

Supporting Information for: “A C—H oxidation approach for streamlining synthesis of chiral polyoxygenated motifs ”

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General Information: The following commercially obtained reagents were used as received; Pd(OAc)₂, (Alfa Aesar), (1R,2R)-(-)-[1,2-Cyclohexanediamino-N,N'-bis(3,5-di-*t*-butylsalicylidene)]Chromium(III)Cl, (1S,2S)-(+)-[1,2-Cyclohexanediamino-N,N'-bis(3,5-di-*t*-butylsalicylidene)]Chromium(III)Cl (Aldrich), 5-hexene-1-ol (Aldrich), methyl 10-undecenoate (Aldrich), benzyl bromide (Aldrich), phenyl boronic acid (Aldrich), Novozyme 435, polymer supported (Aldrich), tetrabutylammonium iodide (Aldrich), sodium hydride (95%) (Aldrich), iron ferricyanide (Strem chemicals), potassium carbonate (Fisher), sodium bicarbonate (Fisher), methanesulfonamide (Aldrich), (DHQD)₂PHAL (Aldrich), hexenoic acid methyl ester (TCI), 2-(2-bromoethyl)-1,3-dioxane (TCI), benzoquinone (Aldrich), acetic acid (Fisher). Lewis acid **2** was prepared from commercially available (1R,2R)-(-)-[1,2-Cyclohexanediamino-N,N'-bis(3,5-di-*t*-butylsalicylidene)]Chromium(III) Chloride as previously described.ⁱ Pd(OAc)₂ was stored in a glove box under an argon atmosphere and weighed out in the air prior to use. Commercially available “White Catalyst” (1,2-Bis(phenylsulfinyl)ethane palladium(II) acetate)(**1**) from Aldrich was found to be equivalent to that prepared freshly by the published procedure.ⁱⁱ Solvents 1,4-dioxane, diethyl ether (Et₂O), tetrahydrofuran (THF), and methylene chloride (CH₂Cl₂) were purified prior to use by passage through a bed of activated alumina (Glass Contour, Laguna Beach, California). Ethyl acetate (EtOAc) (Sure/Seal) was obtained from Sigma-Aldrich and used as received. All allylic oxidation reactions were run under air. Achiral gas chromatographic (GC) analyses were performed on Agilent Technologies 6890N Series instrument equipped with FID detectors using a HP-5 (5%-Phenyl)-methylpolysiloxane column (30m, 0.32mm, 0.25μm). Chiral gas chromatographic (GC) analyses were performed on an Agilent Technologies 5890A Series instrument equipped with an FID detector using a J&W Scientific β-cyclodextrin column (30m, 0.25mm, 0.25μm). HPLC analysis was performed on an Agilent Technologies 1100 HPLC system with a model 1100 Quaternary Pump, Diode Array Detector, Thermostat, and Autosampler. Thin-layer chromatography (TLC) was conducted with E. Merck silica gel 60 F254 precoated plates (0.25 mm) and visualized with UV, potassium permanganate, and ceric ammonium molybdate staining. Flash column chromatography was performed as described by Still et al.ⁱⁱⁱ using EM reagent silica gel 60 (230-400 mesh). ¹H NMR spectra were recorded on a Varian Unity 400 (400 MHz) or a Varian Unity 500 (500 MHz), or a Varian Unity Inova 500NB spectrometer and are reported in ppm using solvent as an internal standard (CDCl₃ at 7.26 ppm). Data reported as: s = singlet, d = doublet, t = triplet, q = quartet, m = multiplet, b = broad; coupling constant(s) in Hz; integration. Proton-decoupled ¹³C- NMR spectra were recorded on a Varian Unity-500 (125 MHz) spectrometer and are reported in ppm using solvent as an internal standard (CDCl₃ at 77.0 ppm). IR spectra were recorded as thin films on NaCl plates on a Perkin-Elmer Spectrum BX and are reported in frequency of absorption (cm⁻¹). High-resolution mass spectra were obtained at the University of Illinois Mass Spectrometry Laboratory.

General Procedure for Asymmetric Branched Allylic Acetoxylation: A vial (8 mL borosilicate) was charged with the following: 1,2-Bis(phenylsulfinyl)ethane palladium(II) acetate(**1**) (10 mol%, 0.10 mmol, 50 mg); (1R,2R)-(-)-[1,2-Cyclohexanediamino-N,N'-bis(3,5-di-*t*-butylsalicylidene)]Chromium(III)F(*R,R*-**2**) (10 mol%, 0.10 mmol, 61.6 mg), 1,4-

benzoquinone (2 equiv., 2.0 mmol, 216 mg), an activated 4Å MS bead (~30 mg), and a Teflon® stir bar. A separate vial (2 mL, borosilicate) was charged with the following: substrate (1.0 mmol), AcOH (1.1 equiv., 63 µL), and EtOAc (200 µL). The liquids were transferred to the solids via pipette and the vial rinsed with EtOAc (3 x 100 µL). After carefully stirring for 48 hrs at room temperature, the reaction mixture was transferred to a separatory funnel with ~3 mL EtOAc and diluted with hexanes (200 mL). The organic layer was rinsed with sat. aq. NaHSO₃ (1 x 50 mL) and 5% aq. K₂CO₃ (2 x 50 mL). *Caution should be taken when combining aqueous layers as carbon dioxide is evolved.* The combined aqueous layers were back extracted with hexanes (100 mL). The combined organic layers were dried (MgSO₄), filtered, and reduced *in vacuo*. The resulting oil was re-dissolved in hexanes (50 mL) and extracted again with 5% aq. K₂CO₃ (3 x 10 mL) to remove residual hydroquinone. The organic layer was again dried (MgSO₄), filtered and reduced *in vacuo* to afford a clean mixture of allylic oxidation products and any unreacted starting material from which the B:L, yield, and conversions were determined (¹H NMR). Unless otherwise noted, this material was taken forward without further purification, though each substrate was isolated at least once for characterization purposes. Reported yields and selectivities are an average of at least two runs. Enantiomeric excess was determined by chiral GC (β-cyclodextrin column), as compared to racemic standards generated through our standard branched oxidation chemistry. Slight variations in B:L ratios and ee's were noted based on batch of Cr catalyst.

General Procedure for Cleavage of Allylic Acetates: To a 25 mL flask containing crude allylic acetate (1 mmol, assumed) was added MeOH (5 mL, 0.2 M) and potassium carbonate (0.276 g, 2 mmol). The reaction was vigorously stirred and monitored via thin layer chromatography (TLC). Upon completion, the reaction was transferred to a separatory funnel with methylene chloride (50 mL). Water (15 mL) was added, and the aqueous layer was extracted with methylene chloride (3 x 50 mL). The combined organics were washed with brine (1 x 10 mL), then dried (MgSO₄), filtered, and reduced *in vacuo*. Products were then purified by standard SiO₂ chromatography. While the branched and linear allylic alcohols were commonly separable, it was found that carrying them forward as a mixture had no detrimental effect as the subsequent resolution acylated the linear alcohol rapidly making its separation from branched alcohol trivial. Individual product yields and characterization are reported below.

General Procedure for Resolution with Novozyme 435: To a flame dried round bottom flask containing allylic alcohol to be resolved (1 equiv.) was added vinyl acetate (0.6M) and Novozyme 435 immobilized on polystyrene beads (33.3 mg/1 mmol). The reaction was stirred vigorously at room temperature for 36 hrs. Upon completion, the solid supported enzyme was removed via filtration. The solid support was rinsed thoroughly with diethyl ether and then the filtrate reduced *in vacuo* and purified via standard SiO₂ chromatography. Enantioselectivities were determined by chiral gas chromatographic analysis on the acetylated derivative of each isolated alcohol. It was found that the recovered solid supported enzyme could be used up to 5 times with little diminishment in activity. Individual yields and selectivities are reported below.

General Procedure for Resolution with S. Carlsberg: The active enzyme for resolution was prepared as previously described by Boren and co-workers.^{iv} To a flame dried round bottom flask containing allylic alcohol to be resolved (1 equiv.) was added isopropenyl valerate^v (1.5 equiv), active S. Carlsberg (36 mg/1 mmol), sodium carbonate (1 equiv.) and THF (0.5M). The

reaction was stirred vigorously at room temperature for 60 hrs. Upon completion, the enzyme was removed via filtration. The enzyme was rinsed thoroughly with diethyl ether and then the filtrate reduced *in vacuo* and purified via standard SiO₂ chromatography. Enantioselectivities were determined by chiral gas chromatographic analysis on the acetylated derivative of each isolated alcohol. Individual yields and selectivities are reported below.

ⁱ S. E. Schaus, J. Brånalt, E. N. Jacobsen. *J. Org. Chem.* **1998**, *63*, 403.

ⁱⁱ J. H. Delcamp, M. C. White, *J. Am. Chem. Soc.* **2006**, *128*, 15076.

ⁱⁱⁱ W. C. Still, M. Kahn, A. Mitra, *J. Org. Chem.* **1978**, *43*, 2923.

^{iv} L. Boren, B. Martin-Matute, Y. Xu, A. Cordova, and J.-E. Backvall; *Chemistry: A European Journal*, **2006**, *12*, 225-232.

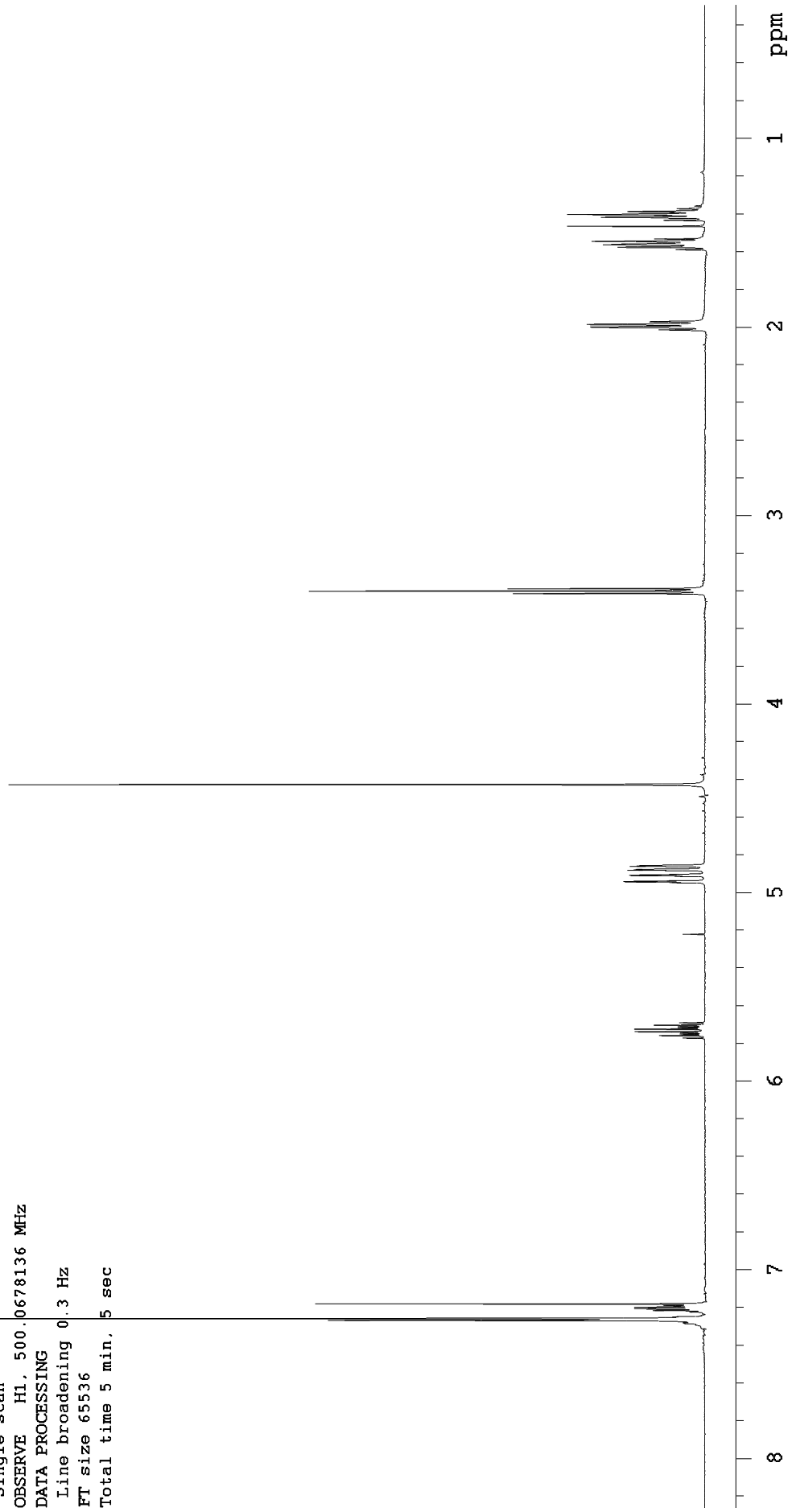
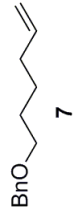
^v Y. F. Wang, J. J. Lalonde, M. Momongan, D. E. Bergbreiter, C. H. Wong *J. Am. Chem. Soc.* **1998**, *110*, 7200.

1-O-Benzyl-5-hexen-1-ol (7)

Pulse Sequence: s2pul

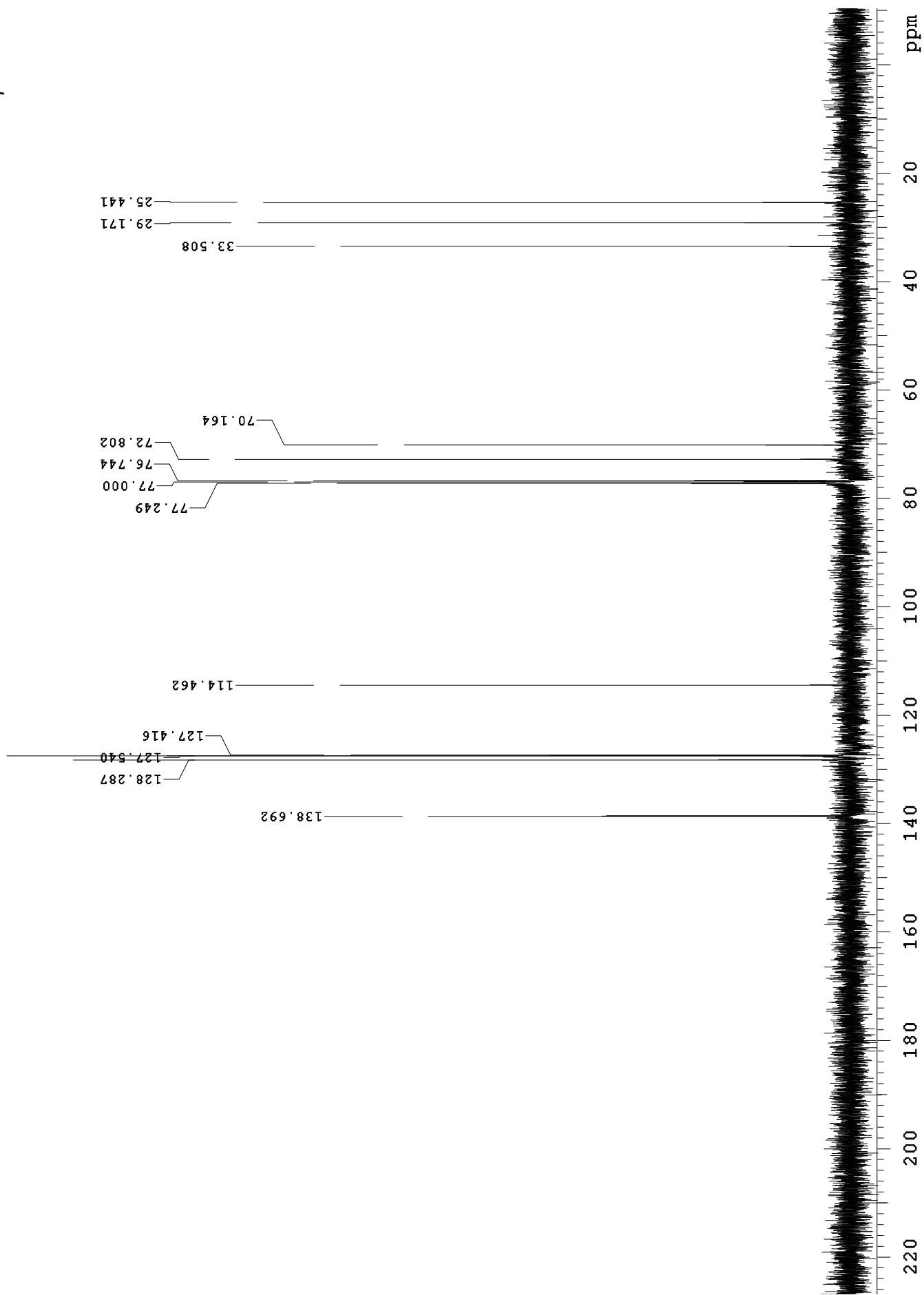
Solvent: CDCl₃
Ambient temperature
File: DJC-7-7-1H
INOVA-500 "sunds1"

Relax. delay 15.000 sec
Pulse 90.0 degrees
Acq. time 4.096 sec
Width 8000.0 Hz
Single scan
OBSERVE H1, 500.0678136 MHz
DATA PROCESSING
Line broadening 0.3 Hz
FT size 65536
Total time 5 min, 5 sec

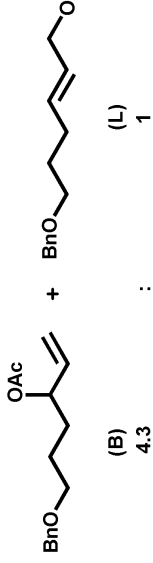


(-)-O-Benzyl-5-hexen-1-ol (7)

Pulse Sequence: s2pul



(4R)-1-O-Benzyl-4-acetoxy-5-hexen-1,4-diol (3-OAc)



Pulse Sequence: s2pul

Solvent: CDCl₃

Ambient temperature

File: DJC-4-301-1

INOVA-500 "sunds1"

