

Aryldihydronaphthalene-type lignans from *Bursera fagaroides* var. *fagaroides* and their antimitotic mechanism of action

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Alvarez id

Sample: BF 4.15x2
Sample ID: s_20120222_03
File: 0703.fid

Pulse Sequence: s2pul

Solvent: cdcl3
Temp. 25.0 C / 298.1 K
Sample #7, Operator: gabby
File: 0703
Mercury-400BB "mercury400"

Relax. delay 1.000 sec
Pulse 45.0 degrees
Acq. time 1.998 sec
Width 3856.5 Hz
8 repetitions
OBSERVE H1, 399.6281478 MHz
DATA PROCESSING
FT size 16384
Total time 0 min, 25 sec

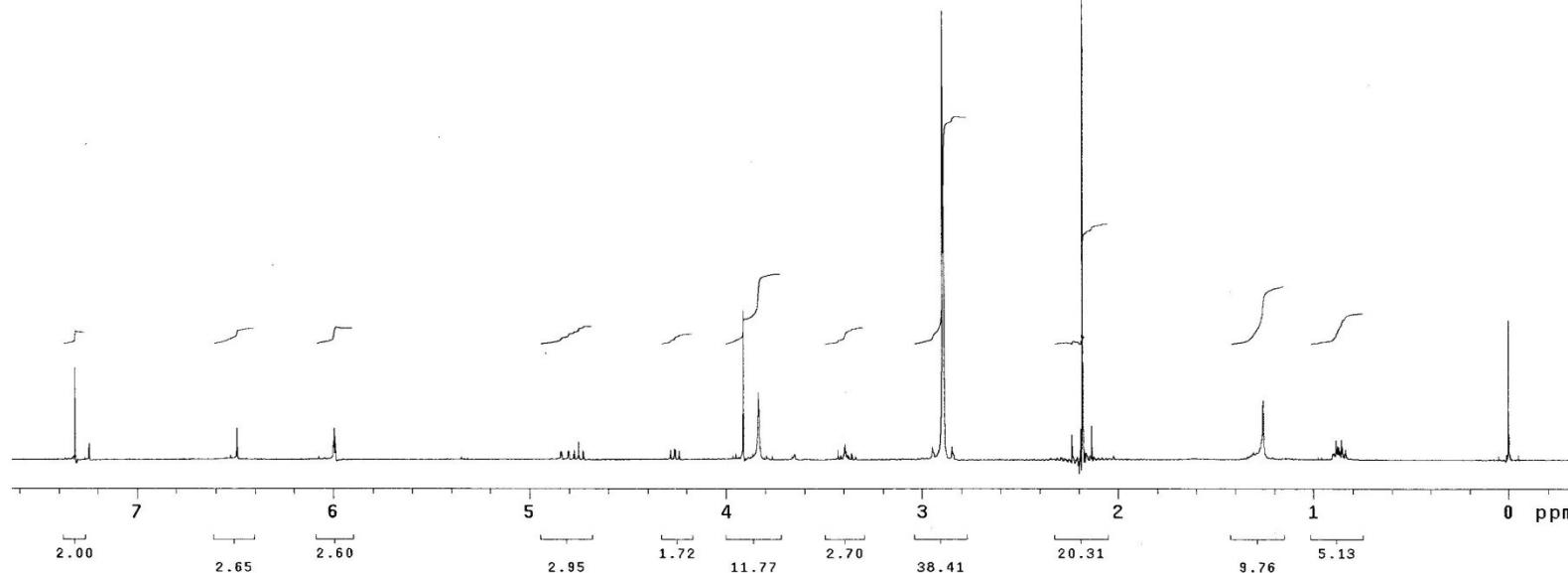


Figure S.1.1 ^1H NMR spectrum of compound 1 (400 MHz, CDCl_3)

¹³C
BF 4.15x2
CDCl₃
100MHz
Alvarez
27-04-12

Pulse Sequence: s2pul

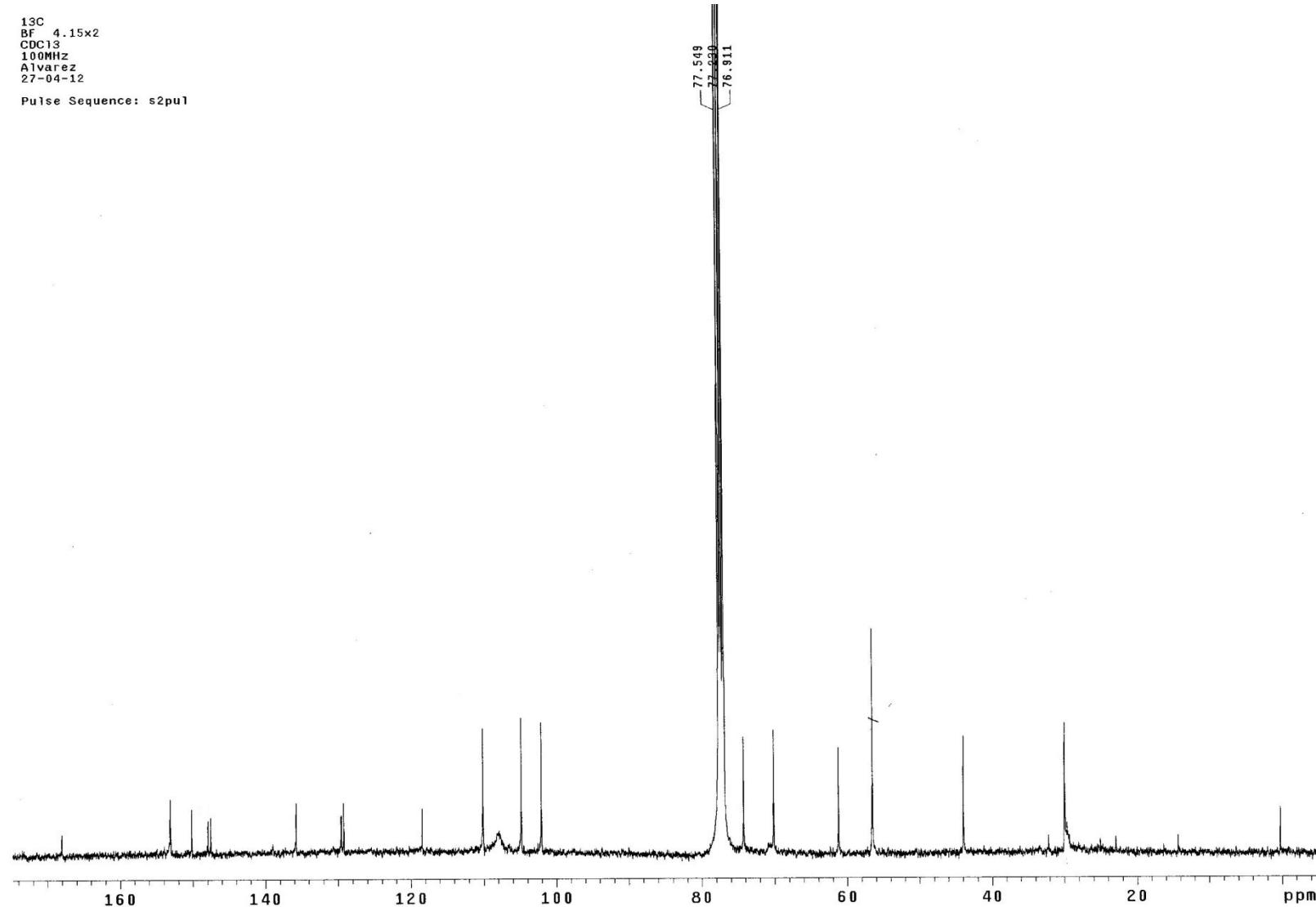


Figure S.1.2. ¹³C NMR spectrum of compound 1 (100 MHz, CDCl₃)

Alvarez id

Sample: BF 4.15x2
Sample ID: s_20120222_03
File: 0702.fid

Pulse Sequence: gHSQC

Solvent: cdcl3
Temp. 25.0 C / 298.1 K
Sample #7, Operator: gaby
File: 0702
Mercury-400BB "mercury400"

Relax. delay 1.301 sec
Acc. time 0.199 sec
Width 3956.5 Hz
2D Width 17083.1 Hz
32 repetitions
2 x 200 increments
OBSERVE H1, 399.6281421 MHz
DECOUPLE C13, 100.4941204 MHz
Power 45 dB
on during acquisition
off during delay
GARP-1 modulated
DATA PROCESSING
Gauss apodization 0.029 sec
F1 DATA PROCESSING
Gauss apodization 0.005 sec
FT size 4096 x 2048
Total time 5 hr, 39 min, 38 sec

