

Phocoenamicins B and C, new antibacterial spirotetronates from the Marine Actinomycete *Micromonospora chaiyaphumensis*

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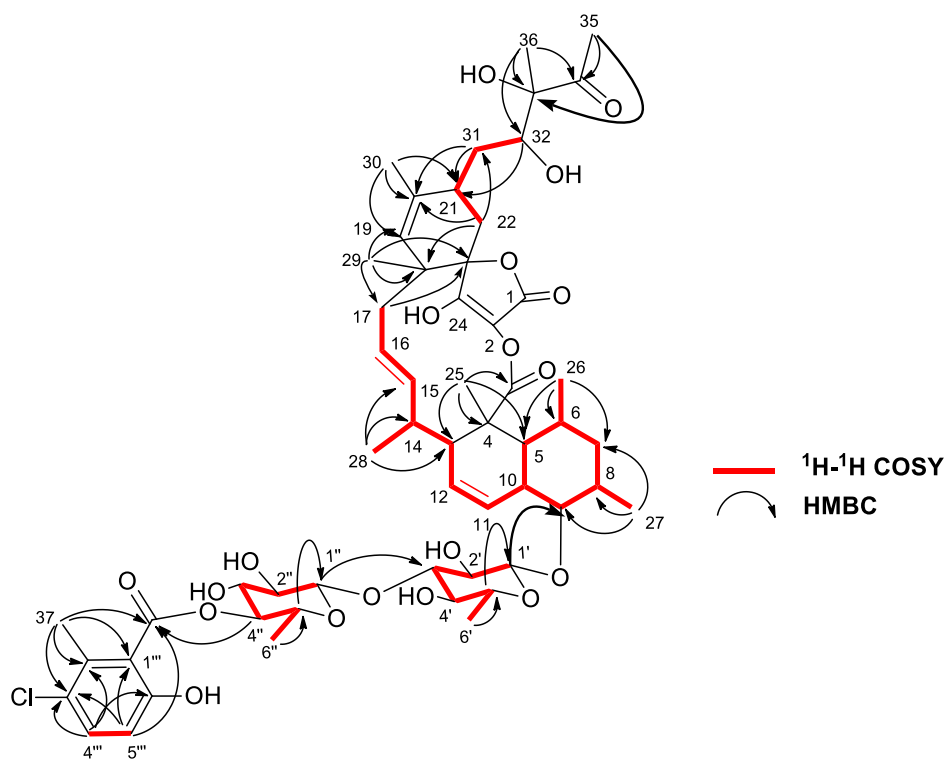
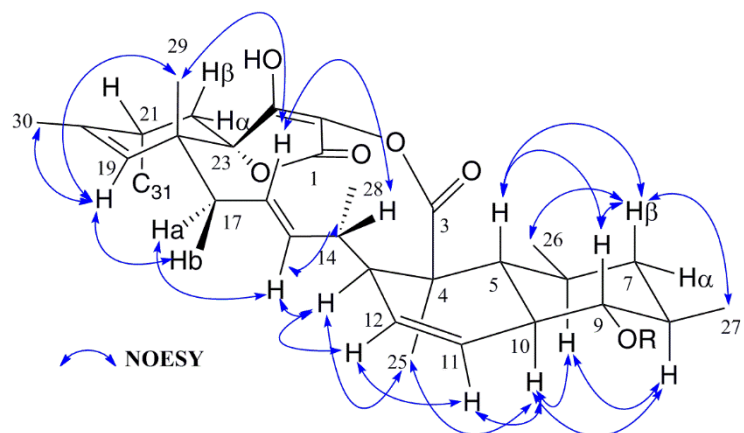
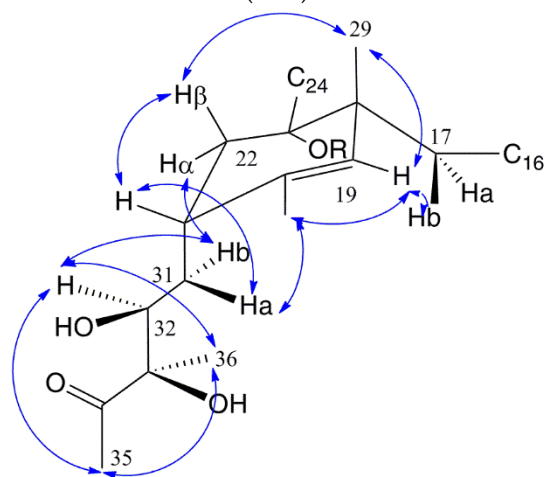


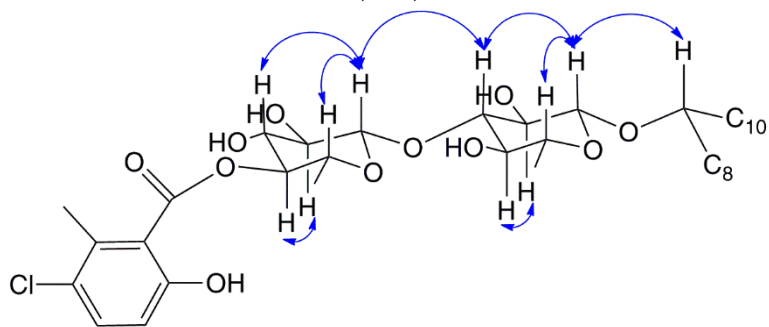
Figure S1. ^1H - ^1H COSY and key HMBC correlations for **2**.



(S2.a)



(S2.b)



(S2.c)

Figure S2. Conformation and configuration of the decalin, cyclohexene and sugars moieties of **2** determined by NOESY data and *J*-based analysis.

Text S3. 16S rRNA gene sequence from strain CA-214671.

cgacttcgtccaatcgccagccccaccttcgacggctcctccacaaggggtggccaccggcttcgggtgttccgacttctgtgacgtgacg
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taggccgagccccatcccaggccgaaaaactttccaccccaaccatgacggtc

Table S4. NMR spectroscopic data (500 Mz, CD₃OD) for phocoenamycin (3).

phocoenamycin (3)		
Position	δ_C , type	δ_H (J in Hz)
1	177.8, C	
2	107.3, C	
3	200.1, C	
4	50.9, C	
5	43.6, CH	1.82, m
6	39.5, CH	1.49, m
7 α	45.7, CH ₂	1.72, m
7 β		1.21, m
8	40.9, CH	1.64, m
9	88.6, CH	3.04, t (9.7)
10	48.4, CH	1.97, m
11	126.0, CH	6.29, d (10.0)
12	126.7, CH	5.57, dd (10.0, 6.2)
13	43.0, CH	2.67, *
14	40.0, CH	2.05, m
15	144.1, CH	5.36, dd (15.1, 9.1)
16	123.2, CH	5.14, dd (15.1, 10.4)
17a	44.0, CH ₂	2.31, m
17b		1.86, m
18	41.2, C	
19	132.1, CH	5.02, s
20	134.6, C	
21	34.3, CH	2.37, m
22 α	30.3, CH ₂	2.28, m
22 β		1.71, m
23	87.5, C	
24	205.7, C	
25	17.0, CH ₃	1.59, s
26	23.7, CH ₃	0.82, brs
27	20.1, CH ₃	1.03, d (6.2)

28	21.5, CH ₃	0.82, brs
29	24.7, CH ₃	1.23, brs
30	22.3, CH ₃	1.74, s
31a		1.95, m
31b	33.7, CH ₂	1.74, m
32	74.0, CH	3.82, d (10.6)
33	83.4, C	
34	215.4, C	
35	25.7, CH ₃	2.24, s
36	22.2, CH ₃	1.22, s
1'	104.0, CH	4.35, d (6.2)
2'	75.3, CH	3.45, m
3'	88.6, CH	3.46, m
4'	75.6, CH	3.11, t (8.7)
5'	72.9, CH	3.22, m
6'	18.4, CH ₃	1.27, d (6.0)
1''	105.4, CH	4.61, d (7.8)
2''	76.0, CH	3.42, t (8.6)
3''	75.3, CH	3.65, t (9.7)
4''	77.9, CH	4.89, *
5''	71.7, CH	3.68, m
6''	18.0, CH ₃	1.36, d (6.2)
1'''	124.3, C	
2'''	135.6, C	
3'''	126.0, C	
4'''	132.4, CH	7.25, d (8.7)
5'''	115.8, CH	6.70, d (8.7)
6'''	155.3, C	
7'''	169.3, C	
37	17.9, CH ₃	2.36, s

*Multiplicity not clear, the signal is overlapped.

