

Supplementary data

New ophiobolins from the deep-sea derived fungus *Aspergillus sp.* WHU0154 and their anti-inflammatory effects

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Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

849 formula(e) evaluated with 2 results within limits (up to 50 closest results for each mass)

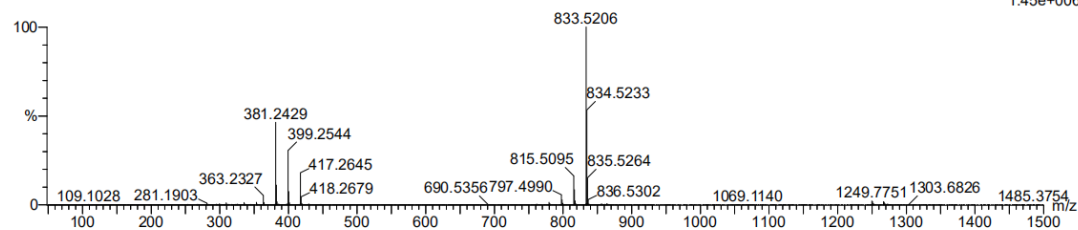
Elements Used:

C: 0-50 H: 0-100 N: 0-10 O: 0-100

0154-78-67-5-5

20191216009 172 (1.388)

1: TOF MS ES+
1.45e+006



Minimum: 5.0 5.0 -1.5
Maximum: 5.0 5.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
417.2645	417.2641	0.4	1.0	7.5	384.9	0.219	80.36	C25 H37 O5
	417.2654	-0.9	-2.2	12.5	386.4	1.628	19.64	C26 H33 N4 O

Fig. S1 HR ESI-Q-TOF-MS spectrum of compound 1

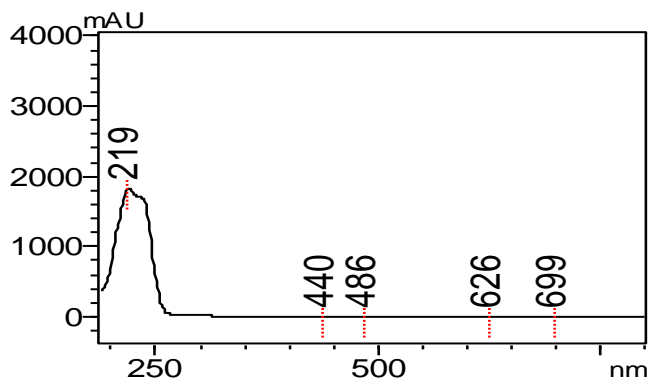


Fig. S2 UV spectrum of compound 1

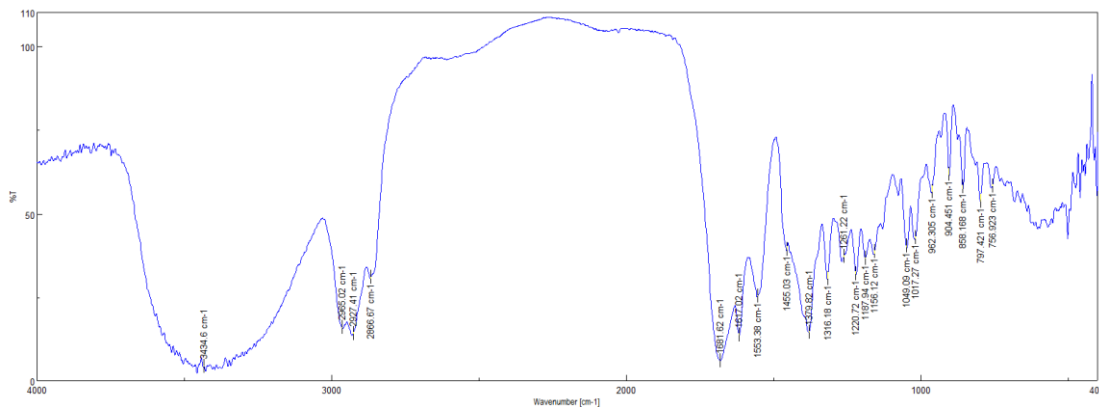


Fig. S3 IR spectrum of compound 1

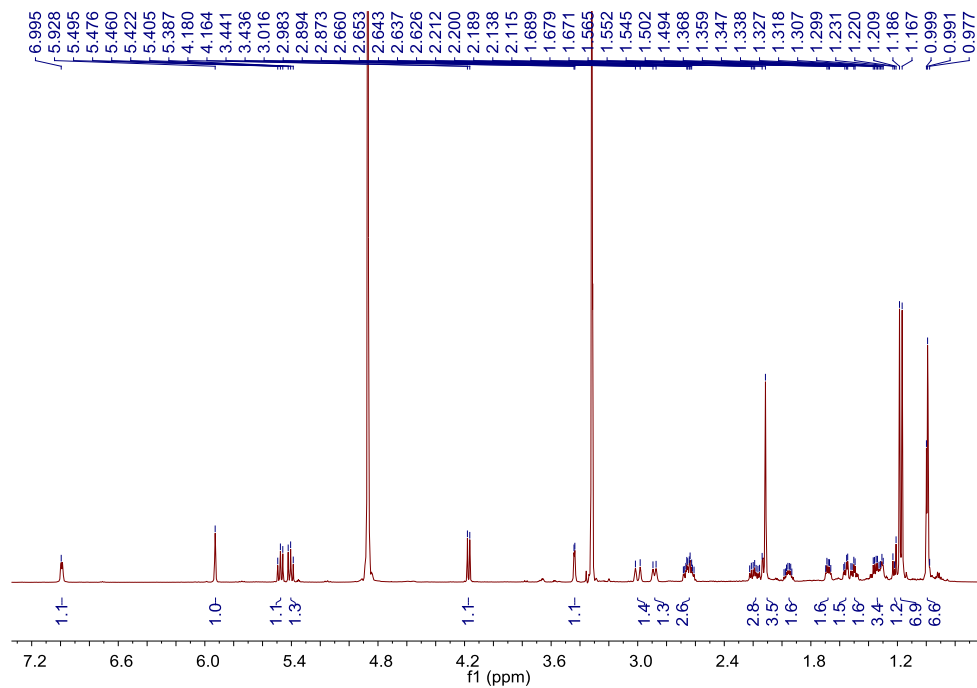


Fig. S4 ^1H -NMR spectrum of compound **1** (in CD_3OD)

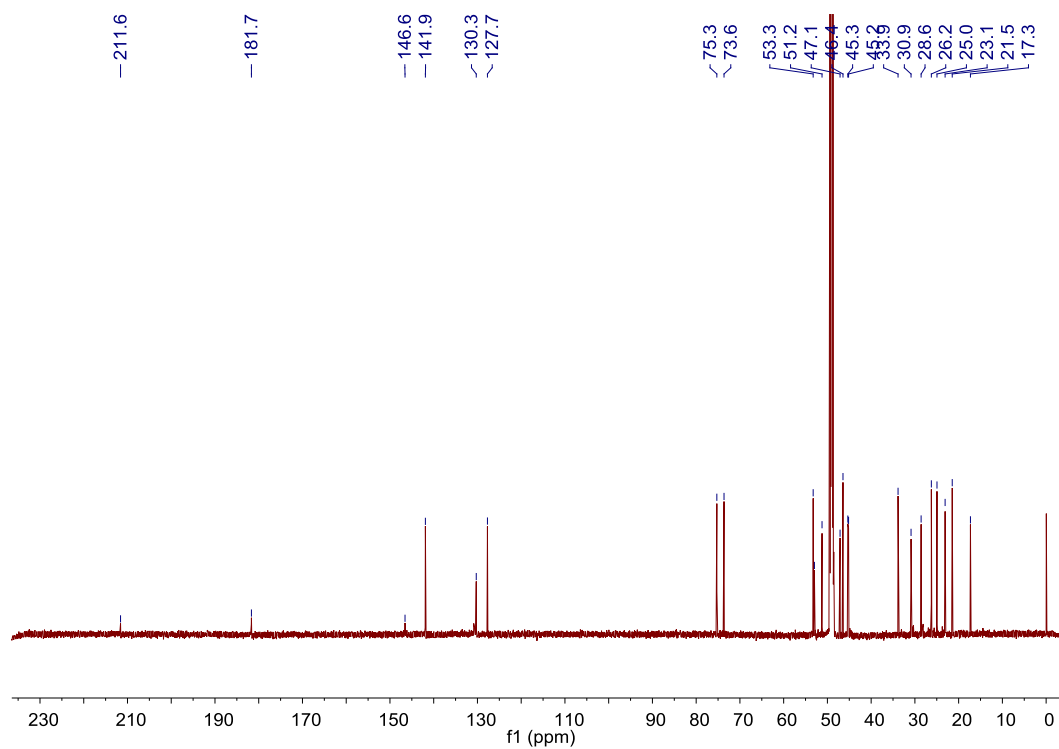


Fig. S5 ^{13}C -NMR spectrum of compound **1** (in CD_3OD)

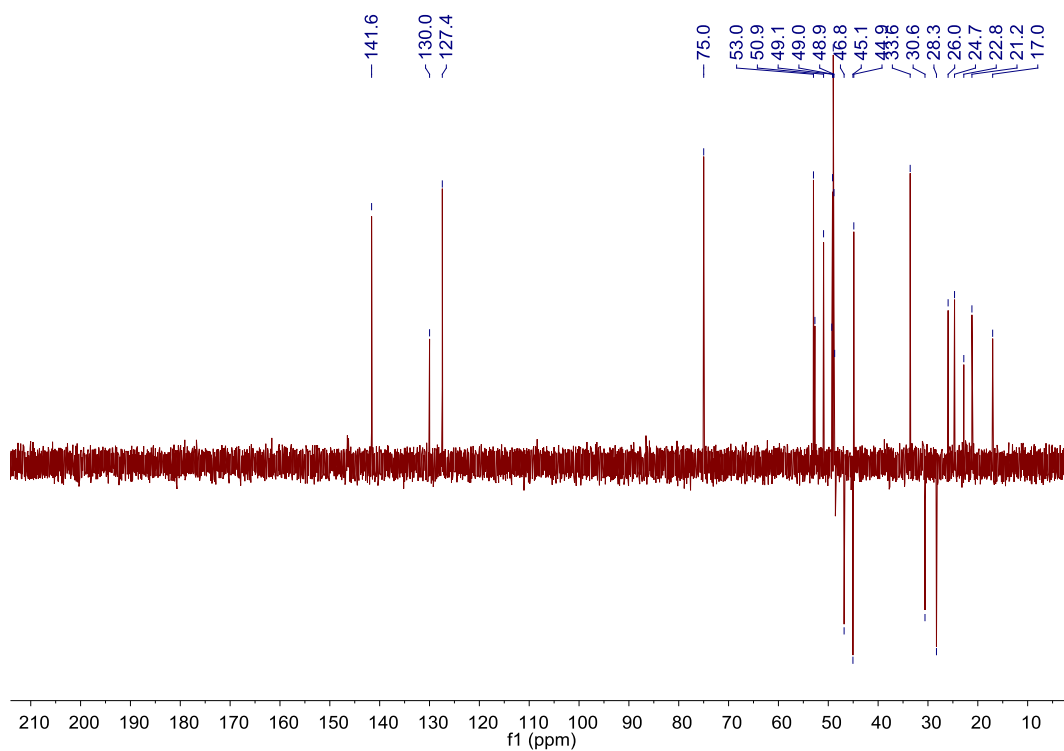


Fig. S6 DEPT135 spectrum of compound **1** (in CD₃OD)

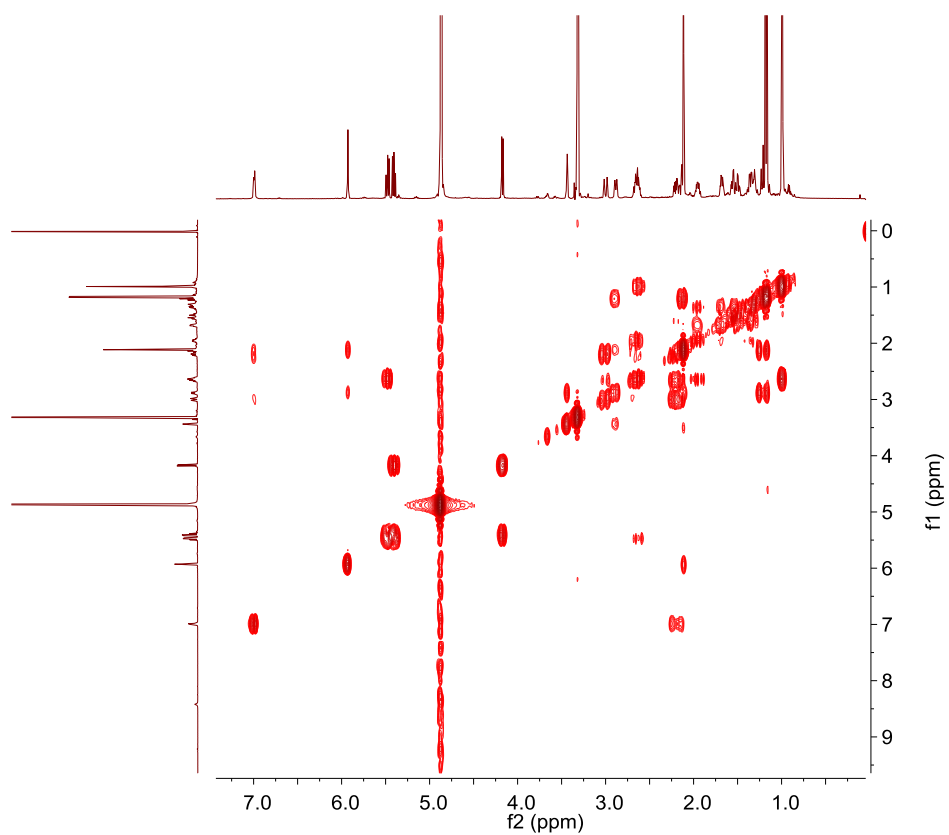


Fig. S7 ¹H-¹H COSY spectrum of compound **1** (in CD₃OD)

