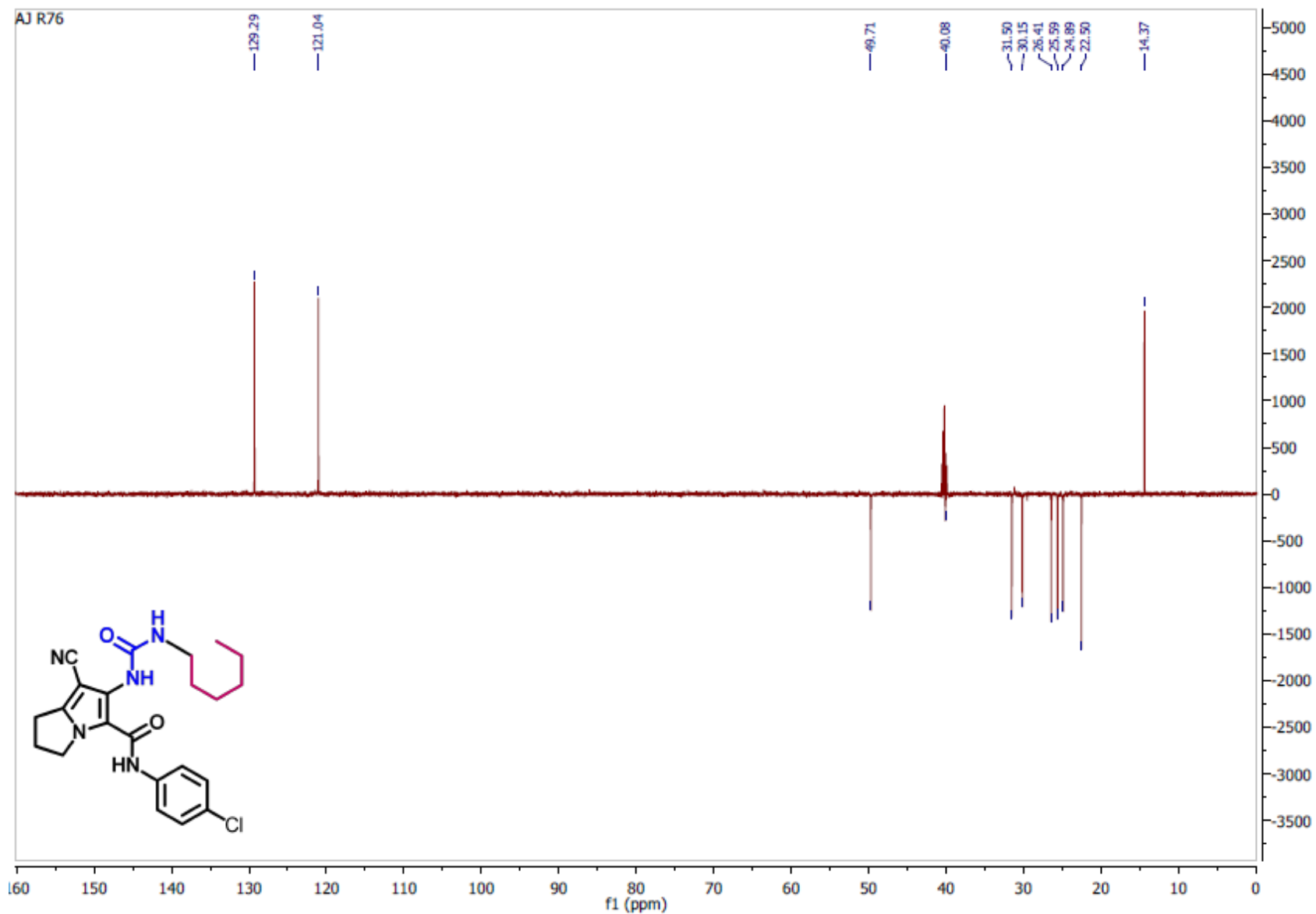


Figure S84. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **17a**.

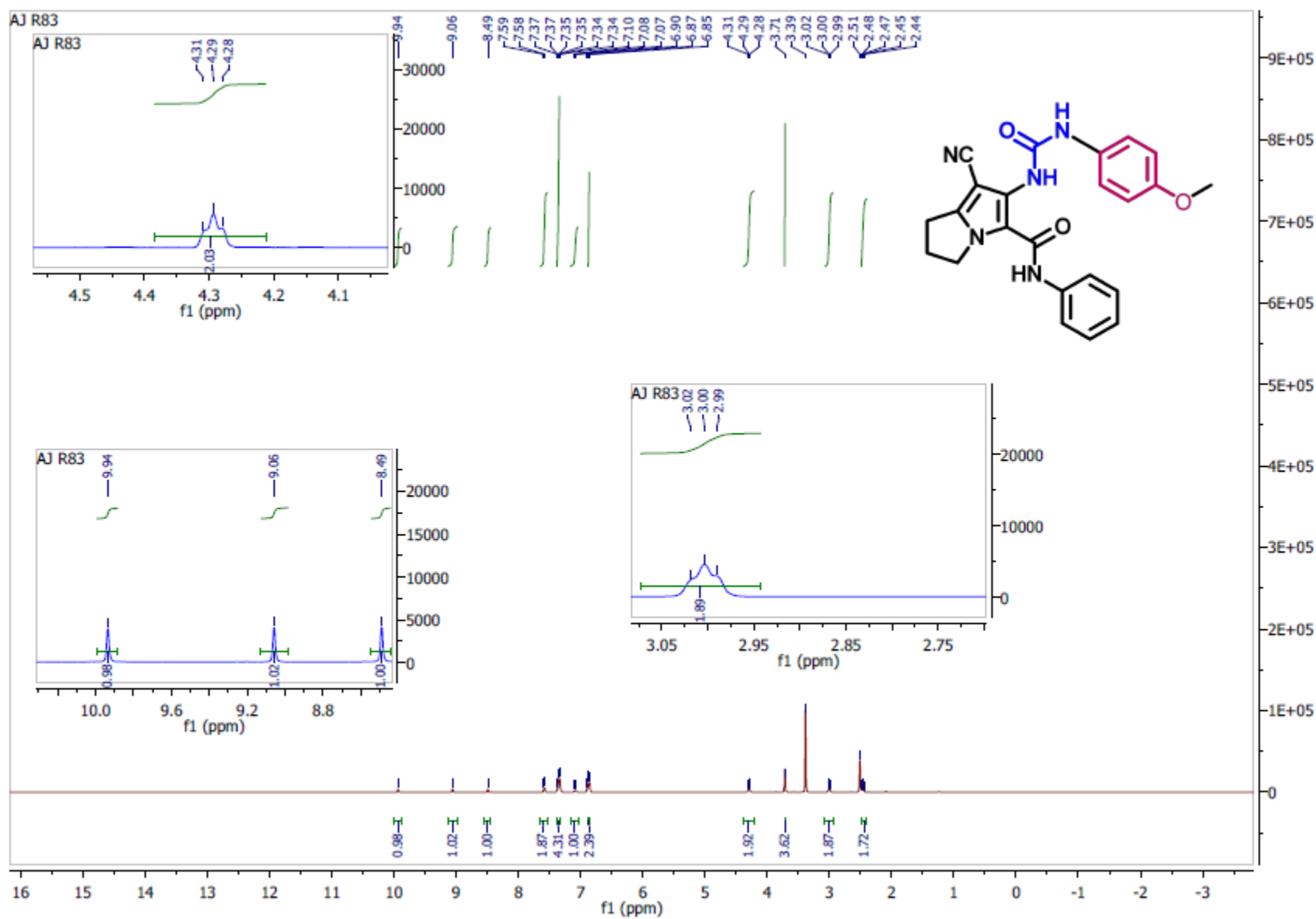


Figure S85. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **17a**.

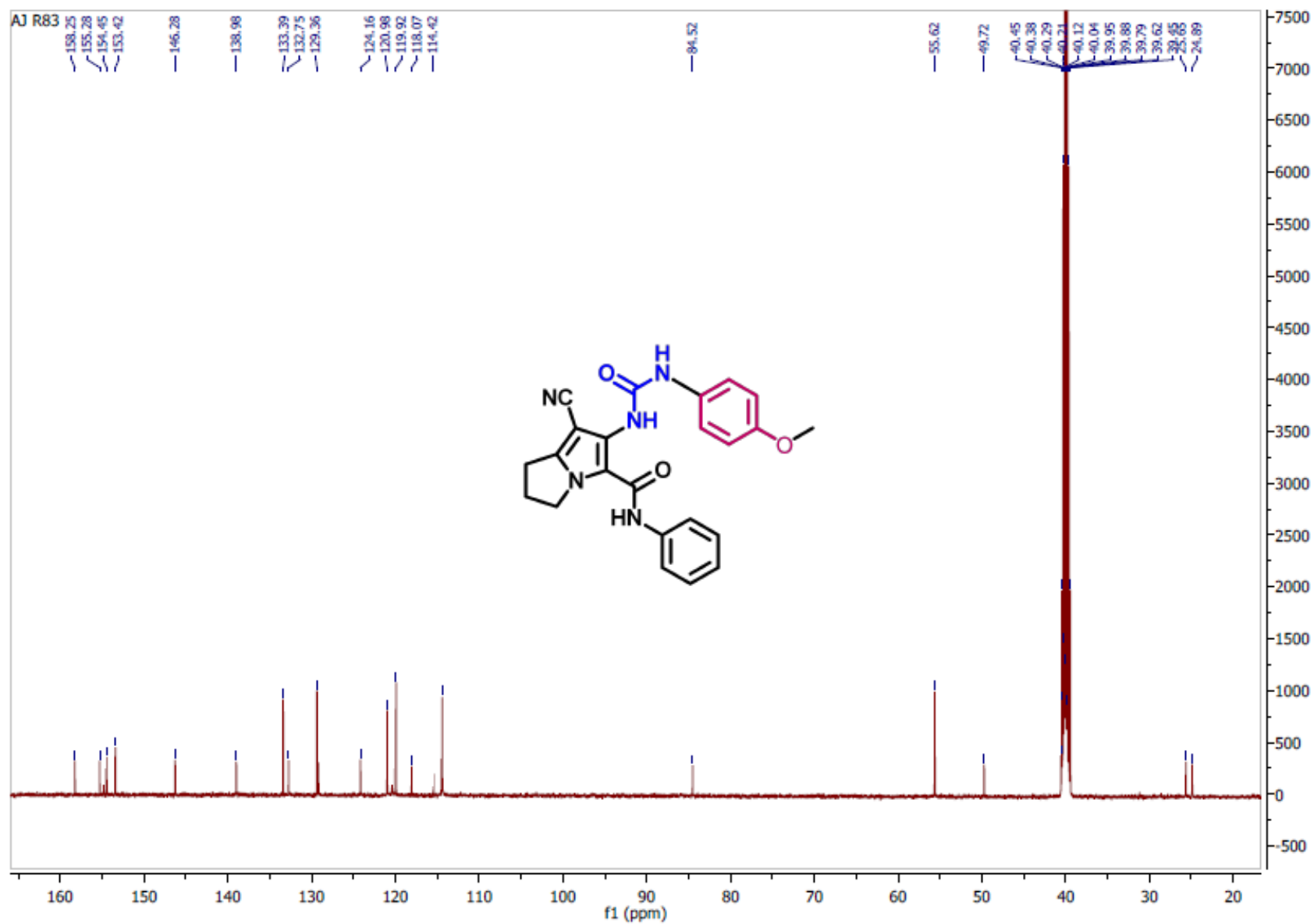


Figure S86. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound 17a.

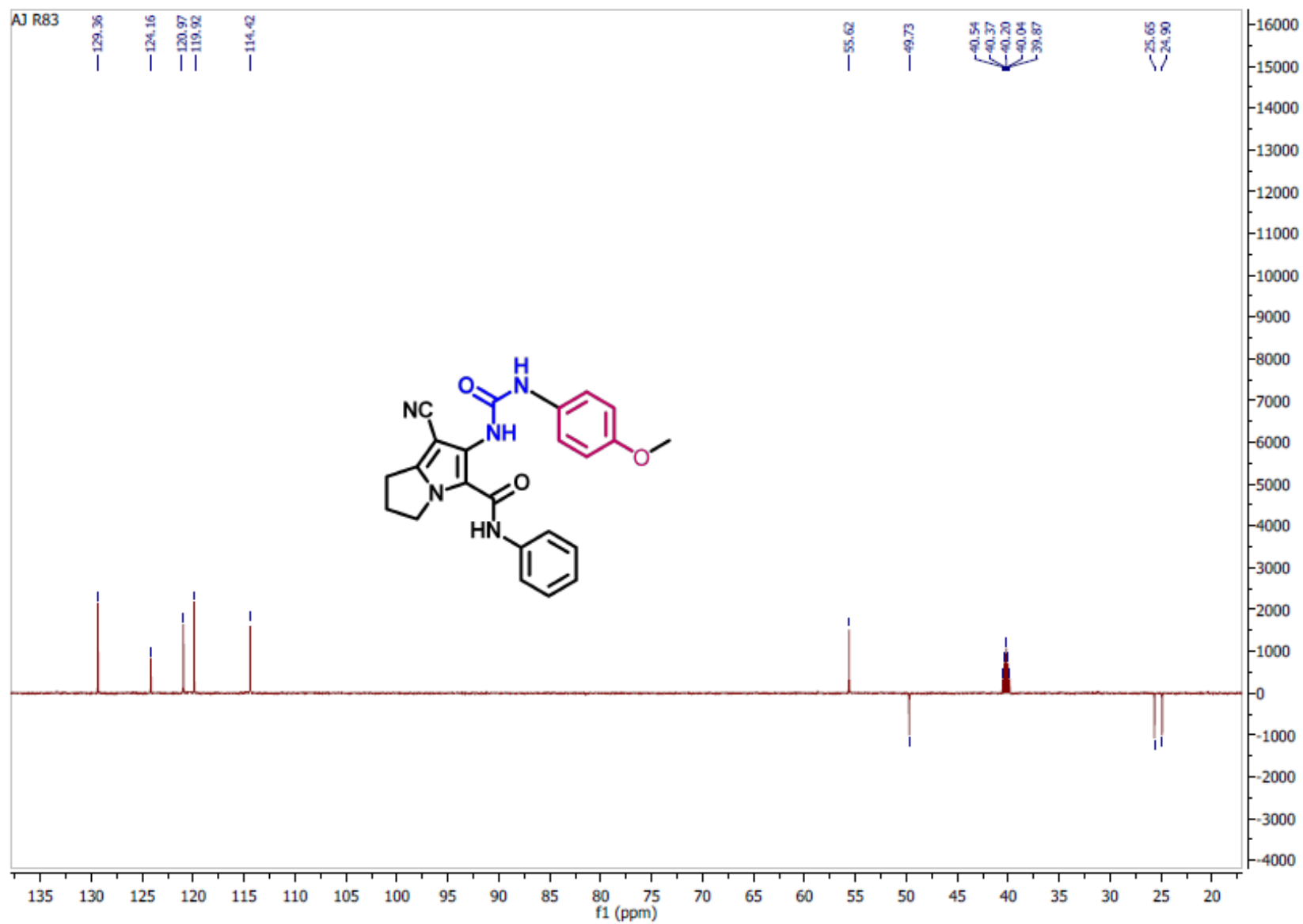


Figure S87. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **17b**

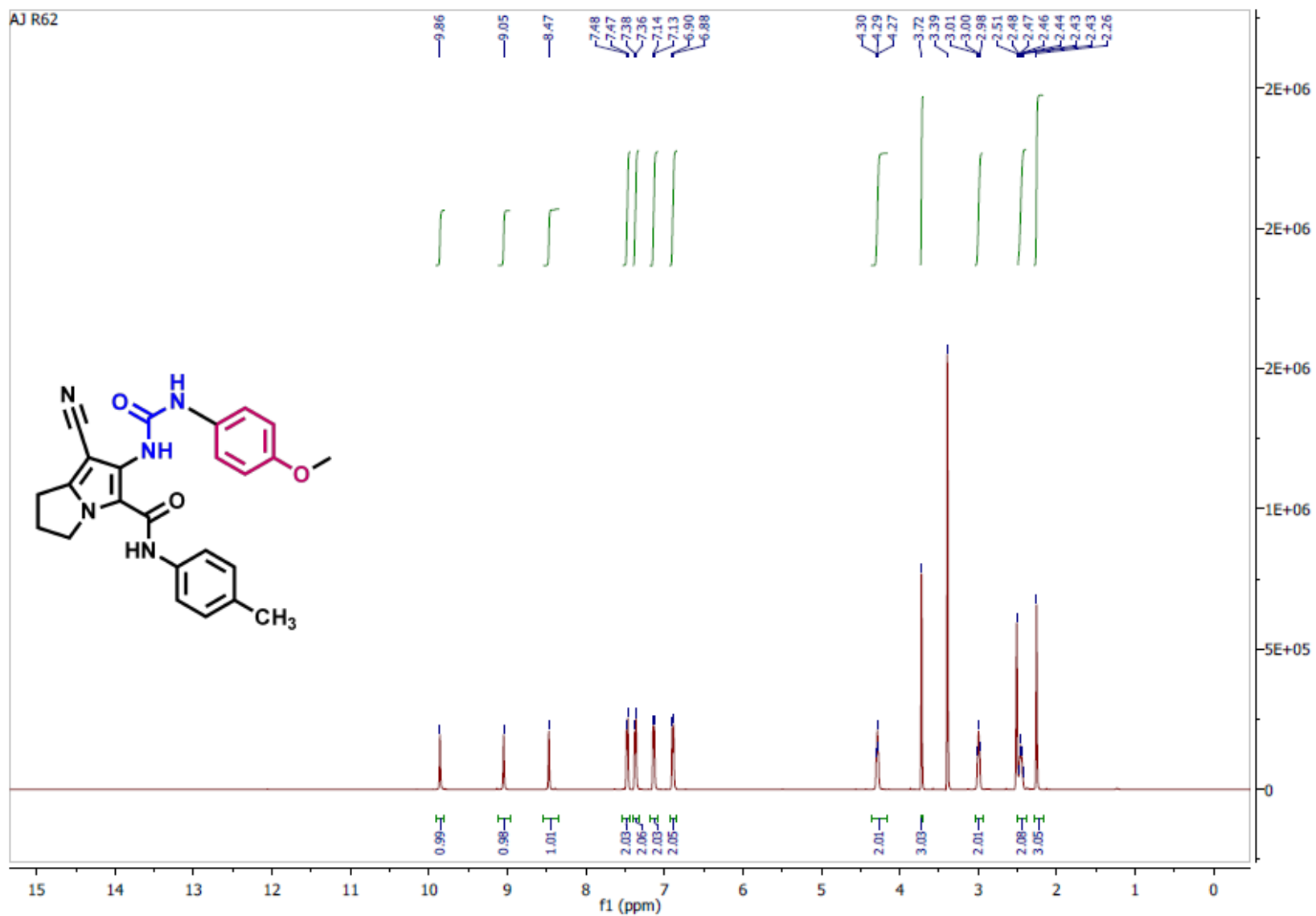


Figure S88. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **17b** (zoom on aliphatic Hs).

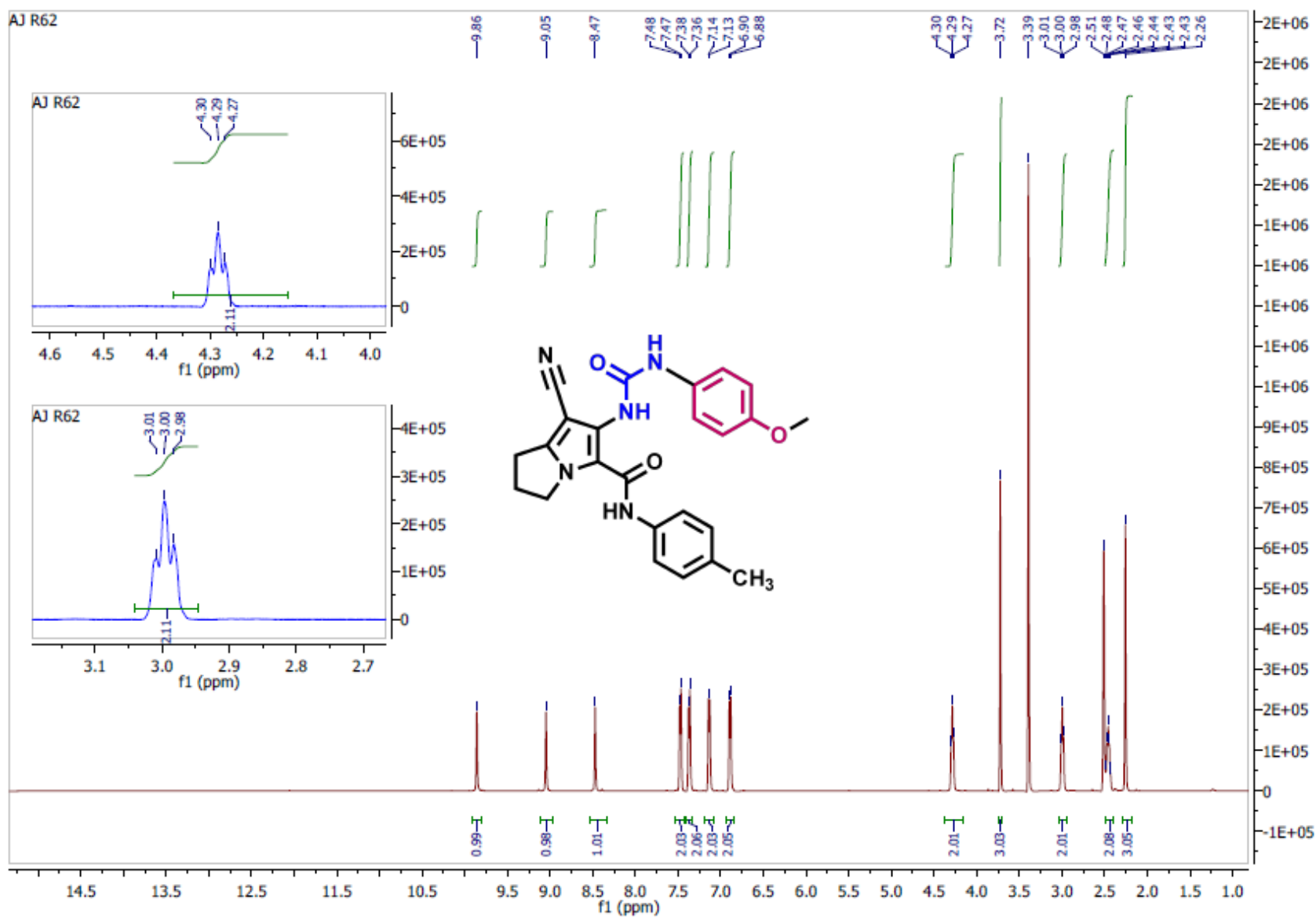


Figure S89. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **17b** (zoom on aliphatic/aromatic Hs).

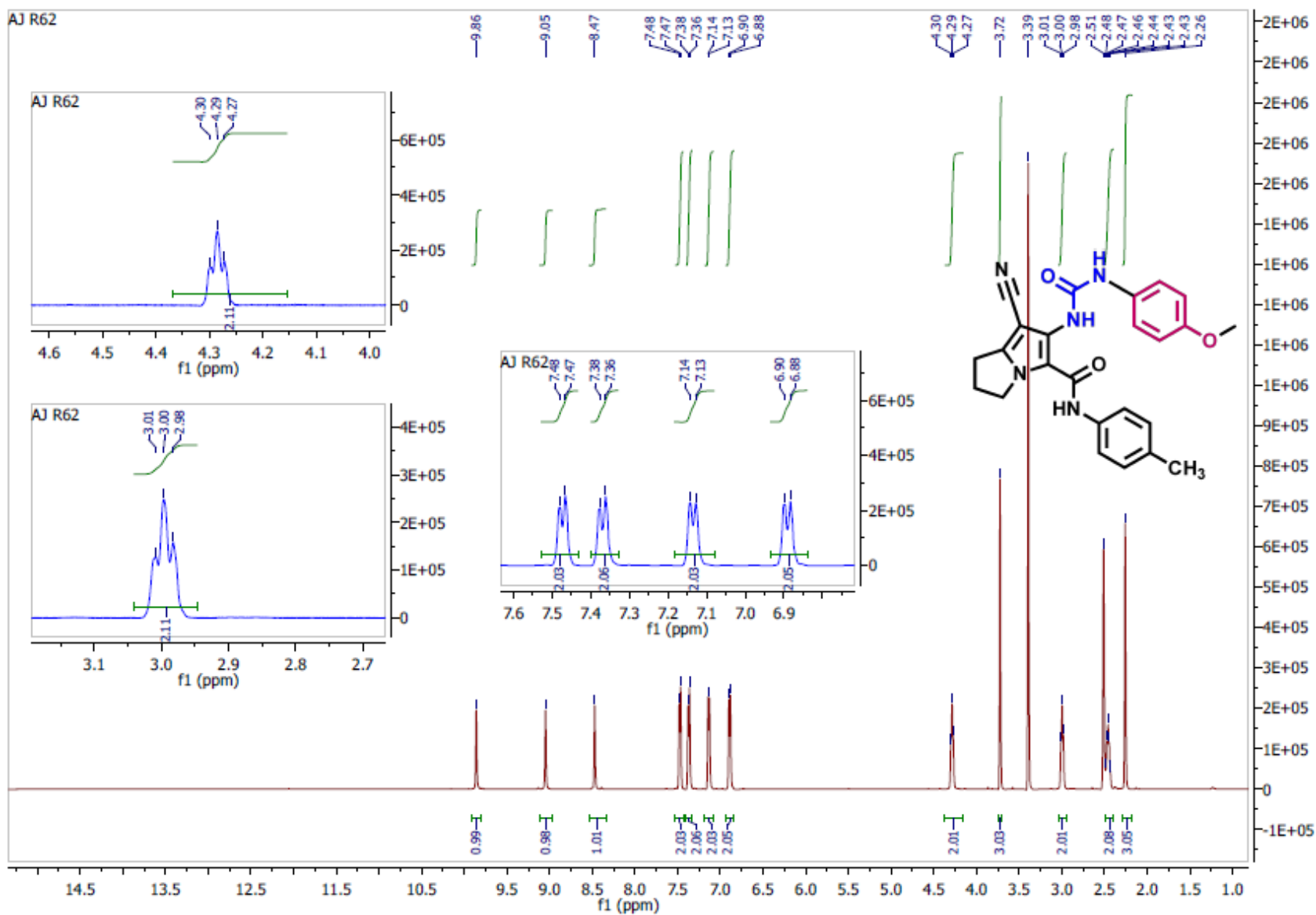


Figure S90. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **17b**.

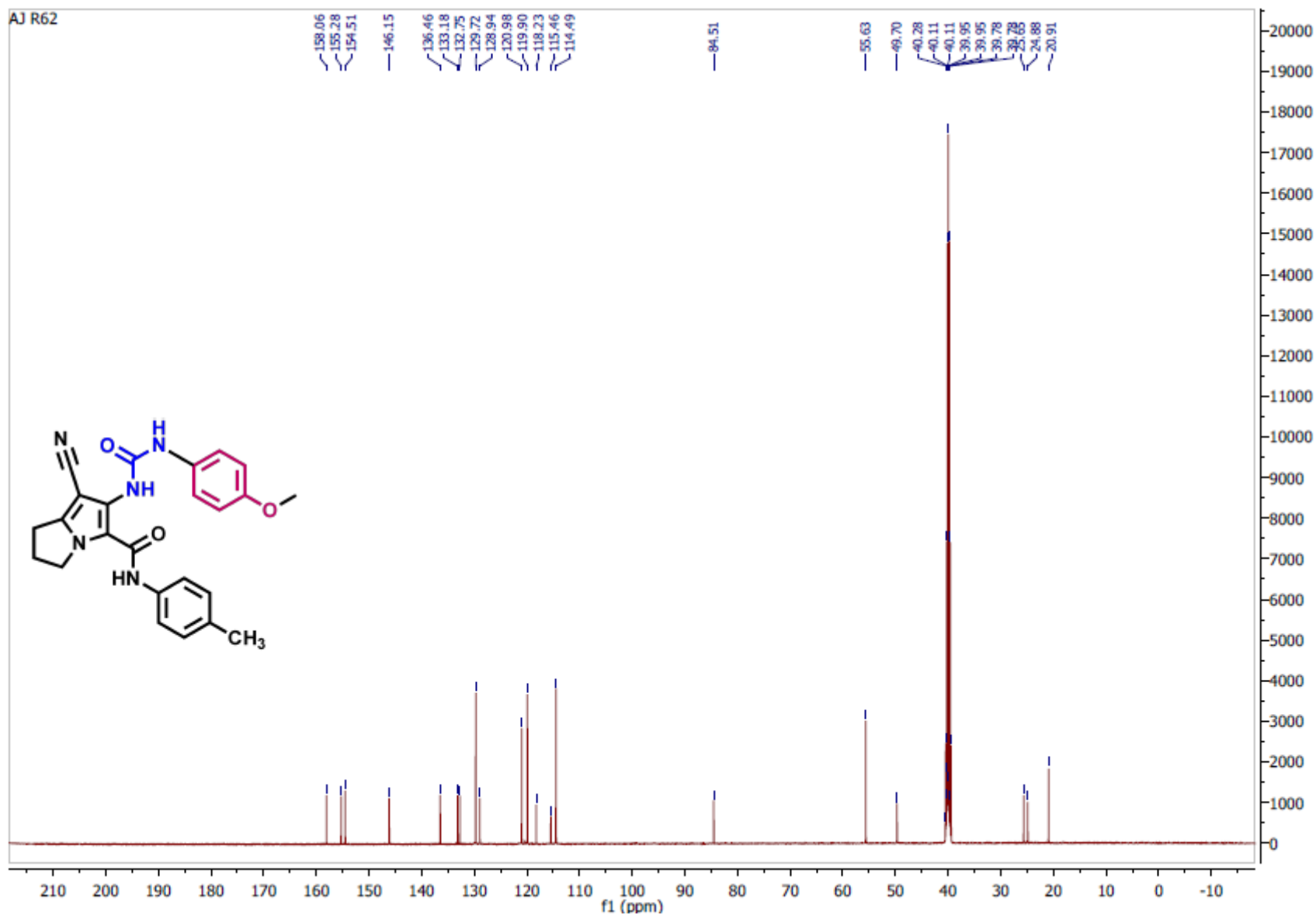


Figure S91. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound **17b**.

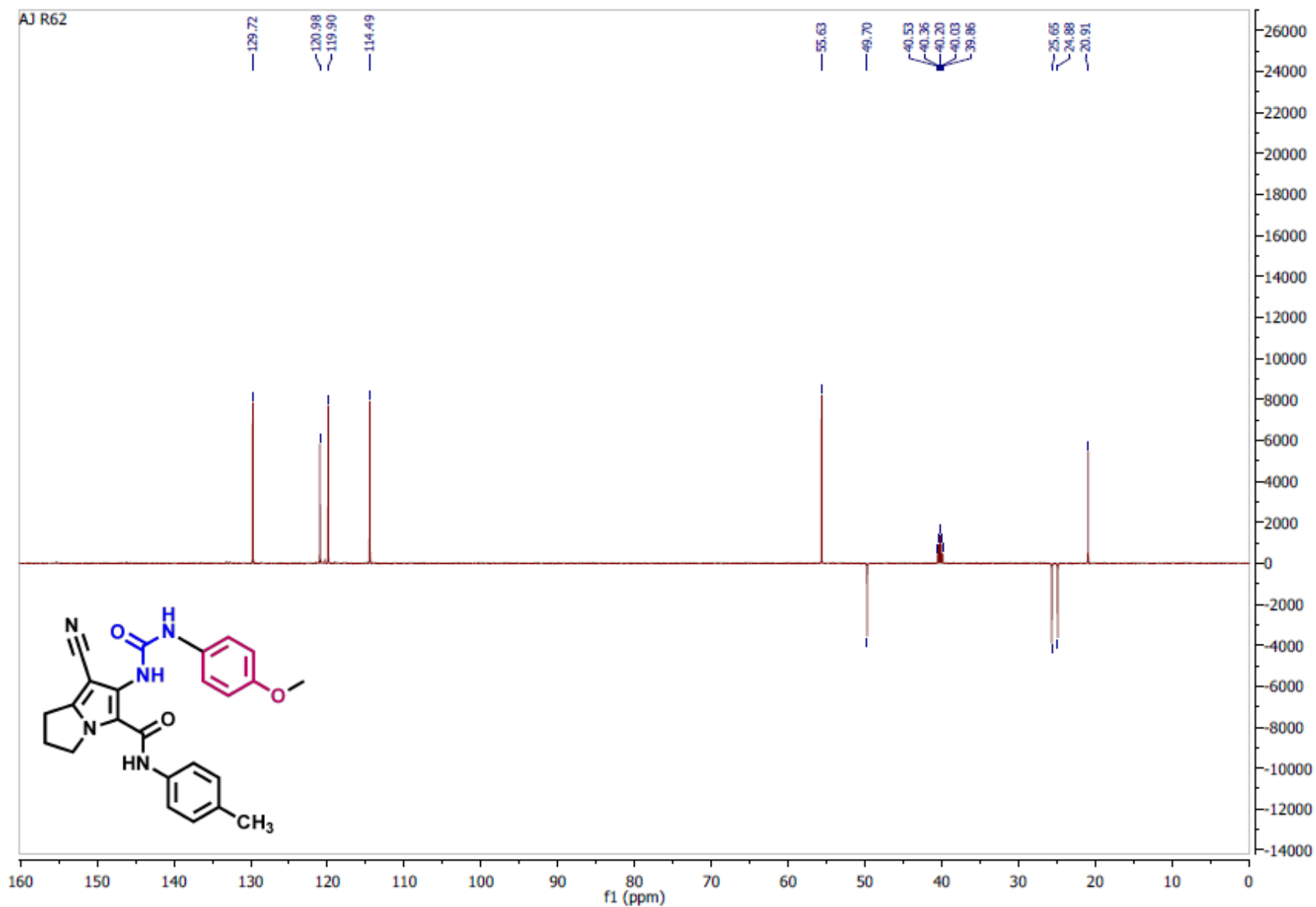


Figure S92. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **18a**

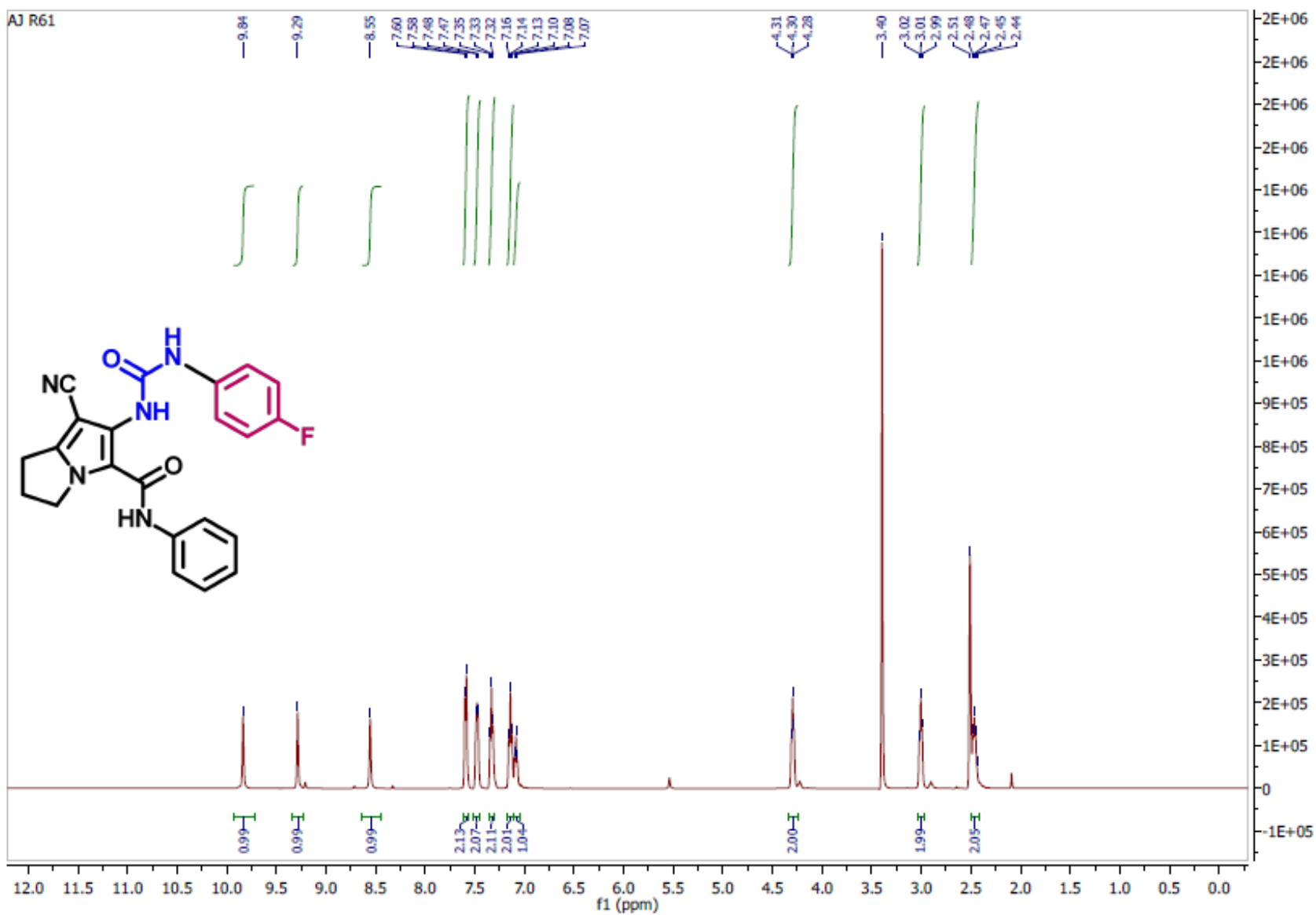


Figure S93. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound **18a** (ZOOM on aliphatic Hs)