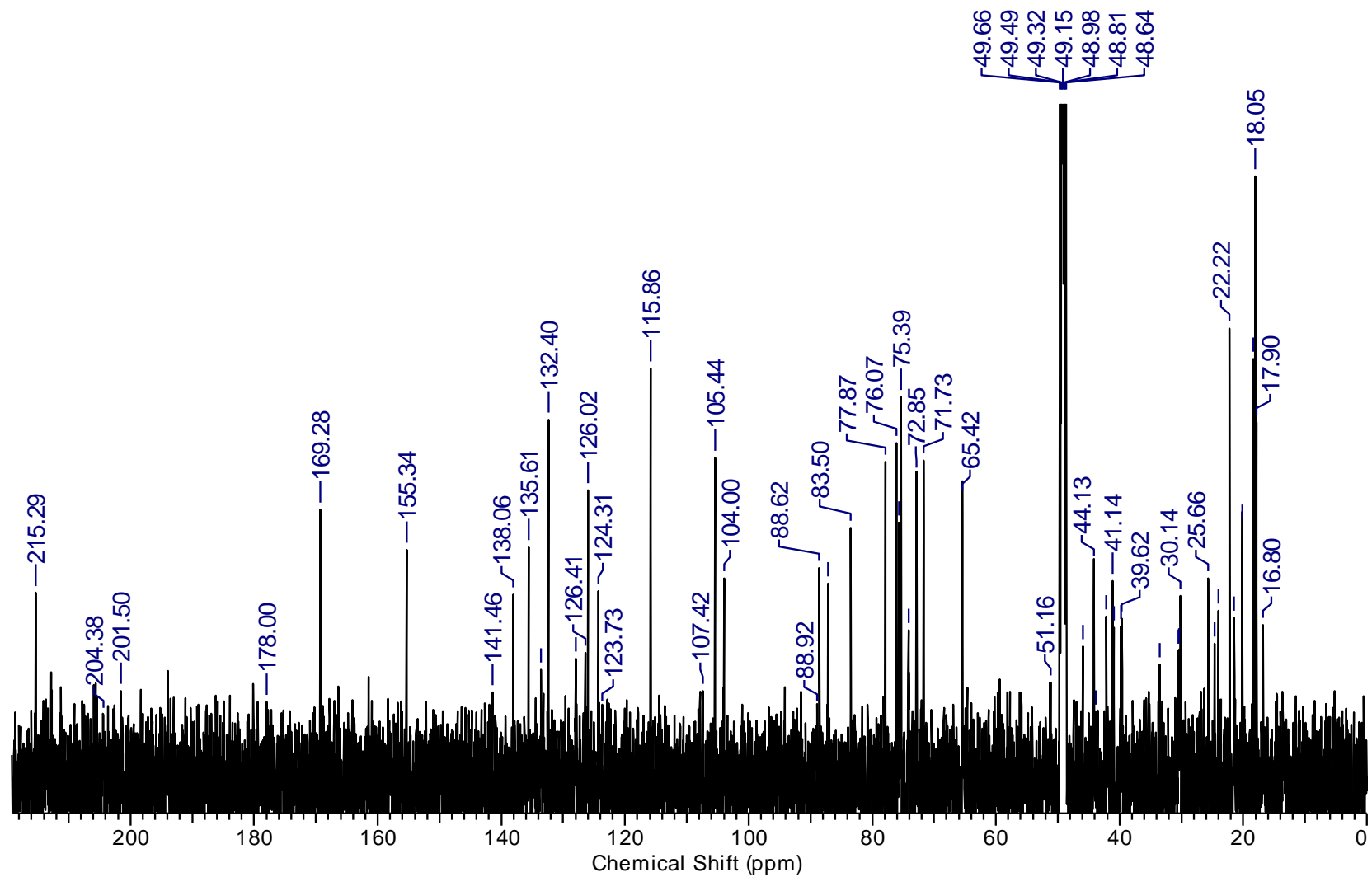


Figure S6. ^{13}C -NMR (125 MHz, methanol- d_4) spectrum of phocoenamicin B (1).

