

Supporting Information (SI) for

Structural Analysis of Lignin-Based Furan Resin

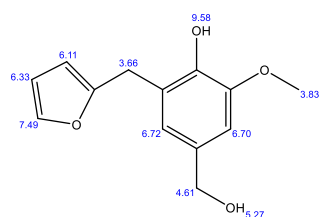
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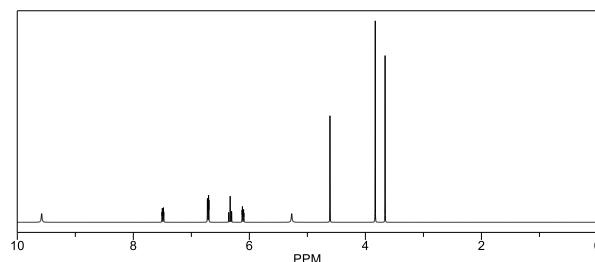
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ChemNMR ¹H Estimation

Estimation quality is indicated by color: **good**, **medium**, **rough**

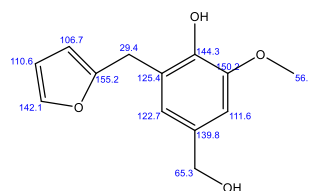


Protocol of the H-1 NMR Prediction (Lib=SU Solvent=DMSO 300 MHz):

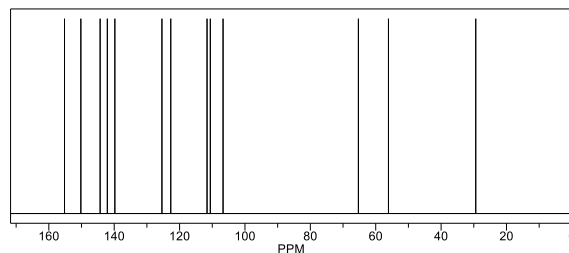
Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
OH	9.58	4.20	alcohol
		4.80	1 -C*R
		0.58	general corrections
OH	5.27	4.20	alcohol
		1.10	1 -C-C*R
		-0.03	general corrections
CH	7.49	7.38	2-furan
		-0.17	1 -C
		0.28	general corrections
CH	6.11	6.30	2-furan
		-0.42	1 -C
		0.23	general corrections
CH	6.33	6.30	2-furan
		-0.12	1 -C
		0.15	general corrections
CH	6.70	7.26	1-benzene
		-0.38	1 -O-C
		-0.17	1 -O
		-0.10	1 -CC*R
		0.00	1 -C-O
		0.09	general corrections
CH	6.72	7.26	1-benzene
		-0.32	1 -O-C
		-0.17	1 -O
		-0.10	1 -CC*R
		0.00	1 -C-O
		0.05	general corrections
CH2	4.61	1.37	methylene
		1.22	1 alpha -1:C*C*C*C*C*1
		2.20	1 alpha -O
		-0.18	general corrections
CH3	3.83	0.86	methyl
		2.87	1 alpha -O-1:C*C*C*C*C*1
		0.10	general corrections
CH2	3.66	1.37	methylene
		1.22	1 alpha -1:C*C*C*C*C*1
		1.07	1 alpha -1:C-O-C=C=C=1

1H NMR Coupling Constant Prediction

shift	atom index	coupling partner, constant and vector
9.58	7	
5.27	17	
7.49	14	
	13	7.5 H-C*C-H
	12	1.5 H-C*CH*C-H
6.11	12	
	13	7.5 H-C*C-H
	14	1.5 H-C*CH*C-H
6.33	13	
	14	7.5 H-C*C-H
	12	7.5 H-C*C-H
6.70	5	
	3	1.5 H-C*C*C-H
6.72	3	
	5	1.5 H-C*C*C-H
4.61	16	
3.83	9	
3.66	10	

ChemNMR ¹³C Estimation

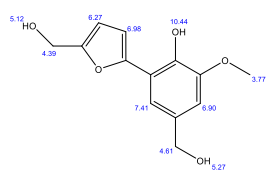
Estimation quality is indicated by color: **good**, **medium**, **rough**



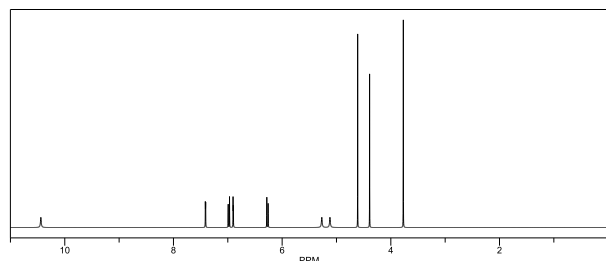
Protocol of the C-13 NMR Prediction: (Lib=S)

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
C	155.2	143.0	2-furan
		9.5	1 -C
		2.7	general corrections
CH	142.1	143.0	2-furan
		-1.8	1 -C
		0.9	general corrections
CH	106.7	109.9	2-furan
		-4.2	1 -C
		1.0	general corrections
CH	110.6	109.9	2-furan
		0.7	1 -C
		0.0	general corrections
C	150.2	128.5	1-benzene
		33.5	1 -O-C
		-12.8	1 -O
		-0.1	1 -C
		0.2	1 -C-O
		0.9	general corrections
C	144.3	128.5	1-benzene
		-14.4	1 -O-C
		28.8	1 -O
		0.7	1 -C
		-1.1	1 -C-O
		1.8	general corrections
C	125.4	128.5	1-benzene
		1.0	1 -O-C
		-12.8	1 -O
		9.2	1 -C
		0.2	1 -C-O
		-0.7	general corrections
C	139.8	128.5	1-benzene
		1.0	1 -O-C
		-7.4	1 -O
		-0.1	1 -C
		12.4	1 -C-O
		5.4	general corrections
CH	111.6	128.5	1-benzene
		-14.4	1 -O-C
		1.4	1 -O
		-3.0	1 -C
		-1.2	1 -C-O
		0.3	general corrections
CH	122.7	128.5	1-benzene
		-7.7	1 -O-C
		1.4	1 -O
		0.7	1 -C
		-1.2	1 -C-O
		1.0	general corrections
CH2	65.3	-2.3	aliphatic
		24.3	1 alpha -1:C*C*C*C*C*1
		49.0	1 alpha -O
		0.3	1 delta -C
		0.3	1 delta -O
		-6.3	general corrections
CH3	56.1	-2.3	aliphatic
		49.0	1 alpha -O
		9.3	1 beta -1:C*C*C*C*C*1
		0.3	1 delta -O
		-0.2	general corrections
CH2	29.4	-2.3	aliphatic
		24.3	1 alpha -1:C*C*C*C*C*1
		17.0	1 alpha -1:C-O-C=C=C=1
		-6.2	1 gamma -O
		0.3	1 delta -C
		0.3	1 delta -O
		-4.0	general corrections

Figure S1. calculated NMR spectra of model **M1**

ChemNMR ¹H Estimation

Estimation quality is indicated by color: **good**, **medium**, **rough**

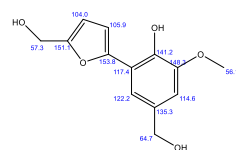


Protocol of the H-1 NMR Prediction (Lib=SU Solvent=DMSO 300 MHz):

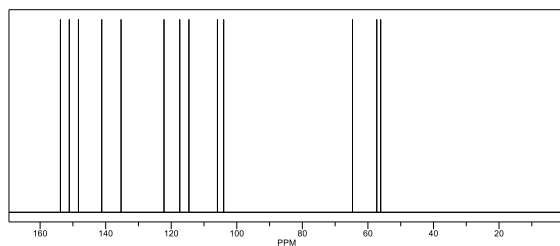
Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
OH	10.44	4.20 4.80 1.44	alcohol 1 -C*R general corrections
OH	5.12	4.20 1.10 -0.18	alcohol 1 -C-C*R general corrections
OH	5.27	4.20 1.10 -0.03	alcohol 1 -C-C*R general corrections
CH	6.98	6.30 7 -0.05 0.73	2-furan 1 unknown substituent(s) 1 -C-O general corrections
CH	6.27	6.30 7 -0.11 0.08	2-furan 1 unknown substituent(s) 1 -C-O general corrections
CH	6.90	7.26 -0.38 -0.17 0.10 0.00	1-benzene 1 -O-C 1 -O 1 -C*R 1 -C-O
CH	7.41	0.09 7.26 -0.32 -0.17 0.34 0.00	general corrections 1-benzene 1 -O-C 1 -O 1 -C*R 1 -C-O
CH2	4.39	0.30 1.37 1.07 2.20 -0.25	general corrections methylene 1 alpha -1:C-O-C=C-C=1 1 alpha -O general corrections
CH2	4.61	1.37 1.22 2.20 -0.18	methylene 1 alpha -1:C-C-C-C-C*1 1 alpha -O general corrections
CH3	3.77	0.86 2.87 0.04	methyl 1 alpha -O-1:C-C-C-C-C*1 general corrections

1H NMR Coupling Constant Prediction

shift	atom index	coupling partner, constant and vector
10.44	7	
5.12	16	
5.27	18	
6.98	11	
6.27	12	12 7.5 H-C*C-H
6.90	5	11 7.5 H-C*C-H
7.41	3	3 1.5 H-C*C-C-H
4.39	15	5 1.5 H-C*C-C-H
4.61	17	
3.77	9	

ChemNMR ¹³C Estimation

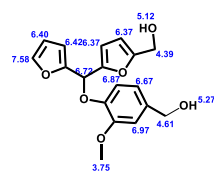
Estimation quality is indicated by color: **good**, **medium**, **rough**



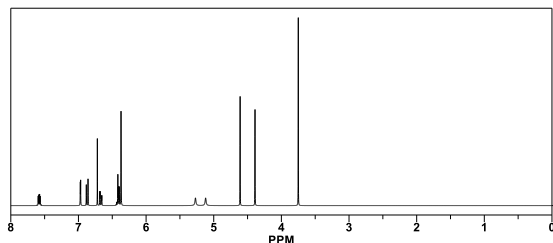
Protocol of the C-13 NMR Prediction: (Lib=S)