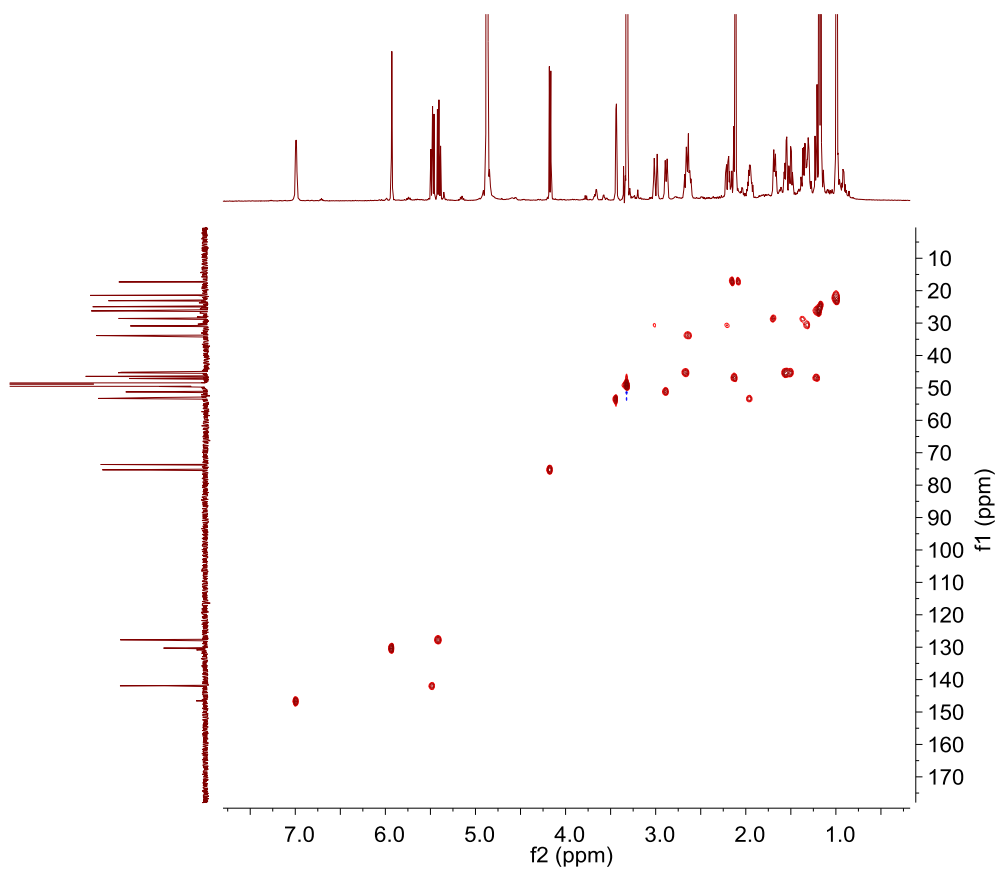


Fig. S8 HSQC spectrum of compound **1** (in CD₃OD)

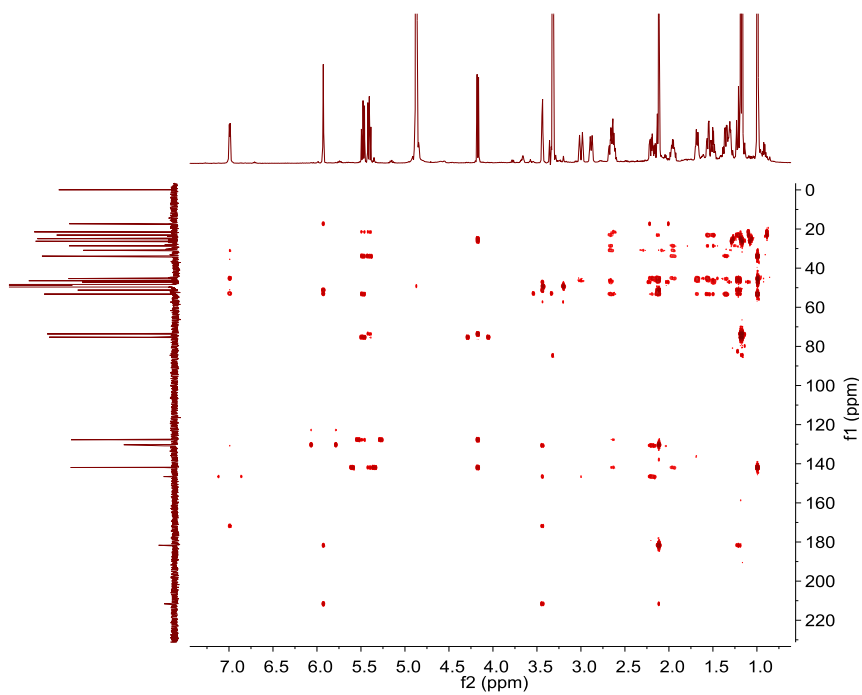


Fig. S9 HMBC spectrum of compound **1** (in CD₃OD).

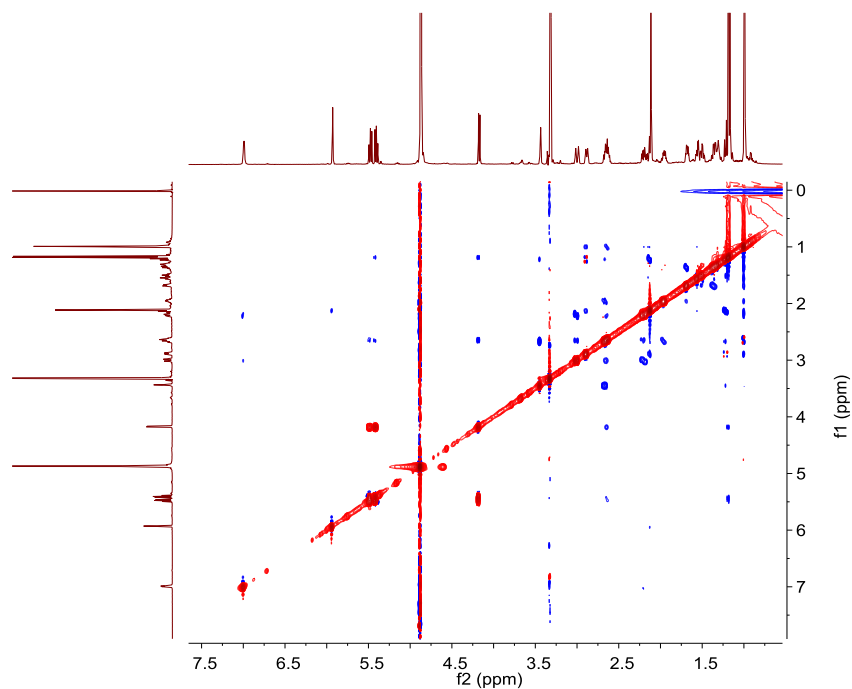


Fig. S10 ROESY spectrum of compound **1** (in CD3OD).

Elemental Composition Report

Page 1

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

728 formula(e) evaluated with 1 results within limits (up to 50 closest results for each mass)

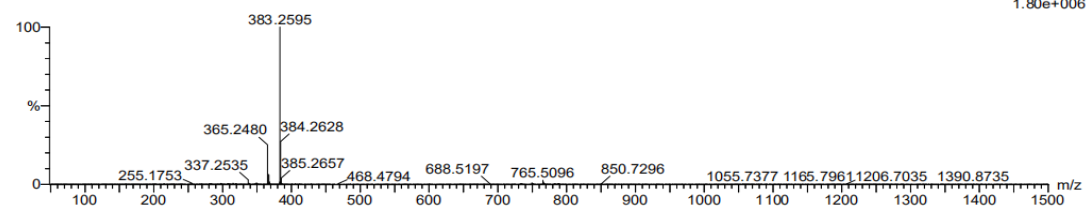
Elements Used:

C: 0-50 H: 0-100 N: 0-10 O: 0-100

0154-6-9-2-7

20191202054 291 (2.345)

1: TOF MS ES+
1.80e+006



Minimum:

Maximum: 5.0 5.0 -1.5

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
383.2595	383.2586	0.9	2.3	8.5	752.1	n/a	n/a	C25 H35 O3

Fig. S11 HR ESI-Q-TOF-MS spectrum of compound **2**

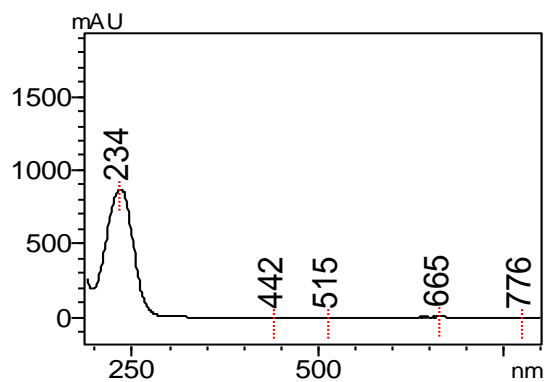


Fig. S12 UV spectrum of compound 2

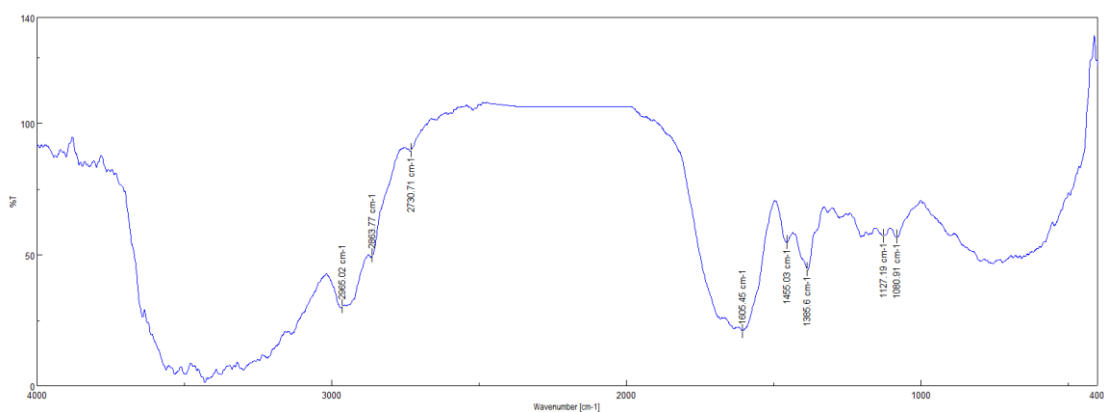


Fig. S13 IR spectrum of compound 2

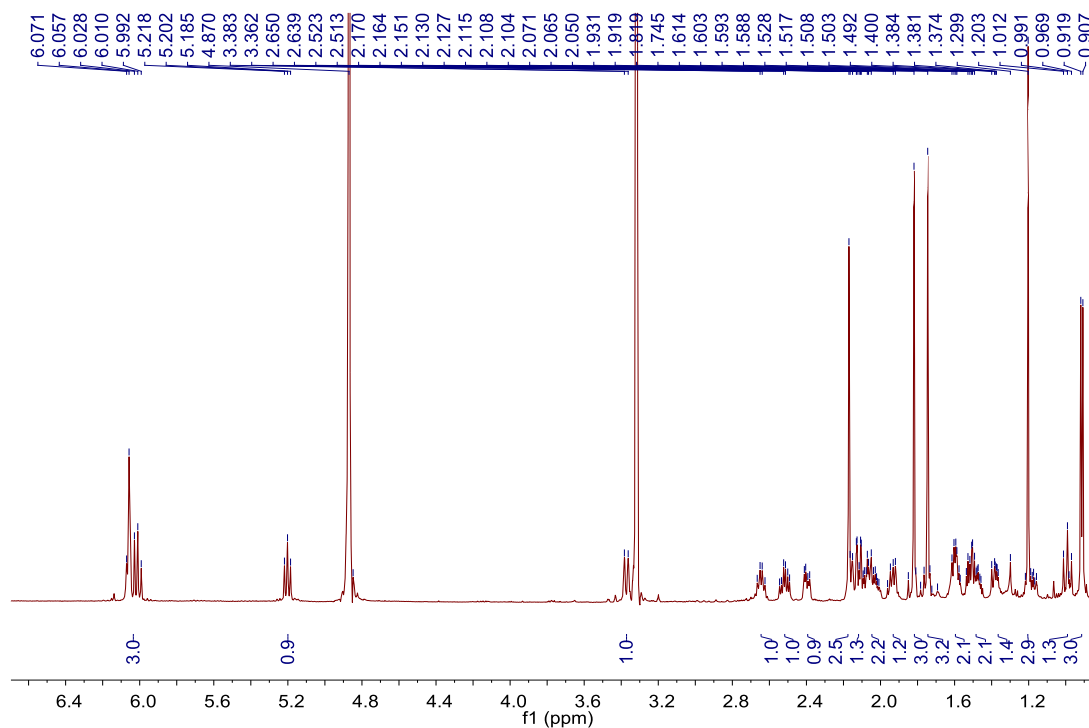


Fig. S14 ¹H-NMR spectrum of compound 2 (in CD₃OD)

