

SUPPLEMENTARY INFORMATION

Chitosan Supported Ionic Liquid, a Multifaceted Catalyst for Streamlined and Efficient Synthesis of Carboxylic, Amino acid and Carbohydrate Esters

Praachi Kakati,^a Satish Kumar Awasthi^{a*}

^aChemical Biology Laboratory, Department of Chemistry, University of Delhi, Delhi-110007, India.

*Corresponding author: satishpna@gmail.com

Table of Contents	Page No.
Materials	S2
Instrumentation	S2
Table S1. Comparison of the present work with previous literature for the esterification of acid to their corresponding esters.	S3-S4
Tale S2. Optimization of reaction conditions for the amount of catalyst using benzoic acid A1 and ethanol B1 as the model substrate. ^a	S4
Gram scale synthesis	S4
Figure S1. Practical application of a few value-added esters synthesized in gram scale.	S5
Figure S2. Plausible mechanism for Chitosan-IL ₆ catalysed selective esterification of glycine amino acid and ethanol to ethyl glycinate.	S6
Figure S3. Plausible mechanism for Chitosan-IL ₆ catalysed selective esterification of acetic anhydride and carbohydrate to carbohydrate esters.	S7
Table S3. Recyclability test of Chitosan-IL ₆ catalyzed esterification of benzoic acid and ethanol to ethyl benzoate C1.	S8
Figure S4. IR, XRD ad SEM image of reused catalyst.	S9
Determination of acidity of Chitosan-IL ₆	S9-S10
Calculation of TOF of Chitosan-IL ₆	S10
Table S4. TOF values Chitosan-IL ₆ for the products C1-C29 ^a	S11-S12

Table S6. TOF values Chitosan-IL ₆ for the products E1-E7 ^a	S13
Figure S5. N ₂ adsorption/desorption isotherm of (a) Chitosan and (b) Chitosan IL ₆ and Pore size distribution of (c) Chitosan and (d) Chitosan IL ₆	S14
¹ H NMR of 1,4-bis(5-carboxypentyl)pyrazine-1,4-dium ([BCPPD][Br])	S15
¹ H NMR, ¹³ C NMR and ESI - MS spectra of compounds (C1-C29), (D1-D12) and (E1-E7).	S16- S119
Supporting References	

Materials

All the reactions were performed in an oven-dried round bottomed flask. Solvents, reagents and chemicals used for reactions in this paper were purchased from Sigma-aldrich/Spectrochem Pvt. Ltd. and used without any further purification unless it is specified otherwise. Double distilled water was used for the preparation of all aqueous solutions. Reactions were monitored by thin-layer chromatography (TLC). TLC was performed using E. Merck pre-coated silica plates (60F-254) with 0.25 mm thickness and visualized using short-wave UV light or developing agents.

Instrumentation

The structural validation of synthesised compounds was based on ¹H NMR, ¹³C NMR, mass spectroscopy. Nuclear magnetic resonance (NMR) was acquired at 400 MHz and 100 MHz for ¹H NMR and ¹³C NMR respectively using a JEOL JNM-ECS 400 spectrometer instrument with DMSO-d₆ and CDCl₃ as solvents. TMS was taken as the reference in NMR, and data were processed with its delta software. Coupling constant (J) is reported in Hertz and chemical shift values are reported in ppm for ¹H NMR, and multiplicities are as follows: s (singlet), d (doublet), dd (doublet), t (triplet) and m (multiplet). ¹³C CP-MAS was obtained through JEOL ECZR at 600 MHz. High resolution mass spectroscopy was generated by XEVO G2-XS QTOF spectrometer, Thermo Fisher Scientific Q Exactive spectrometer, Impact HD (Bruker) ESI QTOF high resolution mass spectrometer.

FT-IR experiments were carried out in the range of 400-4000 cm⁻¹ (Thermo Scientific; Model: INCOLET iS50) spectrometer. The powder X-ray diffraction (PXRD) studies were performed on Bruker diffractometer (D8 Discover) at room temperature and 2θ range 0°–100° (scanning rate = 2°/min, λ = 0.15406 nm, 40 kV, 40 mA). The thermal stability of ionic liquid was determined using a PerkinElmer Pyris diamond TGA/ differential thermal analyser. For obtaining the data, the sample was heated from room temperature to 1000 °C in N₂ atmosphere at a heating rate of 10 °C min⁻¹ and gas flow of 200 mL min⁻¹. Carl Zeiss, India (Jeol Japan Mode: JSM 6610LV) was used to obtain the SEM images.

Table S1. Comparison of the present work with previous literature for the esterification of acid to their corresponding esters.

Entry	Catalyst	Solvent	Temperature	Time	Yield(%)	Ref
1	Triphosgene	DCM	40°C	2h	95%	S ¹
2	TMSCl	-	RT	24h	89%	S ²
3	[Pd(cinnamyl)Cl] ₂ , IBnF·HBr	1,4-dioxane	100°C	36h	75%	S ³
4	Pd(PPh ₃) ₄		140°C	27h	64%	S ⁴
5	(CN-OA-m)	DMSO	-40°C, white LED	14h	92%	S ⁵
6	Sulfated Zirconium Catalyst	-	60 °C	6h	75%	S ⁶
7	silicotungstic acid; H ₄ SiW ₁₂ O ₄₀ ·nH ₂ O(ST A)	-	98 °C(reflux)	4h	91%	S ⁷
8	N,N'- diisopropylcarbodiimi de	Water	RT	4h	92%	S ⁸
9	XtalFluor-E	TFE in CH ₂ Cl ₂	RT	16h	84%	S ⁹
10	[Ir(cod)Cl] ₂	CH ₂ Cl ₂	RT	12h	88%	S ¹⁰
11	PPh ₃ /I ₂	Acetonitrile	Reflux, MW	30 mins	93%	S ¹¹
12	Silica-IL	Cyclohexane	Reflux, 93 °C	3h	86%	S ¹²
13	[Bmim][dca]	-	60 °C	120h	65%	S ¹³
14	IL	-	85 °C	3h	91.5%	S ¹⁴
15	Chitosan-IL₆	-	RT	30	96%	P.

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Table S2. Optimization of reaction conditions for the amount of catalyst using benzoic acid **A1** and ethanol **B1** as the model substrate.^a

Entry	Catalyst (mg)	Yield(%)^b
1	Chitosan-IL ₆ (5mg)	92
2	Chitosan-IL₆ (10mg)	96
3	Chitosan-IL ₆ (15mg)	96
4	Chitosan-IL ₆ (20mg)	96

^aReaction conditions: Benzoic acid A1 (1 mmol), ethanol B1 (1 mmol), catalyst, room temperature for 30 mins. ^bIsolated yield.

Gram scale synthesis.

For the practical application of the produced esters, we synthesised a few value-added compounds on a gram scale (Figure S1) ^{15,16}. The reactions were conducted at a scale of 100 mmol using 50 mg of the catalyst. The reactions proceeded easily with yields of 85% for diisopropyl azodicarboxylate over a period of 6 hours, and 86% for methyl nicotinate over a period of 5 hours. Considering esters of amino acids, methyl cysteinate produced a 93% yield over the course of 8 hours, whereas glucose pentaacetate produced 92% yield over the course of 12 hours. These results demonstrate the proposed protocol's practicality and operational simplicity even at higher reaction scales.

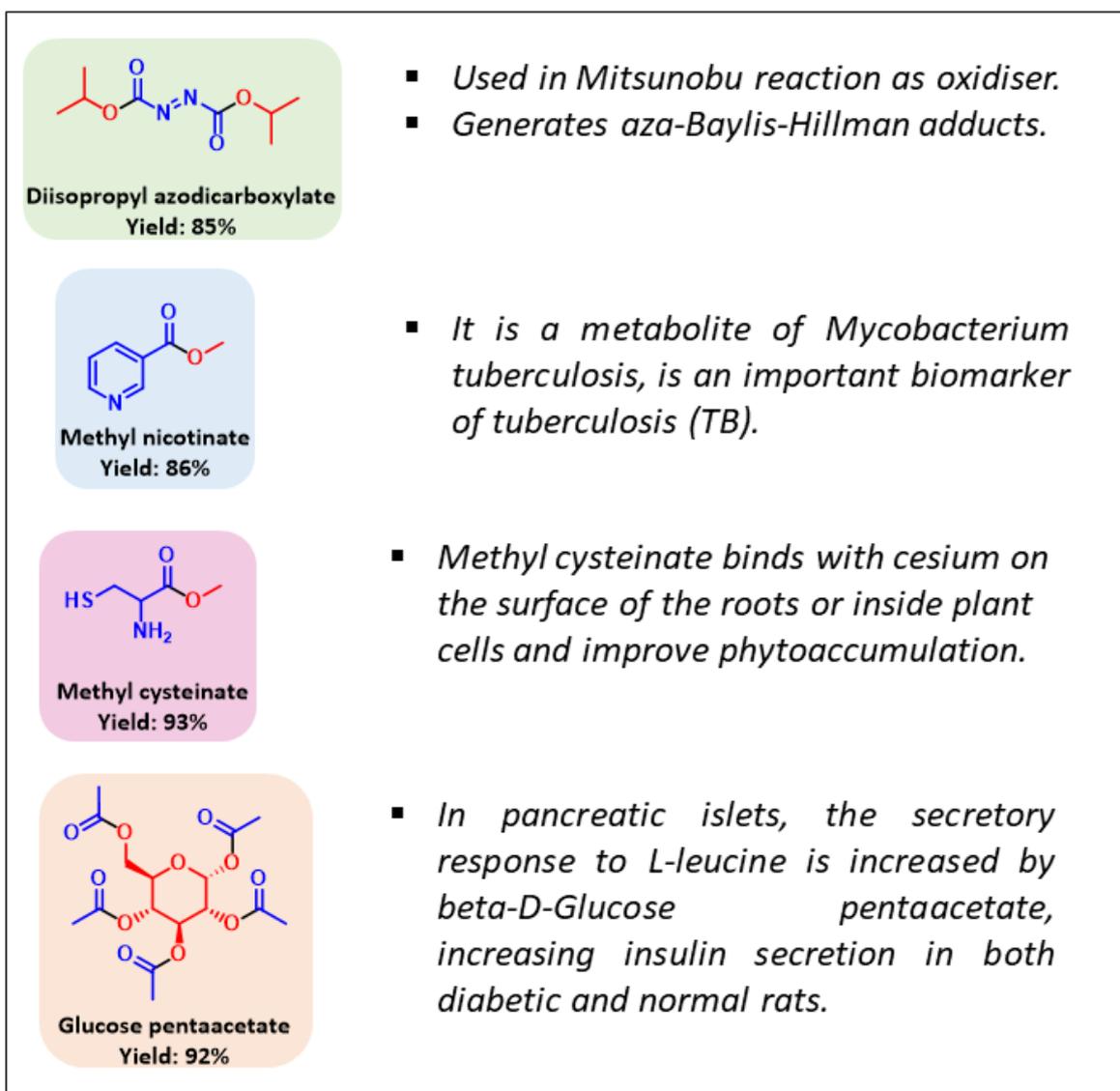


Figure S1. Practical application of a few value-added esters synthesized in gram scale.

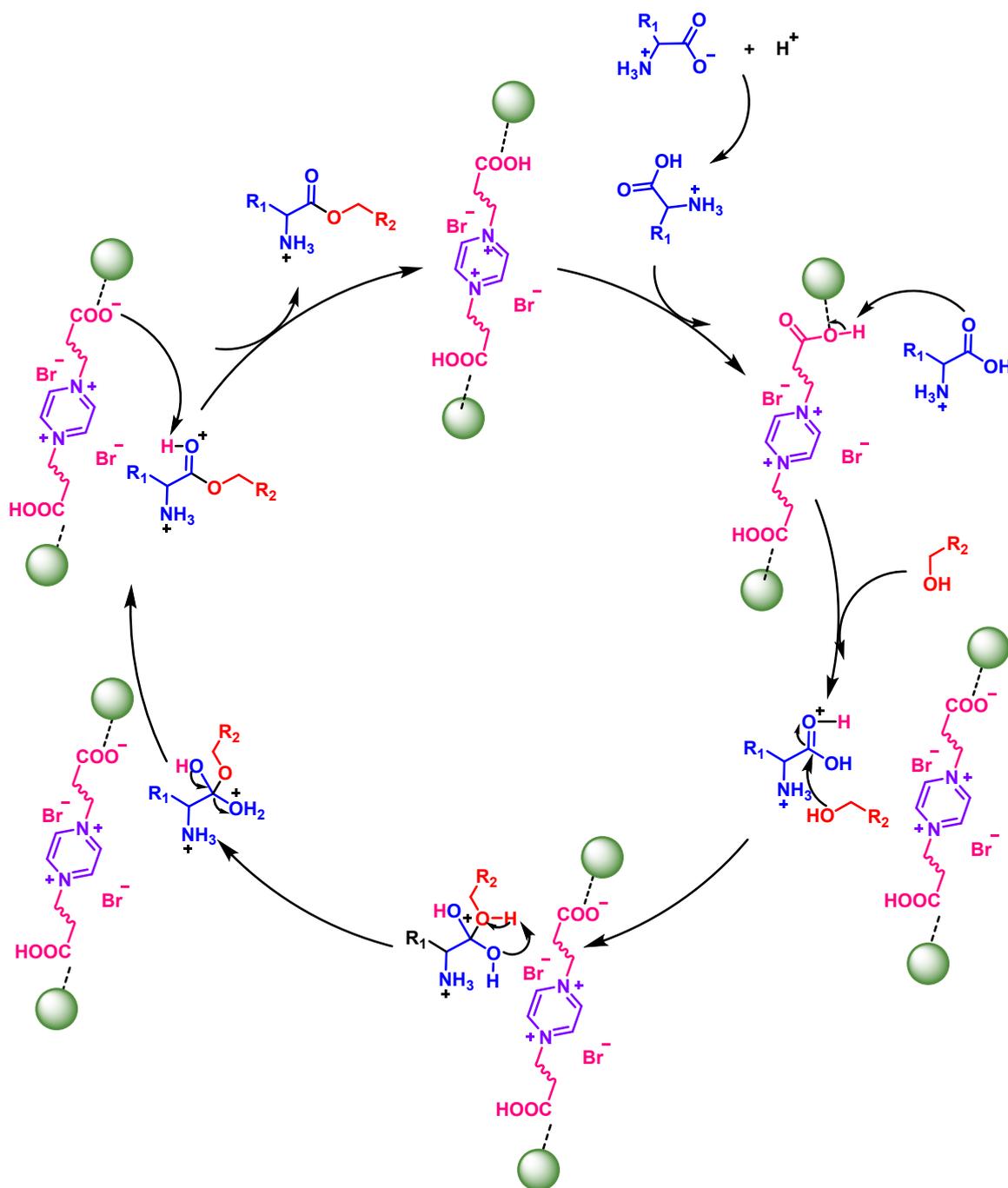


Figure S2. Plausible mechanism for Chitosan-IL₆ catalysed selective esterification of glycine amino acid and ethanol to ethyl glycinate.

To capture an insight into the reaction route of Chitosan-IL₆ catalysed esterification of amino acid we hereby present a feasible mechanism (Figure S2) from theoretical dimensions. We here consider glycine D1 and ethanol B to obtain ethyl glycinate E1. At the starting acetic acid protonates the oxygen of glycine which is in zwitter ion form. The oxygen of carbonyl then extracts a proton from the carboxylic group of Chitosan-IL₆, which is followed by the interaction of ethanolic OH with the carbonyl C of glycine. This leads to proton exchange between glycine and ethanol molecules and followed by the elimination of water molecule.

Finally, the desorption of glycinate from Chitosan-IL₆ results in the synthesis of the desired product and continuing the active use of the catalyst in further cycles.

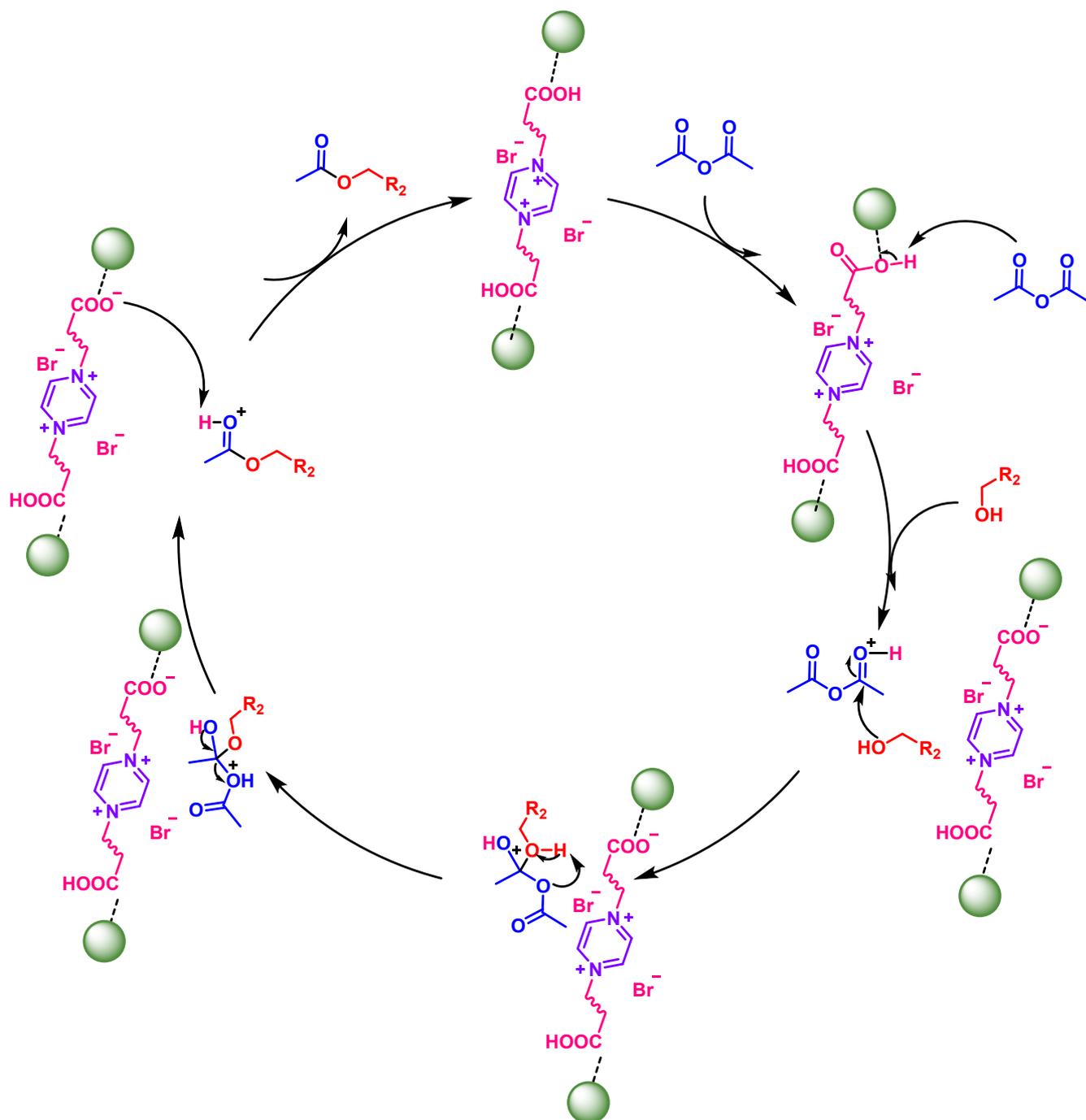


Figure S3. Plausible mechanism for Chitosan-IL₆ catalysed selective esterification of acetic anhydride and carbohydrate to carbohydrate esters.

Figure S3 outlines a plausible mechanistic pathway for the Chitosan-IL₆ catalysed esterification of carbohydrates based on the discussed controlled reactions. The oxygen of

one of the carbonyl C interacts with the ionic liquid first, followed by the interaction of OH of carbohydrate. This causes a proton transfer from carbohydrate OH to acetic anhydride's oxygen. This is followed by the loss of acetic acid and final desorption of the ester moiety from the catalyst to obtain the desired product.

Table S3. Recyclability test of Chitosan IL₆ catalyzed esterification of benzoic acid and ethanol to ethyl benzoate **C1**.^a

Catalytic run	Yield(%) ^b
1	96
2	96
3	96
4	96
5	96
6	95
7	95
8	94
9	93
10	93

^aReaction conditions: Carboxylic acid (1 mmol), alcohol (1 mmol), Chitosan-IL₆ (10 mg) under neat conditions at RT for appropriate time. ^bIsolated yield.

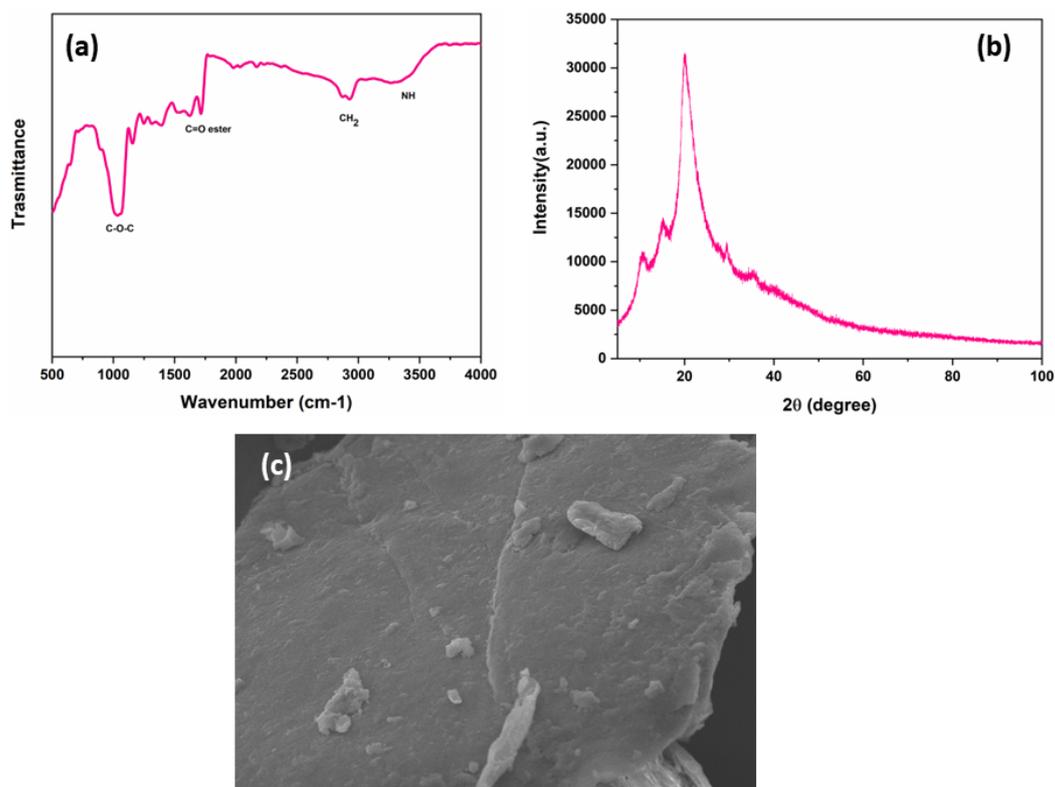


Figure S4. IR (a), XRD (b) and SEM(c) image of reused catalyst.

Determination of acidity of Chitosan-IL₆

The density of total acidic sites on Chitosan-IL₆ was calculated by back acid–base titration¹⁷. First, 100 mg of Chitosan-IL₆ was added to 10 mL of freshly prepared 0.05 N NaOH solution and the resulting mixture was stirred for 3 h at RT. Subsequently, the mixture was centrifuged at 8000 rpm for 2 min and washed two times with double distilled water. The filtrate containing excess NaOH solution was then back titrated with freshly prepared 0.1 N HCl solution till neutralization point, monitored by using phenolphthalein indicator to evaluate the total concentration of acidic sites in Chitosan-IL₆.

Calculation of acidic strength of Chitosan-IL₆

It was found that 1.3 mL of 0.1 N HCl was required to reach the neutralization point.

$$V_{\text{NaOH}} \times S_{\text{NaOH}} = V_{\text{HCl}} \times S_{\text{HCl}}$$

$$V_{\text{NaOH}} \times 0.05 = 1.3 \times 0.1$$

$$V_{\text{NaOH}} = 2.6 \text{ mL}$$

Therefore, the volume of NaOH required to neutralize the acidic sites in Chitosan-IL₆ = (10-2.6) mL = 7.4 mL.

$$V_{\text{NaOH}} \times S_{\text{NaOH}} = V_{\text{Chitosan-IL}_6} \times S_{\text{Chitosan-IL}_6}$$

$$7.4 \times 0.05 = 10 \times S_{\text{Chitosan-IL}_6}$$

$$S_{\text{Chitosan-IL}_6} = 0.037 \text{ N}$$

The equivalent weight of carboxylic acid group (-COOH) is 45.

That is, 1000 mL of 1 N Chitosan-IL₆ would contain 45 g free carboxylic acid sites.

So, 10 mL of 0.037 N Chitosan-IL₆ solution contains 0.01665 g free carboxylic acid sites.

0.01665 g free carboxylic acid sites = 0.37 mmol free carboxylic acid sites.

100 mg sample of Chitosan-IL₆ contains 0.37 mmol free carboxylic acid.

Thus, 1000 mg sample of Chitosan-IL₆ would contain 3.7 mmol free carboxylic acid sites.

That is, total acid sites in Chitosan-IL₆ = 3.7 mmol g⁻¹.

Calculation of TOF of Chitosan-IL₆

1000 mg (1g) of Chitosan-IL₆ contains 3.7 mmol acid sites

To determine TOF¹⁸ of Chitosan-IL₆ we considered the model reaction of benzoic acid **A1** (1mmol) and ethanol **B1** (1mmol) in the presence of 10 mg of catalyst (Chitosan-IL₆) at RT to yield ethyl benzoate **C1**.

As the yield of this product is 96%,

The turn over number (**TON**) of C1 is = $\frac{\text{mmol of product}}{\text{mmol of active sites in catalyst}}$

1g Chitosan-IL₆ has 3.7 mmol active sites, so 10 mg of Chitosan-IL₆ has = 0.037 mmol active sites.

$$\text{TON} = \frac{0.96}{0.037} = 25.94 \text{ (as the yield of product is 96\%, mmol of product} = 0.96)$$

And turn over frequency (TOF) of C1 is = $\frac{TON}{Reaction\ time} = \frac{25.94}{30/60}h^{-1} = 51.88\ h^{-1}$

Table S4. TOF values Chitosan-IL₆ for the products C1-C29^a

Entry	Product Code	Yield (%) ^b	Time (h)	TOF (h ⁻¹)
1	C1	96	0.5	51.88
2	C2	95	0.58	44.26
3	C3	94	0.58	43.80
4	C4	92	0.58	42.87
5	C5	98	0.42	63.06
6	C6	97	0.42	62.42
7	C7	97	0.42	62.42
8	C8	99	0.33	81.08
9	C9	98	0.33	80.26
10	C10	96	0.5	51.88
11	C11	97	0.42	62.42
12	C12	97	0.42	62.42
13	C13	89	0.66	36.44
14	C14	90	0.66	36.85
15	C15	93	0.66	38.08
16	C16	85	0.83	27.68
17	C17	89	0.83	28.98
18	C18	90	0.75	32.43

19	C19	92	0.75	33.15
20	C20	84	0.83	27.35
21	C21	90	0.83	29.31
22	C22	89	0.83	28.98
23	C23	88	0.83	28.65
24	C24	86	0.83	28.00
25	C25	86	0.83	28.00
26	C26	87	0.83	28.33
27	C27	90	0.83	29.31
28	C28	85	0.83	27.68
29	C29	89	0.83	28.98

^aReaction conditions: Carboxylic acid (1 mmol), alcohol (1mmol), Chitosan-IL₆ (10 mg) under neat conditions at RT for appropriate time. ^bIsolated yield

Table S5. TOF values Chitosan-IL₆ for the products **D1-D12**^a

Entry	Product Code	Yield (%) ^b	Time (h)	TOF (h ⁻¹)
1	D1	94	3.5	7.26
2	D2	96	3	8.65
3	D3	94	3.5	7.26
4	D4	96	3	8.65
5	D5	95	3.5	7.33
6	D6	95	3.5	7.33
7	D7	96	3.5	7.41

8	D8	90	3	8.11
9	D9	99	2	13.38
10	D10	98	2	13.24
11	D11	98	2	13.24
12	D12	95	3	8.56

^aReaction conditions: Amino acid (1 mmol), alcohol (1mmol), Chitosan-IL₆ (10 mg), acetic acid (3 drops) under neat conditions at RT for appropriate time. ^bIsolated yield.

Table S6. TOF values Chitosan-IL₆ for the products **E1-E7**^a

Entry	Product Code	Yield (%) ^b	Time (h)	TOF (h ⁻¹)
1	E1	98	1	26.49
2	E2	96	1	25.95
3	E3	96	1	25.95
4	E4	95	1	25.67
5	E5	93	0.83	30.28
6	E6	92	0.83	29.96
7	E7	90	2	12.16

^aReaction conditions: Acetic anhydride (4-5 mmol), carbohydrate (1 mmol), Chitosan-IL₆ (10 mg) under neat conditions at RT for appropriate time. ^bIsolated yield.

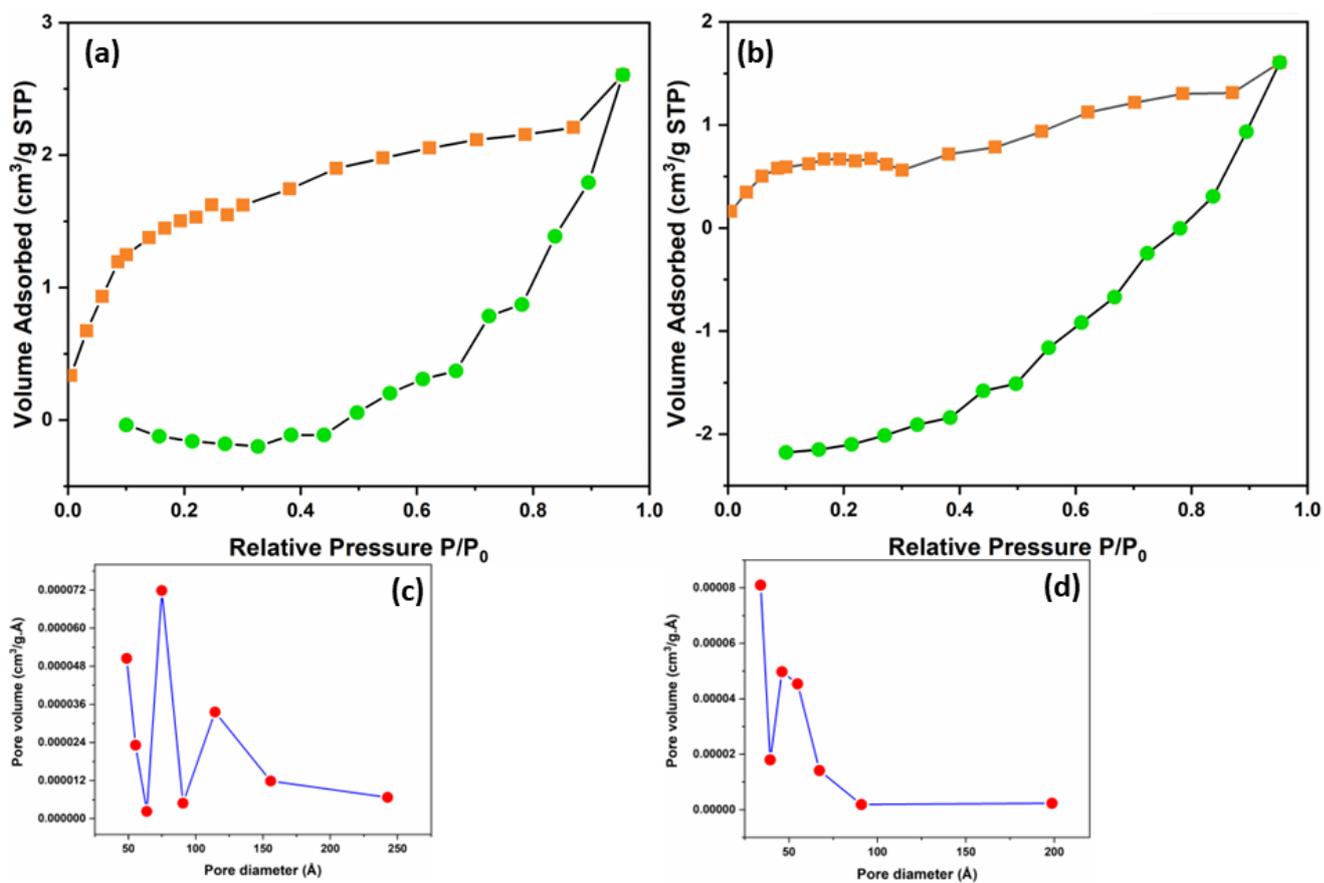
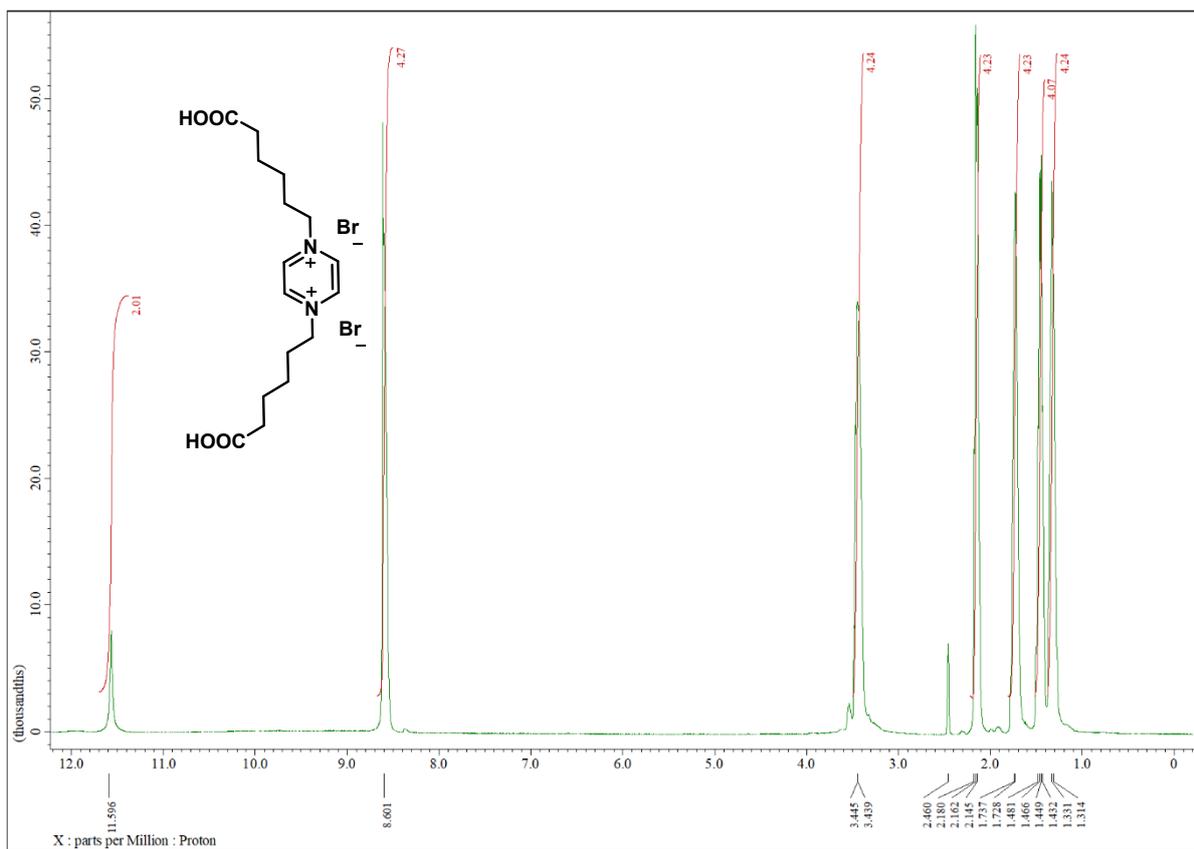


Figure S5. N₂ adsorption/desorption isotherm of (a) Chitosan and (b) Chitosan IL₆ and Pore size distribution of (c) Chitosan and (d) Chitosan IL₆

¹H NMR, ¹³C NMR and MS spectra of all compounds.



FigureS6. ^1H NMR spectrum of 1,4-bis(5-carboxypentyl)pyrazine-1,4-dium ([BCPPD][Br])

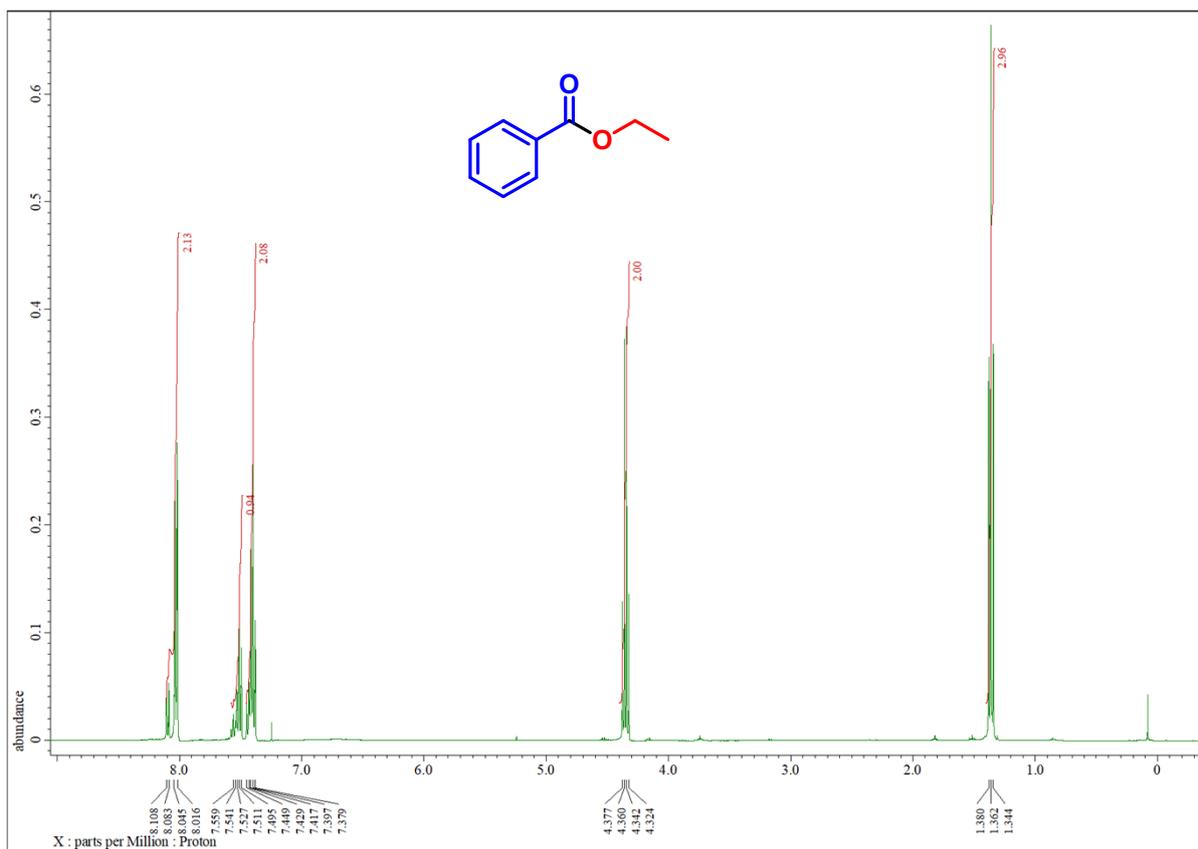


Figure S7. ^1H NMR spectrum of Ethyl benzoate C1.

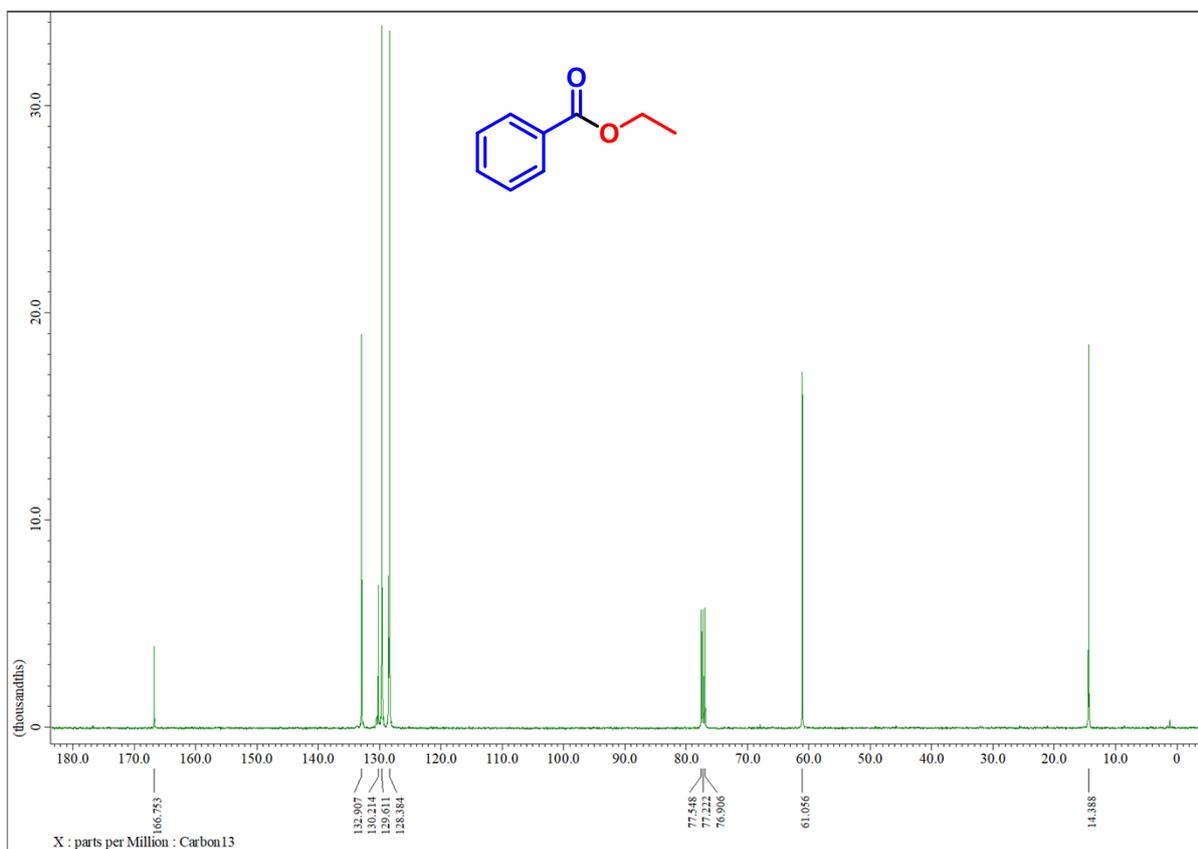


Figure S8. ^{13}C NMR spectrum of Ethyl benzoate **C1**.

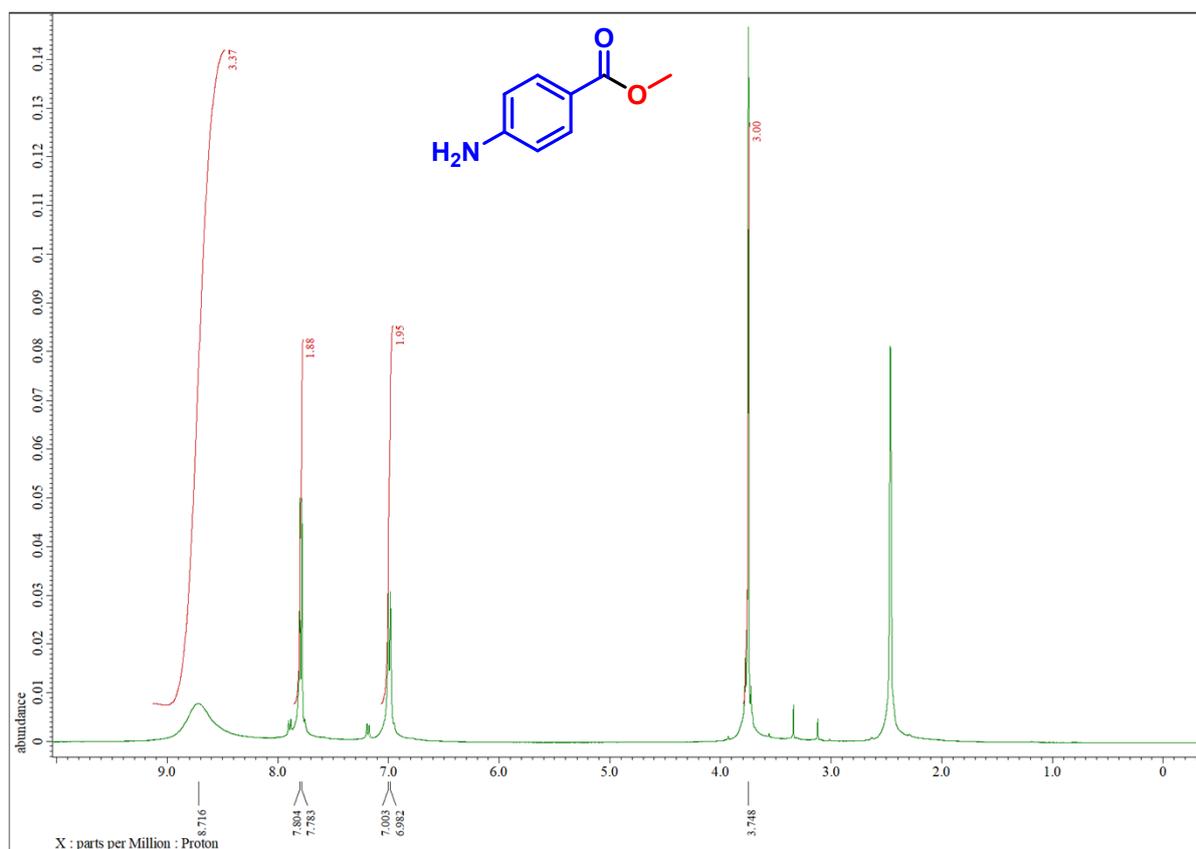


Figure S9. ^1H NMR spectrum of Methyl 4-aminobenzoate **C2**.

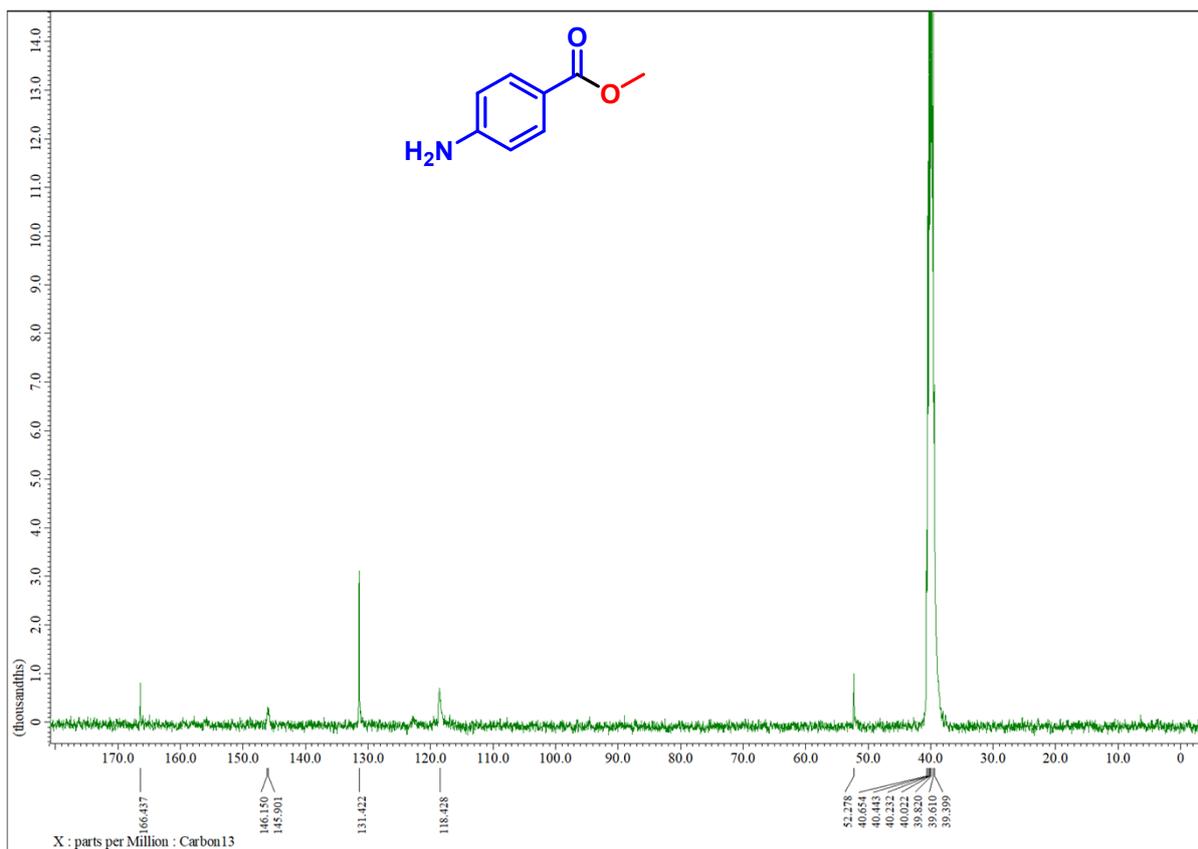


Figure S10. ^{13}C NMR spectrum of Methyl 4-aminobenzoate C2.

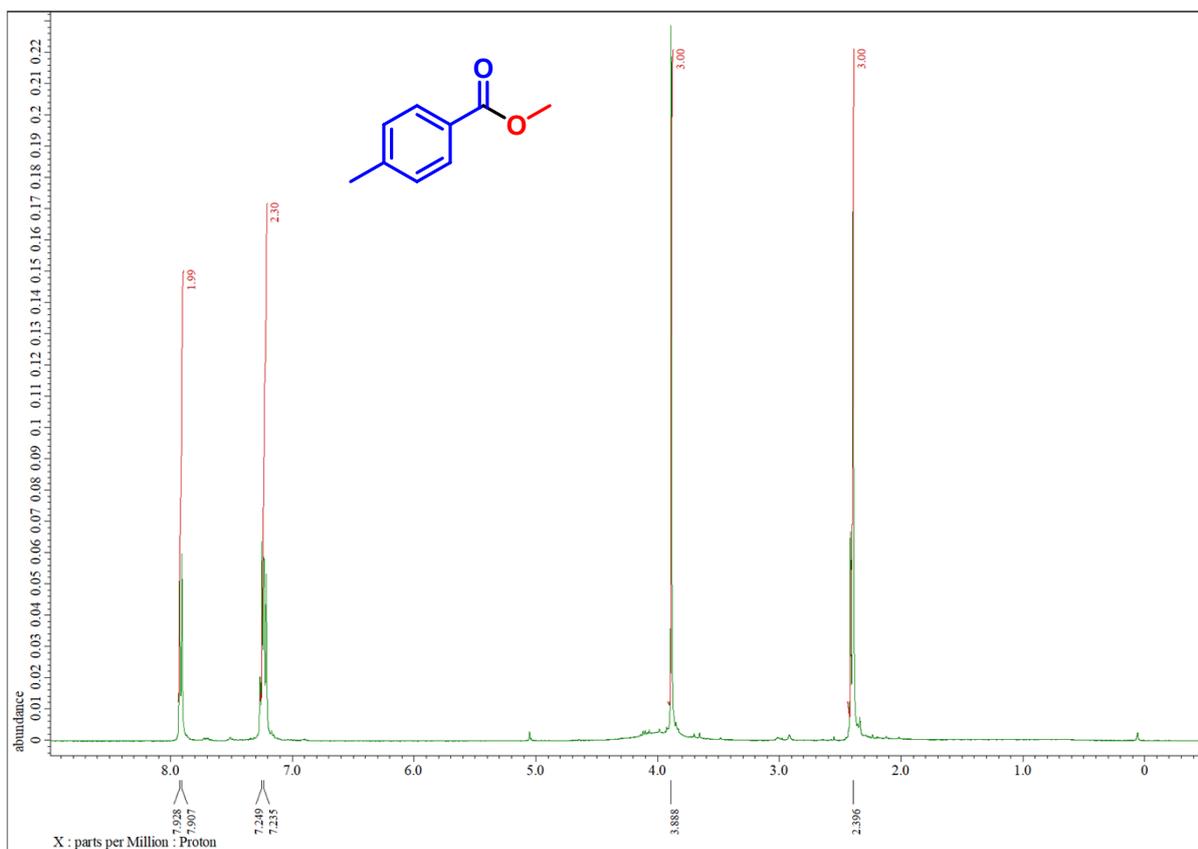


Figure S11. ^1H NMR spectrum of Methyl 4-methylbenzoate **C3**.

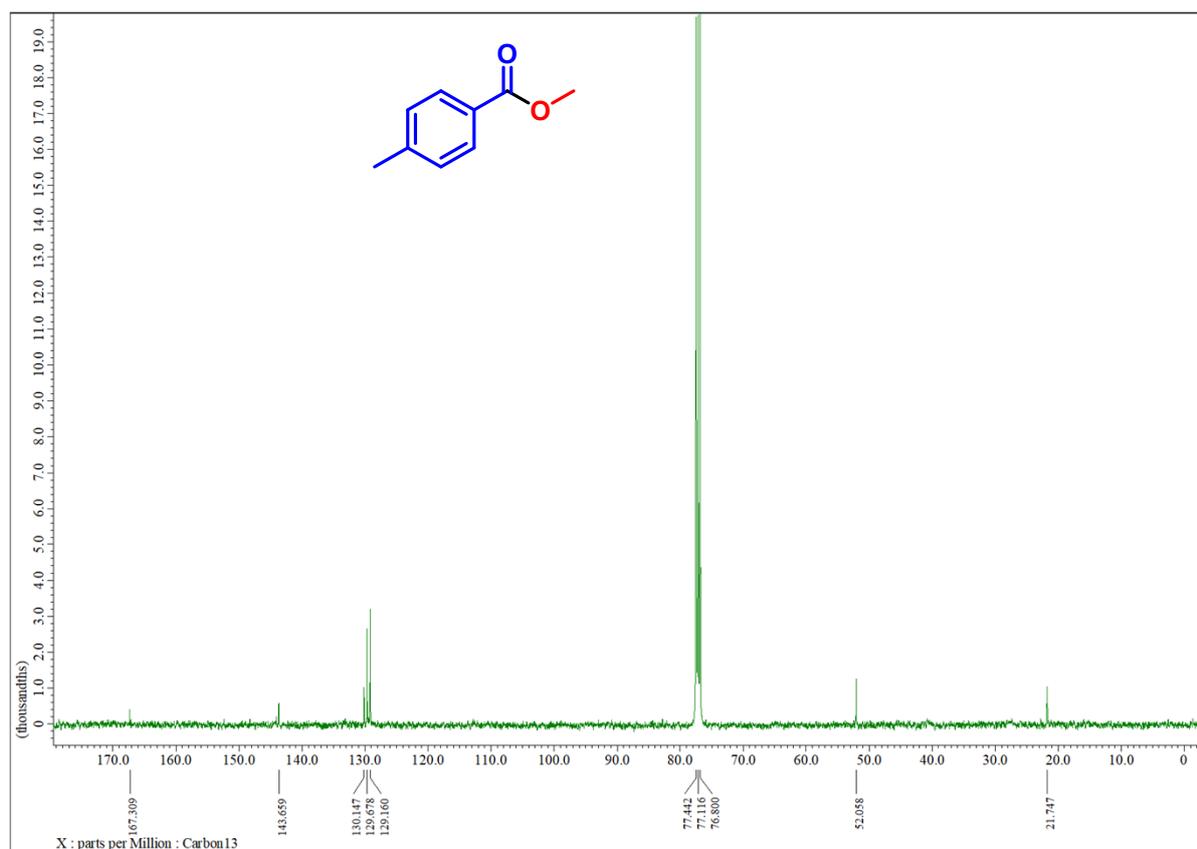


Figure S12. ^{13}C NMR spectrum of Methyl 4-methylbenzoate **C3**.

