

“Fruiting Liquid” of Mushroom-Forming Fungi,
A Novel Source of Bioactive Compounds :
Fruiting-Body Inducer and HIF and Axl Inhibitors

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Supporting Information

“Fruiting Liquid” of Mushroom-Forming Fungi; Novel Source of Bioactive Compounds –Fruiting-Body Inducer, and HIF- and Axl-Inhibitor–

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Table of Contents

Figure S1. “Fruiting liquid” of mushroom-forming fungi.

Figure S2. ^1H NMR, ^{13}C NMR and DEPT spectra for compound **1** (CDCl_3).

Figure S3. HMQC spectrum for compound **1** (CDCl_3).

Figure S4. COSY spectrum for compound **1** (CDCl_3).

Figure S5. HMBC spectrum for compound **1** (CDCl_3).

Figure S6. NOESY spectrum for compound **1** (CDCl_3).

Supplementary Table 1. The ^{13}C and ^1H of **1** accordance.

Figure S7. ^1H NMR, ^{13}C NMR and DEPT spectra for compound **2** (CD_3OD).

Figure S8. HMQC spectrum for compound **2** (CD_3OD).

Figure S9. COSY spectrum for compound **2** (CD_3OD).

Figure S10. HMBC spectrum for compound **2** (CD_3OD).

Figure S11. ^1H NMR, ^{13}C NMR and DEPT spectra for compound **3** (CD_3OD).

Figure S12. HMQC spectrum for compound **3** (CD_3OD).

Figure S13. COSY spectrum for compound **3** (CD_3OD).

Figure S14. HMBC spectrum for compound **3** (CD_3OD).

Figure S15. ^1H NMR, ^{13}C NMR and DEPT spectra for compound **4** (CD_3OD).

Figure S16. HMQC spectrum for compound **4** (CD_3OD).

Figure S17. COSY spectrum for compound **4** (CD_3OD).

Figure S18. HMBC spectrum for compound **4** (CD_3OD).

Figure S19. ^1H NMR and ^{13}C NMR spectra for compound **6** (CDCl_3).

Figure S20. ^1H NMR and ^{13}C NMR spectra for compound **7** (CDCl_3).

Figure S21. ^1H NMR and ^{13}C NMR spectra for compound **10** (CDCl_3).

Figure S22. ^1H NMR spectra for natural **3** and synthetic **3** (CD_3OD).

Figure S23. ^{13}C NMR spectra for natural **3** and synthetic **3** (CD_3OD).

Figure S24. ^1H NMR spectra for compounds **3** and **11** (CD_3OD).

Figure S25. ^{13}C NMR spectra for compounds **3** and **11** (CD_3OD).

Figure S26. ^1H NMR spectra for compounds **4** and **12** (CD_3OD).

Figure S27. ^{13}C NMR spectra for compounds **4** and **12** (CD_3OD).

Supplementary Table 2. Crystallographic data of **2** and **3**.

Figure S28. Effect of compounds **2** and **3** on HIF activation *in vitro*.

Figure S29. Effect of compounds **2** to **4** on expressions of Axl and immune checkpoints (PD-L1 and PD-L2) on lung cancer cell line A549 cells.

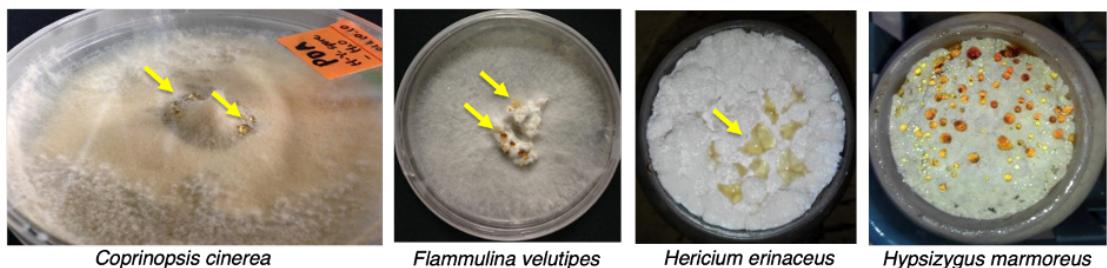


Figure S1. “Fruiting liquid” of mushroom-forming fungi.

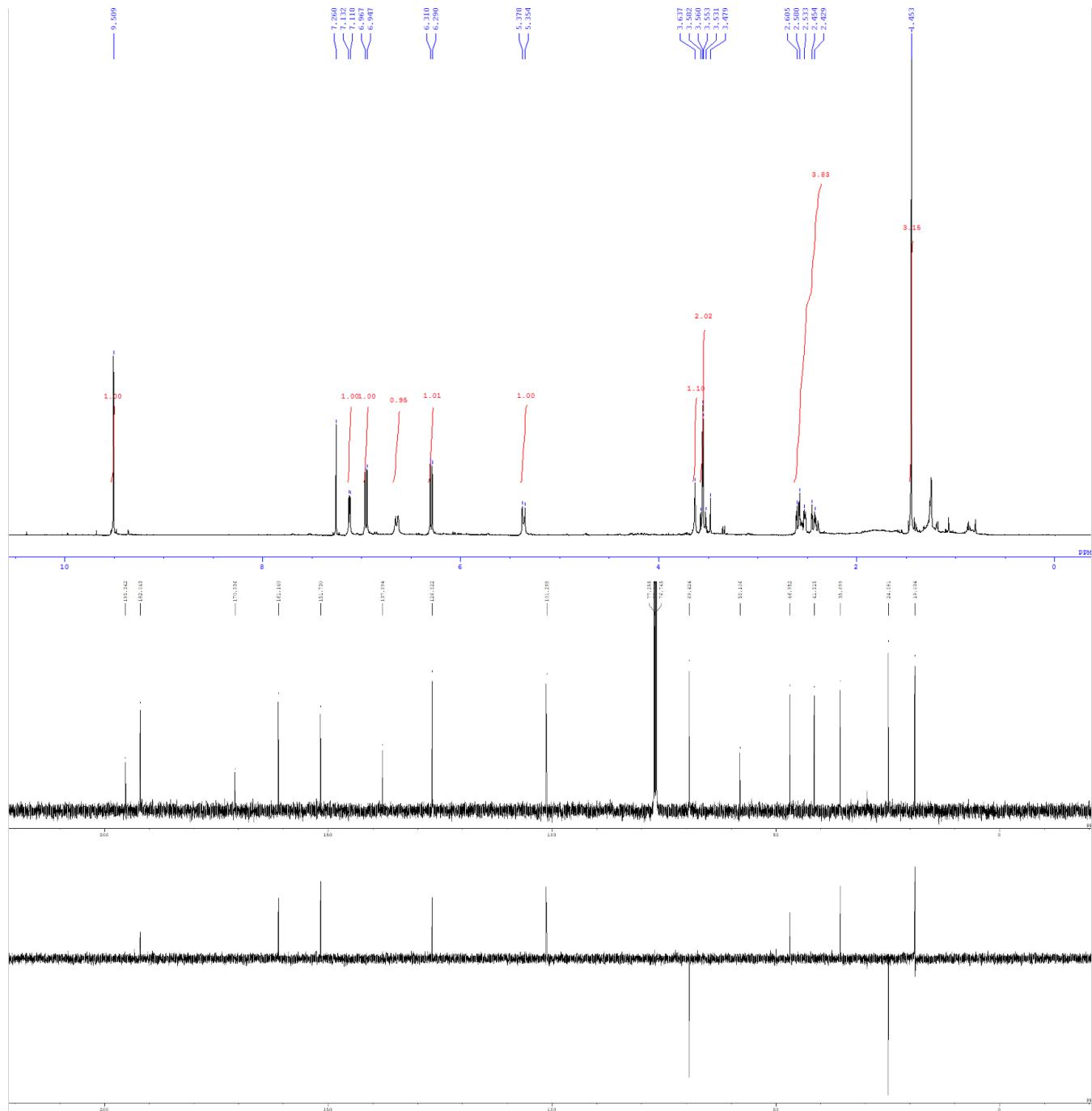


Figure S2. ¹H NMR, ¹³C NMR and DEPT spectra for compound 1 (CDCl₃).

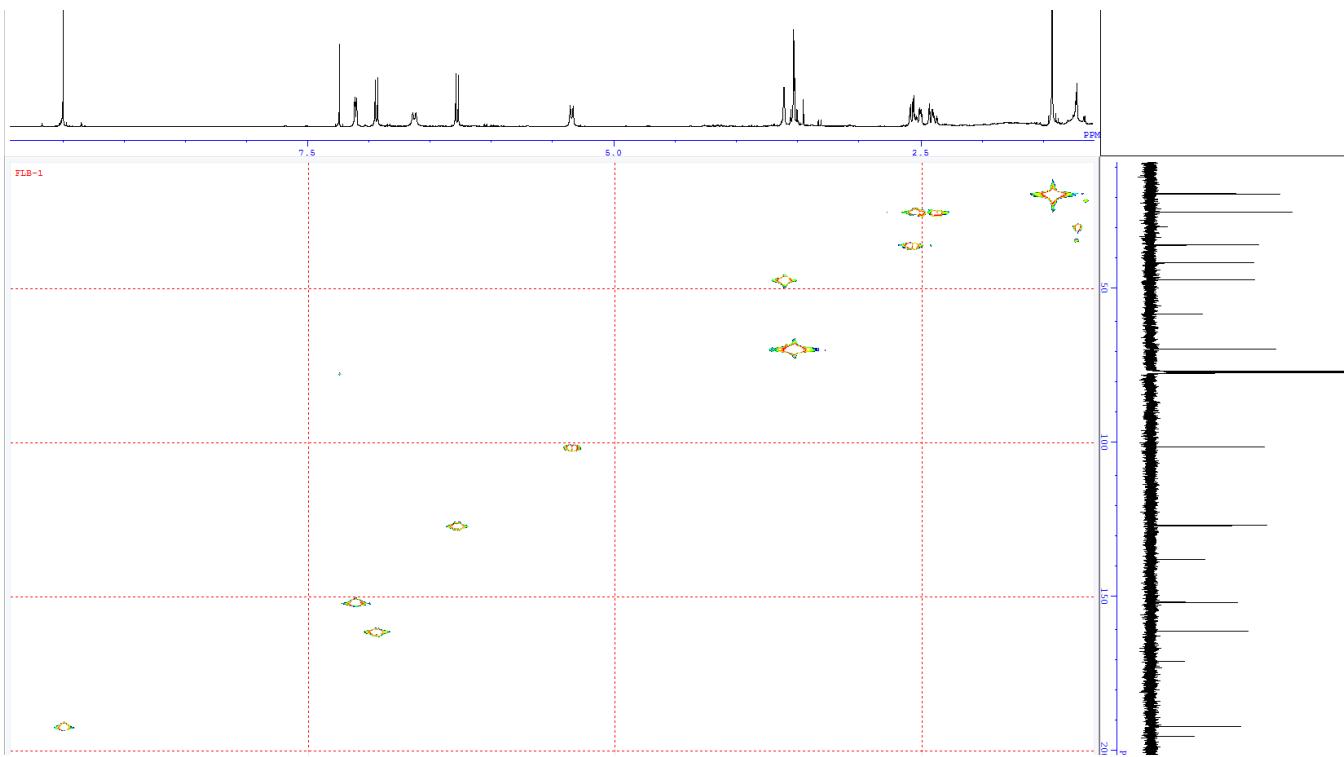


Figure S3. HMQC spectrum for compound 1 (CDCl_3).