Relax. delay 20.000 sec

Pulse 45.0 degrees

Acq. time 4.096 sec

Width 8000.0 Hz

4 repetitions

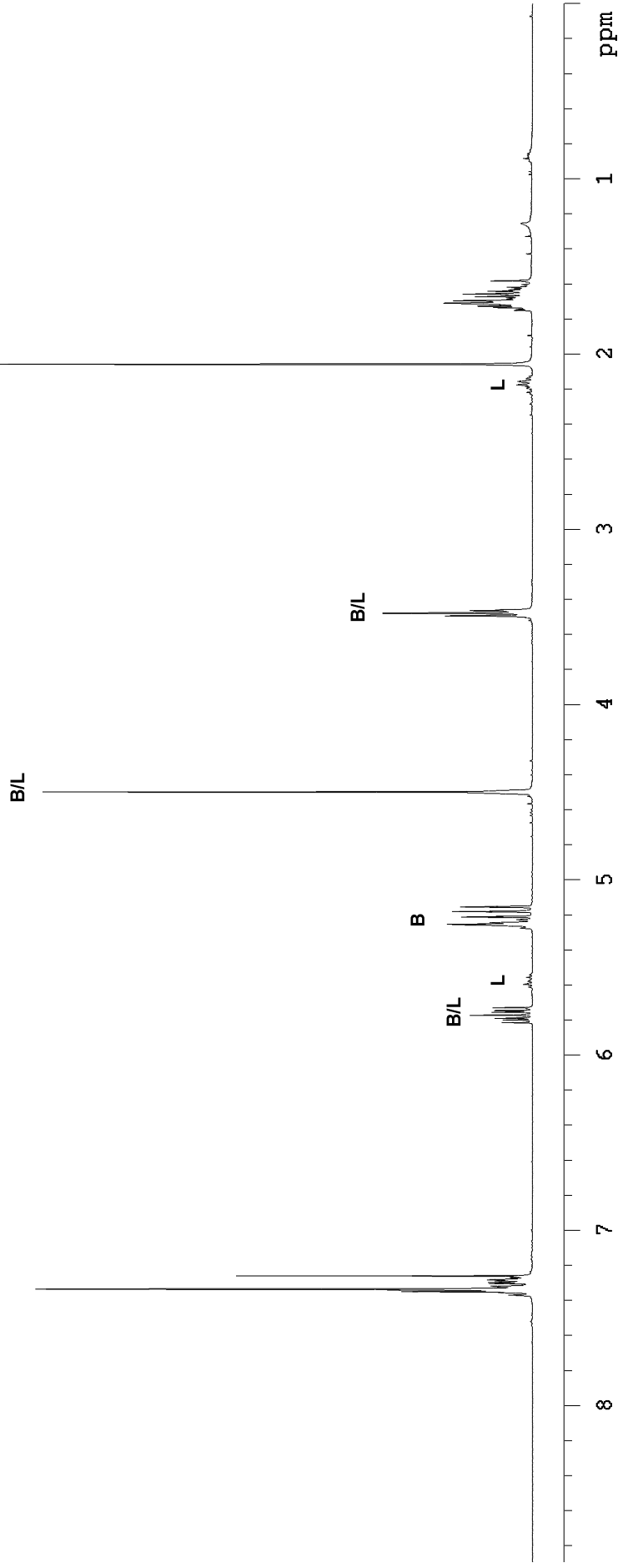
OBSERVE H1, 399.9486781 MHz

DATA PROCESSING

Line broadening 0.3 Hz

FT size 65536

Total time 1 min, 36 sec



(4R)-1-O-Benzyl-4-acetoxy-5-hexen-1,4-diol (3-OAc)

Pulse Sequence: s2pul

Solvent: CDCl₃

Ambient temperature

User: 1-14-87

File: DJC-4-301-13C

INOVA-500 "sunds1"

Relax. delay 5.000 sec

Pulse 53.3 degrees

Acq. time 1.024 sec

Width 32000.0 Hz

28 repetitions

OBSERVE C13, 125.5817522 MHz

DECOUPLE H1, 499.4315638 MHz

Power 44 dB

continuously on

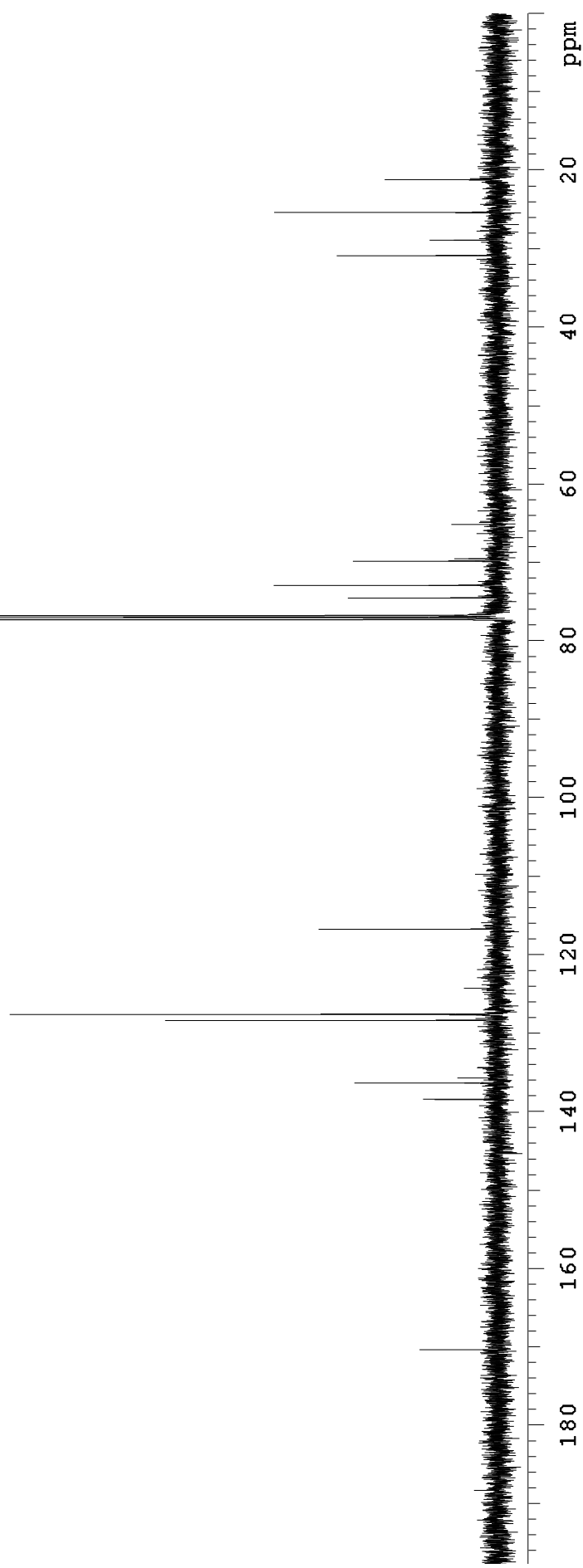
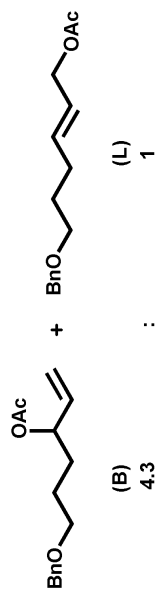
WALTZ-16 modulated

DATA PROCESSING

Line broadening 1.0 Hz

FT size 65536

Total time 1 hr, 40 min, 32 sec



4R)-1-O-Benzyl-5-hexen-1,4-diol ((-)-3)

Pulse Sequence: s2pul

Solvent: CDCl₃

Ambient temperature

File: DJC-6-161-1H

INOVA-500 "sunds1"

Relax. delay 15.000 sec

Pulse 90.0 degrees

Acq. time 4.096 sec

Width 8000.0 Hz

2 repetitions

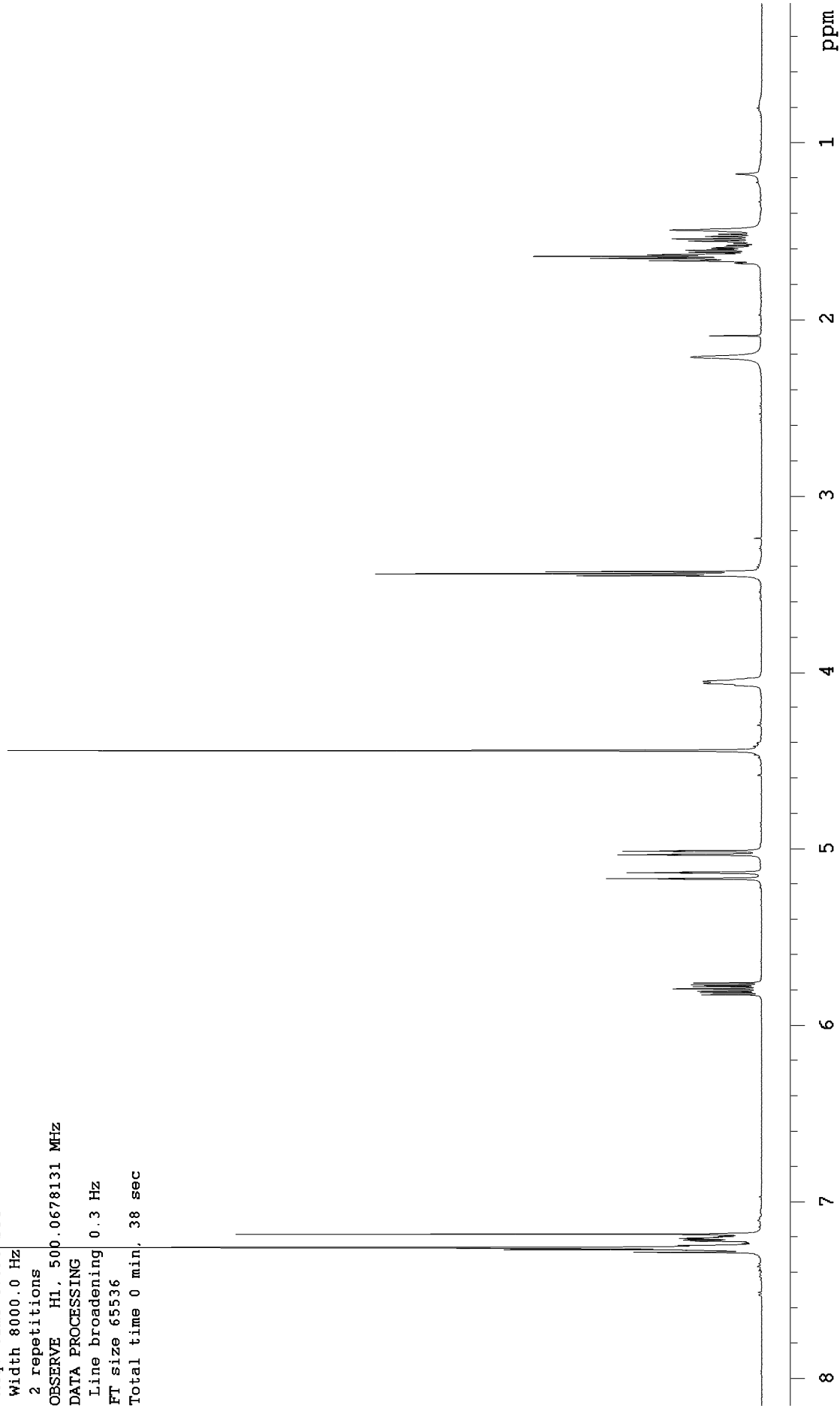
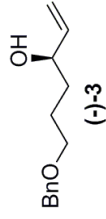
OBSERVE H1, 500.0678131 MHz

DATA PROCESSING

Line broadening 0.3 Hz

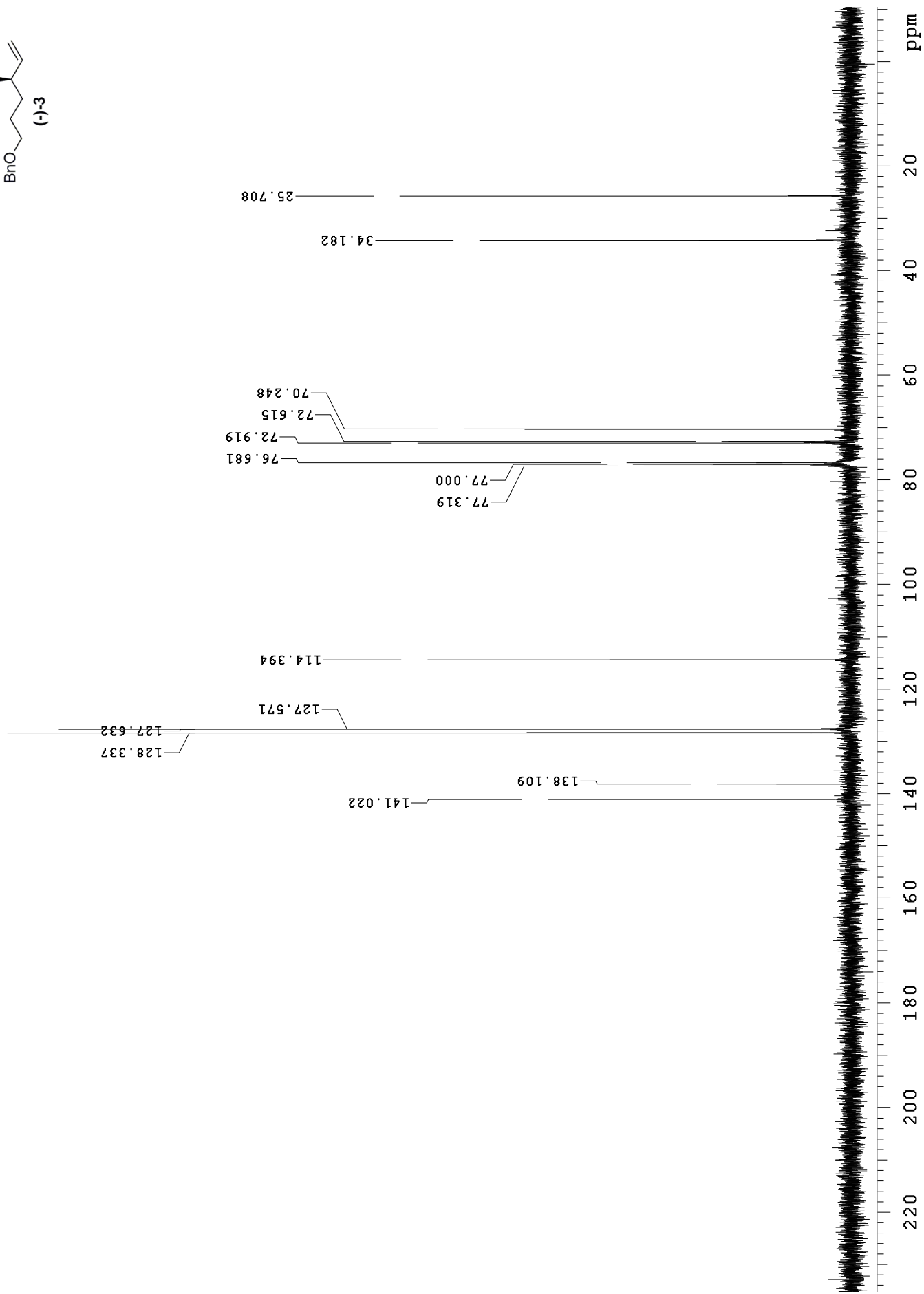
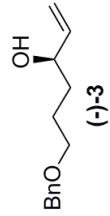
FT size 65536

Total time 0 min, 38 sec

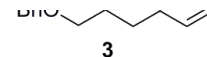


4R)-1-O-Benzyl--5-hexen-1,4-diol ((-)-3)

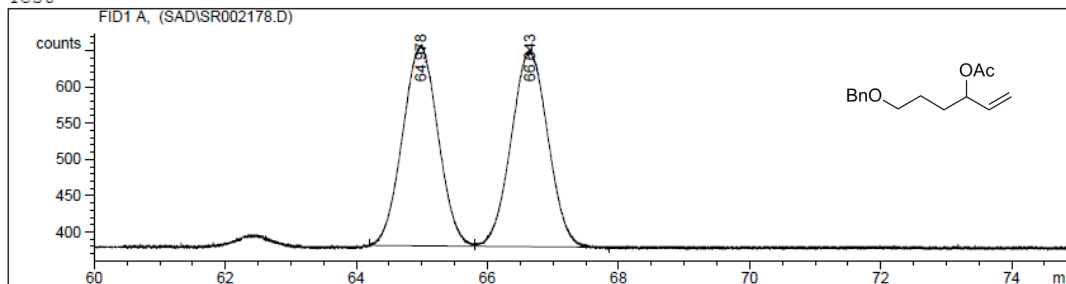
Pulse Sequence: s2pul



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 Acq. Instrument : Chiral GC Inj Volume : Manually
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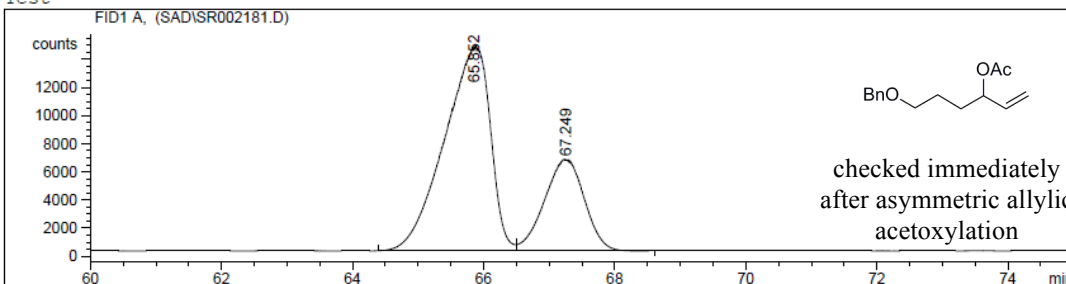
Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	64.978	BV	0.6006	1.03758e4	274.75067	49.70793
2	66.643	VP	0.6327	1.04977e4	268.16049	50.29207

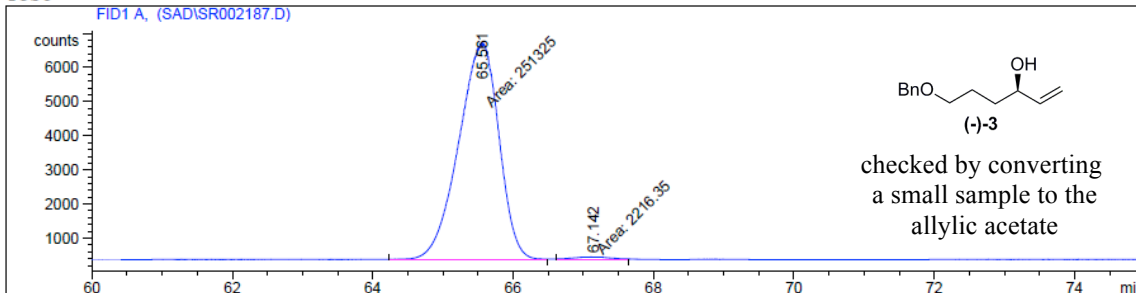
Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	65.852	BV	0.7832	7.09046e5	1.43813e4	72.04422
2	67.249	VB	0.6715	2.75136e5	6478.43213	27.95578