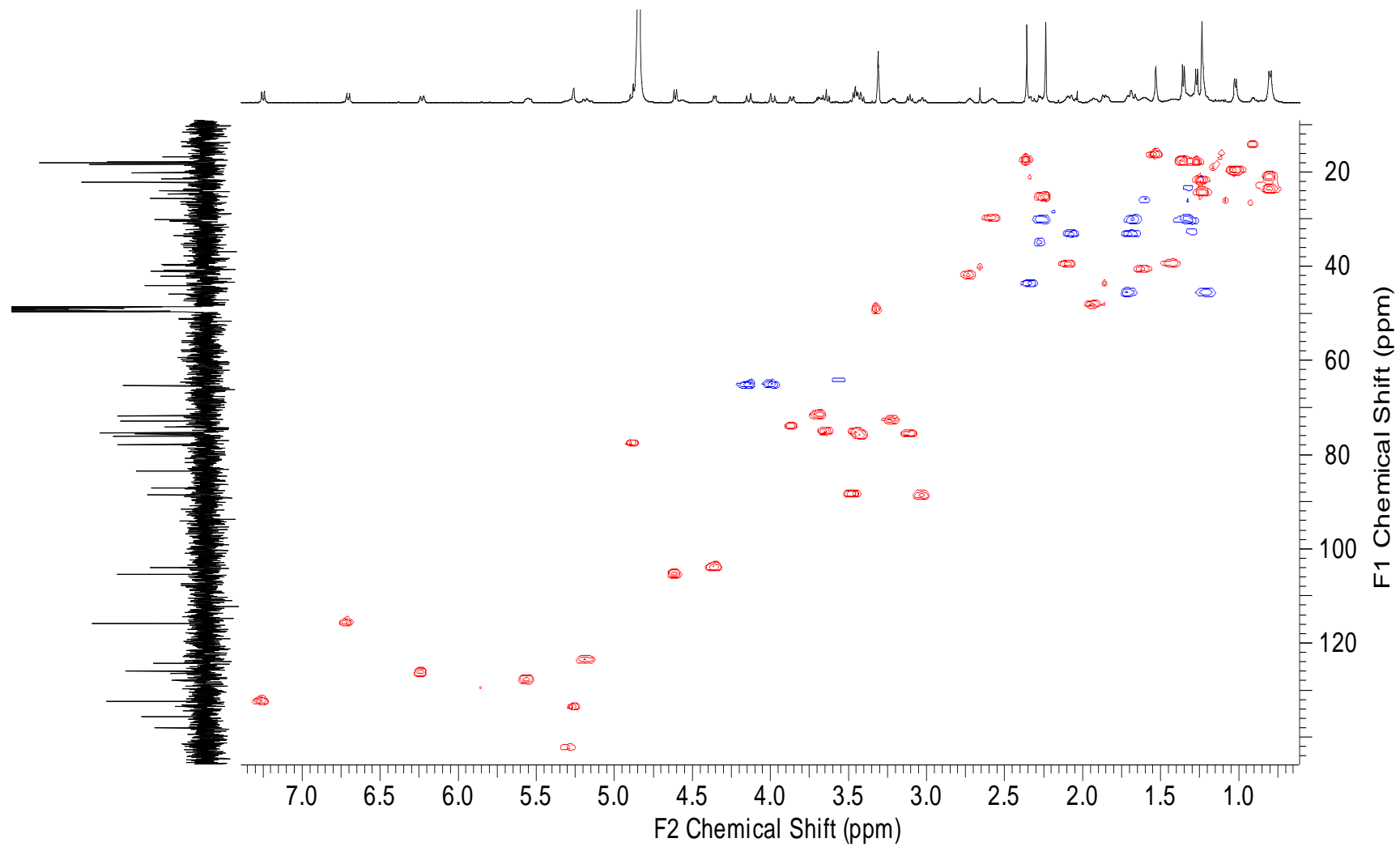


Figure S7. HSQC (methanol-*d*₄) spectrum of phocoenamycin B (1).

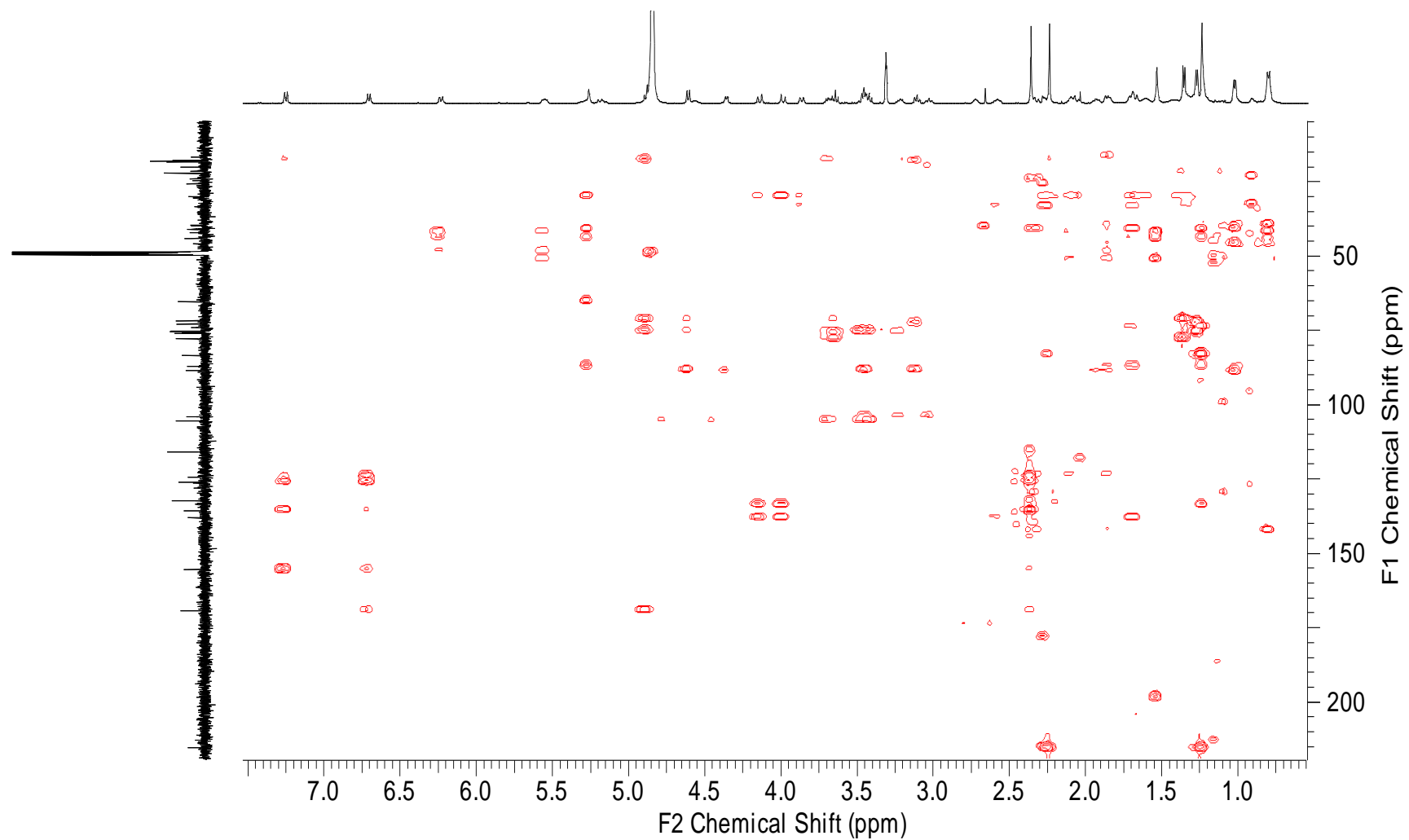


Figure S8. HMBC (methanol- d_4) spectrum of phocoenamycin B (1).