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
C	153.8	143.0 11.0 -0.2 0.0	2-furan 1 -1:C-C-C-C-C*1 1 -C-O general corrections
C	151.1	143.0 -1.0 11.2 -2.1	2-furan 1 -1:C-C-C-C-C*1 1 -C-O general corrections
CH	105.9	109.9 1.7 0.8 -6.5	2-furan 1 -1:C-C-C-C-C*1 1 -C-O general corrections
CH	104.0	109.9 -4.9 -2.0 1.0	2-furan 1 -1:C-C-C-C-C*1 1 -C-O general corrections
C	148.3	128.5 33.5 -12.8 0.5 0.2 -1.6	1-benzene 1 -O-C 1 -O 1 -C*R 1 -C-O general corrections
C	141.2	128.5 -14.4 28.8 -1.5 -1.1 0.9	1-benzene 1 -O-C 1 -O 1 -C*R 1 -C-O general corrections
C	117.4	128.5 -1.0 -12.8 8.0 0.2 -7.5	1-benzene 1 -O-C 1 -O 1 -C*R 1 -C-O general corrections
C	135.3	128.5 1.0 -7.4 0.5 12.4 0.3	1-benzene 1 -O-C 1 -O 1 -C*R 1 -C-O general corrections
CH	114.6	128.5 -14.4 1.4 0.0 -1.2 0.3	1-benzene 1 -O-C 1 -O 1 -C*R 1 -C-O general corrections
CH	122.2	128.5 -7.7 1.4 -1.5 -1.2 2.7	1-benzene 1 -O-C 1 -O 1 -C*R 1 -C-O general corrections
CH2	57.3	-2.3 17.0 49.0 0.3 -6.7	aliphatic 1 alpha -1:C-O-C-C-C=1 1 alpha -O 1 delta -1:C-C-C-C-C*1 general corrections
CH2	64.7	-2.3 24.3 49.0 -0.3 0.3 -6.3	aliphatic 1 alpha -1:C-C-C-C-C*1 1 alpha -O 1 delta -1:C-O-C-C-C=1 1 delta -O general corrections
CH3	56.1	-2.3 49.0 9.3 0.3 -0.2	aliphatic 1 alpha -O 1 beta -1:C-C-C-C-C*1 1 delta -O general corrections

Figure S2. calculated NMR spectra of model M2

ChemNMR ¹H Estimation

Estimation quality is indicated by color: good, medium, rough

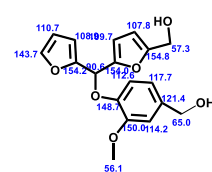


Protocol of the H-1 NMR Prediction (Lib=SU Solvent=DMSO 300 MHz):

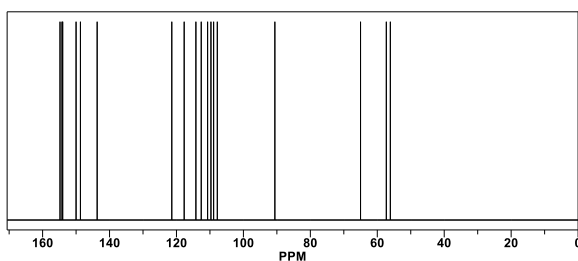
Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
OH	5.12	4.20	alcohol
		1.10	1 -C-C*R
		-0.18	general corrections
OH	5.27	4.20	alcohol
		1.10	1 -C-C*R
		-0.03	general corrections
CH	7.58	7.38	2-furan
		-0.08	1 -C-O
		0.28	general corrections
CH	6.37	6.30	2-furan
		-0.11	1 -C-O
		-0.05	1 -C-O
		0.23	general corrections
CH	6.42	6.30	2-furan
		-0.11	1 -C-O
		0.23	general corrections
CH	6.37	6.30	2-furan
		-0.05	1 -C-O
		-0.11	1 -C-O
		0.23	general corrections
CH	6.40	6.30	2-furan
		-0.05	1 -C-O
		0.15	general corrections
CH	6.97	7.26	1-benzene
		0.00	1 -O-C
		-0.38	1 -O-C
		0.00	1 -O-C
		0.03	1 -C-O
		-0.04	general corrections
CH	6.67	7.26	1-benzene
		0.00	1 -O-C
		-0.32	1 -O-C
		0.00	1 -C-O
		-0.27	general corrections
CH	6.72	1.50	methine
		3.02	2 alpha -1:C*O*C*C*C*1
		2.20	1 alpha -O-1:C*C*C*C*C*1
CH2	4.39	1.37	methylene
		1.07	1 alpha -1:C-O-C=C-C=1
		2.20	1 alpha -O
		-0.25	general corrections
CH2	4.61	1.37	methylene
		1.22	1 alpha -1:C*C*C*C*C*1
		2.20	1 alpha -O
		-0.18	general corrections
CH3	3.75	0.86	methyl
		2.87	1 alpha -O-1:C*C*C*C*C*1
		0.02	general corrections

1H NMR Coupling Constant Prediction

shift	atom index	coupling partner, constant and vector
5.12	13	
5.27	24	
7.58	3	
		2 7.5 H-C*C-H
		1 1.5 H-C*CH*C-H
6.37	11	
		10 7.5 H-C*C-H
6.42	1	
		2 7.5 H-C*C-H
		3 1.5 H-C*CH*C-H
6.37	10	
		11 7.5 H-C*C-H
6.40	2	
		3 7.5 H-C*C-H
		1 7.5 H-C*C-H
6.97	17	
		19 1.5 H-C*C*C-H
6.87	20	
		19 7.5 H-C*C-H
6.67	19	
		20 7.5 H-C*C-H
		17 1.5 H-C*C*C-H
6.72	6	
4.39	12	
4.61	23	
3.75	22	

ChemNMR ¹³C Estimation

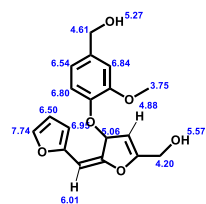
Estimation quality is indicated by color: good, medium, rough



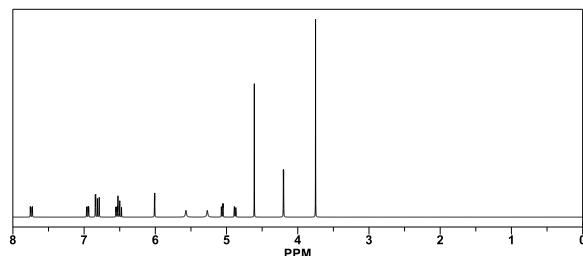
Protocol of the C-13 NMR Prediction: (Lib=S)

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
C	154.0	143.0	2-furan
		11.2	1 -C-O
		-0.2	1 -C-O
C	154.2	143.0	2-furan
		11.2	1 -C-O
C	154.8	143.0	2-furan
		-0.2	1 -C-O
		11.2	1 -C-O
		0.8	general corrections
CH	143.7	143.0	2-furan
		-0.2	1 -C-O
		0.9	general corrections
CH	109.7	109.9	2-furan
		-2.0	1 -C-O
		0.8	1 -C-O
		1.0	general corrections
CH	108.9	109.9	2-furan
		-2.0	1 -C-O
		1.0	general corrections
CH	107.8	109.9	2-furan
		0.8	1 -C-O
		-2.0	1 -C-O
		-0.9	general corrections
CH	110.7	109.9	2-furan
		0.8	1 -C-O
		0.0	general corrections
C	148.7	128.5	1-benzene
		33.5	1 -O-C
		-14.4	1 -O-C
		-1.1	1 -C-O
		2.2	general corrections
C	150.0	128.5	1-benzene
		-14.4	1 -O-C
		33.5	1 -O-C
		0.2	1 -C-O
		2.2	general corrections
C	121.4	128.5	1-benzene
		-7.7	1 -O-C
		1.0	1 -O-C
		12.4	1 -C-O
		-12.8	general corrections
CH	114.2	128.5	1-benzene
		1.0	1 -O-C
		-14.4	1 -O-C
		-1.2	1 -C-O
		0.3	general corrections
CH	112.6	128.5	1-benzene
		-14.4	1 -O-C
		1.0	1 -O-C
		0.2	1 -C-O
		2.7	general corrections
CH	117.7	128.5	1-benzene
		1.0	1 -O-C
		-7.7	1 -O-C
		-1.2	1 -C-O
		-2.9	general corrections
CH	90.6	-2.3	aliphatic
		34.0	2 alpha -1:C-O-C=C-C=1
		49.0	1 alpha -O
		9.3	1 beta -1:C*C*C*C*C*1
		0.3	1 delta -C
		0.3	1 delta -O
CH2	57.3	-2.3	aliphatic
		17.0	1 alpha -1:C-O-C=C-C=1
		49.0	1 alpha -O
		0.3	1 delta -C
		-6.7	general corrections
CH2	65.0	-2.3	aliphatic
		24.3	1 alpha -1:C*C*C*C*C*1
		49.0	1 alpha -O
		0.3	1 delta -O
		-6.3	general corrections
CH3	56.1	-2.3	aliphatic
		49.0	1 alpha -O
		9.3	1 beta -1:C*C*C*C*C*1
		0.3	1 delta -O
		-0.2	general corrections

Figure S3. calculated NMR spectra of model M3

ChemNMR ¹H Estimation

Estimation quality is indicated by color: **good**, **medium**, **rough**

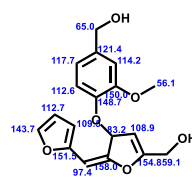


Protocol of the ¹H-1 NMR Prediction (Lib=SU Solvent=DMSO 300 MHz):

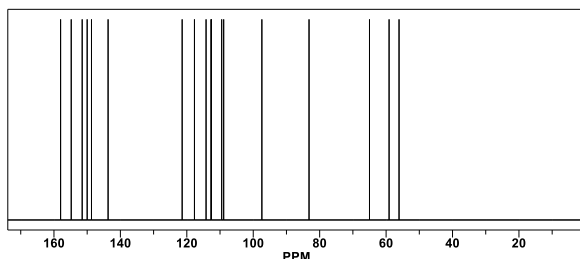
Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
OH	5.57	4.20	alcohol
		0.75	1 -CC=C
		0.62	general corrections
OH	5.27	4.20	alcohol
		1.10	1 -C-C*R
		-0.03	general corrections
CH	7.74	7.38	2-furan
		?	1 unknown substituent(s)
		0.36	general corrections
CH	5.06	1.50	methine
		1.36	2 alpha -C=C
		2.20	1 alpha -O-1:C*C*C*C*C*C*1
CH	6.95	6.30	2-furan
		?	1 unknown substituent(s)
		0.65	general corrections
CH	6.50	6.30	2-furan
		?	1 unknown substituent(s)
		0.20	general corrections
CH	6.84	7.26	1-benzene
		-0.13	1 -O-CC=C
		-0.38	1 -O-C
		0.00	1 -C-O
		0.09	general corrections
CH	6.80	7.26	1-benzene
		-0.45	1 -O-CC=C
		0.00	1 -O-C
		0.03	1 -C-O
		-0.04	general corrections
CH	6.54	7.26	1-benzene
		-0.13	1 -O-CC=C
		-0.32	1 -O-C
		0.00	1 -C-O
		-0.27	general corrections
CH2	4.20	1.37	methylene
		0.63	1 alpha -C=C
		2.20	1 alpha -O
CH2	4.61	1.37	methylene
		1.22	1 alpha -1:C*C*C*C*C*C*1
		2.20	1 alpha -O
		-0.18	general corrections
CH3	3.75	0.86	methyl
		2.87	1 alpha -O-1:C*C*C*C*C*C*1
		0.02	general corrections
H	4.88	5.25	1-ethylene
		-1.00	1 -O-C=C trans
		-0.01	1 -C-O cis
		0.64	1 -C-O gem
H	6.01	5.25	1-ethylene
		-0.60	1 -O-C=C cis
		-0.02	1 -C-O trans
		1.38	1 -C*R gem