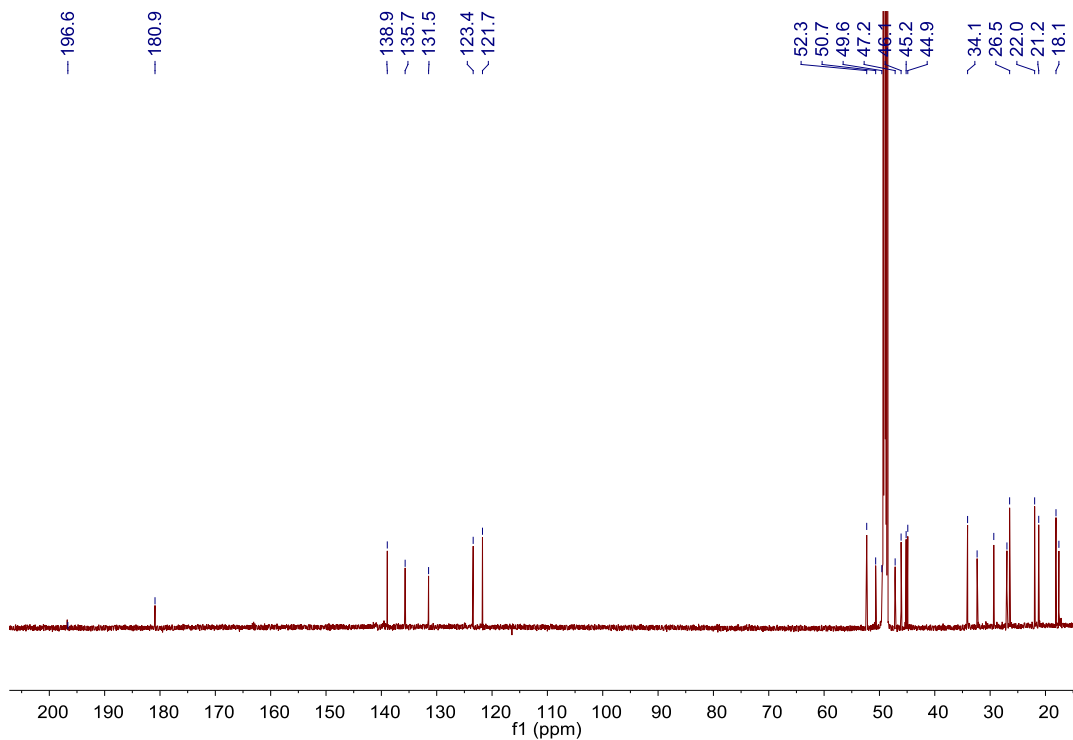


Fig. S15 ^{13}C -NMR spectrum of compound **2** (in CD_3OD)

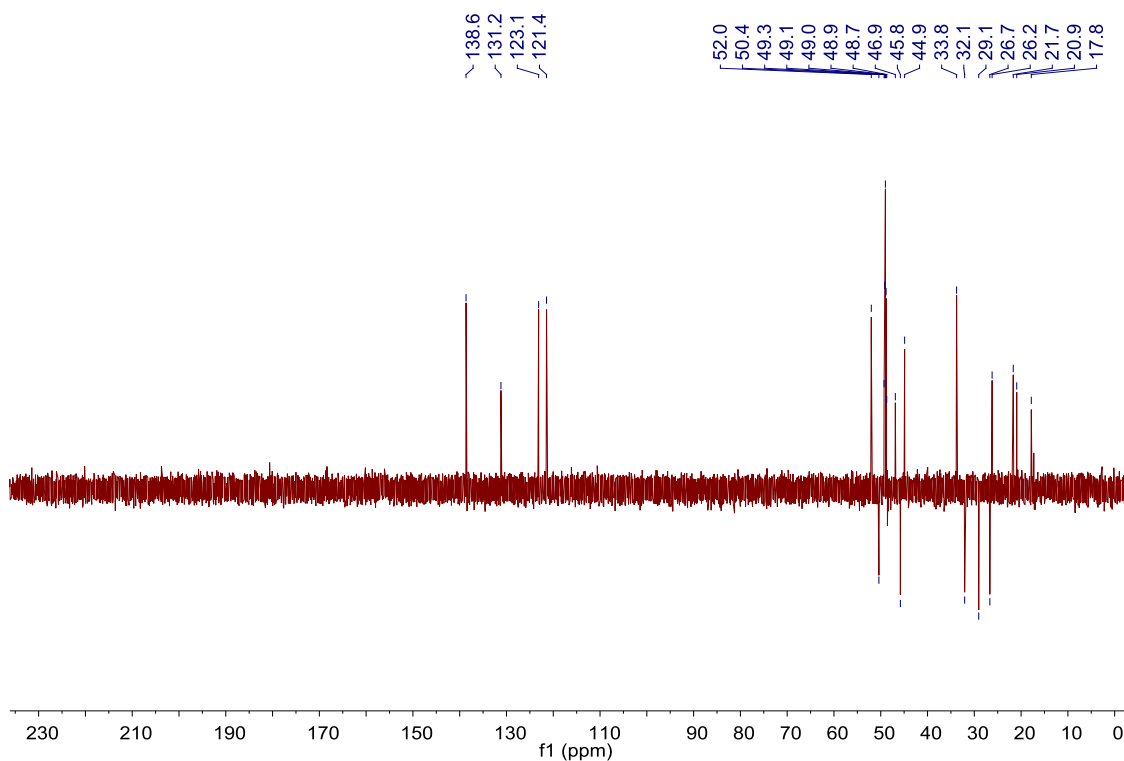


Fig. S16 DEPT135 spectrum of compound **2** (in CD_3OD)

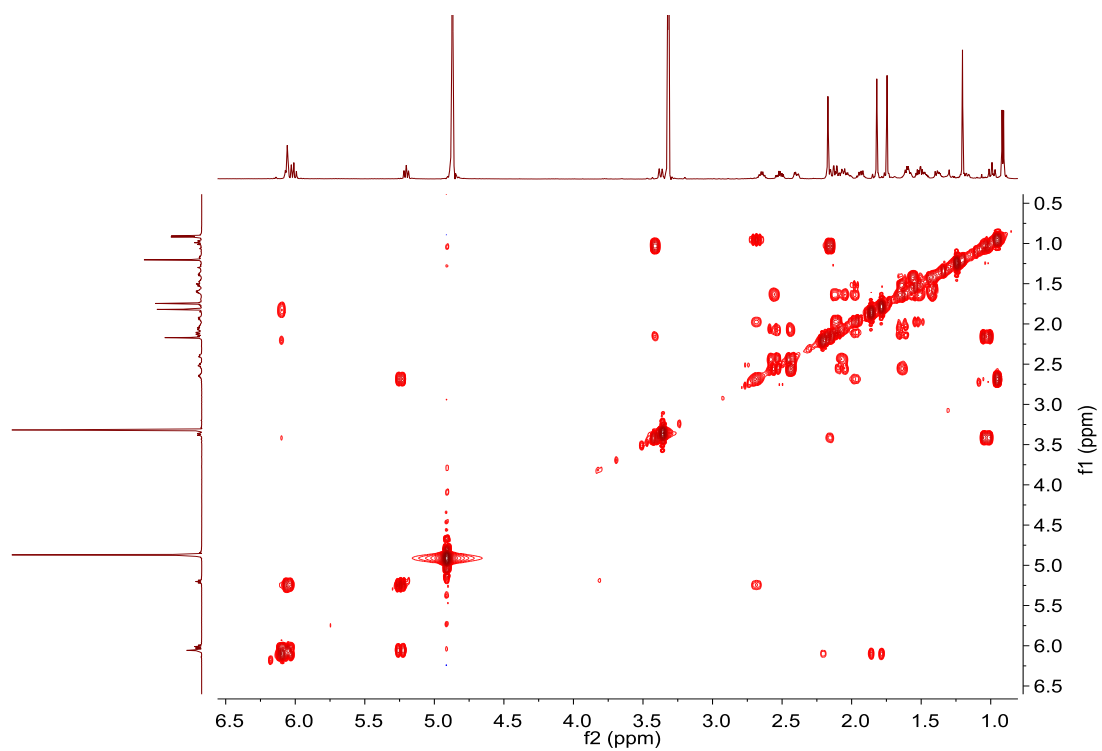


Fig. S17 ^1H - ^1H COSY spectrum of compound **2** (in CD_3OD)

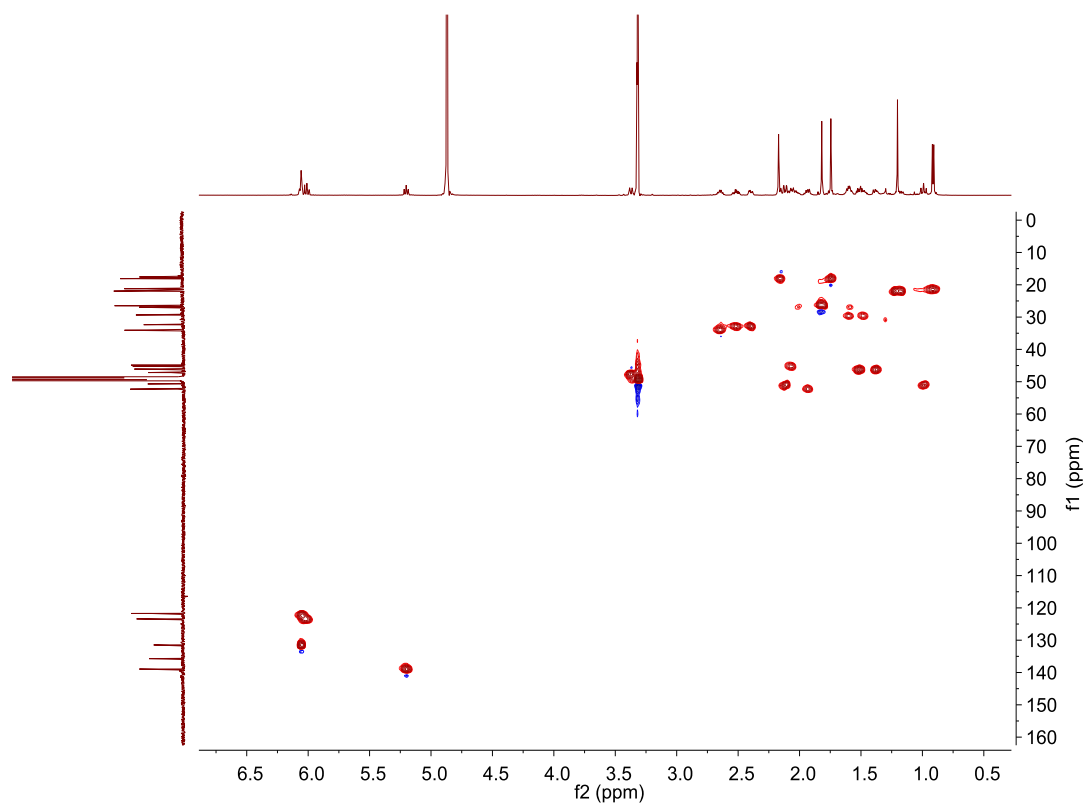


Fig. S18 HSQC spectrum of compound **2** (in CD_3OD)

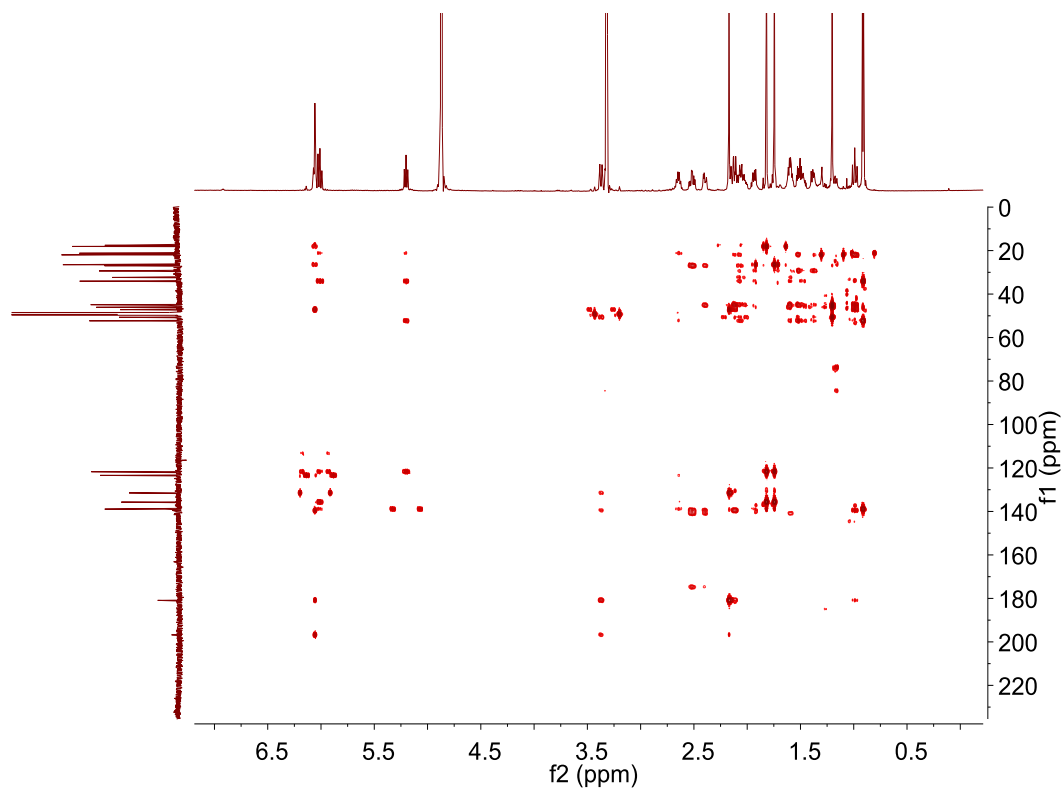


Fig. S19 HMBC spectrum of compound **2** (in CD₃OD)

