

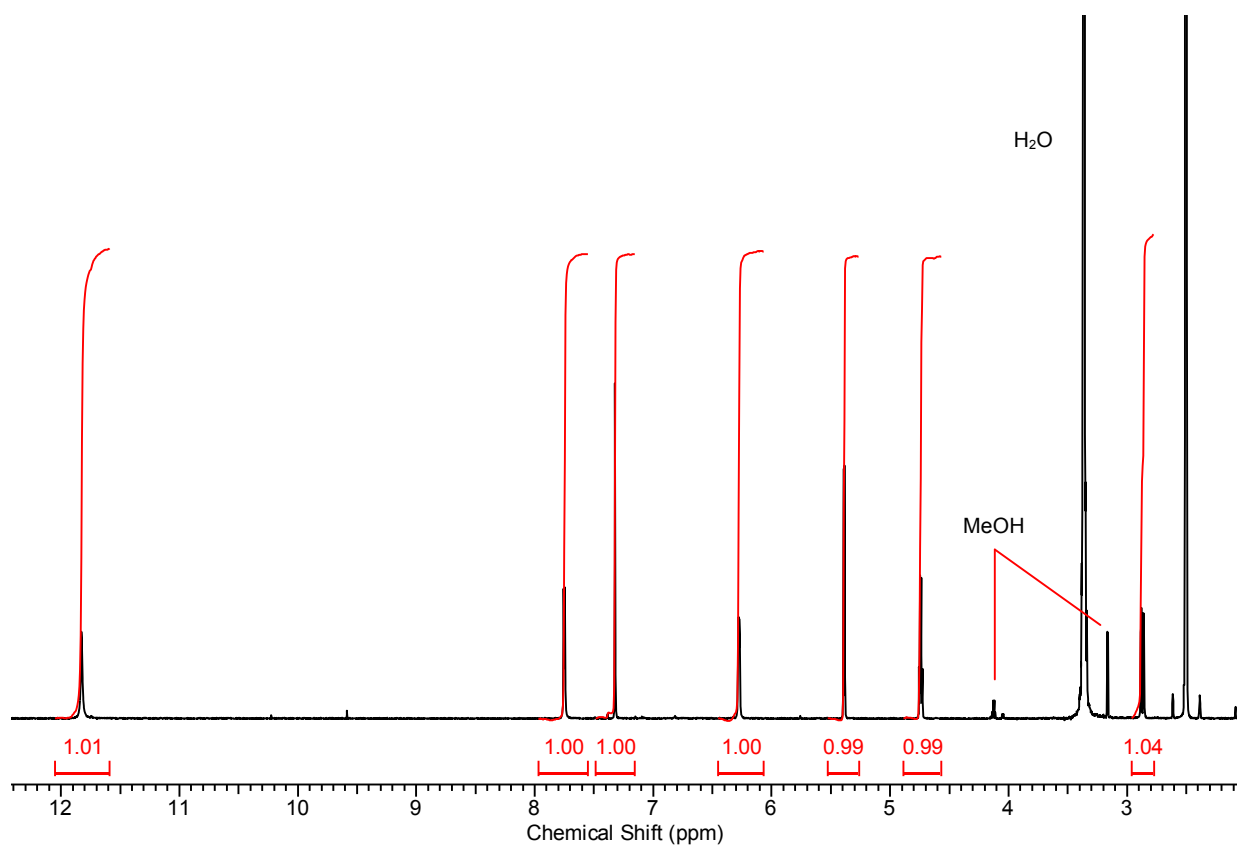
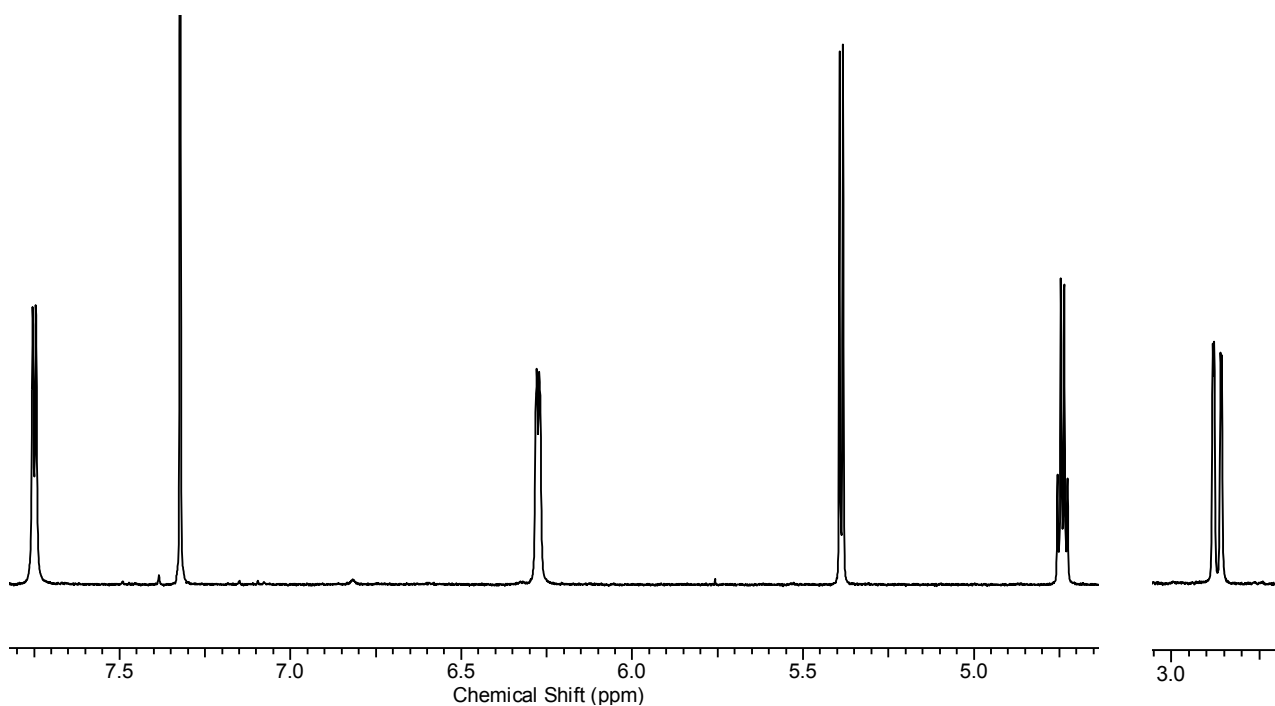
Supplementary information

Spectra

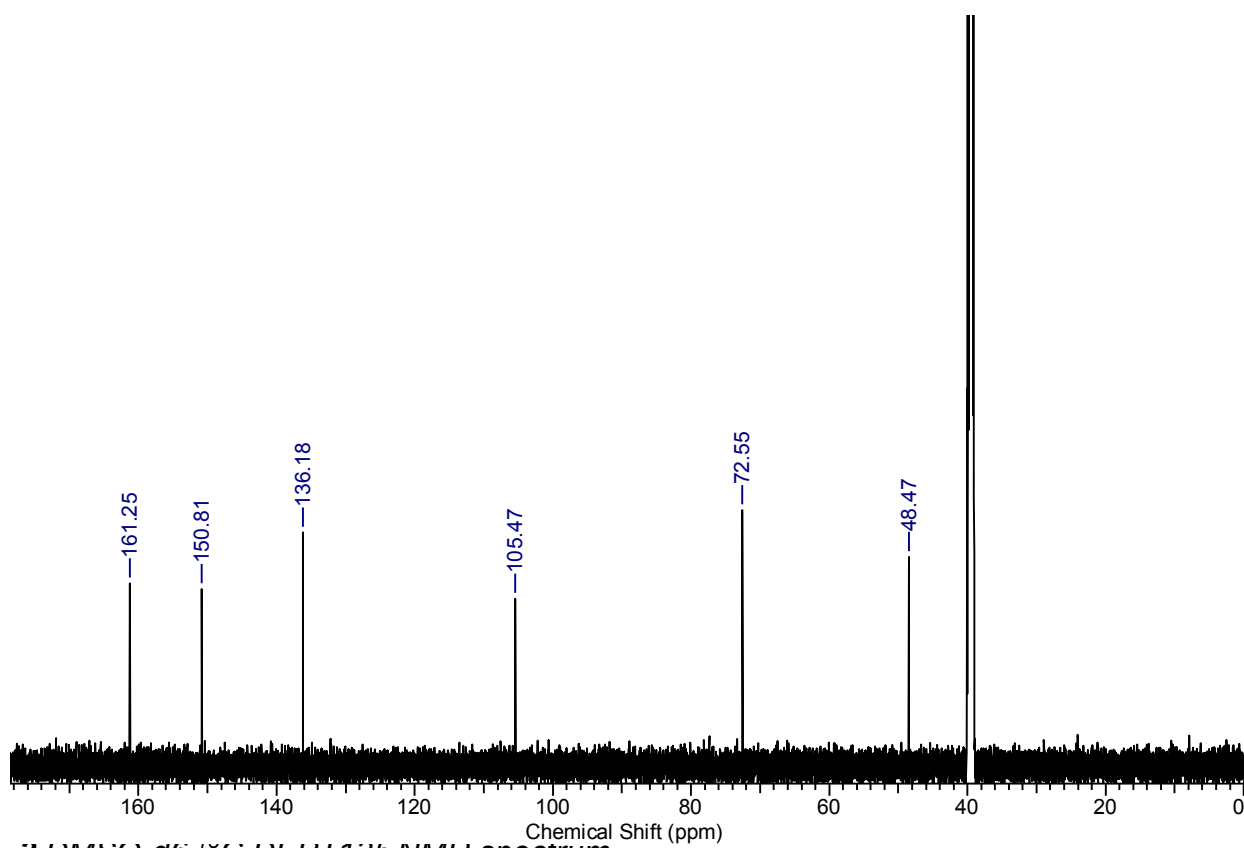
<i>Azepinomycin (2)</i>	2
<i>N1-(beta-D-ribofuranosyl)-(8R)-azepinomycin (5_A)</i>	8
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Crystallography