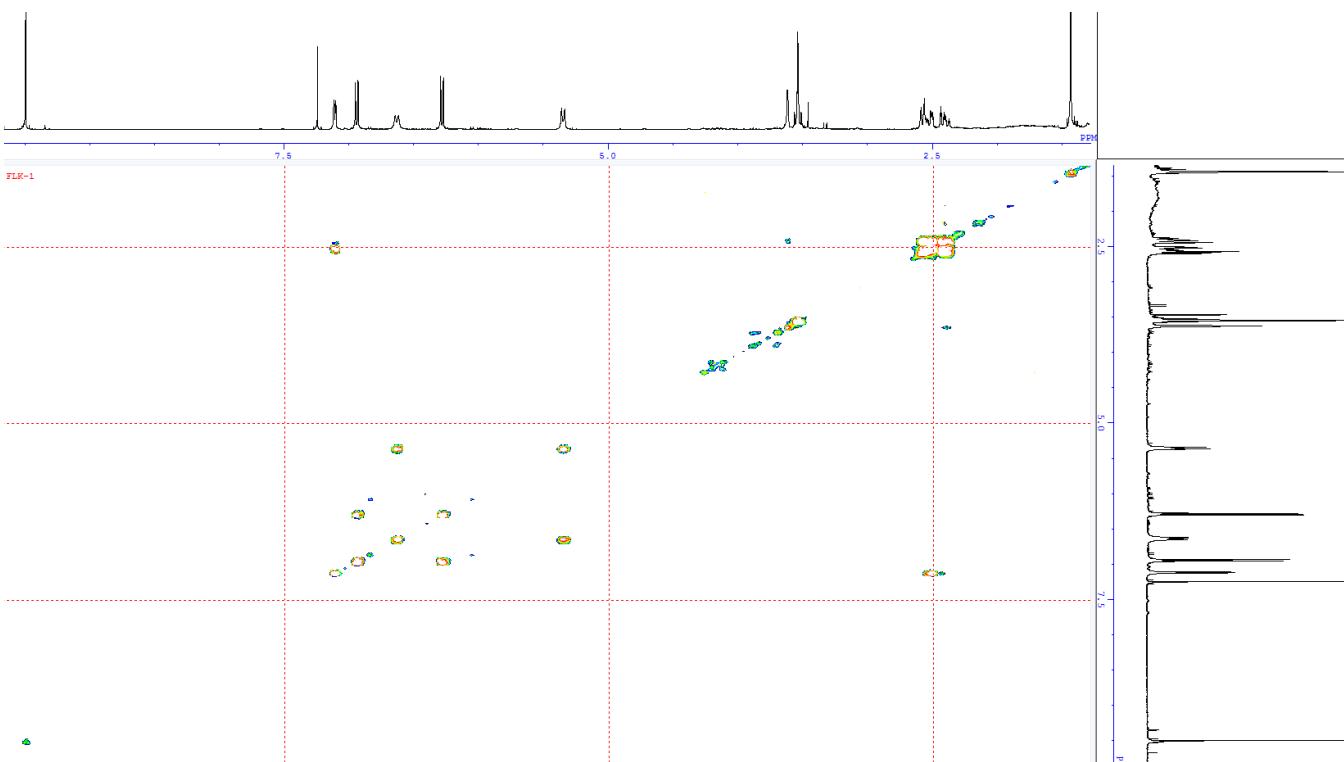


Figure S4. COSY spectrum for compound 1 (CDCl_3).

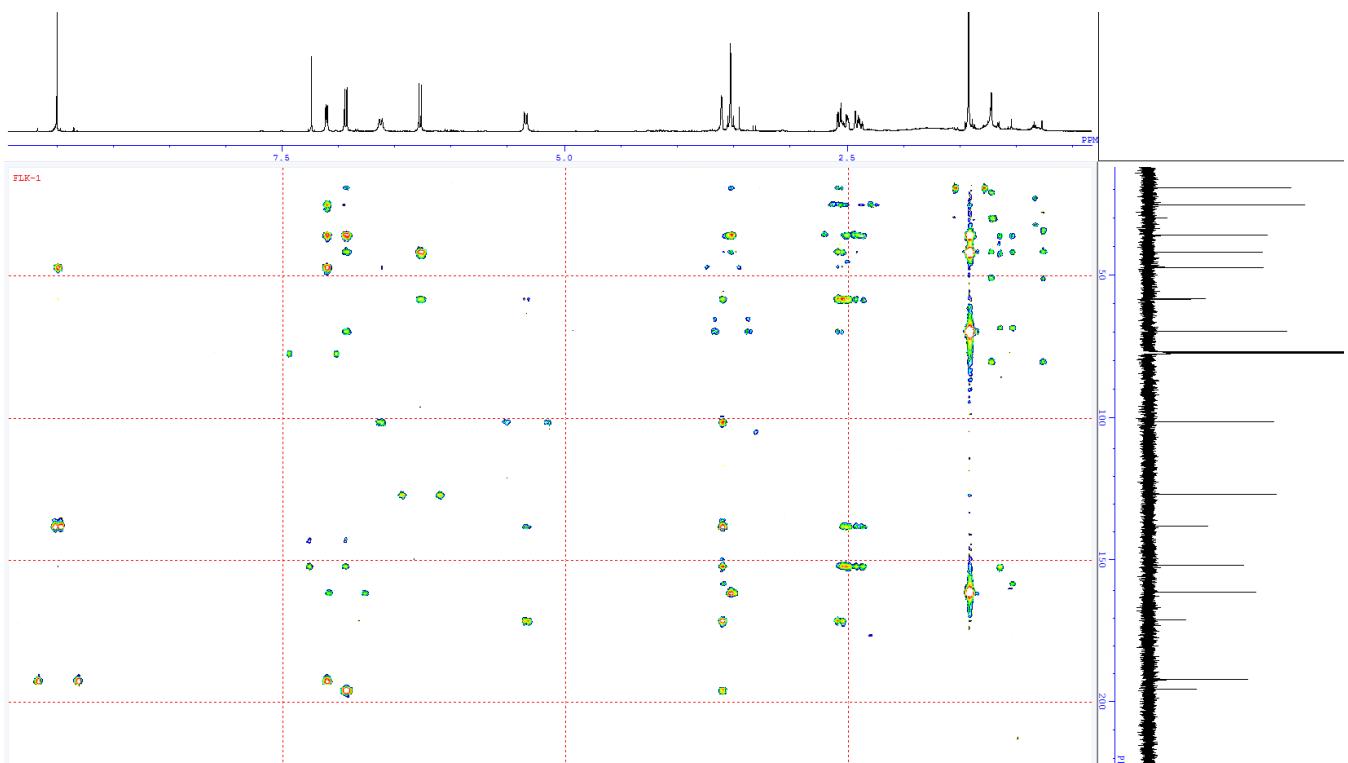


Figure S5. HMBC spectrum for compound **1** (CDCl_3).

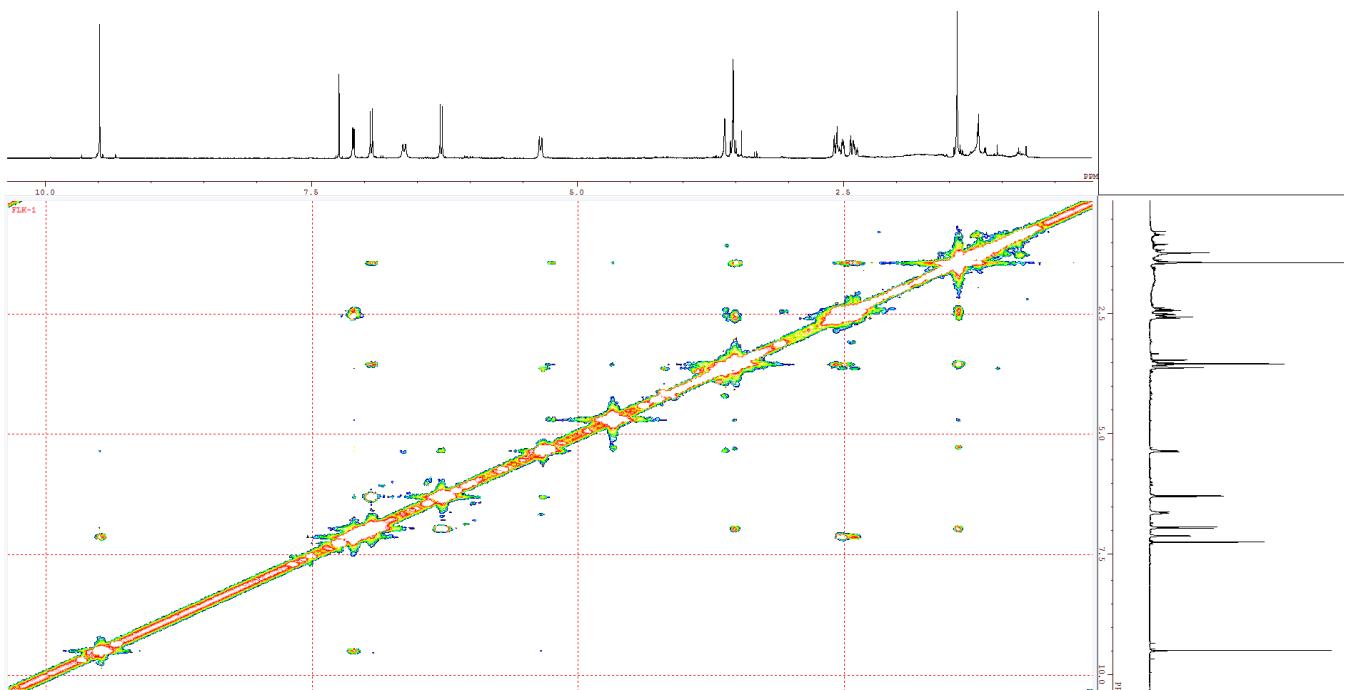
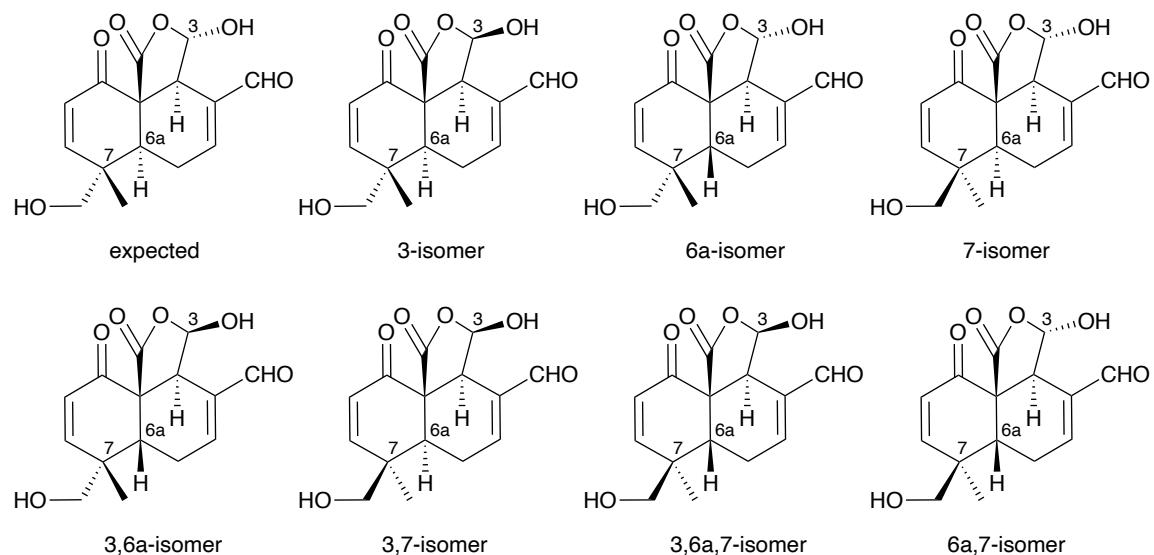


Figure S6. NOESY spectrum for compound **1** (CDCl_3).

Table S1. The ^{13}C and ^1H of **1** accordance.



	^{13}C			^1H			$^{13}\text{C}+^1\text{H}$
	RMSD (ppm)	max absolute (ppm)	DP4	RMSD (ppm)	max absolute (ppm)	DP4	DP4
expected	1.6	3.1	94.1%	0.15	0.28	100.0%	100.0%
3-isomer	2.1	5.0	5.9%	0.40	1.23	0.0%	0.0%
6a-isomer	3.8	12.2	0.0%	0.29	0.58	0.0%	0.0%
7-isomer	3.8	9.2	0.0%	0.31	0.57	0.0%	0.0%
3,6a-isomer	3.1	6.8	0.0%	0.42	1.13	0.0%	0.0%
3,7-isomer	3.6	9.0	0.0%	0.55	1.50	0.0%	0.0%
3,6a,7-isomer	4.4	7.9	0.0%	0.41	0.93	0.0%	0.0%
6a,7-isomer	3.0	6.6	0.0%	0.32	0.76	0.0%	0.0%