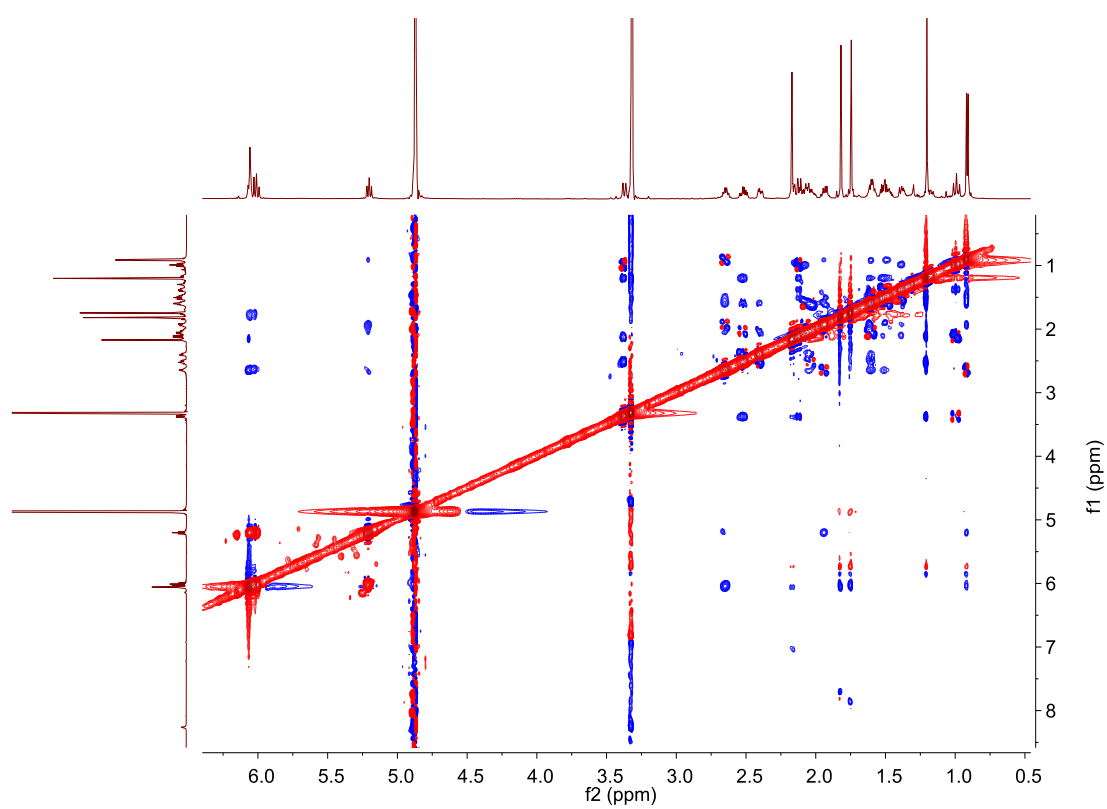


Fig. S20 ROESY spectrum of compound **2** (in CD₃OD).

Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

89 formula(e) evaluated with 1 results within limits (up to 50 best isotopic matches for each mass)

Elements Used:

C: 0-500 H: 0-1000 O: 0-200

0154-78-8-2

20191104042 255 (2.053)

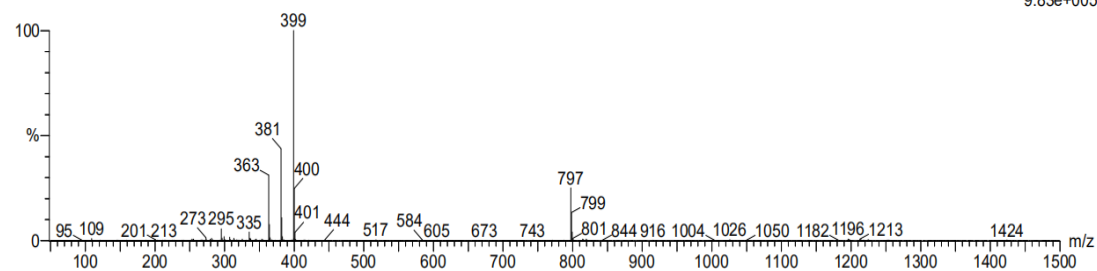
1: TOF MS ES+
9.83e+005

Fig. S21 HR ESI-Q-TOF-MS spectrum of compound 3

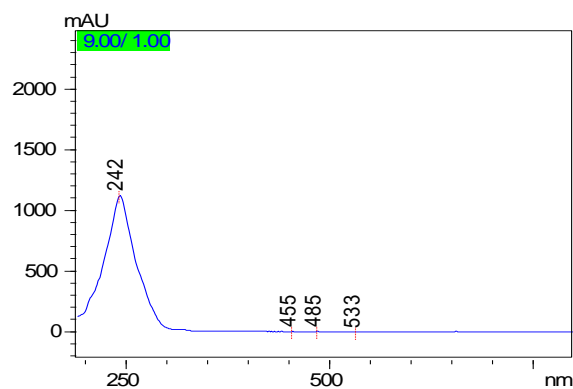


Fig. S22 UV spectrum of compound 3

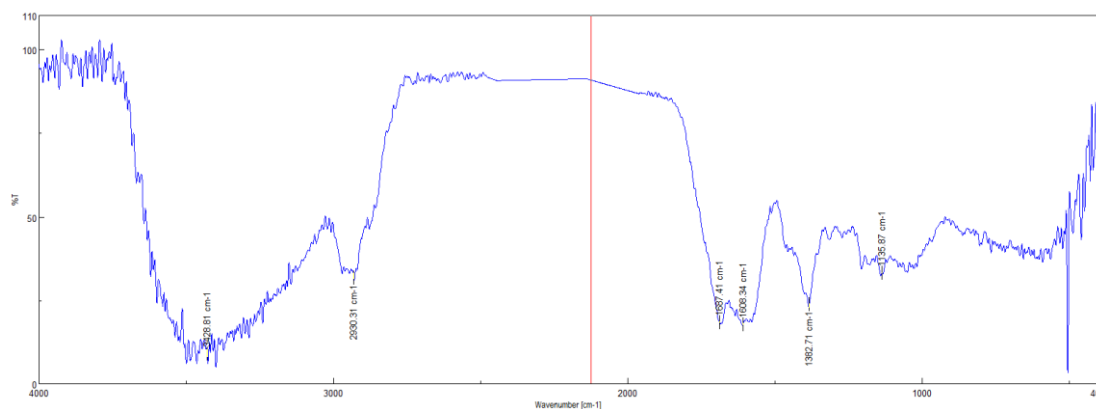


Fig. S23 IR spectrum of compound 3

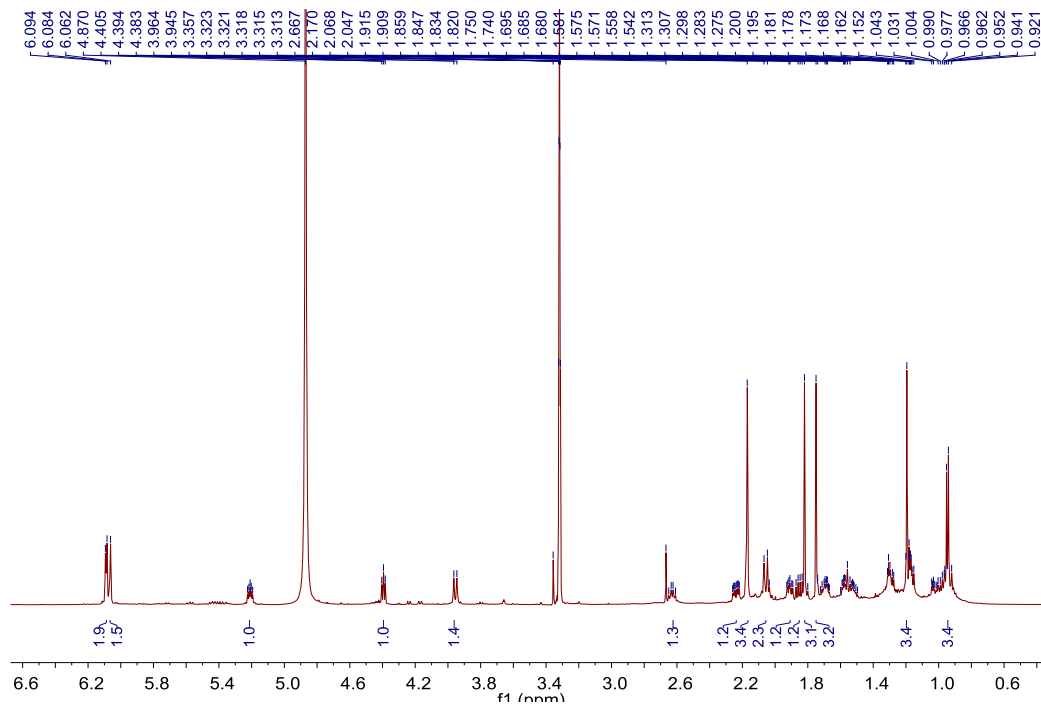


Fig. S24 ^1H -NMR spectrum of compound **3** (in CD_3OD)

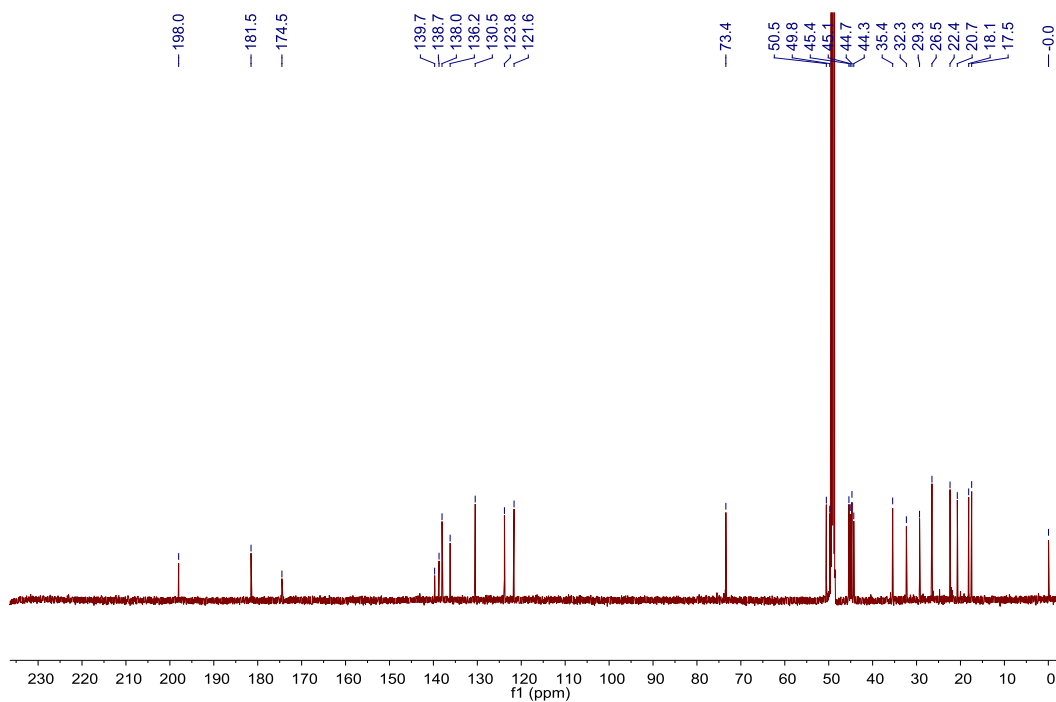


Fig. S25 ^{13}C -NMR spectrum of compound **3** (in CD_3OD)

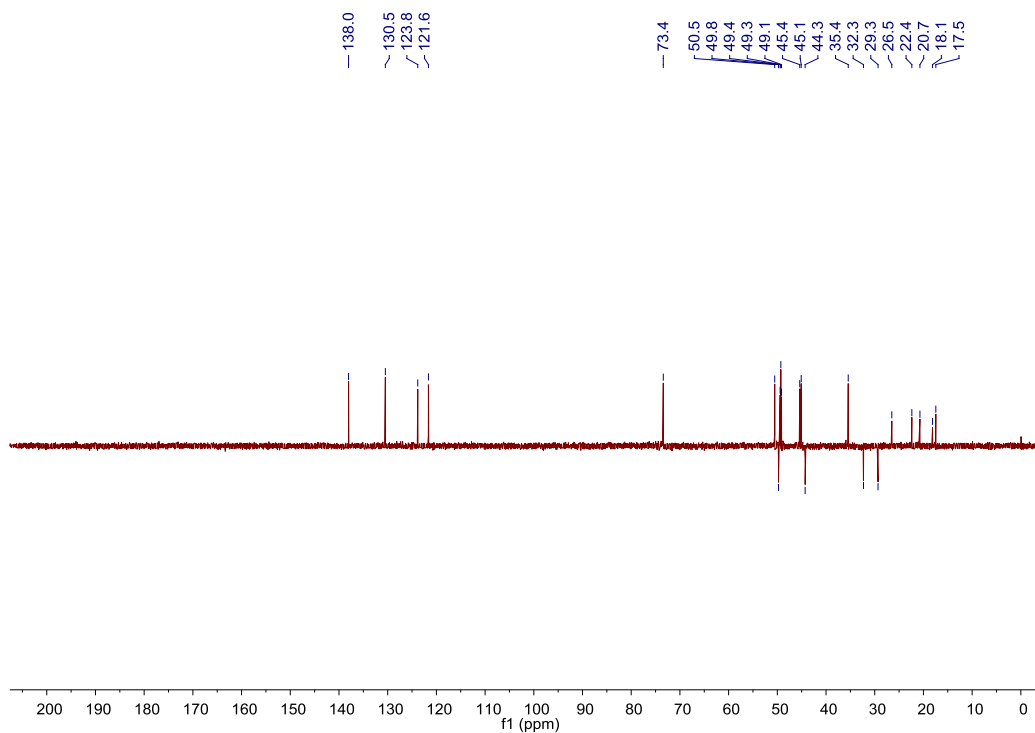


Fig. S26 DEPT135 spectrum of compound **3** (in CD₃OD)