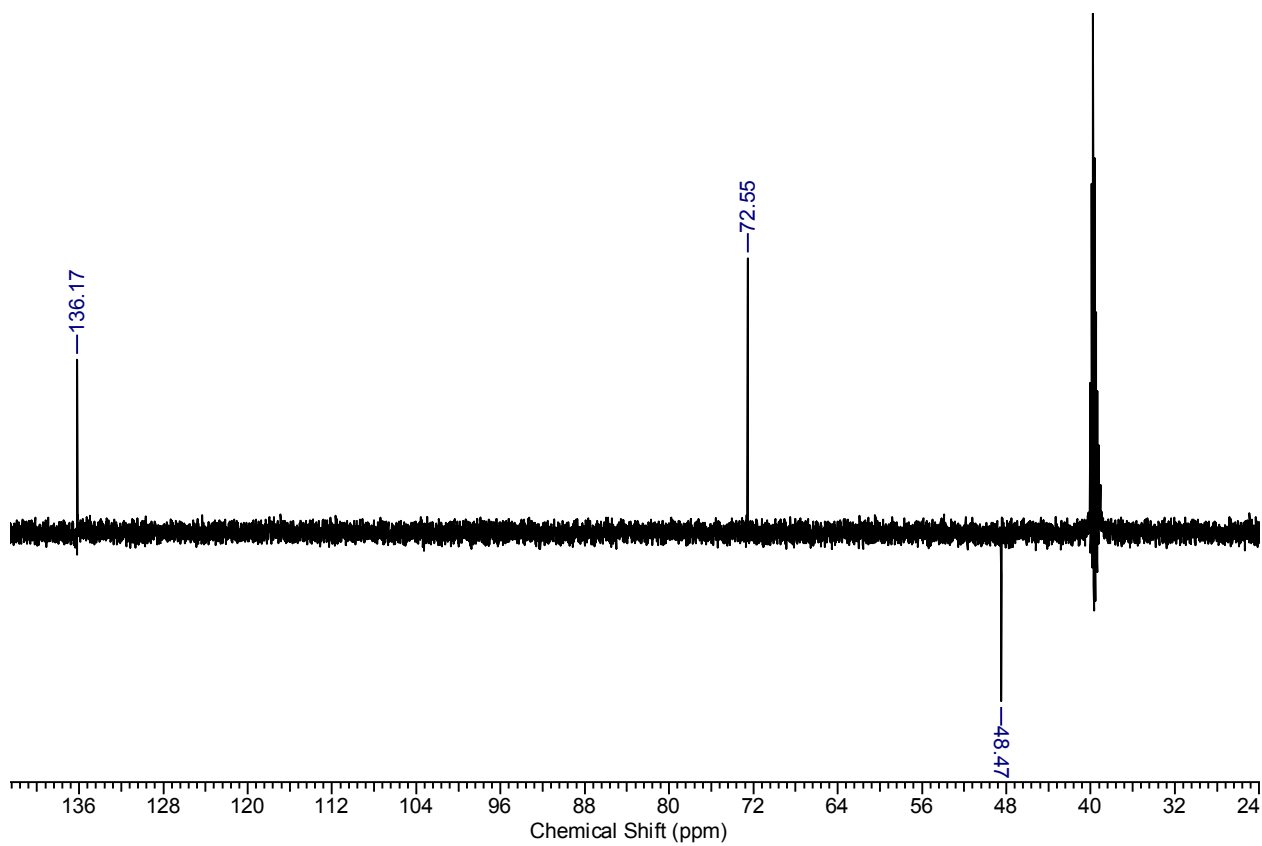
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Spectra**2 DMSO-d₆ ¹H NMR spectrum****2 DMSO-d₆ ¹H NMR spectrum expanded**