RMSD: root mean square of deviation

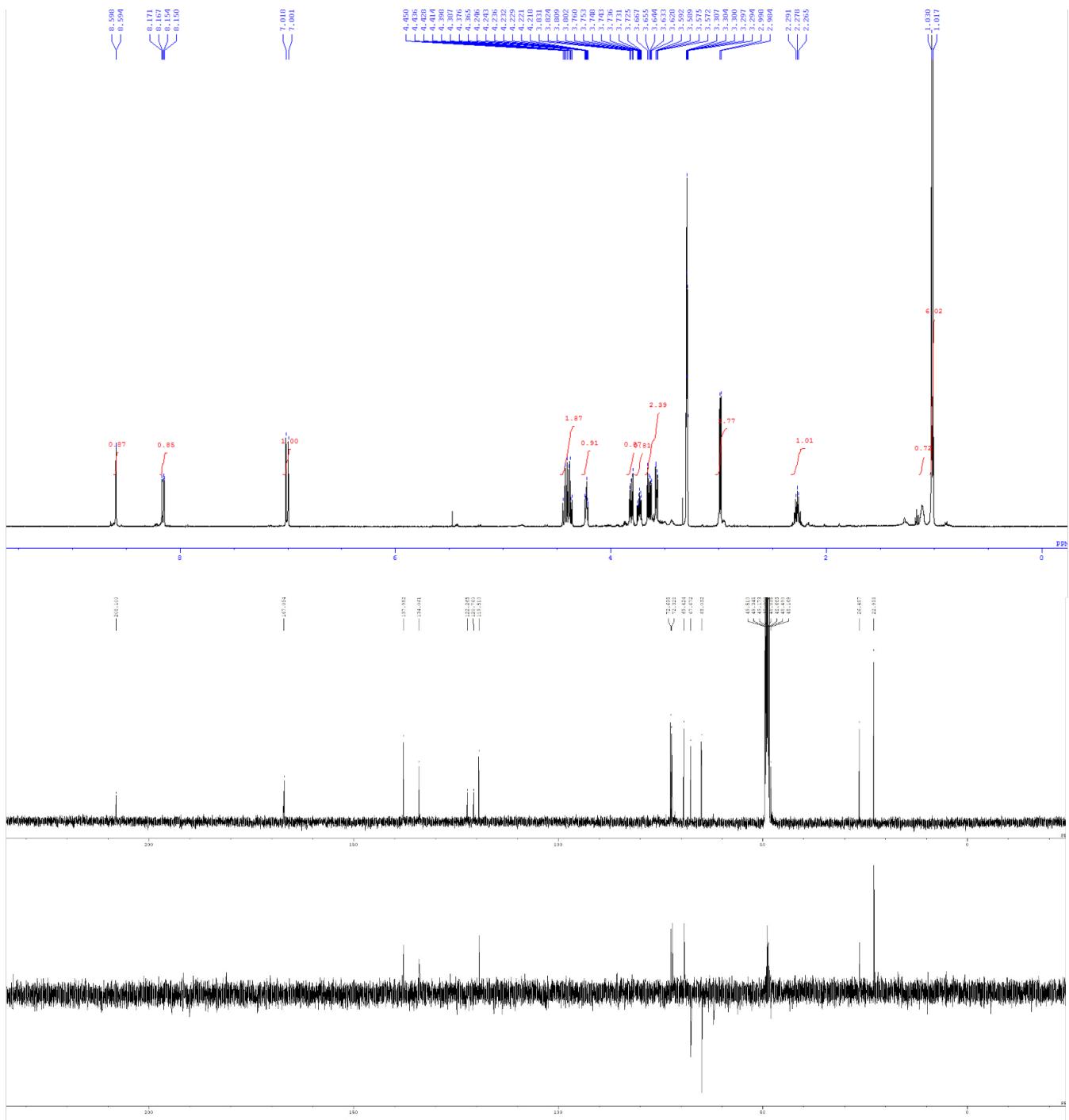


Figure S7. ^1H NMR, ^{13}C NMR and DEPT spectra for compound **2** (CD_3OD).

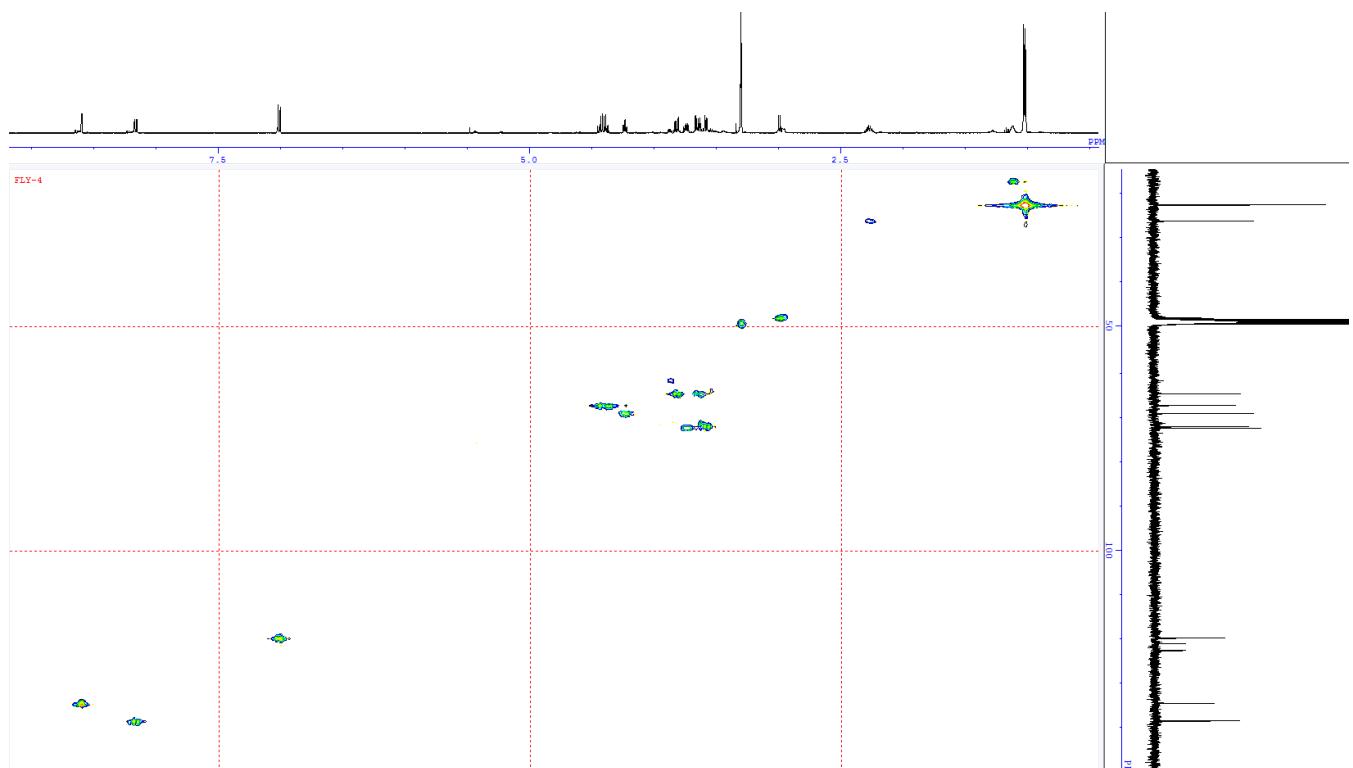


Figure S8. HMQC spectrum for compound **2** (CD_3OD).

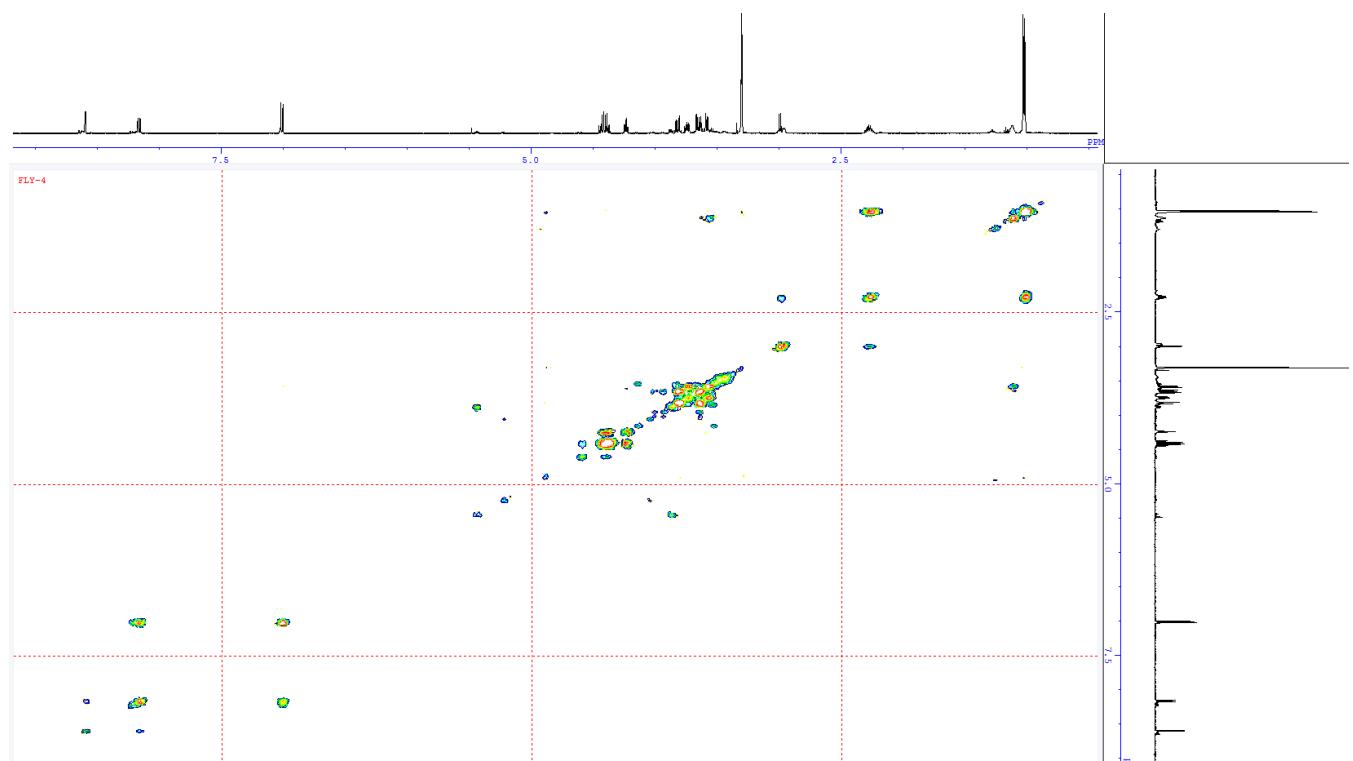


Figure S9. COSY spectrum for compound **2** (CD_3OD).

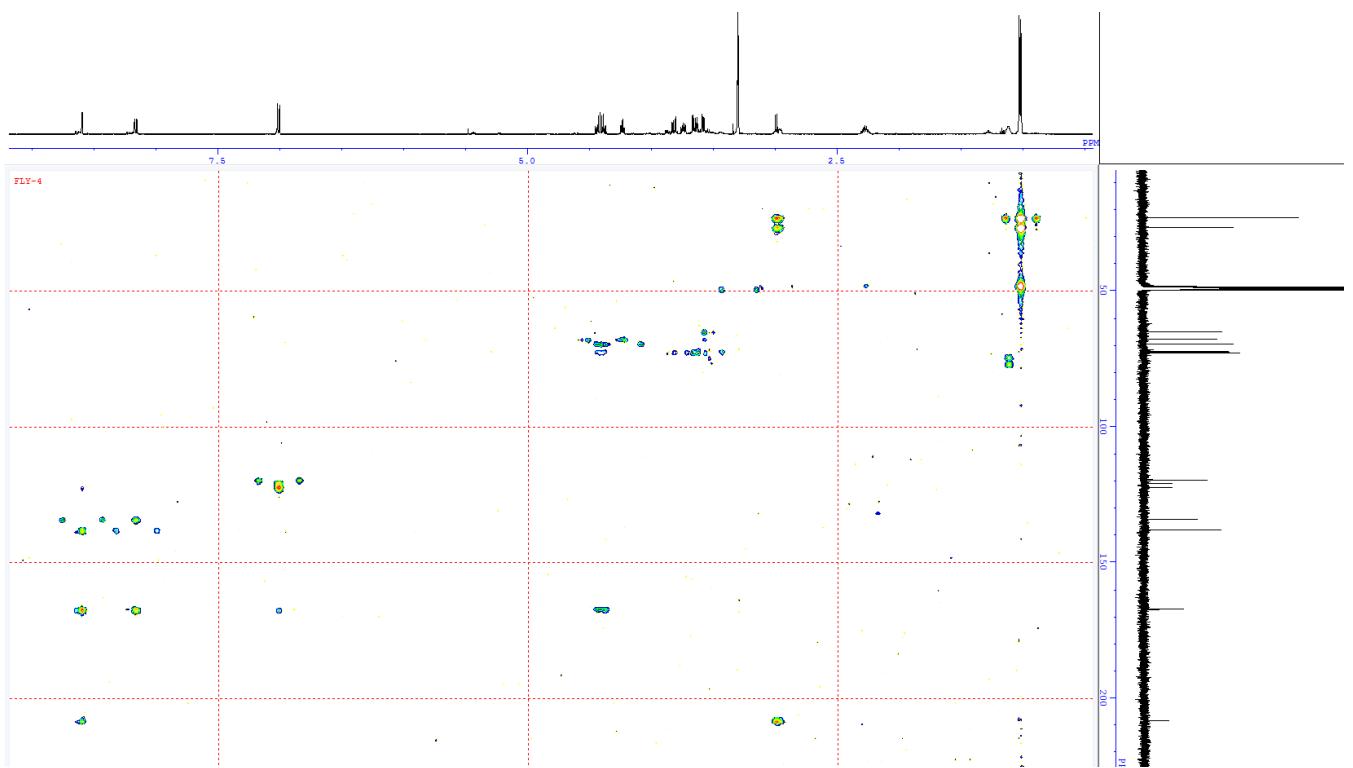


Figure S10. HMBC spectrum for compound **2** (CD_3OD).