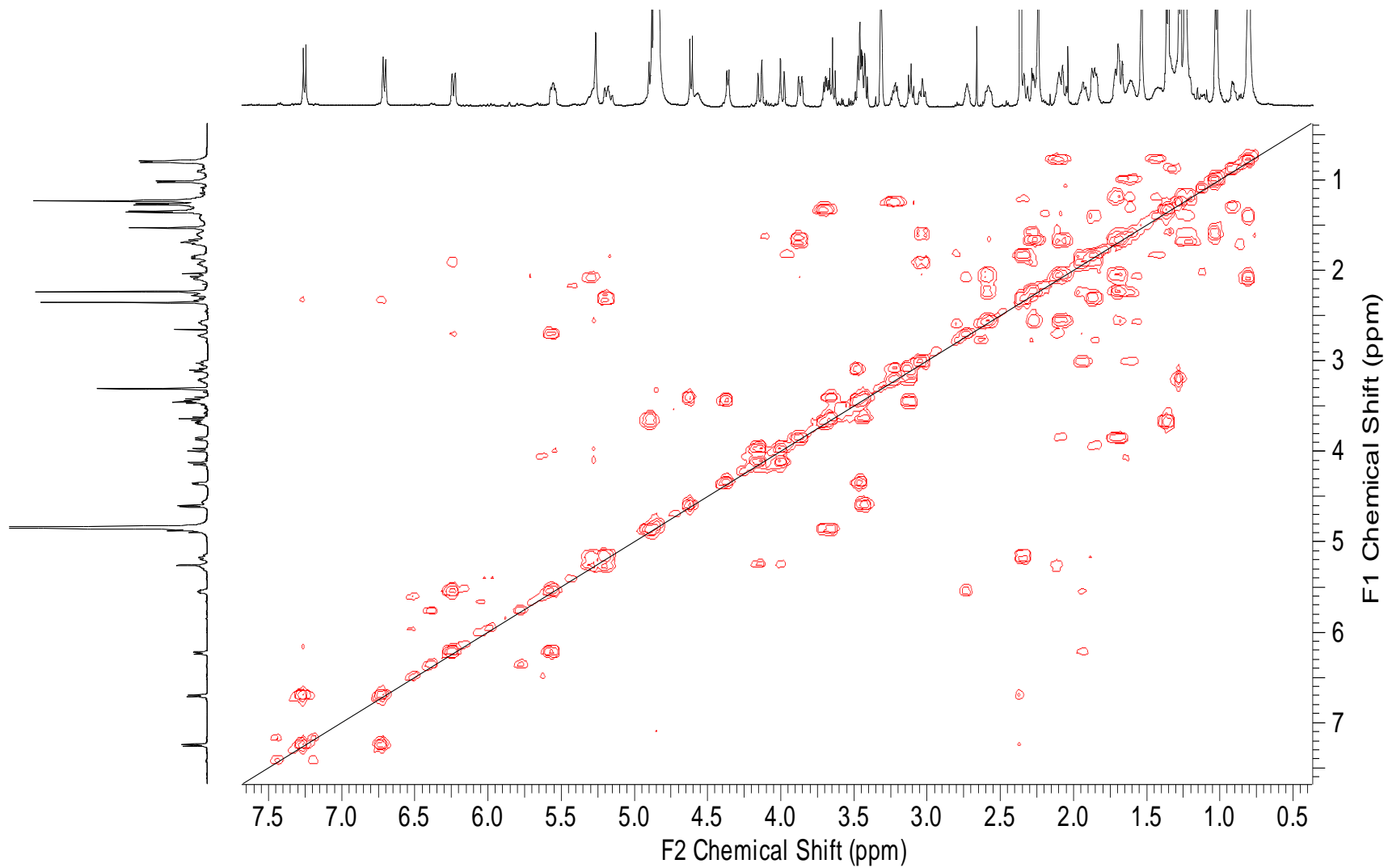


Figure S9. COSY (methanol-*d*₄) spectrum of phocoenamycin B (**1**).

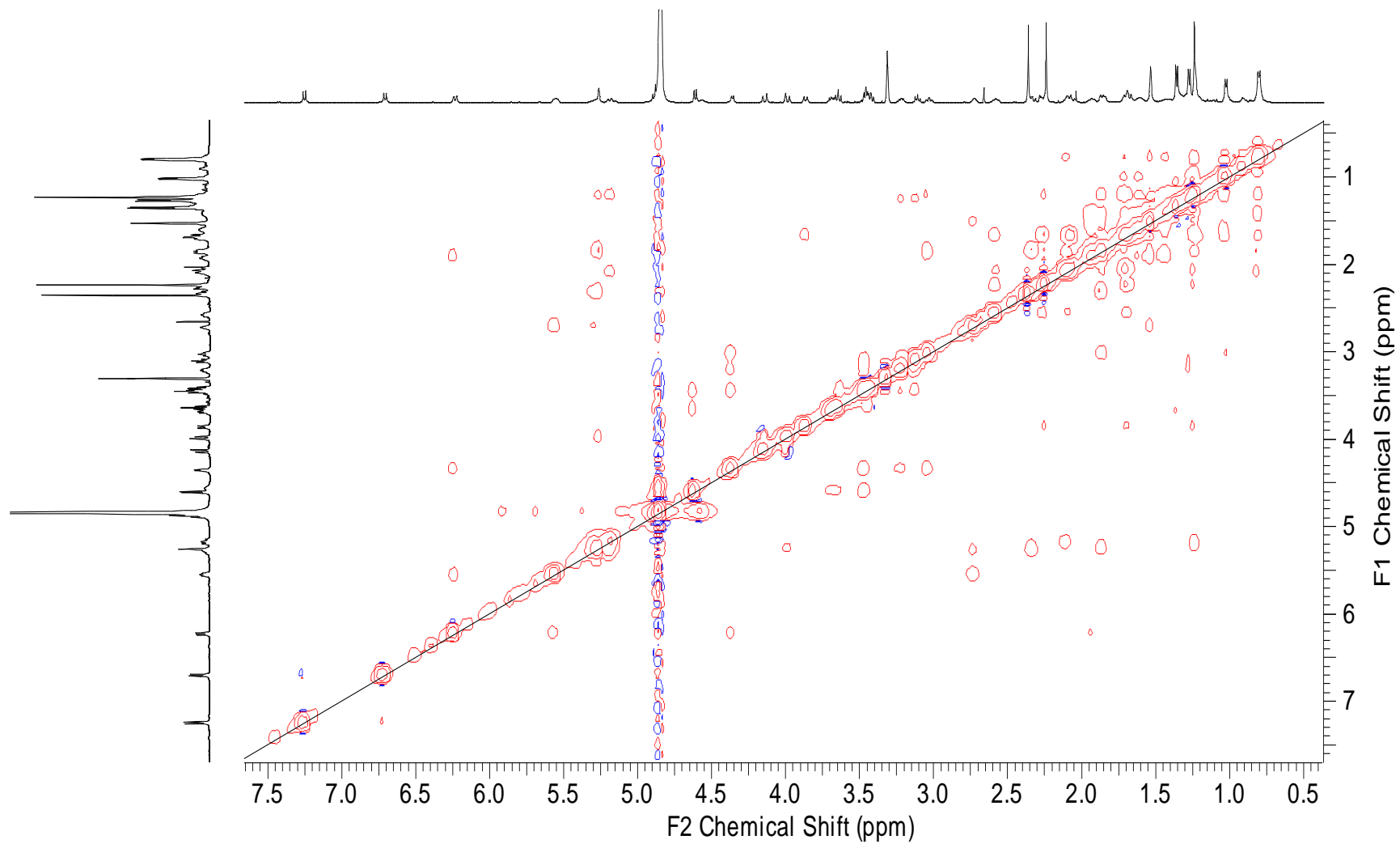


Figure S10. NOESY (methanol- d_4) spectrum of phocoenamycin B (1).