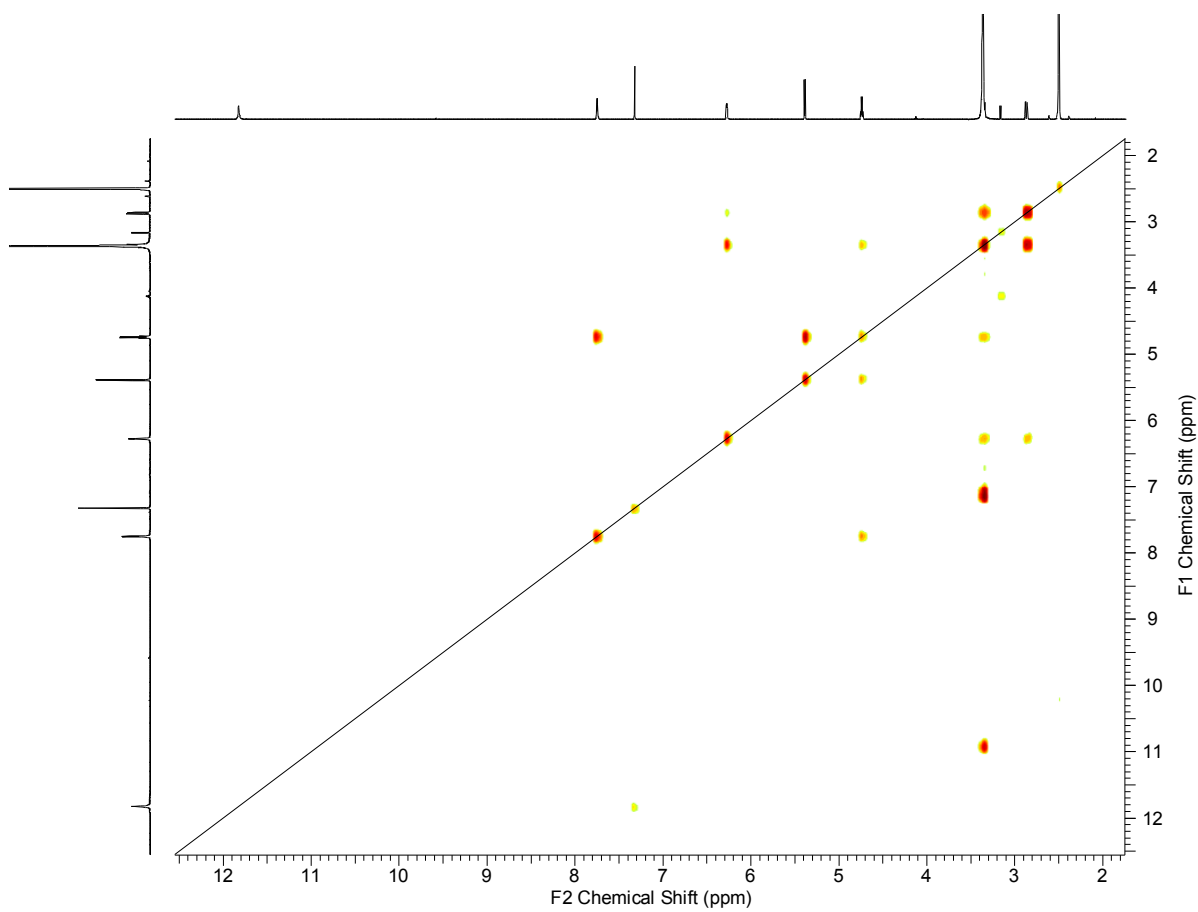
2 DMSO-d₆ ¹³C NMR spectrum



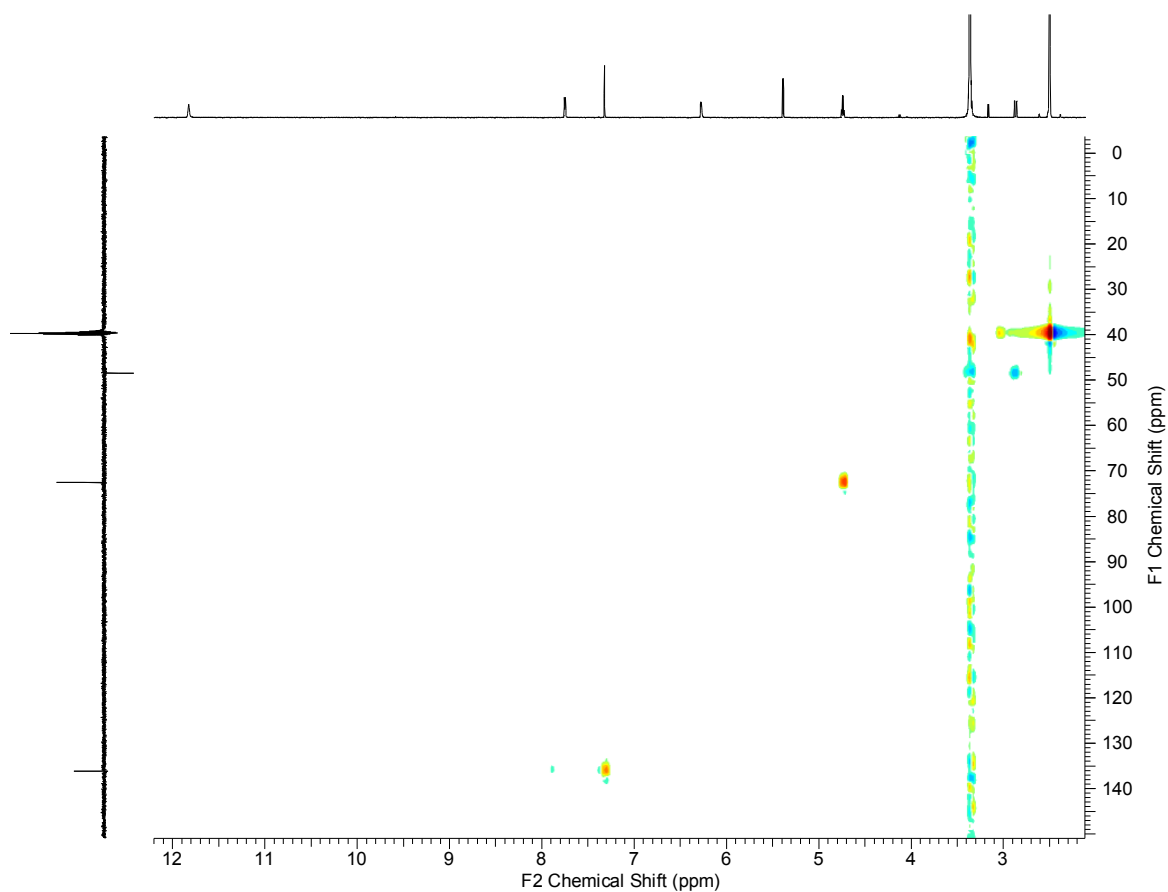
2 DMSO-d₆ ¹³C DEPT 135 NMR spectrum



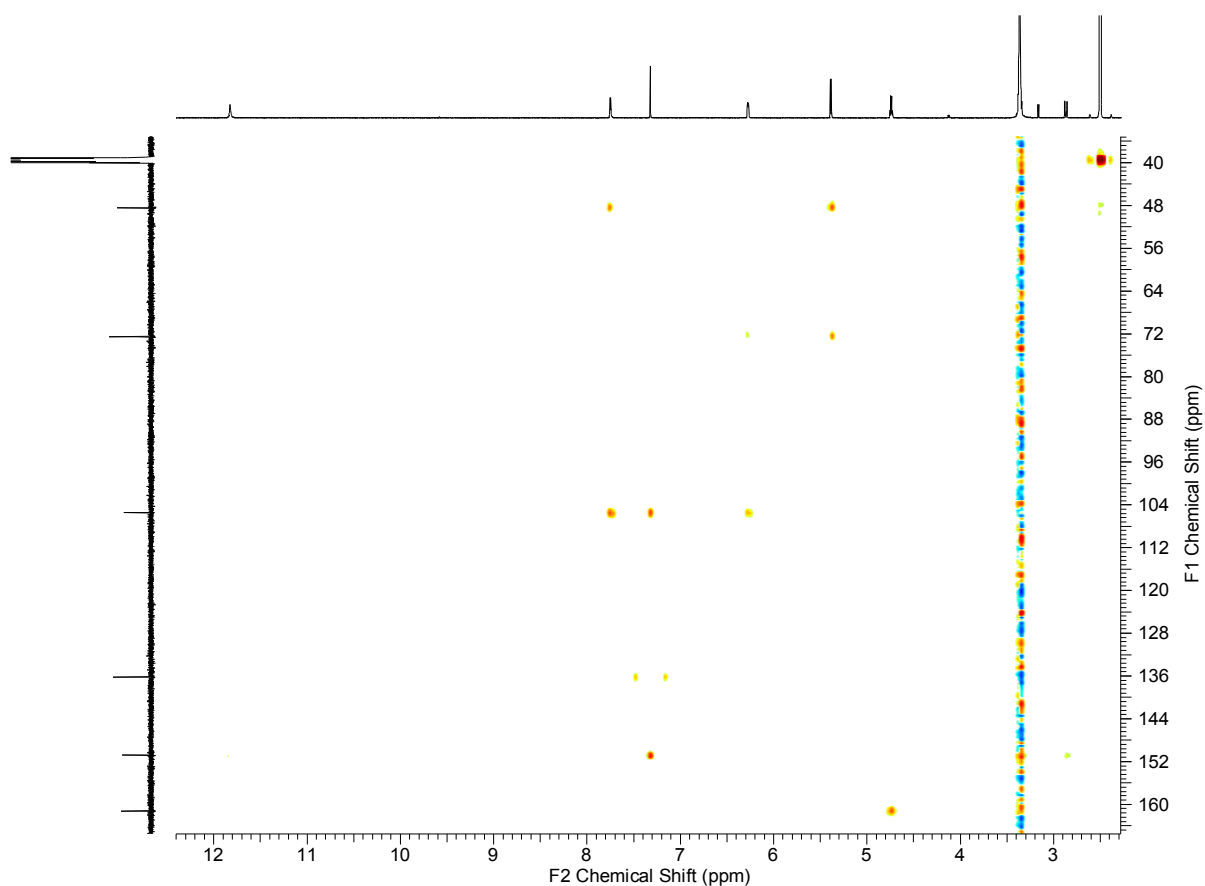