1H NMR Coupling Constant Prediction

shift	atom index	coupling partner, constant and vector
5.57	13	
5.27	24	
7.74	3	
	2	7.5 H-C*C-H
	1	1.5 H-C*CH*C-H
5.06	11	
	25	6.2 H-C-C(sp2)-H
	26	-1.0 H-C>C=C*H
6.95	1	
	2	7.5 H-C*C-H
	3	1.5 H-C*CH*C-H
6.50	2	
	3	7.5 H-C*C-H
	1	7.5 H-C*C-H
6.84	17	
	19	1.5 H-C*C*C-H
6.80	20	
	19	7.5 H-C*C-H
6.54	19	
	20	7.5 H-C*C-H
	17	1.5 H-C*C*C-H
4.20	12	
	25	-1.0 H-CH>C=C<H
4.61	23	
3.75	22	
4.88	25	
	11	6.2 H-C(sp2)-C-H
	12	-1.0 H>C=C<CH-H
6.01	26	
	11	-1.0 H>C=C<C-H

ChemNMR ¹³C Estimation

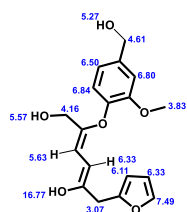
Estimation quality is indicated by color: **good**, **medium**, **rough**



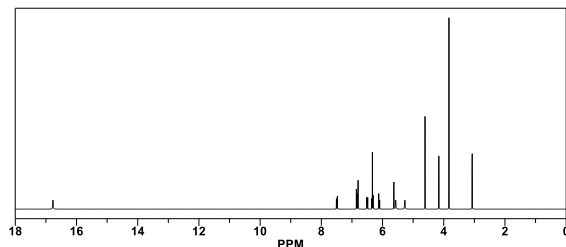
Protocol of the ¹³C-13 NMR Prediction: (Lib=S)

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
C	154.8	123.3	1-ethylene
		14.2	1 -C-O
		25.5	1 -OC=C
		-8.2	1 -C-O
C	151.5	143.0	2-furan
		12.3	1 -C=C
		-2.8	general corrections
CH	143.7	143.0	2-furan
		2.0	1 -C=C
		-1.3	general corrections
C	158.0	123.3	1-ethylene
		14.2	1 -C-O
		25.5	1 -OC=C
		-5.0	1 -C*R
CH	83.2	-2.3	aliphatic
		39.0	2 alpha -C=C
		49.0	1 alpha -O
		9.3	1 beta -1:C*C*C*C*C*C*1
		10.1	1 beta -O
		-5.0	1 gamma -1:C-O-C=C-C=1
		-2.5	1 gamma -C
		0.6	2 delta -O
		-15.0	general corrections
CH	108.9	123.3	1-ethylene
		-8.2	1 -C-O
		-30.2	1 -OC=C
		14.2	1 -C-O
		9.8	general corrections
CH	109.5	109.9	2-furan
		1.9	1 -C=C
		-2.3	general corrections
CH	112.7	109.9	2-furan
		2.8	1 -C=C
		0.0	general corrections
C	148.7	128.5	1-benzene
		33.5	1 -O-C
		-14.4	1 -O-C
		-1.1	1 -C-O
		2.2	general corrections
C	150.0	128.5	1-benzene
		-14.4	1 -O-C
		33.5	1 -O-C
		0.2	1 -C-O
		2.2	general corrections
C	121.4	128.5	1-benzene
		-7.7	1 -O-C
		1.0	1 -O-C
		12.4	1 -C-O
		-12.8	general corrections
CH	114.2	128.5	1-benzene
		1.0	1 -O-C
		-14.4	1 -O-C
		-1.2	1 -C-O
		0.3	general corrections
CH	112.6	128.5	1-benzene
		-14.4	1 -O-C
		1.0	1 -O-C
		0.2	1 -C-O
		-2.7	general corrections
CH	117.7	128.5	1-benzene
		1.0	1 -O-C
		-7.7	1 -O-C
		-1.2	1 -C-O
		-2.9	general corrections
CH2	59.1	-2.3	aliphatic
		19.5	1 alpha -C=C
		49.0	1 alpha -O
		10.1	1 beta -O
		-2.1	1 gamma -C=C
		-2.5	1 gamma -C
		0.3	1 delta -O
		-12.9	general corrections
CH2	65.0	-2.3	aliphatic
		24.3	1 alpha -1:C*C*C*C*C*C*1
		49.0	1 alpha -O
		0.3	1 delta -O
		-6.3	general corrections
CH3	56.1	-2.3	aliphatic
		49.0	1 alpha -O
		9.3	1 beta -1:C*C*C*C*C*C*1
		0.3	1 delta -O
		-0.2	general corrections
CH	97.4	123.3	1-ethylene
		-8.2	1 -C-O
		-30.2	1 -OC=C
		12.5	1 -C*R

Figure S4. calculated NMR spectra of model M4

ChemNMR ¹H Estimation

Estimation quality is indicated by color: **good**, **medium**, **rough**

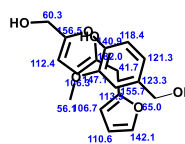


Protocol of the H-1 NMR Prediction (Lib=SU Solvent=DMSO 300 MHz):

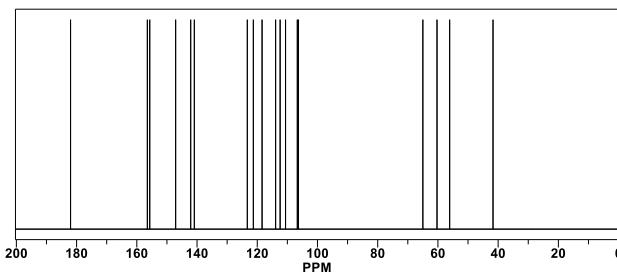
Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
OH	5.27	4.20	alcohol
		1.10	1 -C-C*R
		-0.03	general corrections
OH	16.77	15.00	enol
		1.77	general corrections
OH	5.57	4.20	alcohol
		0.75	1 -C=C
		0.62	general corrections
CH	7.49	7.38	2-furan
		-0.17	1 -C
		0.28	general corrections
CH	6.11	6.30	2-furan
		-0.42	1 -C
		0.23	general corrections
CH	6.33	6.30	2-furan
		-0.12	1 -C
		0.15	general corrections
CH	6.80	7.26	1-benzene
		-0.17	1 -O
		-0.38	1 -O-C
		0.00	1 -C-O
		0.09	general corrections
CH	6.84	7.26	1-benzene
		-0.53	1 -O
		0.00	1 -O-C
		0.03	1 -C-O
		0.08	general corrections
CH	6.50	7.26	1-benzene
		-0.17	1 -O
		-0.32	1 -O-C
		0.00	1 -C-O
		-0.27	general corrections
CH2	4.61	1.37	methylene
		1.22	1 alpha -1:C*C*C*C*C*1
		2.20	1 alpha -O
		-0.18	general corrections
CH2	4.16	1.37	methylene
		0.63	1 alpha -C=C
		2.20	1 alpha -O
		-0.04	general corrections
CH3	3.83	0.86	methyl
		2.87	1 alpha -O-1:C*C*C*C*C*1
		0.10	general corrections
CH2	3.07	1.37	methylene
		1.07	1 alpha -1:C-O-C=C=1
		0.63	1 alpha -C=C
H	5.63	5.25	1-ethylene
		-0.85	1 -O-1:C*C*C*C*C*1 trans
		-0.01	1 -C-O cis
		1.24	1 -C=C gem
H	6.33	5.25	1-ethylene
		-0.22	1 -C cis
		?	1 unknown substituent(s)
		1.24	1 -C=C gem
		0.06	general corrections

1H NMR Coupling Constant Prediction

shift	atom index	coupling partner, constant and vector
5.27	24	
16.77	12	
5.57	22	
7.49	3	
		2 7.5 H-C*C-H
		1 1.5 H-C*CH*C-H
6.11	1	
		2 7.5 H-C*C-H
		3 1.5 H-C*CH*C-H
6.33	2	
		3 7.5 H-C*C-H
		1 7.5 H-C*C-H
6.80	15	
		17 1.5 H-C*C*C-H
6.84	18	
		17 7.5 H-C*C-H
6.50	17	
		18 7.5 H-C*C-H
		15 1.5 H-C*C*C-H
4.61	23	
4.16	21	
		25 -1.0 H-CH>C=C<H
3.83	20	
3.07	6	
		26 -1.0 H-CH>C=C<H
5.63	25	
		21 -1.0 H>C=C<CH-H
6.33	26	
		6 -1.0 H>C=C<CH-H

ChemNMR ¹³C Estimation

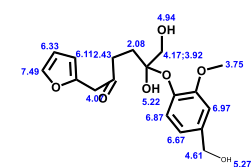
Estimation quality is indicated by color: **good**, **medium**, **rough**



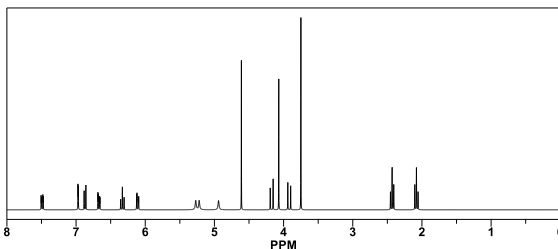
Protocol of the C-13 NMR Prediction: (Lib=S)

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
C	155.7	143.0	2-furan
		9.5	1 -C
		3.2	general corrections
CH	142.1	143.0	2-furan
		-1.8	1 -C
		0.9	general corrections
CH	106.7	109.9	2-furan
		-4.2	1 -C
		1.0	general corrections
CH	110.6	109.9	2-furan
		0.7	1 -C
		0.0	general corrections
C	140.9	128.5	1-benzene
		28.2	1 -O-C=C
		-14.4	1 -O-C
		-1.1	1 -C-O
		-0.3	general corrections
C	147.1	128.5	1-benzene
		-11.5	1 -O-C=C
		33.5	1 -O-C
		0.2	1 -C-O
		-3.6	general corrections
C	123.3	128.5	1-benzene
		-5.8	1 -O-C=C
		1.0	1 -O-C
		12.4	1 -C-O
		-12.8	general corrections
CH	113.9	128.5	1-benzene
		0.7	1 -O-C=C
		-14.4	1 -O-C
		-1.2	1 -C-O
		0.3	general corrections
CH	118.4	128.5	1-benzene
		-11.5	1 -O-C=C
		1.0	1 -O-C
		0.2	1 -C-O
		0.2	general corrections
CH	121.3	128.5	1-benzene
		0.7	1 -O-C=C
		-7.7	1 -O-C
		-1.2	1 -C-O
		1.0	general corrections
CH2	65.0	-2.3	aliphatic
		24.3	1 alpha -1:C*C*C*C*C*1
		49.0	1 alpha -O
		0.3	1 delta -O
		-6.3	general corrections
C	156.5	123.3	1-ethylene
		14.2	1 -C-O
		26.0	1 -O-1:C*C*C*C*C*1
		-7.0	1 -C=C
C	182.0	123.3	1-ethylene
		9.4	1 -C
		52.7	1 -O
		-7.0	1 -C=C
		3.6	general corrections
CH2	60.3	-2.3	aliphatic
		19.5	1 alpha -C=C
		49.0	1 alpha -O
		10.1	1 beta -O
		-2.6	1 gamma -1:C*C*C*C*C*1
		-2.1	1 gamma -C=C
		-11.3	general corrections
CH3	56.1	-2.3	aliphatic
		49.0	1 alpha -O
		9.3	1 beta -1:C*C*C*C*C*1
		0.3	1 delta -O
		-0.2	general corrections
CH2	41.7	-2.3	aliphatic
		17.0	1 alpha -1:C-O-C=C=1
		19.5	1 alpha -C=C
		10.1	1 beta -O
		-2.1	1 gamma -C=C
		-0.5	general corrections
CH	112.4	123.3	1-ethylene
		-8.2	1 -C-O
		-28.4	1 -O-1:C*C*C*C*C*1
		13.6	1 -C=C
		12.1	general corrections
CH	106.3	123.3	1-ethylene
		-7.4	1 -C
		-35.3	1 -O
		13.6	1 -C=C
		12.1	general corrections