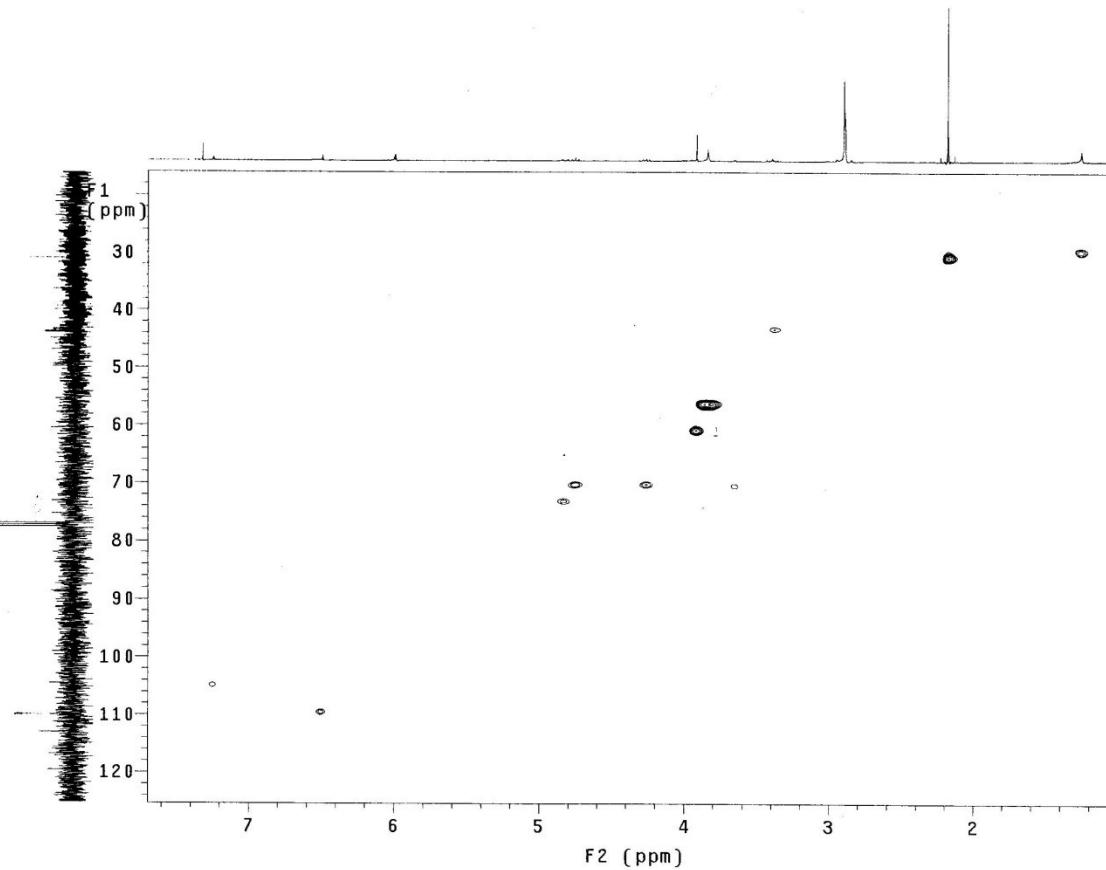


Figure S.1.3. HSQC spectrum of compound 1 (CDCl_3)

Alvarez id
Sample: BF 4.15x2
Sample ID: s_20120222_03
File: 0706.fid

Pulse Sequence: gHMBC
Solvent: cdc13
Temp: 25.0 C / 298.1 K
Sample ID Operator: gabry
File: 0706
Mercury-400BB "mercury400"

Relax. delay 1.500 sec
Mixing 0.080 sec
Acq time 0.128 sec
Width 3856.5 Hz
2D Width 2438.2 Hz
64 repetitions
200 increments
OBSERVE H1 399.6281436 MHz
DATA PROCESSING
Sine bell 0.055 sec
F1 DATA PROCESSING
Sine bell 0.005 sec
FT size 2048 x 2048
Total time 6 hr, 13 min, 23 sec

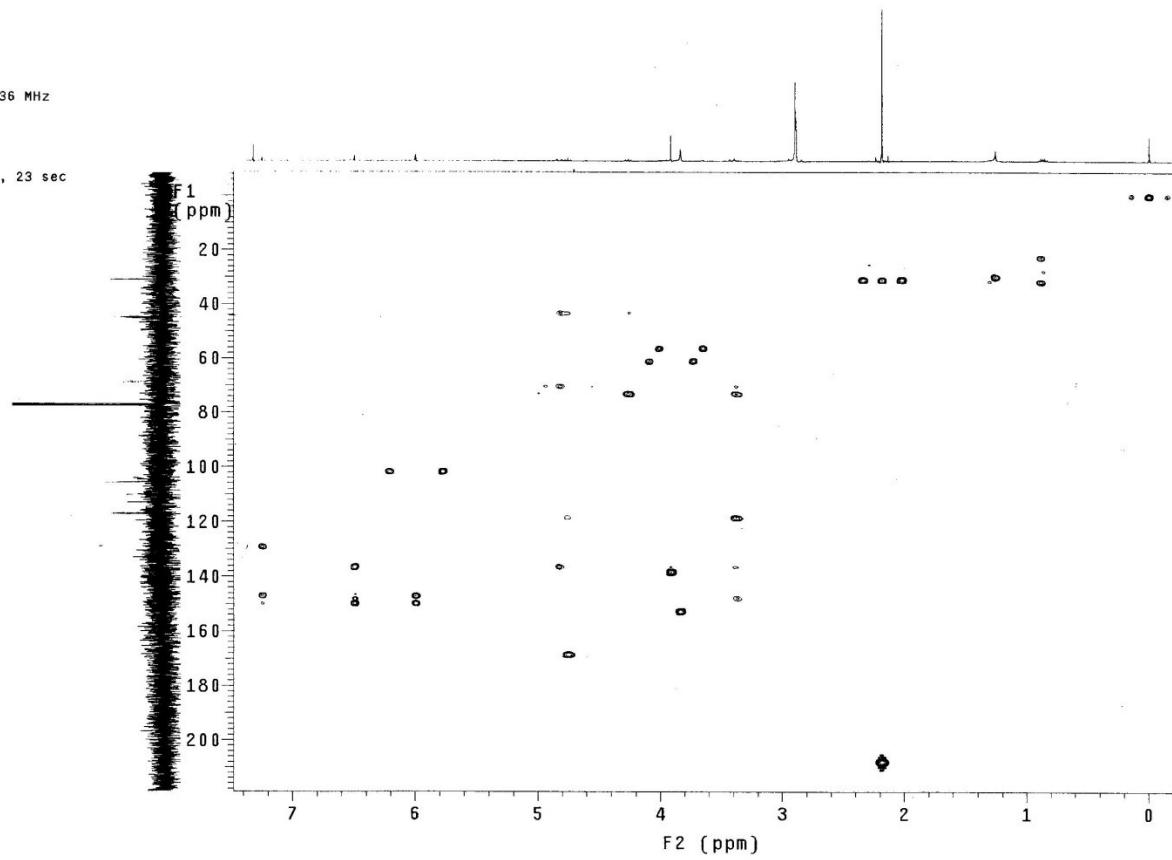


Figure S.1.4. HMBC spectrum of compound 1(CDCl_3).

[Mass Spectrum]
 Data : LAB008 Date : 07-Mar-2012 01:38
 Sample: BF 4.5 Operator name Ing.Victoria Labastida G.
 Note : Dra.Laura Alvarez/Andres Centro de Investigaciones Químicas UAE
 Inlet : Direct Ion Mode : FAB+
 Spectrum Type : Normal Ion [MF-Linear]
 RT : 0.13 min Scan# : (2,3)
 BP : m/z 69.0000 Int. : 2.49
 Output m/z range : 40.0000 to 800.0000 Cut Level : 0.00 %

