

Supplementary Materials

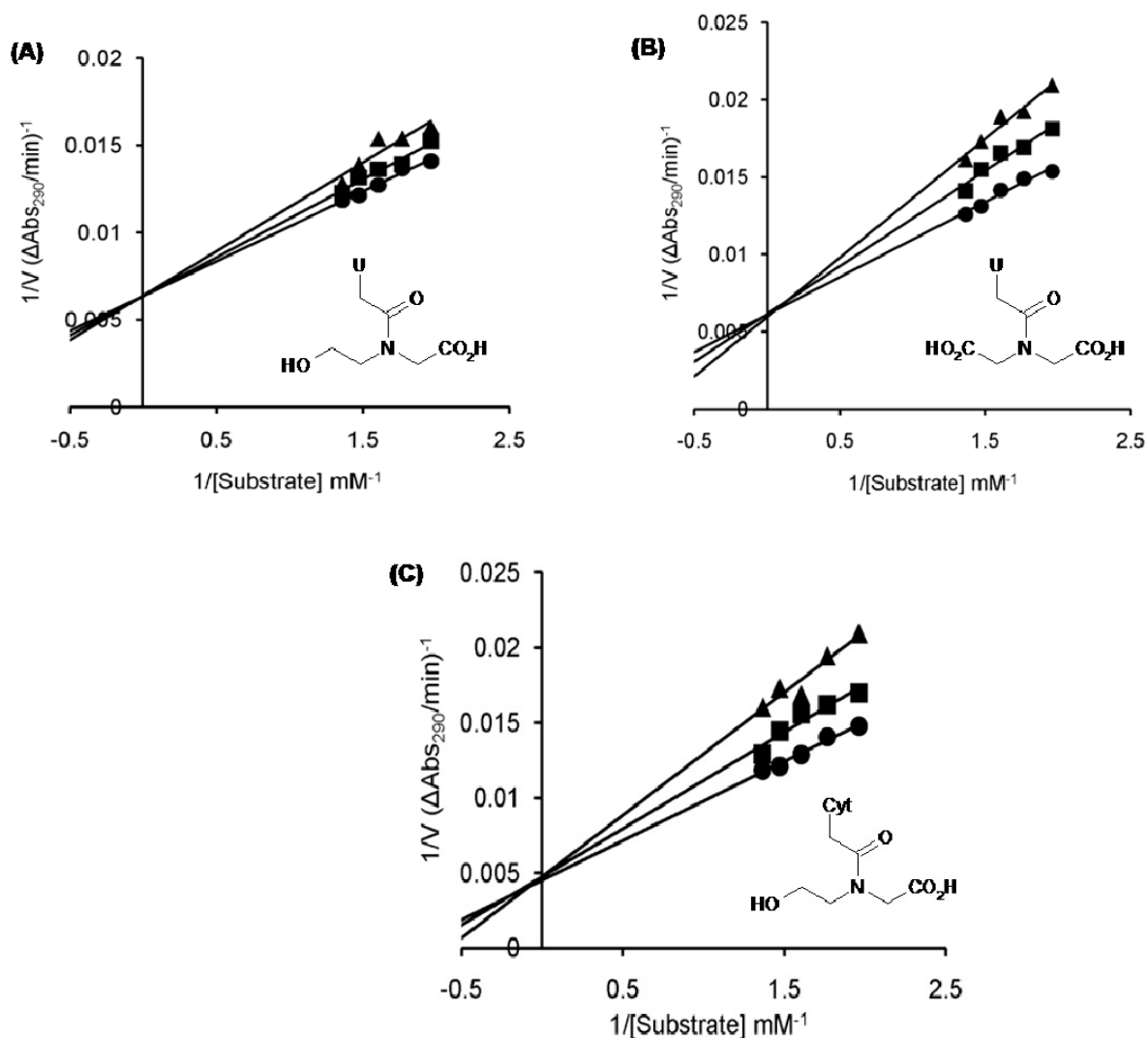


Figure S1. Lineweaver–Burk plot for inhibition of RNase A by (A) U-ol-acid (8) of 0.12 (▲), 0.04 (■), or 0 (●) mM, with 2',3'-cCMP concentrations of 0.73–0.51 mM and RNase A concentration of 9.8 μM . (B) U-di-acid (10) of 0.165 (▲), 0.055 (■), or 0 (●) mM, with 2',3'-cCMP concentrations of 0.73–0.51 mM and RNase A concentration of 10.2 μM . (C) C-ol-acid (18) of 0.20 (▲), 0.10 (■), or 0 (●) mM, with 2',3'-cCMP concentrations of 0.72–0.50 mM and RNase A concentration of 9.9 μM .

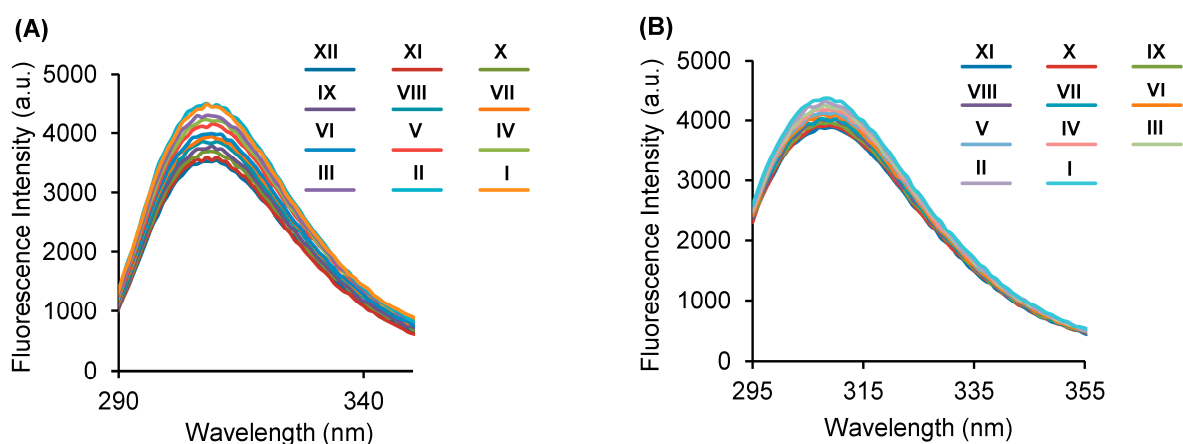


Figure S2. Fluorescence quenching spectra of RNase A in presence of (A) C-di-acid (21) and (B) U-di-acid (10). The increasing order of concentrations of the inhibitors (in μM) for each curve is as follows: (A) I = 0, II = 0.26, III = 0.39, IV = 0.53, V = 0.65, VI = 0.78, VII = 0.91, VIII = 1.04, IX = 1.16, X = 1.29, XI = 1.41, XII = 1.54; (B) I = 0, II = 0.26, III = 0.39, IV = 0.53, V = 0.65, VI = 0.78, VII = 0.91, VIII = 1.04, IX = 1.16, X = 1.29, XI = 1.41.

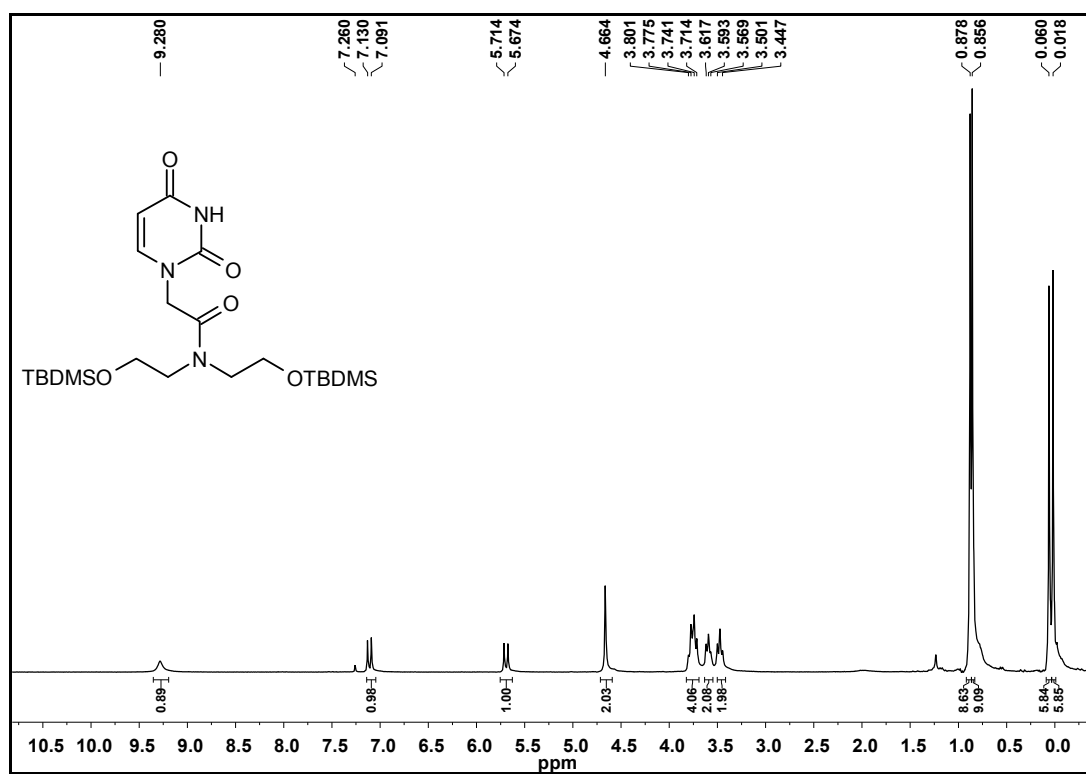
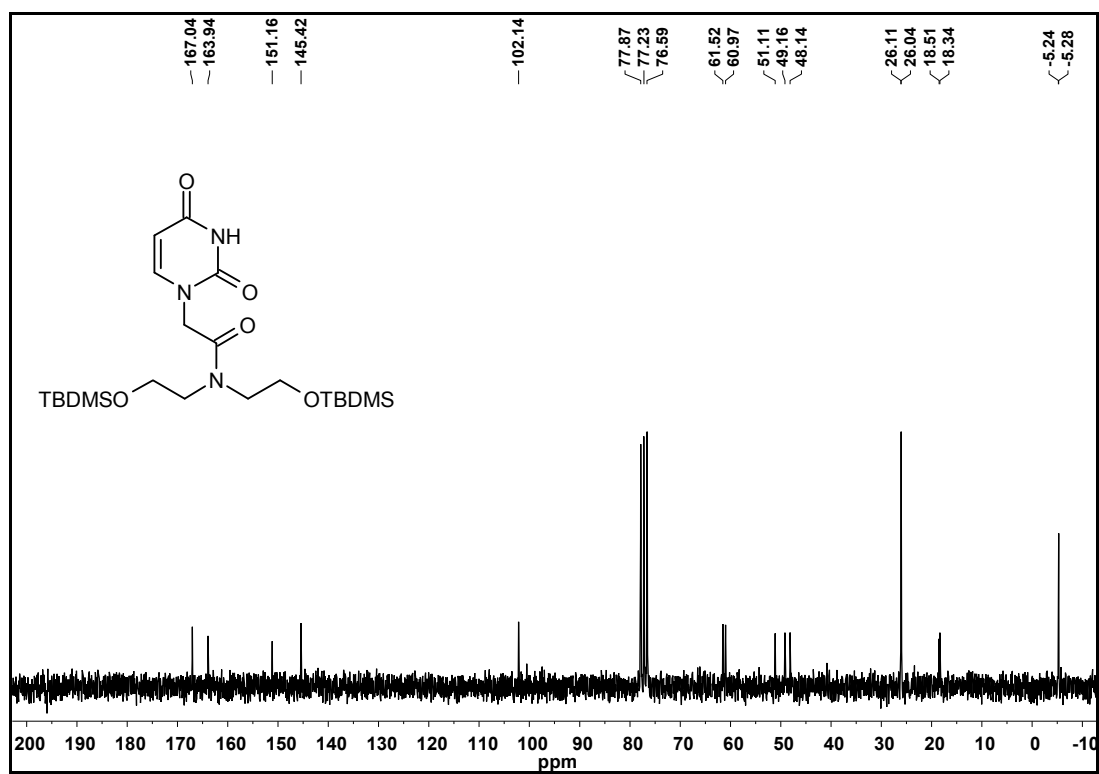
Table S1. Binding constants (K_b) and the number of binding sites (n) for the interactions of U-di-acid (10) and C-di-acid (21) with RNase A.

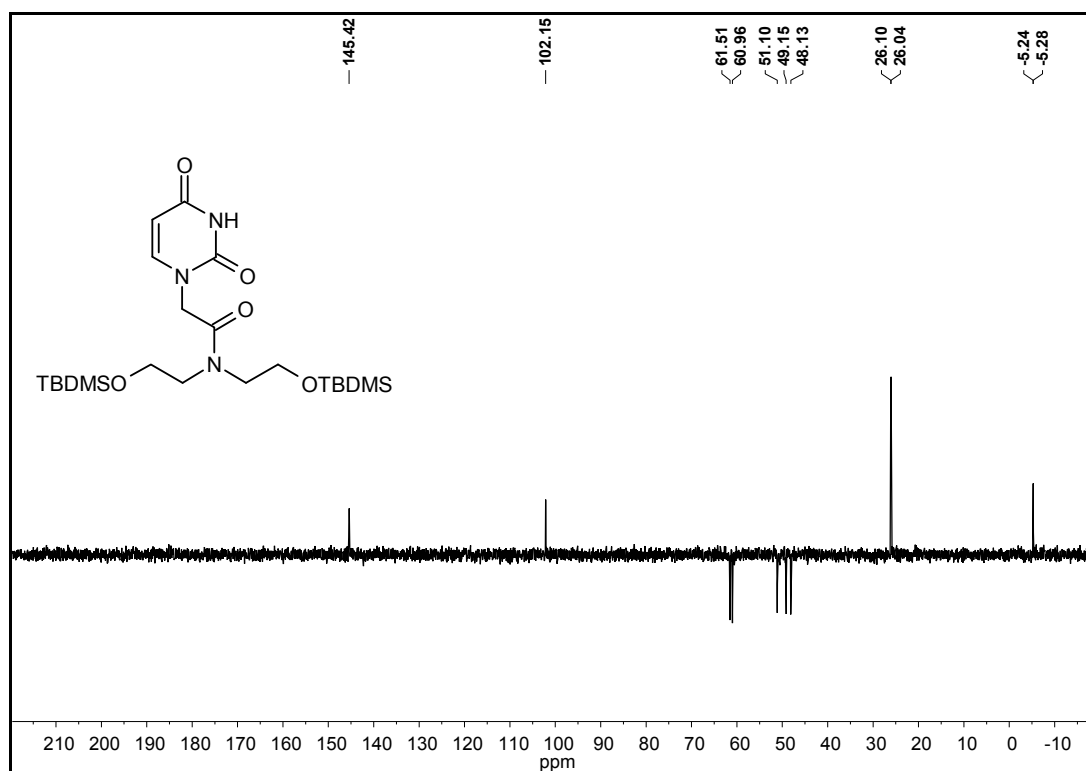
Inhibitor	K_b (M^{-1})	No. of Binding Sites (n)
U-di-acid (10)	$7.64 \pm 0.1 \times 10^4$	0.959
C-di-acid (21)	$1.44 \pm 0.02 \times 10^5$	1.257

Table S2. Distances of polar contacts between inhibitors and amino acid residues of RNase A (1FS3).