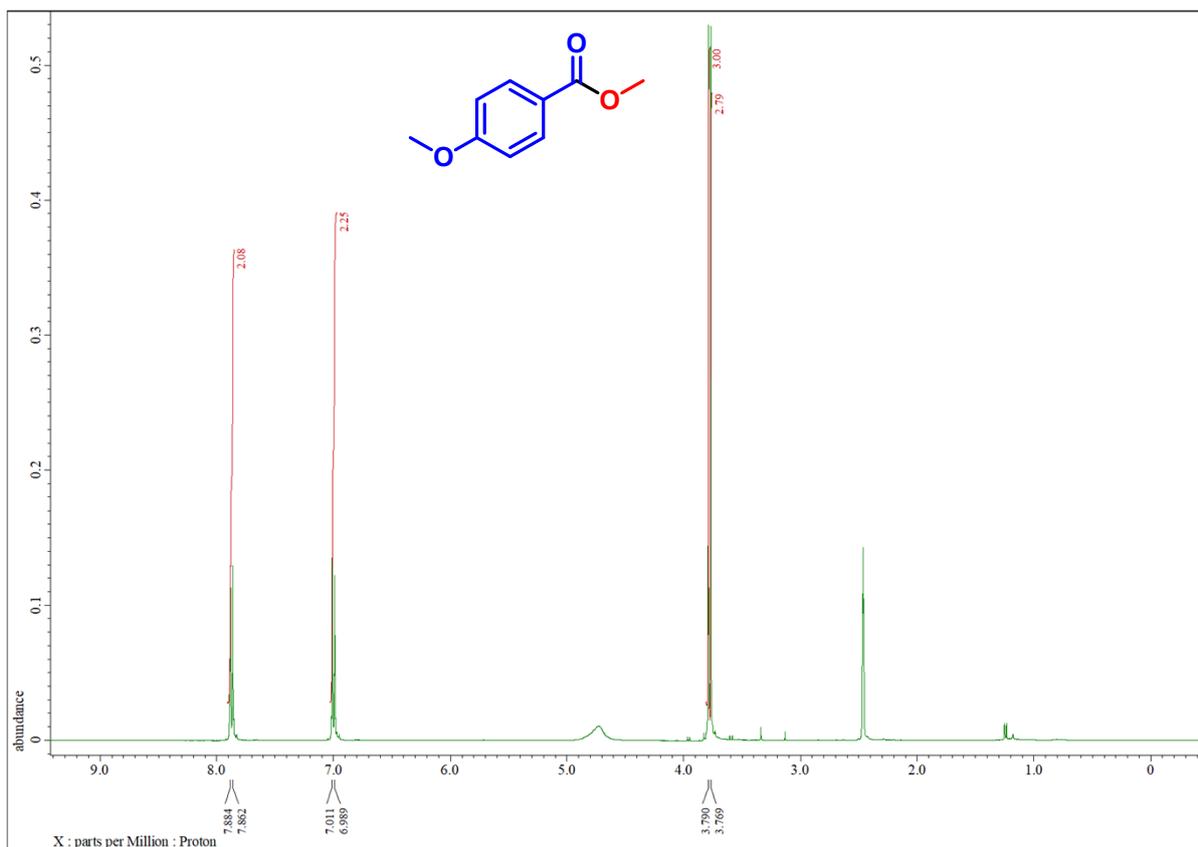


Figure S13. ^1H NMR spectrum of Methyl 4-methoxybenzoate C4.

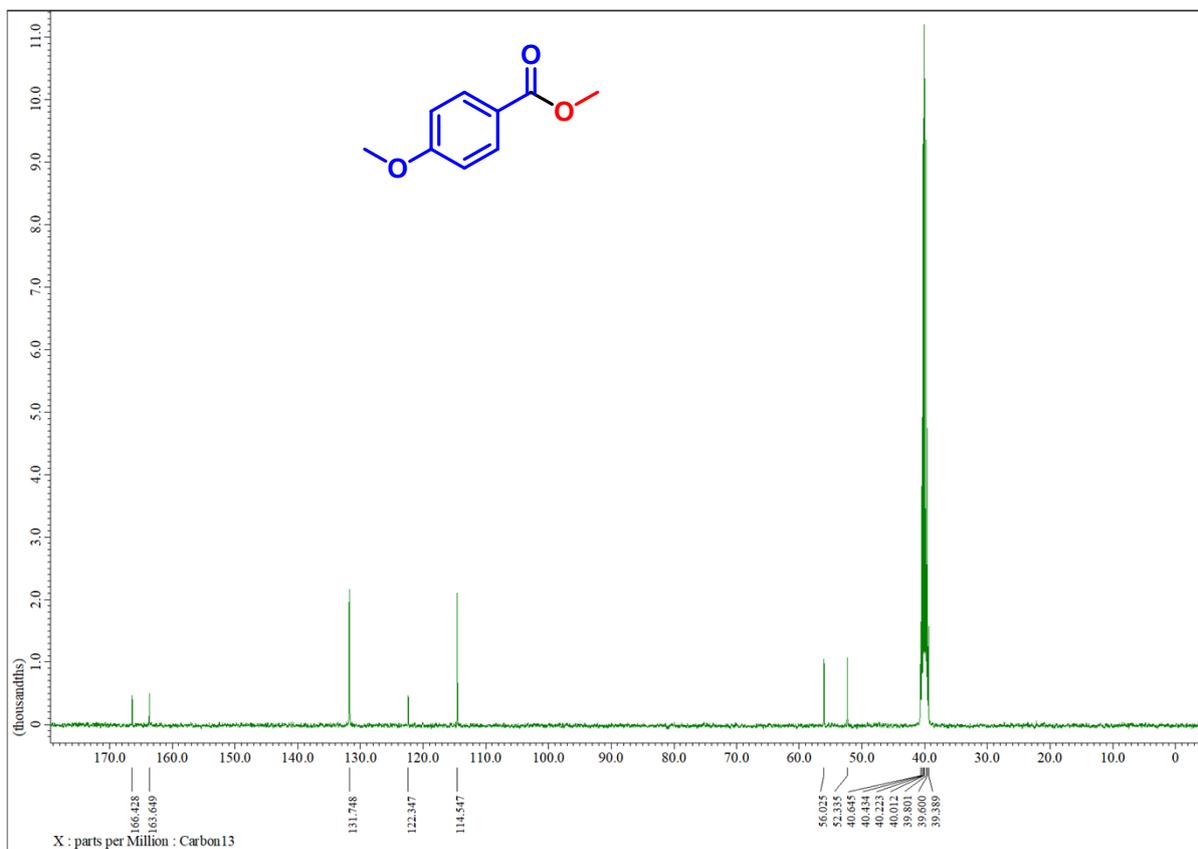


Figure S14. ^{13}C NMR spectrum of Methyl 4-methoxybenzoate C4.

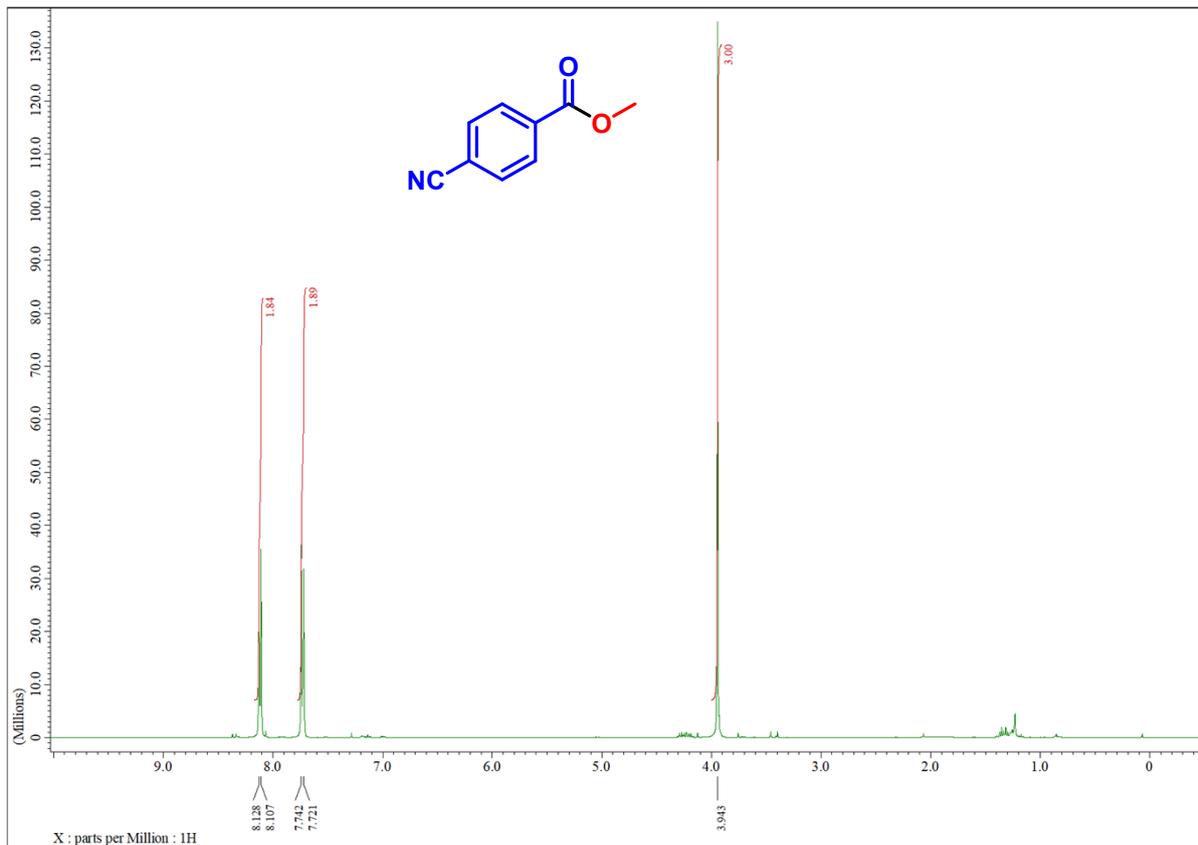


Figure S15. ¹H NMR spectrum of Methyl 4-cyanobenzoate **C5**.

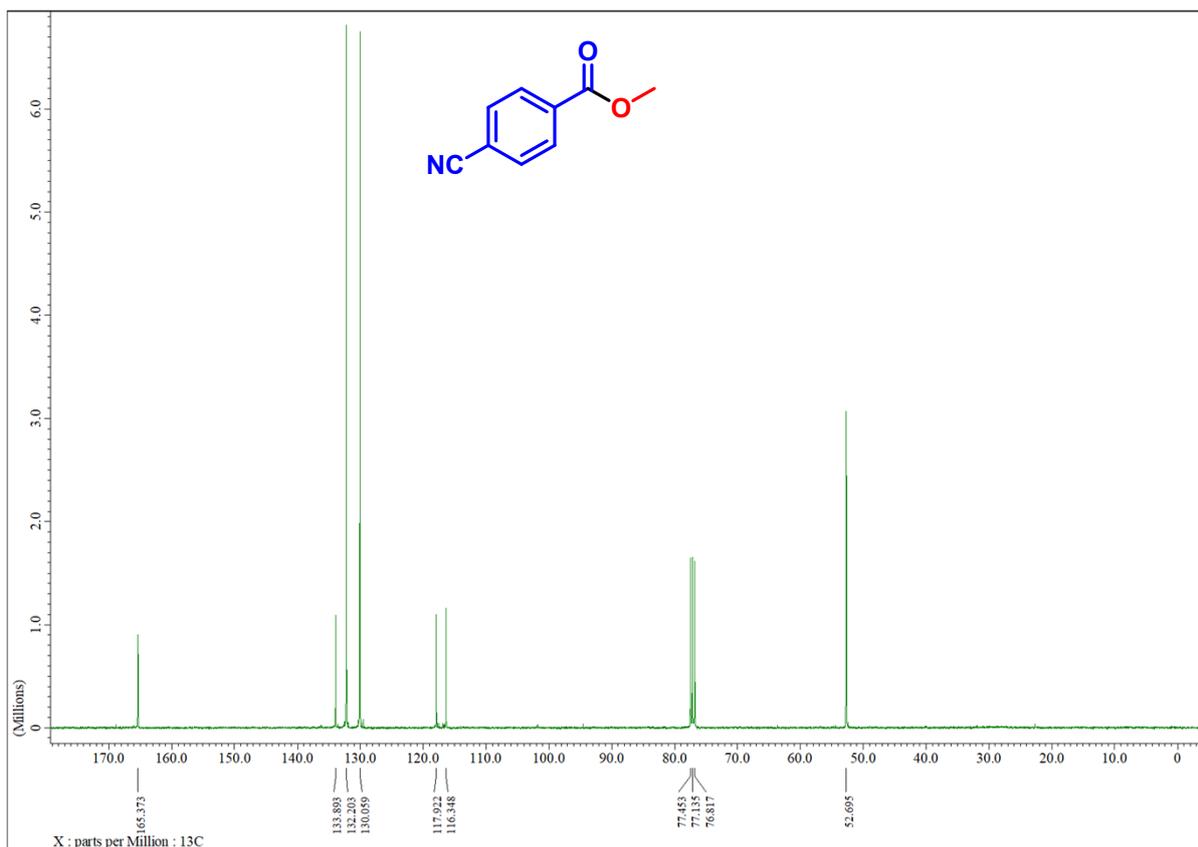


Figure S16. ¹³C NMR spectrum of Methyl 4-cyanobenzoate C5.

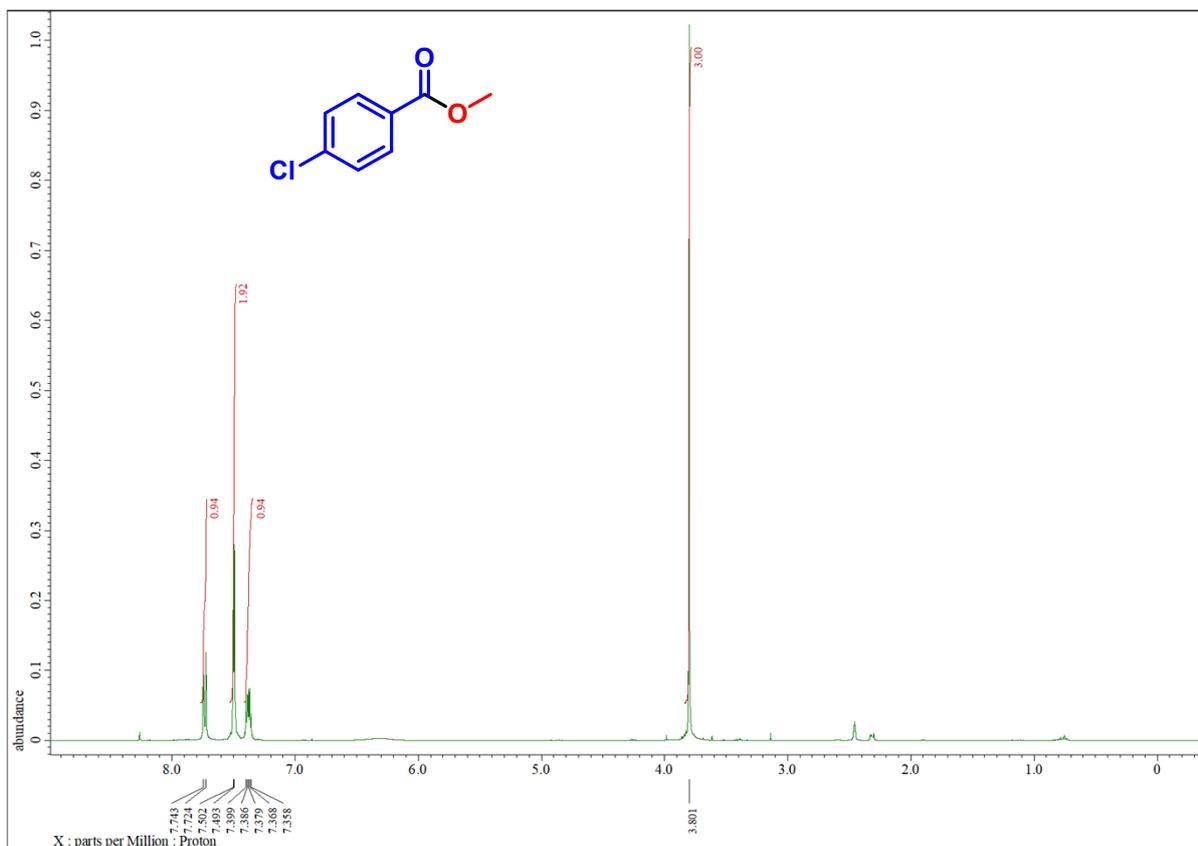


Figure S17. ^1H NMR spectrum of Methyl 4-chlorobenzoate **C6**.

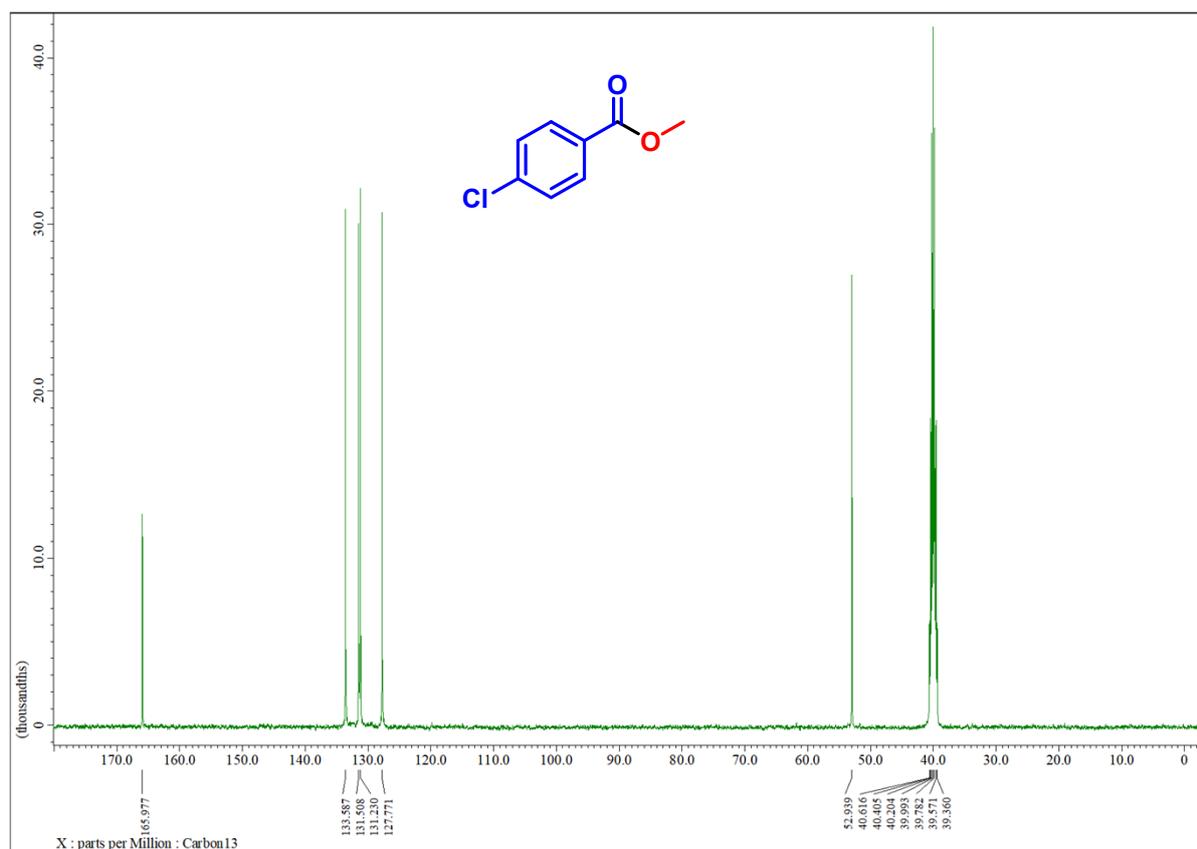


Figure S18. ^{13}C NMR spectrum of Methyl 4-chlorobenzoate **C6**.

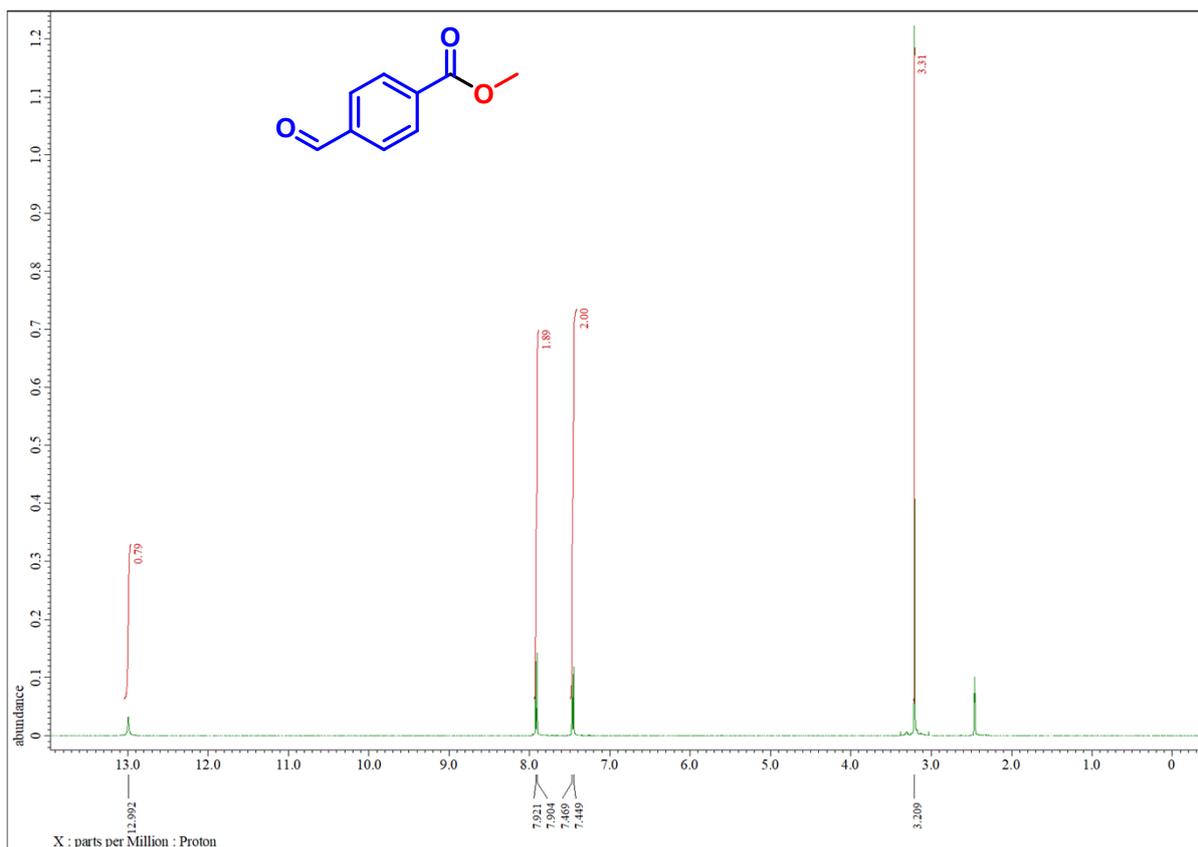


Figure S19. ^1H NMR spectrum of Methyl 4-formylbenzoate **C7**.

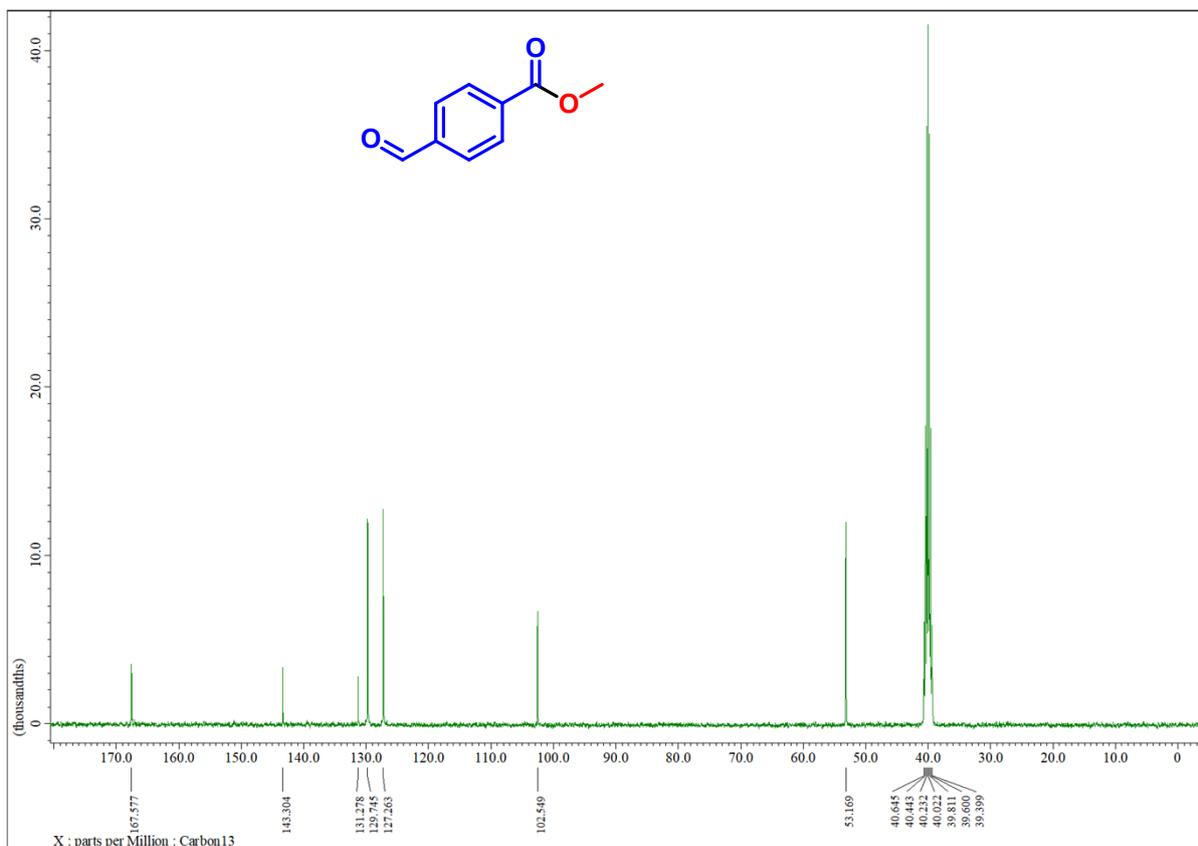


Figure S20. ^{13}C NMR spectrum of Methyl 4-formylbenzoate **C7**.

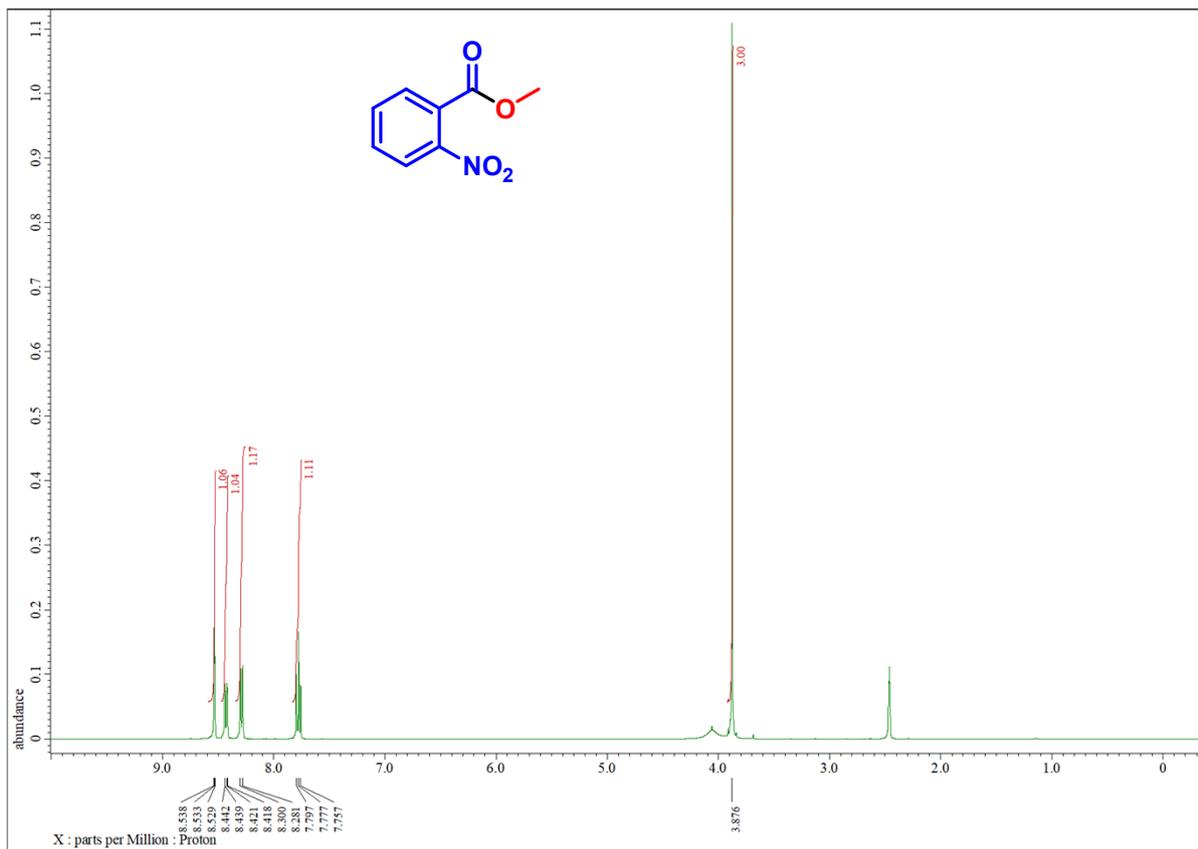


Figure S21. ¹H NMR spectrum of Methyl 2-nitrobenzoate **C8**.

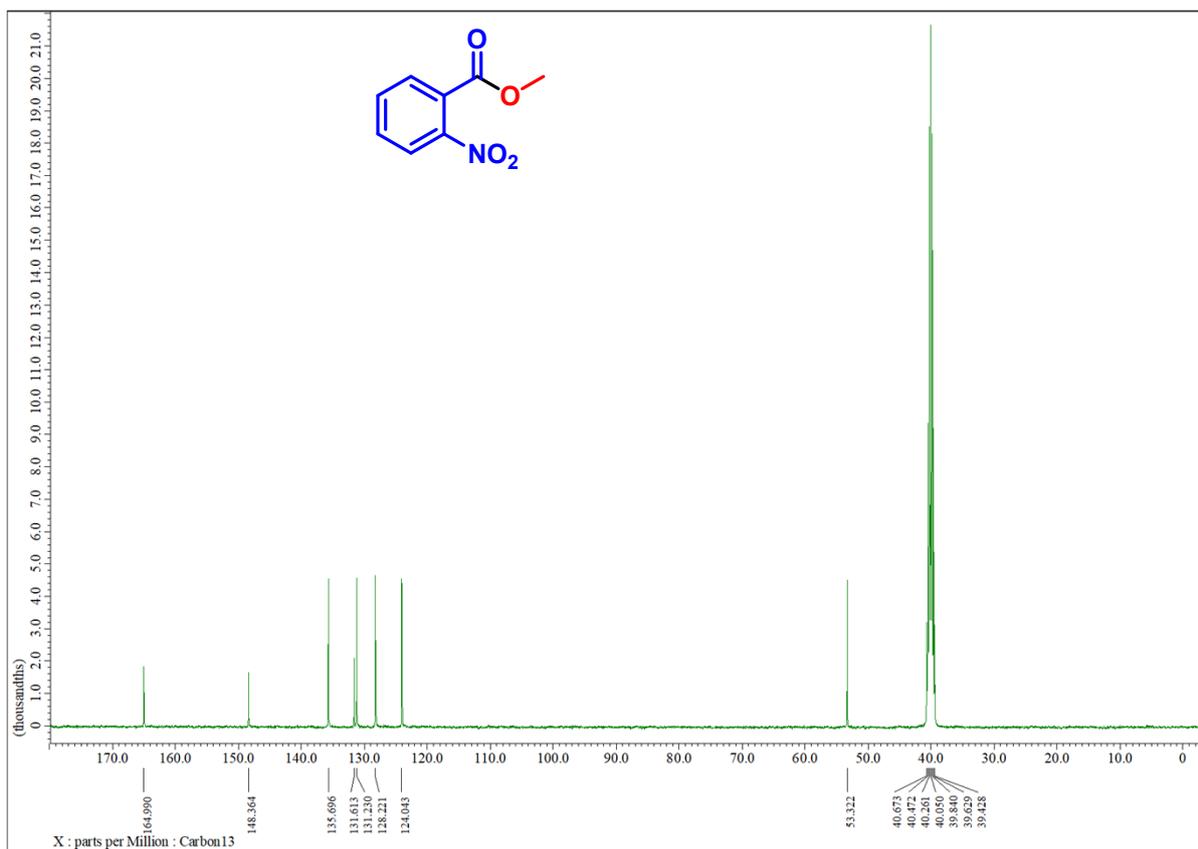


Figure S22. ¹³C NMR spectrum of Methyl 2-nitrobenzoate **C8**.

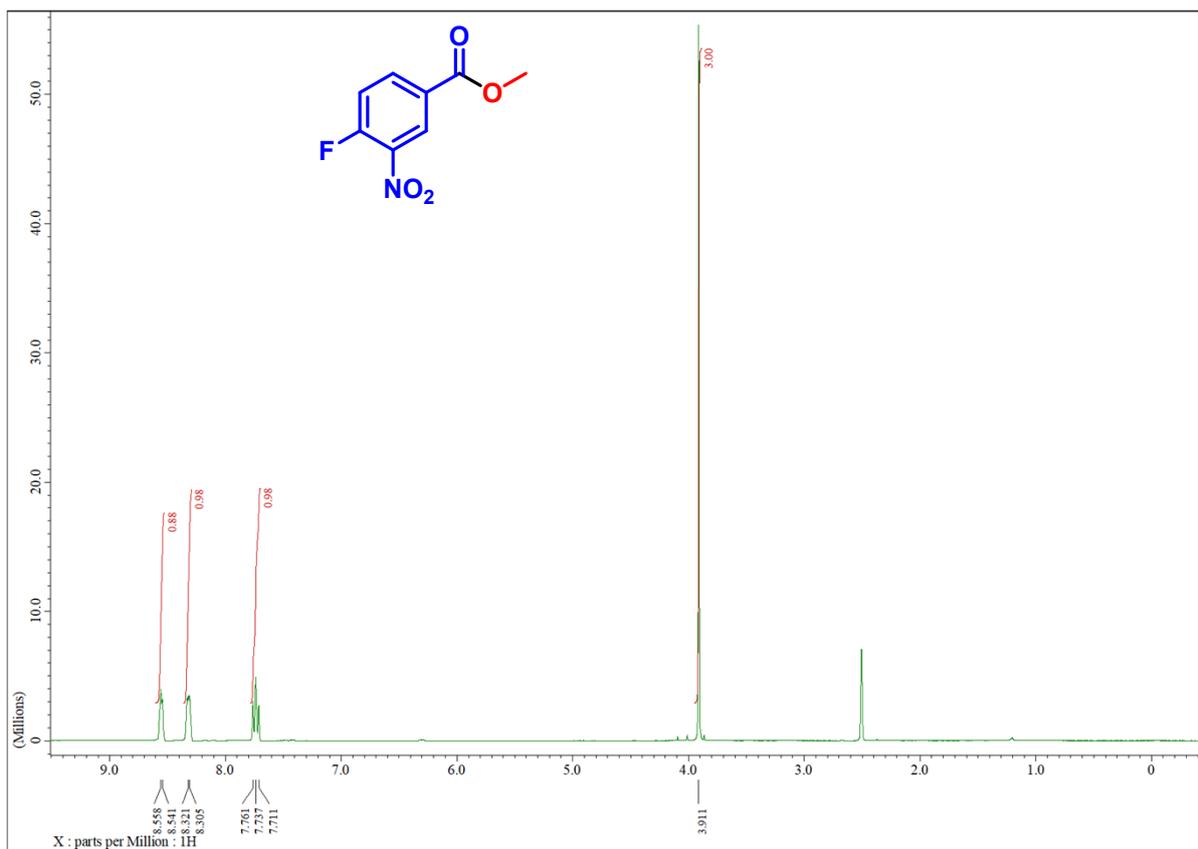


Figure S23. ¹H NMR spectrum of Methyl 4-fluoro-3-nitrobenzoate **C9**.

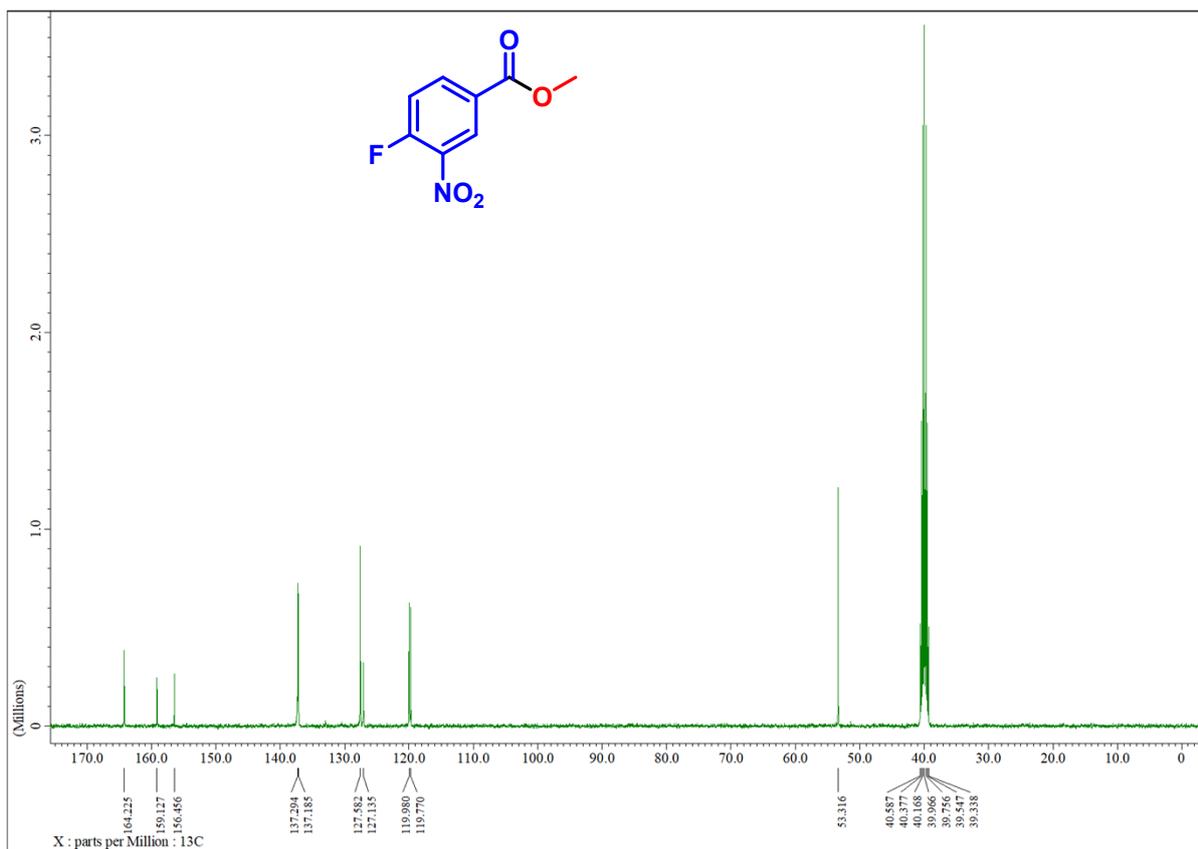


Figure S24. ^{13}C NMR spectrum of Methyl 4-fluoro-3-nitrobenzoate **C9**.

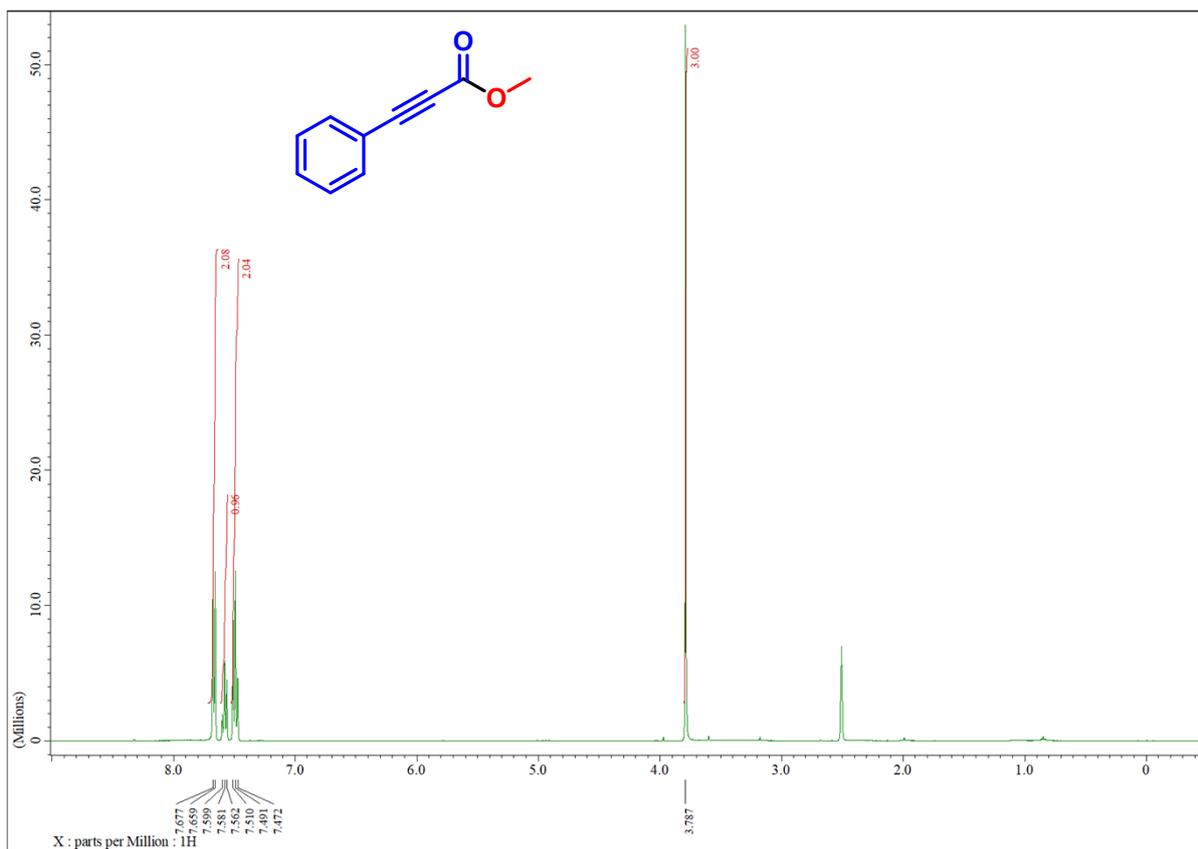


Figure S25. ¹H NMR spectrum of Methyl 3-phenylpropiolate **C10**.

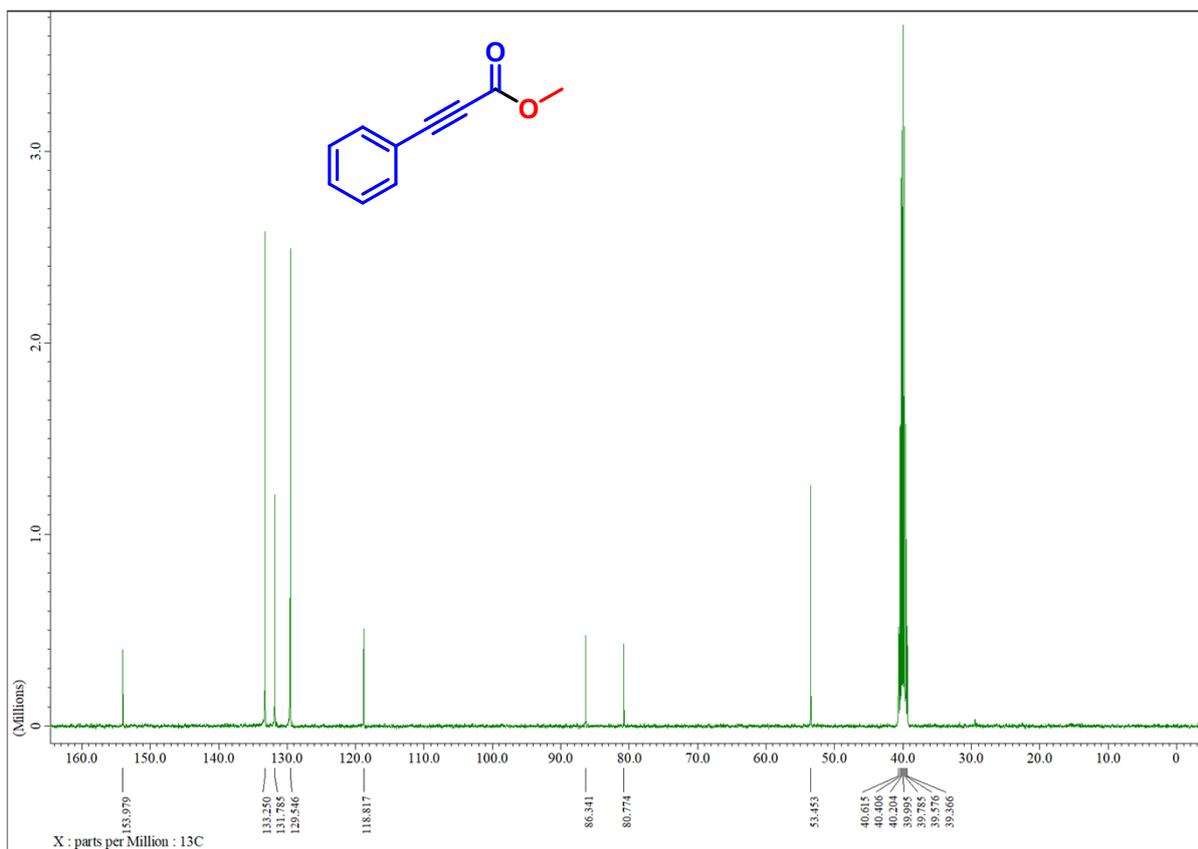


Figure S26. ^{13}C NMR spectrum of Methyl 3-phenylpropiolate C10.

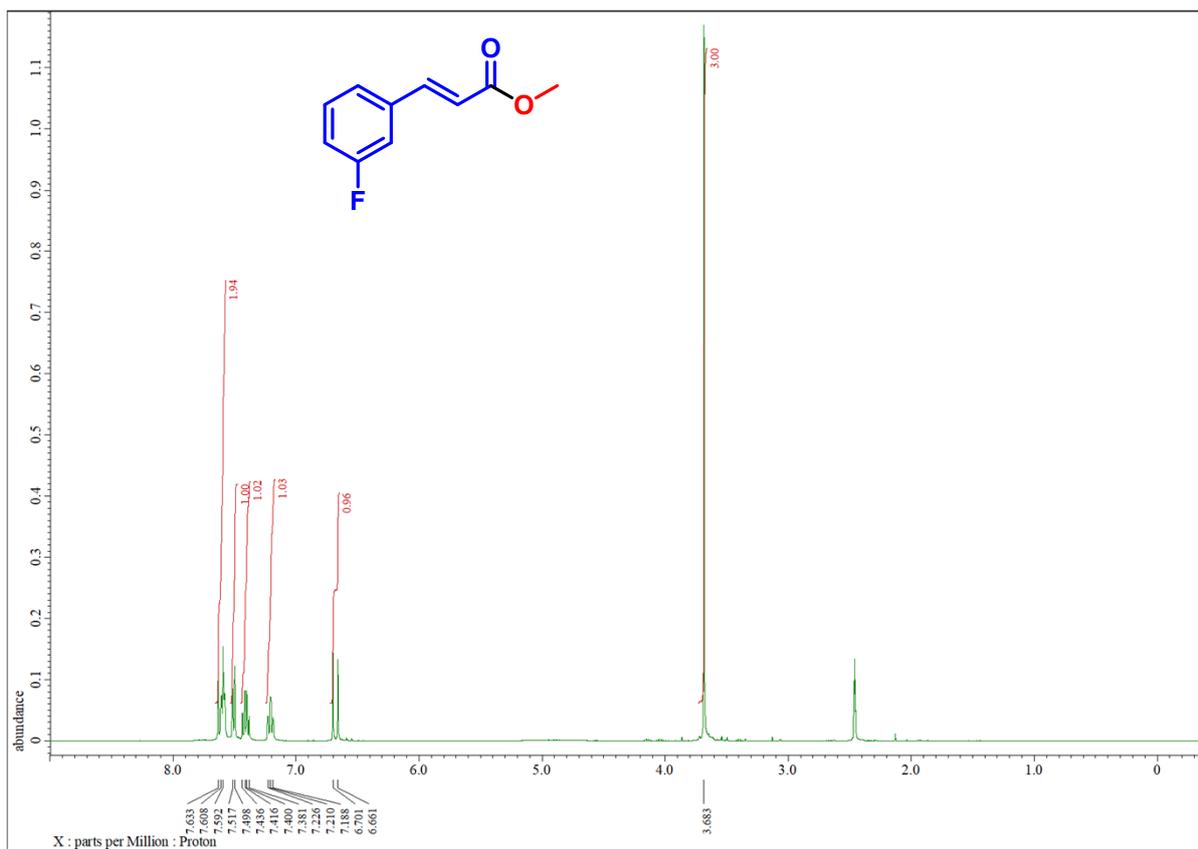


Figure S27. ¹H NMR spectrum of Methyl (E)-3-(3-fluorophenyl)acrylate **C11**.

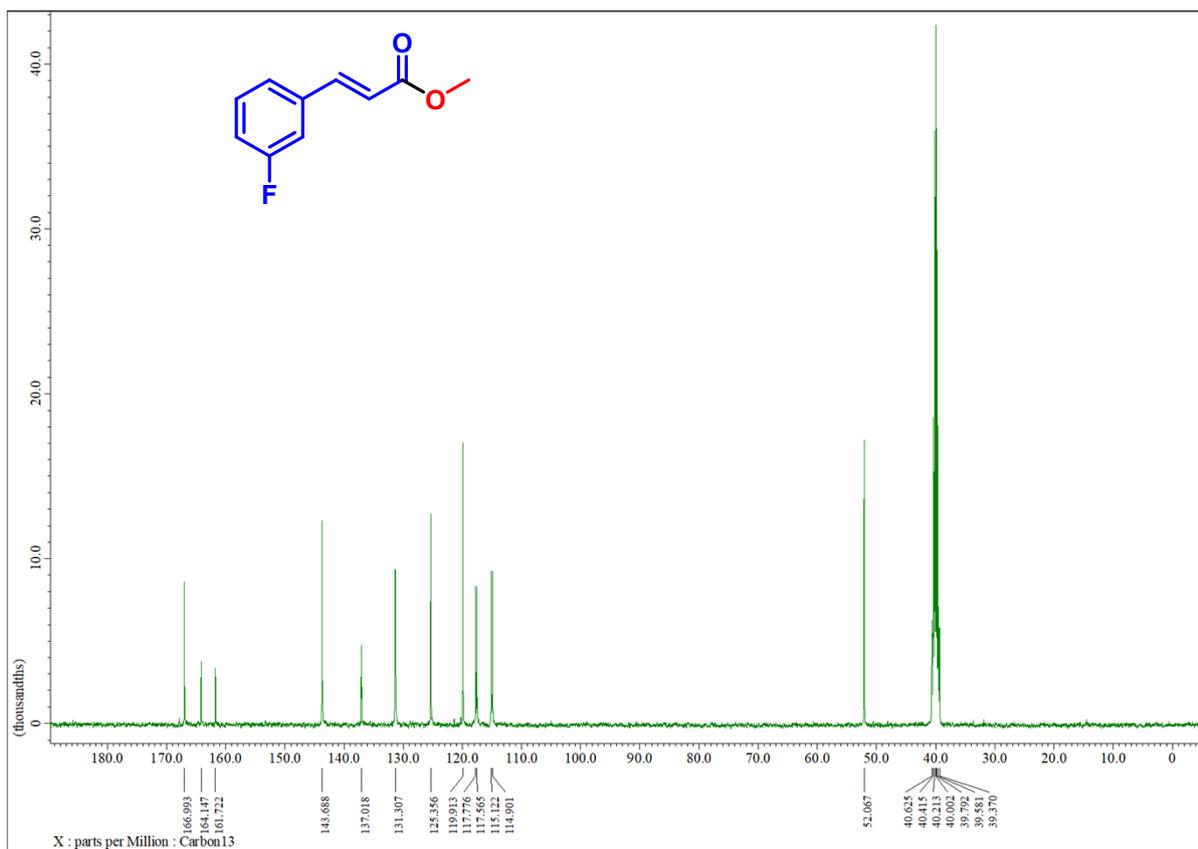


Figure S28. ^{13}C NMR spectrum of Methyl (E)-3-(3-fluorophenyl)acrylate **C11**.

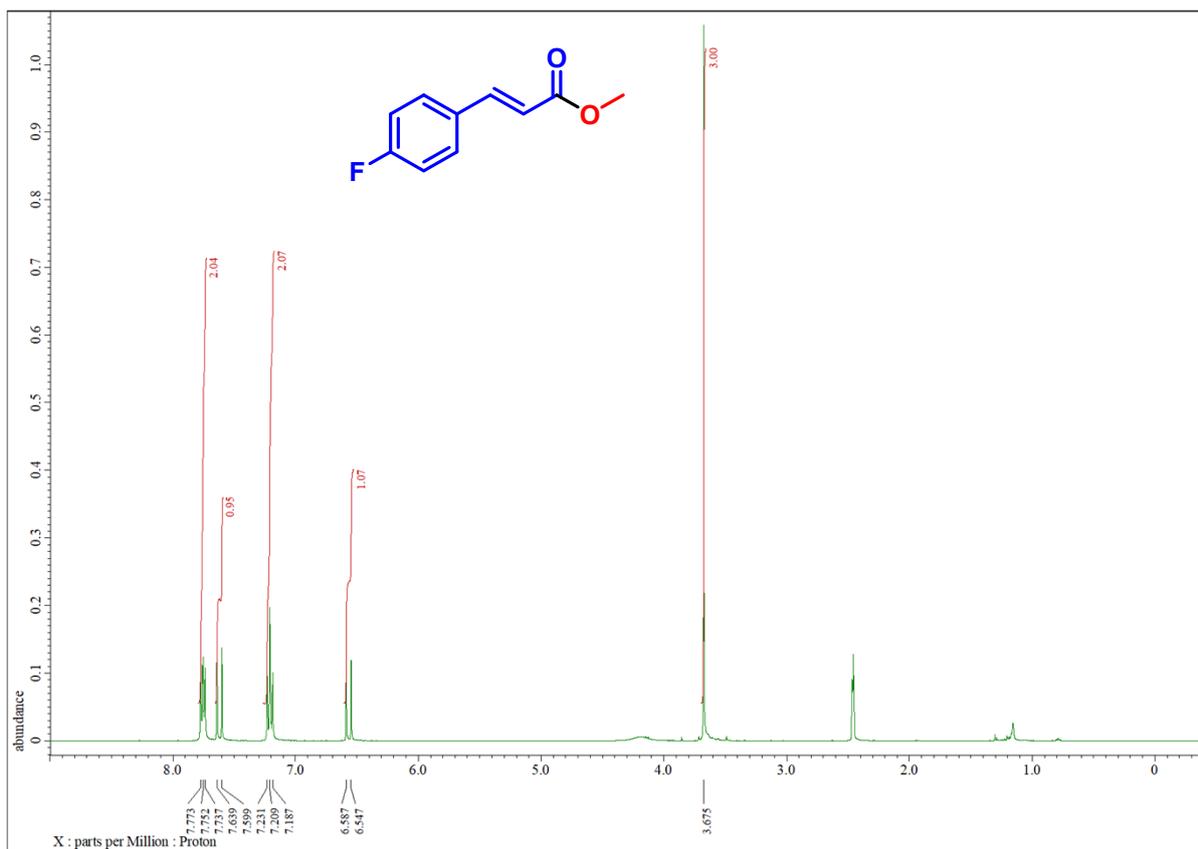


Figure S29. ¹H NMR spectrum of Methyl (E)-3-(4-fluorophenyl)acrylate **C12**.

