

Supporting Information Section

Structures, Biogenesis, and Biological Activities of Pyrano[4,3-*c*]isochromene-4-one Derivatives from the Fungus *Phellinus igniarius*:

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List of Supporting Information

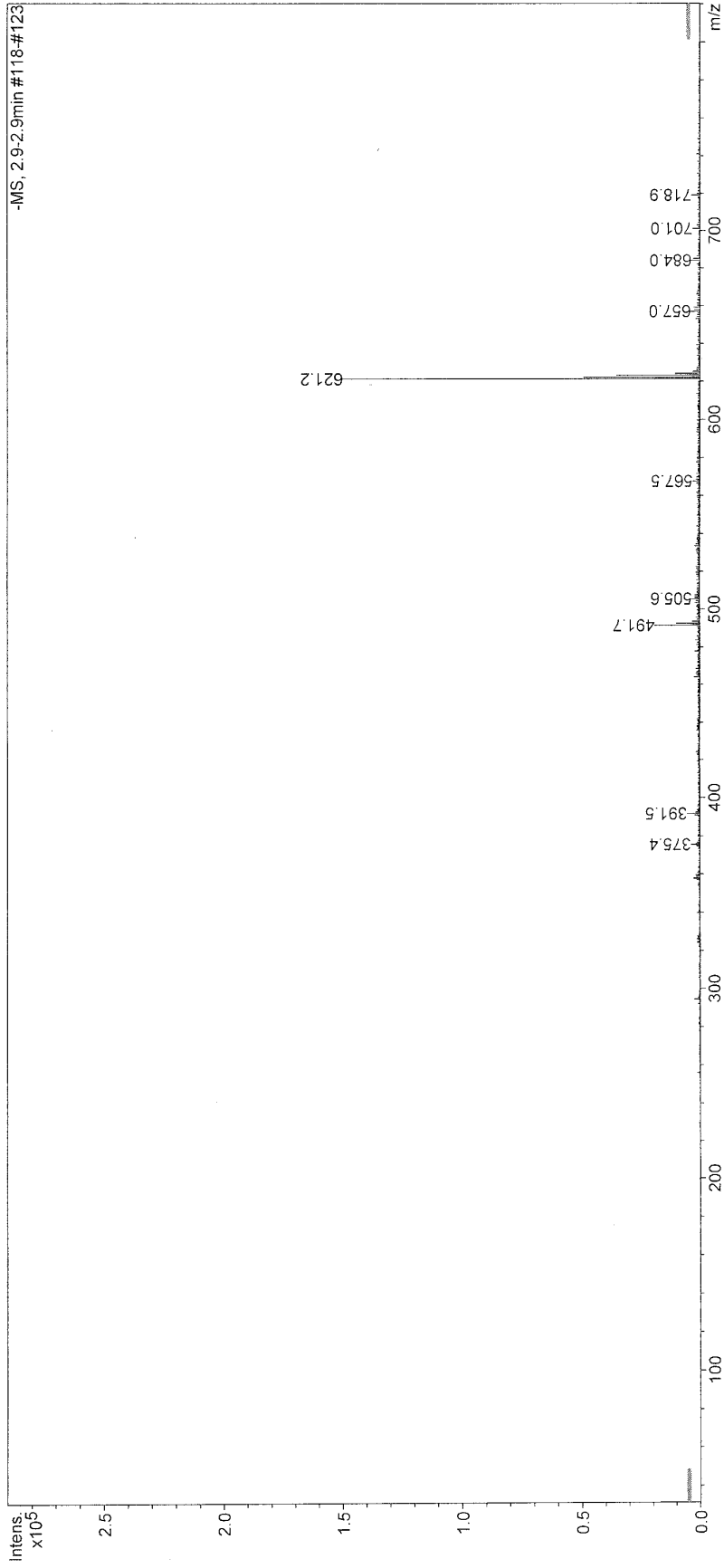
Page	Contents
S1	The negative mode ESIMS spectrum of Phelligrudin H (1)
S2	The HR-MALDI-FTMS spectrum of Phelligrudin H (1)
S3	IR spectrum of Phelligrudin H (1)
S4	UV spectrum of Phelligrudin H (1)
S5	The ¹ H NMR spectrum of Phelligrudin H (1)
S6	Regional enlarged ¹ H NMR spectrum of Phelligrudin H (1)
S7	The ¹³ C NMR spectrum of Phelligrudin H (1)
S8	The DEPT spectrum of Phelligrudin H (1)
S9	¹ H- ¹ H COSY spectrum of Phelligrudin H (1)
S10	HSQC spectrum of Phelligrudin H (1)
S11	HMBC spectrum of Phelligrudin H (1)
S12	Regional enlarged HMBC spectrum of Phelligrudin H (1)
S13	Regional enlarged HMBC spectrum of Phelligrudin H (1)
S14	The negative mode ESIMS spectrum of Phelligrudin I (2)
S15	The HR-MALDI-FTMS spectrum of Phelligrudin I (2)
S16	IR spectrum of Phelligrudin I (2)
S17	UV spectrum of Phelligrudin I (2)
S18	The ¹ H NMR spectrum of Phelligrudin I (2)
S19	Regional enlarged ¹ H NMR spectrum of Phelligrudin I (2)
S20	The ¹³ C NMR spectrum of Phelligrudin I (2)
S21	The DEPT spectrum of Phelligrudin I (2)
S22	¹ H- ¹ H COSY spectrum of Phelligrudin I (2)
S23	HSQC spectrum of Phelligrudin I (2)
S24	HMBC spectrum of Phelligrudin I (2)
S25	Regional enlarged HMBC spectrum of Phelligrudin I (2)
S26	Regional enlarged HMBC spectrum of Phelligrudin I (2)
S27	Regional enlarged HMBC spectrum of Phelligrudin I (2)
S28	IR Spectrum of Phelligrudin J (3)
S29	(-)-ESI-MS of Phelligrudin J (3)
S30	HRESI of Phelligrudin J (3)
S31	¹ H NMR Spectrum of Phelligrudin J (3)
S32	¹³ C NMR Spectrum of Phelligrudin J (3)
S33	gHMBC Spectrum of Phelligrudin J (3)
S34	Table of ¹ H and ¹³ C NMR assignments for Phelligrudins H-J (1-3)
S35	Scheme S1. Biogenetic scheme for phelligrudins H (1) and I (2) involving phelligrudin D, and phelligrudin J (3)
S36	Scheme S2. Biogenetic scheme for phelligrudins H (1) and I (2) involving hypholomine B and 3,14'-bihispidinyl.
S37	Scheme S3. Biogenetic scheme for davallialactone (4), phelligrudin F, and inoscavin A from hispidin and hispilone.

Display Report - Selected Window Selected Analysis

Analysis Name: E13C35D.d
Method: Copy of TEST.MS
Sample Name: E13C35D
Analysis Info:

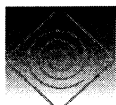
Instrument: LC-MSD-Trap-SL
Operator: Administrator

Print Date: 2006-3-24 8:12:52
Acq. Date: 2004-2-17 14:59:52



ESI-MS of Phelligrudin H (1)

Instrument:



IonSpec 4.7 Tesla FTMS

Card Serial Number: I041198

Sample Serial Number: SH-W6 C155

Operator: Hua Qin Date: 2004/08/03

Operation Mode: MALDI/DHB

Elemental Composition Search Report:

Target Mass:

Target m/z = 623.0808 ± 0.003
Charge = +1

Possible Elements:

Element:	Exact Mass:	Min:	Max:
C	12.000000	0	100
H	1.007825	0	100
O	15.994915	0	30

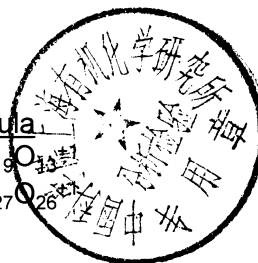
Additional Search Restrictions:

DBE Limit Mode = Both Integer and Half-Integer
Minimum DBE = 0

Search Results:

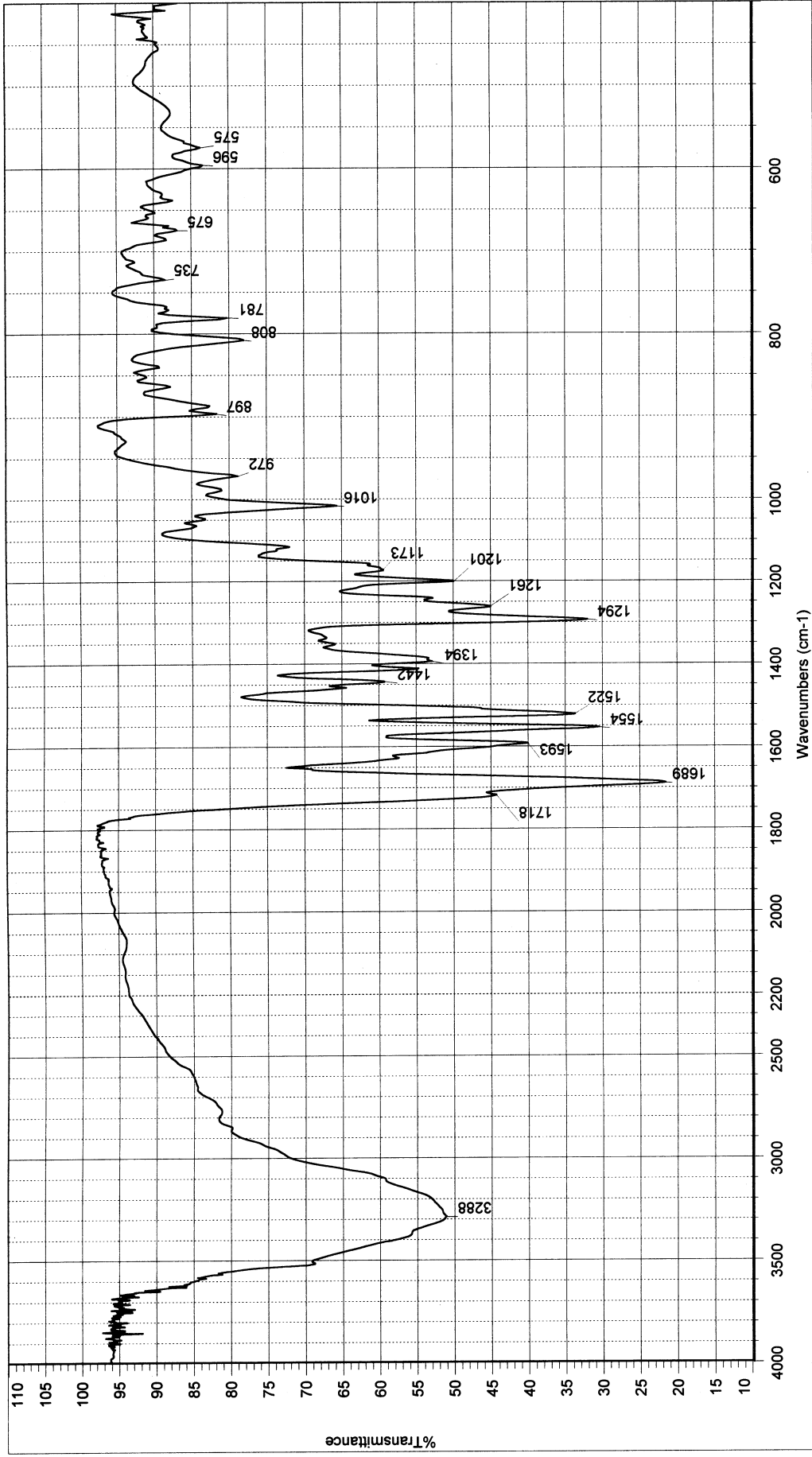
Number of Hits = 2

m/z	Delta m/z	DBE	Formula
623.08202	-0.00122	24.5	C ₃₃ H ₁₉ O ₁₃
623.07852	0.00228	2.5	C ₁₅ H ₂₇ O ₂₆



HR-MS of Phelligrudin H (1)

IR spectrum of Phelligrudin H (1)



Date: Wed Nov 10 10:02:00 2004

Sample Name: C155 (KBr)

Scans: 64

检测单位: 国家药物及代谢产物分析研究中心

Resolution: 4.000

检测仪器: 美国尼高力公司傅立叶变换红外光谱仪: IMPACT - 400

THERMO SPECTRONIC ~ VISION32 SOFTWARE V1.25

Batch Information - scan003

Batch Type Scan Operator Name (None Entered)
Instrument ID 110514 Aborted No

Results Table - scan003

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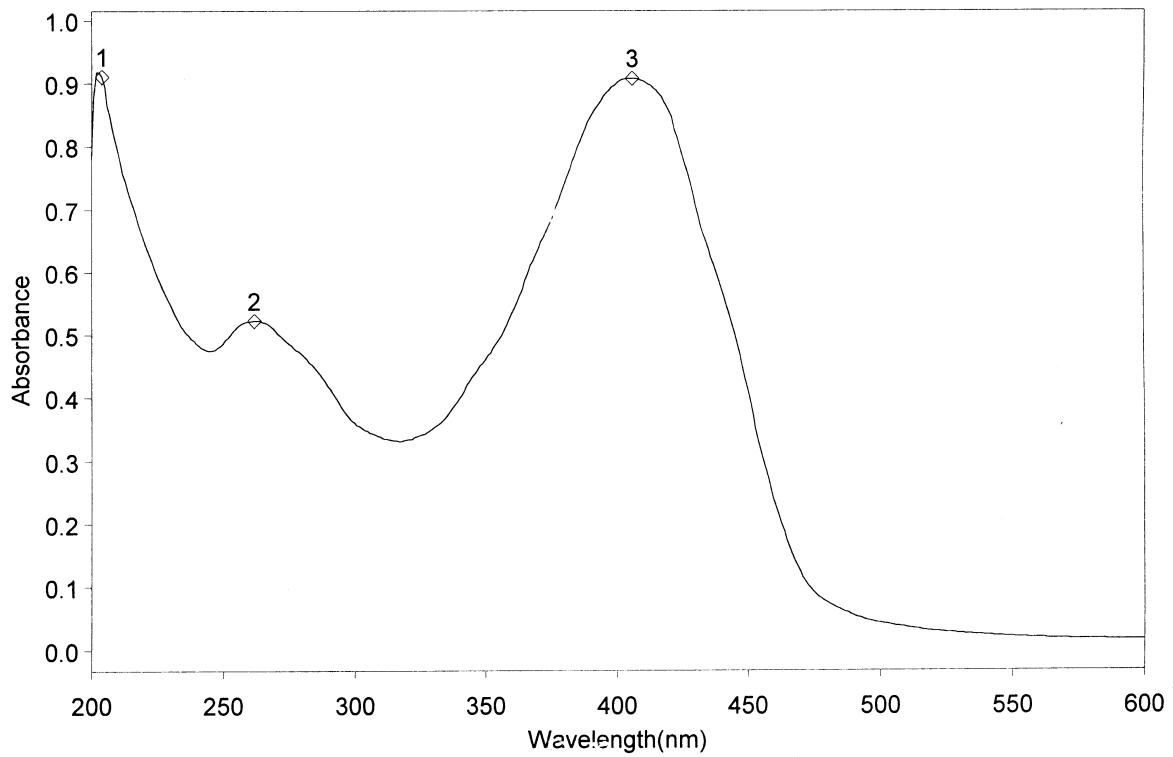
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2	Cycle01	nm	204.0	262.0	406.0
3	Manual	A	.911	.522	.908

All calculations have been performed to double precision as defined by ANSI/IEEE STD 754-1985 but have been rounded for display purposes.