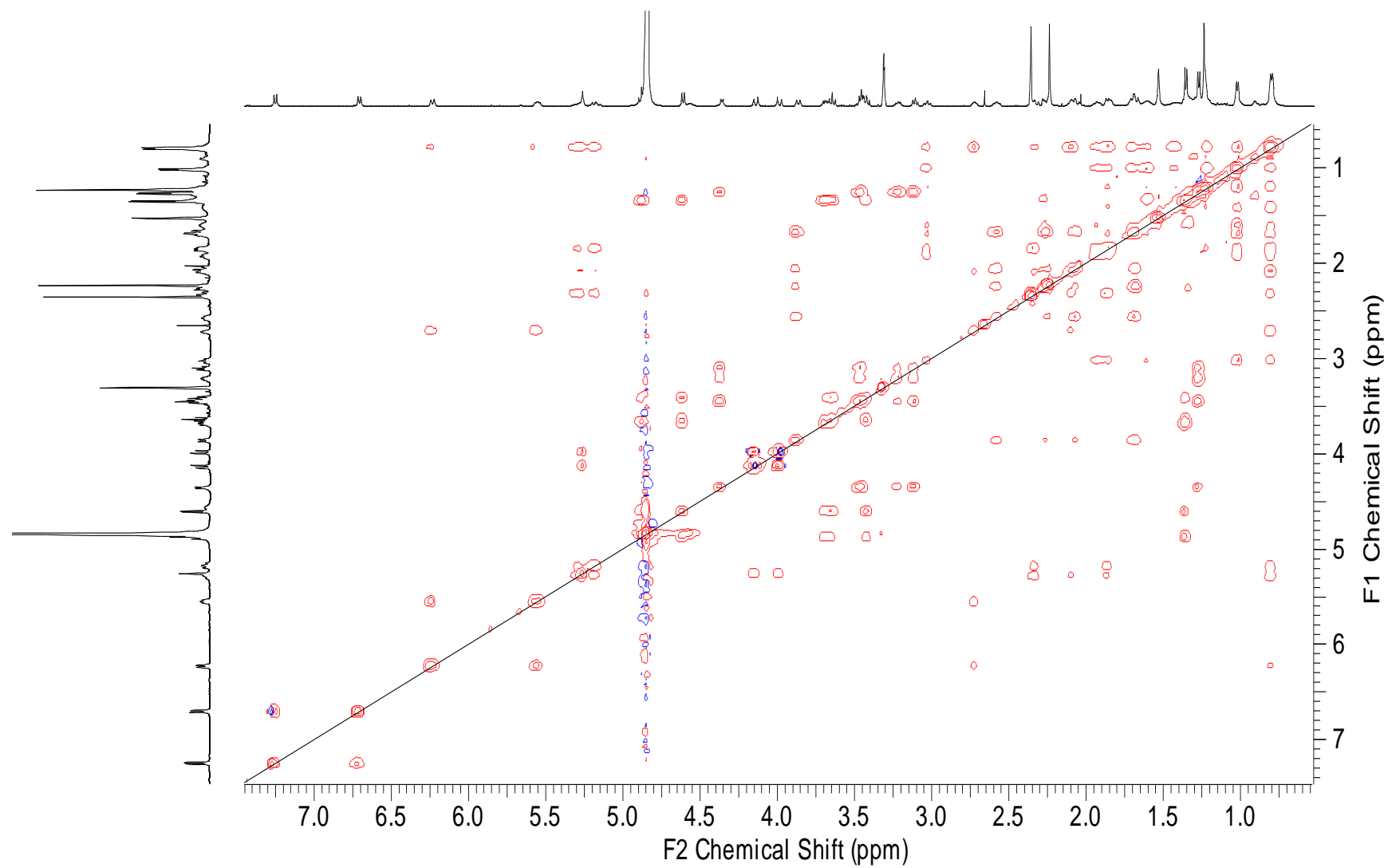


Figure S11. TOCSY (methanol- d_4) spectrum of phocoenamycin B (**1**).

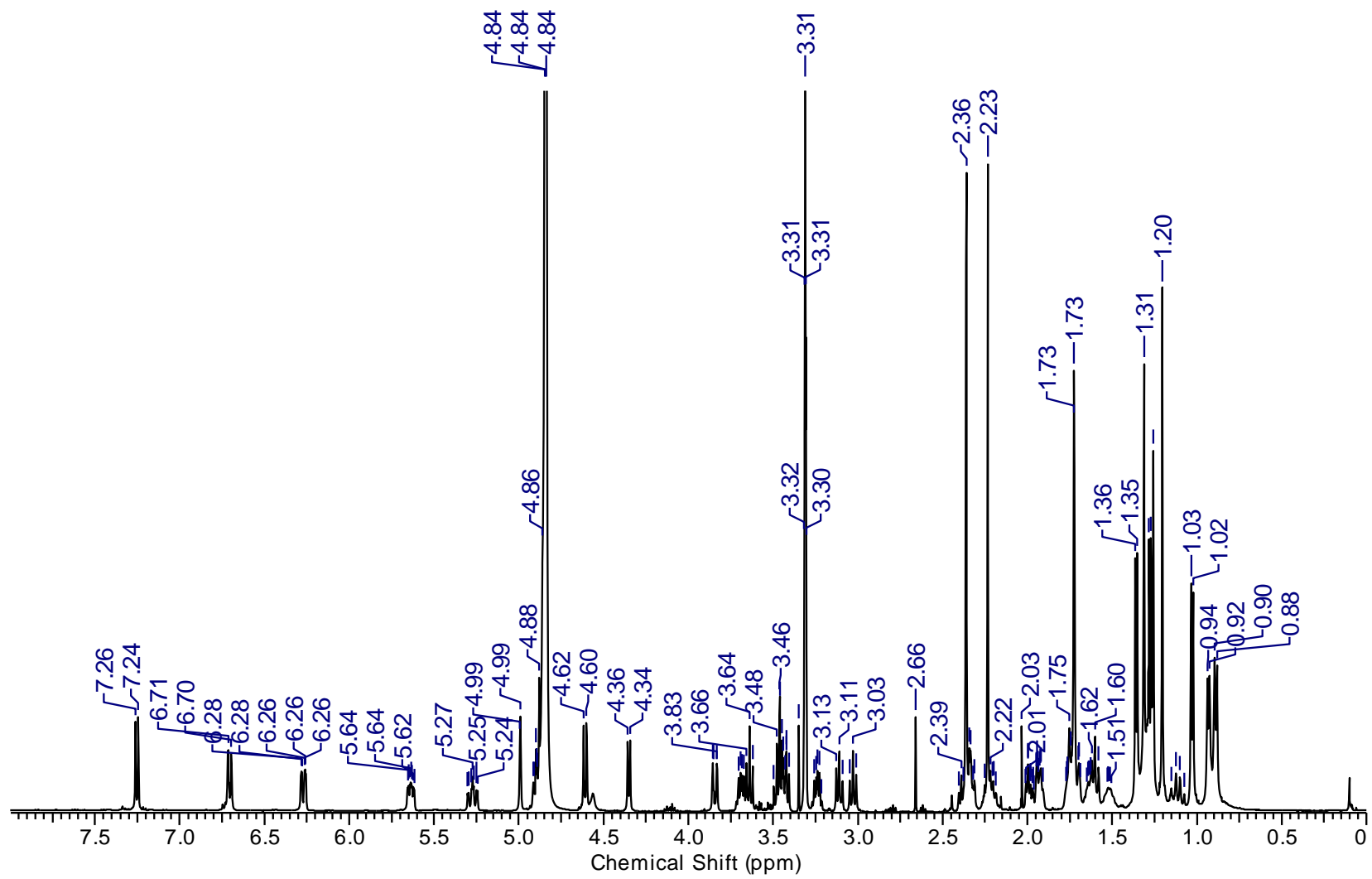


Figure S12. $^1\text{H-NMR}$ (500 MHz, methanol- d_4) spectrum of phocoenamycin C (2).