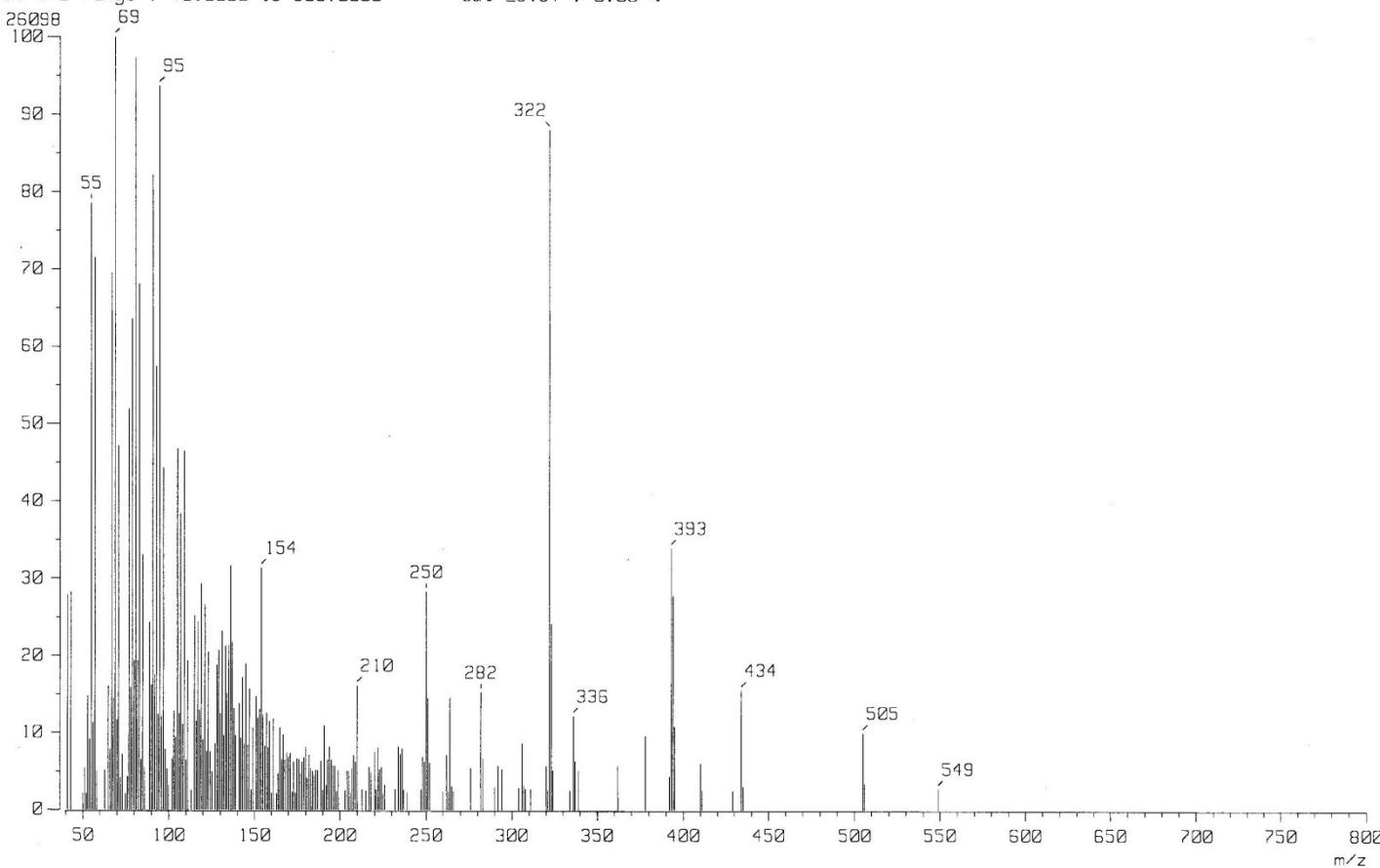


Figure S.1.5. Mass spectrum of compound 1.

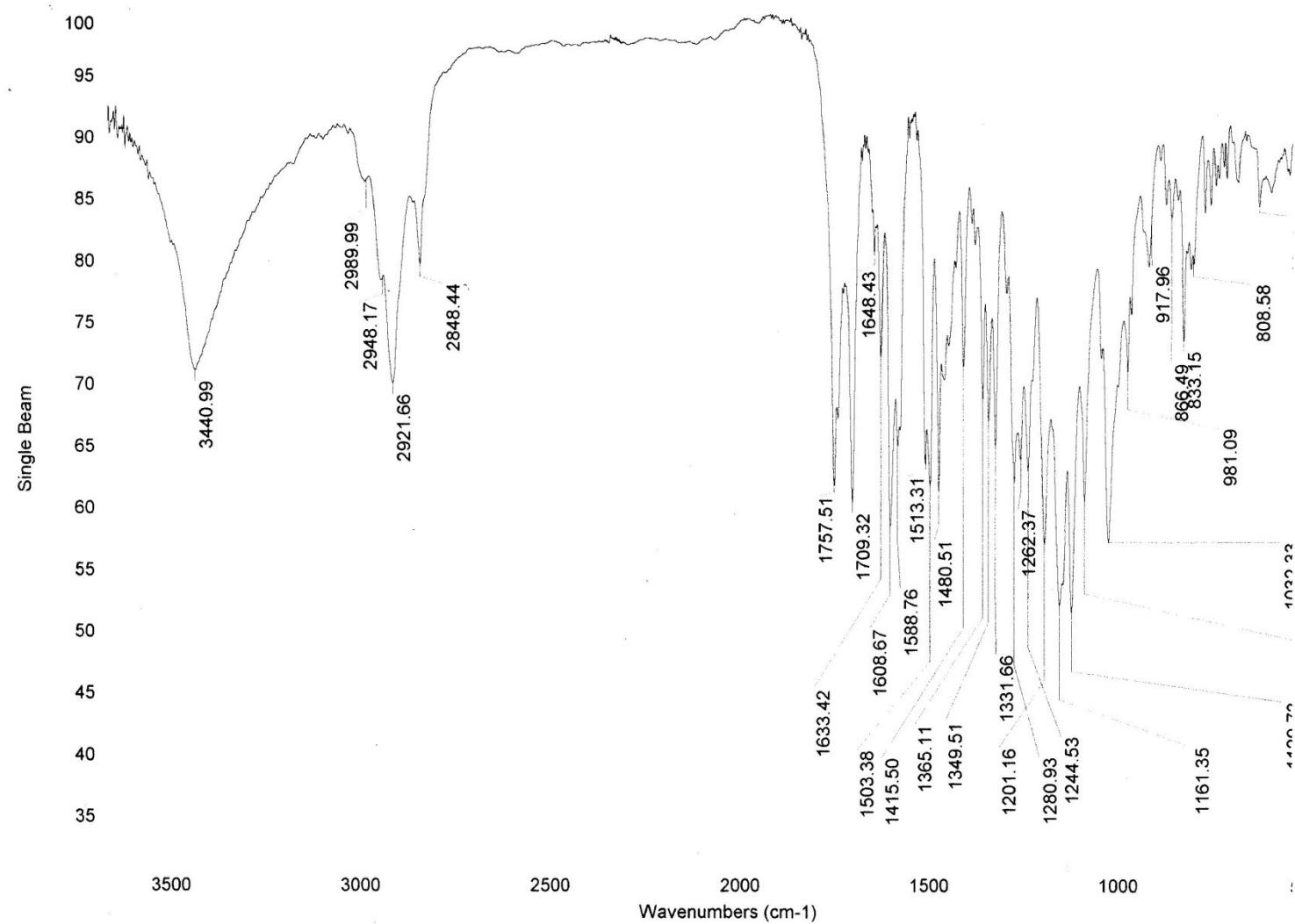


Figure S.1.6. IR spectrum of compound **1**.

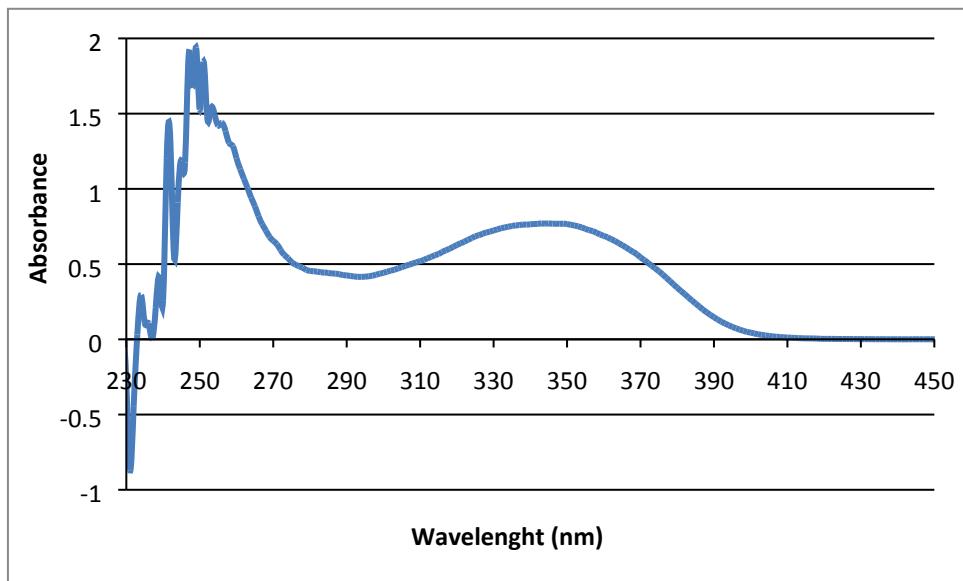


Figure S.1.7. UV spectrum of compound 1

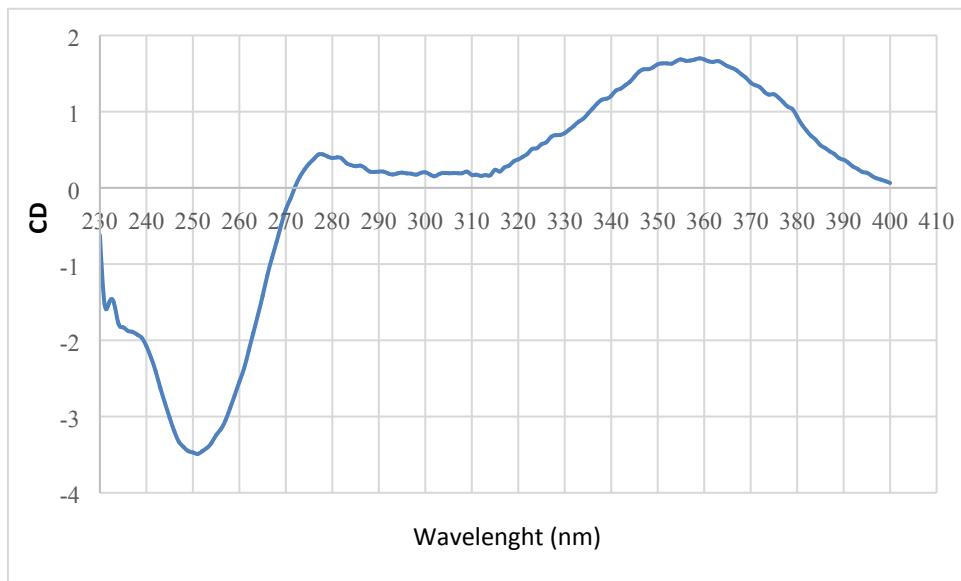


Figure S.1.8. CD spectrum of compound 1

AcPdx
1H-RMN
400MHz-id
CDCl₃-DMSO_{d6}
Alvarez
27-02-14
MARC

File: /home/walkup/vnmrsys/data/ene-feb-14/Alvarez/AcPdx-1H.fid
Pulse Sequence: s2pul