C155

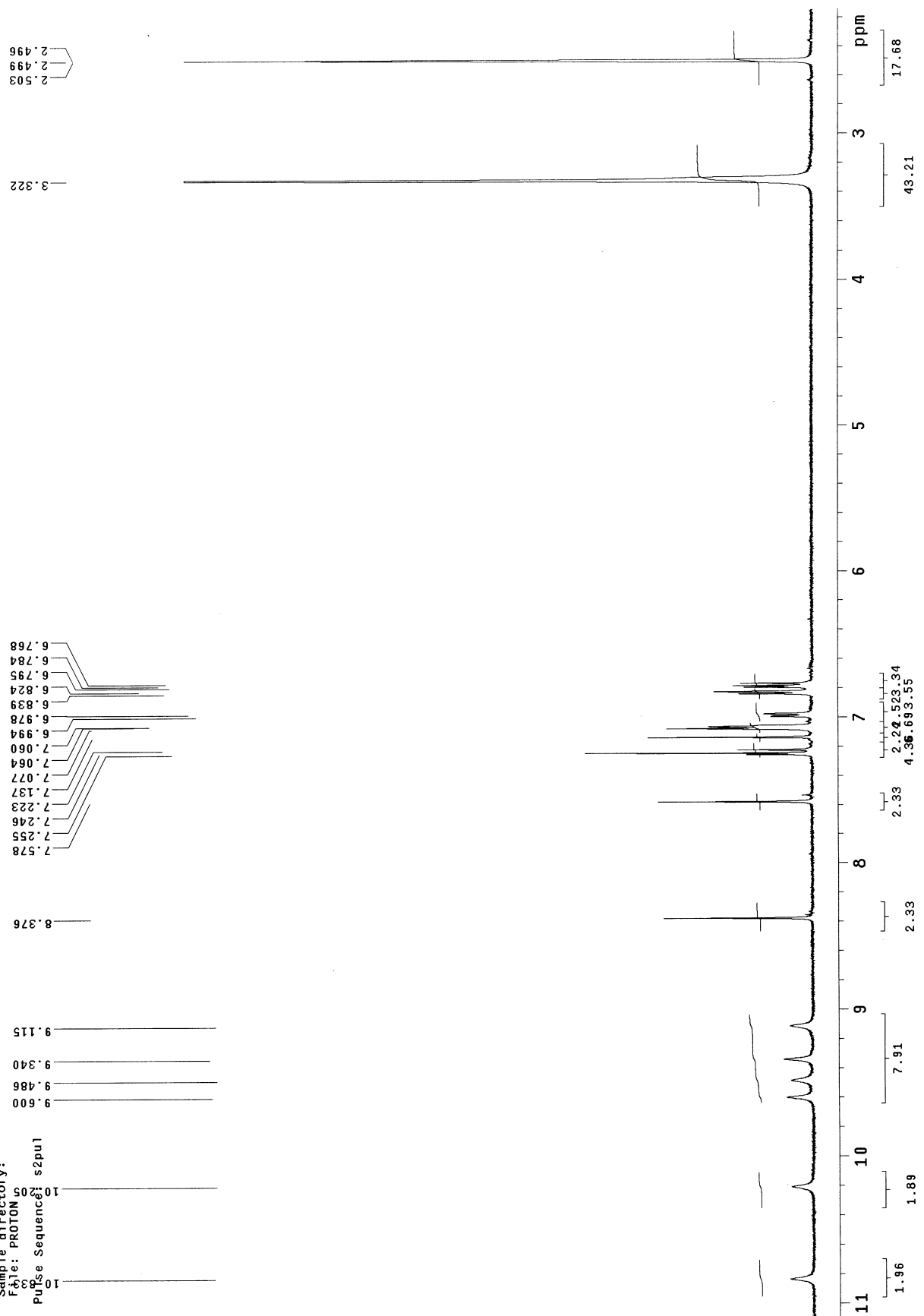
Description MeOH溶液

C155,Cycle01



¹H NMR spectrum of Phelligrudin H (1)

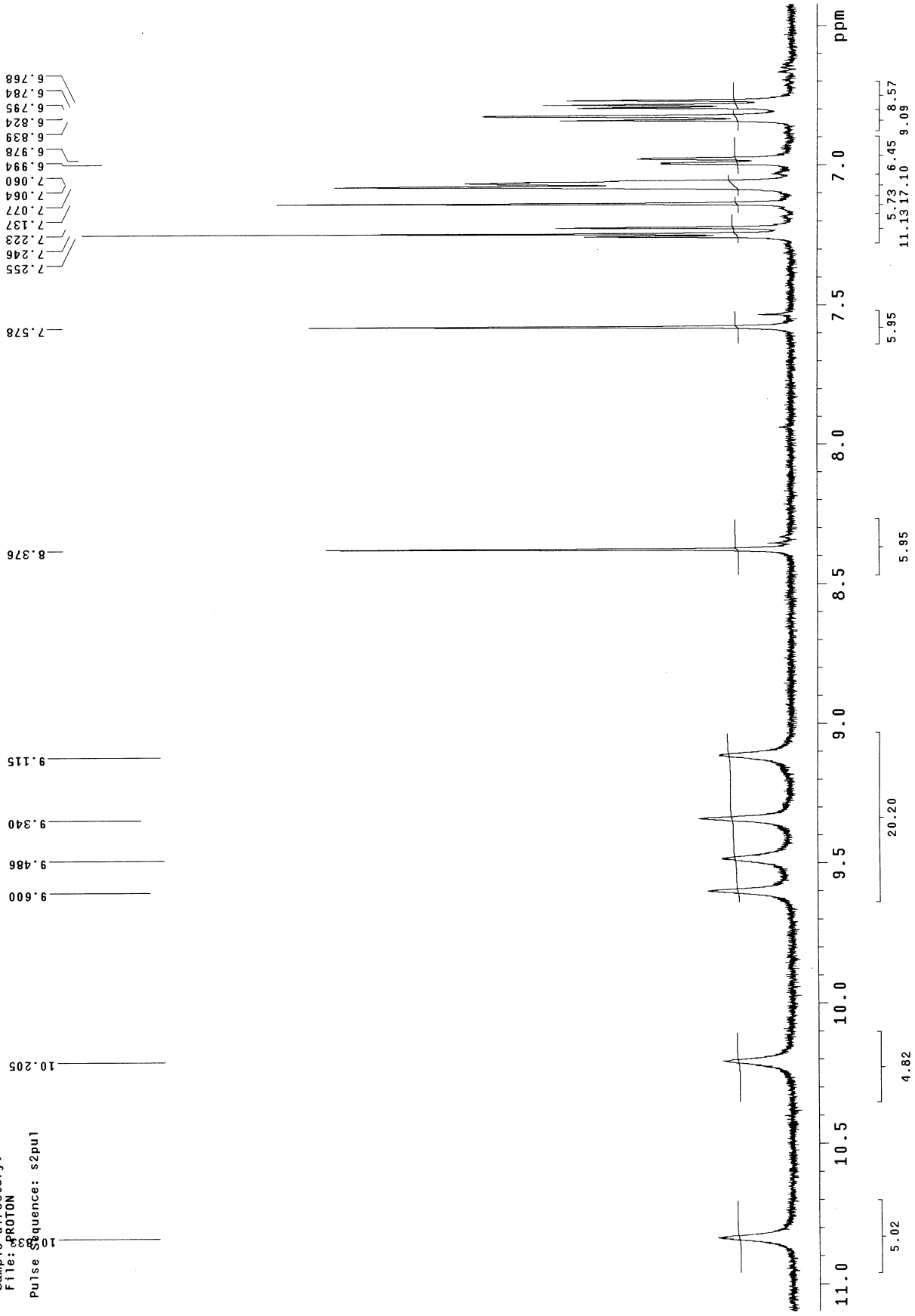
INOVA-500 1H-NMR C155 IN DMSO
 Archive directory: /export/home/vnmr1/vnmrSYS/data
 Sample directory:
 File: PROT0N
 Pulse Sequence: s2pu1



Regional enlarged ¹H NMR spectrum of Phelligridin H (1)

• INOVA-500 1H-NMR C155 IN DMSO

Archive directory: /export/home/vnmr1/vnmrSYS/data
Sample directory:
File: PROTON
Pulse Sequence: s2pu1



¹³C NMR spectrum of Phelli gridin H (1)

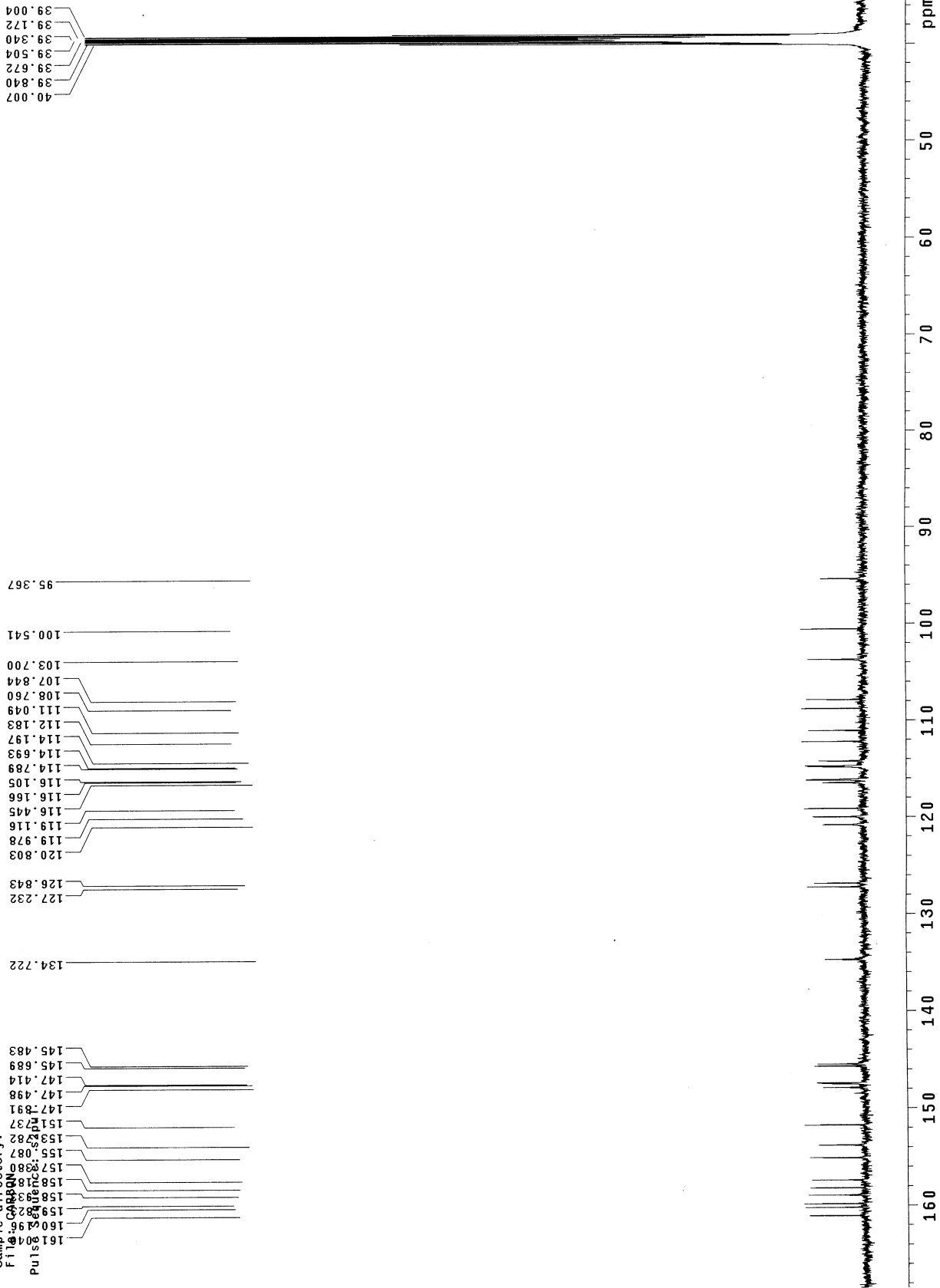
INOVA-500 13C-NMR C155 IN DMSO 06.01

Archive directory: /export/home/vnmr1/vnmrsys/data

Sample directory:

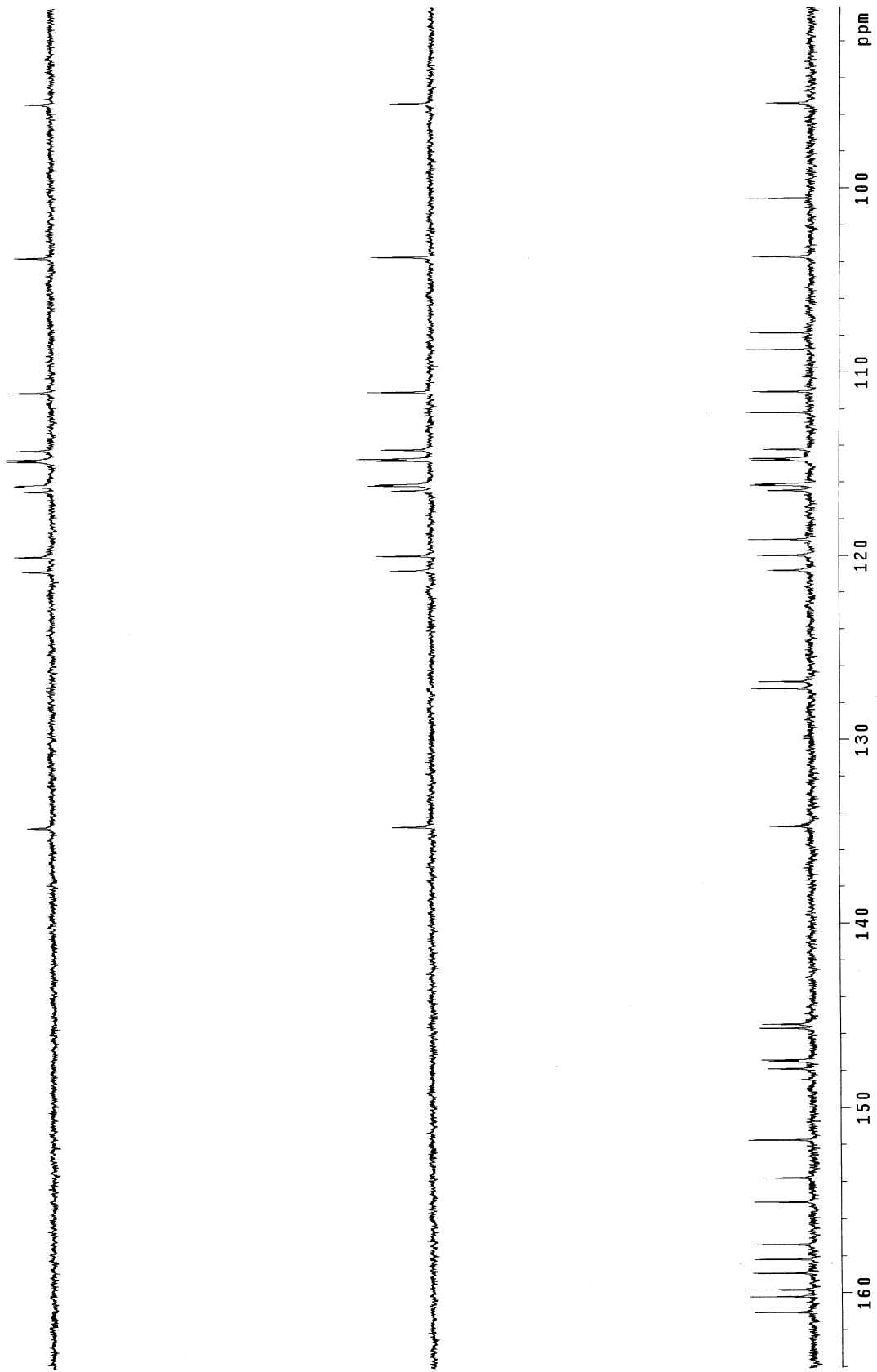
File: C:\ABON

Pul: 500



DEPT spectrum of Phelligrudin H (1)

INOVA-500 DEPT-NMR C155 IN DMSO 06.01
Archive directory: /export/home/vnmr1/vnmrSYS/data
Sample directory:
File: CARBON
Pulse Sequence: DEPT

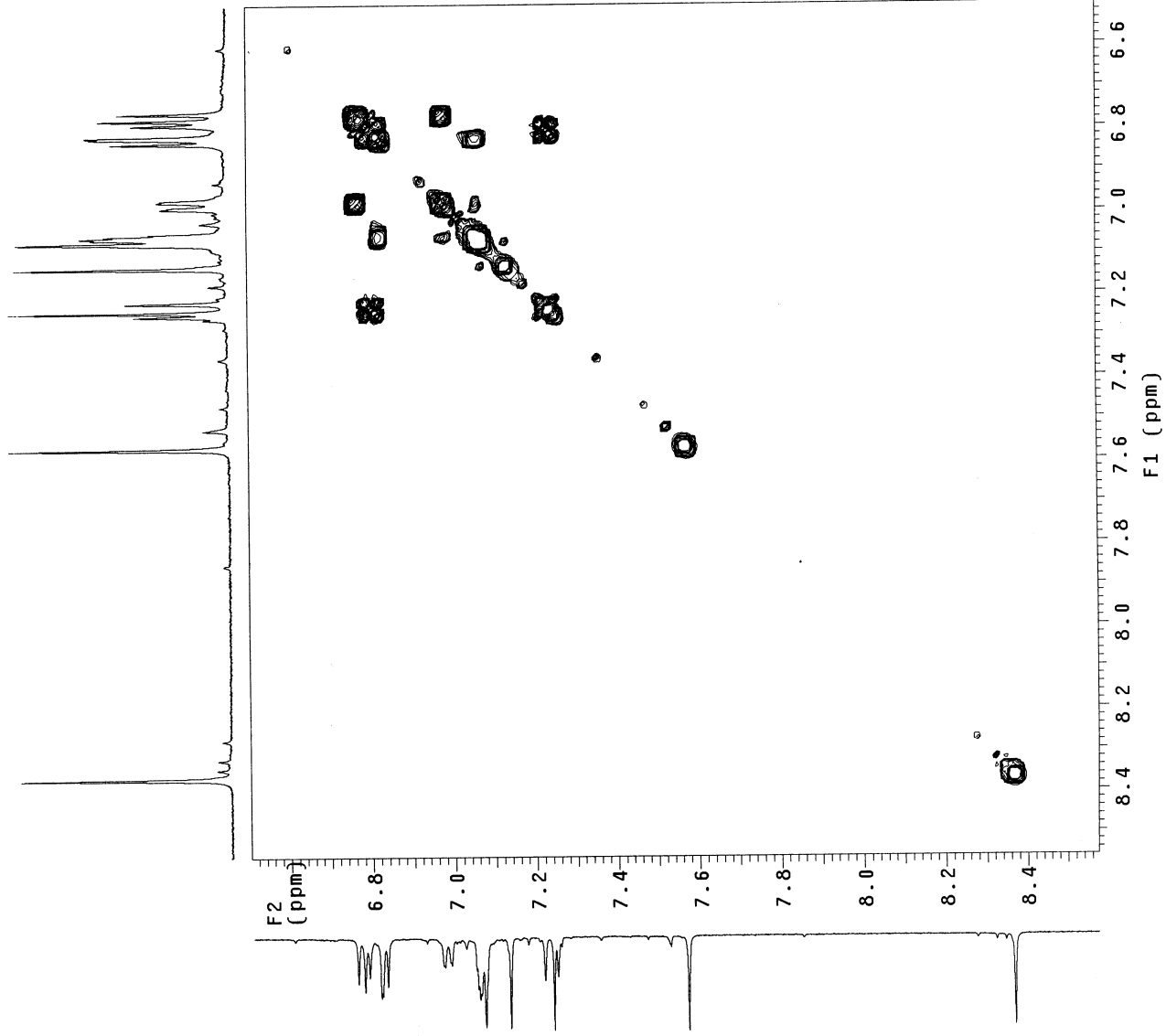


¹H-¹H COSY spectrum of Phelligrudin H (1)