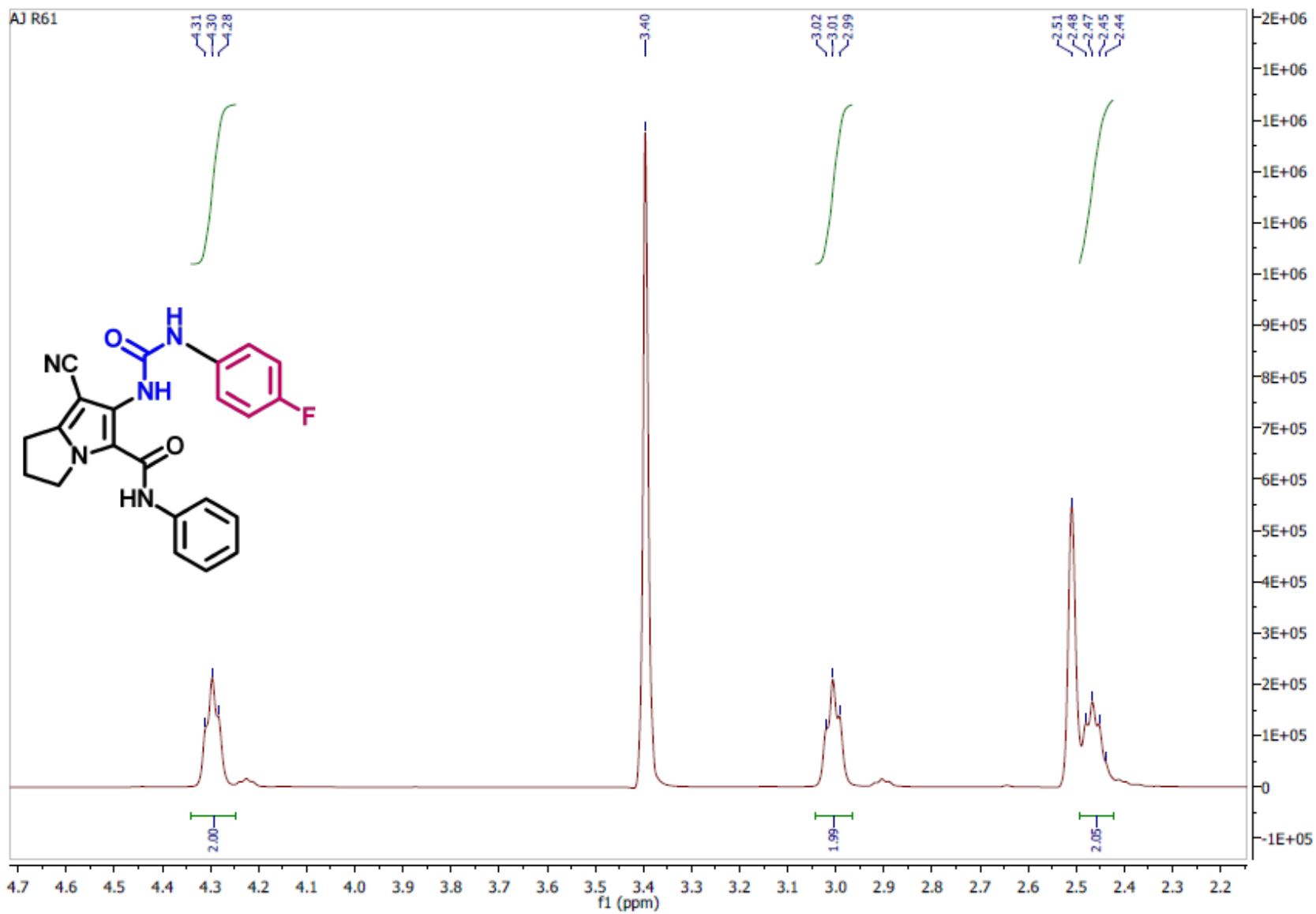


Figure S94. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **18a** (**ZOOM on NHs & aromatic Hs**)

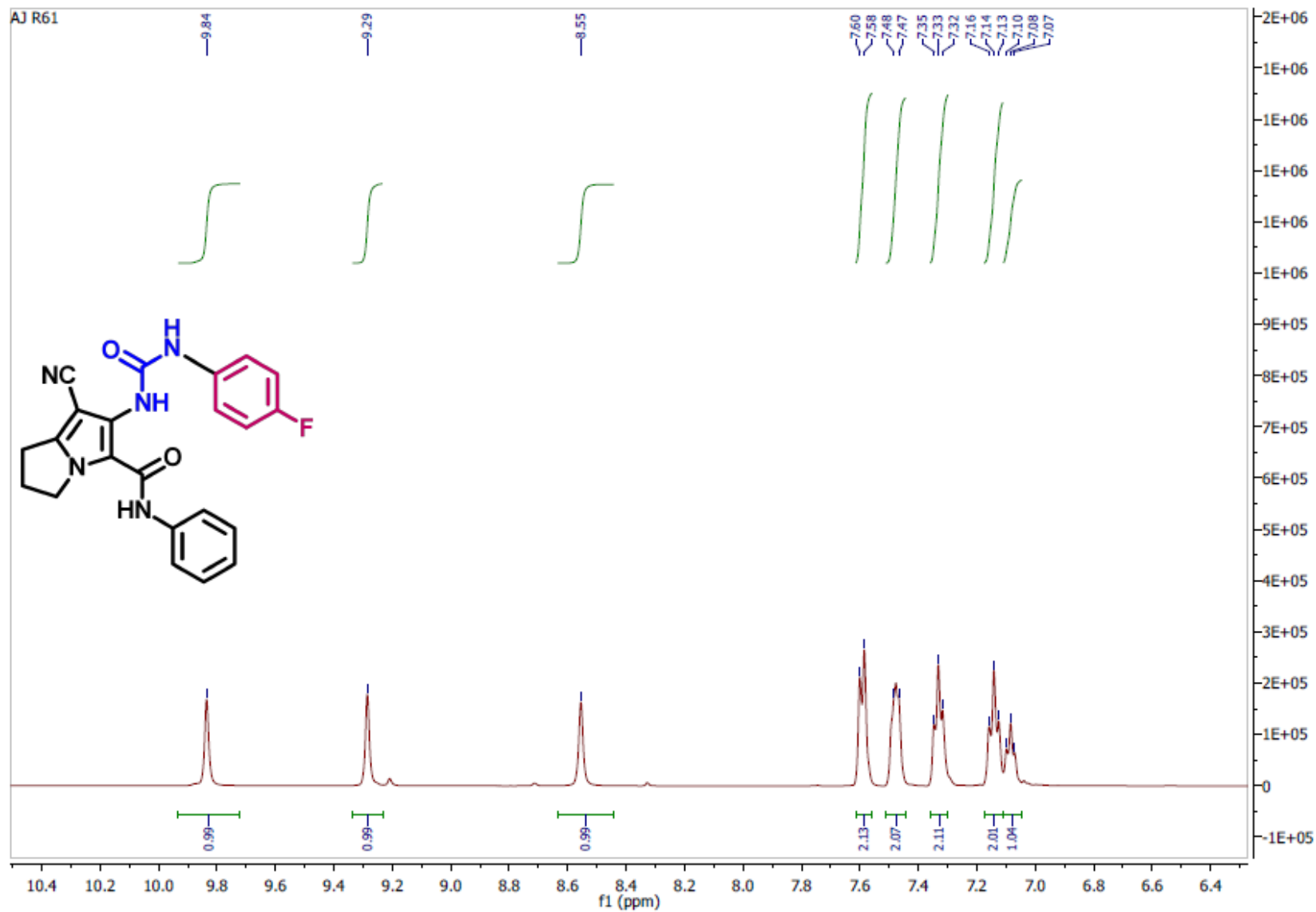


Figure S95. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **16a** (ZOOM on aromatic Hs)

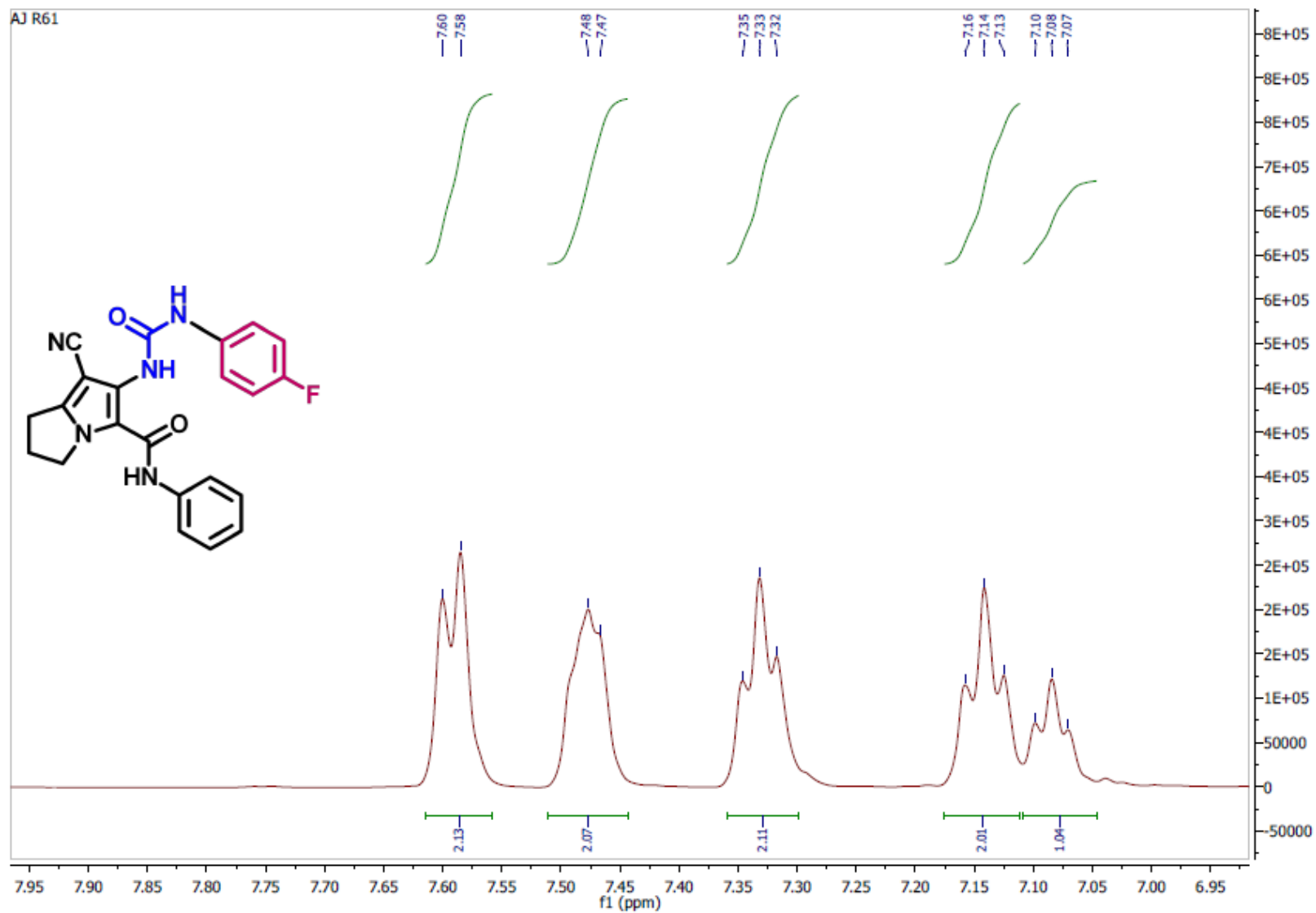


Figure S96. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18a**

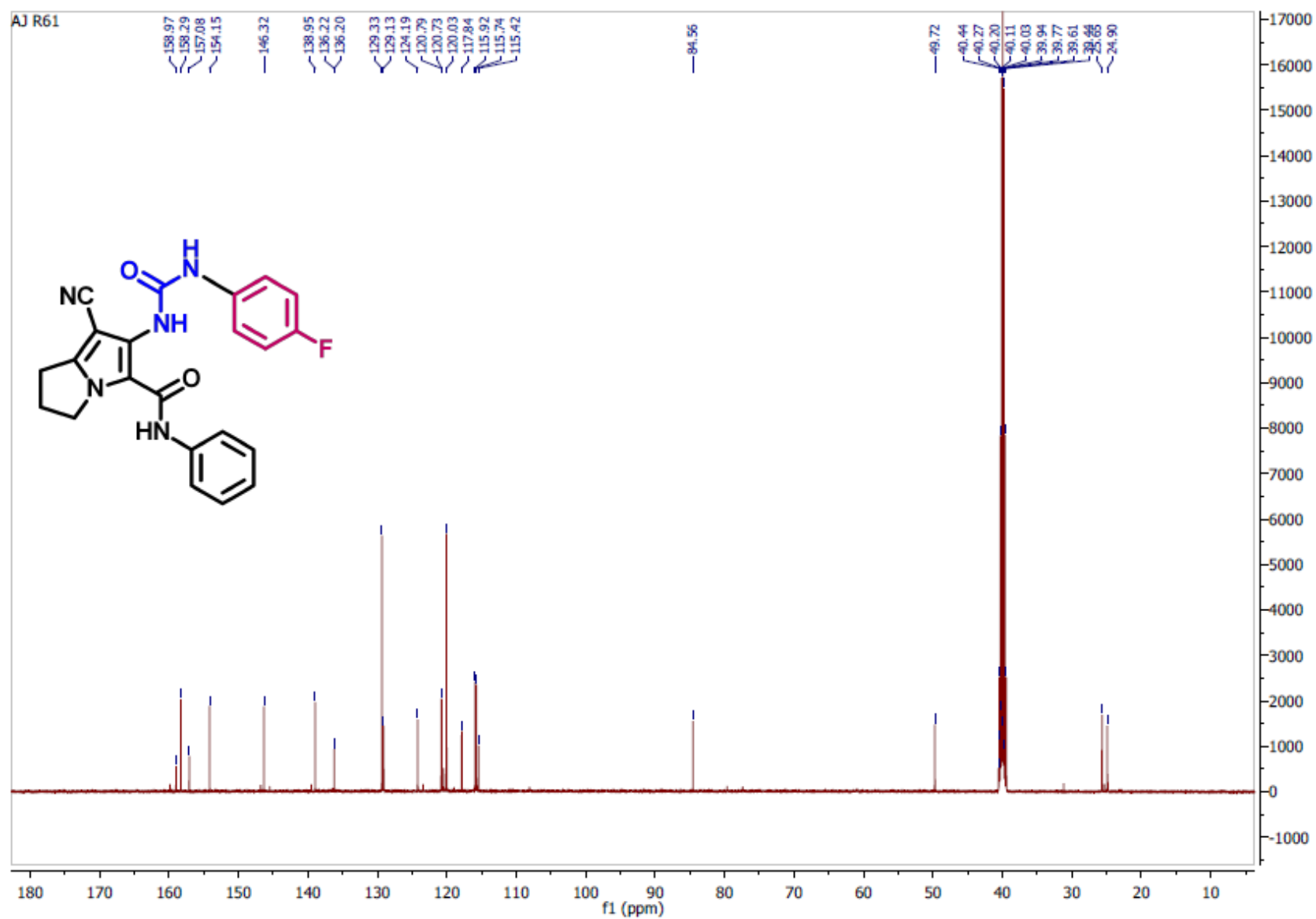


Figure S97. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18a** (ZOOM on aliphatic Cs)

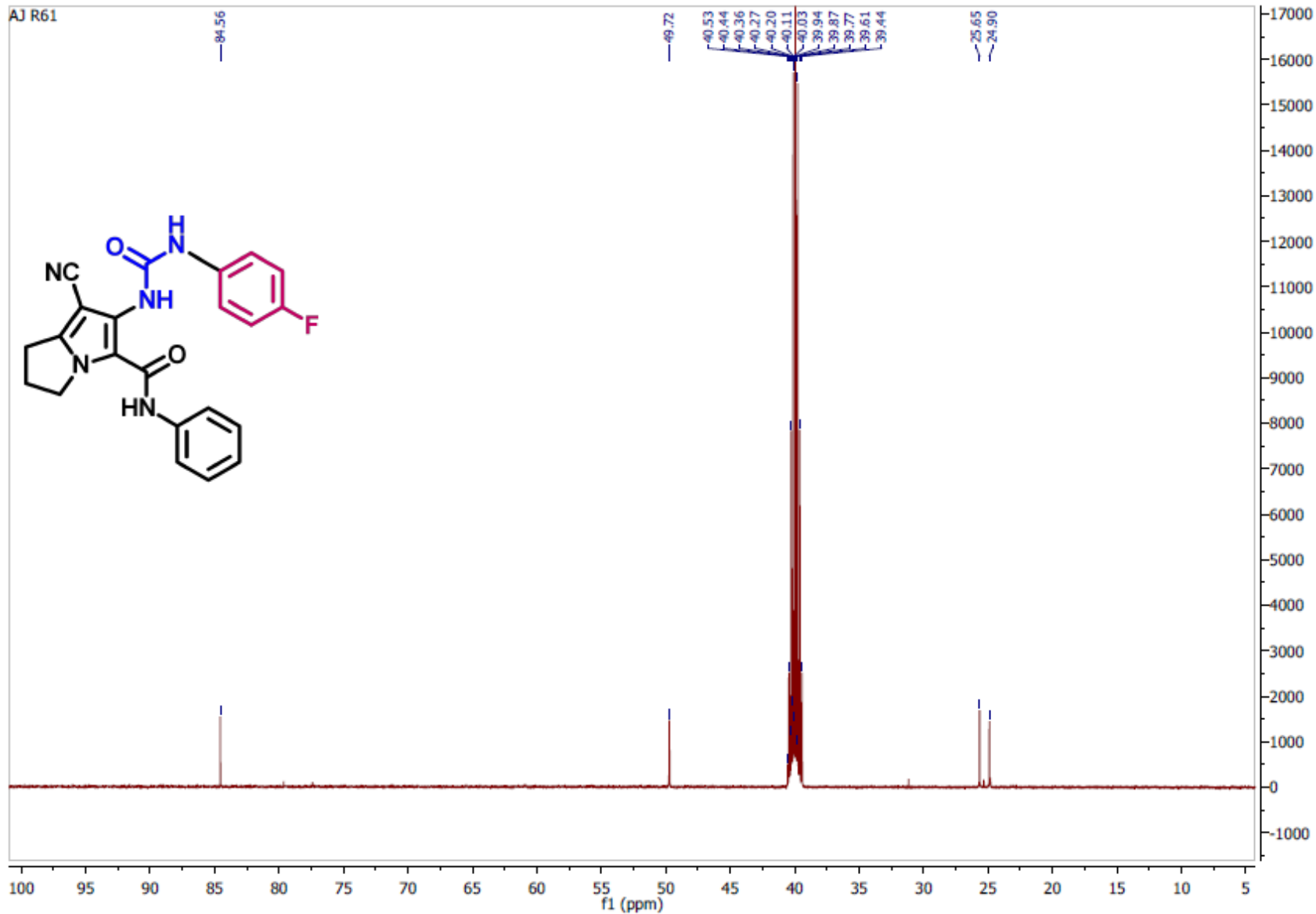


Figure S98. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18a** (**ZOOM on aromatic Cs**)

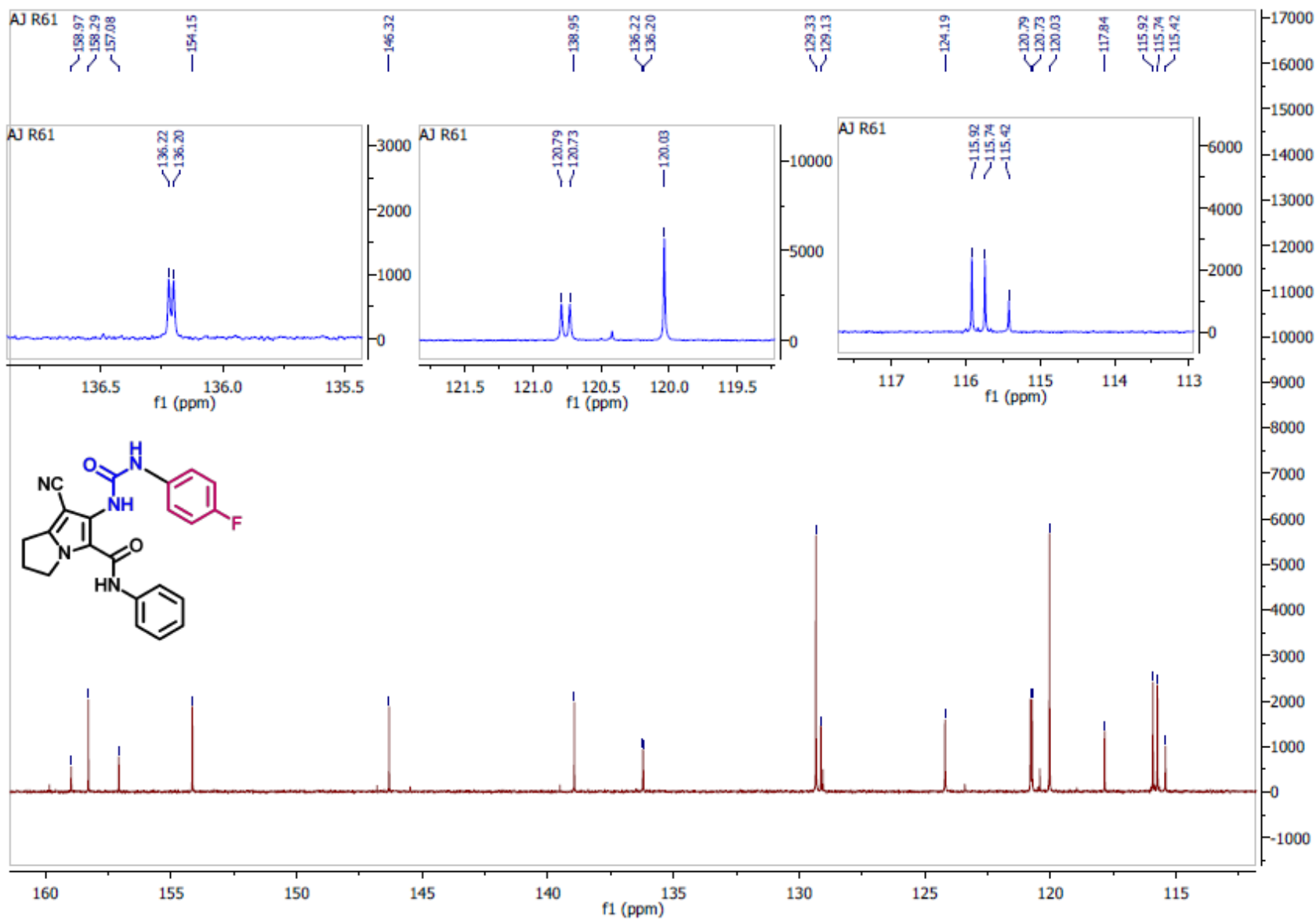


Figure S99. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound **18a**.

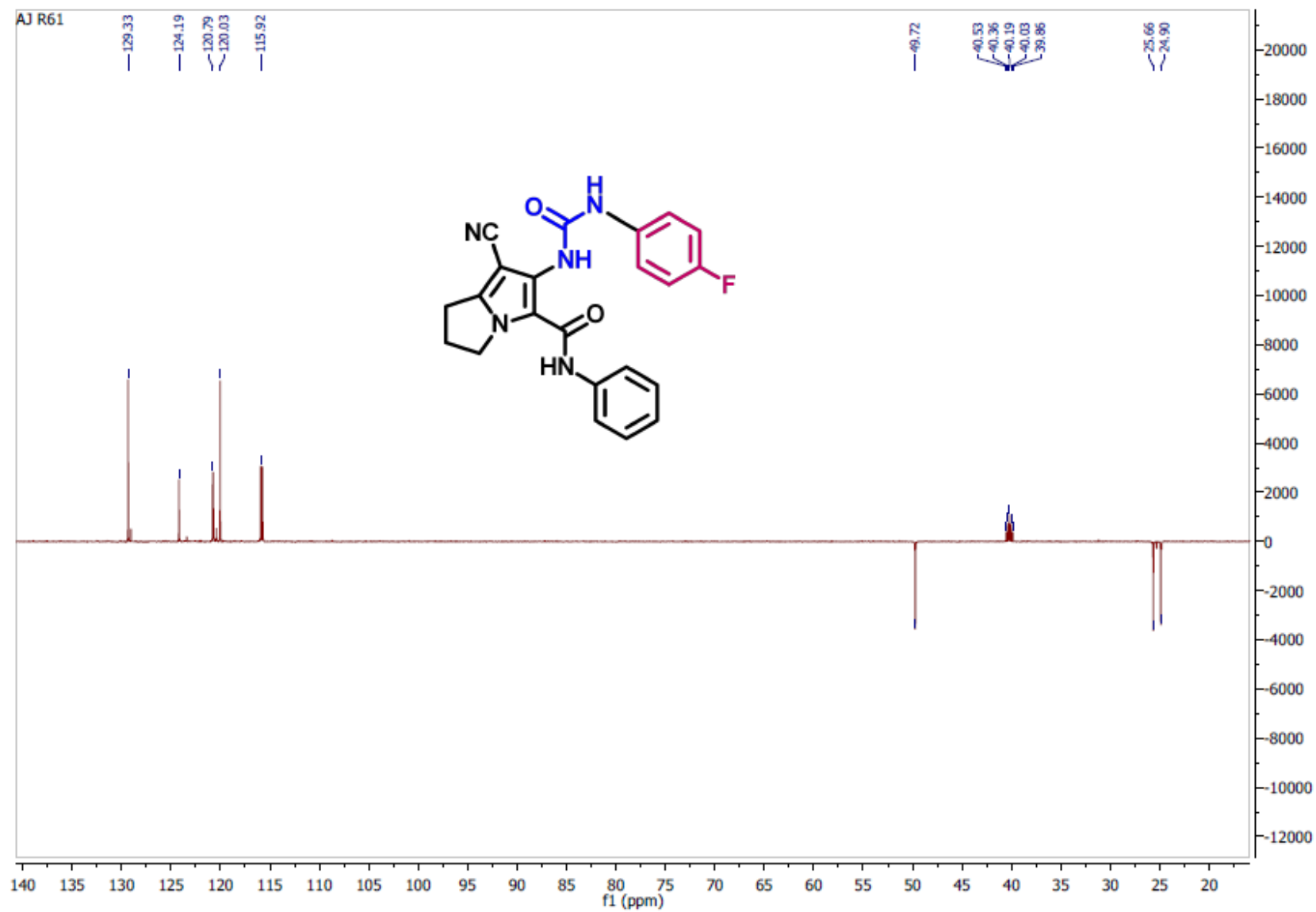


Figure S100. DEPT C^{135} spectrum of compound **18a** (zoom, aliphatic CH_2 groups)



Figure S101. DEPT C¹³⁵ spectrum of compound **18a** (zoom, aromatic CH groups)

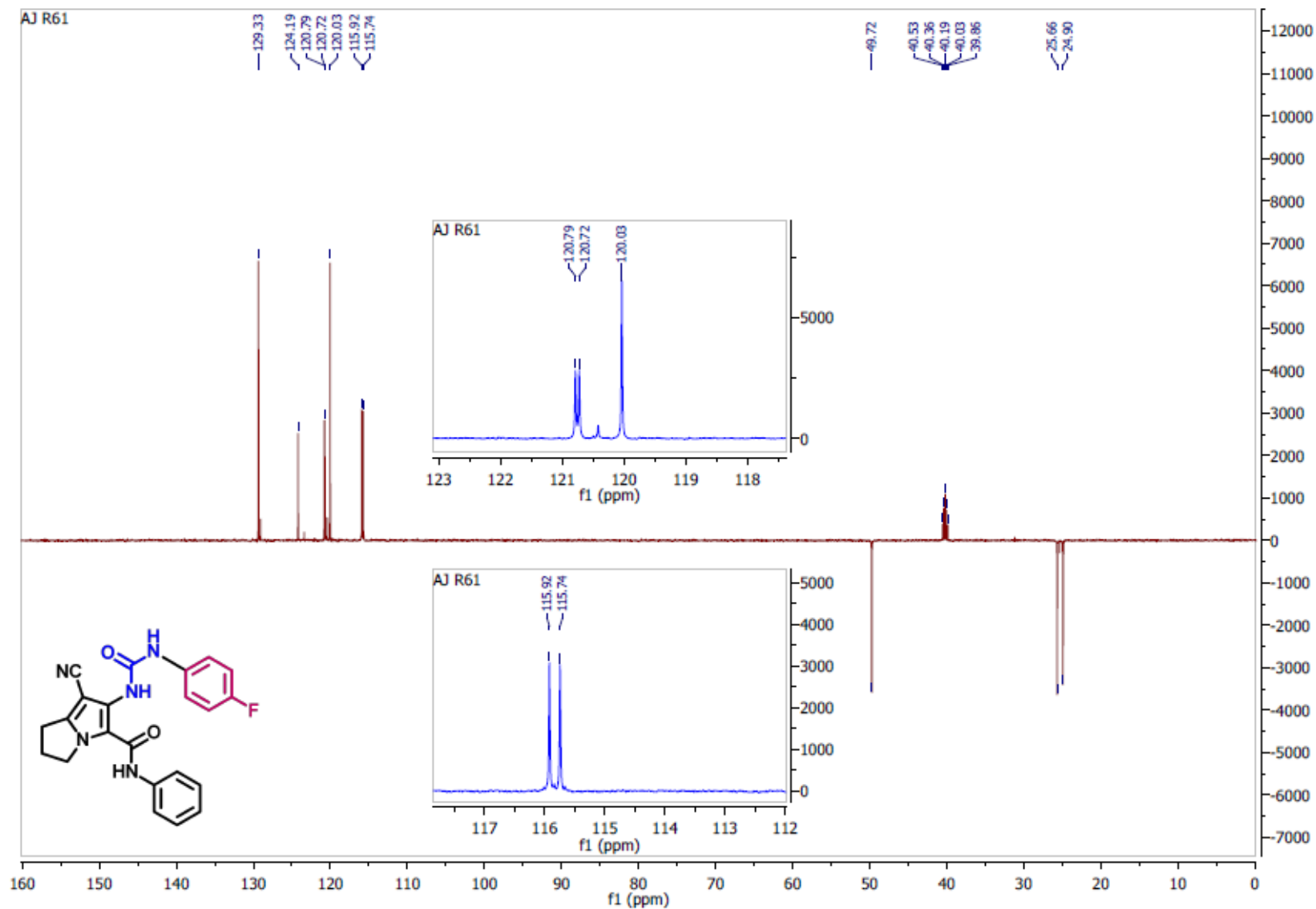


Figure S102. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound **18b**

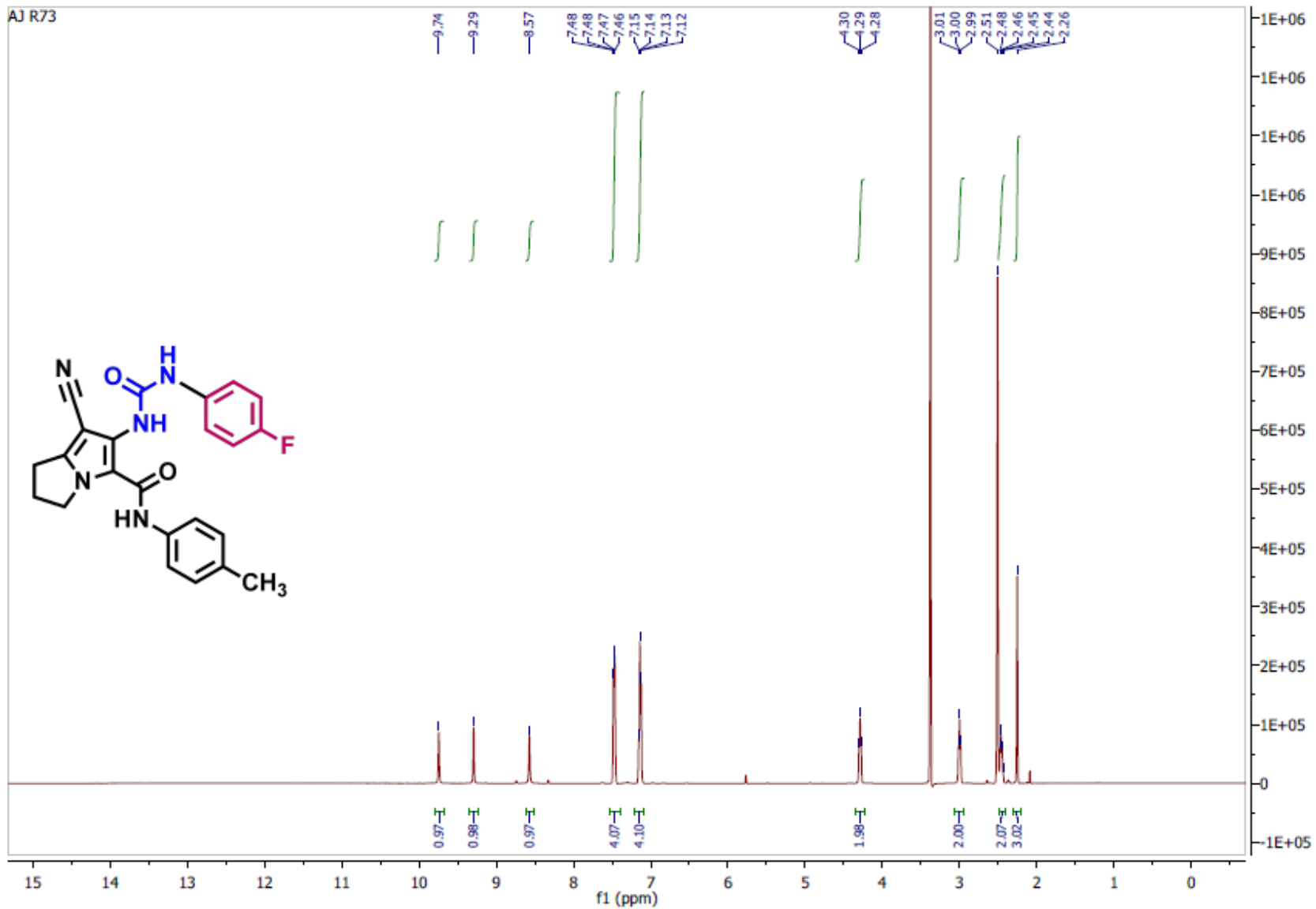


Figure S103. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **18b** (zoom on aliphatic/aromatic Hs)

