

## Supplementary data

# New Eunicellin-type Diterpenes from the Panamanian Octocoral *Briareum asbestinum*

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**Table S1.** Viability of THP-1 human macrophages treated with different concentrations of diterpenes.

**Figure S1.** Briarellin T,  $^1\text{H}$  NMR spectrum

```

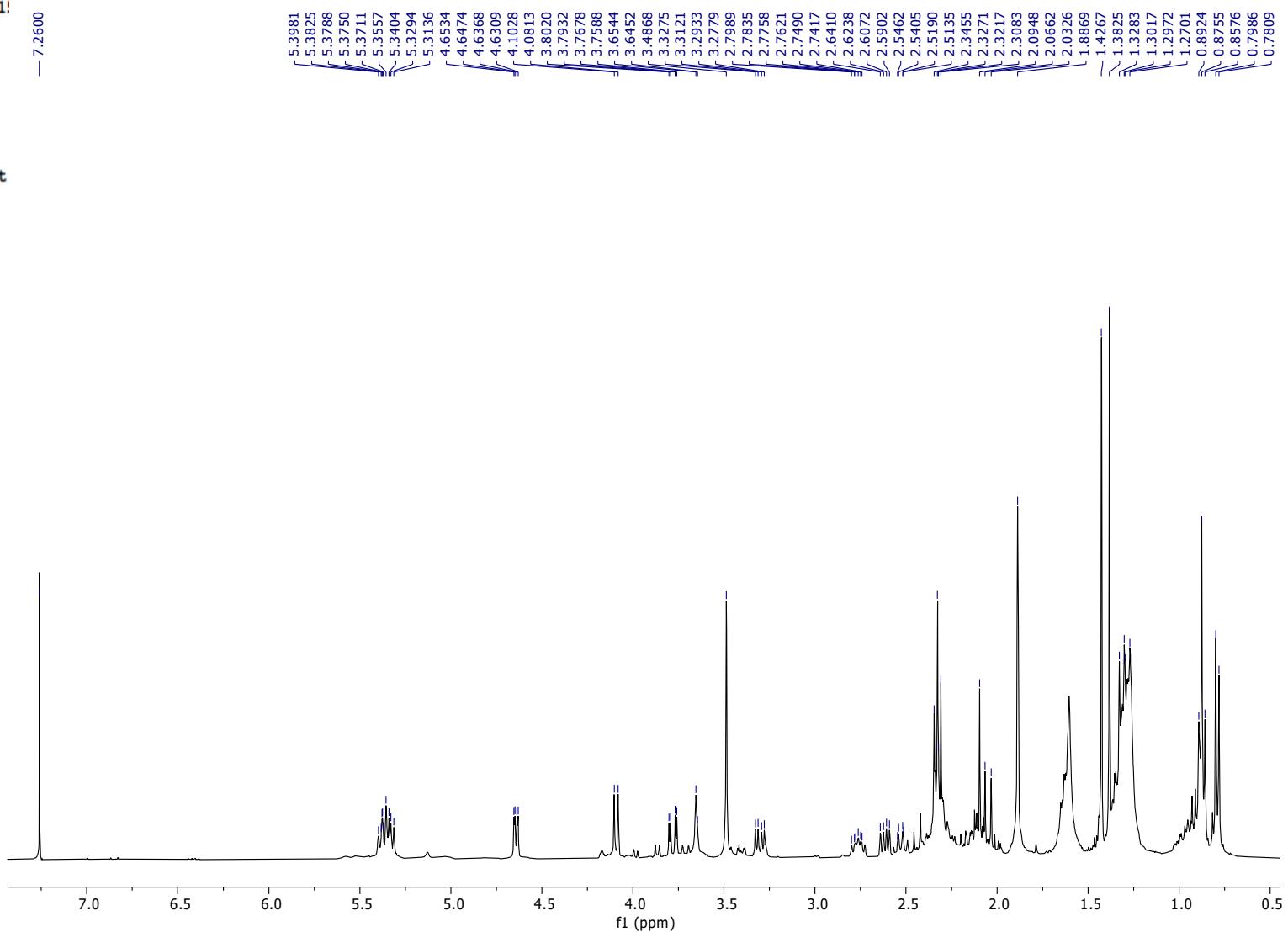
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Revision_Time  = 31-JUL-2015 07:40:50

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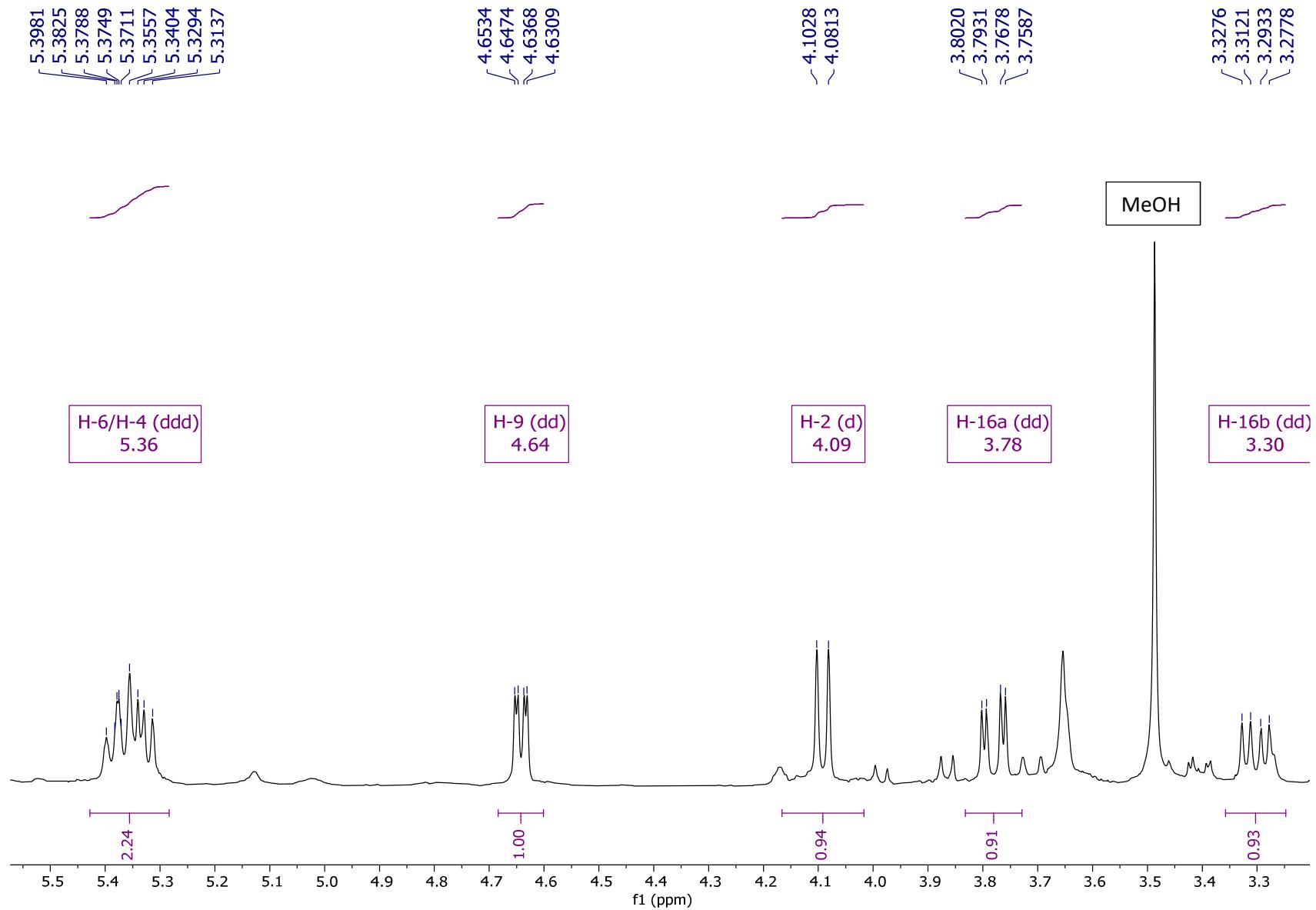
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X_Points       = 16384
