

Supporting information  
for  
**Synthesis of pyrrolo[1,2-*a*]pyrimidine  
enantiomers via domino ring closure retro  
Diels-Alder protocol**

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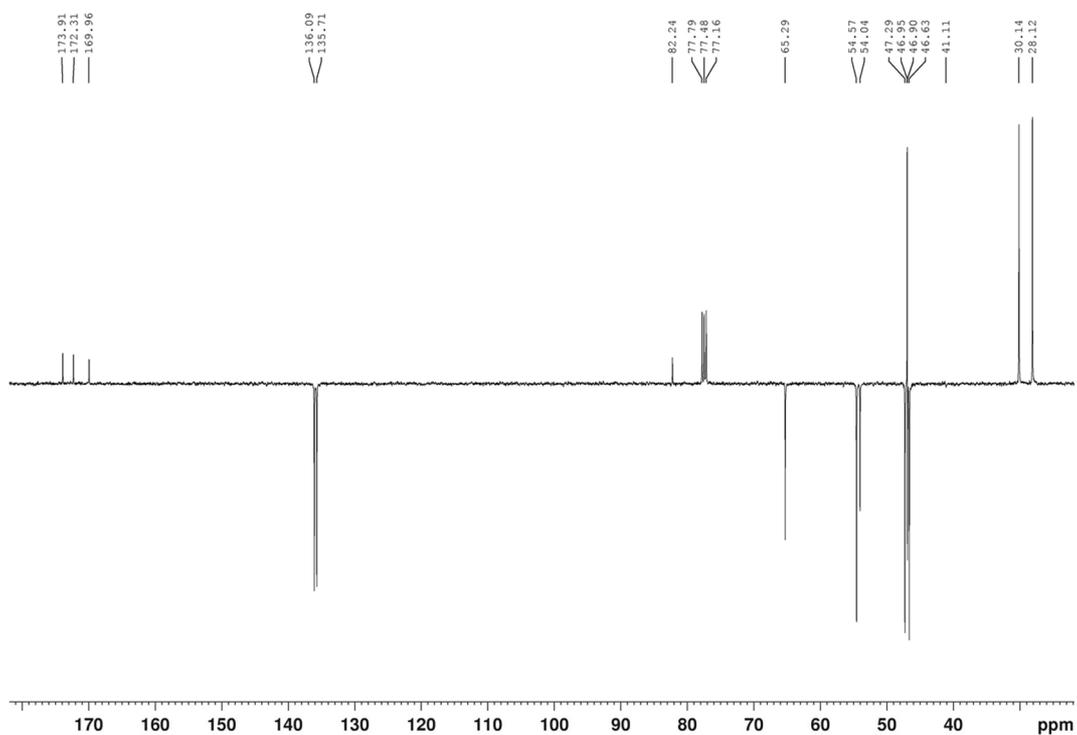
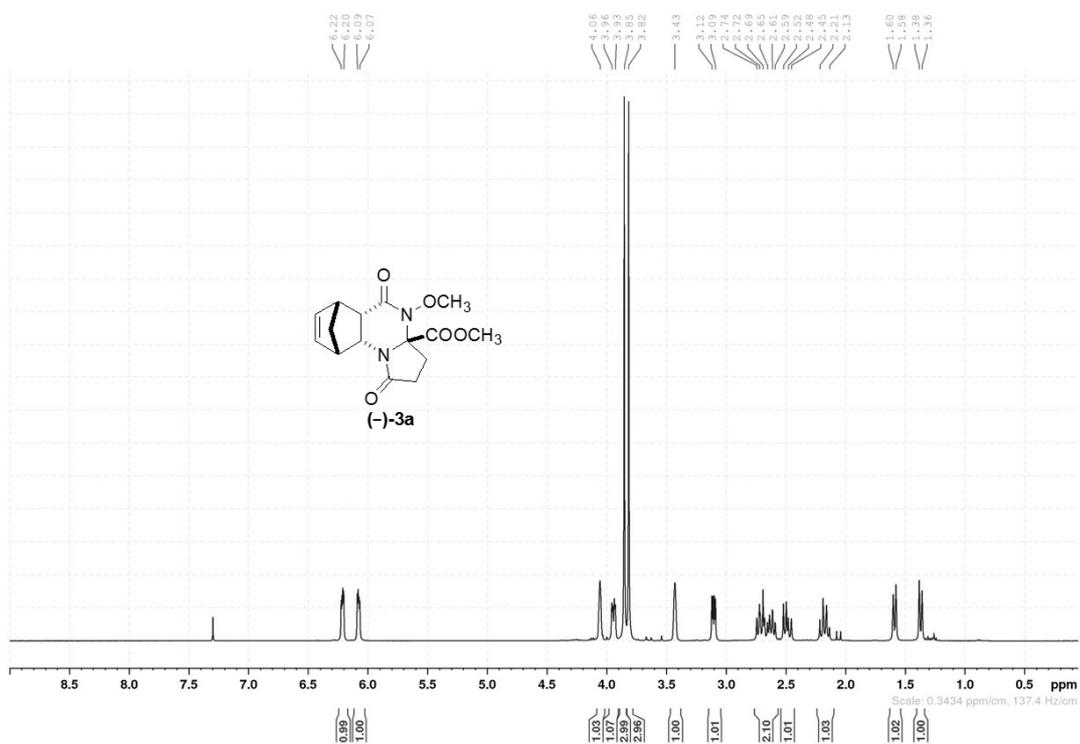
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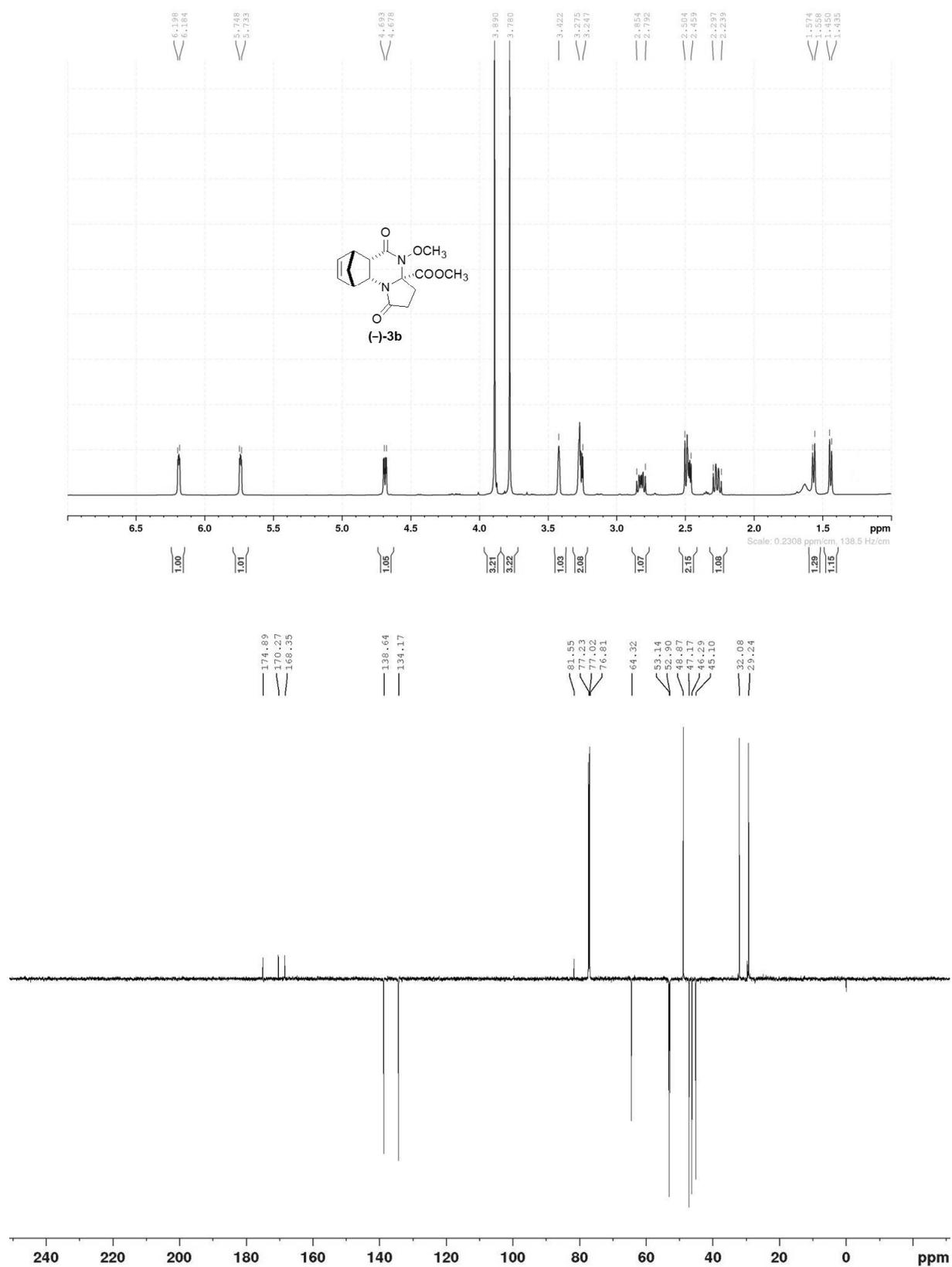
**Contents**