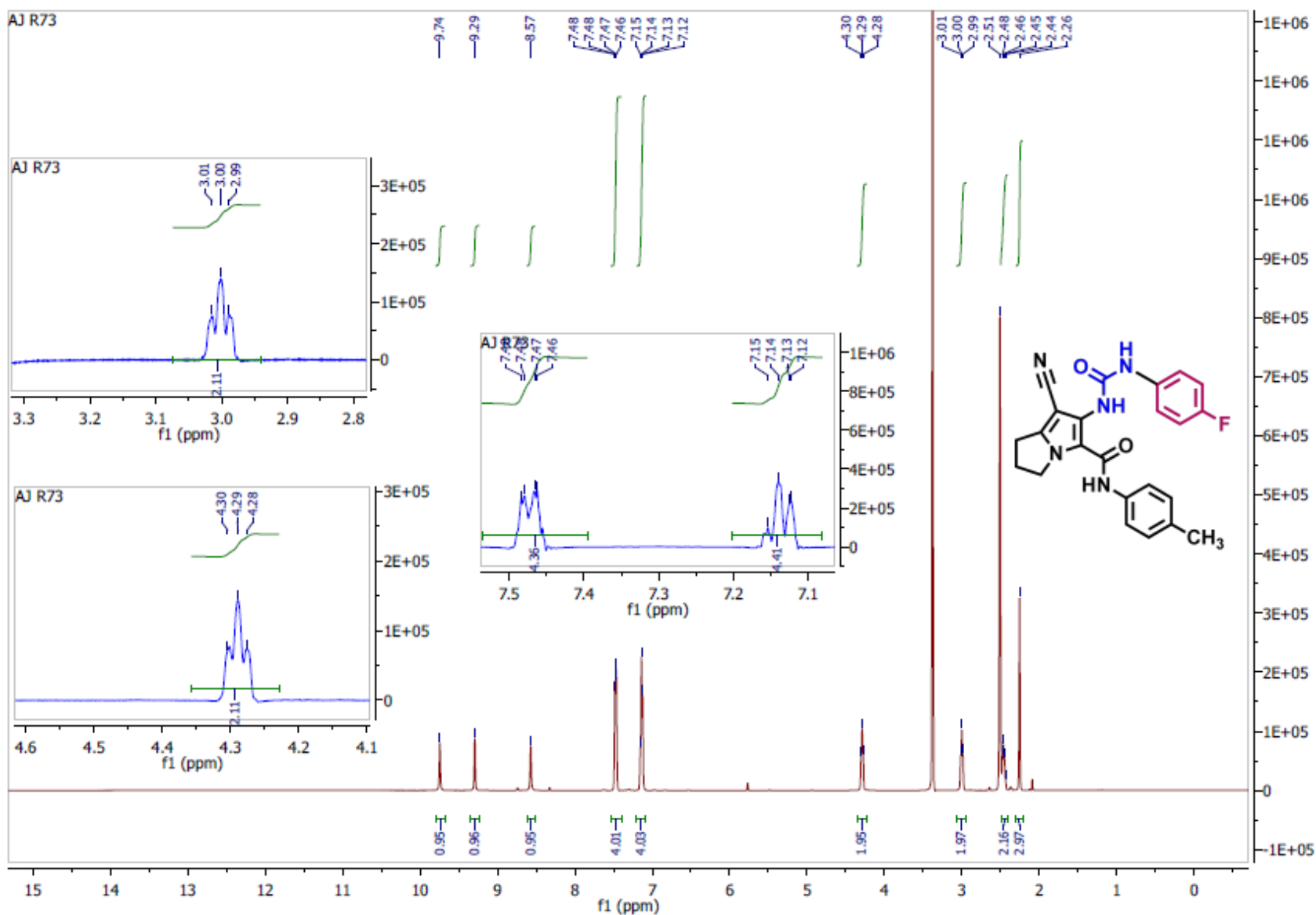


Figure S104. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18b**

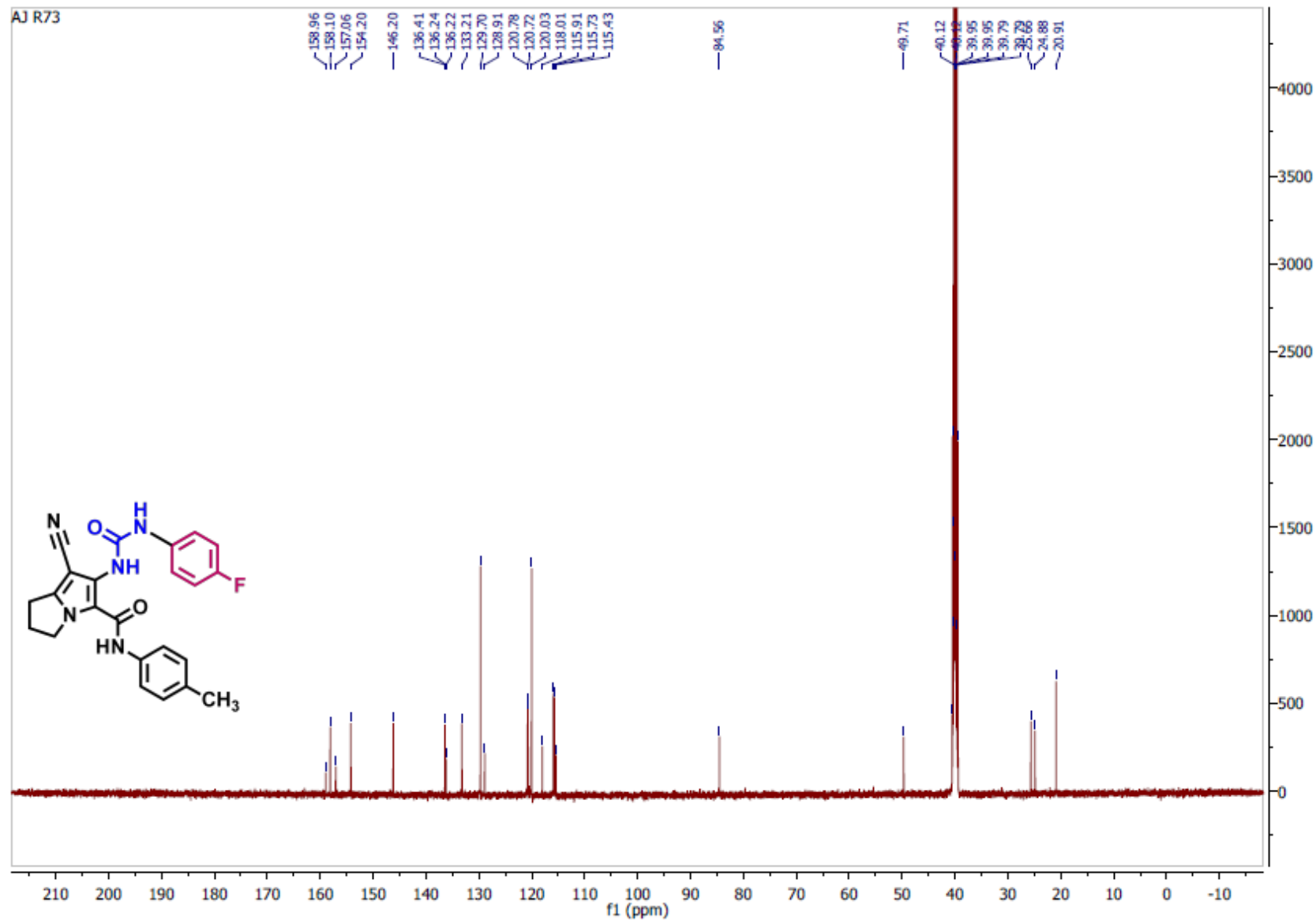


Figure S105. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18b** (zoom on aliphatic Cs)

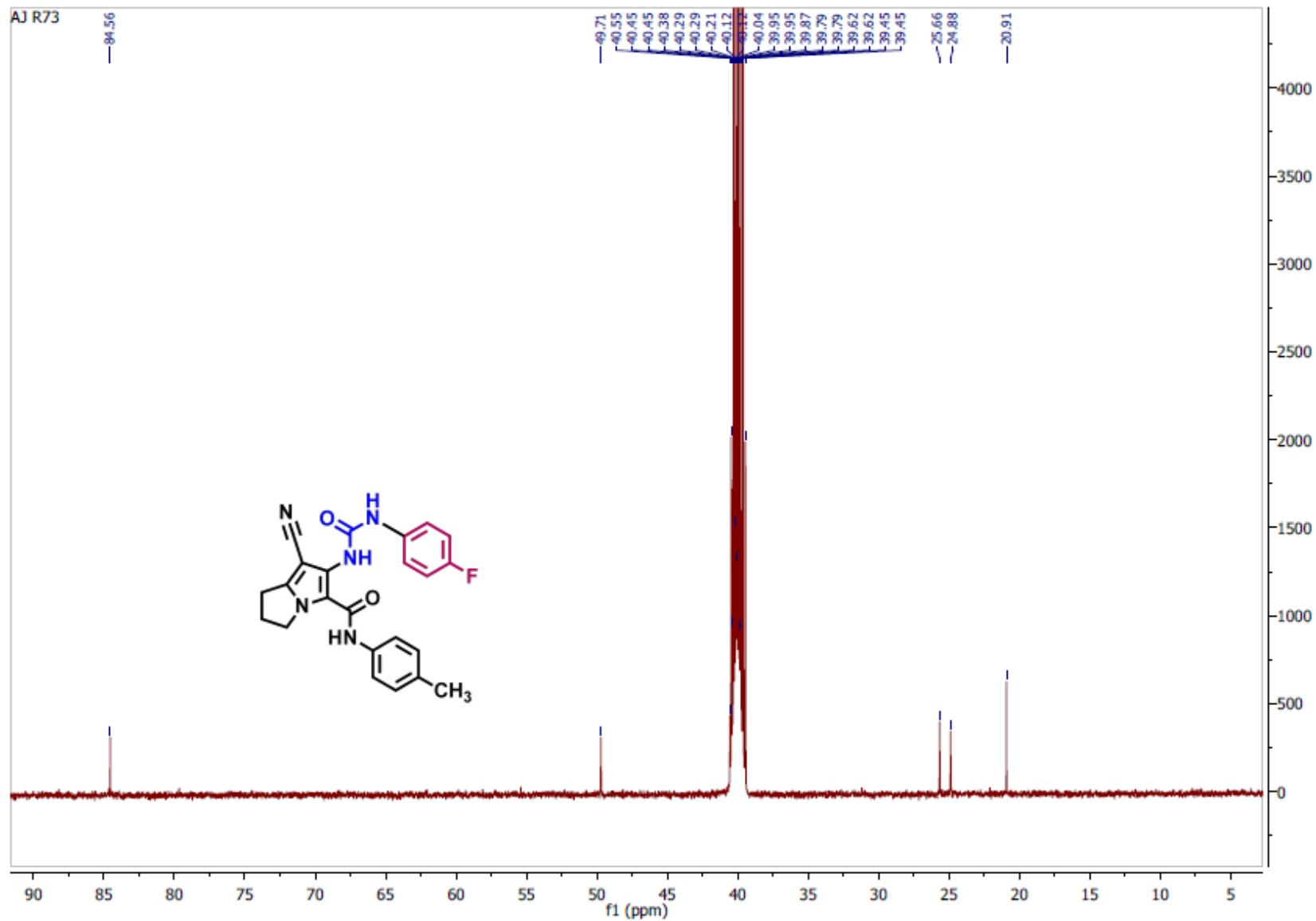


Figure S106. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18b** (zoom on aromatic Cs)

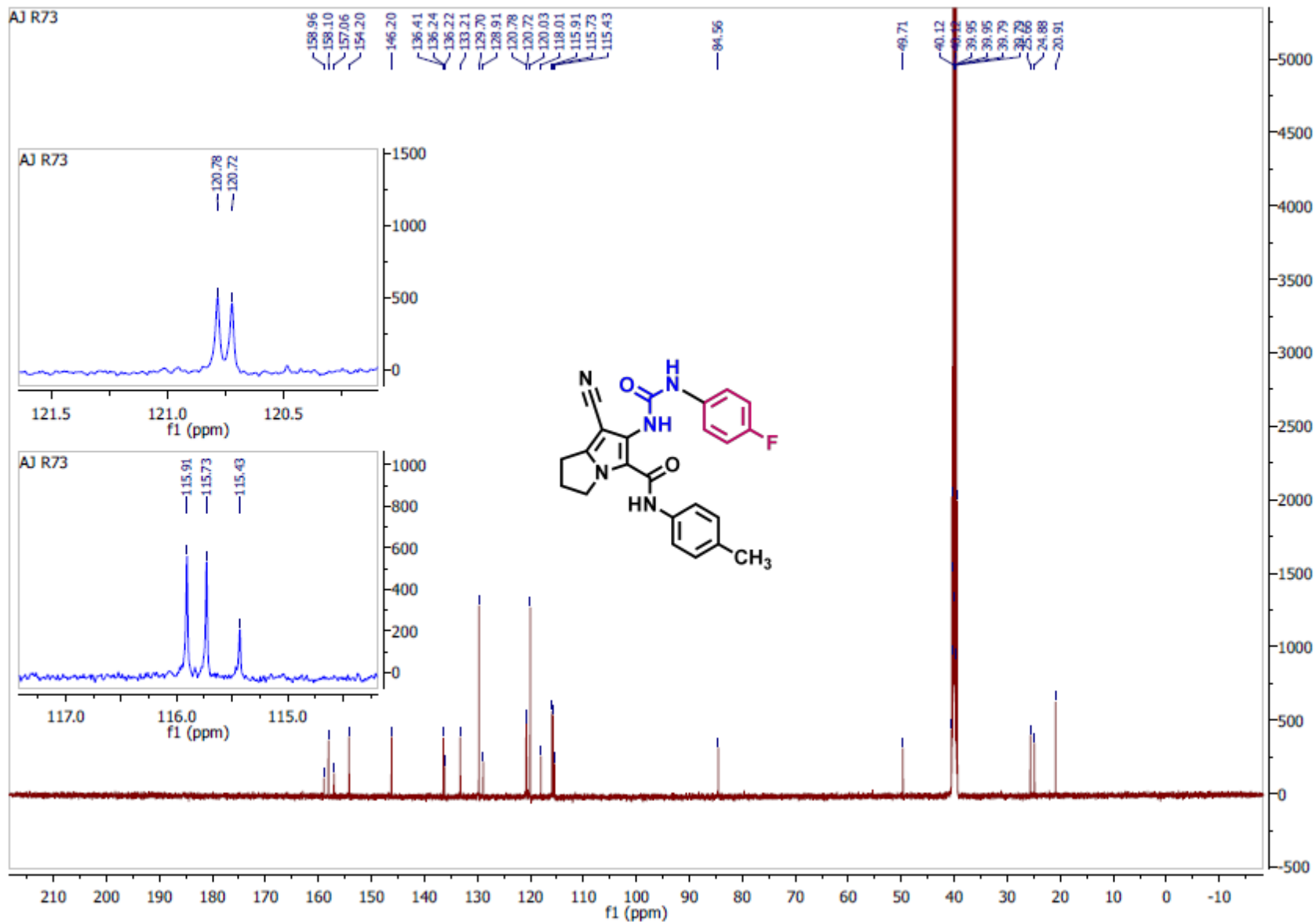


Figure S107. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18b** (zoom on aromatic Cs)

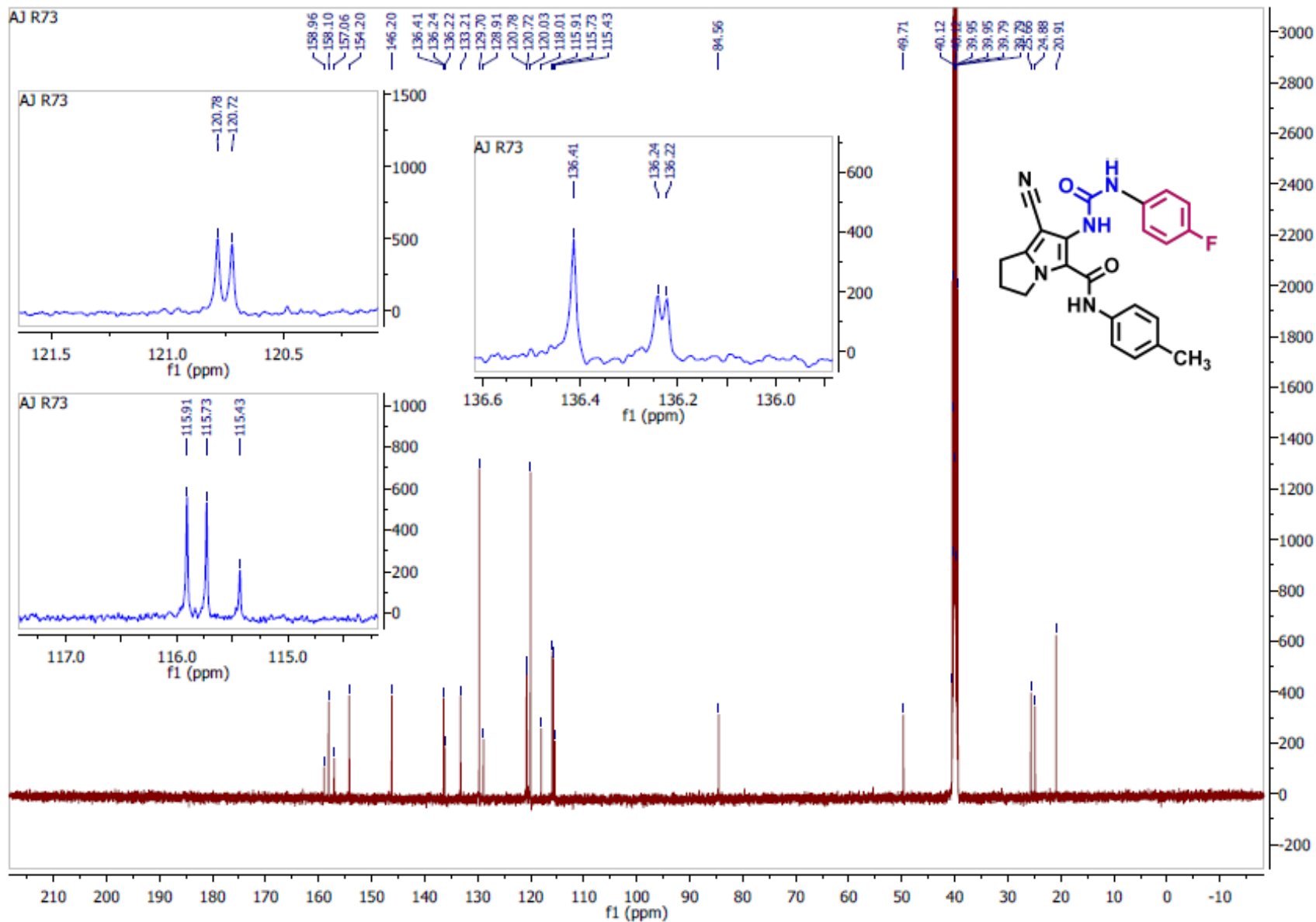


Figure S108. DEPT C^{135} (DMSO, 125 MHz, δ ppm) of compound **18b**.

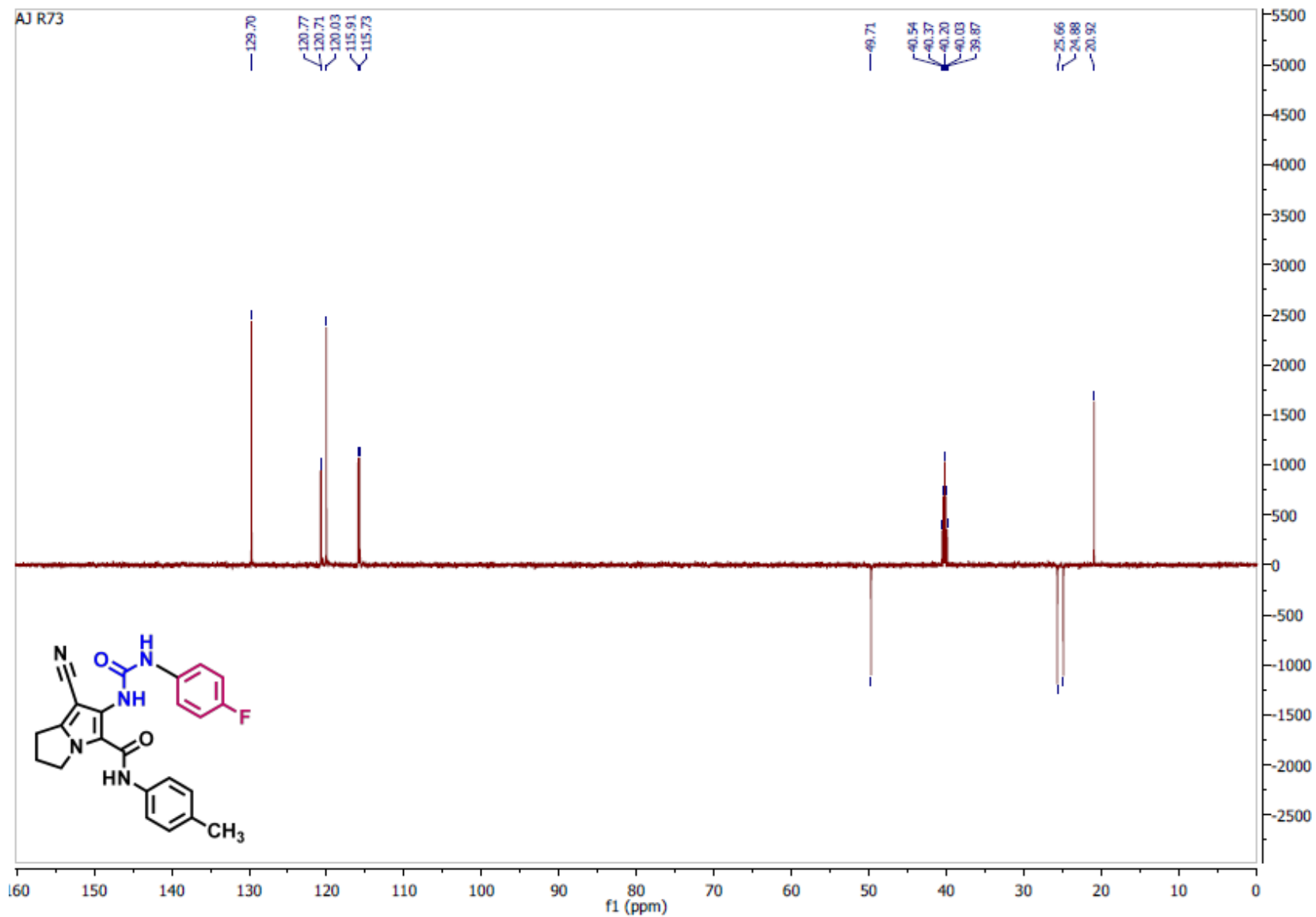


Figure S109. DEPT C¹³⁵ spectrum of compound **18b** (zoom on aromatic Cs)

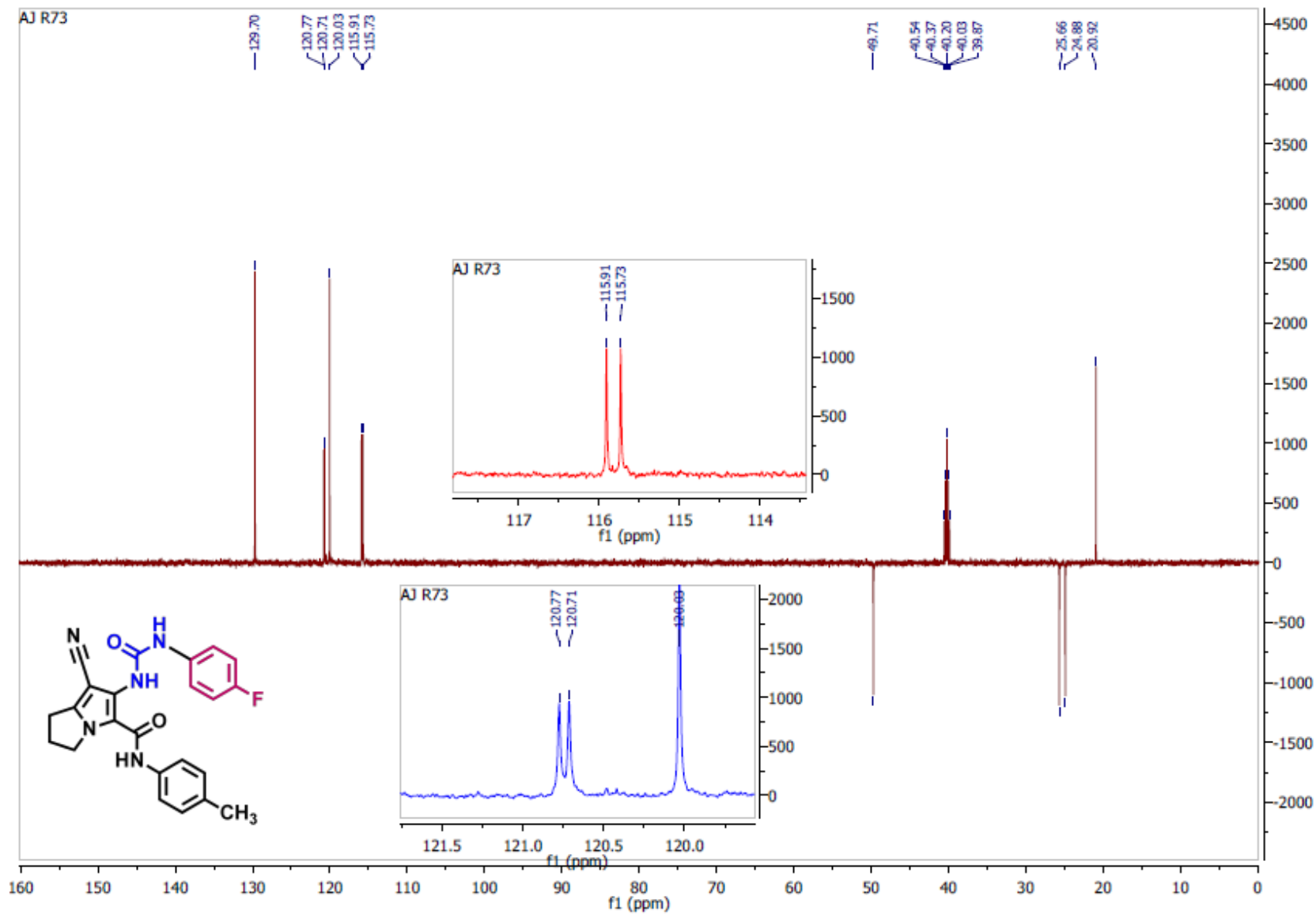


Figure S110. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **18c**.

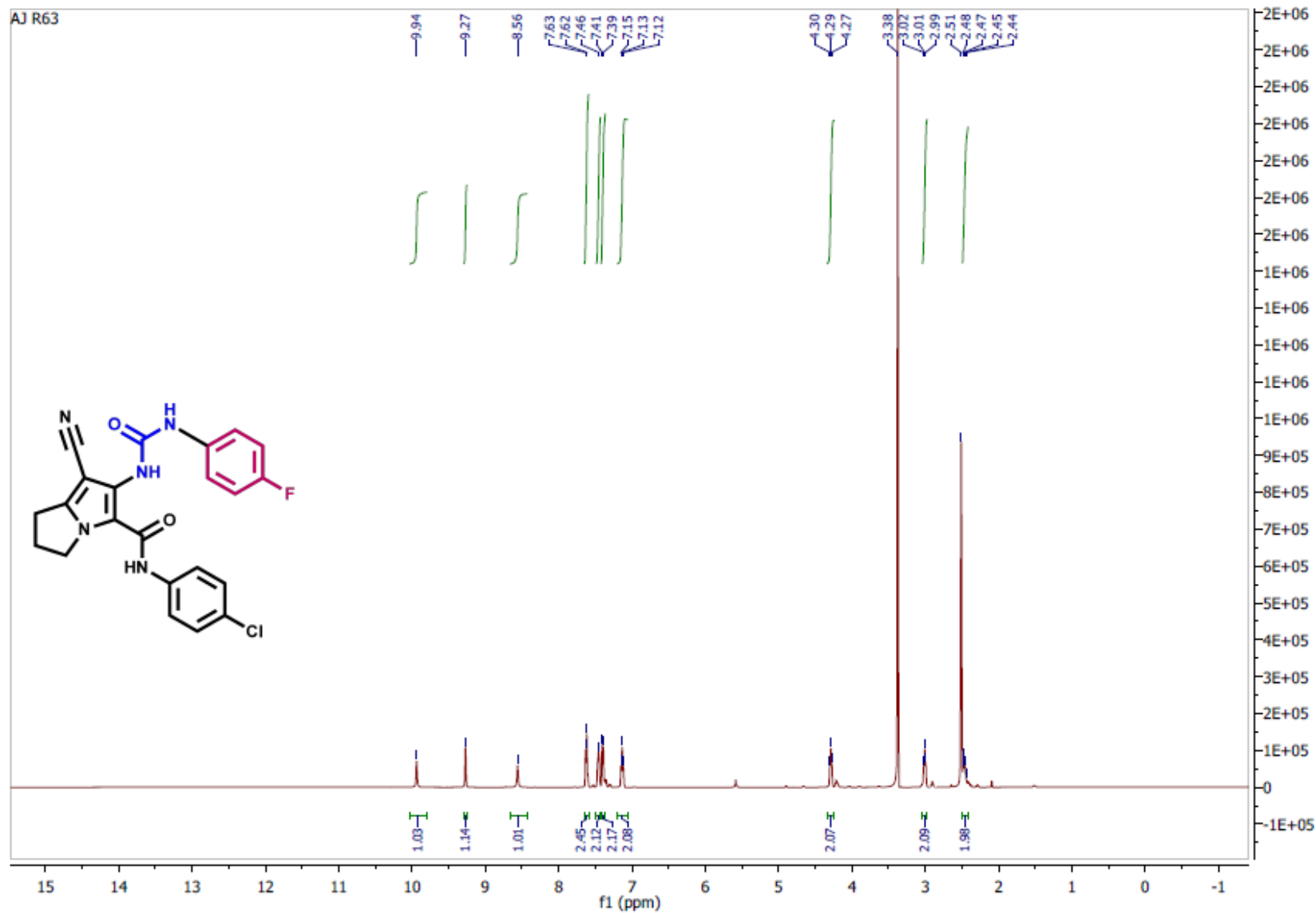


Figure S111. $^1\text{H-NMR}$ (DMSO, 500 MHz, δ ppm) spectrum of compound **18c** (zoom on aromatic Hs)

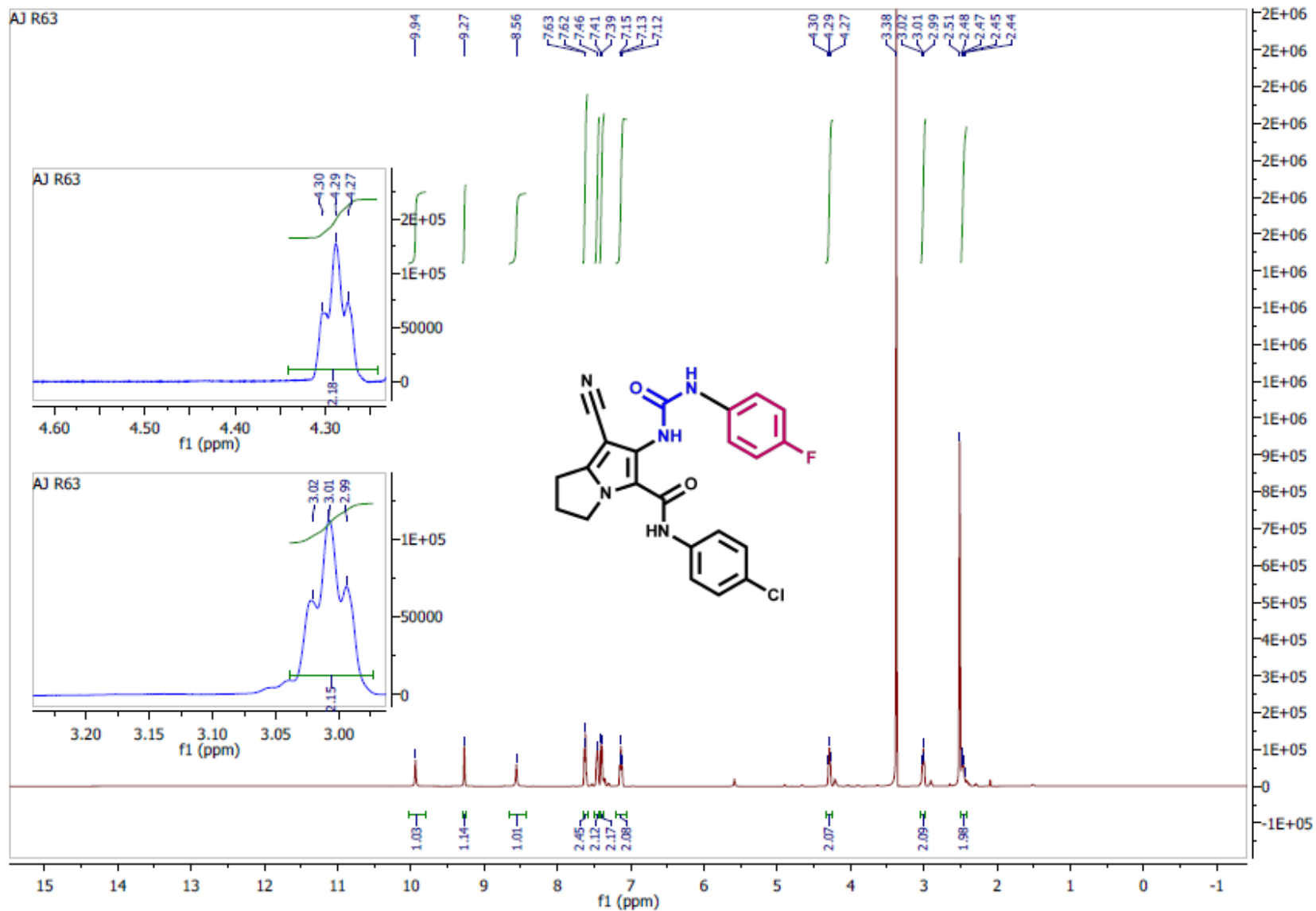


Figure S112. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound **18c** (zoom on aliphatic/aromatic Hs)