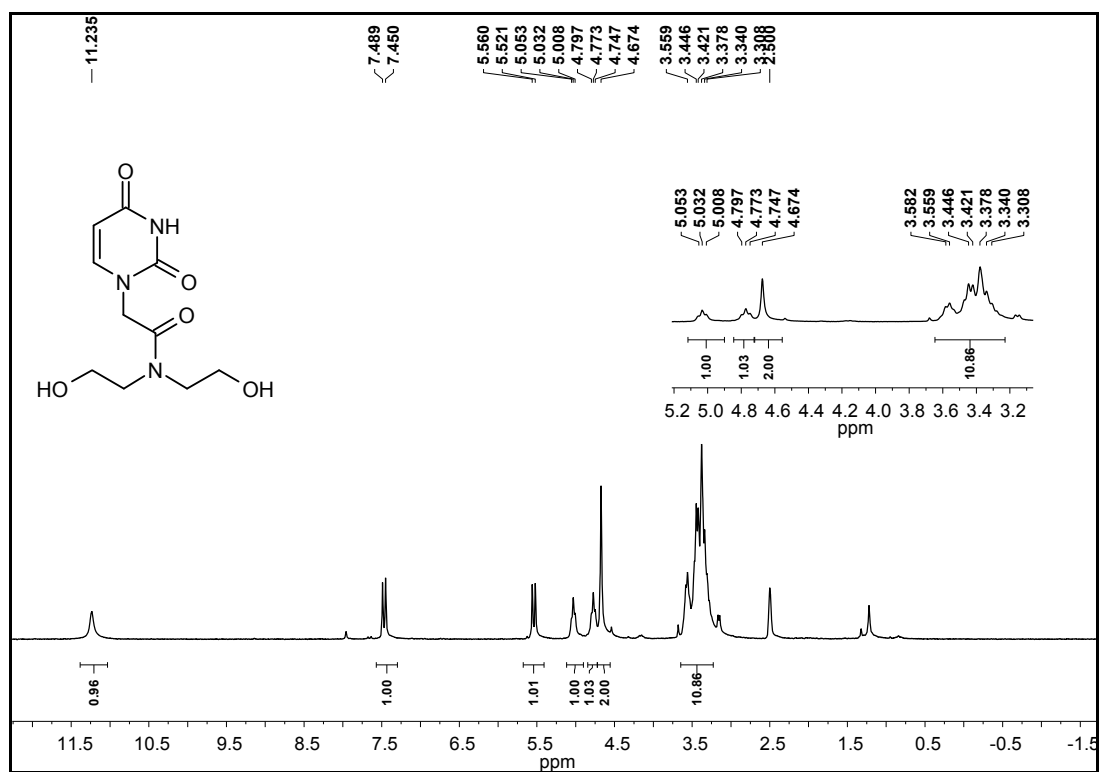
1FS3	U-ol-Acid (8)	C-ol-Acid (18)	U-di-Acid (10)	C-di-Acid (21)
Arg39 NH2	2.72 Å [O of C4 C=O of nucleobase]			3.2 Å [N3 of nucleobase]
Arg39 NH1	2.91 Å [O of C4 C=O of nucleobase]			3.3 Å [O of C2 C=O of nucleobase]
Lys41 CA				3.0 Å [N3 of nucleobase]
Lys41 N ζ		2.4 Å [N3 of nucleobase]	3.1 Å [C=O of amide]	
Lys7 N ζ	2.17 Å [OH of CH ₂ OH]	2.6 Å [OH of acid]		2.8 Å [C=O of amide]
His12 Ne2	3.21 Å [C=O of acid]	3.0 Å [C=O of amide] 2.1 Å [OH of CH ₂ OH]	2.6 Å [C=O of acid]	3.1 Å [C=O of acid]
Gln11 Ne2	2.58 Å [C=O of amide]	2.8 Å [O of C2 C=O of nucleobase] 2.9 Å [OH of acid] 2.2 Å [OH of CH ₂ OH]	2.9 Å [C=O of acid]	2.6 Å [C=O of acid]
Gln11 O ϵ 1			2.2 Å [OH of acid]	
His119 N δ 1	2.57 Å [OH of acid]		1.9 Å [OH of acid]	1.9 Å [OH of acid]
Phe120 CA				3.2 Å [C=O of acid]
Phe120 N	2.83 Å [C=O of acid]	3.2 Å [C=O of amide]	2.8 Å [C=O of acid]	
Val43 O		2.0 Å [NH ₂ of nucleobase]	1.9 Å [N3 of nucleobase]	

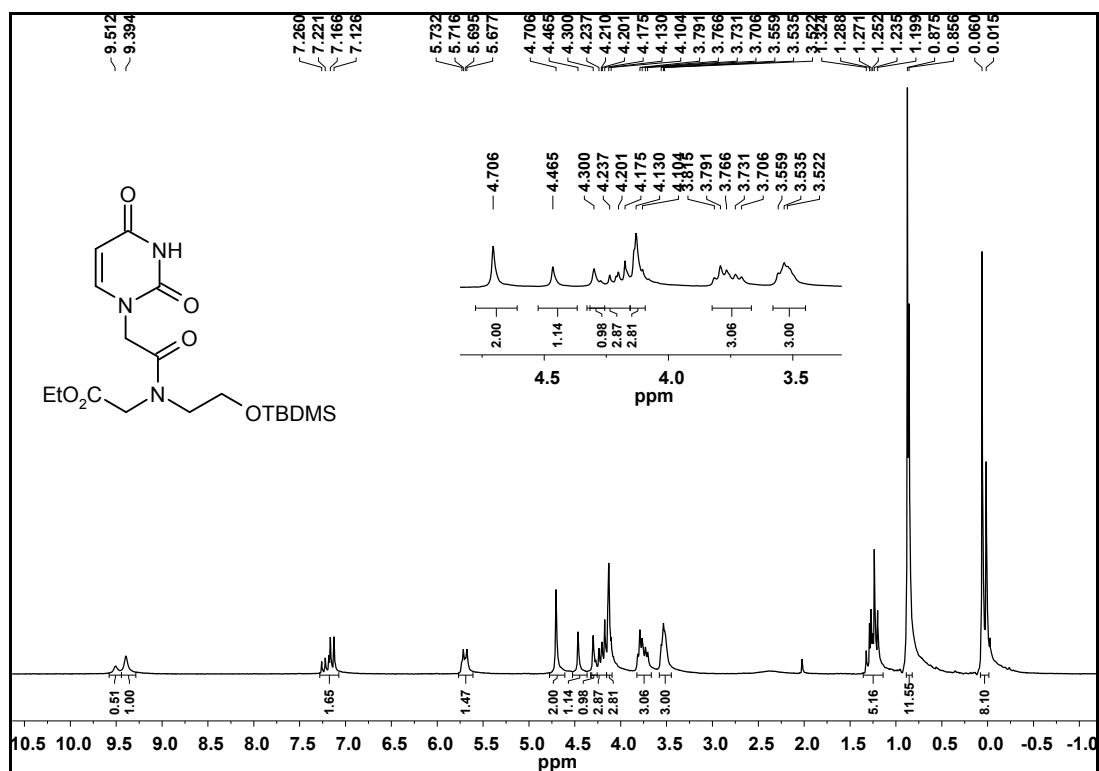
NMR Spectra

¹H-NMR of compound 4.¹³C-NMR of compound 4.

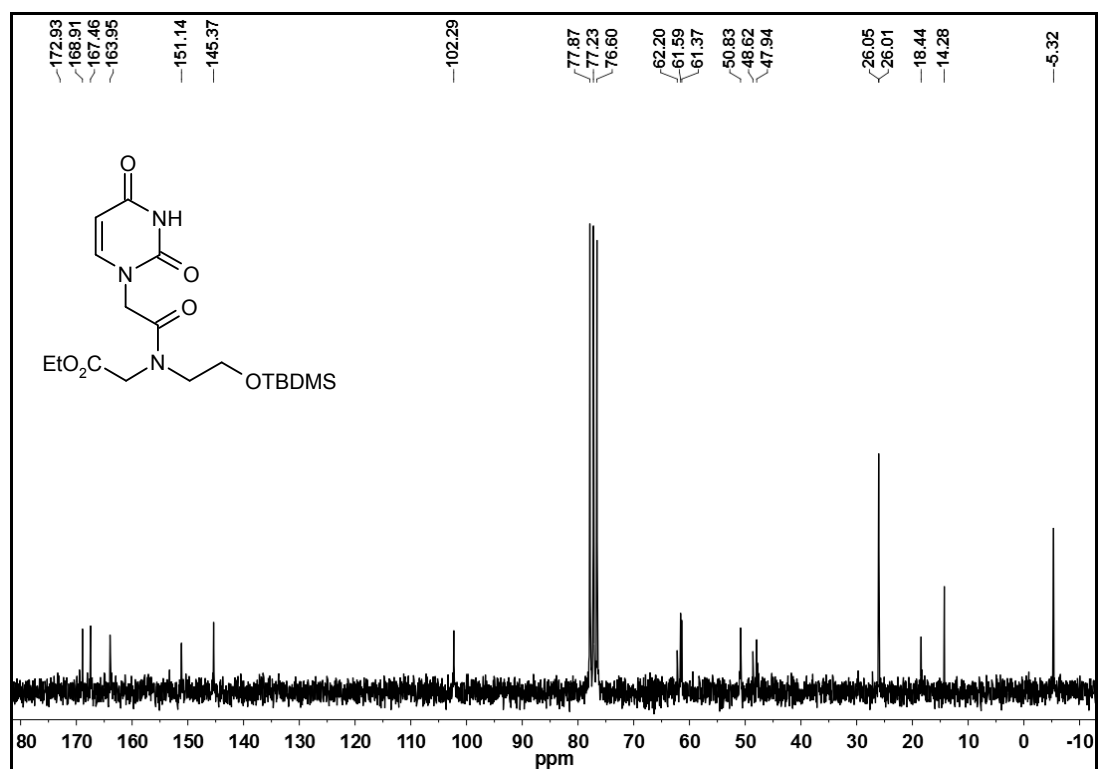


DEPT 135 NMR of compound 4.

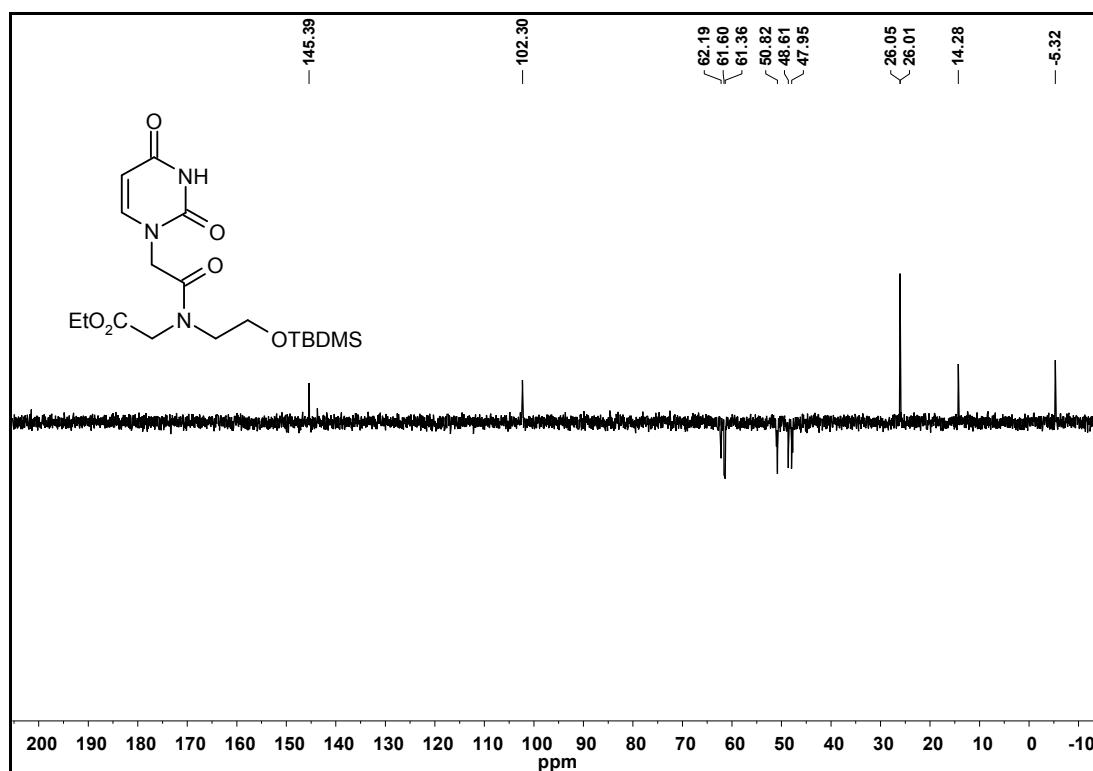
¹H-NMR of compound 5.



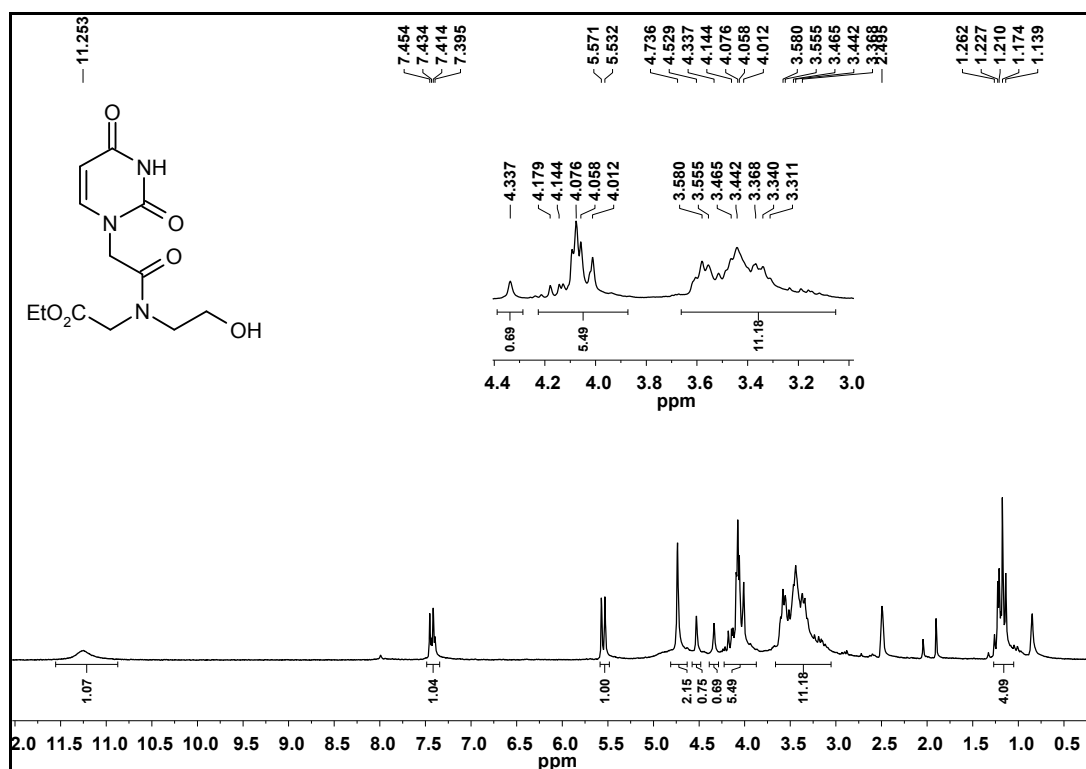
¹H-NMR of compound 6.

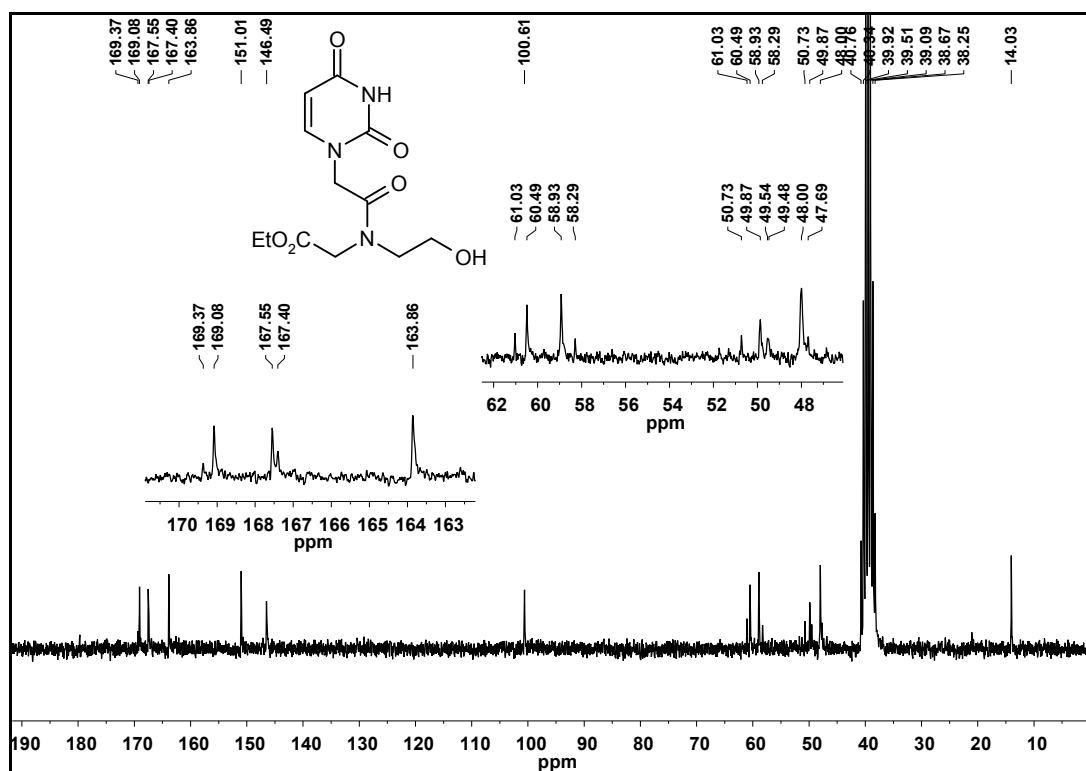


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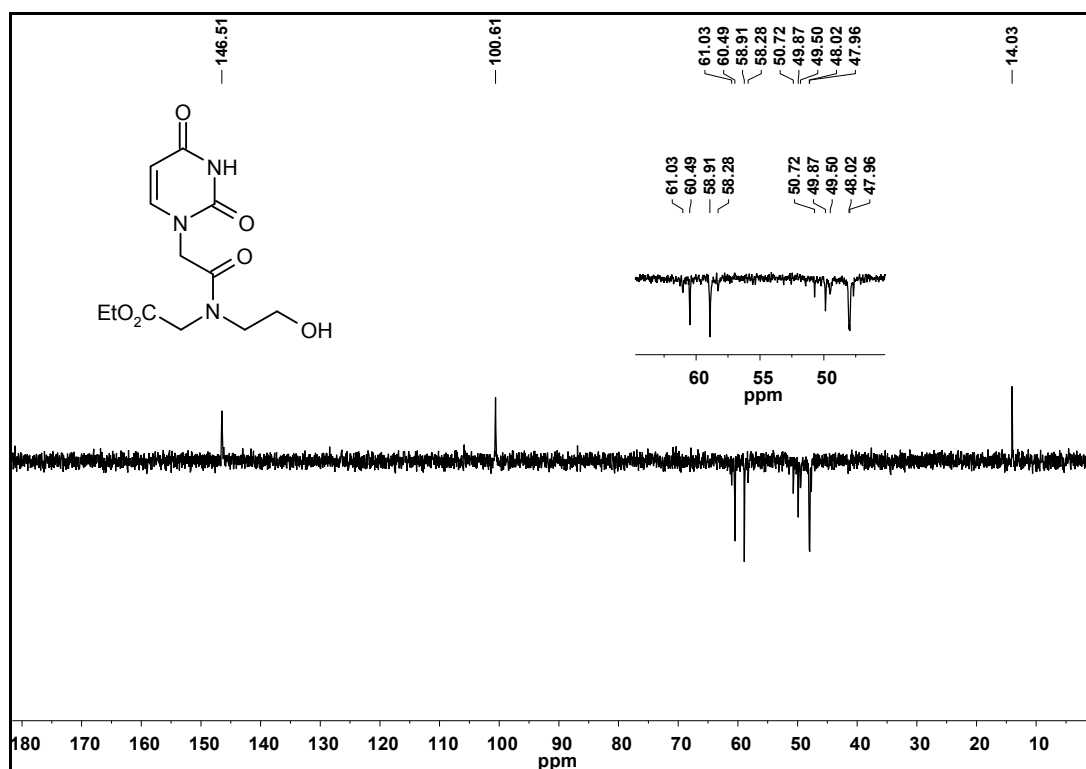


DEPT 135 NMR of compound 6.