<sup>1</sup> H-NMR and <sup>13</sup> C-NMR spectra.....	2
HPLC chromatograms.....	10
Table for X-ray crystallography data for (±)- <b>3a</b> , (±)- <b>4a</b> and (±)- <b>6a</b> ..	12

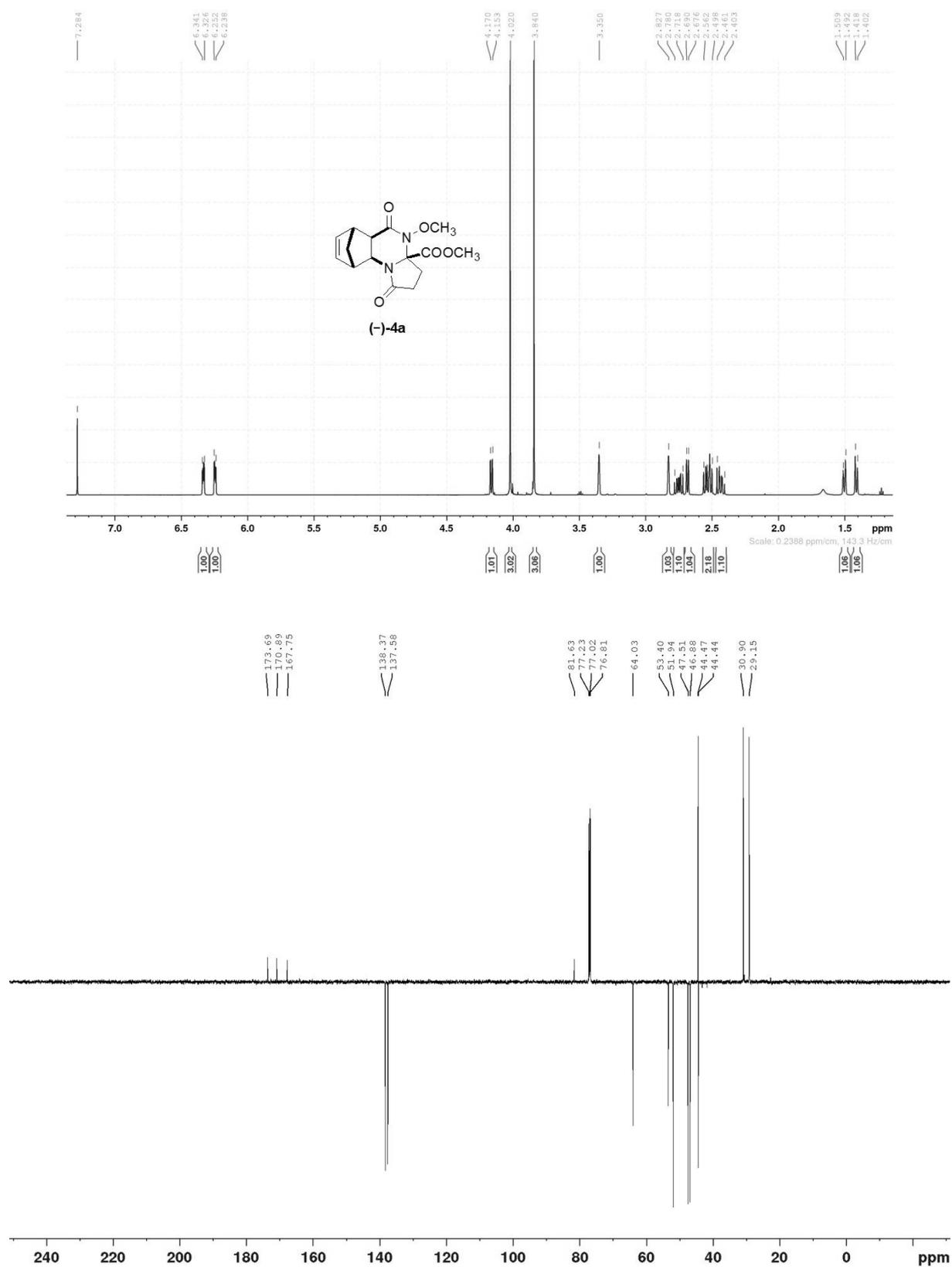
**(3a*S*,5a*S*,6*R*,9*S*,9a*R*)-Methyl 4-methoxy-1,5-dioxo-1,2,3,3a,4,5,5a,6,9a-decahydro-6,9-methanopyrrolo[1,2-*a*]quinazoline-3a-carboxylate [(-)-3a]:**