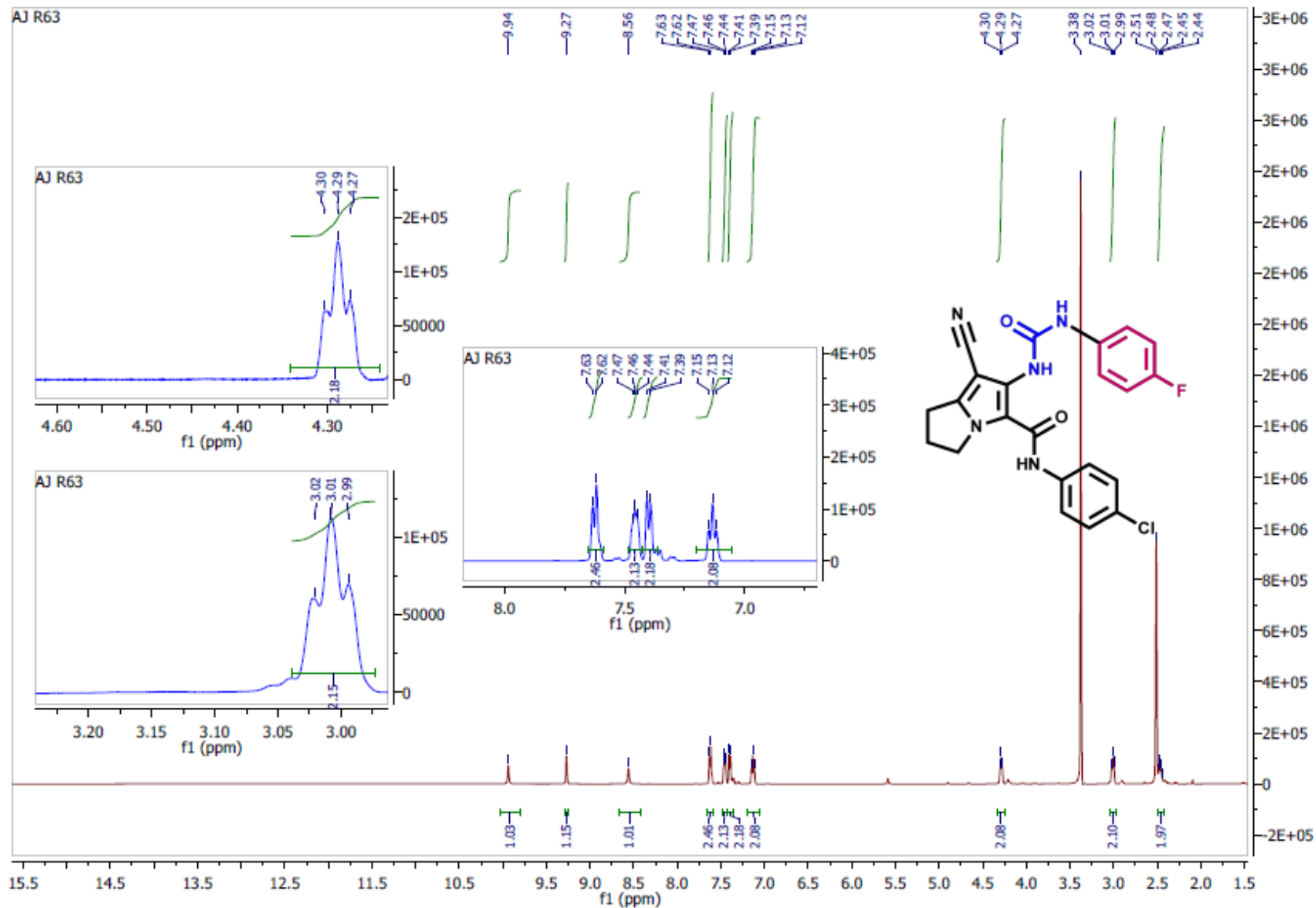


Figure S113. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18c**

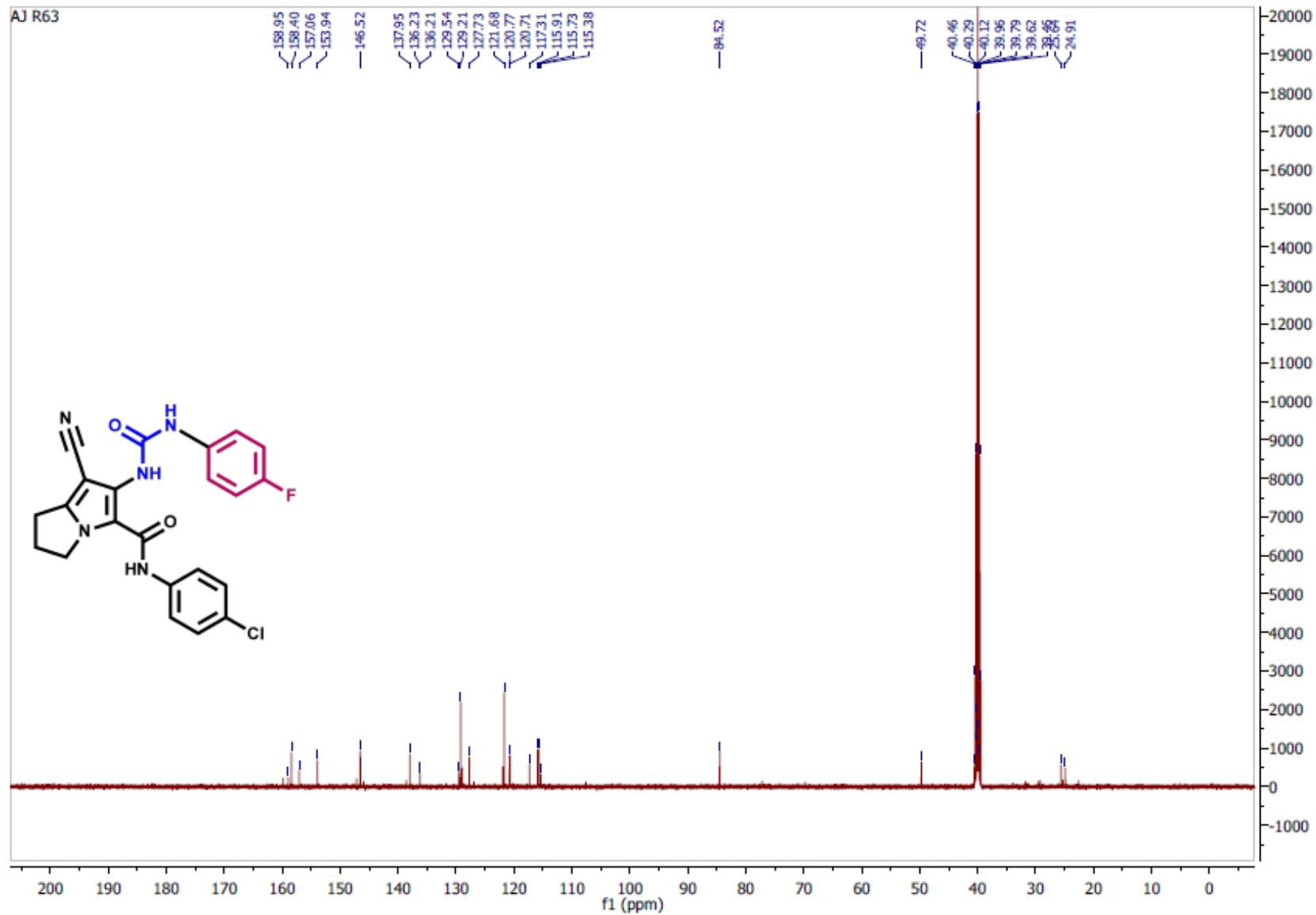


Figure S114. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18c** (zoom on aromatic C-F Cs)

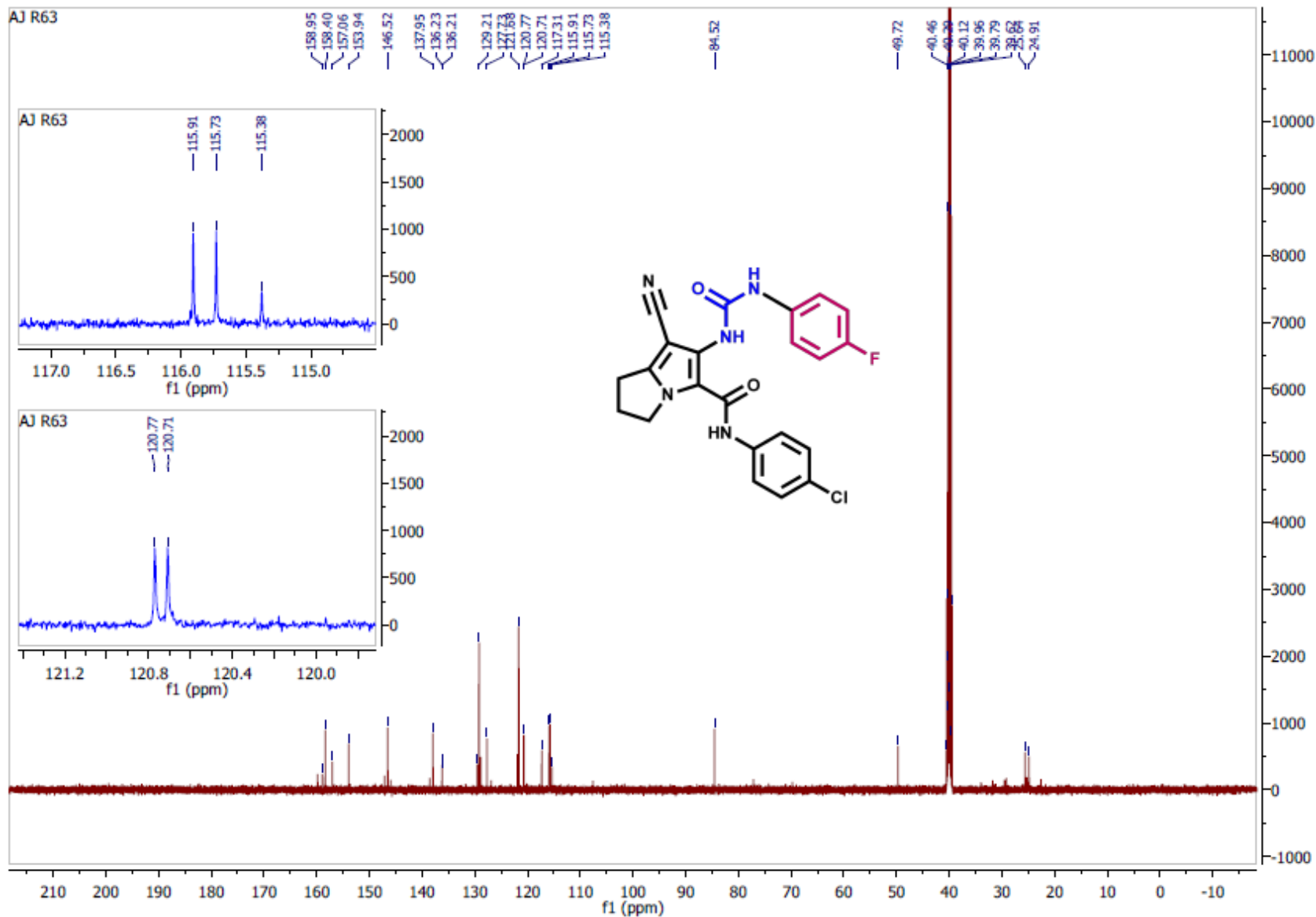


Figure S115. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **18c** (zoom on aromatic C-F Cs)

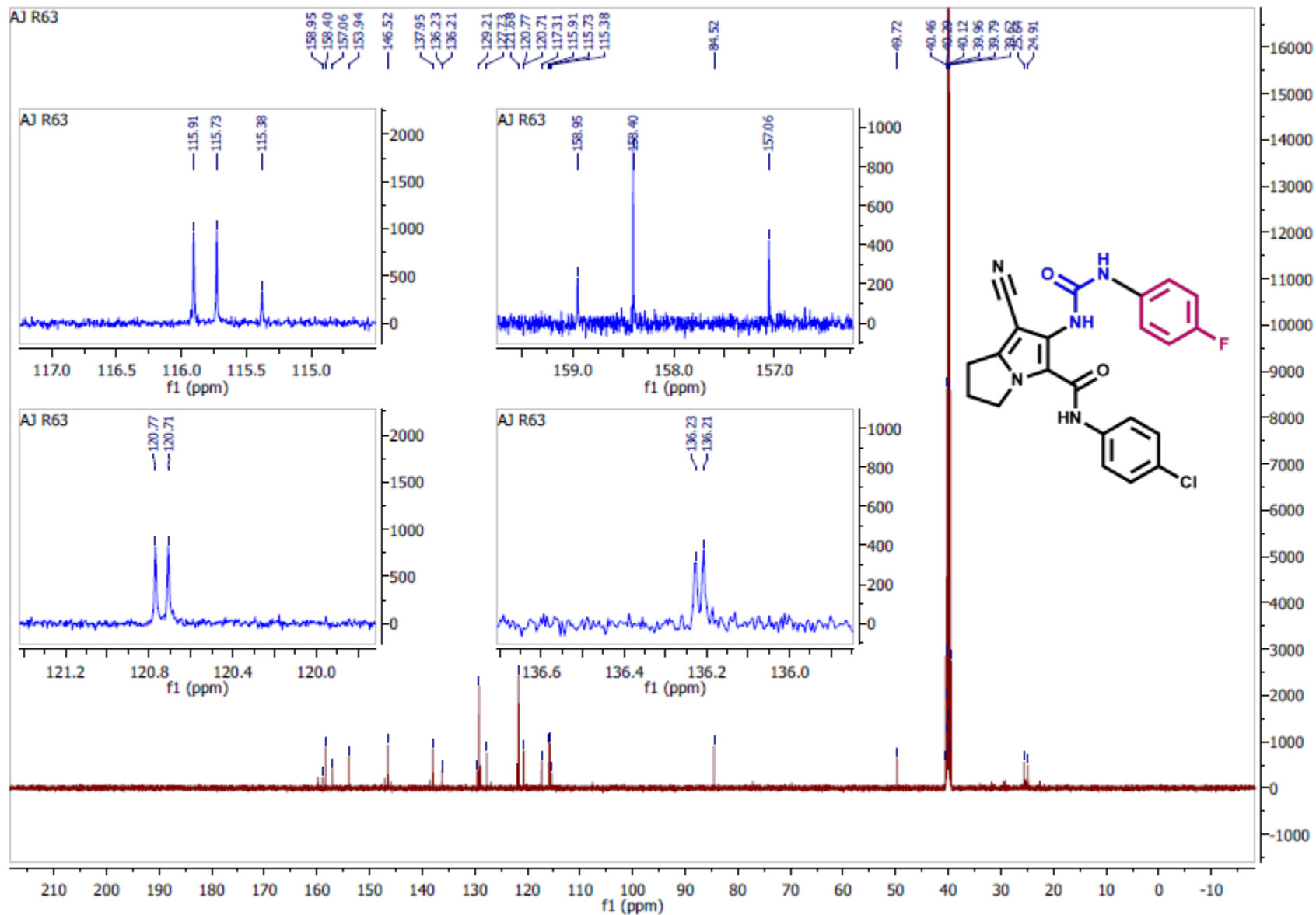


Figure S116. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound **18c**

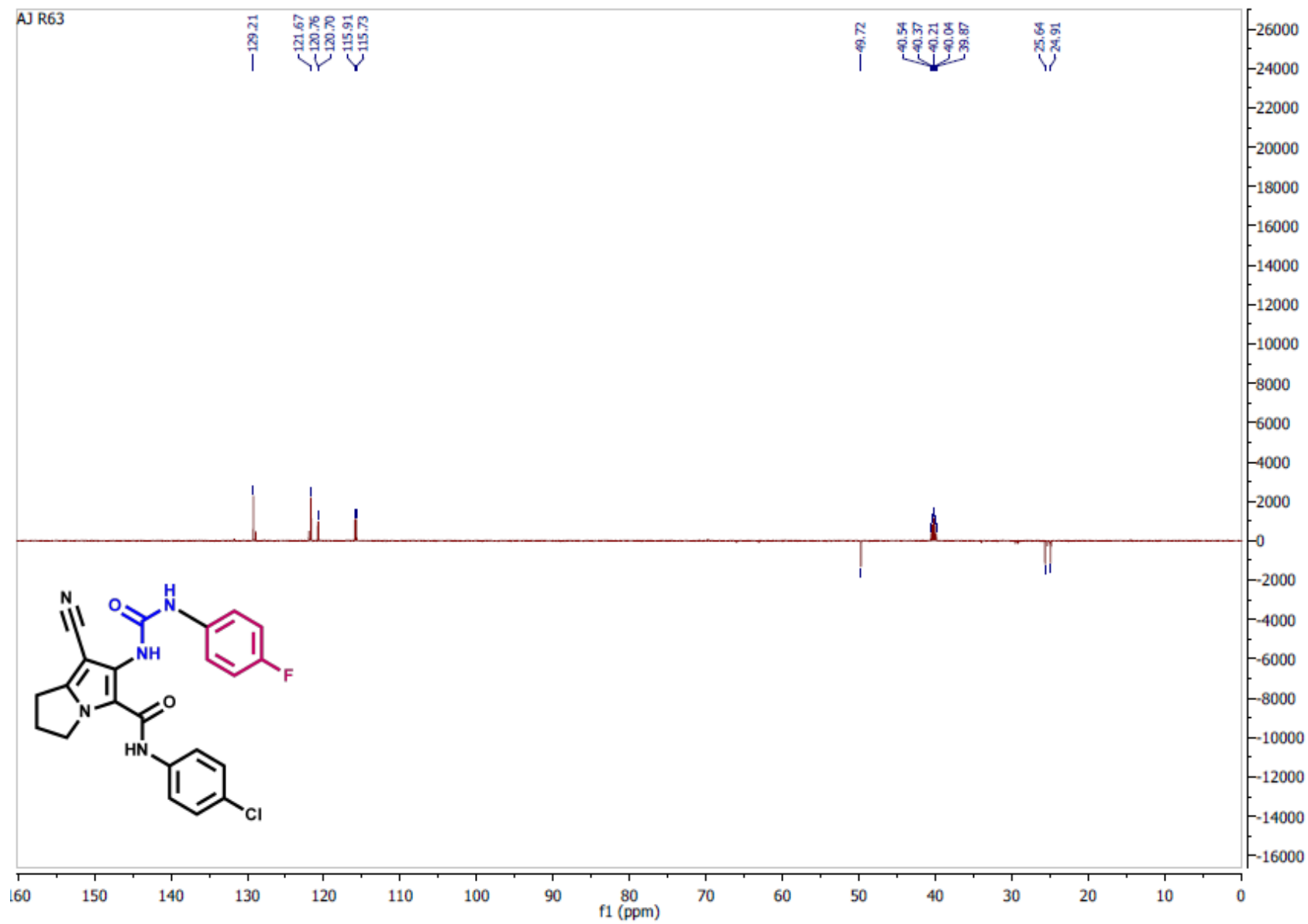


Figure S117. DEPT C^{135} (DMSO, 125 MHz, δ ppm) of compound **18c** (zoom on aromatic C-F Cs)

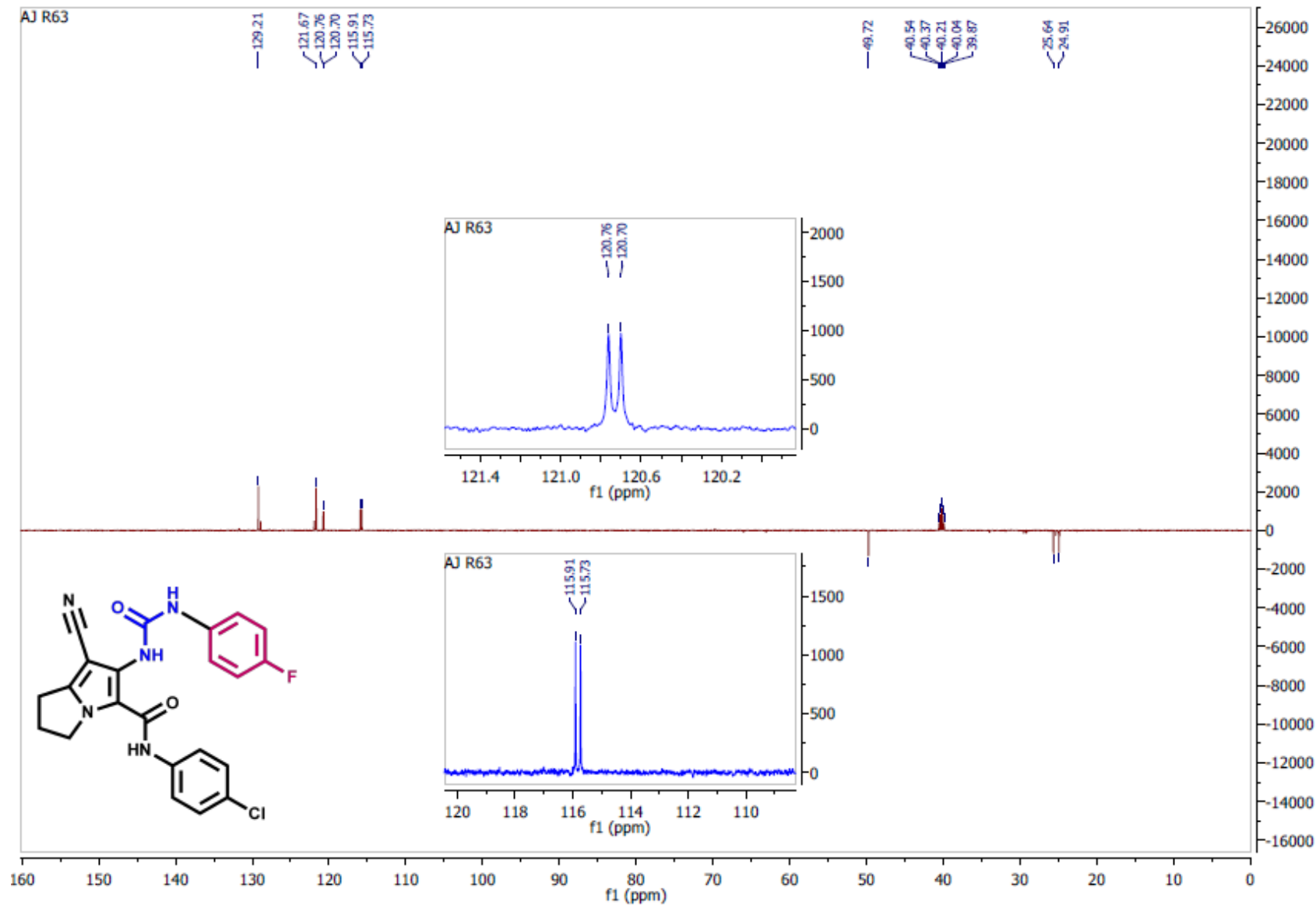


Figure S118. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound **19c**

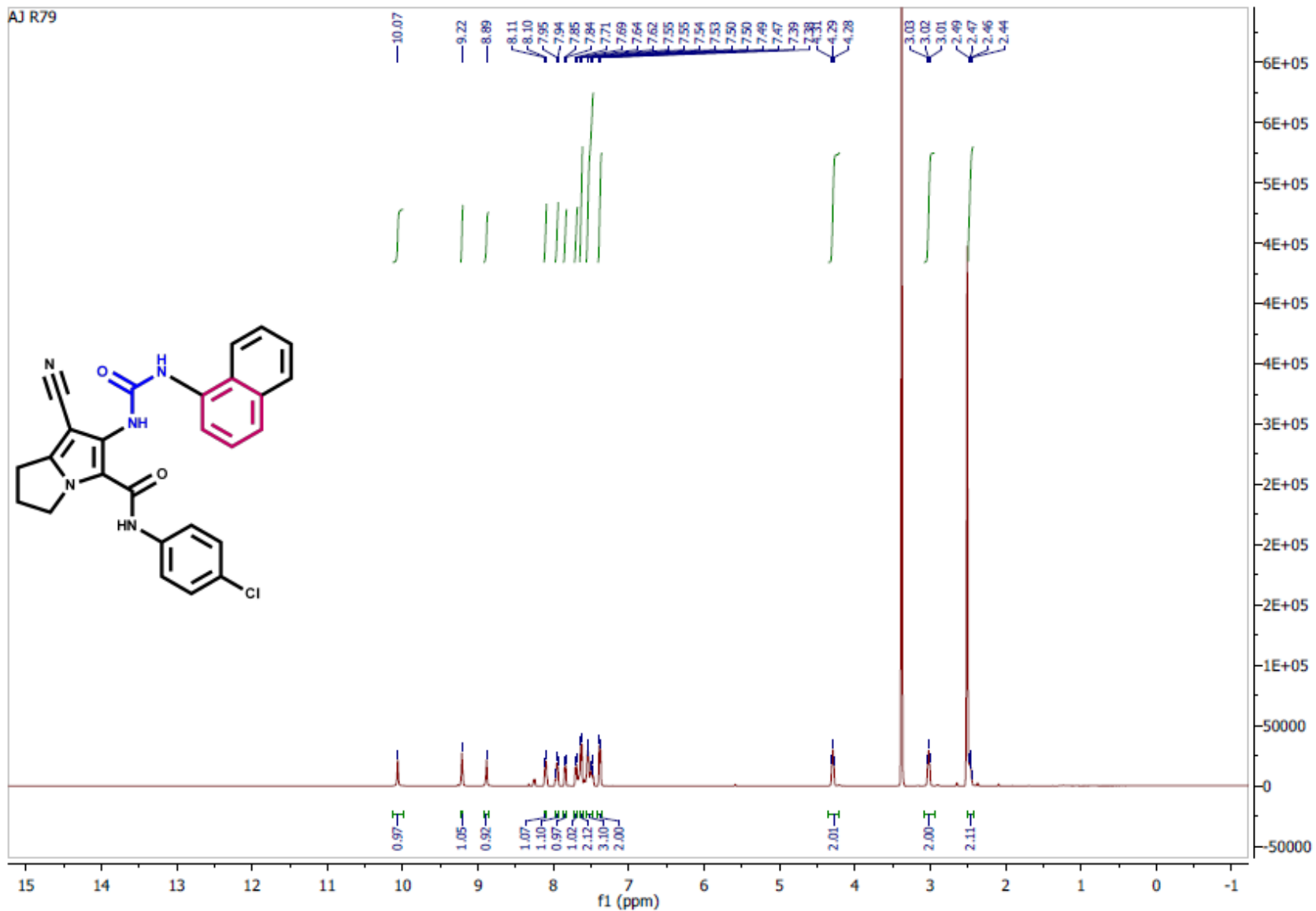


Figure S119. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound 19c (zoom on aliphatic Hs)

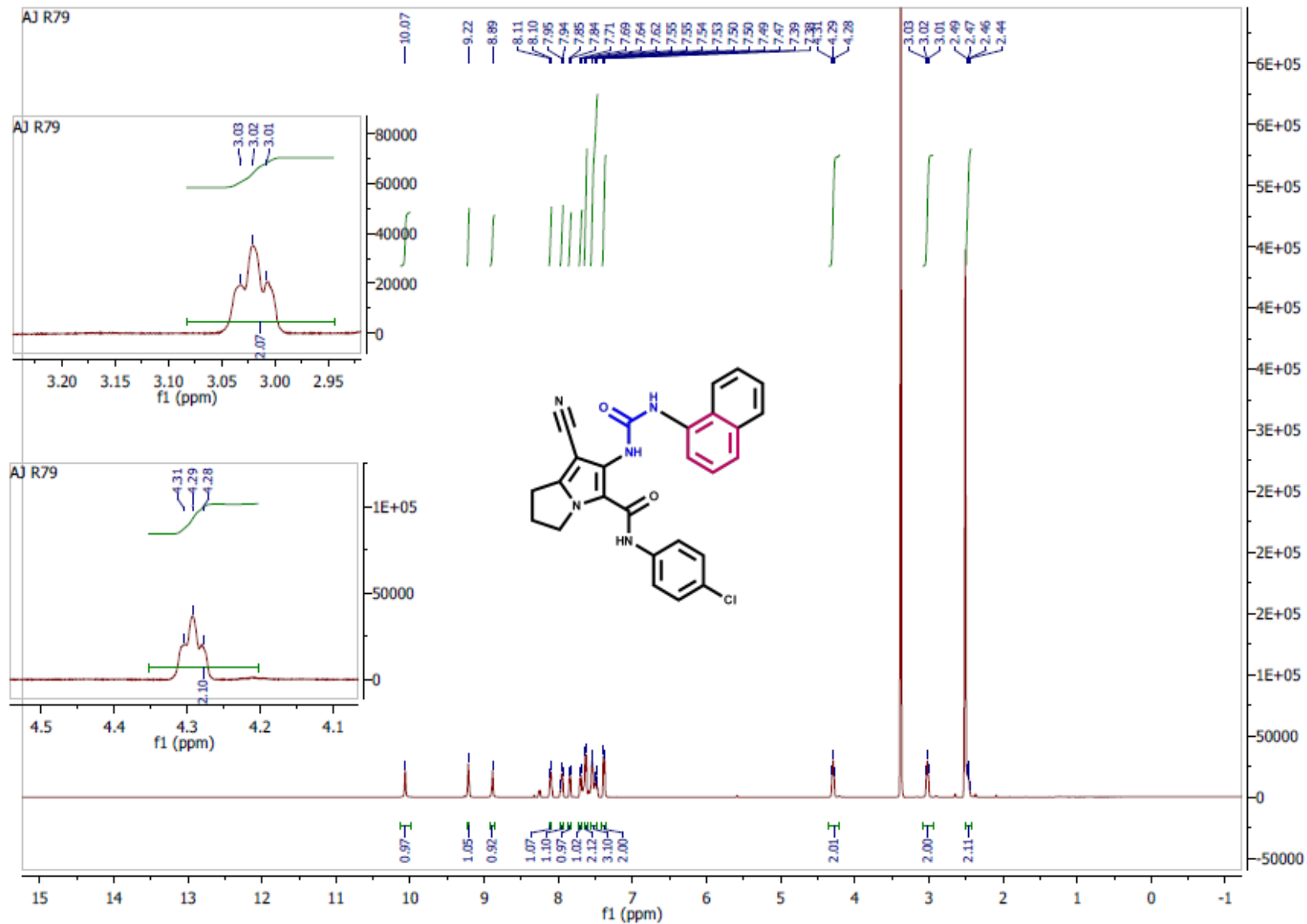


Figure S120. ¹H-NMR (DMSO, 500 MHz, δ ppm) spectrum of compound **19c** (zoom on aromatic Hs)

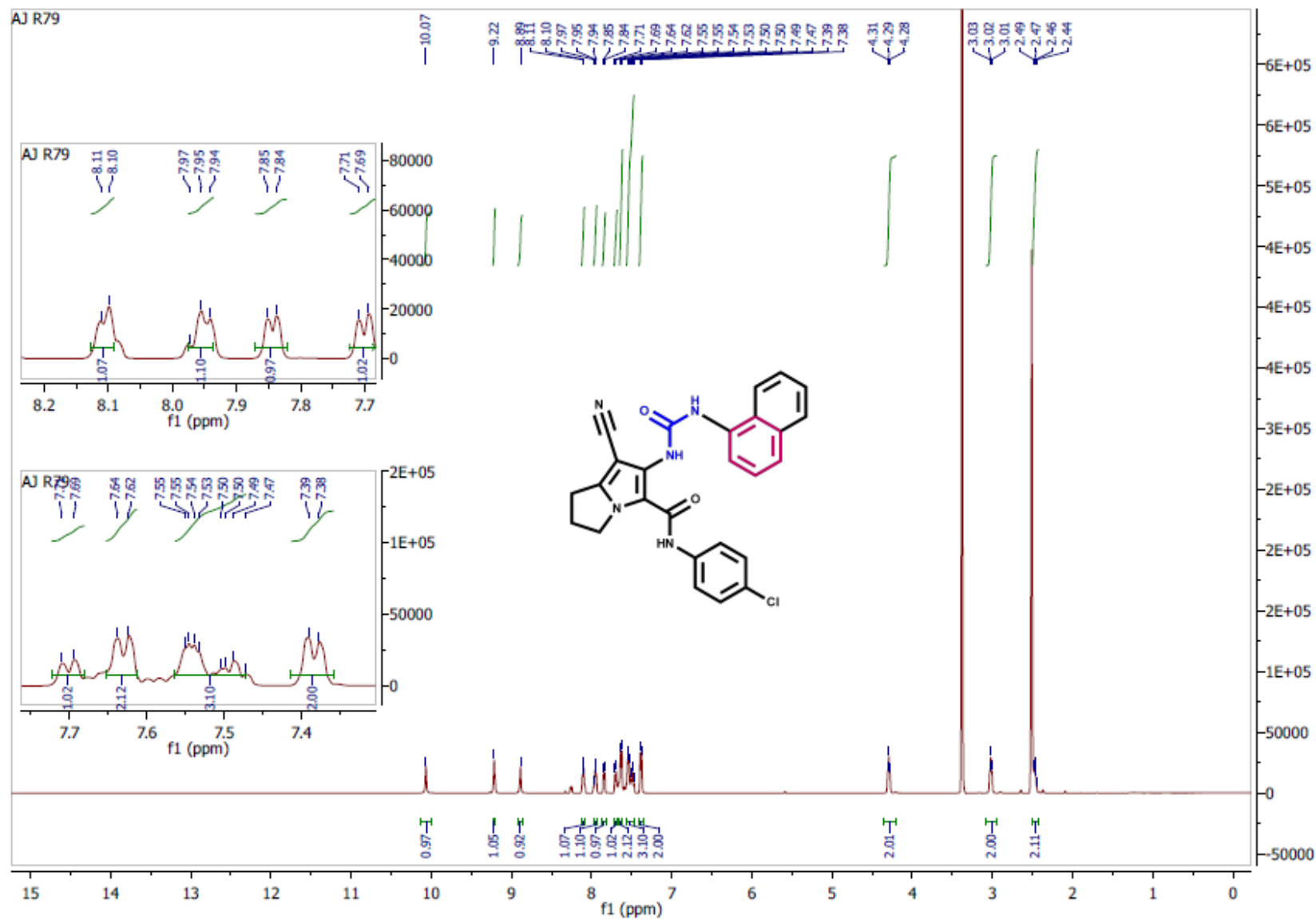


Figure S121. ^{13}C -NMR (DMSO, 125 MHz, δ ppm) spectrum of compound **19c**

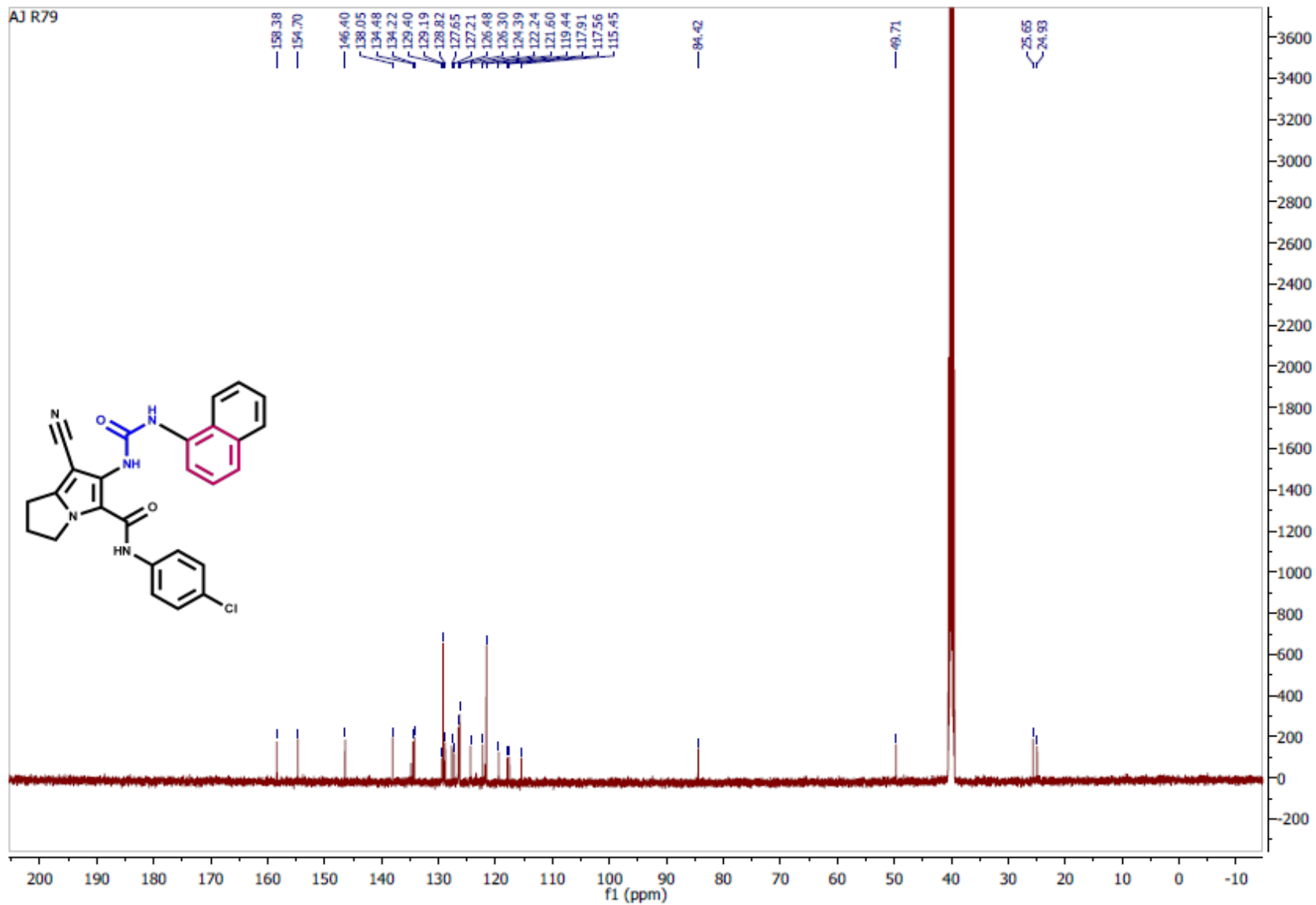
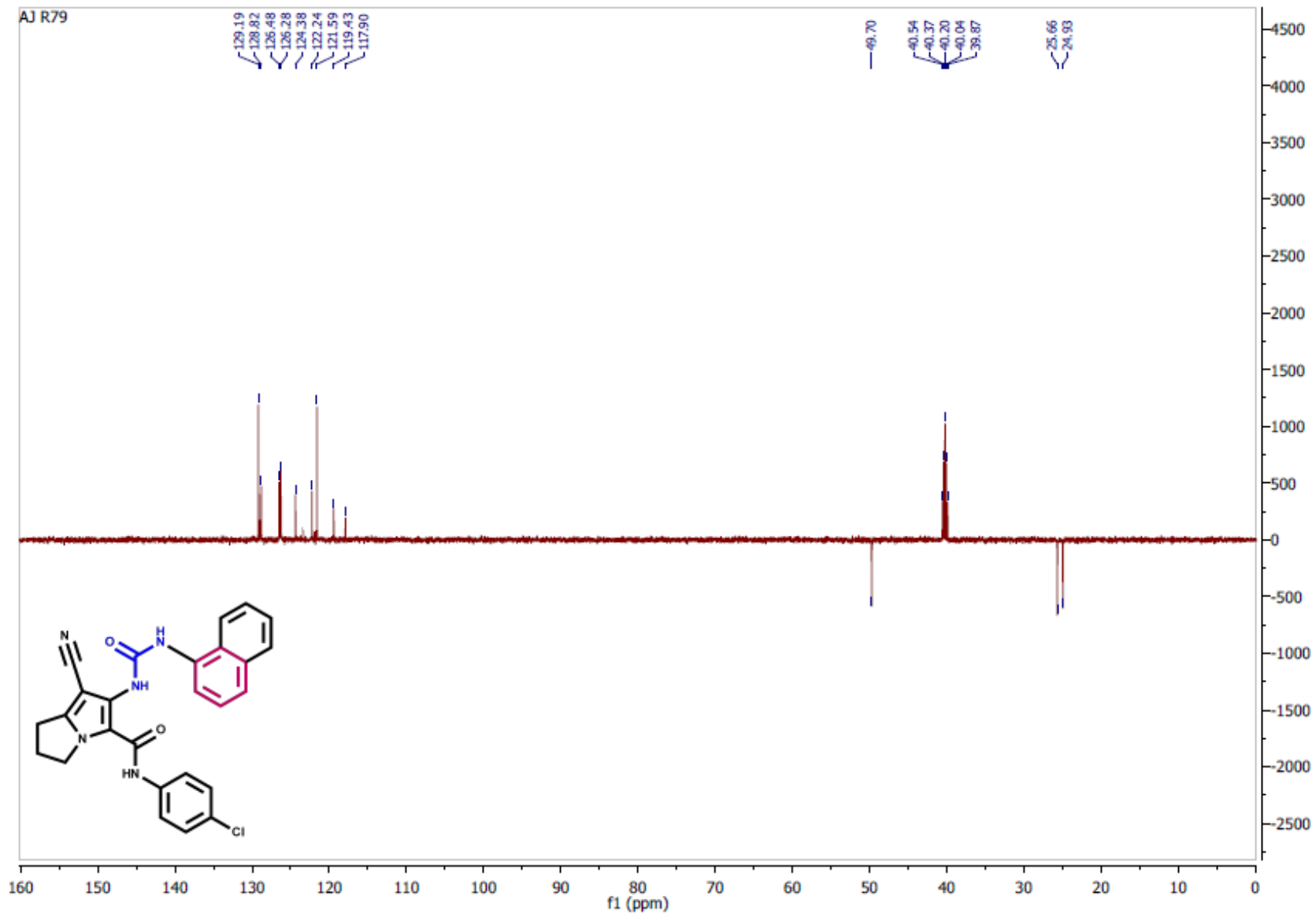


Figure S122. DEPT C¹³⁵ (DMSO, 125 MHz, δ ppm) of compound **19c**

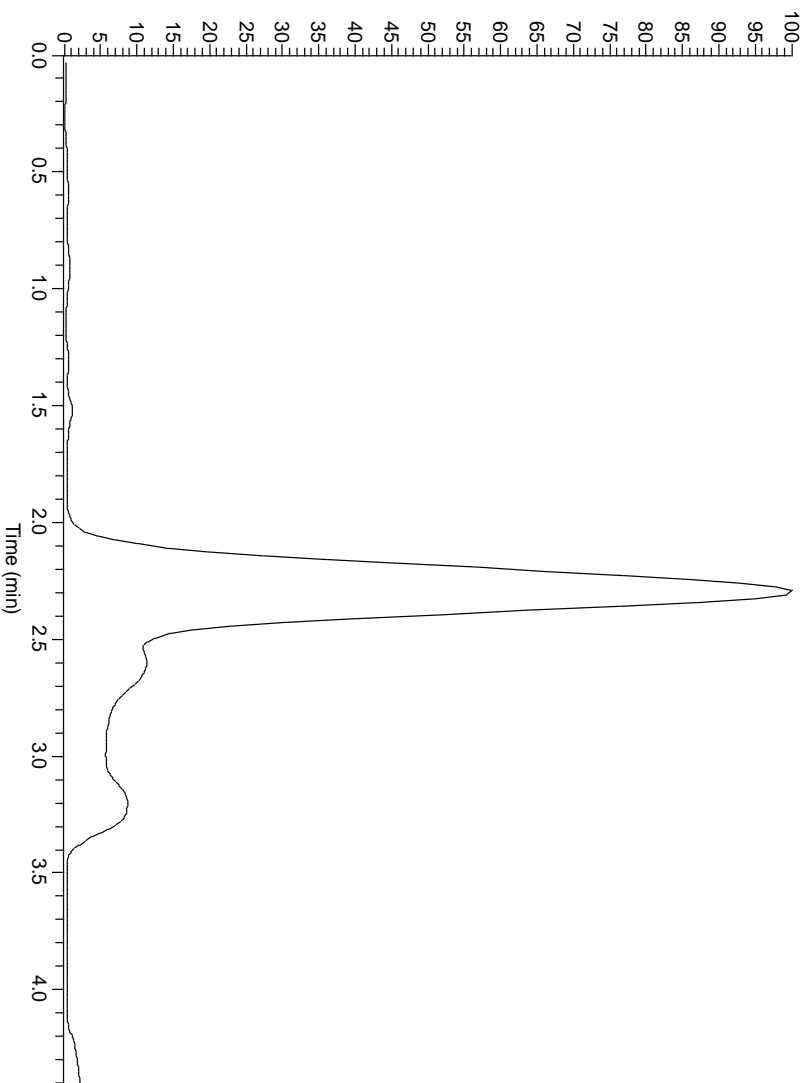


Mass Spectra

Mass spectra were recorded on Shimadzu GCMS QP5050A spectrometer, at 70 eV (EI) at the regional center for mycology and biotechnology, Al-Azhar University.