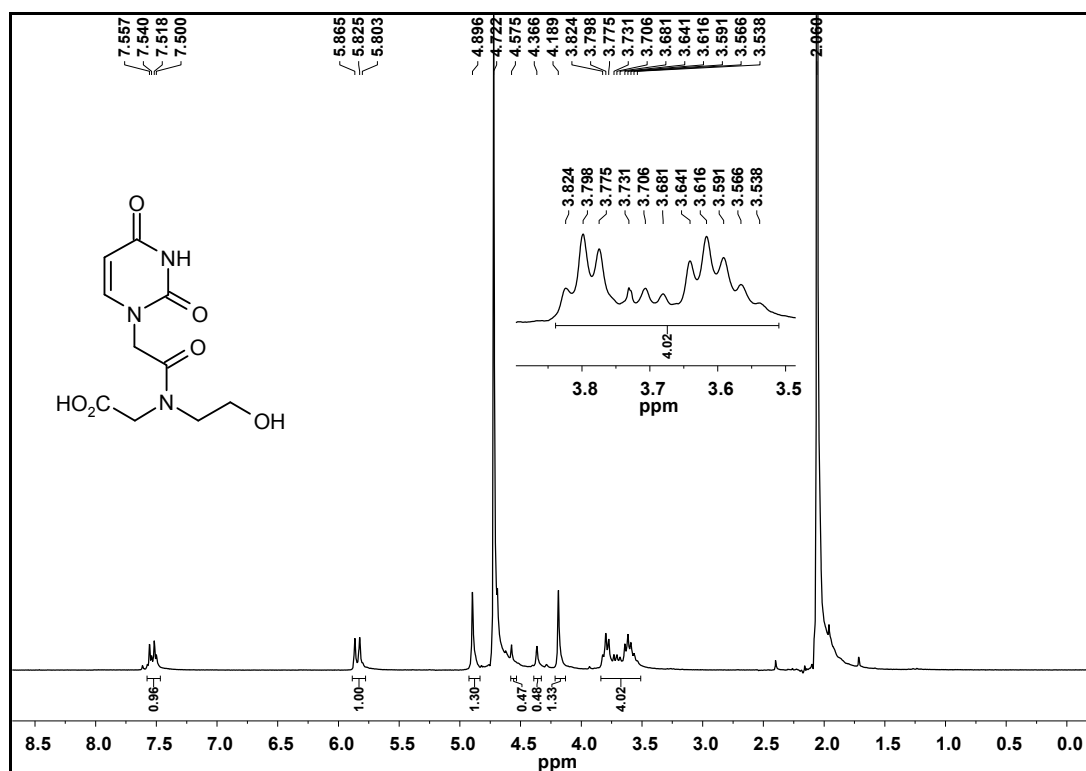
¹H-NMR of compound 7.



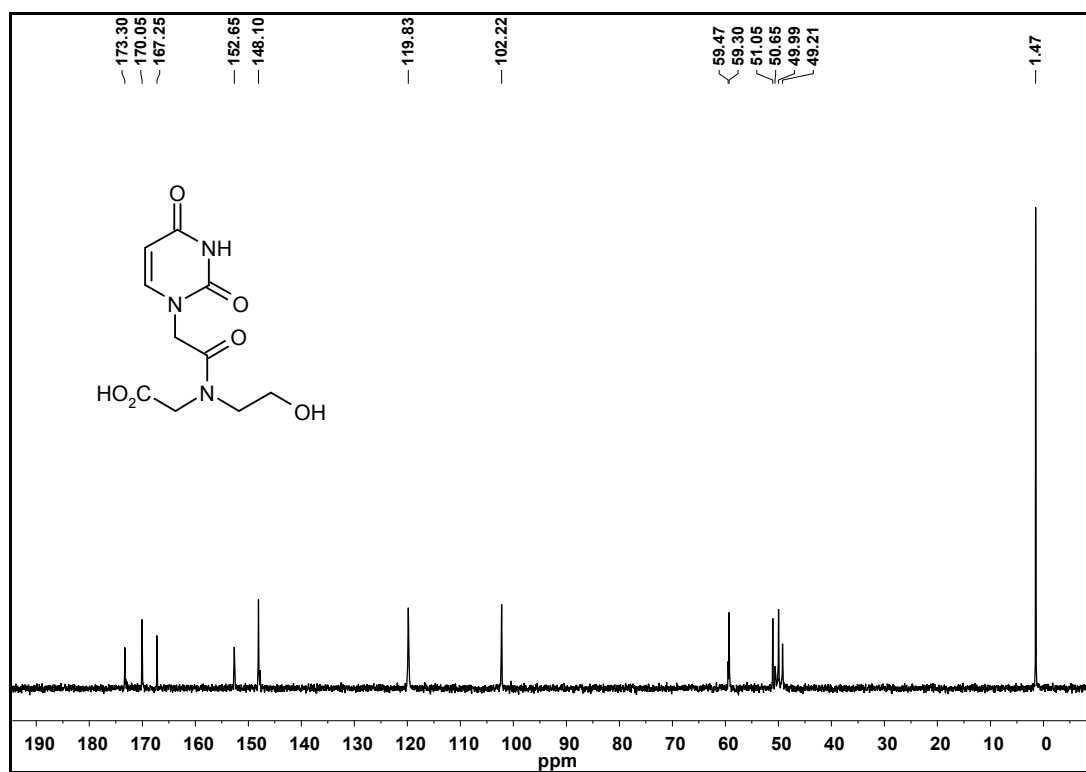
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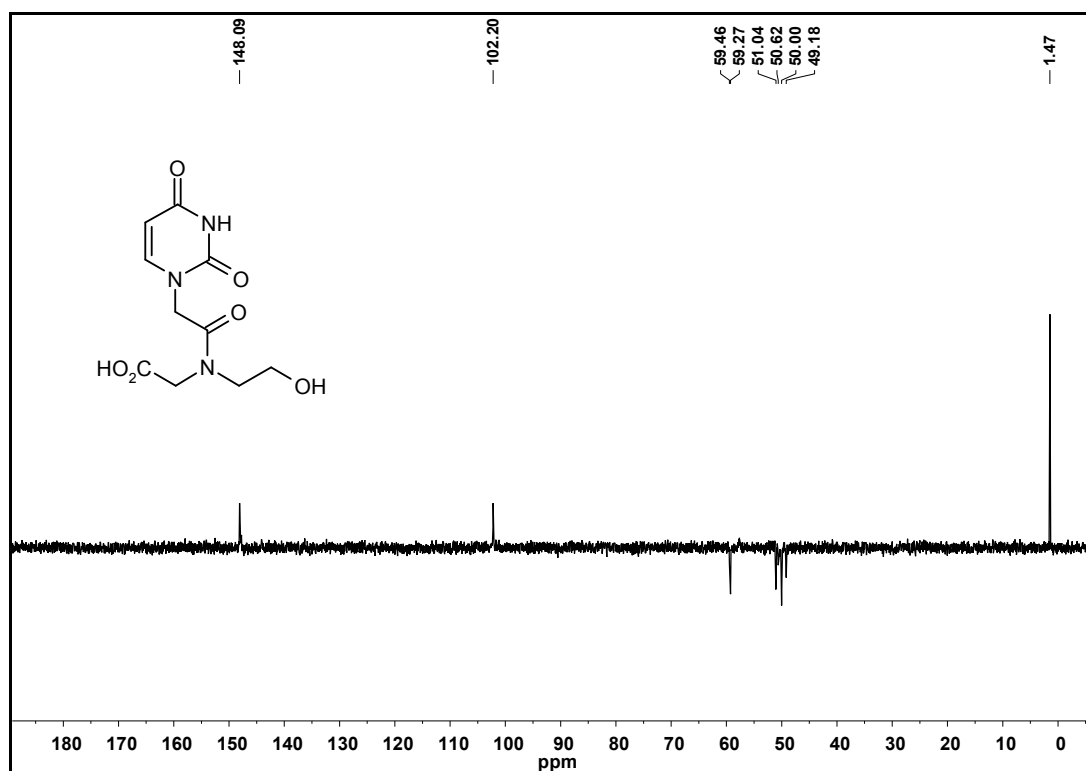
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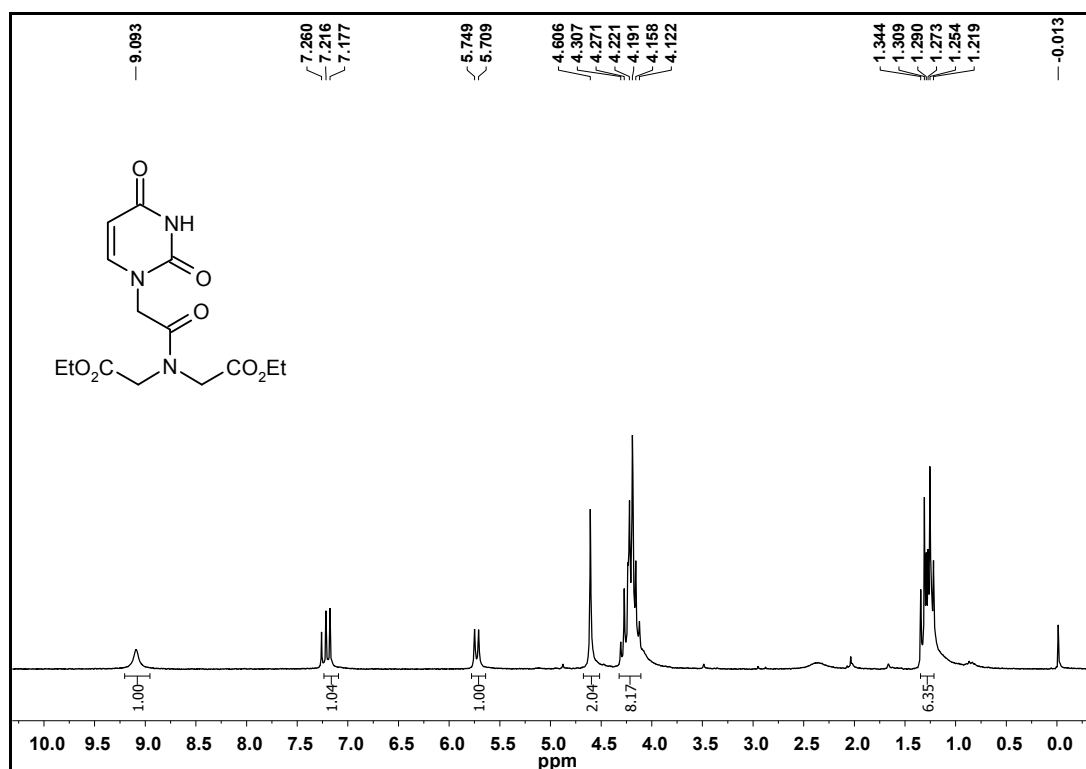
¹H-NMR of compound 8.

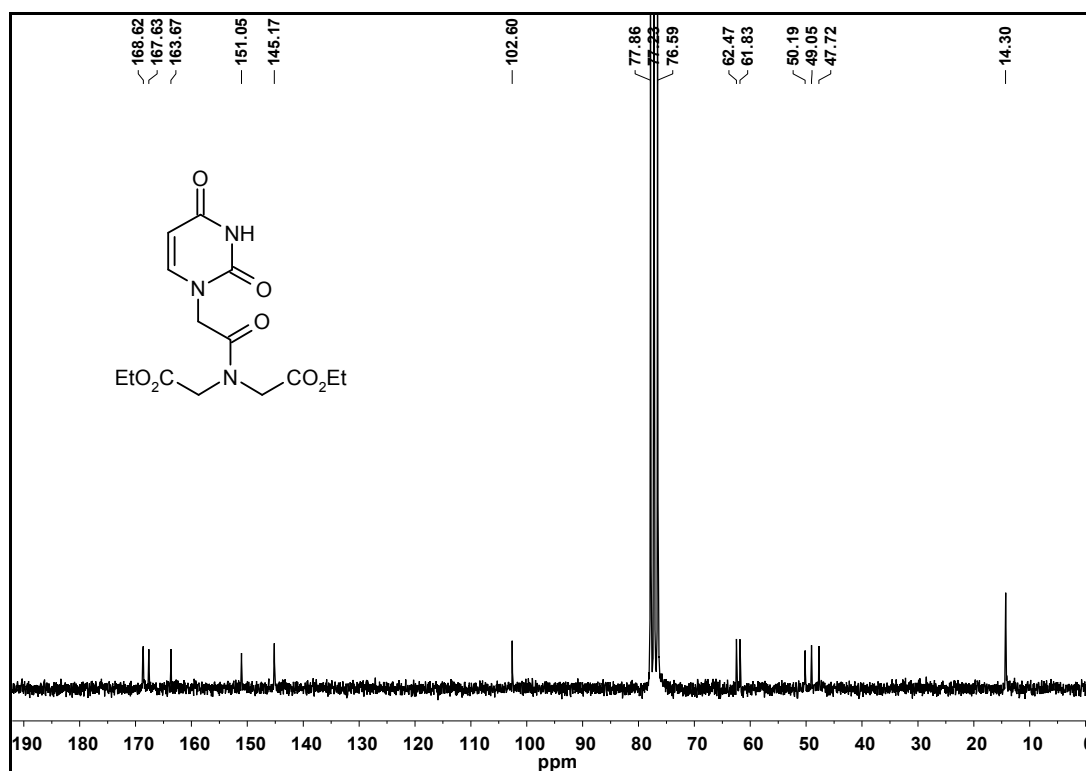


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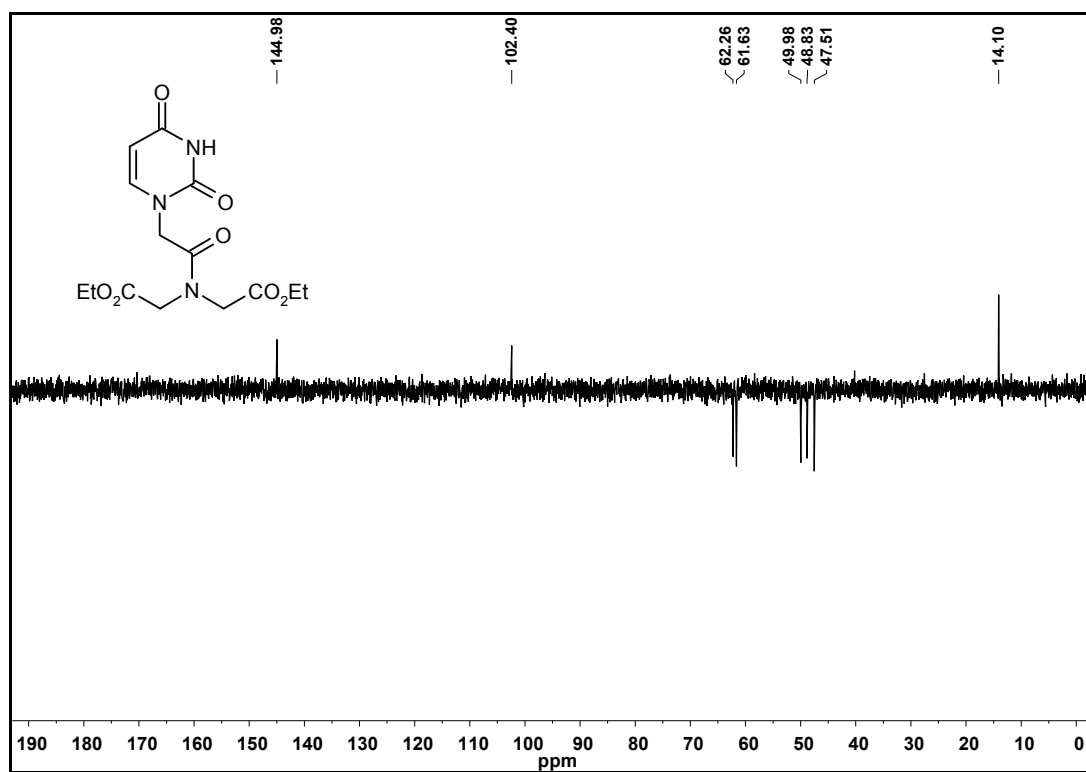


DEPT 135 NMR of compound 8.