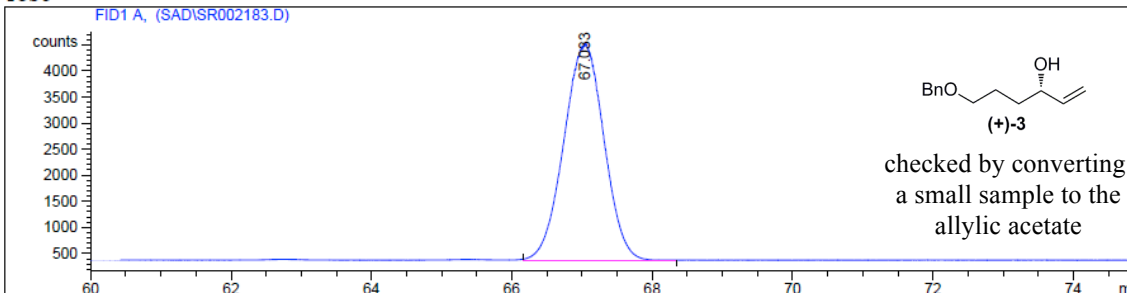
Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	65.561	MM	0.6618	2.51325e5	6329.74902	99.12584
2	67.142	MM	0.5854	2216.35474	63.09792	0.87416

Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
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ethyl (9R)-9-acetoxyundec-10-enoate (9-OAc)

Sequence: s2pul

Solvent: CDCl3

Temp: 19.0 C / 292.1 K

File: DJC-4-145-2

WA-500 "sunds1"

Relax. delay 15.000 sec

ulse 49.2 degrees

Acq. time 4.096 sec

Resolution 8000.0 Hz

Repetitions

Frequency 499.4298992 MHz

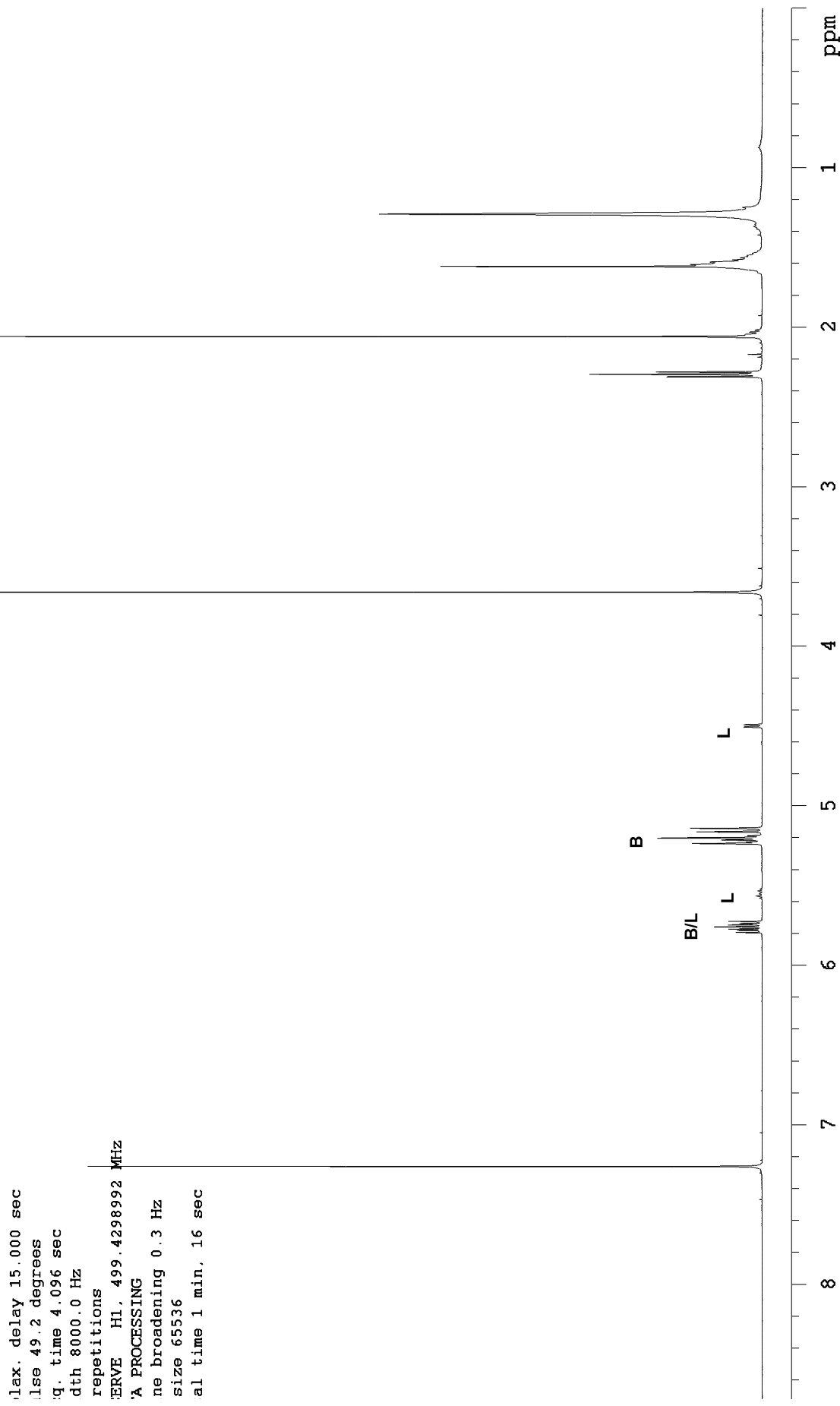
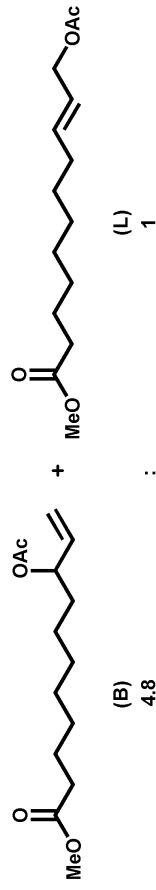
Mode: 1D

Temperature 19.000 C

Time 1 min, 16 sec

File size 65536

Acquisition time 1 min, 16 sec



ethyl (9R)-9-acetoxyundec-10-eneoate (9-OAc)

Use Sequence: s2pul

solvent: CDCl3

ambient temperature

acq. time 1-14-87

file: DJC-4-145-13C

DATA-500 "sunds1"

relax. delay 5.000 sec

pulse 53.3 degrees

acq. time 1.024 sec

width 32000.0 Hz

5 repetitions

SERVE C13, 125.5817522 MHz

COUPLE H1, 499.4315638 MHz

power 44 dB

continuously on

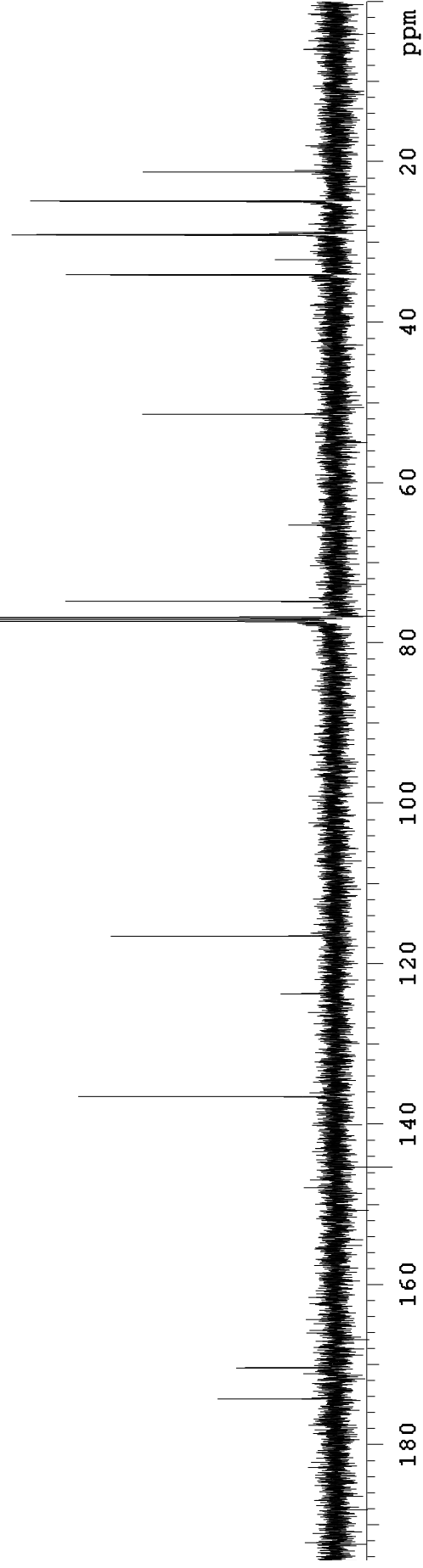
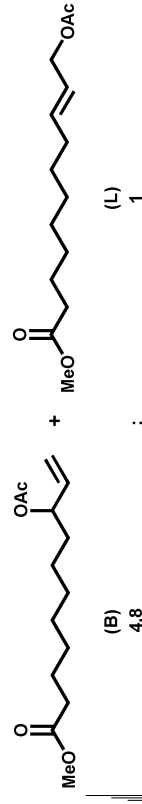
ALTZ-16 modulated

FA PROCESSING

line broadening 1.0 Hz

size 65536

total time 1 hr, 40 min, 32 sec



ethyl (9*R*)-9-hydroxyundec-10-enoate ((-)-9)

Use Sequence: s2pul

solvent: CDCl3

ambient temperature

file: DJC-6-233-1H

AVA-500 "sunds1"

relax. delay 20.000 sec

pulse 90.0 degrees

acq. time 4.096 sec

width 8000.0 Hz

repetitions

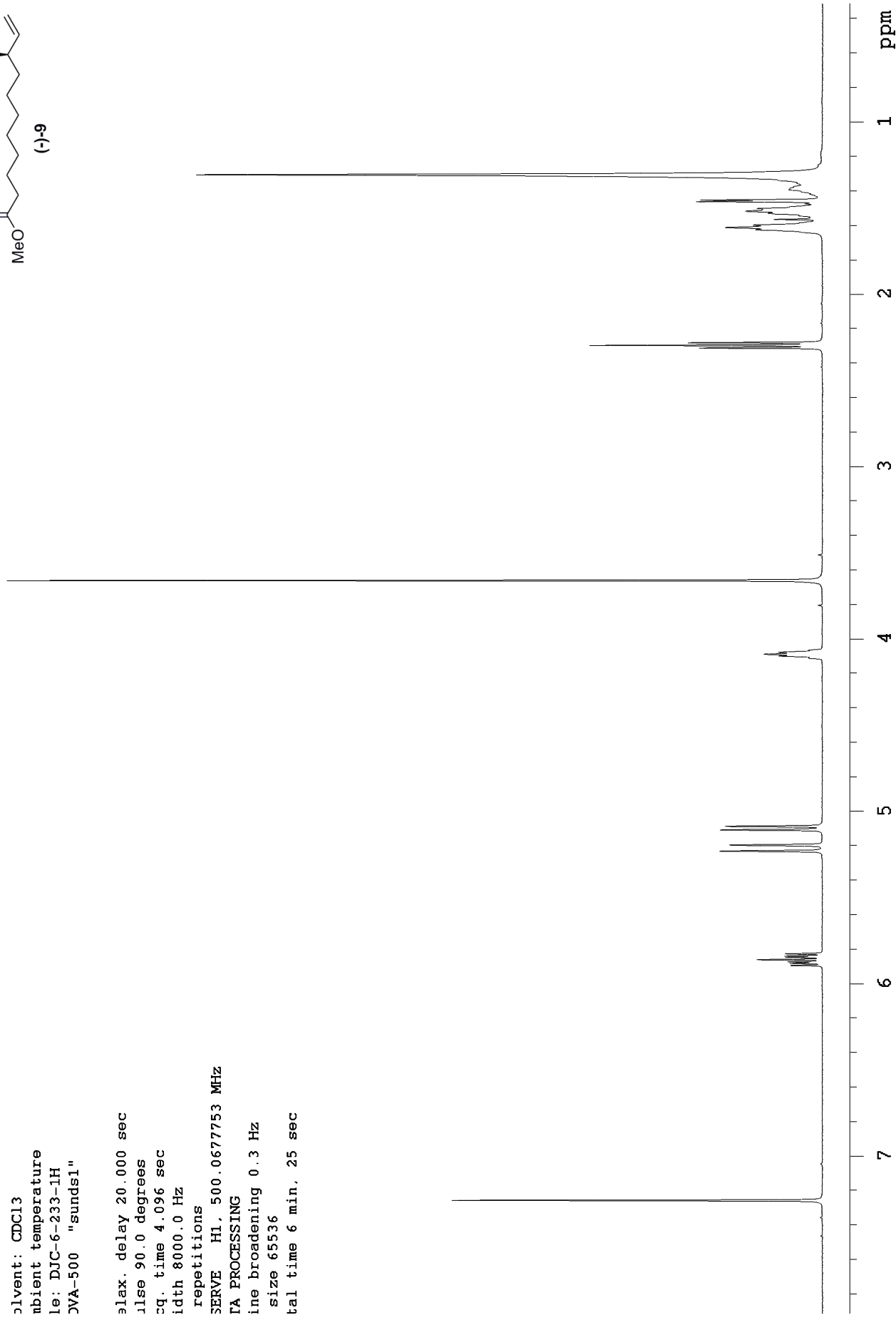
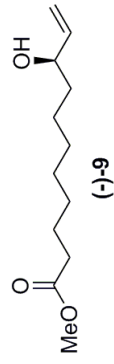
SERVE H1, 500.0677753 MHz

1A PROCESSING

line broadening 0.3 Hz

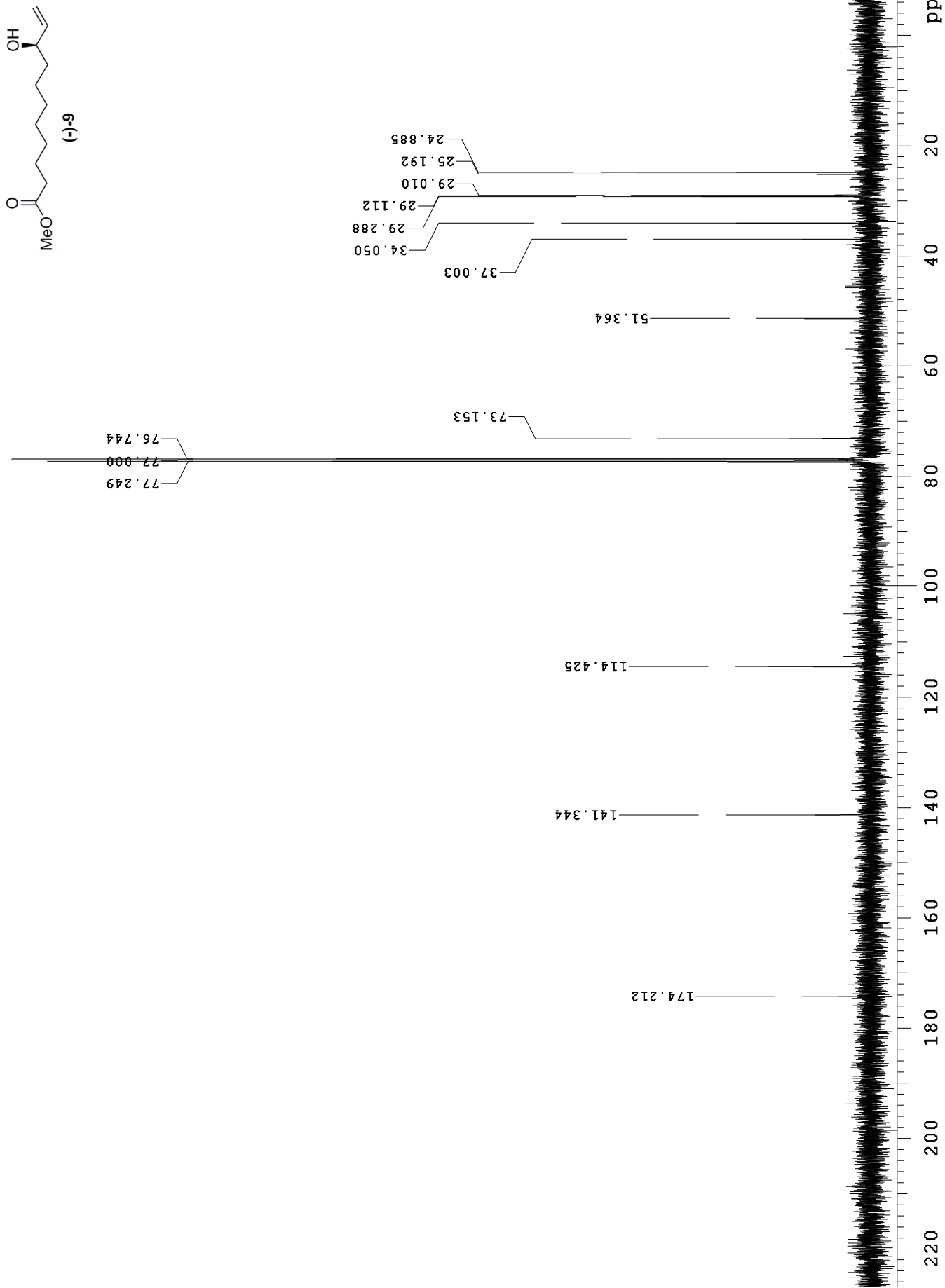
size 65536

total time 6 min., 25 sec

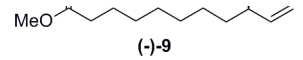


Methyl (9*R*)-9-hydroxyundec-10-enoate ((-)-9)