Figure S5. calculated NMR spectra of model M5

ChemNMR ¹H Estimation

Estimation quality is indicated by color: **good**, **medium**, **rough**

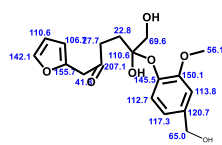


Protocol of the H-1 NMR Prediction (Lib=SU Solvent=DMSO 300 MHz):

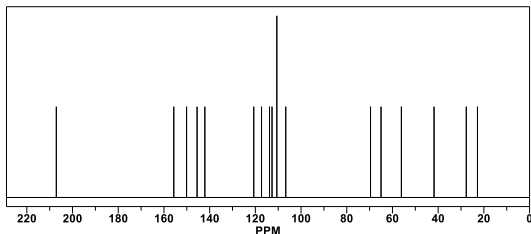
Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
OH	5.22	4.20	alcohol
		1.20	1 -CCO
		-0.18	general corrections
OH	5.27	4.20	alcohol
		1.10	1 -C-C*R
		-0.03	general corrections
OH	4.94	4.20	alcohol
		1.20	1 -CCO
		-0.46	general corrections
CH	7.49	7.38	2-furan
		-0.17	1 -C
		0.28	general corrections
CH	6.11	6.30	2-furan
		-0.42	1 -C
		0.23	general corrections
CH	6.33	6.30	2-furan
		-0.12	1 -C
		0.15	general corrections
CH	6.97	7.26	1-benzene
		0.00	1 -O-C
		-0.38	1 -O-C
		0.00	1 -C-O
		0.09	general corrections
CH	6.87	7.26	1-benzene
		-0.38	1 -O-C
		0.00	1 -O-C
		0.03	1 -C-O
		-0.04	general corrections
CH	6.67	7.26	1-benzene
		0.00	1 -O-C
		-0.32	1 -O-C
		0.00	1 -C-O
		-0.27	general corrections
CH2	4.61	1.37	methylene
		1.22	1 alpha -1:C*C*C*C*C*C*1
		2.20	1 alpha -O
		-0.18	general corrections
CH2	4.17,3.915000	1.37	methylene
		2.20	1 alpha -O
		0.38	1 beta -O-1:C*C*C*C*C*C*1
		0.15	1 beta -O
		-0.06	1 beta -C
CH3	3.75	0.86	methyl
		2.87	1 alpha -O-1:C*C*C*C*C*C*1
		0.02	general corrections
CH2	4.07	1.37	methylene
		1.07	1 alpha -1:C-O-C=C-C=1
		1.12	1 alpha -C(=O)-C
		0.51	general corrections
CH2	2.43	1.37	methylene
		1.12	1 alpha -C(=O)-C
		-0.06	1 beta -C
CH2	2.08	1.37	methylene
		0.38	1 beta -O-1:C*C*C*C*C*C*1
		0.15	1 beta -O
		-0.06	1 beta -C
		0.24	1 beta -C(=O)-C

1H NMR Coupling Constant Prediction

shift	atom index	coupling partner	constant and vector
5.22	8		
5.27	25		
4.94	23		
7.49	3		
		2 7.5 H-C*C-H	
		1 1.5 H-C*CH*C-H	
6.11	1		
		2 7.5 H-C*C-H	
		3 1.5 H-C*CH*C-H	
6.33	2		
		3 7.5 H-C*C-H	
		1 7.5 H-C*C-H	
6.97	18		
		16 1.5 H-C*C*C-H	
6.87	15		
		16 7.5 H-C*C-H	
6.67	16		
		15 7.5 H-C*C-H	
		18 1.5 H-C*C*C-H	
4.61	24		
4.04	22	diastereotopic -12.4 H-C-H	
3.75	21		
4.07	6		
2.43	11		
		10 7.1 H-CH-CH-H	
2.08	10		
		11 7.1 H-CH-CH-H	

ChemNMR ¹³C Estimation

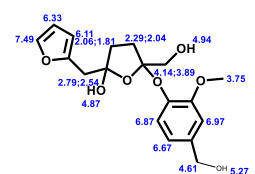
Estimation quality is indicated by color: **good**, **medium**, **rough**



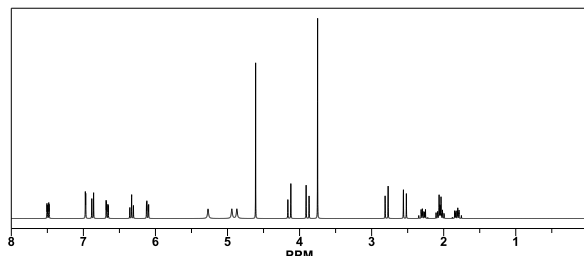
Protocol of the C-13 NMR Prediction: (Lib=S)

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
C	155.7	143.0	2-furan
		9.5	1 -C
		3.2	general corrections
CH	142.1	143.0	2-furan
		-1.8	1 -C
		0.9	general corrections
CH	106.7	109.9	2-furan
		-4.2	1 -C
		1.0	general corrections
CH	110.6	109.9	2-furan
		0.7	1 -C
		0.0	general corrections
C	145.5	128.5	1-benzene
		30.3	1 -OCC
		-14.4	1 -O-C
		-1.1	1 -C-O
		2.2	general corrections
C	150.1	128.5	1-benzene
		-14.3	1 -OCC
		33.5	1 -O-C
		0.2	1 -C-O
		2.2	general corrections
C	120.7	128.5	1-benzene
		-8.4	1 -OCC
		1.0	1 -O-C
		12.4	1 -C-O
		-12.8	general corrections
CH	113.8	128.5	1-benzene
		0.6	1 -OCC
		-14.4	1 -O-C
		-1.2	1 -C-O
		0.3	general corrections
CH	112.7	128.5	1-benzene
		-14.3	1 -OCC
		1.0	1 -O-C
		0.2	1 -C-O
		-2.7	general corrections
CH	117.3	128.5	1-benzene
		0.6	1 -OCC
		-7.7	1 -O-C
		-1.2	1 -C-O
		-2.9	general corrections
C	207.1	193.0	1-carbonyl
		7.6	1 -C-C
		6.5	1 -C
		0.0	general corrections
C	110.6	-2.3	aliphatic
		18.2	2 alpha -C
		98.0	2 alpha -O
		9.3	1 beta -1:C*C*C*C*C*C*1
		9.4	1 beta -C
		10.1	1 beta -O
		-2.7	1 gamma -C(=O)-C
		0.3	1 delta -O
		-29.7	general corrections
CH2	65.0	-2.3	aliphatic
		24.3	1 alpha -1:C*C*C*C*C*C*1
		49.0	1 alpha -O
		0.3	1 delta -O
		-6.3	general corrections
CH2	69.6	-2.3	aliphatic
		9.1	1 alpha -C
		49.0	1 alpha -O
		9.4	1 beta -C
		20.2	2 beta -O
		-2.6	1 gamma -1:C*C*C*C*C*C*1
		-2.5	1 gamma -C
		0.0	1 delta -C(=O)-C
		-10.7	general corrections
CH3	56.1	-2.3	aliphatic
		49.0	1 alpha -O
		9.3	1 beta -1:C*C*C*C*C*C*1
		0.3	1 delta -O
		-0.2	general corrections
CH2	41.8	-2.3	aliphatic
		17.0	1 alpha -1:C-O-C=C-C=1
		29.3	1 alpha -C(=O)-C
		-2.5	1 gamma -C
		0.3	1 delta -C
CH2	27.7	-2.3	aliphatic
		29.3	1 alpha -C(=O)-C
		9.1	1 alpha -C
		9.4	1 beta -C
		-5.0	1 gamma -1:C-O-C=C-C=1
		-2.5	1 gamma -C
		-12.4	2 gamma -O
		0.3	1 delta -1:C*C*C*C*C*C*1
		0.3	1 delta -O
		1.5	general corrections
CH2	22.8	-2.3	aliphatic
		18.2	2 alpha -C
		0.5	1 beta -C(=O)-C
		9.4	1 beta -C
		20.2	2 beta -O
		-2.6	1 gamma -1:C*C*C*C*C*C*1
		-6.2	1 gamma -O
		-0.3	1 delta -1:C-O-C=C-C=1
		-14.1	general corrections

Figure S6. calculated NMR spectra of model M6

ChemNMR ¹H Estimation

Estimation quality is indicated by color: **good**, **medium**, **rough**

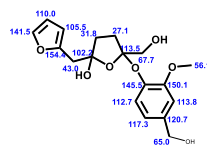


Protocol of the H-1 NMR Prediction (Lib=SU Solvent=DMSO 300 MHz):

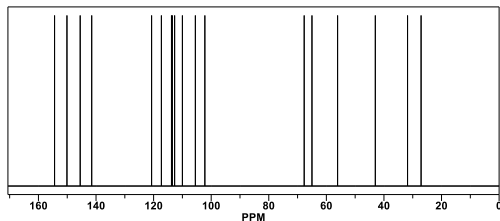
Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
OH	4.87	4.20	alcohol
		0.85	1 -CCC*R
		-0.18	general corrections
OH	4.94	4.20	alcohol
		1.20	1 -CCO
		-0.46	general corrections
OH	5.27	4.20	alcohol
		1.10	1 -C-C*R
		-0.03	general corrections
CH	7.49	7.38	2-furan
		-0.17	1 -C
		0.28	general corrections
CH	6.11	6.30	2-furan
		-0.42	1 -C
		0.23	general corrections
CH	6.33	6.30	2-furan
		-0.12	1 -C
		0.15	general corrections
CH2	2.29,2.045000	1.85	tetrahydrofuran
		0.38	1 beta -O-1:C*C*C*C*1 from methylene
		-0.06	1 beta -C from methylene
CH2	2.06,1.815000	1.85	tetrahydrofuran
		0.15	1 beta -O from methylene
		-0.06	1 beta -C from methylene
CH	6.97	7.26	1-benzene
		0.00	1 -O-C
		-0.38	1 -O-C
		0.00	1 -C-O
		0.09	general corrections
CH	6.87	7.26	1-benzene
		-0.38	1 -O-C
		0.00	1 -O-C
		0.03	1 -C-O
		-0.04	general corrections
CH	6.67	7.26	1-benzene
		0.00	1 -O-C
		-0.32	1 -O-C
		0.00	1 -C-O
		-0.27	general corrections
CH2	4.14,3.895000	1.37	methylene
		2.20	1 alpha -O
		0.38	1 beta -O-1:C*C*C*C*1
		0.13	1 beta -O-C
		-0.06	1 beta -C
CH2	4.61	1.37	methylene
		1.22	1 alpha -1:C*C*C*C*1
		2.20	1 alpha -O
		-0.18	general corrections
CH3	3.75	0.86	methyl
		2.87	1 alpha -O-1:C*C*C*C*1
		0.02	general corrections
CH2	2.79,2.535000	1.37	methylene
		1.07	1 alpha -1:C-O-C=C=1
		0.13	1 beta -O-C
		0.15	1 beta -O
		-0.06	1 beta -C