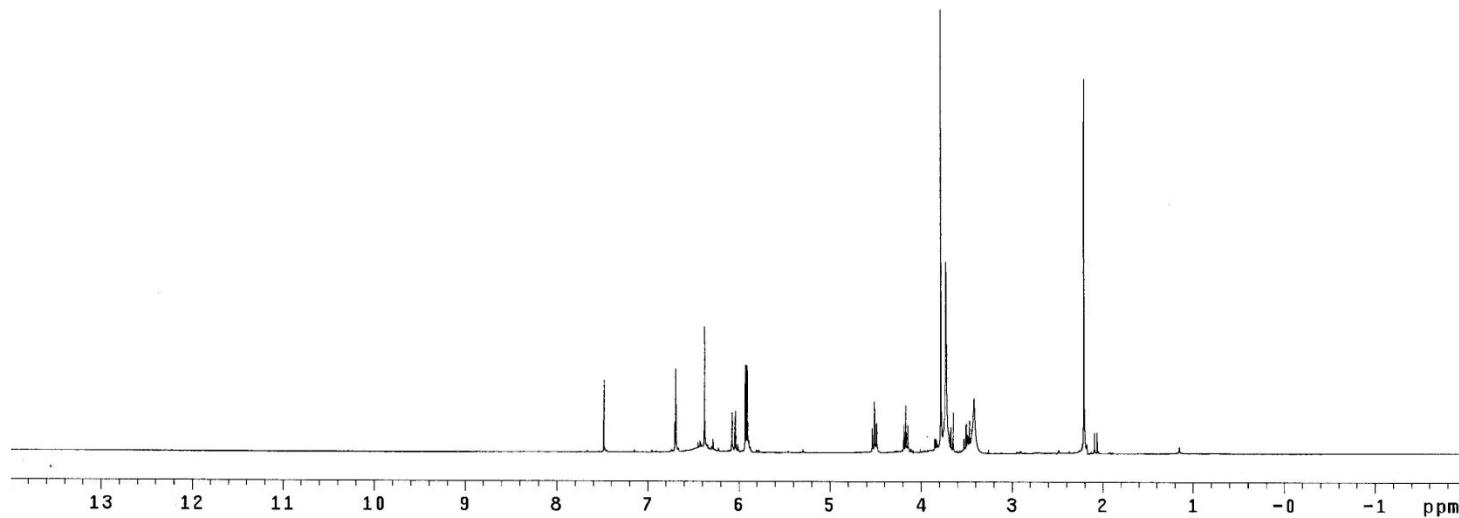


Figure S.2.1. ¹H NMR spectrum of compound 2 (400 MHz, CDCl₃)

AcPdx
13C-RMN
100MHz-1d
CDC13-DMS0d6
Alvarez
27-02-14
MARC

File: exp
Pulse Sequence: s2pul

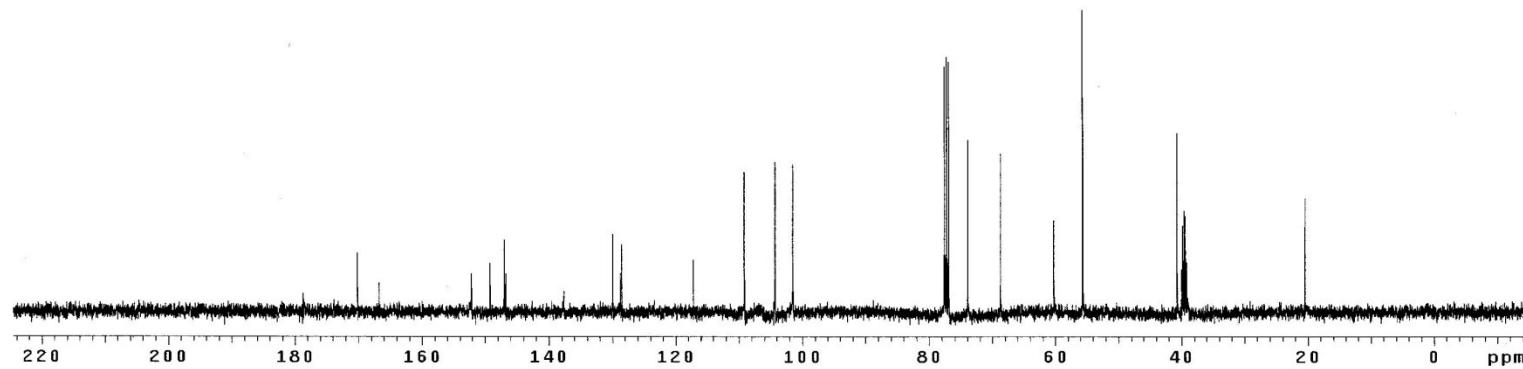


Figure S.2.2. ¹³C NMR spectrum of compound 2 (100 MHz, CDCl₃).

```

AcPdx
1H-RMN
400MHz-id
CDC13-DMSOd6
Alvarez
27-02-14
MARC

File: exp
Pulse Sequence: gHSQC
Solvent: dmso
Ambient temperature
Operator: walkup
INOVA-400 "localhost.localdomain"

Relax, delay 1.000 sec
Aqc. time 0.199 sec
Width 3800.7 Hz
2D Width 17101.3 Hz
32 repetitions
2 x 128 increments
OBSERVE H1, 400.0691404 MHz
DECOUPLE C13, 100.6050102 MHz
Power 41 dB
on during acquisition
off during delay
GARP-1 modulated
DATA PROCESSING
Gauss apodization 0.092 sec
F1 DATA PROCESSING
Gauss apodization 0.007 sec
F1 size 4096 x 2048
Total time 2 hr, 51 min, 31 sec

```

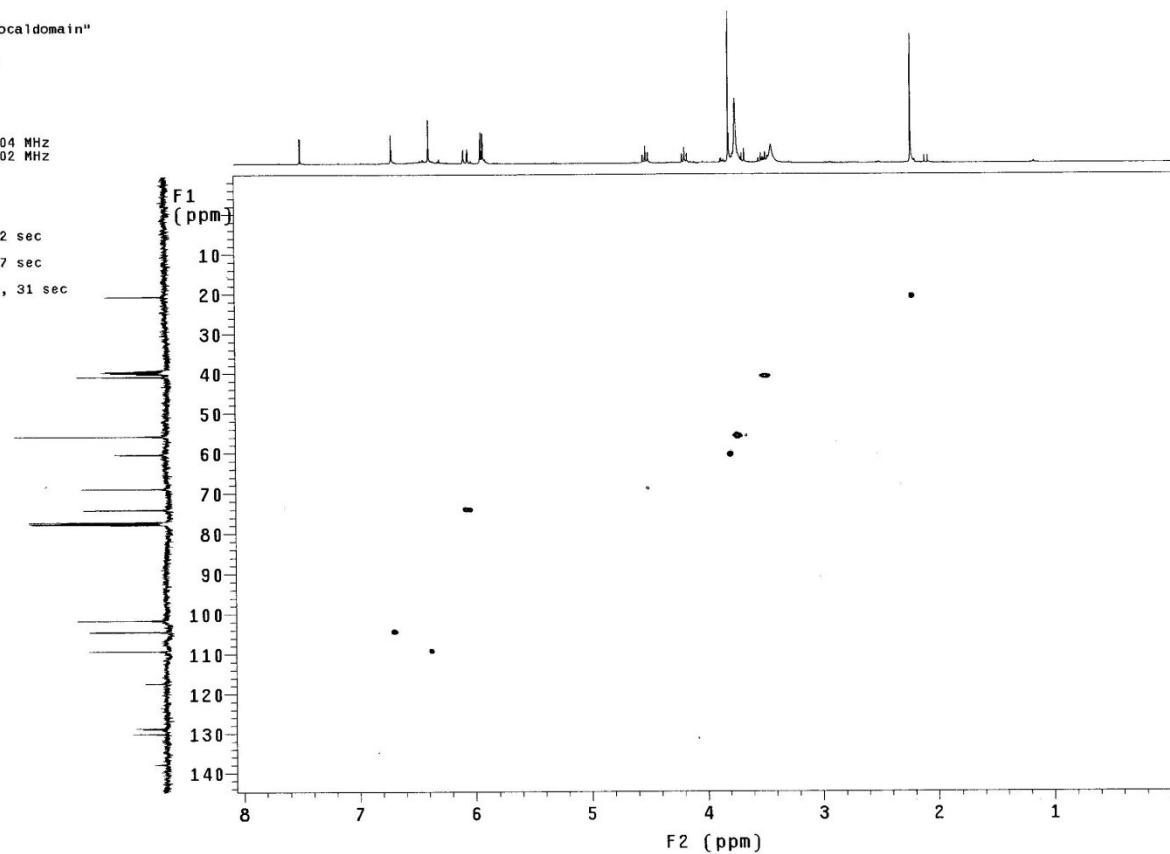


Figure S.2.3. HSQC spectrum of compound **2** (CDCl_3).