RT: 0.00 - 4.40 SM: 15G

NL:
1.33E6
TIC MS
ahmed-
nahmoud-
1s



ahmed-nahmoud-1s #135 RT: 2.28 AV: 1 SB: 2 4.40 , 4.40 NL: 3.90E5
T: (0,0) + c EI Full ms [40.00-1000.00]

Figure S123. Mass spectrum of compound 16a

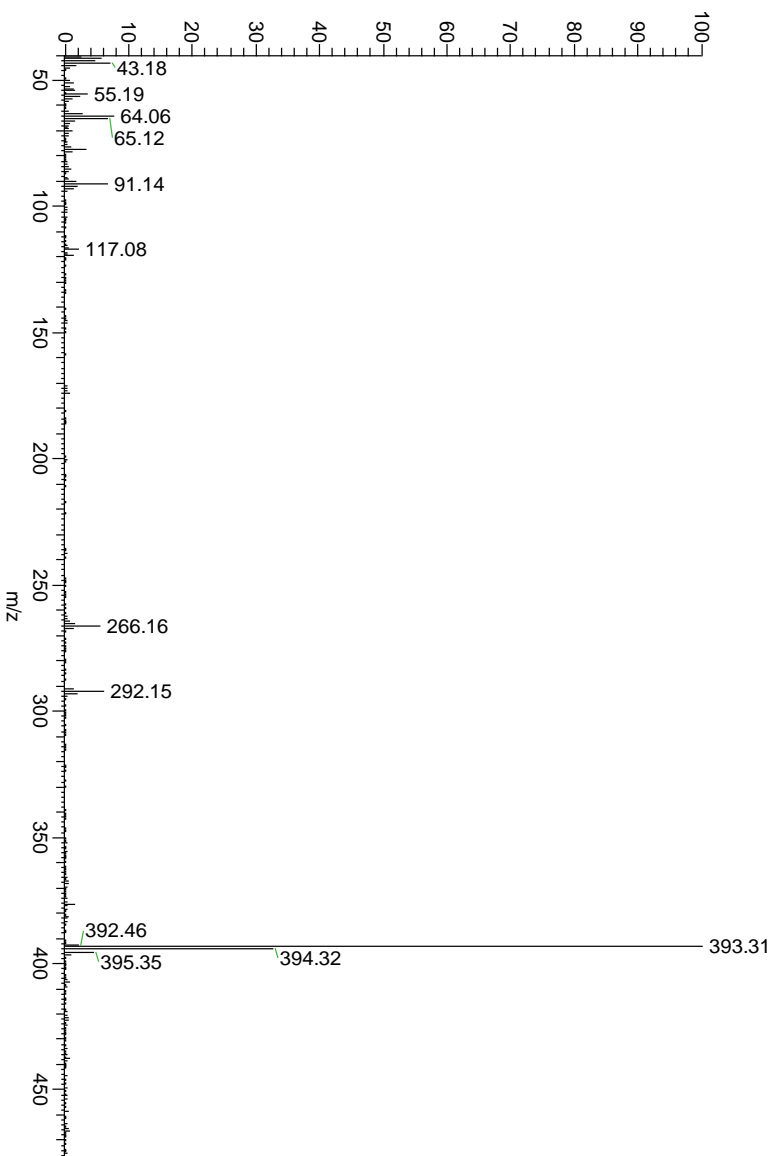
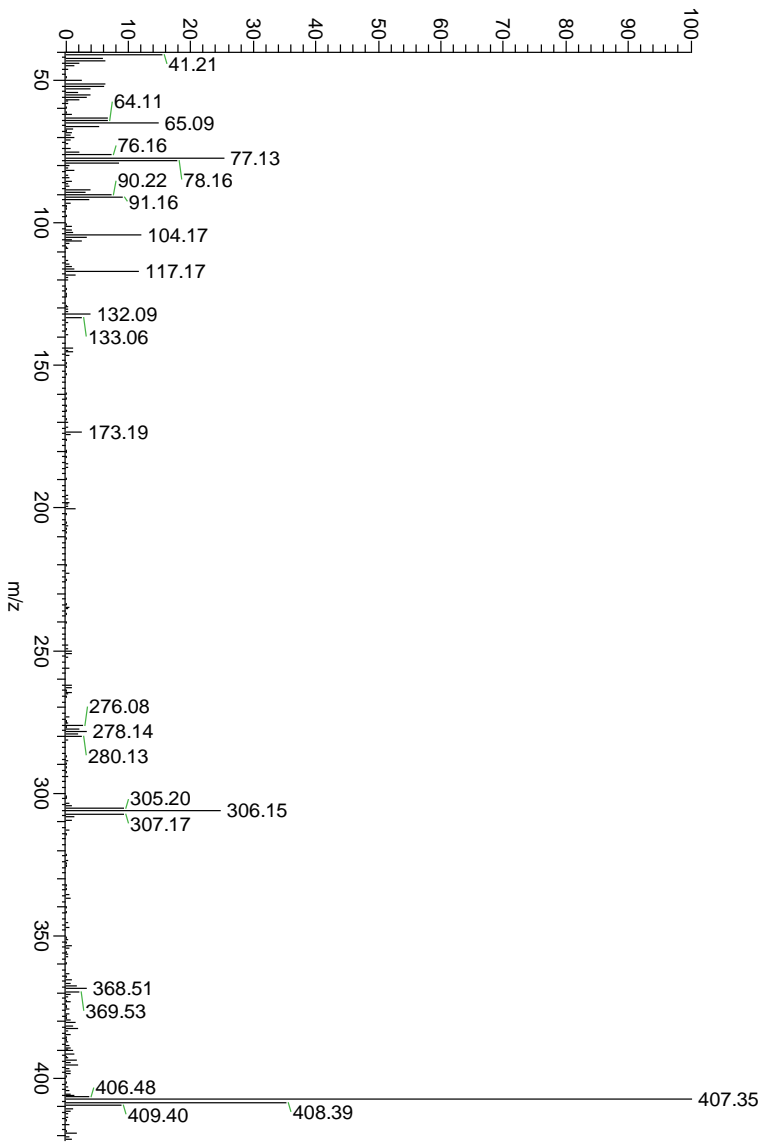
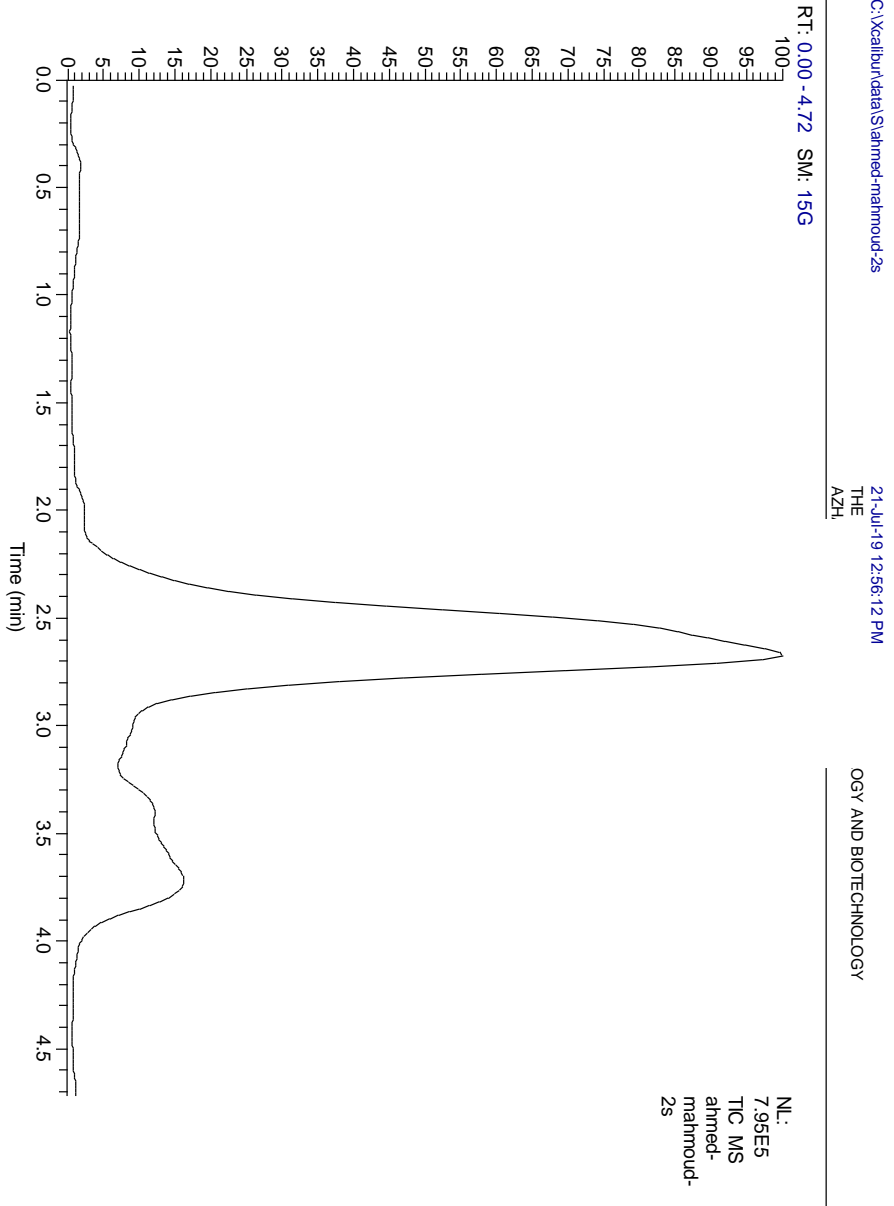


Figure S124. Mass spectrum of compound 16b

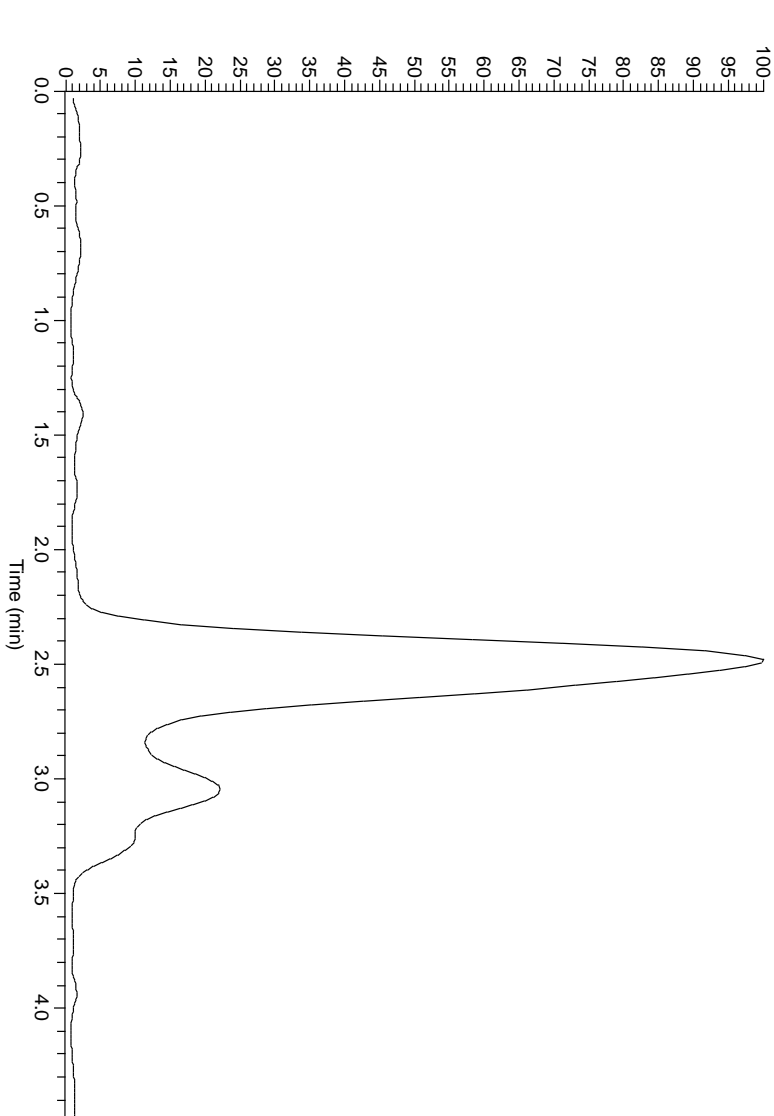


ahmed-mahmoud-2s #154 RT: 2.59 AV: 1 SB: 2 4.55, 4.59 NL: 1.07E5
T: (0.0) + c EI Full ms [40.00-1000.00]



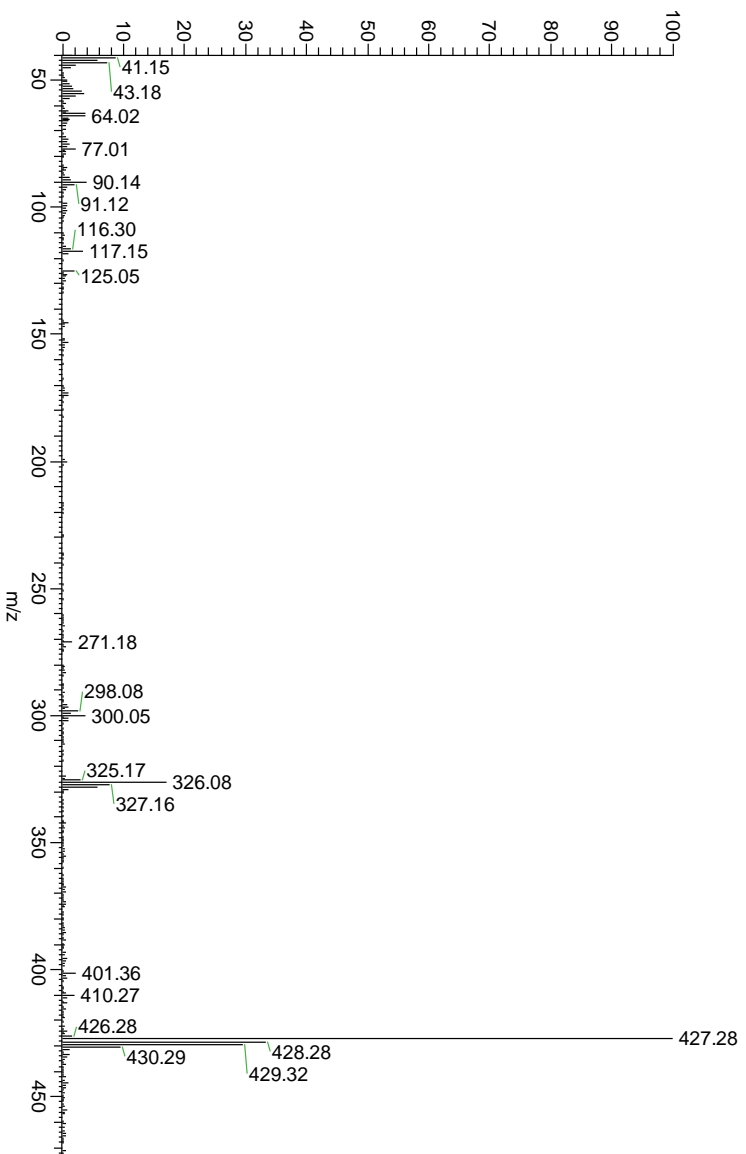
RT: 0.00 - 4.49 SM: 15G

NL:
9.98E5
TIC MS
ahmed-
mahmoud-
3s



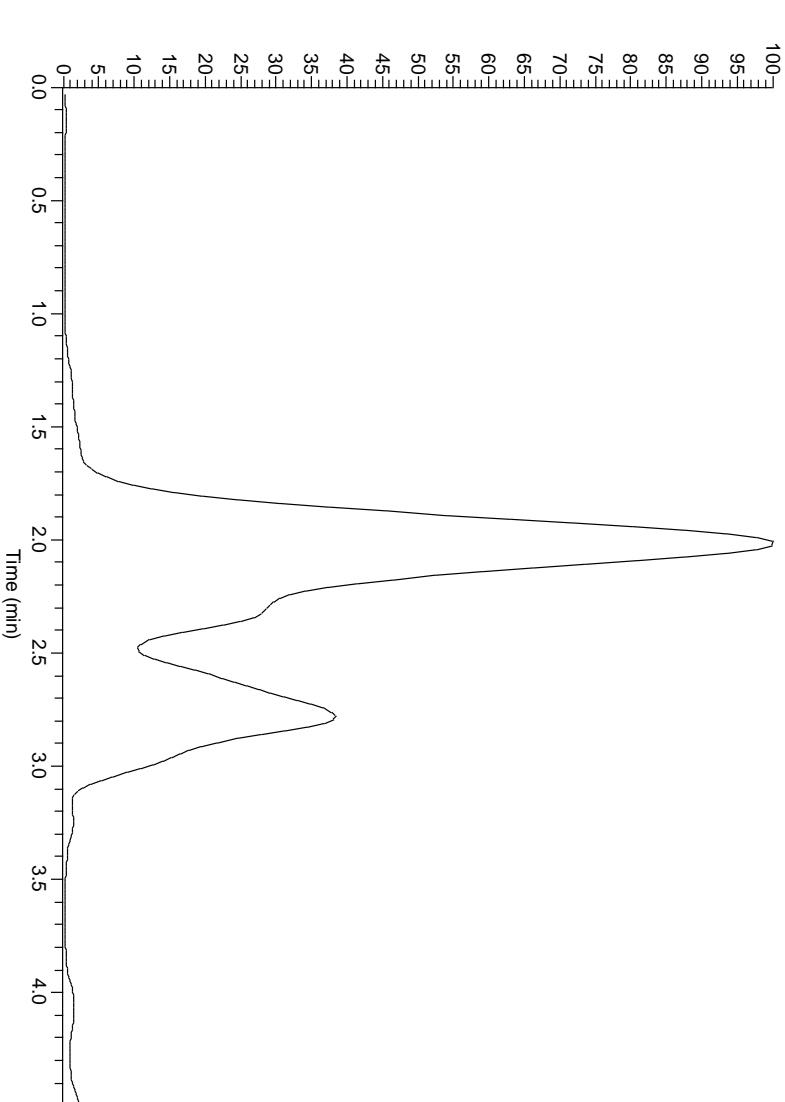
ahmed-mahmoud-3s #148 RT: 2.49 AV: 1 SB: 2.449, 4.49 NL: 2.39E5
T: (0.0) + c EI Full ms [40.00-1000.00]

Figure S125. Mass spectrum of compound 16c



RT: 0.00 - 4.49 SM: 15G

NL:
1.27E6
TIC MS
ahmed-
mahmoud-
4s



ahmed-mahmoud-4s #165 RT: 2.78 AV: 1 SB: 2 4.49 , 4.49 NL: 5.53E4
T: (0.0) + c EI Full ms [40.00-1000.00]

Figure S126. Mass spectrum of compound 17a

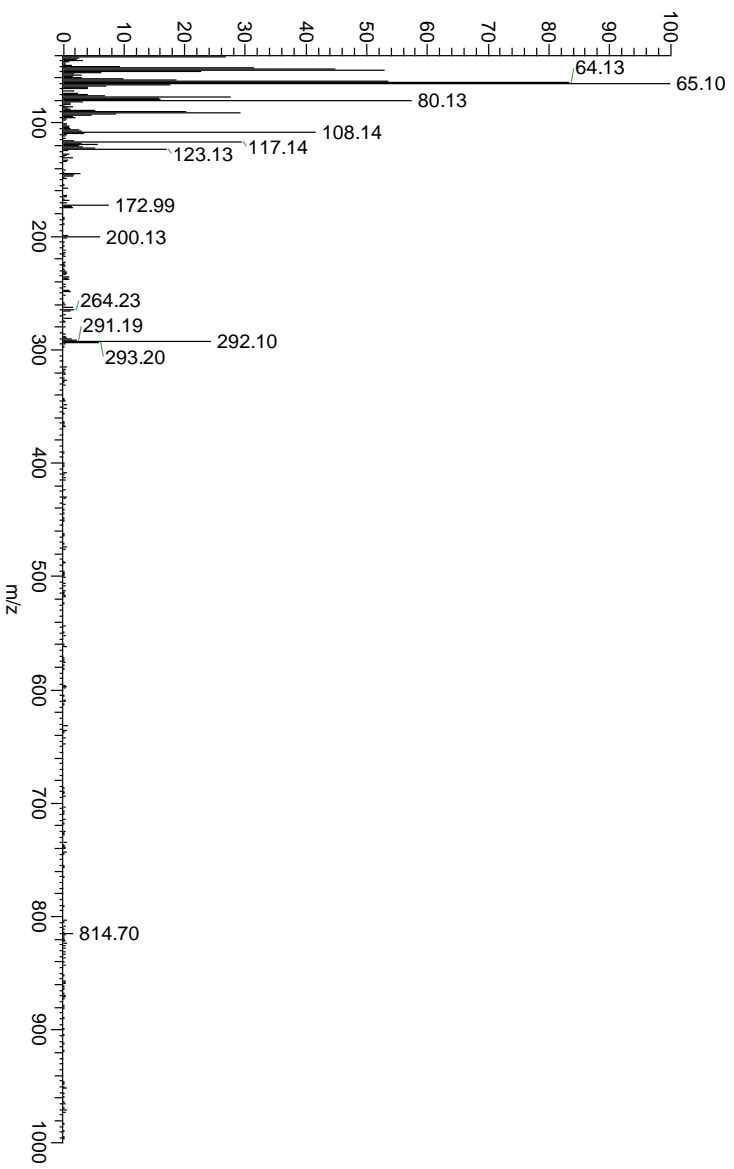
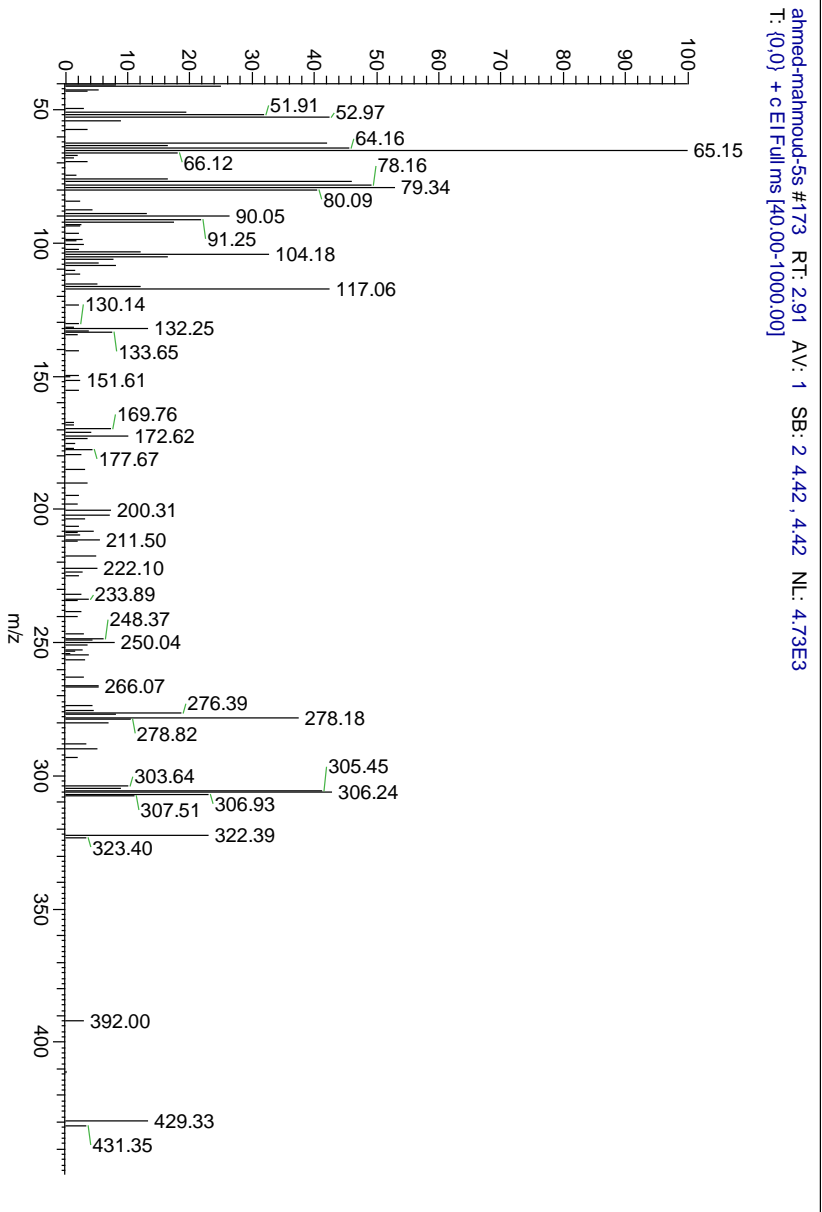
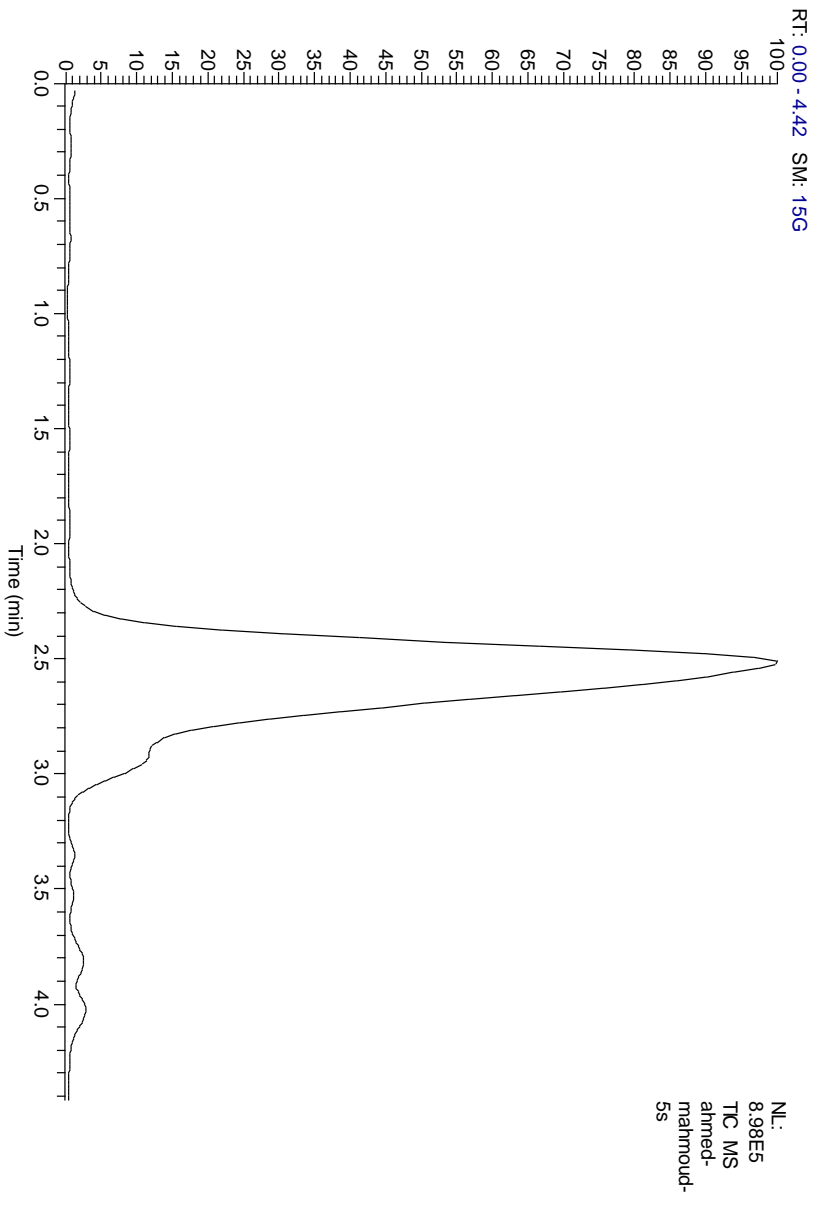


Figure S127. Mass spectrum of compound 17b



ahmed-mahmoud-5s #173 RT: 2.91 AV: 1 SB: 2 4.42, 4.42 NL: 4.73E3
T: (0.0) + c EI Full ms [40.00-1000.00]



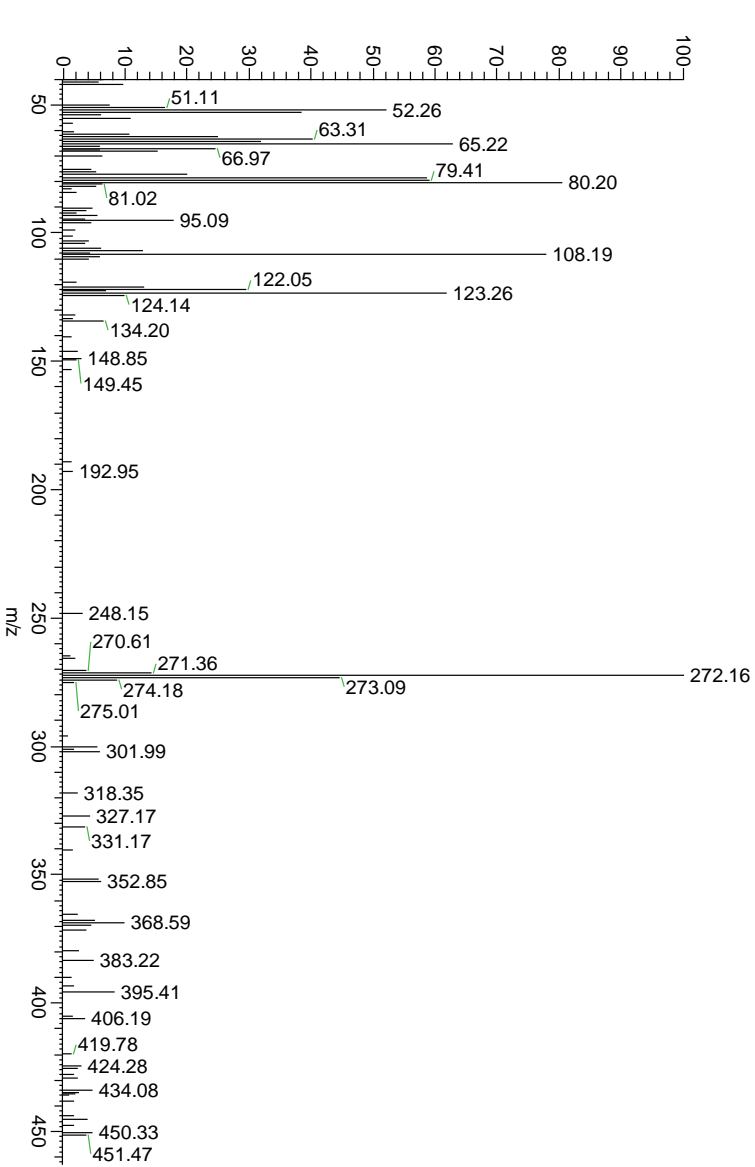
C:\Xcalibur\data\ahmed-mahmoud-5s

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NL:
8.98E5
TIC MS
ahmed-
mahmoud-
5s

Figure S128. Mass spectrum of compound 17c



ahmed-mahmoud-6s #127 RT: 2.14 AV: 1 SB: 2 4.47, 4.47 NL: 9.20E3
T: (0.0) + c EI Full ms [40.00-1000.00]