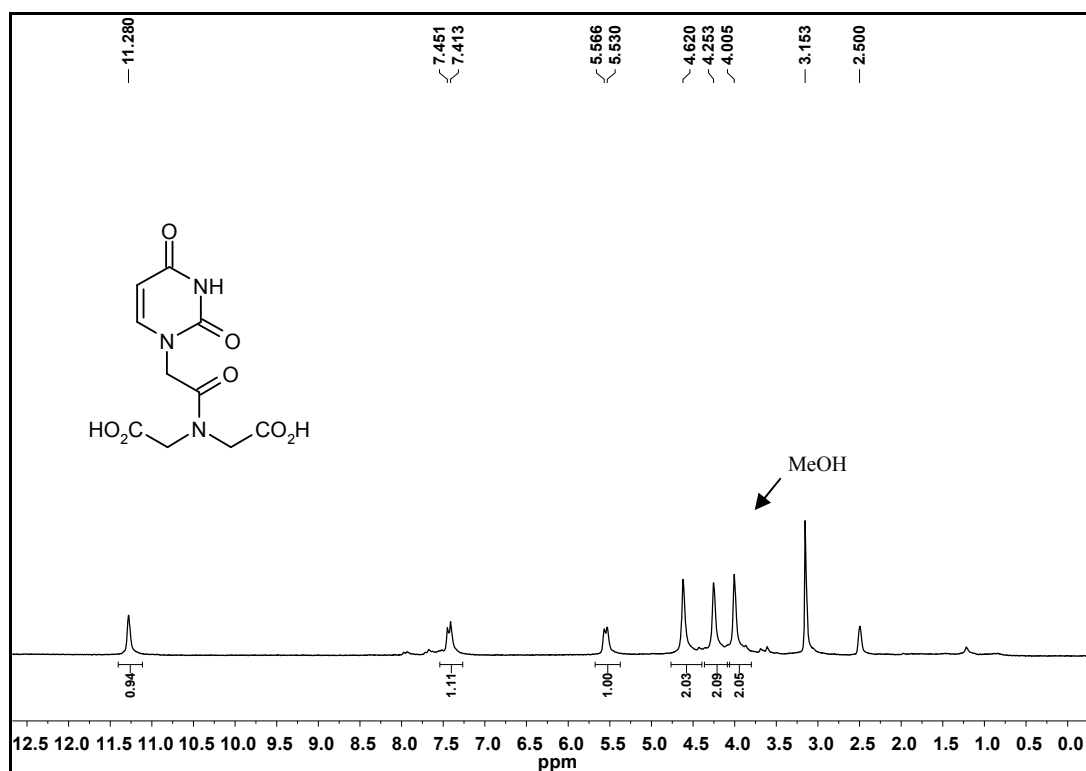
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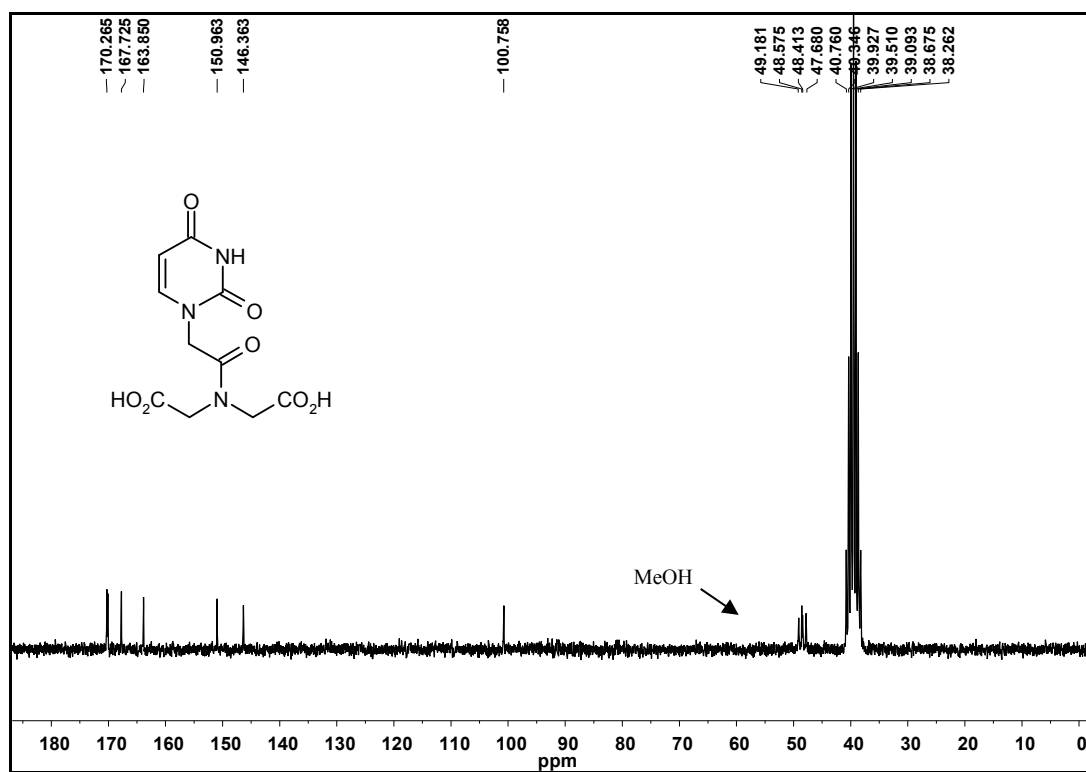
¹³C-NMR of compound **9**.



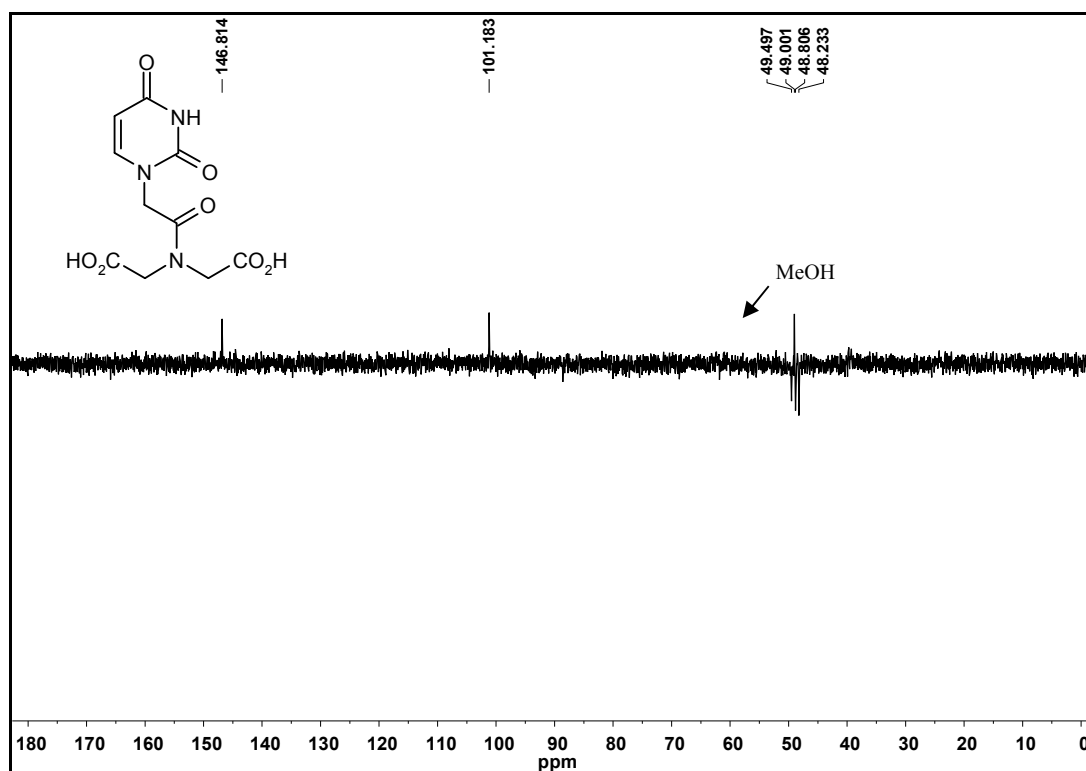
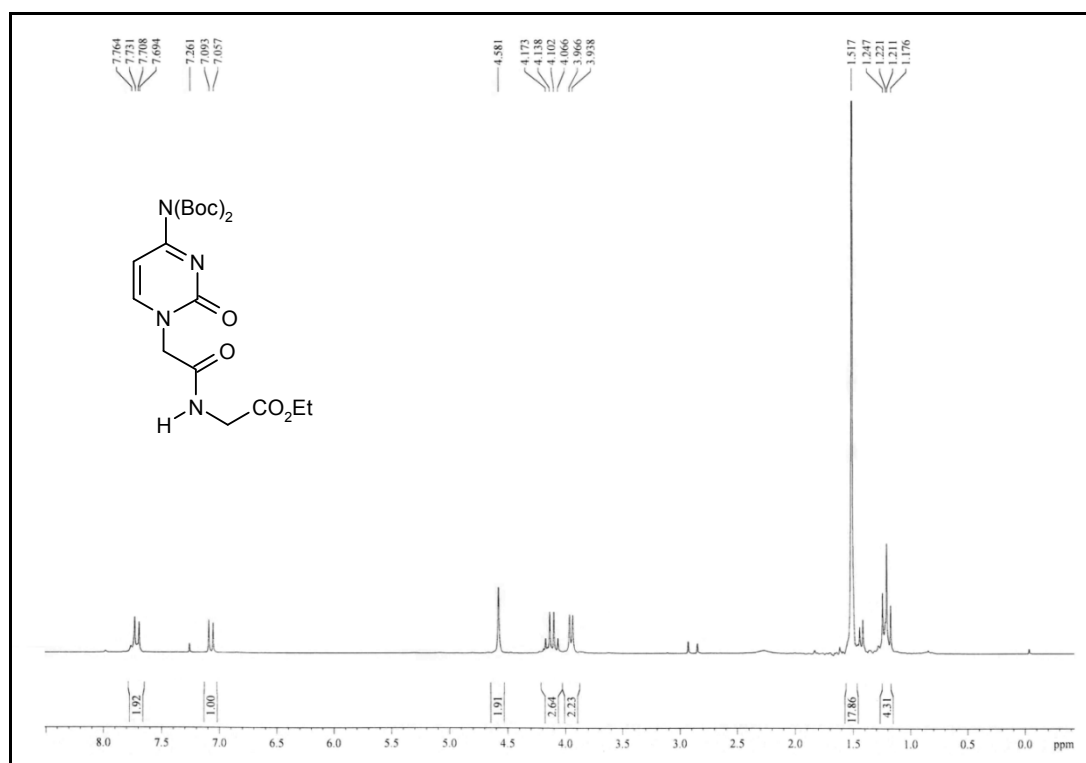
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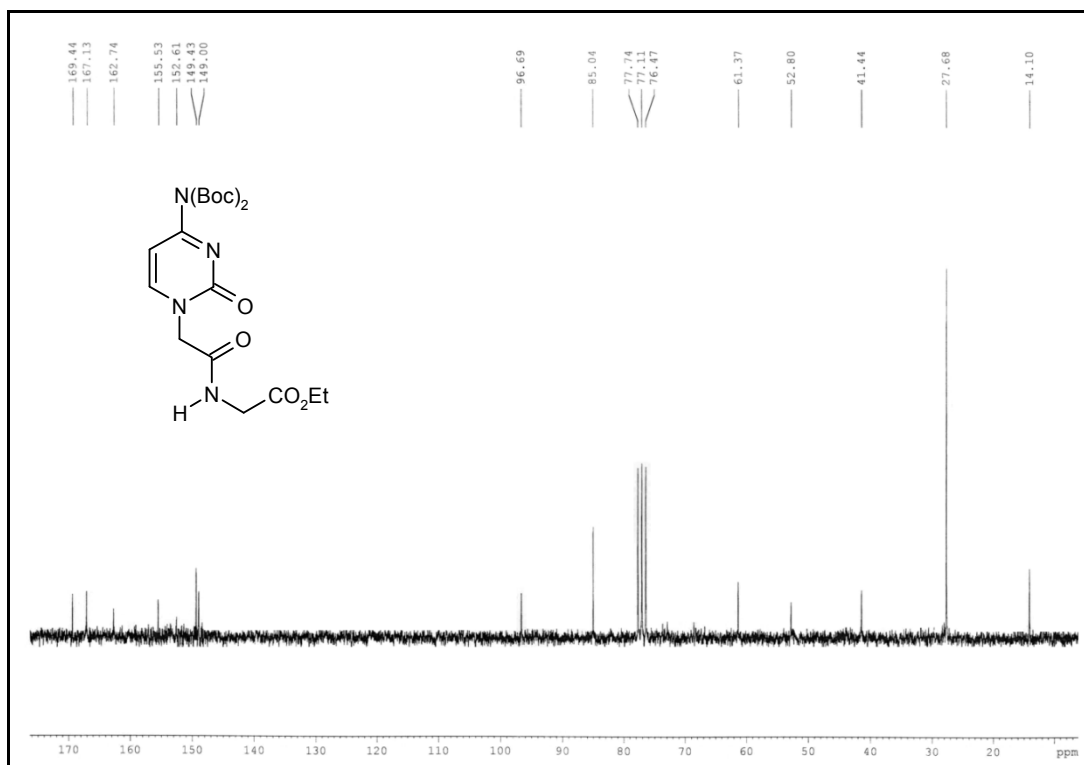


¹H-NMR of compound **10**.

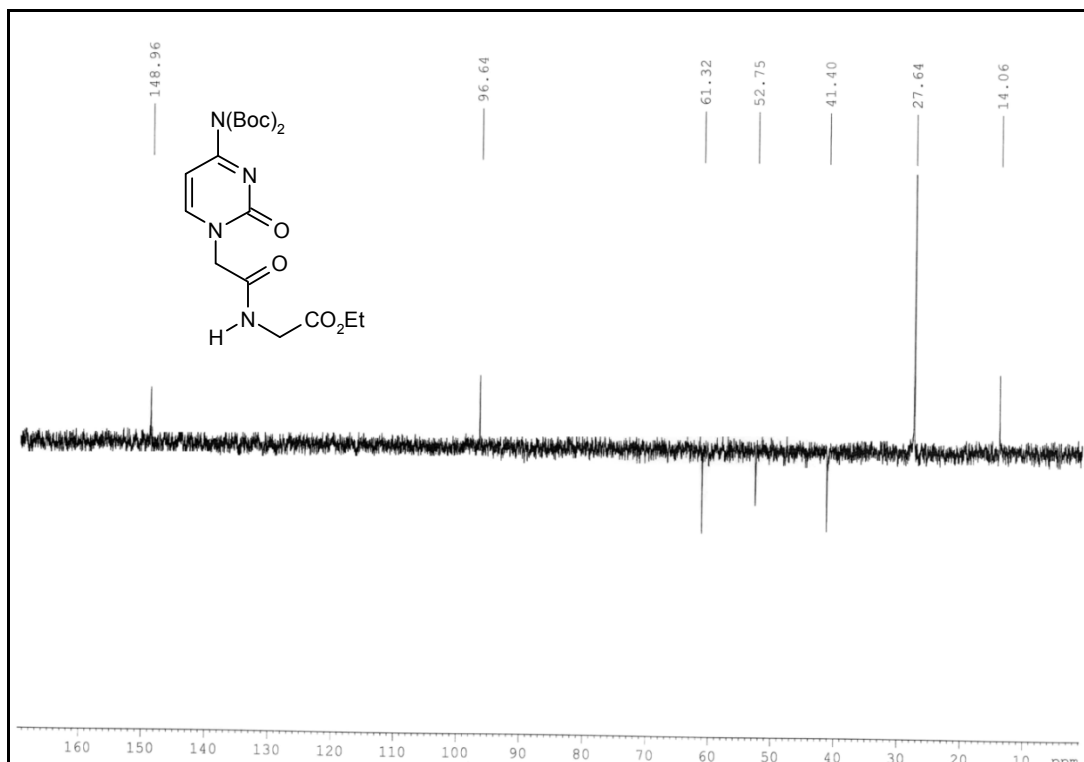


¹³C-NMR of compound **10**.