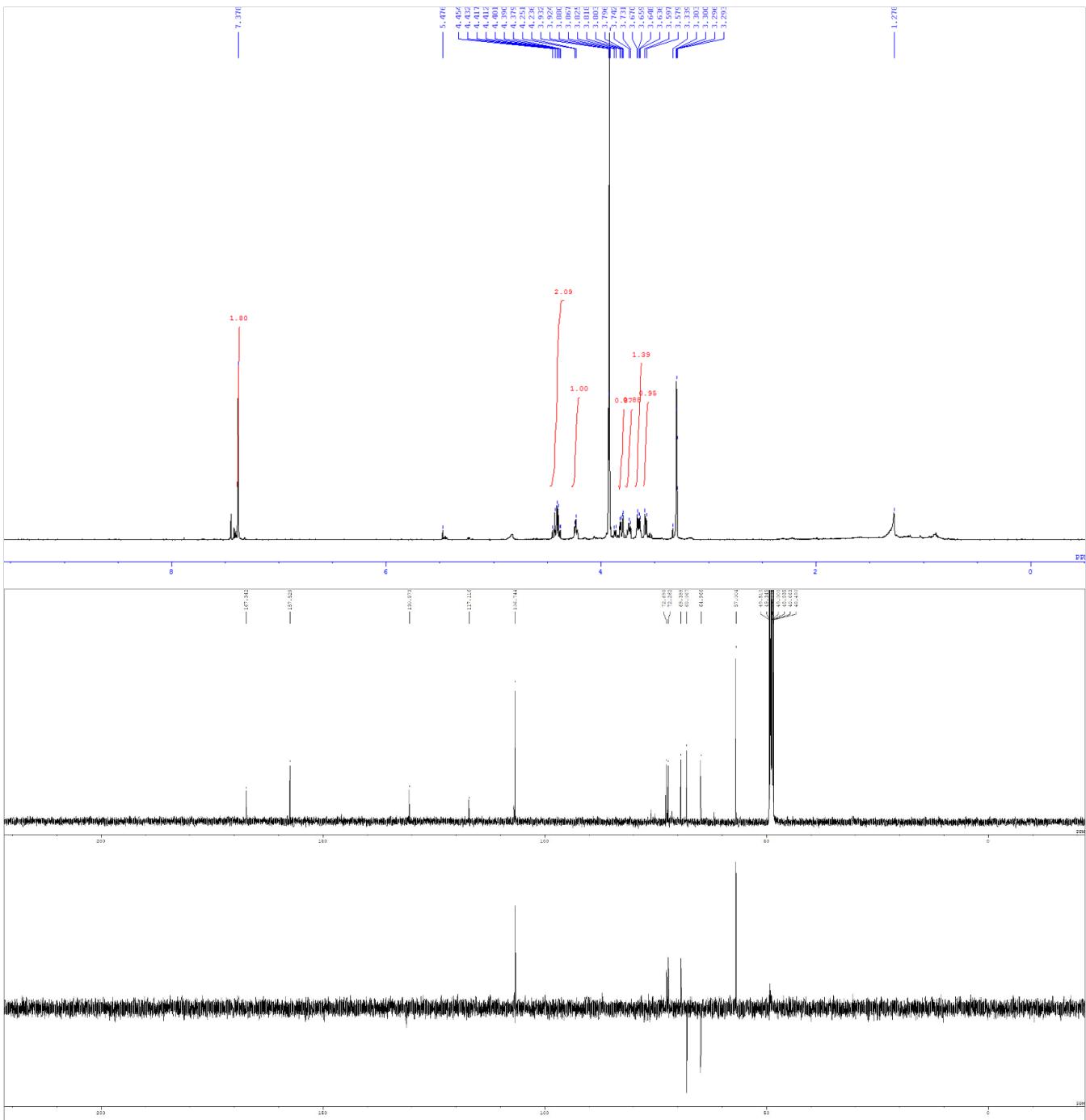


Figure S11. ^1H NMR, ^{13}C NMR and DEPT spectra for compound 3 (CD_3OD).

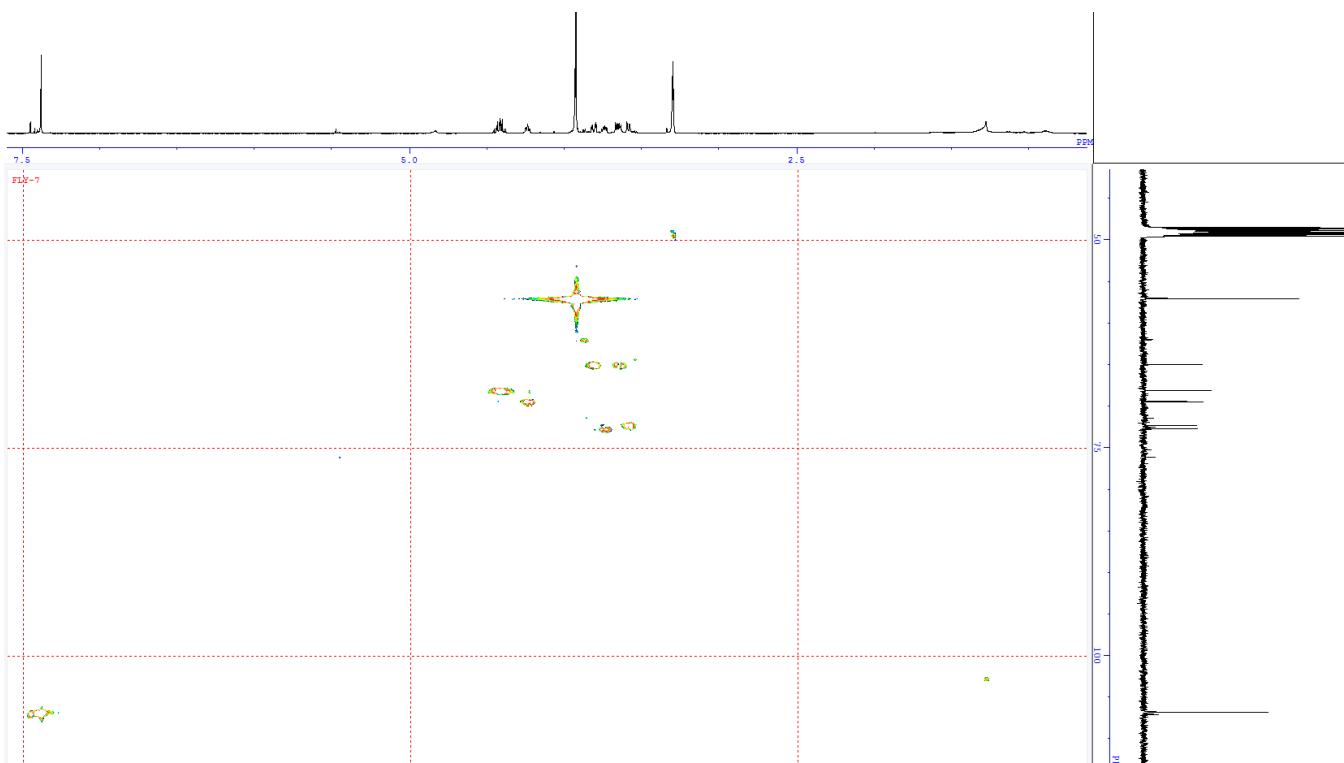


Figure S12. HMQC spectrum for compound 3 (CD_3OD).

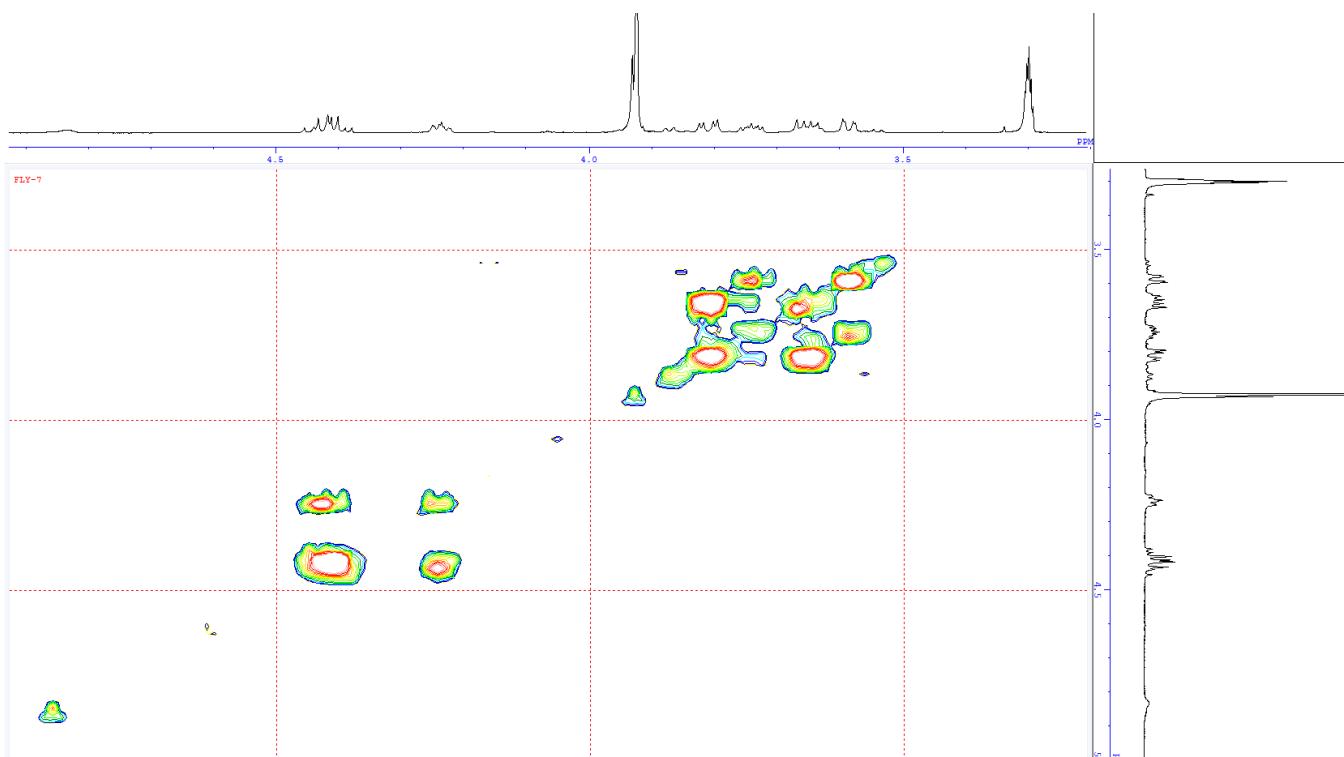


Figure S13. COSY spectrum for compound 3 (CD_3OD).

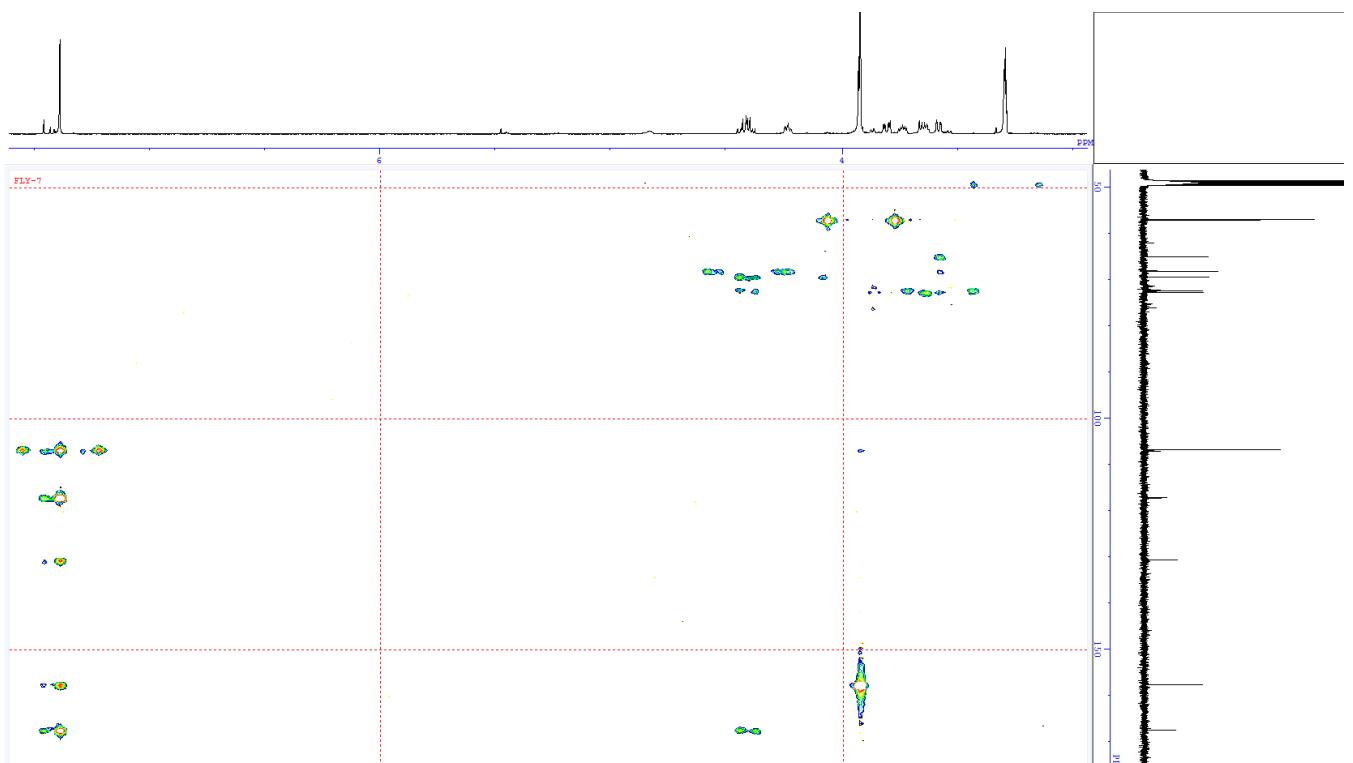


Figure S14. HMBC spectrum for compound 3 (CD_3OD).