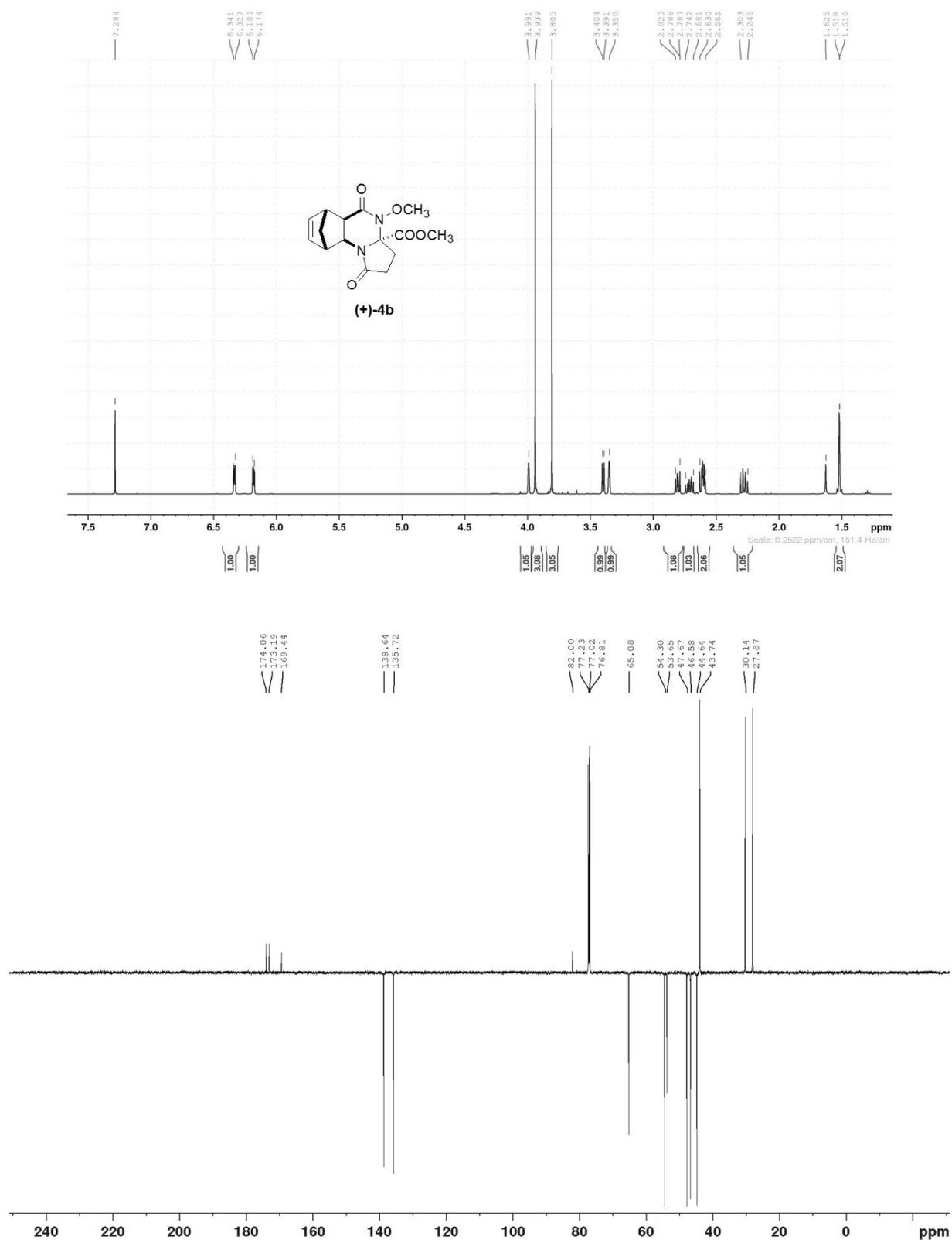
**(3a*R*,5a*S*,6*R*,9*S*,9a*R*)-Methyl 4-methoxy-1,5-dioxo-1,2,3,3a,4,5a,6,9a-decahydro-6,9-methanopyrrolo[1,2-*a*]quinazoline-3a-carboxylate [(-)-3b]**



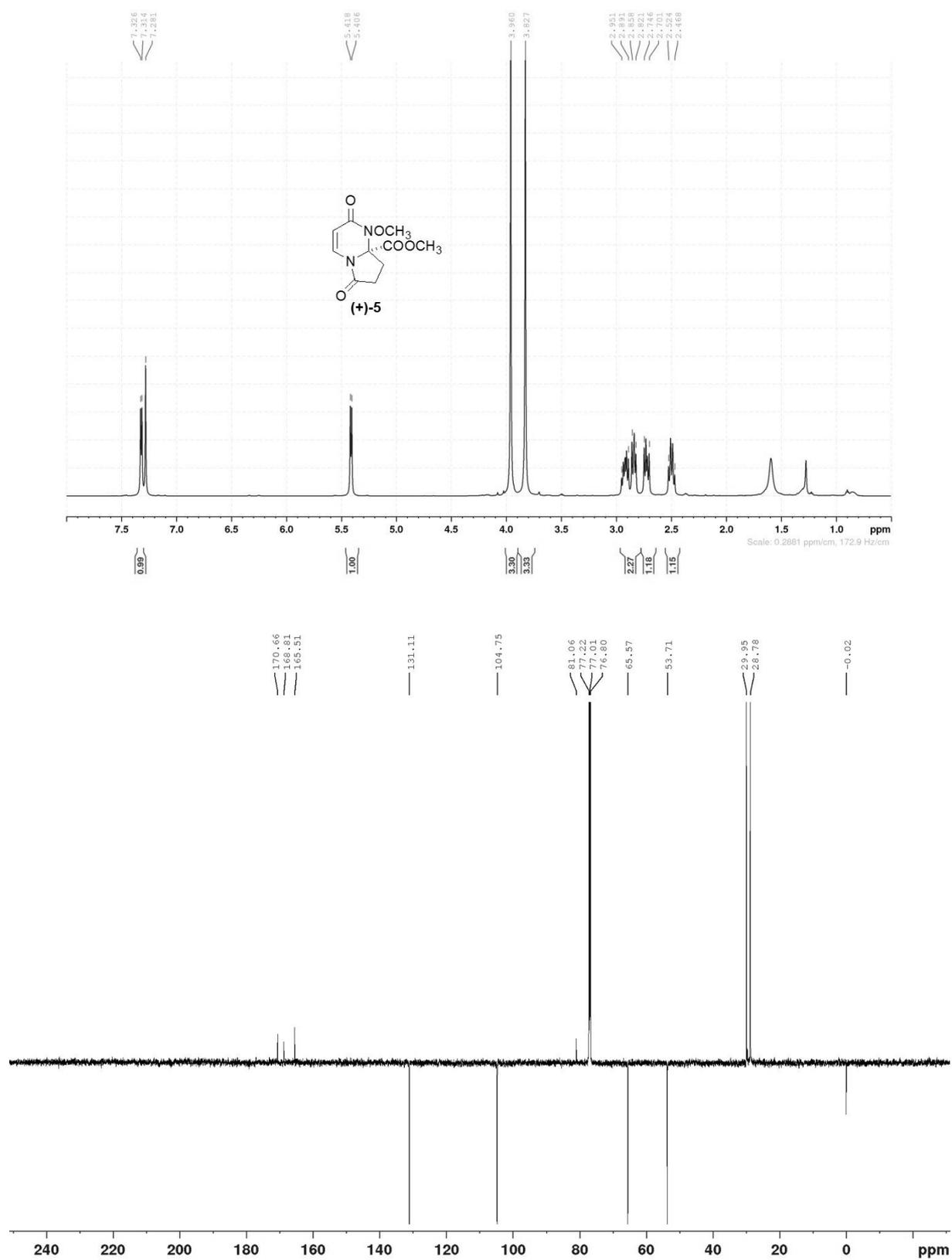
**(3a*S*,5a*R*,6*R*,9*S*,9a*S*)-Methyl 4-methoxy-1,5-dioxo-1,2,3,3a,4,5a,6,9a-decahydro-6,9-methanopyrrolo[1,2-*a*]quinazoline-3a-carboxylate [(-)-4a]:**