X_Prescans    = 0
X_Resolution   = 0.36613771[Hz]
X_Sweep        = 5.99880024[kHz]
Clipped        = FALSE
Scans          = 32
Total_Scans    = 32

Relaxation_Delay = 4[s]
Recvrx_Gain     = 16
Temp_Get        = 21.9[dC]
X_90_Width      = 9.81[us]
X_Acq_Time      = 2.7312128[s]
X_Angle          = 45[deg]
X_Pulse          = 4.905[us]
Initial_Wait    = 1[s]
Unblank_Time    = 2[us]

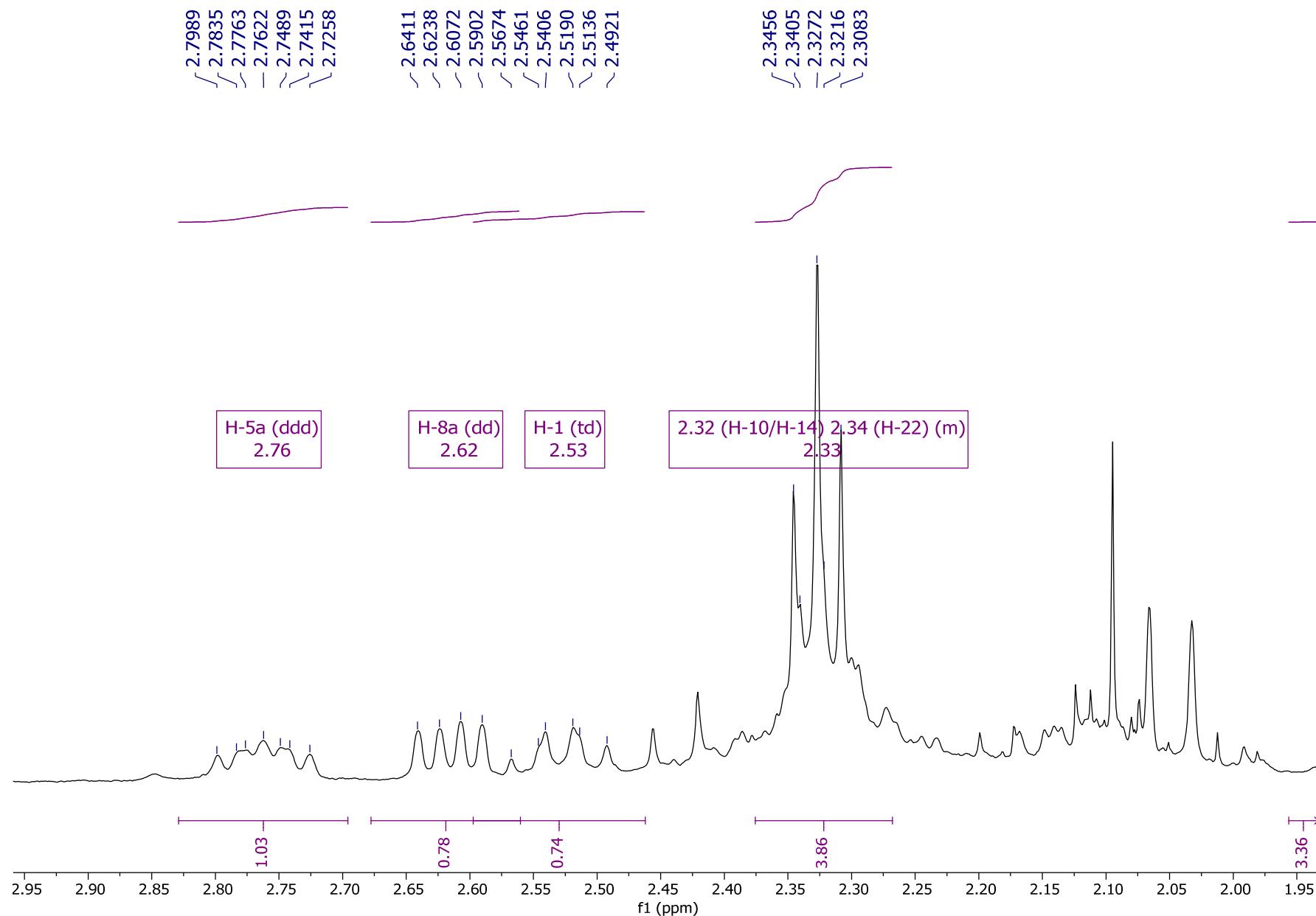
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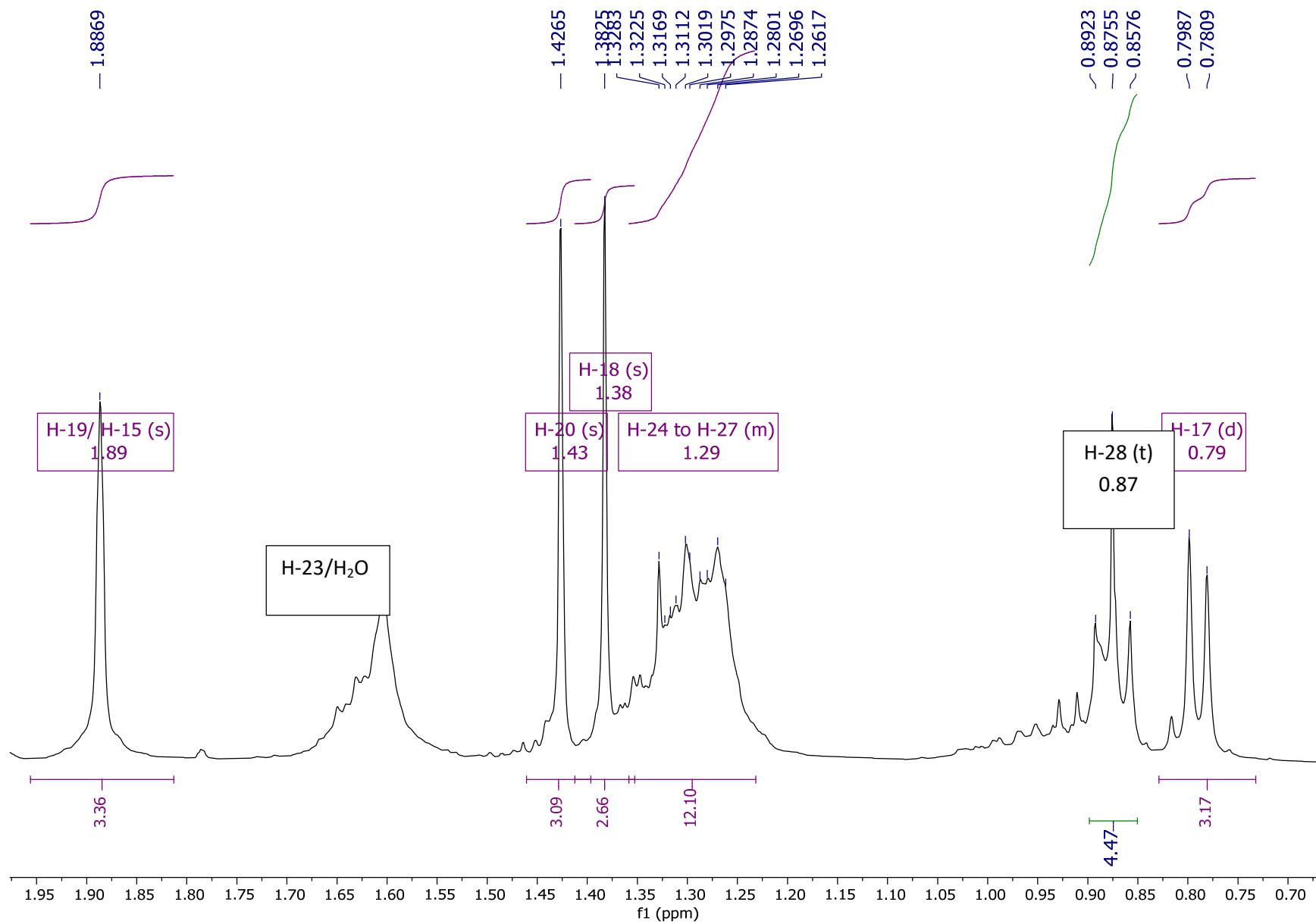
**Figure S1.** Briarellin T,  $^1\text{H}$  NMR spectrum (expanded)



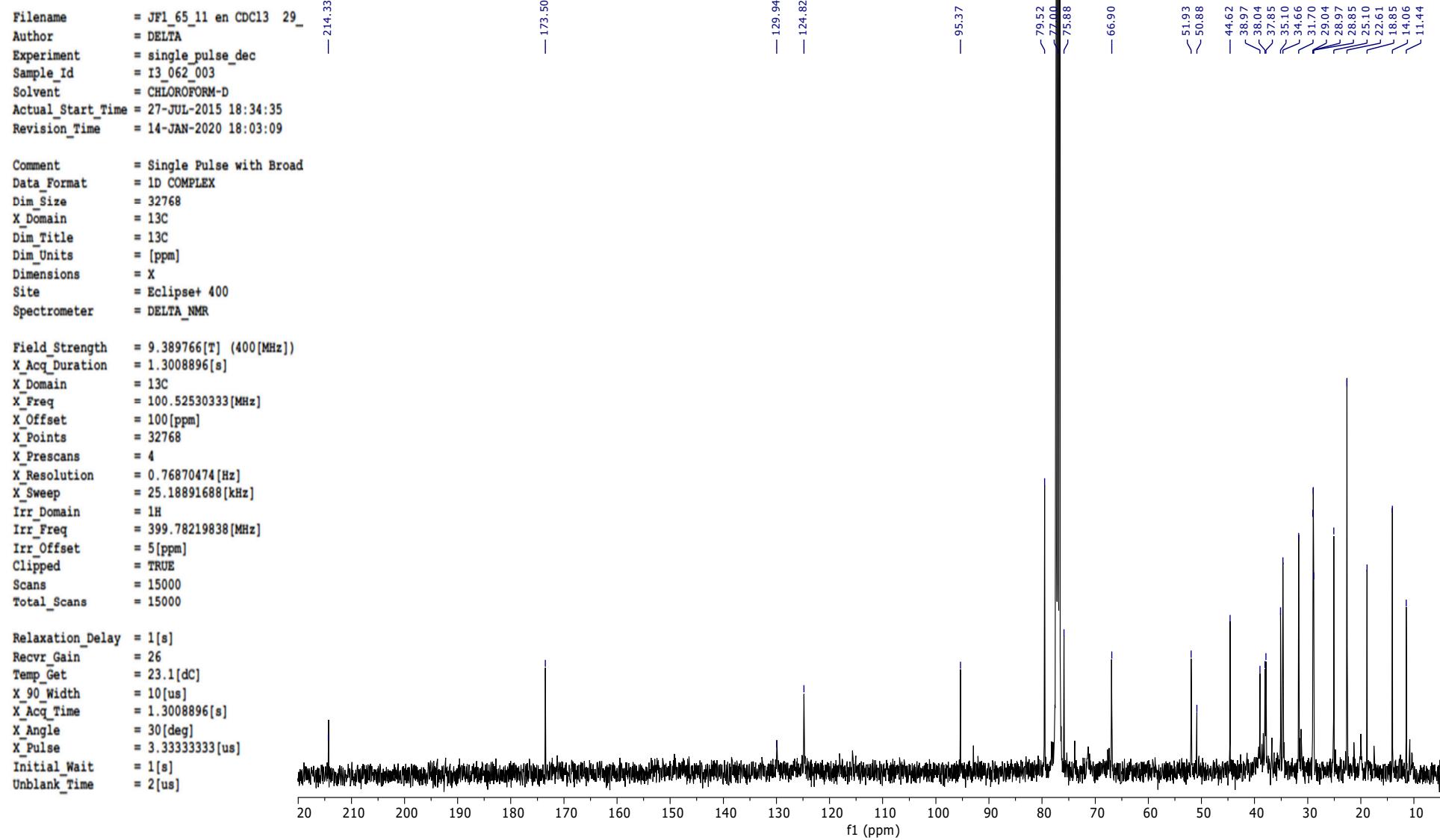
