

Supporting Information

Induction of diverse secondary metabolites in *Aspergillus fumigatus* by microbial co-culture

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Table S1. NMR spectroscopic data for **9** and **10** (CDCl₃, 400 MHz at 298 K):

Atom	9			10			¹³ C NMR of related compounds		
	δ _H , mult. (<i>J</i> in Hz)	δ _C , mult	δ _C , mult ^a	δ _H , mult. (<i>J</i> in Hz)	δ _C , mult	δ _H , mult ^b (<i>J</i> in Hz)	δ _C , mult ^b	δ _C , mult ^c	δ _C , mult ^d
2		186.6, C	187.3, C*		184.4, C		186.8, C	186.8, C	187.6, C
3		113.4, C	113.6, C		113.3, C		111.4, C	111.6, C	111.3, C
4		197.7, C	198.0, C		198.1, C		199.8, C	196.7, C	200.0, C
5		91.8, C	92.3, C		88.5, C		88.2, C	92.4, C	88.2, C
6		166.1, C	167.2, C*		167.9, C		167.5, C	166.6, C	167.7, C
8		90.0, C	92.0, C		95.4, C		97.0, C	91.1, C	97.1, C
9	4.63, s	74.1, CH	75.0, CH	4.78, s	76.5, CH	4.62 (d, 6.2)	75.6, CH	75.0, CH	75.7, CH
10	4.69 (d, 5.7)	70.5, CH	69.7, CH	4.56 (d, 6.4)	70.3, CH	4.48 (t, 5.4)	70.4, CH	72.0, CH	72.8, CH
11	4.32 (dd, 5.2, 9.2)	77.6, CH	77.3, CH	4.28 (dd, 6.5, 9.1)	77.6, CH	4.17 (dd, 5.5, 9.3)	77.1, CH	68.3, CH	68.3, CH
12	5.33 (ddt, 10.9, 9.5, 1.3)	124.1, CH	125.1, CH	5.27 (t, 10.7)	124.7, CH	5.31 (t, 10.9)	125.6, CH	128.4, CH	129.0, CH
13	5.83, m	140.1, CH	138.4, CH	5.82, m	140.3, CH	5.62, m	136.9, CH	134.0, CH	133.8, CH
14	2.11, m	21.4, CH ₂	20.8, CH ₂	2.10, m	21.4, CH ₂	20.4, m	20.7, CH ₂	20.6, CH ₂	20.7, CH ₂
						1.97, m			
15	0.96 (t, 7.5)	14.3, CH ₃	13.1, CH ₃	0.98 (t, 7.5)	14.3, CH ₃	0.87 (t, 7.6)	14.1, CH ₃	14.1, CH ₃	14.2, CH ₃
16	1.75, s	5.8, CH ₃	4.3, CH ₃	1.77, s	5.8, CH ₃	1.64, s	5.6, CH ₃	5.6, CH ₃	5.7, CH ₃
17		194.6, C	195.8, C		193.3, C		194.7, C	196.4, C	194.8, C
18		132.5, C	133.5, C		134.0, C		134.8, C	133.6, C	134.7, C
19/23	8.29(dd, 1.2, 8.5)	130.7, CH	130.2, CH	8.21 (d, 7.9)	129.9, CH	8.11 (d, 7.7)	129.6, CH	130.2, CH	129.7, CH
20/22	7.49 (t, 8.0)	128.9, CH	128.1, CH	7.47 (t, 7.8)	128.7, CH	7.52 (t, 7.8)	128.3, CH	128.4, CH	128.4, CH
21	7.46 (t, 7.6)	134.8, CH	133.7, CH	7.60 (t, 7.2)	134.0, CH	7.63 (t, 7.6)	133.1, CH	133.6, CH	133.2, CH
8-OMe	3.38, s	51.9, CH ₃	51.0, CH ₃	3.29, s	51.6, CH ₃	3.12, s	51.4, CH ₃	51.7, CH ₃	51.5, CH ₃
11-OMe	3.31, s	56.8, CH ₃	55.4, CH ₃	3.25, s	56.7, CH ₃	3.15, s	55.7, CH ₃	---	---
9-OH						6.09 (d, 6.2)			
10-OH						5.75 (d, 5.8)			
NH-7	7.67, br s			7.40, br s		9.89, s			

^a Reported ¹³C NMR data of 11-*O*-methylpseurotin A (**9**) in CD₃OD.⁴⁴ ^b NMR data of **7.10** in DMSO-*d*₆. ^c Reported ¹³C NMR data of pseurotin A in acetone-*d*₆.⁴⁵ ^d Reported ¹³C NMR data of pseurotin A₂ in DMSO-*d*₆.⁴⁶ *The ¹³C shifts were interchanged as they were wrongly reported.

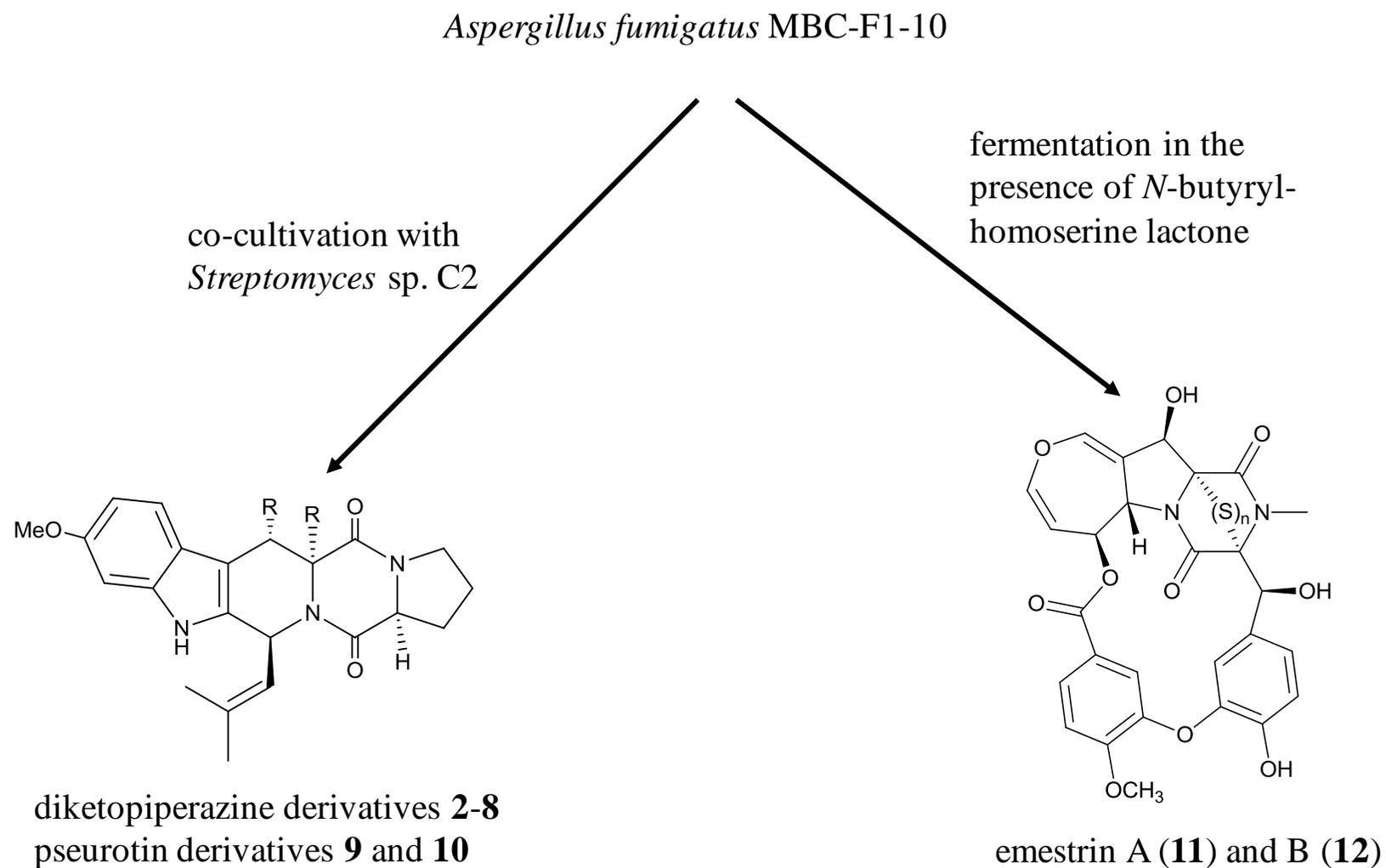


Fig. S1. Induction of cryptic biogenetic pathways in *Aspergillus fumigatus* MBC-F1-10 by different strategies

atd51 #36 RT: 0.41 AV: 1 NL: 2.12E7
F: FTMS + p ESI Full ms [100.00-2000.00]