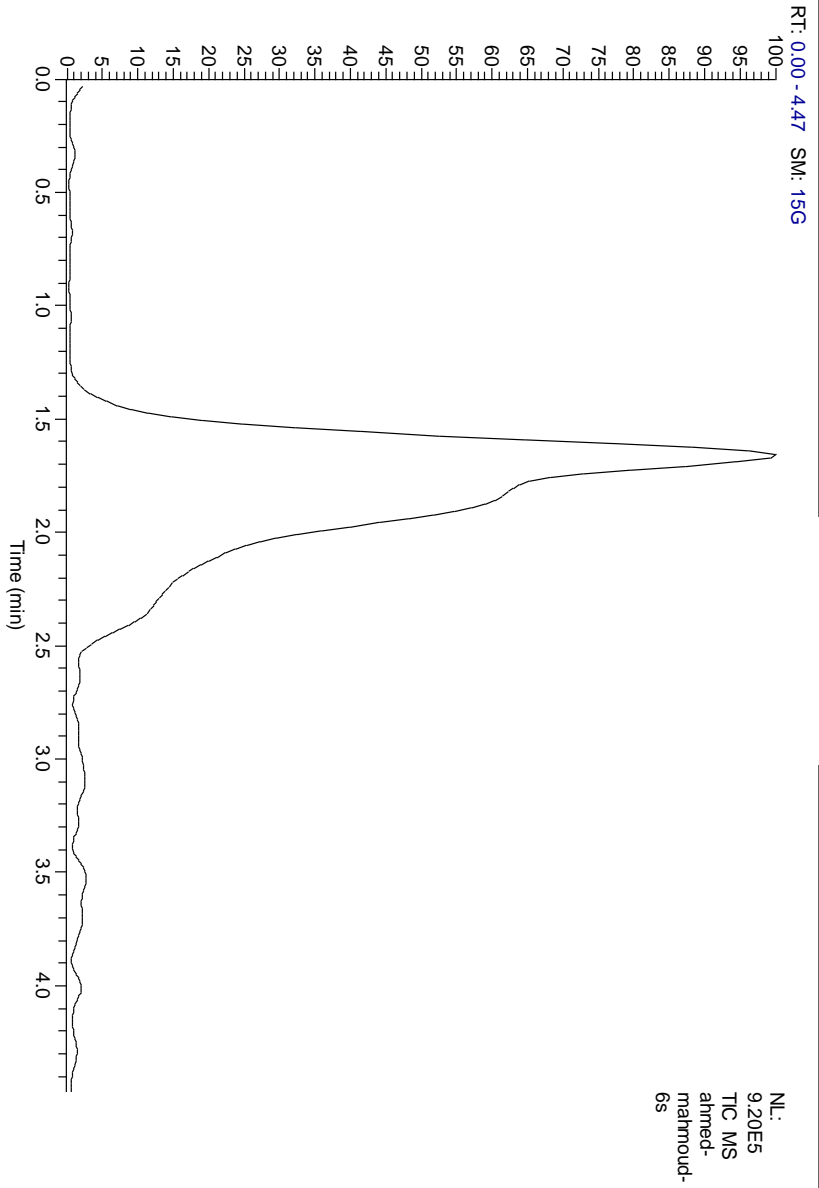
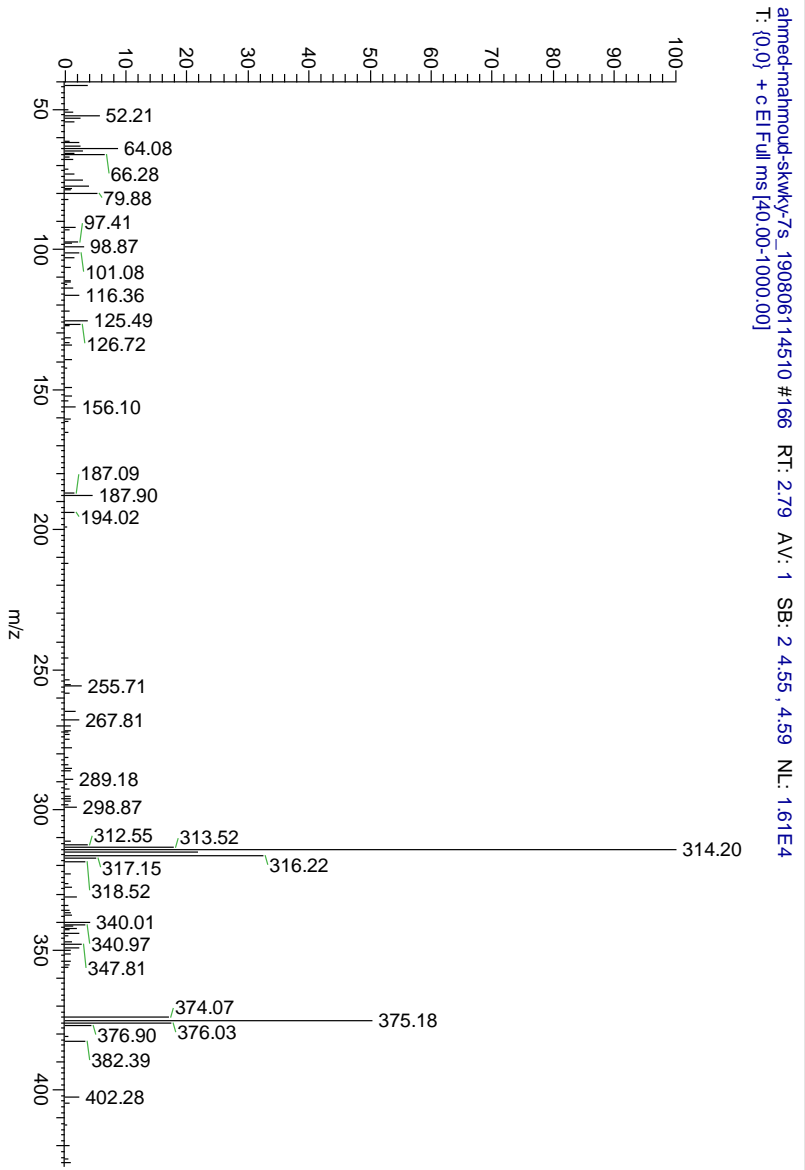
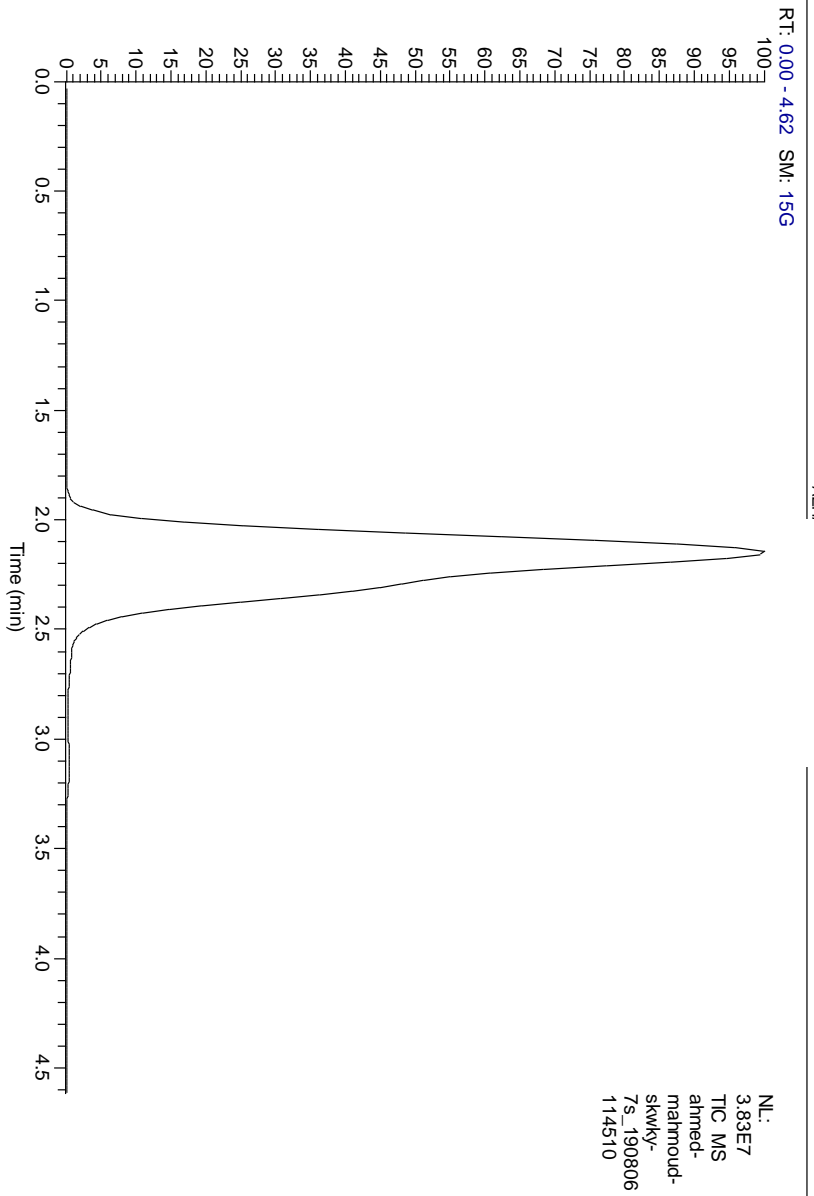


Figure S129. Mass spectrum of compound 18a



ahmed-mahmoud-skwy-7s_190806114510 #166 RT: 2.79 AV: 1 SB: 2 4.55, 4.59 NL: 1.61E4
T: {0.0} + c EI Full ms [40.00-1000.00]



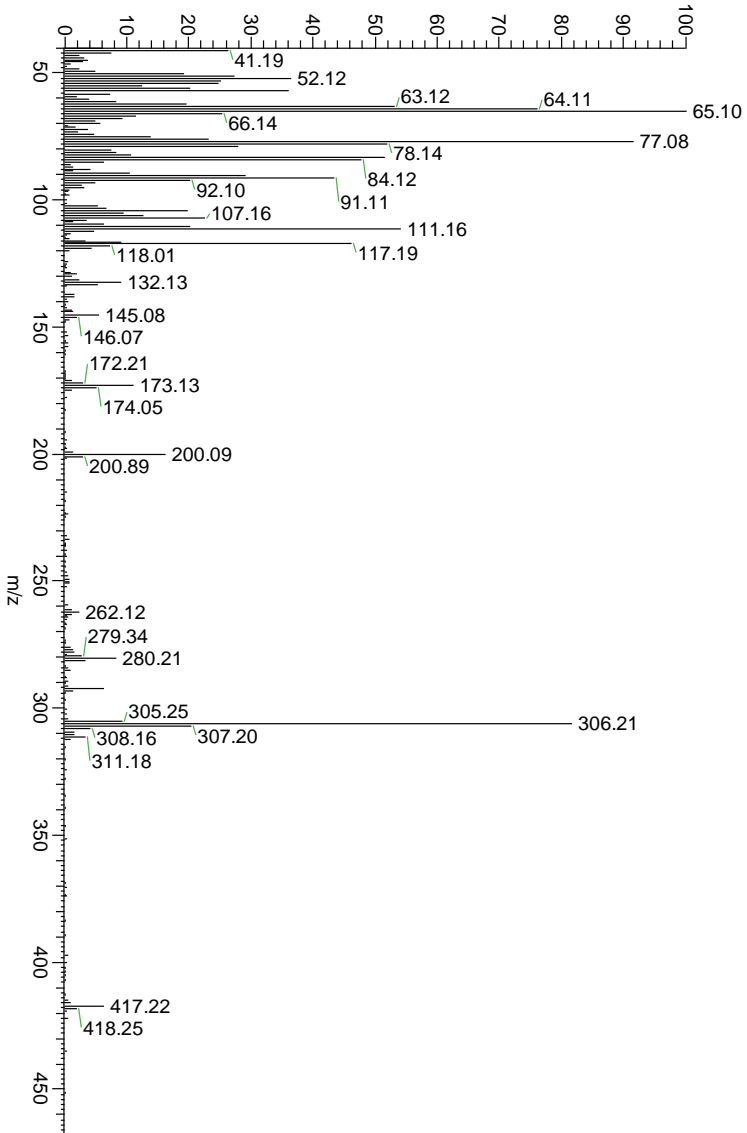
RT: 0.00 - 4.62 SM: 15G

ahmed-mahmoud-skwy-7s_190806114510
06-Aug-19 11:45:10 AM
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NL:
3.83E7
TIC MS
ahmed-
mahmoud-
skwy-
7s_190806
114510

Figure S130. Mass spectrum of compound 18b



ahmed-mahmoud-skwky-8s #183 RT: 3.08 AV: 1 SB: 2 4.49 4.49 NL: 1.16E5
T: (0.0) + c EI Full ms (40.00-1000.00)

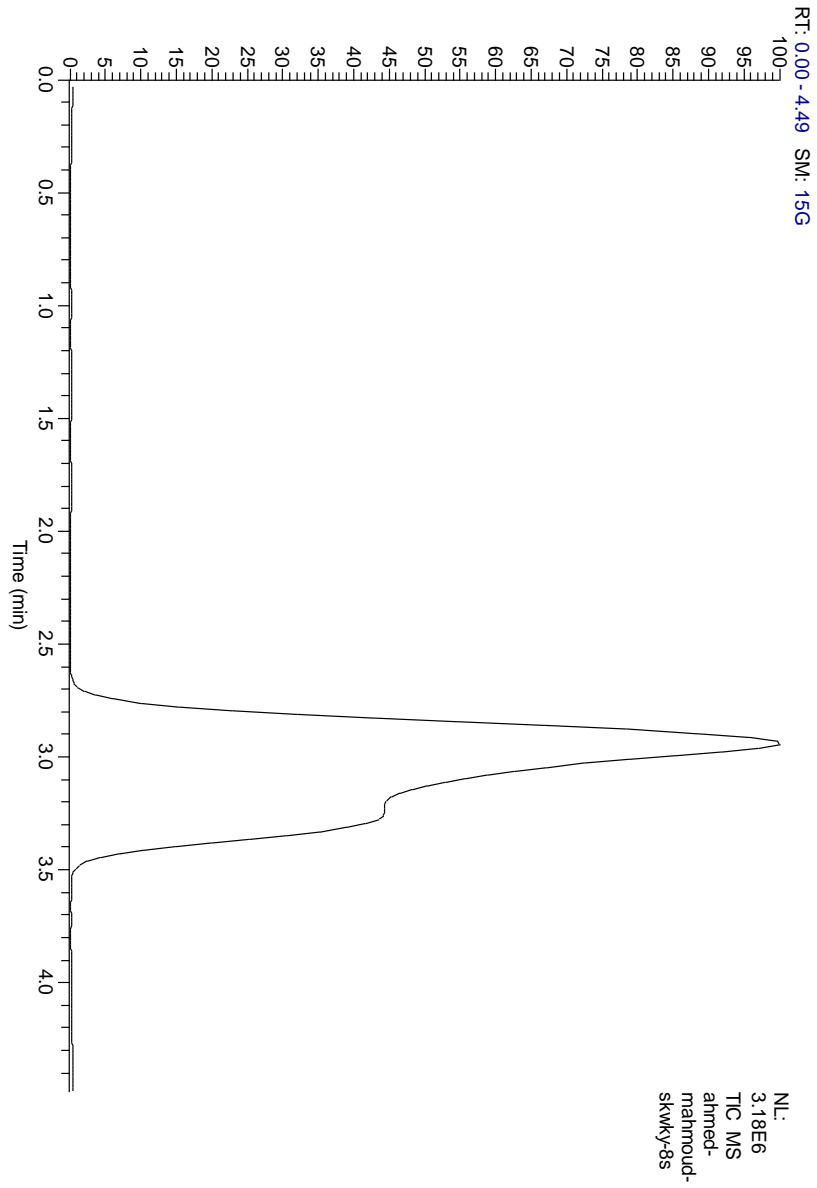
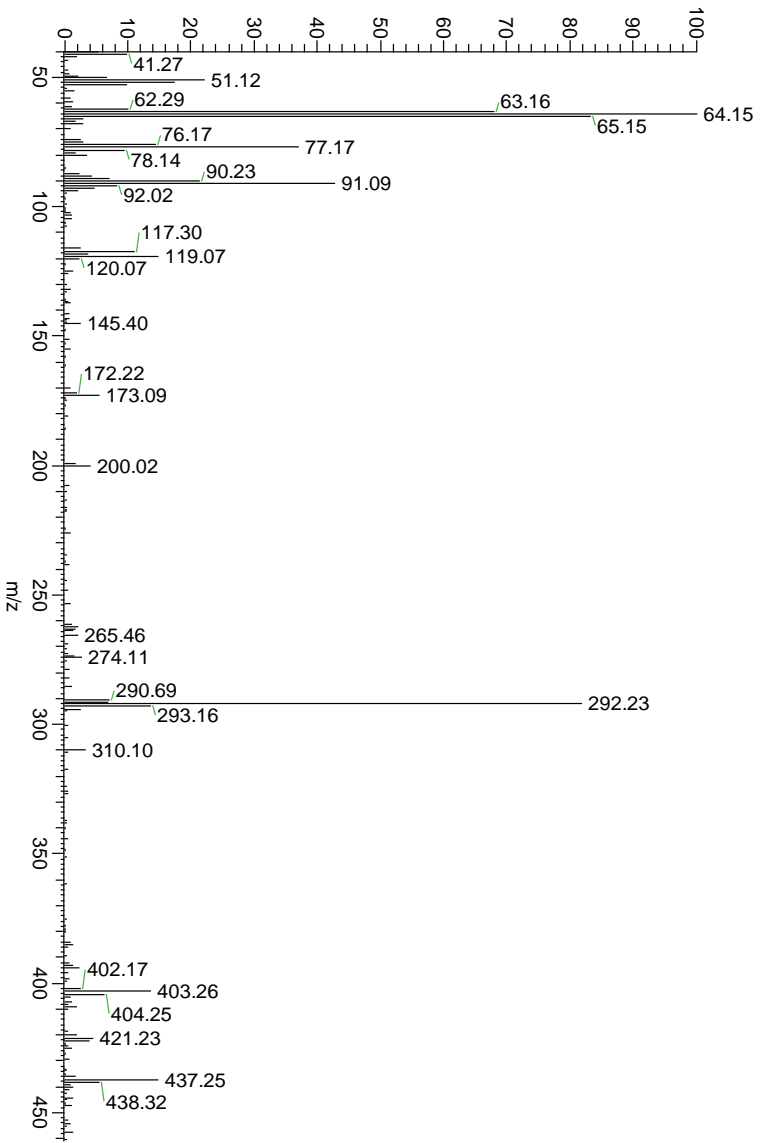
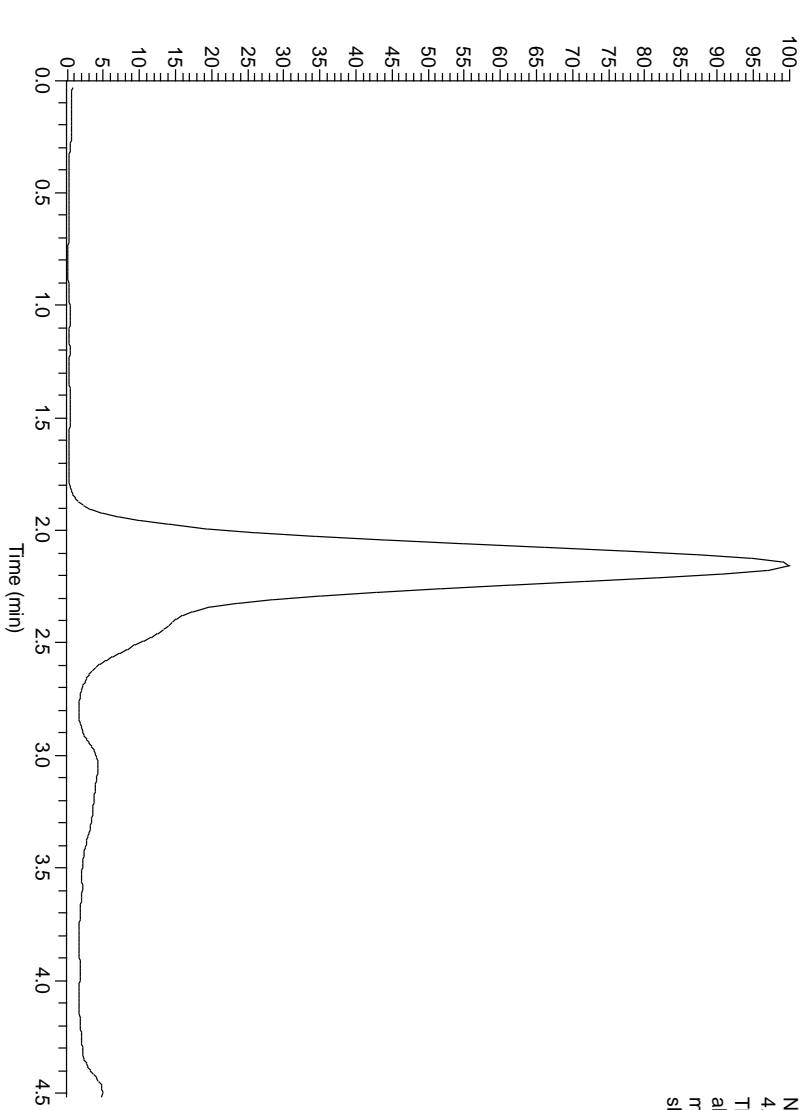


Figure S131. Mass spectrum of compound 18c



ahmed-mahmoud-skwy-9s #144 RT: 2.43 AV: 1 SB: 2 4.52 4.52 NL: 5.54E4
T: (0.0) + c EI Full ms [40.00-1000.00]

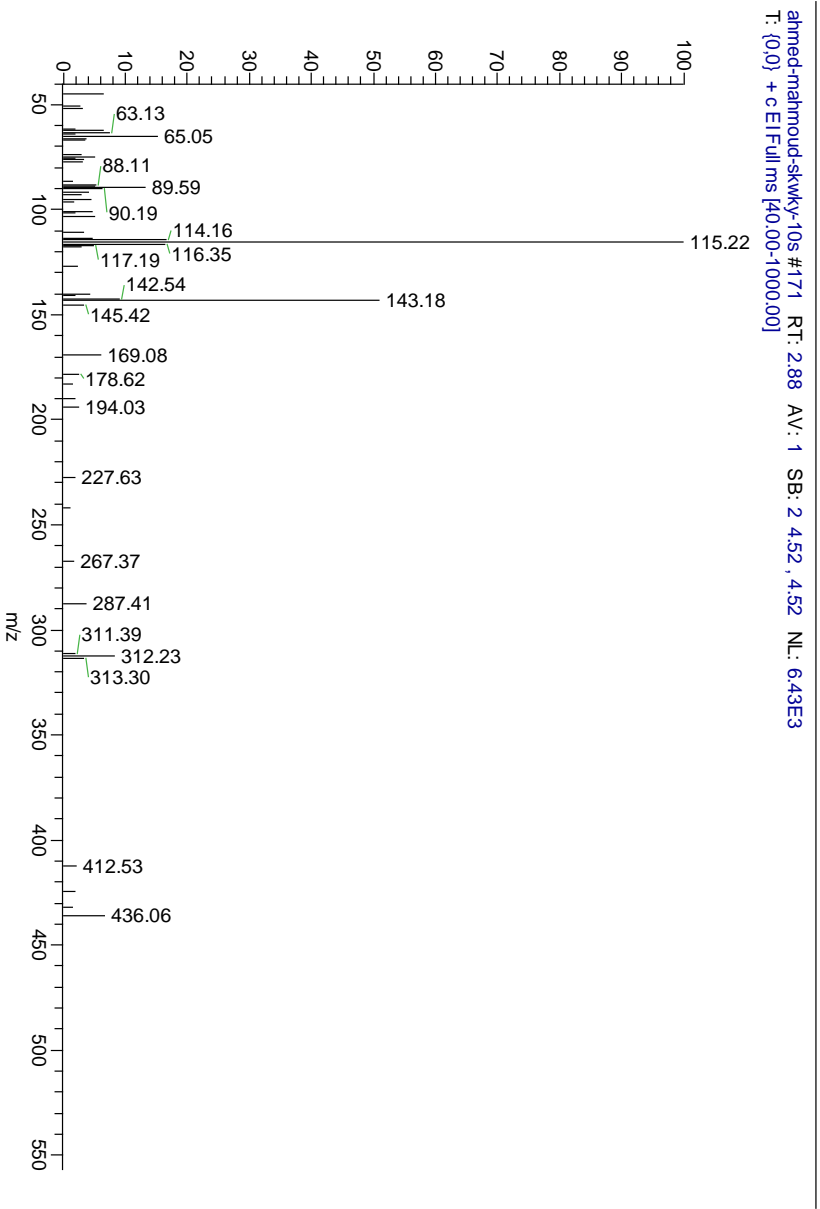


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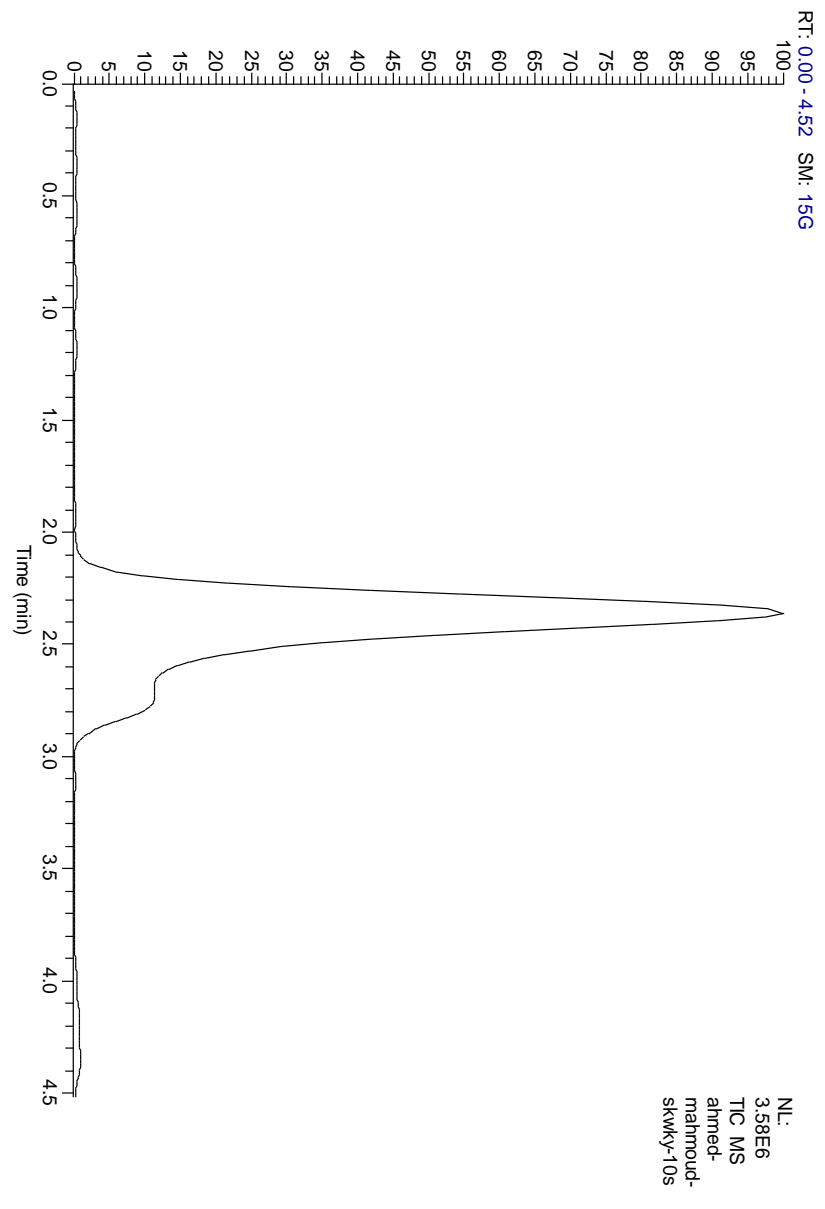
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Figure S132. Mass spectrum of compound 19a

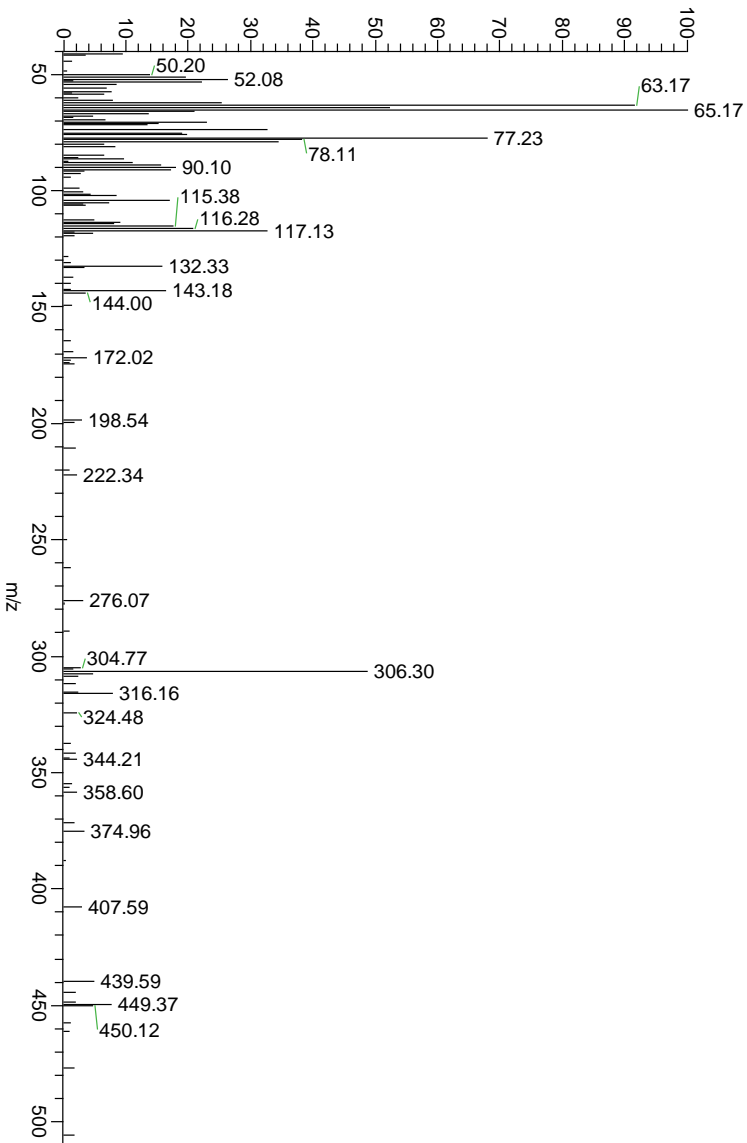


ahmed-mahmoud-skwy-10s #171 RT: 2.88 AV: 1 SB: 2 4.52, 4.52 NL: 643E3
T: (0.0) + c E I Full ms [40.00-1000.00]

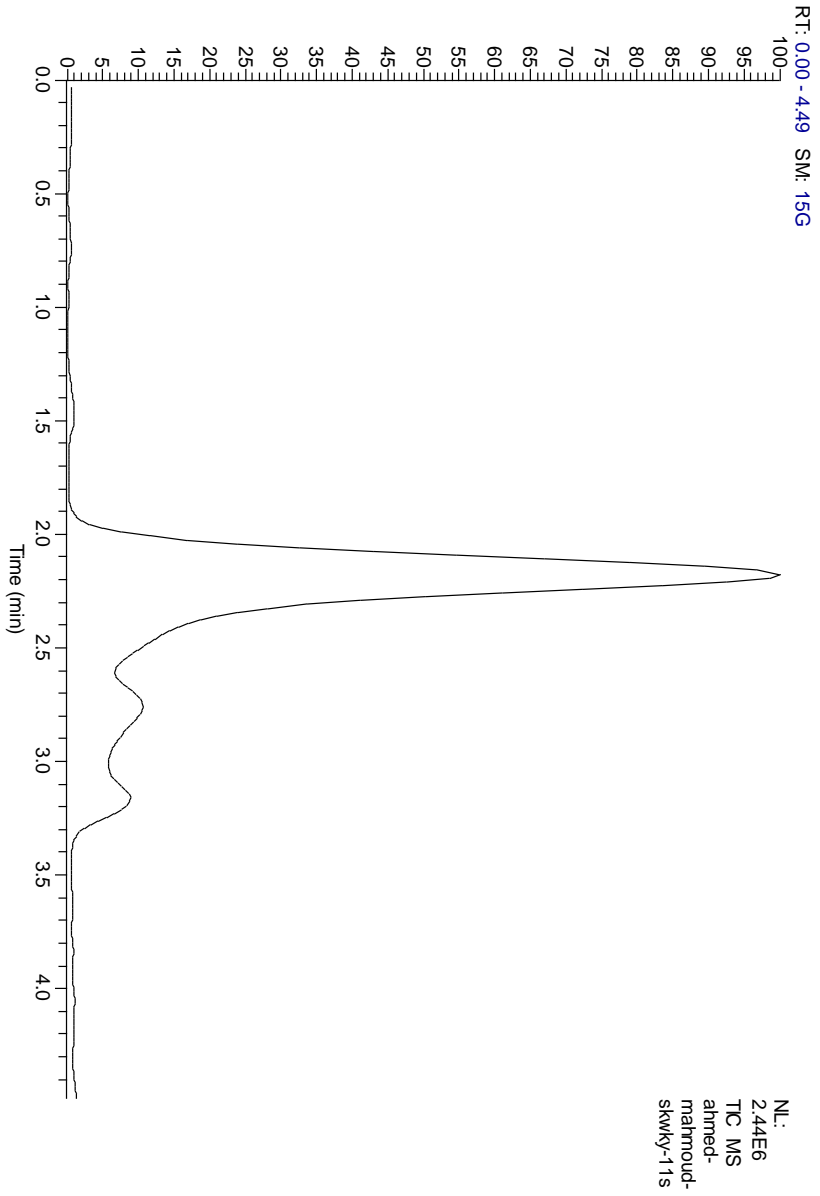


RT: 0.00 - 4.52 SM: 15G
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04-Aug-19 1:43:58 PM
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NL:
3.58E6
TIC MS
ahmed-
mahmoud-
skwy-10s

Figure S133. Mass spectrum of compound 19b



ahmed-nahmoud-skwy-11s #181 RT: 3.05 AV: 1 SB: 2.449, 4.49 NL: 1.05E4
T: (0.0) + c EI Full ms [40.00-1000.00]



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NL:
2.44E6
TIC MS
ahmed-
nahmoud-
skwy-11s