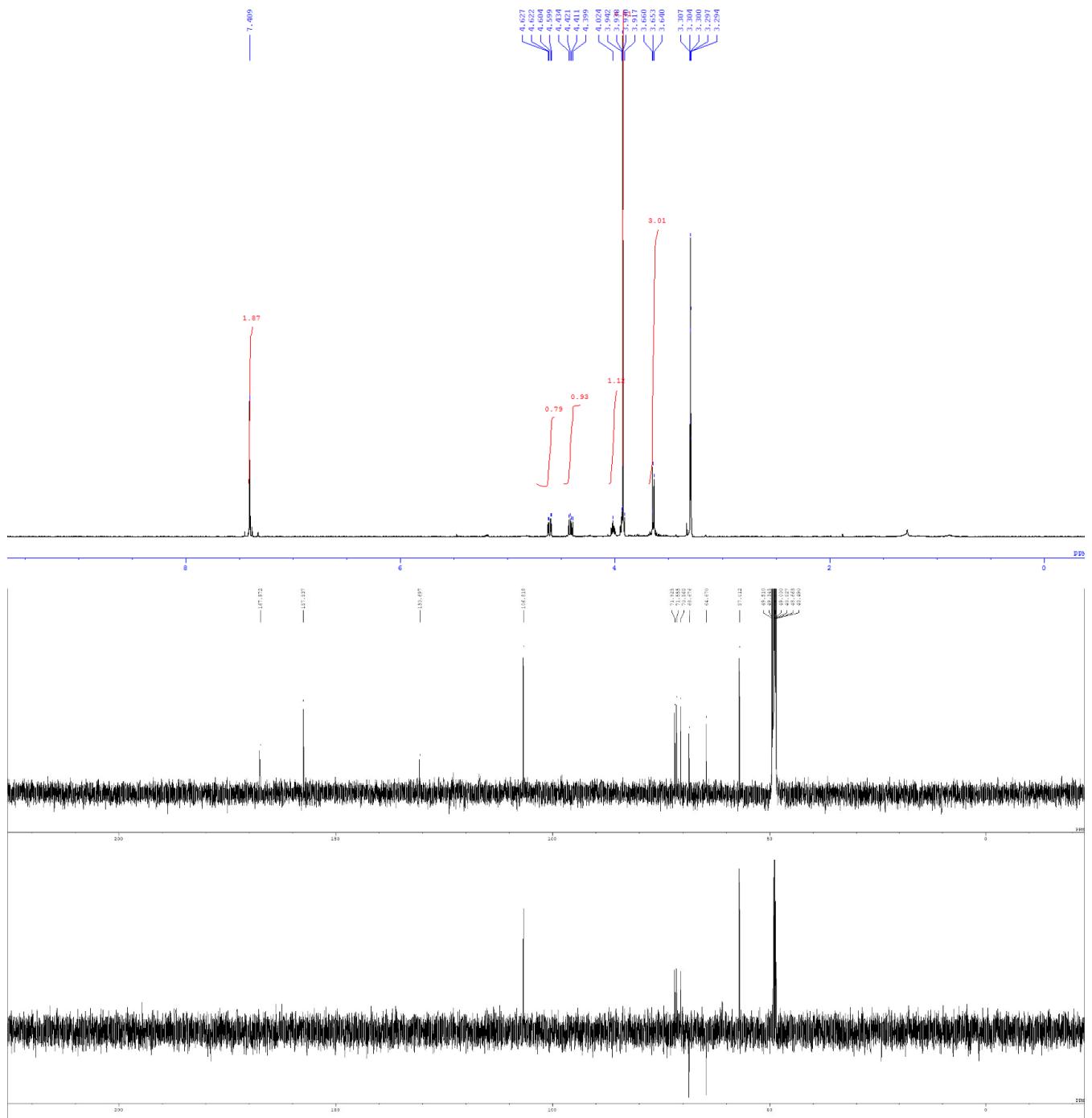


Figure S15. ^1H NMR, ^{13}C NMR and DEPT spectra for compound 4 (CD_3OD).

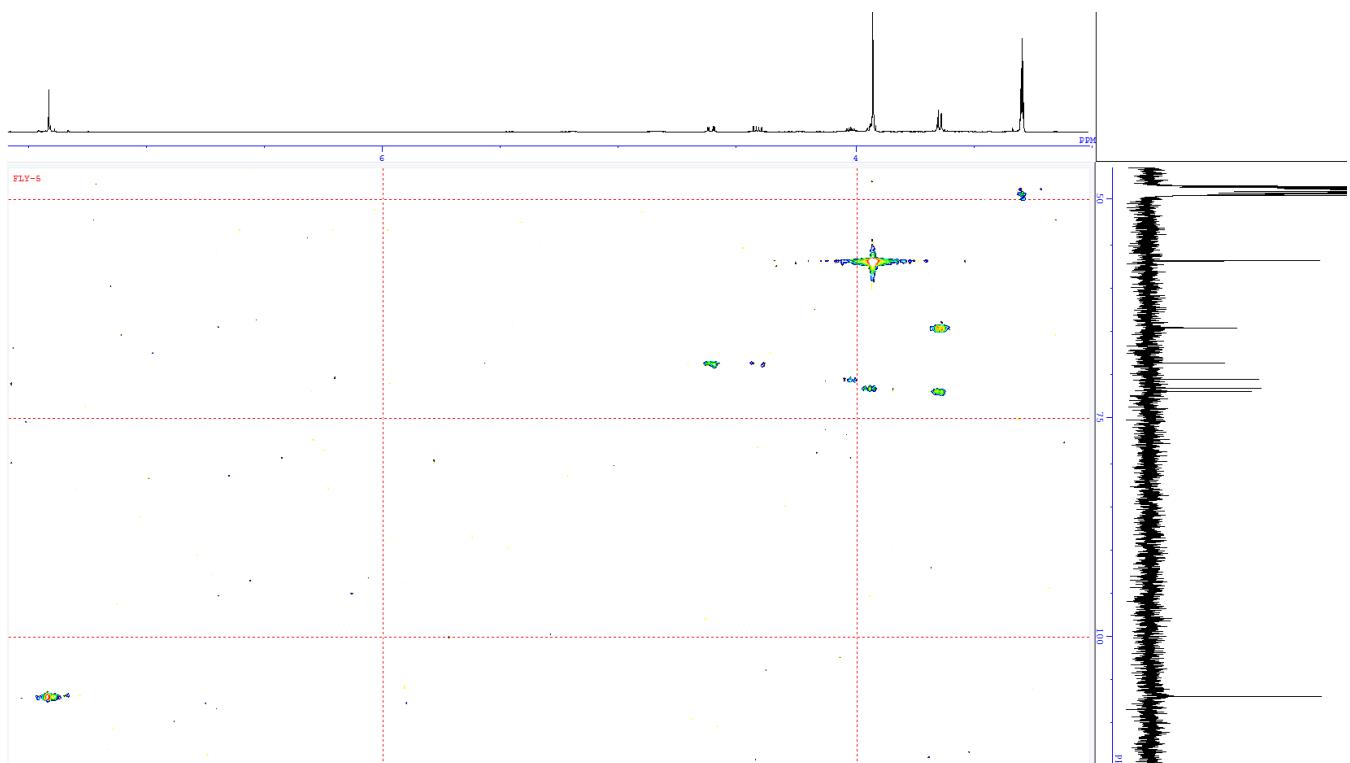


Figure S16. HMQC spectrum for compound **4** (CD_3OD).

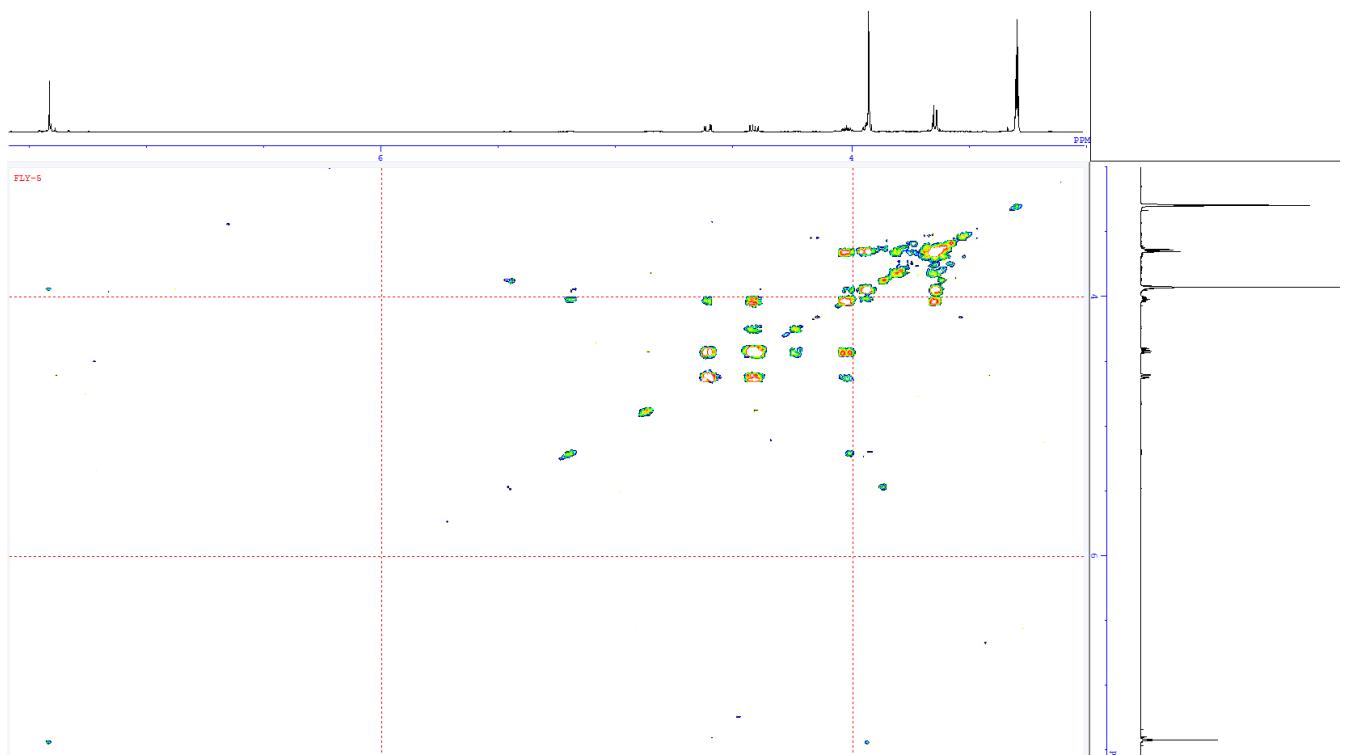


Figure S17. COSY spectrum for compound **4** (CD_3OD).

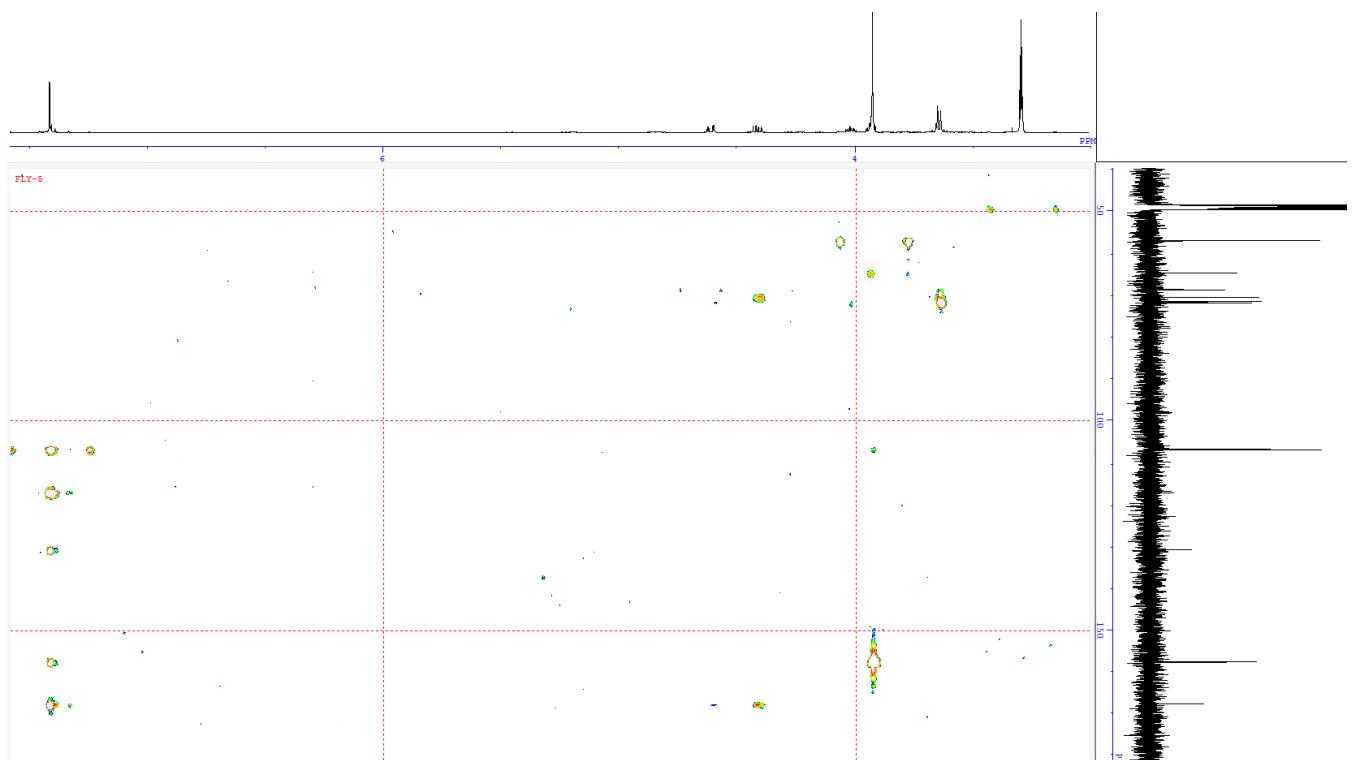


Figure S18. HMBC spectrum for compound 4 (CD_3OD).