AcPdx
1H-RMN
400MHz-id
CDC13-DMS0d6
Alvarez
04-03-14
MARC

File: exp

Pulse Sequence: gHMBC

Solvent: dmso

Ambient temperature

Operator: walkup

INNOVA-400 "localhost.localdomain"

Relax. delay 1.000 sec
Mixing 0.080 sec
Acq. time 0.128 sec
Width 3000.5 Hz
2D Width 24140.0 Hz
64 repetitions
200 increments
OBSERVE H1, 400.0691354 MHz
DATA PROCESSING
Sine bell 0.064 sec
F1 DATA PROCESSING
Sine bell 0.008 sec
FT size 2048 2048
Total time 4 hr, 20 min, 29 sec

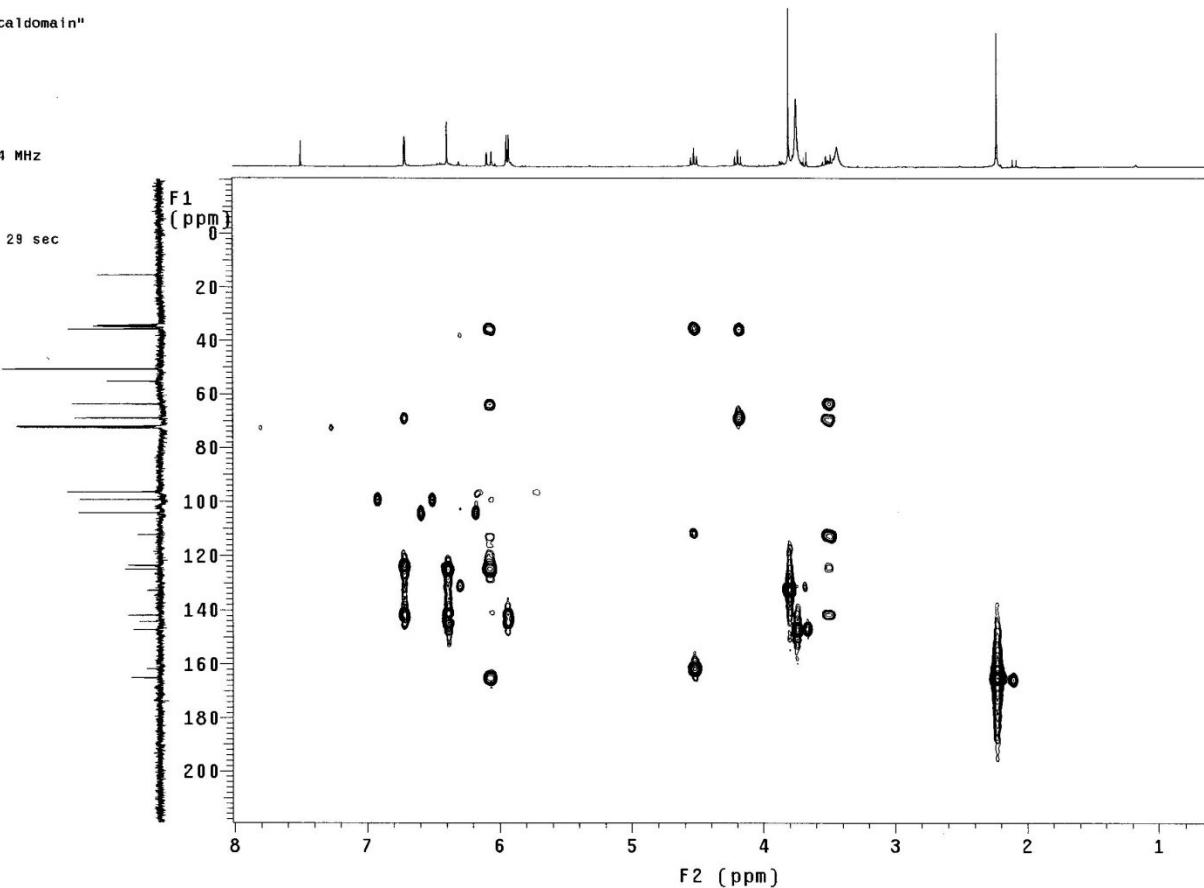


Figure S.2.4. HMBC spectrum of compound 2(CDCl_3)

[Mass Spectrum]
Data : LAB029 Date : 11-Dec-2015 11:58
Sample: Comp A(2) Operator Gabriela Vargas Instrument: MStation JMS-7
Note : Dra.Laura Alvarez/Mayra Centro de Investigaciones Químicas URCM
Inlet : Direct Ion Mode : PFB+
Spectrum Type : Normal Ion DMF-Linear
RT : 0.31 min Scan# : 2
BP : m/z 55.0000 Int. : 4.68
Output m/z range : 0.0000 to 682.1958 Cut Level : 0.00 %

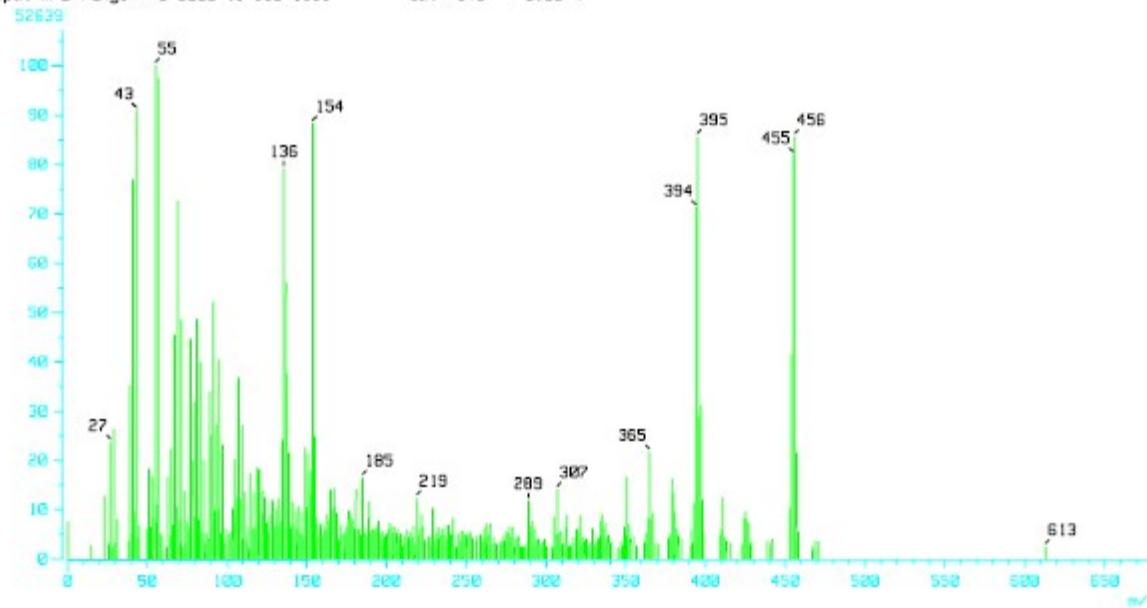


Figure S.2.5. Mass spectrum of compound 2.