2 DMSO-d₆ ¹H-¹H COSY NMR spectrum



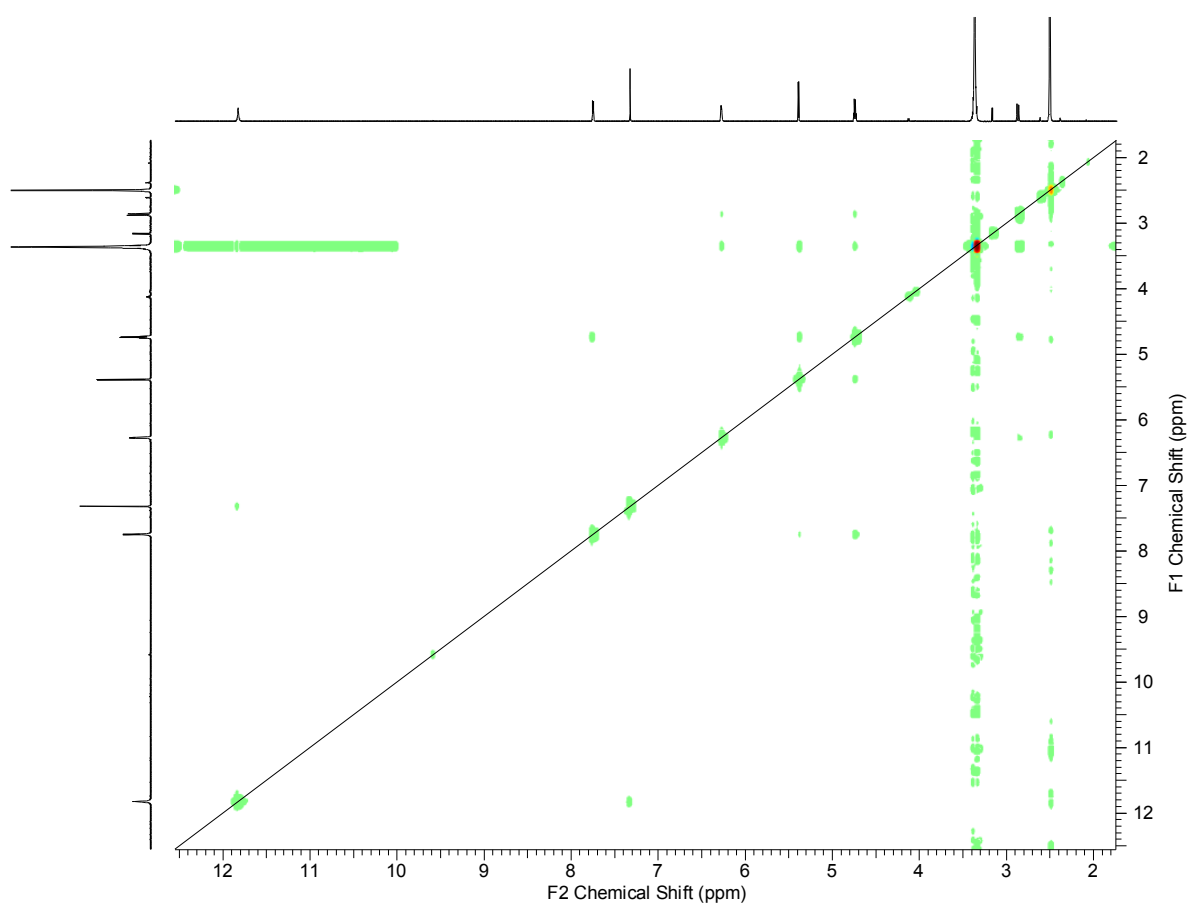
2 DMSO-d₆ ¹H-¹³C HSQC NMR spectrum

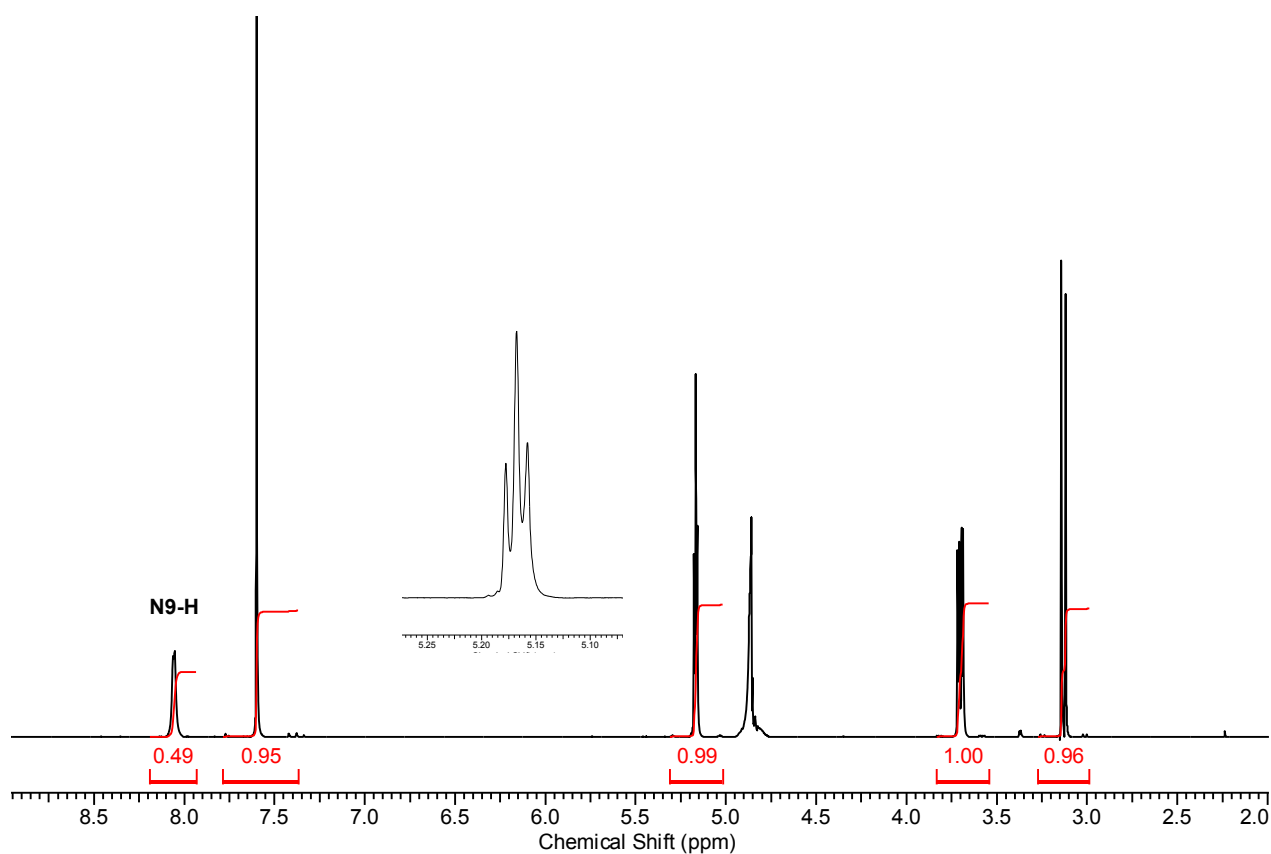
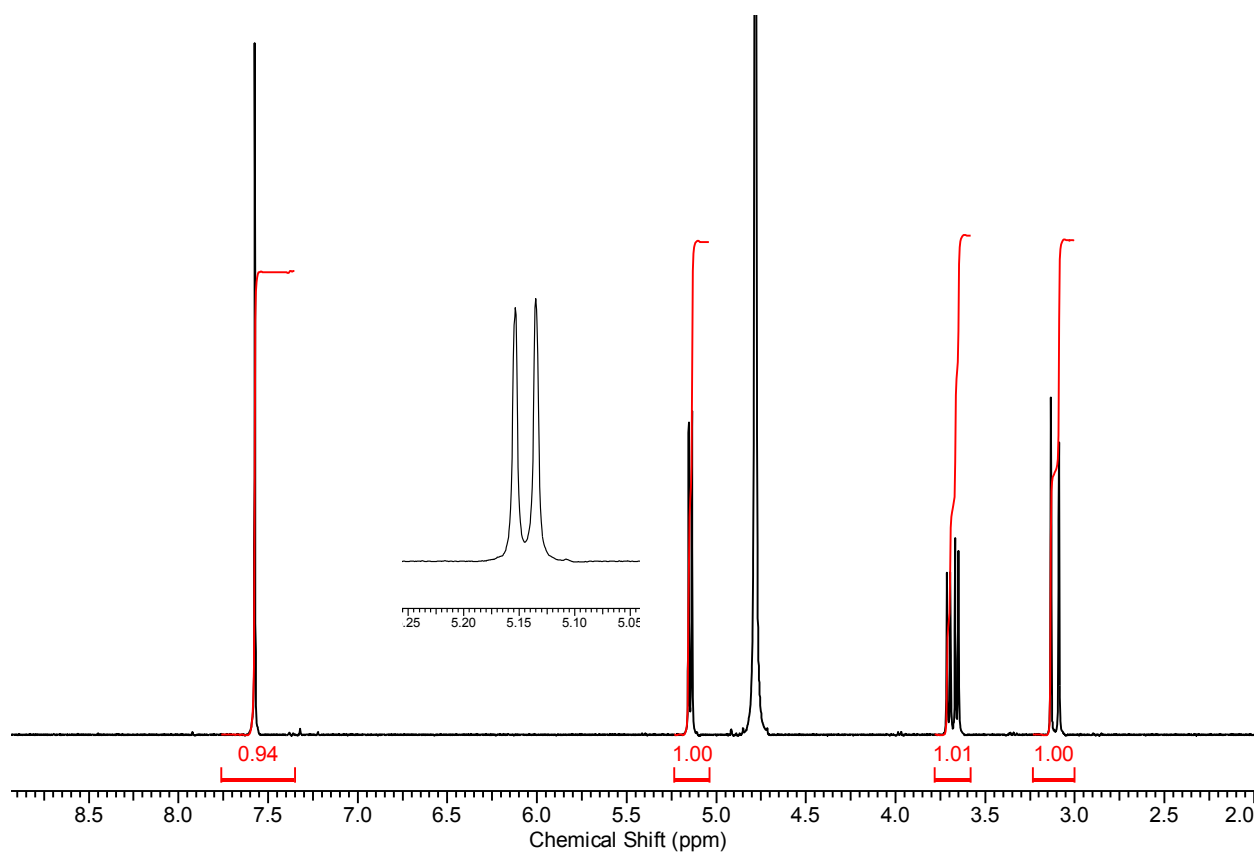