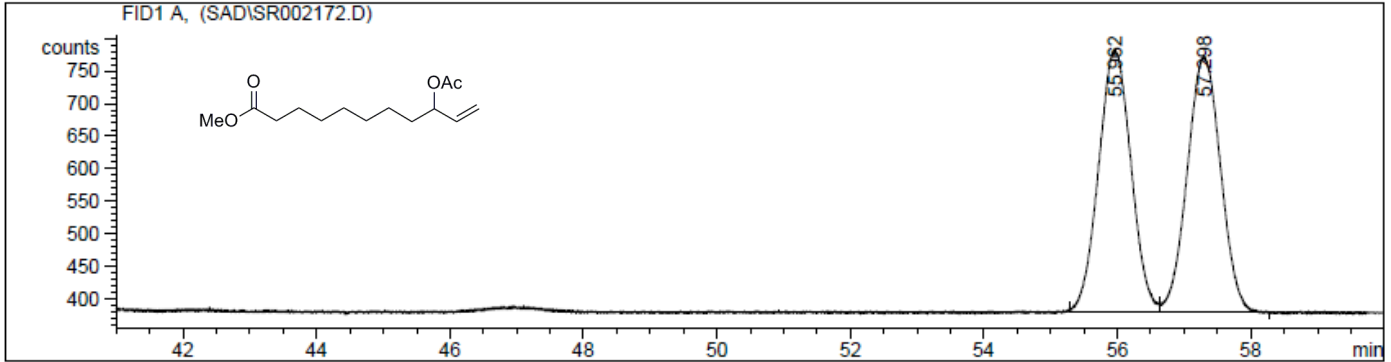
Pulse Sequence: s2pul



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 Acq. Operator : sad
 Acq. Instrument : Chiral GC Inj Volume : Manually
 Acq. Method : C:\HPCHEM\2\METHODS\SUB120DC.M
 Last changed : 3/2/2010 12:47:09 PM by sad
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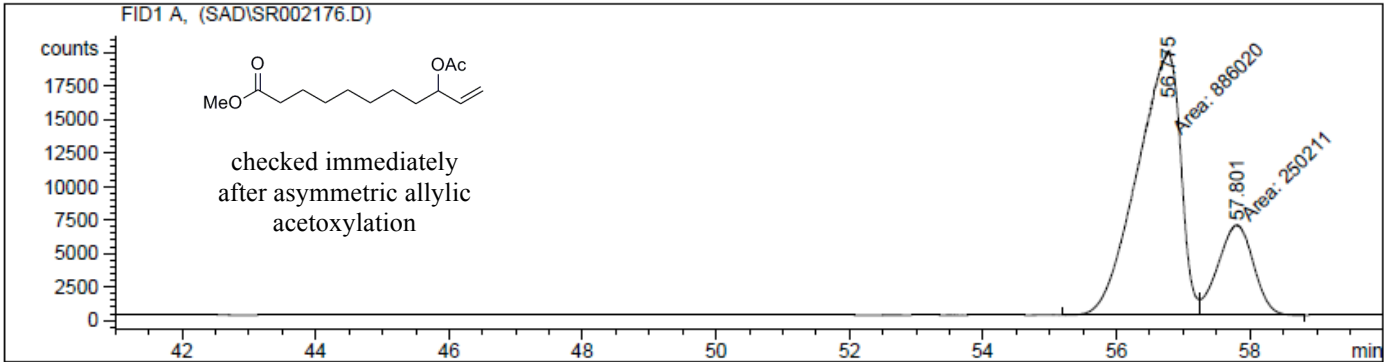
Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	55.962	BV	0.5416	1.36569e4	400.73901	49.89569
2	57.298	VB	0.5529	1.37140e4	391.25912	50.10431

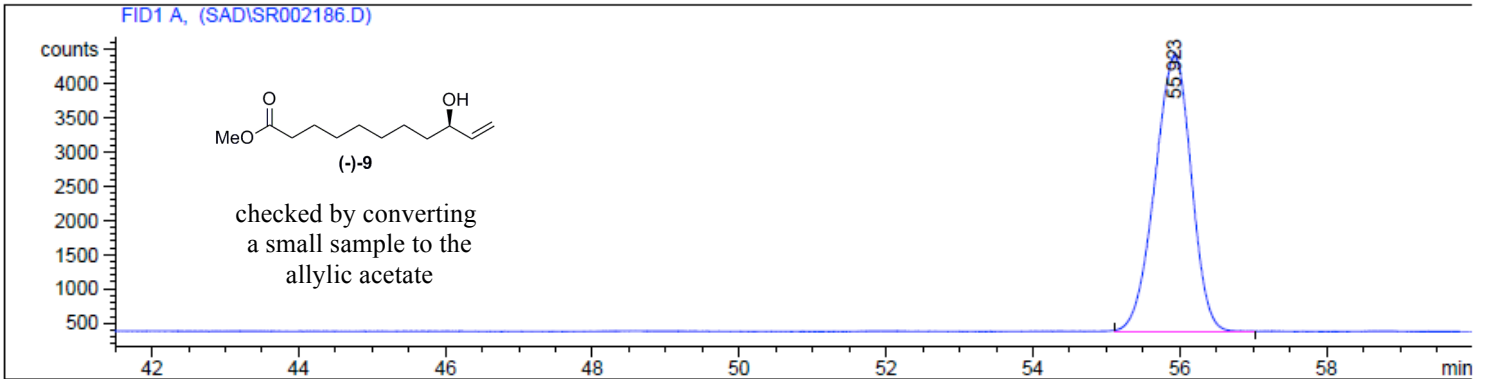
Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	56.775	MF	0.7438	8.86020e5	1.98544e4	77.97884
2	57.801	FM	0.6203	2.50211e5	6723.03027	22.02116

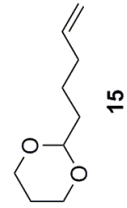
Test



Signal 1: FID1 A,

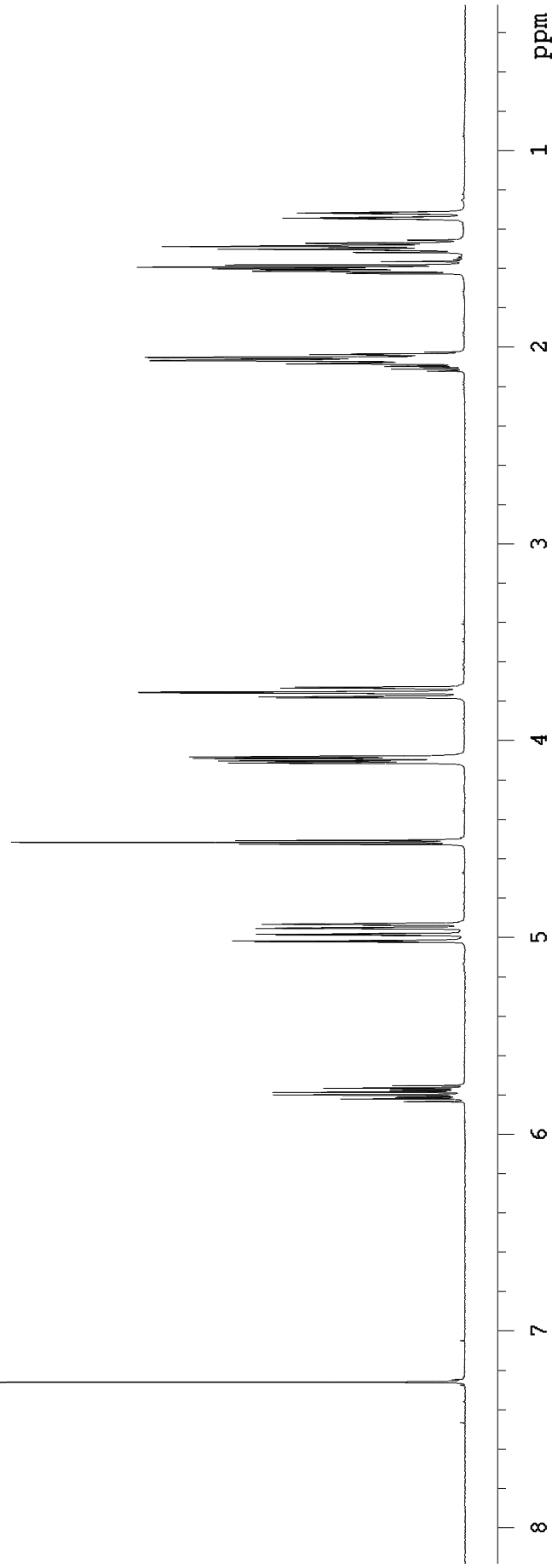
Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	55.923					

Pent-4-en-1-yl)-1,3-dioxane (15)



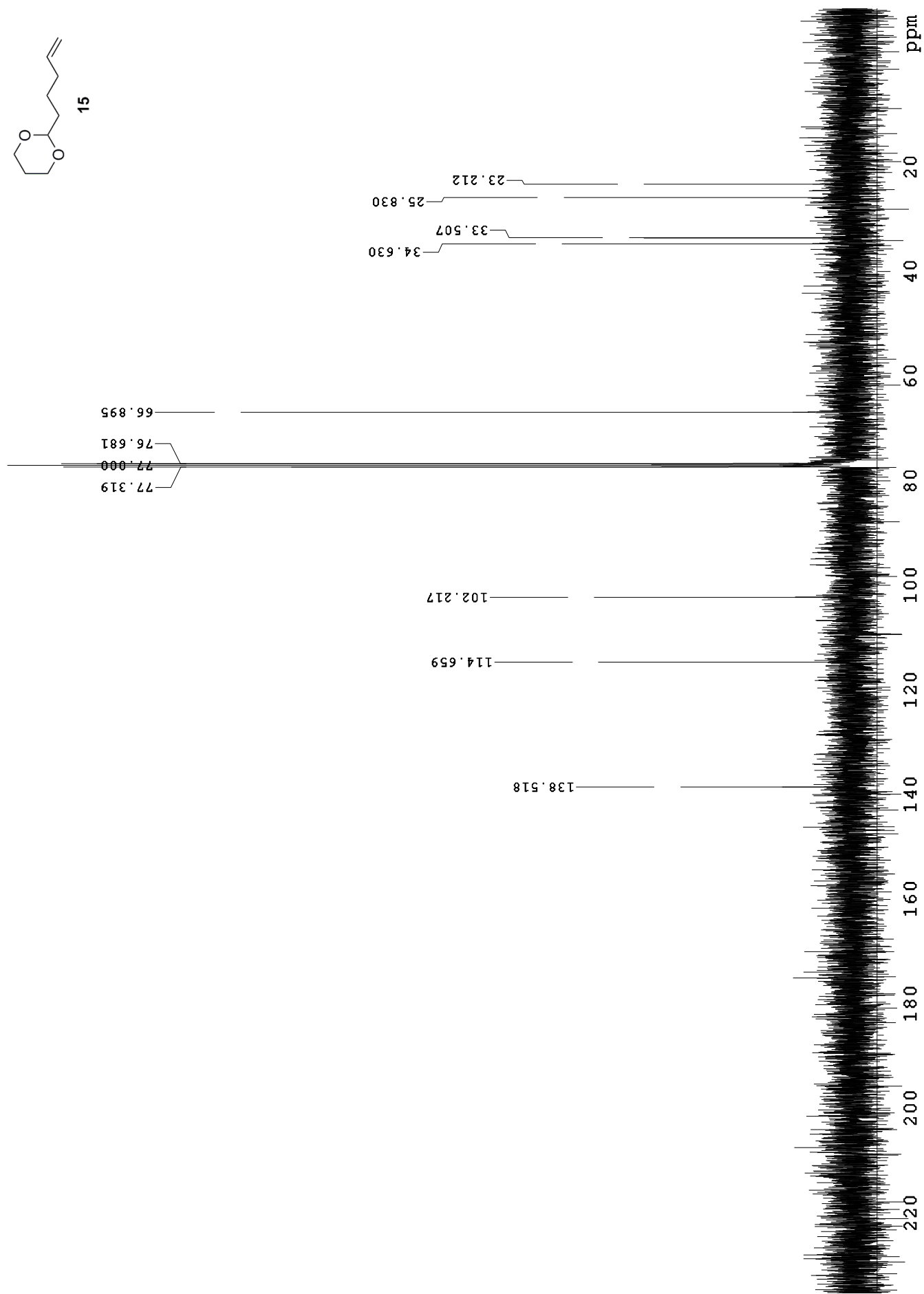
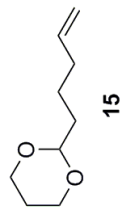
se Sequence: s2pul
 solvent: CDCl3
 ambient temperature
 e: DJC-6-193-IH
 WA-500 "sunds1"

lax. delay 30.000 sec
 lse 90.0 degrees
 q. time 4.096 sec
 dth 8000.0 Hz
 ngle scan
 ERVE H1, 500.0677753 MHz
 'A PROCESSING
 ne broadening 0.3 Hz
 size 65536
 al time 4 min, 32 sec



2-(Pent-4-en-1-yl)-1,3-dioxane (15)

Pulse Sequence: s2pul



2-((3*R*)-Pent-3-acetoxy-4-en-1-yl-3-ol)-1,3-dioxane (13-OAc)

Pulse Sequence: s2pul

Solvent: CDCl₃

Ambient temperature

File: DJC-9-144-1H

INOVA-500 "sunds1"