**Figure S1.** Briarellin T,  $^1\text{H}$  NMR spectrum (expanded)



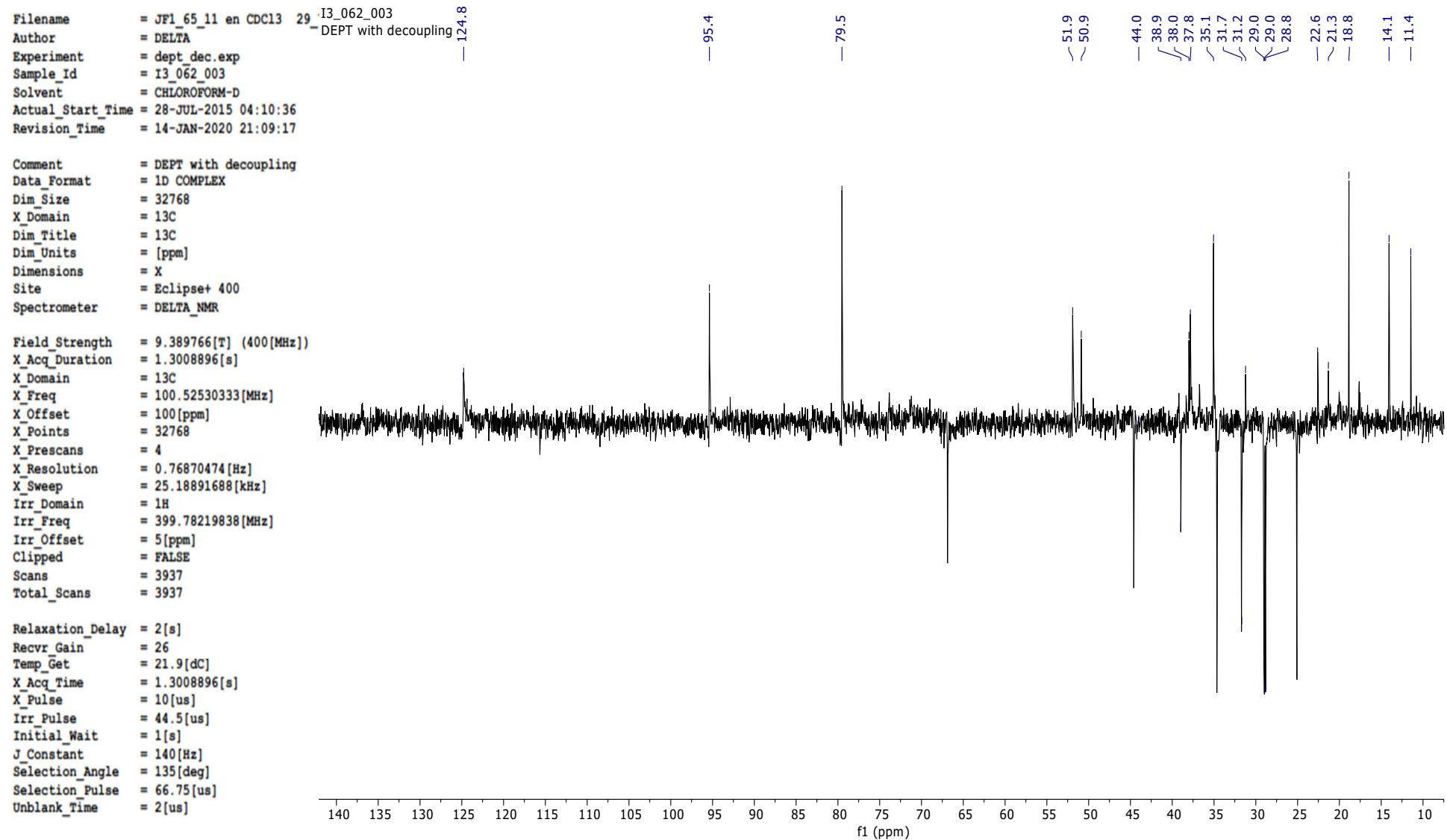
**Figure S1.** Briarellin T,  $^1\text{H}$  NMR spectrum (expanded)



**Figure S2.** Briarellin T,  $^{13}\text{C}$  NMR spectrum



**Figure S3.** Briarellin T, DEPT-135 spectrum



**Figure S4.** Briarellin T, COSY spectrum