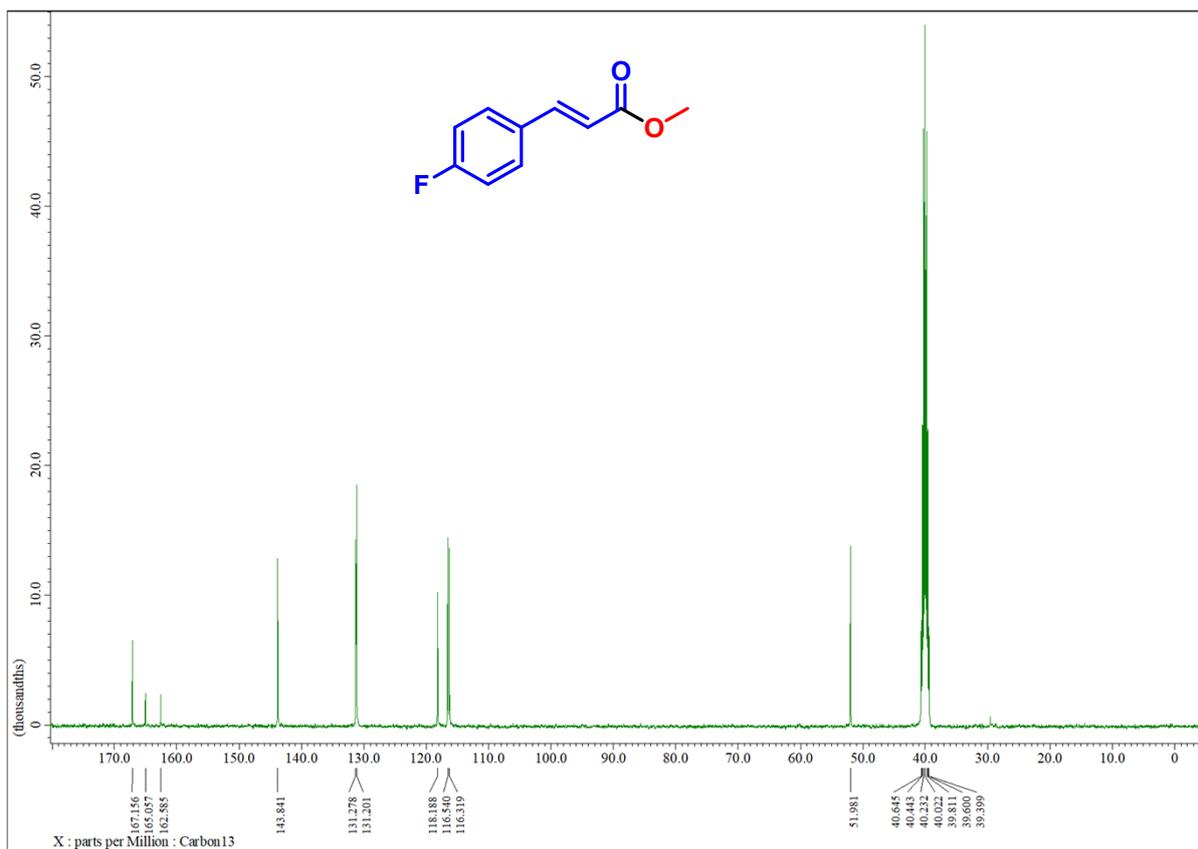


Figure S30. ^{13}C NMR spectrum of Methyl (E)-3-(4-fluorophenyl)acrylate **C12**.

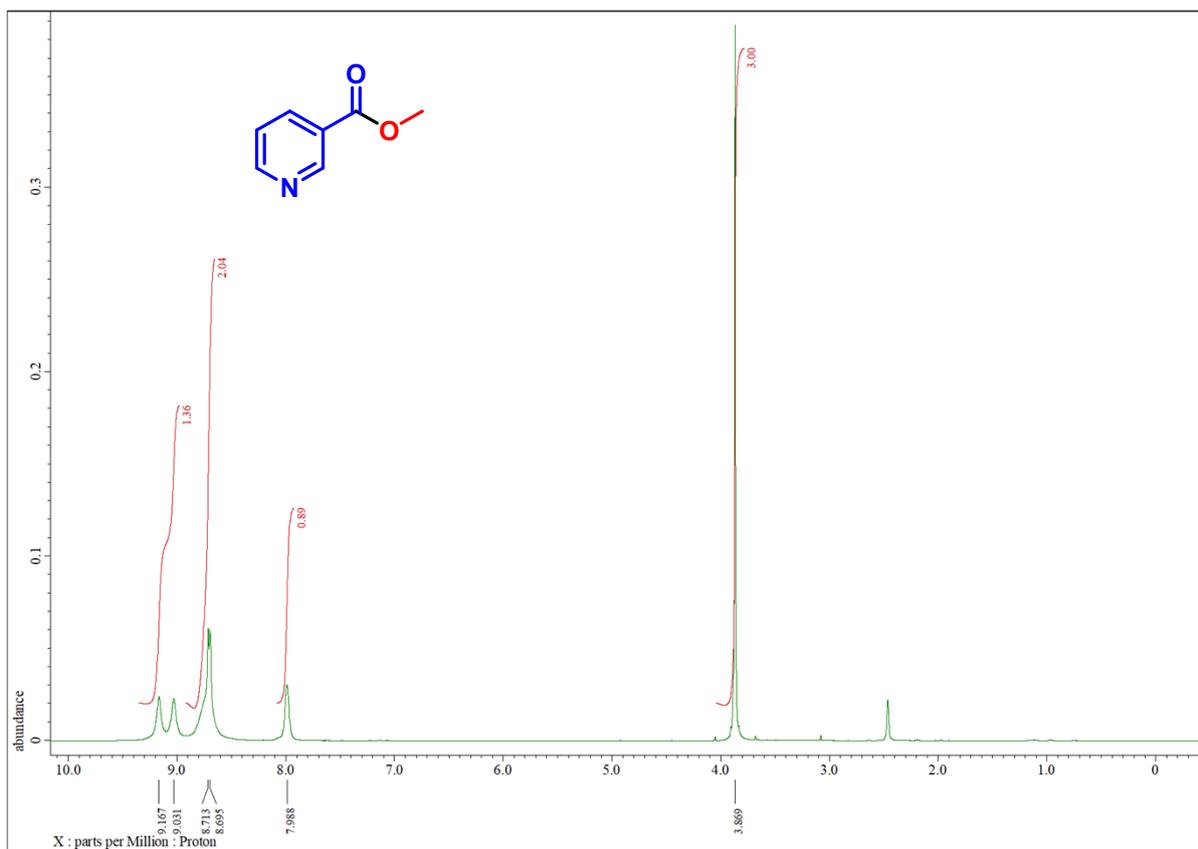


Figure S31. ¹H NMR spectrum of Methyl nicotinate **C13**.

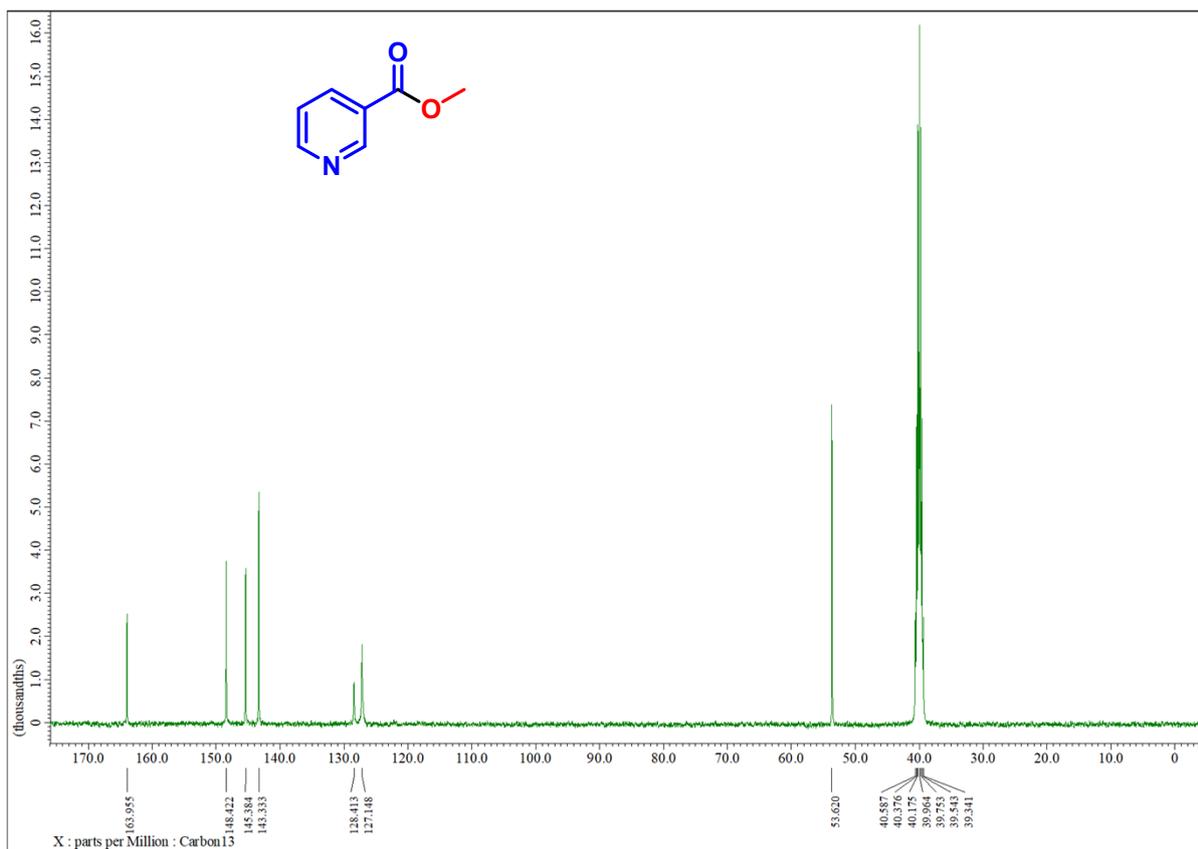


Figure S32. ^{13}C NMR spectrum of Methyl nicotinate C13.

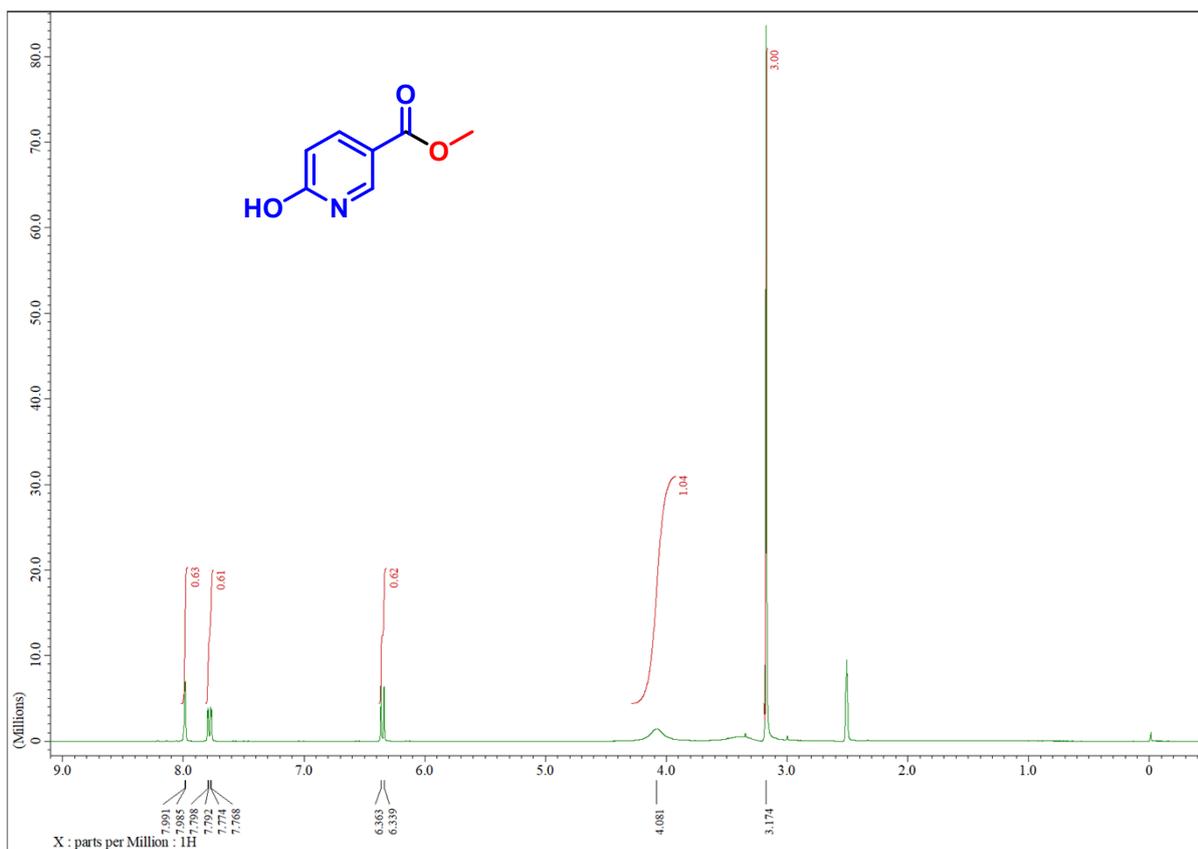


Figure S33. ^1H NMR spectrum of Methyl 6-hydroxynicotinate **C14**.

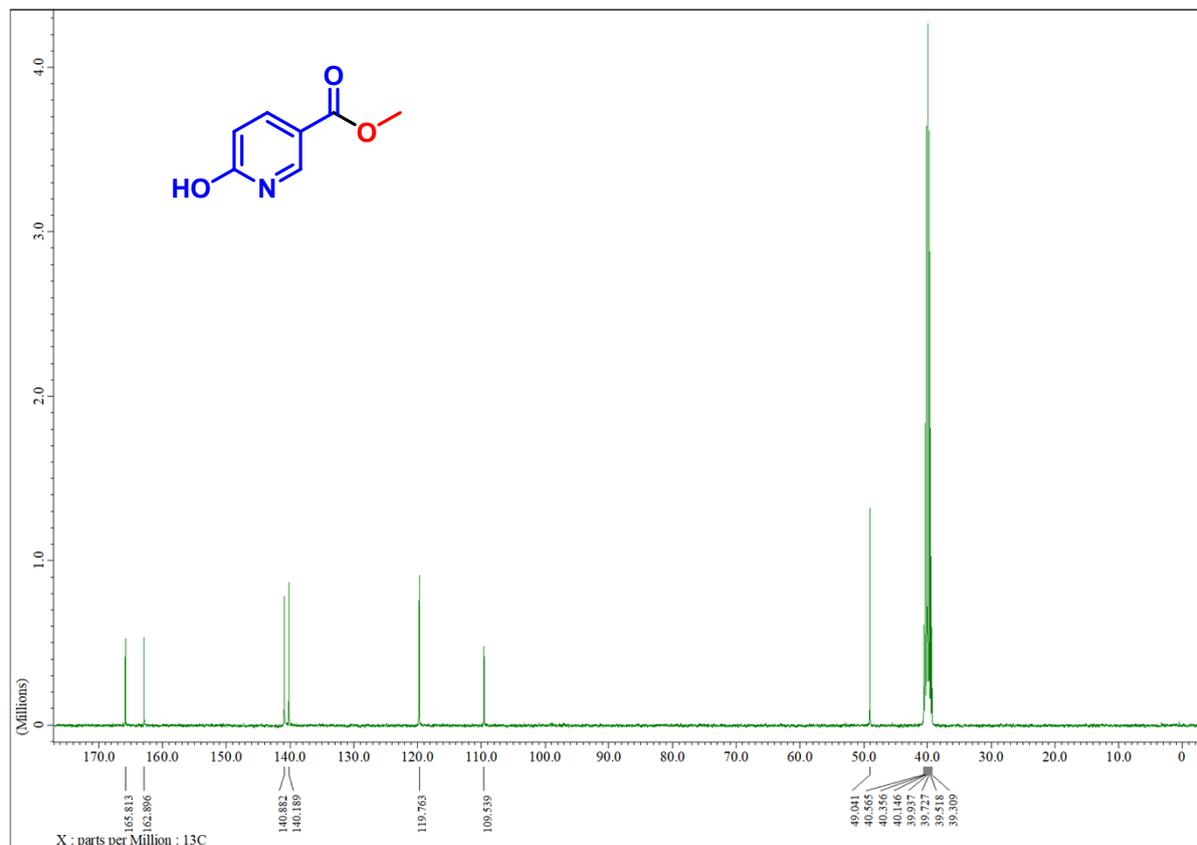


Figure S34. ^{13}C NMR spectrum of Methyl 6-hydroxynicotinate **C14**.

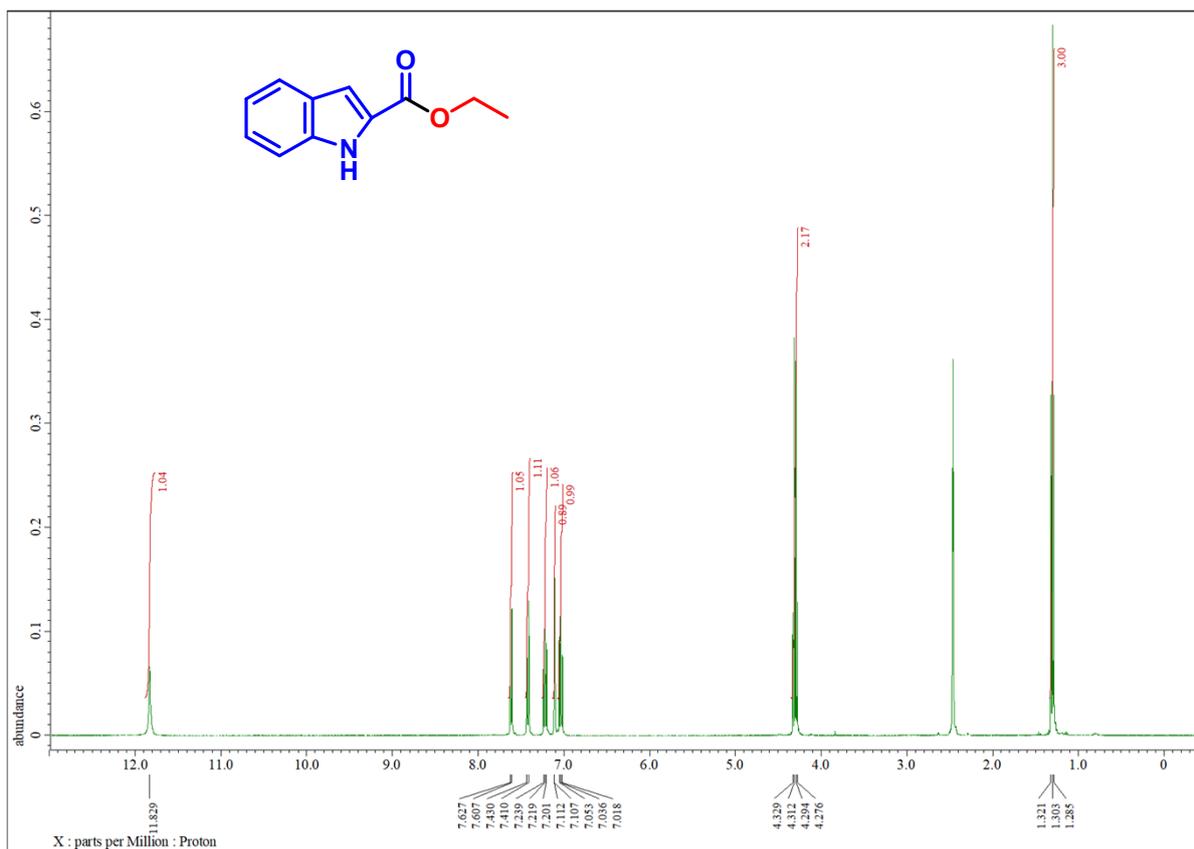


Figure S35. ¹H NMR spectrum of Ethyl 1H-indole-2-carboxylate **C15**.

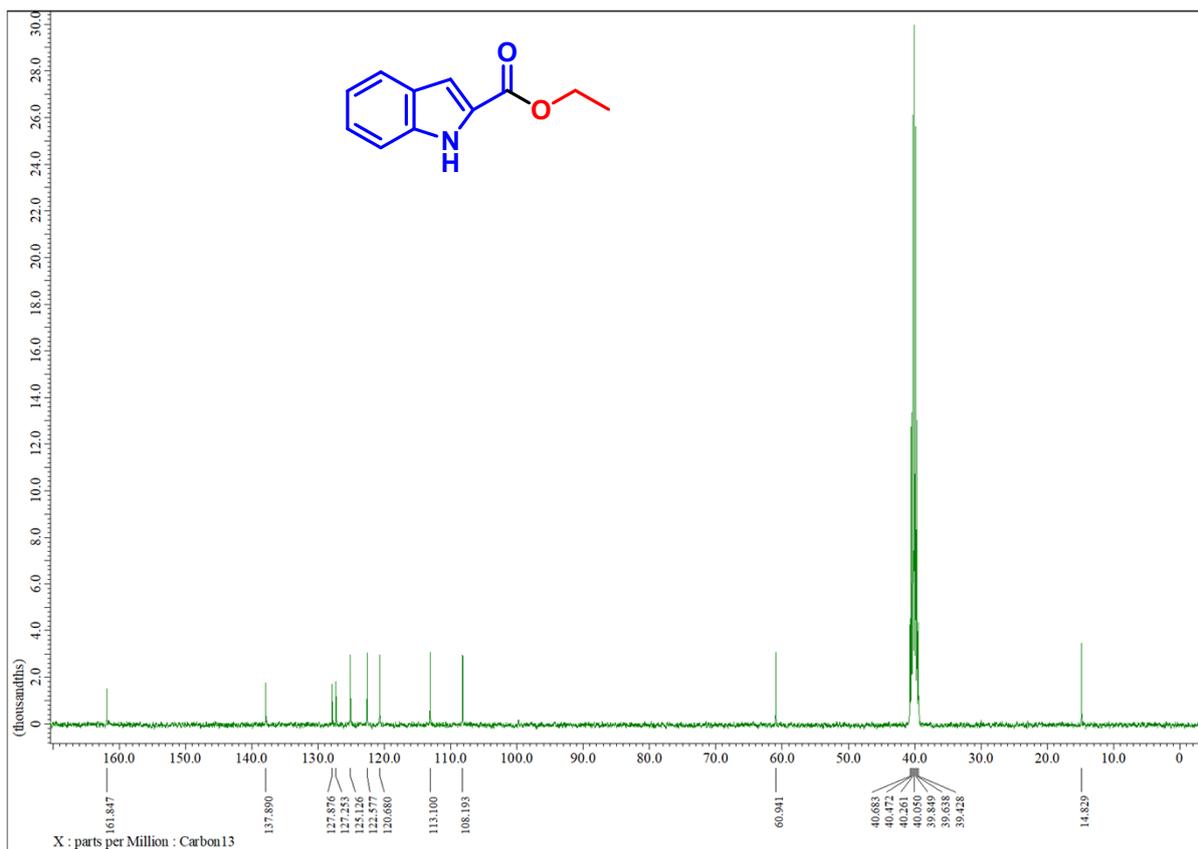


Figure S36. ^{13}C NMR spectrum of Ethyl 1H-indole-2-carboxylate **C15**.

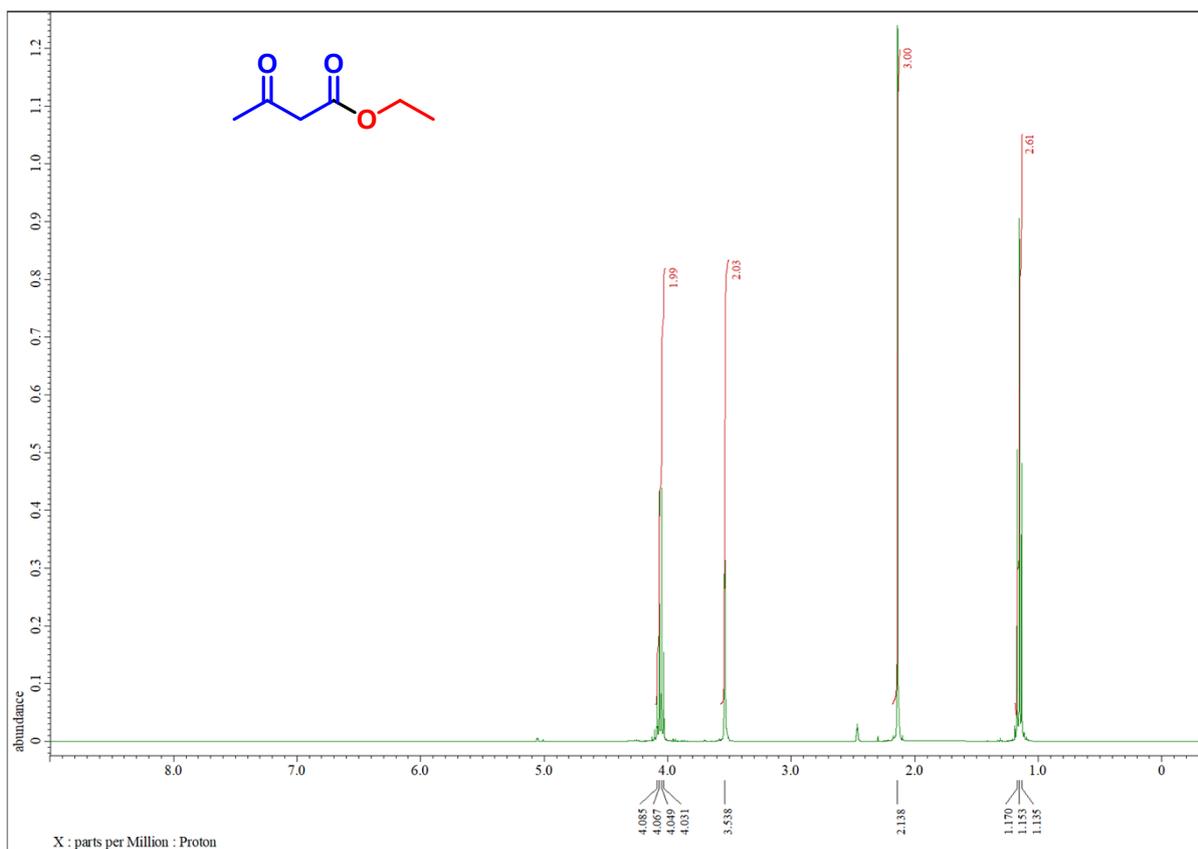


Figure S37. ^1H NMR spectrum of Ethyl 3-oxobutanoate **C16**.

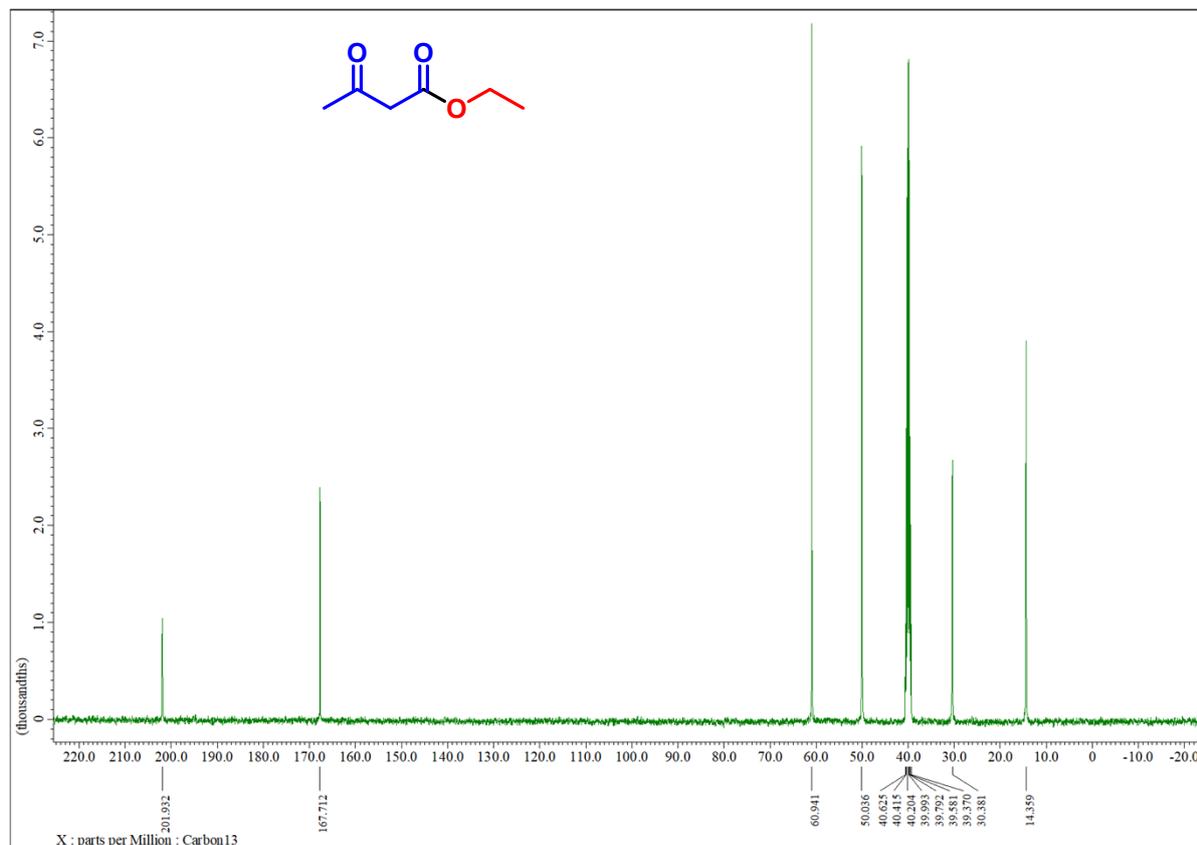


Figure S38. ^{13}C NMR spectrum of Ethyl 3-oxobutanoate **C16**.

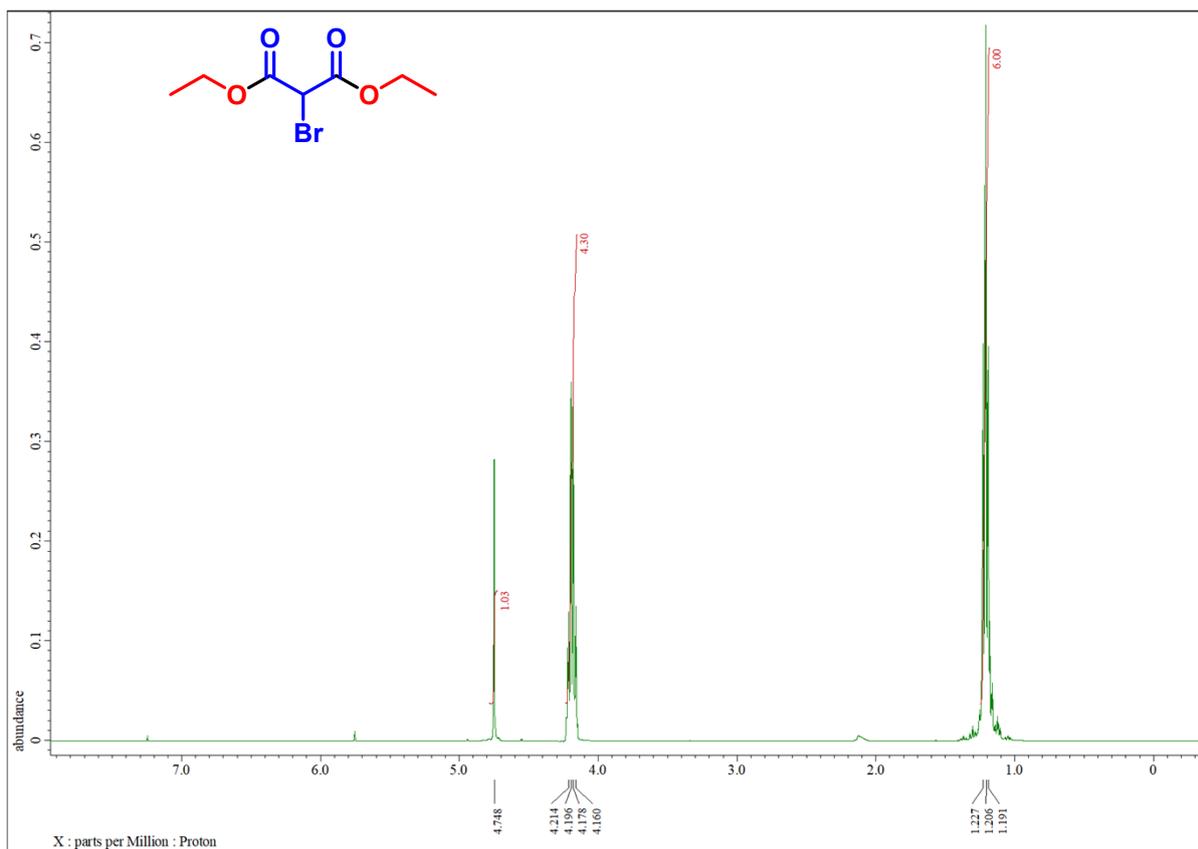


Figure S39. ¹H NMR spectrum of Diethyl 2-bromomalonate **C17**.

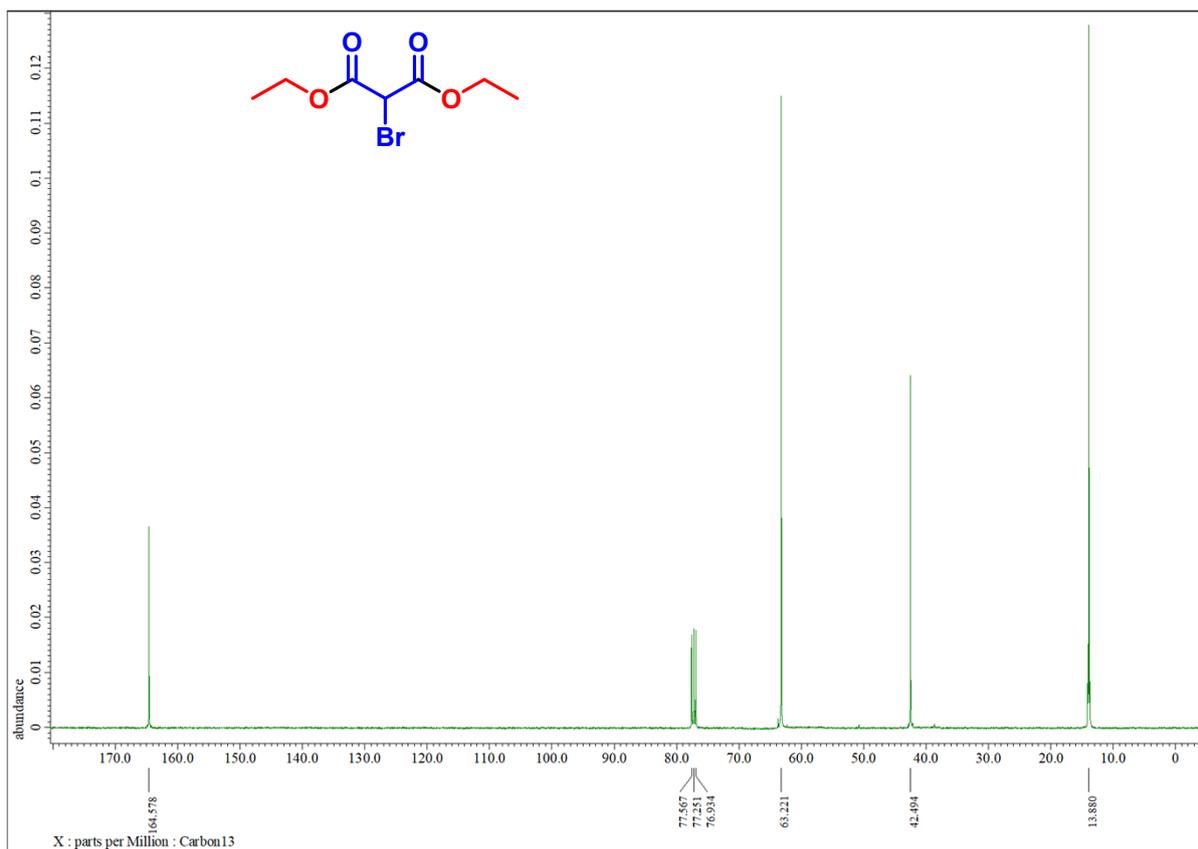


Figure S40. ^{13}C NMR spectrum of Diethyl 2-bromomalonate **C17**.

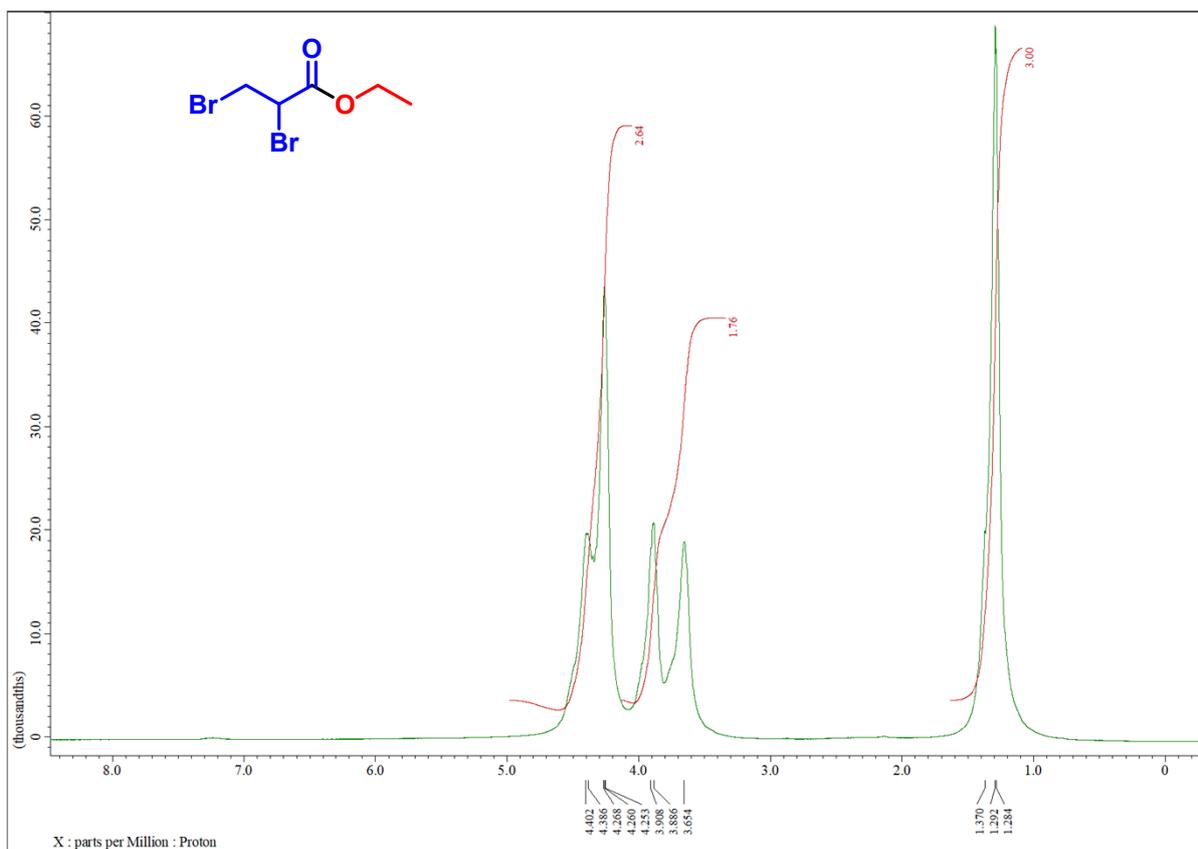


Figure S41. ^1H NMR spectrum of Ethyl 2,3-dibromopropanoate **C18**.

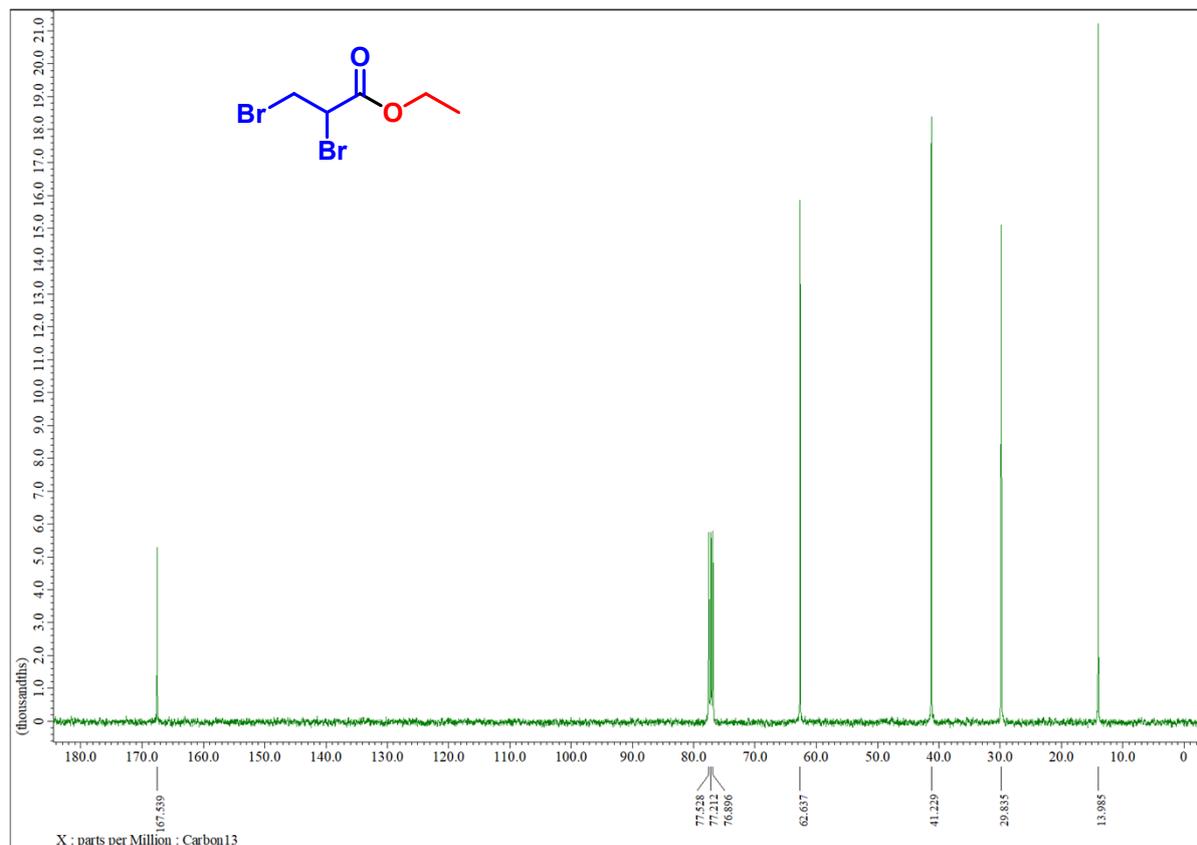


Figure S42. ^{13}C NMR spectrum of Ethyl 2,3-dibromopropanoate **C18**.

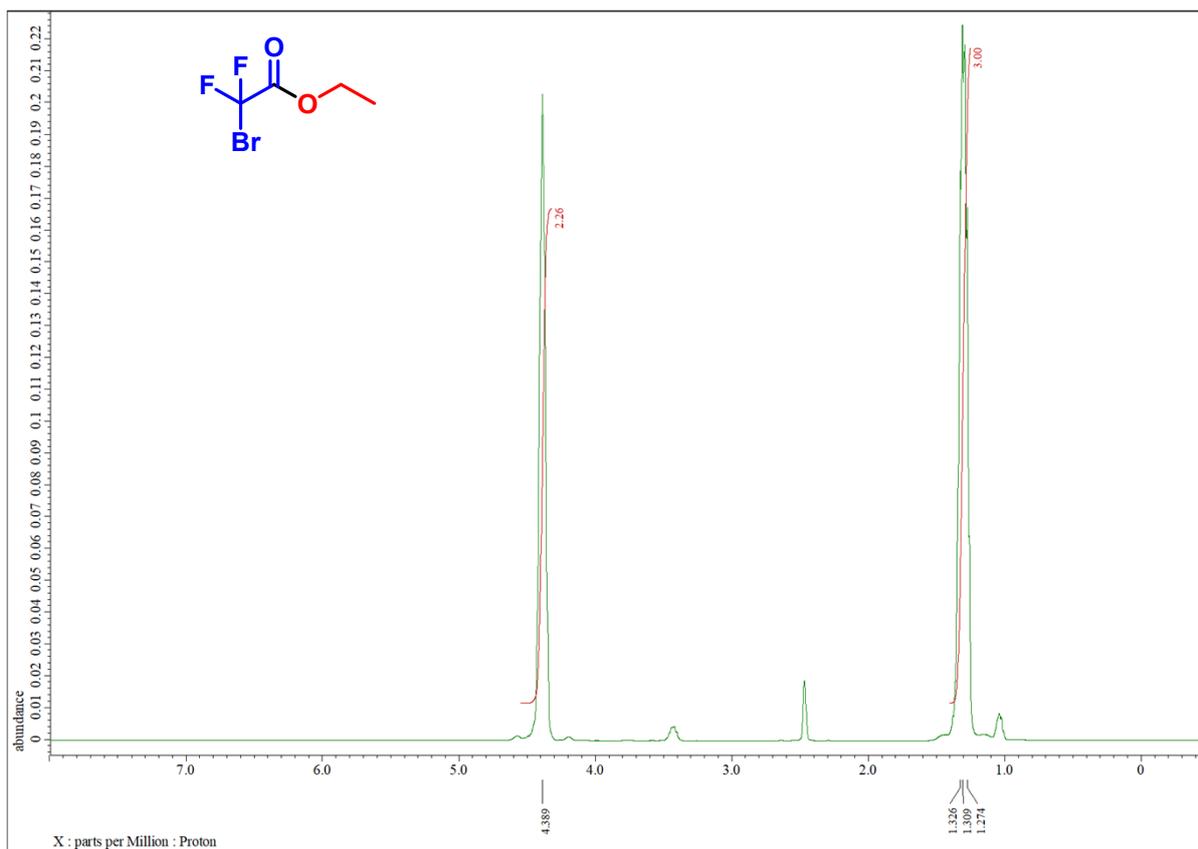


Figure S43. ¹H NMR spectrum of Ethyl 2-bromo-2,2-difluoroacetate **C19**.

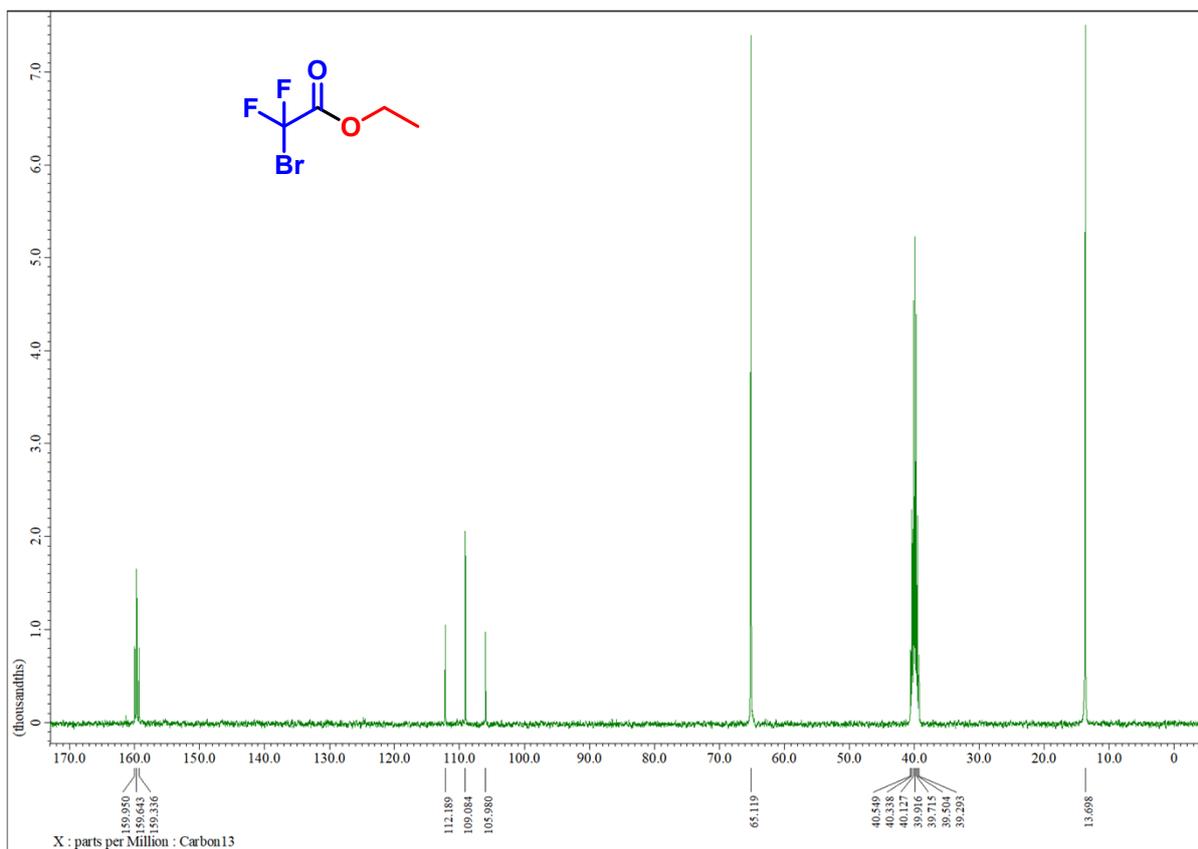


Figure S44. ^{13}C NMR spectrum of Ethyl 2-bromo-2,2-difluoroacetate **C19**.

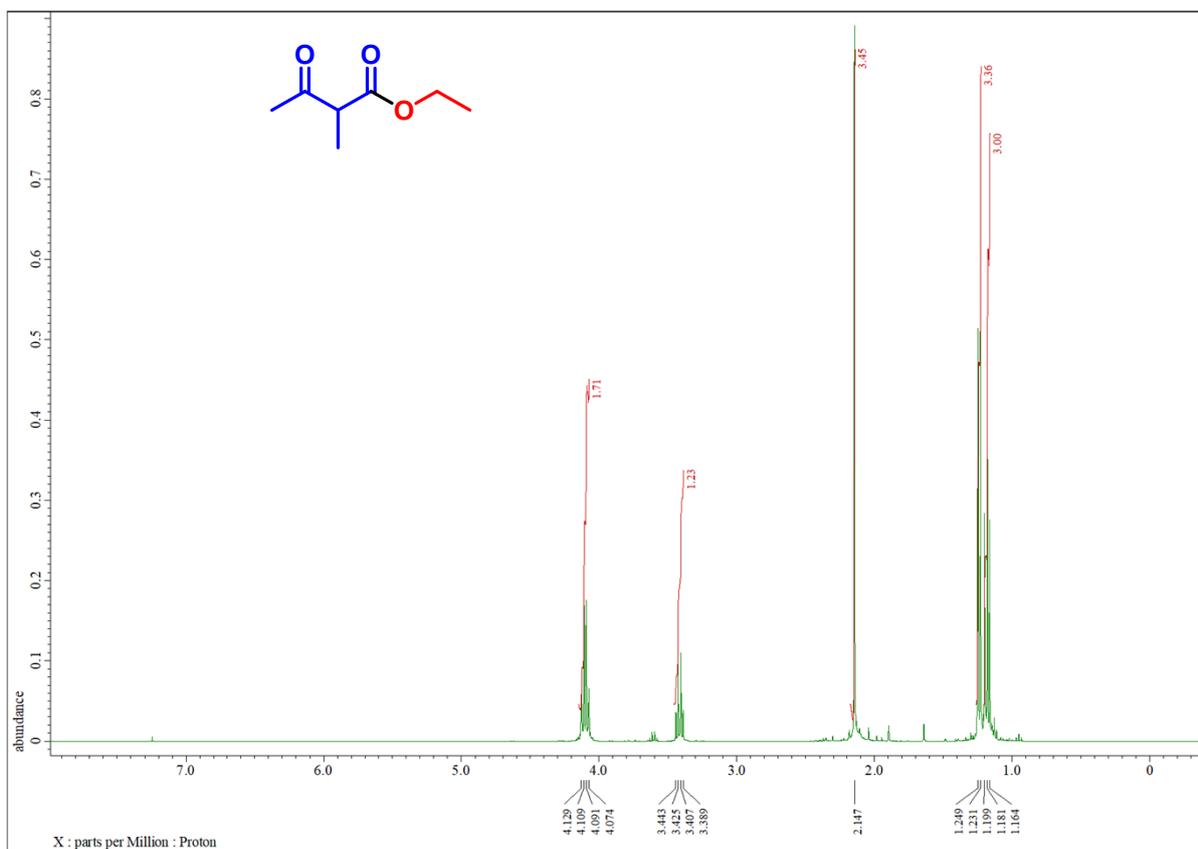


Figure S45. ^1H NMR spectrum of Ethyl 2-methyl-3-oxobutanoate **C20**.

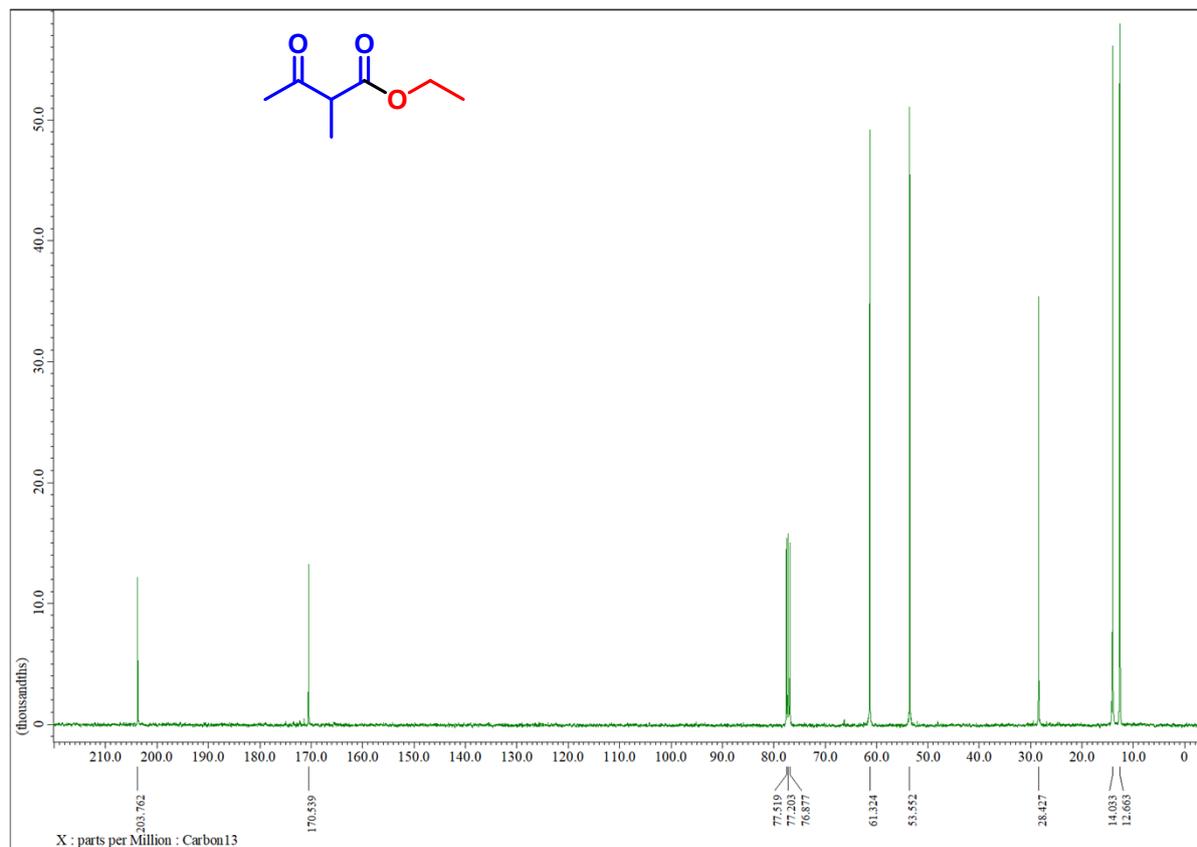


Figure S46. ^{13}C NMR spectrum of Ethyl 2-methyl-3-oxobutanoate **C20**.

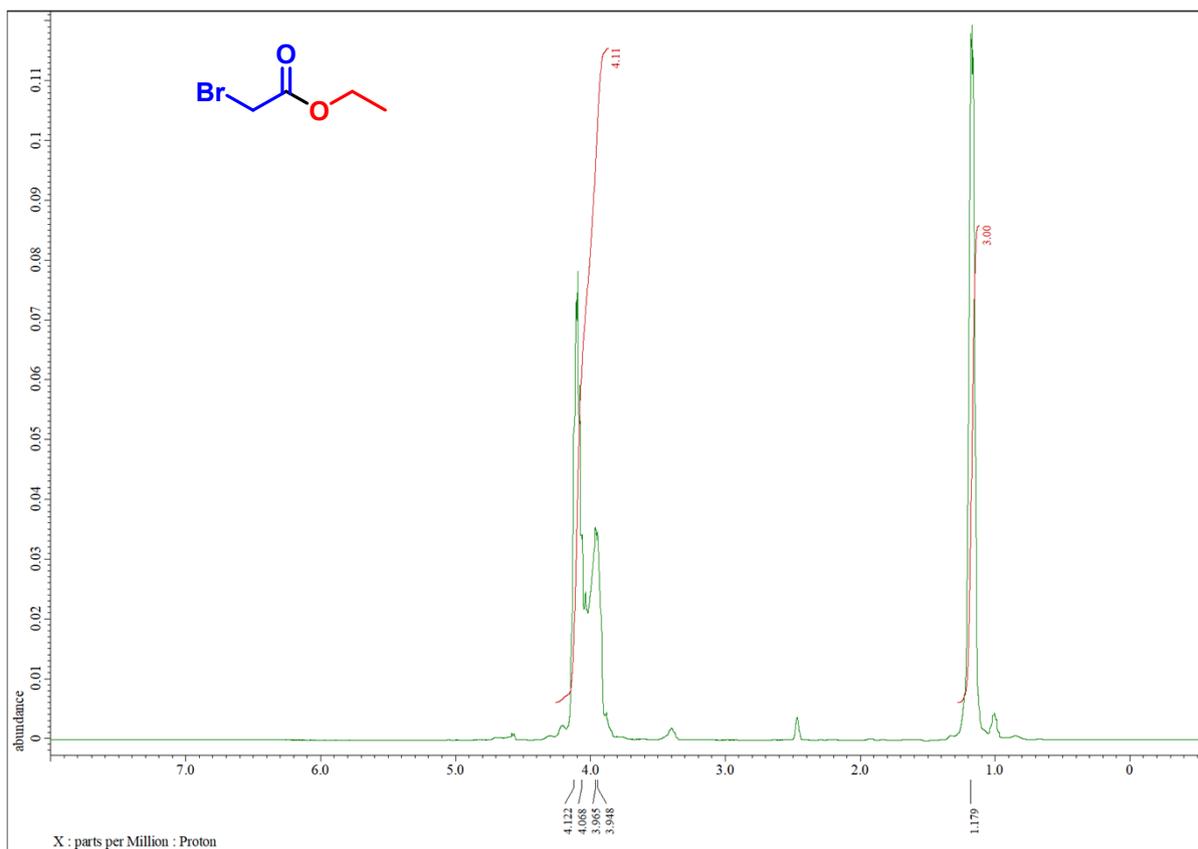


Figure S47. ^1H NMR spectrum of Ethyl 2-bromoacetate **C21**.