1H NMR Coupling Constant Prediction

shift	atom index	coupling partner, constant and vector
4.87	11	
4.94	23	
5.27	25	
7.49	3	
		2 7.5 H-C*C-H
		1 1.5 H-C*CH*C-H
6.11	1	
		2 7.5 H-C*C-H
		3 1.5 H-C*CH*C-H
6.33	2	
		3 7.5 H-C*C-H
		1 7.5 H-C*C-H
2.17	9	diastereotopic -12.4 H-C-H
		10 7.1 H-CH-CH-H
1.94	10	diastereotopic -12.4 H-C-H
		9 7.1 H-CH-CH-H
6.97	18	
		16 1.5 H-C*C*H
6.87	15	
		16 7.5 H-C*C-H
6.67	16	
		15 7.5 H-C*C-H
		18 1.5 H-C*C*H
4.02	22	diastereotopic -12.4 H-C-H
4.61	24	
3.75	21	
2.66	6	diastereotopic -12.4 H-C-H

ChemNMR ¹³C Estimation

Estimation quality is indicated by color: **good**, **medium**, **rough**



Protocol of the C-13 NMR Prediction: (Lib=S)

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
C	113.5	-8.5	tetrahydrofuran
		18.2	2 alpha -C from aliphatic
		98.0	2 alpha -O from aliphatic
		9.3	1 beta -1:C*C*C*C*1 from aliphatic
		18.8	2 beta -C from aliphatic
		10.1	1 beta -O from aliphatic
		-2.5	1 gamma -C from aliphatic
		-6.2	1 gamma -O from aliphatic
		-0.3	1 delta -1:C-O-C=C=1 from aliphatic
		0.3	1 delta -O from aliphatic
		-23.7	general corrections
C	102.2	-8.5	tetrahydrofuran
		18.2	2 alpha -C from aliphatic
		98.0	2 alpha -O from aliphatic
		7.5	1 beta -1:C-O-C=C=1 from aliphatic
		18.8	2 beta -C from aliphatic
		-2.5	1 gamma -C from aliphatic
		-6.2	1 gamma -O from aliphatic
		0.3	1 delta -1:C*C*C*C*1 from aliphatic
		0.3	1 delta -O from aliphatic
		-23.7	general corrections
C	154.4	143.0	2-furan
		13.6	1 -CCCC
		-2.2	general corrections
CH	141.5	143.0	2-furan
		-2.4	1 -CCCC
CH	105.5	109.9	general corrections
		109.9	2-furan
		-5.4	1 -CCCC
CH	110.0	109.9	general corrections
		109.9	2-furan
		0.1	1 -CCCC
		0.0	general corrections
CH2	27.1	-11.2	tetrahydrofuran
		18.2	2 alpha -C from aliphatic
		18.8	2 beta -C from aliphatic
		20.2	2 beta -O from aliphatic
		-2.6	1 gamma -1:C*C*C*C*1 from aliphatic
		-2.5	1 gamma -C from aliphatic
		-12.4	2 gamma -O from aliphatic
		-0.3	1 delta -1:C-O-C=C=1 from aliphatic
		-1.1	general corrections
CH2	31.8	-11.2	tetrahydrofuran
		18.2	2 alpha -C from aliphatic
		18.8	2 beta -C from aliphatic
		20.2	2 beta -O from aliphatic
		-5.0	1 gamma -1:C-O-C=C=1 from aliphatic
		-2.5	1 gamma -C from aliphatic
		-6.2	1 gamma -O from aliphatic
		0.3	1 delta -1:C*C*C*C*1 from aliphatic
		0.3	1 delta -O from aliphatic
		-1.1	general corrections
C	145.5	128.5	1-benzene
		30.3	1 -OCC
		-14.4	1 -O-C
		-1.1	1 -C-O
		2.2	general corrections
C	150.1	128.5	1-benzene
		-14.3	1 -OCC
		33.5	1 -O-C
		0.2	1 -C-O
		2.2	general corrections
C	120.7	128.5	1-benzene
		-8.4	1 -OCC
		1.0	1 -O-C
		12.4	1 -C-O
		-12.8	general corrections
CH	113.8	128.5	1-benzene
		0.6	1 -OCC
		-14.4	1 -O-C
		-1.2	1 -C-O
		0.3	general corrections
CH	112.7	128.5	1-benzene
		-14.3	1 -OCC
		1.0	1 -O-C
		0.2	1 -C-O
		-2.7	general corrections
CH	117.3	128.5	1-benzene
		0.6	1 -OCC
		-17.7	1 -O-C
		-1.2	1 -C-O
		-2.9	general corrections
CH2	67.7	-2.3	aliphatic
		9.1	1 alpha -C
		49.0	1 alpha -O
		9.4	1 beta -C
		20.2	2 beta -O
		-2.6	1 gamma -1:C*C*C*C*1
		-5.0	2 gamma -C
		0.3	1 delta -C
		0.3	1 delta -O
		-10.7	general corrections
CH2	65.0	-2.3	aliphatic
		24.3	1 alpha -1:C*C*C*C*1
		49.0	1 alpha -O
		0.3	1 delta -O
		-6.3	general corrections
CH3	56.1	-2.3	aliphatic
		49.0	1 alpha -O
		9.3	1 beta -1:C*C*C*C*1
		0.3	1 delta -O
		-0.2	general corrections
CH2	43.0	-2.3	aliphatic
		17.0	1 alpha -1:C-O-C=C=1
		9.1	1 alpha -C
		9.4	1 beta -C
		20.2	2 beta -O
		-5.0	2 gamma -C
		0.3	1 delta -C
		0.3	1 delta -O
		-6.0	general corrections

Figure S7. calculated NMR spectra of model M7

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
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Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
OR	4.94	4.20	alcohol
		1.20	1 -CCO
		-0.46	general corrections
OR	5.27	4.20	alcohol
		1.10	1 -C-C*H
		-0.03	general corrections
OR	5.27	4.20	alcohol
		1.10	1 -C-C*H
		-0.03	general corrections
CH	7.49	7.38	2-furan
		-0.17	1 -C
		0.28	general corrections
CH	6.11	6.30	2-furan
		-0.42	1 -C
		0.23	general corrections
CH	6.33	6.30	2-furan
		-0.12	1 -C
		0.13	general corrections
CH2 2.29, 2.045000		1.85	tetrahydrofuran
		0.38	1 beta -0.1.C*CH*CH*CH*CH*1 from methylene
		-0.06	1 beta -C from methylene
CH2 2.29, 2.045000		1.85	tetrahydrofuran
		0.38	1 beta -0.1.C*CH*CH*CH*CH*1 from methylene
		-0.06	1 beta -C from methylene
CH	6.97	7.26	1-benzene
		0.00	1 -O-C
		-0.38	1 -O-C
		0.00	1 -C-O
		0.09	general corrections
CH	6.87	7.26	1-benzene
		-0.38	1 -O-C
		0.00	1 -O-C
		0.03	1 -C-O
		-0.04	general corrections
CH	6.97	7.26	1-benzene
		0.00	1 -O-C
		-0.38	1 -O-C
		0.00	1 -C-O
		0.09	general corrections
CH	6.67	7.26	1-benzene
		0.00	1 -O-C
		-0.32	1 -O-C
		0.00	1 -C-O
		-0.27	general corrections
CH	6.87	7.26	1-benzene
		-0.38	1 -O-C
		0.00	1 -O-C
		0.03	1 -C-O
		-0.04	general corrections
CH	6.67	7.26	1-benzene
		0.00	1 -O-C
		-0.32	1 -O-C
		0.00	1 -C-O
		-0.27	general corrections
CH2 4.14, 3.895000		2.20	methylene
		2.20	1 alpha -O
		0.38	1 beta -0.1.C*CH*CH*CH*CH*1
		0.13	1 beta -C
		-0.06	1 beta -C
CH2 4.61		1.37	methylene
		1.22	1 alpha -0.1.C*CH*CH*CH*1
		0.20	1 alpha -O
		-0.18	general corrections
CH2 4.61		1.37	methylene
		1.22	1 alpha -0.1.C*CH*CH*CH*1
		0.20	1 alpha -O
		-0.18	general corrections
CH3 3.75		0.86	methyl
		0.87	1 alpha -0.1.C*CH*CH*CH*CH*1
		0.02	general corrections
CH3 3.75		0.86	methyl
		0.87	1 alpha -0.1.C*CH*CH*CH*CH*1
		0.02	general corrections
CH2 3.02, 2.765000		1.37	methylene
		1.07	1 alpha -0.1.C-O-CH-C-CH-1
		0.38	1 beta -0.1.C*CH*CH*CH*CH*1
		0.13	1 beta -O-C
		-0.06	1 beta -C

```
shift  atom index  coupling partner, constant and vector
```

5.94	31		
5.27	33		
5.27	35		
7.49	3		
		2	7.5 H-C ⁺ C-H
		1	1.5 H-C ⁺ CH ⁺ C-H
6.11	1		
		2	7.5 H-C ⁺ C-H
		3	1.5 H-C ⁺ CH ⁺ C-H
6.33	2		
		3	7.5 H-C ⁺ C-H
		1	7.5 H-C ⁺ C-H
2.17	9	diastereotopic	-12.4 H-C-H
		10	7.1 H-CH-CH-H
2.17	10	diastereotopic	-12.4 H-C-H
		9	7.1 H-CH-CH-H
6.97	25		
		23	1.5 H-C ⁺ C ⁺ C-H
6.87	22		
		23	7.5 H-C ⁺ C-H
6.97	16		
		14	1.5 H-C ⁺ C ⁺ C-H
6.67	23		
		22	7.5 H-C ⁺ C-H
		25	1.5 H-C ⁺ C ⁺ C-H
6.87	13		
		14	7.5 H-C ⁺ C-H
6.67	14		
		13	7.5 H-C ⁺ C-H
		16	1.5 H-C ⁺ C ⁺ C-H
4.02	30	diastereotopic	-12.4 H-C-H
4.61	32		
4.62	34		
3.75	19		
3.75	28		
2.89	6	diastereotopic	-12.4 H-C-H

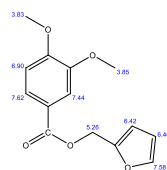
The chemical structure of compound 6 is shown with its ¹³C NMR peaks labeled as follows:

- ¹³C NMR peaks (ppm): 149.8, 119.8, 105.5, 79.3, 77.4, 67.7, 154.9, 111.5, 113.8, 112.360.4, 117.3, 112.3, 107.7, 105.1, 100.0, 113.8, 107.7, 105.0, 117.3, 113.8, 107.7.