```

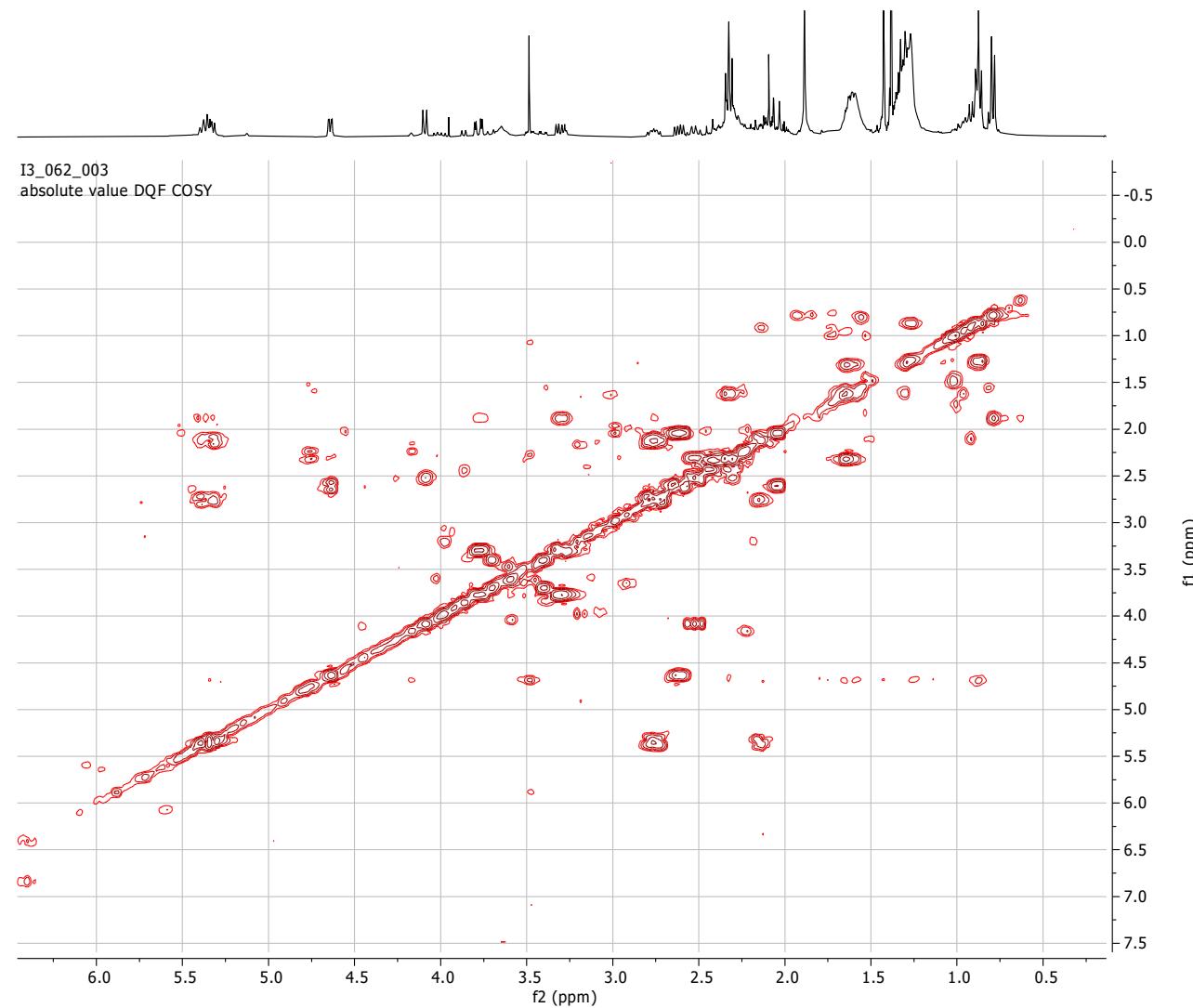
Filename      = JF1_65_11_CDCl3 30_7_1
Author        = DELTA
Experiment    = dqf_cosy.exp
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Revision_Time = 23-OCT-2018 12:47:02

Comment       = absolute value DQF COSY
Data_Format   = 2D REAL REAL
Dim_Size      = 512, 1024
X_Domain     = 1H
Y_Domain     = 1H
Dim_Title    = 1H 1H
Dim_Units    = [ppm] [ppm]
Dimensions   = X Y
Site          = Eclipse+ 400
Spectrometer = DELTA_NMR

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X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 4.68335[ppm]
X_Points       = 512
X_Prescans    = 4
X_Resolution  = 8.8217028[Hz]
X_Sweep        = 4.51671183[kHz]
Y_Domain      = 1H
Y_Freq         = 399.78219838[MHz]
Y_Offset       = 4.68335[ppm]
Y_Points       = 256
Y_Prescans    = 0
Y_Resolution  = 17.6434056[Hz]
Y_Sweep        = 4.51671183[kHz]
Clipped       = FALSE
Scans          = 16
Total_Scans   = 4096

Relaxation_Delay = 1.5[s]
Recvr_Gain      = 15
Temp_Get         = 21.6[dC]
X_Acq_Time      = 0.1133568[s]
X_Pulse          = 9.81[us]
Y_Acq_Time      = 56.6784[ms]
Initial_Wait    = 1[s]
T1              = 1[us]
Unblank_Time   = 2[us]

```



**Figure S5.** Briarellin T, HSQC spectrum