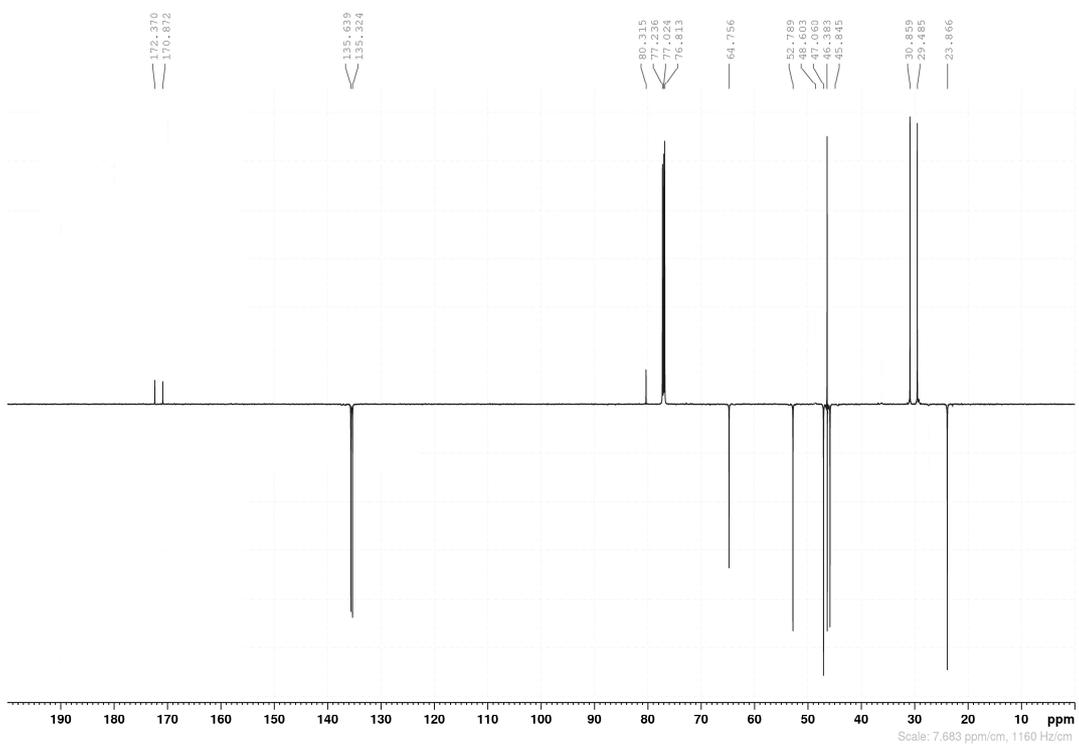
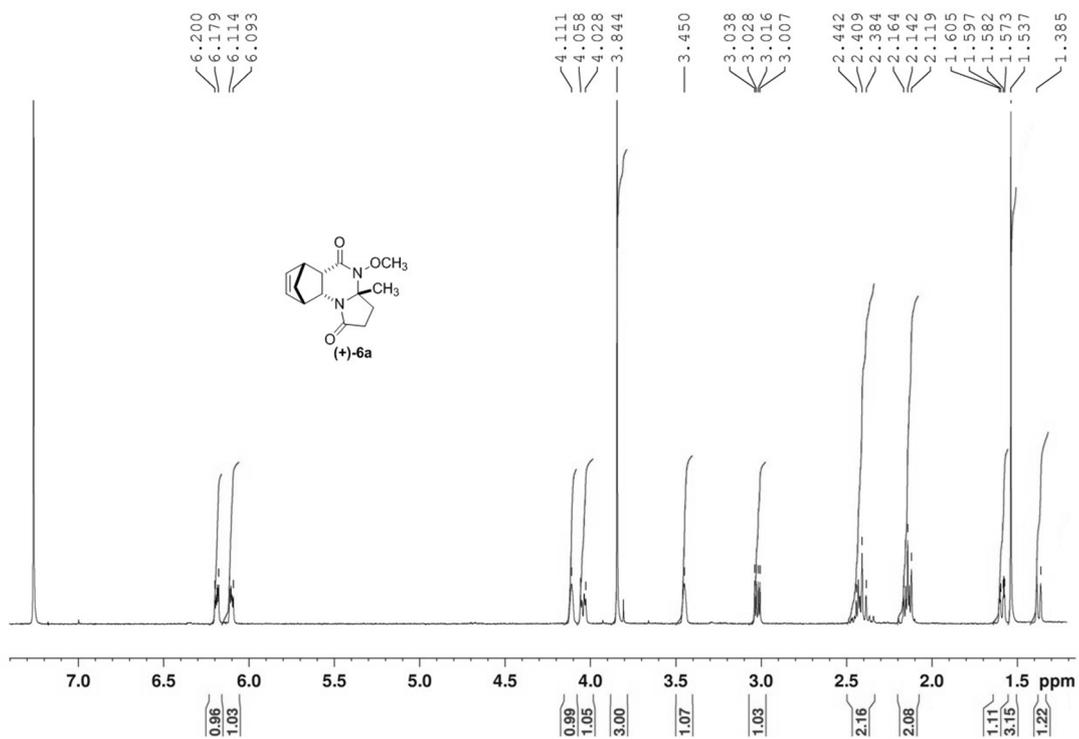
**(3a*R*,5a*R*,6*R*,9*S*,9a*S*)-Methyl 4-methoxy-1,5-dioxo-1,2,3,3a,4,5a,6,9a-decahydro-6,9-methanopyrrolo[1,2-*a*]quinazoline-3a-carboxylate [(+)-4b]:**