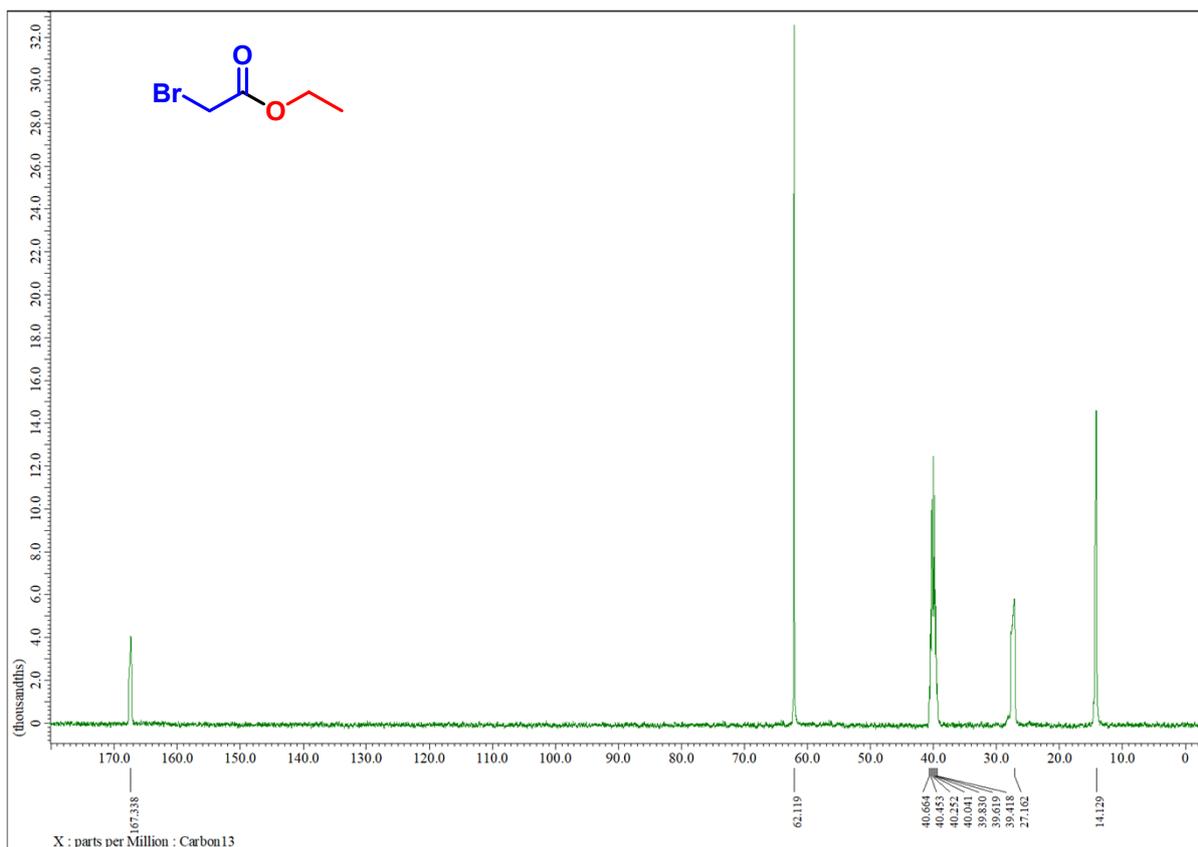


Figure S48. ^{13}C NMR spectrum of Ethyl 2-bromoacetate C21.

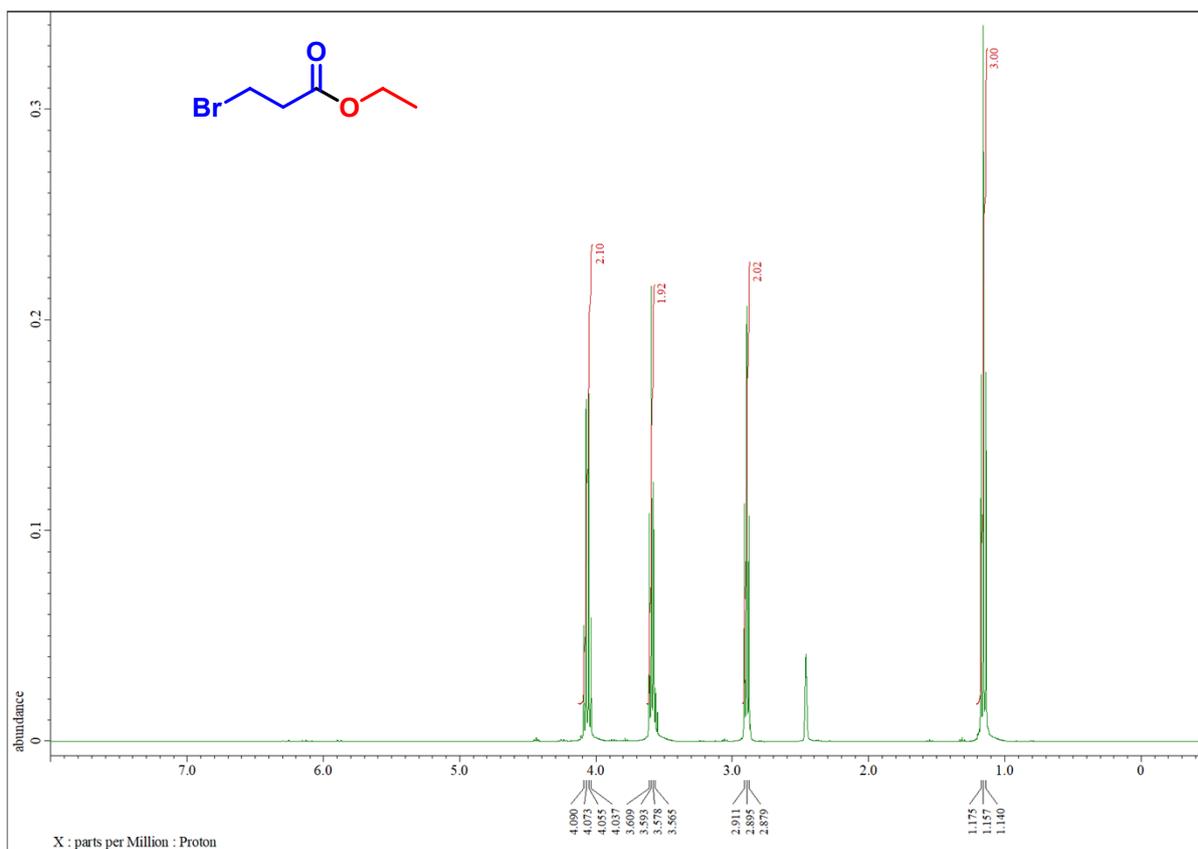


Figure S49. ^1H NMR spectrum of Ethyl 3-bromopropanoate **C22**.

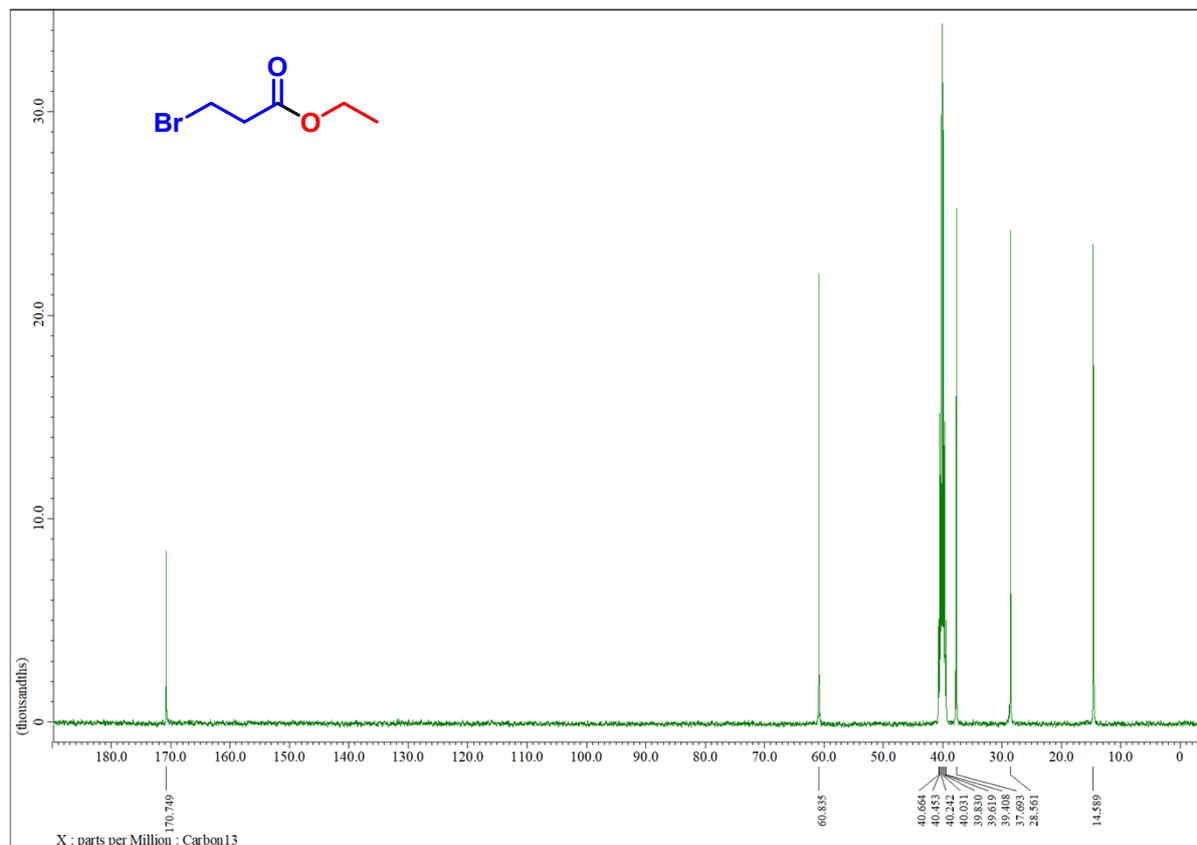


Figure S50. ^{13}C NMR spectrum of Ethyl 3-bromopropanoate **C22**.

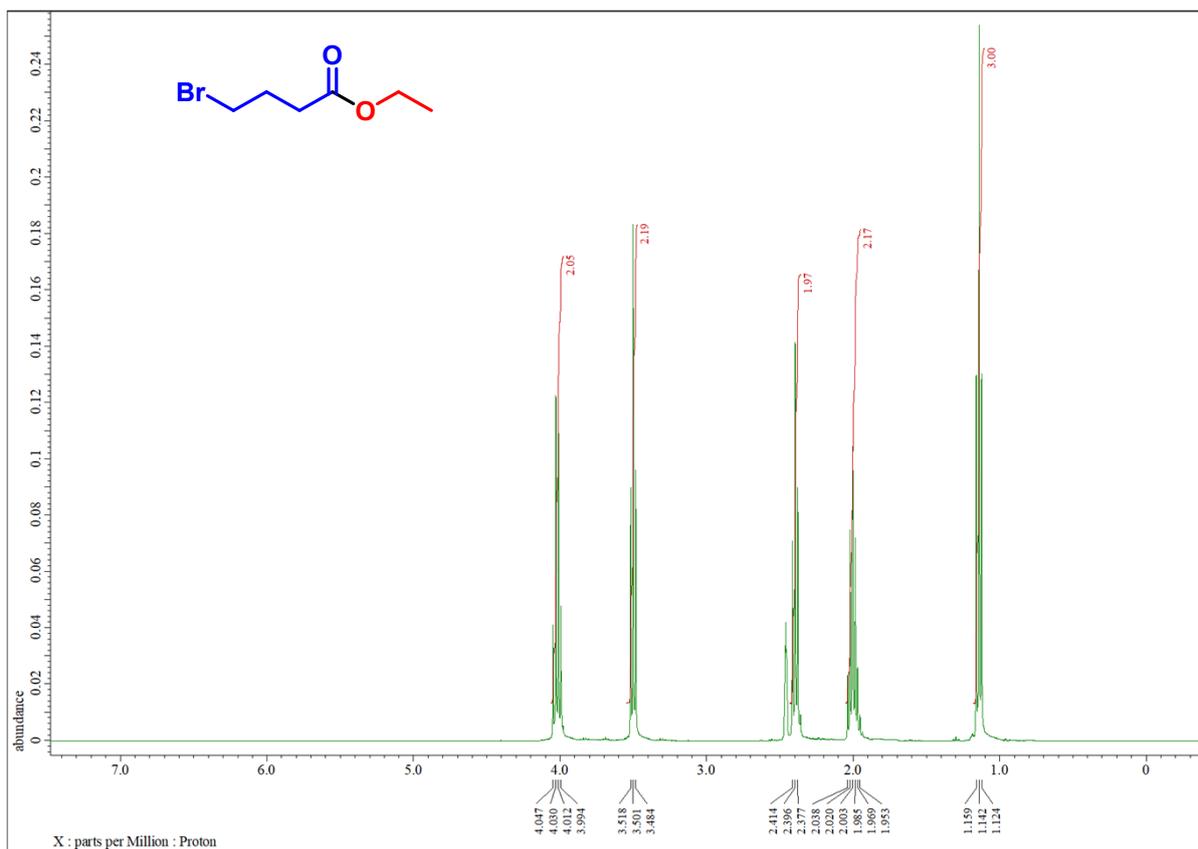


Figure S51. ¹H NMR spectrum of Ethyl 4-bromobutanoate **C23**.

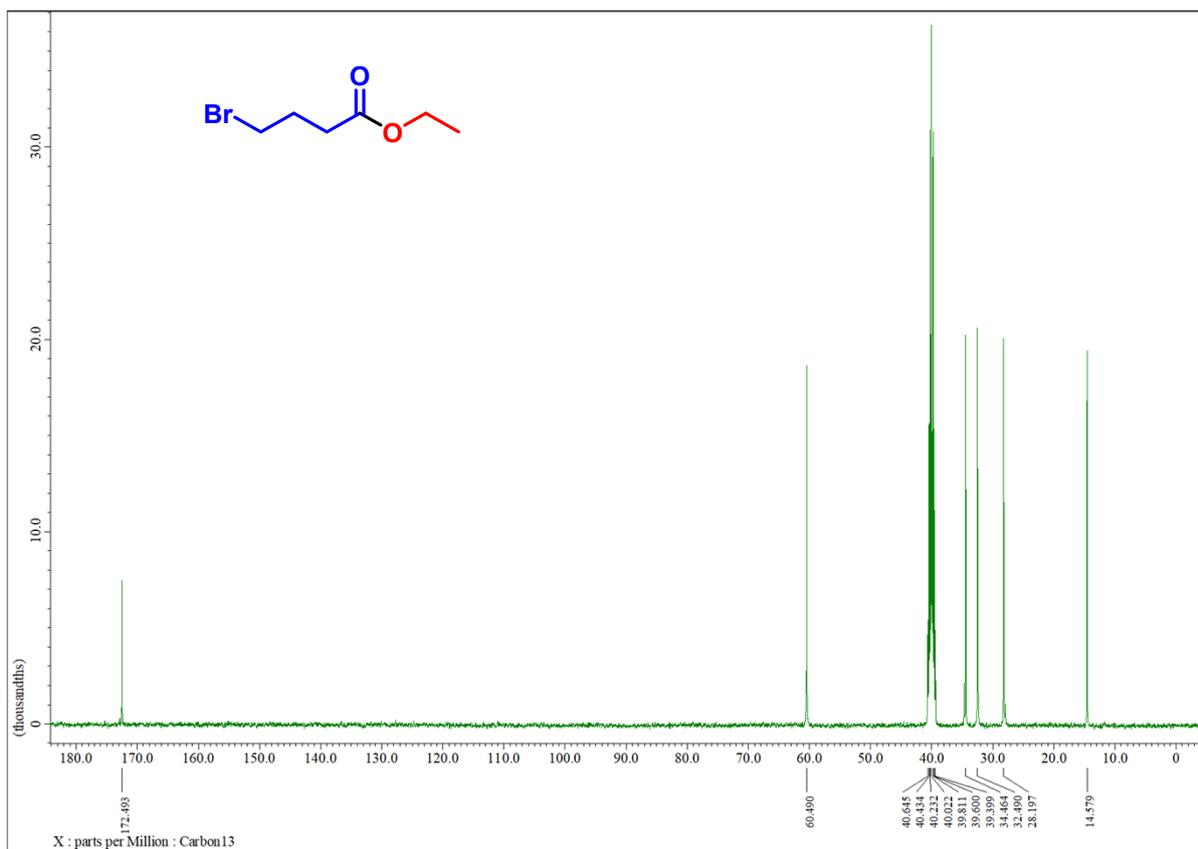


Figure S52. ^{13}C NMR spectrum of Ethyl 4-bromobutanoate C23.

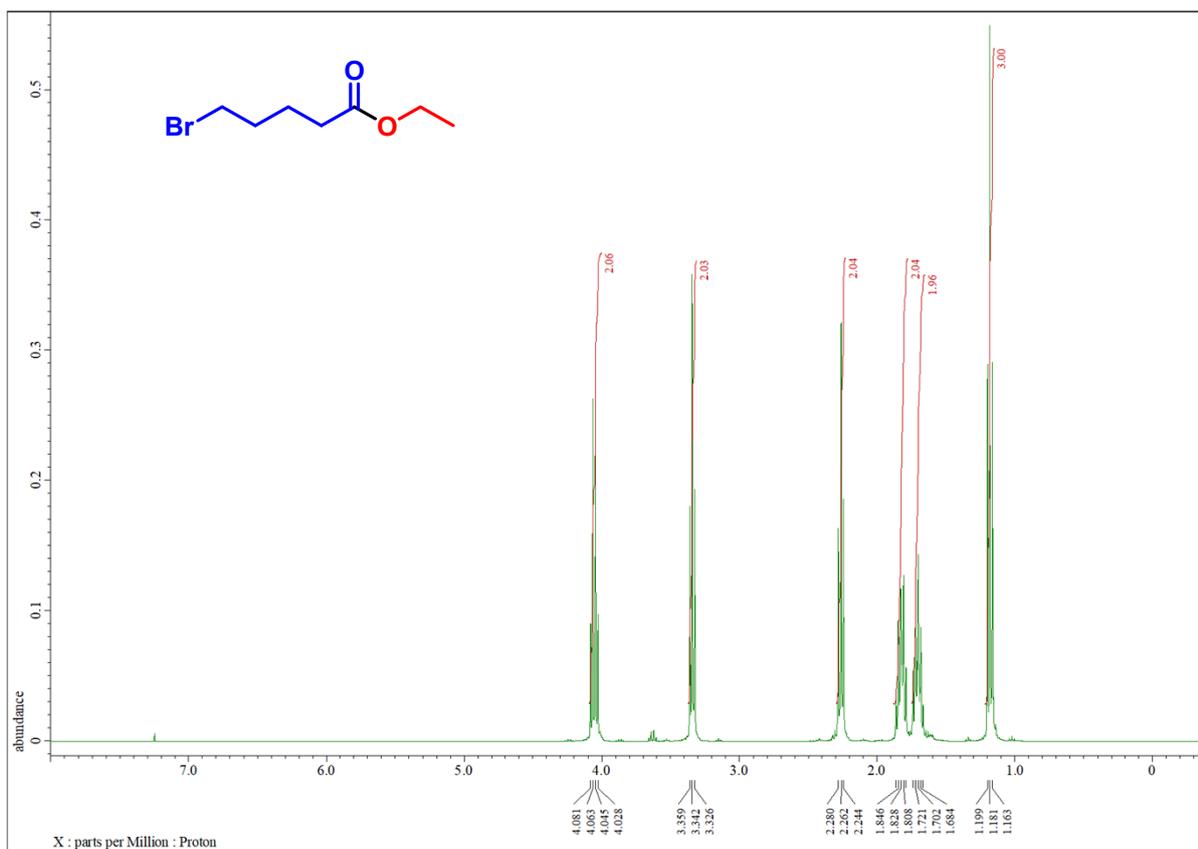


Figure S53. ^1H NMR spectrum of Ethyl 5-bromopentanoate **C24**.

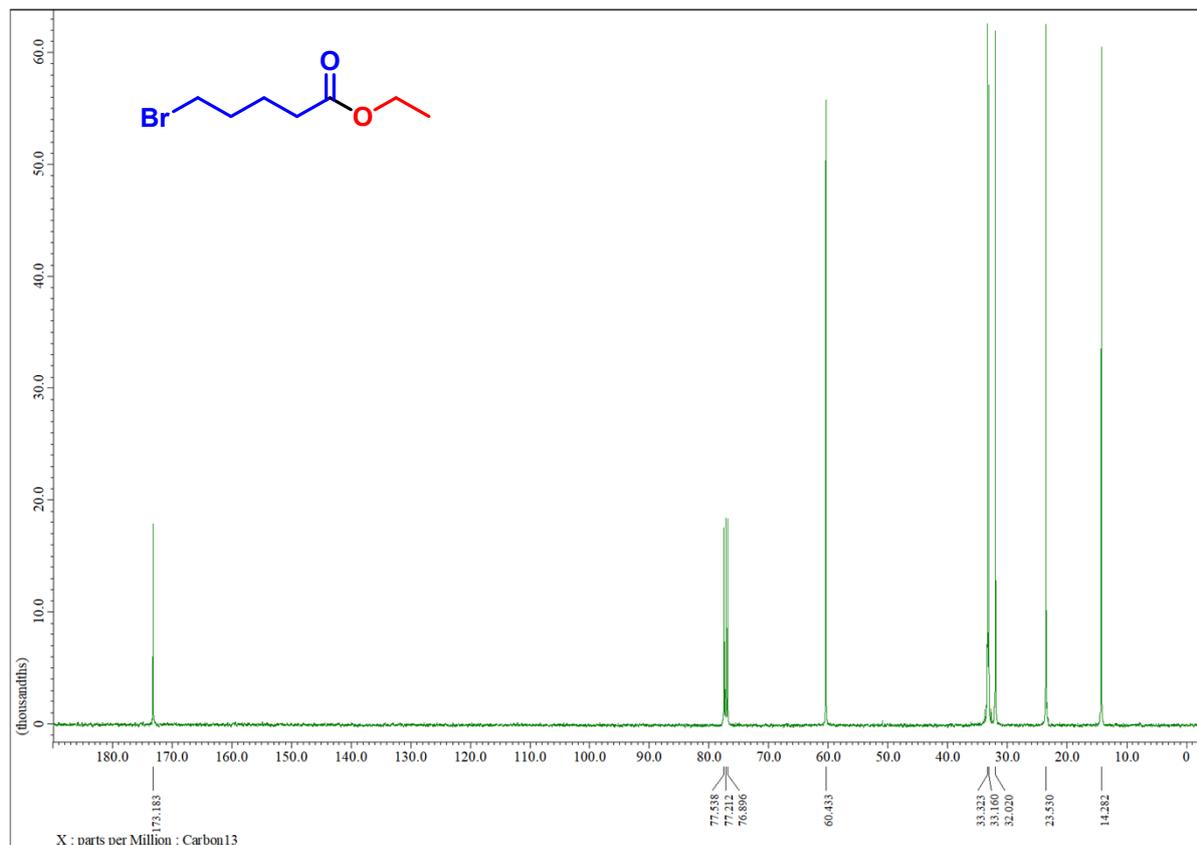


Figure S54. ^{13}C NMR spectrum of Ethyl 5-bromopentanoate **C24**.

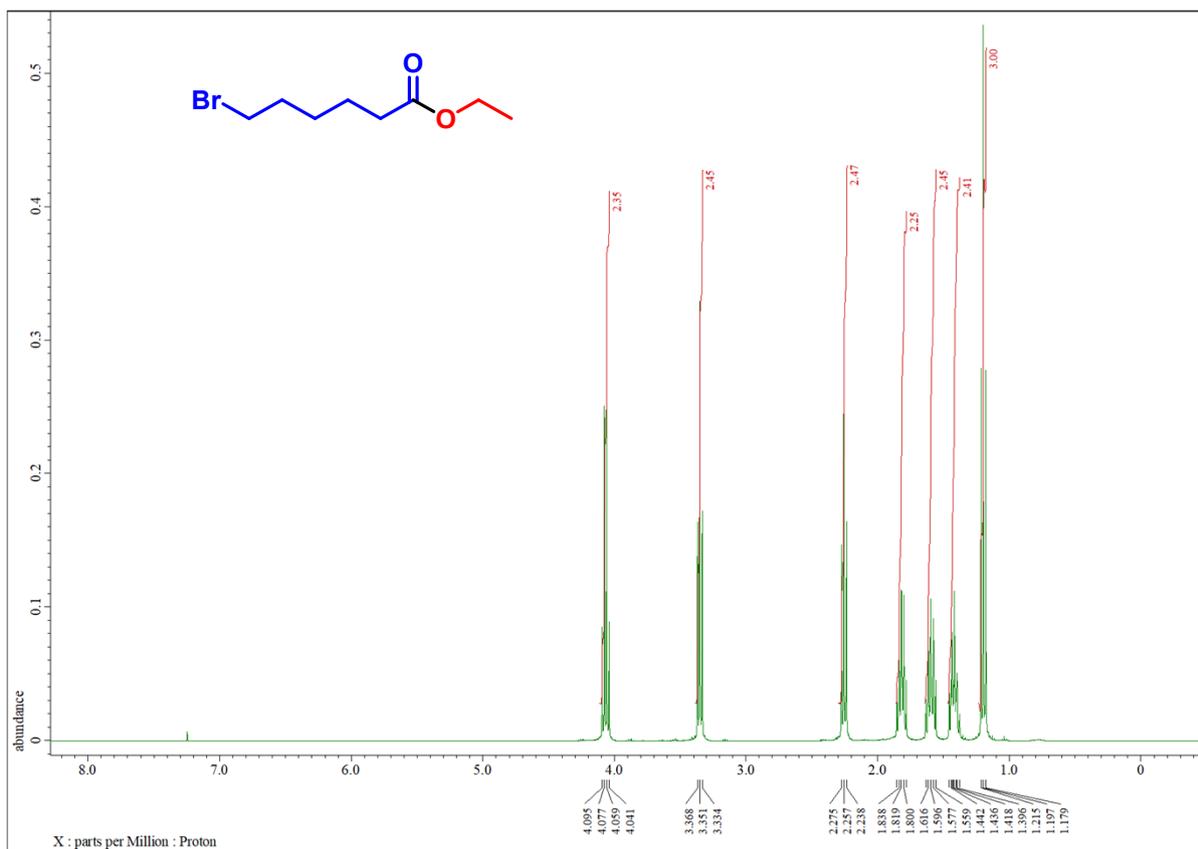


Figure S55. ¹H NMR spectrum of Ethyl 6-bromohexanoate **C25**.

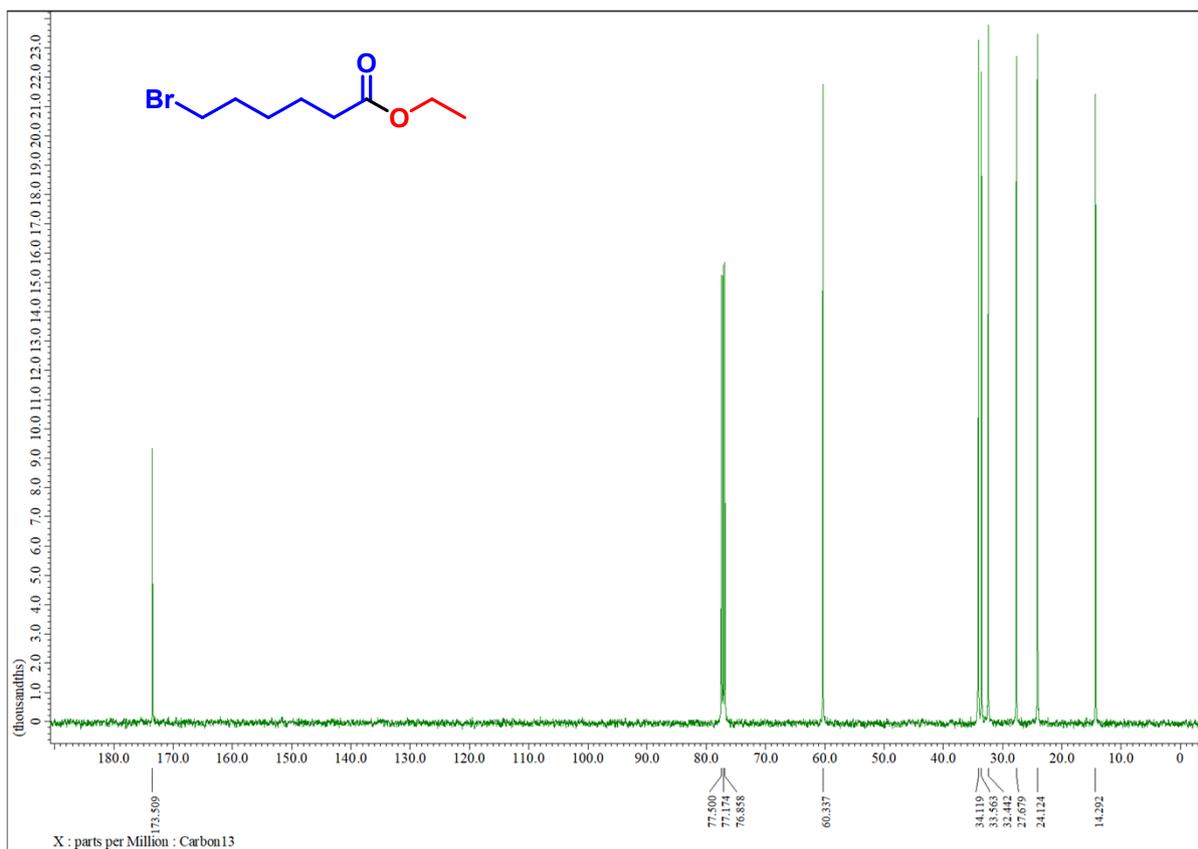


Figure S56. ¹³C NMR spectrum of Ethyl 6-bromohexanoate C25.

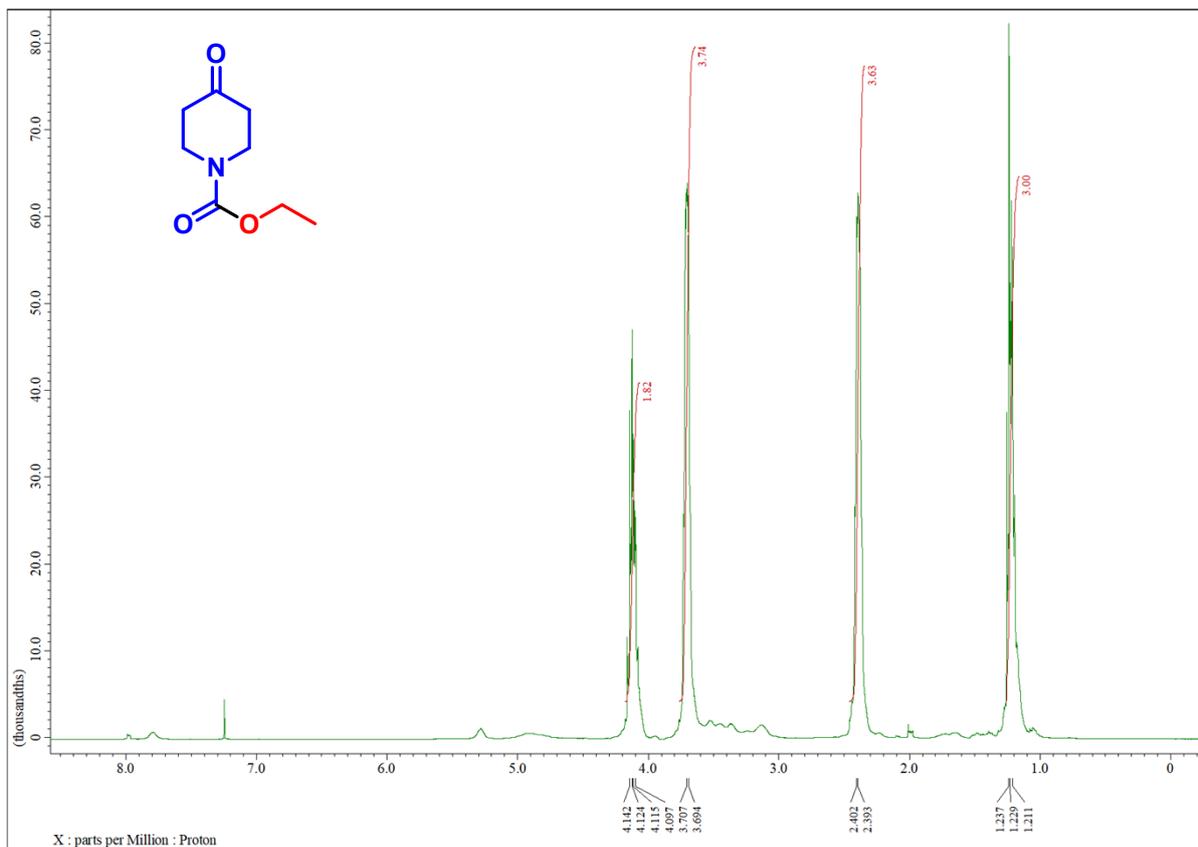


Figure S57. ^1H NMR spectrum of Ethyl 4-oxopiperidine-1-carboxylate **C26**.

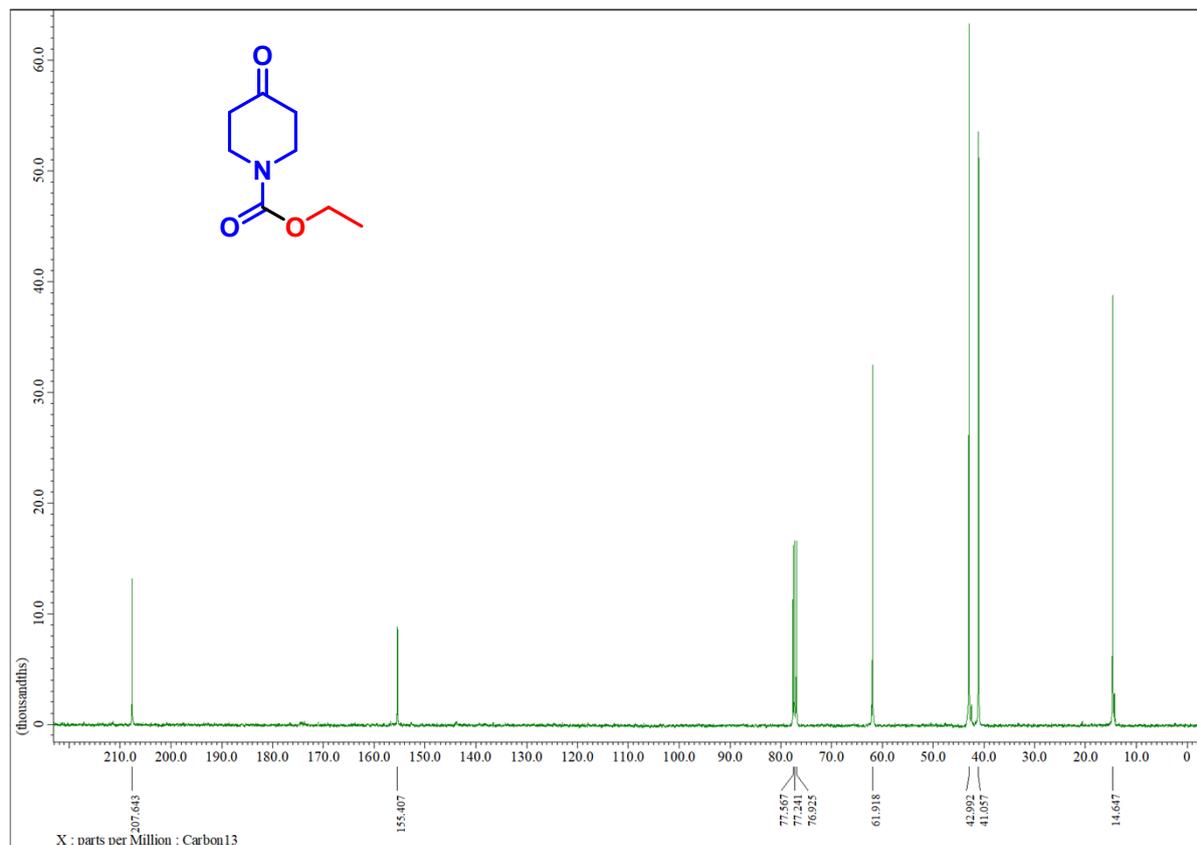


Figure S58. ^{13}C NMR spectrum of Ethyl 4-oxopiperidine-1-carboxylate **C26**.

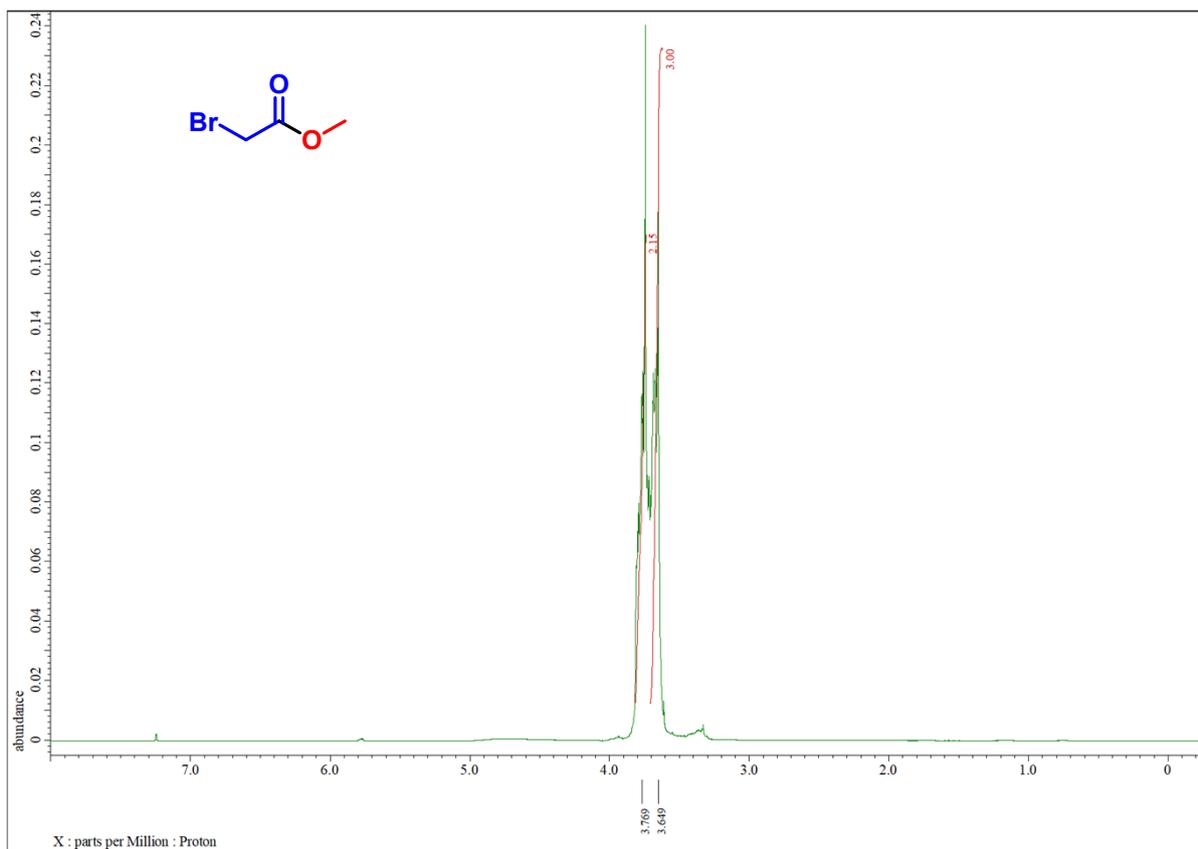


Figure S59. ¹H NMR spectrum of Methyl 2-bromoacetate **C27**.

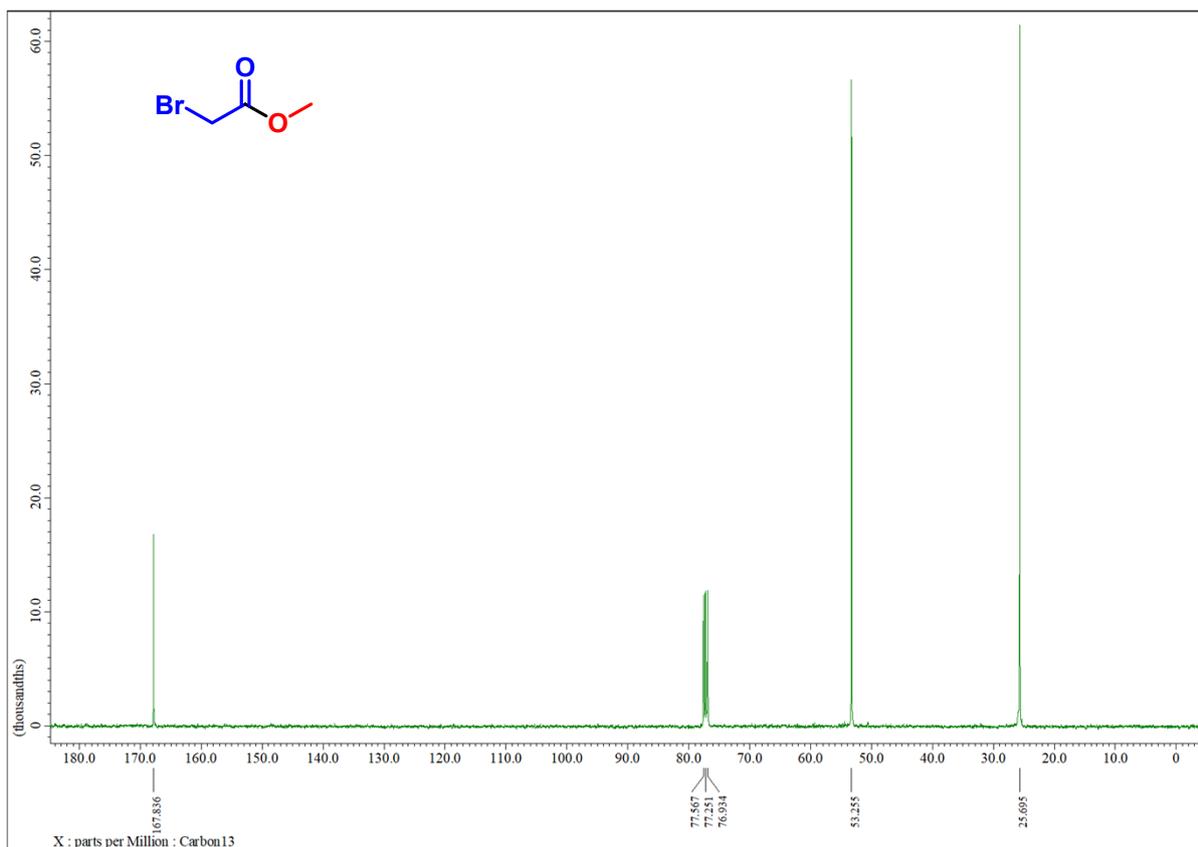


Figure S60. ^{13}C NMR spectrum of Methyl 2-bromoacetate C27.

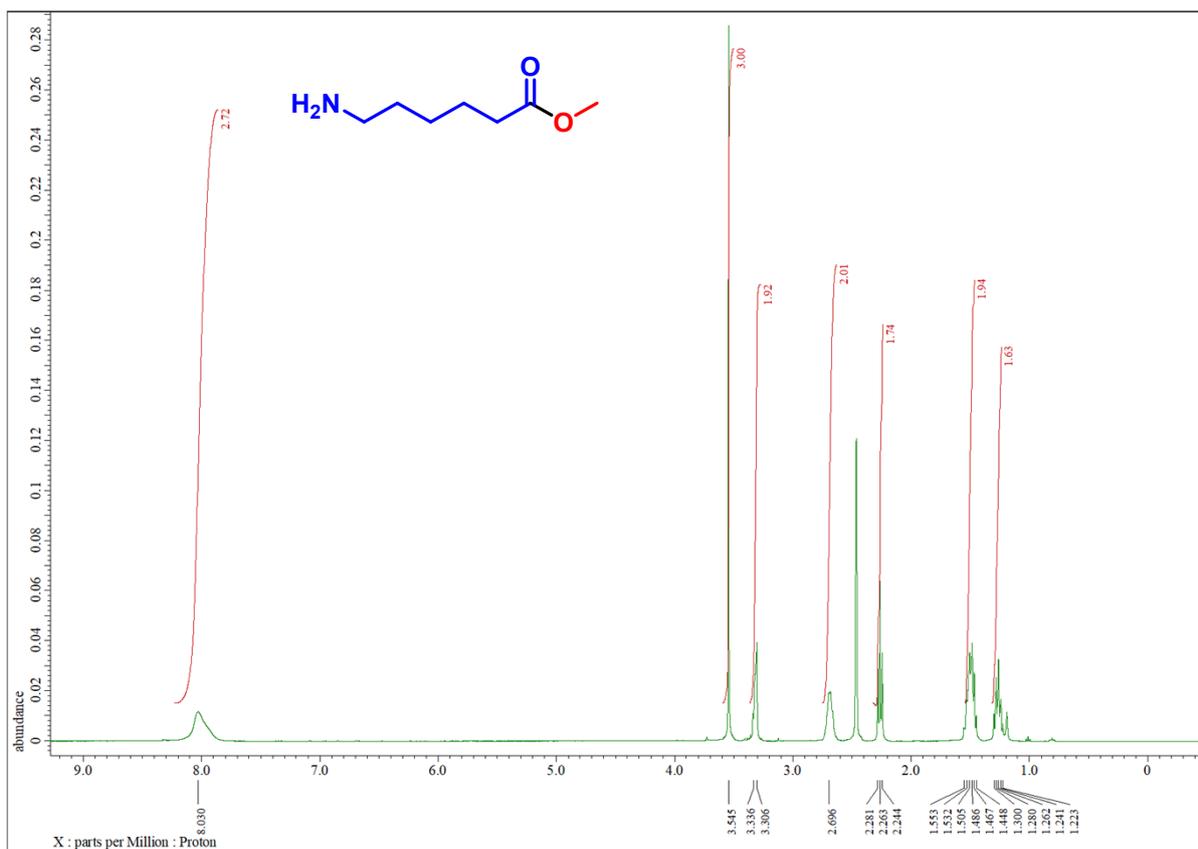


Figure S61. ^1H NMR spectrum of Methyl 6-aminohexanoate **C28**.

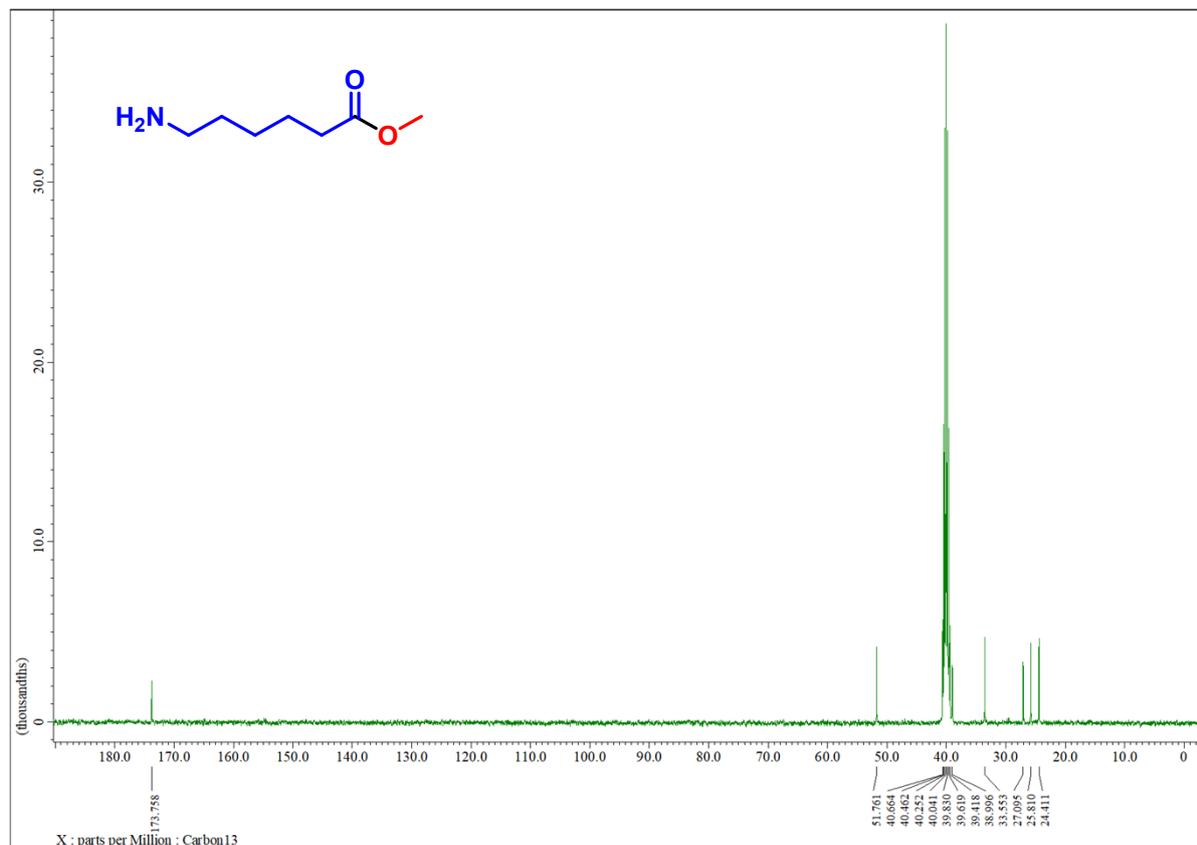


Figure S62. ^{13}C NMR spectrum of Methyl 6-aminohexanoate **C28**.

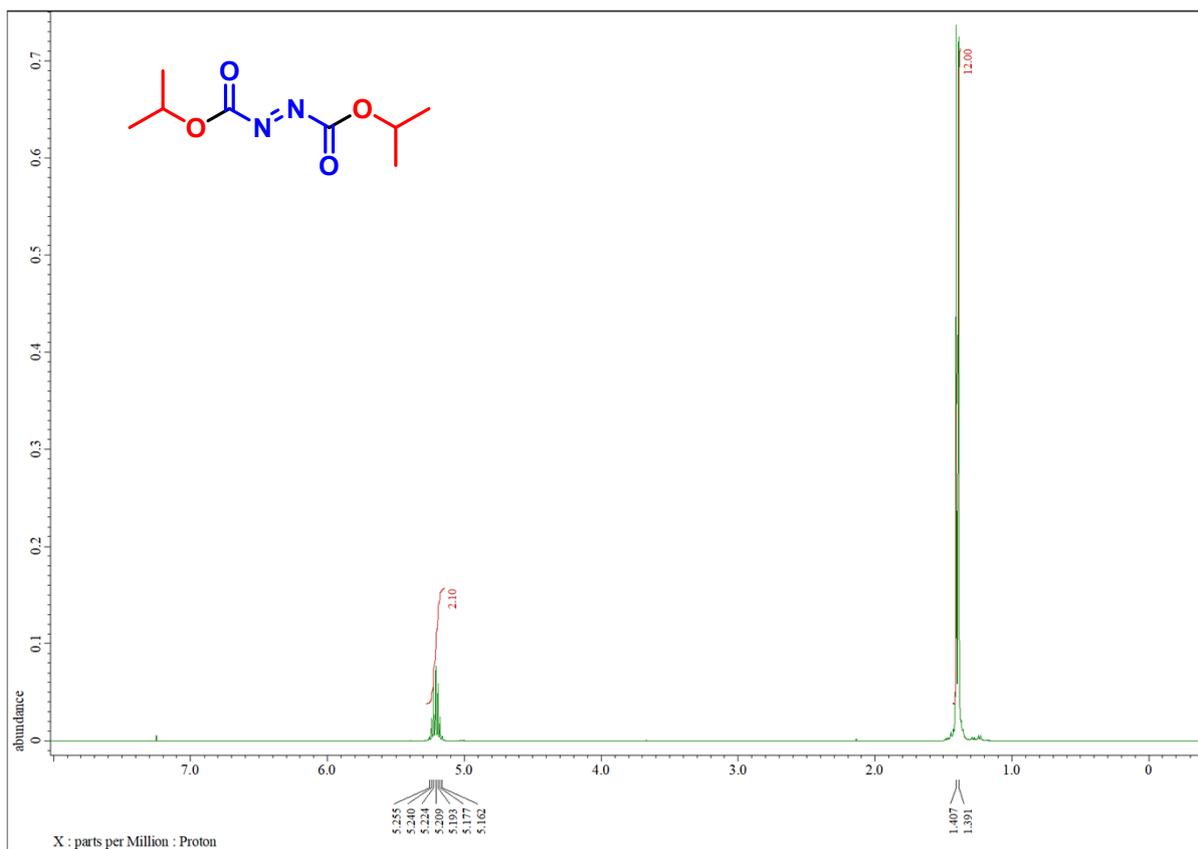


Figure S63. ¹H NMR spectrum of Diisopropyl (E)-diazene-1,2-dicarboxylate **C29**.

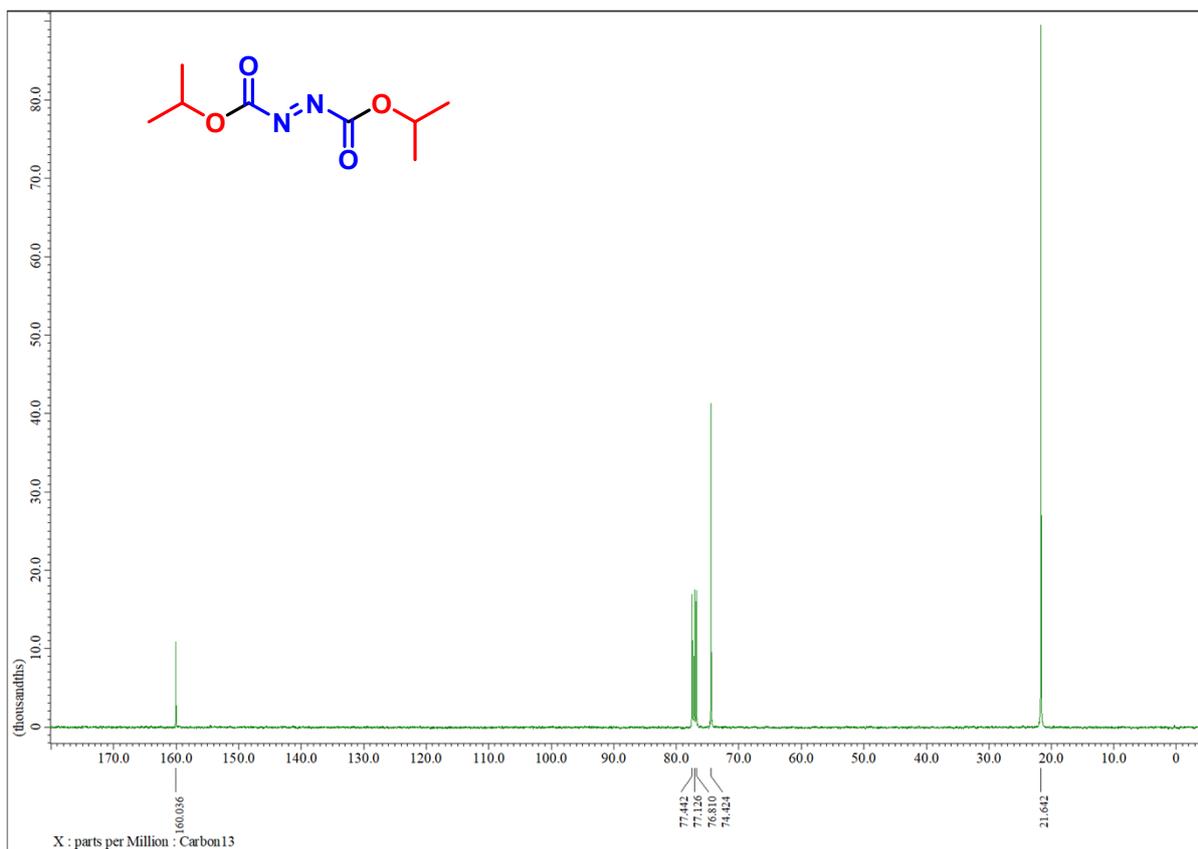


Figure S64. ¹³C NMR spectrum of Di-isopropyl (E)-diazene-1,2-dicarboxylate **C29**.