Relax. delay 20.000 sec

Pulse 61.6 degrees

Acq. time 4.665 sec

Width 7024.9 Hz

2 repetitions

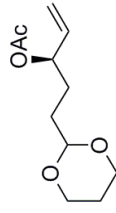
OBSERVE H1, 499.6906570 MHz

DATA PROCESSING

Line broadening 0.3 Hz

FT size 65536

Total time 13 min. 9 sec



B

B/L

L

8

7

6

5

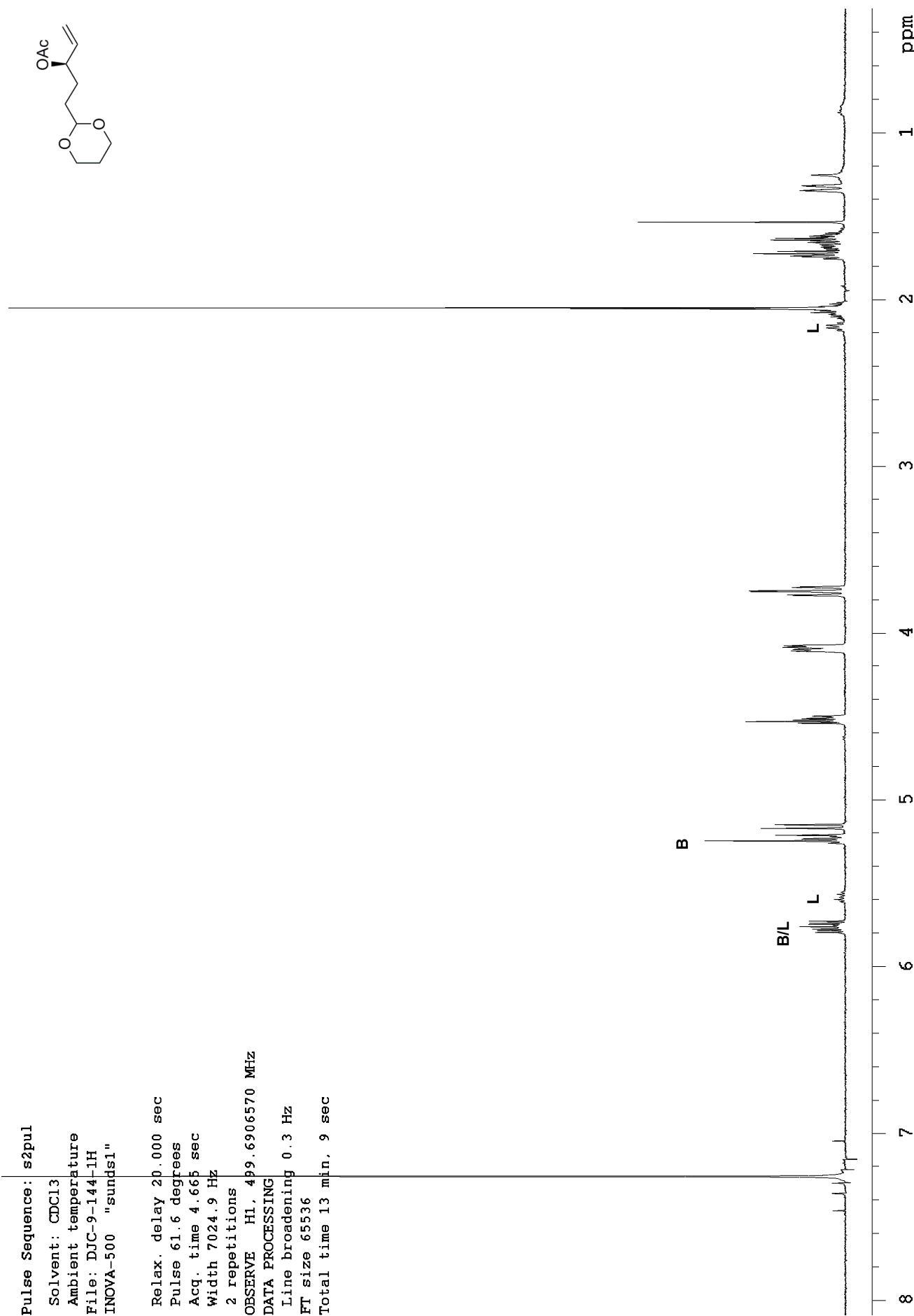
4

3

2

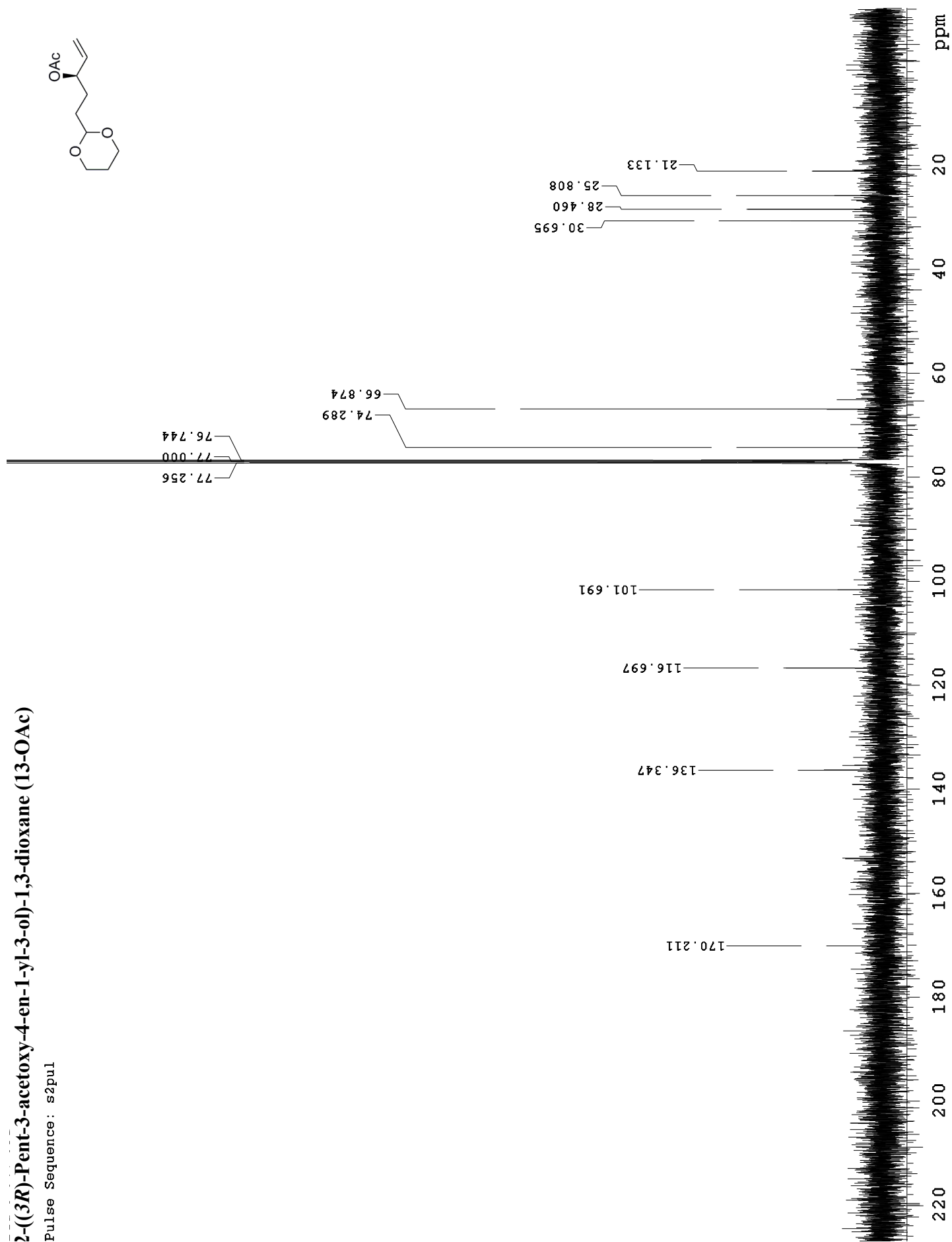
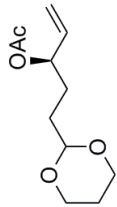
1

ppm



2-((3R)-Pent-3-acetoxy-4-en-1-yl-3-ol)-1,3-dioxane (13-OAc)

Pulse Sequence: s2pul



2-(((3R)-Pent-4-en-1-yl-3-ol)-1,3-dioxane ((-)-13)

Pulse Sequence: s2pul

Solvent: CDCl3

Ambient temperature

File: DJC-6-171

INOVA-500 "sunds1"

Relax. delay 10.000 sec

Pulse 45.0 degrees

Acq. time 4.096 sec

Width 8000.0 Hz

4 repetitions

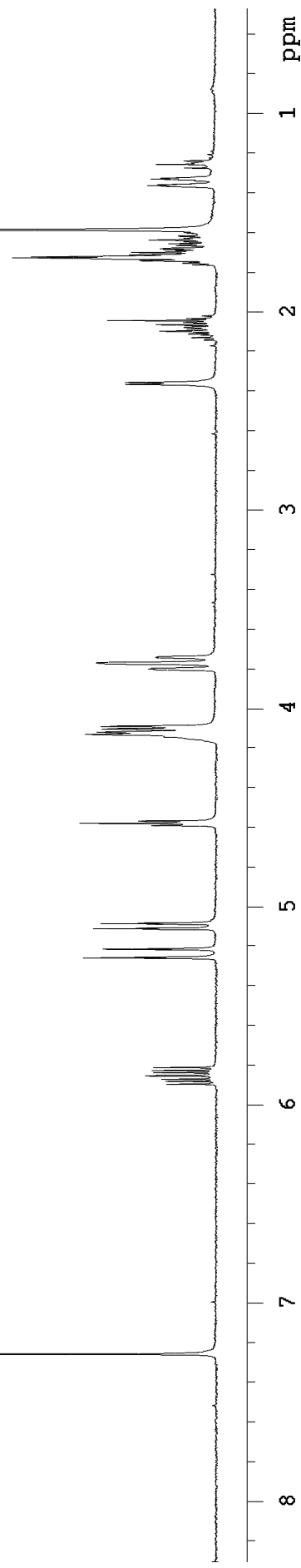
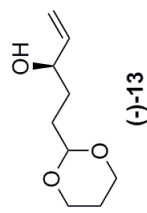
OBSERVE H1, 399.9486778 MHz

DATA PROCESSING

Line broadening 0.3 Hz

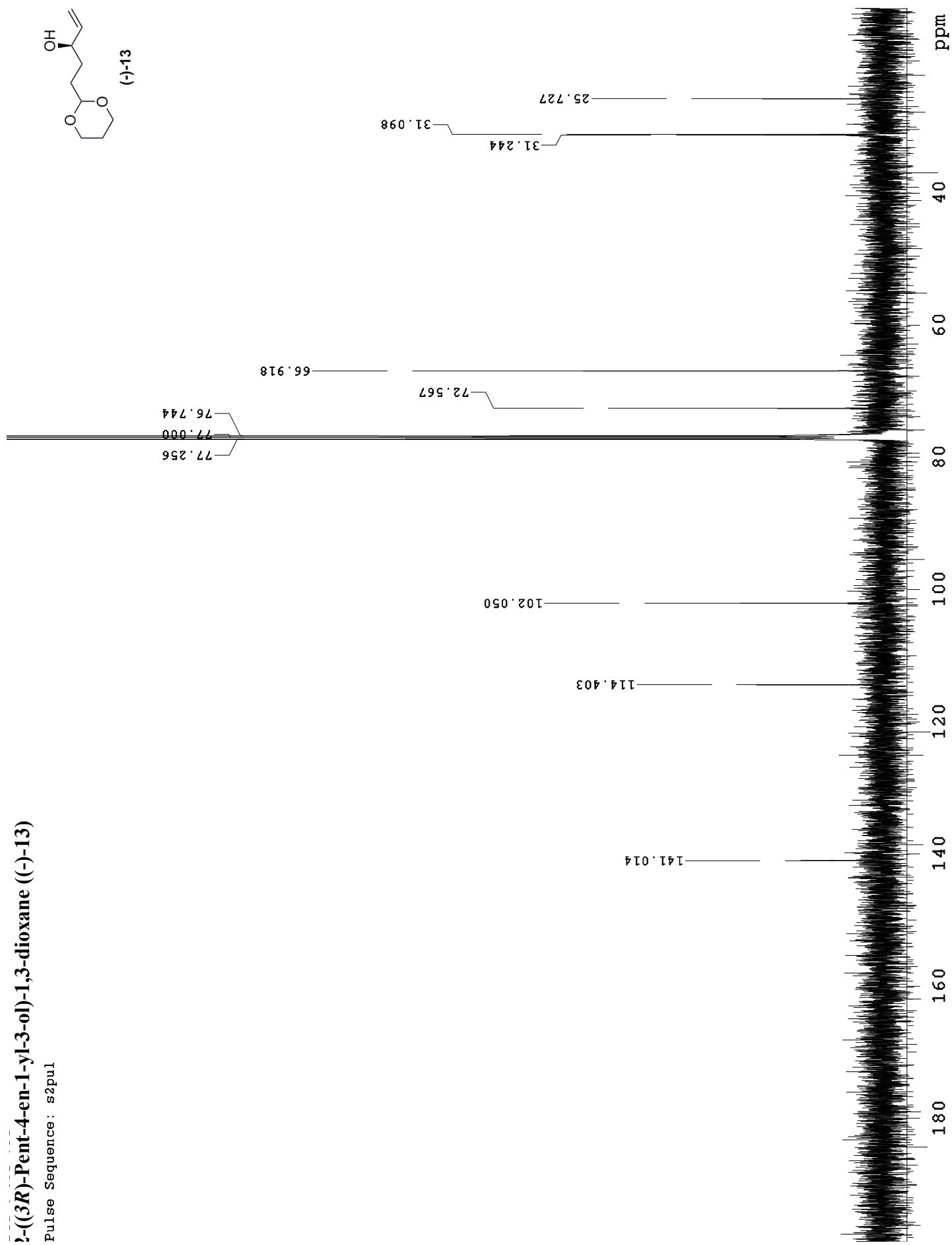
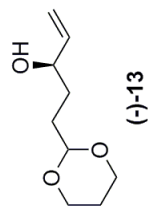
FT size 65536

Total time 3 min. 45 sec

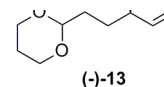


2-(((3R)-Pent-4-en-1-yl-3-ol)-1,3-dioxane ((-)-13)

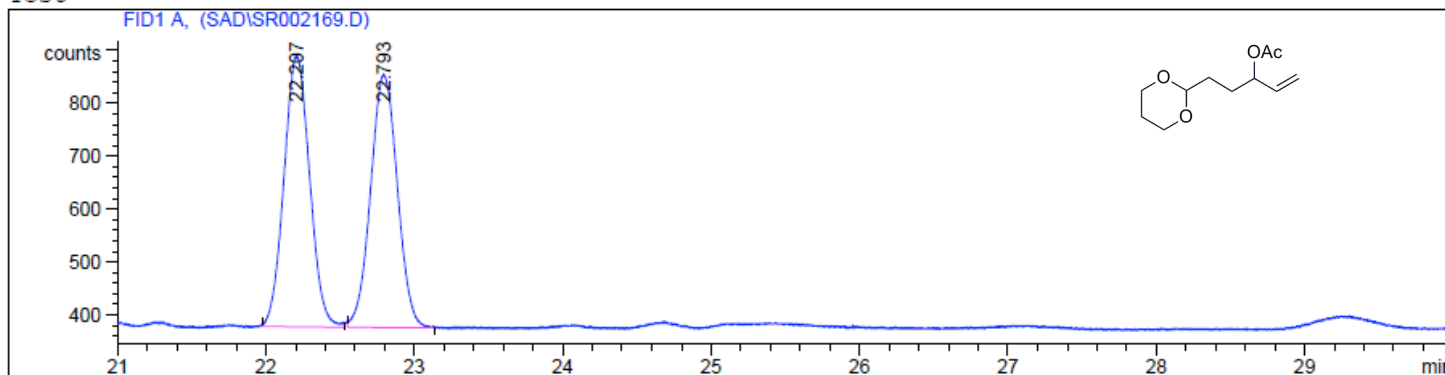
Pulse Sequence: s2pul



Acq. Operator : sad
 Acq. Instrument : Chiral GC Inj Volume : Manually
 Acq. Method : C:\HPCHEM\2\METHODS\SUB110DC.M
 Last changed : 10/14/2010 9:46:00 PM by sad
 Analysis Method : C:\HPCHEM\2\METHODS\SUB110DC.M
 Last changed : 10/15/2010 12:55:17 PM by sad
 (modified after loading)



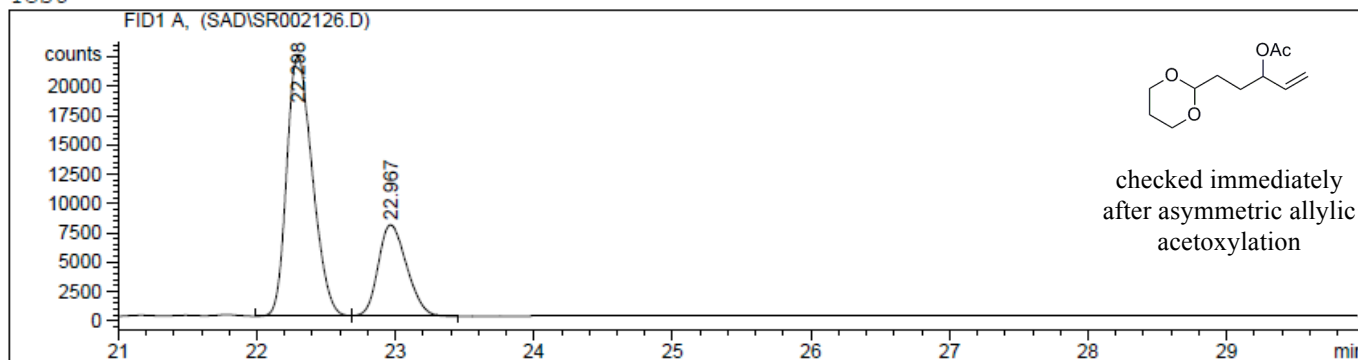
Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	22.207	BB	0.1731	6155.46387	512.69556	50.65797
2	22.793	BP	0.1840	5995.56445	479.18893	49.34203

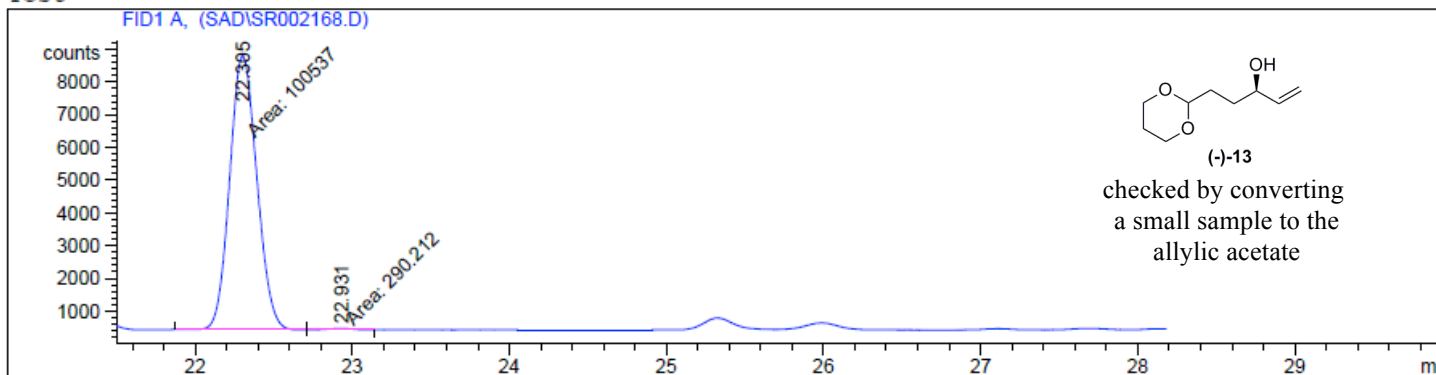
Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	22.298	VV	0.1882	2.85705e5	2.21882e4	71.93395
2	22.967	VB	0.2060	1.11472e5	7787.25928	28.06605

Test



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
--------	---------------	------	-------------	---------------	-----------------	--------

4*R*, 5*E*)-Methyl 4-acetoxy-6-phenylhex-5-enoate (18)

Pulse Sequence: s2pul

Solvent: CDCl₃

Ambient temperature

File: DJC-9-35-1H

INOVA-500 "sunds1"

Relax. delay 30.000 sec

Pulse 90.0 degrees

Acq. time 4.096 sec

Width 8000.0 Hz

2 repetitions

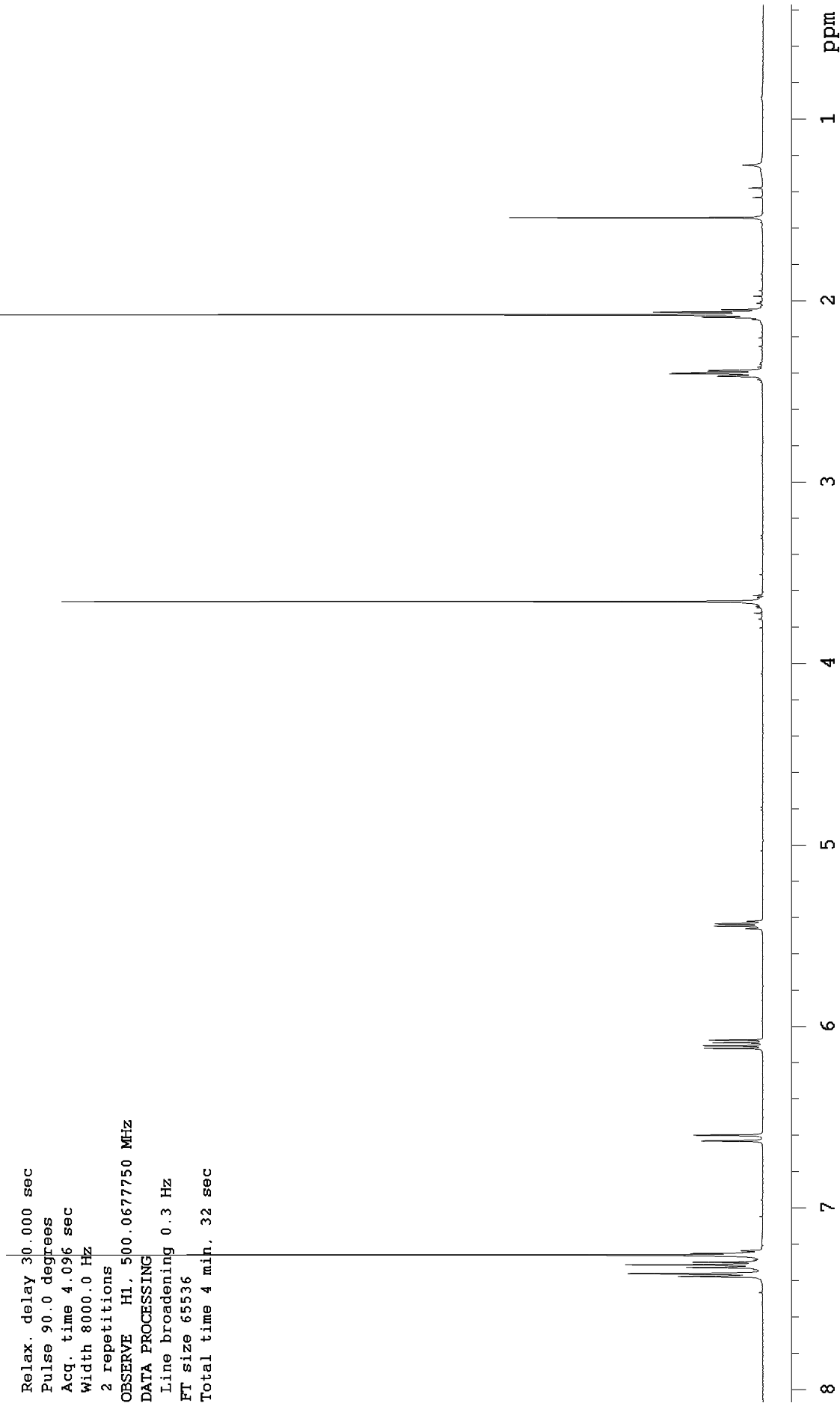
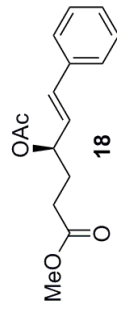
OBSERVE H1, 500.0677750 MHz