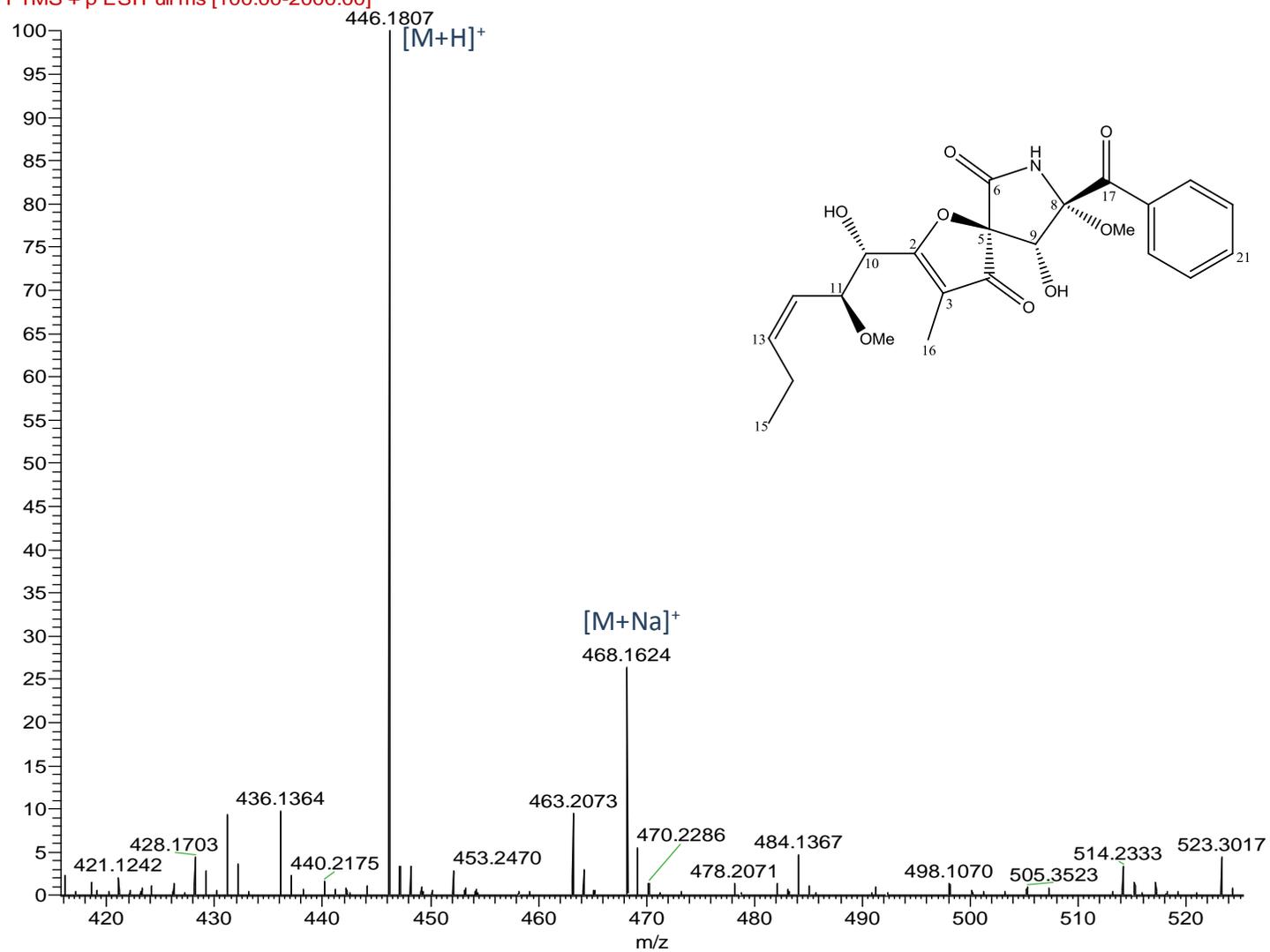


Fig. S2. HRESIMS spectrum of **9**

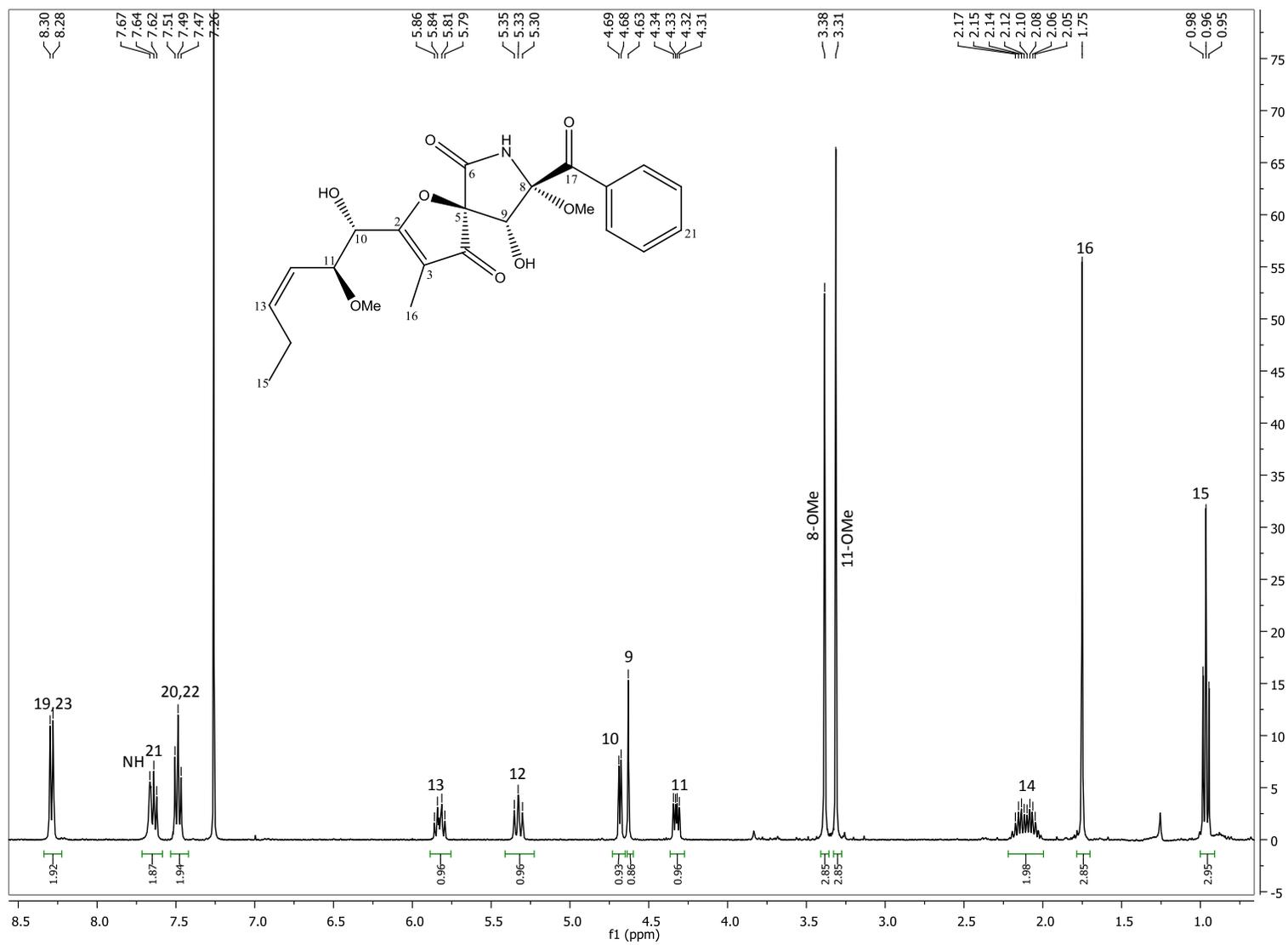


Fig. S3. ^1H NMR spectrum of **9**

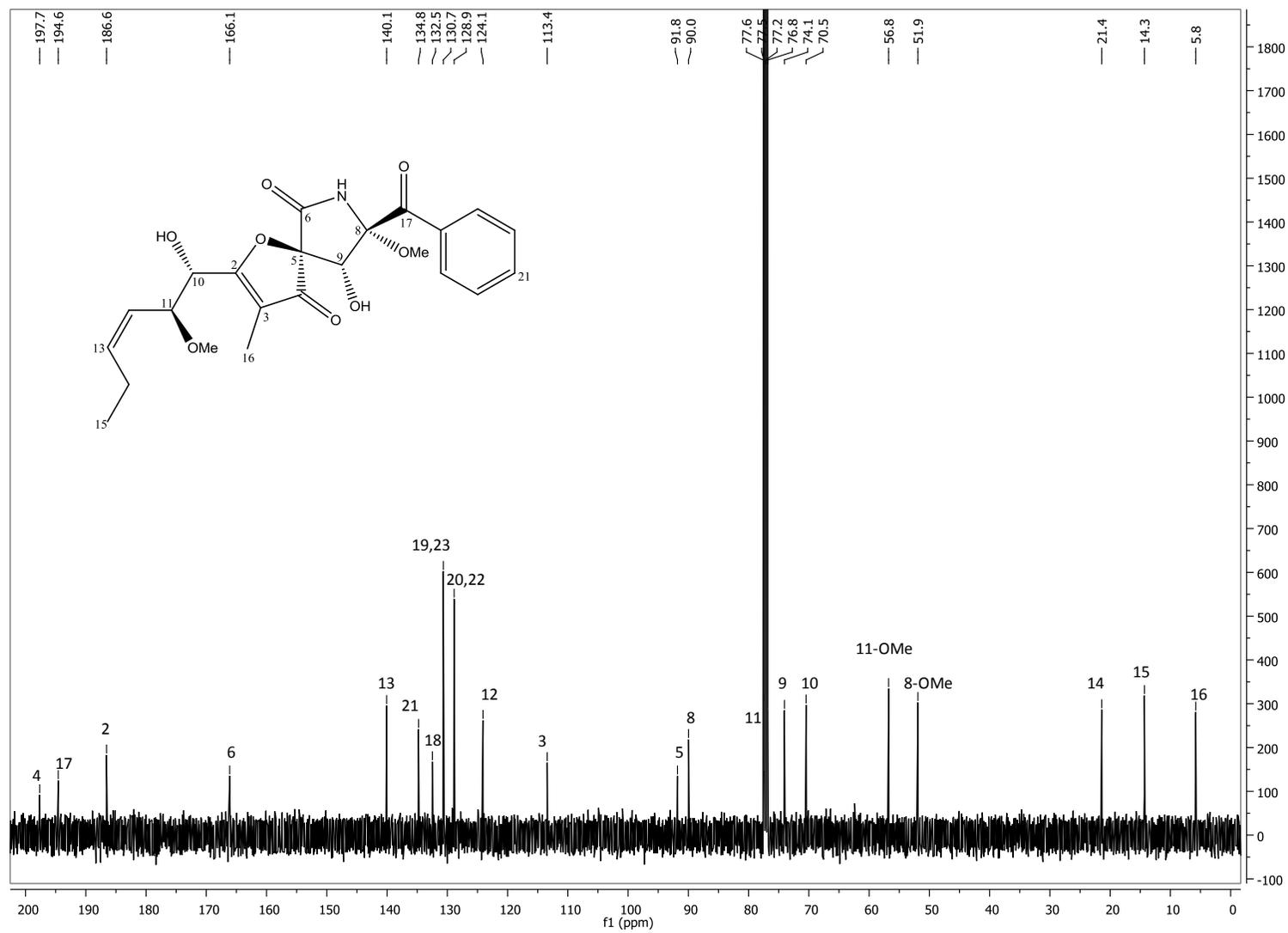