2 DMSO-d₆ ¹H-¹³C HMBC NMR spectrum

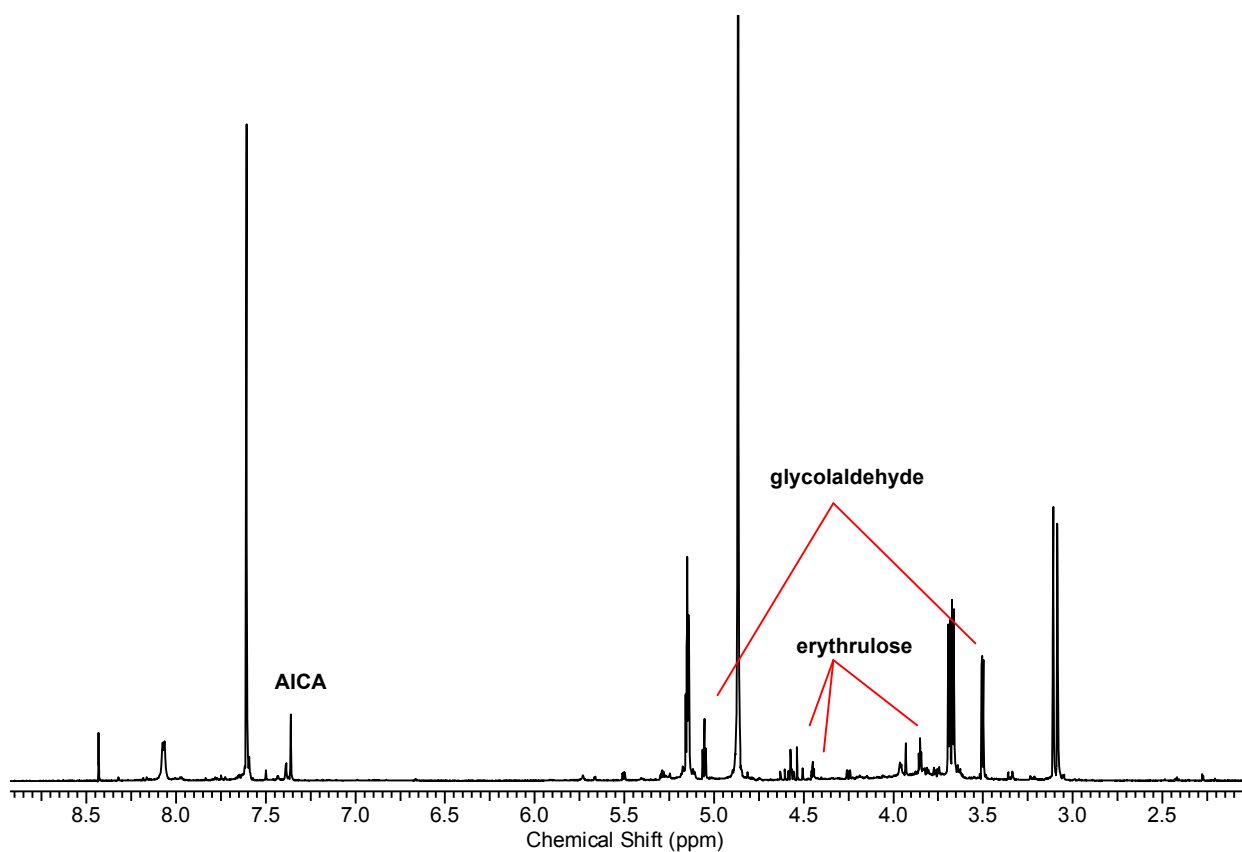


2 DMSO-d₆ ¹H-¹H NOESY NMR spectrum

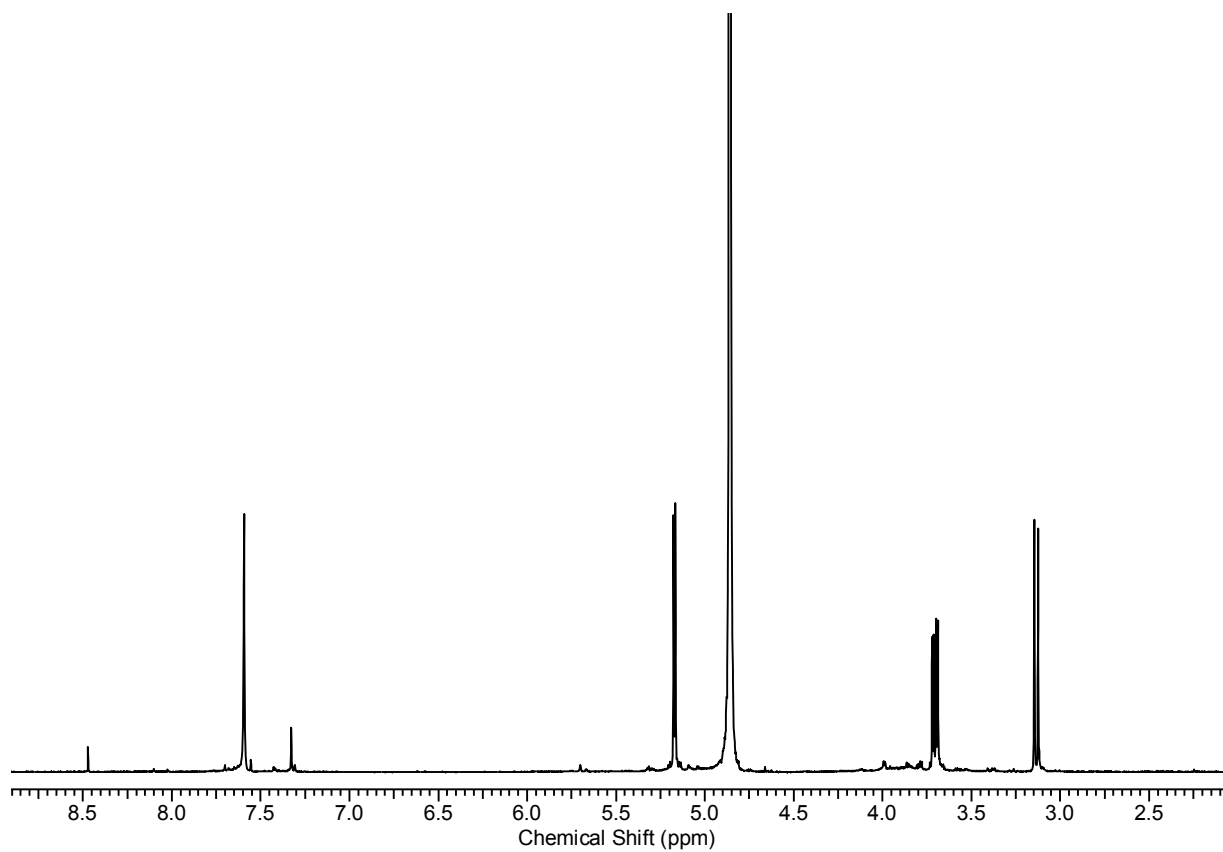


2 10% D₂O/H₂O ¹H NMR spectrum**2** D₂O ¹H NMR spectrum

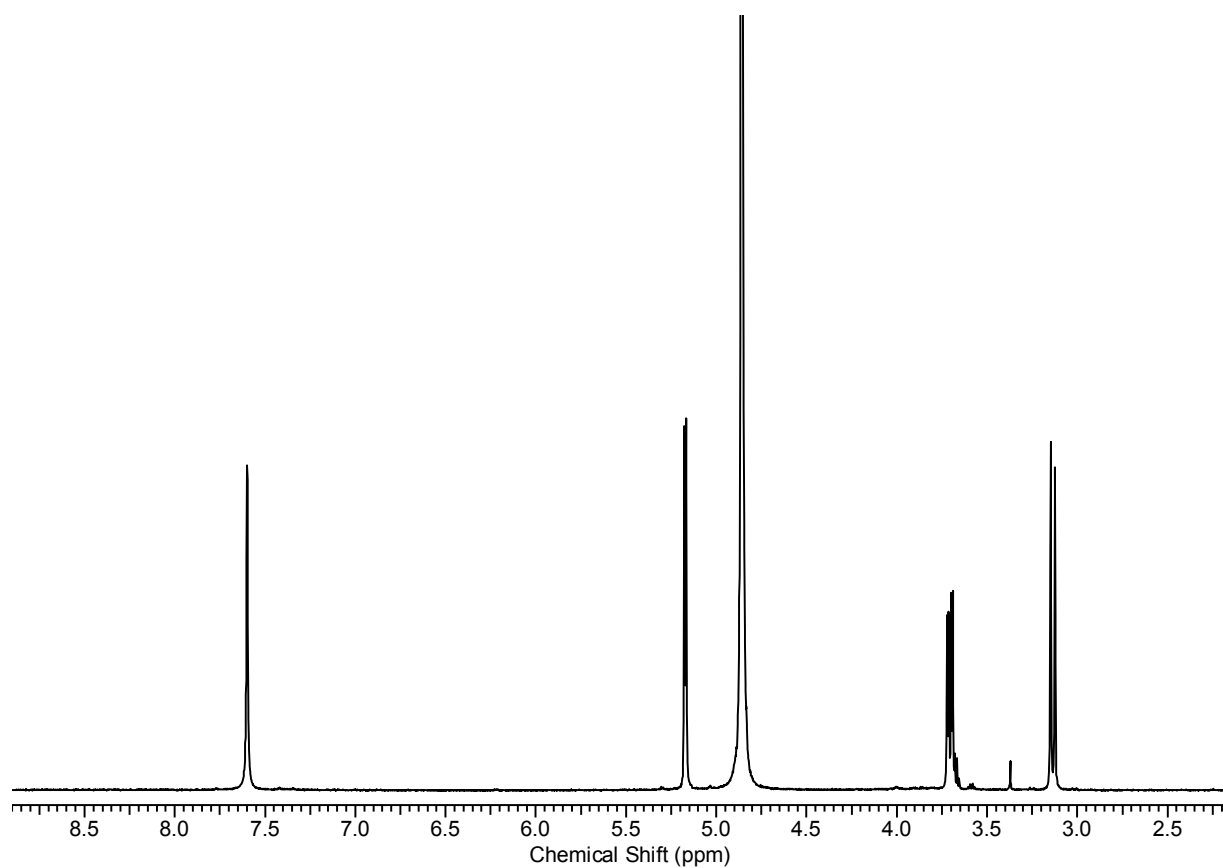
2 Reaction mixture, 28 h, 10% D₂O/H₂O ¹H NMR spectrum



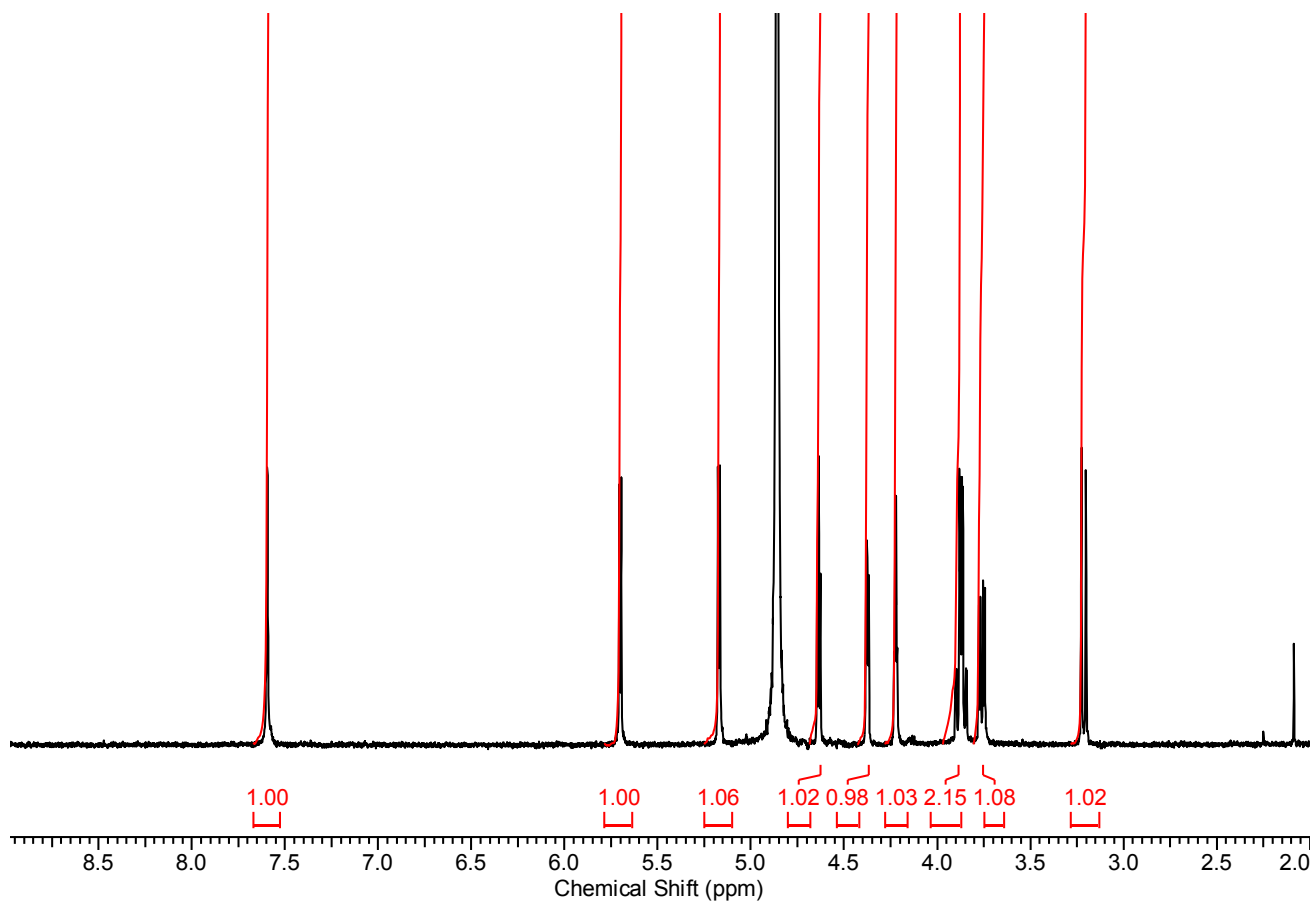
2 Crude product after Dowex exchange and lyophilisation, D₂O ¹H NMR spectrum



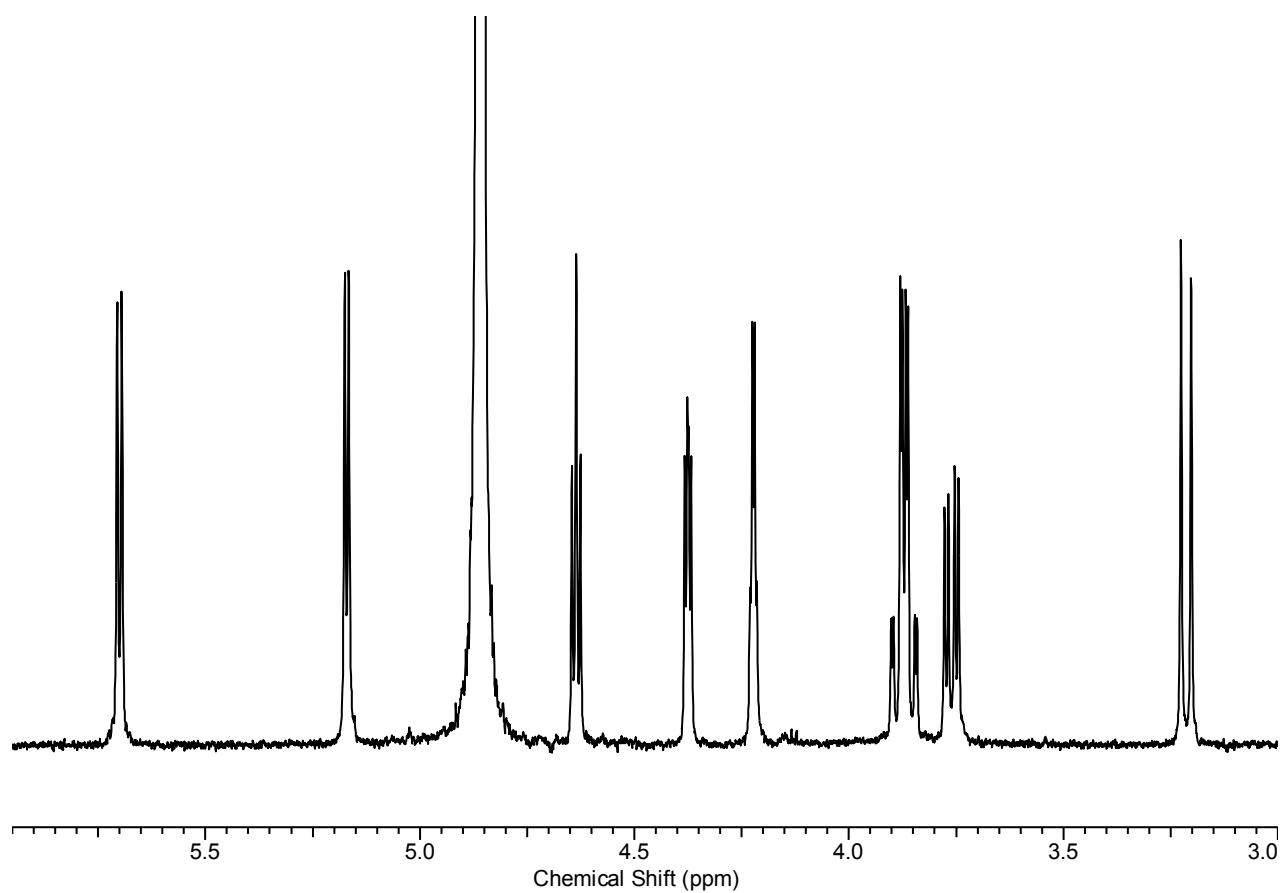
2 after precipitation from H₂O/EtOH, D₂O ¹H NMR spectrum