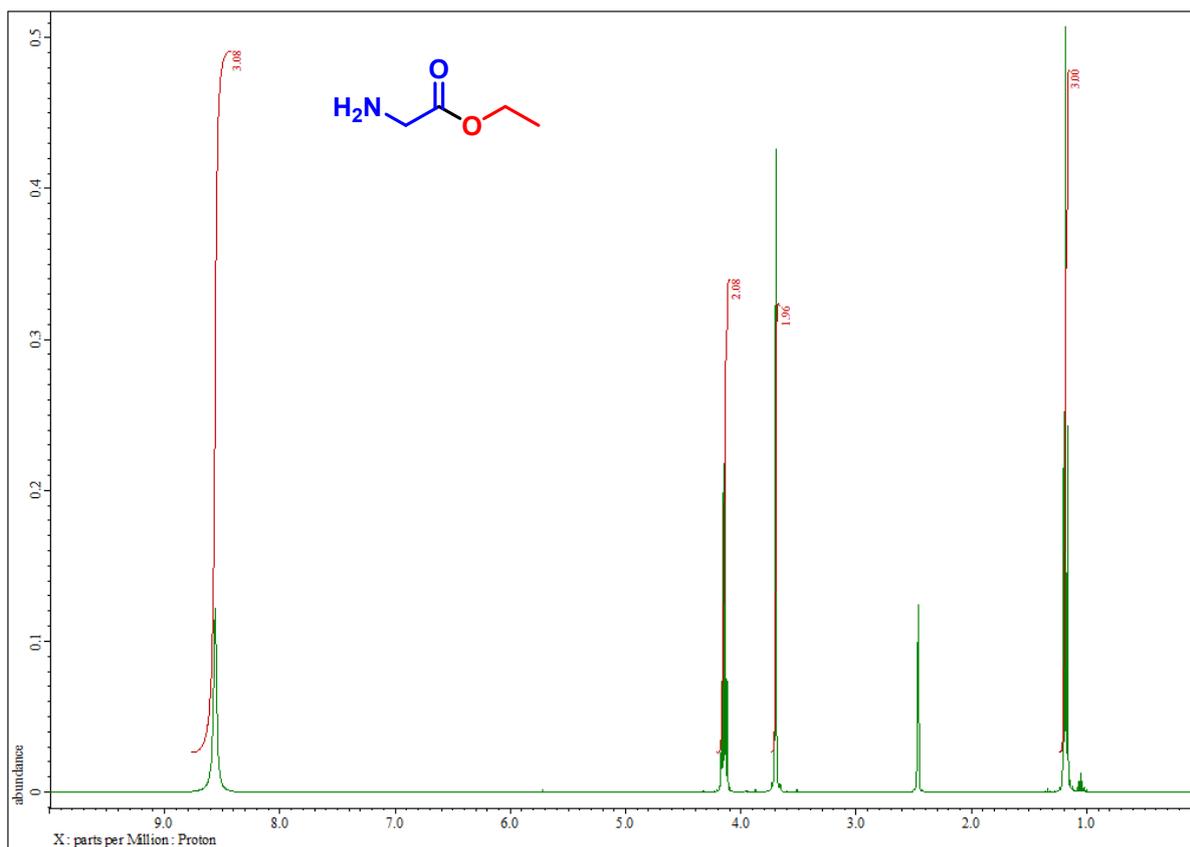


Figure S65. ¹H NMR spectrum of Ethyl glycinate **D1**.

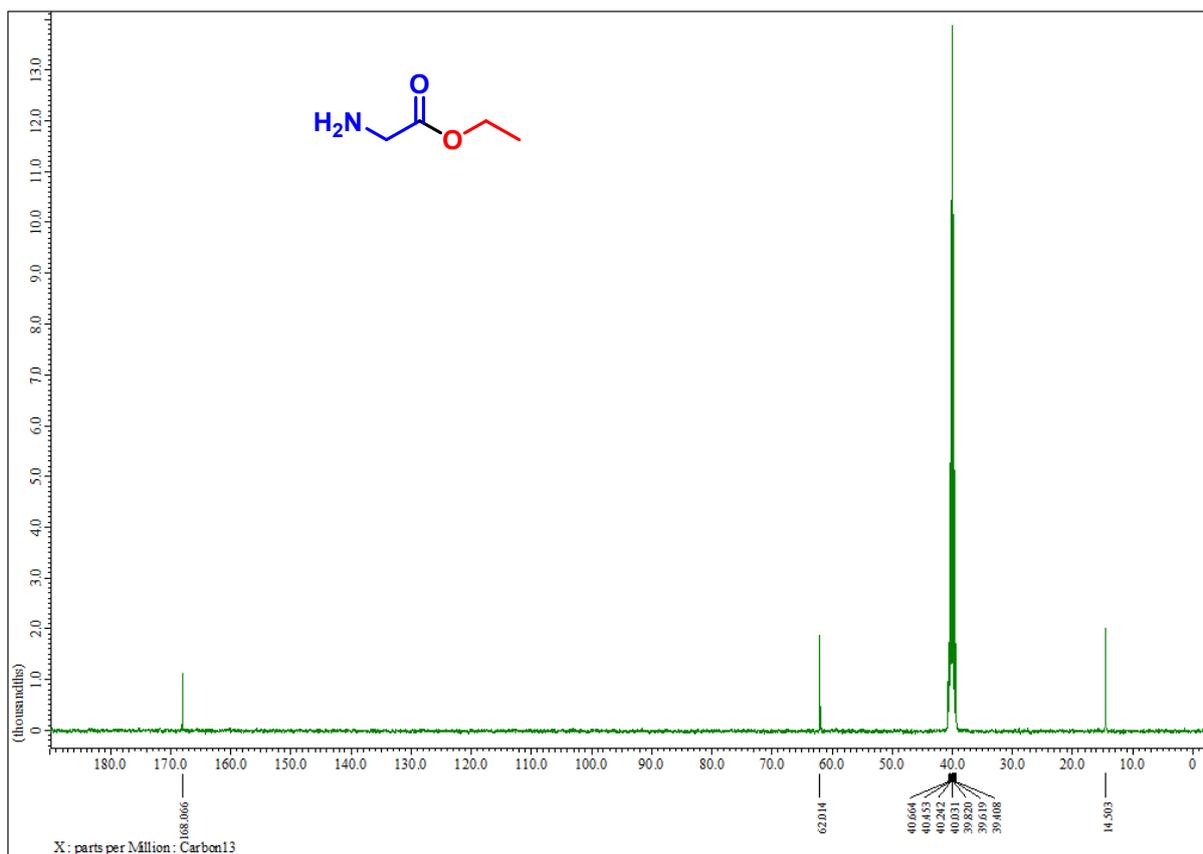


Figure S66. ^{13}C NMR spectrum of Ethyl glycinate D1.

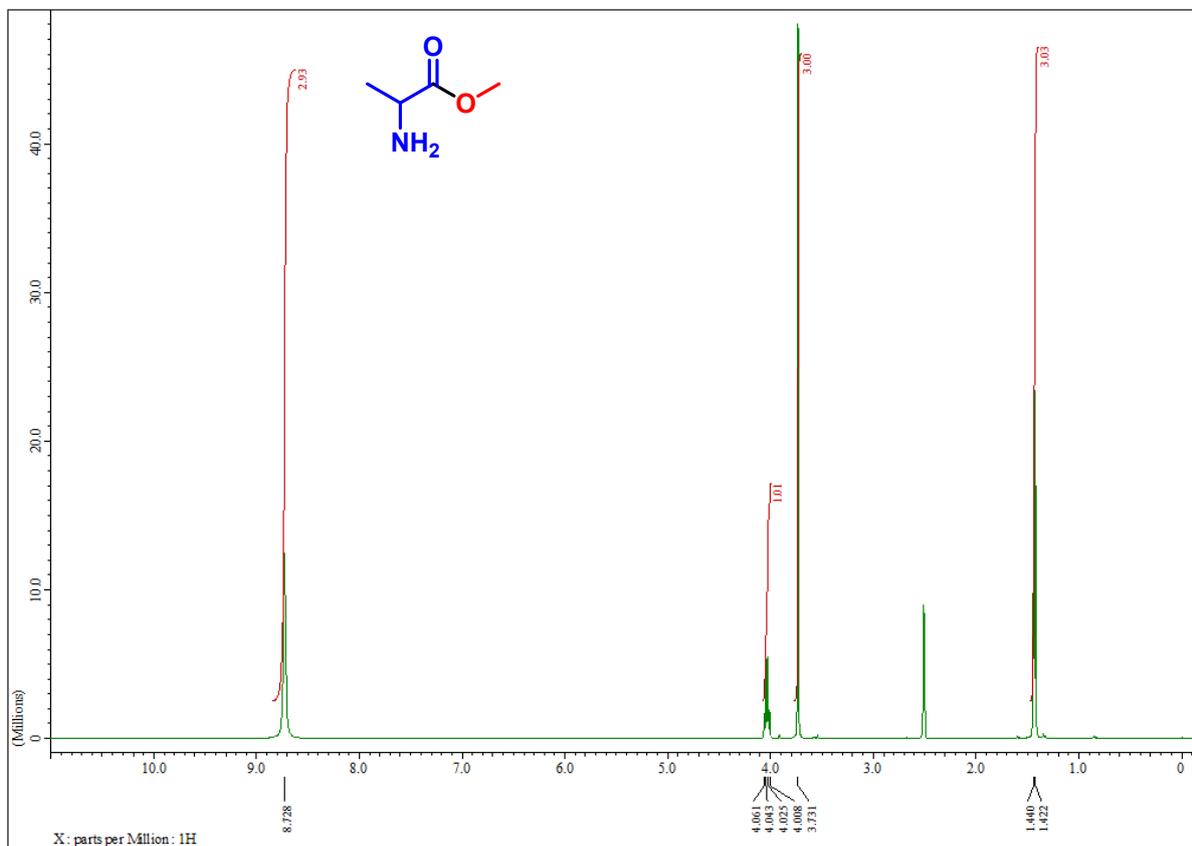


Figure S67. ^1H NMR spectrum of Methyl alaninate **D2**.

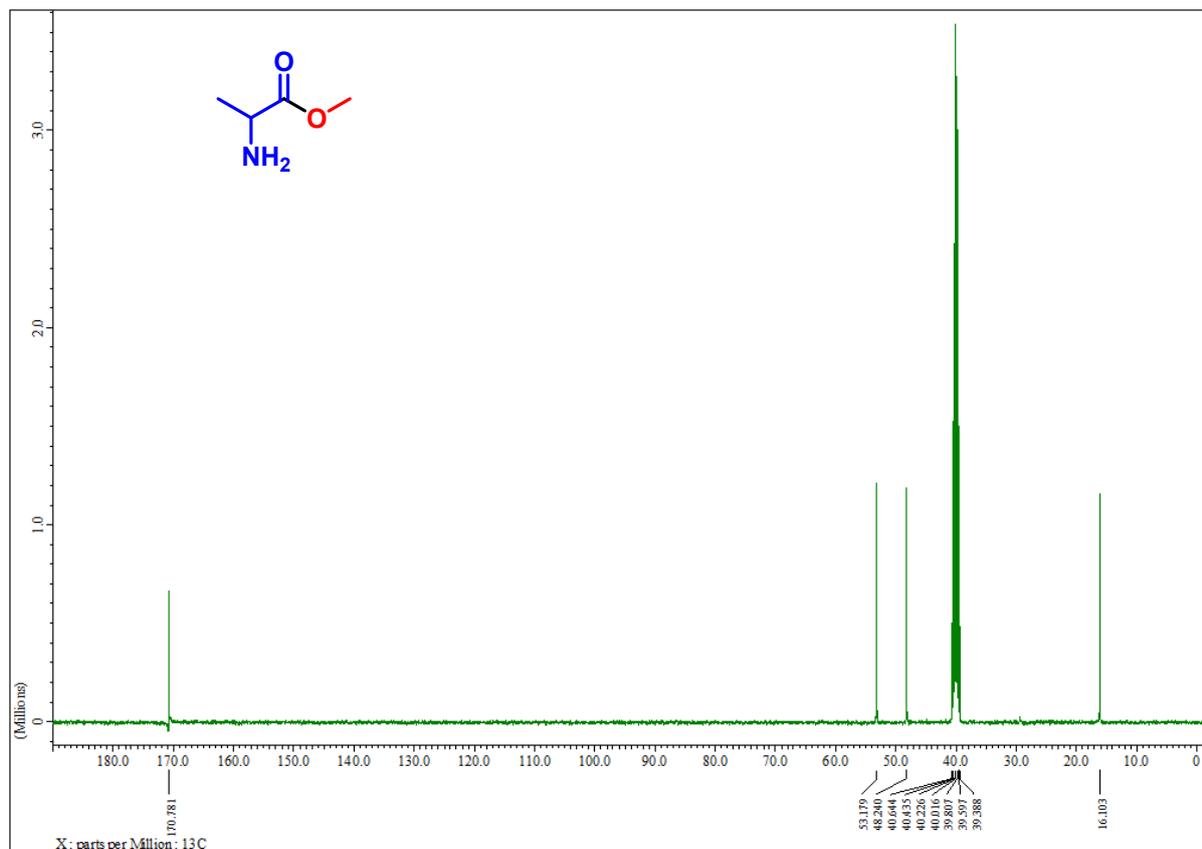


Figure S68. ^{13}C NMR spectrum of Methyl alaninate **D2**.

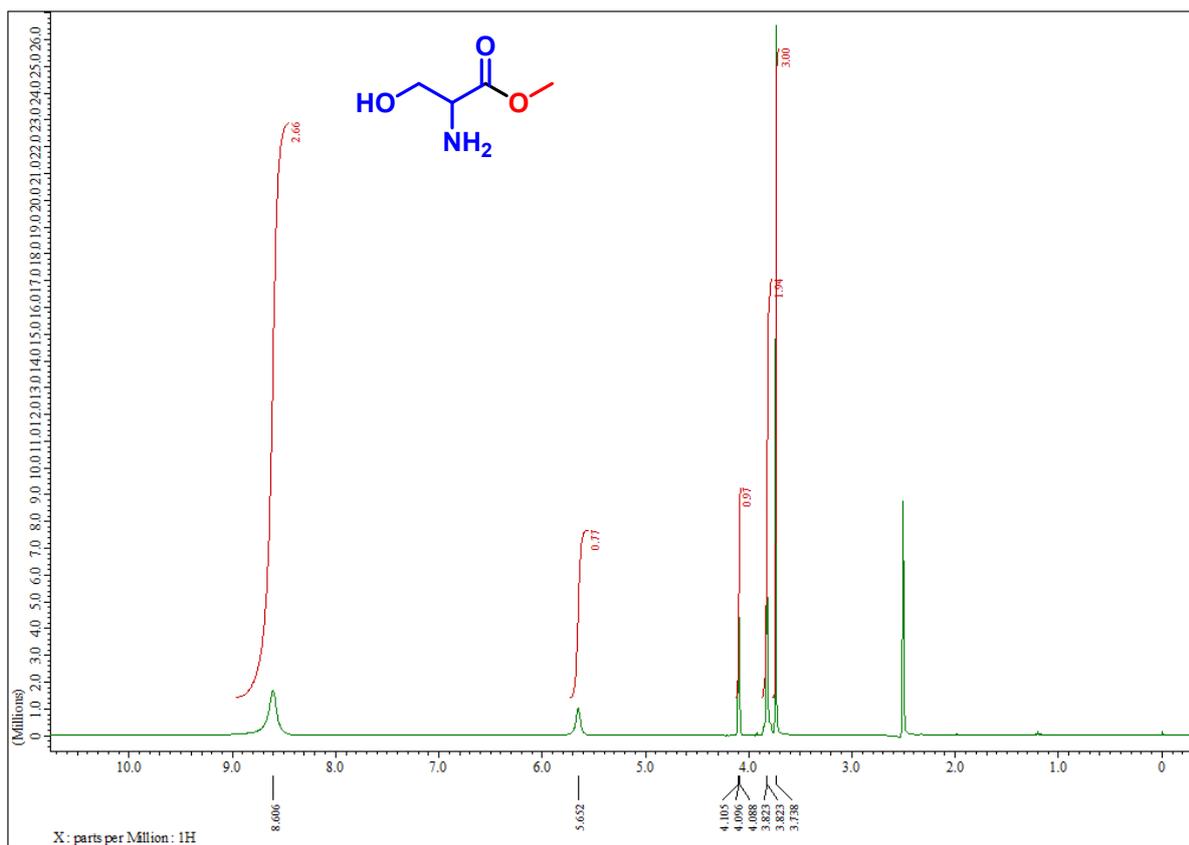


Figure S69. ¹H NMR spectrum of Methyl serinate **D3**.

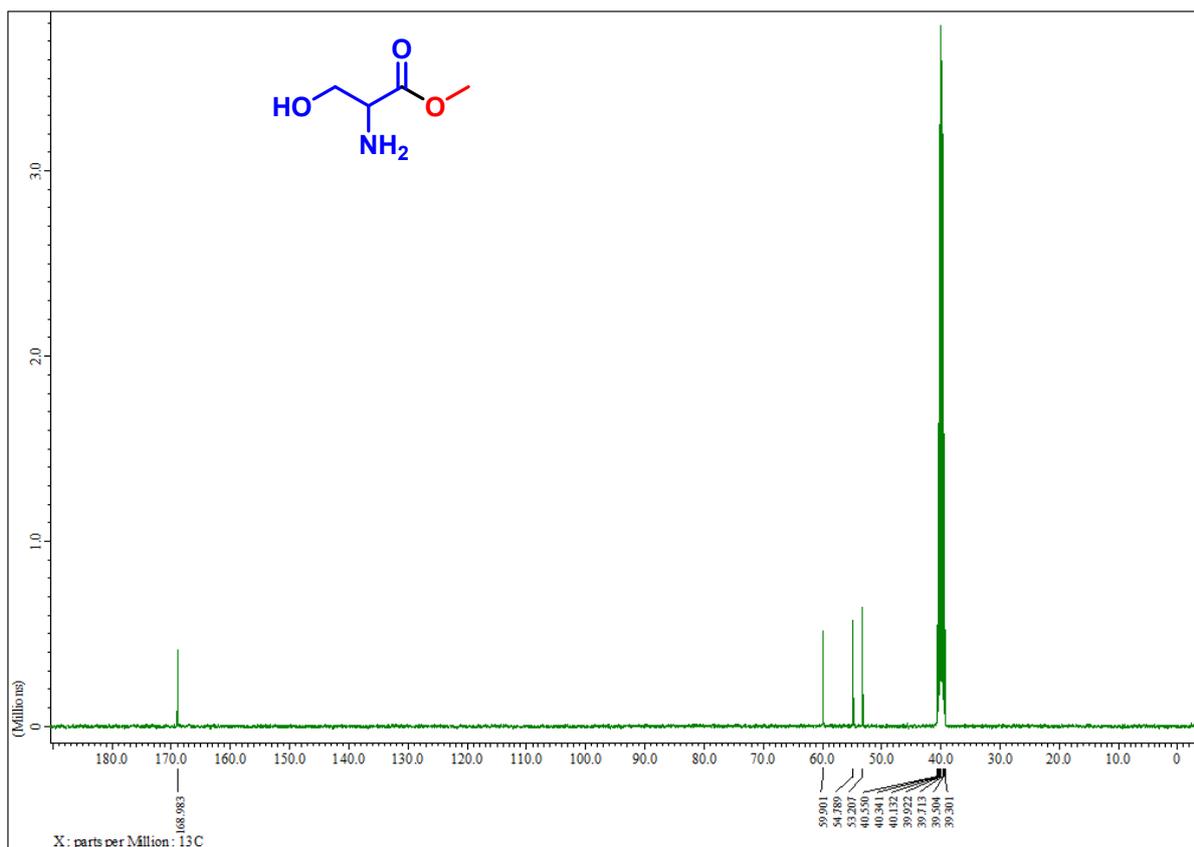


Figure S70. ^{13}C NMR spectrum of Methyl serinate D3.

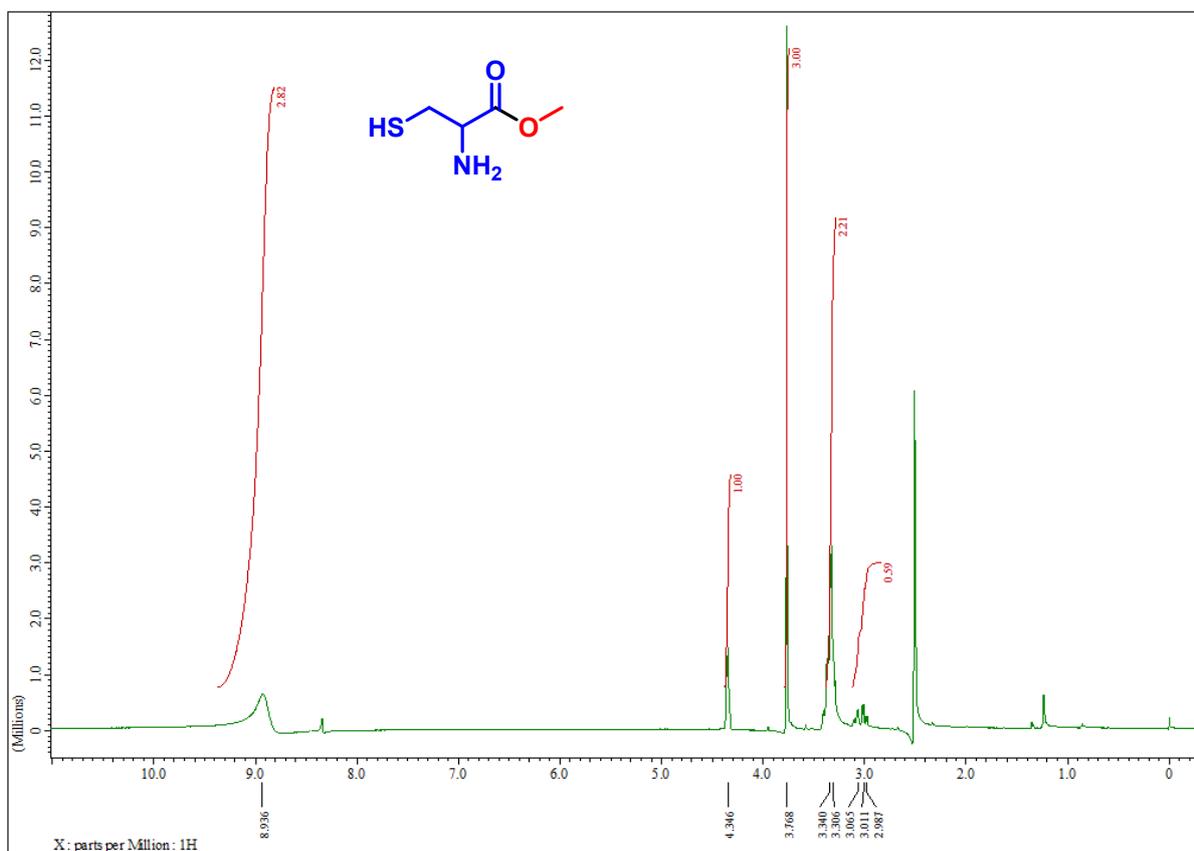


Figure S71. ^1H NMR spectrum of Methyl cysteinate **D4**.

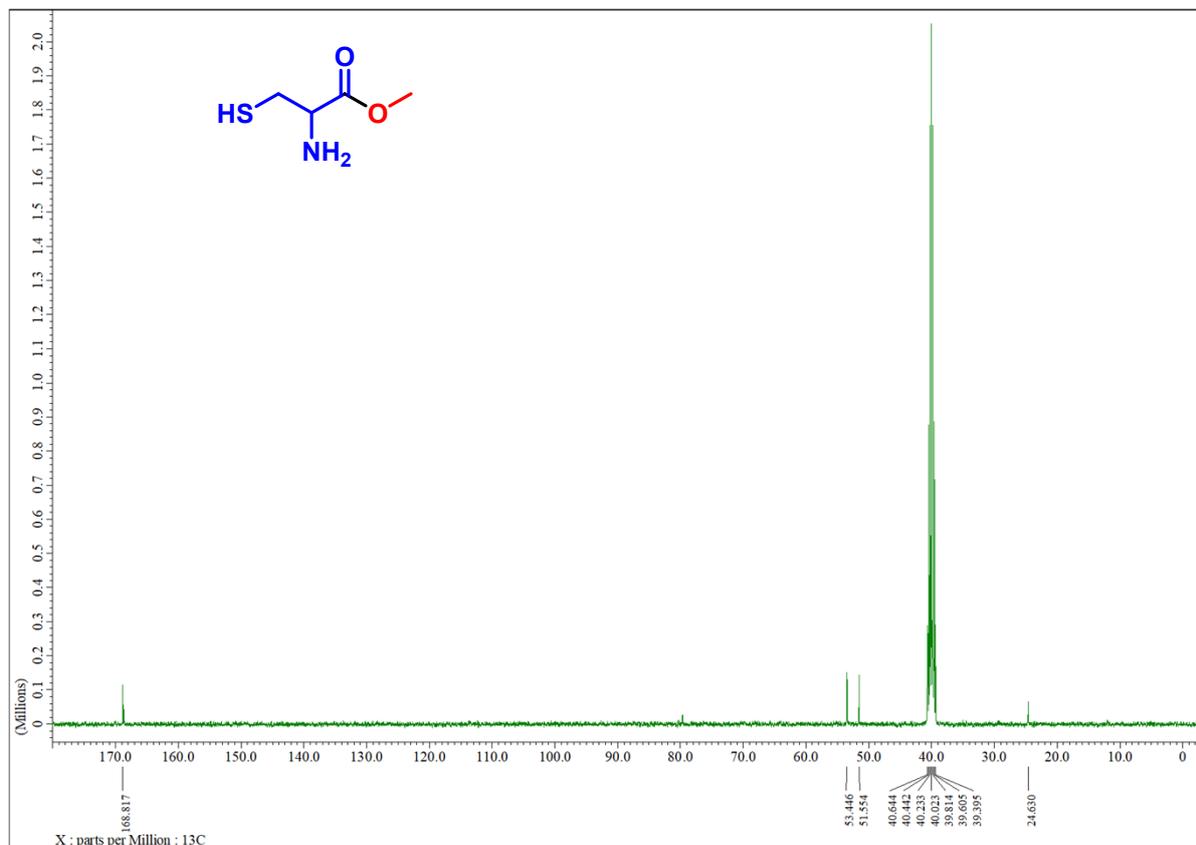


Figure S72. ^{13}C NMR spectrum of Methyl cysteinate **D4**.

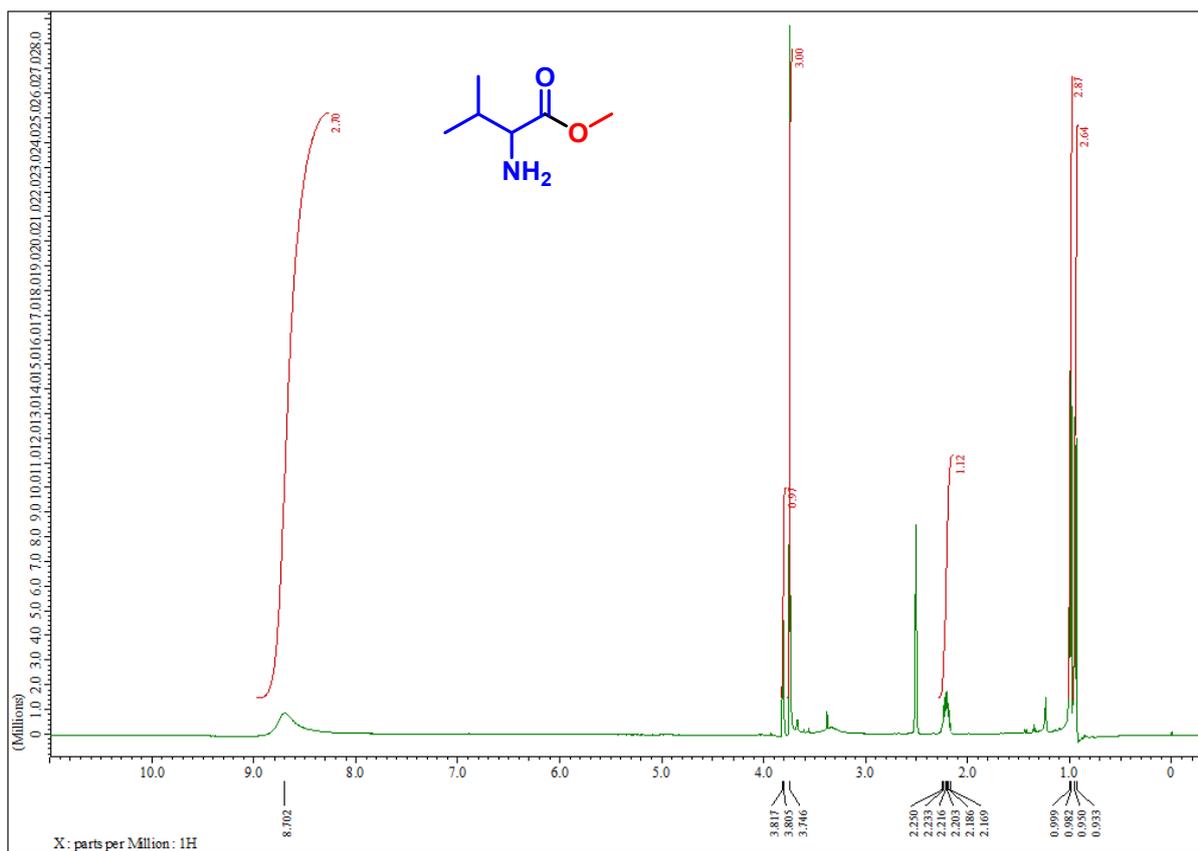


Figure S73. ¹H NMR spectrum of Methyl valinate **D5**.

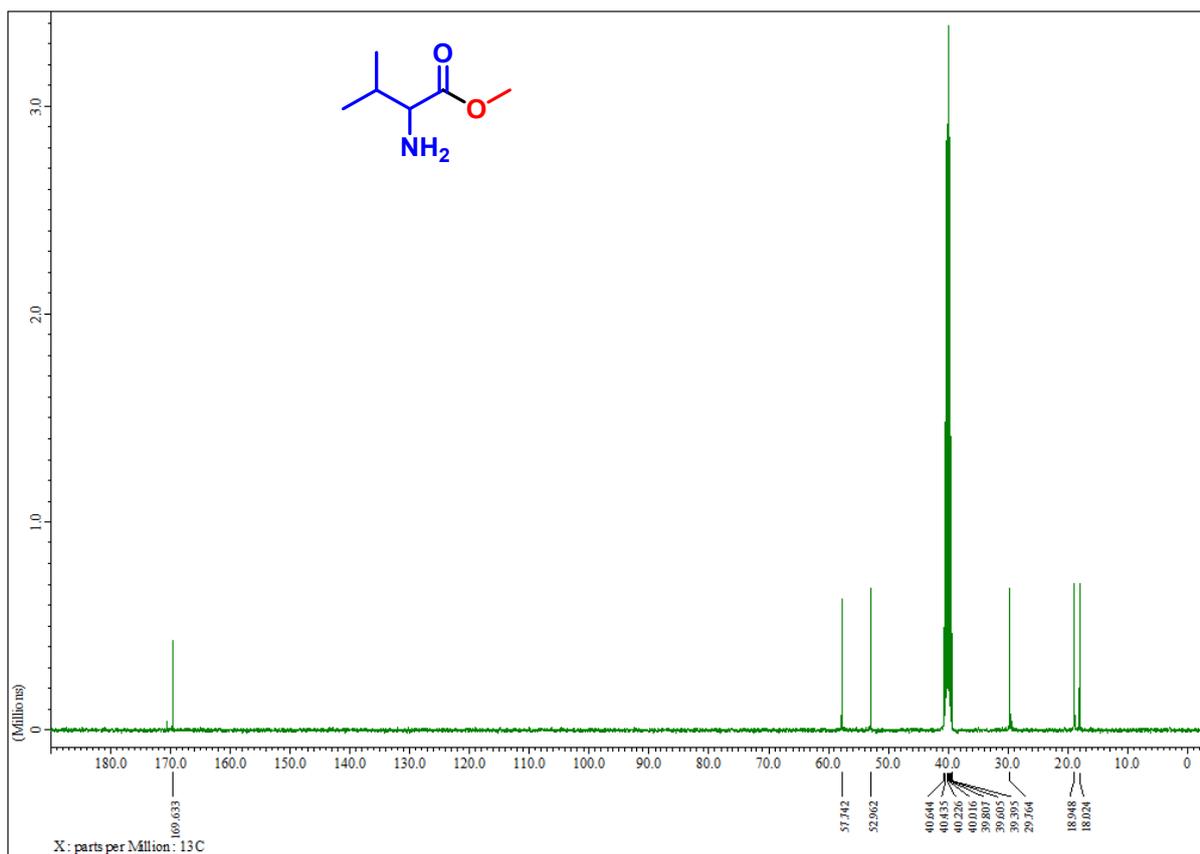


Figure S74. ^{13}C NMR spectrum of Methyl valinate **D5**.

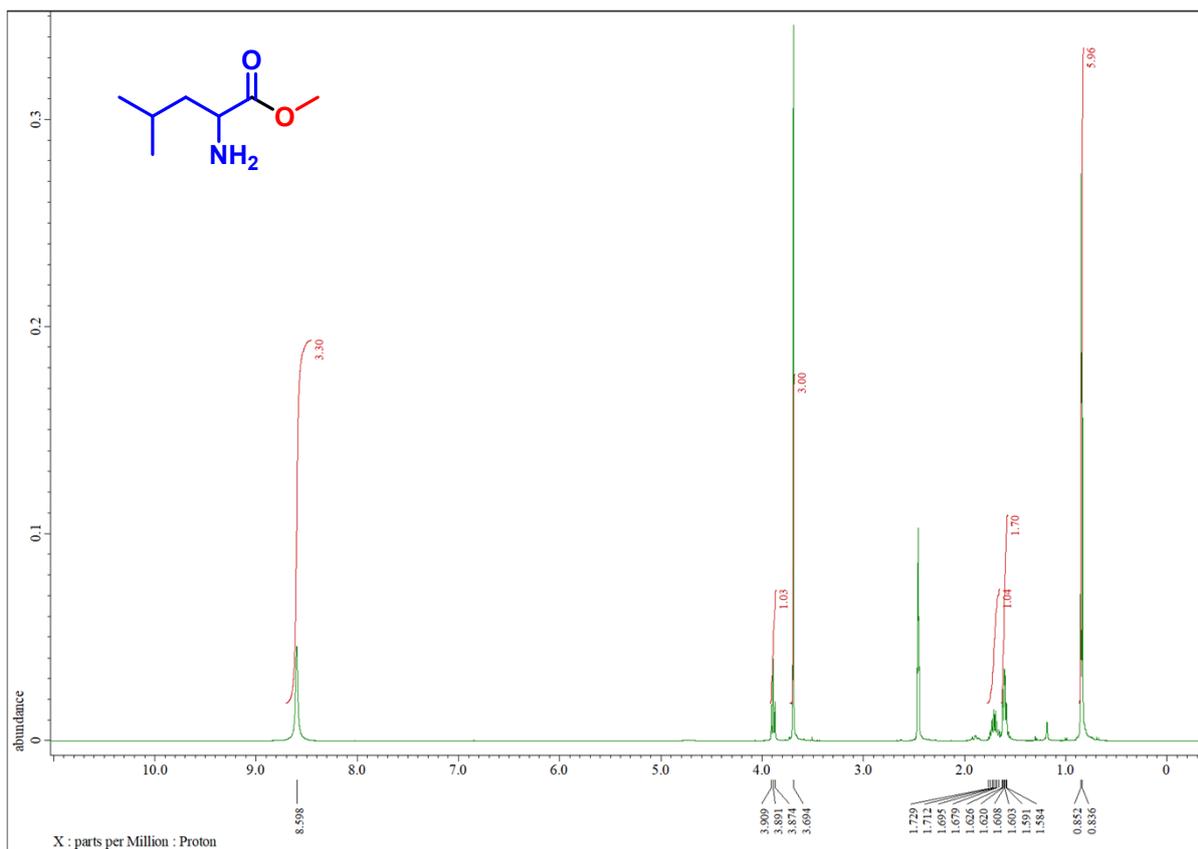


Figure S75. ^1H NMR spectrum of Methyl leucinate **D6**.

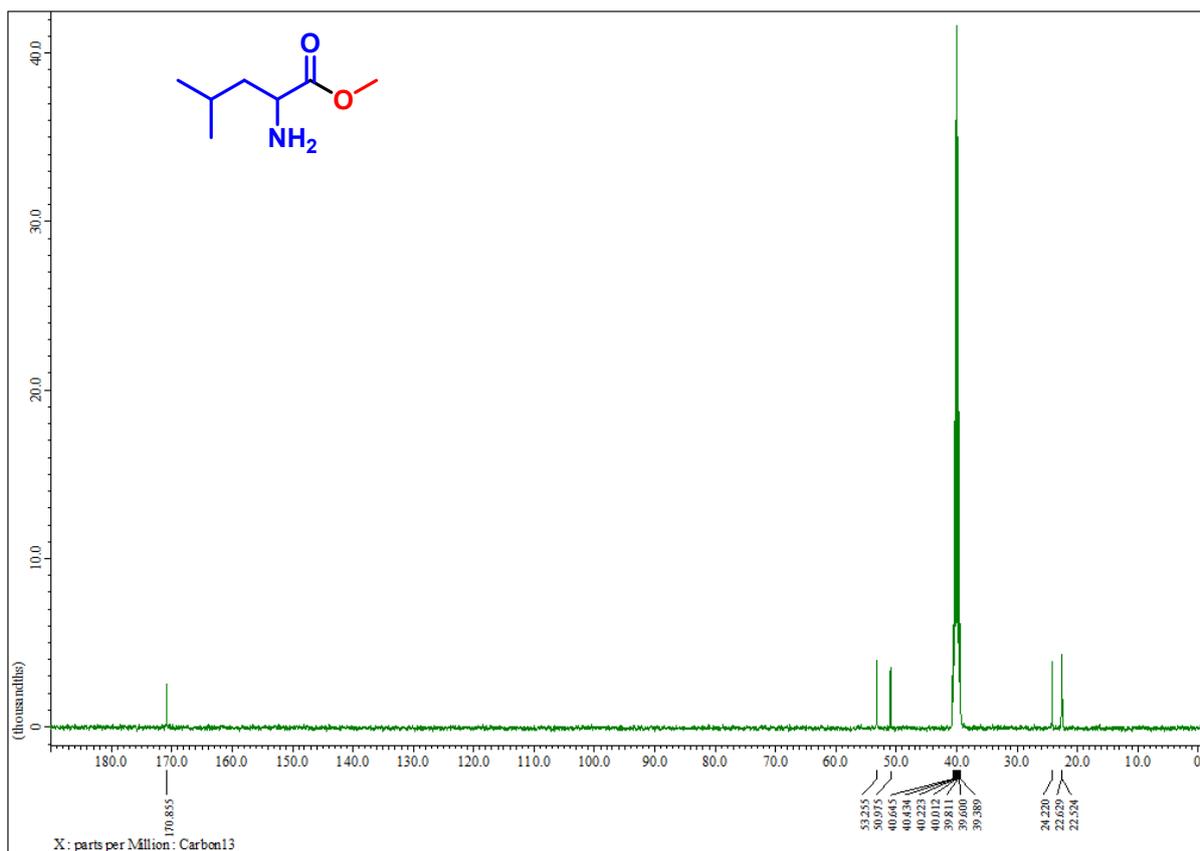


Figure S76. ^{13}C NMR spectrum of Methyl leucinate D6.

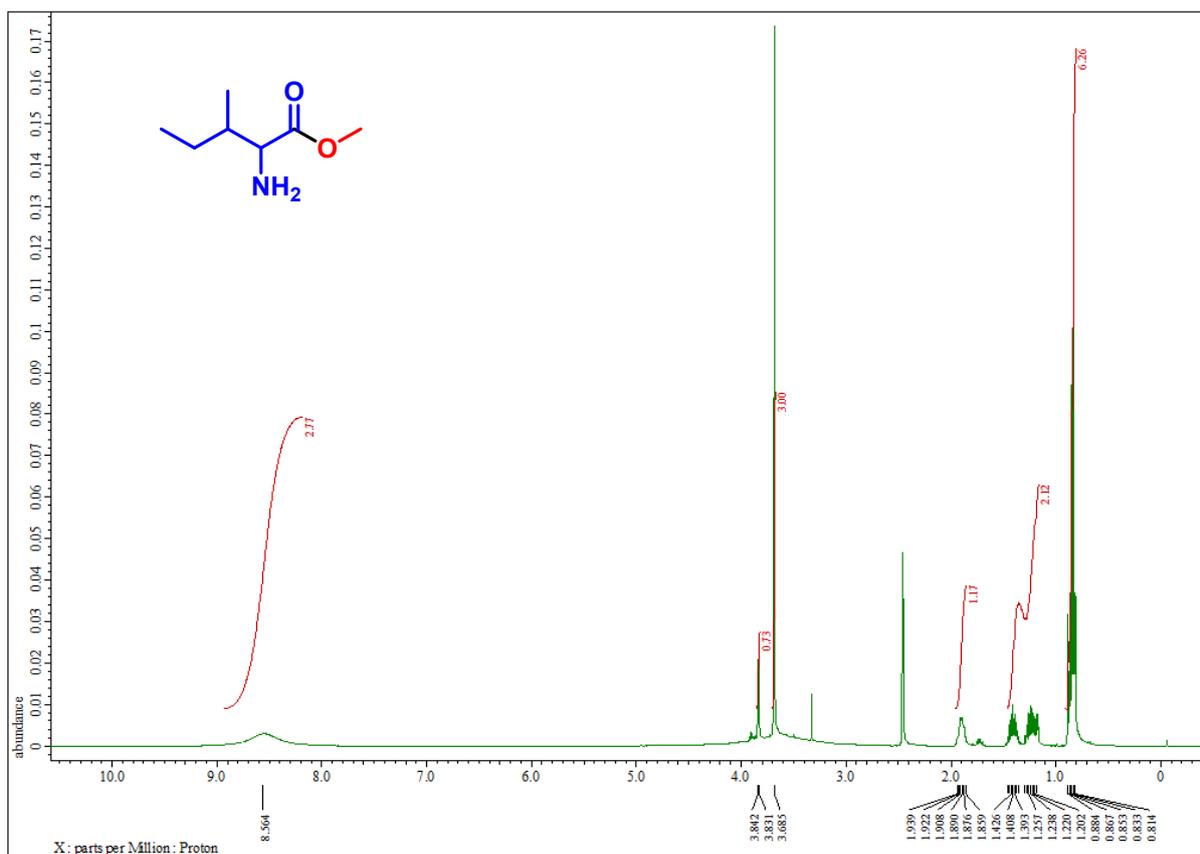


Figure S77. ^1H NMR spectrum of Methyl 2-amino-3-methylpentanoate **D7**.

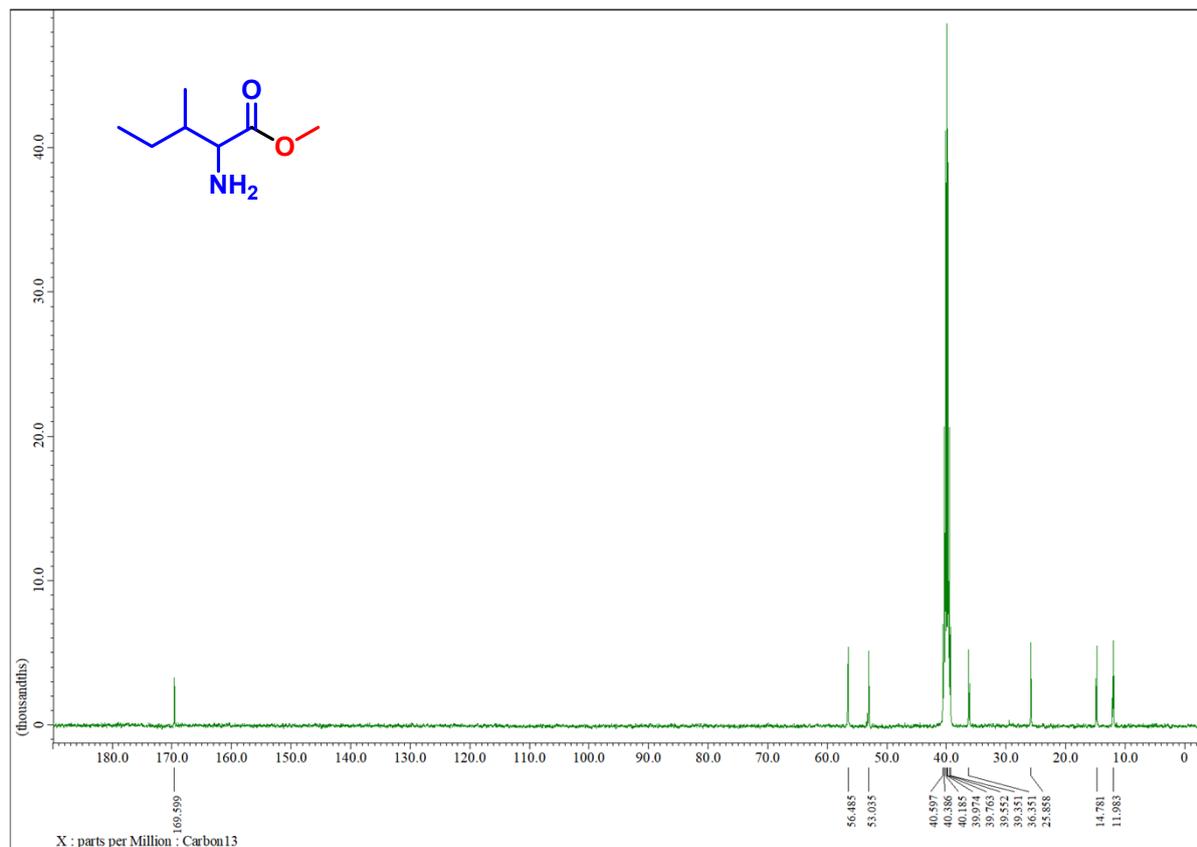


Figure S78. ^{13}C NMR spectrum of Methyl 2-amino-3-methylpentanoate **D7**.

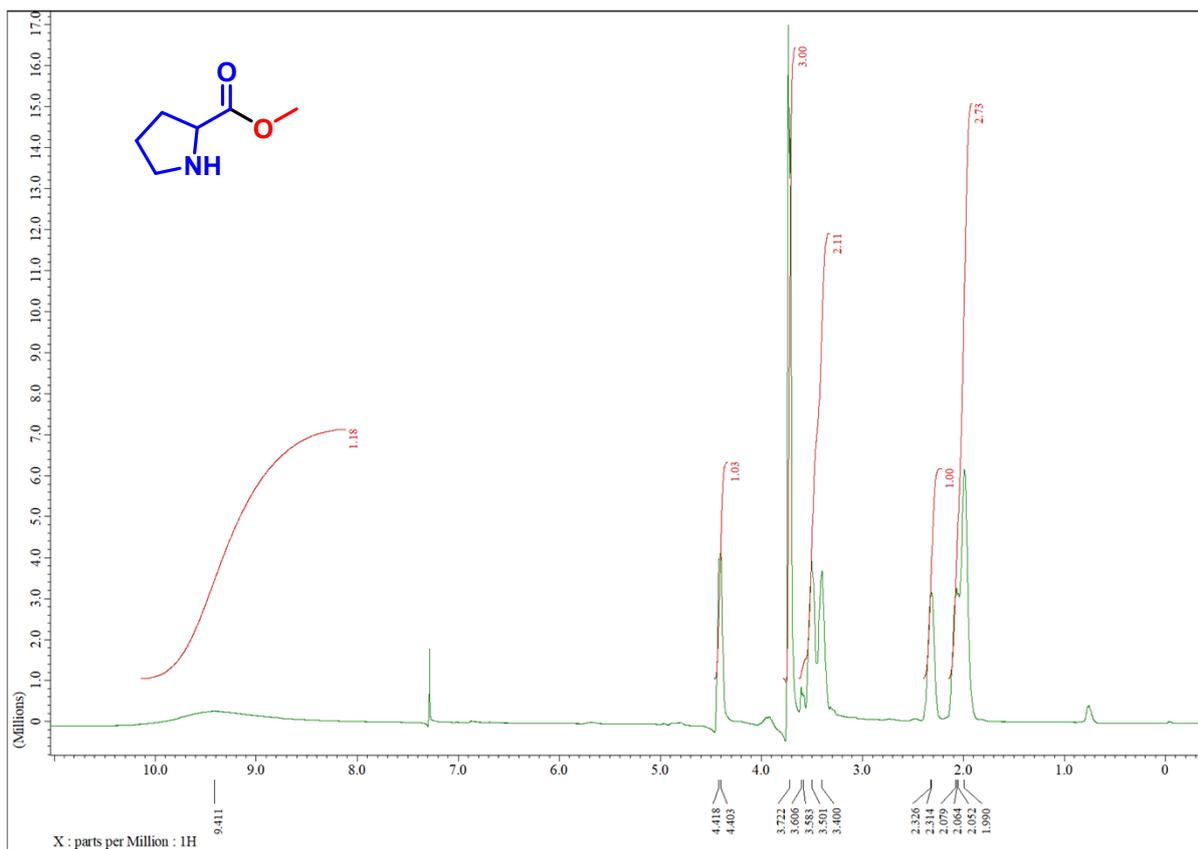


Figure S79. ¹H NMR spectrum of Methyl prolinatate **D8**.

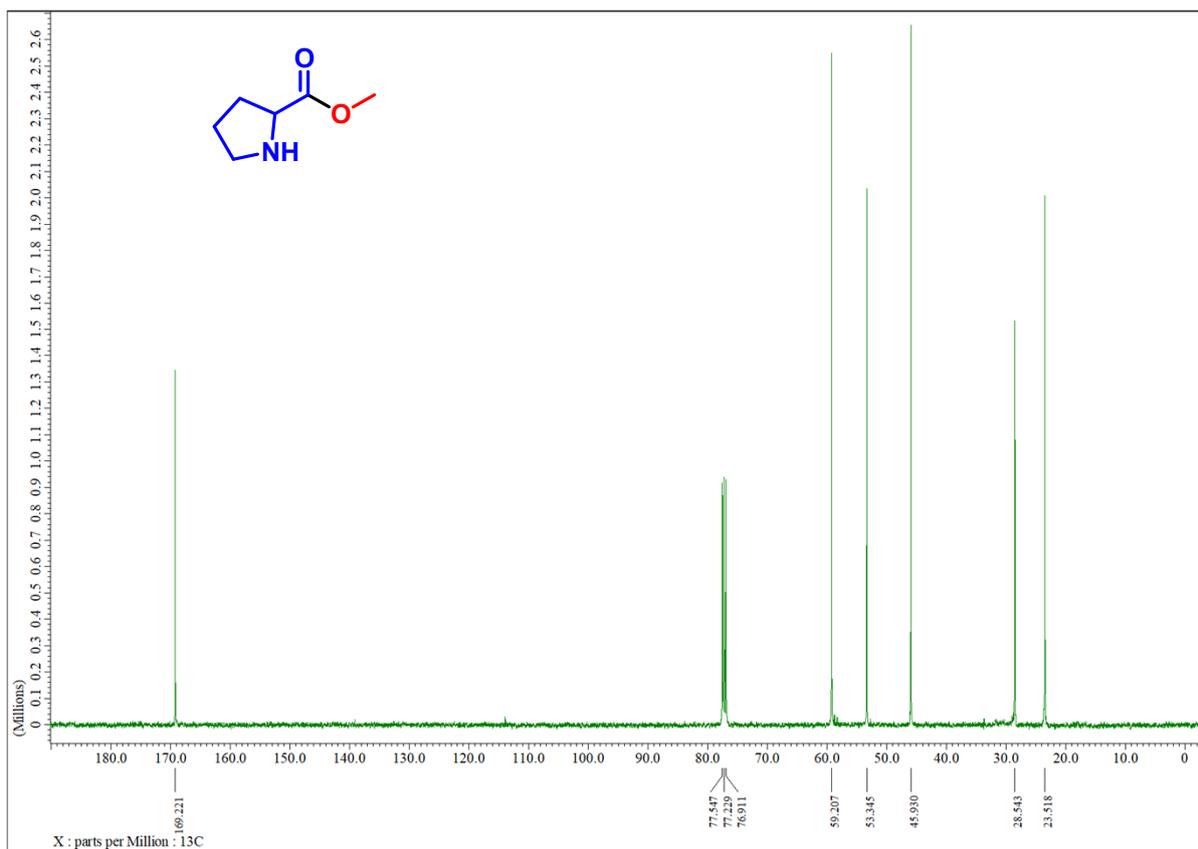


Figure S80. ¹³C NMR spectrum of Methyl proline D8.

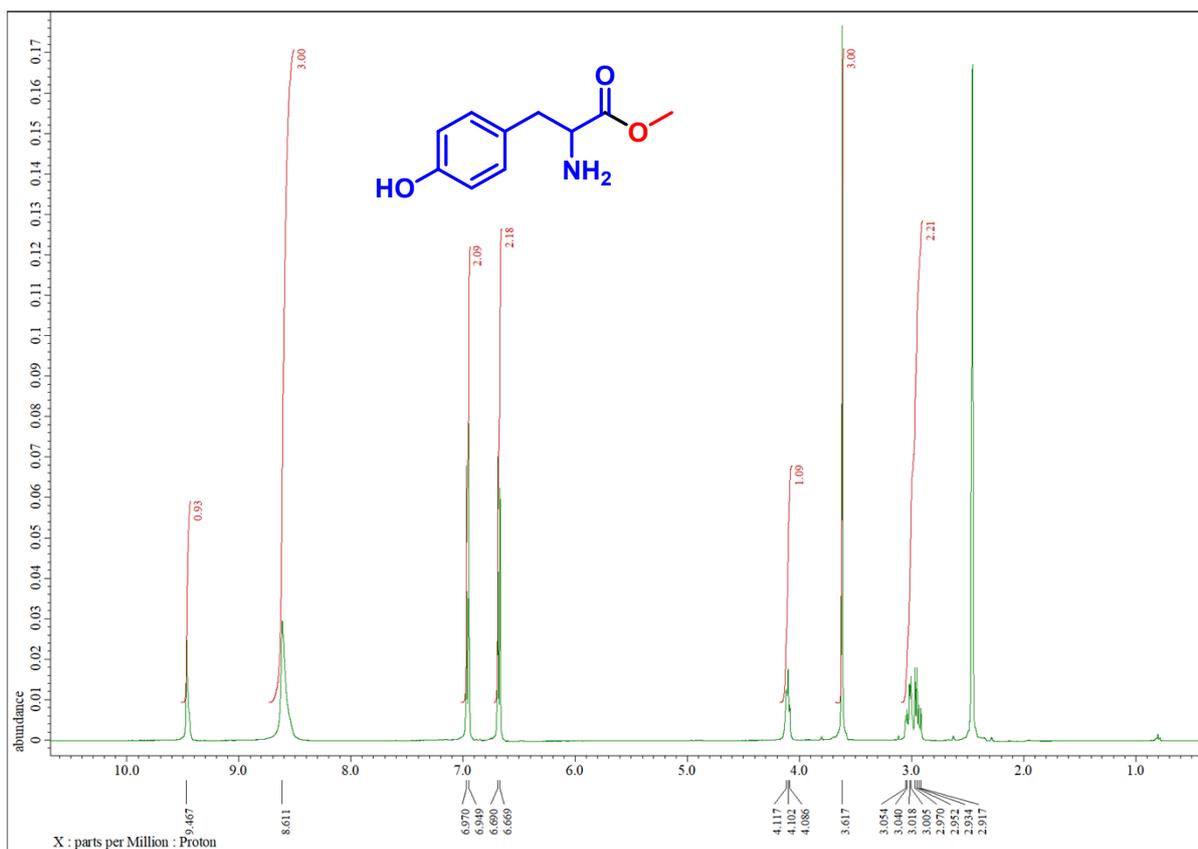


Figure S81. ^1H NMR spectrum of Methyl tyrosinate **D9**.