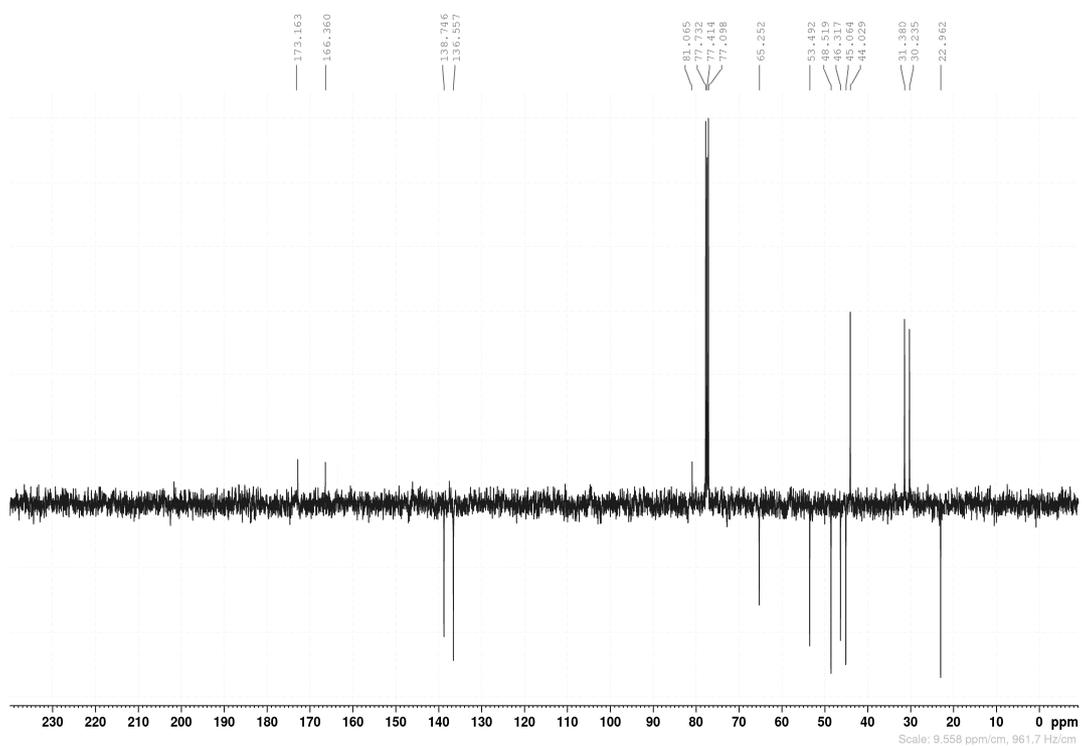
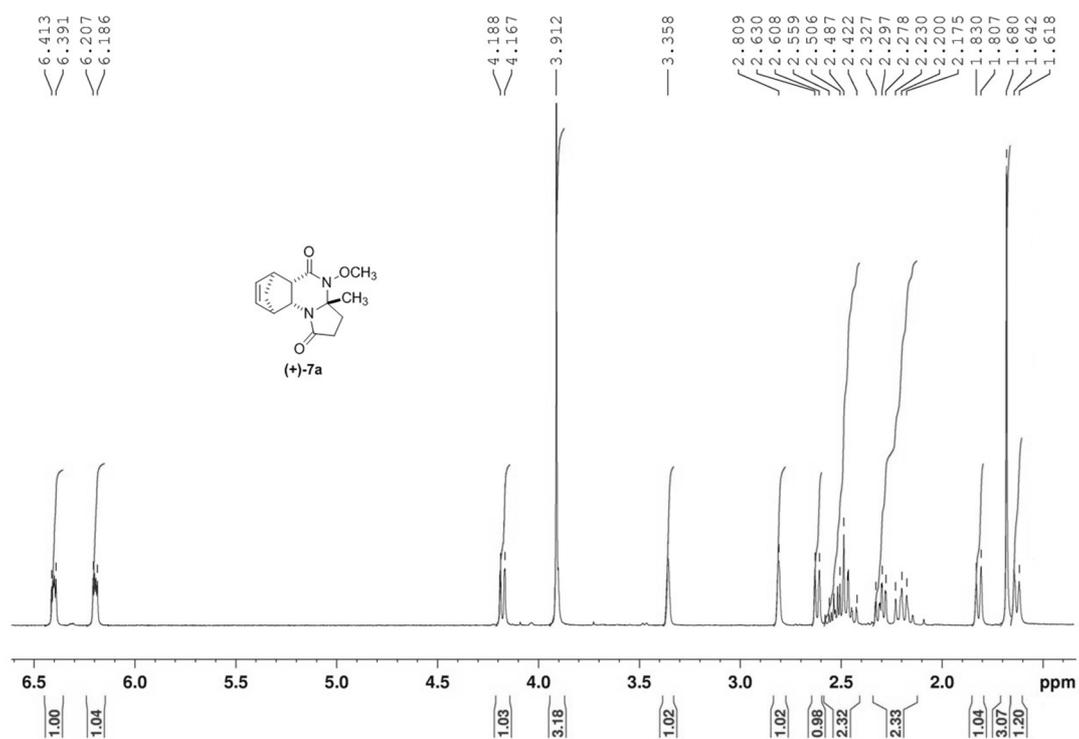
**(R)-methyl 1-methoxy-2,6-dioxo-1,2,6,7,8,8a-hexahydropyrrolo[1,2-a]pyrimidine-8a-carboxylate [(+)-5]:**