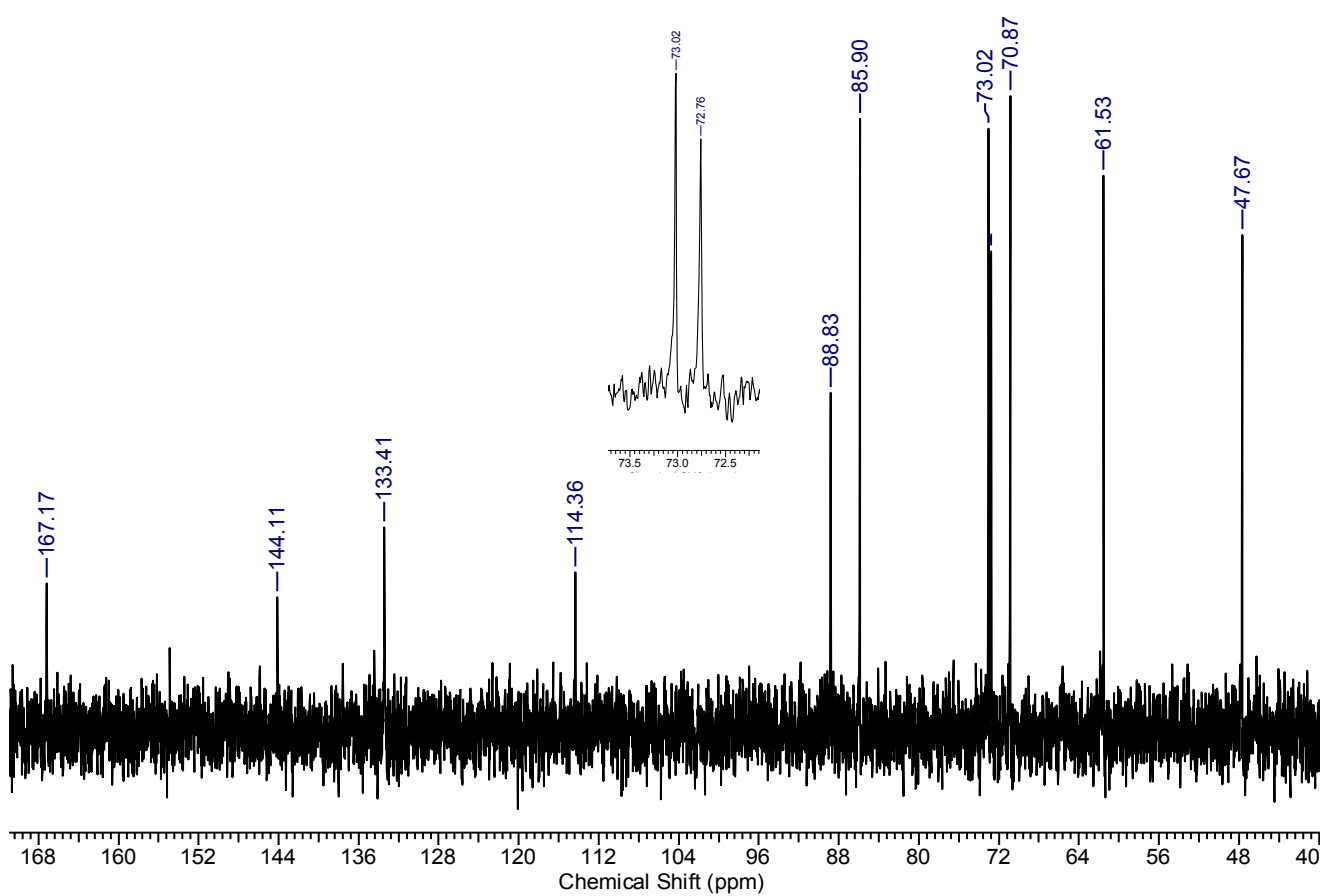
5A D₂O ¹H NMR spectrum



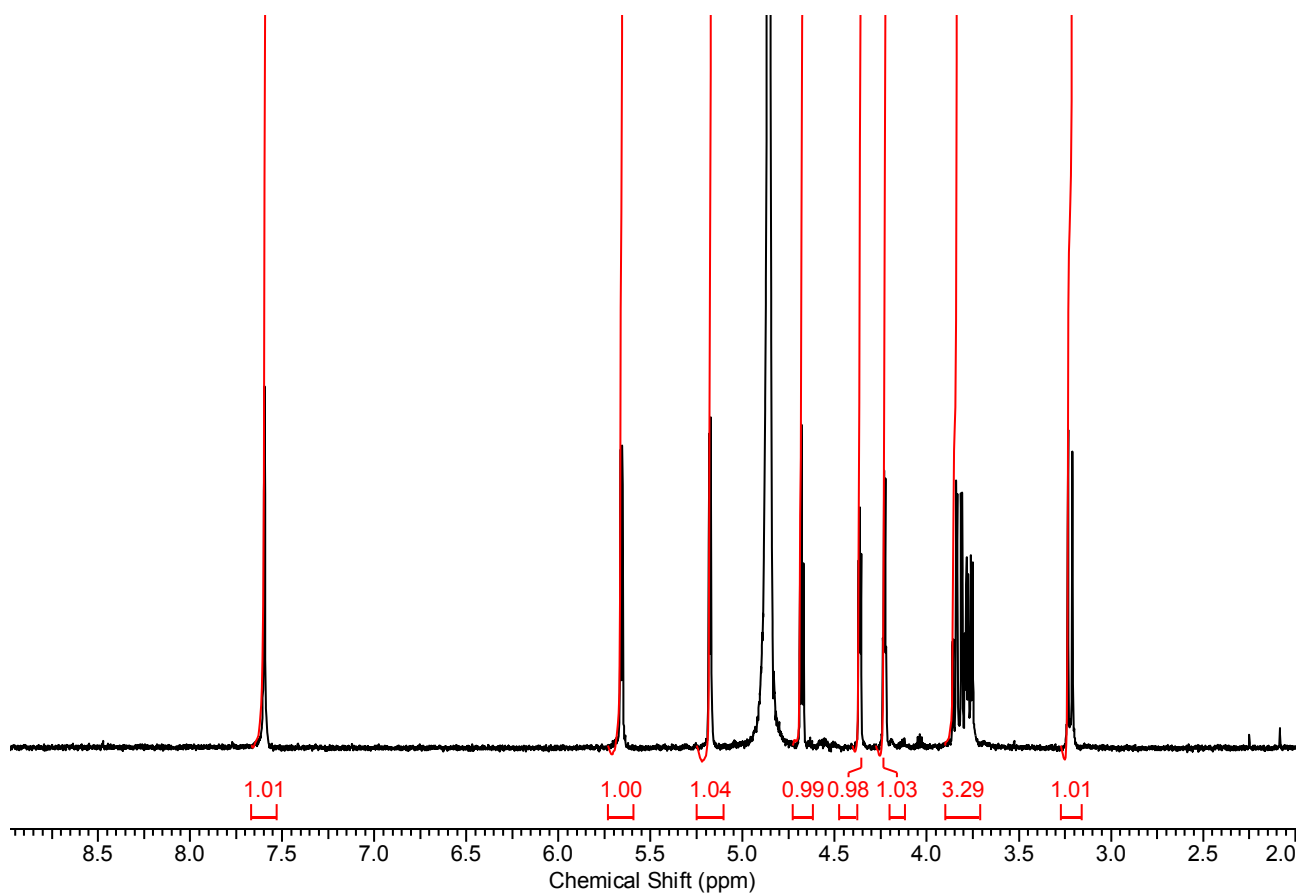
5A D₂O ¹H NMR spectrum expanded



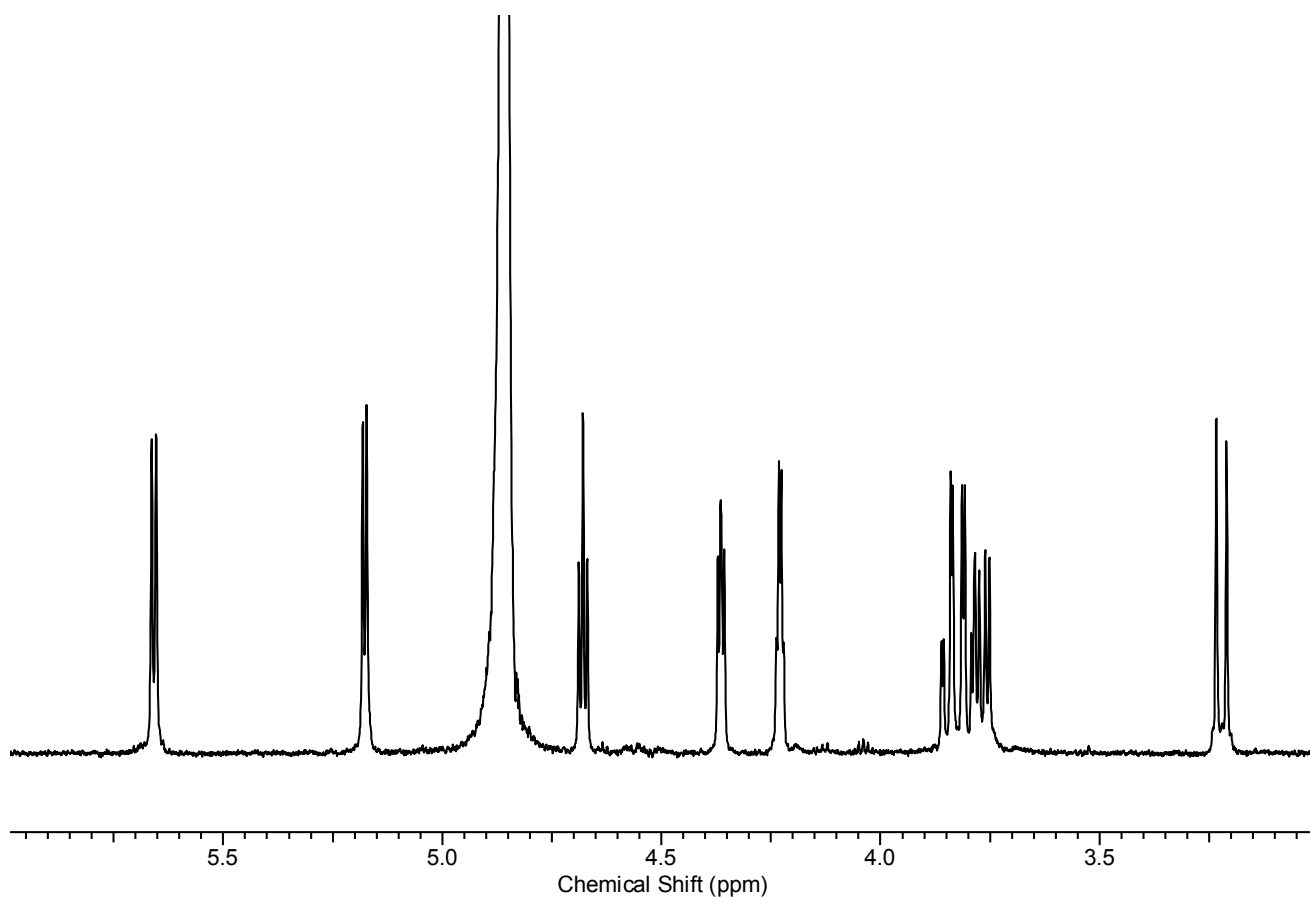
5A D₂O ¹³C NMR spectrum

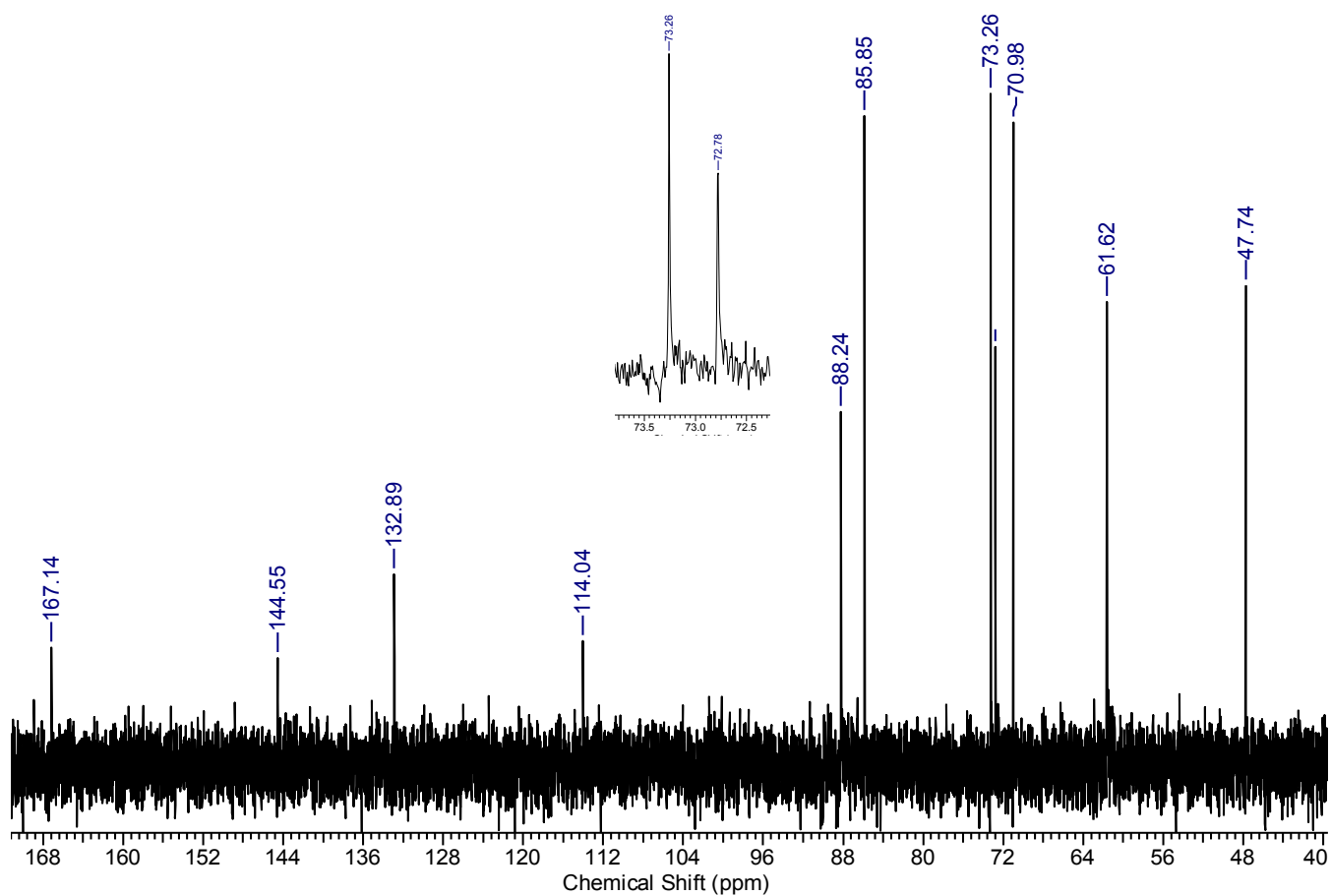