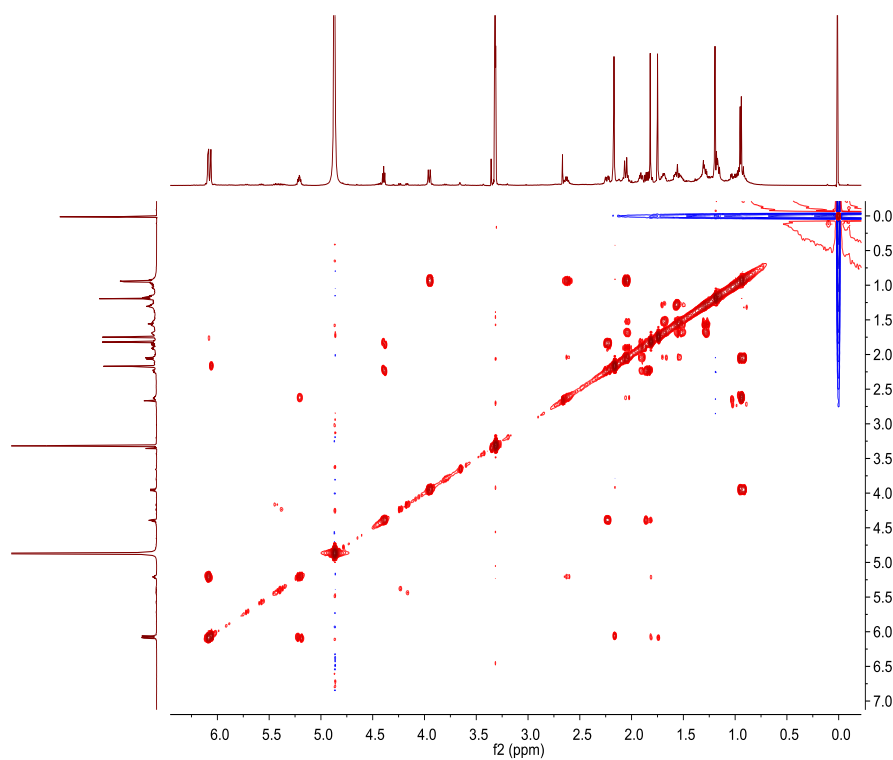


Fig. S27 ¹H-¹H COSY spectrum of compound **3** (in CD₃OD)

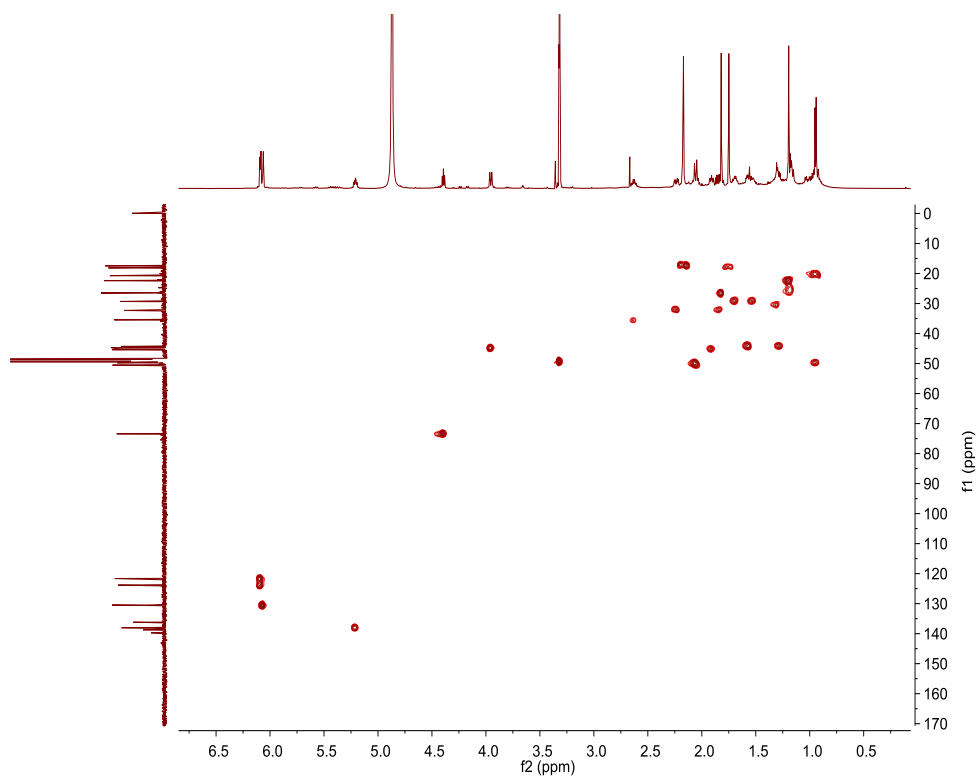


Fig. S28 HSQC spectrum of compound **3** (in CD₃OD)

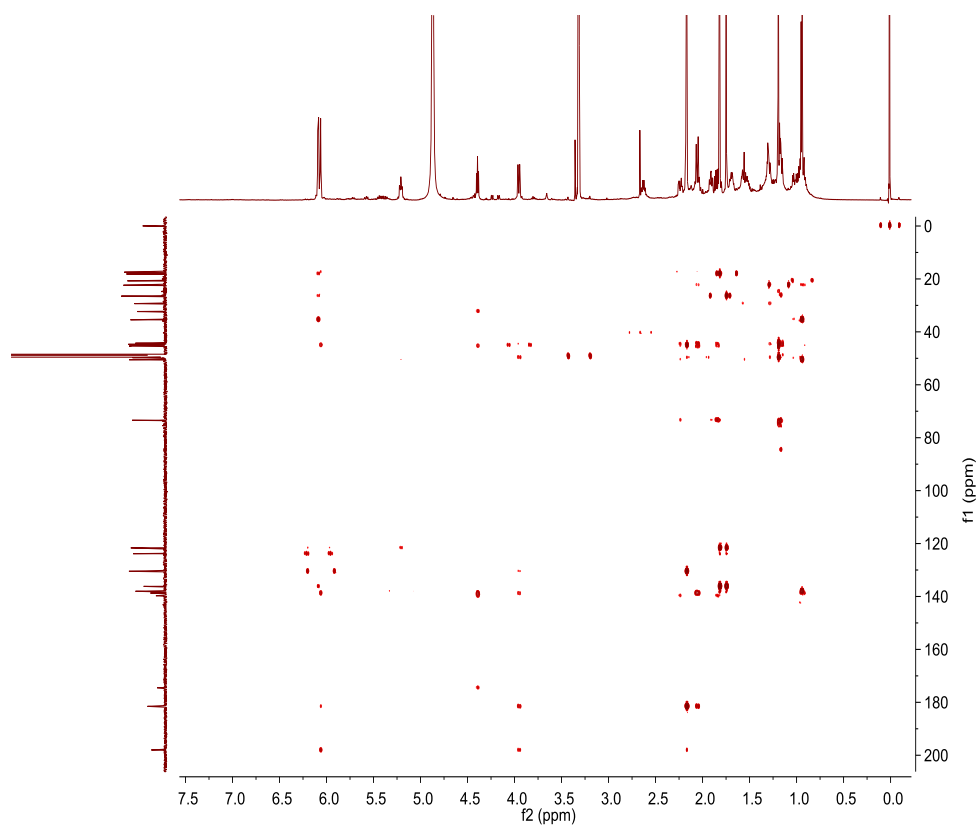


Fig. S29 HMBC spectrum of compound **3** (in CD₃OD).

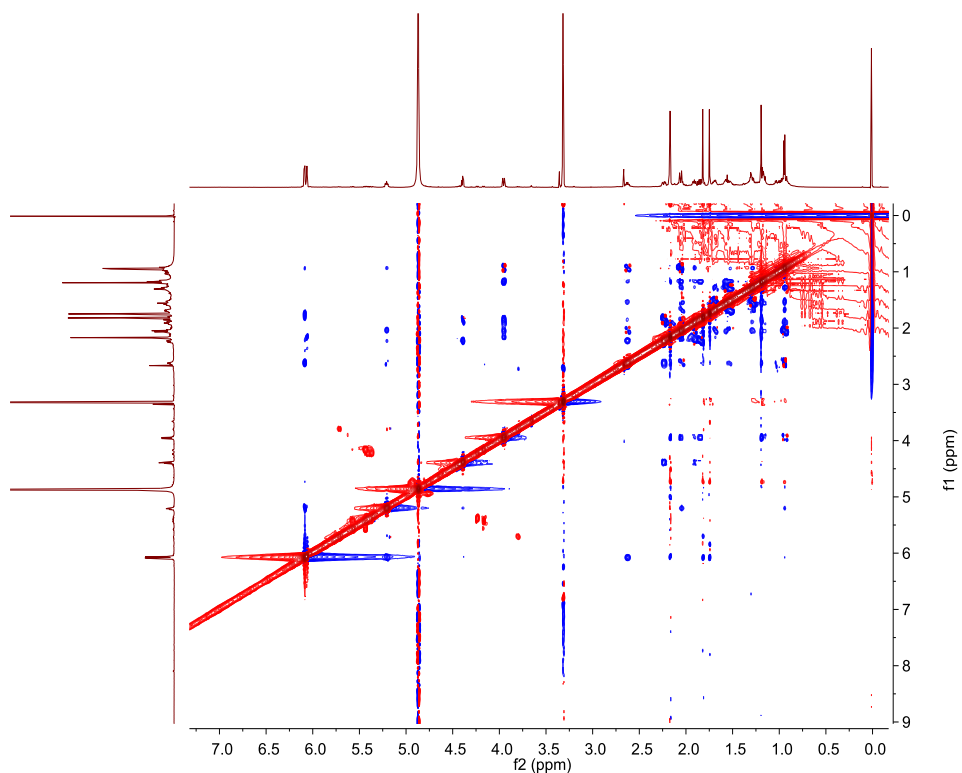


Fig. S30 ROESY spectrum of compound **3** (in CD₃OD).

Elemental Composition Report

Page 1

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

2369 formula(e) evaluated with 7 results within limits (up to 50 best isotopic matches for each mass)

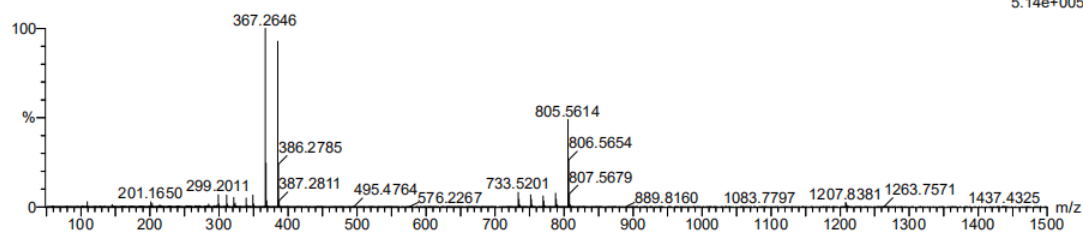
Elements Used:

C: 0-50 H: 0-100 N: 0-10 O: 0-100

0154-78-8-4

20191104041 255 (2.053)

1: TOF MS ES+
5.14e+005



Minimum: 5.0 5.0 -1.5
Maximum: 5.0 5.0 50.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
805.5614	805.5605	0.9	1.1	18.5	417.0	1.874	15.35	C47 H69 N10 O2
	805.5592	2.2	2.7	13.5	415.3	0.199	81.97	C46 H73 N6 O6
	805.5618	-0.4	-0.5	12.5	418.9	3.819	2.19	C50 H77 O8
	805.5578	3.6	4.5	8.5	420.5	5.399	0.45	C45 H77 N2 O10
	805.5650	-3.6	-4.5	4.5	424.2	9.084	0.01	C39 H77 N6 O11
	805.5610	0.4	0.5	0.5	424.7	9.620	0.01	C34 H77 N8 O13
	805.5637	-2.3	-2.9	-0.5	423.8	8.753	0.02	C38 H81 N2 O15