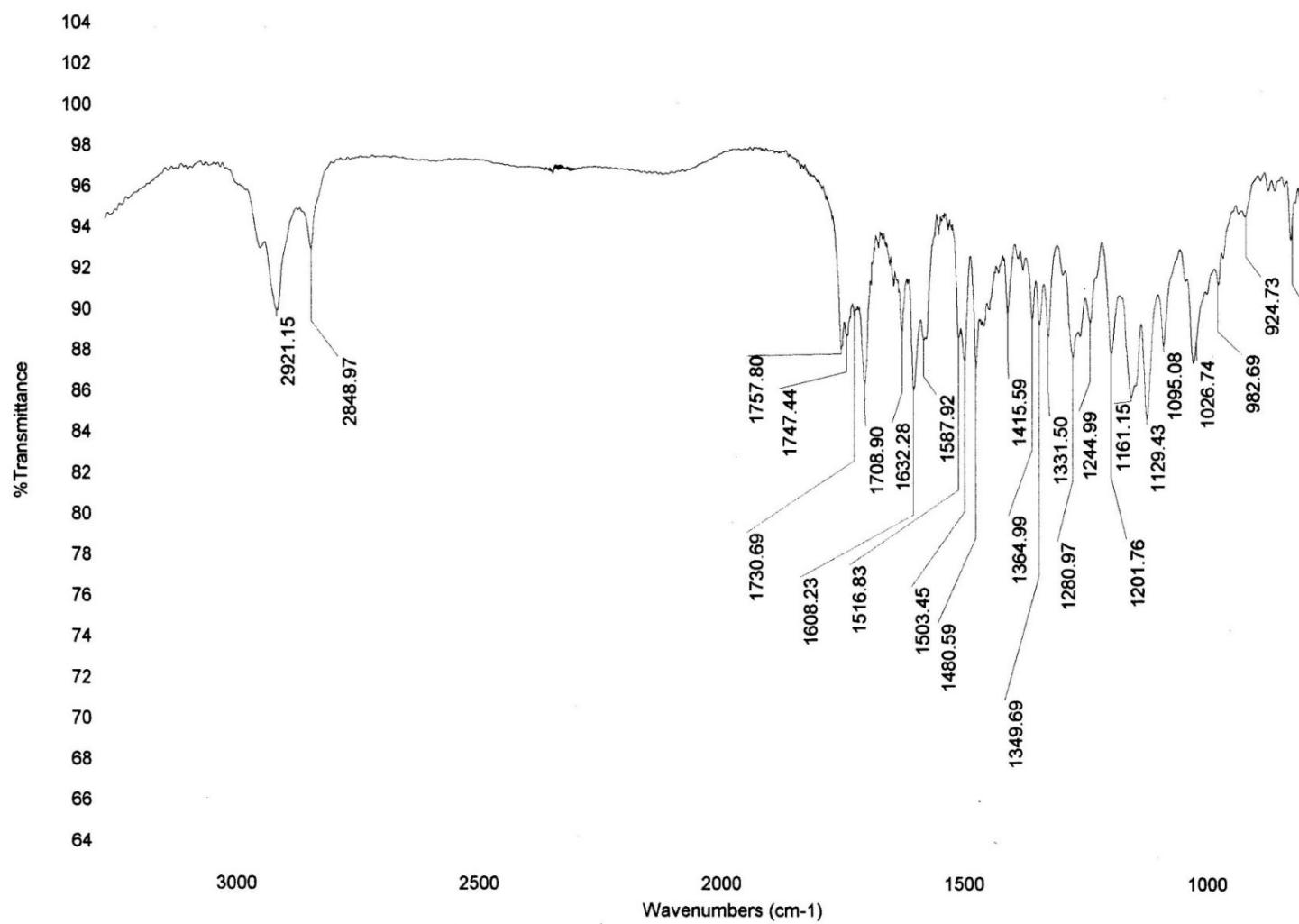


Figure S.2.6. IR spectrum of compound 2.

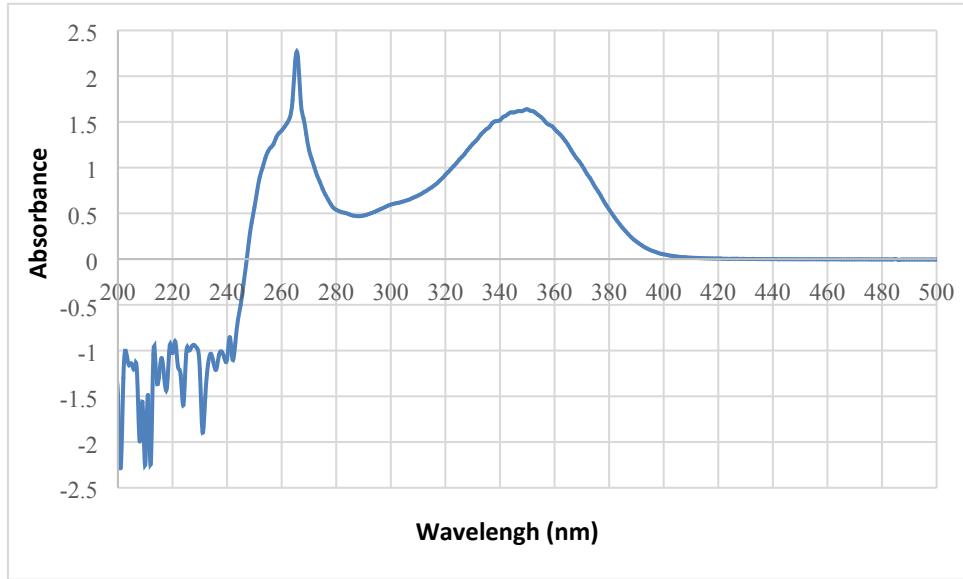


Figure S.2.7. UV spectrum of compound **2**.

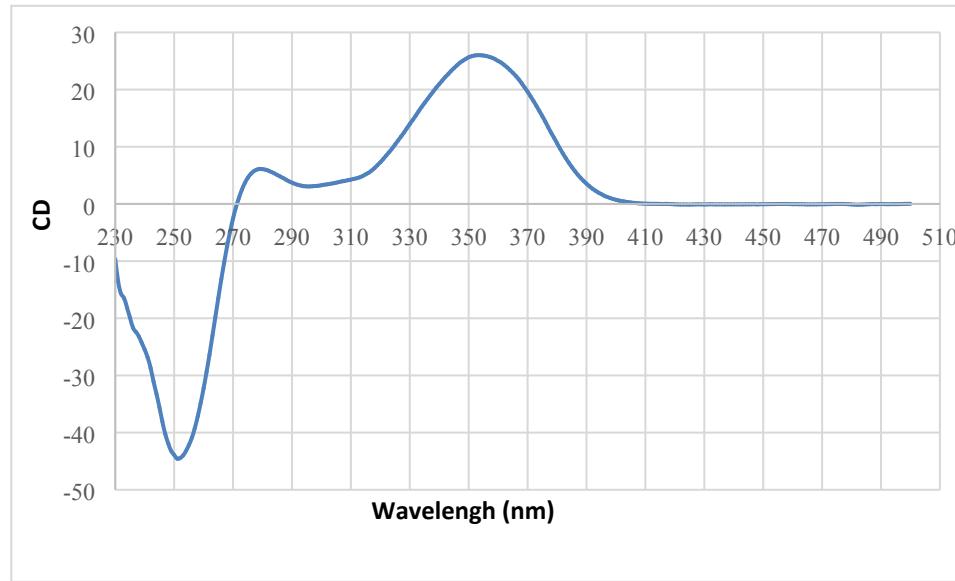


Figure S.2.8. CD spectrum of compound **2**.

¹H
BF1.4
CDCl₃
400MHz
Alvarez
17-02-12

Pulse Sequence: s2pu1

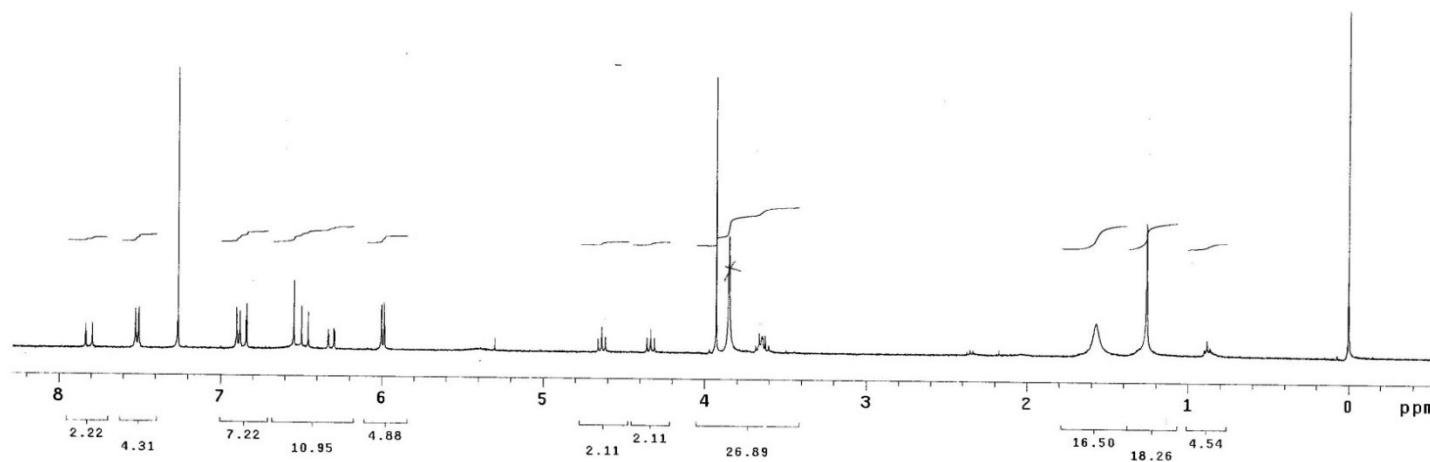


Figure S.3.1. ¹H NMR spectrum of compound 3 (400 MHz, CDCl₃)