Fig. S4. ^{13}C NMR spectrum of **9**

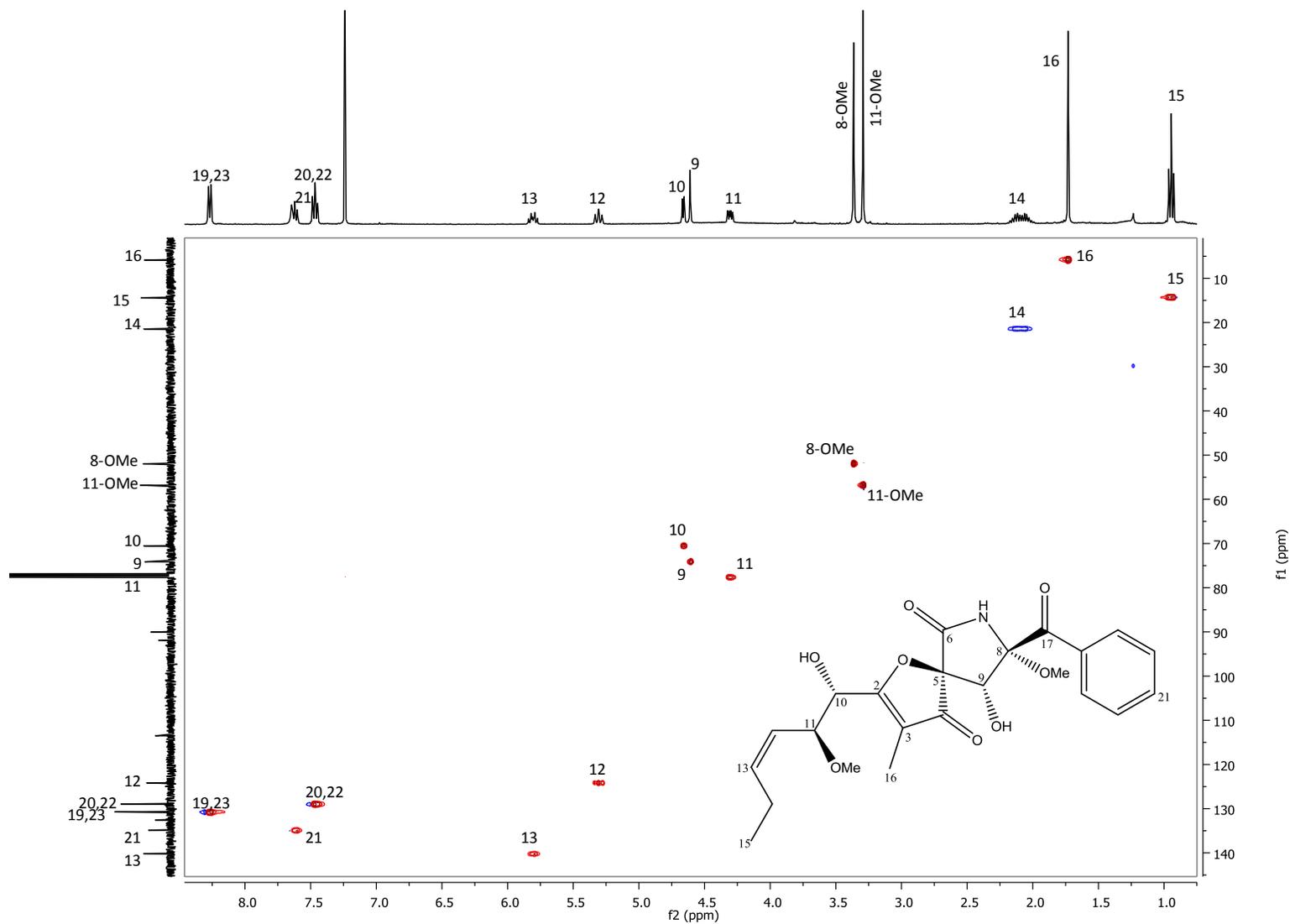


Fig. S5. HSQC spectrum of **9**

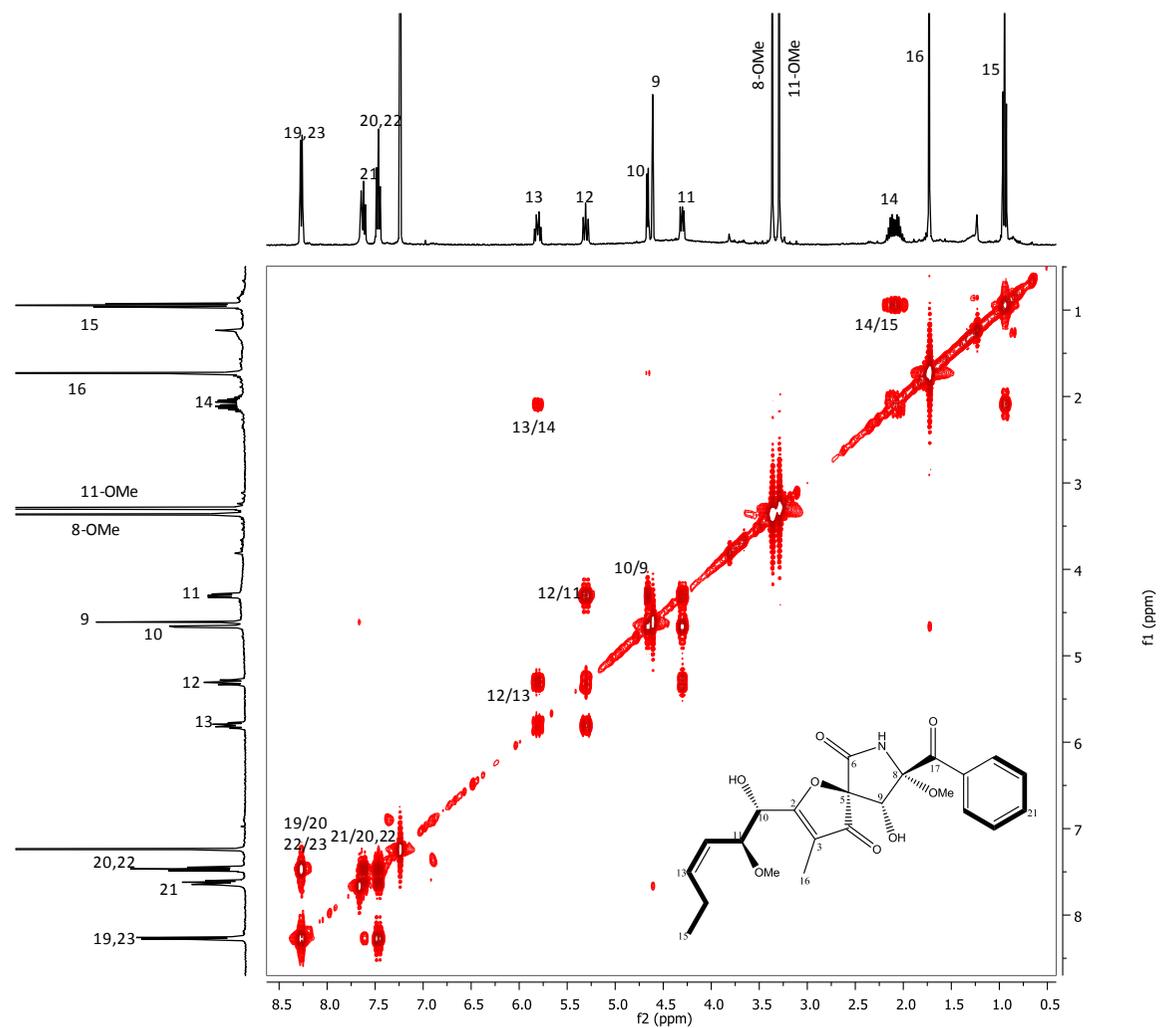


Fig. S6. COSY spectrum of **9**

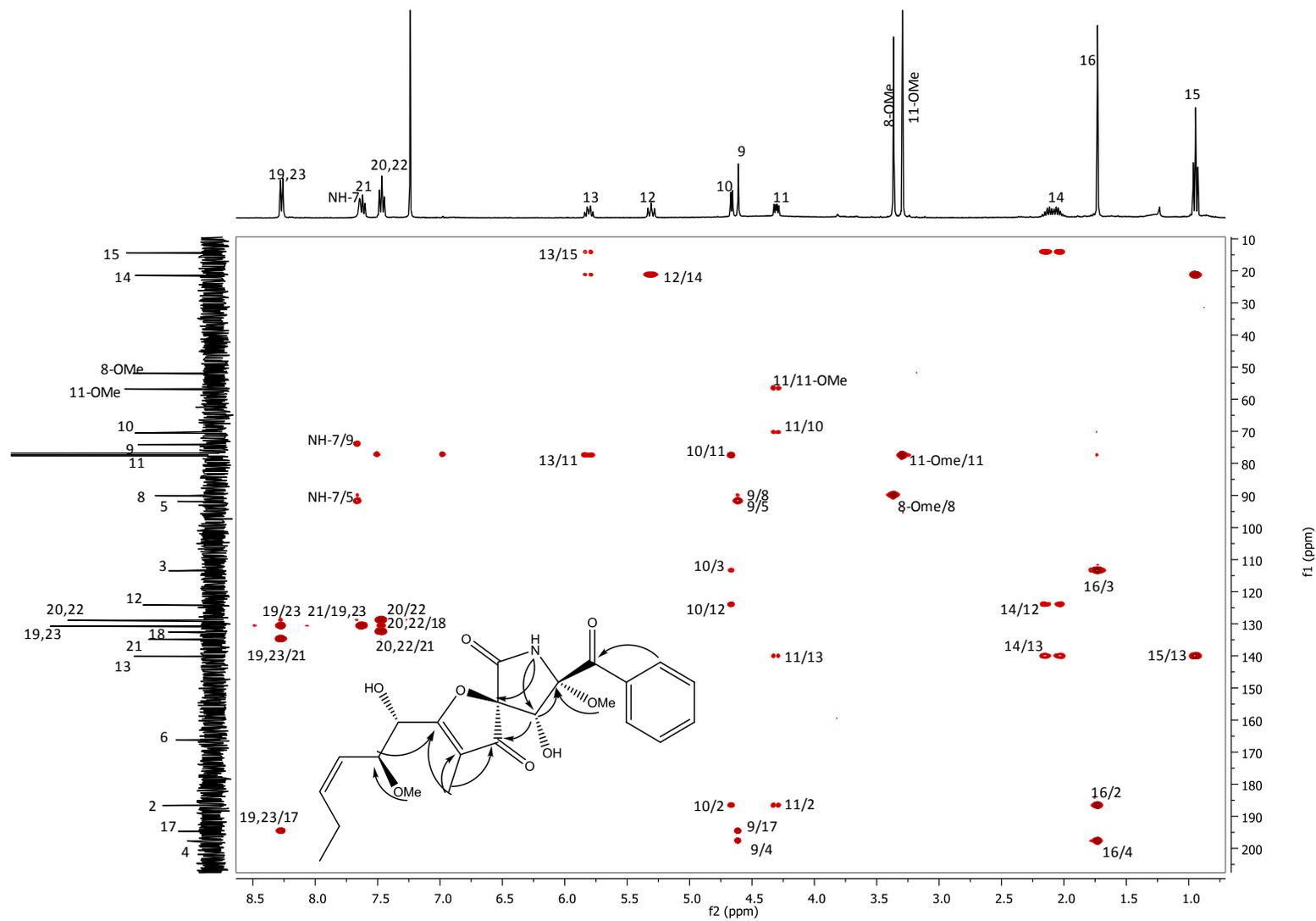