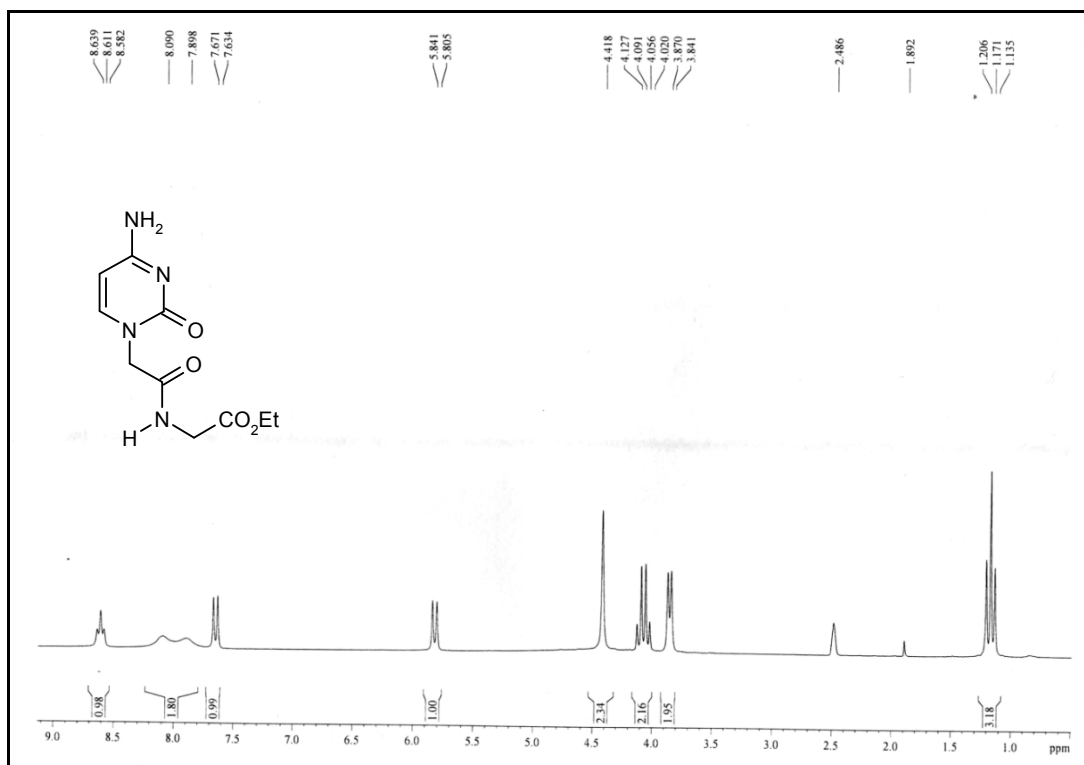
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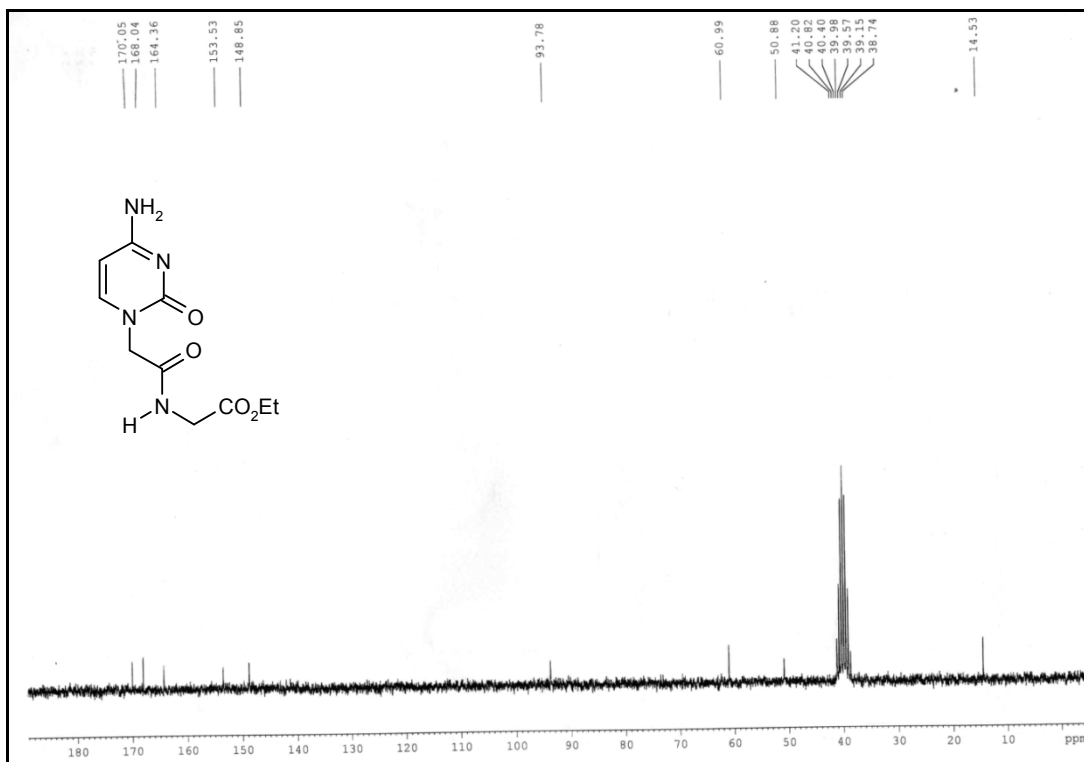
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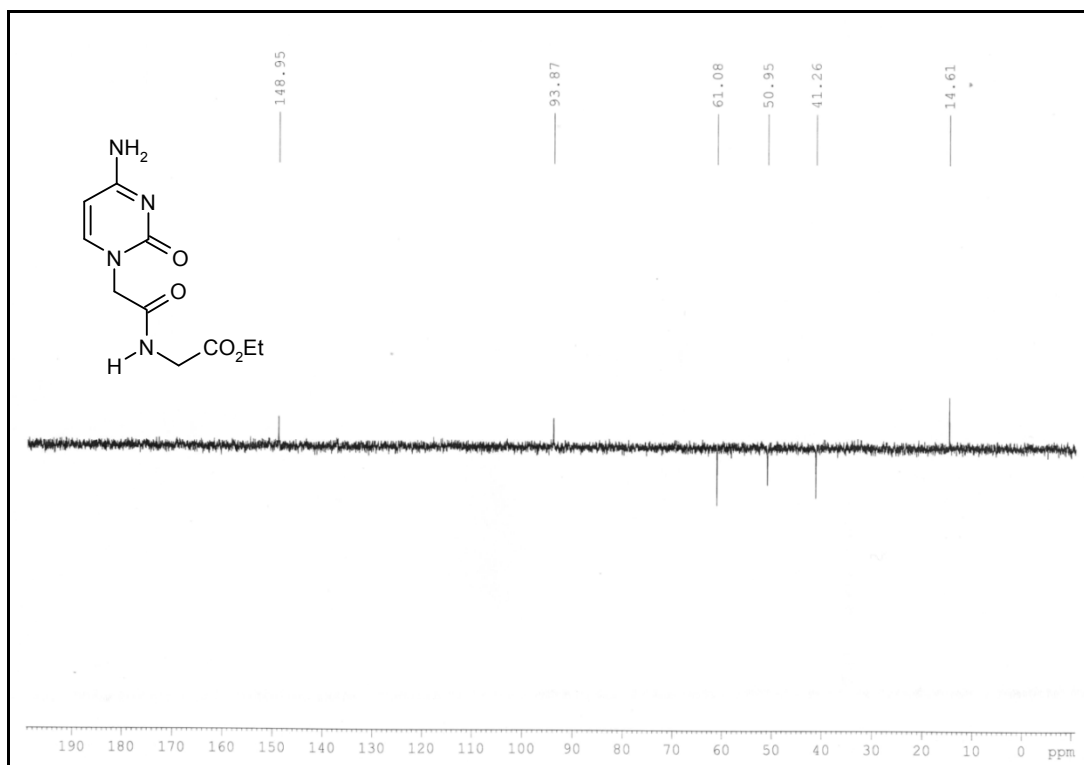
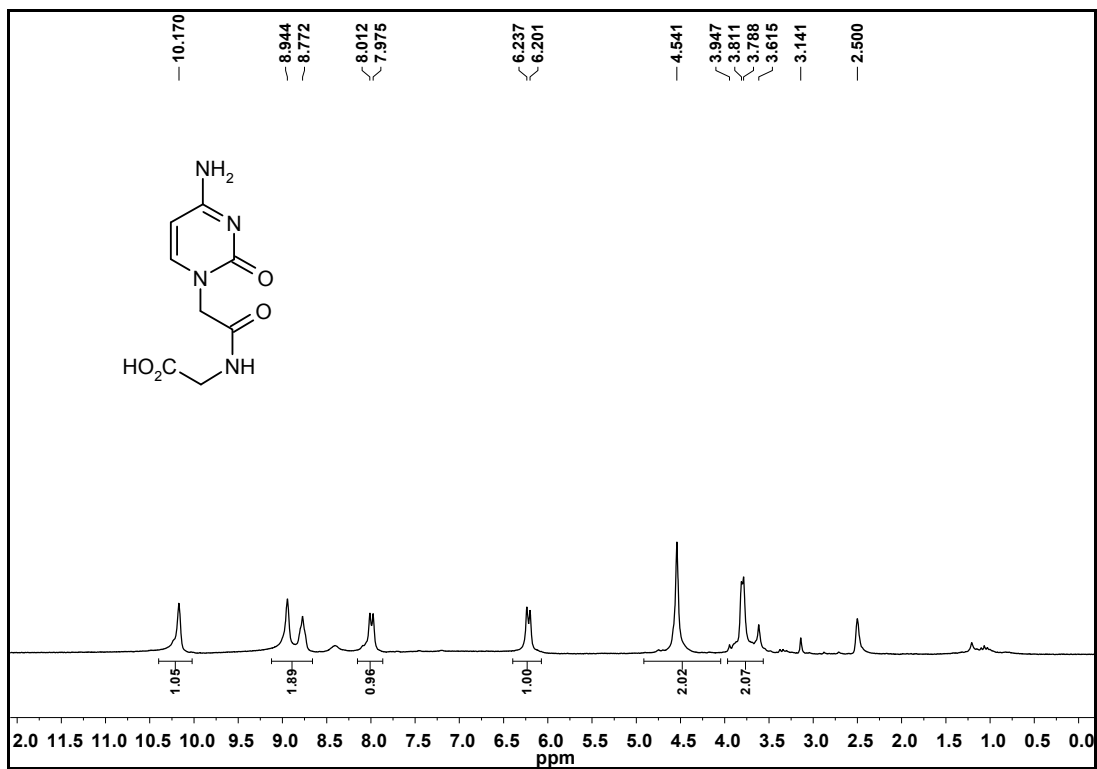
DEPT 135 NMR of compound **12**.

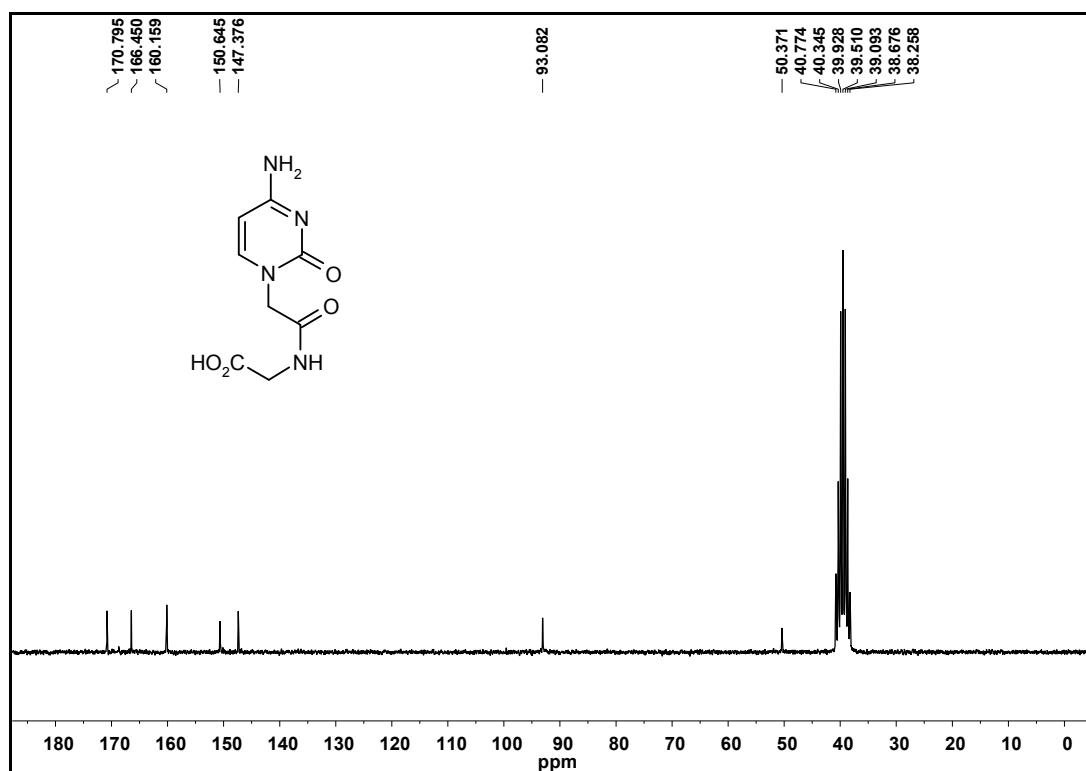


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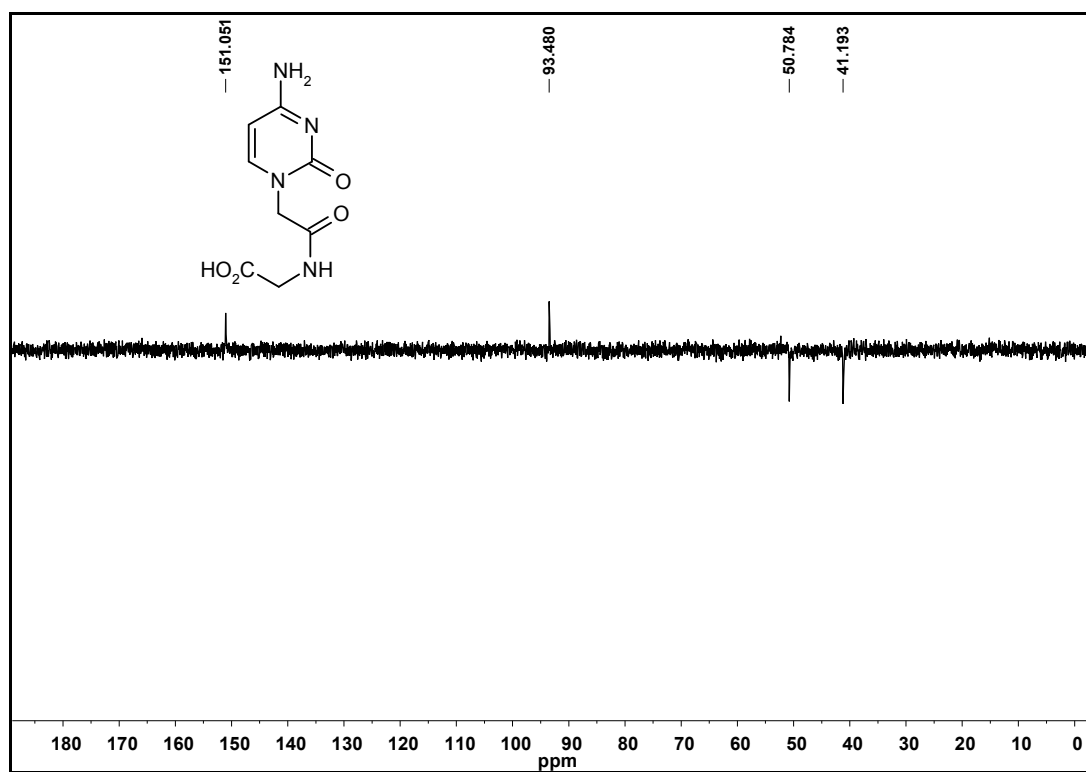