¹³C
BF1.4
CDCl₃
100MHz
Alvarez
17-02-12
Pulse Sequence: s2pul

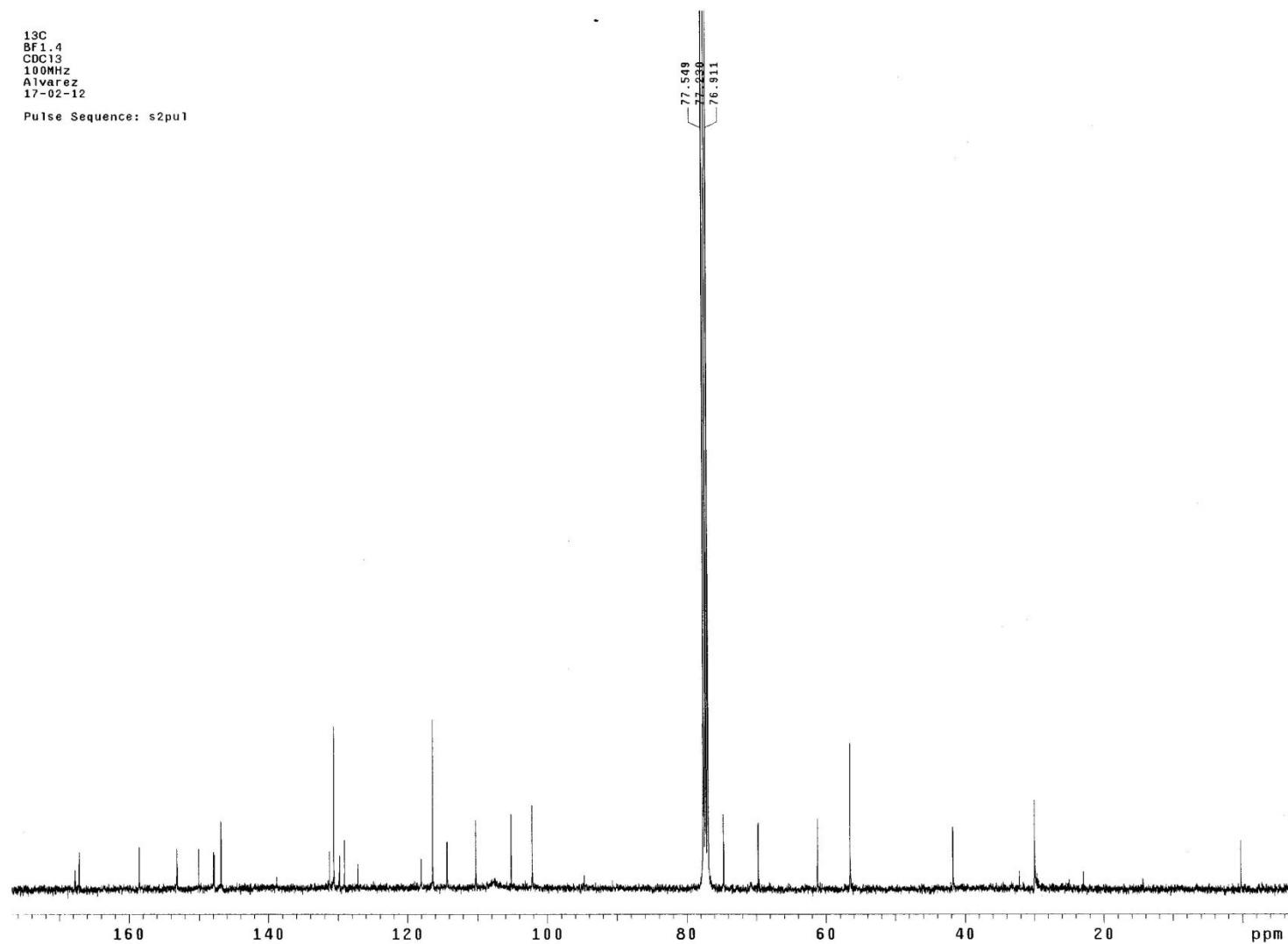


Figure S.3.2. ¹³C NMR spectrum of compound 3 (100 MHz, CDCl₃)

```

Alvarez id
Sample: BF 1.4
Sample ID: s_20120220_08
File: 0510.fid
Pulse Sequence: gHSQC
Solvent: cdc13
Temp. 25.0 C / 298.1 K
Sample #5, Operator: gaby
File: 0510
Mercury-400BB "mercury400"

Relax. delay 1.301 sec
Acq. time 0.199 sec
Width 4040.4 Hz
2D Width 17083.1 Hz
32 repetitions
2 x 200 increments
OBSERVE H1, 399.6281725 MHz
DECOUPLE C13, 100.4941204 MHz
Power 45 dB
On during acquisition
off during delay
GARP-1 modulated
DATA PROCESSING
Gauss apodization 0.039 sec
F1 DATA PROCESSING
Gauss apodization 0.005 sec
FT size 4096 x 4096
Total time 5 hr, 35 min, 36 sec

```

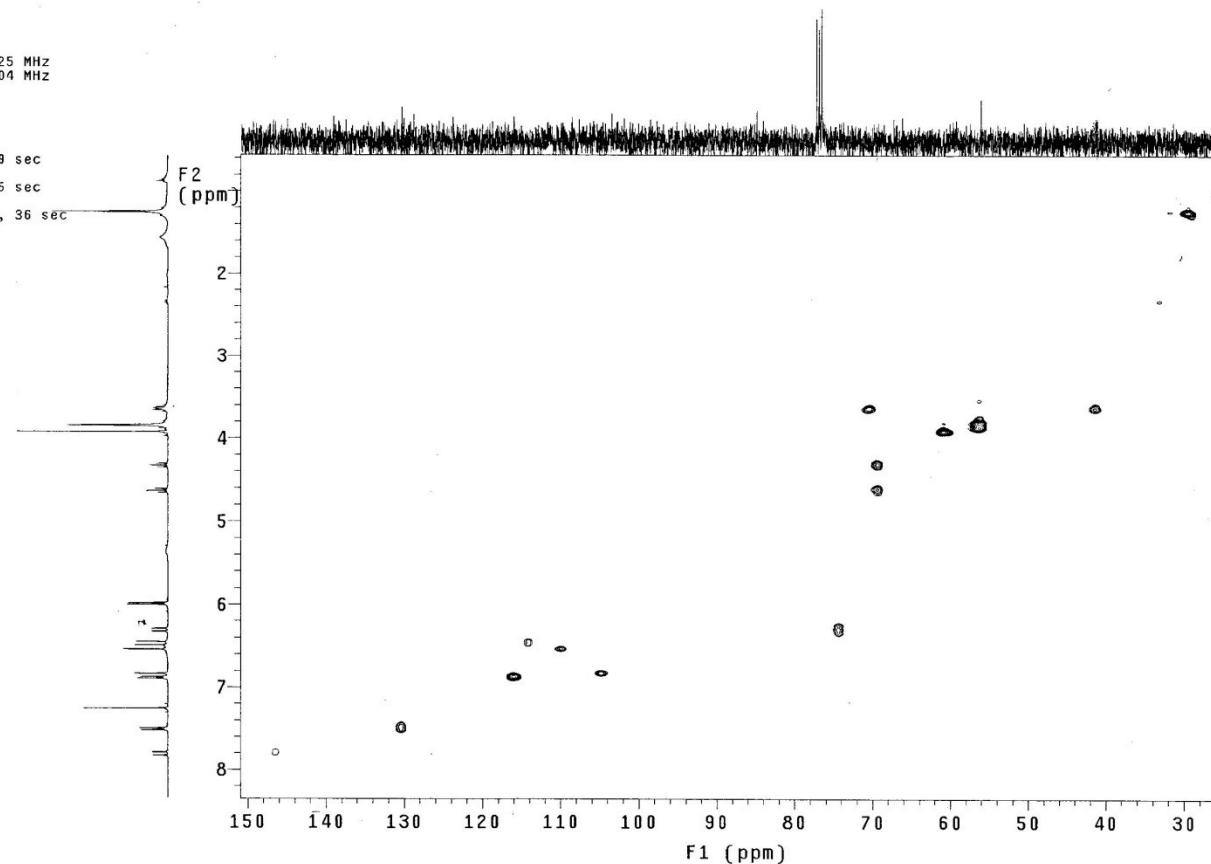


Figure S.3.3. HSQC spectrum of compound 3 (CDCl_3).