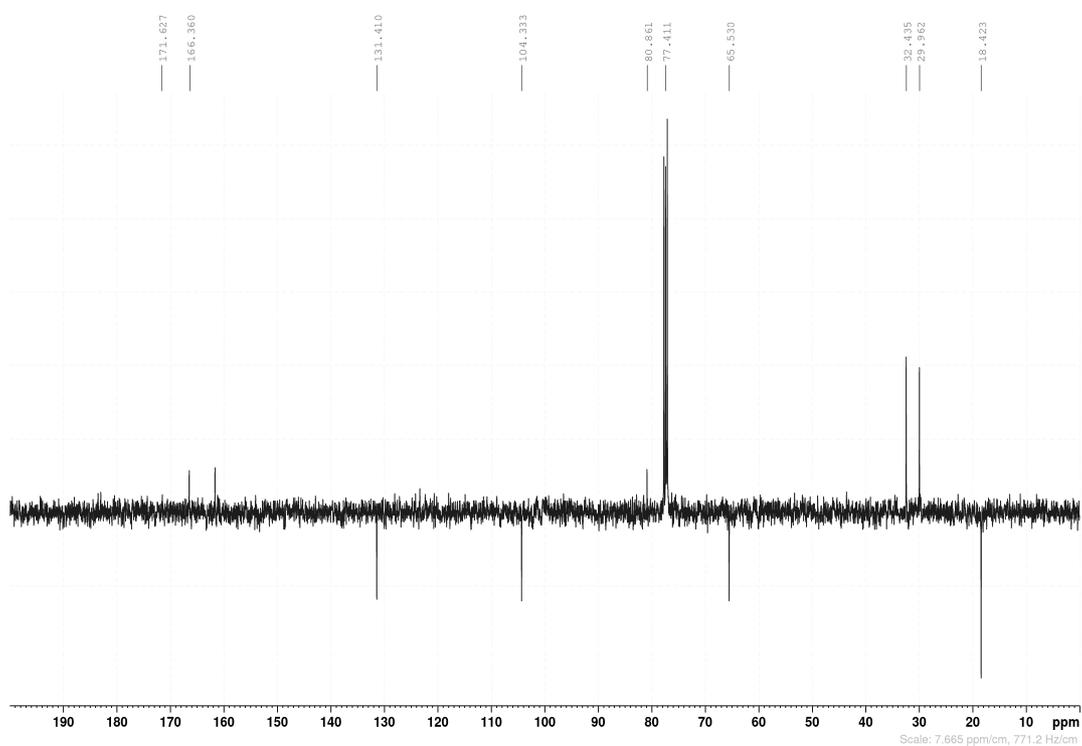
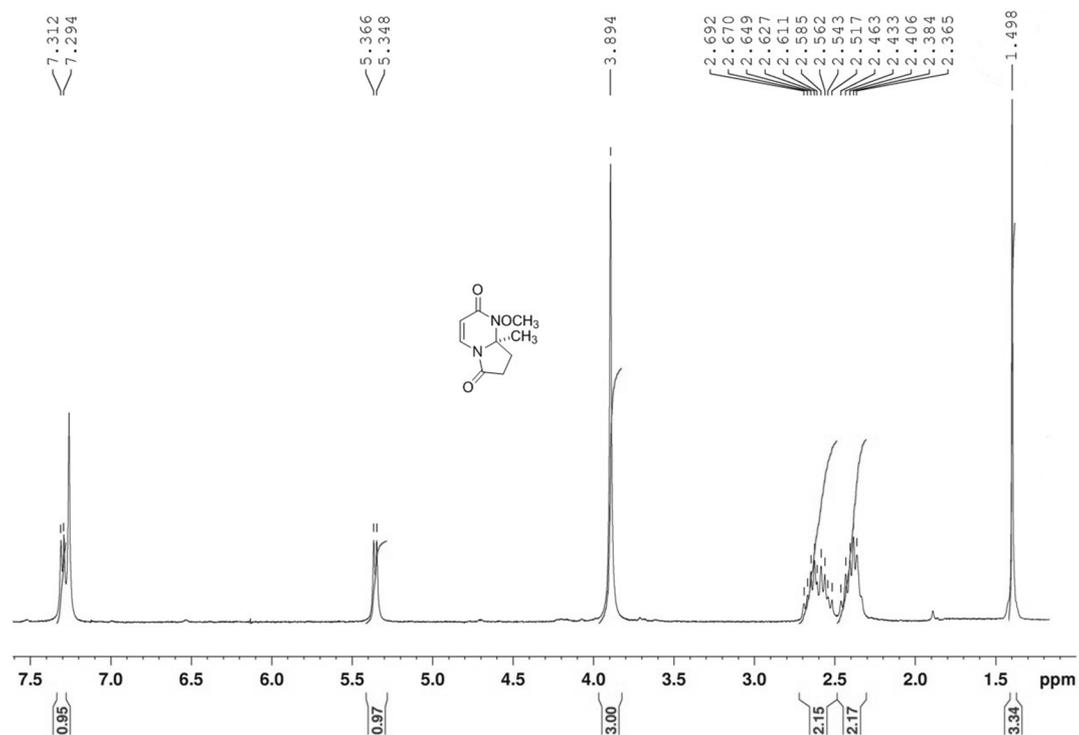
**(3*R*,5*aS*,6*R*,9*S*,9*aR*)-4-methoxy-3*a*-methyl-2,3,3*a*,4,5*a*,6,9*a*-octahydro-6,9methanopyrrolo[1,2-*a*]quinazoline-1,5-dione [(+)-6*a*]:**



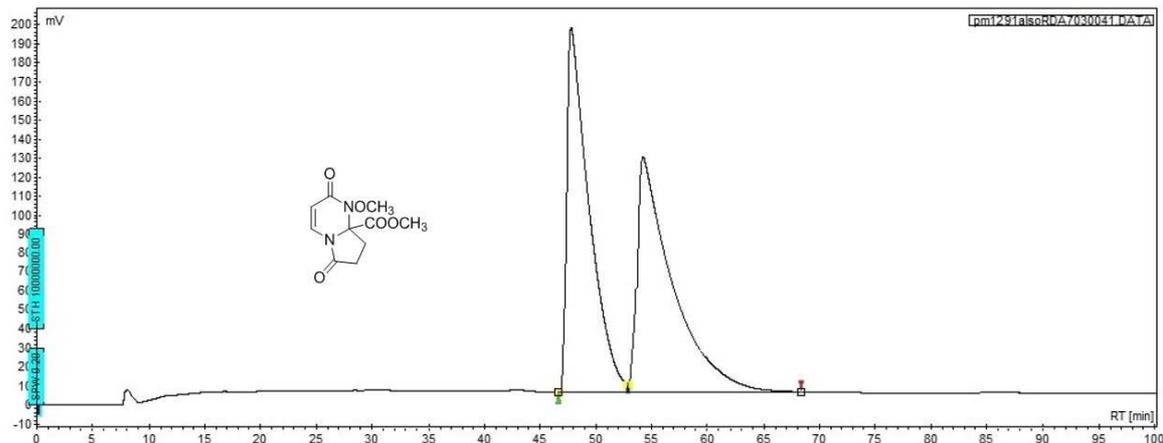
**(3*R*,5*S*,6*S*,9*R*,9*aR*)-4-methoxy-3*a*-methyl-2,3,3*a*,4,5*a*,6,9*a*-octahydro-6,9-methanopyrrolo[1,2-*a*]quinazoline-1,5-dione [(+)-7*a*]:**