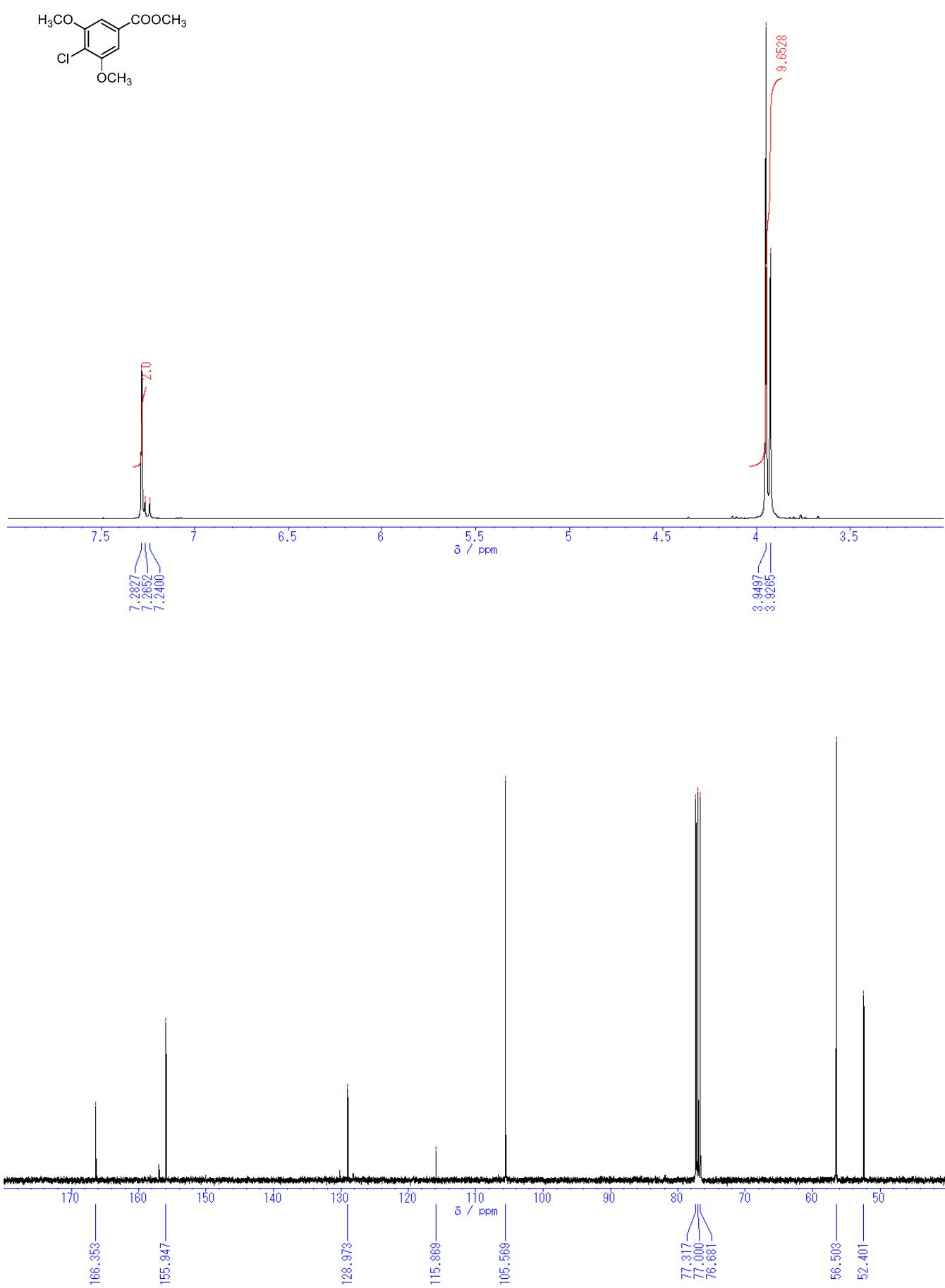


Figure S19. ^1H NMR and ^{13}C NMR spectra for compound 6 (CDCl_3).

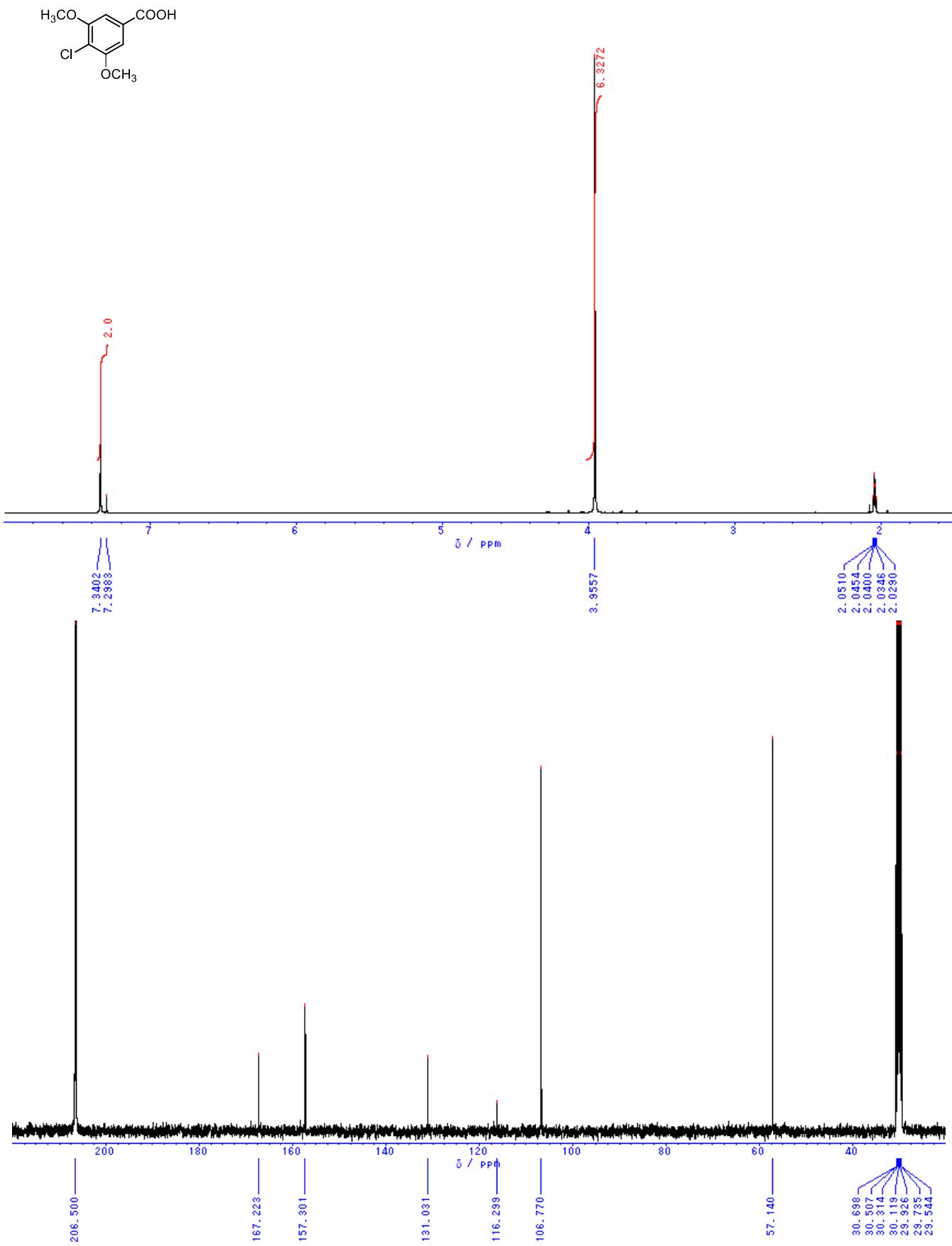


Figure S20. ^1H NMR and ^{13}C NMR spectra for compound 7 (CDCl_3).

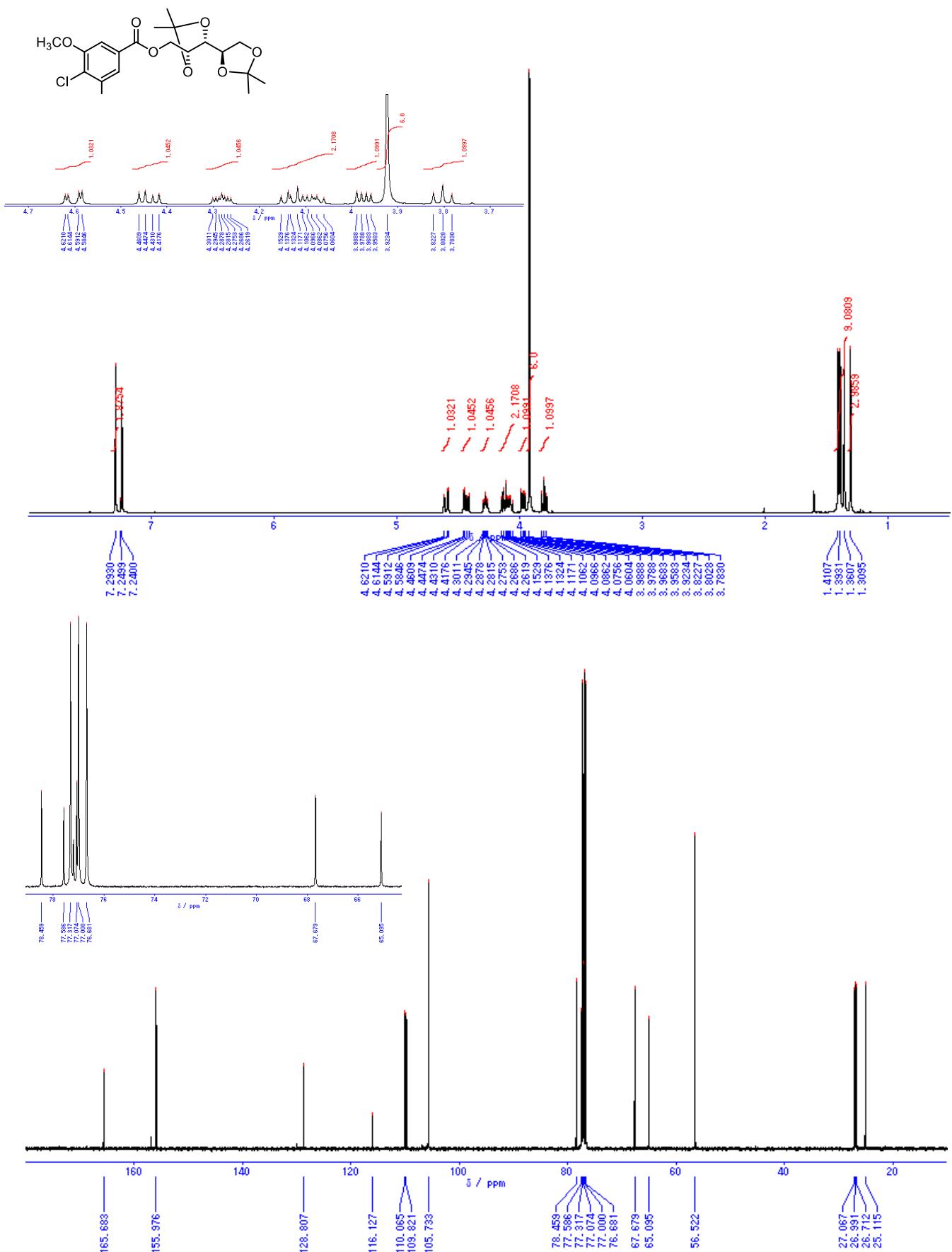


Figure S21. ^1H NMR and ^{13}C NMR spectra for compound **10** (CDCl_3).