INDVA-501 gCOSY B96G IN DMSO 04.04.30

File: PROTON

Solvent: DMSO
Temp. 25.0 C / 298.1 K
INDVA-500 "NMR501"
Relax. delay 1.000 sec
Acq. time 0.229 sec
Width 4473.3 Hz
2D Width 4473.3 Hz
2 repetitions
300 increments
OBSERVE F1 99.7474033 MHz
DATA PROCESSING 14
SINE DE 1.000 sec
F1 DATA PROCESSING
F1 line 01 C. 033
F1 size 2048 x 2048
Total time 13 min, 5 sec

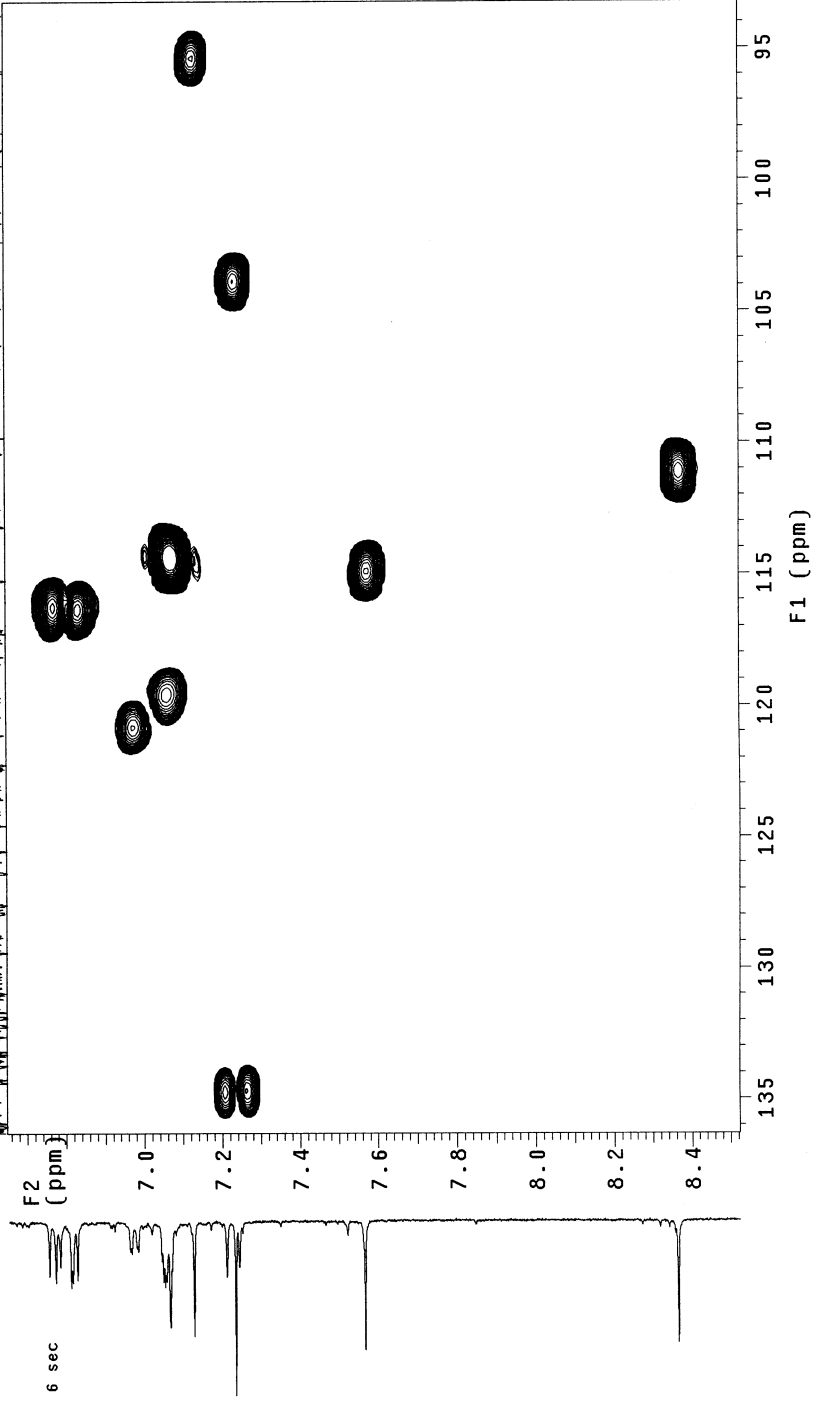


HSQC spectrum of Phelligrudin H

INOVA-501 gHSQC B96G IN DMSO 04.05.07

Solvent: DMSO
Temp. 25.0 C / 298.1 K
User: 1-14-87
File: HSQCDS0507-B96G
INOVA-500 "NMR501"

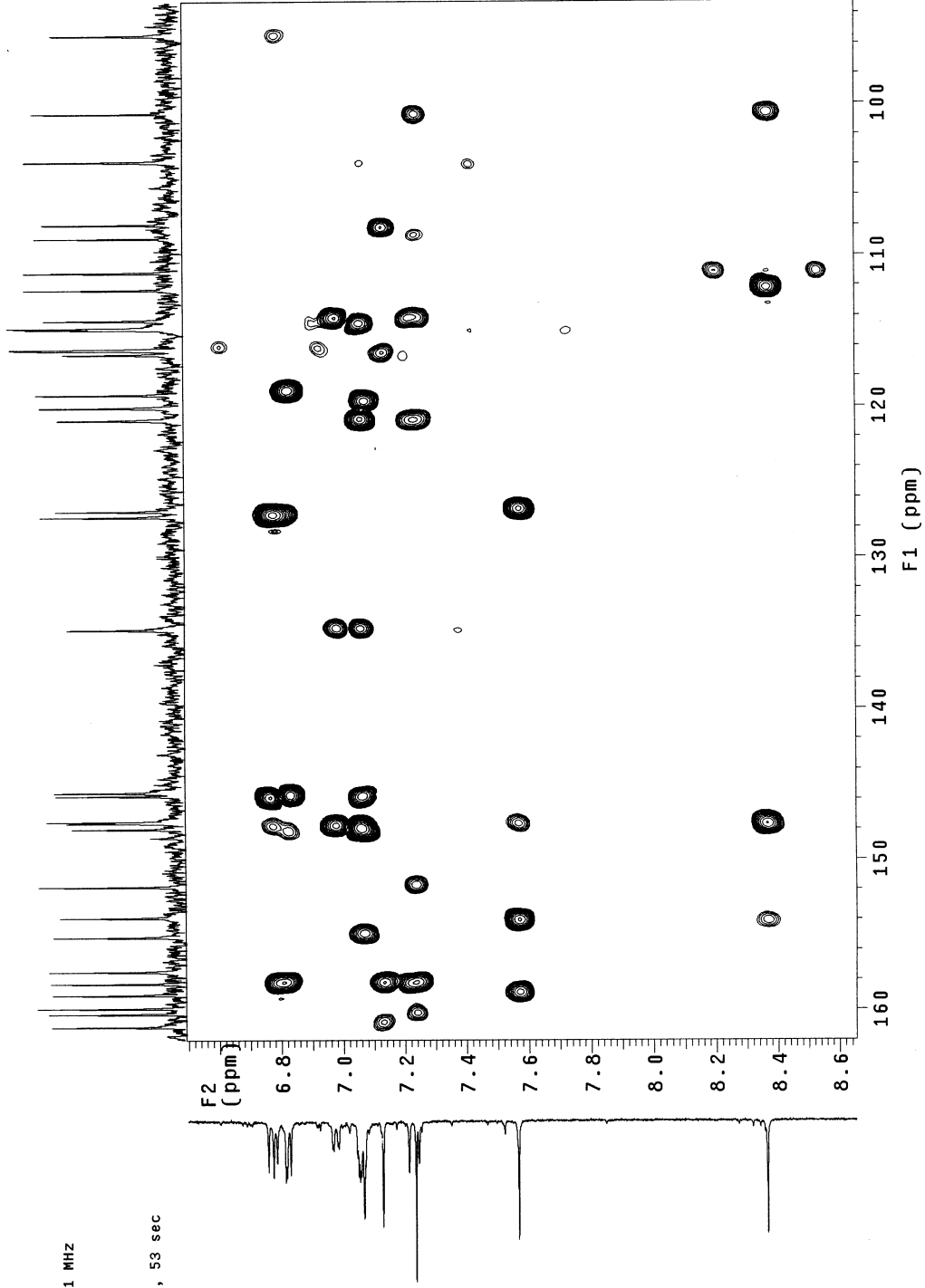
Relax. delay 1.000 sec
Acq. time 0.220 sec
Width 4650.1 Hz
2D Width 17609.5 Hz
128 repetitions
160 increments
OBSERVE H1, 499.7474031 MHz
DECOUPLE C13, 125.6742122 MHz
Power 48 dB
on during acquisition
off during delay
GARP-1 modulated
DATA PROCESSING
Sine bell 0.030 sec
F1 DATA PROCESSING
Sine bell 0.004 sec
FT size 2048 x 8192
Total time 7 hr, 22 min, 6 sec



HMBC spectrum of Phelligrudin H (1)

Solvent: DMSO
Temp. 25.0 C / 298.1 K
User: 1-14-87
File: HMBCDMSO0507-B96G
INOVA-500 "NMR501"

Relax. delay 1.000 sec
Acq. time 0.214 sec
Width 4782.4 Hz
2D Width 22624.4 Hz
192 repetitions
320 increments
OBSERVE H1 499.7474031 MHZ
DATA PROCESSING
Sine bell 0.032 sec
F1 DATA PROCESSING
Sine bell 0.007 sec
F1 size 2048 x 8192
Total time 22 hr, 14 min, 53 sec

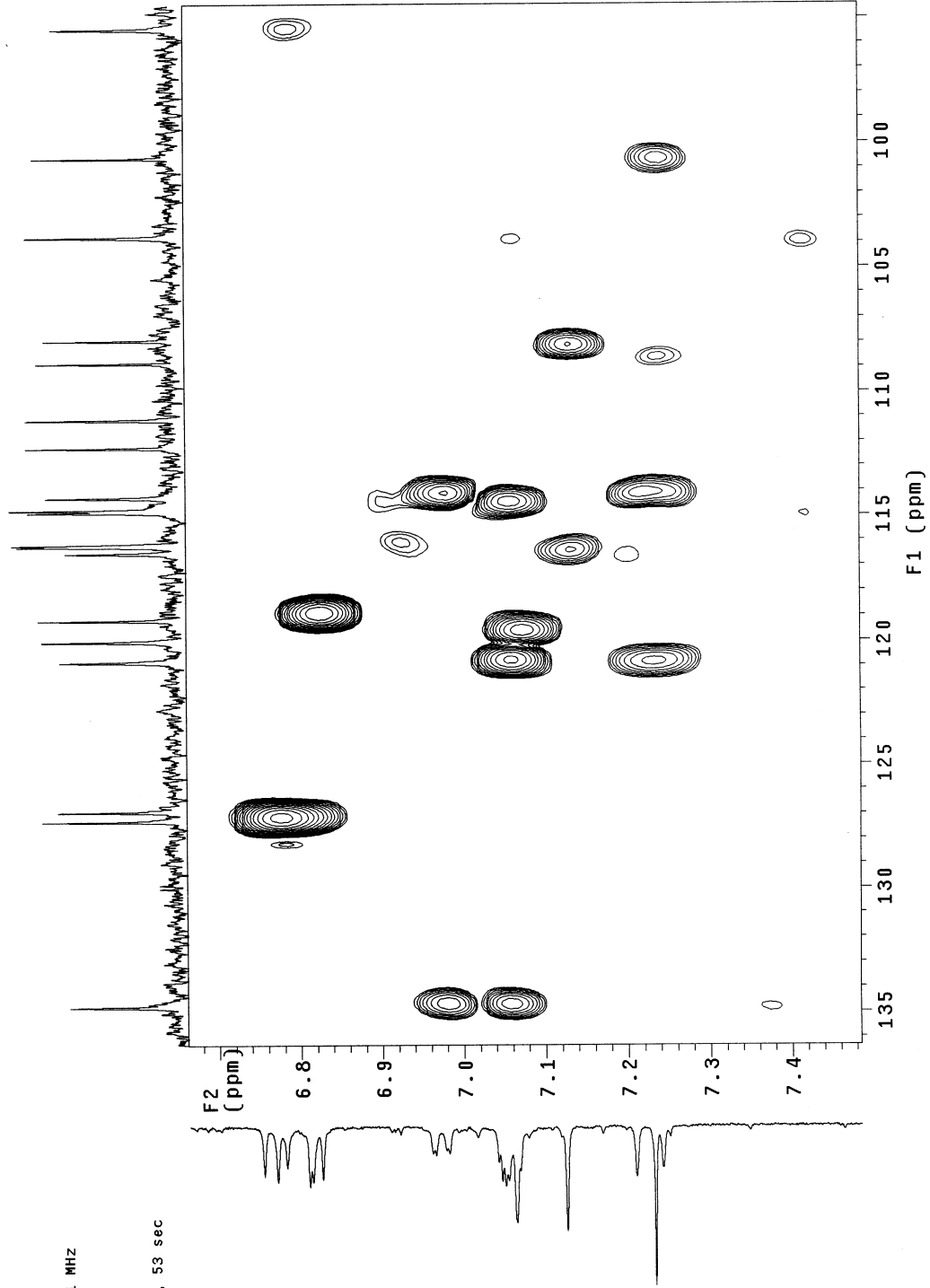


INOVA-501 gHMBC B96G IN DMSO 04.05.07

Regional enlarged HMBC spectrum of Phelligrudin H (1)

Solvent: DMSO
Temp.: 25.0 C / 298.1 K
User: i-14-87
File: HMBCDMSO0507-B96G
INOVA-500 "NMR501"

Relax. delay: 1.000 sec
Acq. time: 0.214 sec
Width: 4782.4 Hz
2D Width: 22624.4 Hz
192 repetitions
320 increments
OBSERVE: H1, 499.7474031 MHz
DATA PROCESSING
Sine bell: 0.032 sec
F1 DATA PROCESSING
Sine bell: 0.007 sec
FT size: 2048 x 8192
Total time: 22 hr, 14 min, 53 sec

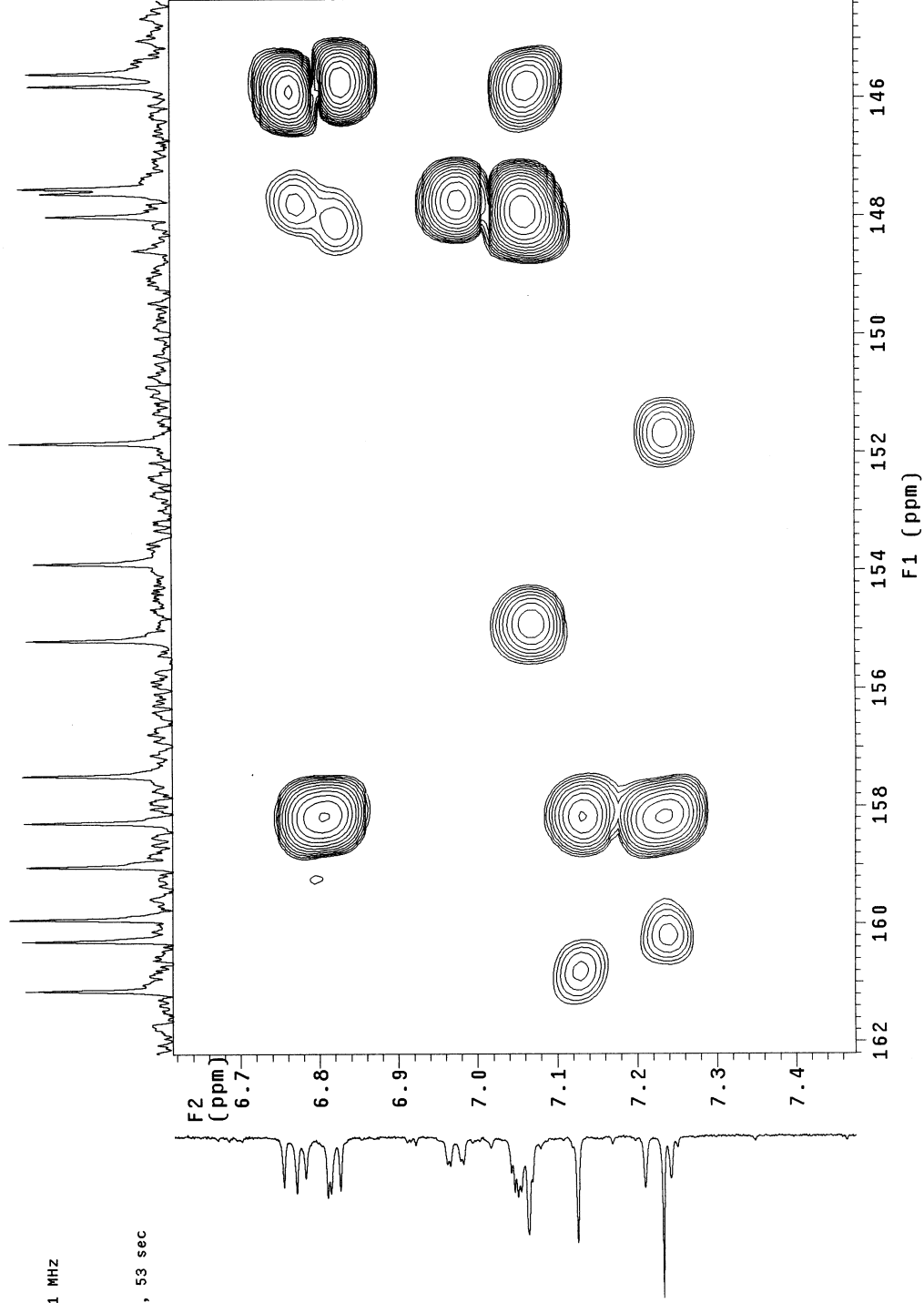


INOVA-501 gHMBC B96G IN DMSO 04.05.07

Regional enlarged HMBC spectrum of Phelligrudin H (1)