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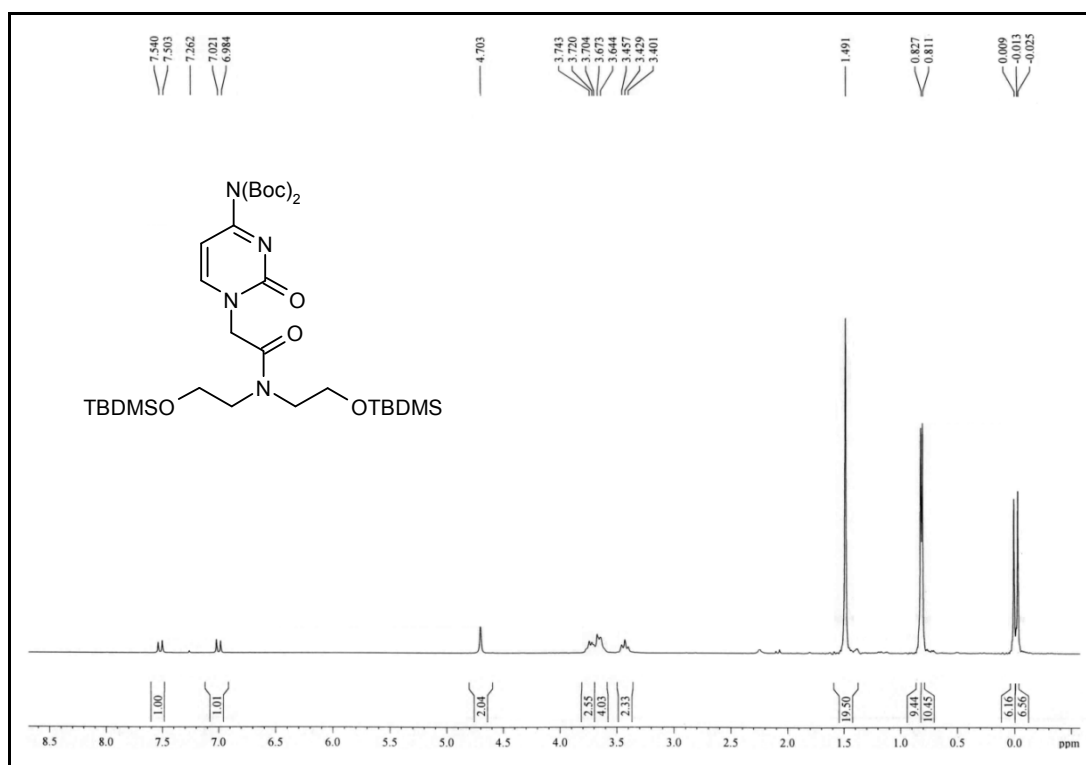
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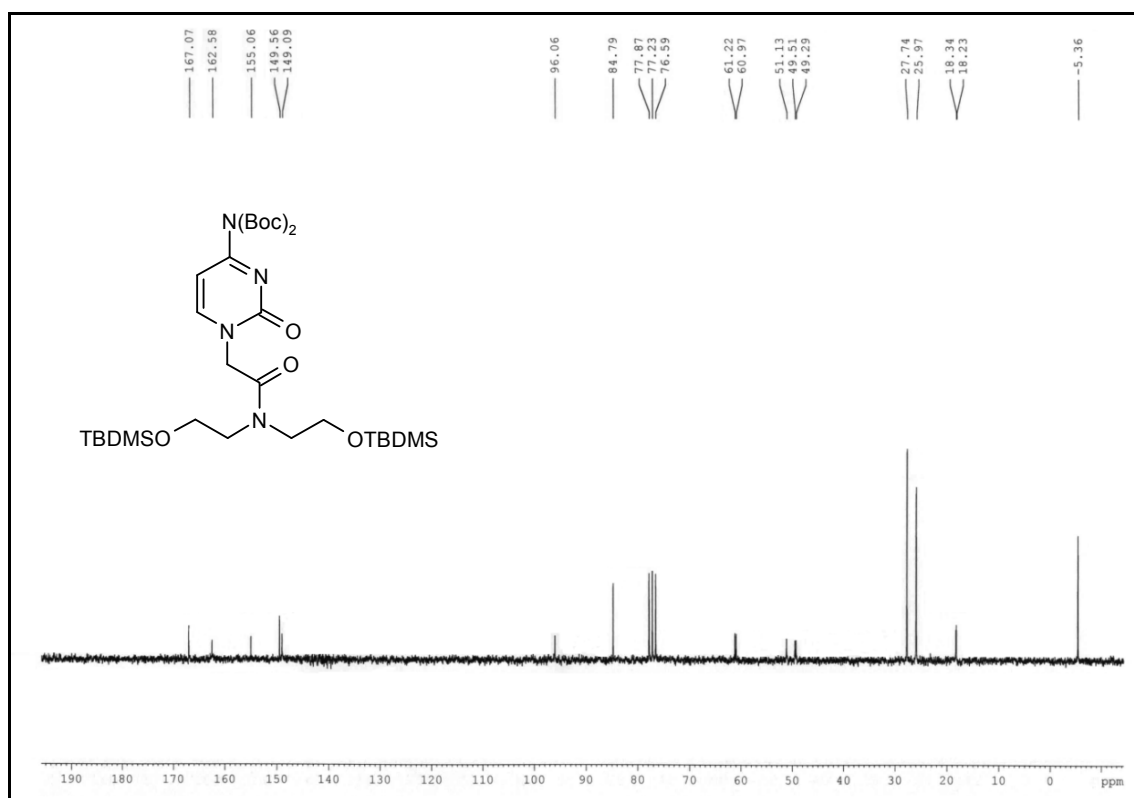
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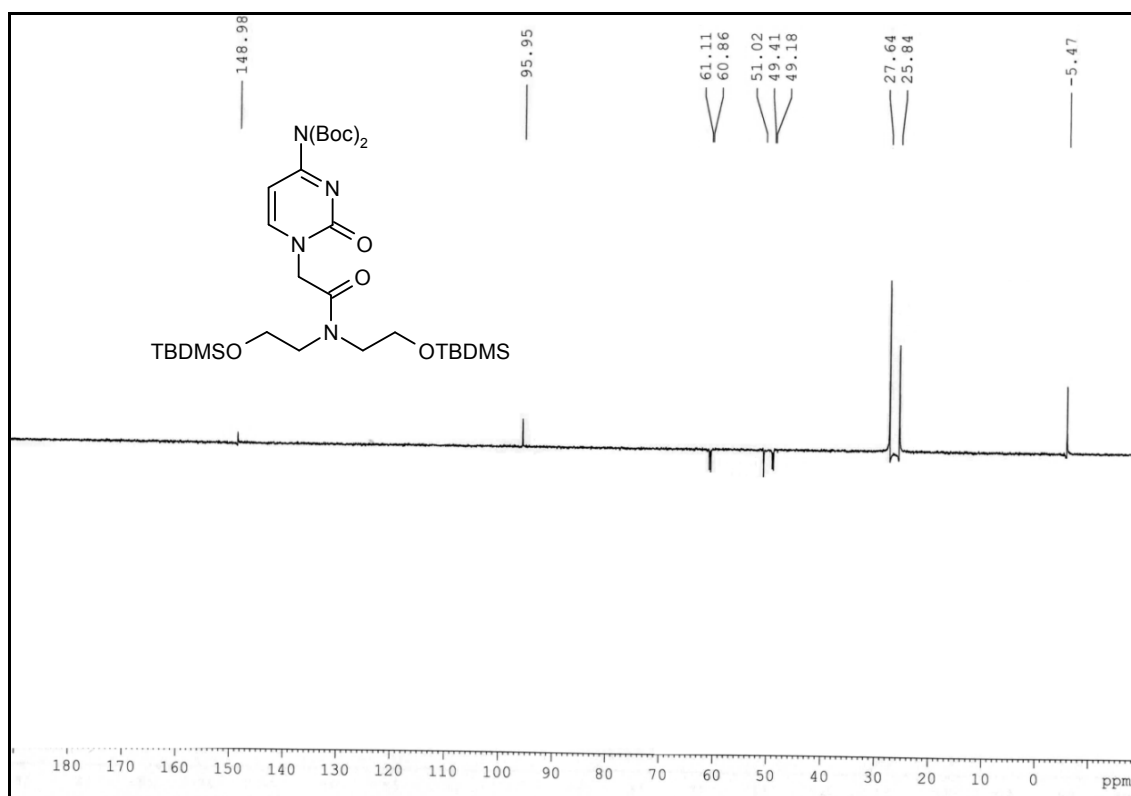
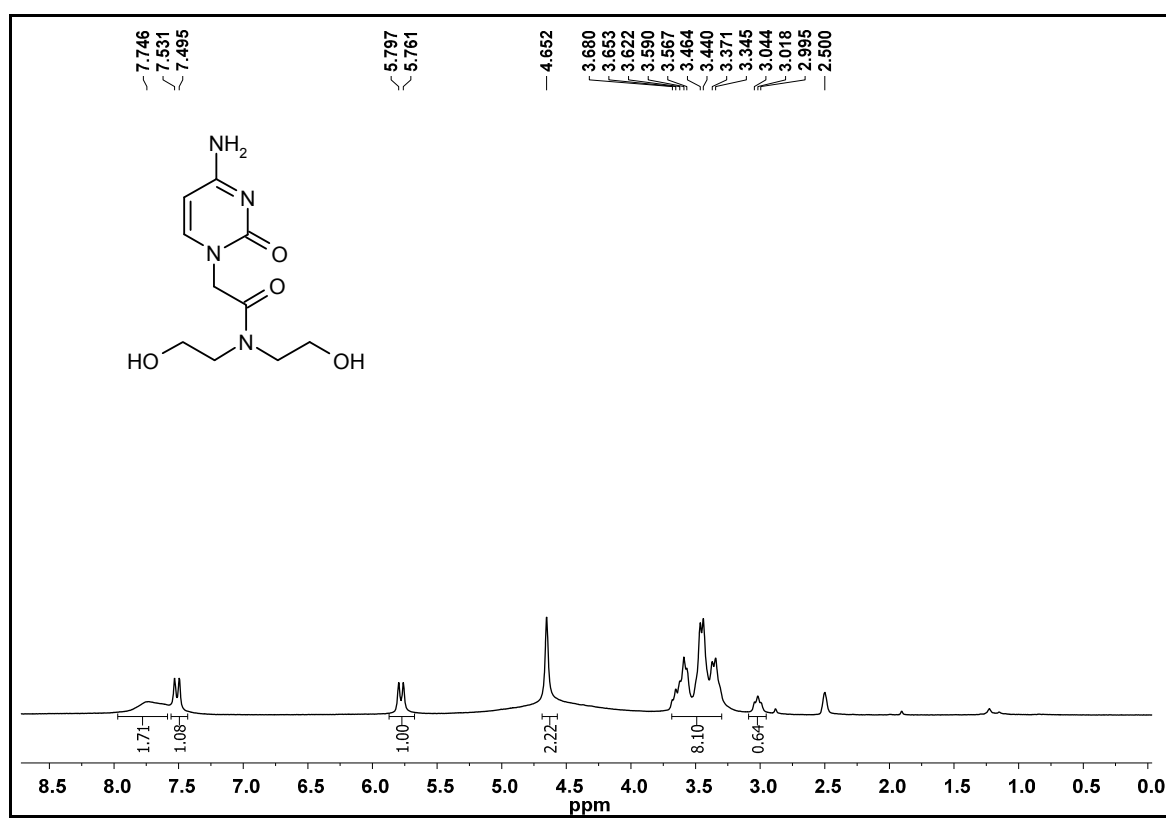
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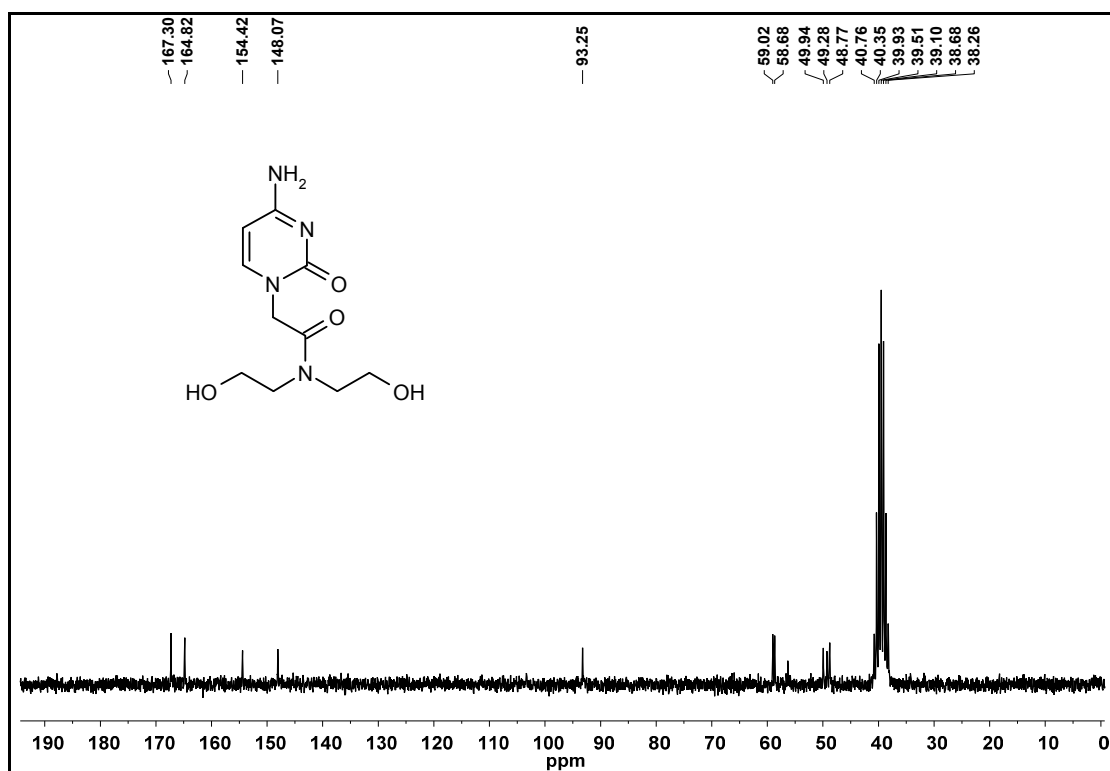


¹H-NMR of compound **15**.

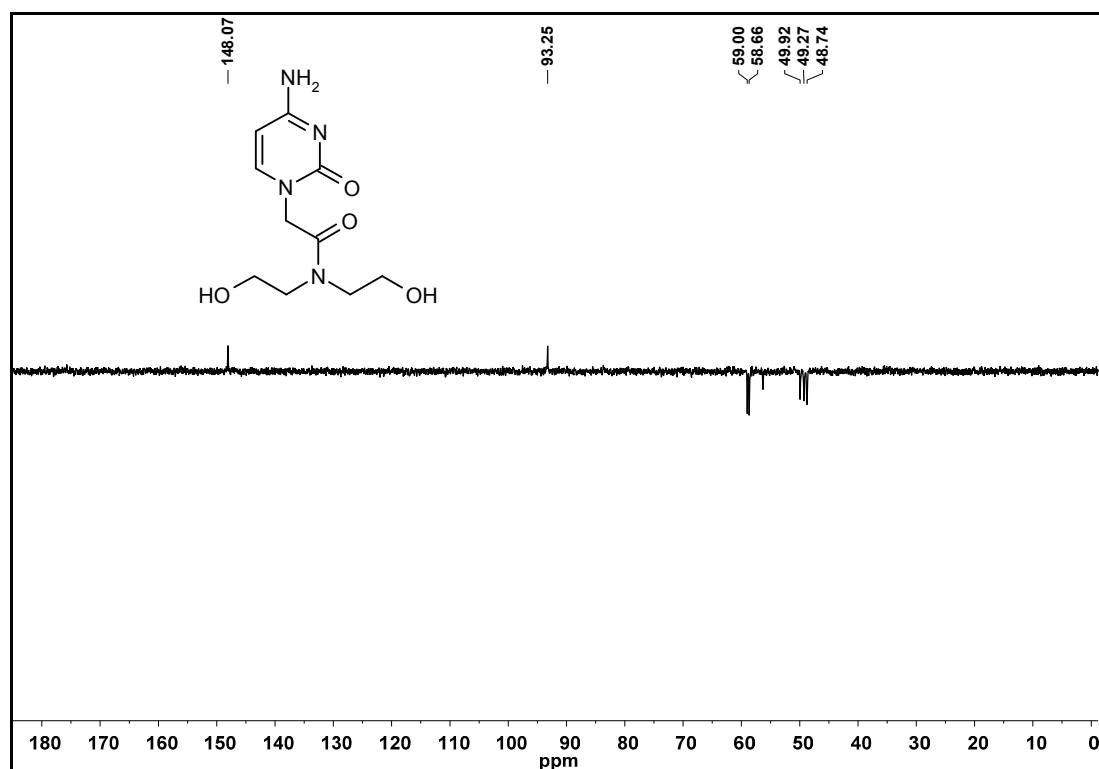


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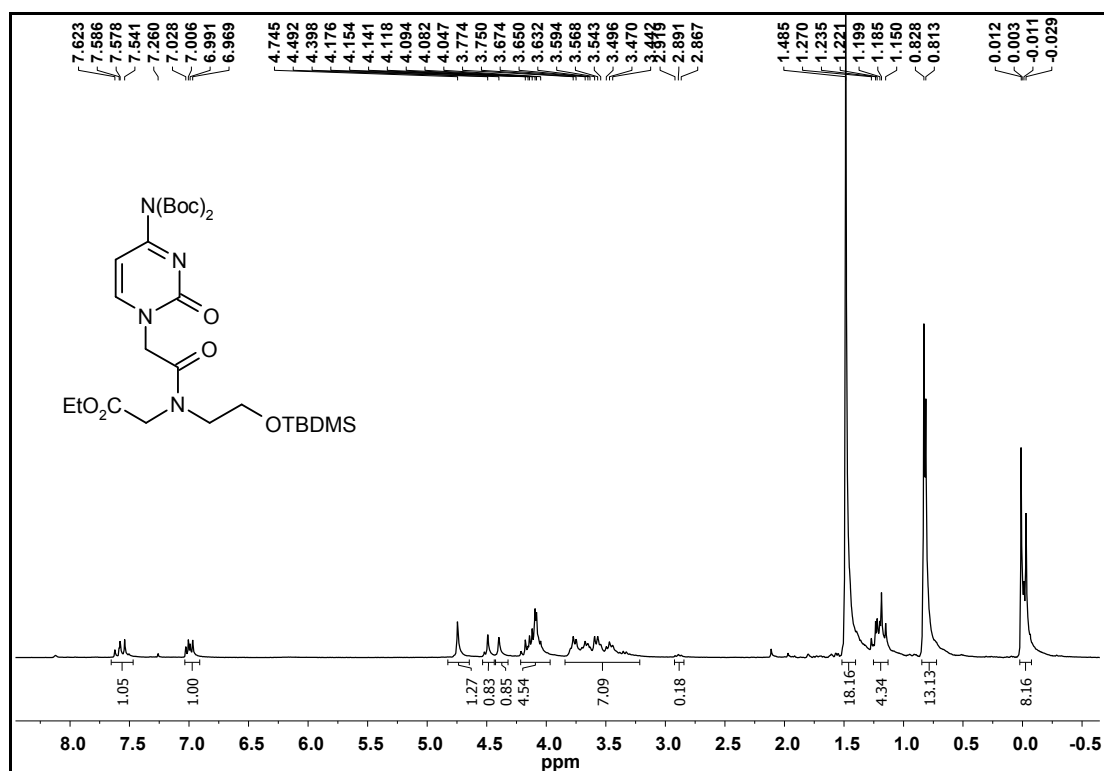
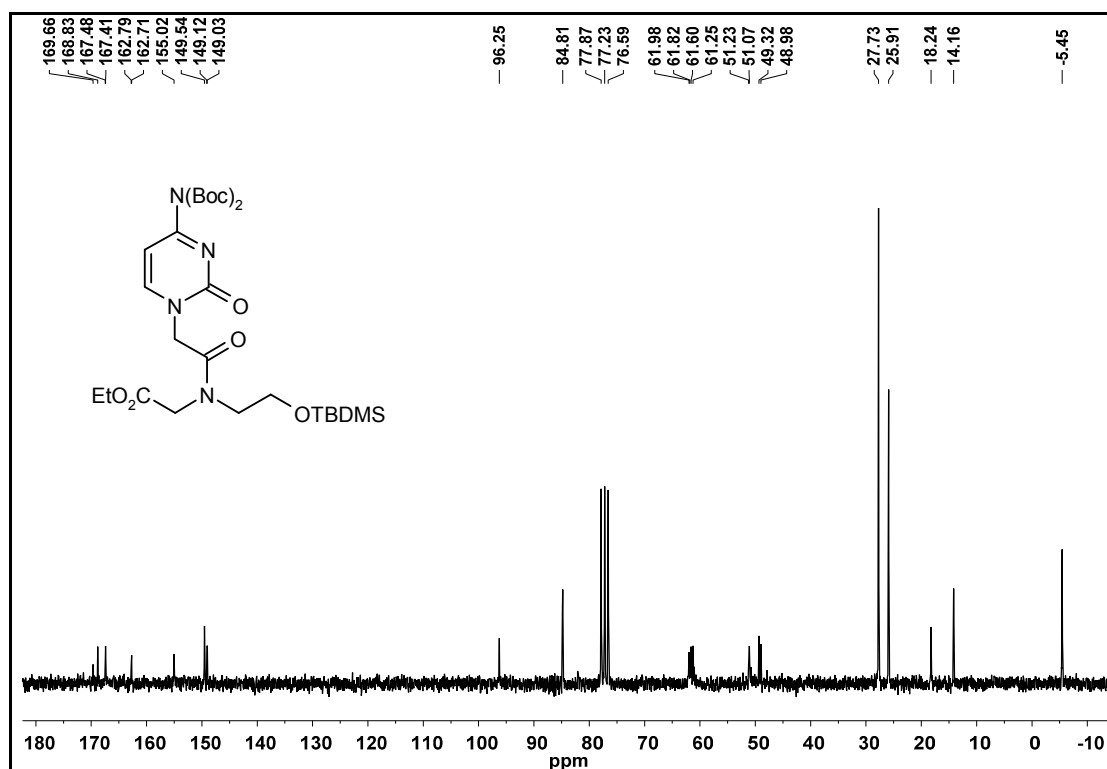
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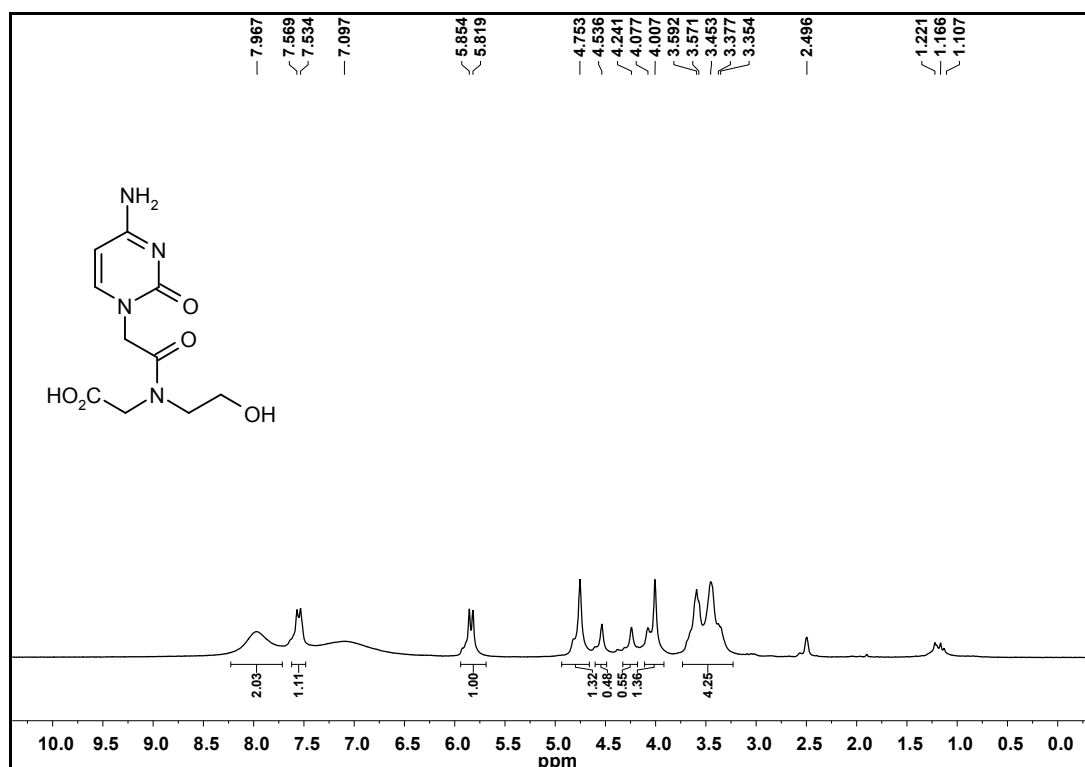
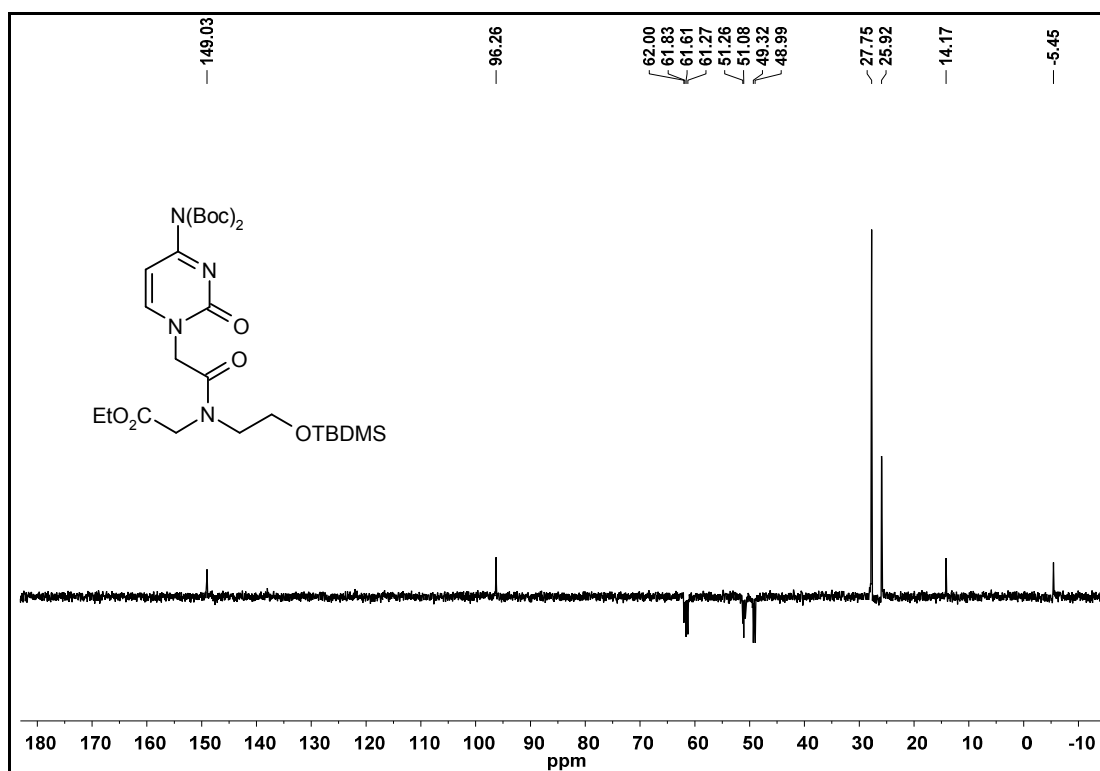