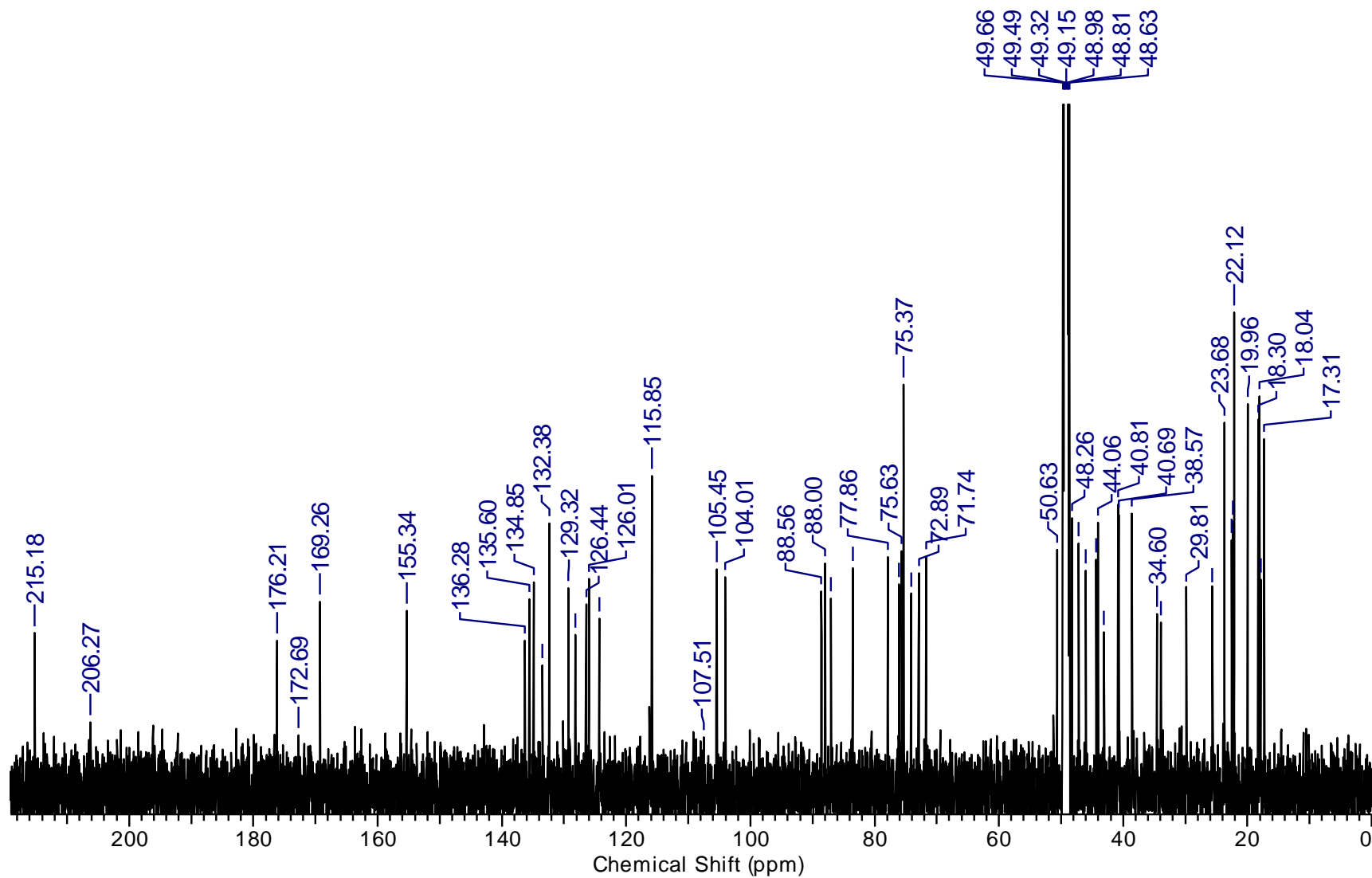


Figure S13. ^{13}C -NMR (125 MHz, methanol- d_4) spectrum of phocoenamicin C (2).

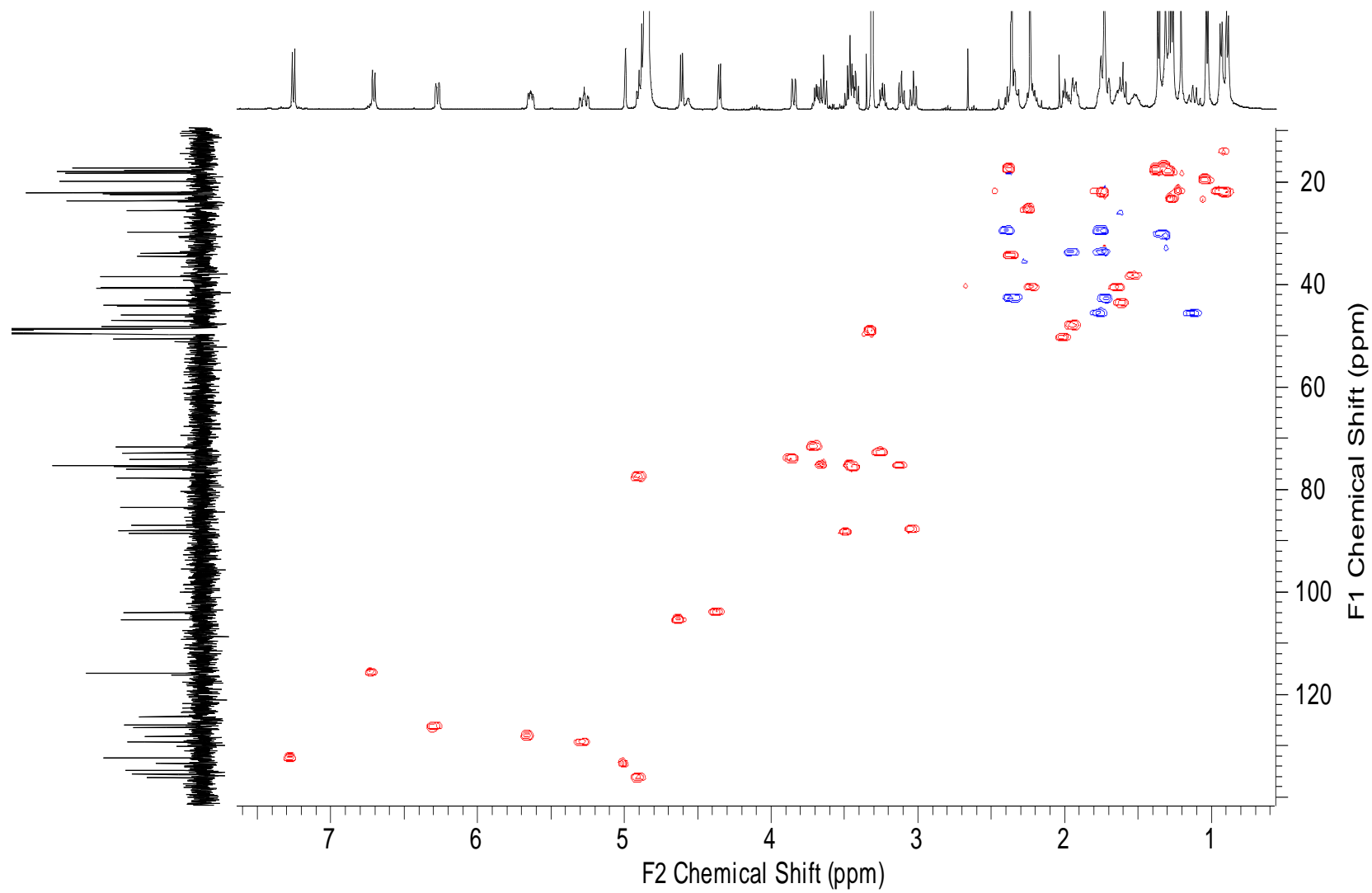


Figure S14. HSQC (methanol-*d*₄) spectrum of phocoenamicin C (2).