Solvent: DMSO
Temp. 25.0 C / 298.1 K
User: 1-14-87
File: HMBCDMSO0507-B96G
INOVA-500 "NMR501"

Relax. delay 1.000 sec
Acq. time 0.214 sec
Width 4782.4 Hz
2D Width 22624.4 Hz
192 repetitions
320 increments
OBSERVE H1, 499.7474031 MHZ
DATA PROCESSING
Sine bell 0.032 sec
F1 DATA PROCESSING
Sine bell 0.007 sec
FT size 2048 x 8192
Total time 22 hr, 14 min, 53 sec

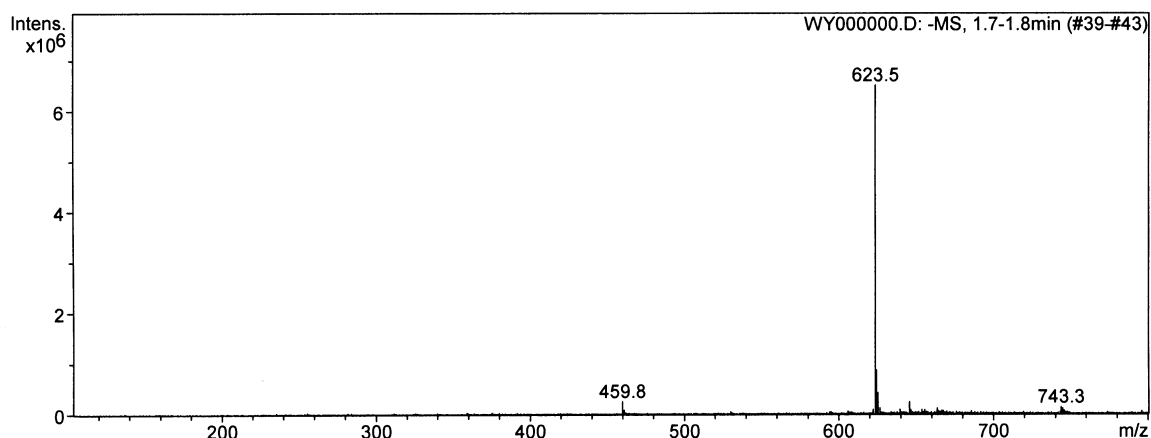


Single Mass Spectrum Deconvolution Report

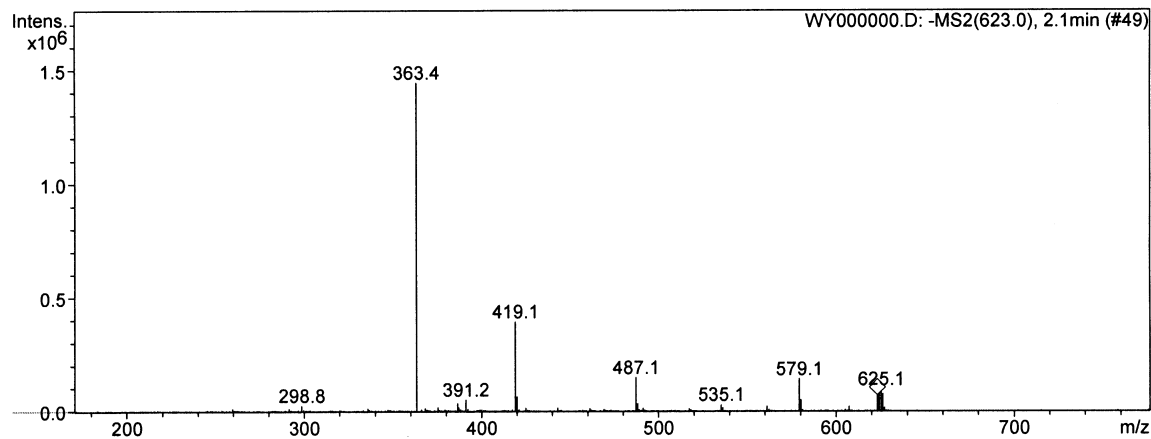
Analysis Name: WY000000.D **Instrument:** LC-MSD-Trap-SL **Print Date:** 11/28/03 17:26:48
Method: Copy of TEST.M **Operator:** Administrator **Acq. Date:** 11/28/03 17:17:24
Sample Name: E13D
Analysis Info:

Acquisition Parameter:

Mass Range Mode	Std/Normal	Trap Drive	61.0	Scan Begin	100 m/z
Ion Polarity	Positive	Skim 1	-40.0 Volt	Scan End	800 m/z
Ion Source Type	ESI	Skim 2	5.0 Volt	Averages	20 Spectra
Dry Temp (Set)	325 °C	Octopole RF Amplitude	150.0 Vpp	Max. Accu Time	300000 µs
Nebulizer (Set)	20.00 psi	Capillary Exit	-137.7 Volt	ICC Target	30000
Dry Gas (Set)	6.00 l/min			Charge Control	on

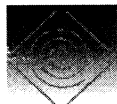


Component	Deconvoluted Mass	Molecule	Absolute Abundance	Relative Abundance
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Component	Deconvoluted Mass	Molecule	Absolute Abundance	Relative Abundance
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Instrument:



IonSpec 4.7 Tesla FTMS

Card Serial Number: I041195

Sample Serial Number: SH-W2 E130

Operator: Hua Qin Date: 2004/08/03

Operation Mode: MALDI/DHB

Elemental Composition Search Report:

Target Mass:

Target m/z = 625.0998 ± 0.003
Charge = +1

Possible Elements:

Element:	Exact Mass:	Min:	Max:
C	12.000000	0	100
H	1.007825	0	100
O	15.994915	0	30

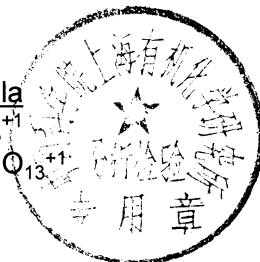
Additional Search Restrictions:

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Minimum DBE = 0

Search Results:

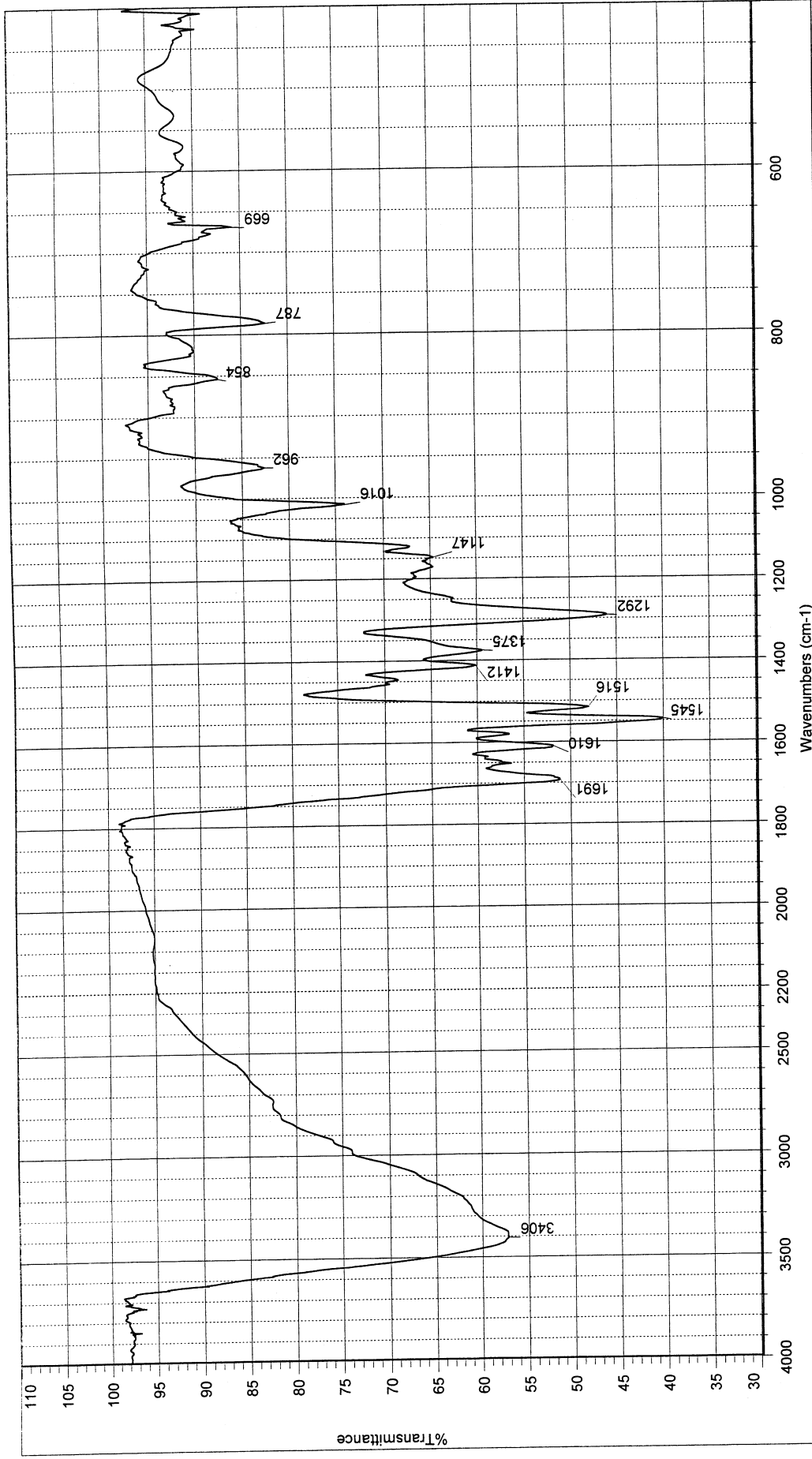
Number of Hits = 2

m/z	Delta m/z	DBE	Formula
625.10118	-0.00138	45.5	C ₅₁ H ₁₃ ⁺¹
625.09767	0.00213	23.5	C ₃₃ H ₂₁ O ₁₃ ⁺¹



HR-MS of Phelligridin I (2)

IR spectrum of Phelligrin I (2)



Date: Thu Apr 01 10:23:52 2004

Sample Name: E13D (KBr)

Scans: 64

检测单位: 国家药物及代谢产物分析研究中心

Resolution: 4.000

检测仪器: 美国尼高力公司傅立叶变换红外光谱仪: IMPACT - 400

THERMO SPECTRONIC ~ VISION32 SOFTWARE V1.25

Batch Information - scan006

Batch Type Scan Operator Name (None Entered)
Instrument ID 110514 Aborted No

Results Table - scan006

Data Mode Absorbance

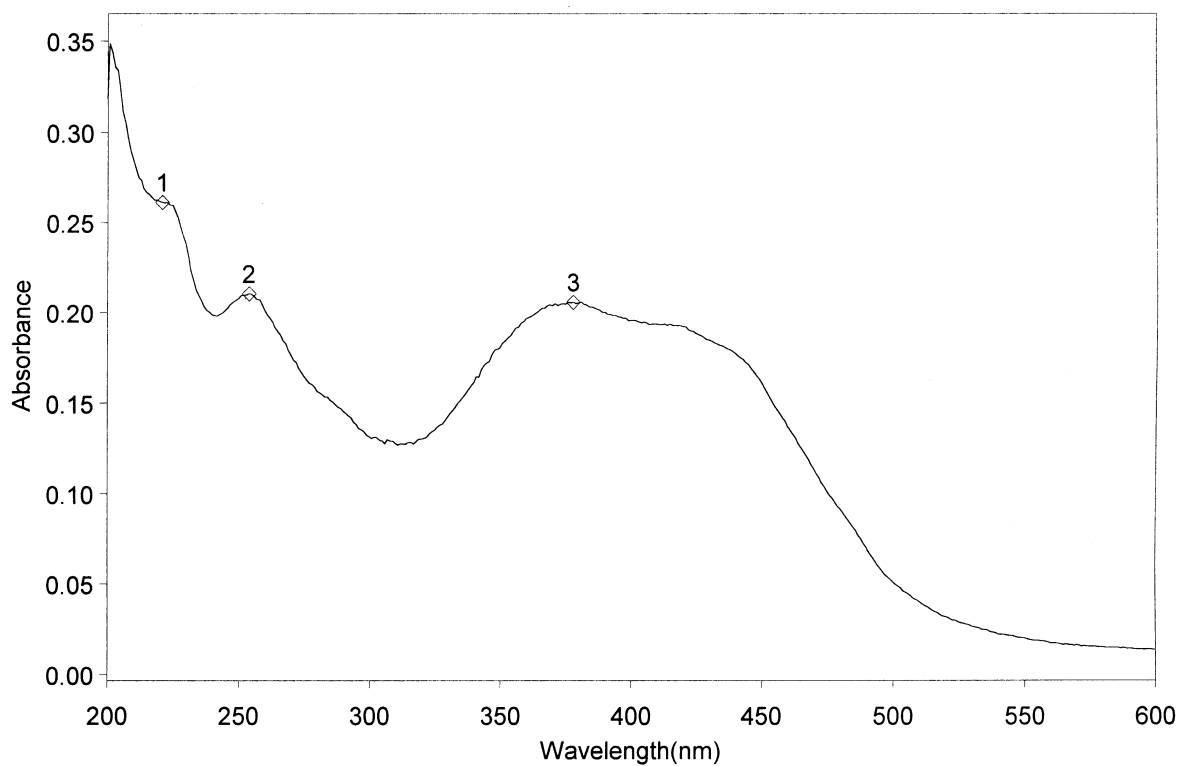
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2	Cycle01	nm	221.0	254.0	378.0
3	Manual	A	.261	.210	.206

All calculations have been performed to double precision as defined by ANSI/IEEE STD 754-1985 but have been rounded for display purposes.

E13C4D

Description 0.074mg/10mlMeOH

E13C4D,Cycle01

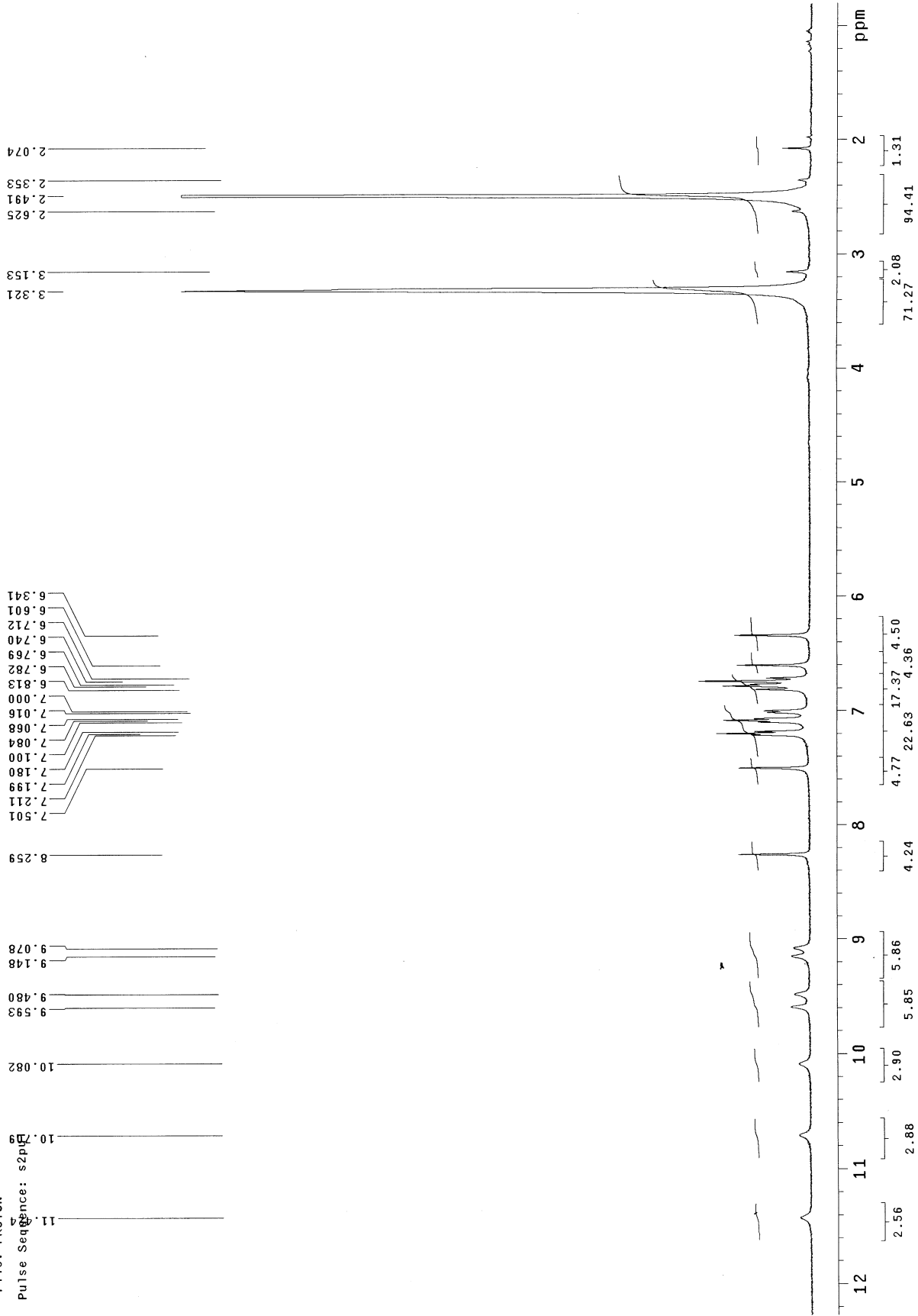


¹H NMR spectrum of Phelligridin I (2)