```

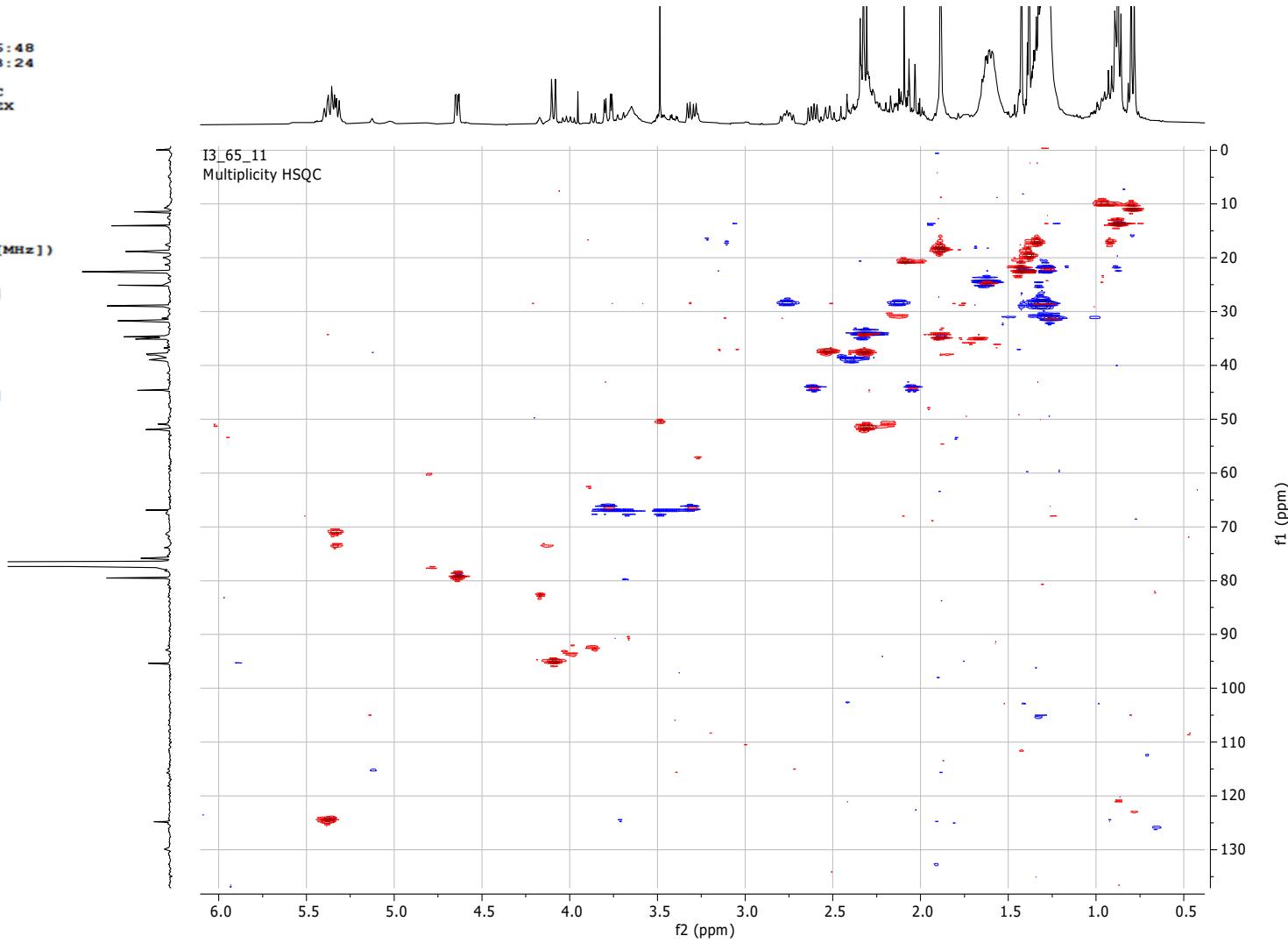
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Author        = DELTA
Experiment    = multiplicity_hsq
Sample_Id     = I3_062_003
Solvent       = CHLOROFORM-D
Actual_Start_Time = 28-JUL-2015 10:56:48
Revision_Time = 14-JAN-2020 21:23:24

Comment       = Multiplicity HSQC
Data_Format   = 2D COMPLEX COMPLEX
Dim_Size      = 2048, 324
X_Domain     = 13C
Y_Domain     = 13C
Dim_Title    = 13C 13C
Dim_Units    = [ppm] [ppm]
Dimensions   = X Y
Site          = Eclipse+ 400
Spectrometer = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acc_Duration = 66.9696[ms]
X_Domain      = 13C
X_Freq         = 100.52530333[MHz]
X_Offset       = 68.30581[ppm]
X_Points       = 1024
X_Prescans    = 4
X_Resolution  = 14.93214832[Hz]
X_Sweep        = 15.29051988[kHz]
Y_Domain      = 13C
Y_Freq         = 100.52530333[MHz]
Y_Offset       = 100[ppm]
Y_Points       = 162
Y_Prescans    = 0
Y_Resolution  = 155.87978551[Hz]
Y_Sweep        = 25.25252525[kHz]
Clipped       = FALSE
Scans          = 16
Total_Scans   = 2579

Relaxation_Delay = 2[s]
Recvr_Gain      = 27
Temp_Get         = 22.1[dC]
X_Acc_Time      = 66.9696[ms]
X_Pulse          = 10[us]
Y_Acc_Time      = 10.1376[ms]
Y_Pulse          = 10[us]
Enhance_Temp    = 12
Enhancement     = 1/6J
Grad_1           = 1[ms]
Grad_1_Amp      = 4[pnt]
Grad_1_Value    = 4[pnt*ms]
Grad_2           = 1[ms]
Grad_2_Amp      = -1[pnt]
Grad_2_Value    = -1[pnt*ms]
Grad_3           = 1[ms]
Grad_3_Amp      = 1[pnt]
Grad_Recover    = 0.1[ms]
Grad_Selection  = 13C = 4:1
Grad_Shape      = square
Grad_Type       = 0
Initial_Wait   = 1[s]
J_Constant     = 140[Hz]
T1              = 1[us]
Tau             = 0.5952381[ms]

```



**Figure S6.** Briarellin T, HMBC spectrum