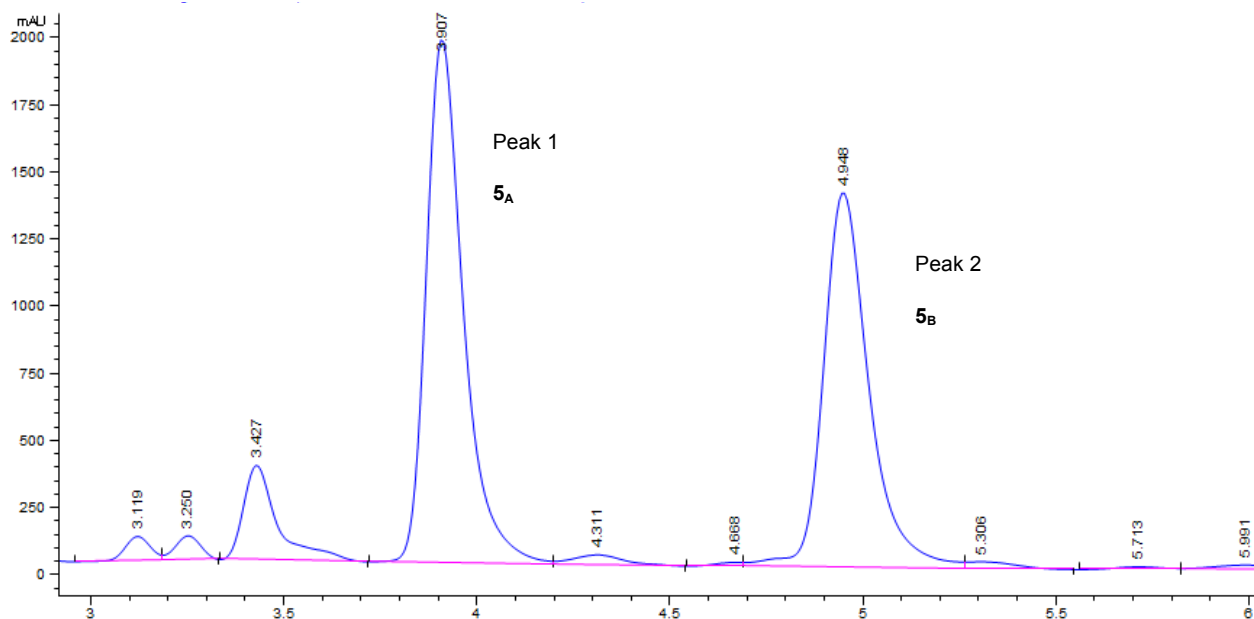


5_B D₂O ¹H NMR spectrum

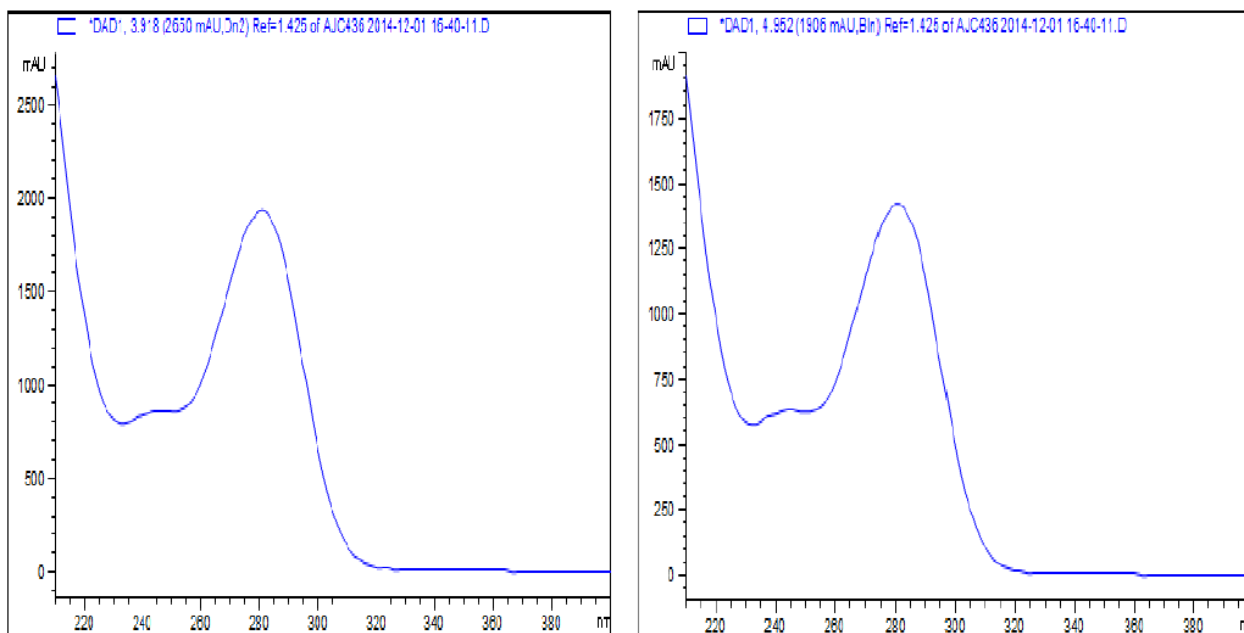


5_B D₂O ¹H NMR spectrum expanded



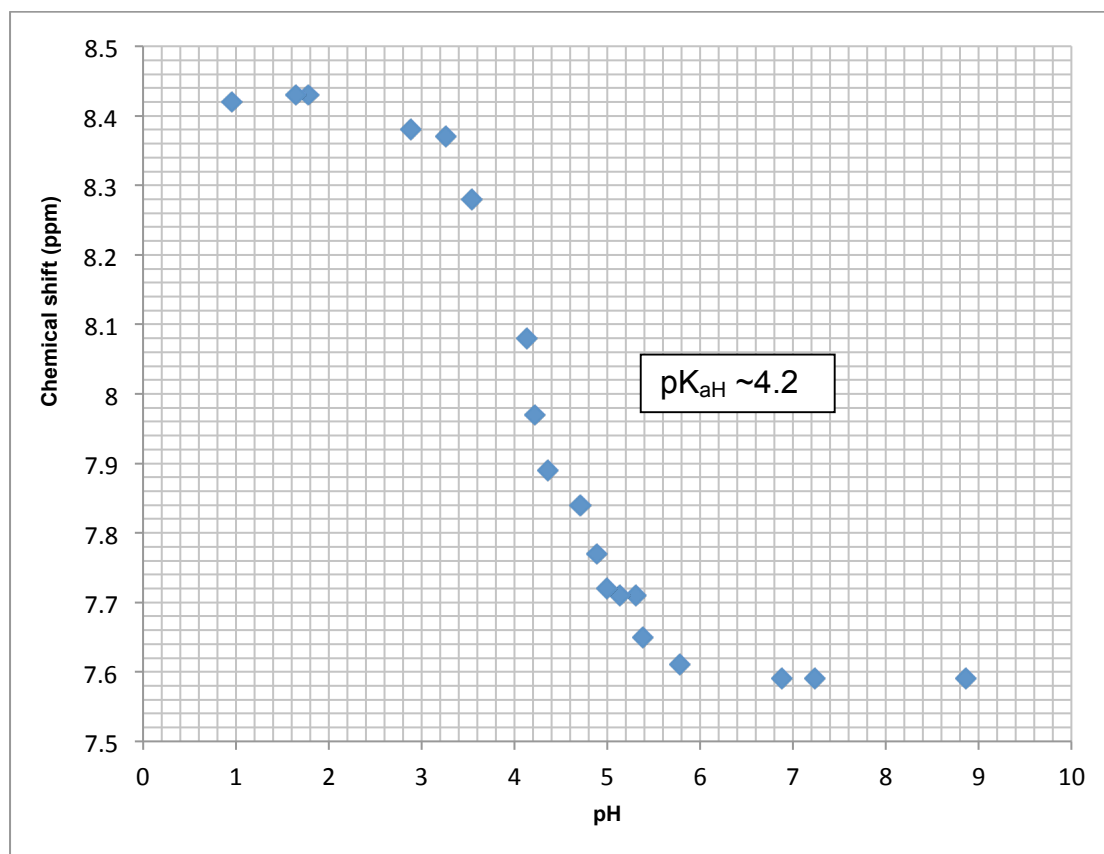
5_B D₂O ¹³C NMR spectrum**HPLC traces****Crude 5 HPLC trace, isocratic water**