INOVA-500 1H-NMR E13D IN DMSO 11.17

Sample directory:
File: PROTON

Pulse Sequence: s2p(2)

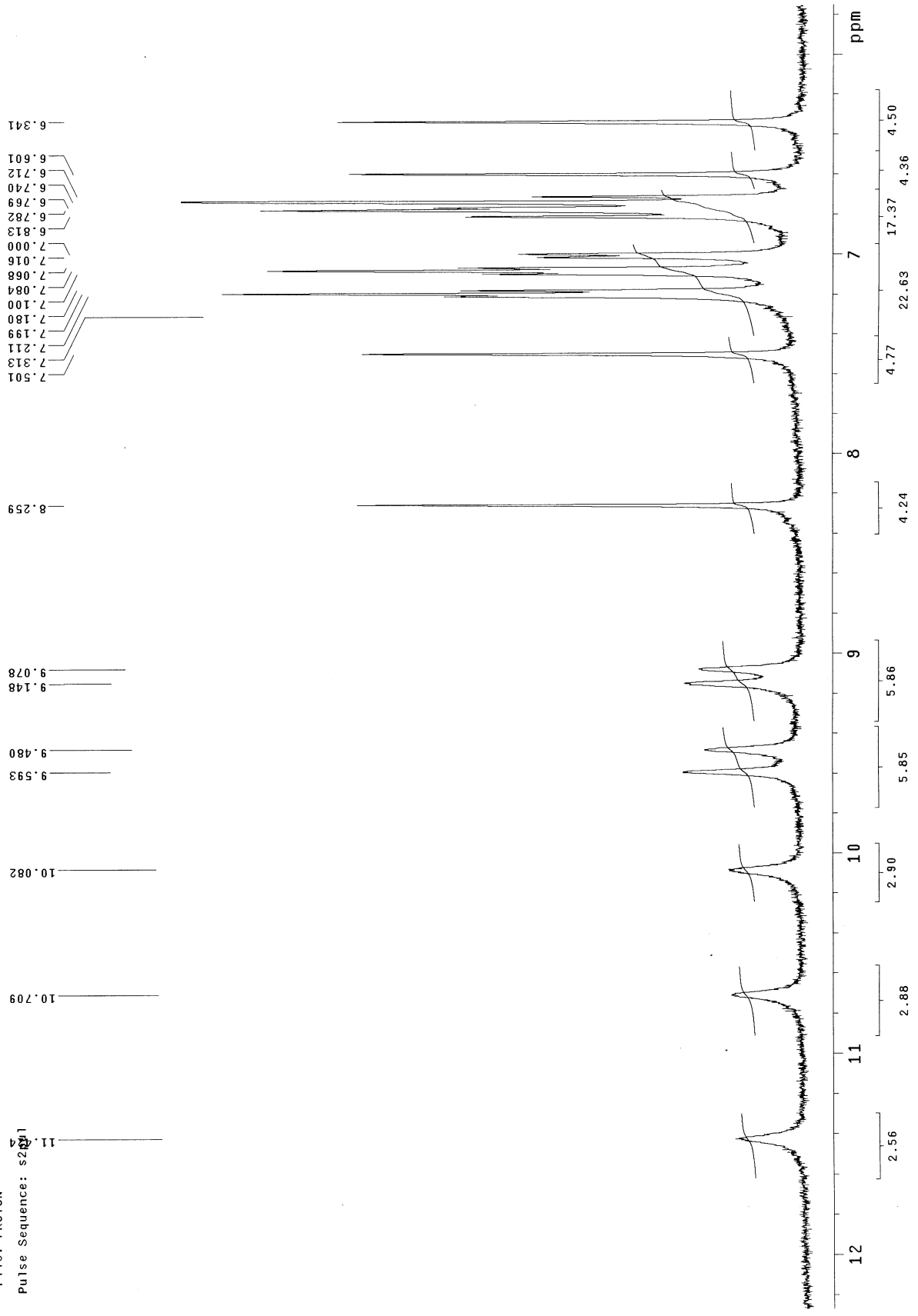


Regional enlarged ¹H NMR spectrum of Phelligridin I (2)

INOVA-500 1H-NMR E13D IN DMSO 11.17

Sample directory:
File: PROTON

Pulse Sequence: s2g1



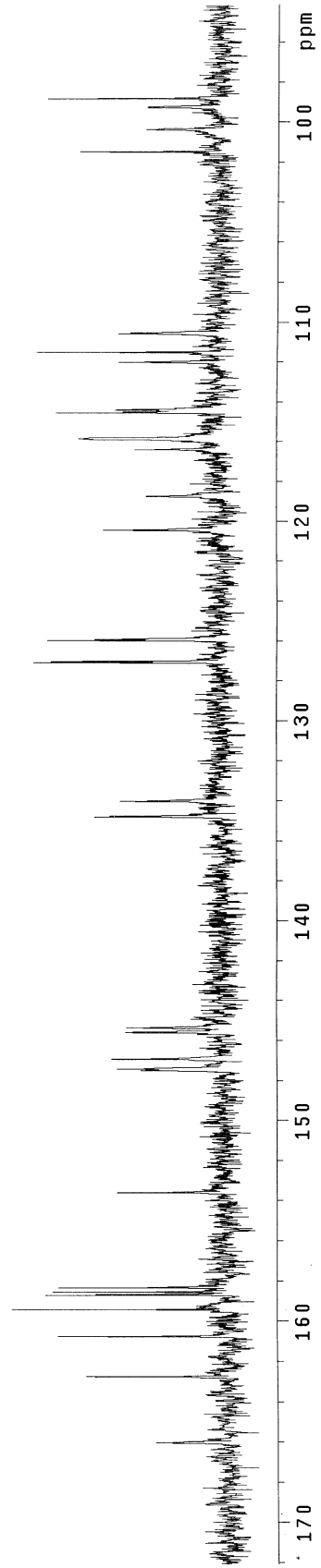
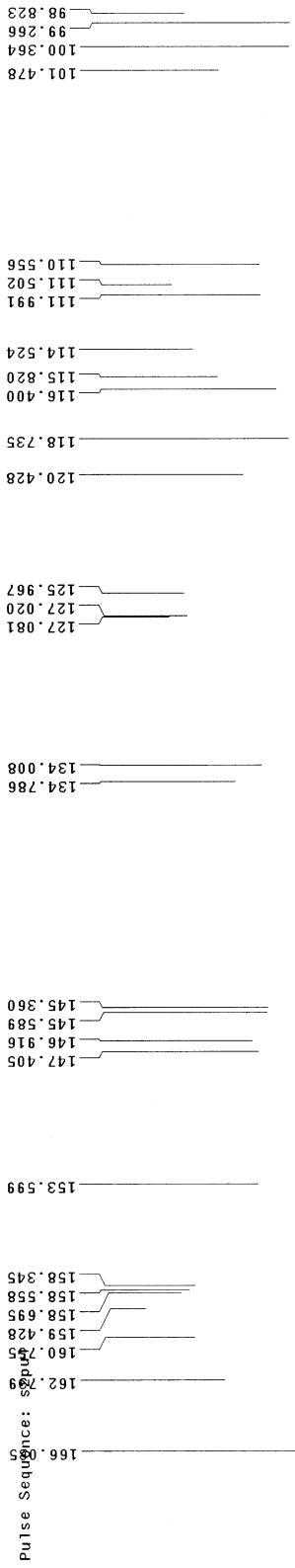
¹³C NMR spectrum of Phelligrudin I (2)

INOVA-500 13C-NMR E13D IN DMSO T=40C

Sample directory:

File: CARBON

Pulse Sequence: zgpg30



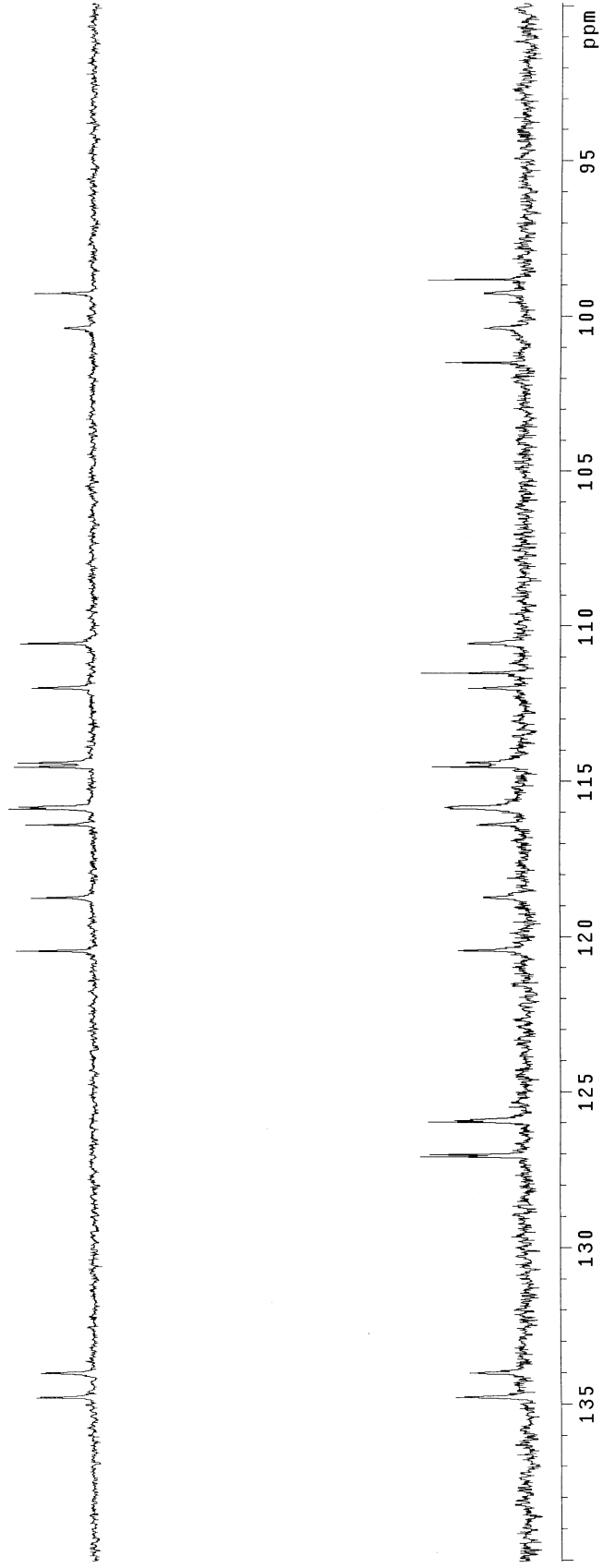
INOVA-500 DEPT-NMR E130 IN DMSO

Sample directory:

File: CARBON

Pulse Sequence: DEPT

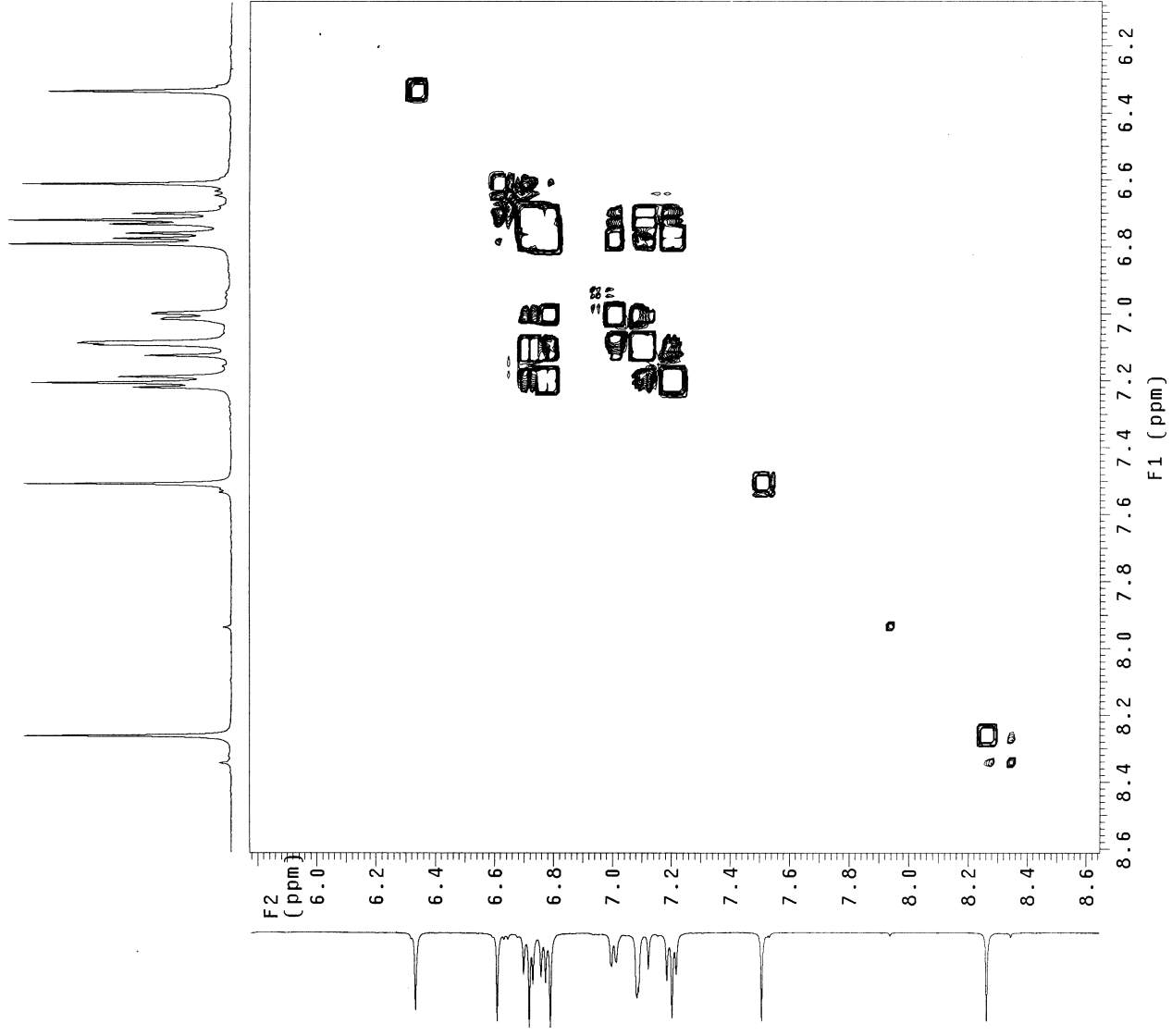
DEPT spectrum of Phelligrudin I (2)



^1H - ^1H COSY spectrum of Phelliigrudin I (2)

INOVA-501 gCOSY E13D IN DMSO 03.11.26

Solvent: DMSO
Temp. 40.0 C / 313.1 K
INOVA-500 "NMR501"
Relax. delay 1.000 sec
Acq. time 0.136 sec
Pulch 5228.8 Hz
Z0 width 5228.8 Hz
4 repetitions
256 increments
OBSERVE F1
DATA PROCESSING 499.7474034 MHZ
S_q sine bell 0.098 sec
F1 DATA PROCESSING
S_q sine bell 0.015 sec
F1 size 2048 x 2048
Total time 21 min, 22 sec