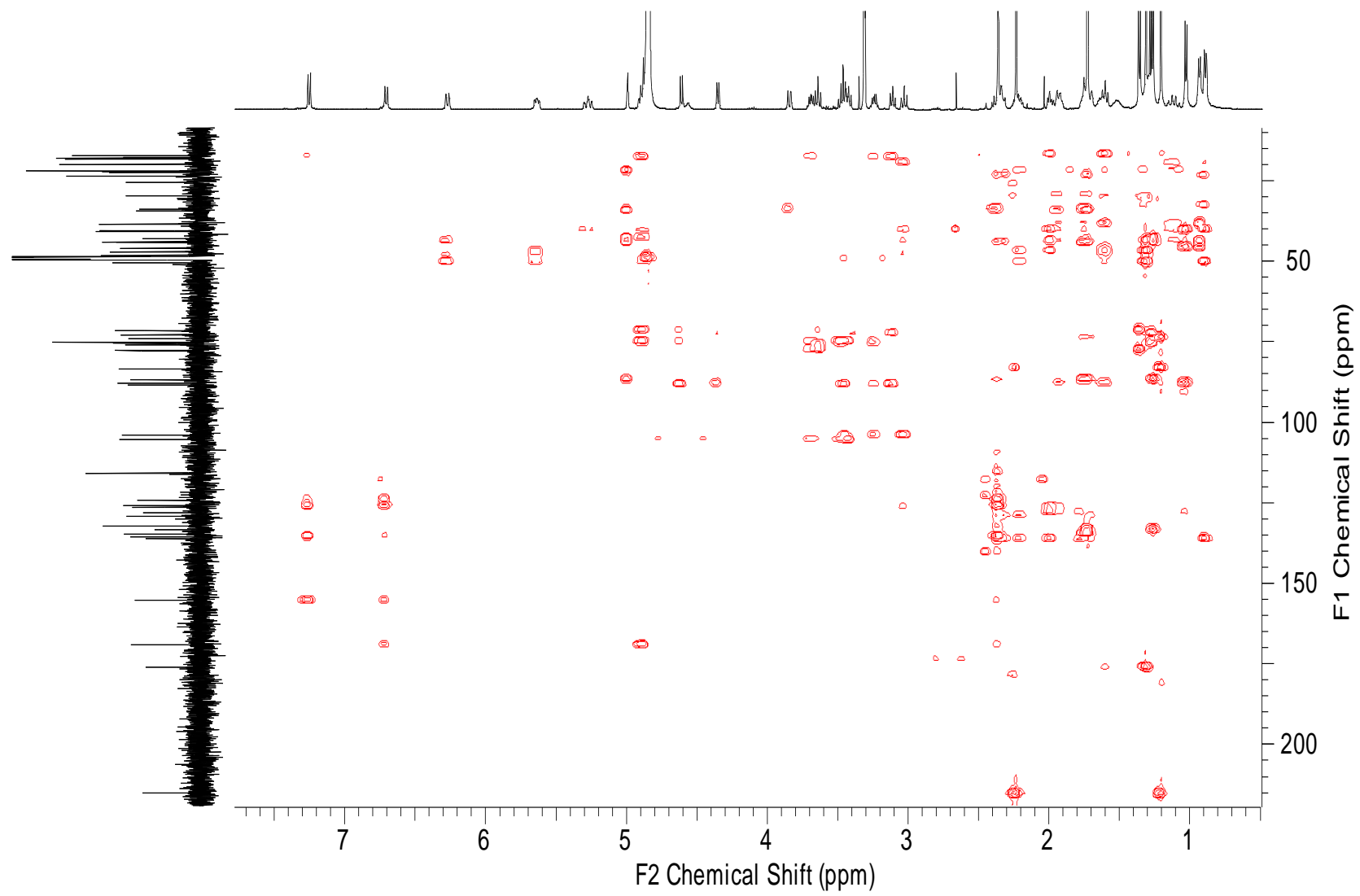


Figure S15. HMBC (methanol-*d*₄) spectrum of phocoenamycin C (2).

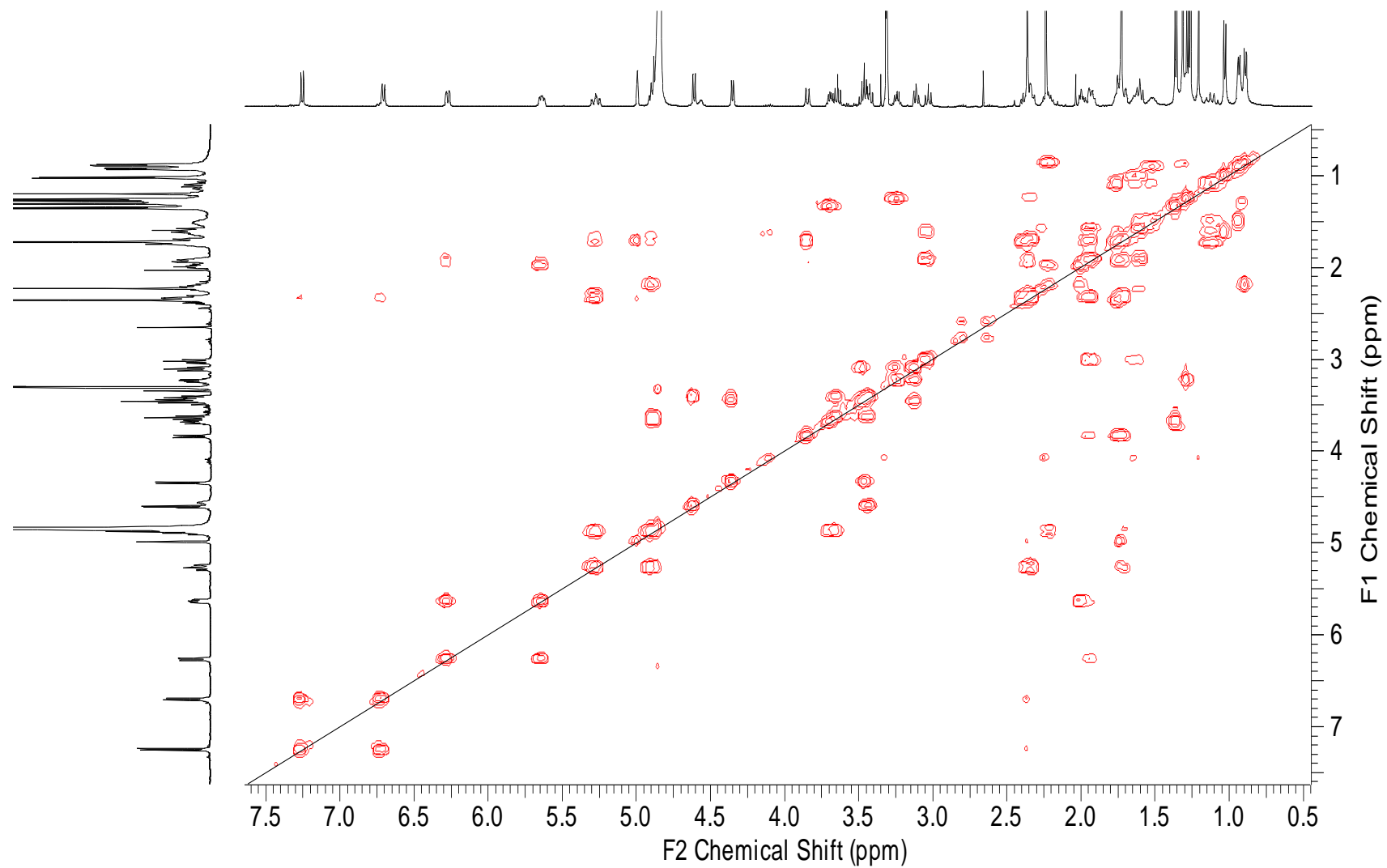


Figure S16. COSY (methanol- d_4) spectrum of phocoenamycin C (2).