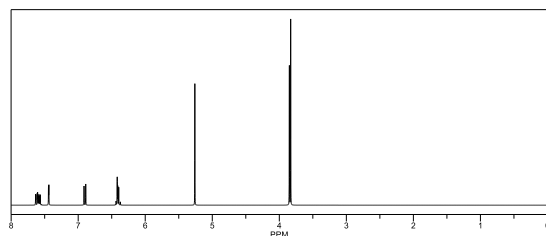
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CR	113.8	Base + Inc.	Comment (ppm rel. to TMS)
	-8.5		tetrahydrofuran
	18.2	2	alpha -C from aliphatic
	90.0	2	alpha -O from aliphatic
	9.3	1	beta -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	18.8	2	beta -O from aliphatic
	10.1	1	beta -O from aliphatic
	-2.5	1	gamma -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-6.2	1	gamma -O from aliphatic
	0.3	1	delta -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-0.3	1	delta -1-C-O-C-O-C-O from aliphatic
	-0.3	1	delta -O from aliphatic
	-23.7		general corrections
CR 111.8	-95.0		tetrahydrofuran
	18.2	2	alpha -C from aliphatic
	9.3	1	beta -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	7.5	1	beta -1-C-O-C-O-C-O from aliphatic
	18.8	2	beta -O from aliphatic
	-2.5	1	gamma -C from aliphatic
	-6.2	1	gamma -O from aliphatic
	0.3	1	delta -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	0.6	1	delta -O from aliphatic
	-23.7		general corrections
CR 154.4	12.4	2	-furan
	13.6	1	-1-C ⁴ OCC
	10.2		general corrections
CR 141.5	143.0		-furan
	-1.4	1	-1-C ⁴ OCC
CR 105.5	105.9		-furan
	-3.4	1	-1-C ⁴ OCC
CR 110.0	105.9		-furan
	0.1	1	-1-C ⁴ OCC
	0.0		general corrections
CR 27.4	-11.2		tetrahydrofuran
	18.2	2	alpha -C from aliphatic
	18.8	2	beta -C from aliphatic
	20.2	2	beta -O from aliphatic
	-2.6	1	gamma -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-2.5	1	gamma -O from aliphatic
	-12.4	2	gamma -O from aliphatic
	0.3	1	delta -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-0.3	1	delta -1-C-O-C-O-C-O from aliphatic
	-0.3		general corrections
CR 29.2	-11.2		tetrahydrofuran
	18.2	2	alpha -C from aliphatic
	18.8	2	beta -C from aliphatic
	20.2	2	beta -O from aliphatic
	-2.6	1	gamma -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-5.0	1	gamma -1-C-O-C-O-C-O from aliphatic
	-2.5	1	gamma -O from aliphatic
	-6.2	1	gamma -O from aliphatic
	0.3	1	delta -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	0.3	1	delta -O from aliphatic
	11.1		general corrections
CR 145.5	128.5		1-benzene
	30.3	1	-OCC
	-14.4	1	-O-C
	-1.1	1	-C-O
	2.2		general corrections
CR 145.5	128.5		1-benzene
	30.3	1	-OCC
	-14.4	1	-O-C
	-1.1	1	-C-O
	2.2		general corrections
CR 150.1	128.5		1-benzene
	-14.3	1	-OCC
	33.5	1	-C-O
	0.2	1	-C-O
	2.2		general corrections
CR 150.1	128.5		1-benzene
	-14.3	1	-OCC
	33.5	1	-O-C
	0.2	1	-C-O
	2.2		general corrections
CR 120.7	-18.4		1-benzene
	-1.0	1	-OCC
	1.0	1	-O-C
	12.4	1	-C-O
	-12.8		general corrections
CR 120.7	-18.4		1-benzene
	-8.4	1	-OCC
	1.0	1	-O-C
	12.4	1	-C-O
	-12.8		general corrections
CR 113.8	128.5		1-benzene
	0.6	1	-OCC
	-14.4	1	-O-C
	-1.2	1	-C-O
	0.3		general corrections
CR 112.7	-14.3		1-benzene
	-14.3	1	-OCC
	1.0	1	-O-C
	0.2	1	-C-O
	2.7		general corrections
CR 113.8	128.5		1-benzene
	0.6	1	-OCC
	-14.4	1	-O-C
	-1.2	1	-C-O
	0.3		general corrections
CR 117.3	128.5		1-benzene
	0.6	1	-OCC
	-7.7	1	-O-C
	-1.2	1	-C-O
	-2.9		general corrections
CR 112.7	-14.3		1-benzene
	-14.3	1	-OCC
	1.0	1	-O-C
	0.2	1	-C-O
	2.7		general corrections
CR 117.3	128.5		1-benzene
	0.6	1	-OCC
	-7.7	1	-O-C
	-1.2	1	-C-O
	-2.9		general corrections
CR 67.7	-2.3		aliphatic
	9.1	1	alpha -C
	49.0	1	alpha -O
	9.4	1	beta -C
	20.2	2	beta -O
	-2.6	1	gamma -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-5.0	2	gamma -C
	0.3	1	delta -C
	0.3	1	delta -O
	-10.7		general corrections
CR2 65.0	-2.3		aliphatic
	9.1	1	alpha -C
	49.0	1	alpha -O
	9.3	1	beta -C
	20.2	2	beta -O
	-2.6	1	gamma -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-5.0	2	gamma -C
	0.3	1	delta -C
	0.3	1	delta -O
	-10.7		general corrections
CR2 65.0	-2.3		aliphatic
	9.1	1	alpha -C
	49.0	1	alpha -O
	9.3	1	beta -C
	20.2	2	beta -O
	-2.6	1	gamma -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-5.0	2	gamma -C
	0.3	1	delta -C
	0.3	1	delta -O
	-10.7		general corrections
CR2 67.7	-2.3		aliphatic
	9.1	1	alpha -C
	49.0	1	alpha -O
	9.4	1	beta -C
	20.2	2	beta -O
	-2.6	1	gamma -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-5.0	2	gamma -C
	0.3	1	delta -C
	0.3	1	delta -O
	-10.7		general corrections
CR2 45.4	-2.3		aliphatic
	9.1	1	alpha -C
	49.0	1	alpha -O
	9.3	1	beta -C
	20.2	2	beta -O
	-2.6	1	gamma -1-C ⁴ C ⁴ C ⁴ C ⁴ C ⁴ from aliphatic
	-5.0	2	gamma -C
	0.3	1	delta -C
	0.3	1	delta -O
	-10.7		general corrections

Figure S8. calculated NMR spectra of model **M8**

ChemNMR ¹H Estimation

Estimation quality is indicated by color: **good**, **medium**, **rough**

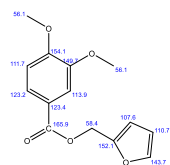


Protocol of the H-1 NMR Prediction (Lib=SU Solvent=DMSO 300 MHz):

Node	Shift	Base + Inc.	Comment (ppm rel. to TMS)
CH 7.58	7.38	2-furan	
	-0.08	1 -C-O	
	0.28	general corrections	
CH 6.42	6.30	2-furan	
	-0.11	1 -C-O	
	0.23	general corrections	
CH 6.40	6.30	2-furan	
	-0.05	1 -C-O	
	0.15	general corrections	
CH 7.44	7.26	1-benzene	
	-0.38	1 -O-C	
	0.00	1 -O-C	
	0.71	1 -C(=O)OC	
	-0.15	general corrections	
CH 6.90	7.26	1-benzene	
	0.00	1 -O-C	
	-0.38	1 -O-C	
	0.11	1 -C(=O)OC	
	-0.09	general corrections	
CH 7.62	7.26	1-benzene	
	-0.32	1 -O-C	
	0.00	1 -O-C	
	0.71	1 -C(=O)OC	
	-0.03	general corrections	
CH2 5.26	1.37	methylene	
	1.07	1 alpha -1:C-O-C=C=1	
	2.92	1 alpha -OC(=O)-1:C*C*C*C*C*1	
	0.10	general corrections	
CH3 3.85	0.86	methyl	
	2.87	1 alpha -O-1:C*C*C*C*C*1	
	0.12	general corrections	
CH3 3.83	0.86	methyl	
	2.87	1 alpha -O-1:C*C*C*C*C*1	
	0.10	general corrections	

1H NMR Coupling Constant Prediction

shift	atom index	coupling partner, constant and vector
7.58	16	17 7.5 H-C*C-H
		18 1.5 H-C*CH*C-H
6.42	18	17 7.5 H-C*C-H
		16 1.5 H-C*CH*C-H
6.40	17	16 7.5 H-C*C-H
		18 7.5 H-C*C-H
7.44	5	3 1.5 H-C*C*C-H
6.90	2	3 7.5 H-C*C-H
7.62	3	2 7.5 H-C*C-H
		5 1.5 H-C*C*C-H
5.26	13	
3.85	10	
3.83	19	