DATA PROCESSING

Line broadening 0.3 Hz

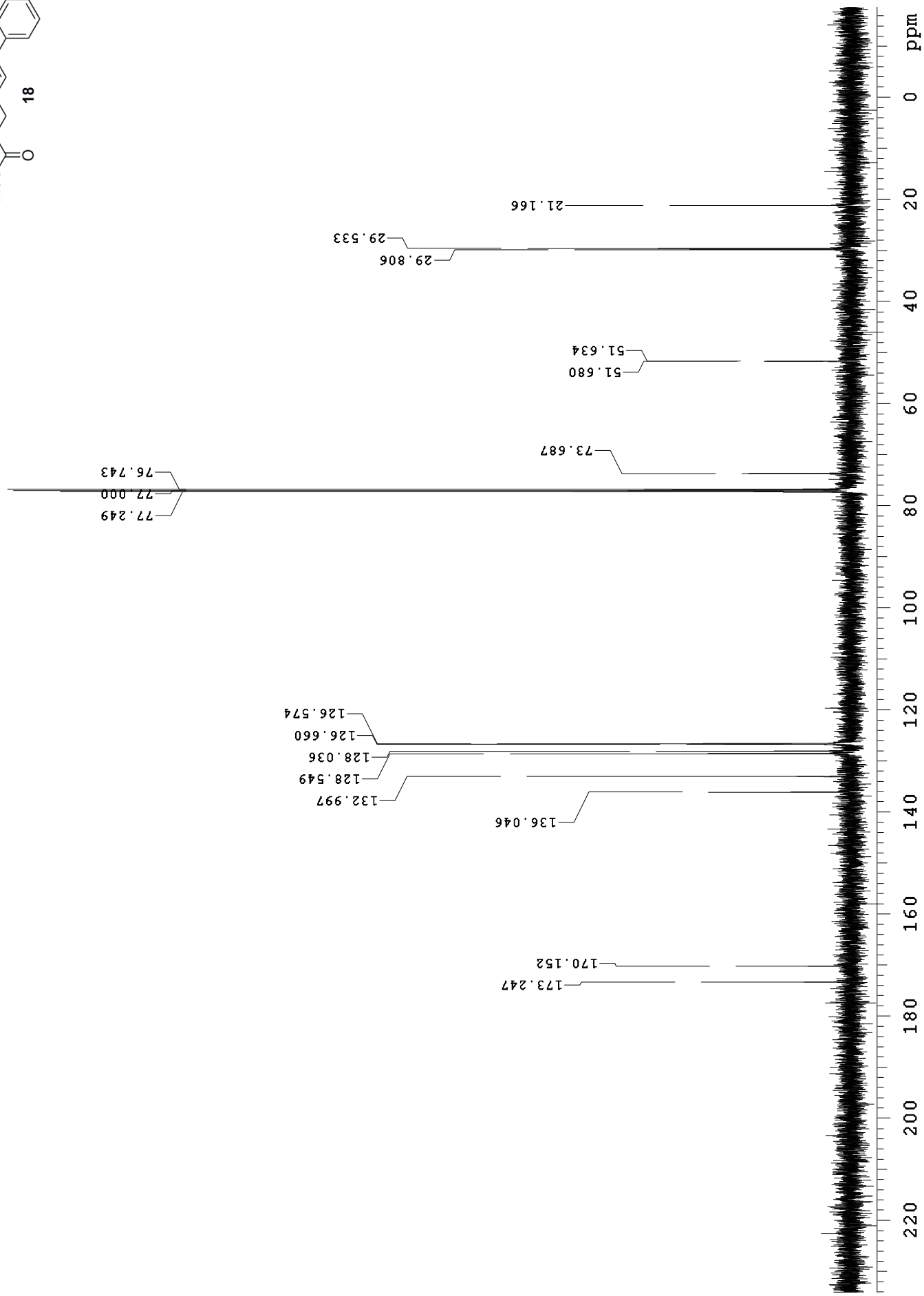
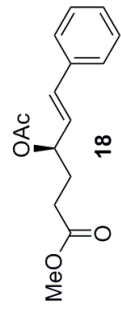
FT size 65536

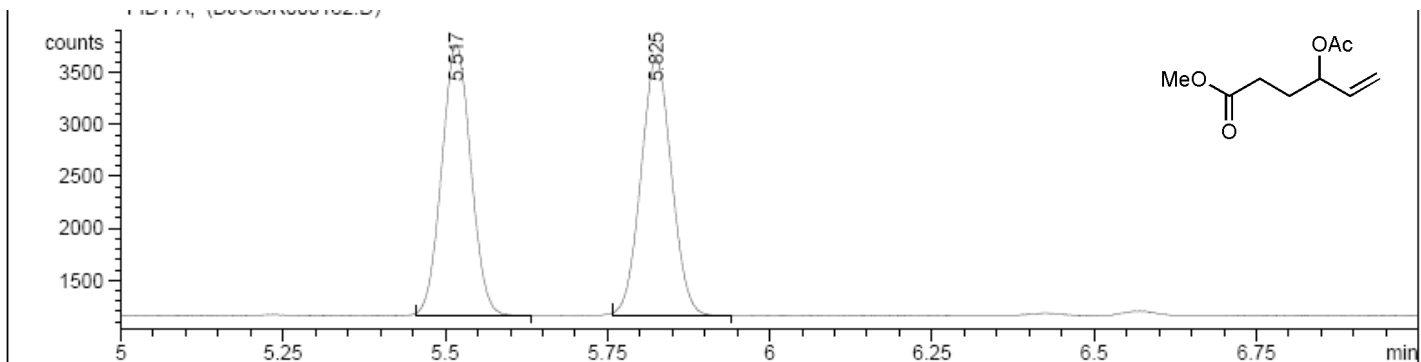
Total time 4 min. 32 sec



4*R*, *E*)-Methyl 4-acetoxy-6-phenylhex-5-enoate (18)

Pulse Sequence: s2pul





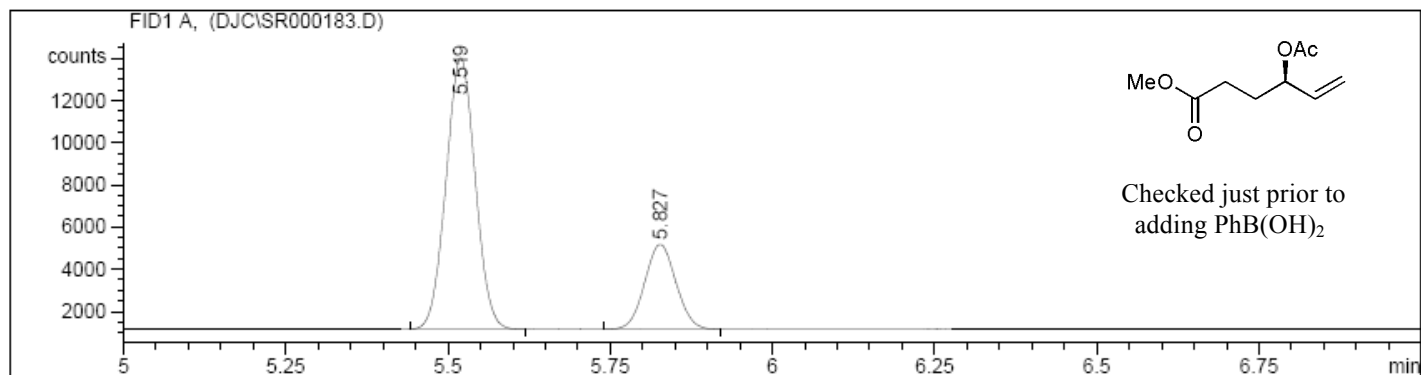
Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	5.517	BB	0.0532	8113.60303	2543.25464	49.97155
2	5.825	BB	0.0554	8122.84131	2403.16333	50.02845

Totals : 1.62364e4 4946.41797

Data File C:\HPCHEM\2\DATA\DJC\SR000183.D

Sample Name: 04-MeEst-R



Signal 1: FID1 A,

Peak #	RetTime [min]	Type	Width [min]	Area counts*s	Height [counts]	Area %
1	5.519	BB	0.0435	4.02791e4	1.28630e4	75.04569
2	5.827	BB	0.0448	1.33937e4	4003.59375	24.95431

Totals : 5.36728e4 1.68666e4

(R, E)-5-styrylidihydrofuran-2(3H)-one (19)

Pulse Sequence: s2p1

Solvent: CDCl₃

Ambient temperature

File: DJC-9-70-1H

INOVA-500 "sunds1"

Relax. delay 30.000 sec

Pulse 90.0 degrees

Acq. time 4.096 sec

Width 8000.0 Hz

2 repetitions

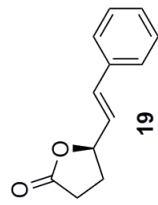
OBSERVE H1, 500.0677750 MHz

DATA PROCESSING

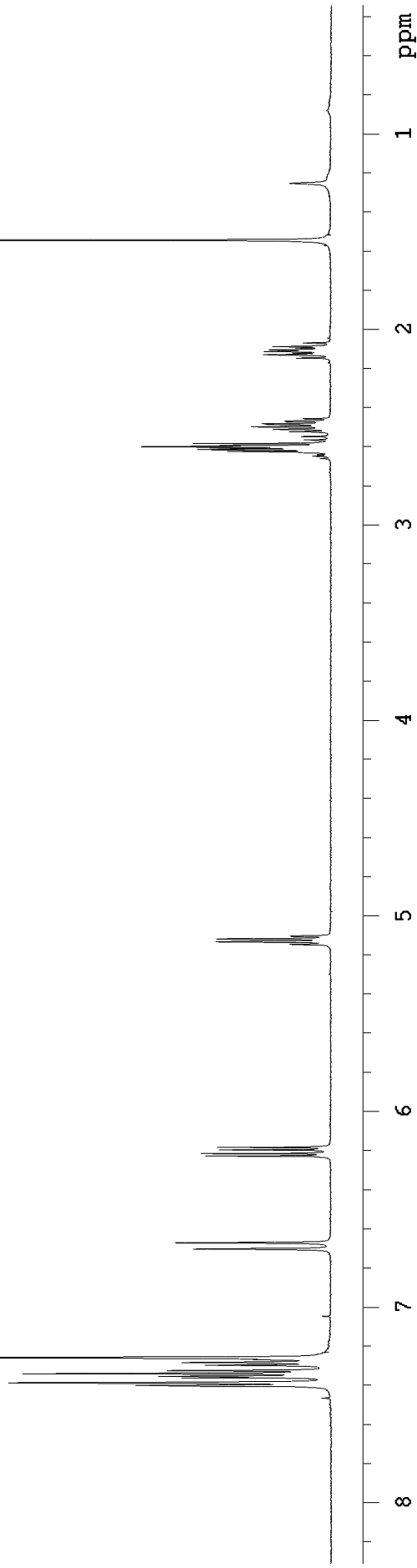
Line broadening 0.3 Hz

FT size 65536

Total time 4 min. 32 sec

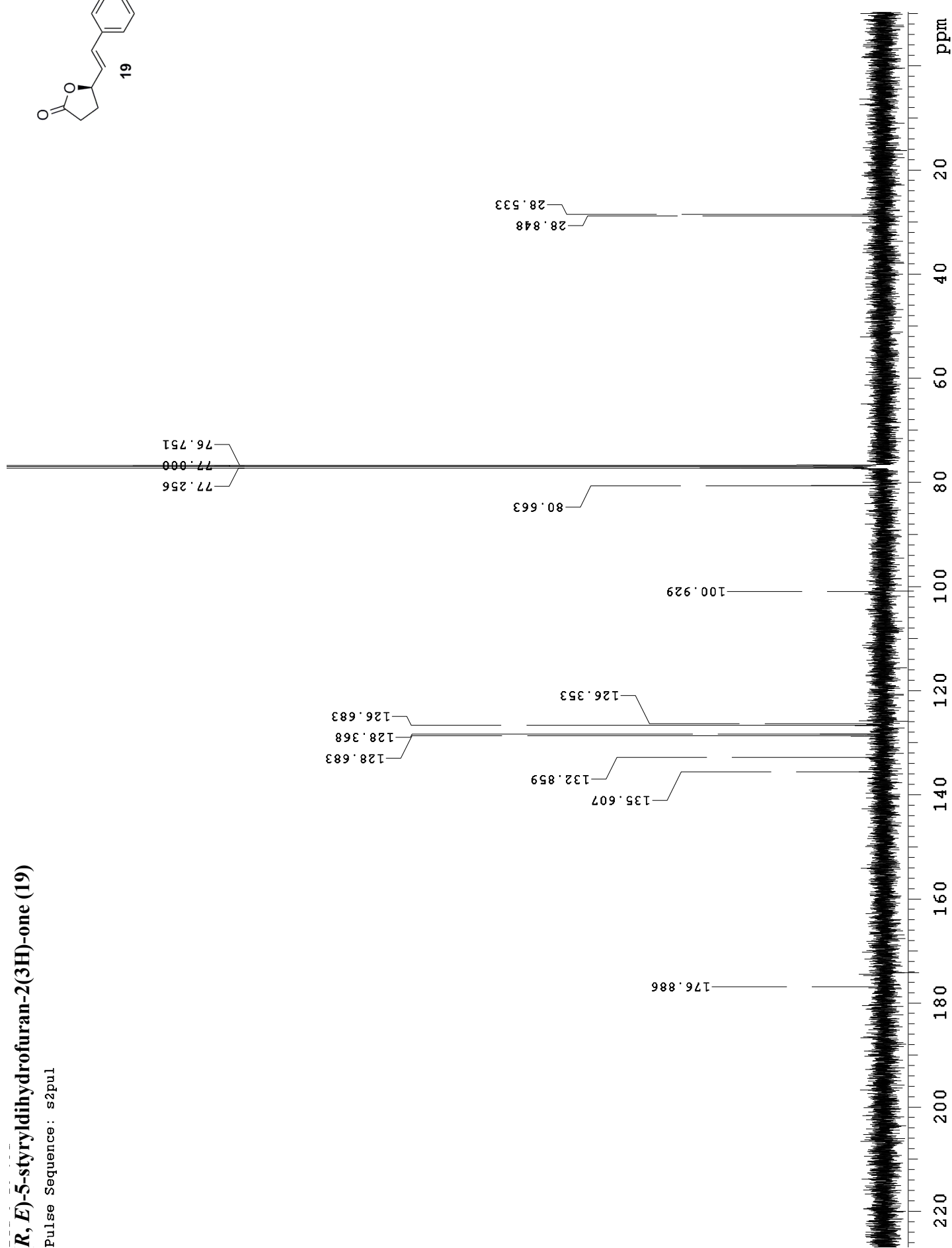
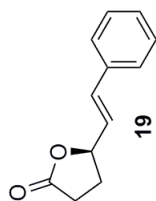


19



***R, E*-5-styrylidihydrofuran-2(3H)-one (19)**

Pulse Sequence: s2pul



(R)-5-((4R,5R)-2,2-dimethyl-5-phenyl-1,3-dioxolan-4-yl)dihydrofuran-2(3H)-one ((-)-20)

Pulse Sequence: s2pul

Solvent: CDCl3

Ambient temperature

File: DJC-9-125

INOVA-500 "sunds1"

Relax. delay 15.000 sec

Pulse 90.0 degrees

Acq. time 4.096 sec

Width 8000.0 Hz

4 repetitions

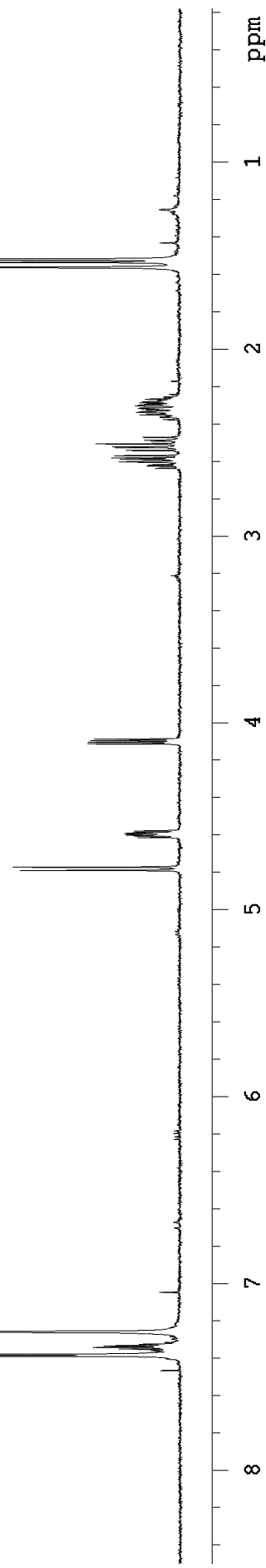
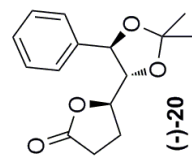
OBSERVE H1, 500.0677753 MHz