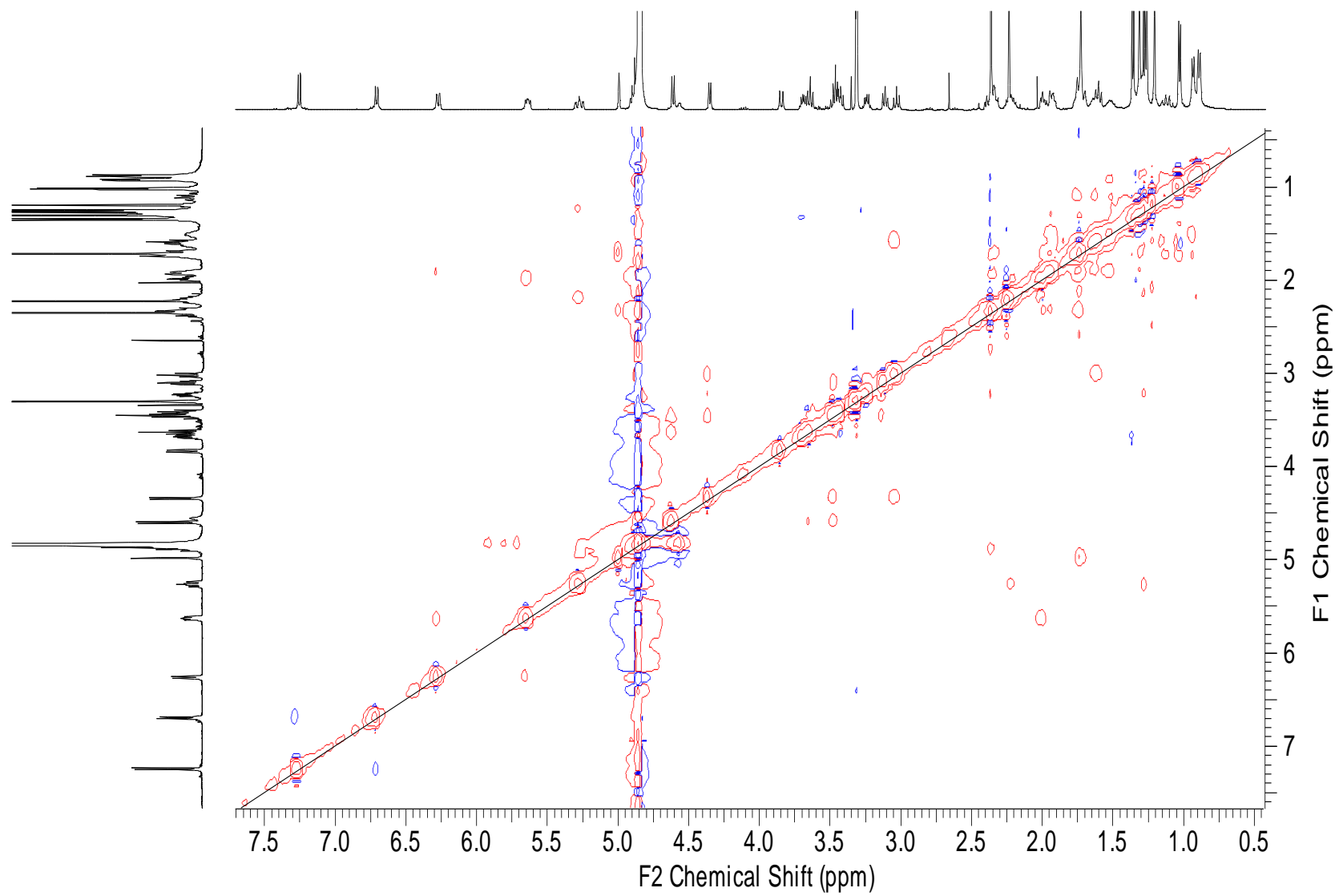


Figure S17. NOESY (methanol-*d*₄) spectrum of phocoenamycin C (2).

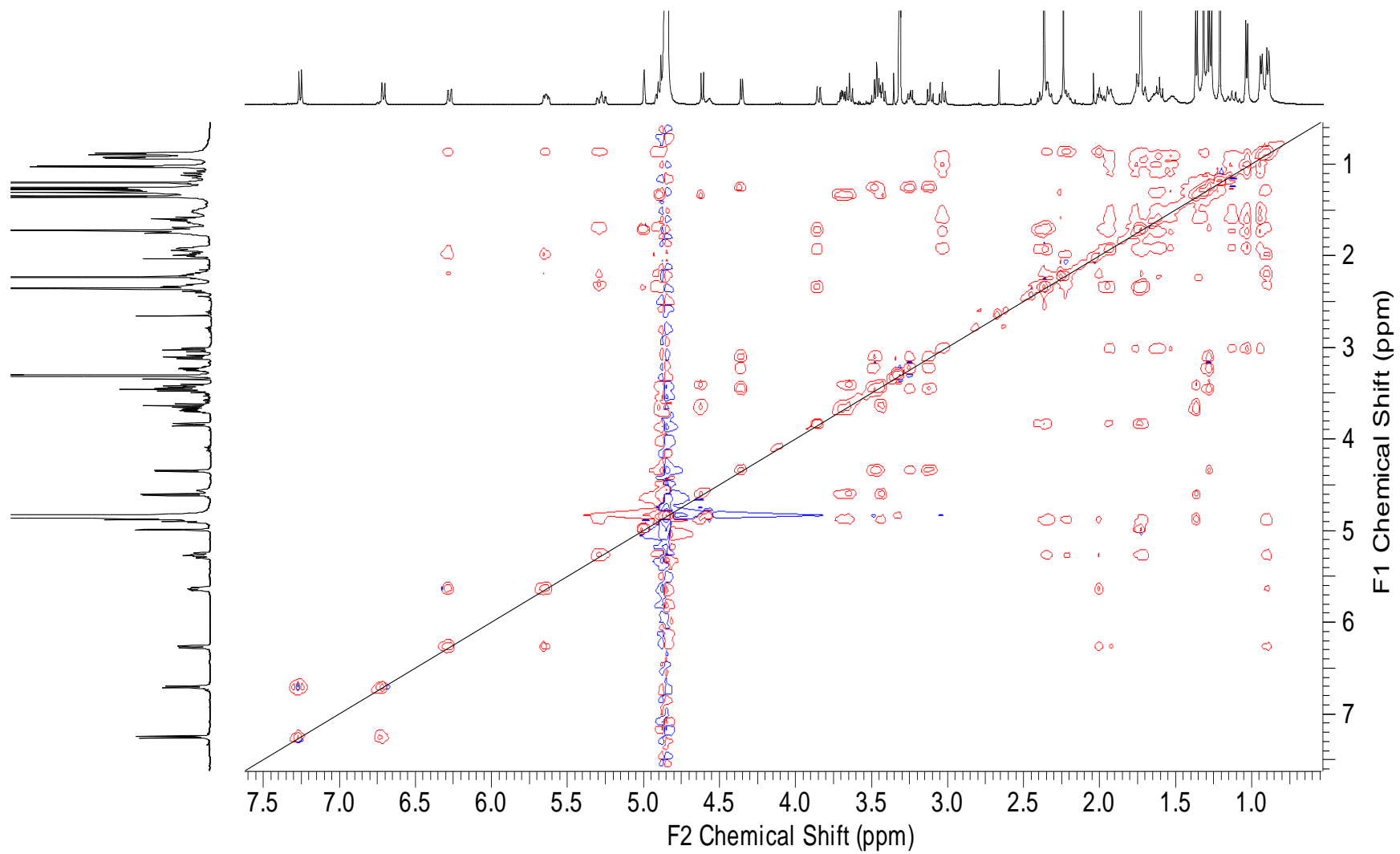


Figure S18. TOCSY (methanol- d_4) spectrum of phocoenamycin C (2).