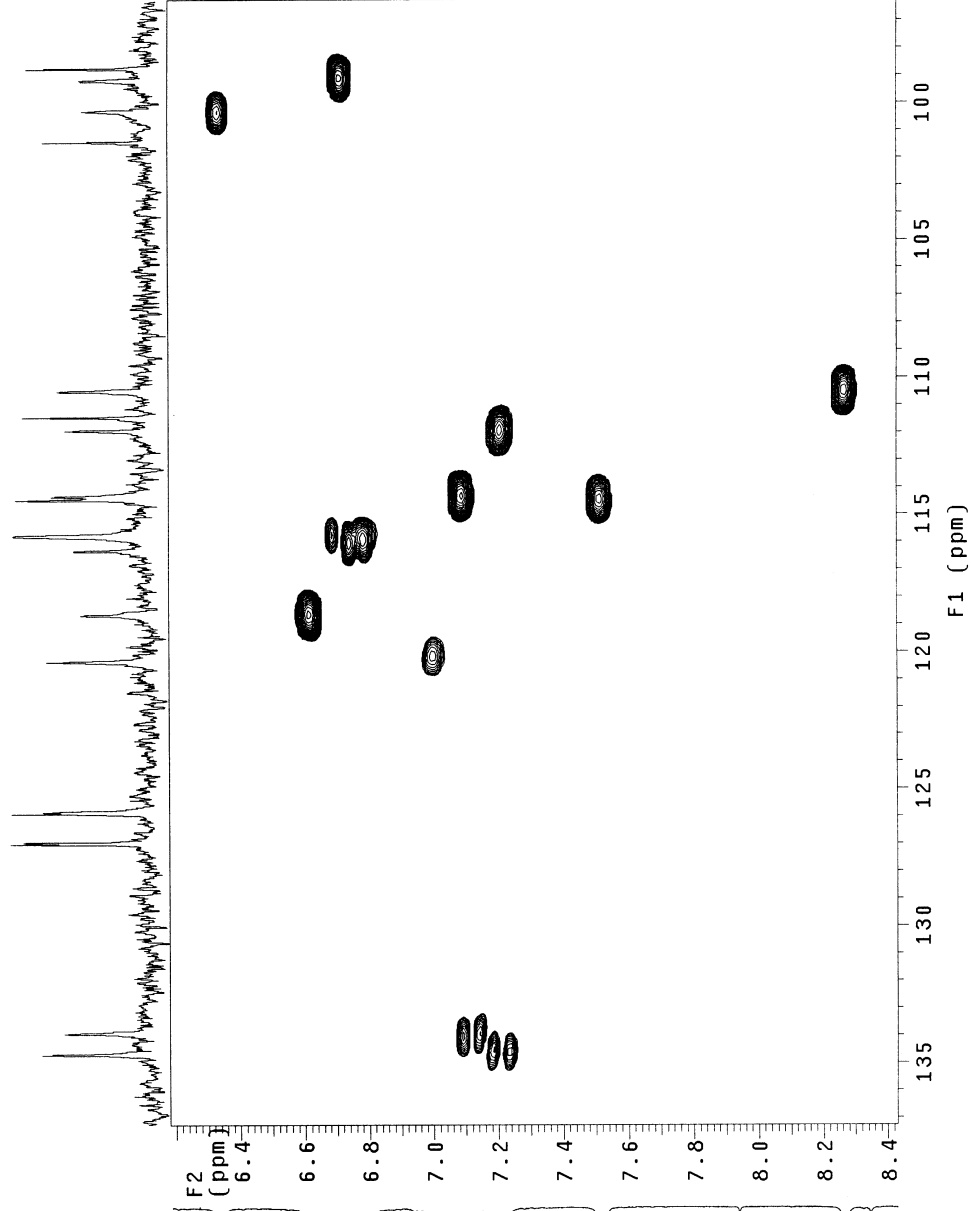
INOVA-501 gHSQC E13D IN DMSO 03.11.26 temp=40C

File: CARBON

HSQC spectrum of Phelligridin I (2)

Solvent: DMSO
Temp. 40.0 C / 313.1 K
User: 1-14-87
INOVA-500 "NMR501"

Relax. delay 1.000 sec
Acq. time 5.446 sec
Width 5046.6 Hz
2D Width 38859.0 Hz
24 repetitions
256 increments
OBSERVE H1, 499.7474034 MHz
DECOUPLE C13, 125.6748279 MHz
Power 48 dB,
on during acquisition
off during delay
GARP-1 modulated
DATA PROCESSING
Sine bell 0.031 sec
F1 DATA PROCESSING
Sine bell 0.005 sec
FT size 2048 x 4096
Total time 2 hr, 10 min, 1 sec



INOVA-501 gHMBC E130 IN DMSO 03.11.26 temp=40C

File: CARBON

HMBC spectrum of Phelligrudin I (2)

Solvent: DMSO
Temp. 40.0 C / 313.1 K
User: j1487
INOVA-500 "NMR501"

Relax. delay 1.000 sec
Acq. time 0.180 sec
Width 5683.4 Hz
2D Width 18859.0 Hz
32 repetitions
256 increments

OBSERVE H1, 499.7474034 MHz

DATA PROCESSING

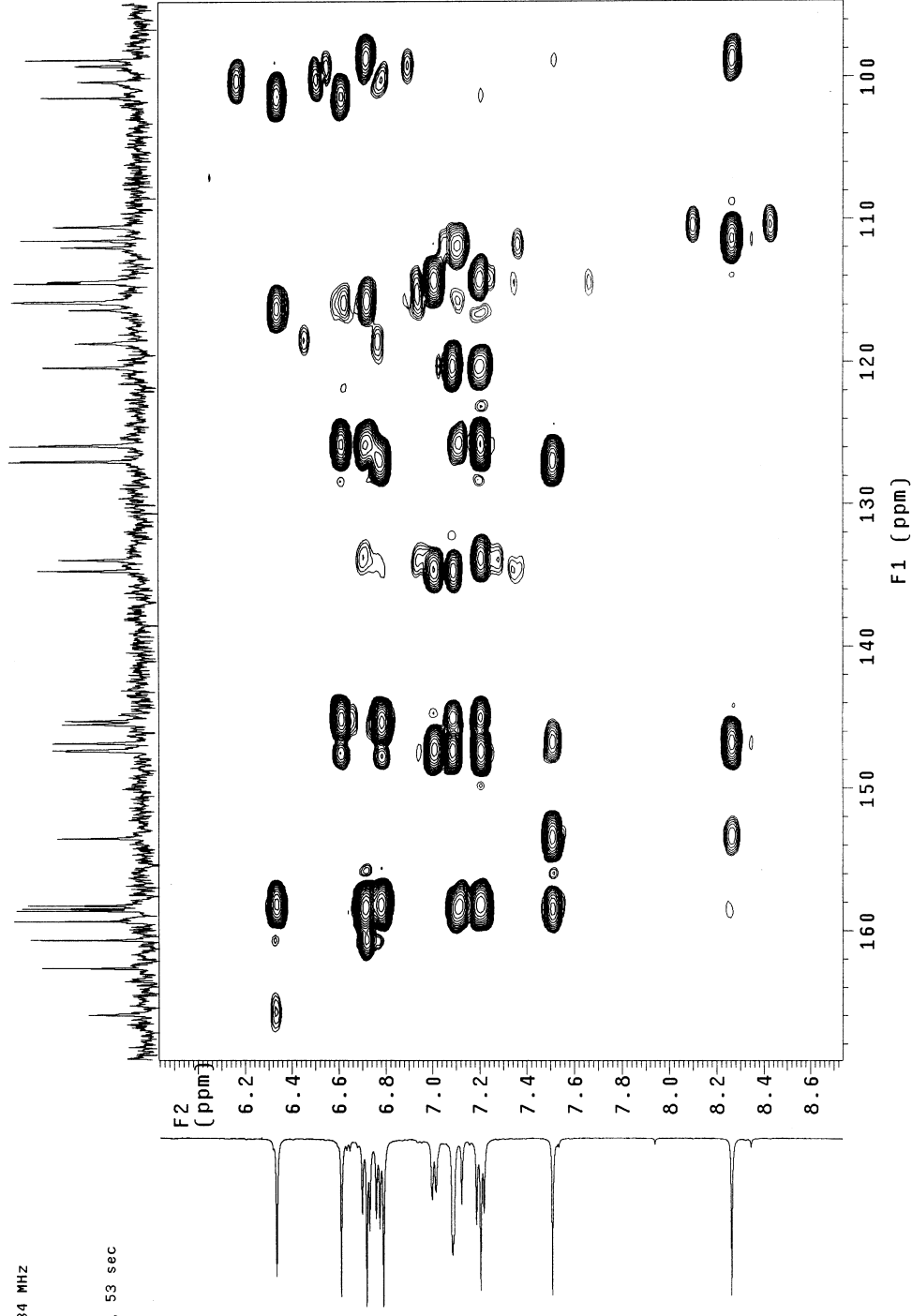
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F1 DATA PROCESSING

Sine bell 0.003 sec

FT size 2048 x 4096

Total time 2 hr, 53 min, 53 sec



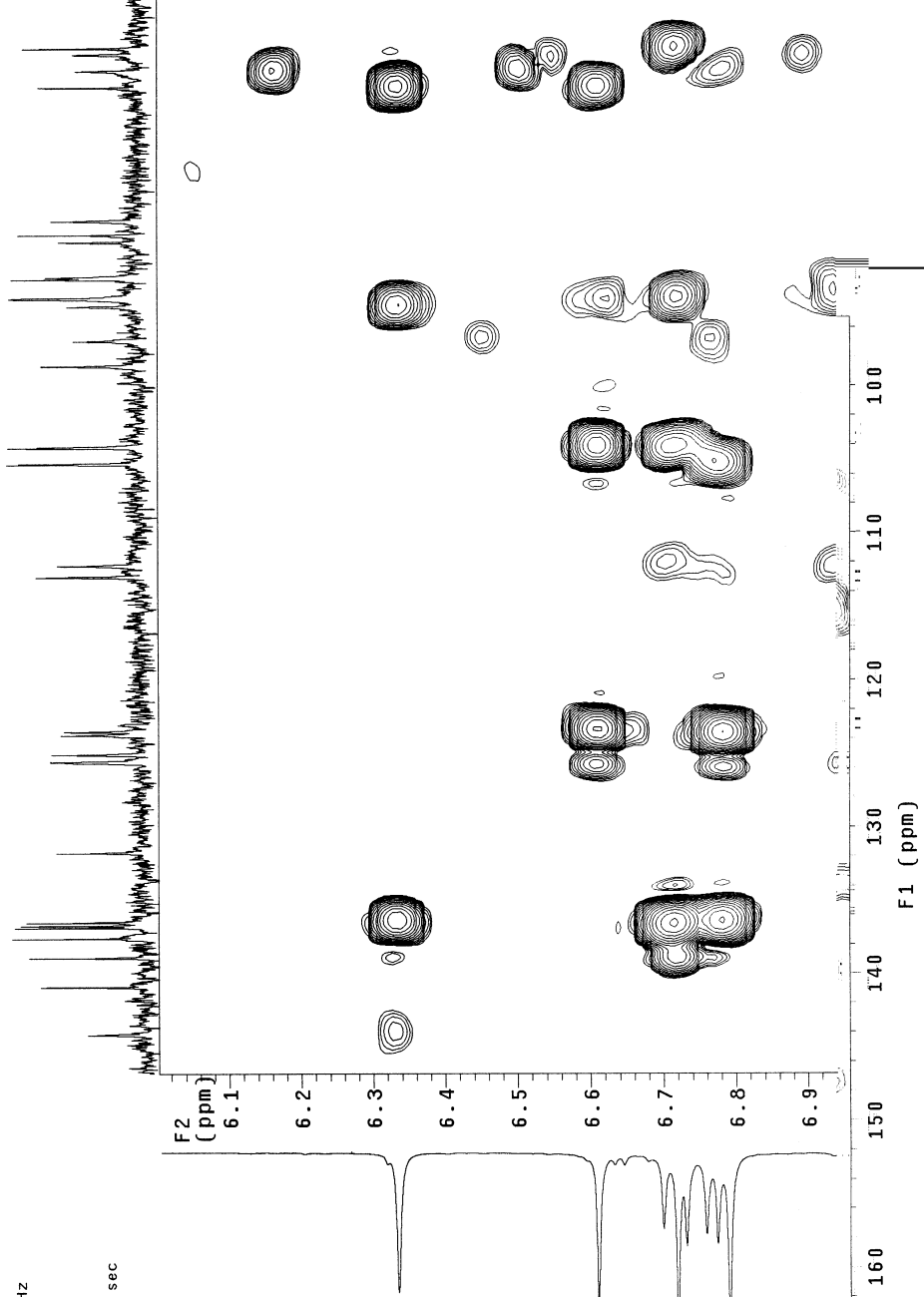
INOVA-501 gHMBC E13D IN DMSO 03.11.20 temp=40C

File: CARBON

Regional enlarged HMBC spectrum of Phelligrudin I (2)

Solvent: DMSO
Temp. 40.0 C / 313.1 K
User: 1-14-87
INOVA-500 "NMR501"

Relax. delay 1.000 sec
Acq. time 0.180 sec
Width 5683.4 Hz
2D Width 16853.0 Hz
32 repetitions
26 increments
OBSERVE F1 499.7474034 MHz
DATA PROCESSING
Sine bell 0.037 sec
F1 DATA PROCESSING
Sine bell 0.003 sec
FT size 2048 x 4096
Total time 2 hr, 53 min, 53 sec



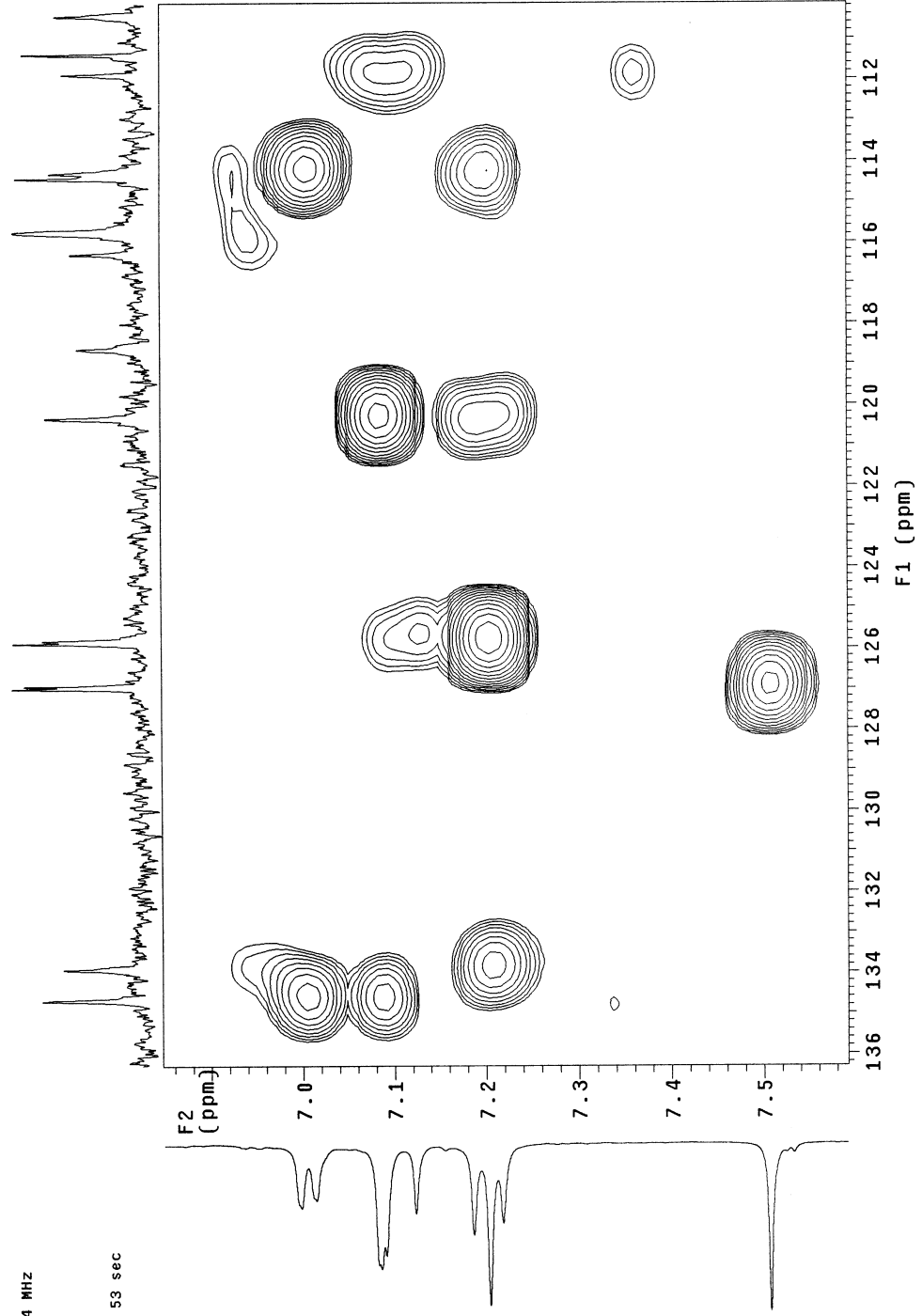
INOVA-501 gHMBC E13D IN DMSO 03.11.26 temp=40C

File: CARBON

Regional enlarged HMBC spectrum of Phelligrudin I (2)

Solvent: DMSO
Temp. 40.0 C / 313.1 K
User: 1-14-87
INOVA-500 "NMR501"

Relax: delay 1.000 sec
Acq: time 150 sec
Width 5683.4 Hz
2D width 18859.0 Hz
32 repetitions
256 increments
OBSERVE H1 499.7474034 MHz
DATA PROCESSING
Sine bell 0.031 sec
F1 DATA PROCESSING
Sine bell 0.004 sec
F1 size 2048 x 4096
Total time 2 hr, 53 min, 53 sec