DATA PROCESSING

Line broadening 0.3 Hz

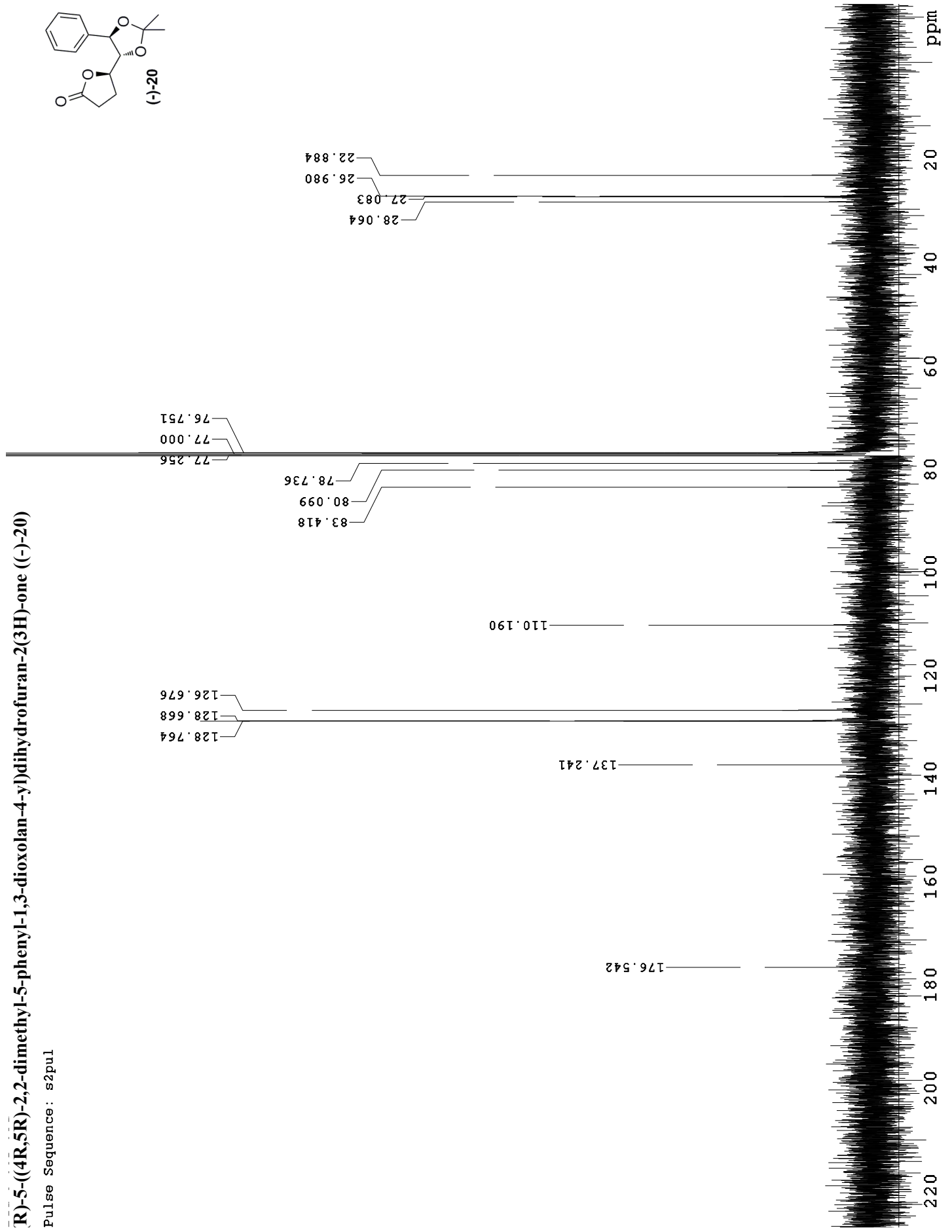
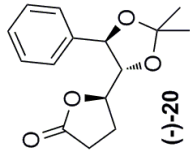
FT size 65536

Total time 2 min, 32 sec



(R)-5-((4R,5R)-2,2-dimethyl-5-phenyl-1,3-dioxolan-4-yl)dihydrofuran-2(3H)-one ((-)-20)

Pulse Sequence: s2pul



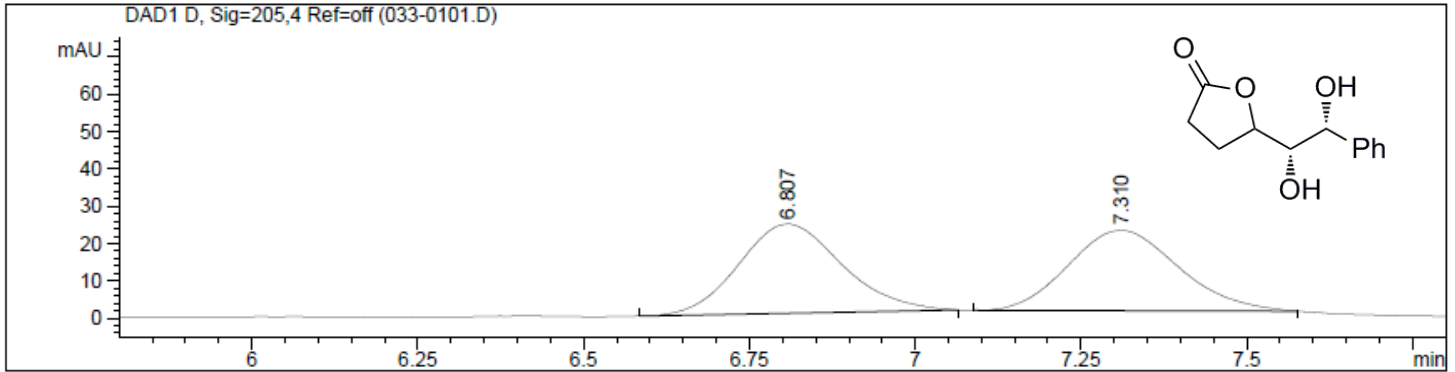
```

=====
Injection Date : 11/23/2010 8:21:02 PM      Seq. Line : 1
Sample Name    : DC-diol-rac                 Location  : Vial 33
Acq. Operator  : HPLC Host                   Inj       : 1
Acq. Instrument: Instrument 1                 Inj Volume: 0.5 µl
Sequence File  : C:\HPCHEM\1\SEQUENCE\STANDARD.S
Acq. Method    : C:\HPCHEM\1\METHODS\COLUMNS\REVERSE\CHIRAL\AD-RH\35ACCN.M
Last changed   : 11/23/2010 8:20:08 PM by HPLC Host
Analysis Method: C:\HPCHEM\1\METHODS\COLUMNS\REVERSE\CHIRAL\AD-RH\35ACCN.M
Last changed   : 11/23/2010 8:33:48 PM by HPLC Host
                (modified after loading)

```

AD-RH
Reverse Phase Chiral
ACCN35%

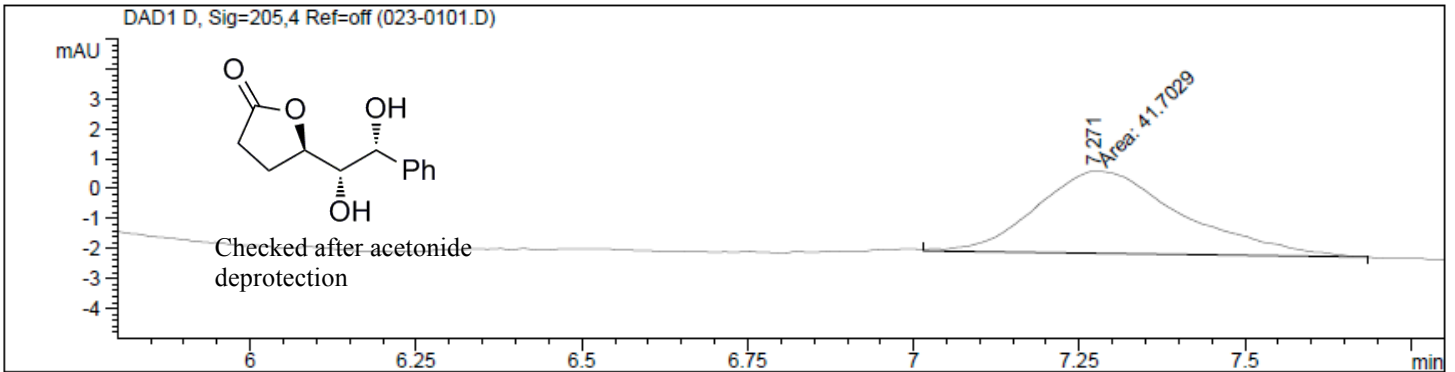
Standard Run



Signal 1: DAD1 D, Sig=205,4 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	6.807	BB	0.1665	258.61758	23.89842	50.9221
2	7.310	BB	0.1793	249.25174	21.55332	49.0779

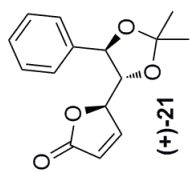
Standard Run



Signal 1: DAD1 D, Sig=205,4 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	7.271	MM	0.2538	41.70292	2.73845	100.0000

R)-5-((4R,5R)-2,2-dimethyl-5-phenyl-1,3-dioxolan-4-yl)furan-2(5H)-one ((+)-21)



Pulse Sequence: s2pul

Solvent: CDCl3

Ambient temperature

File: DJC-9-122-1H

INOVA-500 "sundsl"

Relax. delay 15.000 sec

Pulse 90.0 degrees

Acq. time 4.096 sec

Width 8000.0 Hz

3 repetitions

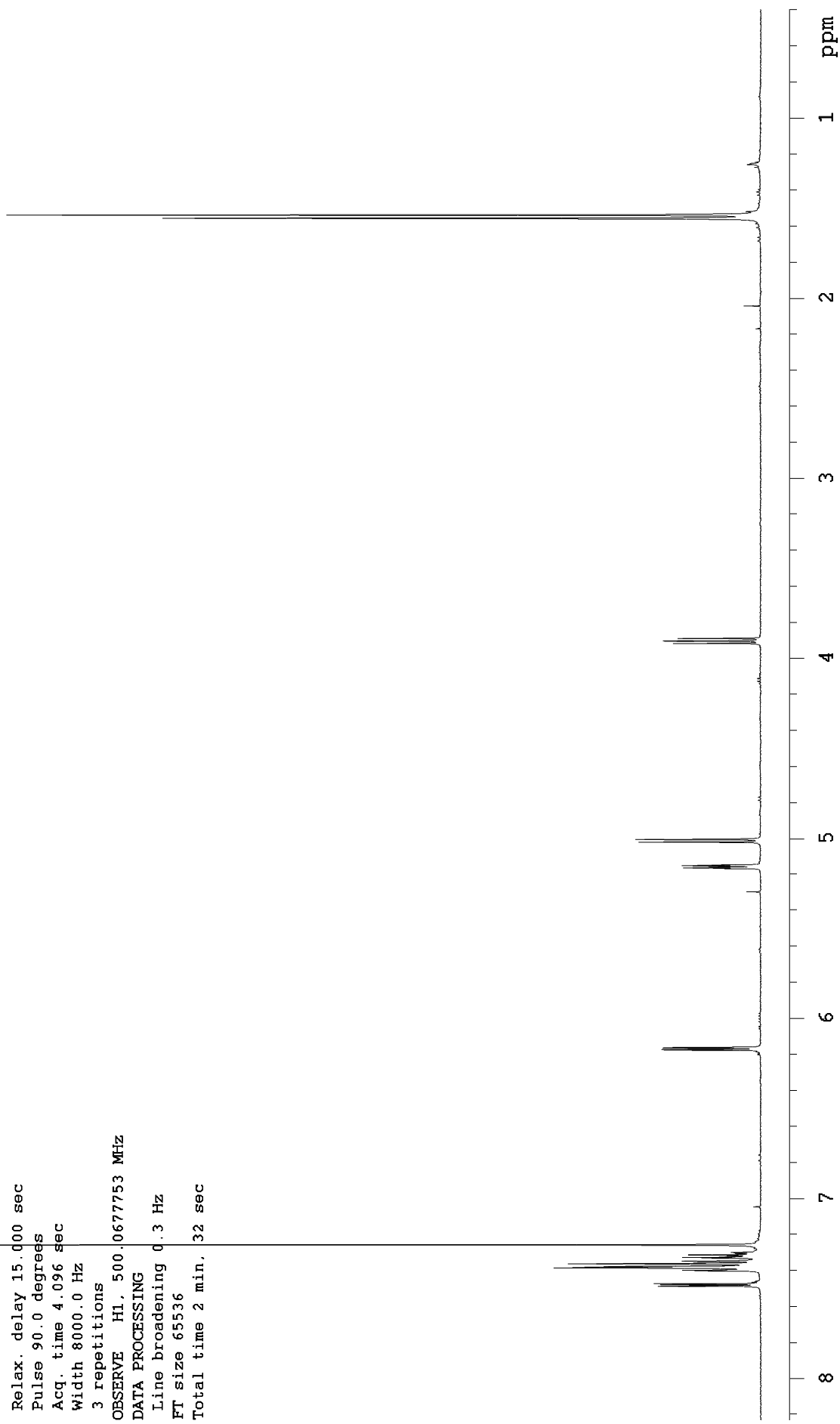
OBSERVE H1, 500.0677753 MHz

DATA PROCESSING

Line broadening 0.3 Hz

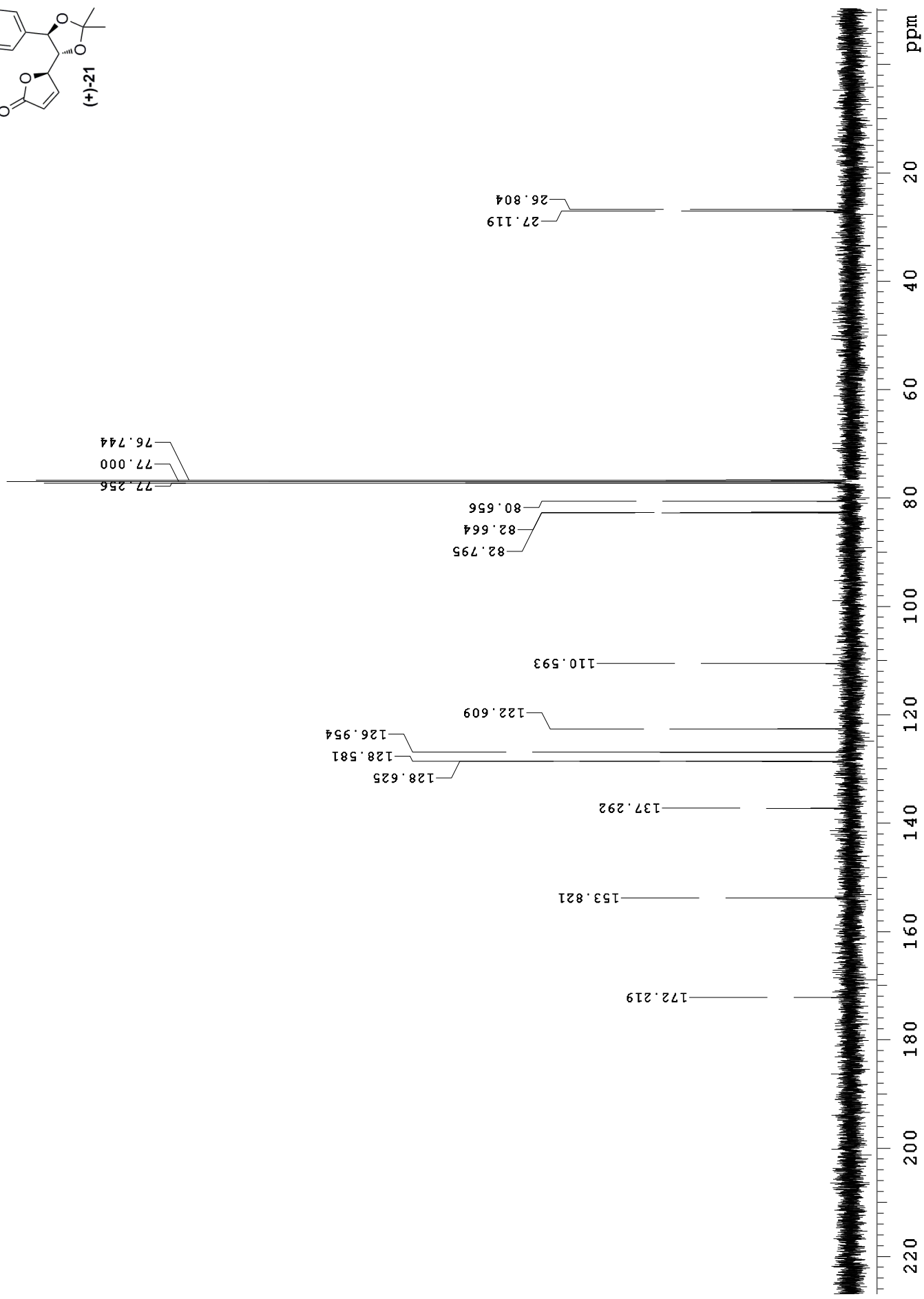
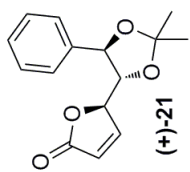
FT size 65536

Total time 2 min. 32 sec

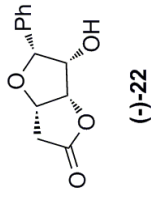


R)-5-((4R,5R)-2,2-dimethyl-1,3-dioxolan-4-yl)furan-2(5H)-one ((+)-21)

Pulse Sequence: s2pul



3aS,5R,6S,6aS)-6-hydroxy-5-phenyltetrahydrofuro[3,2-b]furan-2(5H)-one((-)-22)



Pulse Sequence: s2pul

Solvent: CDCl3

Ambient temperature

File: DJC-9-78-2

INOVA-500 "sunds1"