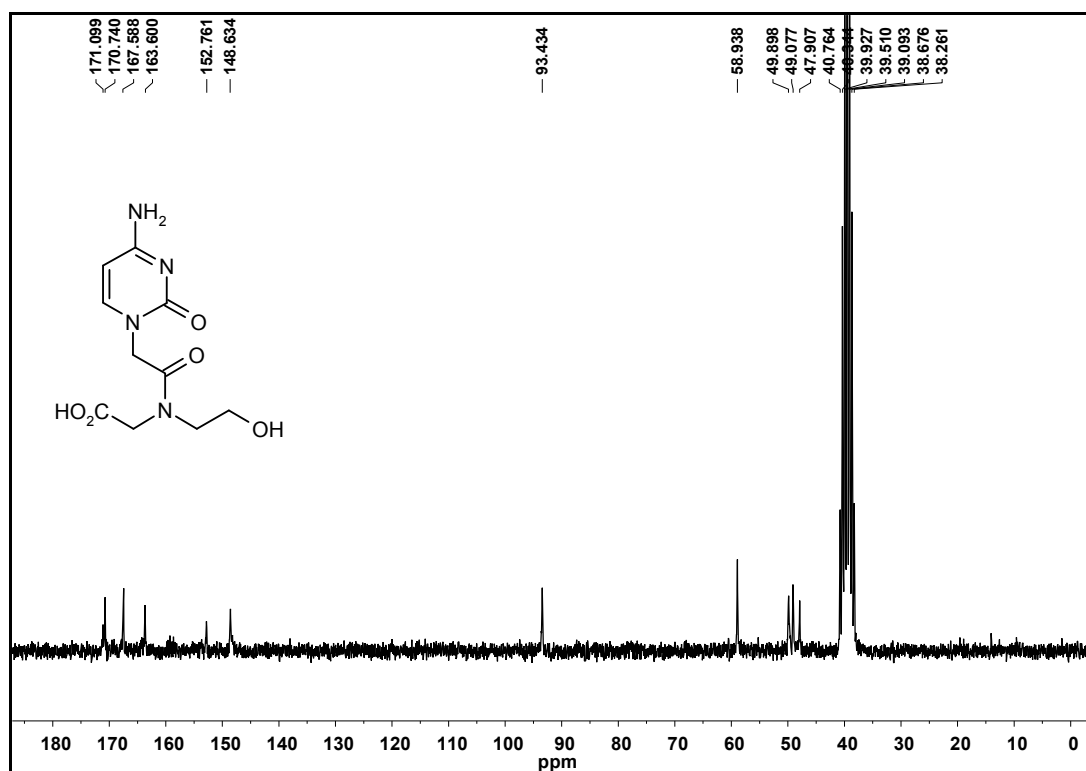
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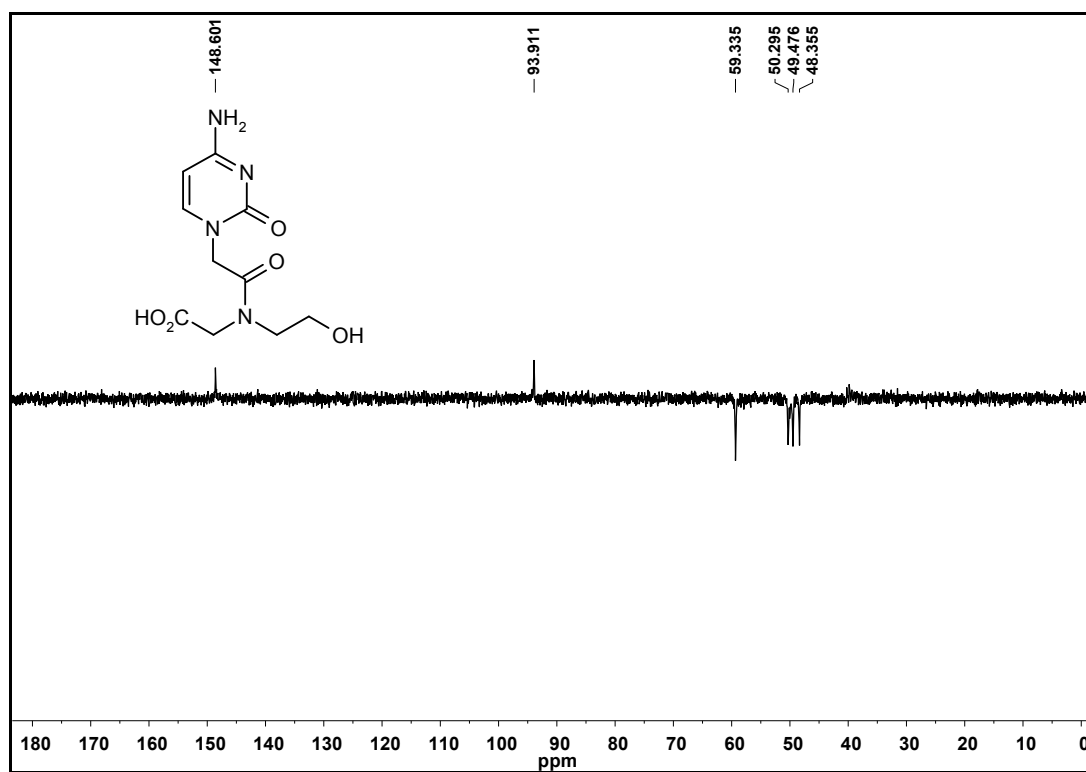
DEPT 135 NMR of compound **16**.

¹H-NMR of compound 17.¹³C-NMR of compound 17.

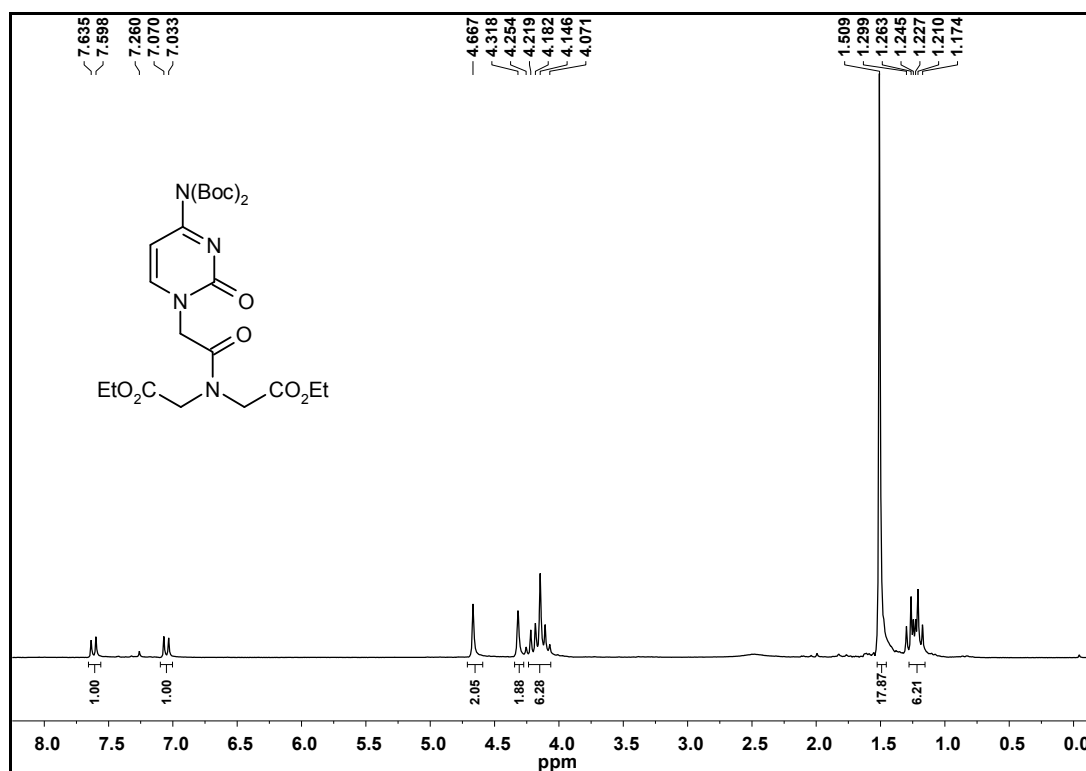




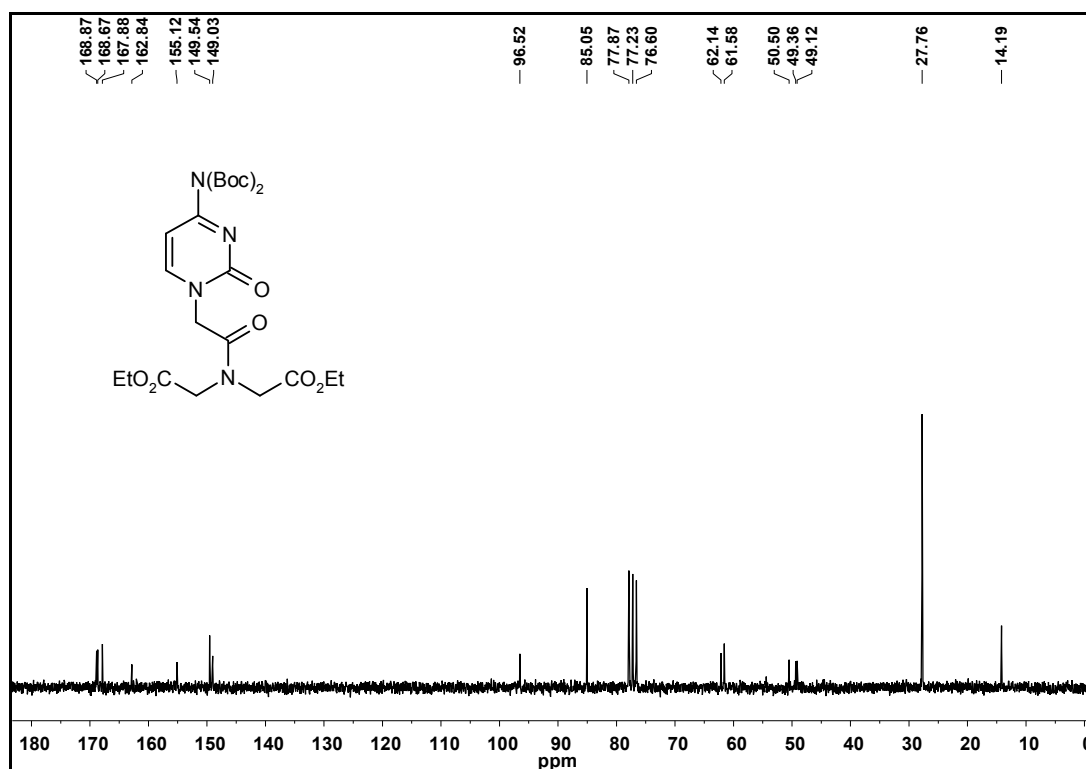
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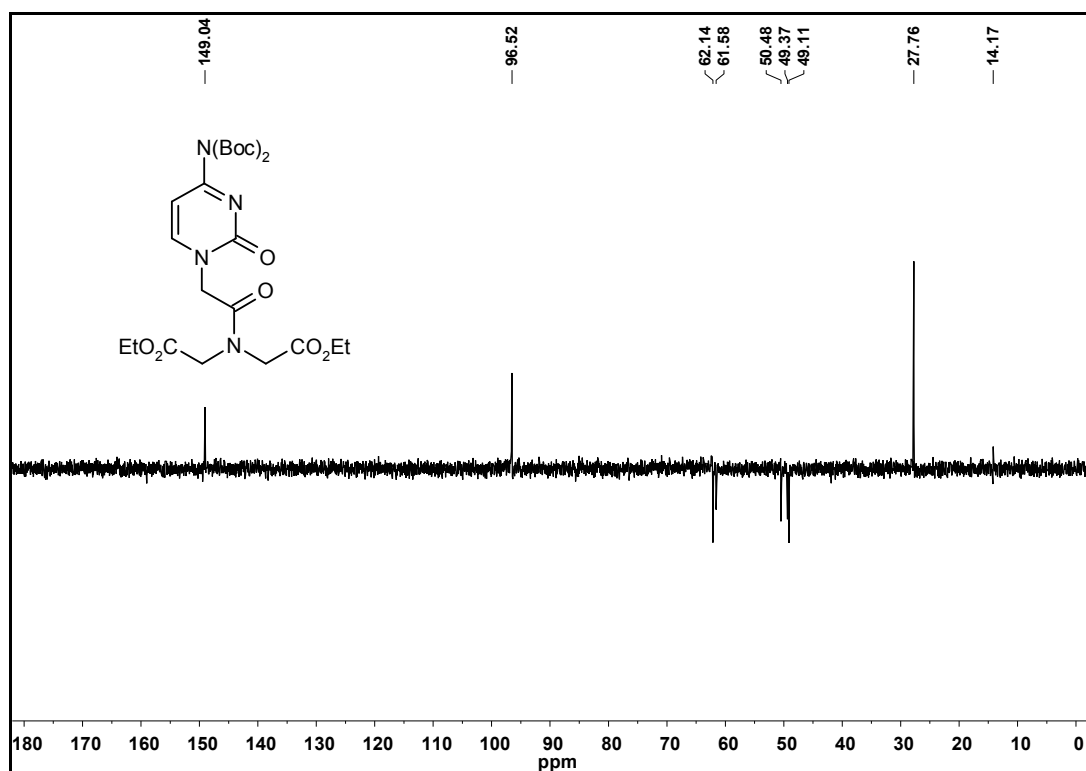
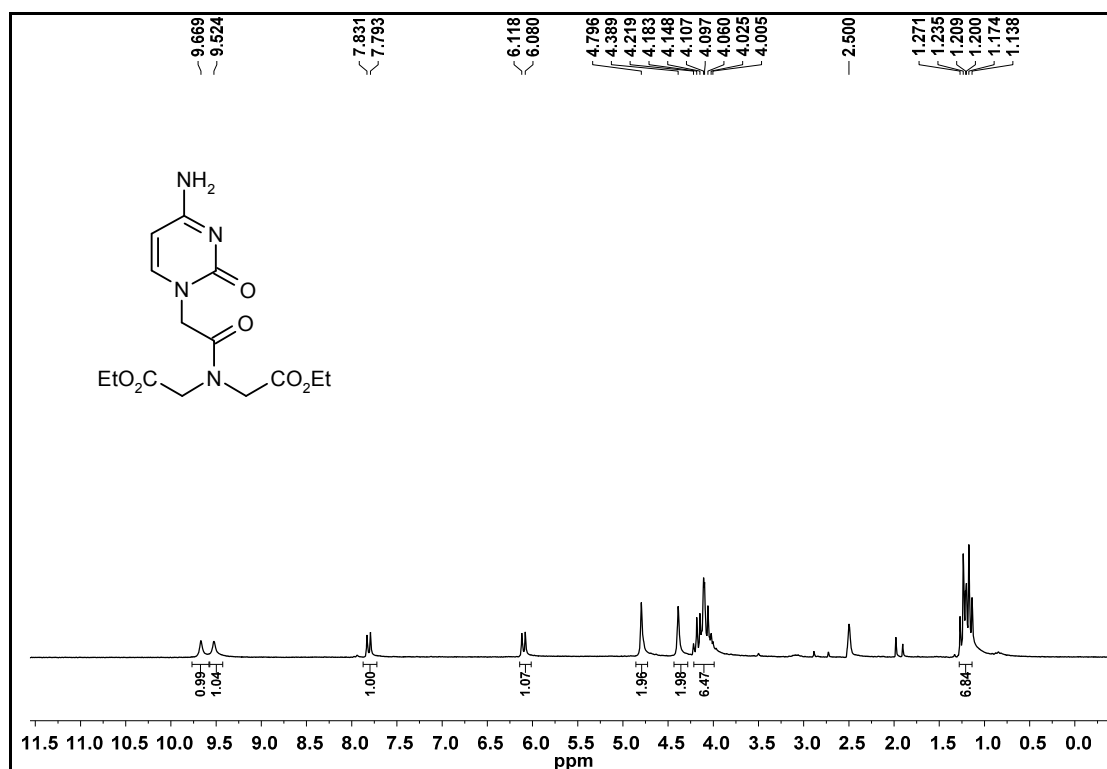
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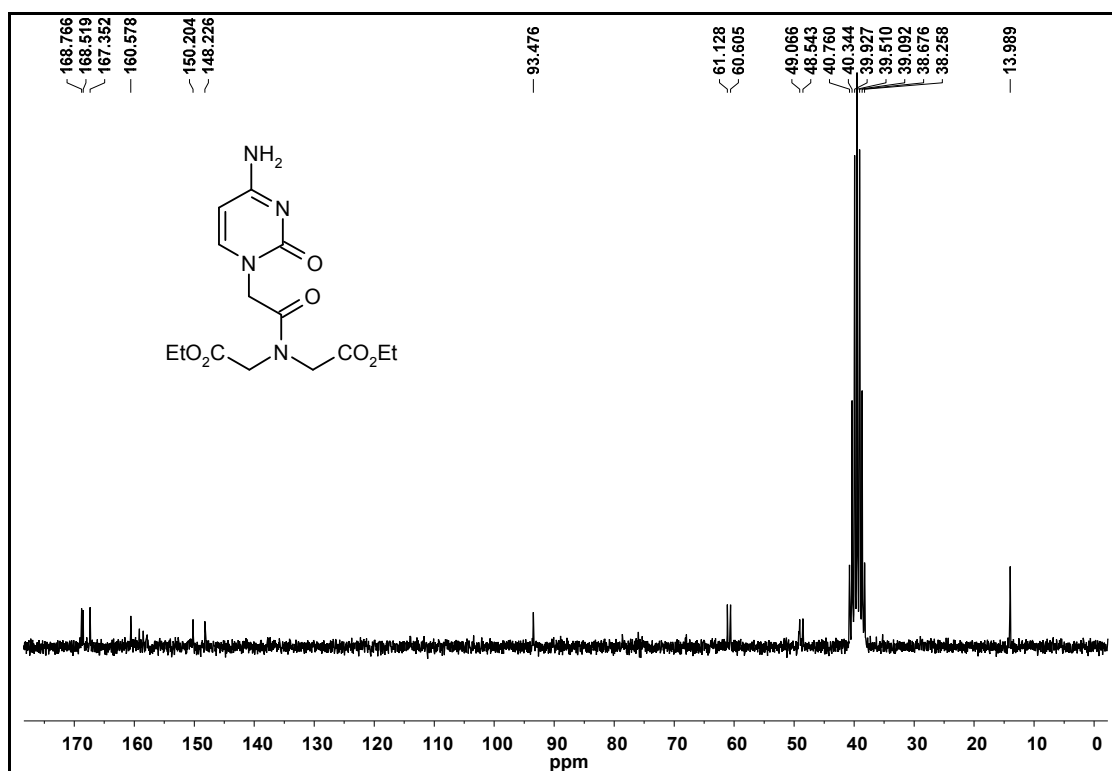