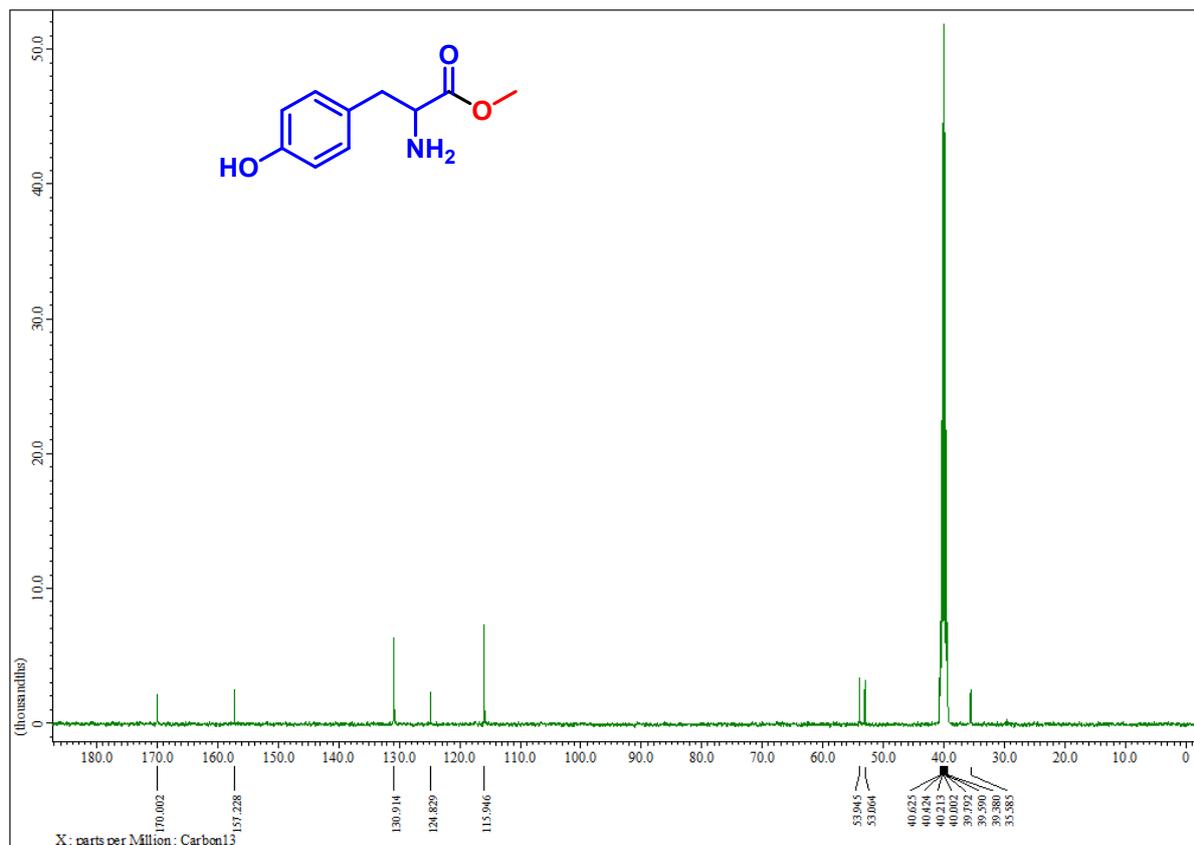


Figure S82. ^{13}C NMR spectrum of Methyl tyrosinate **D9**.

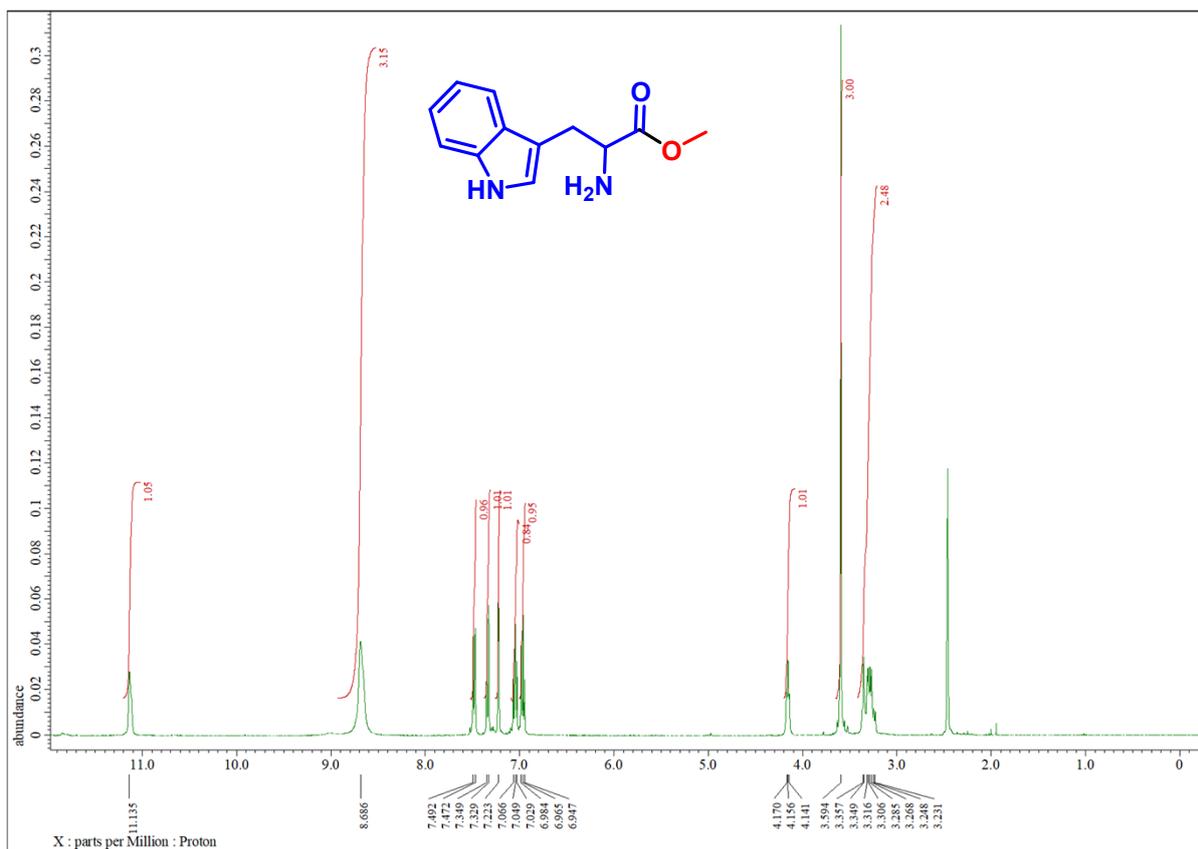


Figure S83. ¹H NMR spectrum of Methyl tryptophanate **D10**.

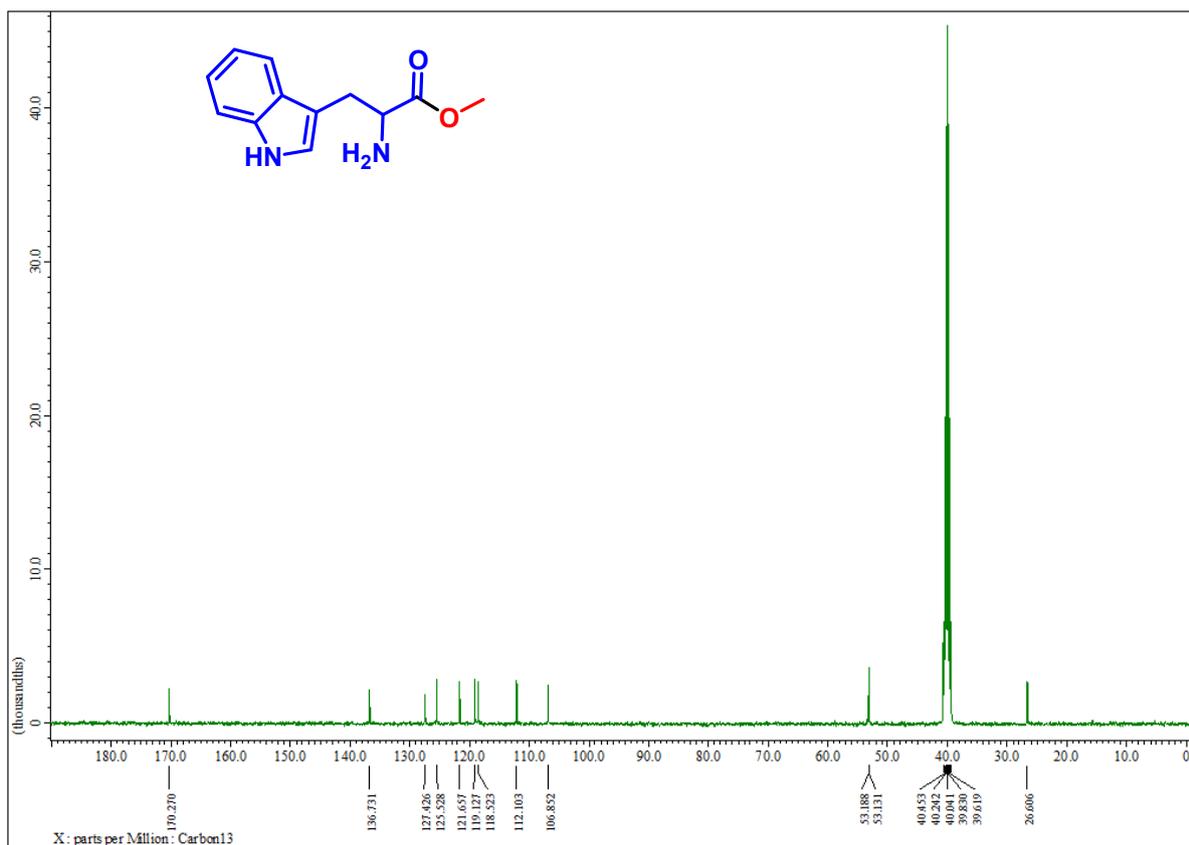


Figure S84. ^{13}C NMR spectrum of Methyl tryptophanate **D10**.

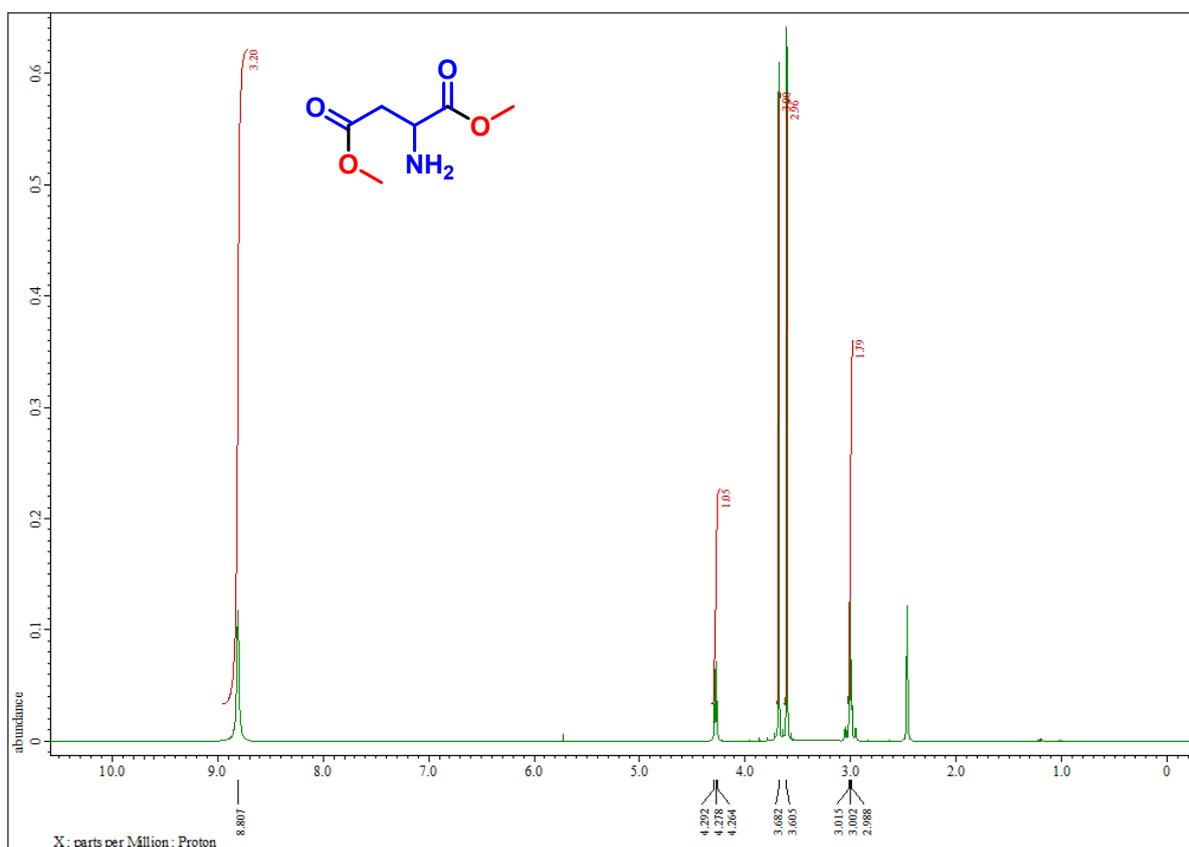


Figure S85. ^1H NMR spectrum of Dimethyl aspartate **D11**.

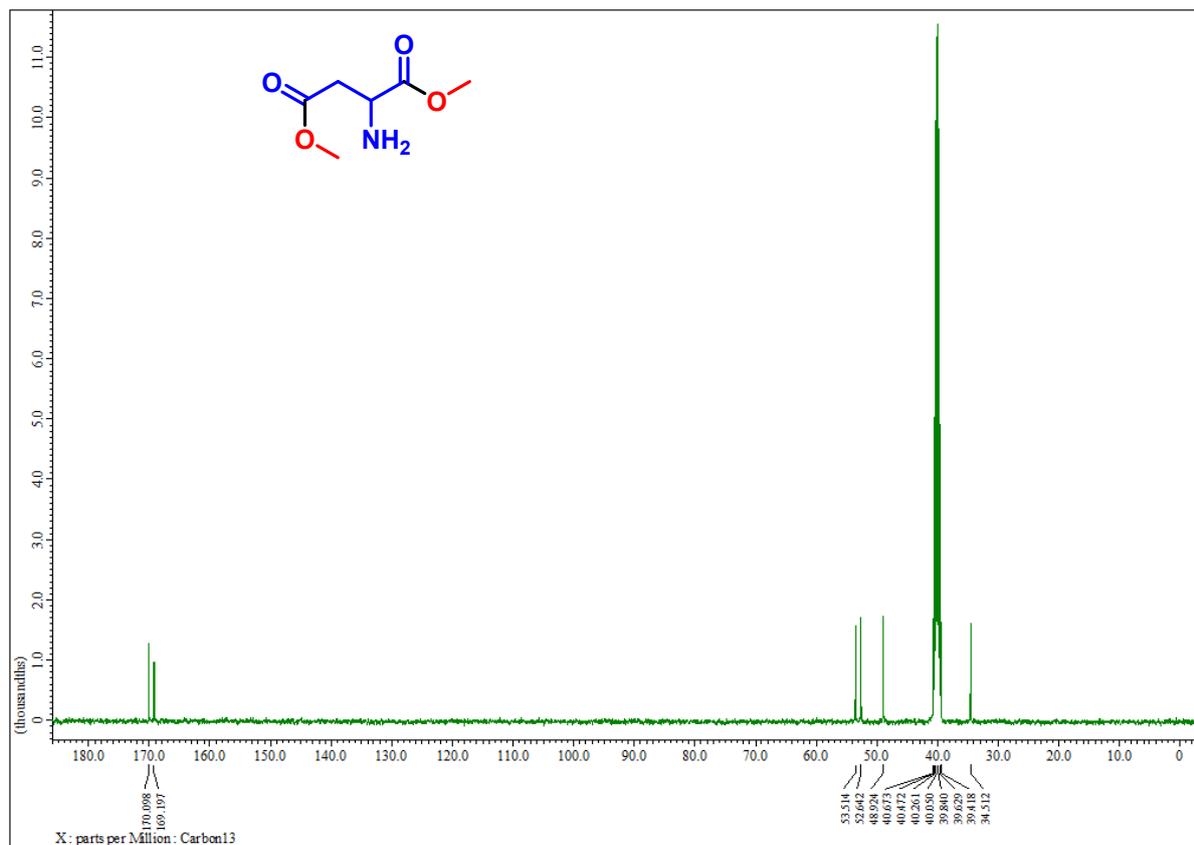


Figure S86. ^{13}C NMR spectrum of Dimethyl aspartate **D11**.

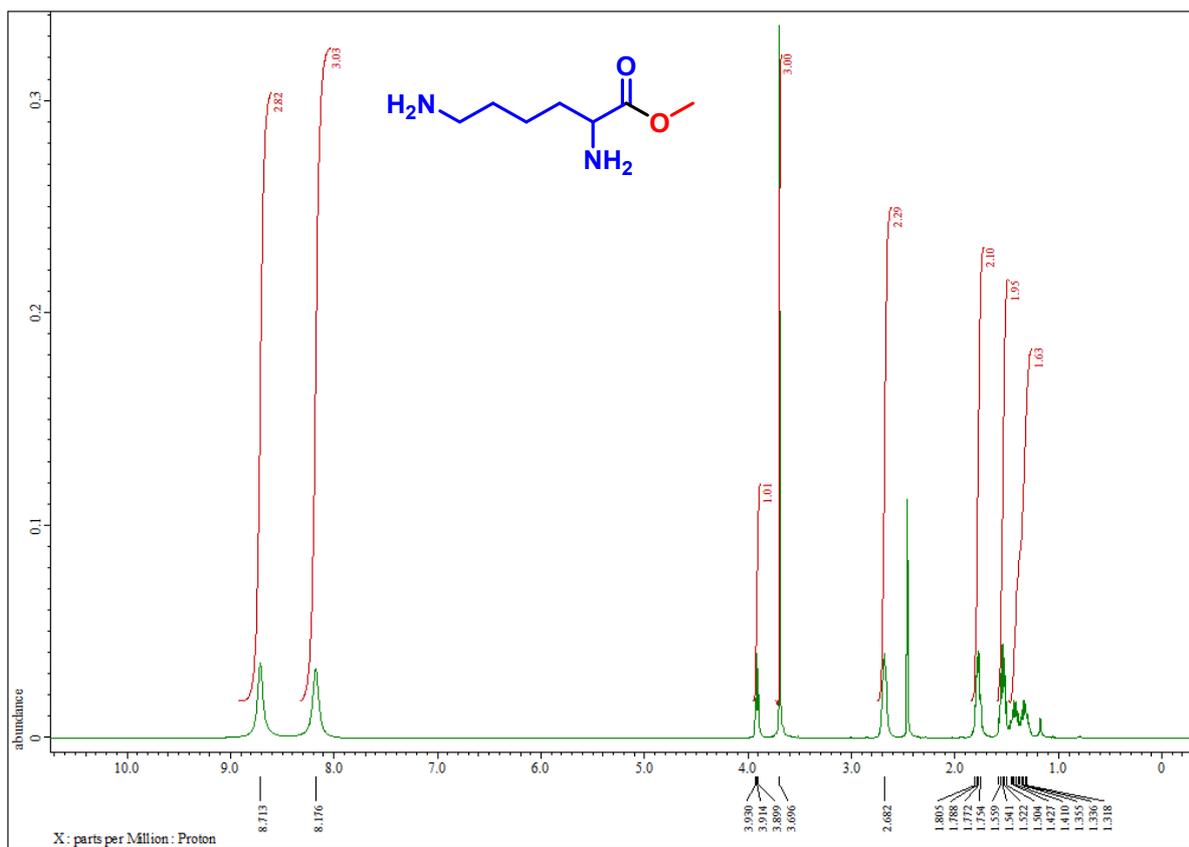


Figure S87. ^1H NMR spectrum of Methyl lysinate D12.

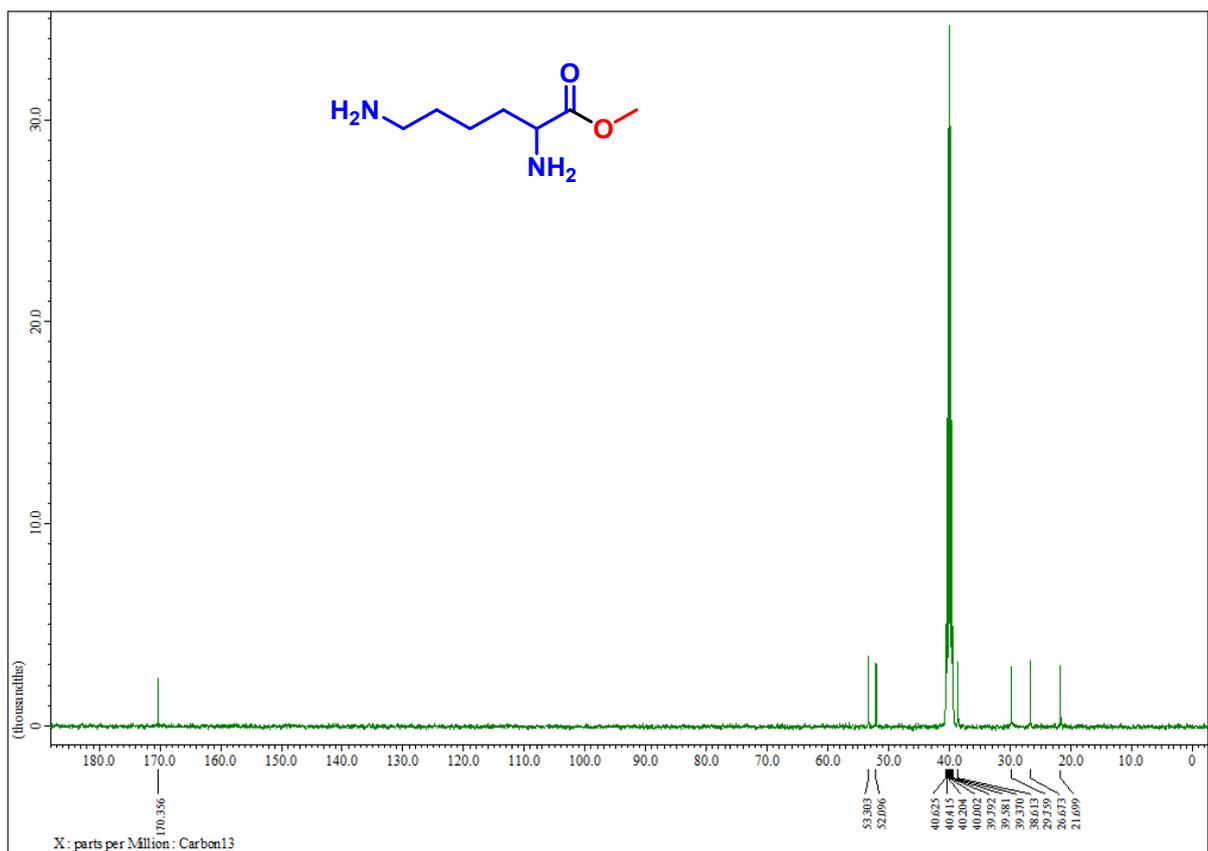


Figure S88. ¹³C NMR spectrum of Methyl lysinate D12.

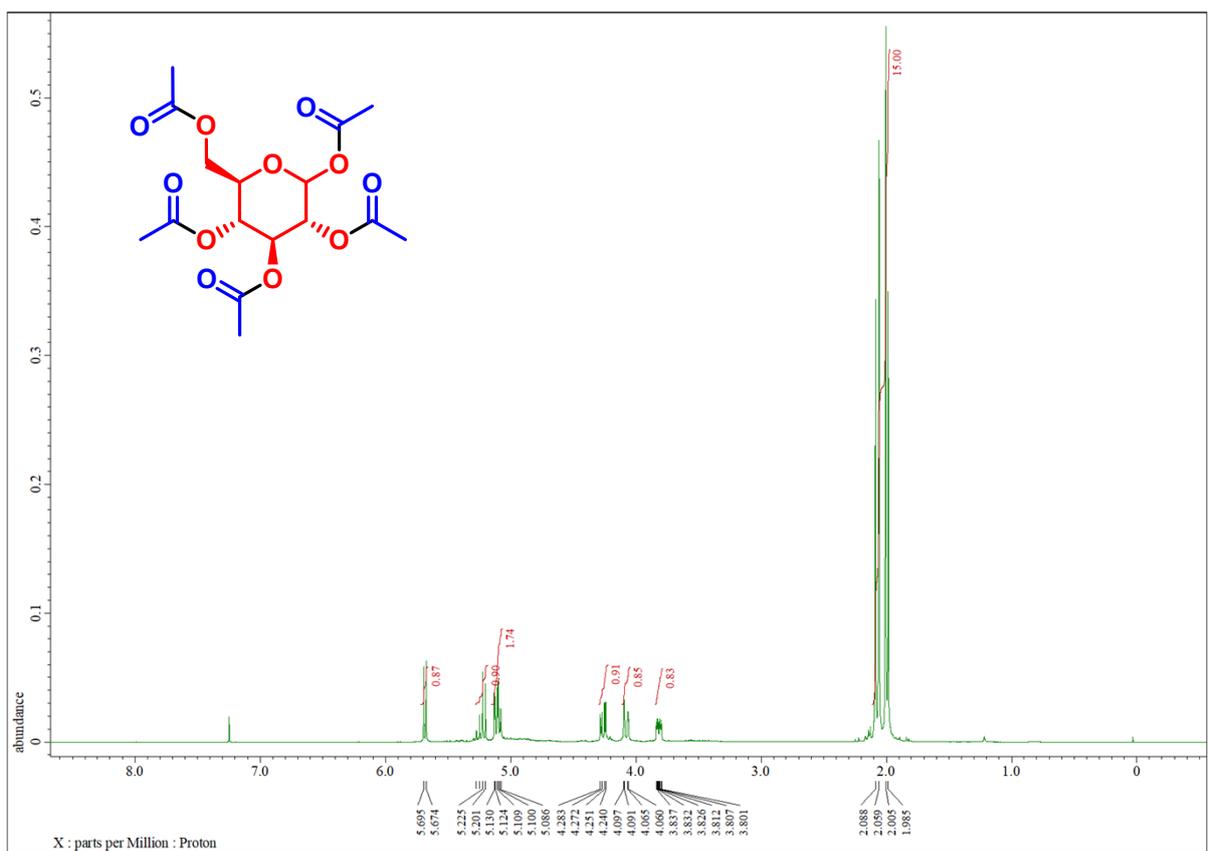


Figure S89. ^1H NMR spectrum of Glucose pentaacetate **E1**.

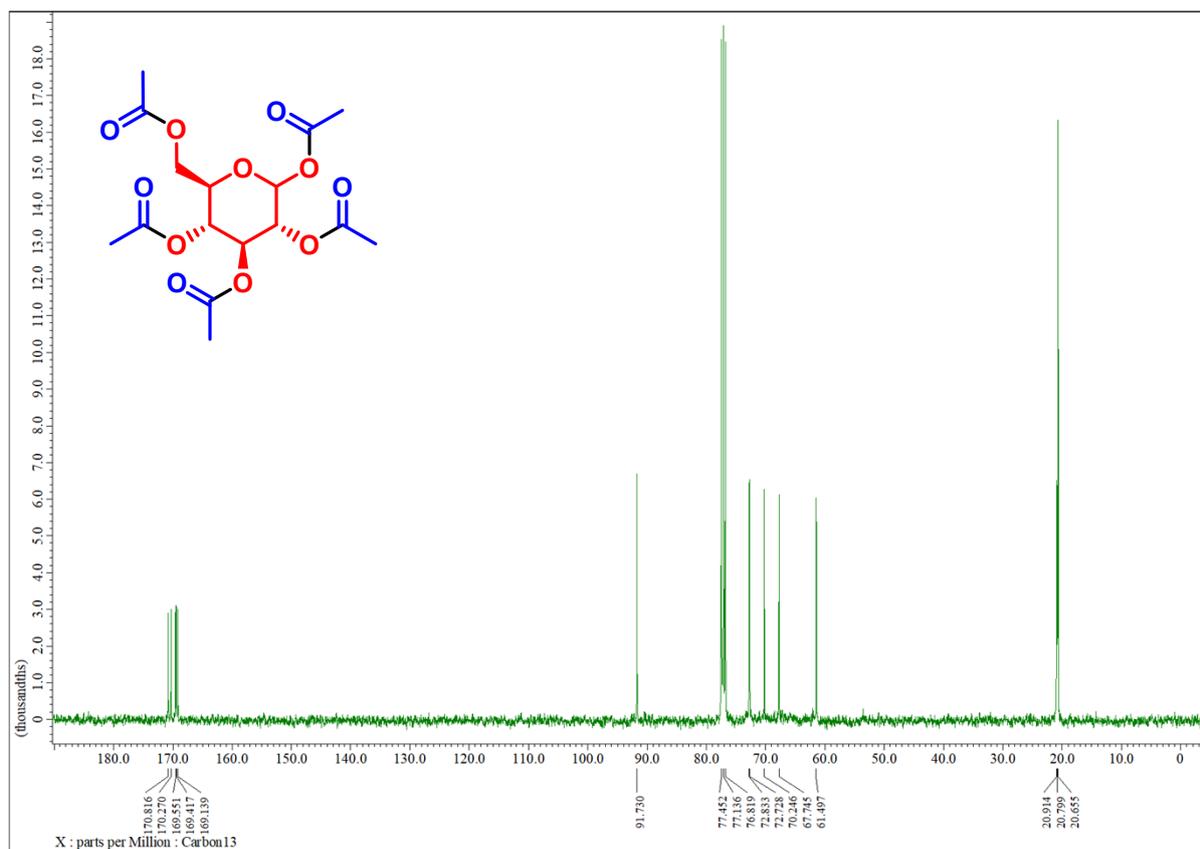


Figure S90. ^{13}C NMR spectrum of Glucose pentaacetate **E1**.

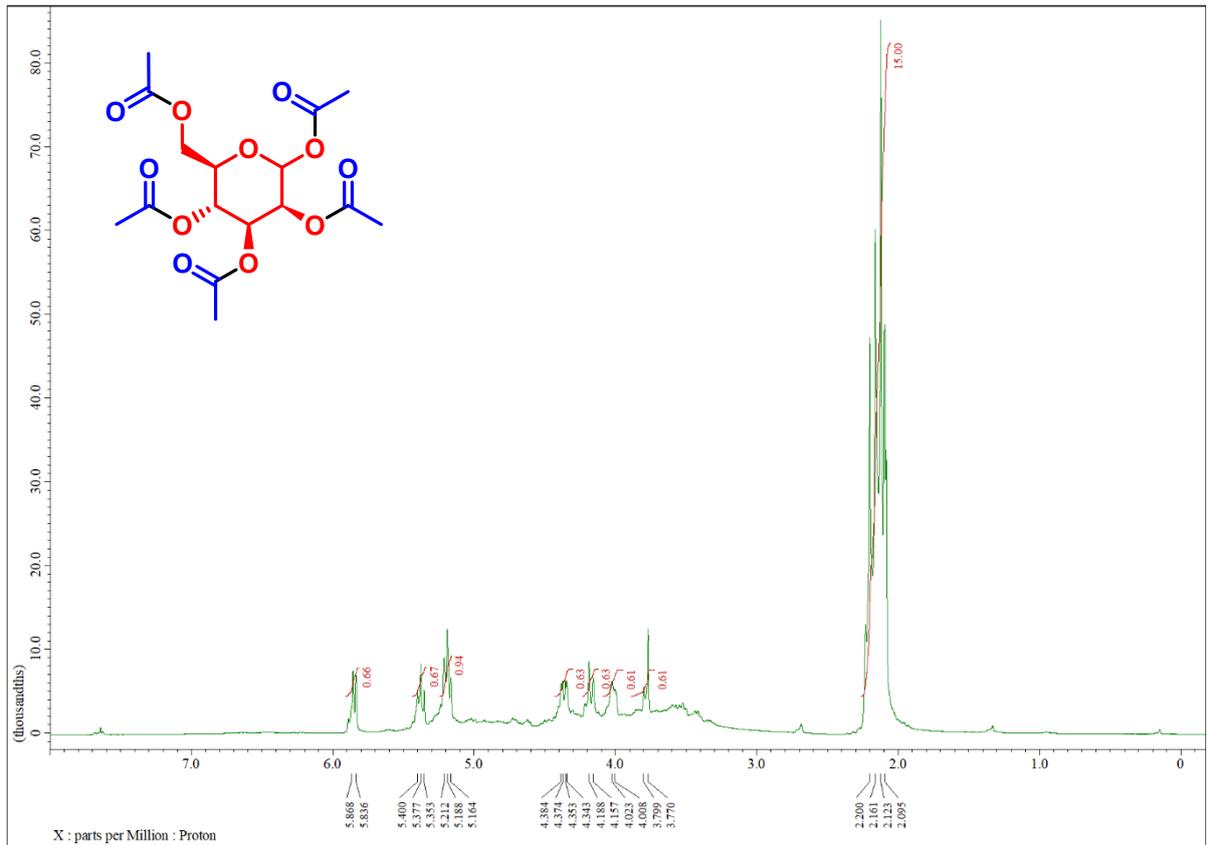


Figure S91. ¹H NMR spectrum of Mannose pentaacetate **E2**.

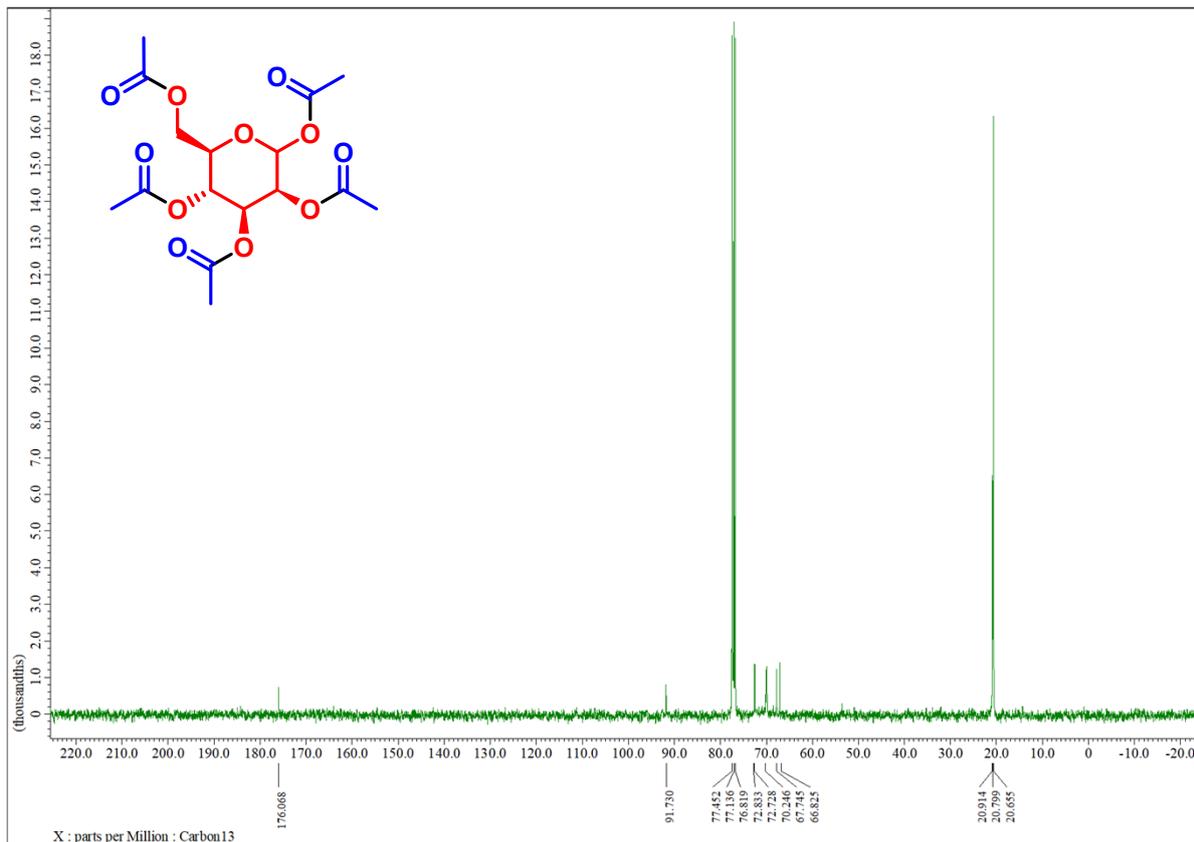


Figure S92. ^{13}C NMR spectrum of Mannose pentaacetate **E2**.

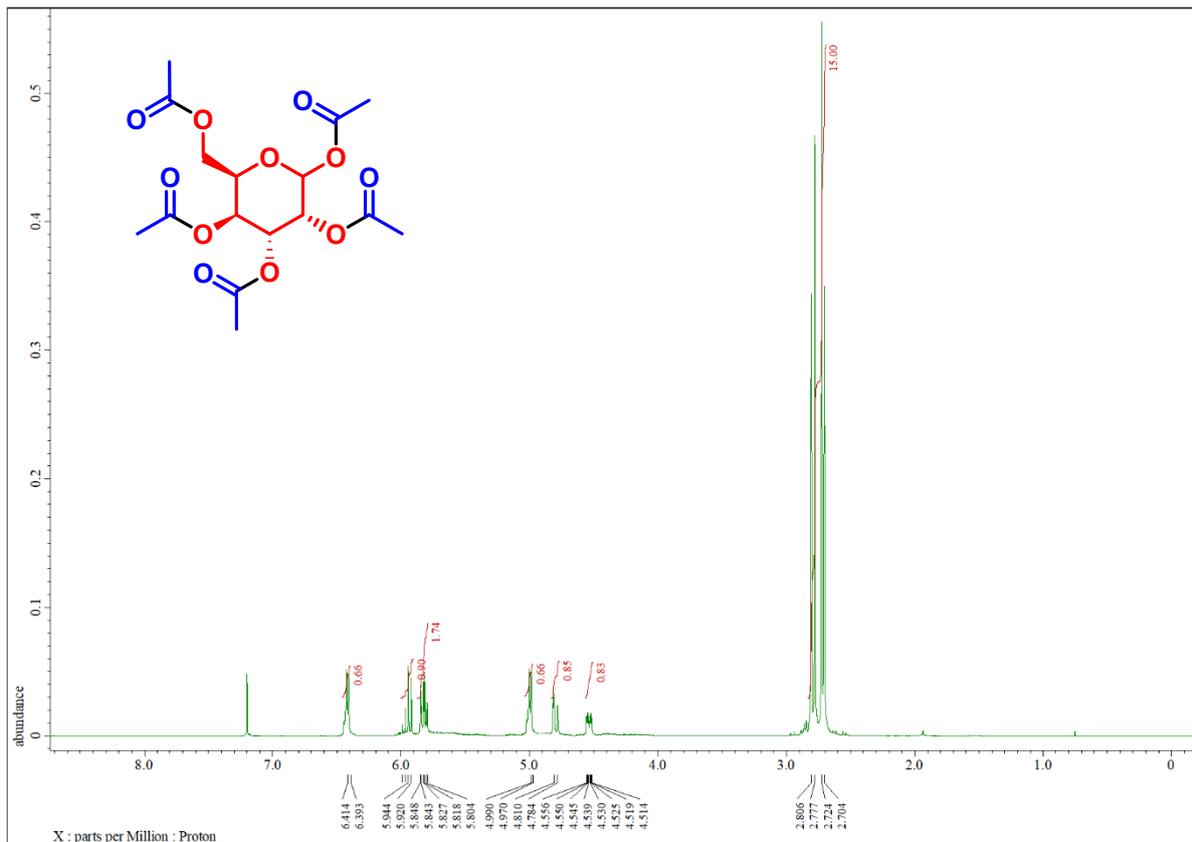


Figure S93. ^1H NMR spectrum of Gulose pentaacetate **E3**.

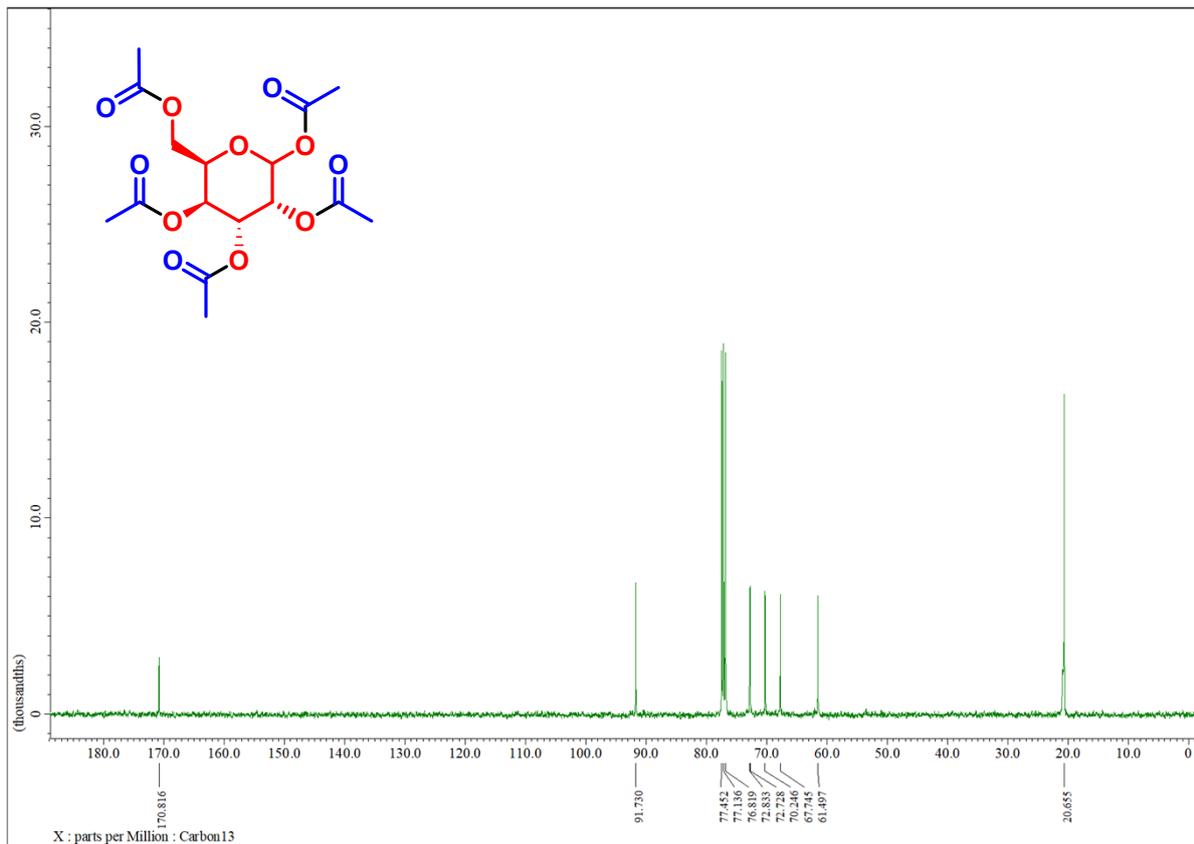


Figure S94. ^{13}C NMR spectrum of Gulose pentaacetate **E3**.

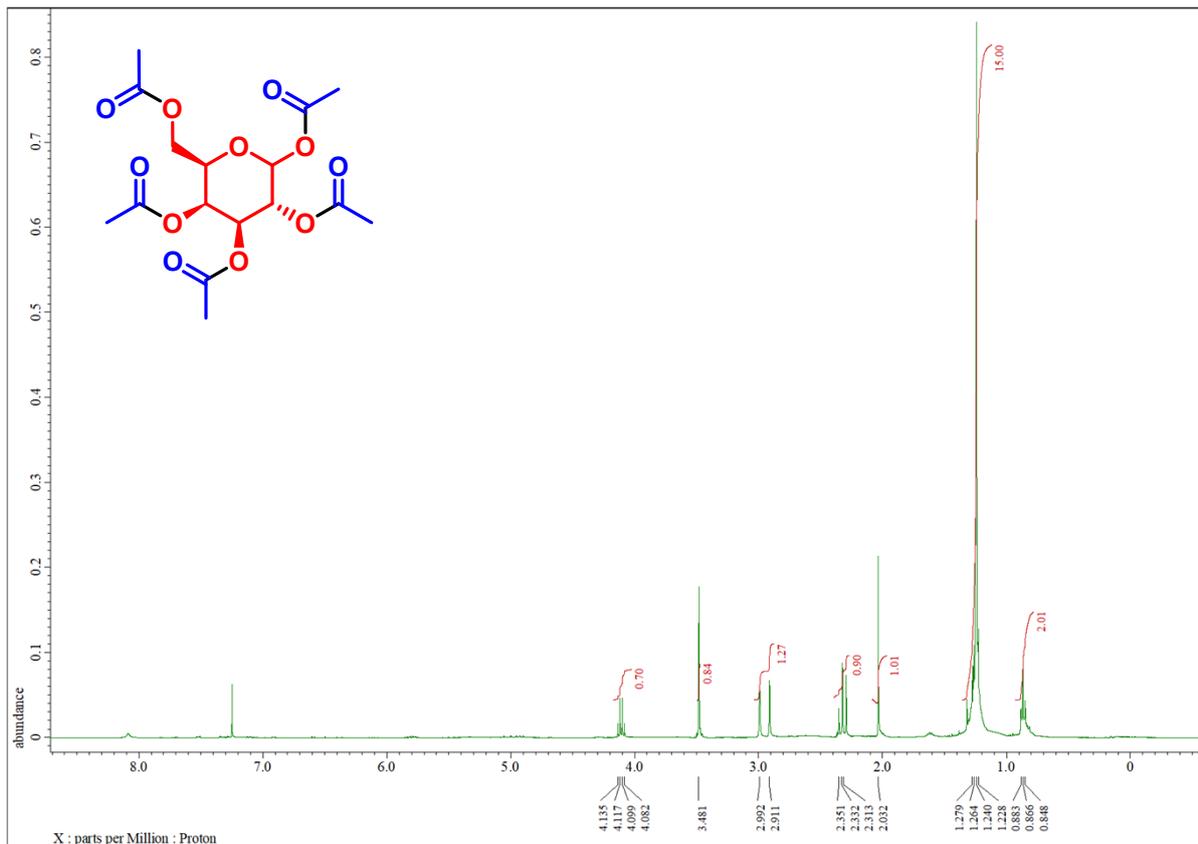


Figure S95. ¹H NMR spectrum of Galactose pentaacetate **E4**.

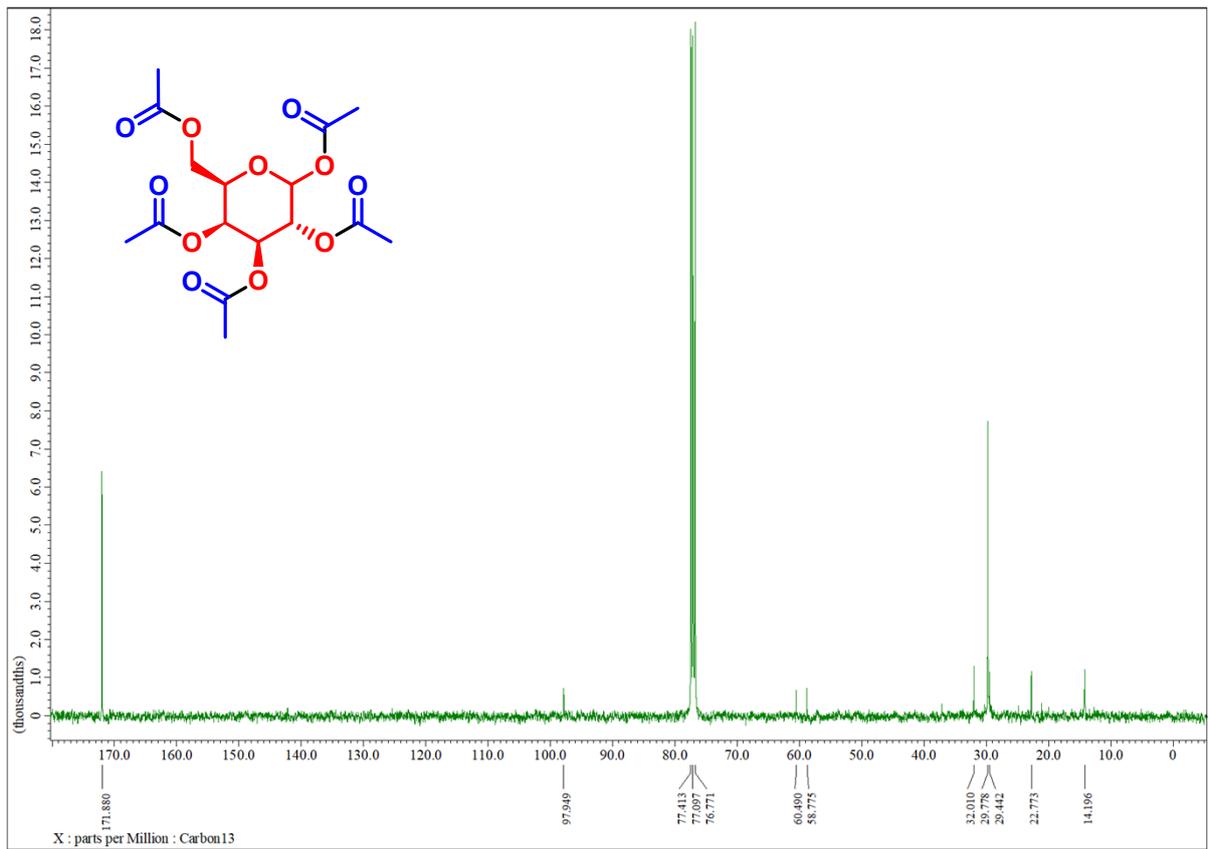


Figure S96. ^{13}C NMR spectrum of Galactose pentaacetate **E4**.

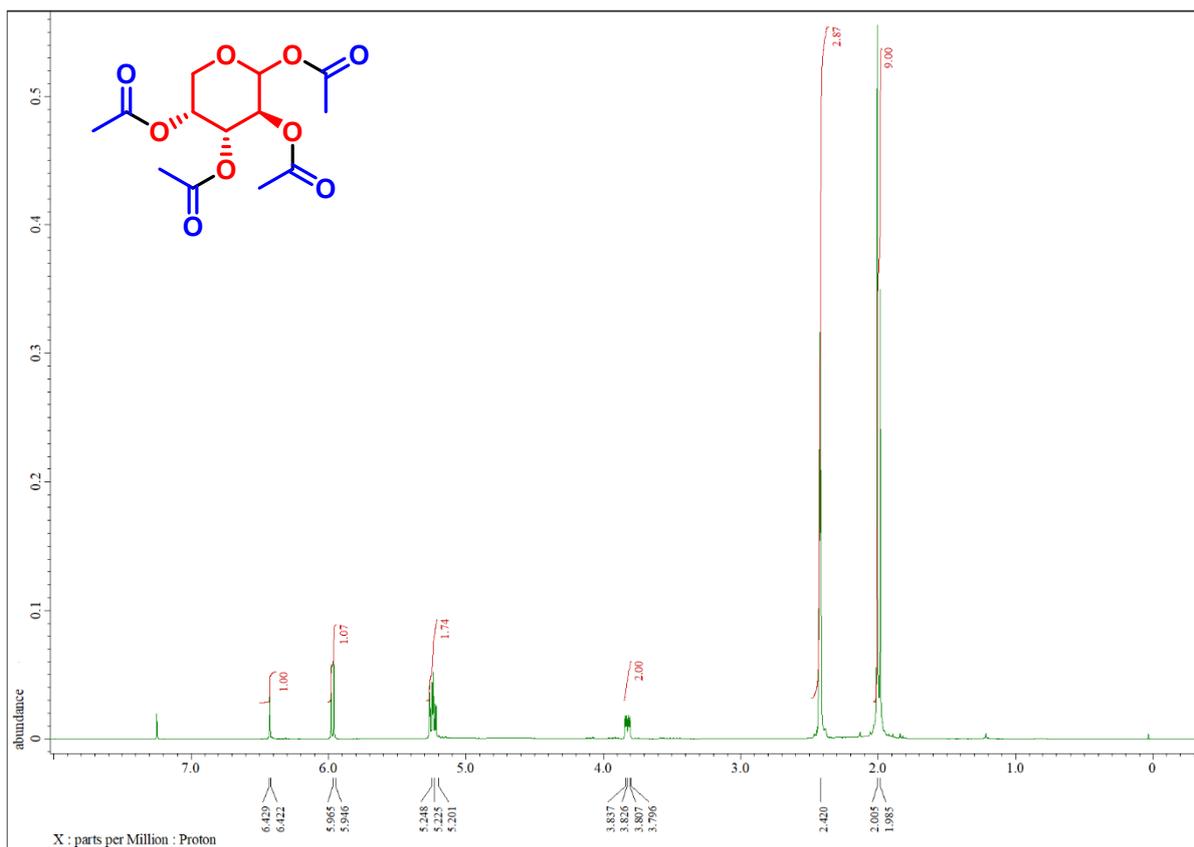


Figure S97. ^1H NMR spectrum of Ribose tetraacetate **E5**.

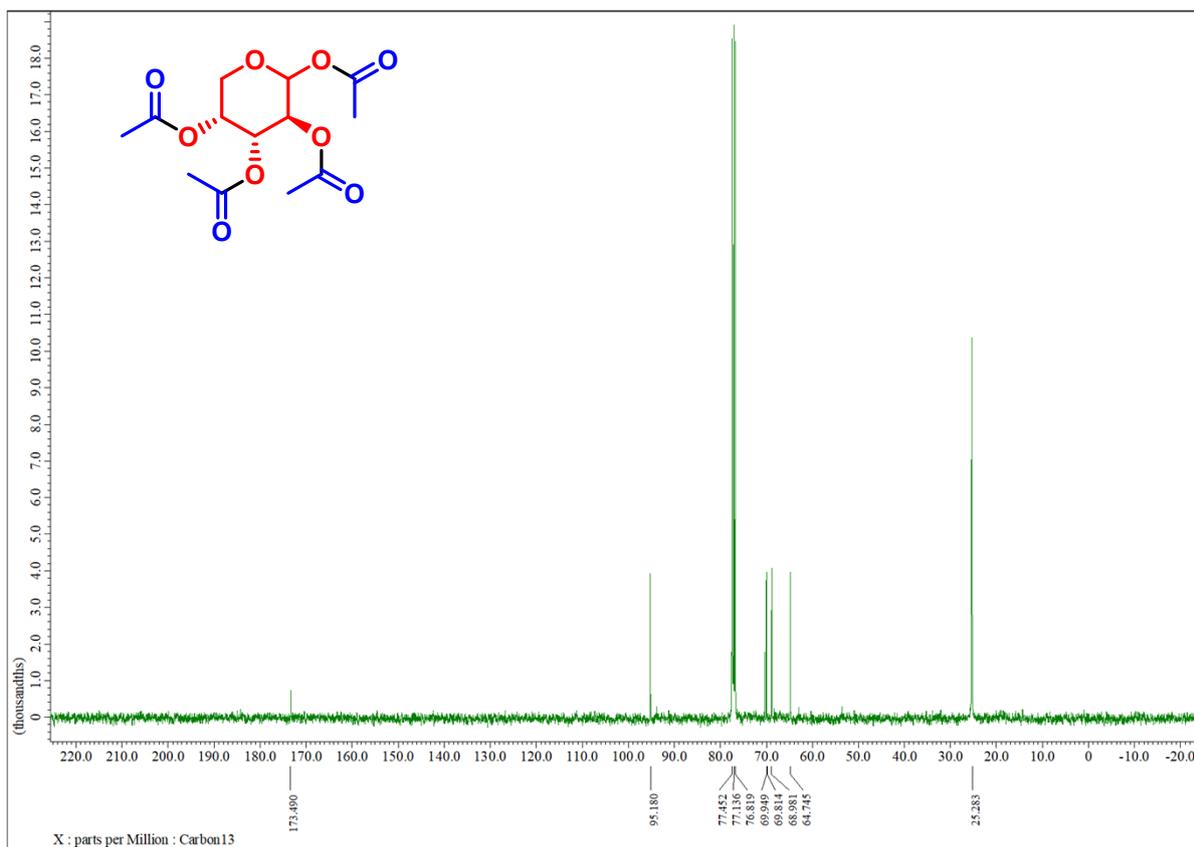


Figure S98. ^{13}C NMR spectrum of Ribose tetraacetate **E5**.