Fig. S31 HR ESI-Q-TOF-MS spectrum of compound **4**

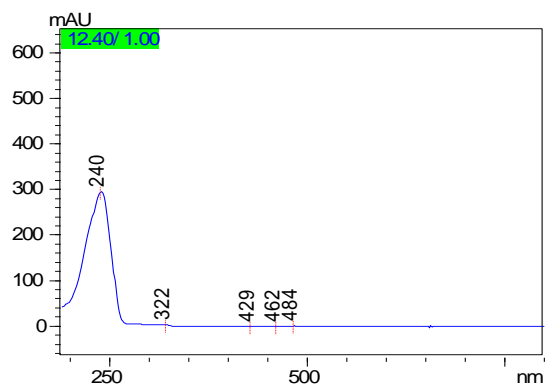


Fig. S32 UV spectrum of compound **4**

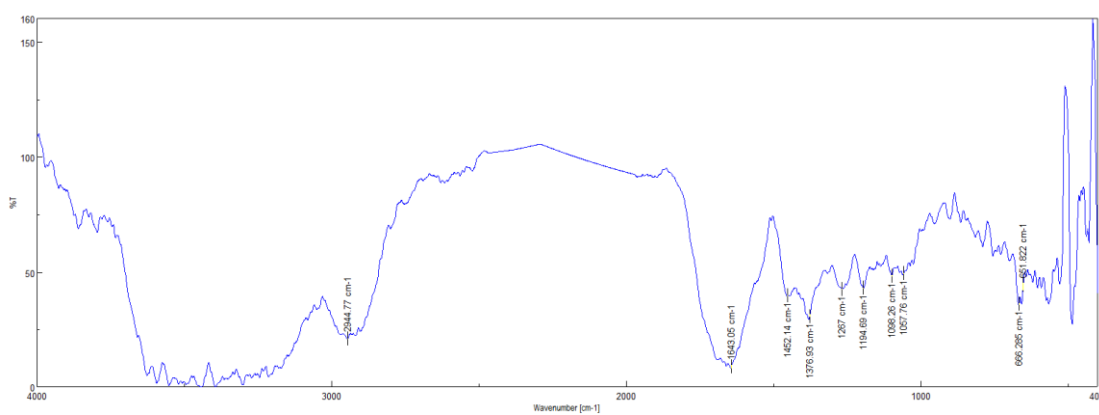


Fig. S33 IR spectrum of compound **4**

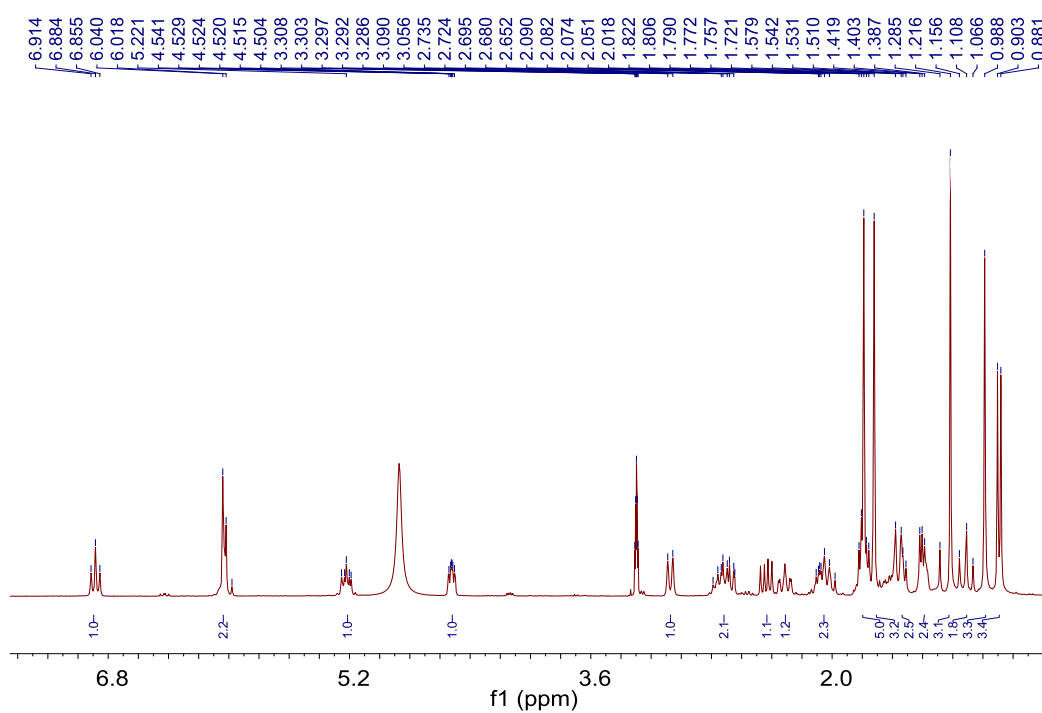


Fig. S34 $^1\text{H-NMR}$ spectrum of compound **4** (in CD_3OD)

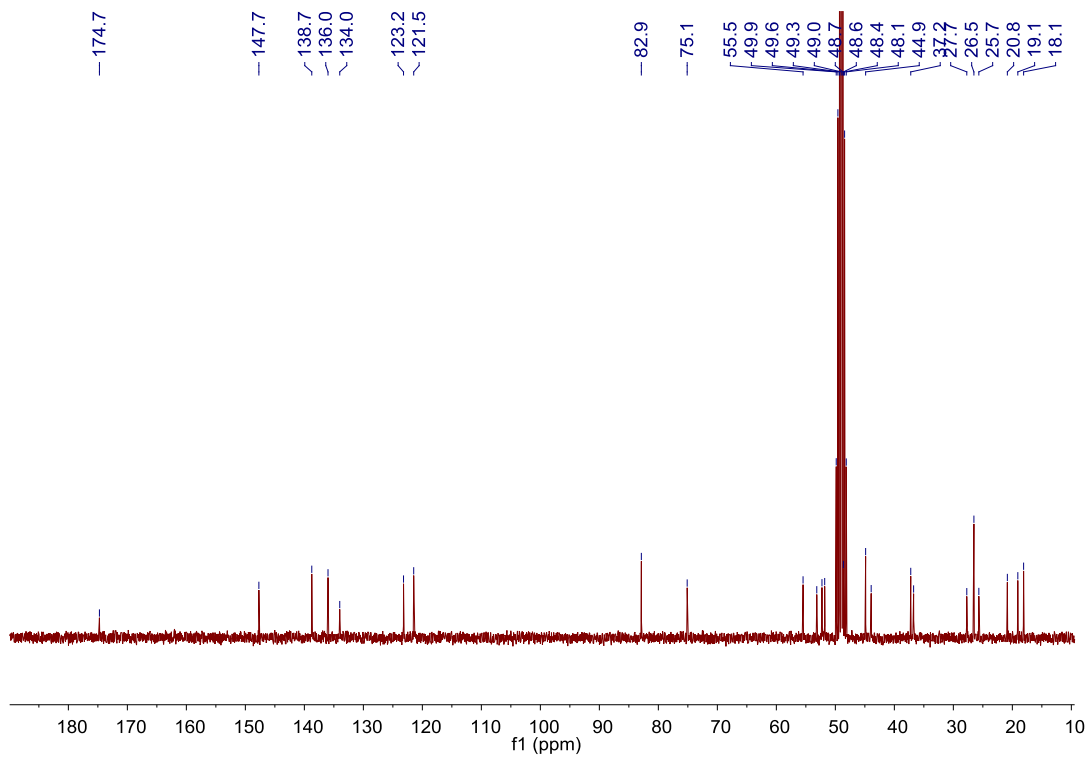


Fig. S35 ^{13}C -NMR spectrum of compound **4** (in CD_3OD)

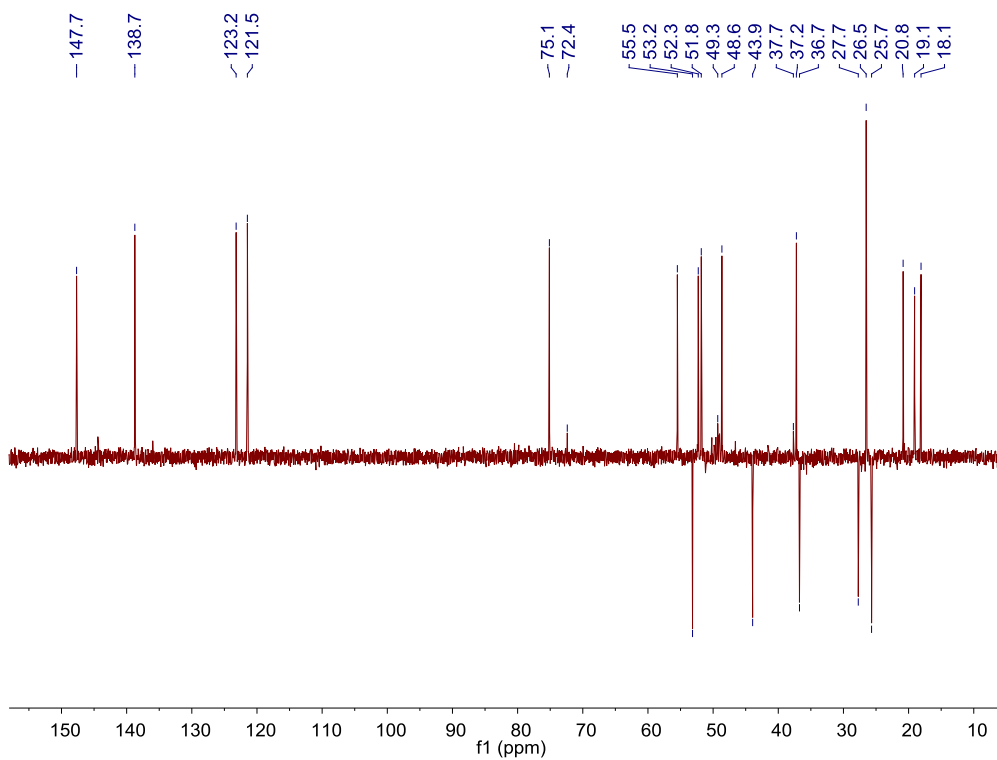


Fig. S36 DEPT135 spectrum of compound **4** (in CD_3OD)

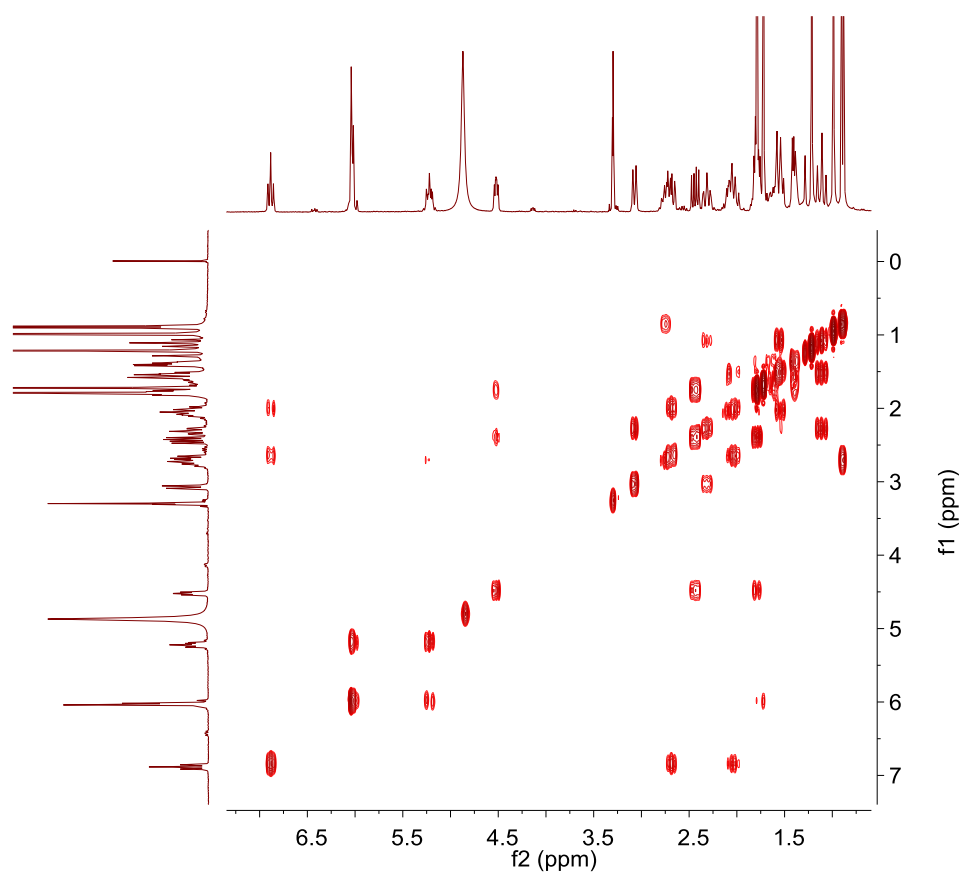


Fig. S37 ^1H - ^1H COSY spectrum of compound **4** (in CD_3OD)

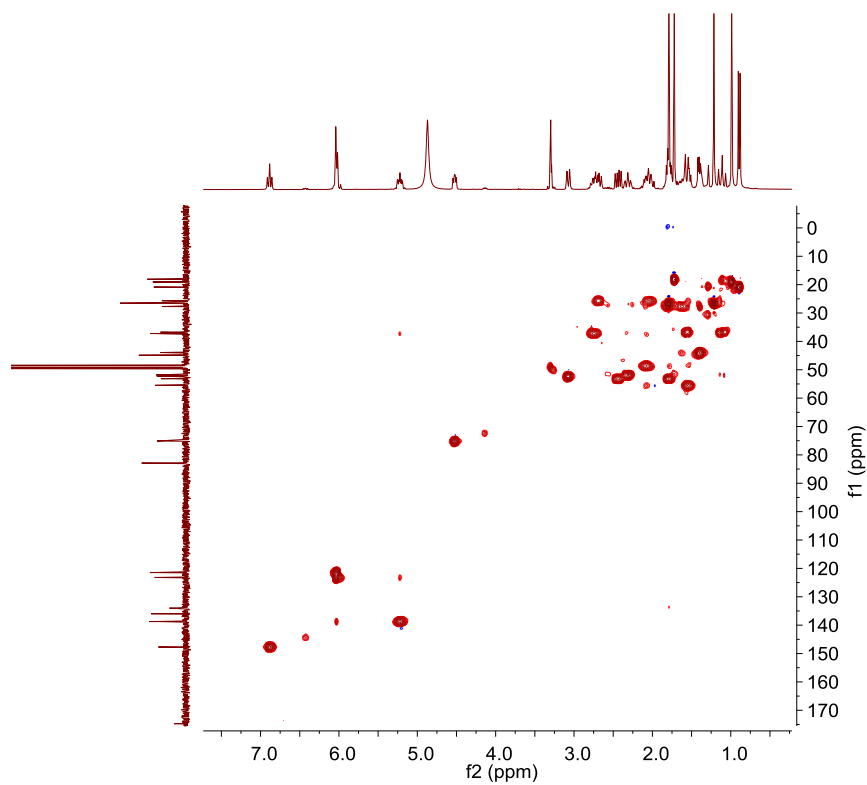


Fig. S38 HSQC spectrum of compound **4** (in CD_3OD)