Fig. S7. HMBC spectrum of **9**

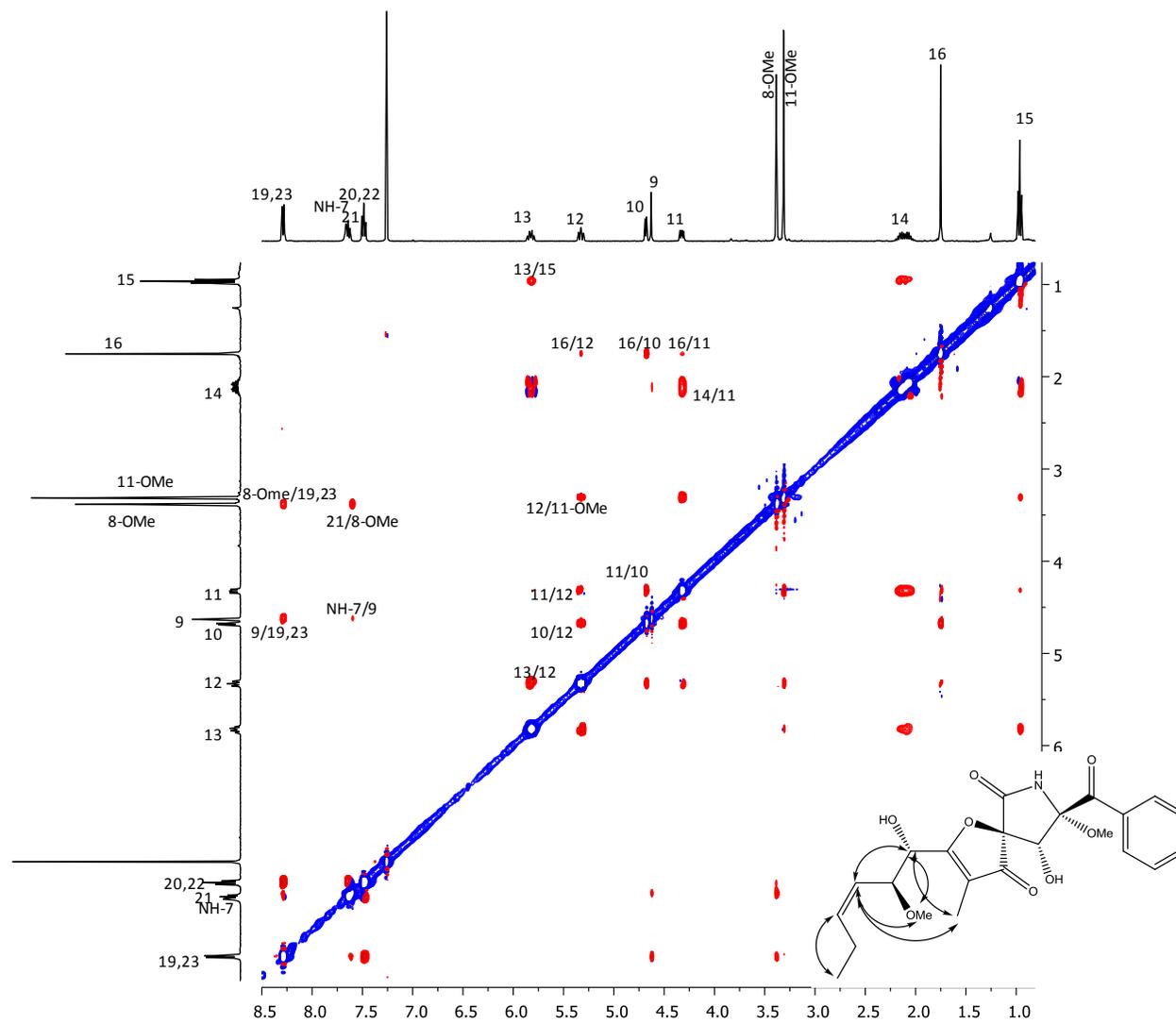


Fig. S8. NOESY spectrum of **9**

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F: FTMS + p ESI Full ms [100.00-2000.00]