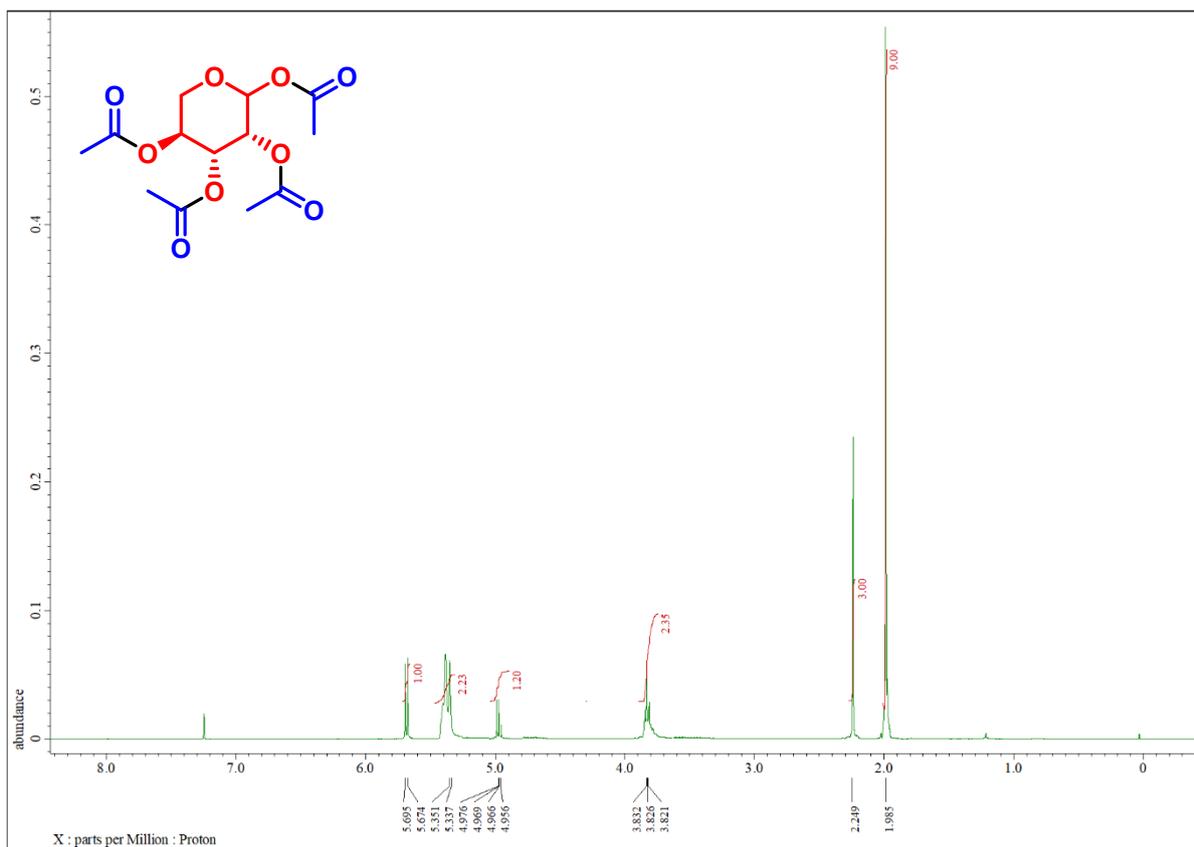


Figure S99. ^1H NMR spectrum of Lyxose tetraacetate **E6**.

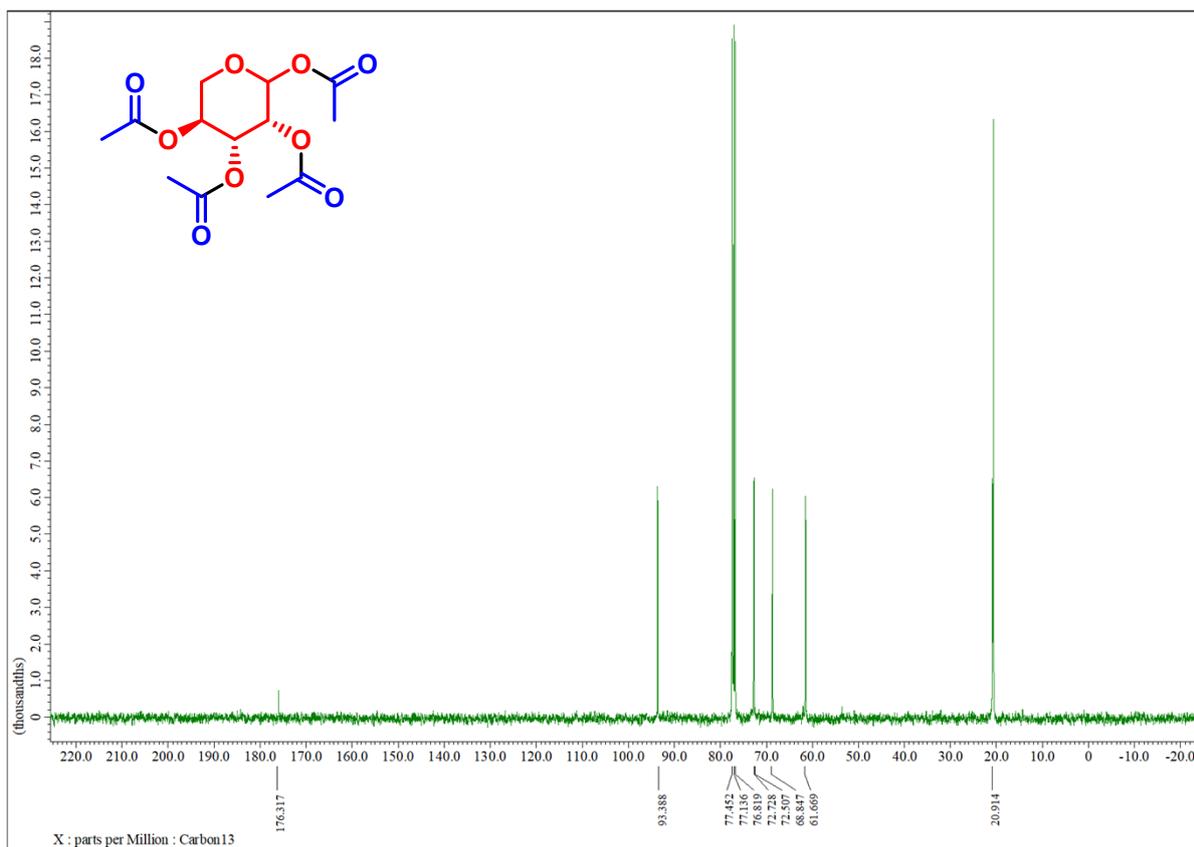


Figure S100. ^{13}C NMR spectrum of Lyxose tetraacetate **E6**.

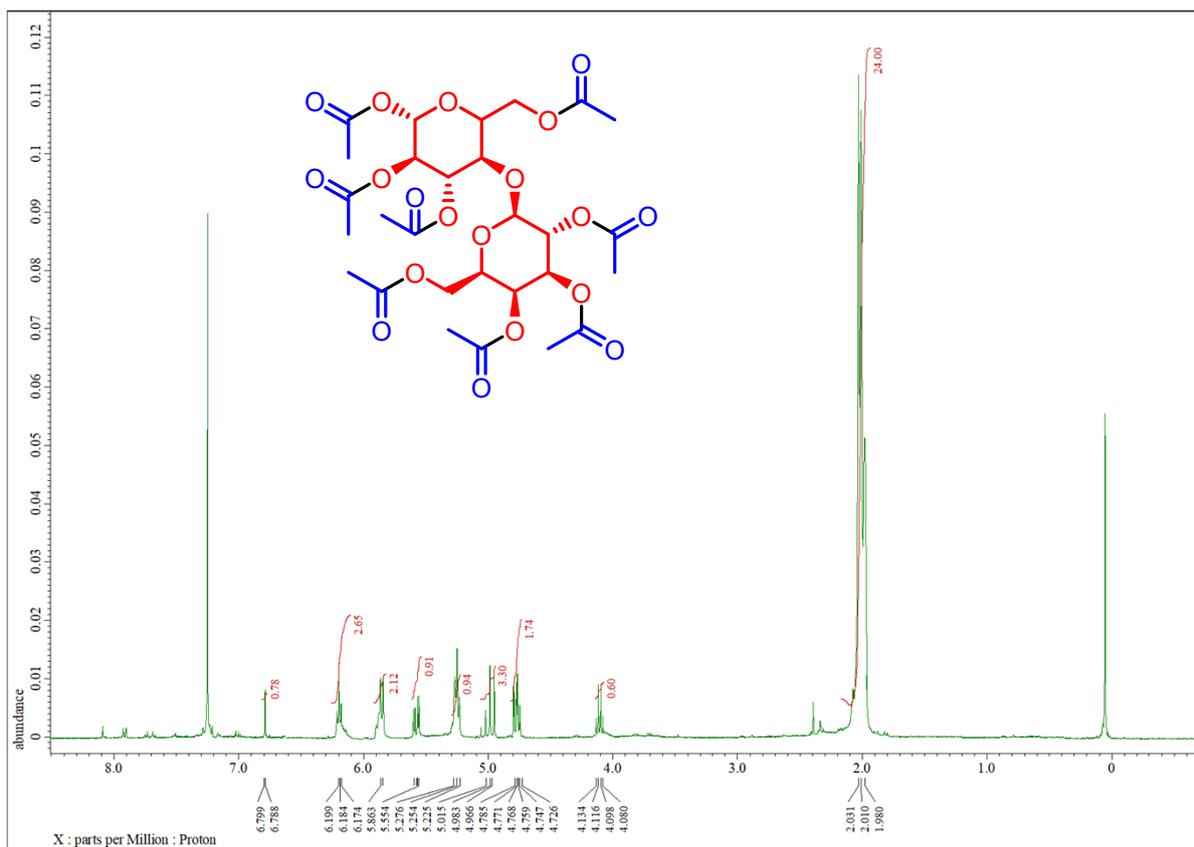


Figure S101. ^1H NMR spectrum of Lactose octaacetate **E7**.

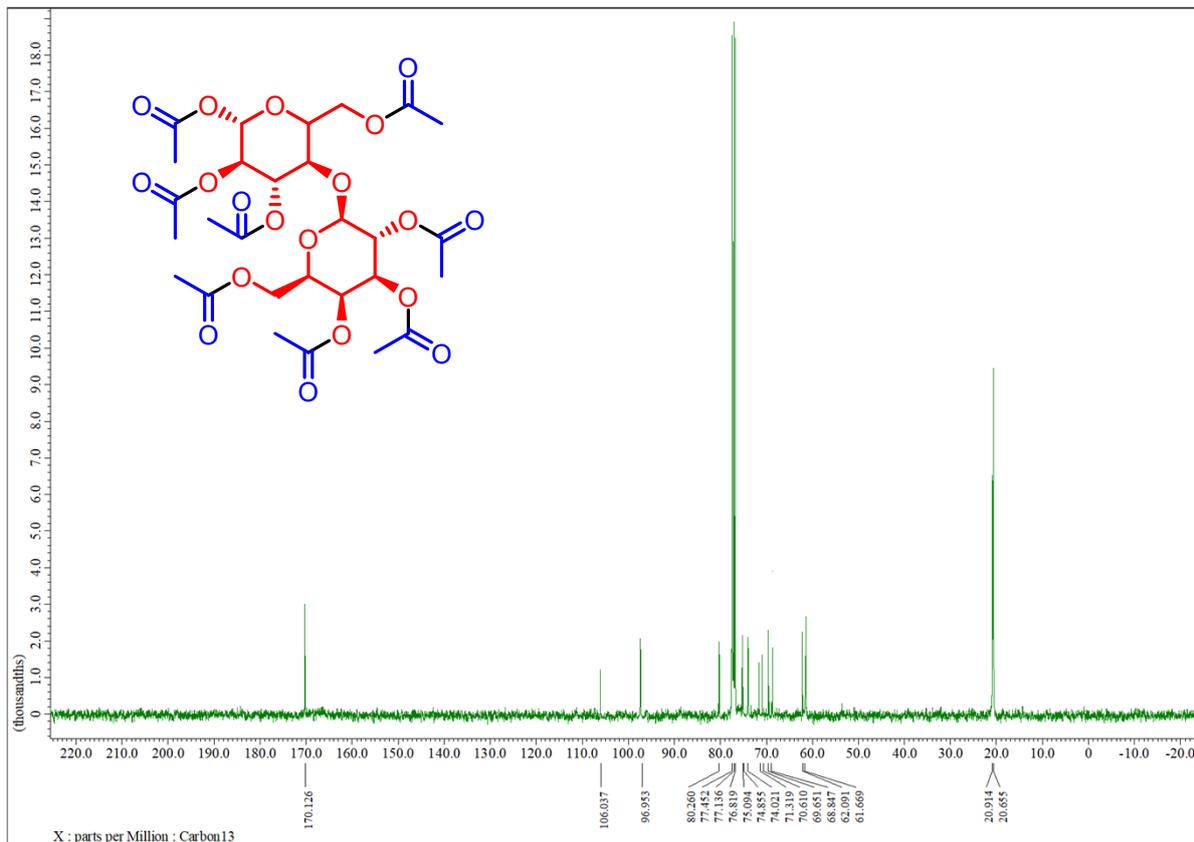


Figure S102. ^{13}C NMR spectrum of Lactose octaacetate E7.

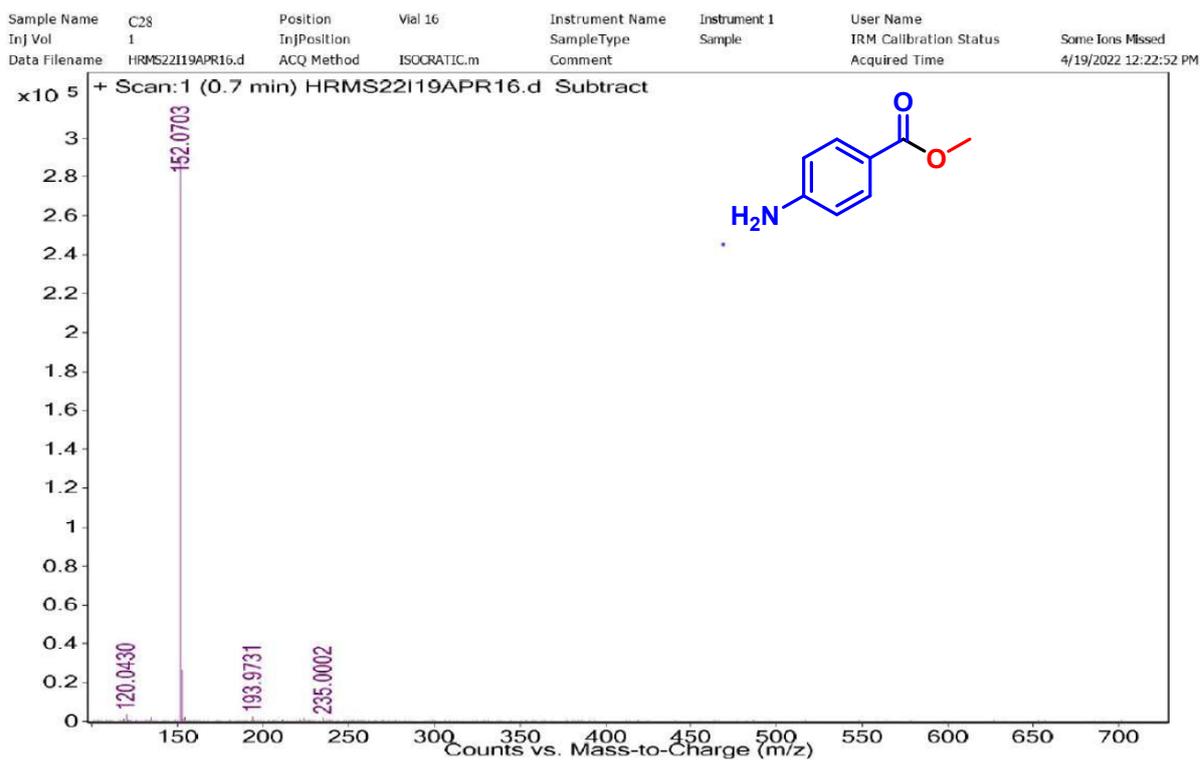


Figure S103. Mass spectrum of Methyl 4-aminobenzoate C2.

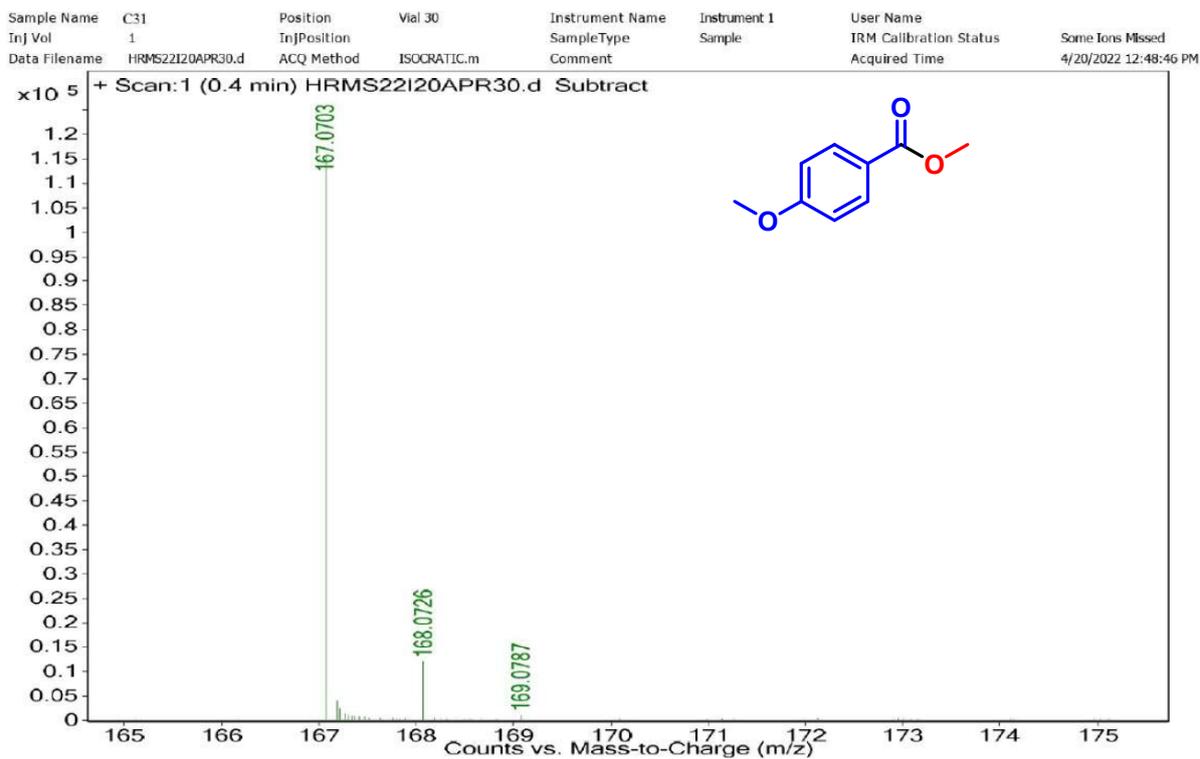


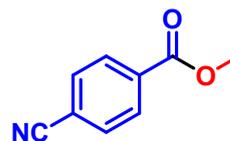
Figure S104. Mass spectrum of Methyl 4-methoxybenzoate C4.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5



Monoisotopic Mass, Even Electron Ions

48 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

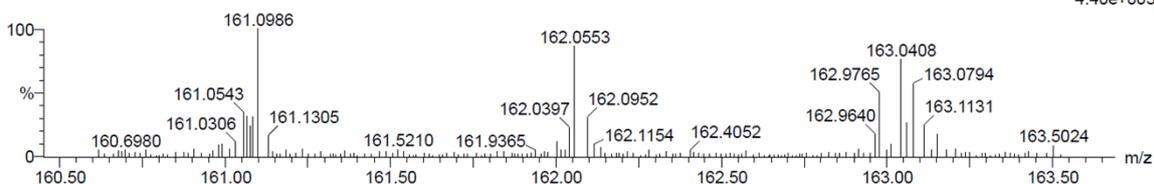
C: 2-20 H: 2-30 N: 0-2 O: 0-4 Br: 0-4

Sample Name : C8
Test Name :
11022022_C8 13 (0.294)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
4.40e+003



Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
162.0553	162.0555	-0.2	-1.2	6.5	838.6	n/a	n/a	C9 H8 N O2

Figure S105. Mass spectrum of Methyl 4-cyanobenzoate C5.

Sample Name : A_35
Test Name :
090522_A_35 9 (0.203)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
1.11e7

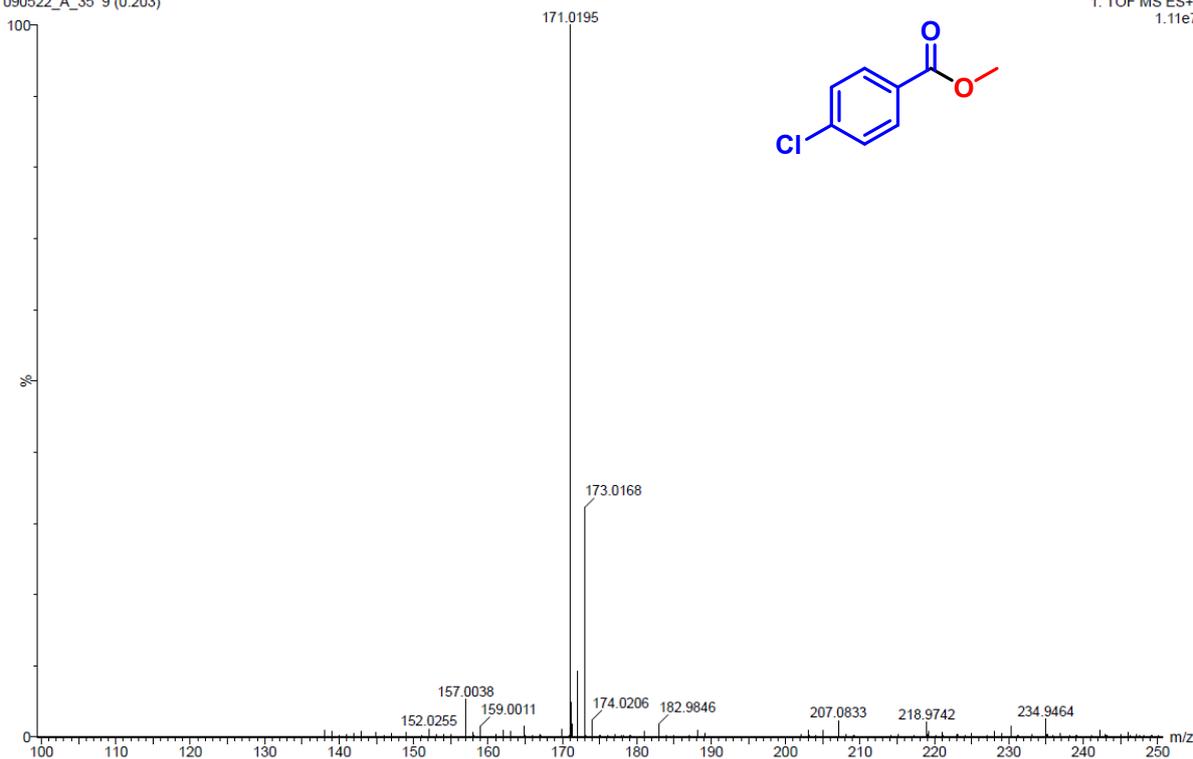


Figure S106. Mass spectrum of Methyl 4-chlorobenzoate C6.

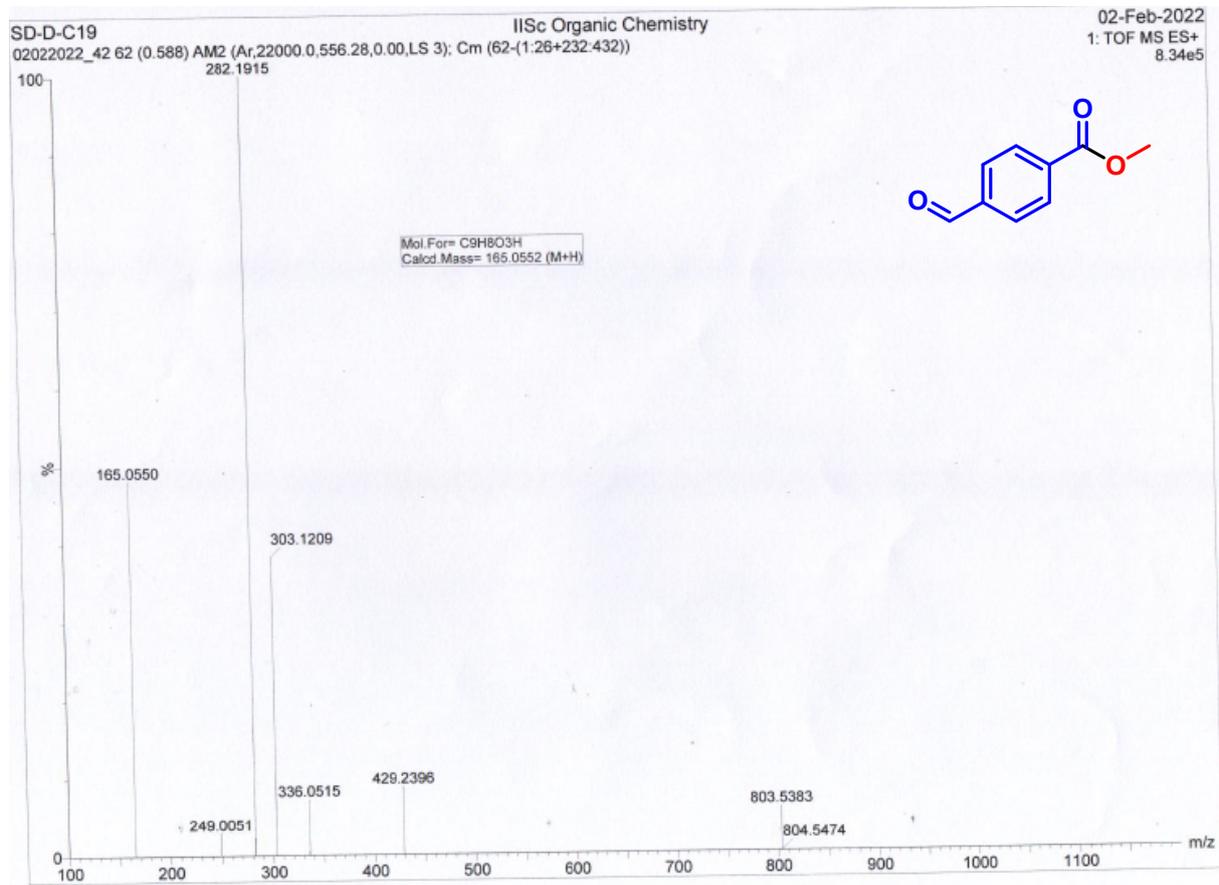


Figure S107. Mass spectrum of Methyl 4-formylbenzoate **C7**.

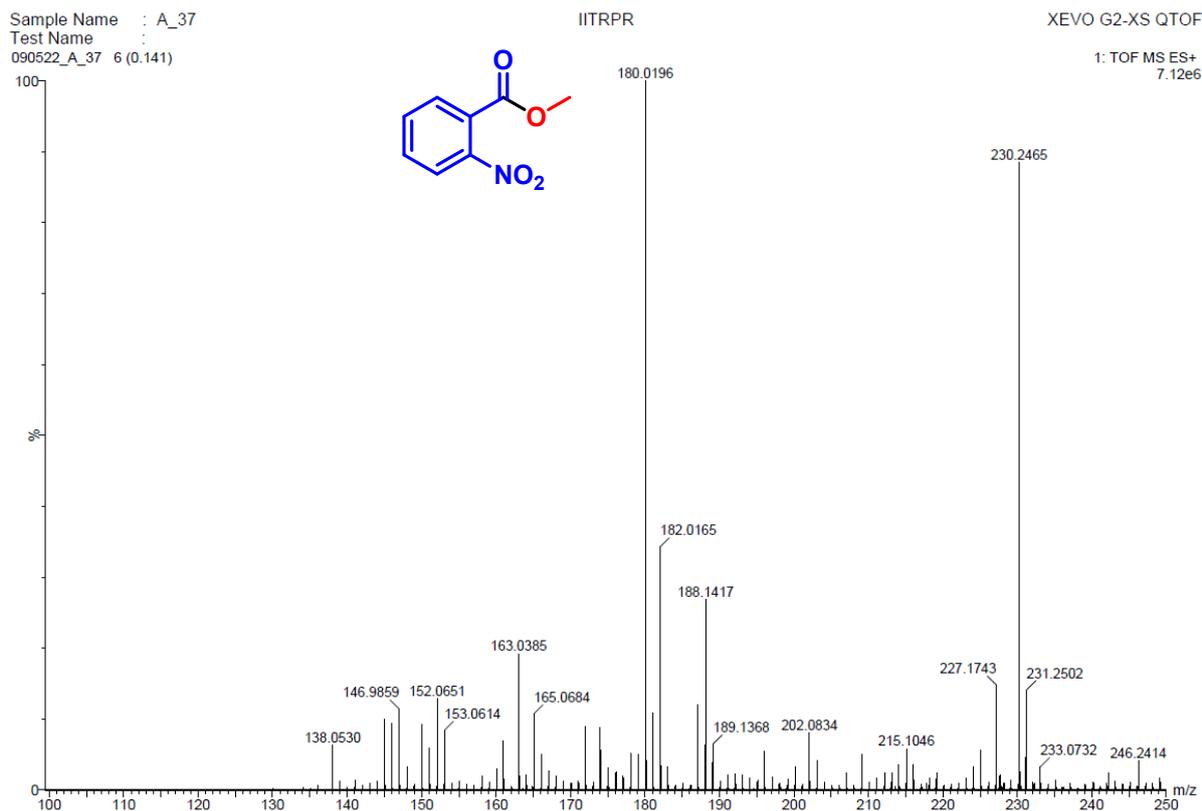


Figure S108. Mass spectrum of Methyl 2-nitrobenzoate **C8**.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

Monoisotopic Mass, Even Electron Ions

20 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 8-8 H: 2-30 N: 0-1 O: 0-4 F: 0-1

Sample Name : C10_NEG

Test Name :

11022022_C10_NEG 6 (0.130)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES-
 1.40e+005

Minimum:
 Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
200.0365	200.0359	0.6	3.0	5.5	1063.3	n/a	n/a	C8 H7 N O4 F

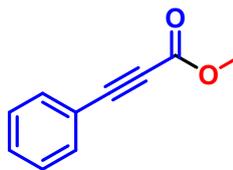
Figure S109. Mass spectrum of Methyl 4-fluoro-3-nitrobenzoate **C9**.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5



Monoisotopic Mass, Even Electron Ions

47 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

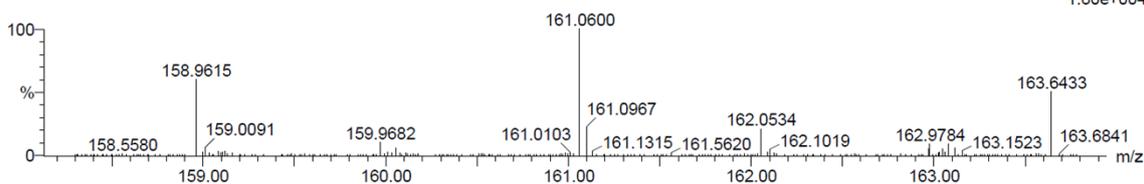
C: 2-20 H: 2-30 N: 0-2 O: 0-4 Br: 0-4

Sample Name : C9
Test Name :
11022022_C9 22 (0.480)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
1.80e+004



Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
161.0600	161.0603	-0.3	-1.9	6.5	1002.5	n/a	n/a	C10 H9 O2

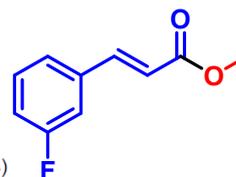
Figure S110. Mass spectrum of Methyl 3-phenylpropiolate C10

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5



Monoisotopic Mass, Even Electron Ions

170 formula(e) evaluated with 2 results within limits (up to 1 closest results for each mass)

Elements Used:

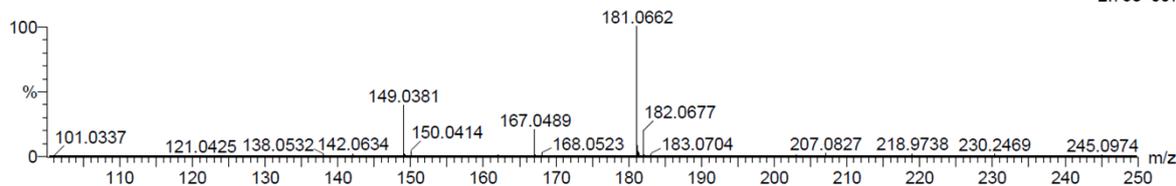
C: 0-50 H: 0-50 N: 0-4 O: 1-10 F: 1-3

Sample Name : A_34
Test Name :
090522_A_34 8 (0.186)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
2.73e+007



Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
181.0662	181.0665	-0.3	-1.7	5.5	2919.7	n/a	n/a	C10 H10 O2 F

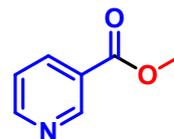
Figure S111. Mass spectrum of Methyl (E)-3-(3-fluorophenyl)acrylate C11.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5



Monoisotopic Mass, Even Electron Ions

51 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-50 H: 0-50 N: 0-4 O: 1-10

Sample Name : A_33

IITRPR

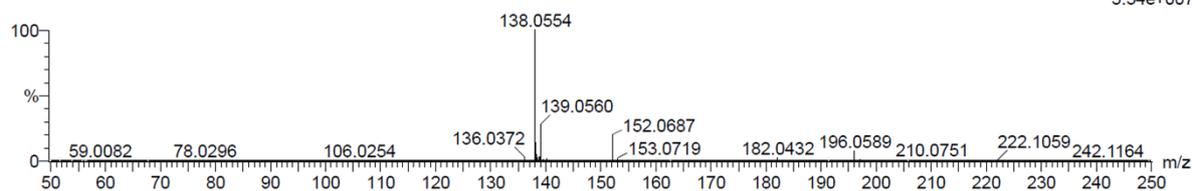
XEVO G2-XS QTOF

Test Name :

090522_A_33 8 (0.186)

1: TOF MS ES+

3.54e+007



Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
138.0554	138.0555	-0.1	-0.7	4.5	3677.0	n/a	n/a	C7 H8 N O2

Figure S112. Mass spectrum of Methyl nicotinate **C13**.

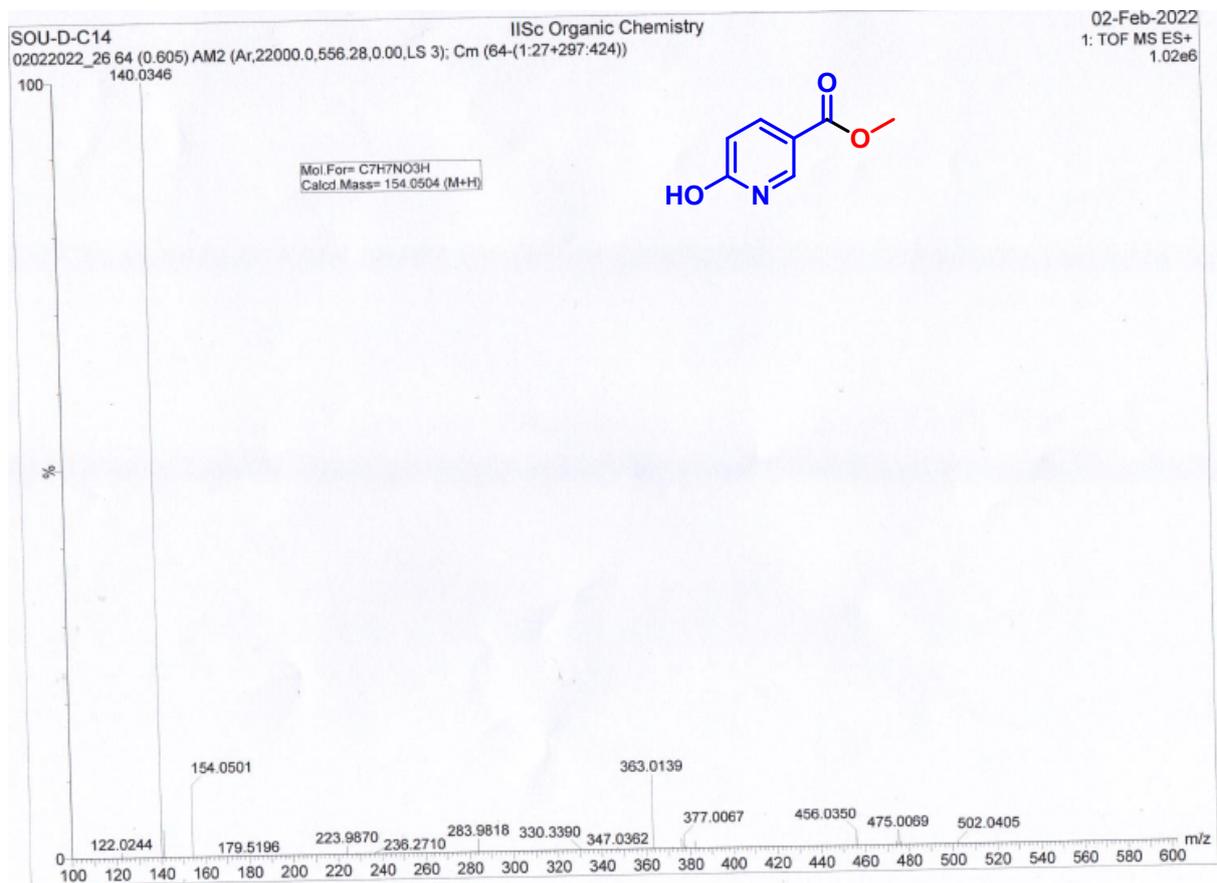


Figure S113. Mass spectrum of Methyl 6-hydroxynicotinate **C14**.

Sample Name	C29	Position	Vial 24	Instrument Name	Instrument 1	User Name	
Inj Vol	1	InjPosition		SampleType	Sample	IRM Calibration Status	Some Ions Missed
Data Filename	HRMS22119APR24.d	ACQ Method	ISOCRATIC.m	Comment		Acquired Time	4/19/2022 12:43:57 PM

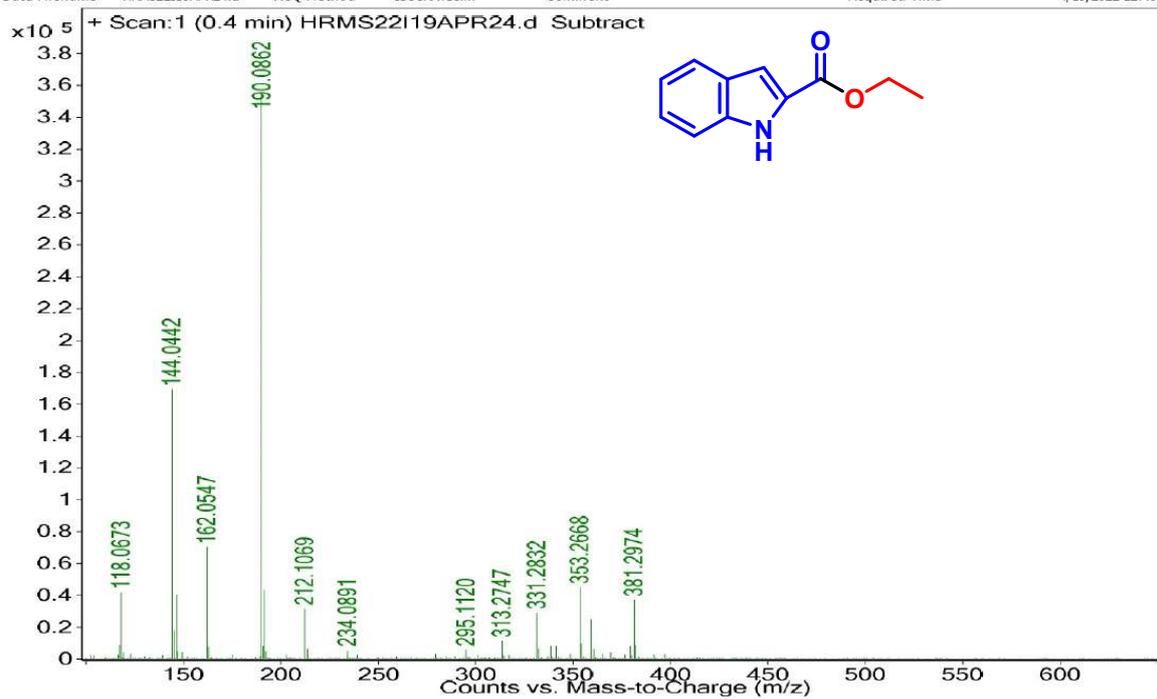


Figure S114. Mass spectrum of Ethyl 1H-indole-2-carboxylate **C15**.

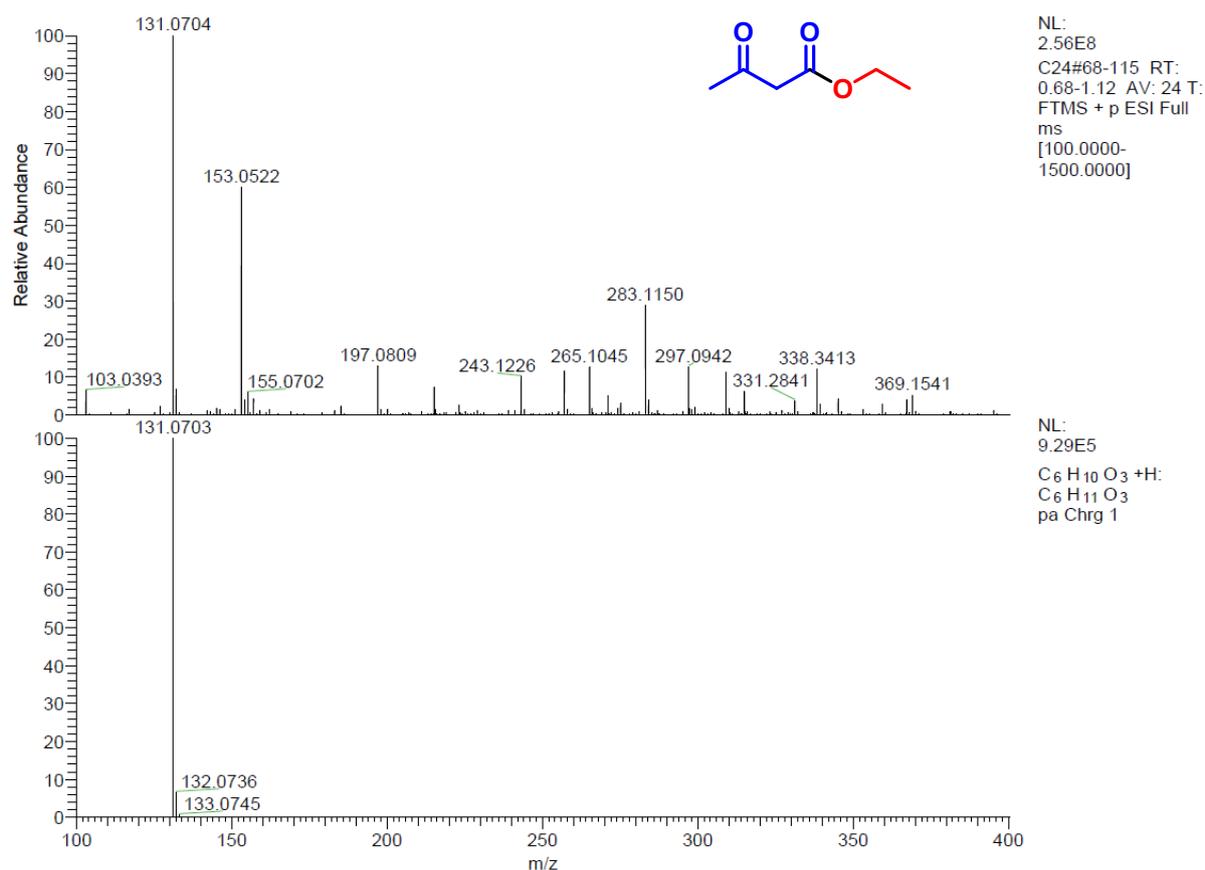


Figure S115. Mass spectrum of Ethyl 3-oxobutanoate **C16**.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5

Monoisotopic Mass, Even Electron Ions

191 formula(e) evaluated with 2 results within limits (up to 1 closest results for each mass)

Elements Used:

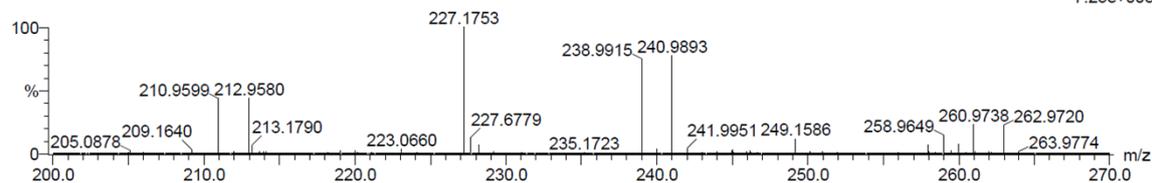
C: 2-20 H: 2-30 N: 0-1 O: 0-4 Br: 0-4 F: 0-3

Sample Name : C1
Test Name :
11022022_C1 39 (0.825)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
7.25e+005



Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
238.9915	238.9919	-0.4	-1.7	1.5	1961.8	n/a	n/a	C ₇ H ₁₂ O ₄ Br

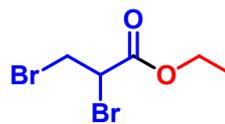
Figure S116. Mass spectrum of Diethyl 2-bromomalonate **C17**.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5



Monoisotopic Mass, Even Electron Ions

303 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

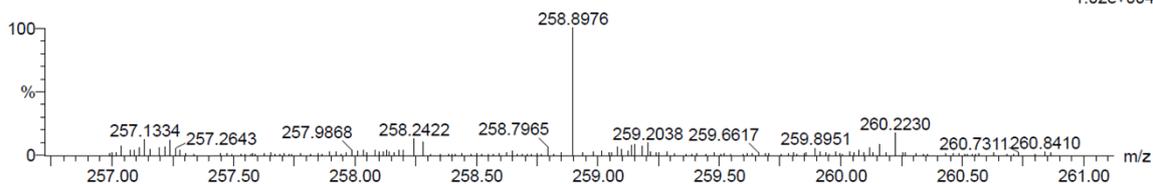
C: 2-20 H: 2-30 N: 0-2 O: 0-4 F: 0-3 Br: 0-4

Sample Name : C4
Test Name :
11022022_C4 14 (0.311)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
1.02e+004

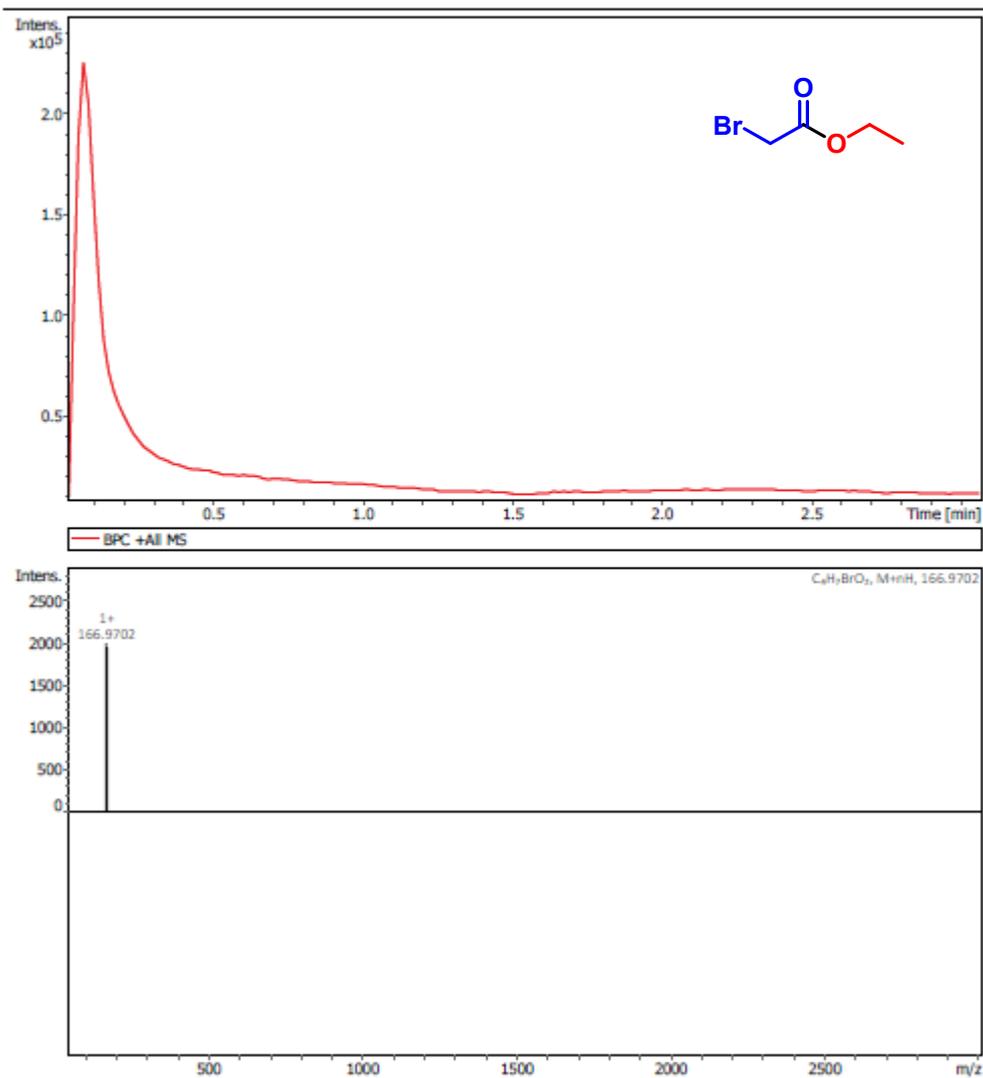


Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
258.8976	258.8969	0.7	2.7	0.5	715.2	n/a	n/a	C5 H9 O2 Br2

Figure S117. Mass spectrum of Ethyl 2,3-dibromopropanoate **C18**.

Analysis Info
Analysis Name C:\Users\ANUPAM MISHRA\AppData\Local\Temp\Temp1_PM.zip\PM\m chem AR-DP-C22_RA3_01_16454.d
Method 7. LCMS tune low MeOH.m
Sample Name m chem AR-DP-C22
Comment
Acquisition Date 18-04-2022 14:51:48
Operator IIT Indore
Instrument micrOTOF-Q



Bruker Compass DataAnalysis 5.1 printed: 23-04-2022 12:22:49 by: IIT INDORE Page 1 of 1

Figure S118. Mass spectrum of Ethyl 2-bromoacetate **C21**.

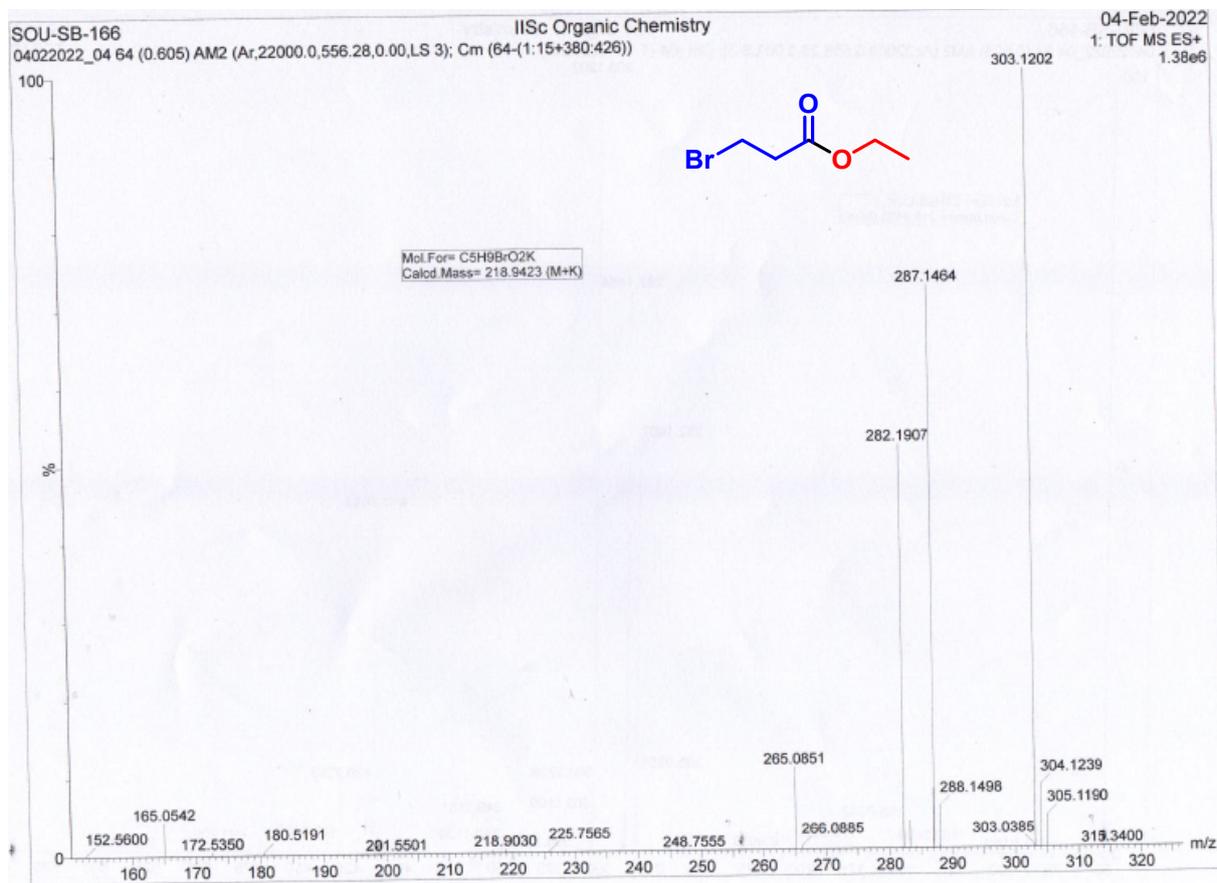
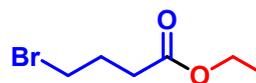


Figure S119. Mass spectrum of Ethyl 3-bromopropanoate **C22**.

Qualitative Compound Report

Data File C23.d **Sample Name** C23
Sample Type Sample **Position** P1-B1
Instrument Name Instrument 1 **User Name**
Acq Method MS Scan.m **Acquired Time** 24-05-2022 11:39:35
IRM Calibration Status Success **DA Method** Default.m
Comment

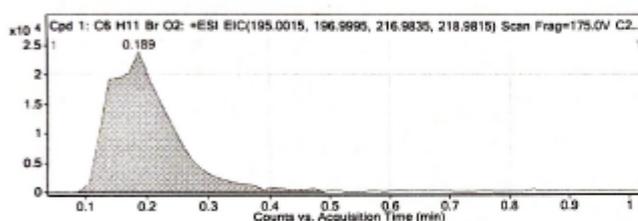


Sample Group Info. 3
Acquisition SW Version 6200 series TOF/ESQ series
 Q-TOF B.05.01 (85125)

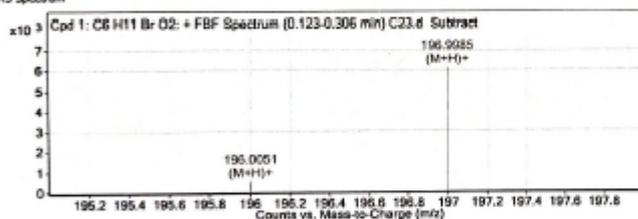
Compound Table

Compound Label	RT	Mass	Abund	Formula	Tgt Mass	Diff (ppm)	HFG Formula	DB Formula
Cpd 1: C6 H11 Br O2	0.189	193.9937	6670	C6 H11 Br O2	193.9942	-3.04	C6 H11 Br O2	C6 H11 Br O2

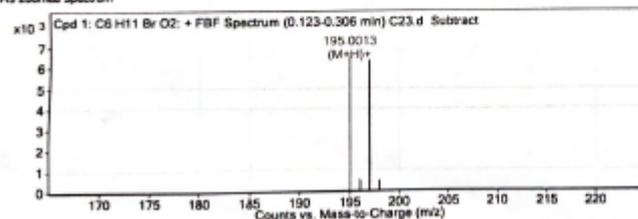
Compound Label	m/z	RT	Algorithm	Mass
Cpd 1: C6 H11 Br O2	195.0013	0.189	Find By Formula	193.9937



MS Spectrum



MS Zoomed Spectrum



MS Spectrum Peak List

m/z	z	Abund	Formula	Ion
195.0013	1	6670.37	C6H12BrO2	(M+H)+
196.0051	1	568.1	C6H12BrO2	(M+H)+
196.9985	1	6155.59	C6H12BrO2	(M+H)+
198.0026	1	365.94	C6H12BrO2	(M+H)+

--- End Of Report ---

Figure S120. Mass spectrum of Ethyl 4-bromobutanoate C23.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5



Monoisotopic Mass, Even Electron Ions

68 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

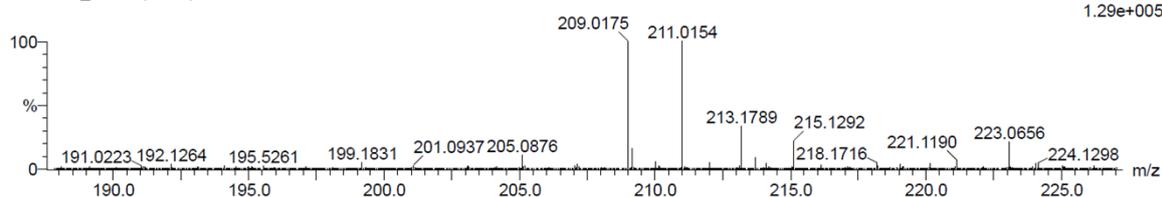
C: 2-20 H: 2-30 N: 0-2 O: 0-4 Br: 0-4

Sample Name : C6
Test Name :
11022022_C6 23 (0.497)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
1.29e+005

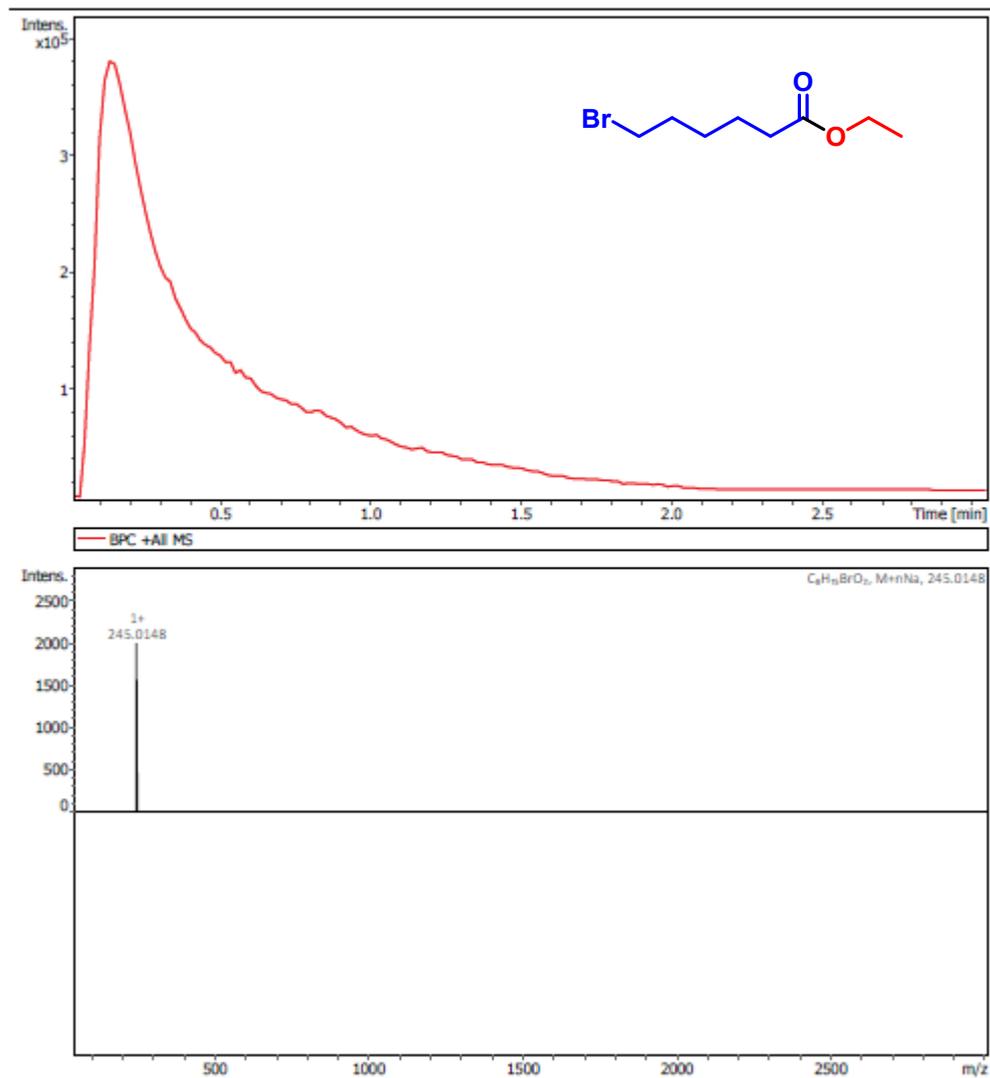


Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
209.0175	209.0177	-0.2	-1.0	0.5	1657.9	n/a	n/a	C7 H14 O2 Br

Figure S121. Mass spectrum of Ethyl 5-bromopentanoate **C24**.