Alvarez id
 Sample: BF 1.4
 Sample ID: s_20120220_08
 File: 0511.fid
 Pulse Sequence: gHMBC
 Solvent: cdc13
 Temp. 25.0 C / 298.1 K
 Sample #5, Operator: gaby
 File: 0511
 Mercury-400BB "mercury400"
 Relax. delay 1.500 sec
 Mixing 0.080 sec
 Acq. time 0.128 sec
 Width 4040.4 Hz
 2D Width 24118.2 Hz
 64 Repetitions
 200 Averages
 OBSERVE H1 399.6281725 MHz
 DATA PROCESSING
 Sine bell 0.050 sec
 F1 DATA PROCESSING
 Sine bell 0.005 sec
 FT size 2048 x 2048
 Total time 6 hr, 13 min, 34 sec

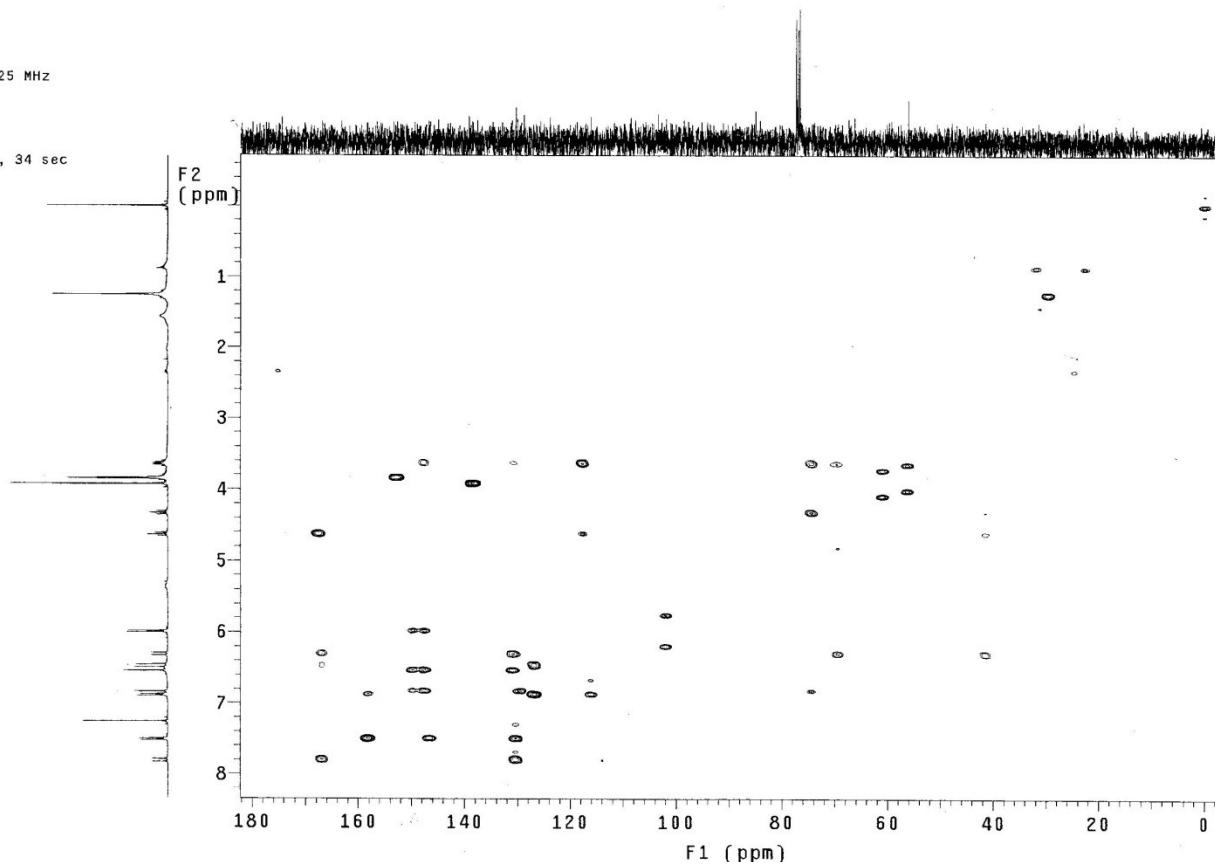


Figure S.3.4. HMBC spectrum of compound 3(CDCl_3)

[Mass Spectrum]
Data : LAB009 Date : 07-Mar-2012 01:45
Sample: BF 1.4 Operator name Ing.Victoria Labastida G.
Note : Dra.Laura Alvarez/Andres Centro de Investigaciones Químicas UAEM
Inlet : Direct Ion Mode : FAB+
Spectrum Type : Normal Ion [MF-Linear]
RT : 0.00 min Scan# : (1,2)
BP : m/z 55.0000 Int. : 8.22
Output m/z range : 40.0000 to 800.0000 Cut Level : 0.00 %

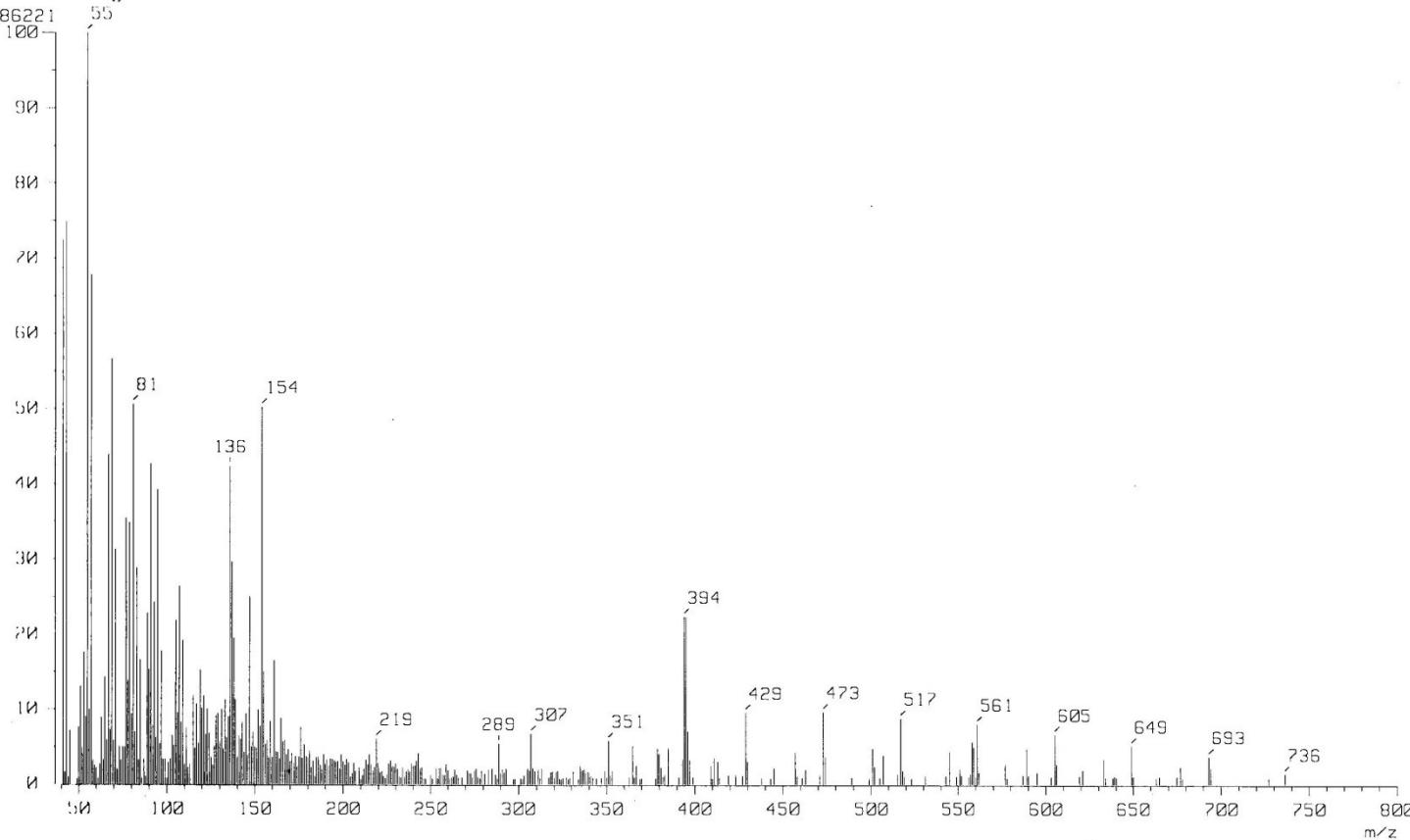


Figure S.3.5. Mass spectrum of compound 3

Table S.1. Quantification of the effect of *B. fagaroides* lignans **1-3** on the cell cycle in zebrafish embryos.

Treatment	Cell cycle activity	H3S10ph fold change	p <0.001	n	N	Student's t-test
DMSO		1.00 ± 0.13		10	1	
Aphidicolin	-	0.23 ± 0.05	*	9	1	7.53E-12
Nocodazole	+	1.80 ± 0.20	*	9	1	9.81E-09
1	+	1.92 ± 0.23	*	10	1	2.41E-09
2	+	2.41 ± 0.19	*	10	1	1.73E-13
3	+	2.57 ± 0.26	*	10	1	1.88E-12

Cell cycle activity; -, denotes decrease, +, increase.

Table S.2. Quantification of the effect of *B. fagaroides* lignans **1-3** on the morphology of zebrafish embryos.

Treatment	Effect on morphology	Circularity	p<0.001	n	N	Student's t-test
DMSO		0.42 ± 0.02		10	1	
Aphidicolin	=	0.38 ± 0.05		9	1	4.90E-02
Nocodazole	+	0.76 ± 0.06	*	9	1	4.80E-12
1	+	0.78 ± 0.07	*	10	1	3.10E-12
2	+	0.73 ± 0.06	*	10	1	2.56E-11
3	+	0.77 ± 0.04	*	10	1	1.68E-15

Circularity, measured circularity. Morphological effect; +, increase in circularity, = without change. n, total number of embryos analyzed. Student's t-test was used to determine the p-value. **1** (7',8'-dehydropodophyllotoxin). **2**, (7',8'-dehydro acetyl podophyllotoxin). **3** (7',8'-dehydro-*trans*-p-coumaroyl podophyllotoxin).