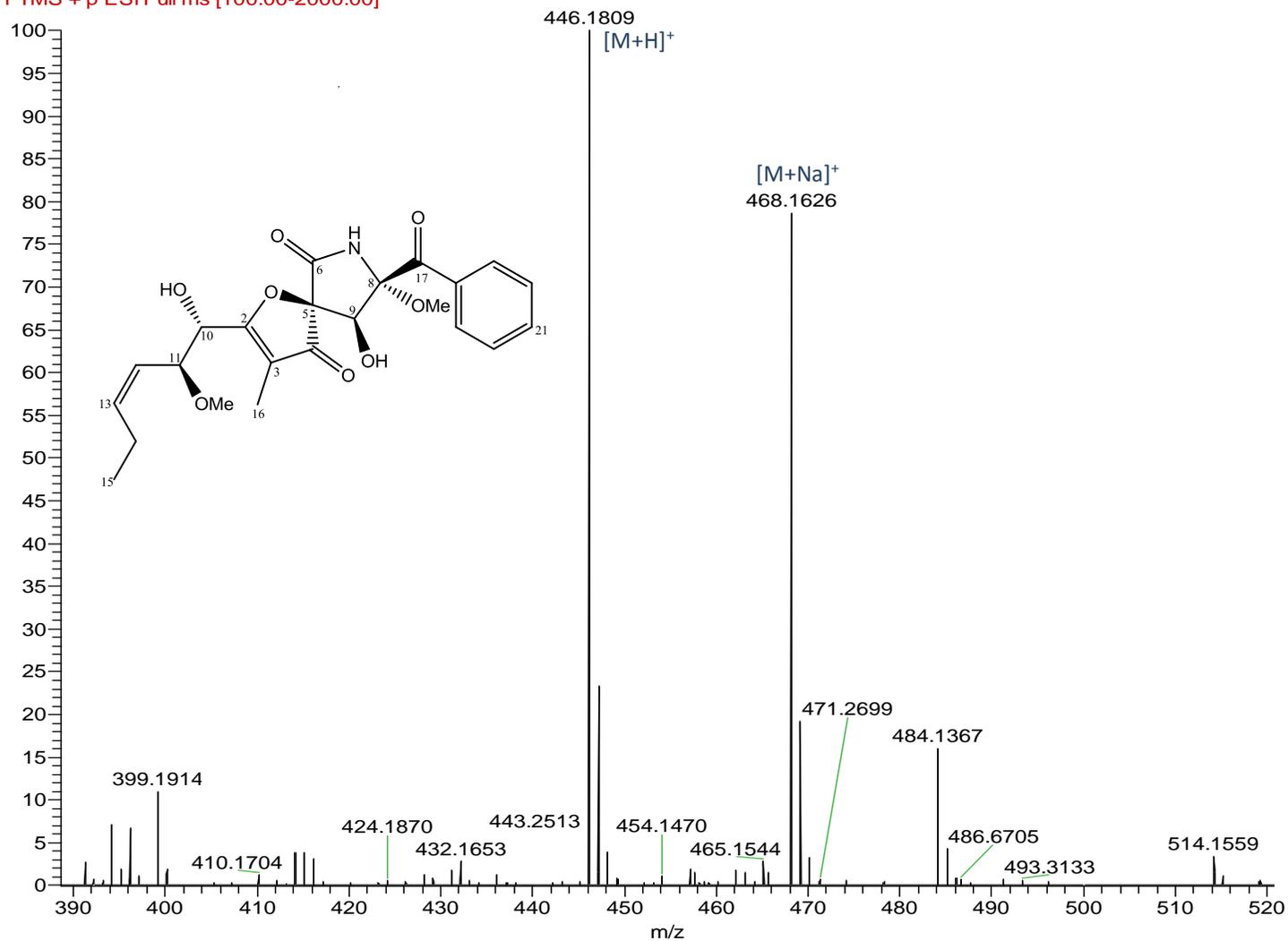


Fig. S9. HRESIMS spectrum of **10**

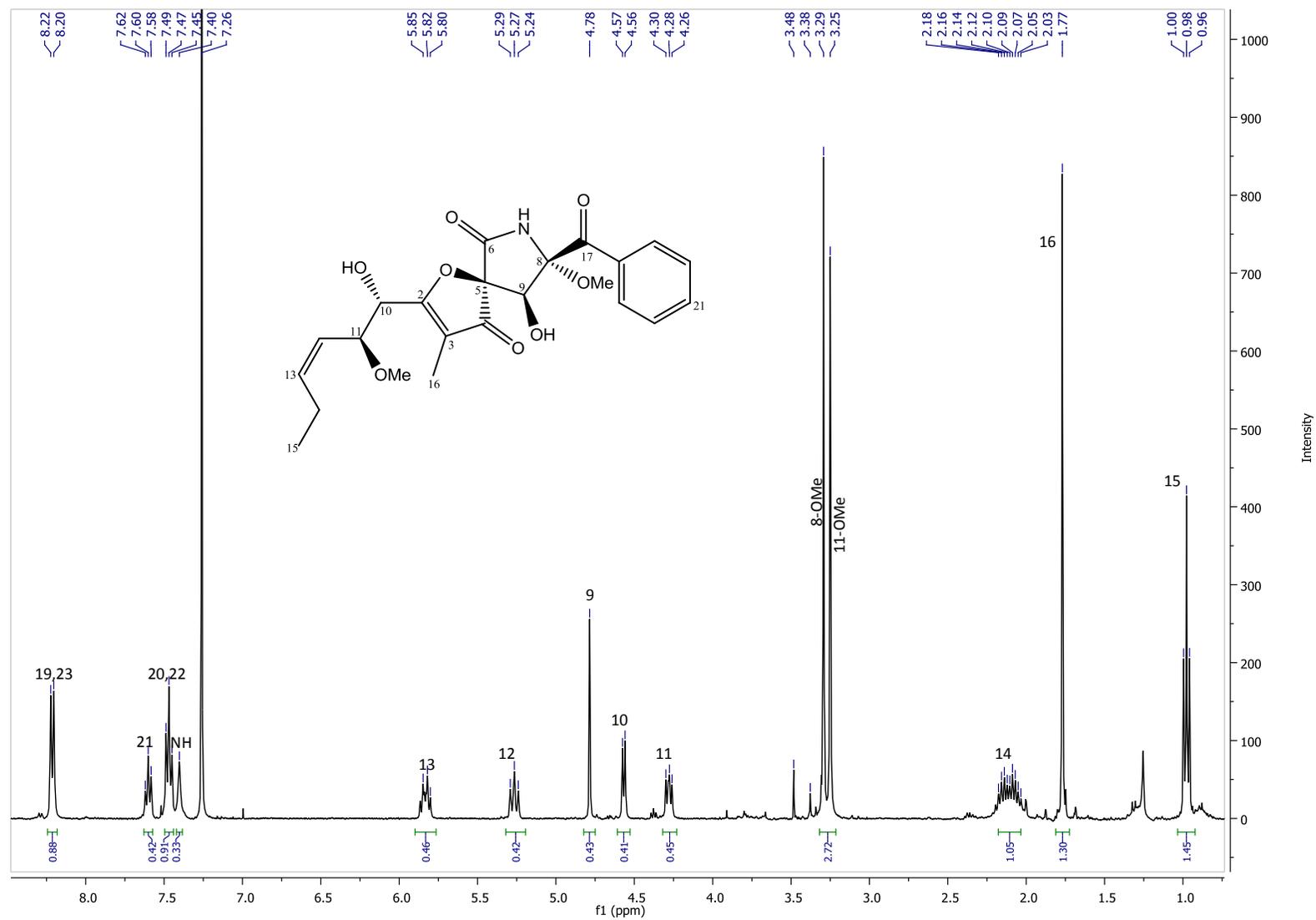


Fig. S10. ^1H NMR spectrum of **10**