Analysis Info
Analysis Name C:\Users\ANUPAM MISHRA\AppData\Local\Temp\Temp1_PM.zip\PM\m chem AR-DP-C23_RA4_01_16455.d
Method 7. LCMS tune low MeOH.m
Sample Name m chem AR-DP-C23
Comment
Acquisition Date 18-04-2022 14:55:51
Operator IIT Indore
Instrument micrOTOF-Q



Bruker Compass DataAnalysis 5.1 printed: 23-04-2022 12:27:28 by: IIT INDORE Page 1 of 1

Figure S122. Mass spectrum of Ethyl 6-bromohexanoate **C25**.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

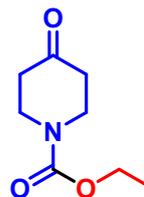
Number of isotope peaks used for i-FIT = 5

Monoisotopic Mass, Even Electron Ions

111 formula(e) evaluated with 2 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 2-20 H: 2-30 N: 0-1 O: 0-4 F: 0-3 Br: 0-4

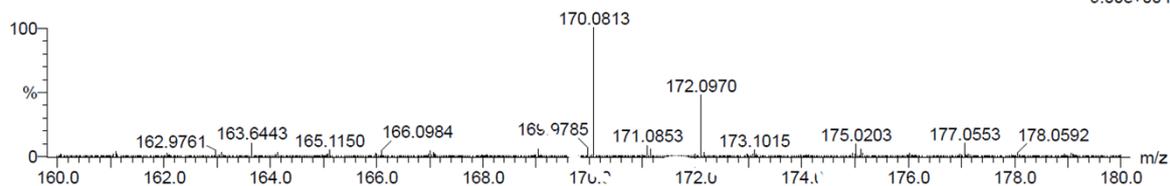


Sample Name : C2
Test Name :
11022022_C2 65 (1.368)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
9.00e+004



Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
172.0970	172.0974	-0.4	-2.3	2.5	1816.7	n/a	n/a	C8 H14 N O3

Figure S123. Mass spectrum of Ethyl 4-oxopiperidine-1-carboxylate **C26**.

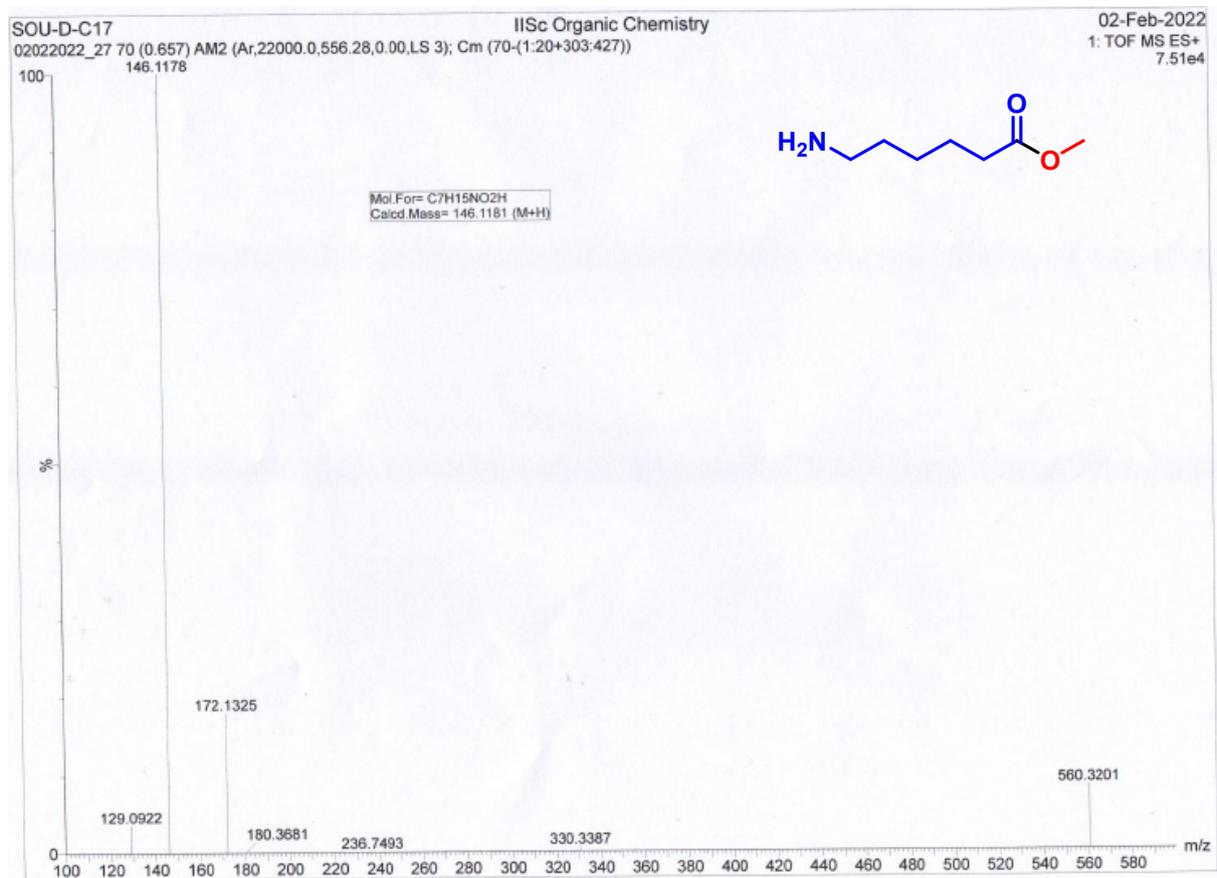
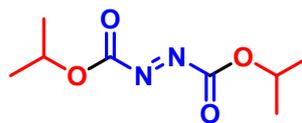


Figure S124. Mass spectrum of Methyl 6-aminohexanoate **C28**.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 5



Monoisotopic Mass, Even Electron Ions

207 formula(e) evaluated with 2 results within limits (up to 1 closest results for each mass)

Elements Used:

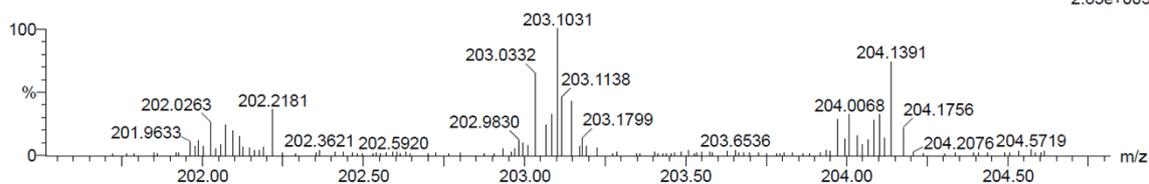
C: 2-20 H: 2-30 N: 0-2 O: 0-4 F: 0-3 Br: 0-4

Sample Name : C5
Test Name :
11022022_C5 27 (0.576)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
2.85e+003



Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
203.1031	203.1032	-0.1	-0.5	2.5	580.5	n/a	n/a	C8 H15 N2 O4

Figure S125. Mass spectrum of Diisopropyl (E)-diazene-1,2-dicarboxylate **C29**.

Sample Name : A_9
Test Name :
090522_A_9 5 (0.124)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
2.29e4

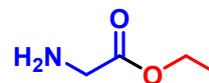
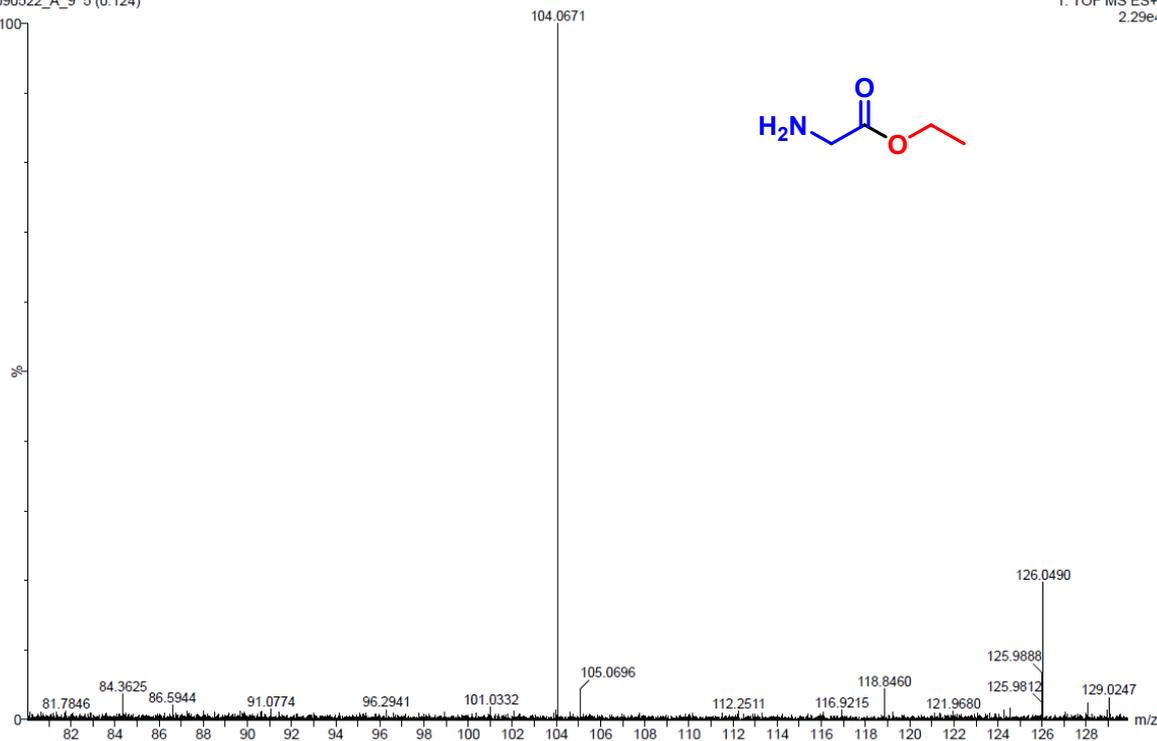


Figure S126. Mass spectrum of Ethyl glycinate **D1**.

Analysis Info
Analysis Name C:\Users\ANUPAM MISHRA\AppData\Local\Temp\Temp1_PM.zip\PM\m chem AR-DP-A2_RA2_01_16453.d
Method 7. LCMS tune low MeOH.m
Sample Name m chem AR-DP-A2
Comment
Acquisition Date 18-04-2022 14:47:51
Operator IIT Indore
Instrument micrOTOF-Q

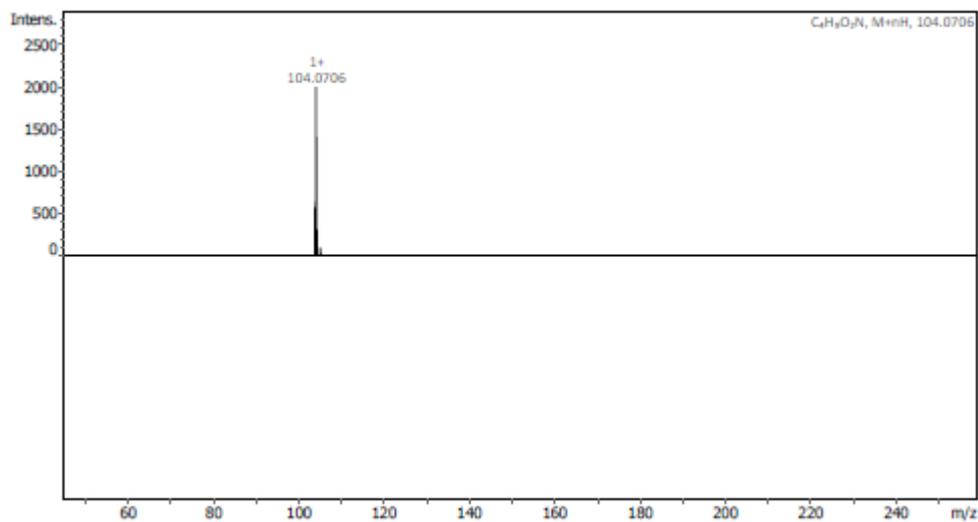
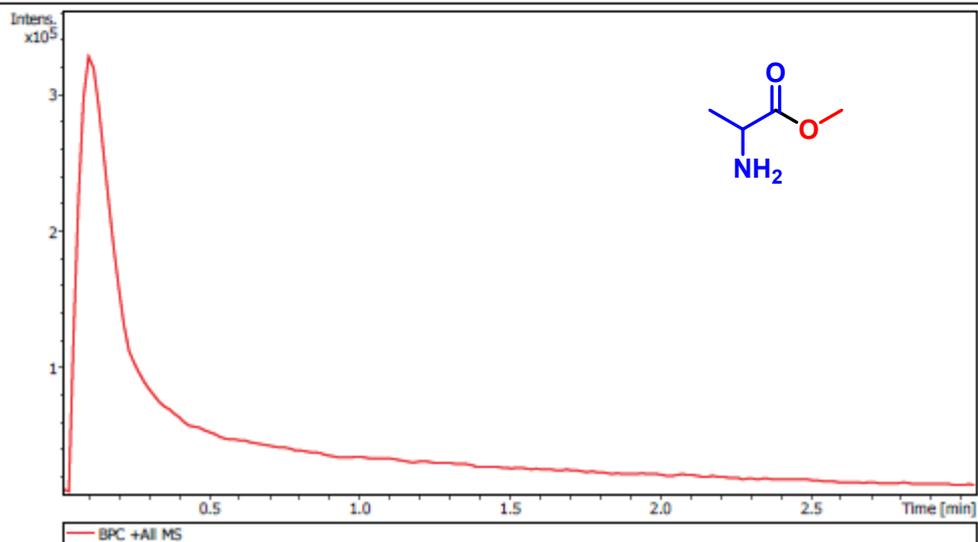


Figure S127. Mass spectrum of Methyl alaninate **D2**.

Sample Name : A_12
Test Name :
090522_A_12 8 (0.186)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
7.79e3

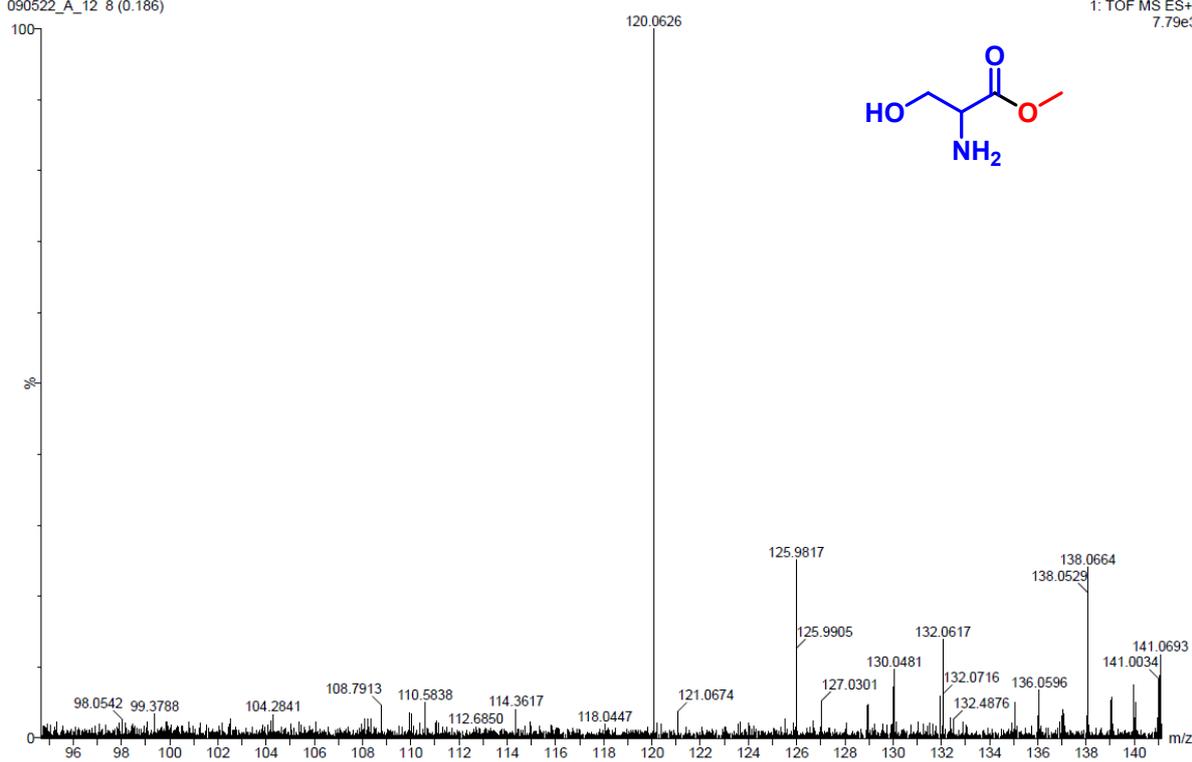


Figure S128. Mass spectrum of Methyl serinate **D3**.

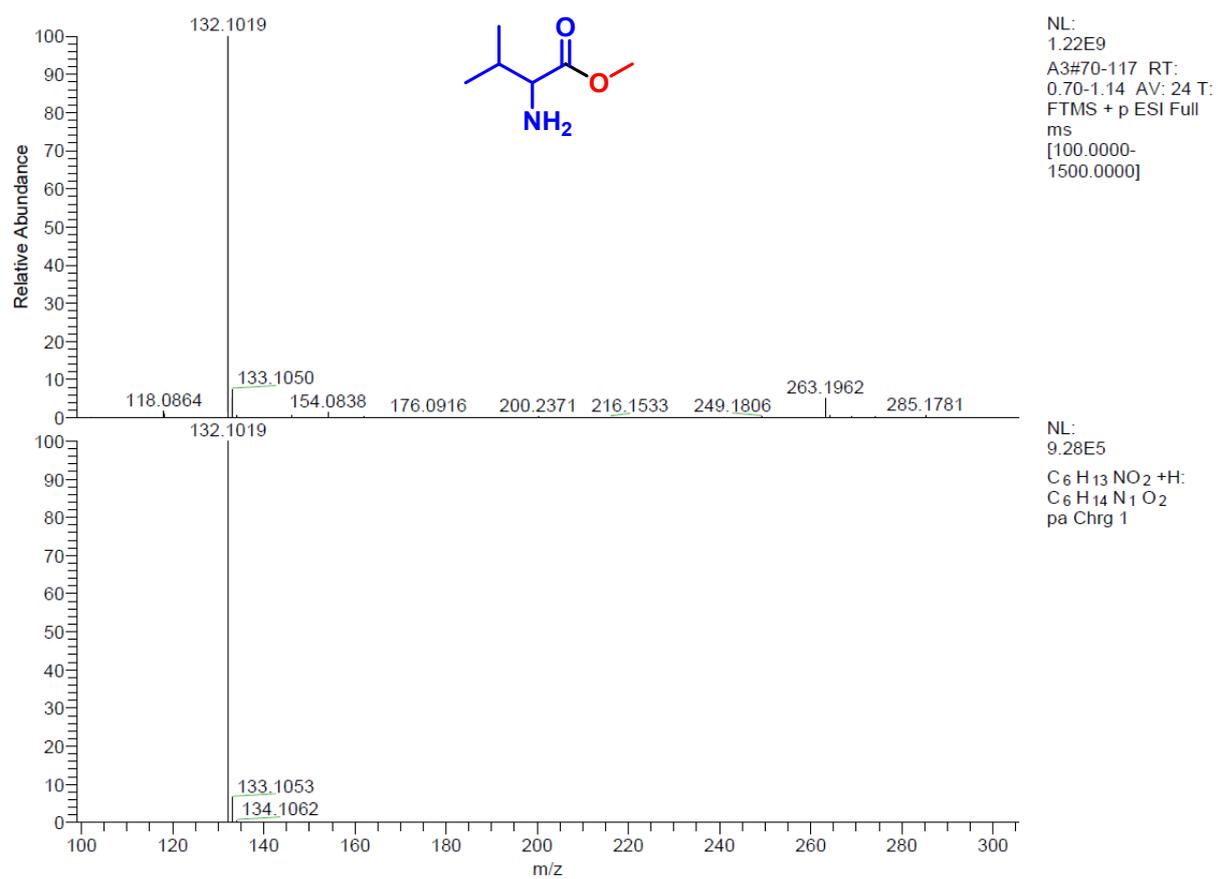


Figure S129. Mass spectrum of Methyl valinate D5.

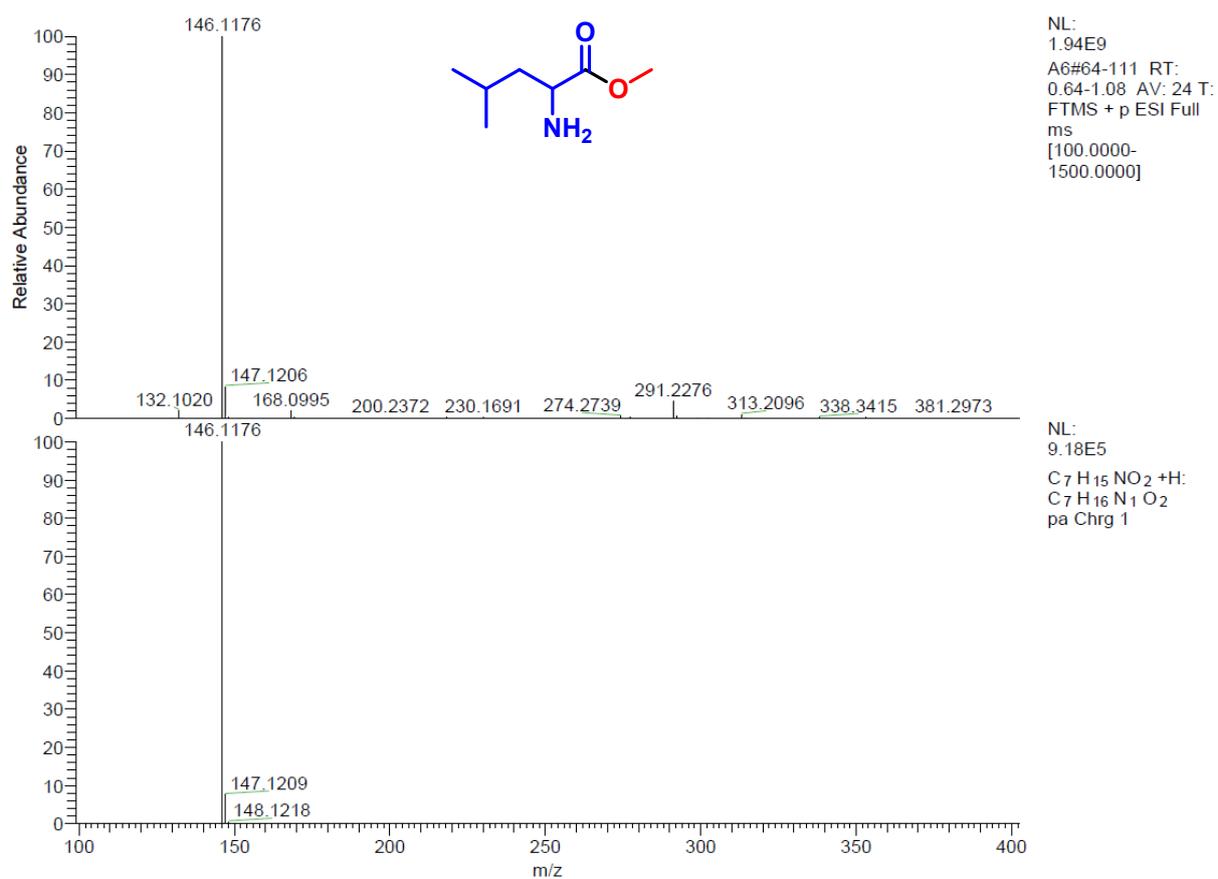


Figure S130. Mass spectrum of Methyl leucinate **D6**.

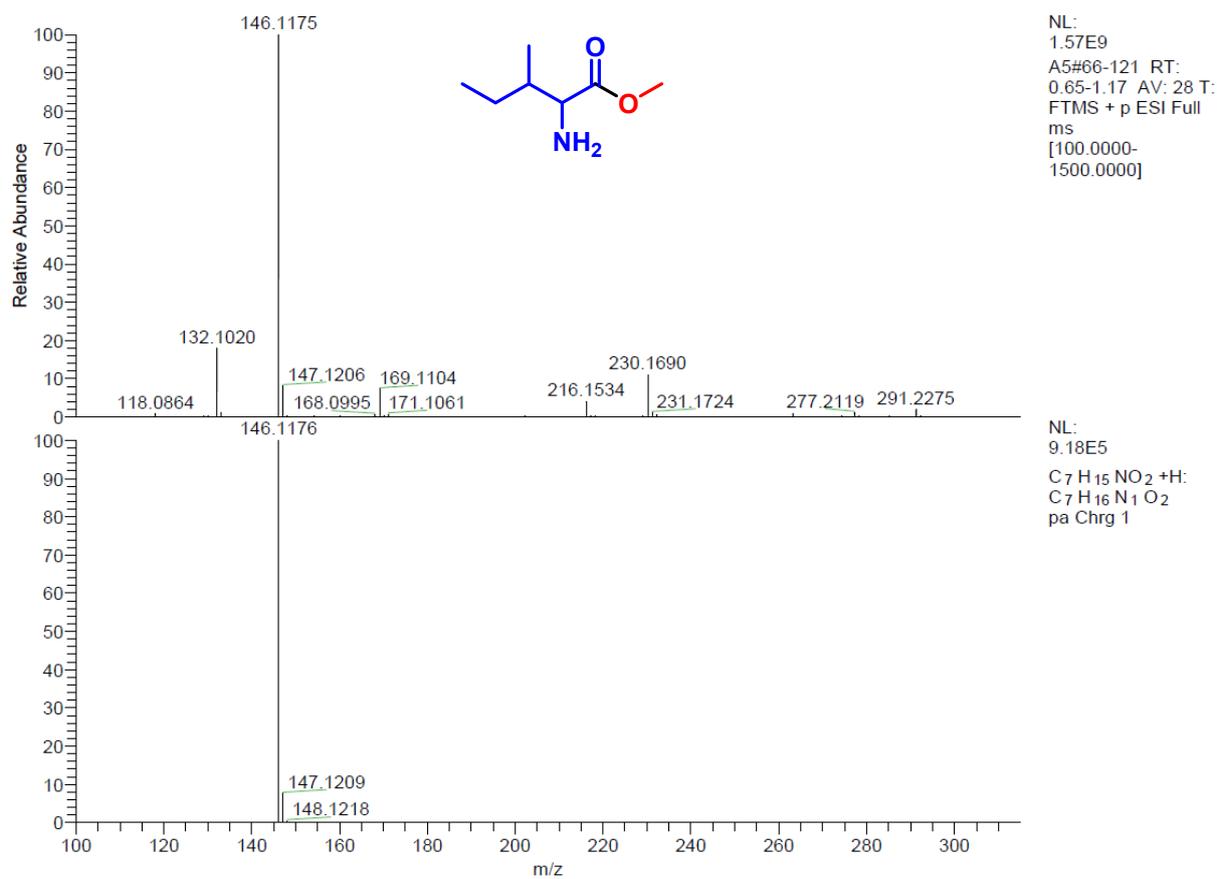


Figure S131. Mass spectrum of Methyl 2-amino-3-methylpentanoate **D7**.

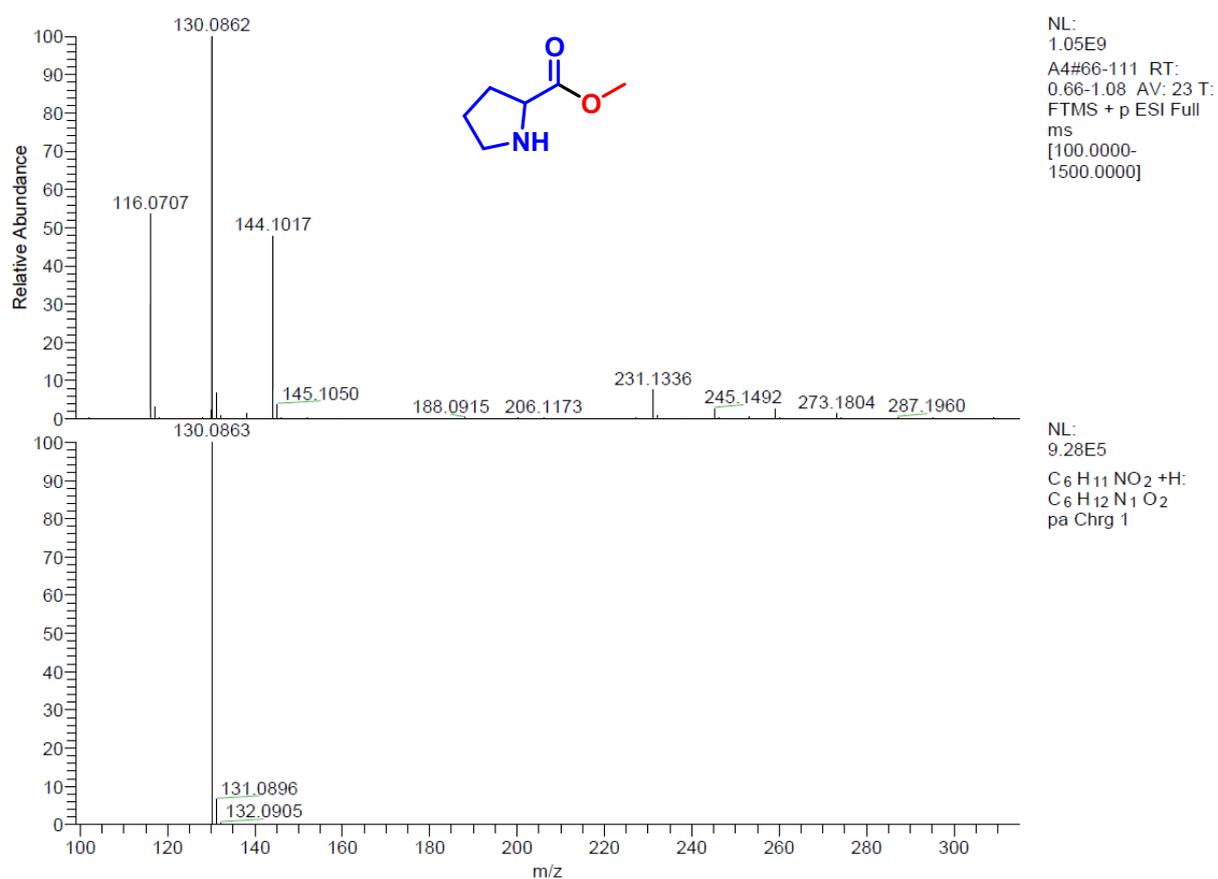


Figure S132. Mass spectrum of Methyl proline **D8**.

Analysis Info
Analysis Name C:\Users\ANUPAM MISHRA\AppData\Local\Temp\Temp1_PM.zip\PM\m chem AR-DP-A1_RA1_01_16452.d
Method 7. LCMS tune low MeOH.m
Sample Name m chem AR-DP-A1
Comment
Acquisition Date 18-04-2022 14:43:50
Operator IIT Indore
Instrument micrOTOF-Q

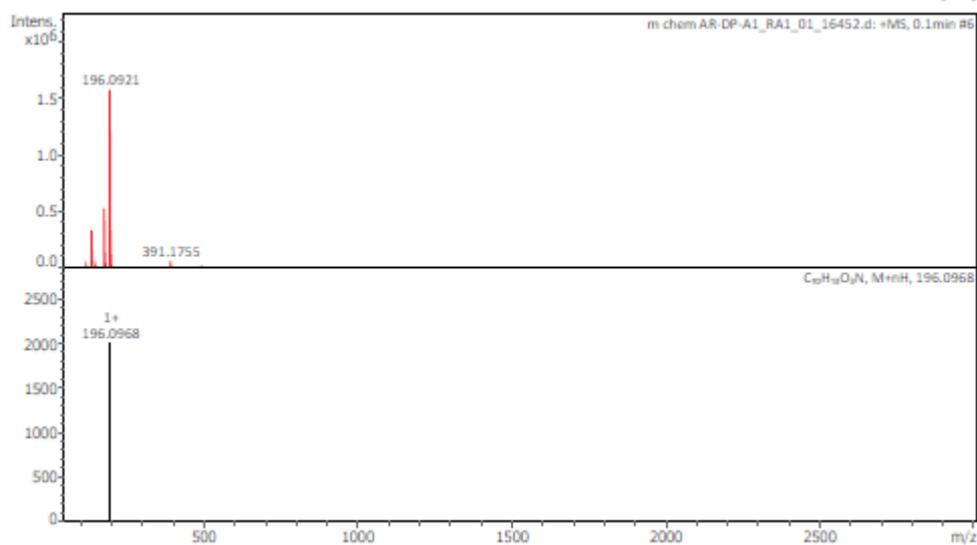
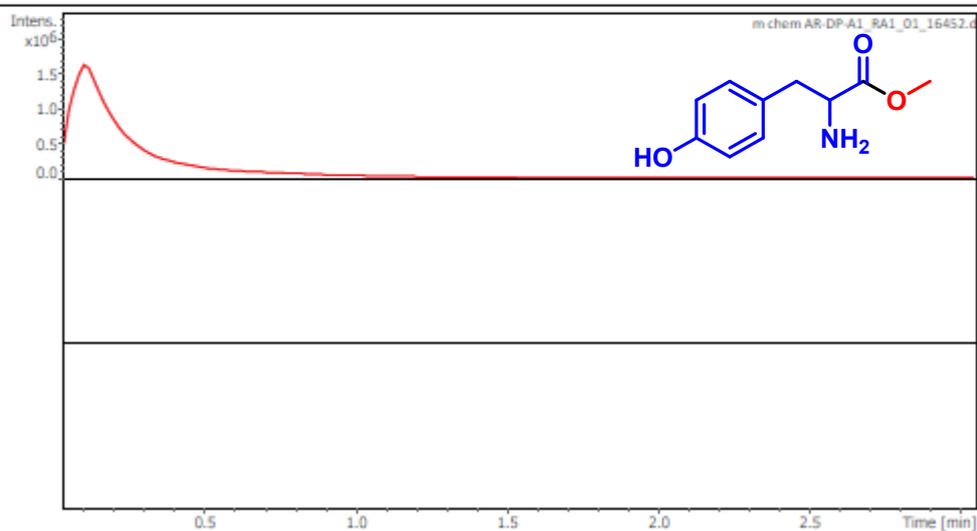


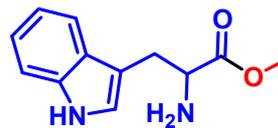
Figure S133. Mass spectrum of Methyl tyrosinate **D9**.

Single Mass Analysis

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 5



Monoisotopic Mass, Even Electron Ions

119 formula(e) evaluated with 1 results within limits (up to 1 closest results for each mass)

Elements Used:

C: 0-50 H: 0-50 N: 0-4 O: 1-10

Sample Name : A_10

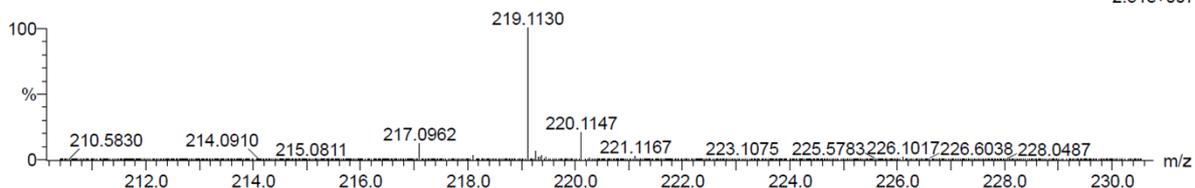
Test Name :

090522_A_10 21 (0.452)

IITRPR

XEVO G2-XS QTOF

1: TOF MS ES+
2.51e+007



Minimum: -1.5
Maximum: 2.0 10.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
219.1130	219.1134	-0.4	-1.8	6.5	2294.1	n/a	n/a	C12 H15 N2 O2

Figure S134. Mass spectrum of Methyl tryptophanate D10.

D:\HR-MS DATA\...A8

04/04/22 18:19:01

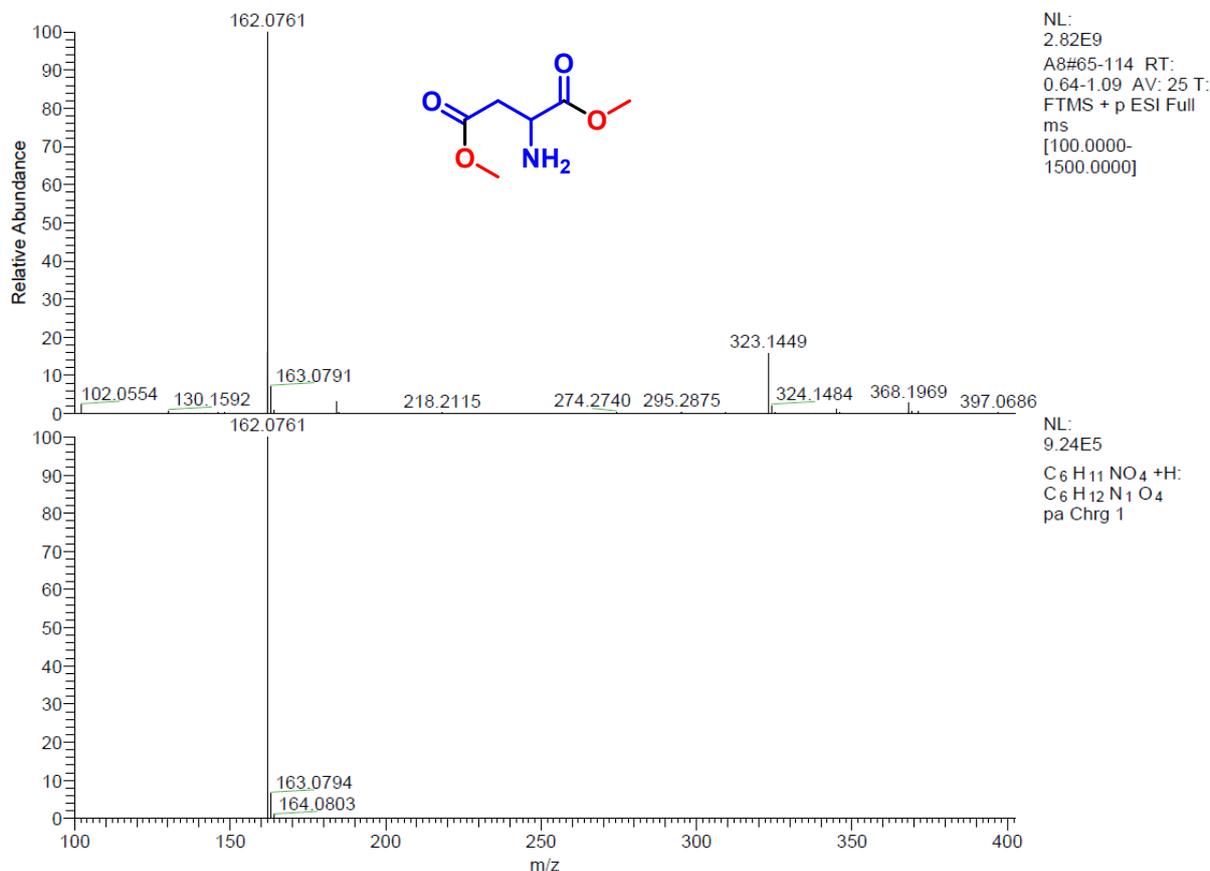


Figure S135. Mass spectrum of Dimethyl aspartate D11.

D:\HR-MS DATA\...A7

04/04/22 18:10:22

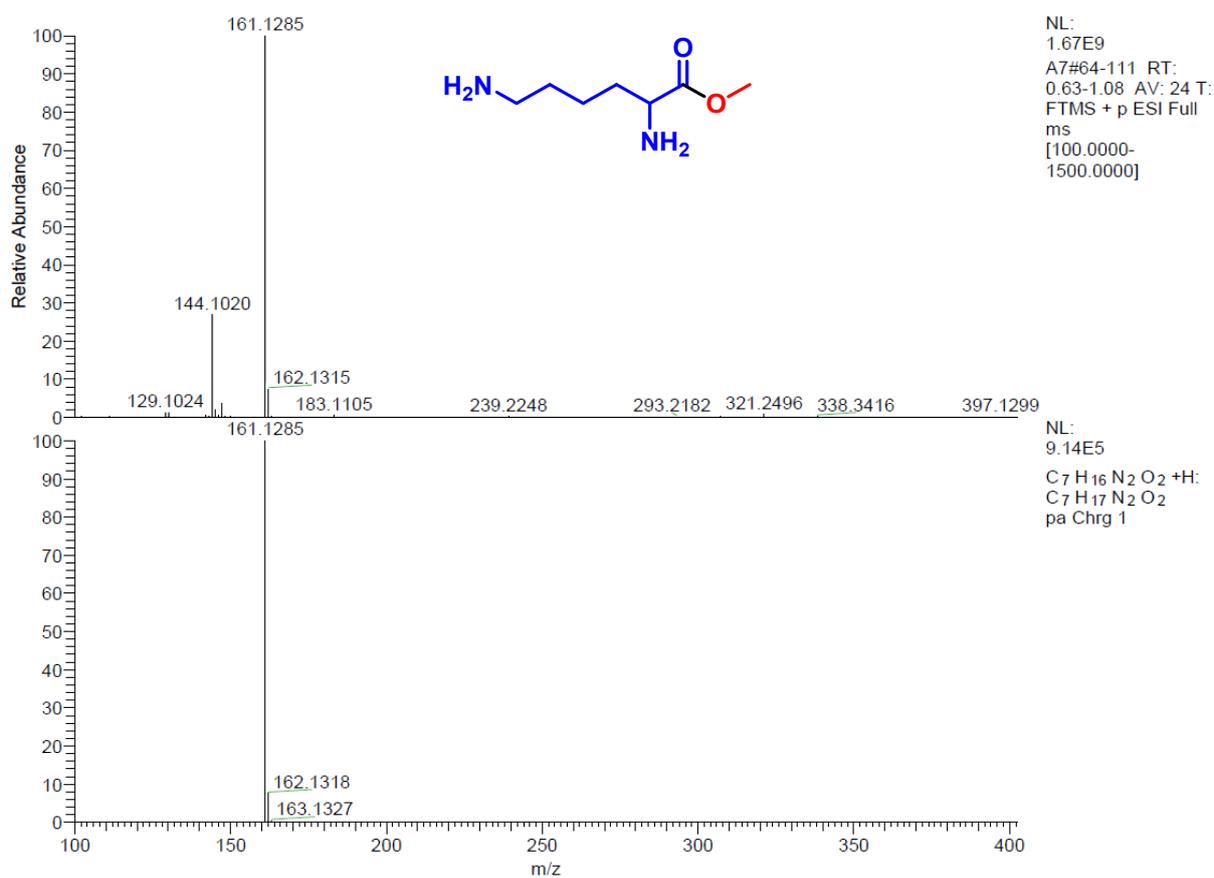


Figure S136. Mass spectrum of Methyl lysinate **D12**.

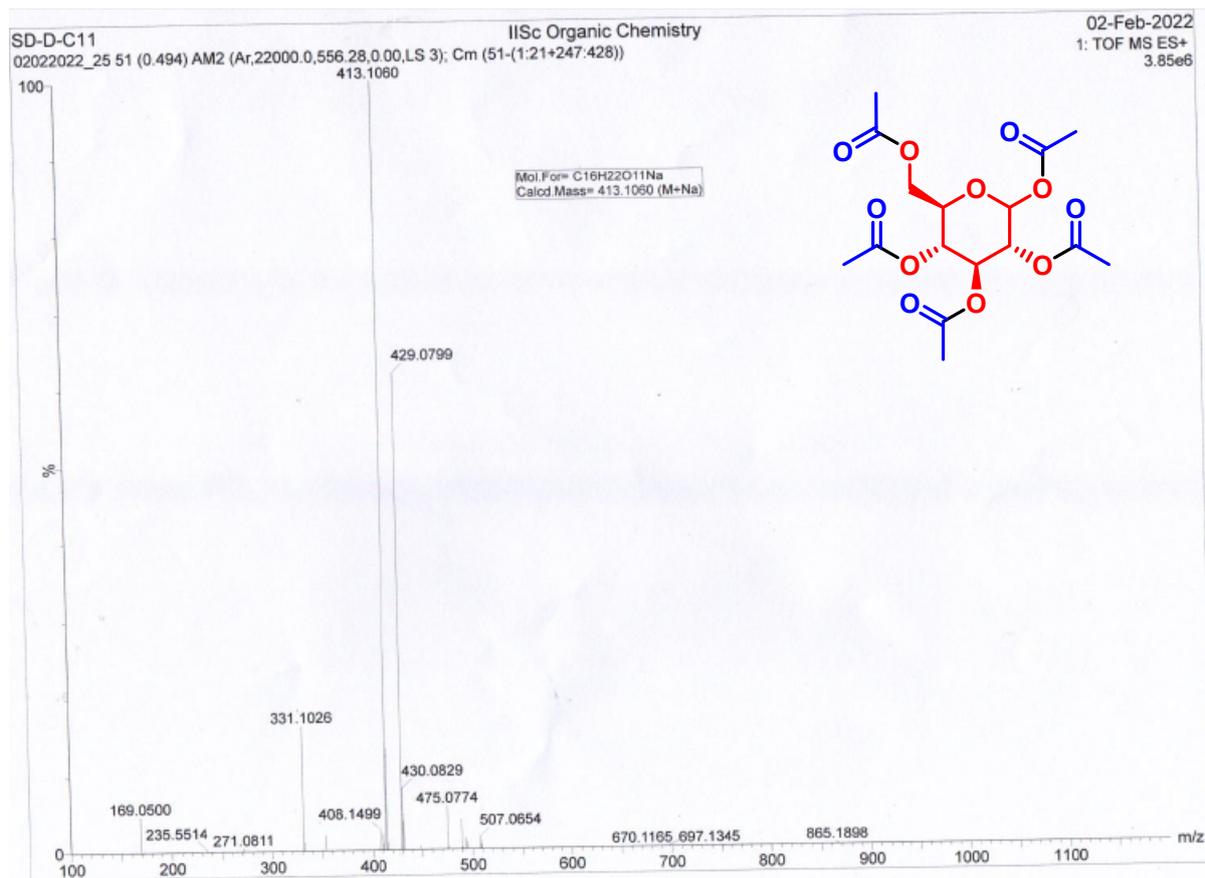


Figure S137. Mass spectrum of Glucose pentaacetate **E1**.

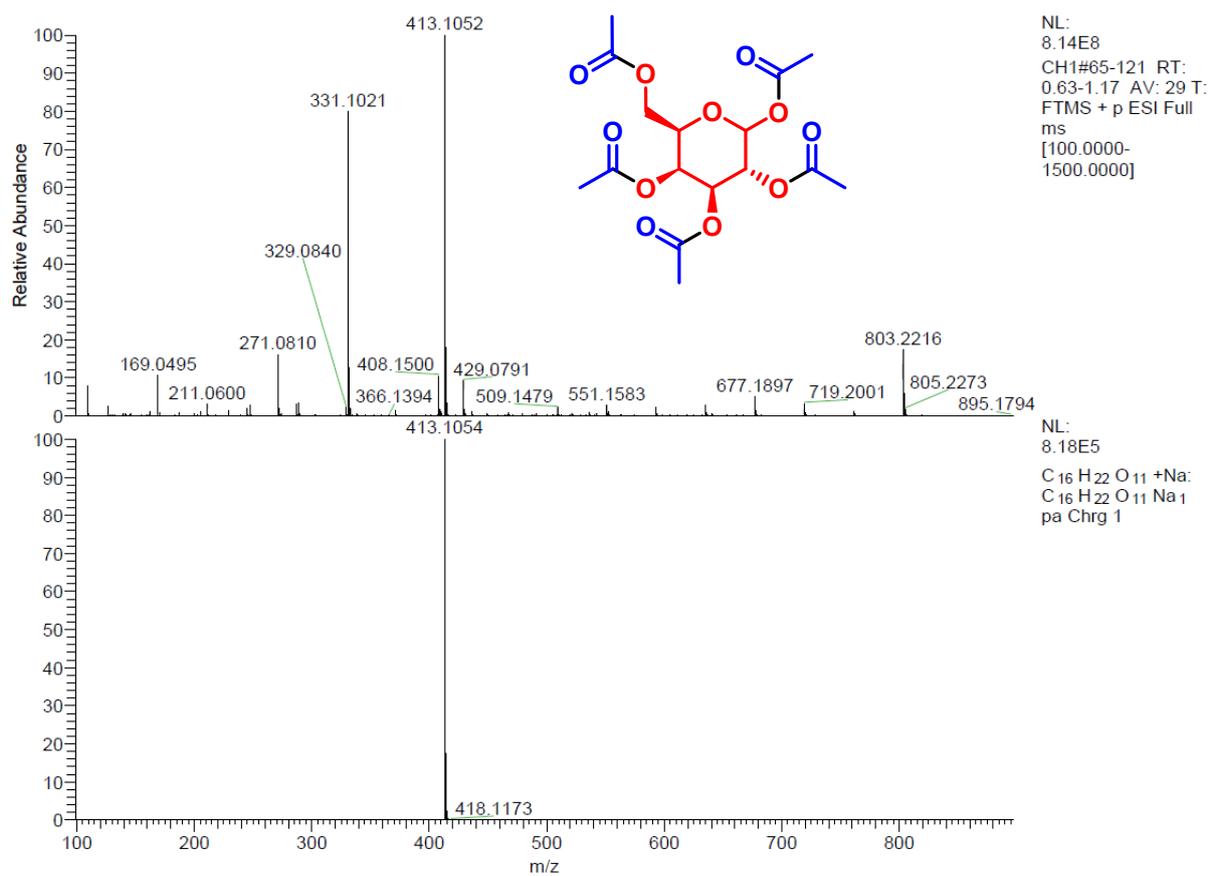
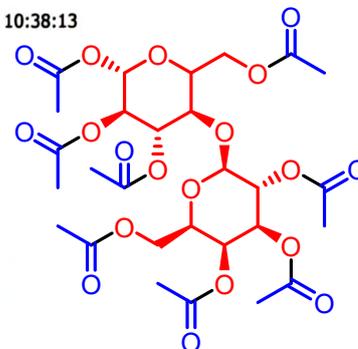


Figure S138. Mass spectrum of Galactose pentaacetate E4.

Qualitative Compound Report

Data File	LAC.d	Sample Name	LAC
Sample Type	Sample	Position	P1-A2
Instrument Name	Instrument 1	User Name	
Acq Method	MS Scan.m	Acquired Time	28-04-2022 10:38:13
IRM Calibration Status	Success	DA Method	Default.m
Comment			

Sample Group		Info.	3
Acquisition SW	6200 series TOF/6500 series		
Version	Q-TOF B.05.01 (B5125)		

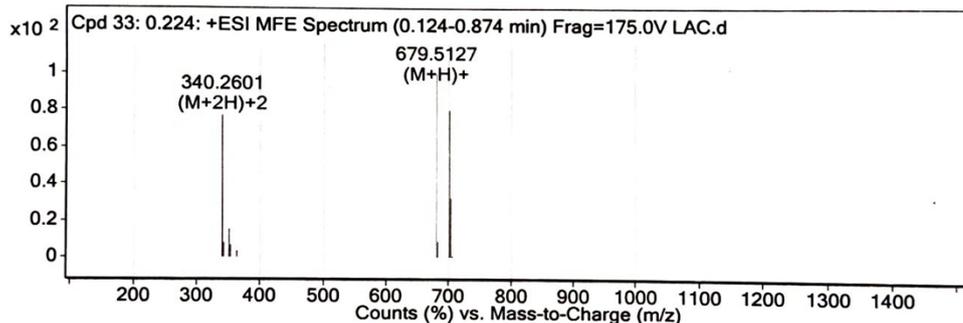


Compound Table

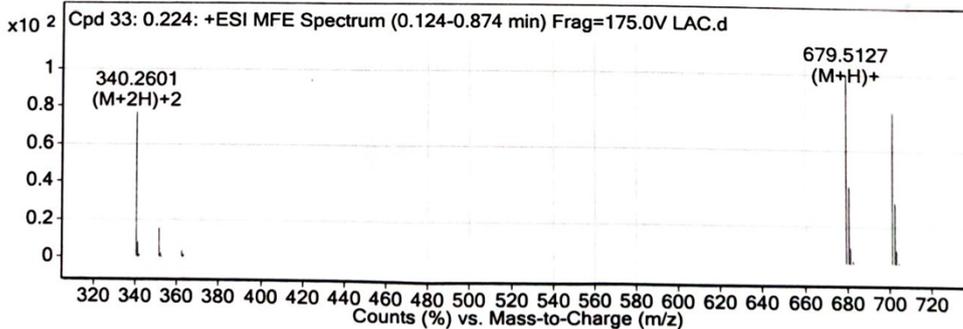
Compound Label	RT	Mass	MFG Formula
Cpd 33: 0.224	0.224	678.5053	<none>

Compound Label	m/z	RT	Algorithm	Mass
Cpd 33: 0.224	679.5127	0.224	Find by Molecular Feature	678.5053

MFE MS Spectrum



MFE MS Zoomed Spectrum



MS Spectrum Peak List

m/z	z	Abund	Ion
340.2601	2	597327.88	(M+2H)+2
340.7615	2	249486.43	(M+2H)+2
341.2632	2	61259.82	(M+2H)+2
351.2492	2	118892.21	(M+H+Na)+2
679.5127	1	778768.94	(M+H)+

Figure S139. Mass spectrum of Lactose octaacetate E7.

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