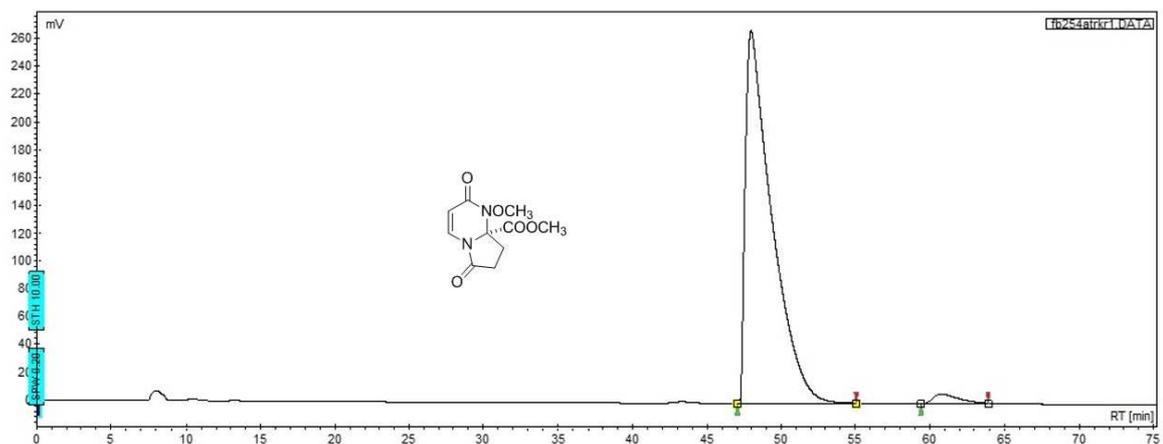
**(S)-1-methoxy-8a-methyl-1,7,8,8a-tetrahydropyrrolo[1,2-a]pyrimidine-2,6-dione [(+)-8]:**



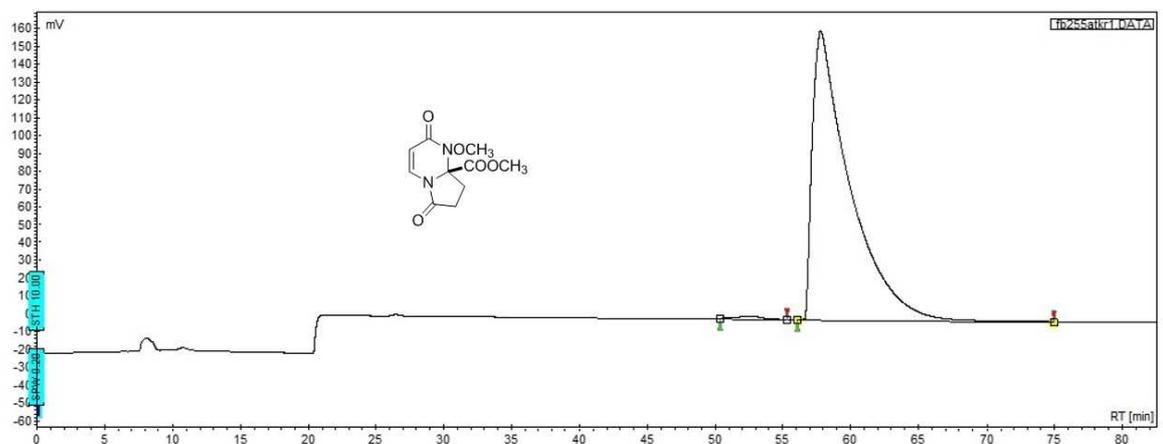
**(RS)-methyl 1-methoxy-2,6-dioxo-1,2,6,7,8,8a-hexahydropyrrolo[1,2-a]pyrimidine-8a-carboxylate [(±)-5]:**



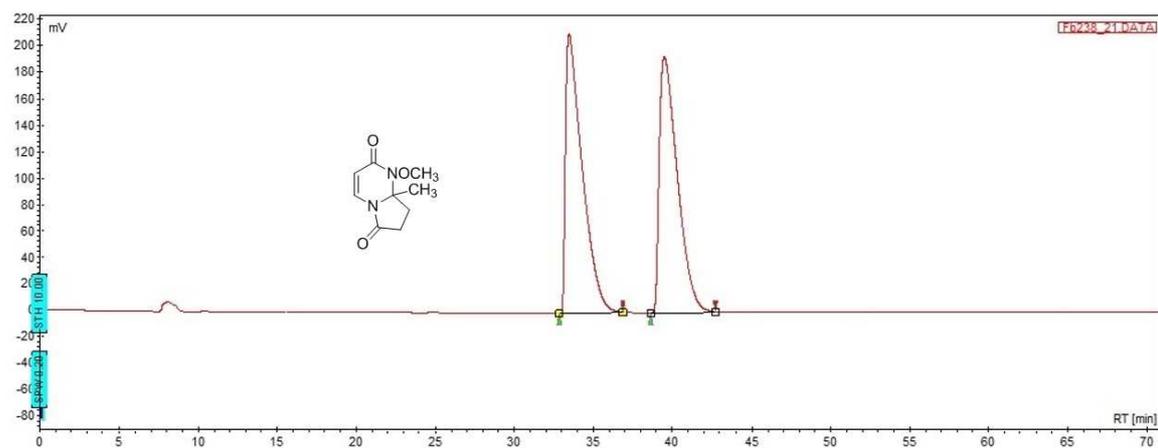
**(R)-methyl 1-methoxy-2,6-dioxo-1,2,6,7,8,8a-hexahydropyrrolo[1,2-a]pyrimidine-8a-carboxylate [(+)-5]:**



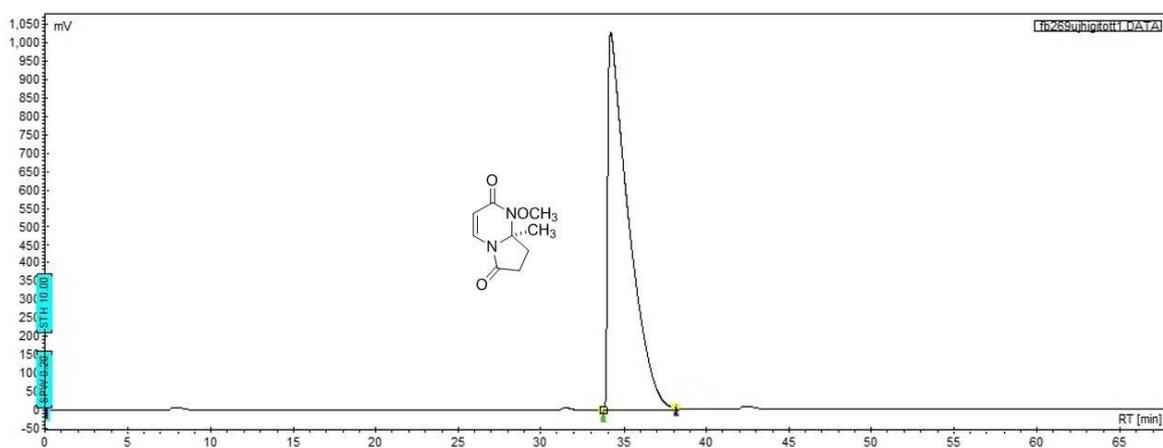
**(S)-methyl 1-methoxy-2,6-dioxo-1,2,6,7,8,8a-hexahydropyrrolo[1,2-a]pyrimidine-8a-carboxylate [(-)-5]:**