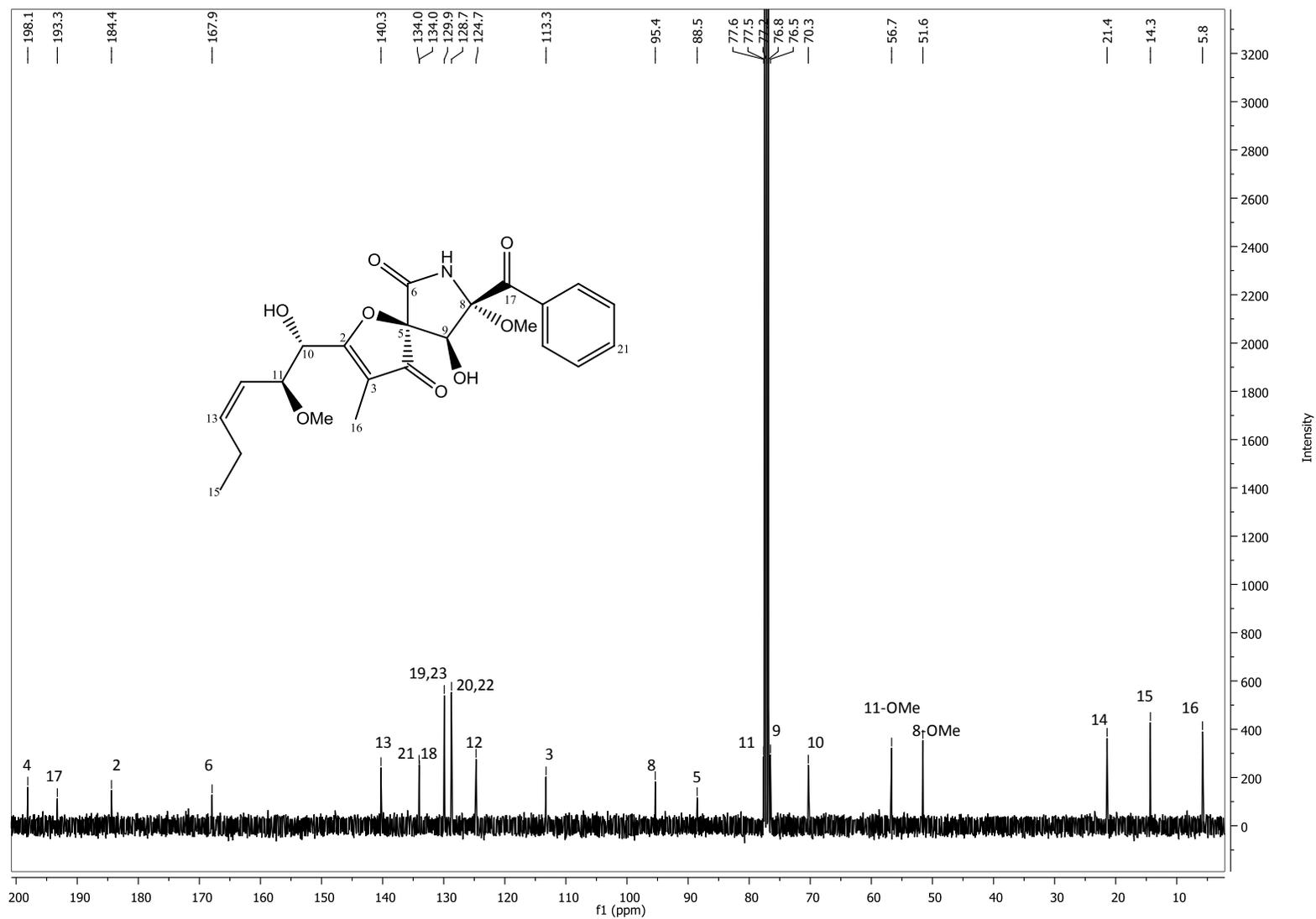


Fig. S11. ^{13}C NMR spectrum of **10**

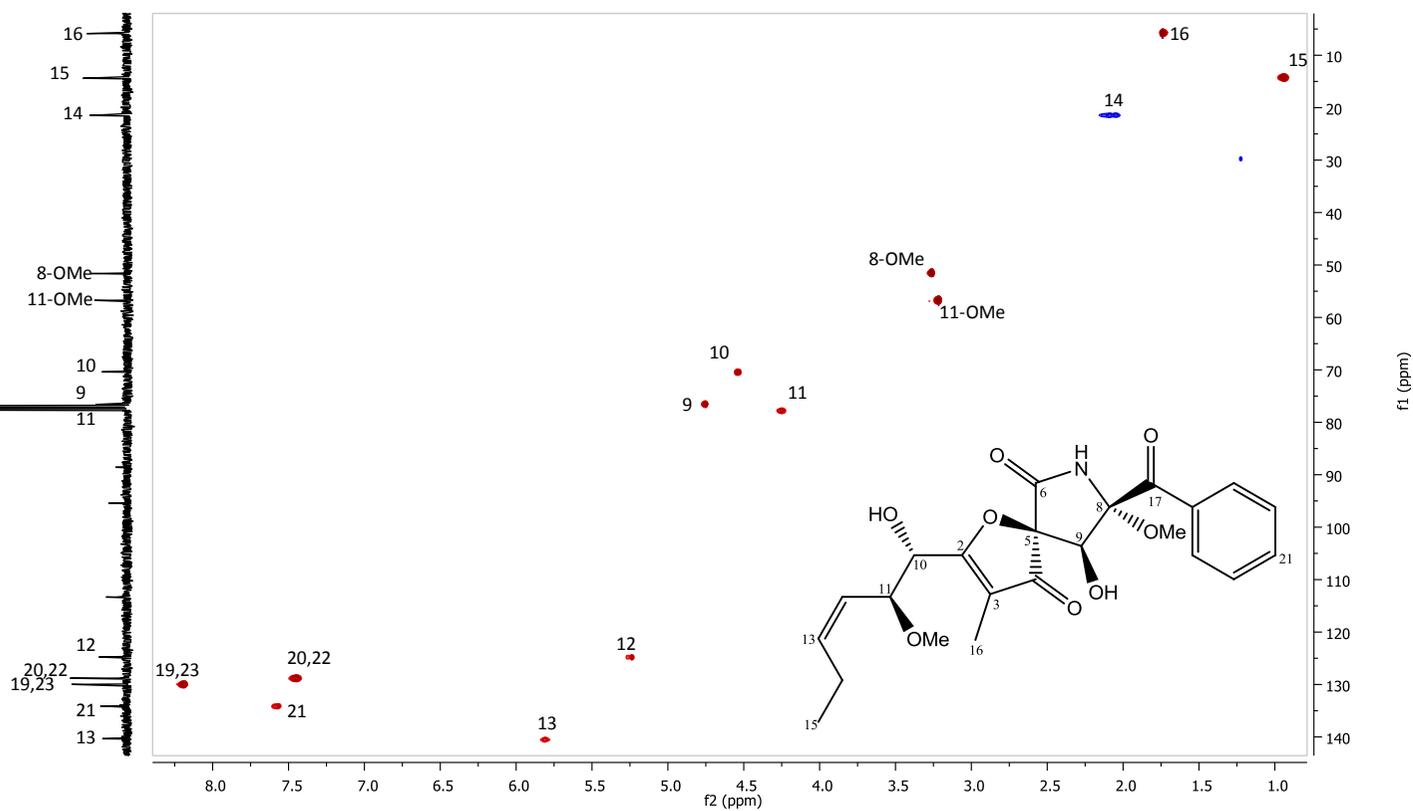
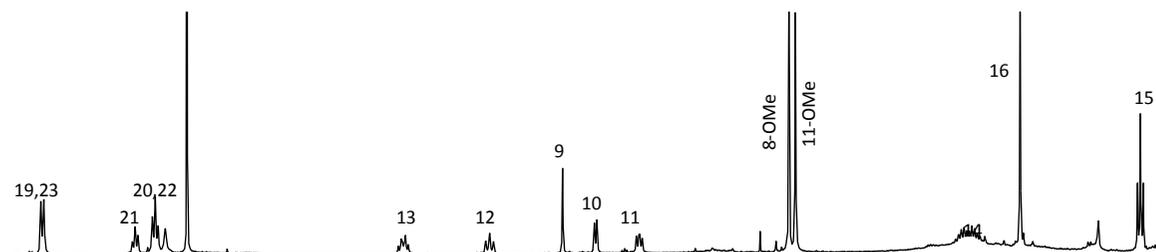