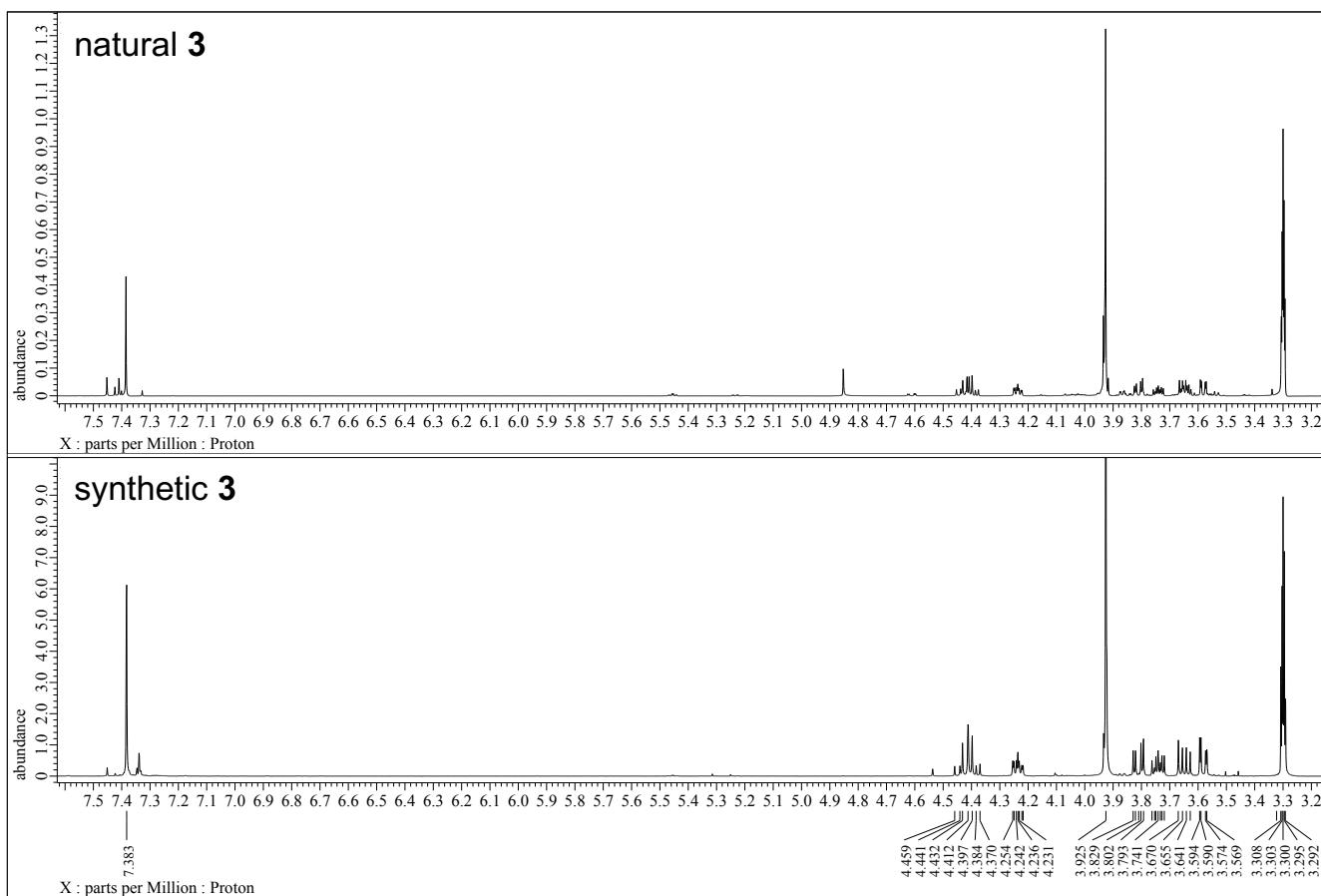


Figure S22. ^1H NMR spectra for natural 3 and synthetic 3 (CD_3OD).

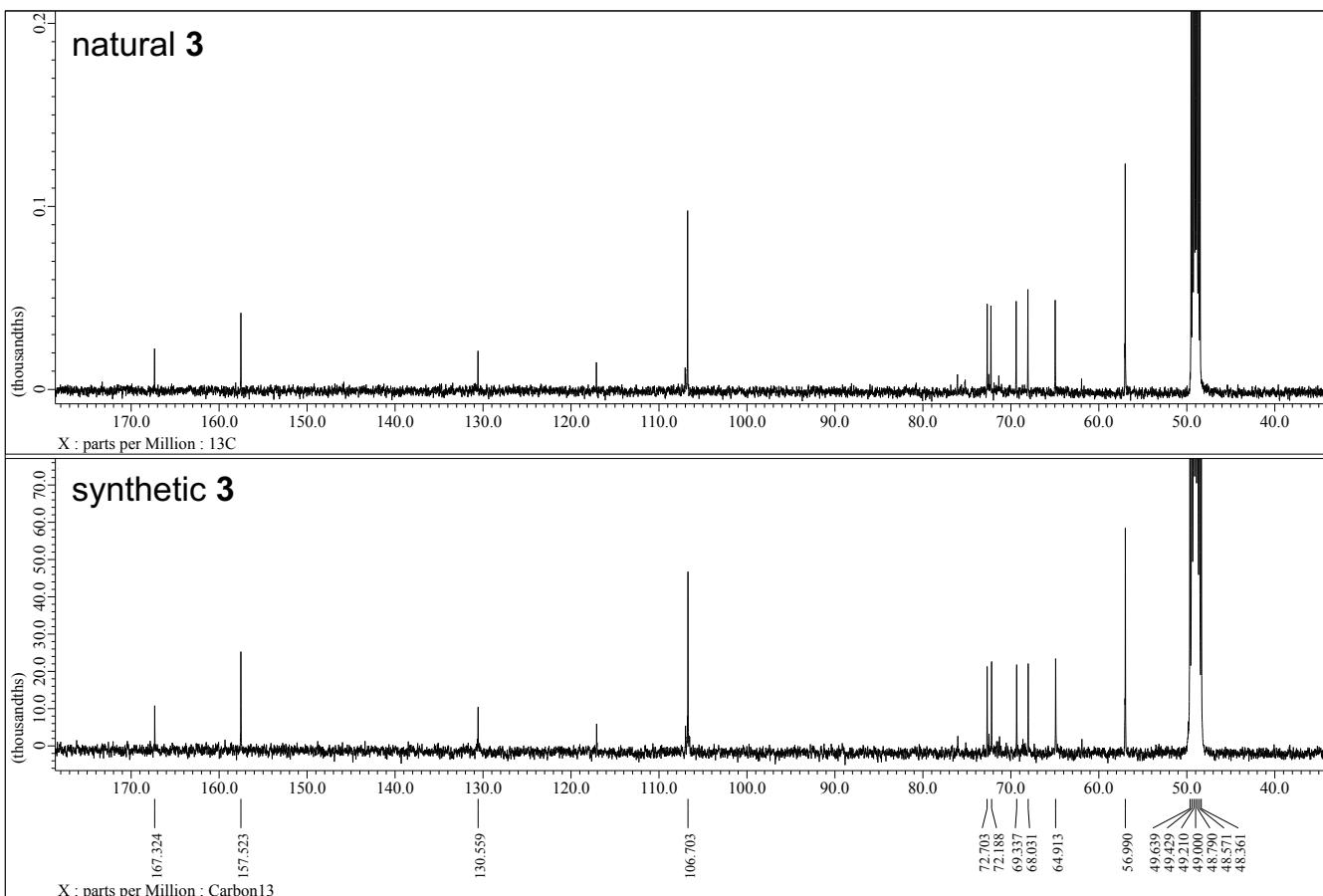


Figure S23. ^{13}C NMR spectra for natural 3 and synthetic 3 (CD_3OD).

Syntheses of 11 and 12

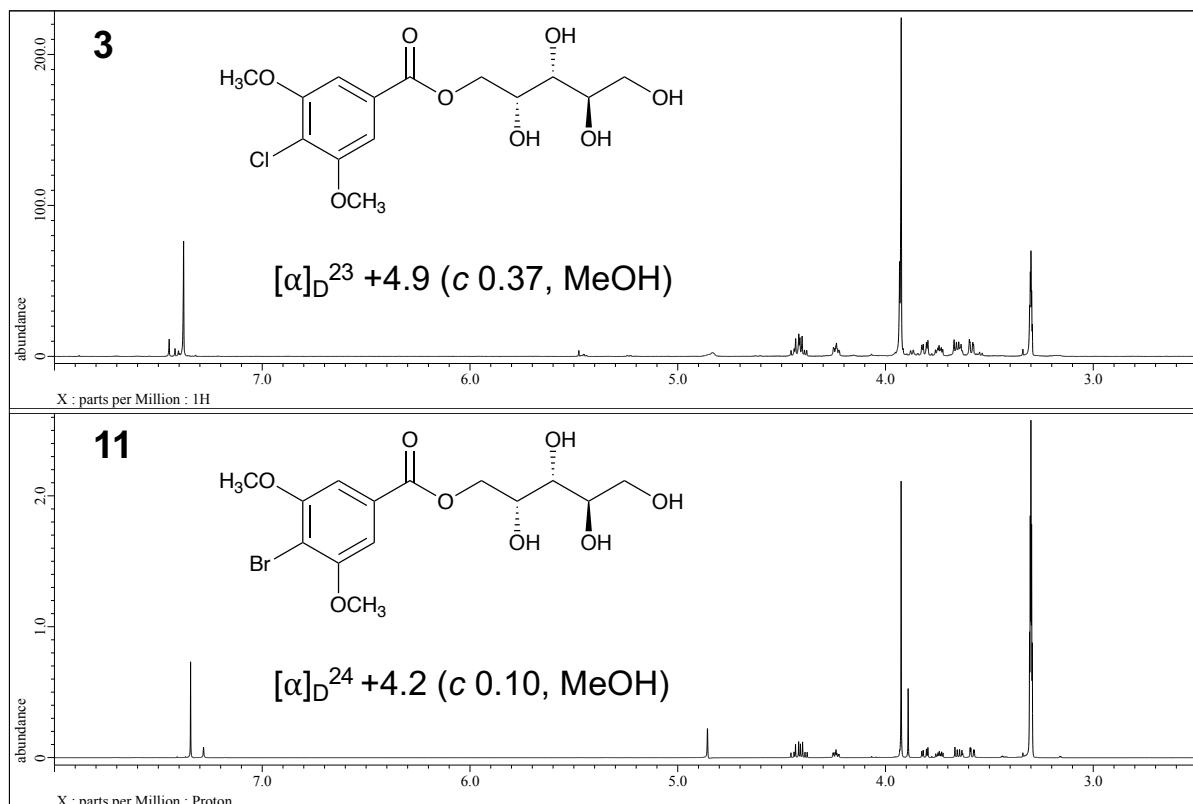
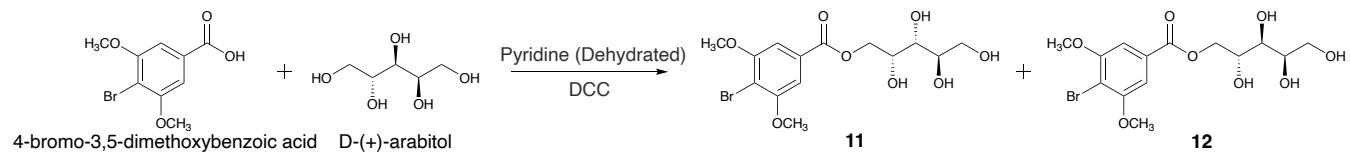


Figure S24. ^1H NMR spectra for compounds 3 and 11 (CD_3OD).