Figure S134. Mass spectrum of compound 19c

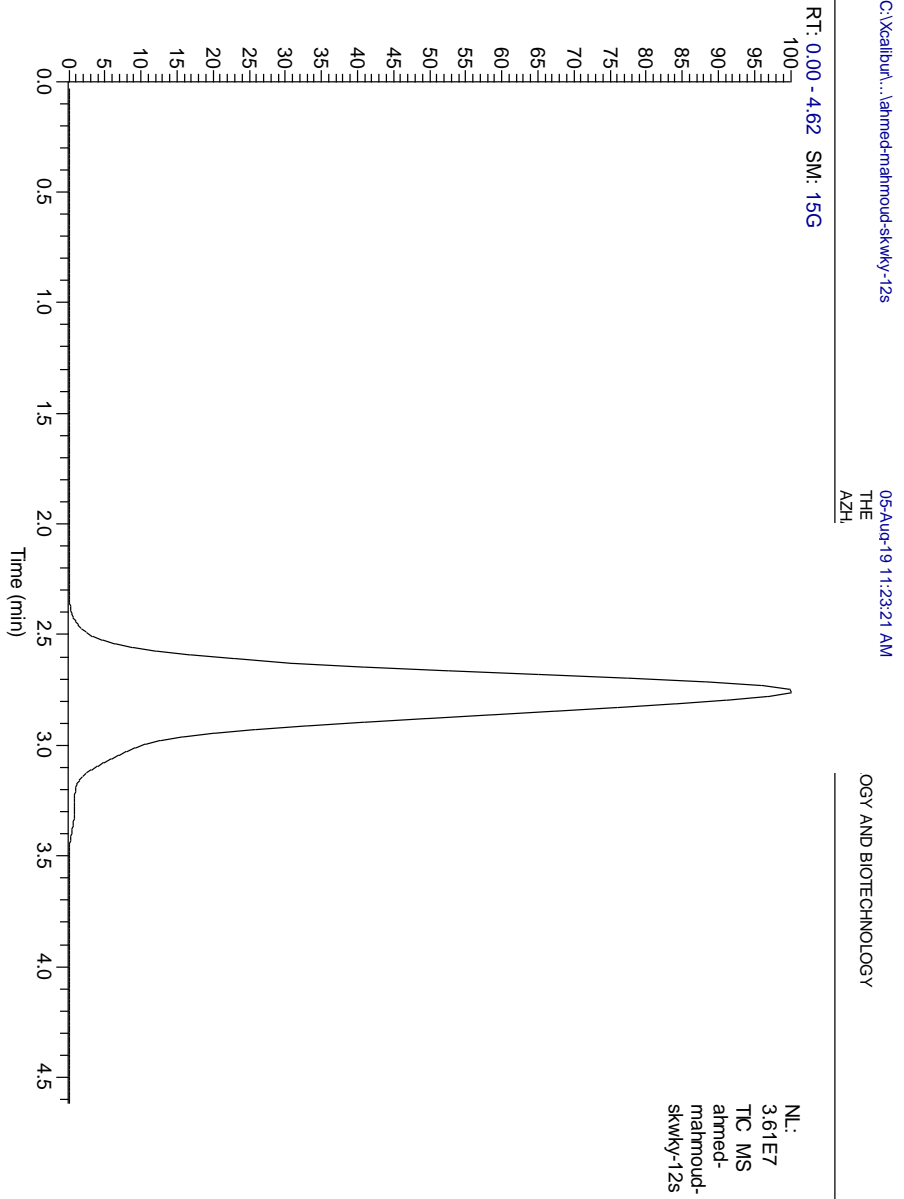
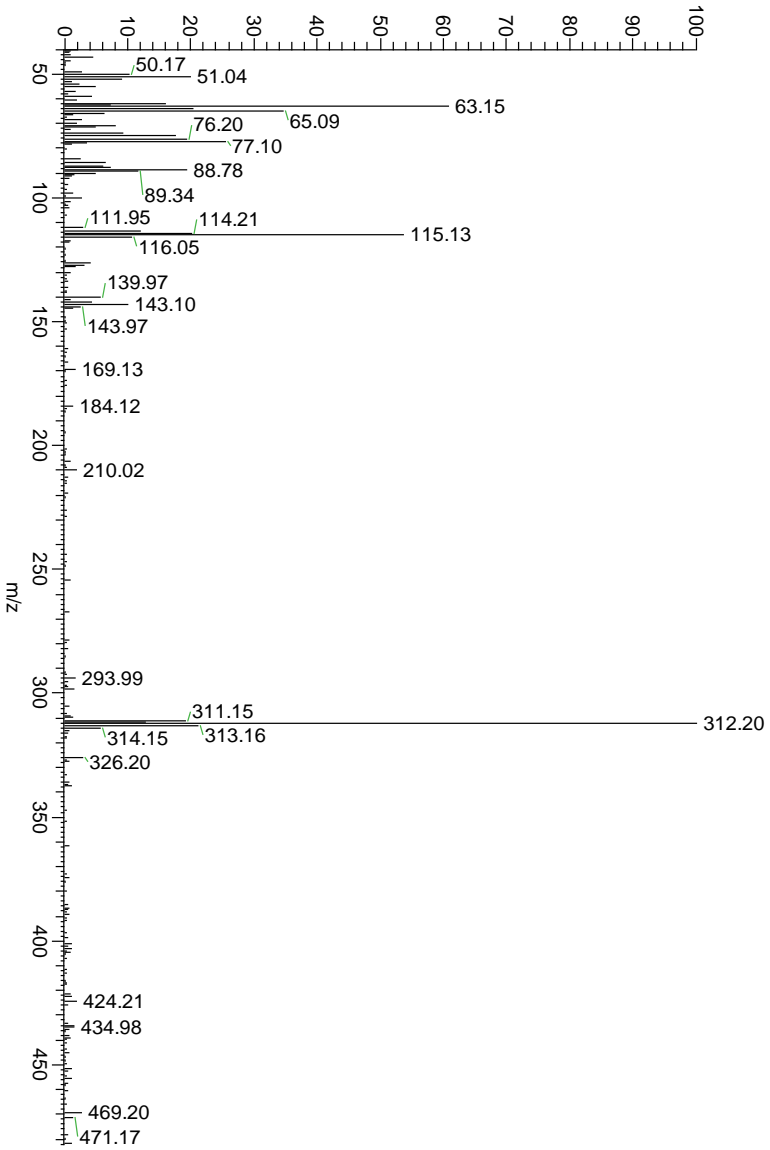
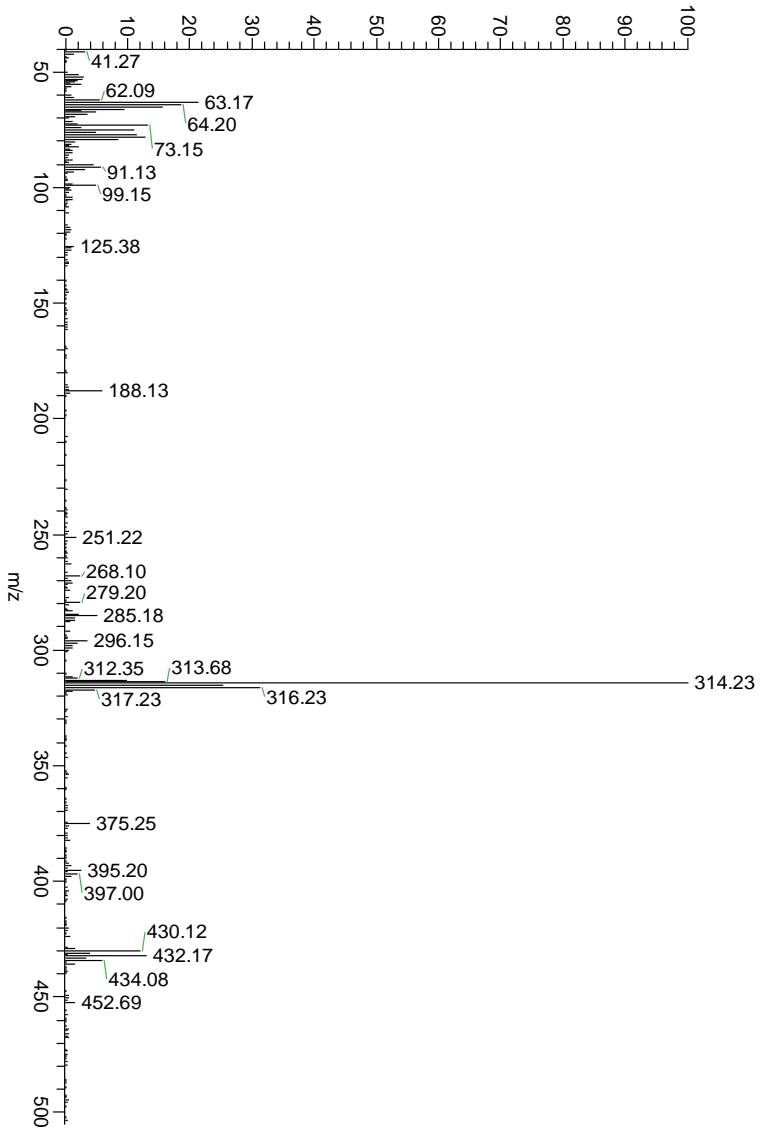


Figure S135. Mass spectrum of compound 20a



ahmed-mahmoud-shwki-17s #144 RT: 2.43 AV: 1 SB: 2 4.52 4.52 NL: 1.38E5
T: (0.0) + c E I Full ms [40.00-1000.00]

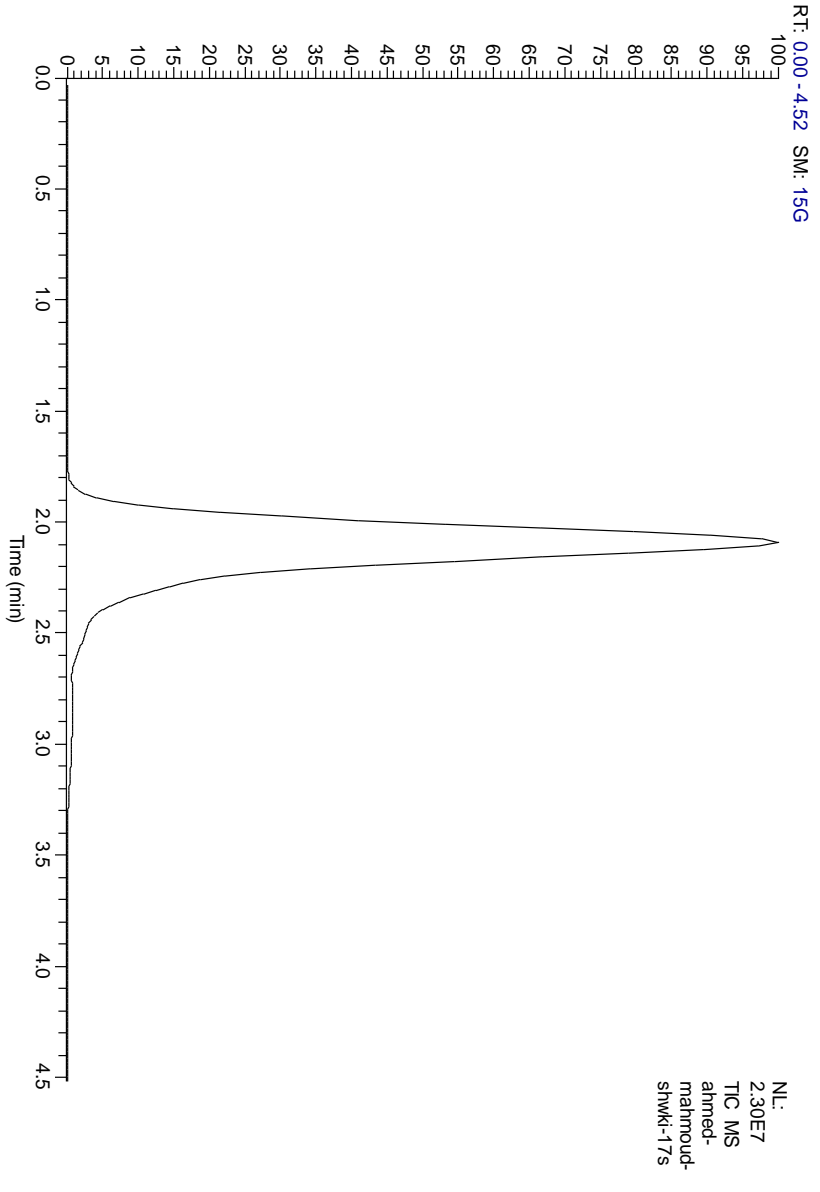
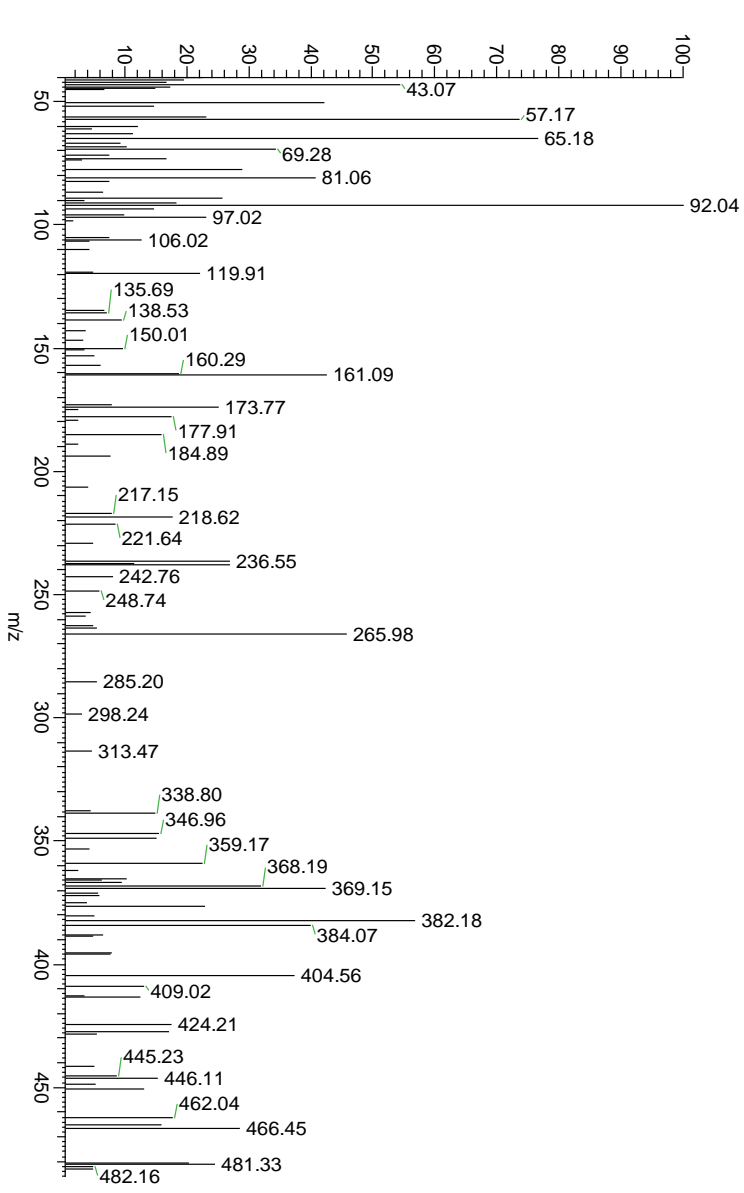
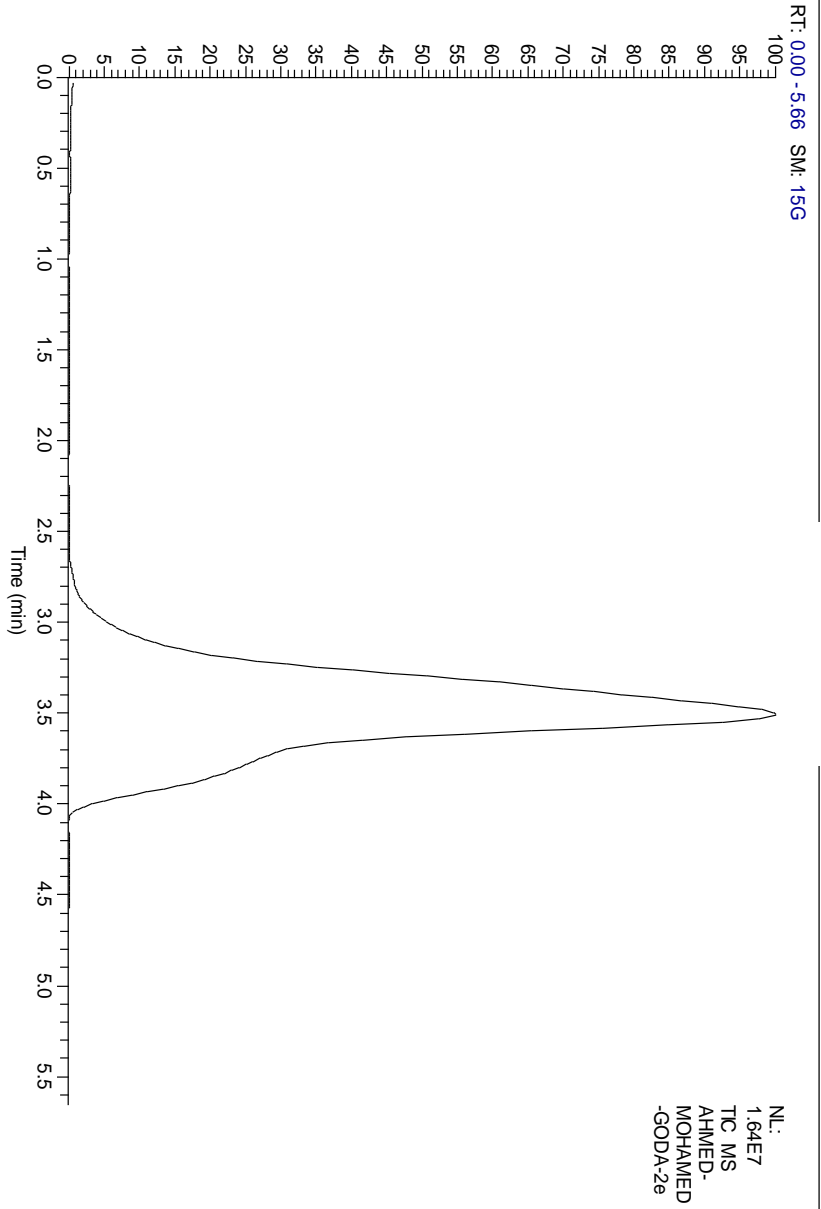


Figure S136. Mass spectrum of compound 20b

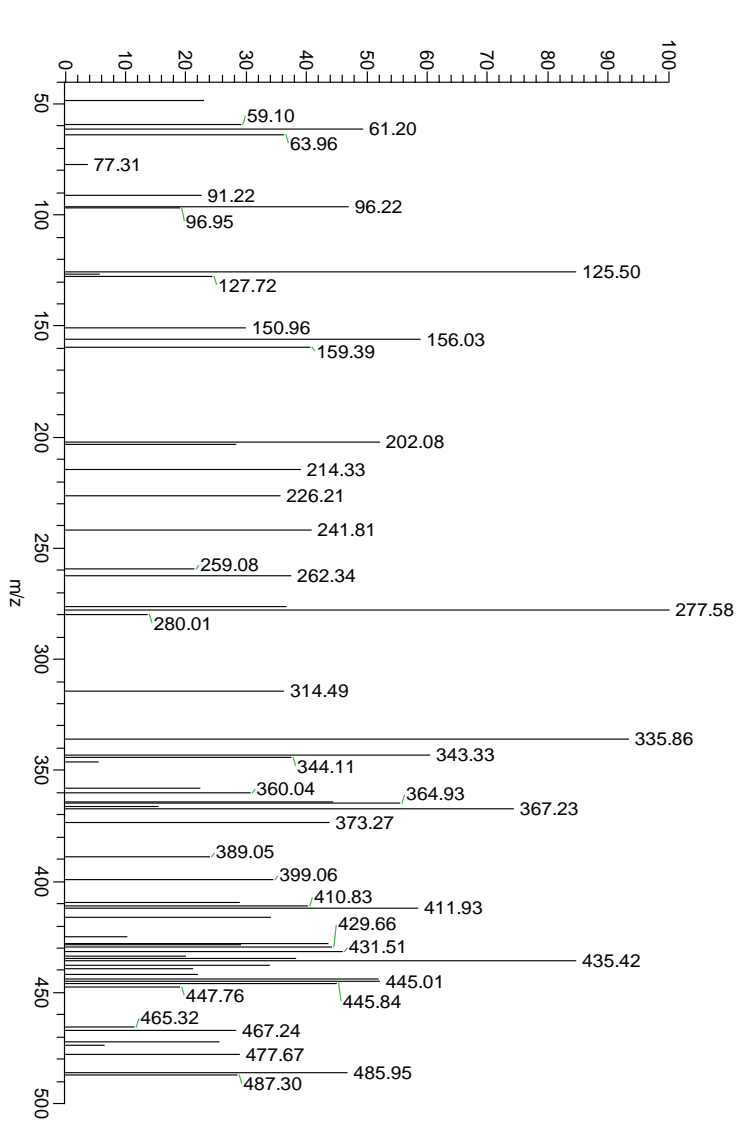


AHMED-MOHAMED-GODA-2e #268 RT: 4.50 AV: 1 NL: 2.29E3
T: {0.0} + cEI Full ms [40.00-1000.00]

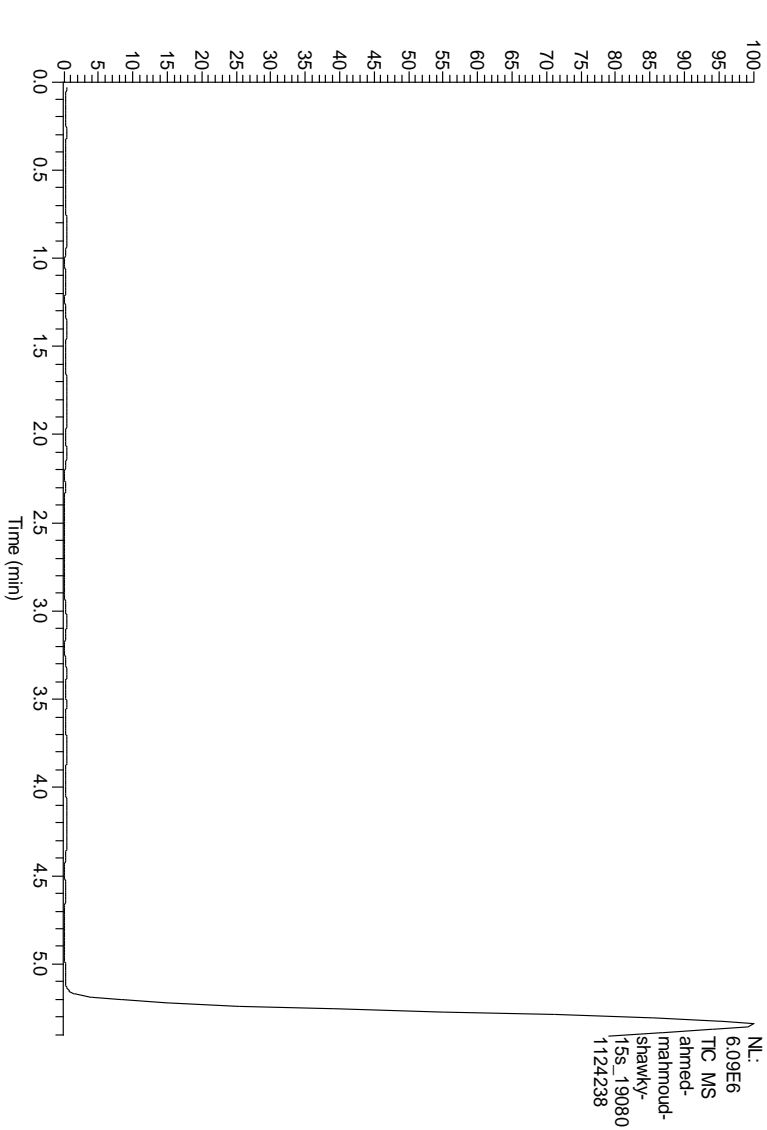


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AMED-GODA-2e
NL: 1.64E7
TIC MS
AHMED-MOHAMED-GODA-2e

Figure S137. Mass spectrum of compound 20b



ahmed-mahmoud-shawky-15s_190801124238 #223 RT: 3.75 AV: 1 SB: 2 4.55, 4.59 NL: 2.94E2
T: {0.0} + c EI Full ms [40.00-1000.00]



RT: 0.00 - 5.41 SM: 15G

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NL:
6.09E6
TIC MS
ahmed-
mahmoud-
shawky-
15s_19080
1124238

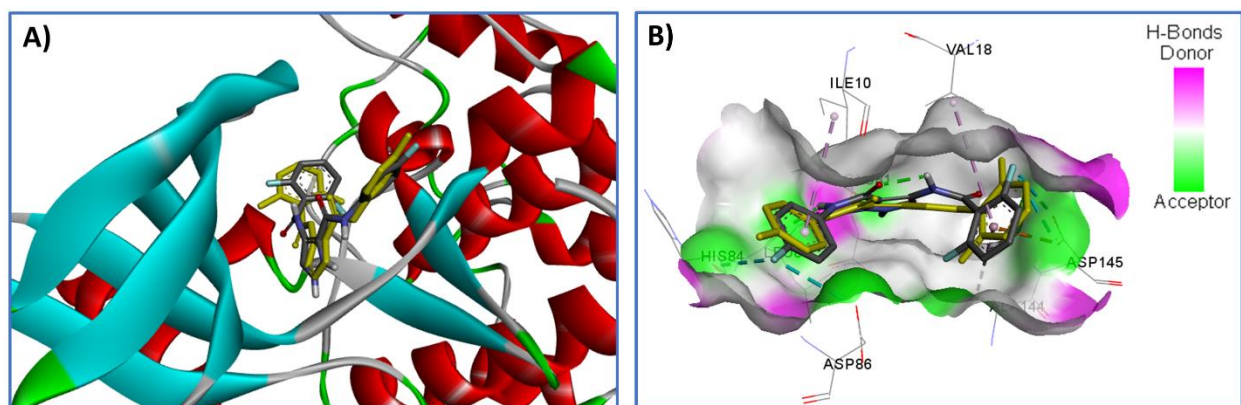


Figure S138. Binding modes/interactions of re-docked/bound LZ9 (compound **3**) into CDK-2 (pdb 2VTP): A) 3D binding mode of re-docked LZ9 (shown as sticks colored by element) into CDK-2 overlaid with the native ligand, LZ9 shown as yellow sticks; B) 3D binding mode of re-docked LZ9 (shown as sticks colored by element) overlaid with the native ligand (shown as yellow sticks) into CDK-2, the native ligand LZ9 shown as yellow sticks, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity.

Table S1. Docking results of the new compounds into CDK-2 (pdb: 2VTP).

Ligand	ΔG_b^a	K_i^b	HBs ^c	Atoms in H-bonding		Length ^d (Å)
				In ligand	In protein	
16a	-8.92	288.31 nM	4	CN	Thr14	2.17
				Urea NH	Lys33	1.95
				Urea NH	Asp145	1.78
				Urea NH	Asp145	1.97
16b	-8.73	398.11 nM	3	Urea NH	Asp86	2.04
				Urea NH	Gln131	2.13
				CONH	Asp145	2.10
16c	-8.60	500.18 nM	4	CN	Lys129	2.84
				CN	Gln131	2.36
				Urea NH	Asn132	2.36
				Urea NH	Asp145	1.84
17a	-9.84	60.97 nM	3	CN	Thr14	2.10
				Urea CO	Lys33	1.75
				Urea NH	Asp145	2.05
17b	-9.65	85.09 nM	2	Urea CO	Leu83	1.92
				CONH	Gln131	2.24
17c	-9.48	111.80 nM	2	CN	Leu83	2.09
				Urea NH	Asp86	2.06
18a	-8.89	301.92 nM	2	CN	Leu83	1.91
				Urea NH	Leu83	2.06
18c	-9.54	101.92 nM	3	Urea NH	Leu83	2.27
				CN	Leu83	2.19
				CONH	Gln131	2.15
19b	-10.41	23.47 nM	3	CN	Leu83	2.17
				Urea NH	Asp86	2.08
				CONH	Gln131	2.81
19c	-10.67	15.06 nM	2	Urea CO	Lys33	2.34
				Urea NH	Asp145	2.04
20b	-10.10	39.21 nM	4	CN	Leu83	2.05
				Thiourea S	Leu83	2.86
				Thiourea NH	Asp86	3.06
				CONH	Gln131	2.39
20c	-9.76	70.64 nM	2	Urea NH	Asp86	2.09
				CONH	Gln131	1.96

^a Binding free energy (kcal/mol);^b Inhibition constant (n/μM)^c HBs, number of hydrogen bonds^d length in angstrom (Å)

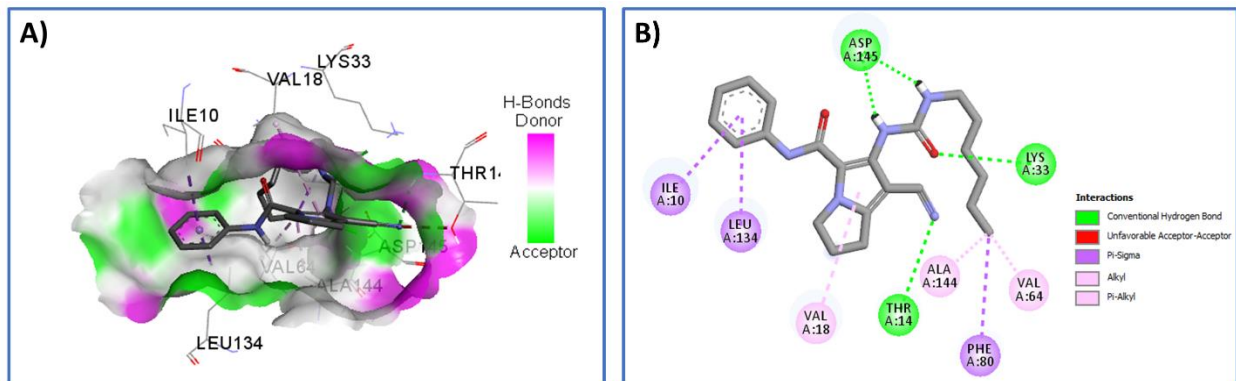


Figure S139. Binding modes/interactions of compound **16a** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **16a** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **16a** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

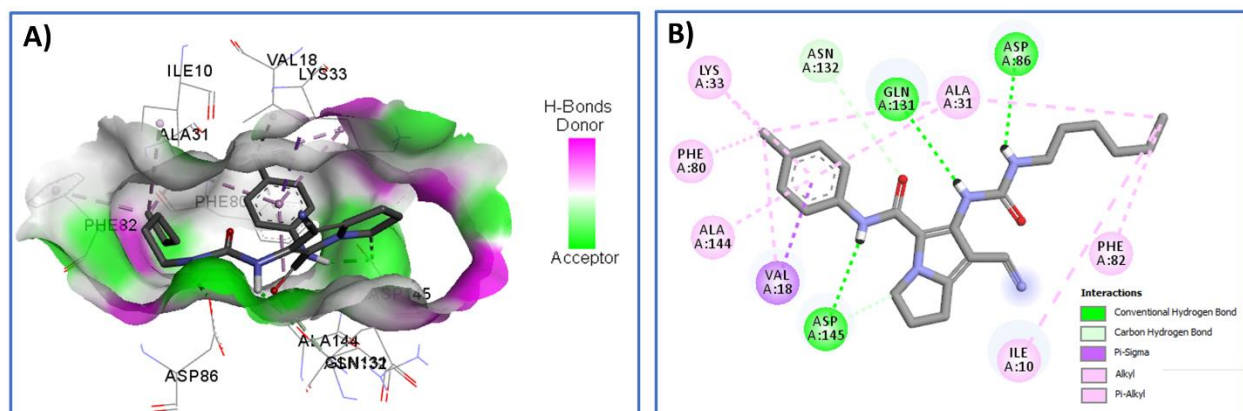


Figure S140. Binding modes/interactions of compound **16b** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **16b** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **16b** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

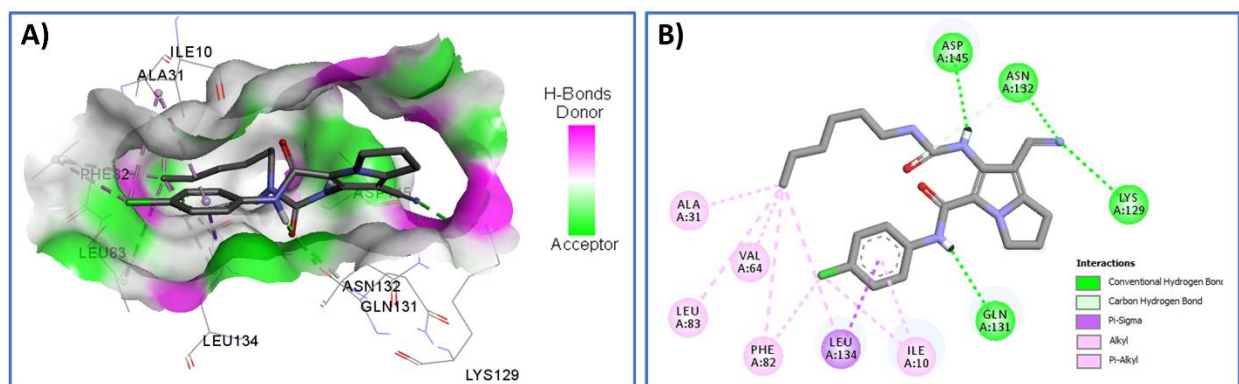


Figure S141. Binding modes/interactions of compound **16c** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **16c** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **16c** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

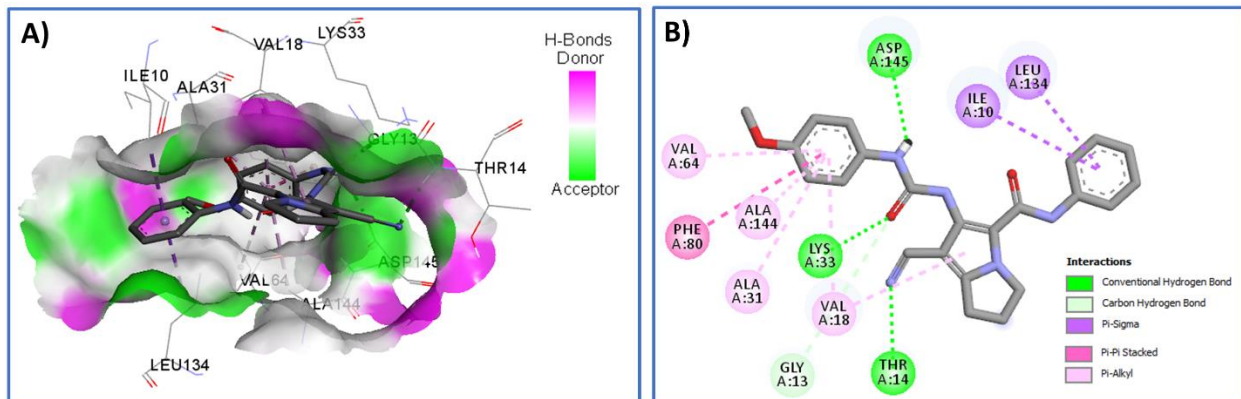


Figure S142. Binding modes/interactions of compound **17a** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **17a** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **17a** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

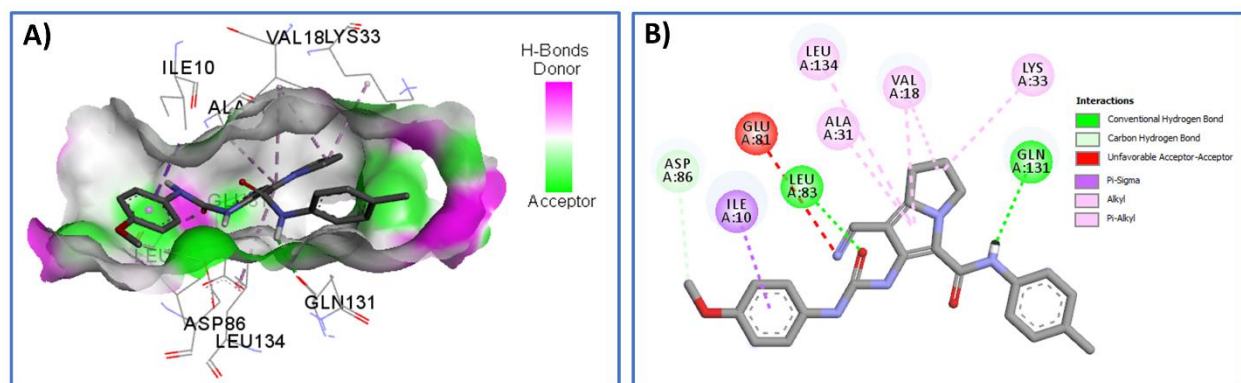


Figure S143. Binding modes/interactions of compound **17b** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **17b** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **17b** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

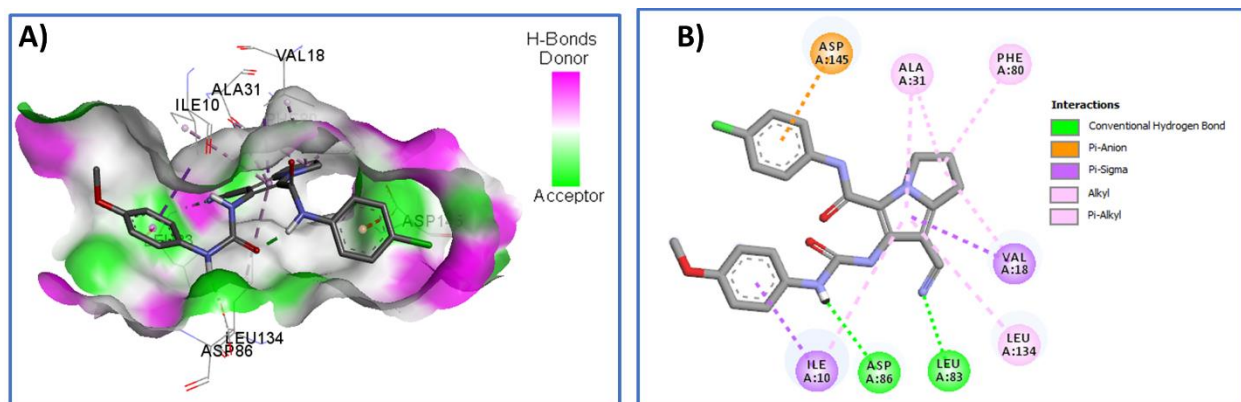


Figure S144. Binding modes/interactions of compound **17c** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **17c** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **17c** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

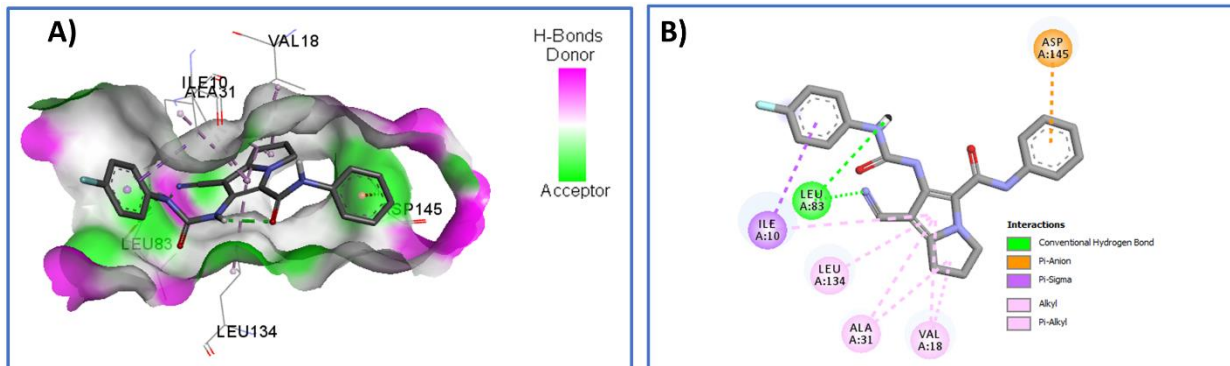


Figure S145. Binding modes/interactions of compound **18a** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **18a** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **18a** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