INOVA-501 gHMBC E13D IN DMSO 03.11.26 temp=40C

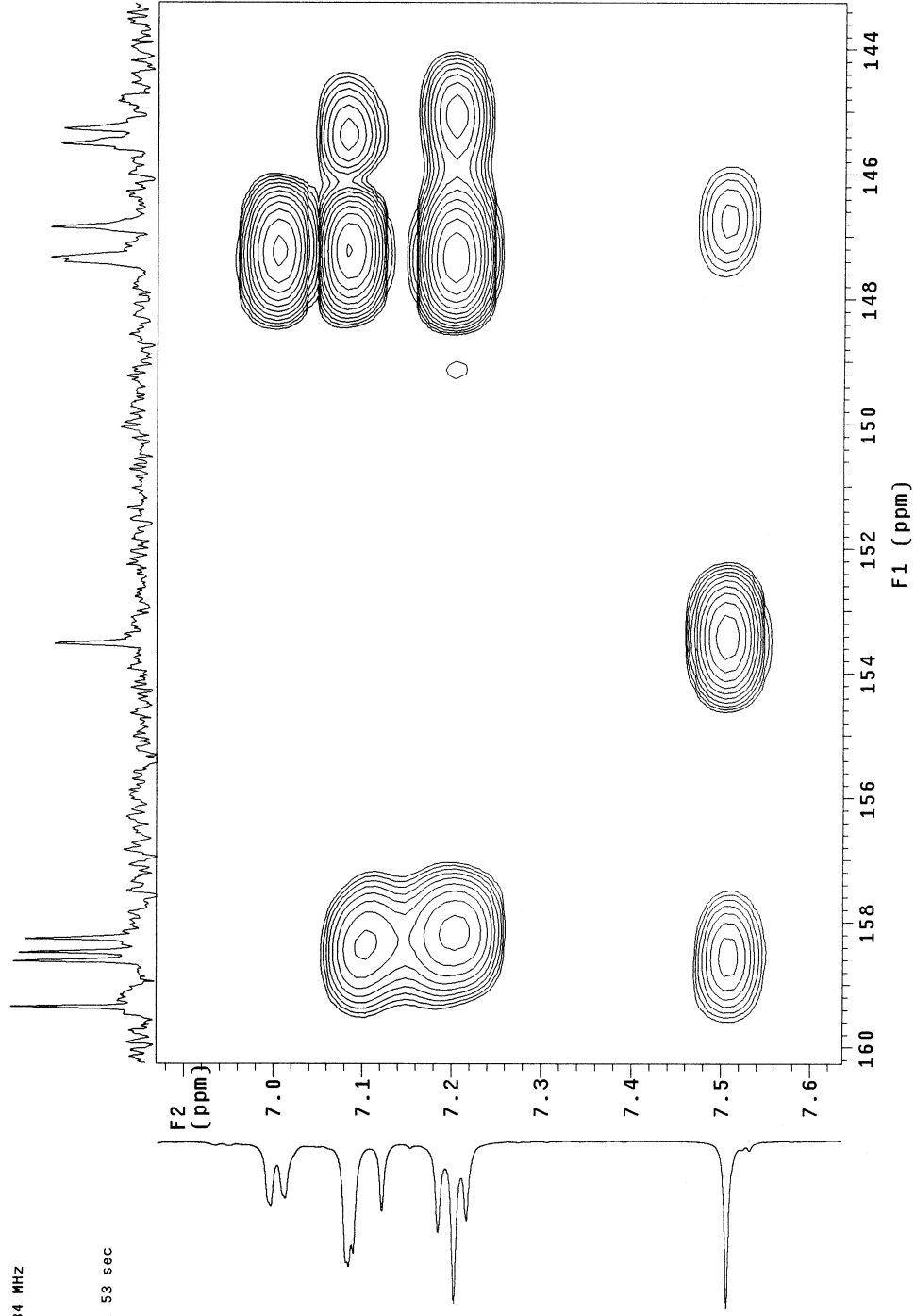
File: CARBON

Regional enlarged HMBC spectrum of Phelligrudin I (2)

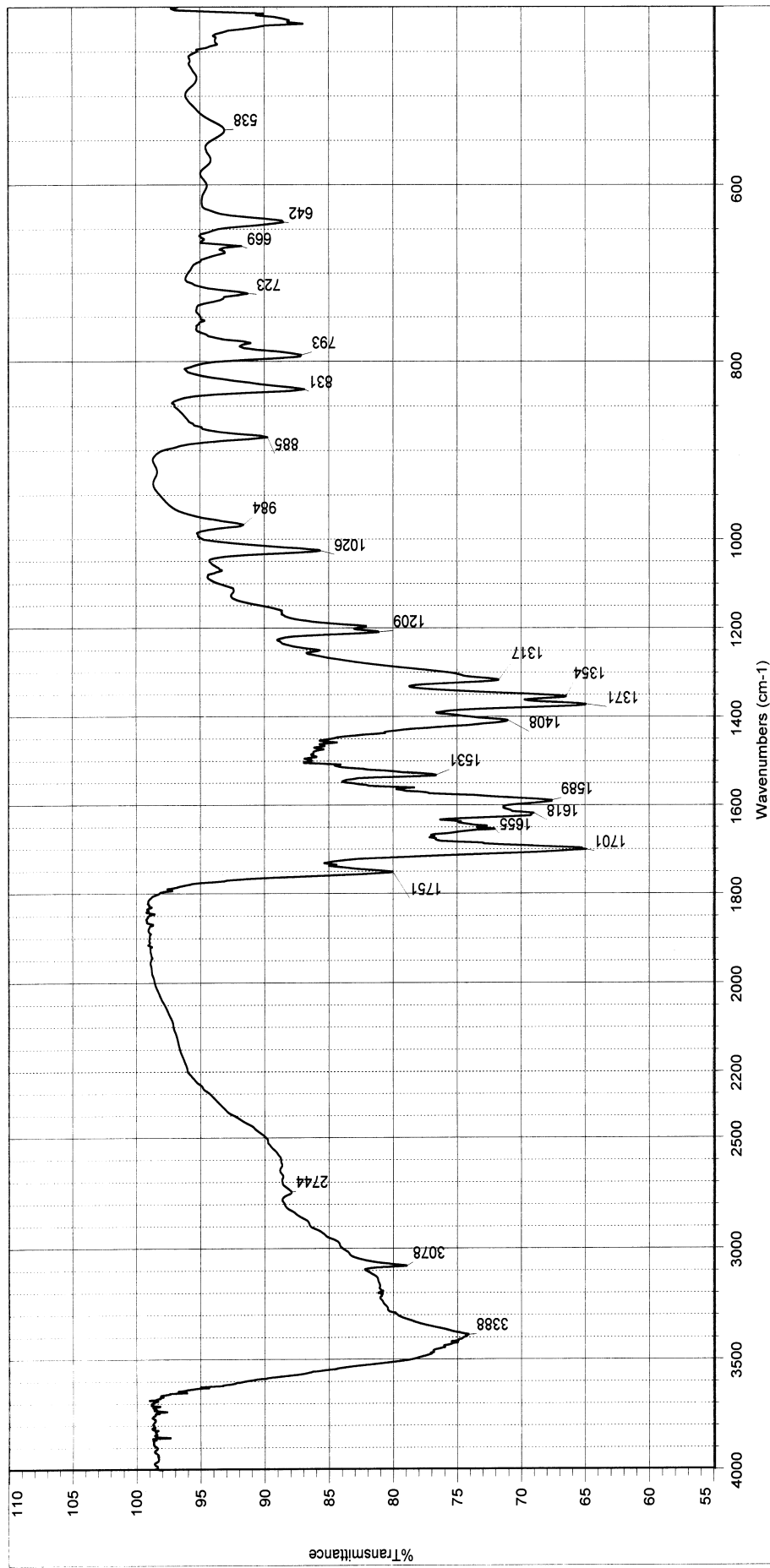
Solvent: DMSO
Temp. 40.0 C / 313.1 K
User: 1-14-87
INOVA-500 "NNR501"

Relax. delay 1.000 sec
Acq. time 0.180 sec
Width 5683.4 Hz
20 Width 18959.0 Hz
32 repetitions
256 increments

OBSERVE H1 499.7474034 MHZ
DATA PROCESSING
Sine bell 0.031 sec
F1 DATA PROCESSING
Sine bell 0.004 sec
FT size 2048 x 4096
Total time 2 hr, 53 min, 53 sec



IR spectrum of Phelliigrudin J (3)



Date: Wed Dec 28 09:12:01 2005

Scans: 64

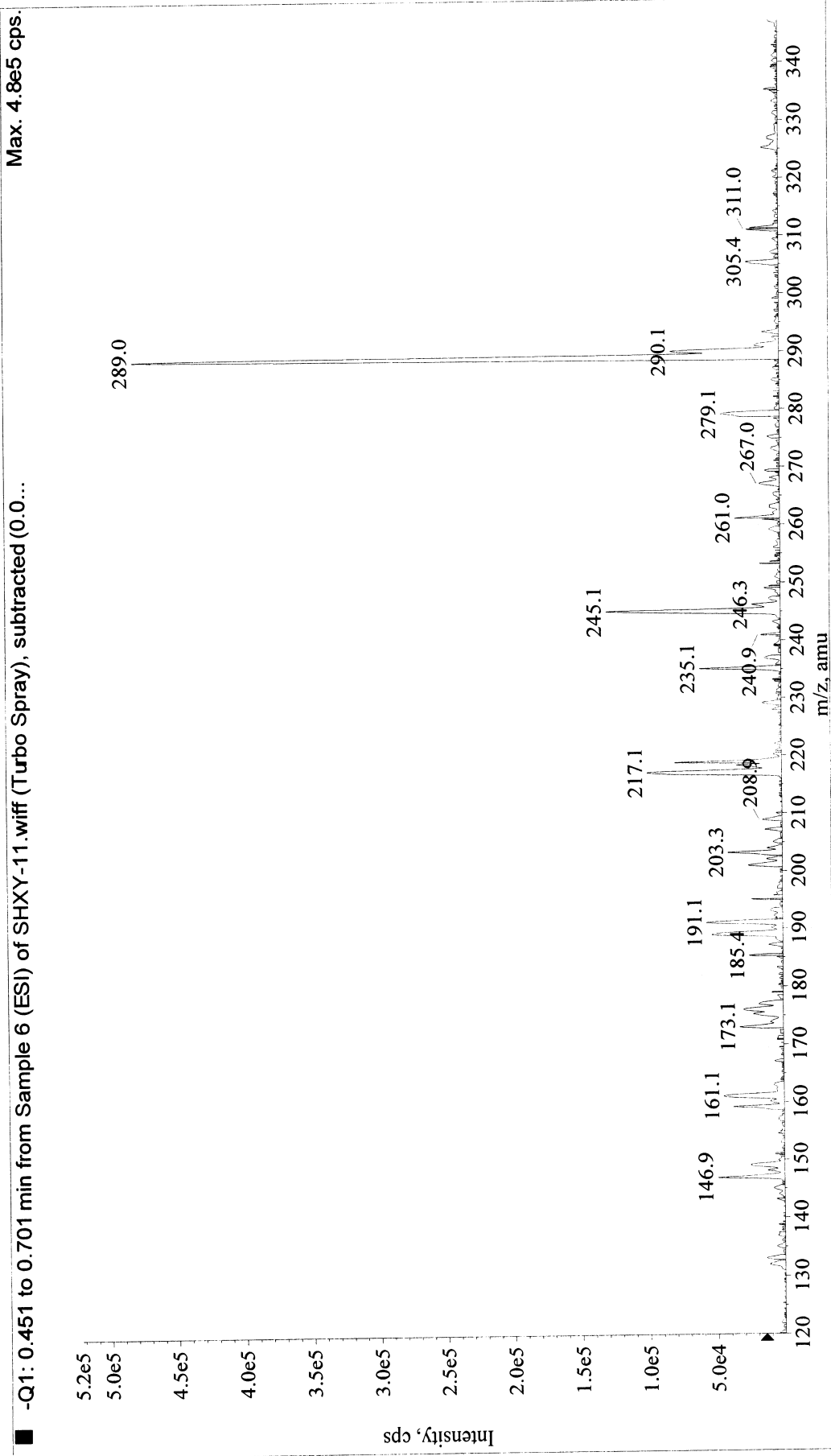
Resolution: 4.000

Sample Name: shxy - 11 (KBr)

检测单位: 国家药物及代谢产物分析研究中心

美国尼高力公司(Nicolet Co.)傅立叶变换红外光谱仪: IMPACT - 400

(-)-ESIMS of PheIigrudin J (3)



(-)-HRESIMS of Phelligridin J (3)

Data:shxy-11
 Sample Name:
 Description:
 Ionization Mode:ESI-
 History:Determine m/z[Peak Detect[Centroid,50,Area],Correct Base[]];Correct Base[5.0%];Average[MS[1] 1.4..2.7]

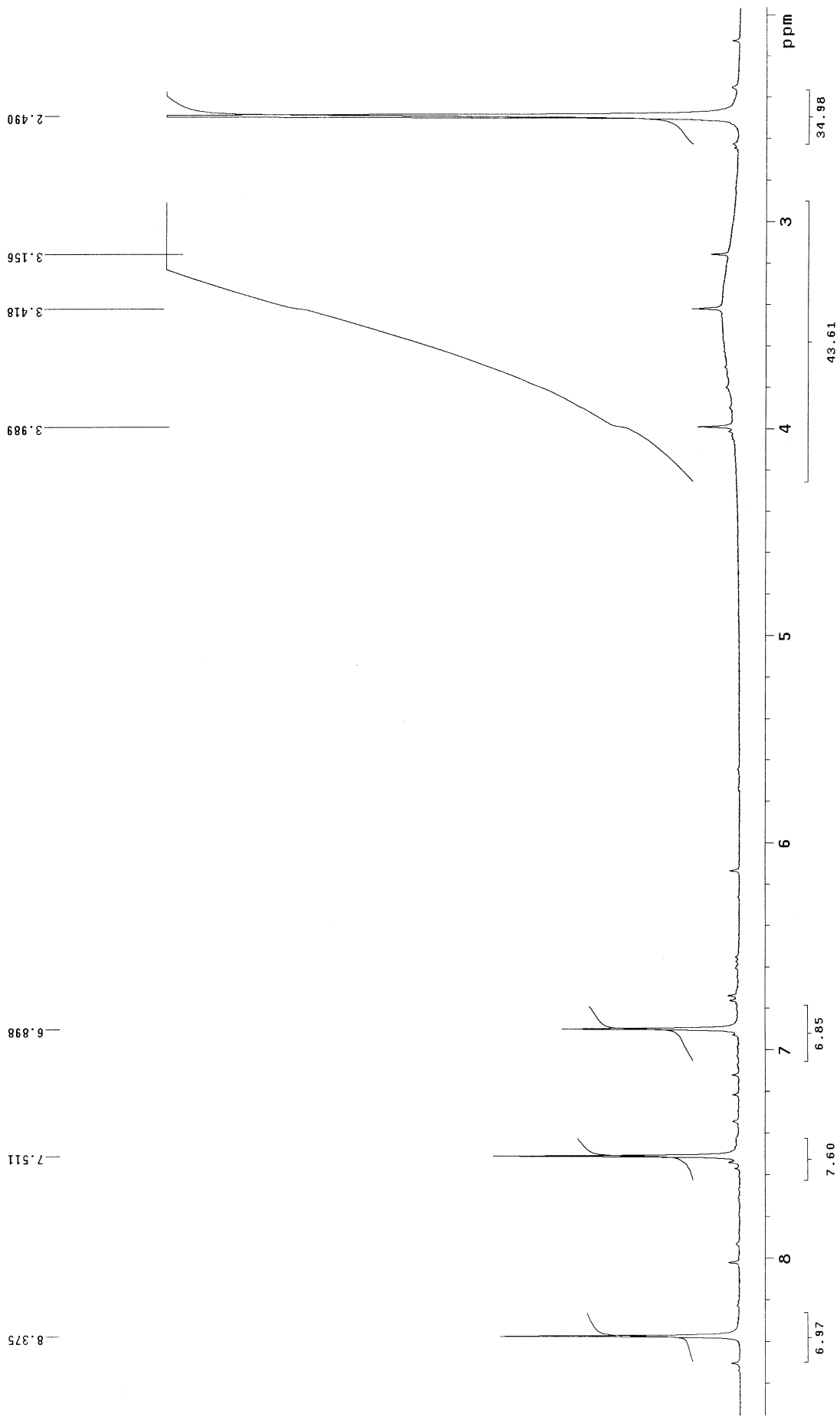
Acquired:12/22/2005 2:01:38 PM
 Operator:Accutof
 Mass Calibration data:TFANa_ESI-_1000
 Created:12/22/2005 2:13:50 PM
 Created by:Accutof

Charge number:1
 Element:¹²C:0 .. 100, ¹H:0 .. 200, ¹⁶O:0 .. 10
 Tolerance:5.00(mmu)
 Unsaturatation Number:-1.5 .. 20.0 (Fraction:Both)

Mass	Calc. Mass	Mass Difference (mmu)	Mass Difference (ppm)	¹² C	¹ H	¹⁶ O	Unsaturatation Number
289.00046	288.99844	2.02	6.98	13	5	8	11.5

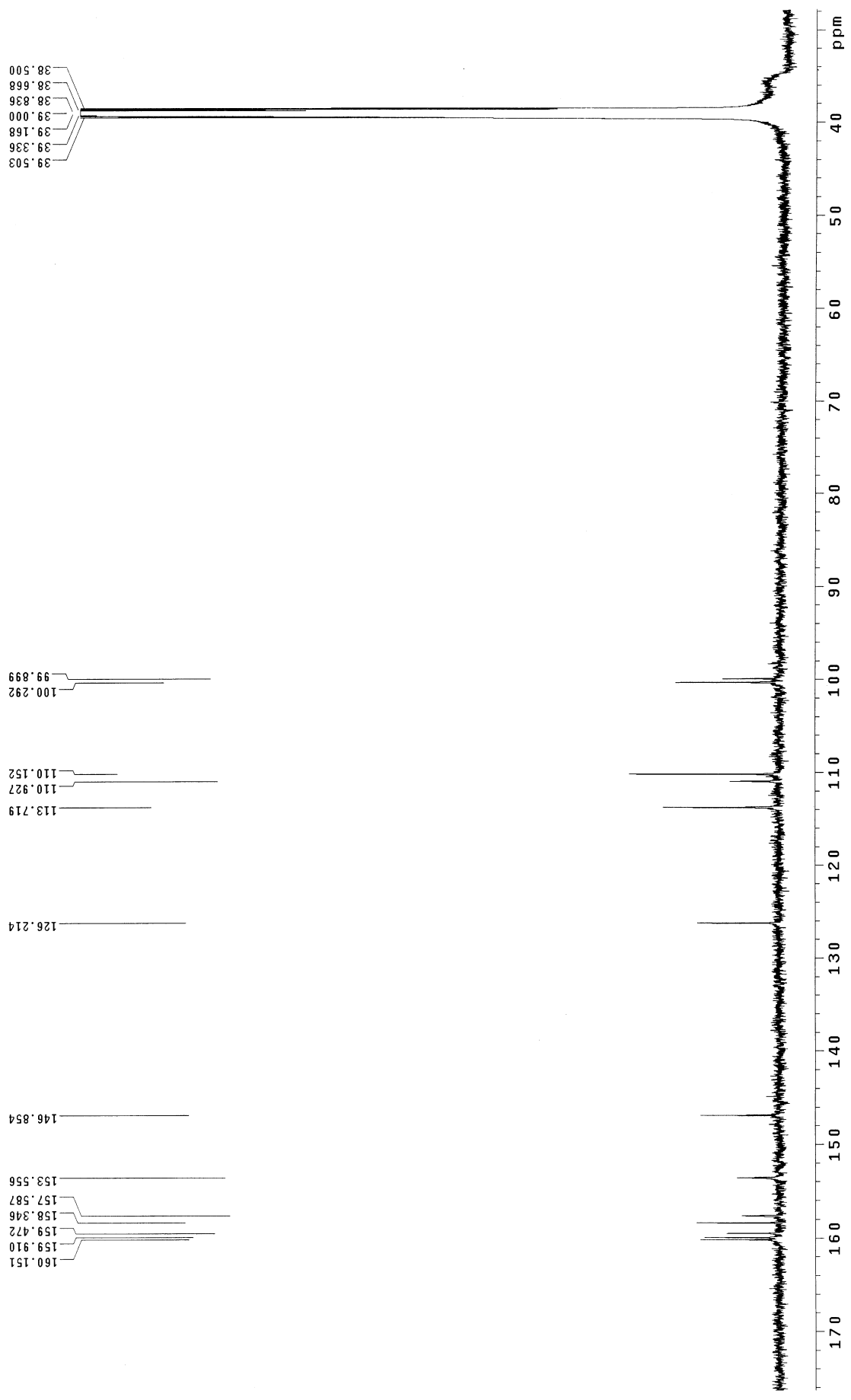
¹H NMR Spectrum of Phelligridin J (3)