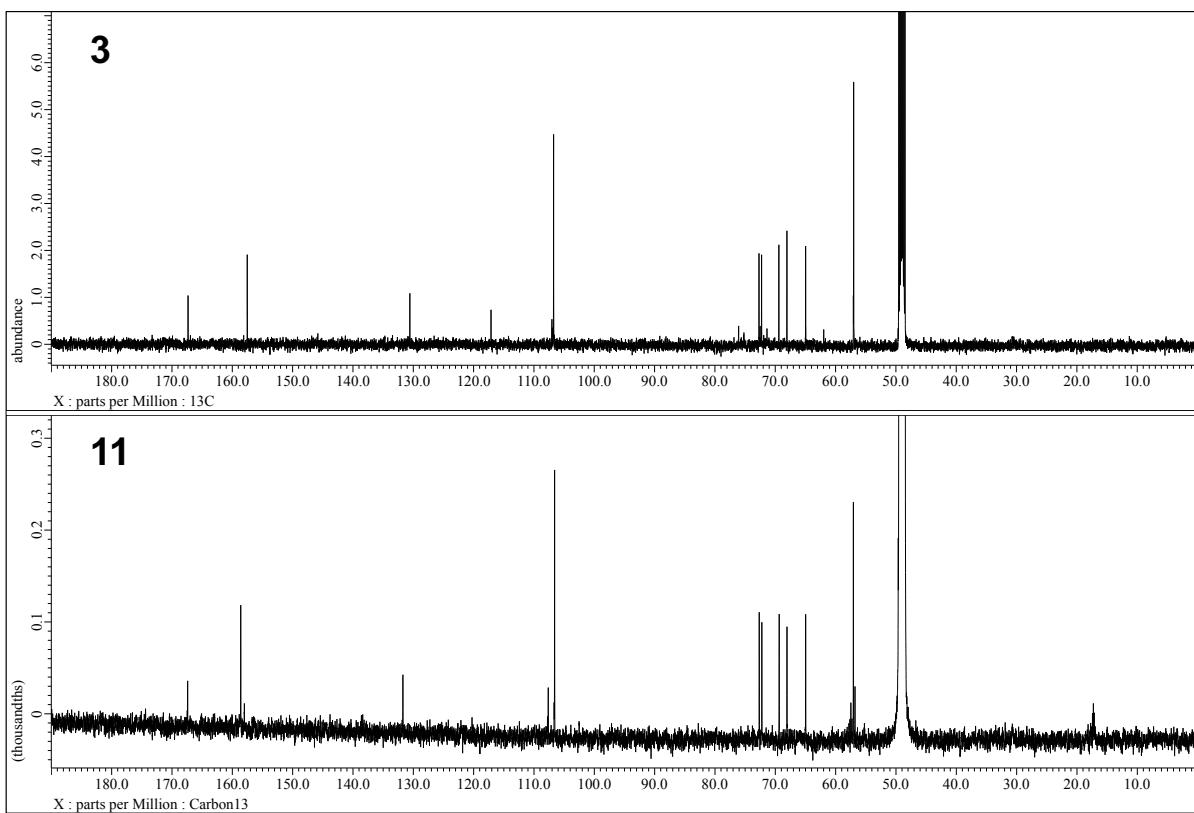


Figure S25. ¹³C NMR spectra for compounds **3** and **11** (CD_3OD).

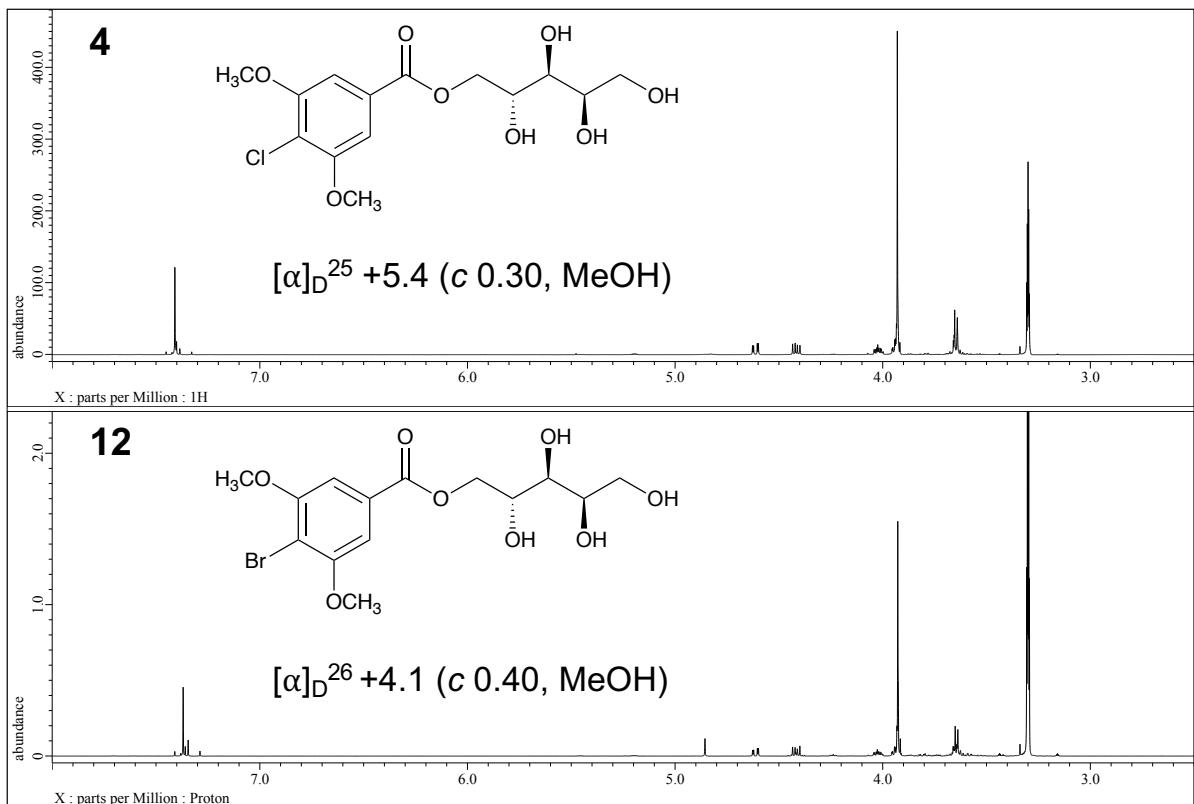


Figure S26. ¹H NMR spectra for compounds **4** and **12** (CD_3OD).

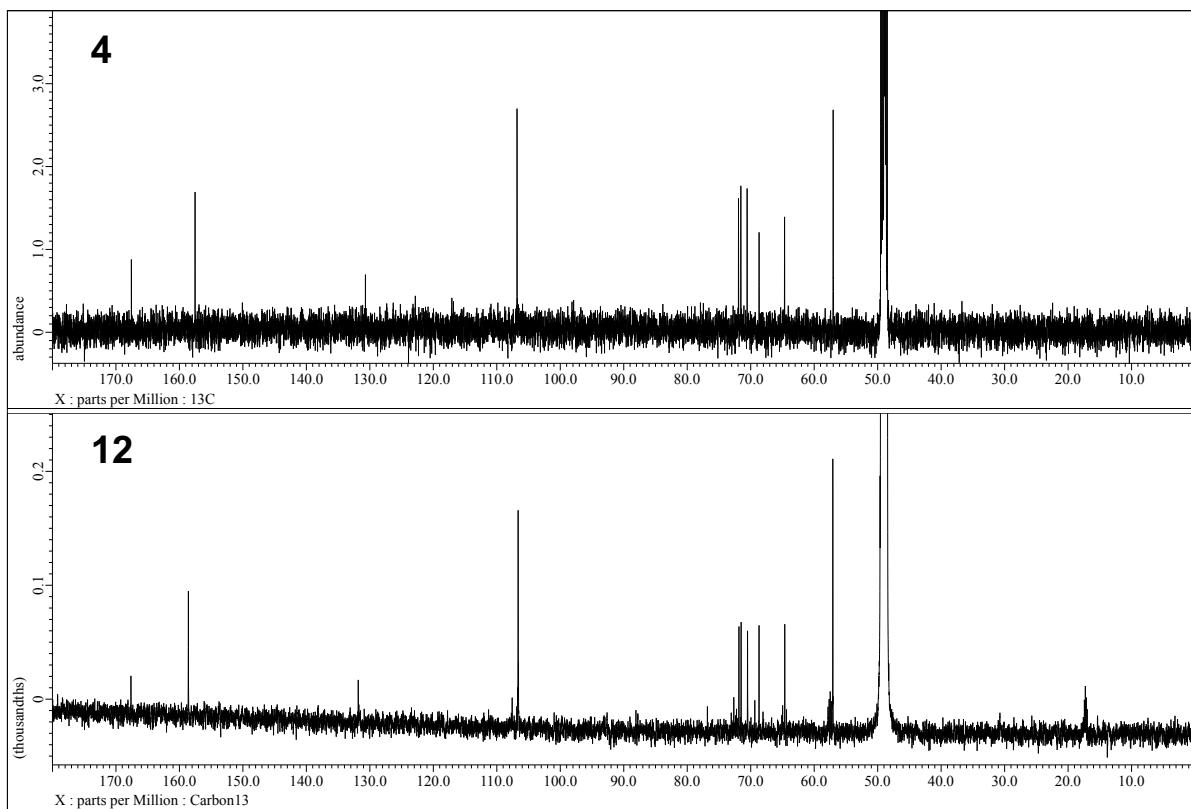


Figure S27. ^{13}C NMR spectra for compounds **4** and **12** (CD_3OD).

Table S2. Crystallographic data of **2** and **3**.

	2	3
Formula	C ₁₇ H ₂₄ O ₈	C ₁₄ H ₁₉ ClO ₈
Formula weight	356.36	350.74
Crystal system	monoclinic	triclinic
Space Group	<i>I</i> 2	<i>P</i> 1
<i>a</i> /Å	10.5226(5)	6.09830(10)
<i>b</i> /Å	5.7292(3)	9.27910(10)
<i>c</i> /Å	28.8051(13)	14.06900(10)
α /°	90	96.4020(10)
β /°	94.548(4)	91.0450(10)
γ /°	90	91.0990(10)
<i>V</i> /Å ³	1731.08(15)	790.842(17)
<i>Z</i>	4	2
<i>D_x</i> , g cm ⁻³	1.367	1.473
<i>F</i> (000)	760.0	368.0
μ /mm ⁻¹	0.919	2.515
Temperature/°C	-100	-100
reflections collected	15769	68656
unique reflections	3371 (<i>R</i> _{int} = 0.0668)	6144 (<i>R</i> _{int} = 0.0442)
parameters refined	245	433
<i>R</i> ₁ (<i>I</i> >2 <i>s</i> (<i>I</i>))	0.0751	0.0269
ωR_2 (all data)	0.2155	0.0765
GOF	1.063	1.118

*R*₁=Σ||*F_o*|-|*F_c*||/Σ|*F_o*|. *wR*₂=[Σ*w*(*F_o*²-*F_c*²)²/Σ*w*(*F_o*²)²]^{1/2}.

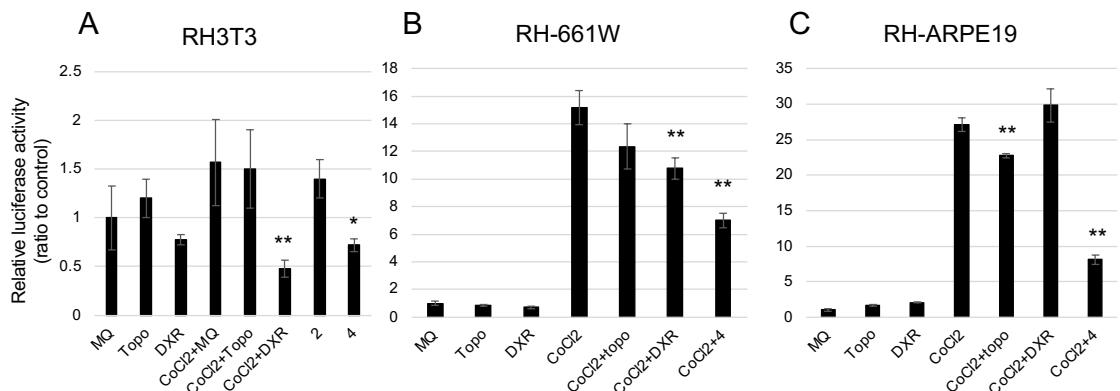


Figure S28. Effect of compounds **2** and **4** on HIF activation *in vitro*. (A) Administration of **2** and **4** suppressed HIF activation in 3T3 cells. (B) Administration of **4** suppressed CoCl₂-induced HIF activation in 661W cells. (C) Administration of **4** suppressed CoCl₂-induced HIF activation in ARPE19 cells. ***p* < 0.01, **p* < 0.05 compared with CoCl₂ without topotecan (Topo) and doxorubicin (DXR), n = 3.

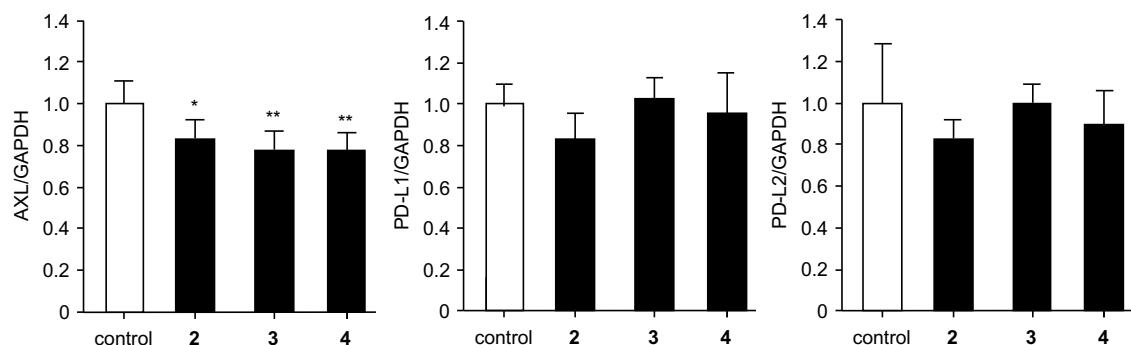


Figure S29. Effect of compounds **2** to **4** on expressions of Axl and immune checkpoints (PD-L1 and PD-L2) on lung cancer cell line A549 cells. Values indicate means with standard deviation from three independent triplicate experiments. Statistical analysis was performed using Fisher's test (***p* < 0.01, **p* < 0.05 vs control, n = 3).