Relax. delay 15.000 sec

Pulse 90.0 degrees

Acq. time 4.096 sec

Width 8000.0 Hz

7 repetitions

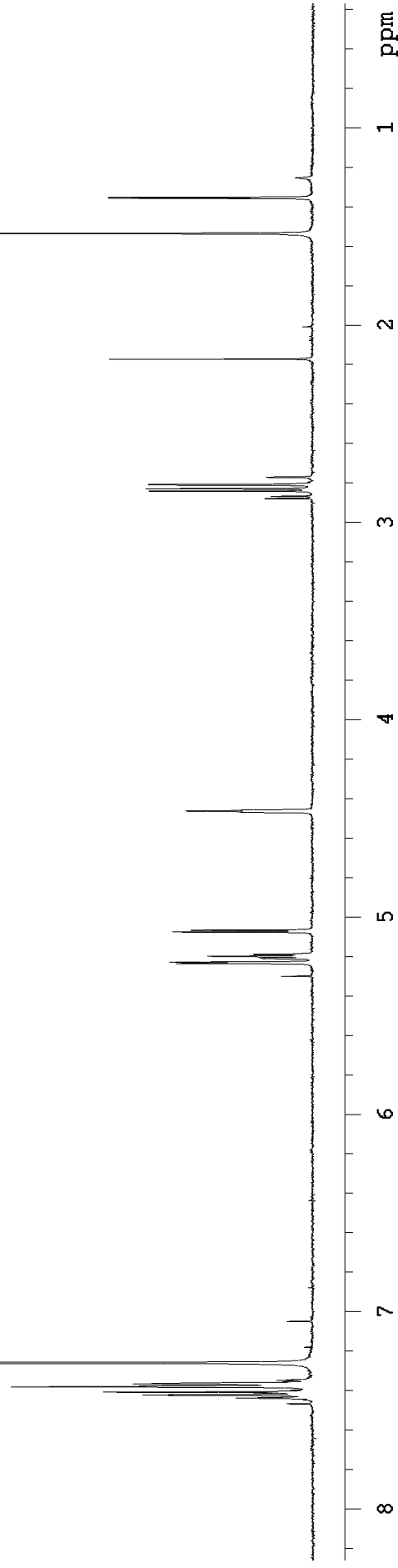
OBSERVE H1, 500.0677748 MHz

DATA PROCESSING

Line broadening 0.3 Hz

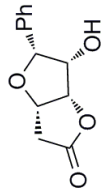
FT size 65536

Total time 2 min, 32 sec

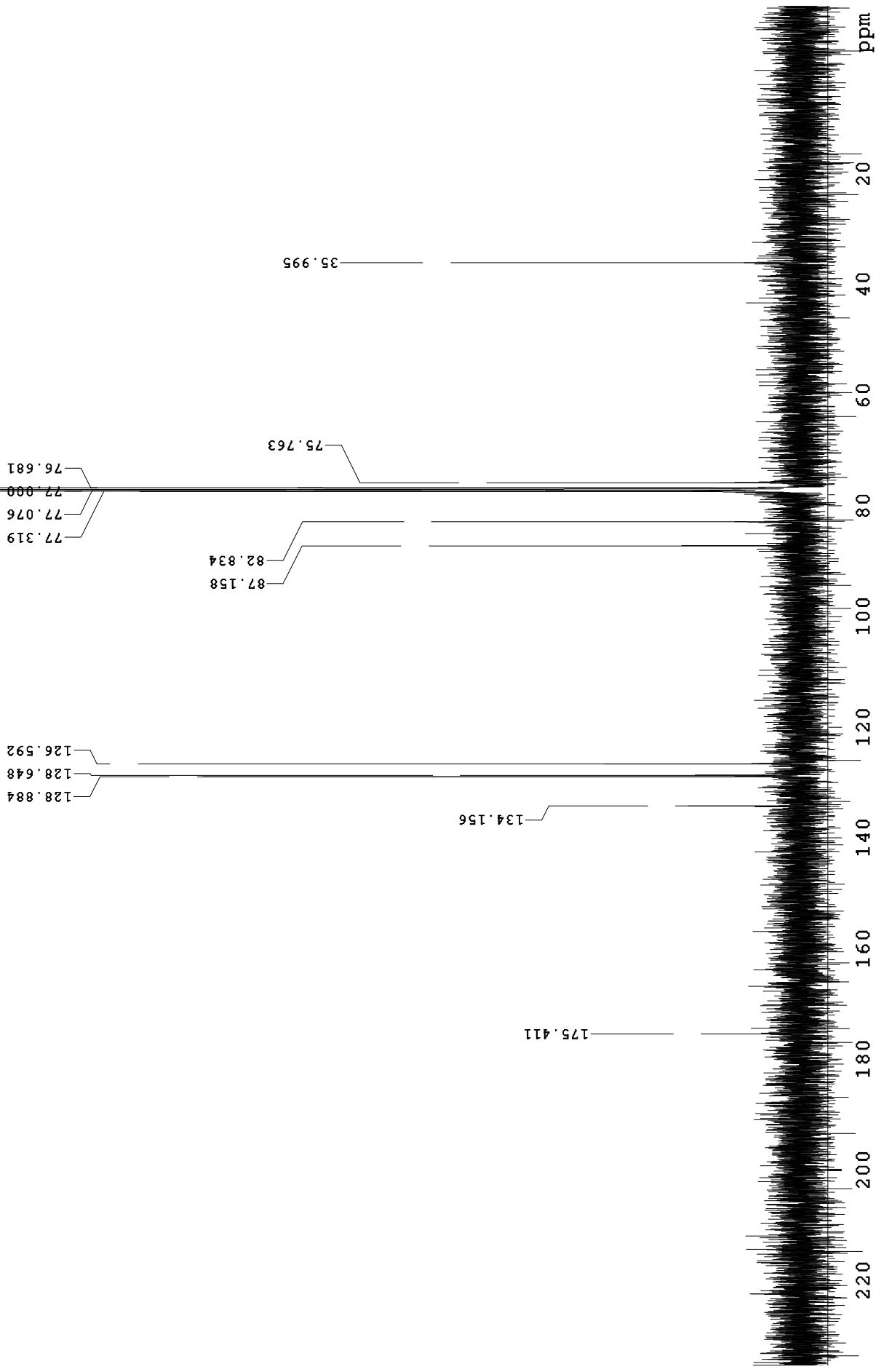


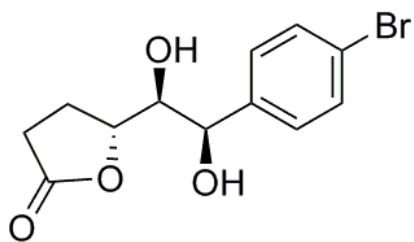
3aS,5R,6S,6aS)-6-hydroxy-5-phenyltetrahydrofuro[3,2-b]furan-2(5H)-one((-)-22)

Pulse Sequence: s2pul

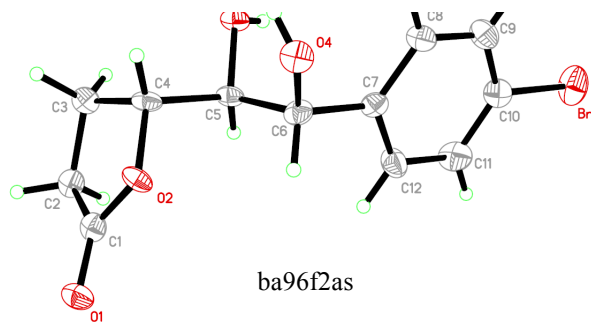


(-)-22





p-Bromophenyl- **20**-diol



ba96f2as

The Model has Chirality at C4	(Verify)	R
The Model has Chirality at C5	(Verify)	S
The Model has Chirality at C6	(Verify)	R

Table 1. Crystal data and structure refinement for ba96f2as.

Identification code	ba96f2as	
Empirical formula	C ₁₂ H ₁₃ Br O ₄	
Molecular weight	301.13	
Temperature	193(2) K	
Wavelength	0.71073 Å	
Crystal system	Monoclinic	
Space group	P 21	
Unit cell dimensions	a = 10.791(2) Å	α = 90°.
	b = 4.9772(8) Å	β = 101.532(12)°.
	c = 11.585(2) Å	γ = 90°.
Volume	609.63(19) Å ³	
Z	2	
Density (calculated)	1.640 Mg/m ³	
Absorption coefficient (000)	3.371 mm ⁻¹	
Crystal size	0.862 x 0.396 x 0.05 mm ³	
Theta range for data collection	1.79 to 25.31°.	
Index ranges	-12 ≤ h ≤ 12, -5 ≤ k ≤ 5, -13 ≤ l ≤ 13	
Reflections collected	8519	
Independent reflections	2168 [R(int) = 0.0829]	
Completeness to theta = 25.31°	99.6 %	
Absorption correction	Integration	
Max. and min. transmission	0.8408 and 0.2253	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	2168 / 1 / 156	
Goodness-of-fit on F ²	0.970	
Final R indices [I > 2σ(I)]	R1 = 0.0476, wR2 = 0.1055	
R indices (all data)	R1 = 0.0767, wR2 = 0.1196	
Absolute structure parameter	0.054(19)	
Largest diff. peak and hole	0.614 and -0.592 e.Å ⁻³	

$U(\text{eq})$ is defined as one third of the trace of the orthogonalized U -tensor.

	x	y	z	U(eq)
r(1)	2483(1)	5824(2)	-957(1)	58(1)
l(1)	110(4)	-3643(8)	6064(3)	36(1)
l(2)	1949(3)	-3189(7)	5495(3)	28(1)
l(3)	4040(3)	1842(7)	4654(3)	27(1)
l(4)	3861(4)	-3368(6)	3542(3)	32(1)
l'(1)	1056(6)	-2341(11)	6078(5)	30(1)
l'(2)	1441(5)	291(12)	6693(5)	33(2)
l'(3)	2834(5)	462(14)	6671(4)	31(1)
l'(4)	2986(5)	-1274(11)	5641(5)	28(1)
l'(5)	2902(5)	281(10)	4484(5)	26(1)
l'(6)	2812(6)	-1543(12)	3396(5)	30(1)
l'(7)	2749(5)	139(10)	2300(5)	28(1)
l'(8)	3799(5)	570(18)	1793(5)	36(1)
l'(9)	3712(6)	2244(13)	815(5)	38(2)
l'(10)	2587(6)	3494(12)	353(5)	37(2)
l'(11)	1521(6)	3065(14)	829(6)	44(2)
l'(12)	1621(6)	1401(12)	1809(5)	39(2)

Table 3. Bond lengths [\AA] and angles [$^\circ$] for ba96f2as.

r(1)-C(10)	1.896(6)
l(1)-C(1)	1.206(7)
l(2)-C(1)	1.350(7)
l(2)-C(4)	1.454(7)
l(3)-C(5)	1.434(6)
l(3)-H(3C)	0.8400
l(4)-C(6)	1.434(7)
l(4)-H(4B)	0.8400
l'(1)-C(2)	1.510(8)
l'(2)-C(3)	1.511(8)
l'(2)-H(2A)	0.9900
l'(2)-H(2B)	0.9900
l'(3)-C(4)	1.508(8)
l'(3)-H(3A)	0.9900
l'(3)-H(3B)	0.9900
l'(4)-C(5)	1.535(8)
l'(4)-H(4A)	1.0000
l'(5)-C(6)	1.541(8)
l'(5)-H(5A)	1.0000
l'(6)-C(7)	1.511(7)
l'(6)-H(6A)	1.0000
l'(7)-C(12)	1.386(8)
l'(7)-C(8)	1.393(7)
l'(8)-C(9)	1.394(9)
l'(8)-H(8A)	0.9500
l'(9)-C(10)	1.374(9)
l'(9)-H(9A)	0.9500
l'(10)-C(11)	1.387(9)
l'(11)-C(12)	1.392(8)
l'(11)-H(11A)	0.9500
l'(12)-H(12A)	0.9500
l(1)-O(2)-C(4)	110.3(4)
l(5)-O(3)-H(3C)	109.5
l(6)-O(4)-H(4B)	109.5
l(1)-C(1)-O(2)	120.8(5)
l(1)-C(1)-C(2)	128.7(5)
l(2)-C(1)-C(2)	110.6(5)
l'(1)-C(2)-C(3)	102.8(5)
l'(1)-C(2)-H(2A)	111.2
l'(3)-C(2)-H(2A)	111.2
l'(1)-C(2)-H(2B)	111.2
l'(3)-C(2)-H(2B)	111.2
l(2A)-C(2)-H(2B)	109.1

'(4)-C(3)-H(3B)	110.9
(3A)-C(3)-H(3B)	108.9
(2)-C(4)-C(3)	105.3(4)
(2)-C(4)-C(5)	108.6(4)
'(3)-C(4)-C(5)	113.9(4)
(2)-C(4)-H(4A)	109.6
'(3)-C(4)-H(4A)	109.6
'(5)-C(4)-H(4A)	109.6
(3)-C(5)-C(4)	104.7(4)
(3)-C(5)-C(6)	110.1(4)
'(4)-C(5)-C(6)	113.6(4)
(3)-C(5)-H(5A)	109.4
'(4)-C(5)-H(5A)	109.4
'(6)-C(5)-H(5A)	109.4
(4)-C(6)-C(7)	110.7(5)
(4)-C(6)-C(5)	110.9(4)
'(7)-C(6)-C(5)	110.2(4)
(4)-C(6)-H(6A)	108.3
'(7)-C(6)-H(6A)	108.3
'(5)-C(6)-H(6A)	108.3
'(12)-C(7)-C(8)	118.6(5)
'(12)-C(7)-C(6)	119.0(5)
'(8)-C(7)-C(6)	122.3(5)
'(7)-C(8)-C(9)	120.4(6)
'(7)-C(8)-H(8A)	119.8
'(9)-C(8)-H(8A)	119.8
'(10)-C(9)-C(8)	119.9(6)
'(10)-C(9)-H(9A)	120.1
'(8)-C(9)-H(9A)	120.1
'(9)-C(10)-C(11)	120.9(6)
'(9)-C(10)-Br(1)	119.7(5)
'(11)-C(10)-Br(1)	119.4(5)
'(10)-C(11)-C(12)	118.7(6)
'(10)-C(11)-H(11A)	120.6
'(12)-C(11)-H(11A)	120.6
'(7)-C(12)-C(11)	121.5(5)
'(7)-C(12)-H(12A)	119.3
'(11)-C(12)-H(12A)	119.3

ymmetry transformations used to generate equivalent atoms:

Table 4. Anisotropic displacement parameters ($\text{\AA}^2 \times 10^3$) for ba96f2as. The anisotropic displacement factor exponent takes the form: $-2\pi^2 [h^2 a^{*2} U^{11} + \dots + 2 h k a^* b^* U^{12}]$

	U11	U22	U33	U23	U13	U12
r(1)	89(1)	42(1)	42(1)	14(1)	8(1)	-5(1)
(1)	31(2)	33(3)	47(2)	5(2)	12(2)	-5(2)
(2)	30(2)	15(2)	42(2)	3(2)	16(2)	-5(2)
(3)	24(2)	15(2)	42(2)	6(2)	7(2)	-4(2)
(4)	40(2)	12(2)	44(2)	4(1)	9(2)	4(2)
'(1)	37(4)	20(3)	32(3)	6(2)	5(3)	1(3)
'(2)	39(3)	25(4)	34(3)	1(2)	5(2)	-1(3)
'(3)	37(3)	16(3)	38(3)	5(3)	7(2)	-8(3)
'(4)	18(3)	21(3)	46(4)	7(2)	5(3)	-3(2)
'(5)	21(3)	18(4)	37(3)	4(2)	6(2)	-4(2)
'(6)	26(3)	28(3)	35(3)	8(2)	-1(3)	2(3)
'(7)	30(3)	19(4)	33(3)	-2(2)	3(2)	-6(2)
'(8)	30(3)	36(4)	42(3)	3(4)	9(2)	0(4)
'(9)	42(4)	33(3)	42(4)	5(3)	15(3)	2(3)
'(10)	44(4)	25(3)	42(4)	-1(3)	8(3)	-4(3)
'(11)	38(4)	37(4)	55(4)	15(3)	2(3)	12(3)
'(12)	39(4)	45(5)	35(3)	10(3)	13(3)	6(3)

	x	y	z	U(eq)
i(3C)	3905	3273	4265	41
i(4B)	4492	-2666	3981	48
i(2A)	1299	268	7512	39
i(2B)	969	1815	6264	39
i(3A)	3352	-230	7414	37
i(3B)	3086	2340	6554	37
i(4A)	3811	-2255	5833	34
i(5A)	2154	1511	4367	31
i(6A)	2016	-2619	3304	37
i(8A)	4580	-282	2116	43
i(9A)	4429	2520	469	46
i(11A)	739	3890	492	53
i(12A)	900	1124	2149	47

Table 6. Torsion angles [°] for ba96f2as.

i(4)-O(2)-C(1)-O(1)	179.7(5)
i(4)-O(2)-C(1)-C(2)	-0.6(6)
i(1)-C(1)-C(2)-C(3)	164.4(6)
i(2)-C(1)-C(2)-C(3)	-15.3(6)
i(1)-C(2)-C(3)-C(4)	24.1(6)
i(1)-O(2)-C(4)-C(3)	16.4(6)
i(1)-O(2)-C(4)-C(5)	-105.9(5)
i(2)-C(3)-C(4)-O(2)	-25.2(5)
i(2)-C(3)-C(4)-C(5)	93.7(6)
i(2)-C(4)-C(5)-O(3)	-172.5(4)
i(3)-C(4)-C(5)-O(3)	70.4(6)
i(2)-C(4)-C(5)-C(6)	-52.4(6)
i(3)-C(4)-C(5)-C(6)	-169.4(4)
i(3)-C(5)-C(6)-O(4)	60.7(5)
i(4)-C(5)-C(6)-O(4)	-56.4(6)
i(3)-C(5)-C(6)-C(7)	-62.3(6)
i(4)-C(5)-C(6)-C(7)	-179.3(5)
i(4)-C(6)-C(7)-C(12)	160.5(5)
i(5)-C(6)-C(7)-C(12)	-76.4(7)
i(4)-C(6)-C(7)-C(8)	-22.5(7)
i(5)-C(6)-C(7)-C(8)	100.6(6)
i(12)-C(7)-C(8)-C(9)	0.2(10)
i(6)-C(7)-C(8)-C(9)	-176.8(6)
i(7)-C(8)-C(9)-C(10)	0.5(10)
i(8)-C(9)-C(10)-C(11)	-1.6(10)
i(8)-C(9)-C(10)-Br(1)	178.8(5)
i(9)-C(10)-C(11)-C(12)	1.9(10)
r(1)-C(10)-C(11)-C(12)	-178.4(5)
i(8)-C(7)-C(12)-C(11)	0.2(9)
i(6)-C(7)-C(12)-C(11)	177.3(6)
i(10)-C(11)-C(12)-C(7)	-1.2(10)

Symmetry transformations used to generate equivalent atoms:

Table 7. Hydrogen bonds for ba96f2as [Å and °].

D-H...A	d(D-H)	d(H...A)	d(D...A)	<(DHA)
i(3)-H(3C)...O(4)#1	0.84	1.87	2.698(5)	170.6
i(4)-H(4B)...O(3)#2	0.84	2.02	2.759(5)	147.0
i(4)-H(4B)...O(3)	0.84	2.46	2.885(5)	112.6

Symmetry transformations used to generate equivalent atoms:

1 x,y+1,z #2 -x+1,y-1/2,-z+1