UV traces for peak 1 and peak 2



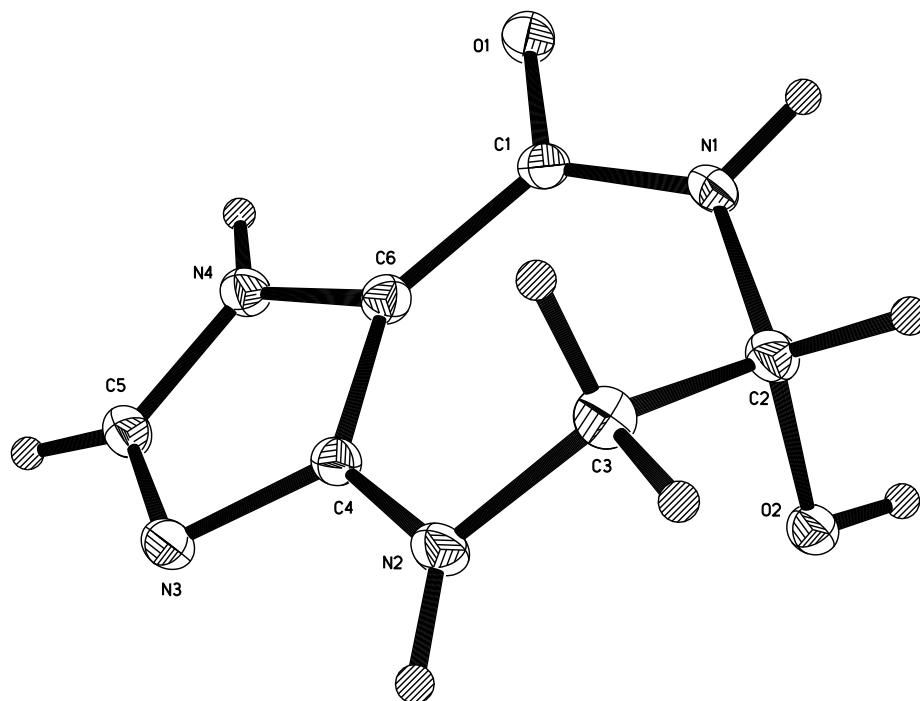
pH titration of azepinomycin (2)

NMR titration curve (measuring shift of C2-H peak vs pH) with calculated pK_{aH} for 2

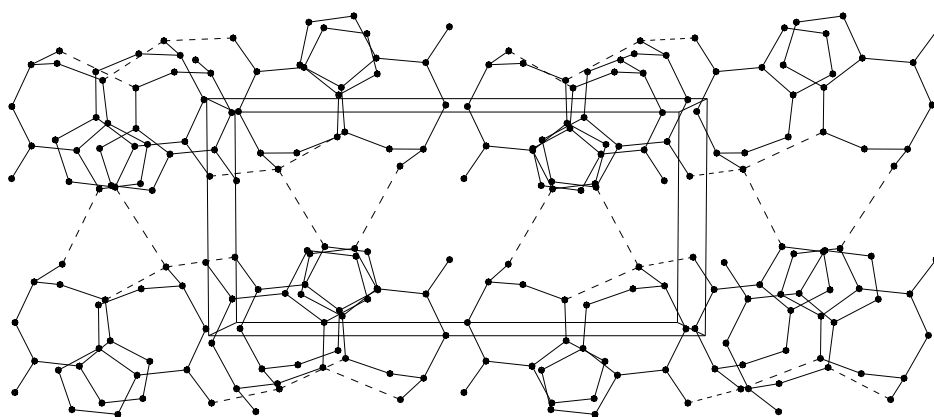


Crystallography

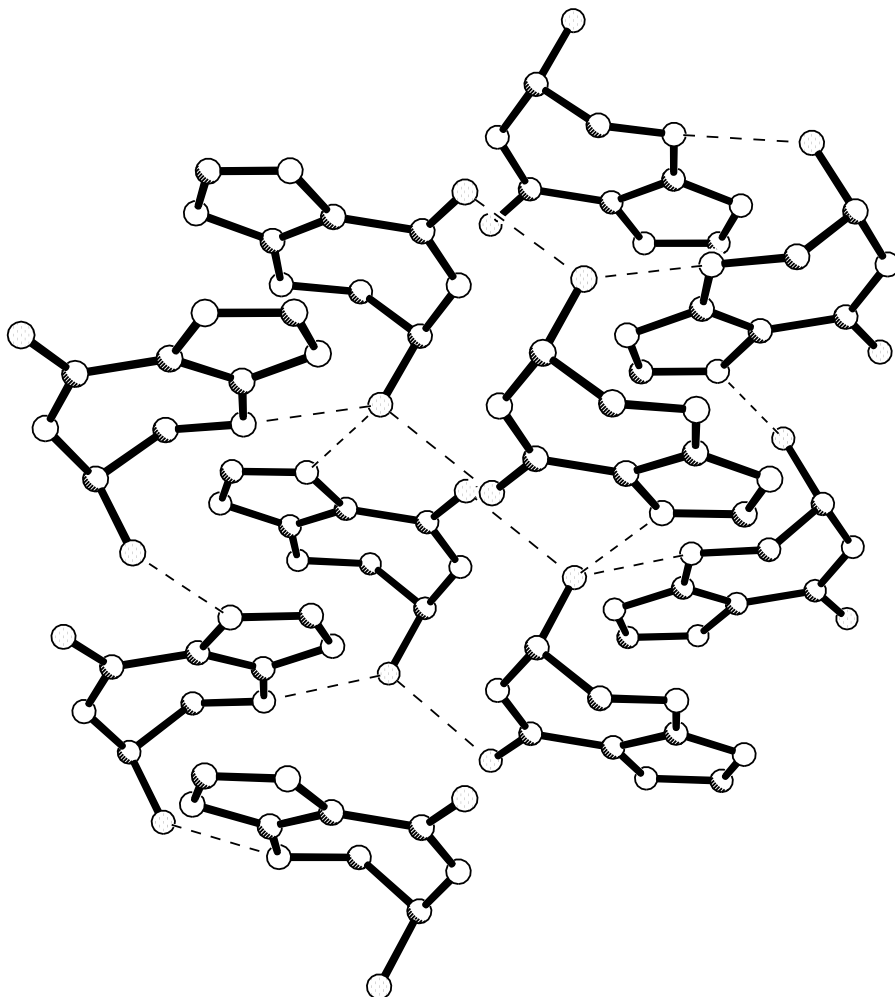
ORTEP plot of azepinomycin (2)



Cell packing of azepinomycin 2, viewed down the c-axis



Intermolecular hydrogen bonding interactions of azepinomyacin 2



Crystal data structure refinement table

Identification code	xstr0044
Empirical formula	C ₆ N ₄ O ₂ H ₈
Formula weight	168.16
Temperature/K	150
Crystal system	monoclinic
Space group	P2 ₁ /c
a/Å	6.68948(7)
b/Å	14.04389(15)
c/Å	7.22458(9)
α/°	90
β/°	93.8750(11)
γ/°	90
Volume/Å ³	677.171(13)
Z	4
ρ _{calc} /mg/mm ³	1.649
m/mm ⁻¹	1.090
F(000)	352.0
Crystal size/mm ³	0.28 × 0.26 × 0.15
Radiation	Cu Kα (λ = 1.5418)
2θ range for data collection	12.606 to 147.344°
Index ranges	-8 ≤ h ≤ 8, -17 ≤ k ≤ 17, -8 ≤ l ≤ 8
Reflections collected	9206
Independent reflections	1328[R(int) = 0.0198]
Data/restraints/parameters	1328/0/110
Goodness-of-fit on F ²	1.088
Final R indexes [I ≥ 2σ (I)]	R ₁ = 0.0383, wR ₂ = 0.1004
Final R indexes [all data]	R ₁ = 0.0389, wR ₂ = 0.1007
Largest diff. peak/hole / e Å ⁻³	0.69/-0.66

Bond lengths table

Atom	Atom	Length/Å	Atom	Atom	Length/Å
O1	C1	1.2558(17)	N3	C5	1.3307(18)
O2	C2	1.4235(16)	N4	C5	1.3338(19)
N1	C1	1.3498(18)	N4	C6	1.3909(17)
N1	C2	1.4477(17)	C1	C6	1.4477(19)
N2	C3	1.4482(18)	C2	C3	1.5326(19)
N2	C4	1.3678(18)	C4	C6	1.3922(18)
N3	C4	1.3680(18)			

Bond angles table

Atom	Atom	Atom	Angle/°	Atom	Atom	Atom	Angle/°
C1	N1	C2	127.73(11)	N1	C2	C3	114.13(11)
C4	N2	C3	117.48(11)	N2	C3	C2	112.87(11)
C5	N3	C4	104.93(11)	N2	C4	N3	120.91(12)
C5	N4	C6	107.52(11)	N2	C4	C6	128.62(13)
O1	C1	N1	120.63(12)	N3	C4	C6	110.47(12)
O1	C1	C6	119.56(12)	N3	C5	N4	112.67(12)
N1	C1	C6	119.80(12)	N4	C6	C1	119.71(12)
O2	C2	N1	112.05(11)	N4	C6	C4	104.40(12)
O2	C2	C3	108.33(10)	C4	C6	C1	135.80(13)