ChemNMR ¹³C Estimation

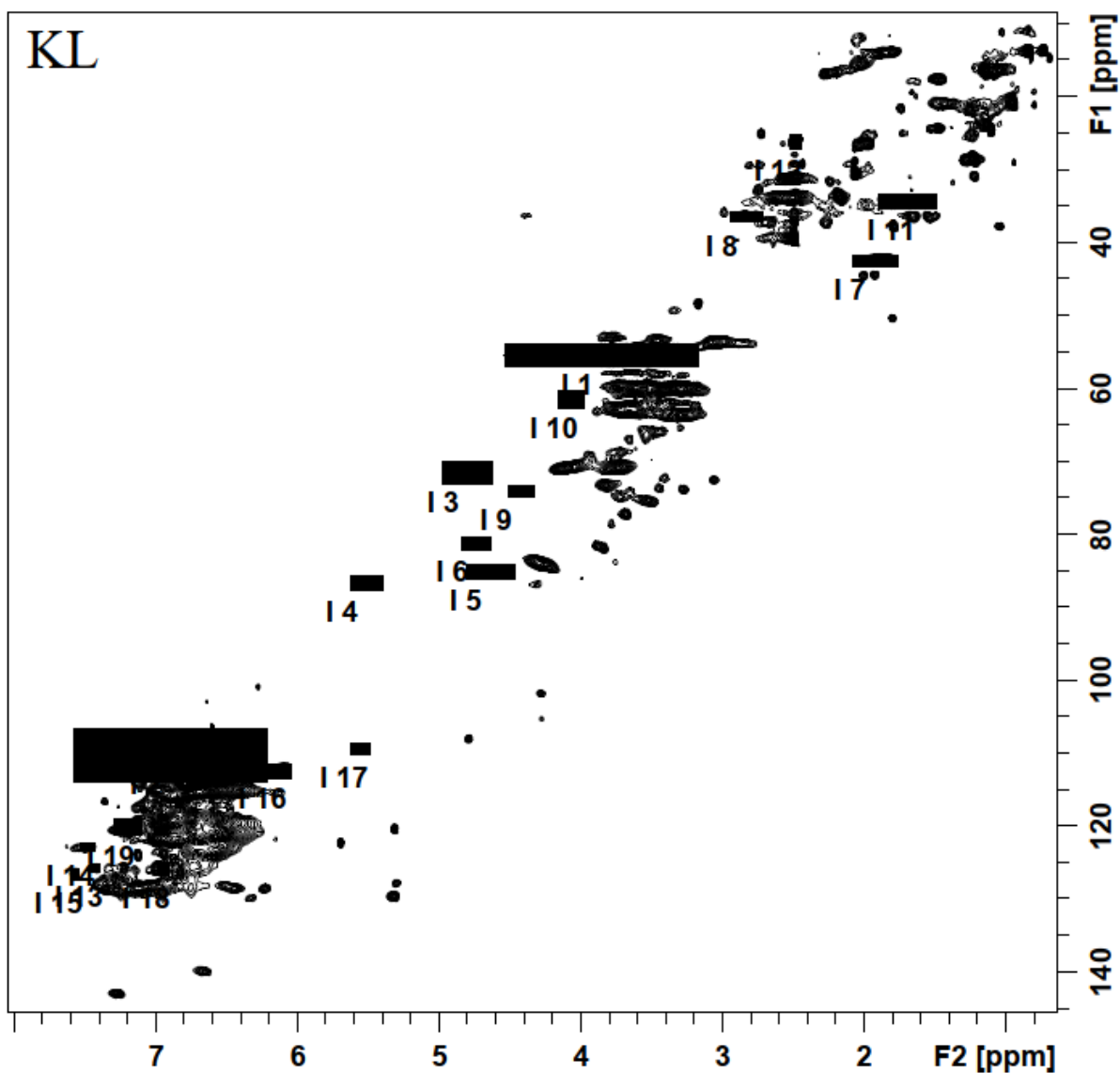


Figure S10. integral of typical structures in KL (nlev = 50, lev0 = 20, toplev = 100 %)

Detailed data set for Figure S10:

Name = KL xzc12-4 ExpNo = 2 ProcNo = 1

Baseline = 0 Noise = 0

#	SI_F1	row1	row2	row1(ppm)	row2(ppm)	abs. Int.	Integral	norm. Int.	Mode
SI_F2	col1	col2	col1(ppm)	col2(ppm)					

Methoxyl (I1)	1024	739	754	56.87234	53.74060	7.4108e+13	7.0675e+07	300	a
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	1024	578	706	4.53892	3.17355				
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G2(I2)	1024	468	503	113.83346	106.57680	2.2984e+13	2.1919e+07	93.043	a
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	1024	294	421	7.57623	6.21993				
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Aa(I3)	1024	663	677	72.88113	70.01336	1.7135e+12	1.6341e+06	6.9364	a
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	1024	537	570	4.98152	4.62657				
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Bα(I4)	1024	593	603	87.71448	85.63781	5.2104e+11	4.969e+05	2.1092 a
	1024	476	498	5.63634	5.39155			
Cα(I5)	1024	601	610	86.03336	84.05559	8.9228e+11	8.5095e+05	3.6121 a
	1024	553	584	4.80405	4.47970			
C'α(I6)	1024	620	628	81.97892	80.19892	3.2991e+11	3.1463e+05	1.3355 a
	1024	550	569	4.84076	4.63881			
SRβ(I7)	1024	804	812	43.21573	41.53674	1.2648e+12	1.2062e+06	5.1202 a
	1024	807	837	2.08733	1.76134			
Arα(I8)	1024	834	840	37.01077	35.76978	3.9495e+11	3.7666e+05	1.5988 a
	1024	727	748	2.94367	2.71499			
AGα(I9)	1024	654	662	74.79440	73.24104	2.9683e+11	2.8307e+05	1.2016 a
	1024	580	597	4.51274	4.33490			
Dγ(I10)	1024	712	723	62.53218	60.29508	7.1084e+11	6.7791e+05	2.8776 a
	1024	613	630	4.16823	3.97984			
DHCAβ(I11)	1024	842	852	35.36743	33.23687	2.0074e+12	1.9144e+06	8.1264 a
	1024	825	863	1.89965	1.48361			
AVβ(I12)	1024	881	890	27.05822	25.14071	2.3597e+11	2.2504e+05	0.95525 a
	1024	766	773	2.52763	2.44913			
V6(I13)	1024	408	416	126.53445	124.84582	2.1618e+11	2.0616e+05	0.87512 a
	1024	302	313	7.49588	7.37565			
VA6(I14)	1024	423	428	123.44588	122.30872	1.6251e+11	1.5498e+05	0.65786 a
	1024	298	307	7.53416	7.43899			
Bi6(I16)	1024	405	411	127.04691	125.78339	1.6579e+11	1.5811e+05	0.67115 a
	1024	292	297	7.60157	7.55002			
E-EEα(I17)	1024	471	480	113.33772	111.31609	5.9059e+11	5.6323e+05	2.3908 a
	1024	421	437	6.21767	6.05112			
Z-EEα(I18)	1024	486	493	110.05257	108.59953	2.5104e+11	2.3941e+05	1.0162 a
	1024	477	489	5.61890	5.49201			
SB1α(I19)	1024	407	416	126.73103	124.83575	7.7581e+11	7.3987e+05	3.1406 a
	1024	349	356	6.99091	6.91557			
SB5β(I20)	1024	433	444	121.17155	118.96039	1.3081e+12	1.2475e+06	5.2952 a
	1024	320	338	7.30020	7.10987			

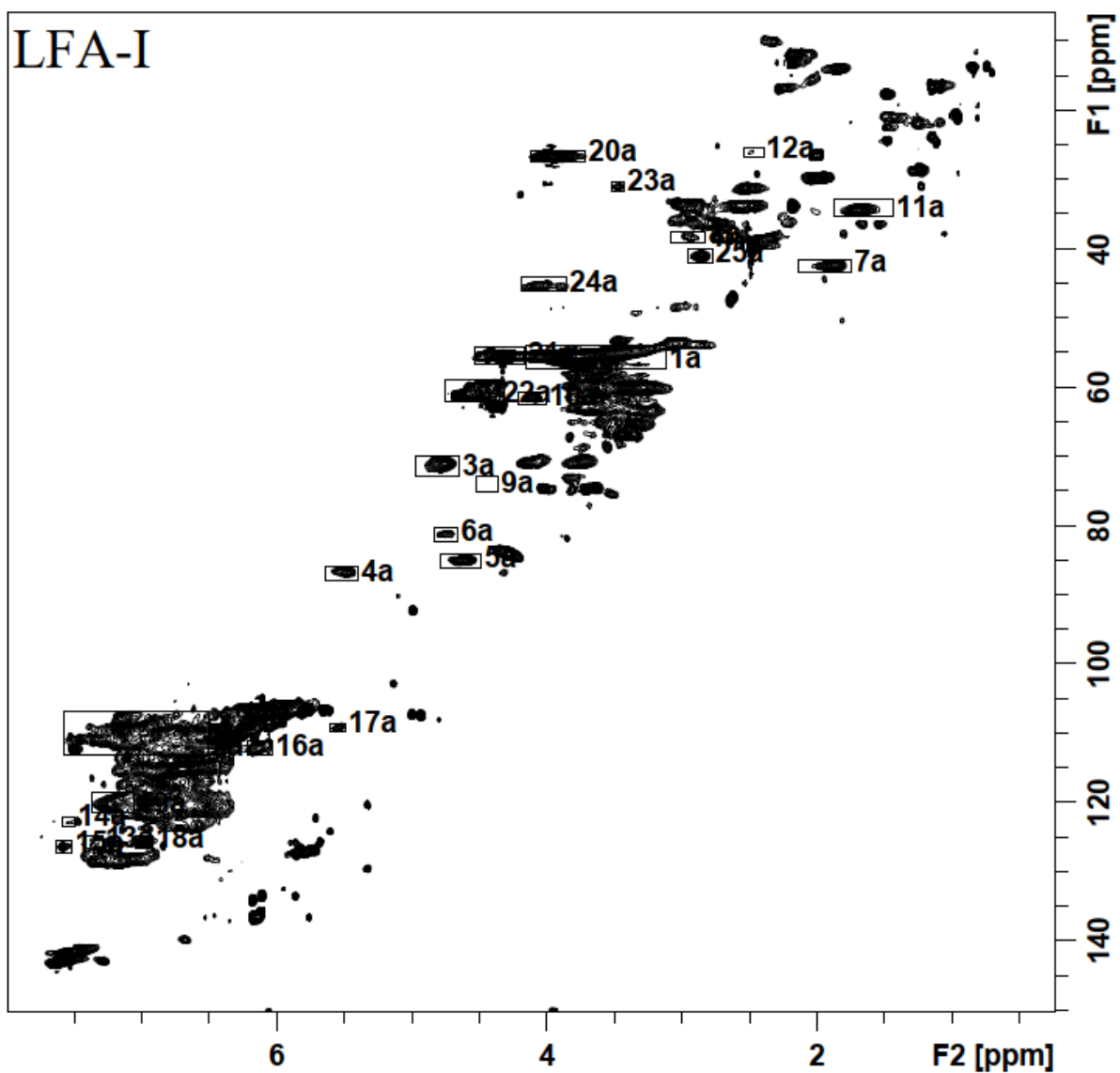


Figure S11. integral of typical aromatic units in LFA-I (nlev = 50, lev0 = 20, toplev = 100 %)

Detailed data set for Figure S11:

Name = LFA-I xzc33-S ExpNo = 2 ProcNo = 1

Baseline = 0 Noise = 0

#	SI_F1	row1	row2	row1(ppm)	row2(ppm)	abs. Int.	Integral	norm. Int.	Mode
	SI_F2	col1	col2	col1(ppm)	col2(ppm)				

Methoxyl(I1)	1024	738	753	57.07990	54.00709	5.9417e+13	5.6665e+07	300	a
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	1024	614	710	4.15006	3.12031				
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G2(I2)	1024	472	501	113.11025	106.89172	1.8077e+13	1.7239e+07	91.27	a
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	1024	294	393	7.57550	6.51197				
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Aα(I3)	1024	664	677	72.67445	70.06174	1.2703e+12	1.2115e+06	6.414 a
	1024	537	567	4.96677	4.65019			
Bα(I4)	1024	592	601	87.80064	85.94424	4.7693e+11	4.5483e+05	2.408 a
	1024	475	497	5.63108	5.39754			
Cα(I5)	1024	601	609	85.94424	84.22535	7.5991e+11	7.2471e+05	3.8368 a
	1024	555	582	4.77474	4.49449			
C'α(I6)	1024	619	627	82.09393	80.51256	2.6982e+11	2.5732e+05	1.3624 a
	1024	550	566	4.83183	4.66576			
SRβ(I7)	1024	804	812	43.29043	41.63521	9.814e+11	9.3594e+05	4.9551 a
	1024	802	838	2.13209	1.75063			
Aα(I8)	1024	824	831	39.02505	37.68814	3.7038e+11	3.5322e+05	1.87 a
	1024	714	737	3.07212	2.83371			
AGα(I9)	1024	653	663	75.12163	73.01823	2.1066e+11	2.009e+05	1.0636 a
	1024	579	594	4.51650	4.36076			
Dγ(II0)	1024	713	721	62.42607	60.77340	4.6521e+11	4.4366e+05	2.3489 a
	1024	609	627	4.19903	4.00735			
DHCAβ(II1)	1024	842	854	35.33900	32.87951	1.6852e+12	1.6071e+06	8.5086 a
	1024	821	867	1.92651	1.43658			
AVβ(II2)	1024	883	889	26.74288	25.40851	1.2242e+11	1.1675e+05	0.6181 a
	1024	765	777	2.53260	2.40511			
V6(II3)	1024	409	415	126.34295	125.13434	1.8565e+11	1.7705e+05	0.93733 a
	1024	309	320	7.41127	7.29067			
VA6(II4)	1024	423	428	123.42866	122.40794	9.5628e+10	91198	0.48283 a
	1024	293	301	7.58153	7.49676			
Bi6(II5)	1024	405	412	127.11896	125.62713	1.9833e+11	1.8914e+05	1.0014 a
	1024	289	299	7.62921	7.51795			
E-EEα(II6)	1024	472	481	113.06442	111.18002	6.0827e+11	5.8009e+05	3.0712 a
	1024	421	438	6.21463	6.02920			
Z-EEα(II7)	1024	488	493	109.76671	108.58895	1.5602e+11	1.4879e+05	0.78776 a
	1024	478	488	5.60535	5.49939			
SB1α(II8)	1024	408	416	126.49082	124.92048	5.5834e+11	5.3247e+05	2.8191 a
	1024	344	355	7.03583	6.91927			

SB5 β (I19) 1024 433 445 121.18516 118.70278 1.2164e+12 1.1601e+06 **6.1417 a**
1024 305 332 7.45604 7.16186

CH2 in pFA(I20) 1024 880 887 27.37992 25.91314 1.4712e+13 1.403e+07 **74.28 a**
1024 617 654 4.11856 3.72374

CH2OH in pFA(I21) 1024 741 752 56.59333 54.14869 8.2265e+12 7.8454e+06 **41.536 a**
1024 578 612 4.52887 4.17275

CH2OCH2 in pFA(I22) 1024 715 729 61.97152 59.16019 4.1691e+12 3.9759e+06 **21.05 a**
1024 558 595 4.74564 4.35081

New peak 2(I23) 1024 860 865 31.53580 30.43572 1.1151e+11 1.0635e+05 **0.56303 a**
1024 673 681 3.51471 3.42955

New peak 3(I24) 1024 791 800 45.95916 44.24791 5.7779e+11 5.5102e+05 **2.9173 a**
1024 611 641 4.18050 3.85535

New peak 4(I25) 1024 811 819 41.92551 40.09203 4.84e+11 4.6157e+05 **2.4437 a**
1024 726 742 2.94957 2.77926

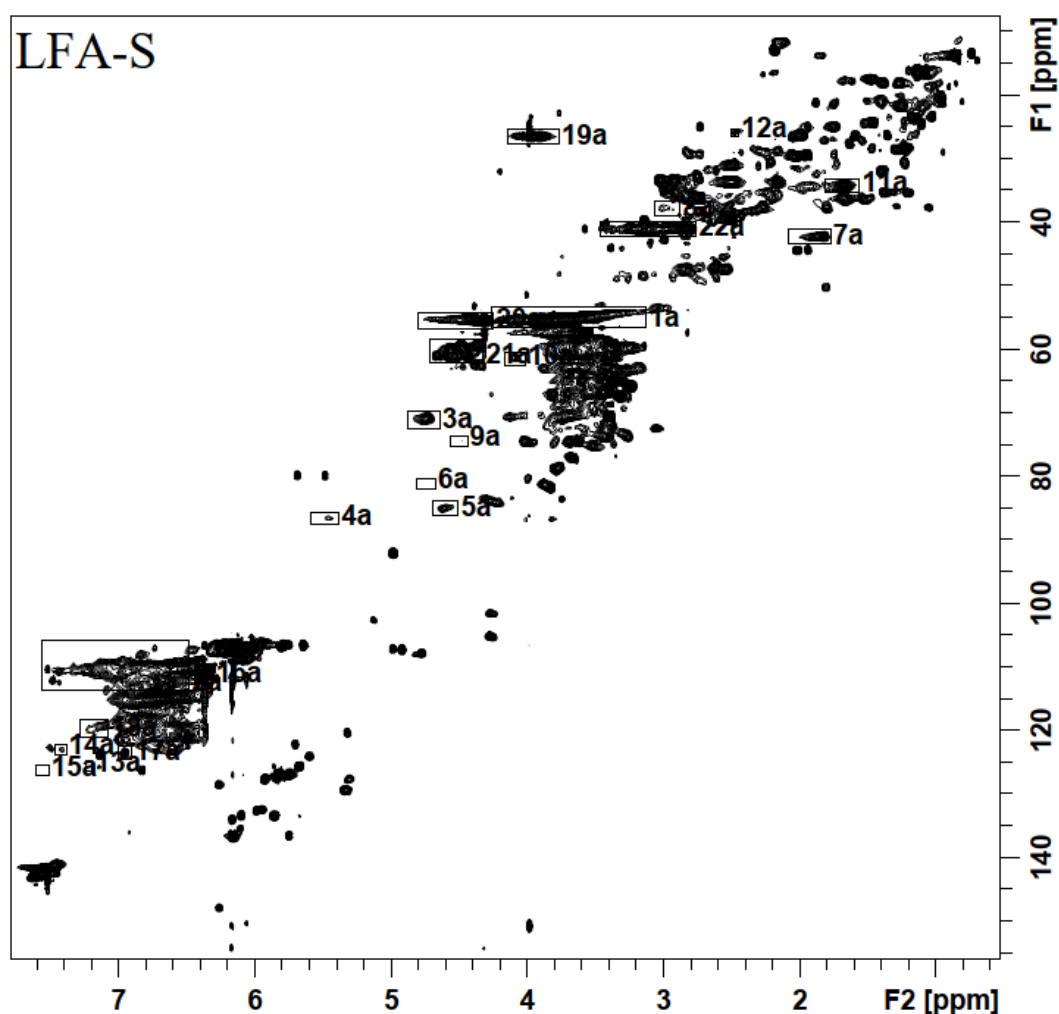


Figure S12. integral of typical aromatic units in LFA-S (nlev = 50, lev0 = 20, toplev = 100 %)

Detailed data set for Figure S12:

Name = LFA-S xzc33-L ExpNo = 2 ProcNo = 1

Baseline = 0 Noise = 0

#	SI_F1	row1	row2	row1(ppm)	row2(ppm)	abs. Int.	Integral	norm. Int.	Mode
SI_F2	col1	col2	col1(ppm)	col2(ppm)					
Methoxyl(I1)	1024	741	755	56.33739	53.46749	3.4298e+13	3.271e+07	300	a
	1024	604	709	4.24803	3.12895				
G2(I2)	1024	472	497	112.91586	107.64885	1.0506e+13	1.0019e+07	91.892	a
	1024	296	384	7.54531	6.60333				
Aα(I3)	1024	665	677	72.34975	69.77652	6.9156e+11	6.5953e+05	6.0489	a
	1024	546	567	4.87054	4.64776				
Bα(I4)	1024	592	601	87.63552	85.69382	2.2353e+11	2.1317e+05	1.9551	a
	1024	477	499	5.61500	5.37349				
Cα(I5)	1024	600	611	86.08216	83.62267	4.1166e+11	3.9259e+05	3.6007	a
	1024	553	583	4.79992	4.47539				
C'α(I6)	1024	618	626	82.14528	80.59537	1.487e+11	1.4181e+05	1.3006	a
	1024	552	568	4.80493	4.63447				
SRβ(I7)	1024	804	814	43.11005	41.08134	7.1584e+11	6.8268e+05	6.2613	a
	1024	816	834	1.98494	1.78934				
Ara(I8)	1024	825	832	38.70461	37.15125	2.5115e+11	2.3951e+05	2.1967	a
	1024	716	729	3.04900	2.91316				
AGα(I9)	1024	652	658	75.17170	73.89760	1.0919e+11	1.0413e+05	0.95505	a
	1024	575	586	4.55936	4.43995				
Dγ(I10)	1024	712	721	62.36367	60.55312	2.6153e+11	2.4941e+05	2.2875	a
	1024	613	626	4.15770	4.02201				
DHCAβ(I11)	1024	843	850	34.99521	33.37225	9.0064e+11	8.5892e+05	7.8777	a
	1024	834	854	1.78934	1.57925				
AVβ(I12)	1024	885	890	26.14826	25.11269	6.6369e+10	63295	0.58051	a
	1024	771	776	2.46033	2.41505				

V6(I13) 1024 408 415 126.43388 124.90428 1.0777e+11 1.0277e+05 **0.9426 a**
 1024 322 330 7.27053 7.18673

VA6(I14) 1024 422 427 123.45518 122.28786 8.1289e+10 77524 **0.71102 a**
 1024 305 311 7.45824 7.38785

Bi6(I15) 1024 405 412 126.89611 125.41914 9.0653e+10 86453 **0.79292 a**
 1024 291 299 7.59788 7.51748

E-EE α (I16) 1024 475 479 112.34071 111.42860 3.8079e+11 3.6315e+05 **3.3307 a**
 1024 400 413 6.43551 6.29877

SB1 α (I17) 1024 417 427 124.42645 122.44919 3.2148e+11 3.0659e+05 **2.8119 a**
 1024 345 356 7.01968 6.91093

SB5 β (I18) 1024 430 444 121.77834 118.70651 6.6448e+11 6.337e+05 **5.8121 a**
 1024 320 341 7.29651 7.06912

CH2 in pFA(I19) 1024 878 888 27.48687 25.41445 1.7224e+13 1.6426e+07 **150.66 a**
 1024 615 649 4.13779 3.76692

CH2OH in pFA(I20) 1024 740 751 56.50079 54.21022 2.1092e+13 2.0115e+07 **184.49 a**
 1024 553 604 4.80005 4.25434

CH2OCH2 in pFA(I21) 1024 714 731 62.06361 58.57322 5.8616e+12 5.5901e+06 **51.27 a**
 1024 561 597 4.70998 4.32852

New peak 7(I22) 1024 809 819 41.99383 40.03048 1.1442e+13 1.0912e+07 **100.08 a**
 1024 678 743 3.45964 2.76029