INOVA-500 1H-NMR SHXY-11 IN DMSO 2005.12.19



¹³C NMR Spectrum of PheIigrudin J (3)

INOVA-500 ¹³C-NMR SHXY-11 IN DMSO



HMBc Spectrum of Phelliigrudin J (3)

Solvent: DMSO
Temp. 25.0 C / 288.1 K
User: 1-14-87
INOVA-500 "NMR500"

Relax. delay 1.000 sec
Acq. time 0.152 sec
Width 3377.1 Hz
2D Width 9358.9 Hz
400 repetitions
128 increments
OBSERVE H1, 500.0948900 MHz
DATA PROCESSING
Sine bell 0.076 sec
F1 DATA PROCESSING
Sine bell 0.007 sec
F1 size 1024 x 4096
Total time 17 hr, 38 min, 56 sec

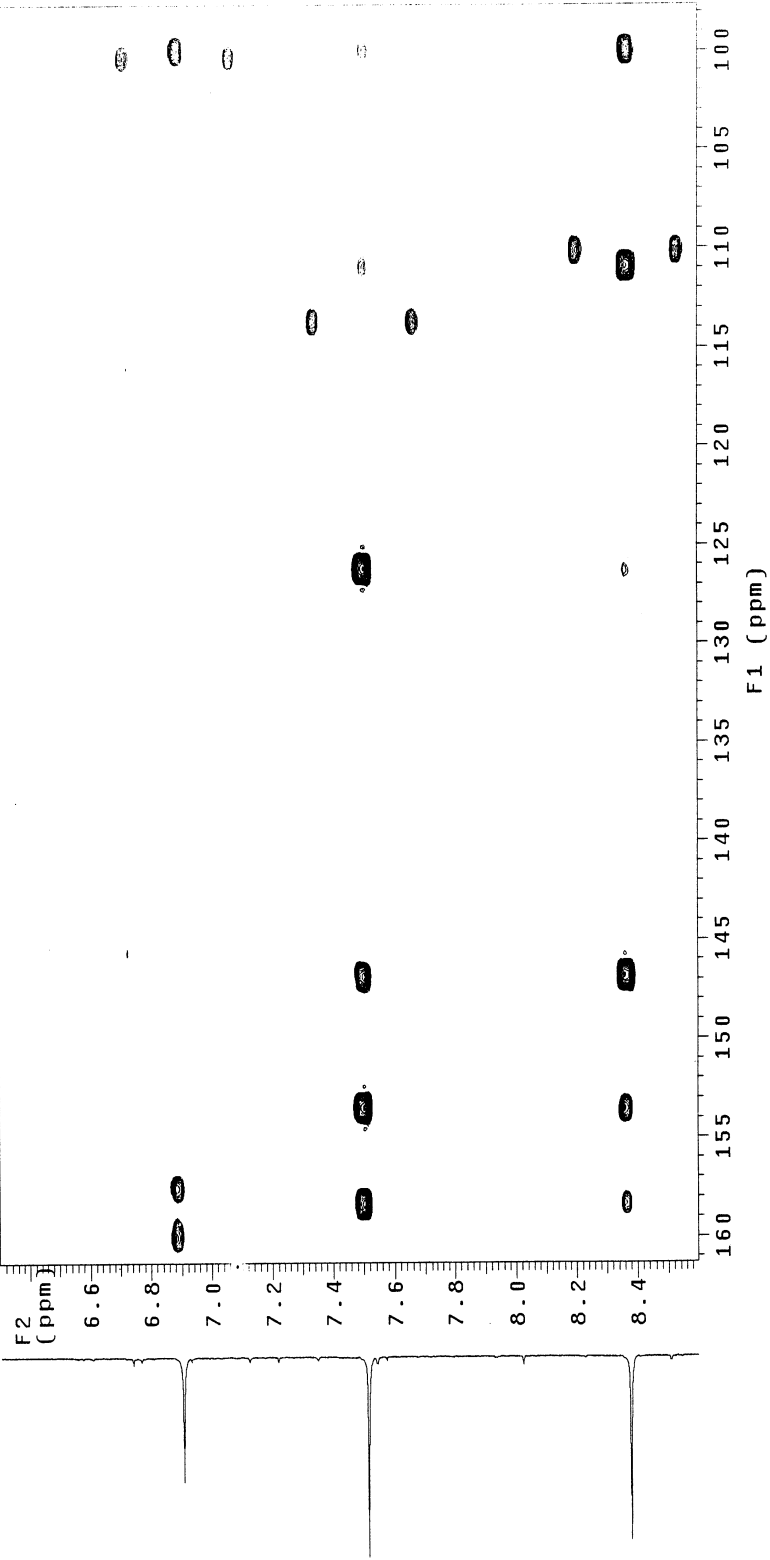


Table 1. ^1H and ^{13}C NMR data of compounds **1-3**. (Recorded in DMSO-d_6)^a

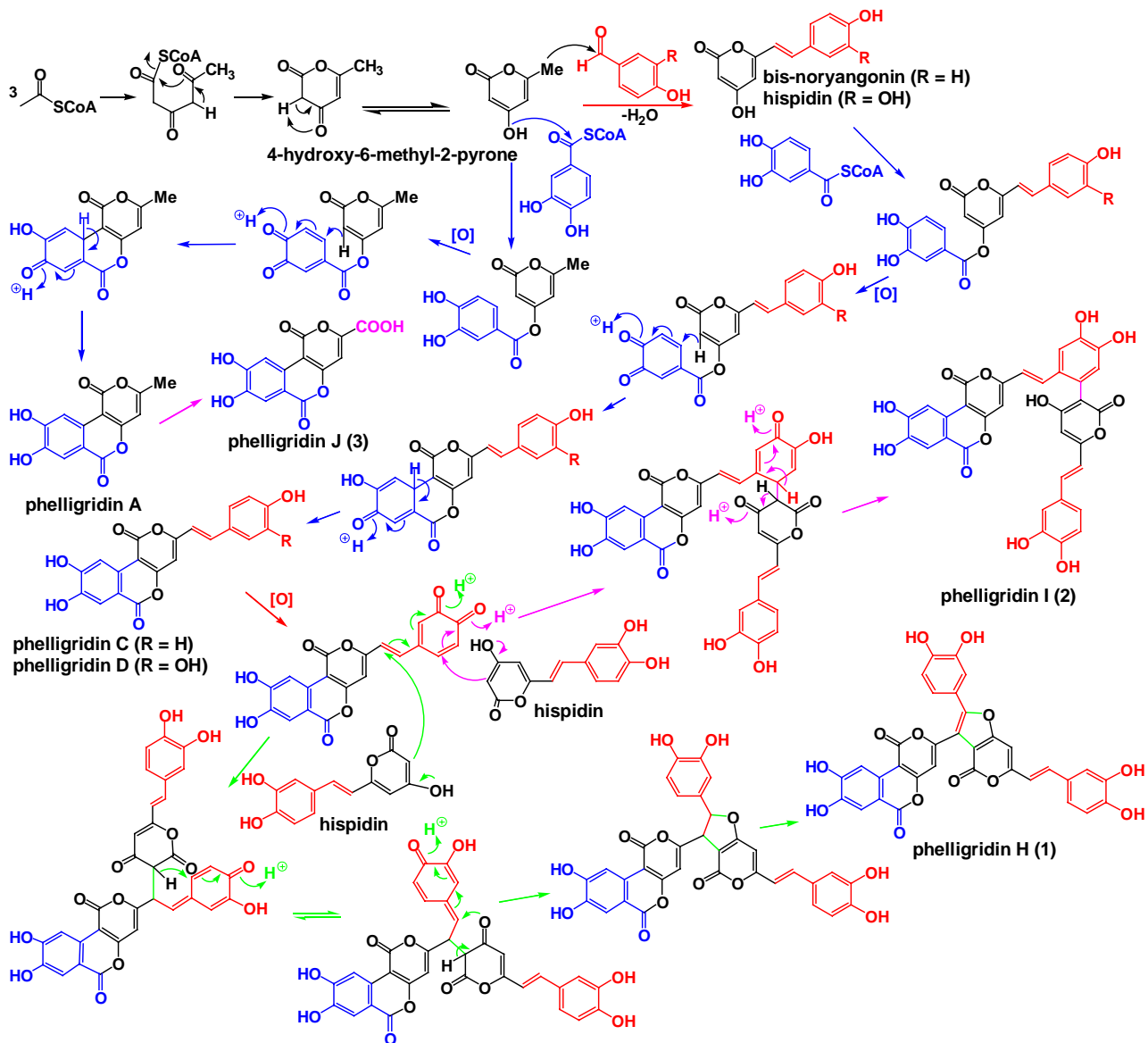
No.	δ_{H} (1)	δ_{C} (1)	δ_{H} (2)	δ_{C} (2)	δ_{H} (3)	δ_{C} (3)
1		159.8 s		159.4 s		159.9 s
3		151.7 s		158.6 s		159.5 s
4	7.25 s	103.7 d	6.74 s	99.3 d	6.90 s	100.3 d
4a		160.2 s		160.8 s		160.2 s
6		158.9 s		158.7 s		158.3 s
6a		112.2 s		111.5 s		110.9 s
7	7.58 s	114.8 d	7.50 s	114.5 d	7.51 s	113.7 d
8		147.4 s		146.9 s		146.9 s
9		153.8 s		153.6 s		153.6 s
10	8.38 s	111.0 d	8.26 s	110.6 d	8.38 s	110.2 d
10a		126.8 s		127.1 s		126.2 s
10b		100.5 s		98.8 s		99.9 s
11						157.6 s
1'		108.8 s	6.73 d (16.0)	115.8 d		
2'		155.1 s	7.08 d (16.0)	134.0 d		
3'		119.1 s		126.0 s		
4'	7.08 d (1.5)	114.7 d	7.20 s	112.0 d		
5'		145.5 s		145.4 s		
6'		147.9 s		147.4 s		
7'	6.83 d (7.5)	116.2 d	6.60 s	118.7 d		
8'	7.07 dd (7.5, 1.5)	120.0 d		126.0 s		
2''		157.4 s		162.7 s		
3''		107.8 s		101.5 s		
4''		161.0 s		166.0 s		
5''	7.14 s	95.4 d	6.34 s	100.4 d		
6''		158.2 s		158.3 s		
7''	6.81 d (16.0)	116.4 d	6.80 d (16.0)	116.4 d		
8''	7.24 d (16.0)	134.7 d	7.20 d (16.0)	134.8 d		
9''		127.2 s		127.0 s		
10''	7.06 d (1.5)	114.2 d	7.08 d (1.5)	114.4 d		
11''		145.7 s		145.6 s		
12''		147.5 s		147.4 s		
13''	6.77 d (8.0)	116.1 d	6.78 d (8.0)	115.8 d		
14''	6.98 dd (8.0, 1.5)	120.8 d	7.01 dd (8.0, 1.5)	120.4 d		

^aNMR data were measured at 500 MHz for proton and at 125 MHz for carbon. Proton coupling constants (J) in Hz are given in parentheses. The assignments were based on DEPT, ^1H - ^1H COSY, HSQC, and HMBC experiments.

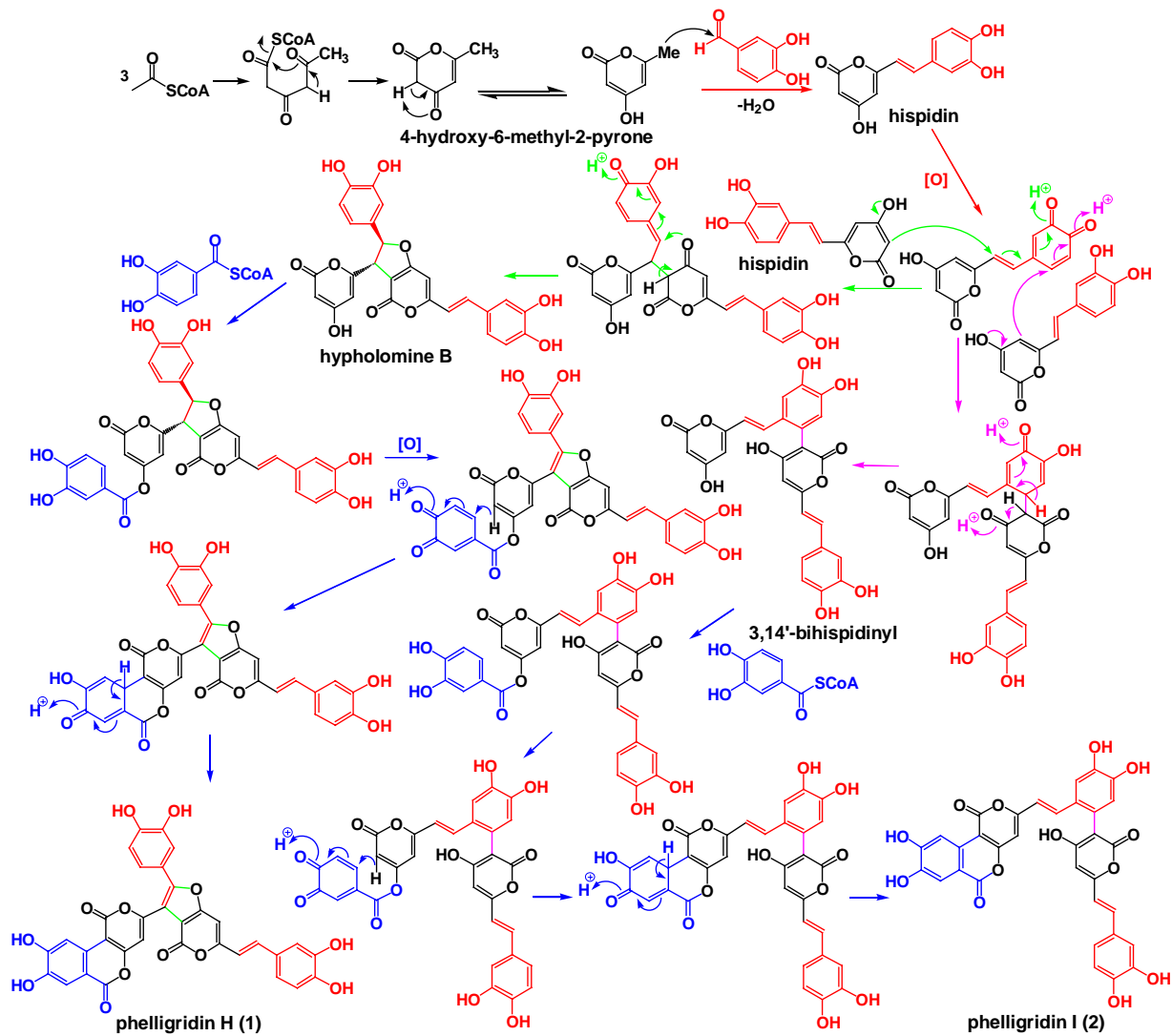
–OH of compound **1**: δ 9.12 (brs, 11''-OH), 9.34 (brs, 5'-OH), 9.49 (brs, 6'-OH), 9.60 (brs, 12''-OH), 10.21 (brs, 8-OH), 10.83 (brs, 9-OH).

–OH of compound **2**: δ 9.08 (brs, 5'-OH), 9.15 (brs, 11''-OH), 9.48 (brs, 6'-OH), 9.59 (brs, 12''-OH), 10.08 (brs, 8-OH), 10.71 (brs, 9-OH), 11.42 (brs, 4''-OH).

Scheme S1. Biogenetic scheme for phelligrindins H (1) and I (2) involving phelligrardin D, and phelligrardin J (3)



Scheme S2. Biogenetic scheme for phelligridins H (**1**) and I (**2**) involving hypholomine B and 3,14'-bihispidinyl.



Scheme S3. Biogenetic scheme for davallialactone (4), phelligrudin F, and inoscavin A from hispidin and hispilone.