¹H-NMR of compound **19**.

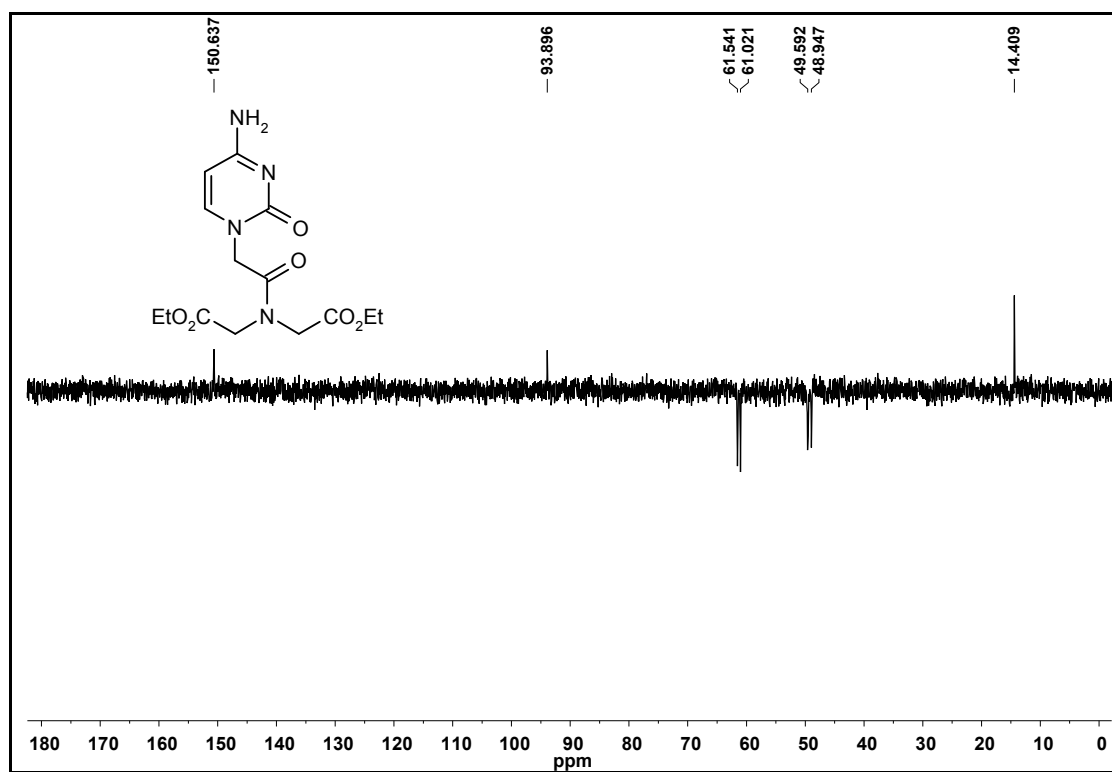


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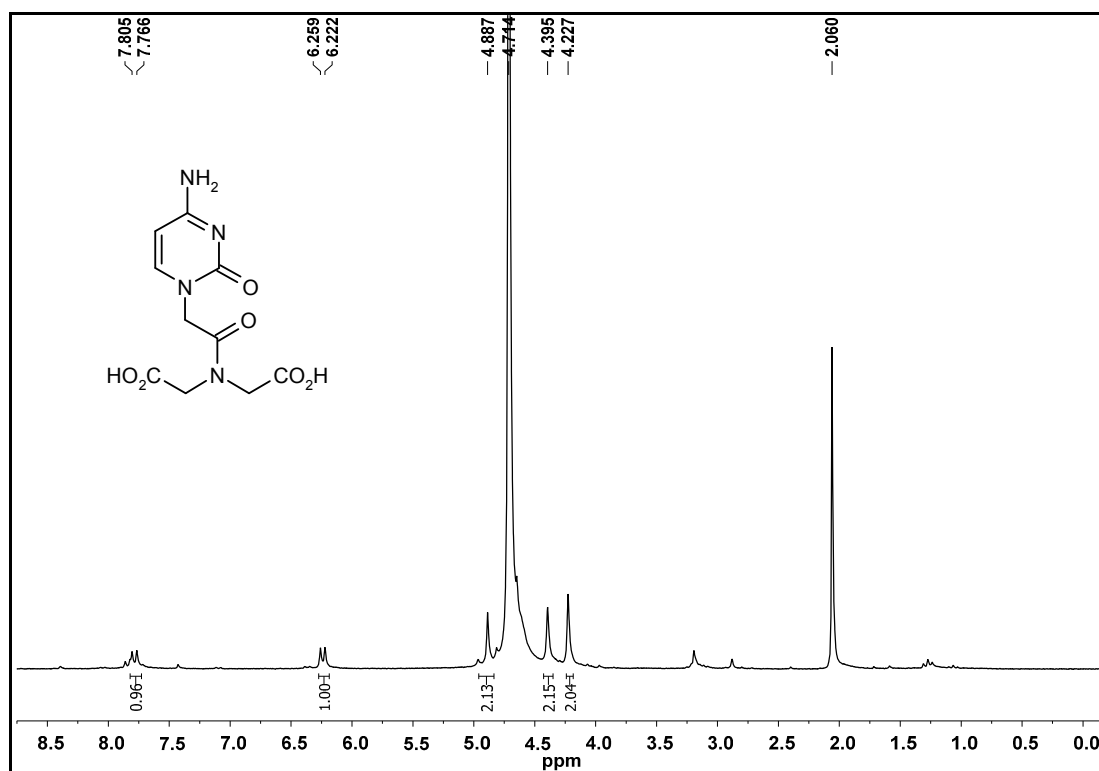
DEPT 135 NMR of compound **19**. ^1H -NMR of compound **20**.



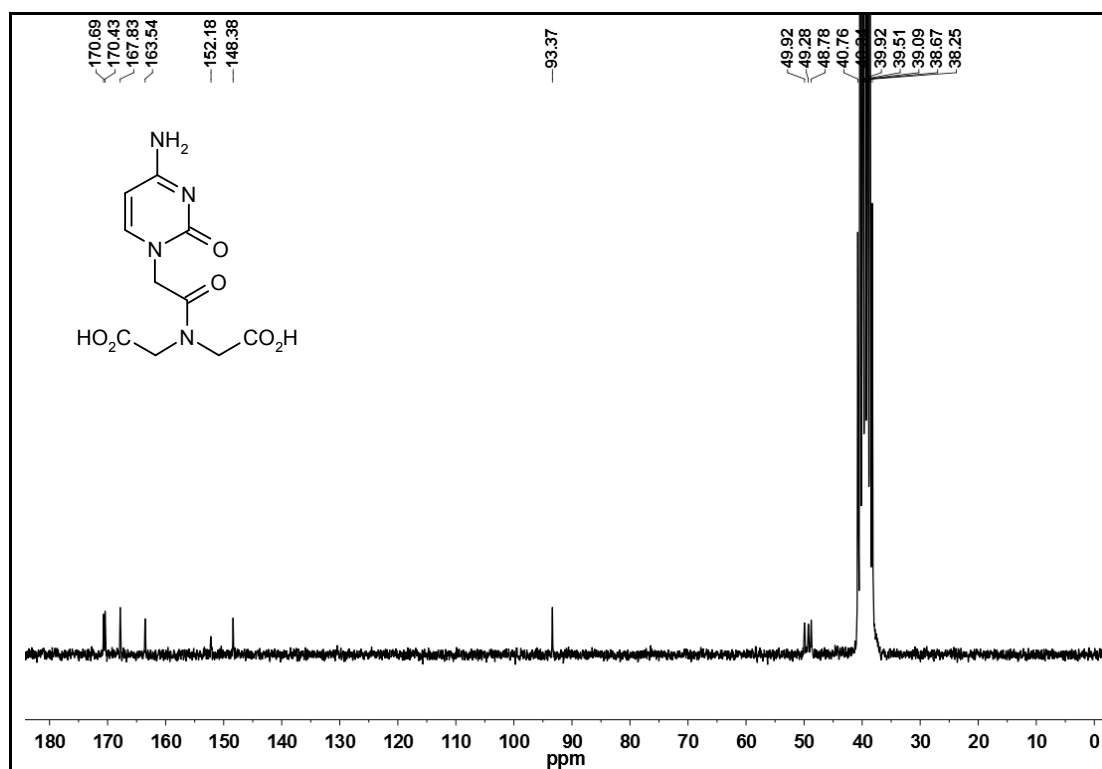
¹³C-NMR of compound **20**.



DEPT 135 NMR of compound **20**.



¹H-NMR of compound **21**.



¹³C-NMR of compound **21**.

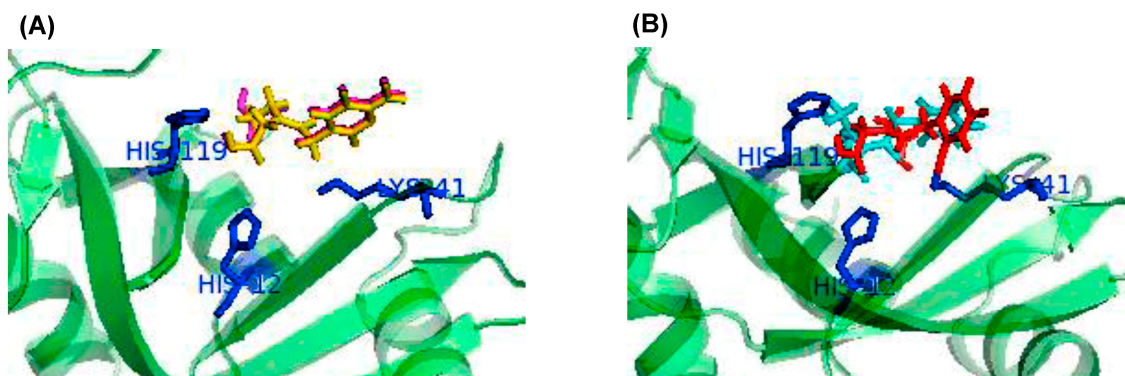
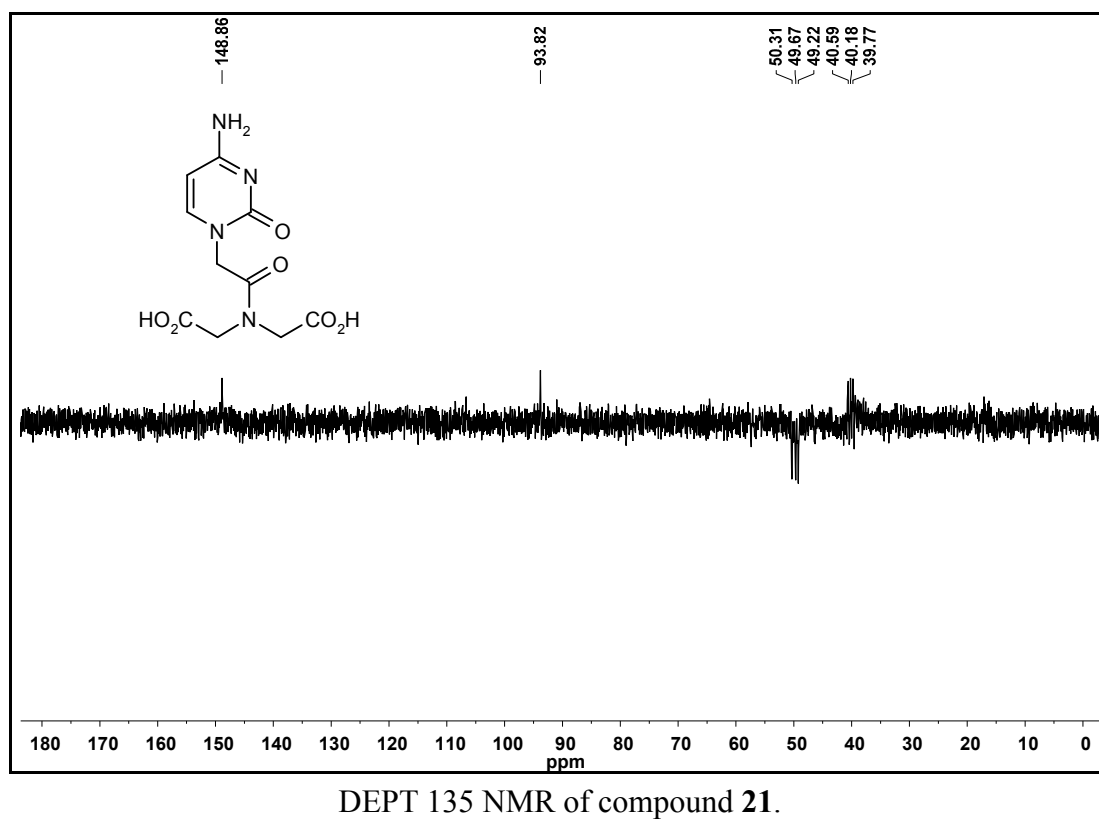


Figure S3. Docked poses of (A) C-di-acid (**21**) (magenta) and C-ol-acid (**18**) (yellow); or (B) U-di-acid (**10**) (red) and U-ol-acid (**8**) (cyan) with RNase A (1FS3).