Fig. S12. HSQC spectrum of **10**

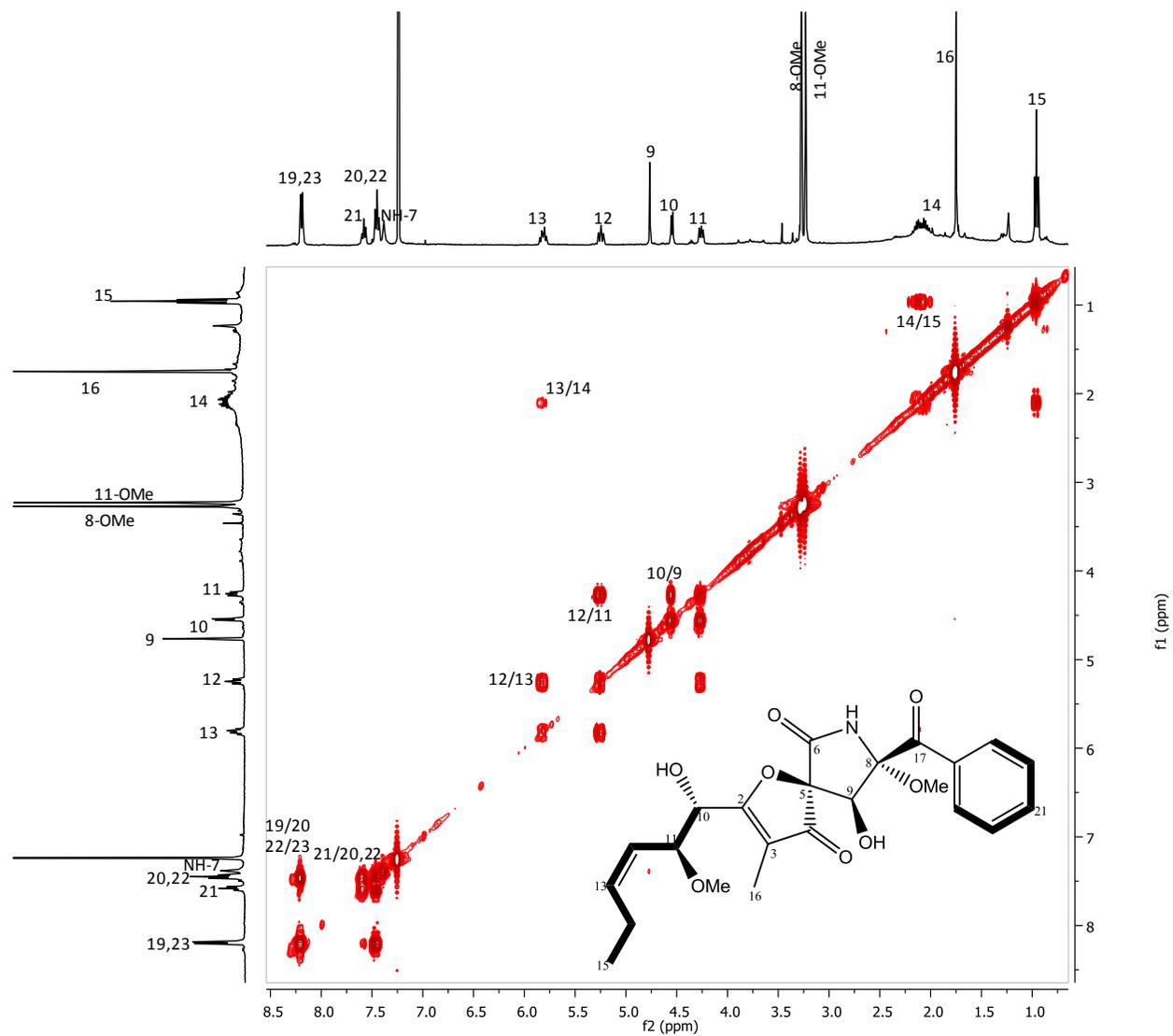


Fig. S13. COSY spectrum of **10**

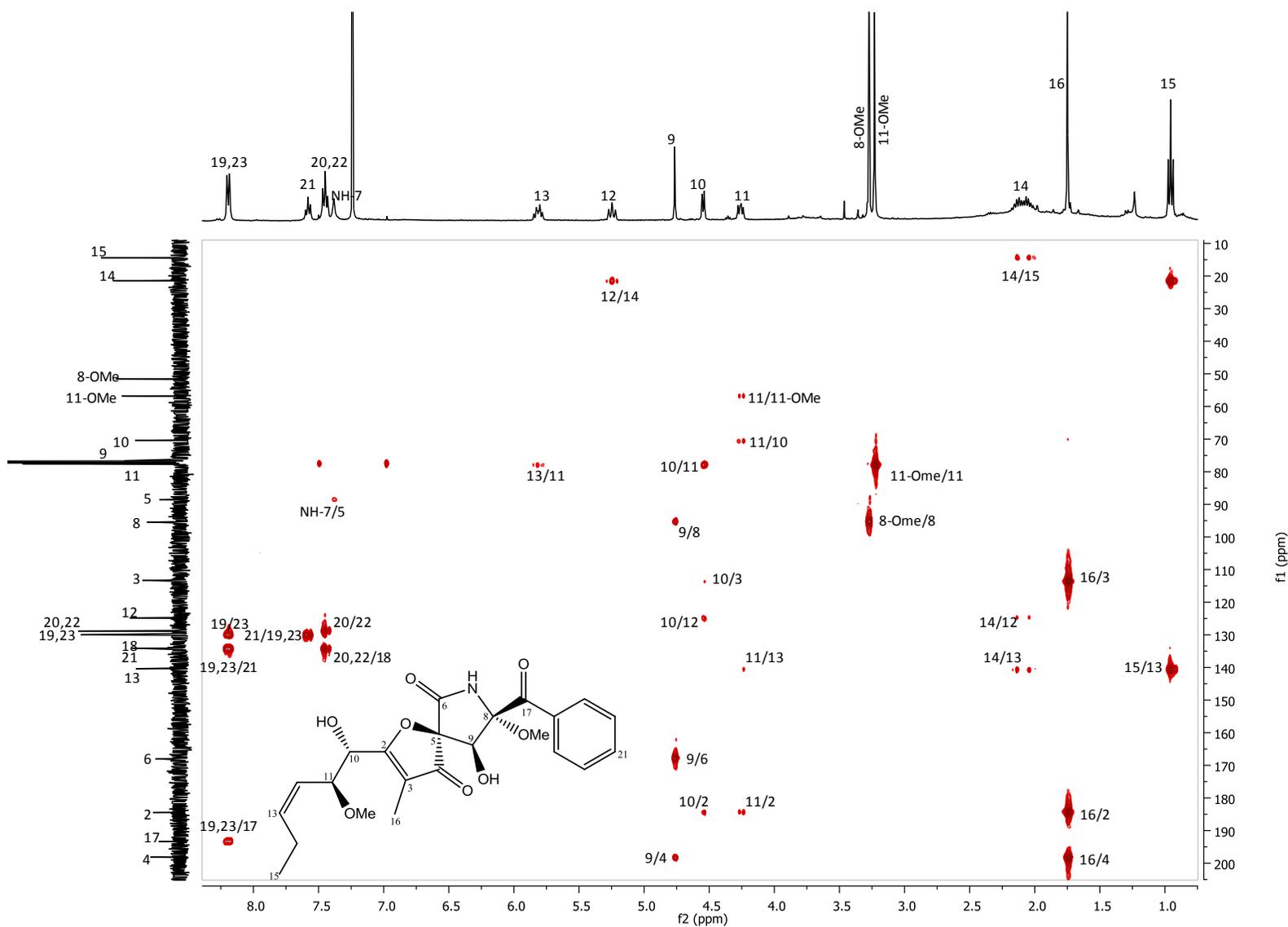


Fig. 14. HMBC spectrum of 10

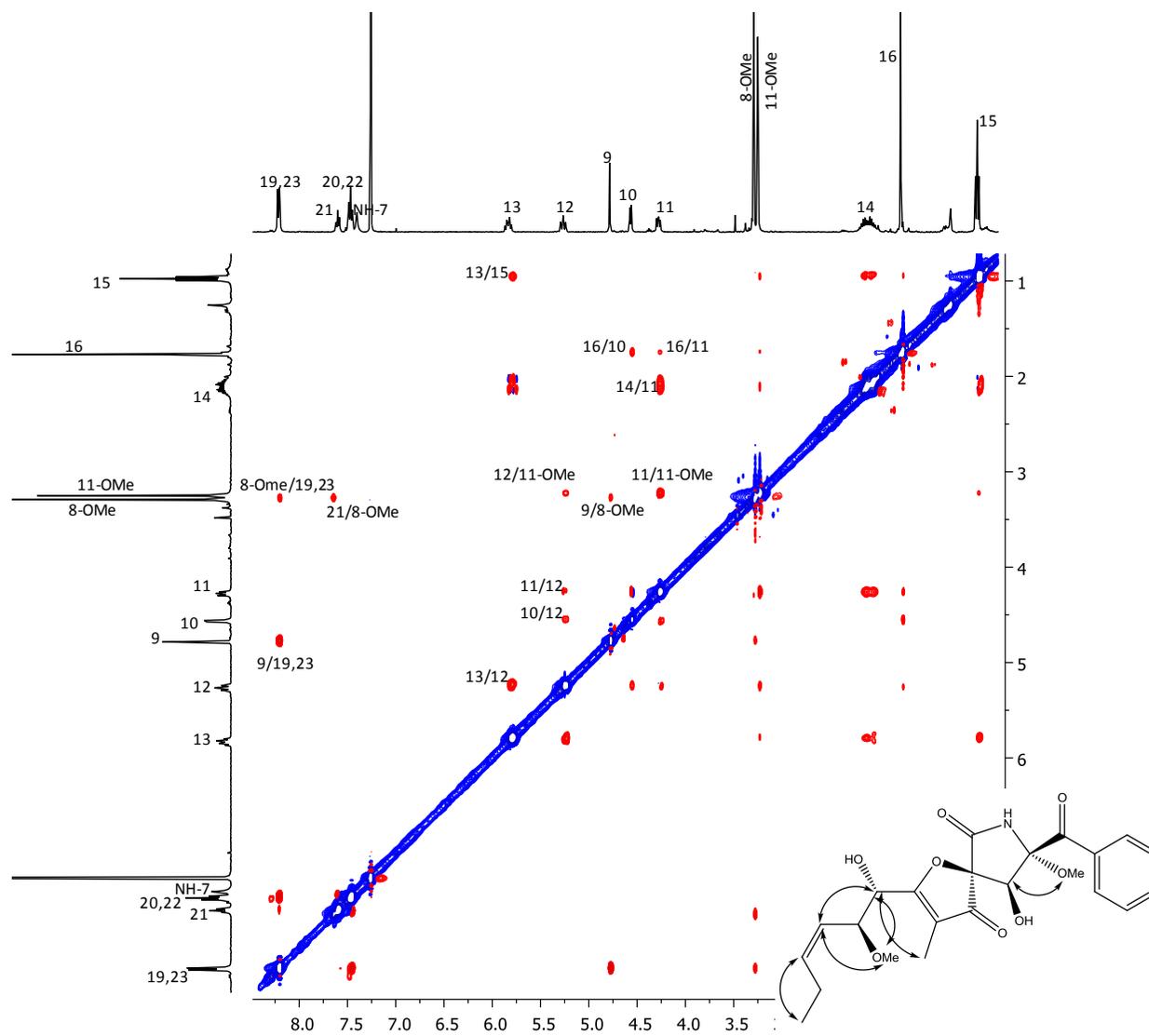