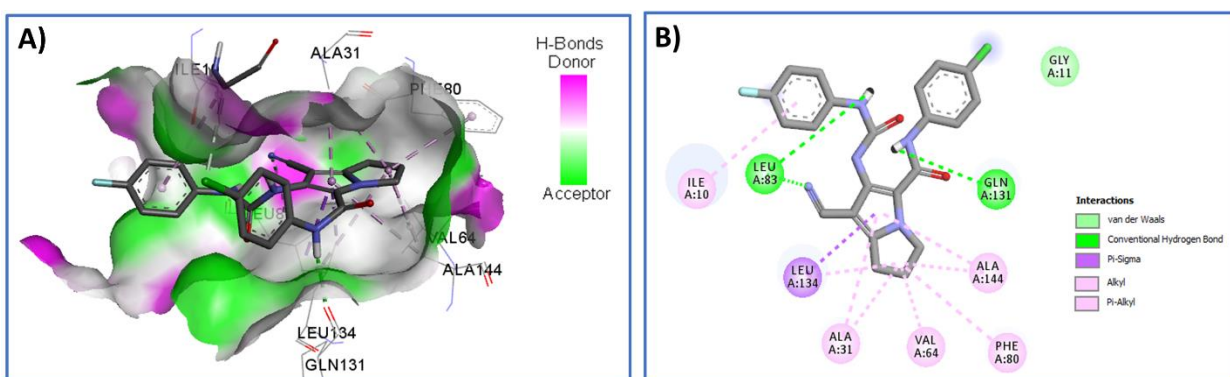


Figure S146. Binding modes/interactions of compound **18c** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **18c** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **18c** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

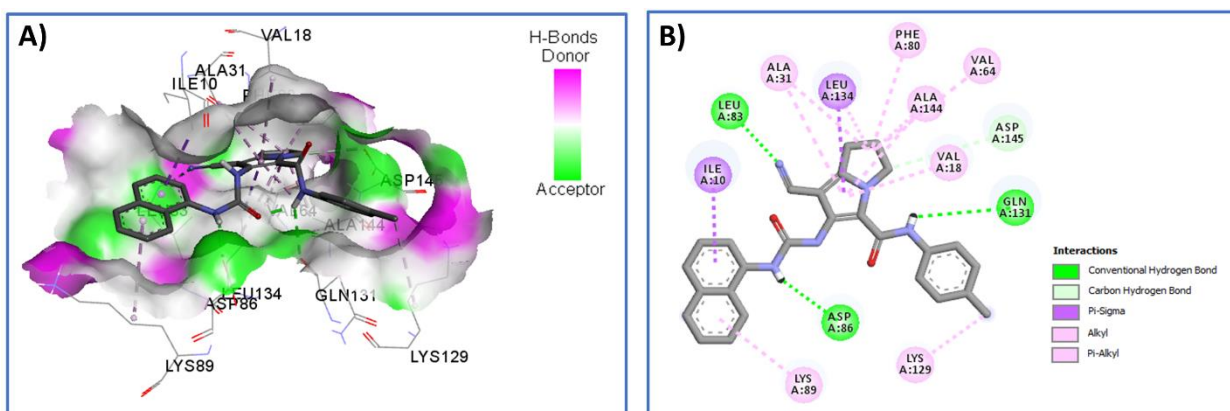


Figure S147. Binding modes/interactions of compound **19b** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **19b** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **19b** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

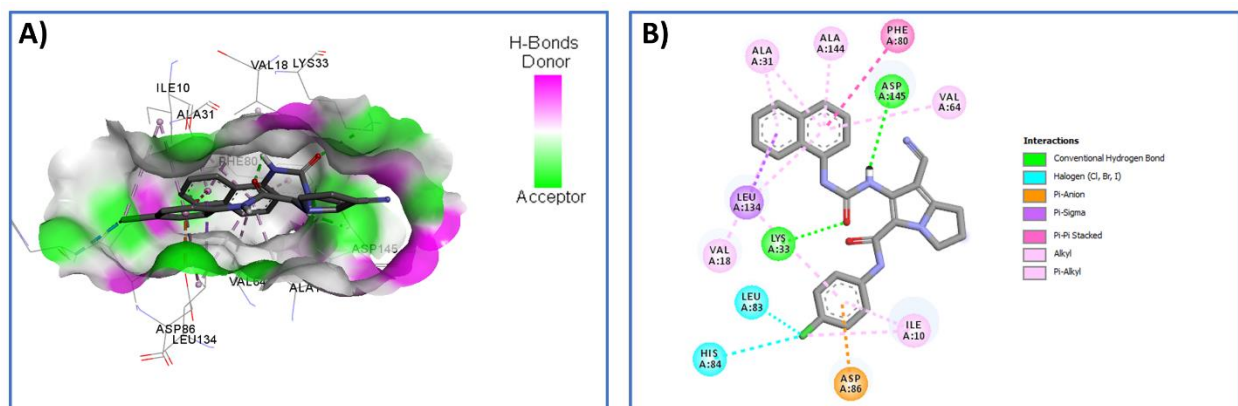


Figure S148. Binding modes/interactions of compound **19c** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **19c** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **19c** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

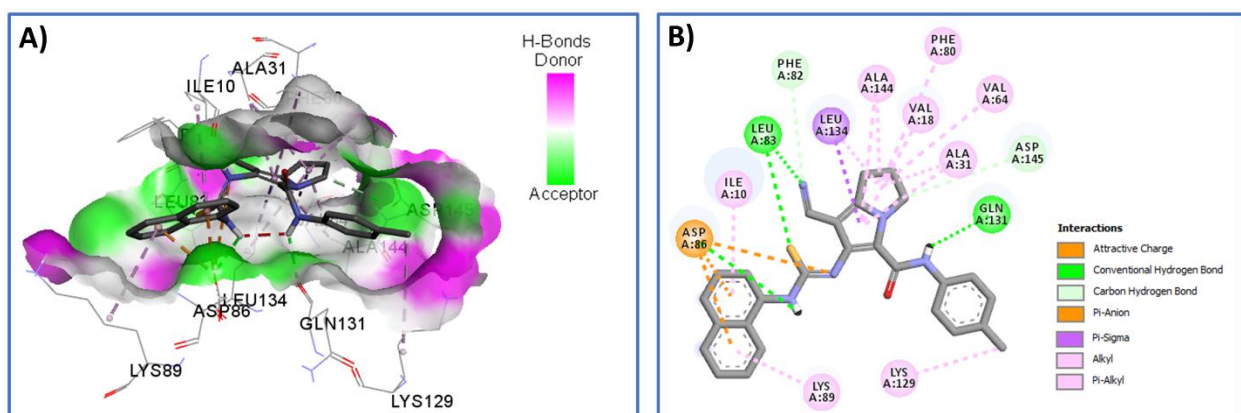


Figure S149. Binding modes/interactions of compound **20b** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **20b** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **20b** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

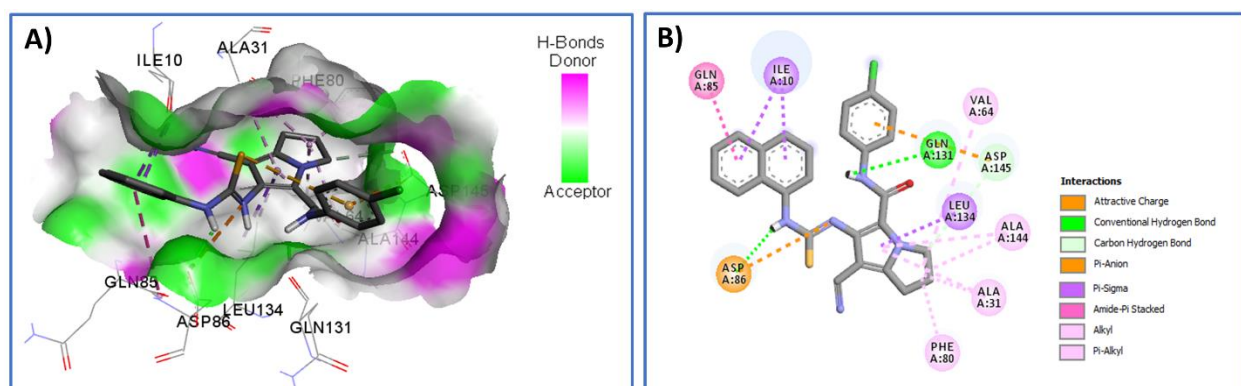


Figure S150. Binding modes/interactions of compound **20c** into CDK-2 (pdb: 2VTP): A) 3D binding mode of compound **20c** into the active site of CDK-2, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity; B) 3D binding mode of compound **20c** into CDK-2 showing different types of hydrogen bonding and hydrophobic interactions.

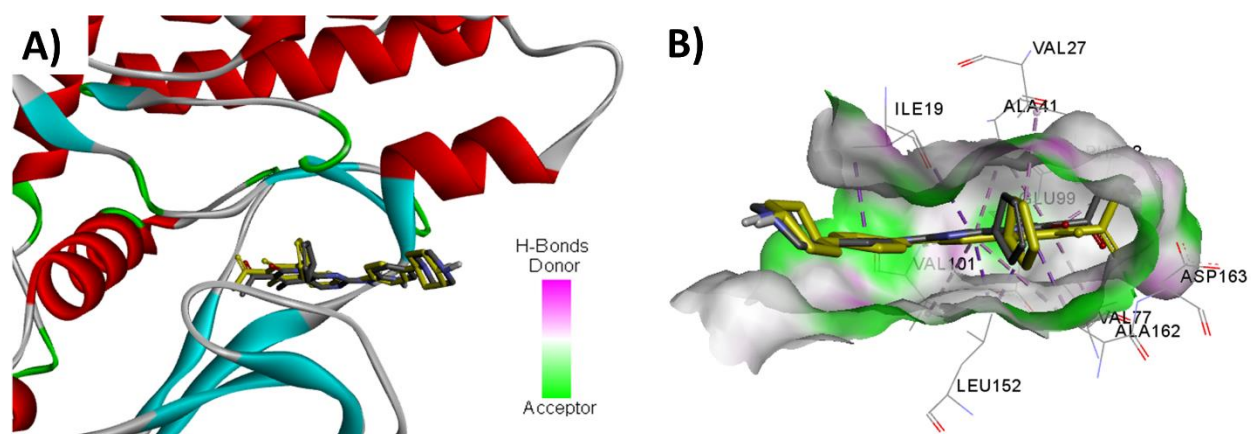


Figure S151. Binding modes/interactions of re-docked/bound palbociclib into CDK-6 (pdb 2EUF): A) 3D binding mode of re-docked palbociclib (shown as sticks colored by element) into CDK-6 overlaid with the native ligand (shown as yellow sticks); B) 3D binding mode of re-docked palbociclib (shown as sticks colored by element) overlaid with the native ligand (shown as yellow sticks) into CDK-6, the native ligand palbociclib shown as yellow sticks, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity.

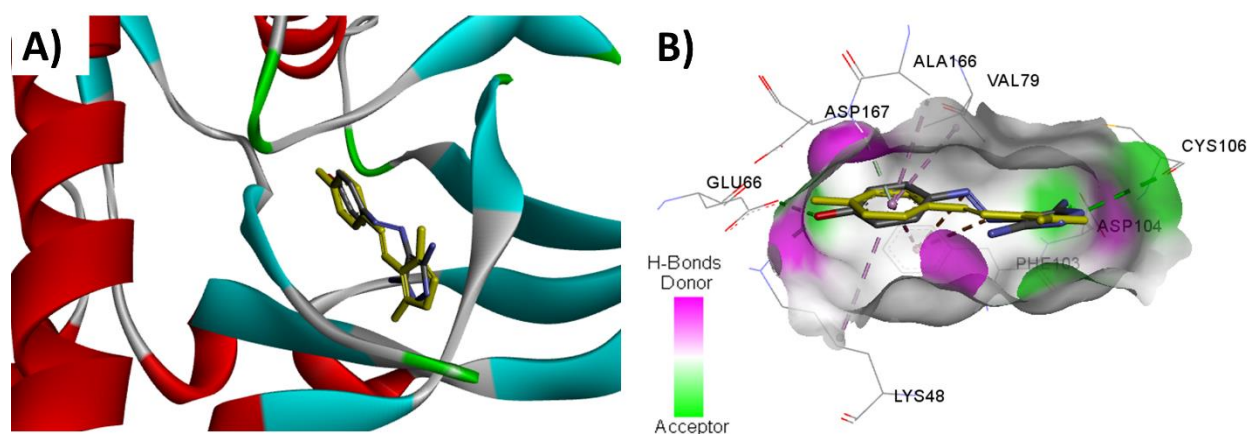
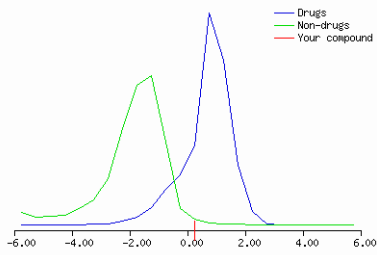
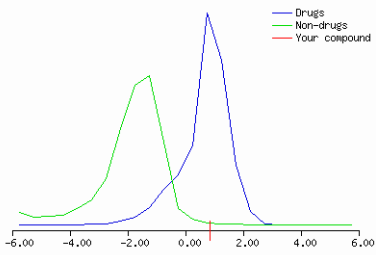
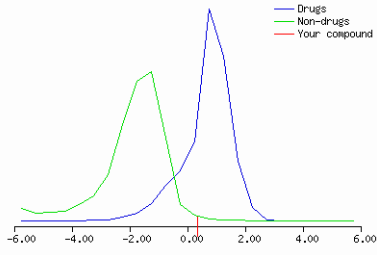
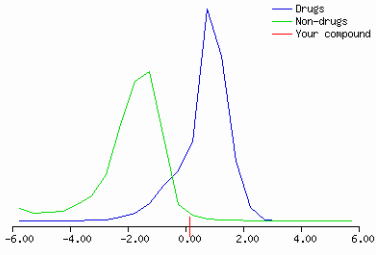
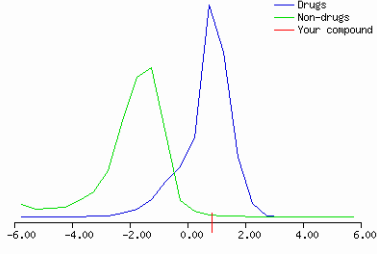
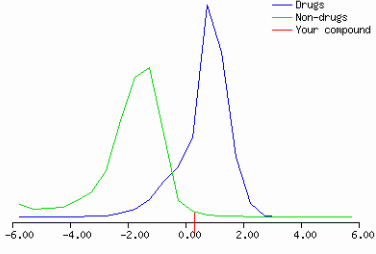
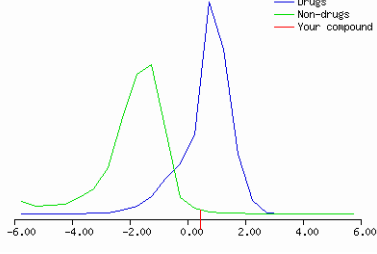
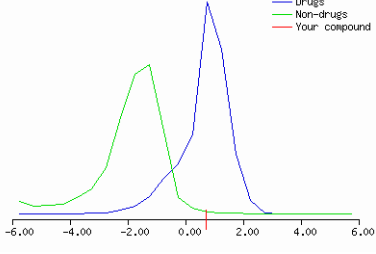
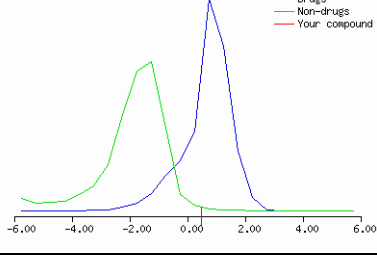
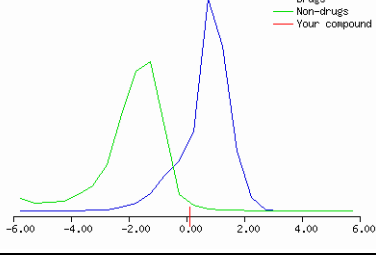


Figure S152. Binding modes/interactions of re-docked/bound palbociclib into CDK-9 (pdb 3TNH): A) 3D binding mode of re-docked CAN508 (shown as sticks colored by element) into CDK-9 overlaid with the native ligand (shown as yellow sticks); B) 3D binding mode of re-docked CAN508 (shown as sticks colored by element) overlaid with the native ligand (shown as yellow sticks) into CDK-9, the native ligand CAN508 shown as yellow sticks, receptor shown as hydrogen-bond surface, hydrogen atoms were omitted for clarity

Table S2. DLS figures and scores of compounds 9 and 16-20a-c.

Comp.		DLS	Comp.		DLS
16a		0.24	18c		0.86
16b		0.35	19a		0.14
16c		0.84	19b		0.30
17a		0.45	19c		0.71
17b		0.48	20a		0.11

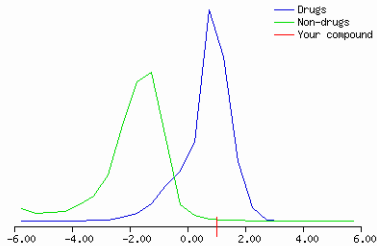
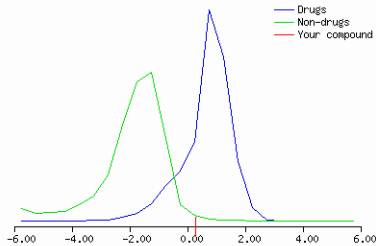
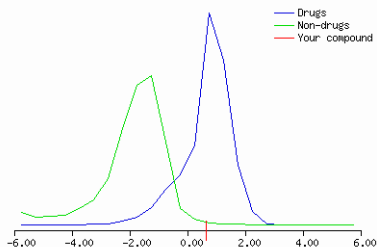
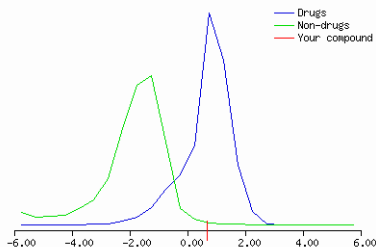
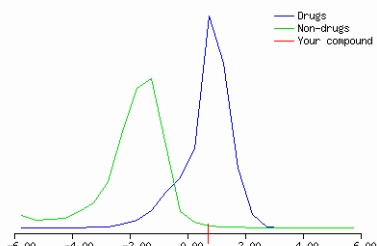
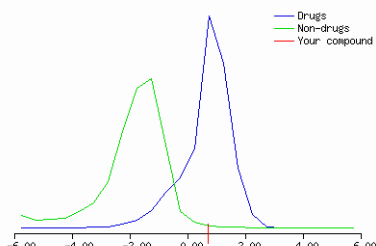
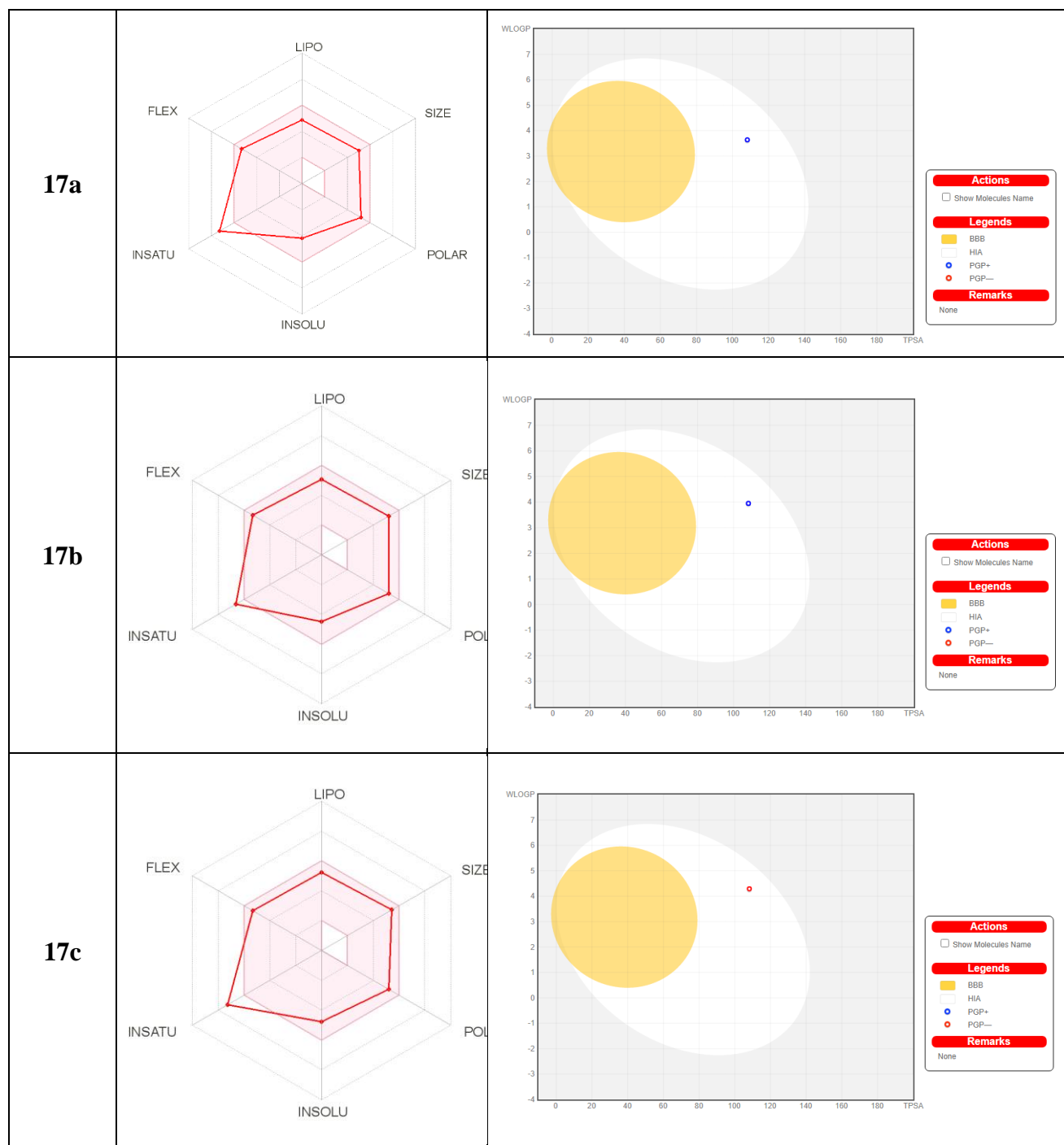
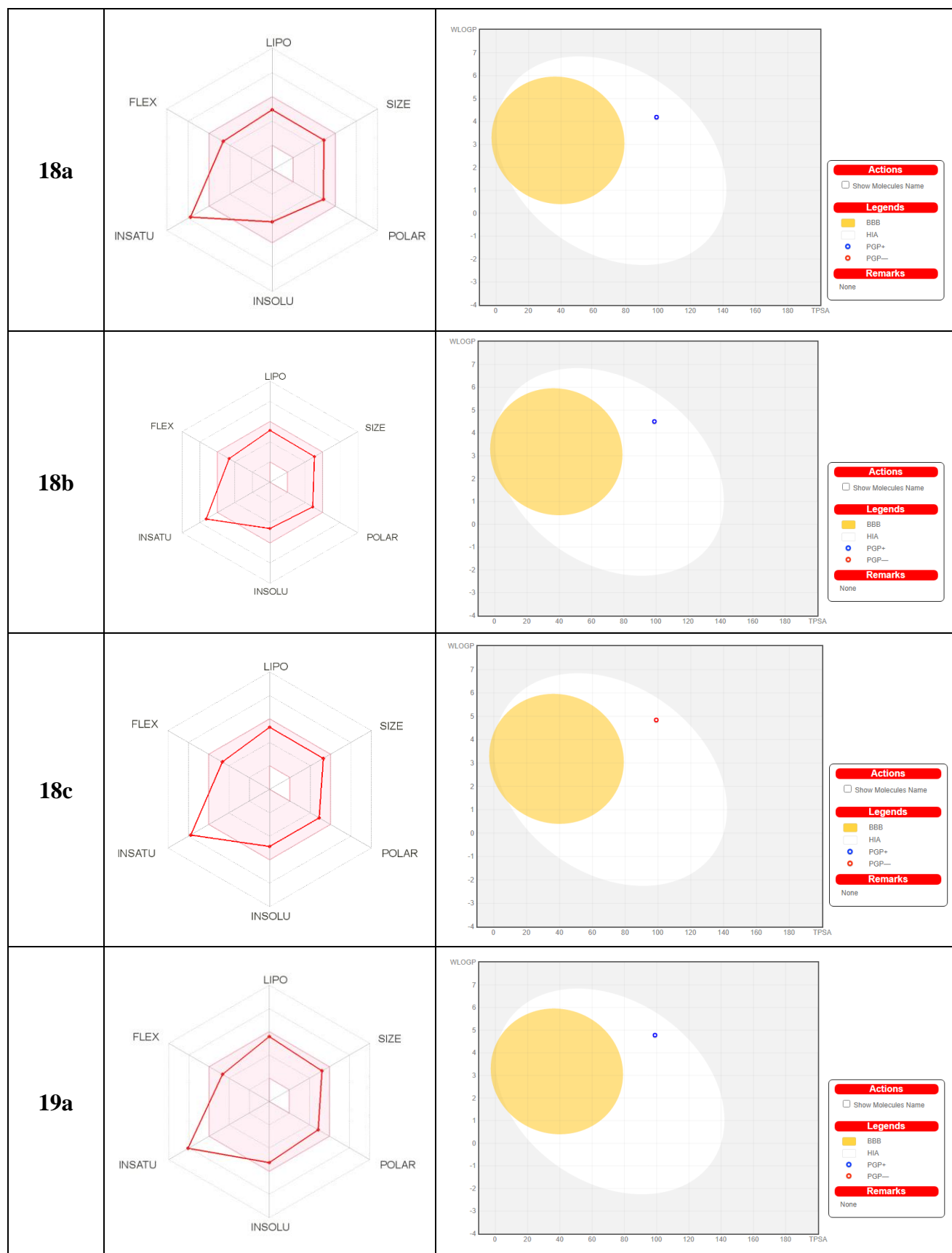
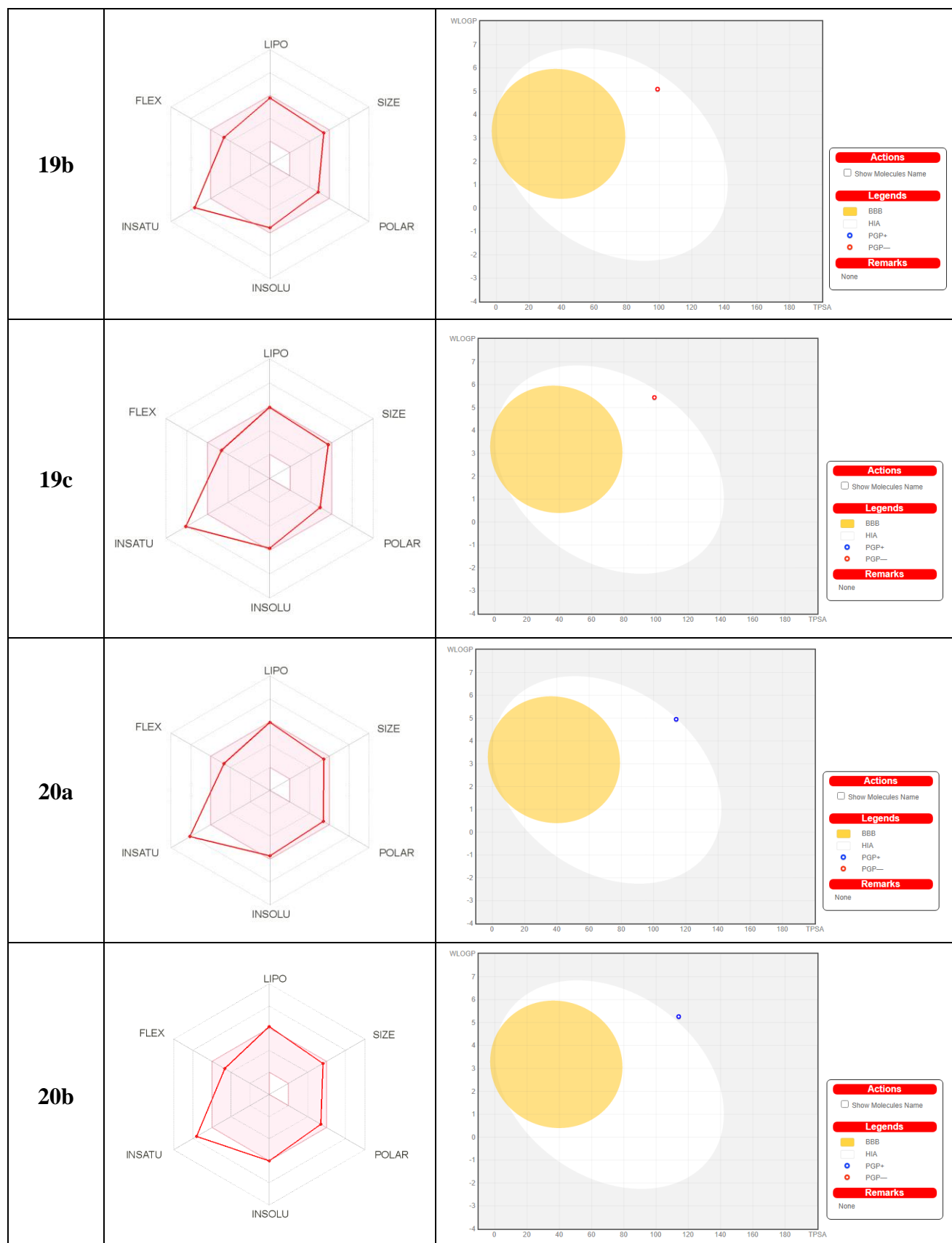
17c	 <p>Density plot for compound 17c. The x-axis ranges from -6.00 to 6.00. The 'Drugs' distribution (blue) is centered at approximately 1.0. The 'Non-drugs' distribution (green) is centered at approximately -1.5. The 'Your compound' distribution (red) is centered at approximately 0.5. The legend indicates: Drugs (blue), Non-drugs (green), Your compound (red).</p>	1.01	20b	 <p>Density plot for compound 20b. The x-axis ranges from -6.00 to 6.00. The 'Drugs' distribution (blue) is centered at approximately 1.0. The 'Non-drugs' distribution (green) is centered at approximately -1.5. The 'Your compound' distribution (red) is centered at approximately 0.5. The legend indicates: Drugs (blue), Non-drugs (green), Your compound (red).</p>	0.28
18a	 <p>Density plot for compound 18a. The x-axis ranges from -6.00 to 6.00. The 'Drugs' distribution (blue) is centered at approximately 1.0. The 'Non-drugs' distribution (green) is centered at approximately -1.5. The 'Your compound' distribution (red) is centered at approximately 0.5. The legend indicates: Drugs (blue), Non-drugs (green), Your compound (red).</p>	0.65	20c	 <p>Density plot for compound 20c. The x-axis ranges from -6.00 to 6.00. The 'Drugs' distribution (blue) is centered at approximately 1.0. The 'Non-drugs' distribution (green) is centered at approximately -1.5. The 'Your compound' distribution (red) is centered at approximately 0.5. The legend indicates: Drugs (blue), Non-drugs (green), Your compound (red).</p>	0.69
18b	 <p>Density plot for compound 18b. The x-axis ranges from -6.00 to 6.00. The 'Drugs' distribution (blue) is centered at approximately 1.0. The 'Non-drugs' distribution (green) is centered at approximately -1.5. The 'Your compound' distribution (red) is centered at approximately 0.5. The legend indicates: Drugs (blue), Non-drugs (green), Your compound (red).</p>	0.71	9	 <p>Density plot for compound 9. The x-axis ranges from -6.00 to 6.00. The 'Drugs' distribution (blue) is centered at approximately 1.0. The 'Non-drugs' distribution (green) is centered at approximately -1.5. The 'Your compound' distribution (red) is centered at approximately 0.5. The legend indicates: Drugs (blue), Non-drugs (green), Your compound (red).</p>	0.71

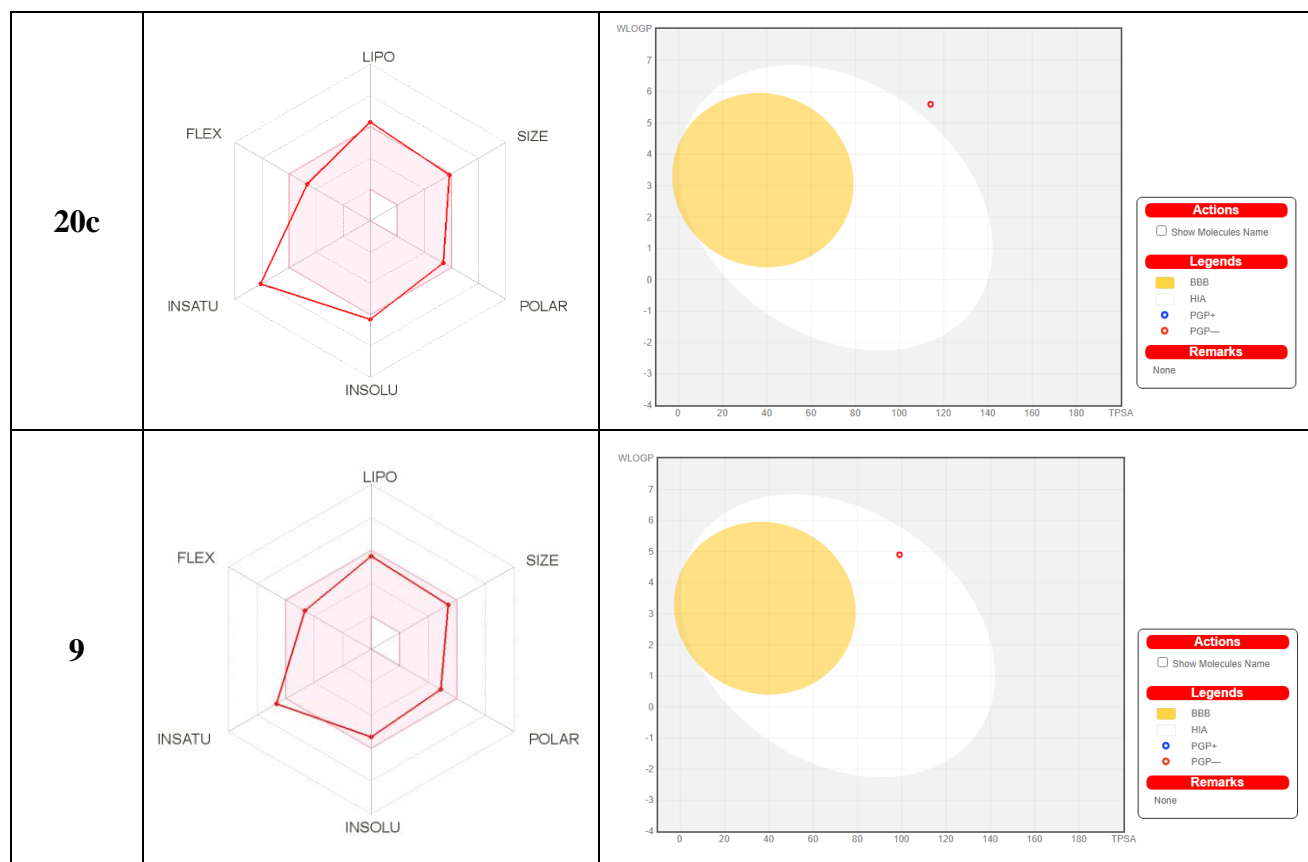
Table S3. The suitable physicochemical space for oral bioavailability and Boiled-Egg models of compounds **9** and **16-20a-c**.

Comp.	Physicochemical space	Boiled-Egg model
16a		
16b		
16c		









The suitable physicochemical space figures and BOILED-Egg's models were calculated using SwissADME (<http://www.swissadme.ch/>)

The colored zone is the suitable physicochemical space for oral bioavailability

- LIPO (lipophilicity) -0.7 to +5.0
- SIZE (MW), 150g/mol to 500g/mol
- POLAR (polarity): $20 \text{ \AA}^2 < \text{TPSA} < 130 \text{ \AA}^2$.
- INSOLU (insolubility): $0 < \text{Log S (ESOL)} < 6$
- INSATU (insaturation): $0.25 < \text{Fraction Csp3} < 1$
- Flex (flexibility): $0 < \text{Num. rotatable bonds} < 9$

BOILED-Egg's models

- Points located in BOILED-Egg's yolk are molecules predicted to passively permeate through the blood-brain barrier (BBB).
- Points located in BOILED-Egg's white are molecules predicted to passively absorbed by the GIT.
- Blue dots are for molecules predicted to be effluated from the CNS by P-glycoprotein.
- Red dots are for molecules predicted not to be effluated from the CNS by P-glycoprotein.