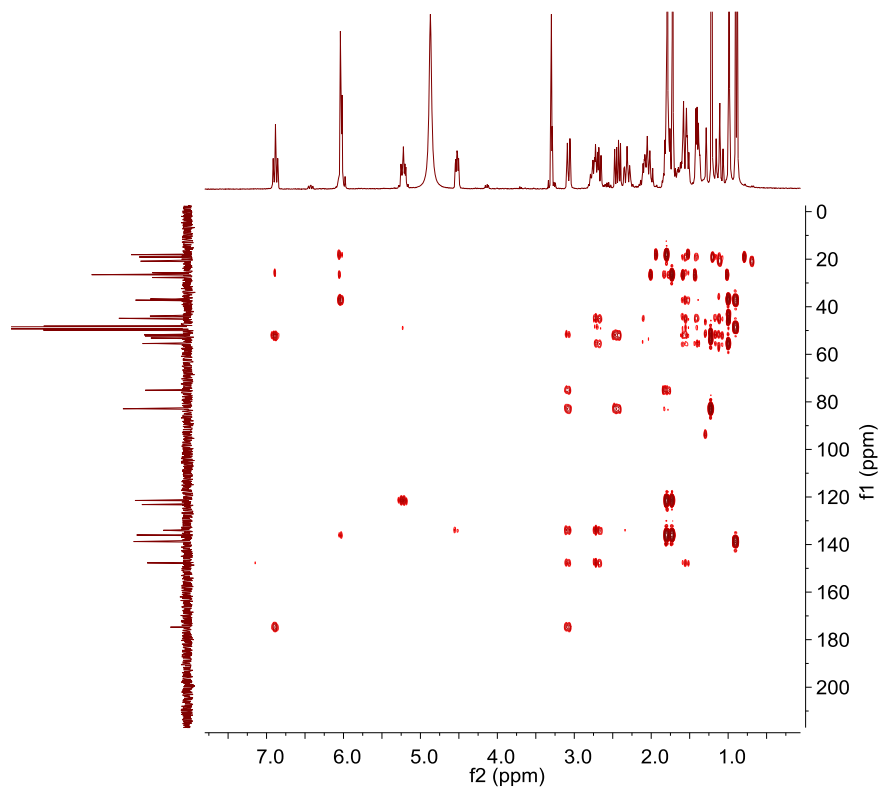


Fig. S39 HMBC spectrum of compound **4** (in CD₃OD).

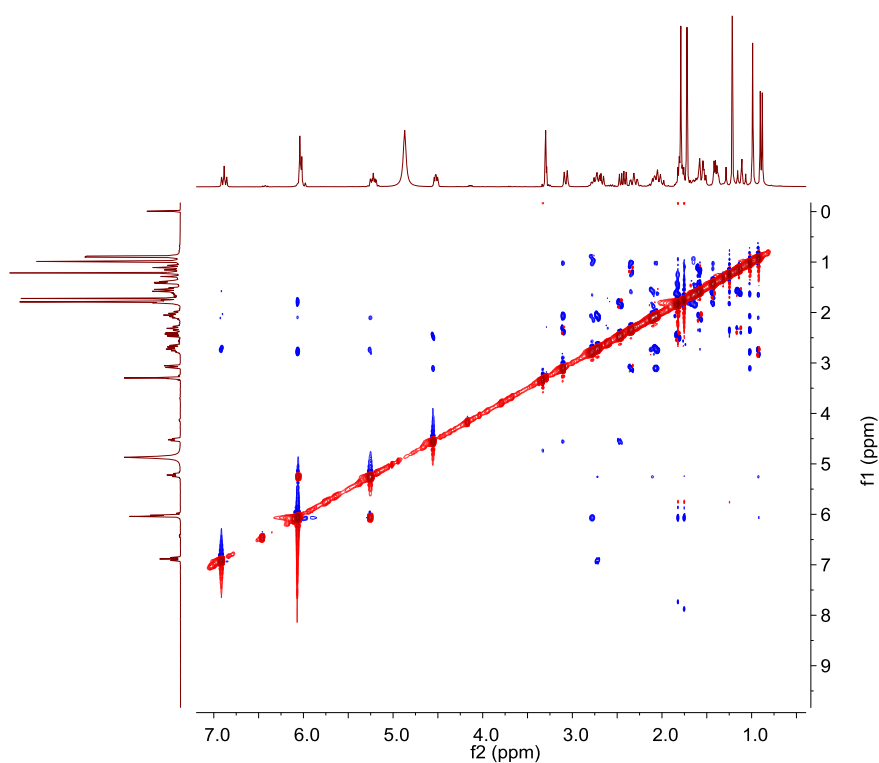


Fig. S40 ROESY spectrum of compound **4** (in CD₃OD).

Single Mass Analysis

Tolerance = 5.0 PPM / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

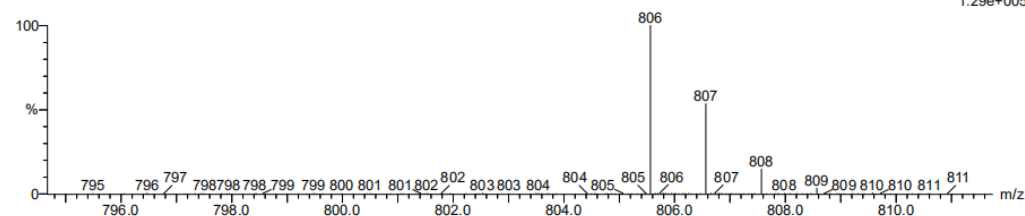
4513 formula(e) evaluated with 11 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-50 H: 0-200 N: 0-20 O: 0-100

0154-78-8-5

20191104039 256 (2.061)

1: TOF MS ES+
1.29e+005

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf (%)	Formula
805.5621	805.5623	-0.2	-0.2	5.5	343.2	7.886	0.04	C35 H73 N12 O9
	805.5618	0.3	0.4	12.5	340.1	4.842	0.79	C50 H77 O8
	805.5610	1.1	1.4	0.5	344.0	8.765	0.02	C34 H77 N8 O13
	805.5637	-1.6	-2.0	10.5	343.0	7.749	0.04	C36 H69 N16 O5
	805.5637	-1.6	-2.0	-0.5	344.4	9.107	0.01	C38 H81 N2 O15
	805.5605	1.6	2.0	18.5	337.2	1.943	14.33	C47 H69 N10 O2
	805.5597	2.4	3.0	6.5	345.9	10.624	0.00	C31 H69 N18 O7
	805.5650	-2.9	-3.6	15.5	343.4	8.100	0.03	C37 H65 N20 O
	805.5650	-2.9	-3.6	4.5	344.5	9.200	0.01	C39 H77 N6 O11

Fig. S41 HR ESI-Q-TOF-MS spectrum of compound 5

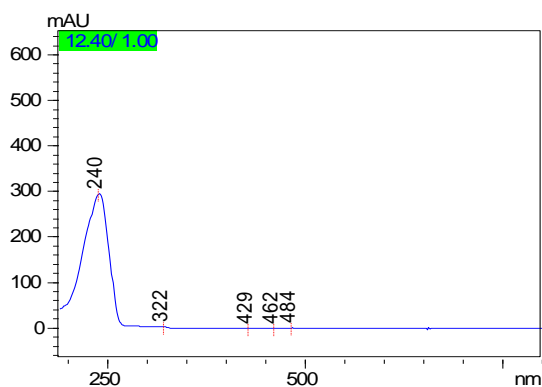


Fig. S42 UV spectrum of compound 5

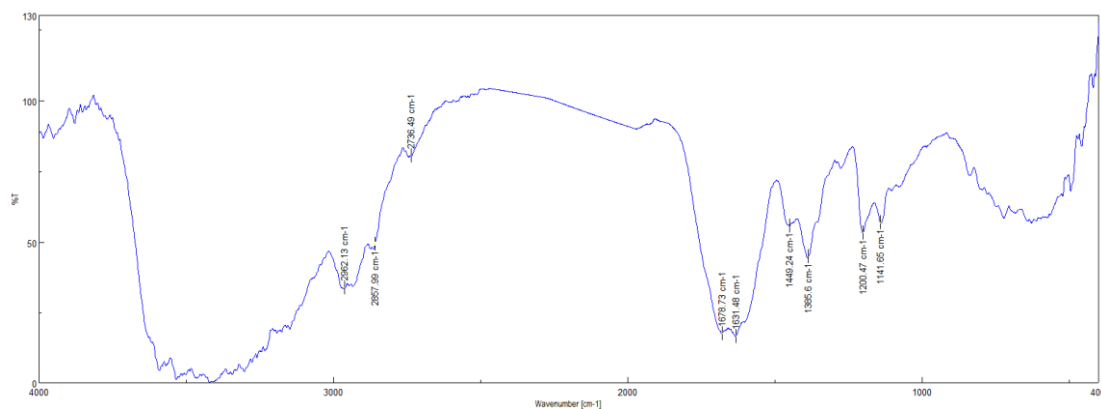


Fig. S43 IR spectrum of compound 5

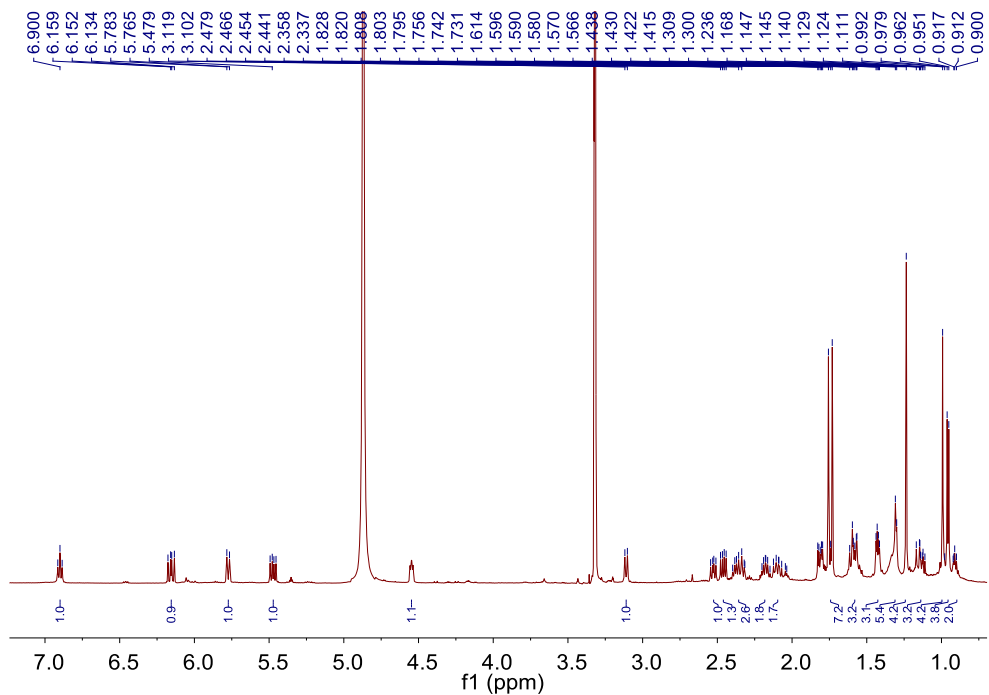


Fig. S44 ^1H -NMR spectrum of compound **5** (in CD_3OD)

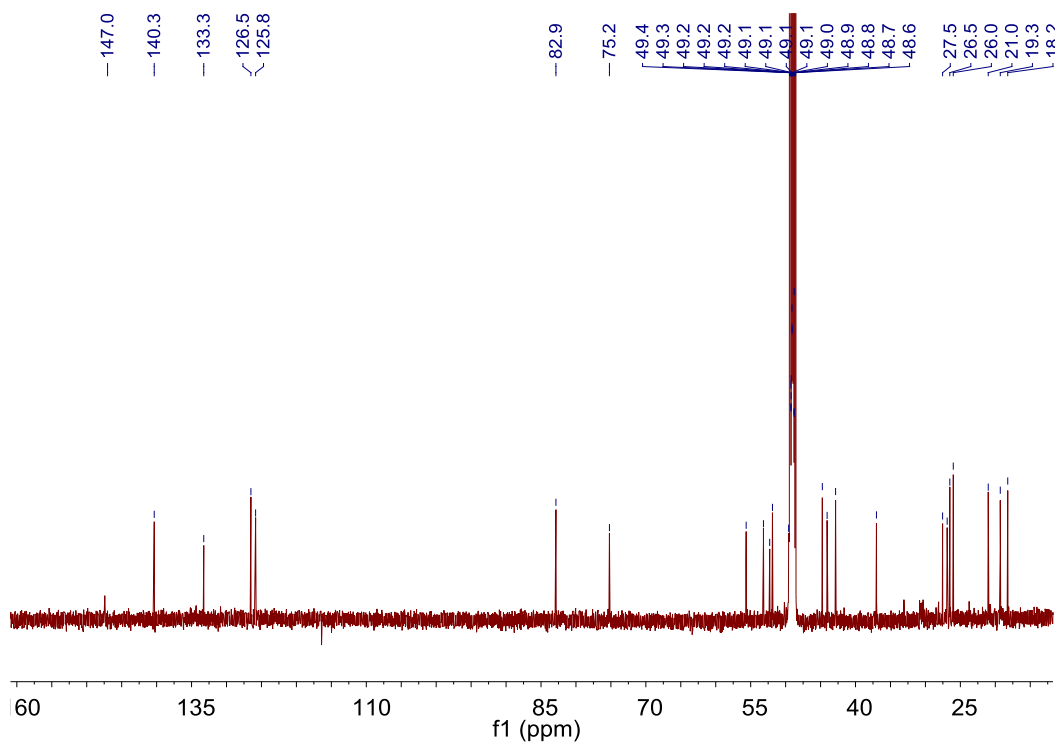


Fig. S45 ^{13}C -NMR spectrum of compound **5** (in CD_3OD)