Fig. S15. NOESY spectrum of 10

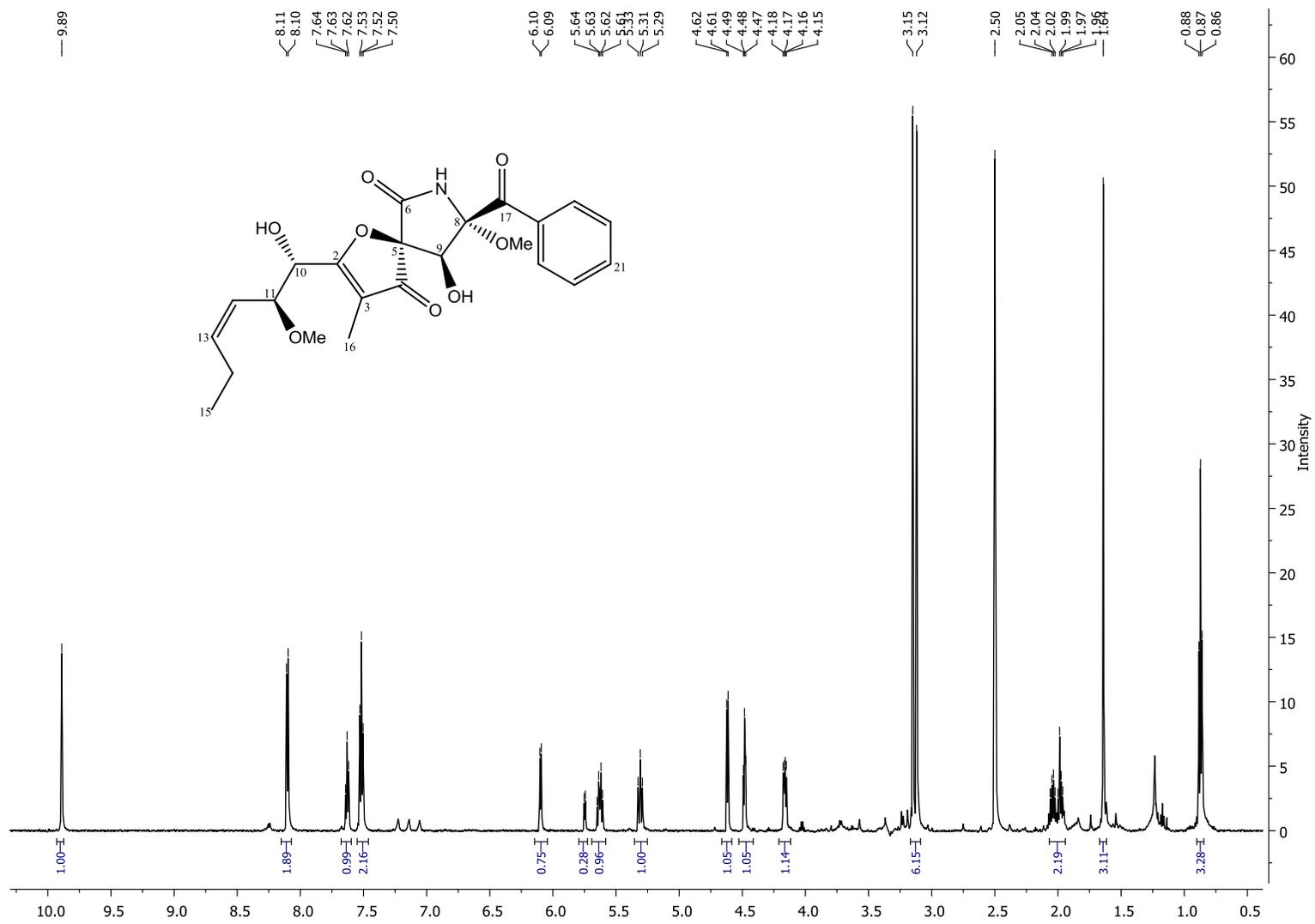


Fig. S16. ¹H NMR spectrum of **10** in DMSO-*d*₆

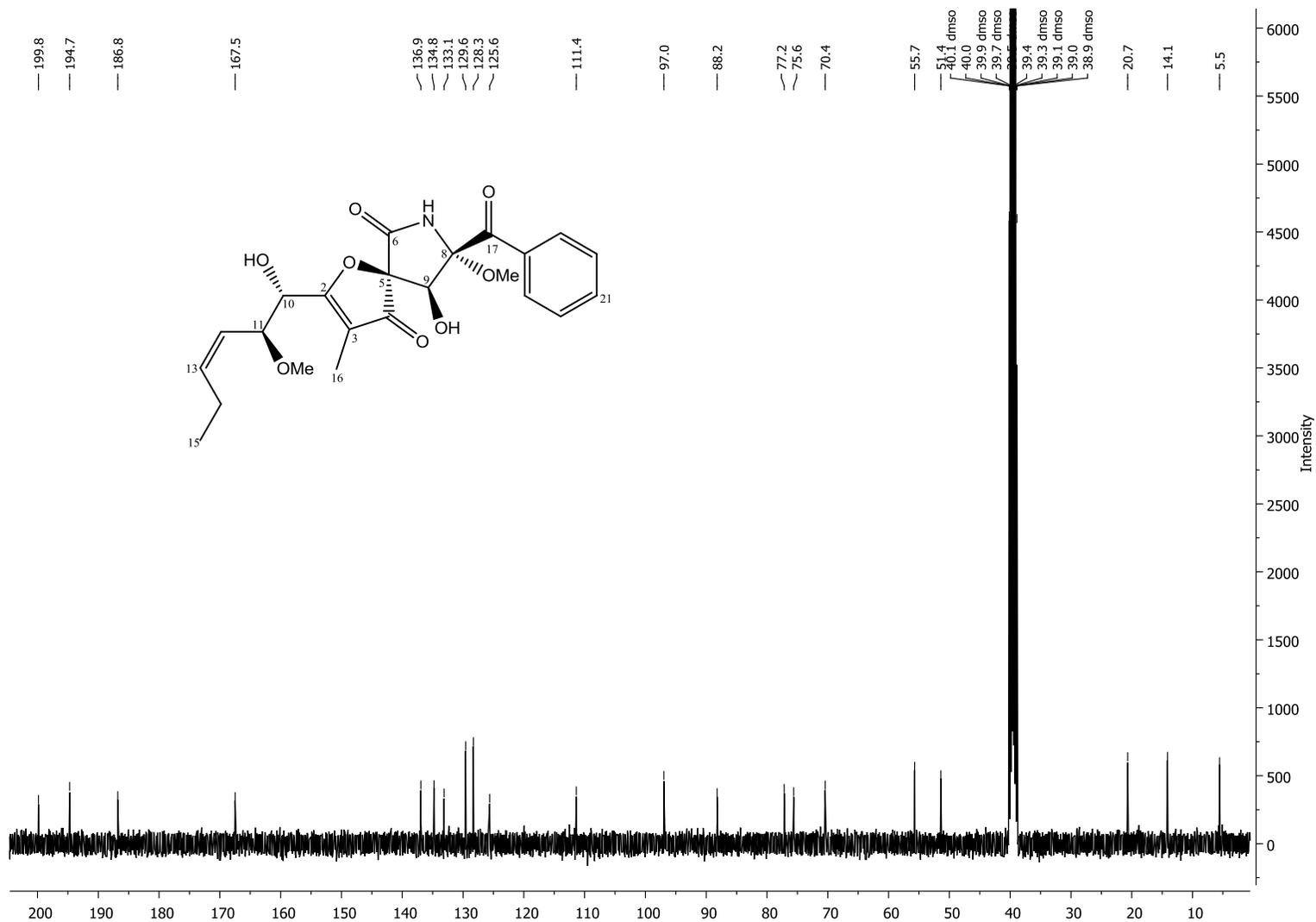


Fig. S17. ^{13}C NMR spectrum of **10** in $\text{DMSO-}d_6$