**(RS)-1-methoxy-8a-methyl-1,7,8,8a-tetrahydropyrrolo[1,2-a]pyrimidine-2,6-dione [(±)-8]:**



**(S)-1-methoxy-8a-methyl-1,7,8,8a-tetrahydropyrrolo[1,2-a]pyrimidine-2,6-dione [(+)-8]:**



**(R)-1-methoxy-8a-methyl-1,7,8,8a-tetrahydropyrrolo[1,2-a]pyrimidine-2,6-dione [(-)-8]:**

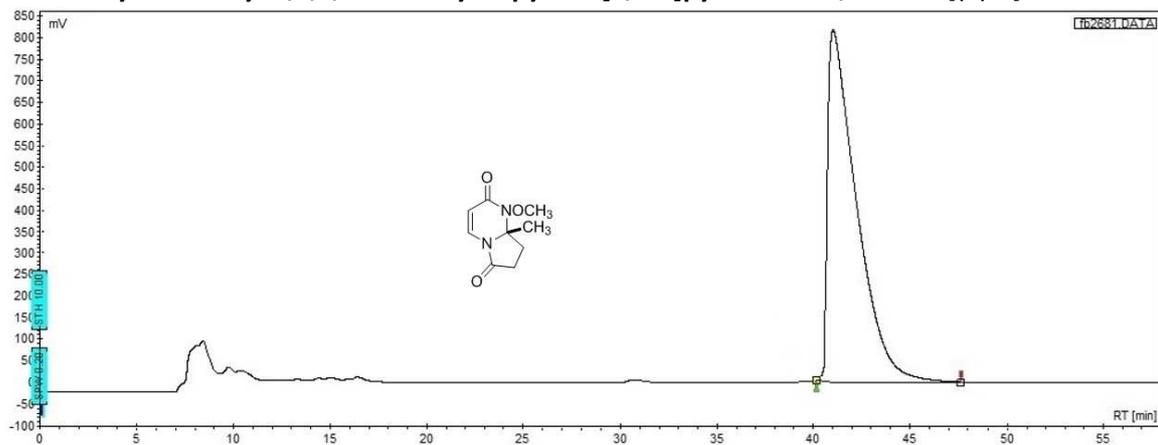


Table for X-ray crystallography data for (±)-**3a**, (±)-**4b** and (±)-**6a**

	(±)- <b>3a</b>	(±)- <b>4b</b>	(±)- <b>6a</b>
Empirical formula	C <sub>15</sub> H <sub>18</sub> N <sub>2</sub> O <sub>5</sub>	C <sub>15</sub> H <sub>18</sub> N <sub>2</sub> O <sub>5</sub>	C <sub>14</sub> H <sub>18</sub> N <sub>2</sub> O <sub>3</sub>
Formula weight	306.31	306.31	262.30
Temperature	120(2) K	120(2) K	120(2) K
Wavelength	1.54184 Å	1.54184 Å	0.71073 Å
Crystal system	Triclinic	Monoclinic	Monoclinic
Space group	P $\bar{1}$	P2 <sub>1</sub> /c	P2 <sub>1</sub> /n
Unit cell dimensions	a = 8.3315(3) Å α = 81.778(3)	a = 16.7015(3) Å α = 90	a = 9.4473(3) Å α = 90°
	b = 8.7788(3) Å β = 82.075(3)	b = 9.4689(3) Å β = 98.6847(17)°	b = 10.0314(2) Å β = 105.075(3)°
	c = 9.8739(3) Å γ = 83.141(2)	c = 18.1845(4) Å γ = 90	c = 13.7952(4) Å γ = 90°
Volume	704.26(4) Å <sup>3</sup>	2842.82(11) Å <sup>3</sup>	1262.36(6) Å <sup>3</sup>
Z	2	8	4
Density (calculated)	1.444 Mg/m <sup>3</sup>	1.431 Mg/m <sup>3</sup>	1.380 Mg/m <sup>3</sup>
Absorption coefficient	0.916 mm <sup>-1</sup>	0.908 mm <sup>-1</sup>	0.098 mm <sup>-1</sup>
F(000)	324	1296	560
Crystal size	0.191 x 0.152 x 0.081 mm <sup>3</sup>	0.702 x 0.498 x 0.299 mm <sup>3</sup>	0.418 x 0.391 x 0.243 mm <sup>3</sup>
Theta range for data collection	4.559 to 76.864°.	4.920 to 76.944°.	3.110 to 29.575°.
Index ranges	-10 ≤ h ≤ 10, -11 ≤ k ≤ 11, -12 ≤ l ≤ 12	-21 ≤ h ≤ 21, -11 ≤ k ≤ 11, -22 ≤ l ≤ 22	-12 ≤ h ≤ 12, -13 ≤ k ≤ 13, -19 ≤ l ≤ 18
Reflections collected	15453	44312	20374
Independent reflections	2963 [R(int) = 0.0573]	5930 [R(int) = 0.0496]	3501 [R(int) = 0.0432]
Completeness to theta	67.684° 99.7 %	67.684° 98.9 %	26.000° 99.8 %

Refinement method	Full-matrix least-squares on F <sup>2</sup>	Full-matrix least-squares on F <sup>2</sup>	Full-matrix least-squares on F <sup>2</sup>
Data / restraints / parameters	2963 / 0 / 201	5930 / 0 / 401	3501 / 0 / 174
Goodness-of-fit on F <sup>2</sup>	1.070	1.072	1.046
Final R indices [I>2sigma(I)]	R1 = 0.0432, wR2 = 0.1170	R1 = 0.0485, wR2 = 0.1283	R1 = 0.0434, wR2 = 0.1110
R indices (all data)	R1 = 0.0439, wR2 = 0.1177	R1 = 0.0488, wR2 = 0.1286	R1 = 0.0517, wR2 = 0.1181
Extinction coefficient	n/a	n/a	n/a
Largest diff. peak and hole	0.350 and -0.280 e.Å <sup>-3</sup>	0.335 and -0.275 e.Å <sup>-3</sup>	0.286 and -0.225 e.Å <sup>-3</sup>