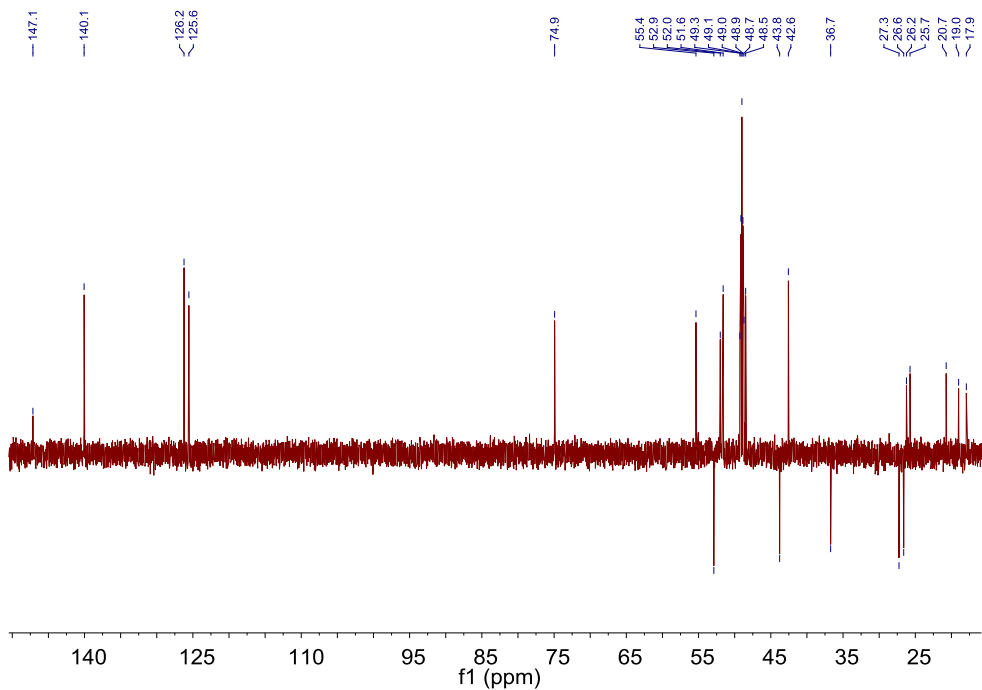


Fig. S46 DEPT135 spectrum of compound **5** (in CD₃OD)

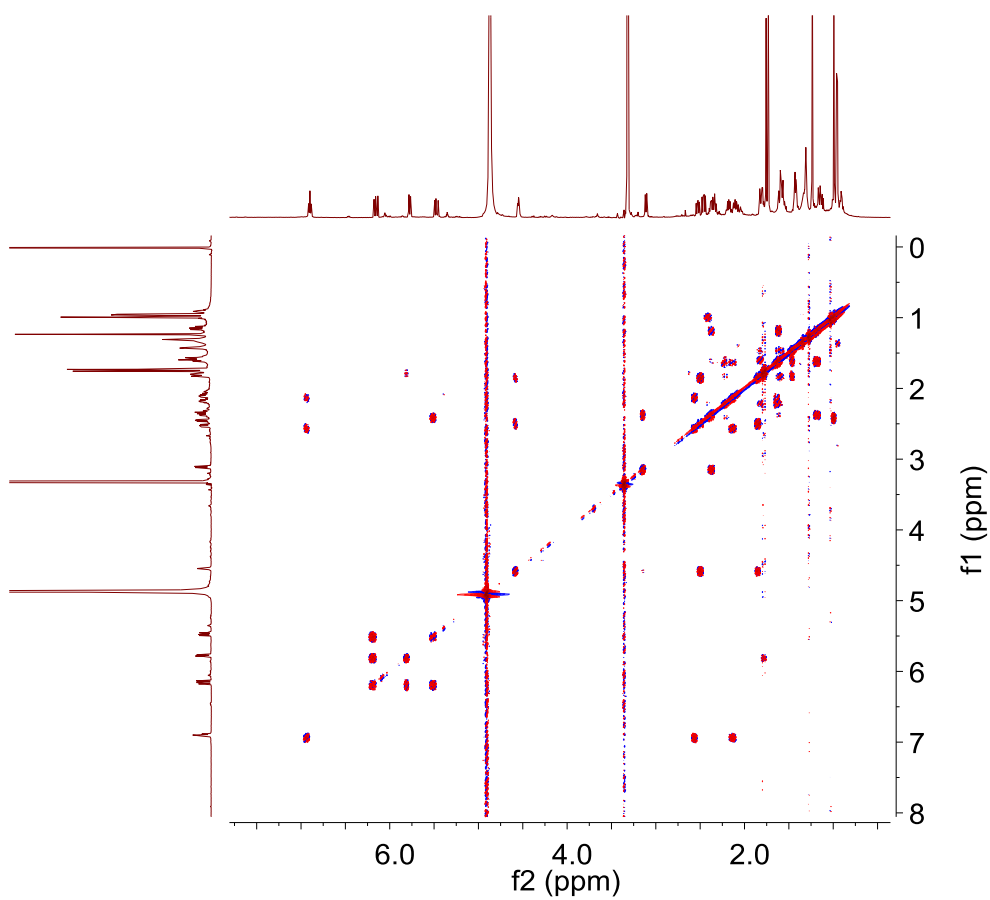


Fig. S47 ¹H-¹H COSY spectrum of compound **5** (in CD₃OD)

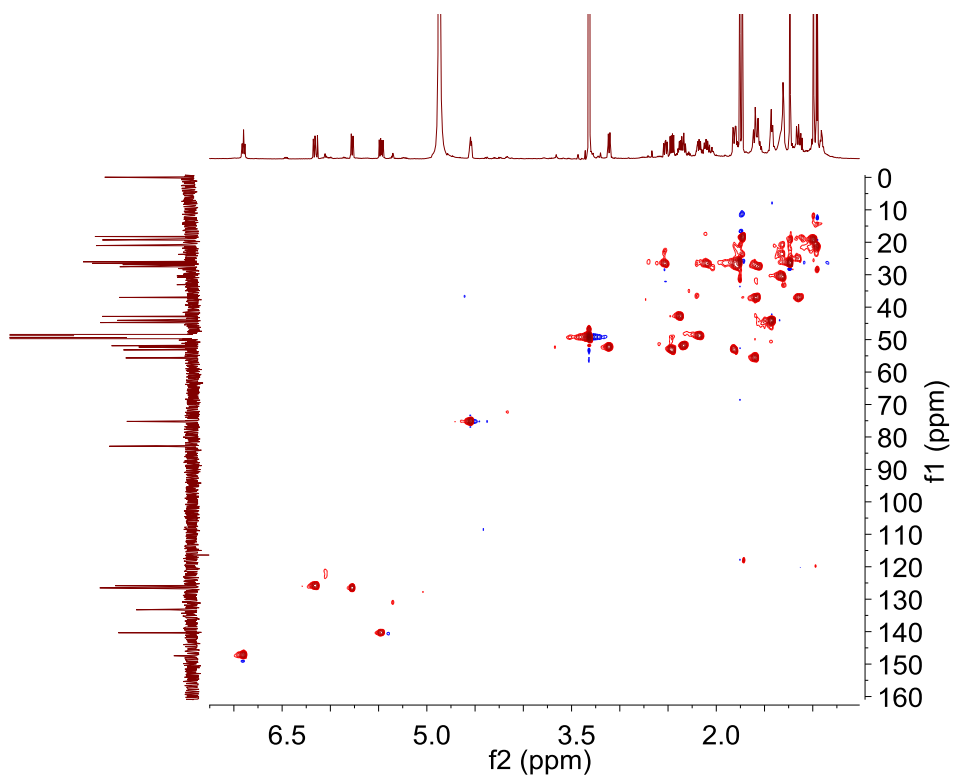


Fig. S48 HSQC spectrum of compound **5** (in CD₃OD).

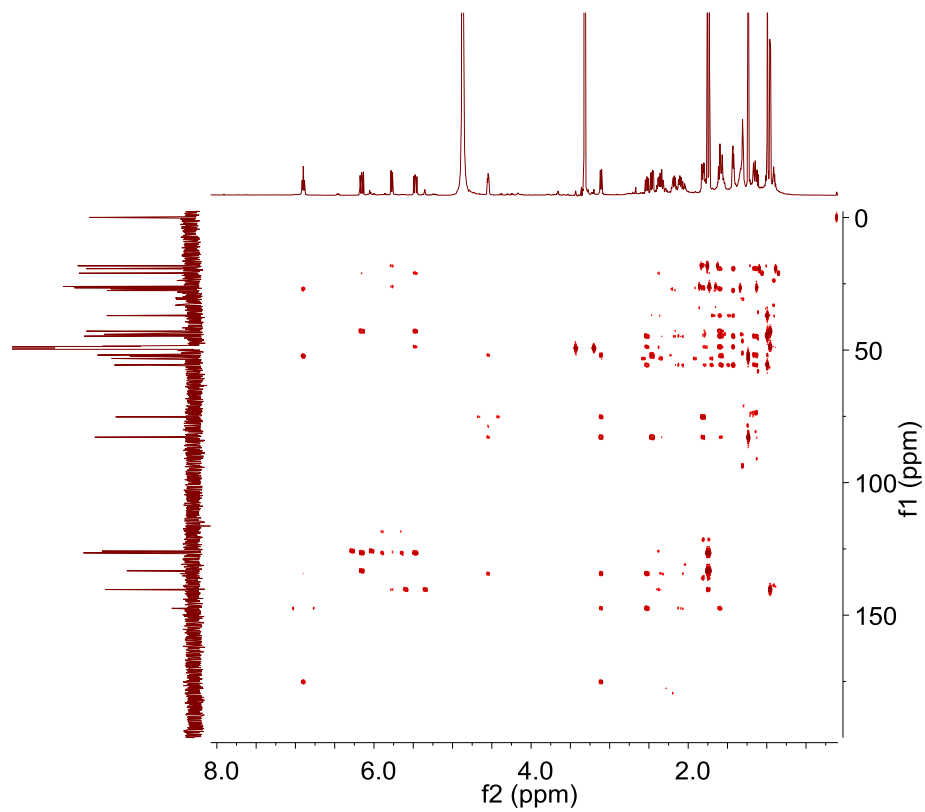


Fig. S49 HMBC spectrum of compound **5** (in CD₃OD).

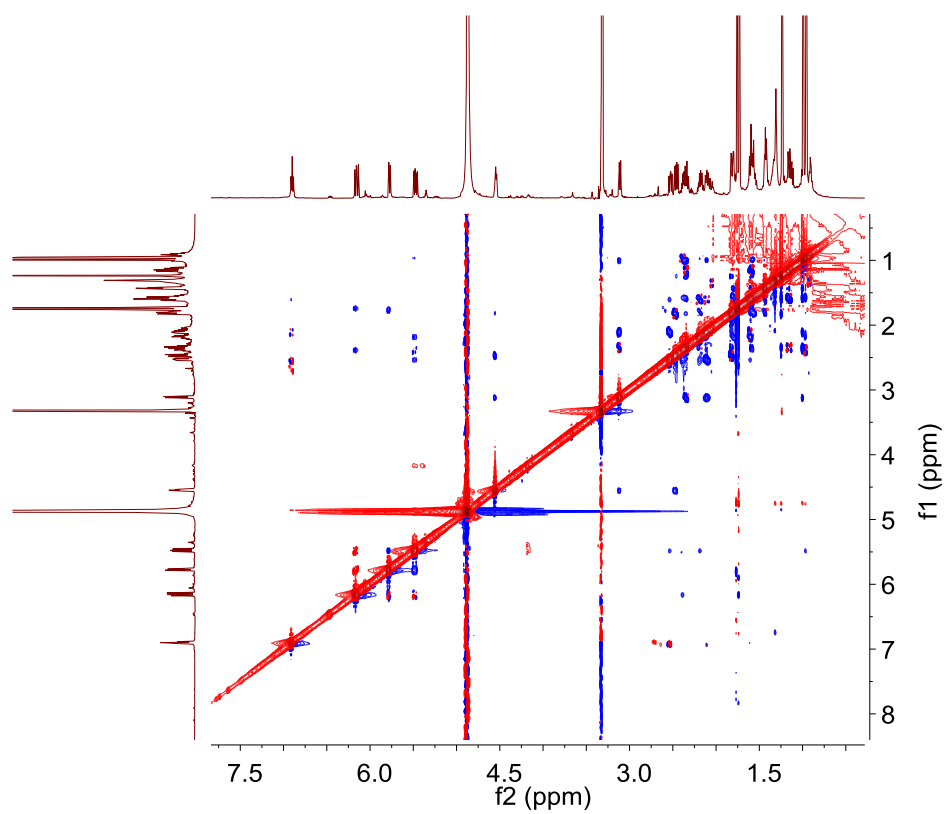


Fig. S50 ROESY spectrum of compound **5** (in CD₃OD).

Fig. S51 Cultural and microscopic morphology of strain WHU0154

a. Cultural characteristics (7 days, front) of strain WHUF0154 on a Potato Dextrose Agar medium at 28 °C. The diameter of the colony is 60~65mm. The front of the colony is white, velvety texture, no exudate, and the center of the colony turns light brown-gray when grow old. The reverse side of the colony is gray.

b. The conidiogenous structures of strain WHU0154 (400×). The conidiophore originates from aerial hyphae, with a diameter of 3.8-5 μm, smooth, and the top is slightly enlarged to form a flask-shaped vesicle with a diameter of 7.5-8.8 μm. The upper part to 3/4 of the surface is fertile; the sporulation structure is double-layered, the stem base is 3-4×2-3μm; the bottle stem is 6-7×3-4μm; the conidia are spherical, with a diameter of 2-3μm, smooth.