```

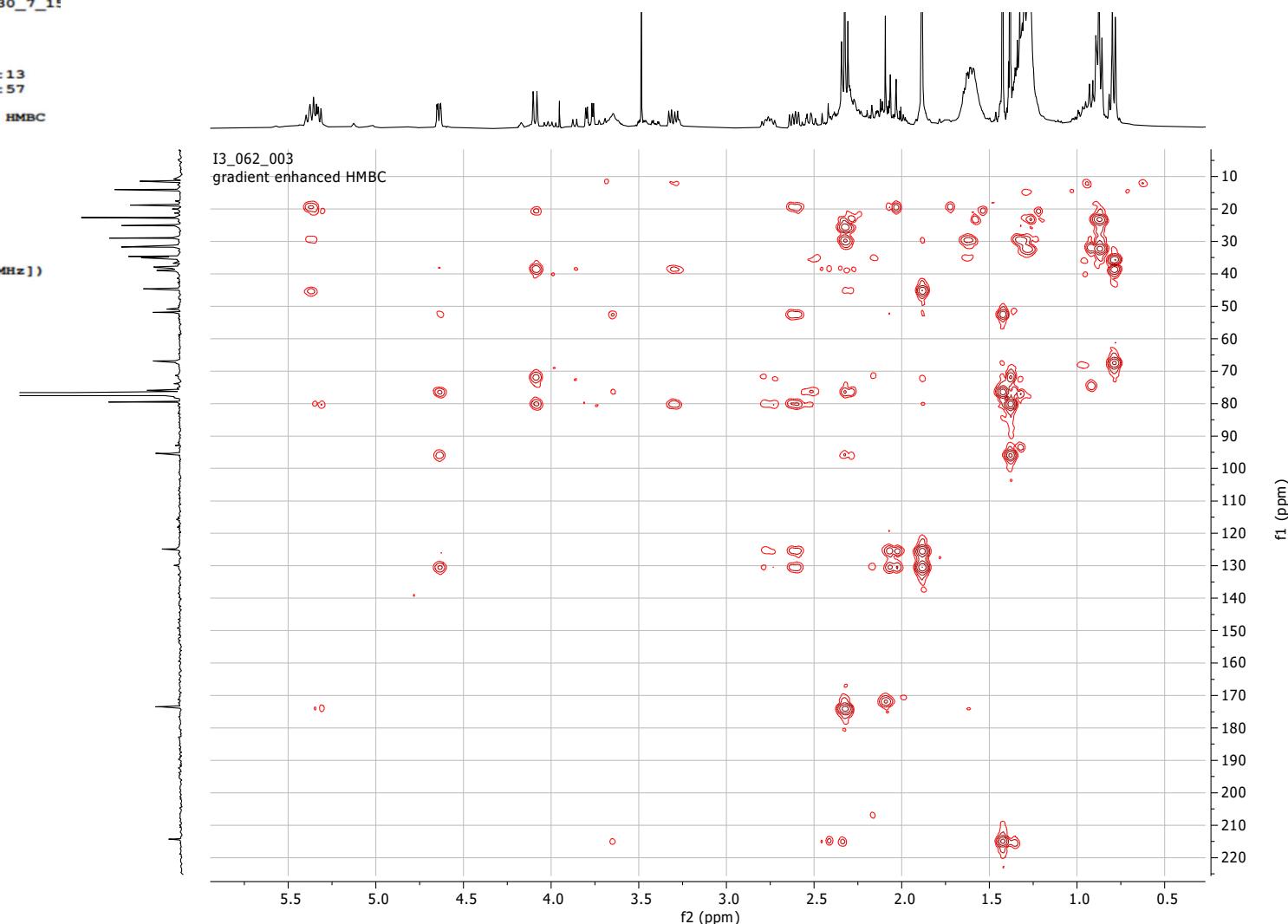
Filename      = JF1_65_11_CDC13 30_7_15
Author        = DELTA
Experiment    = hmbo_pfg_s.exp
Sample_Id     = I3_062_003
Solvent       = CHLOROFORM-D
Actual_Start_Time = 29-JUL-2015 00:40:13
Revision_Time = 31-JUL-2015 07:55:57

Comment       = gradient enhanced HMBC
Data_Format   = 2D REAL REAL
Dim_Size      = 1024, 512
X_Domain     = 1H
Y_Domain     = 13C
Dim_Title    = 1H 13C
Dim_Units    = [ppm] [ppm]
Dimensions   = X Y
Site          = Eclipse+ 400
Spectrometer = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 0.2267136[s]
X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 4.68335[ppm]
X_Points       = 1024
X_Prescans    = 4
X_Resolution  = 4.4108514[Hz]
X_Sweep        = 4.51671183[kHz]
Y_Domain      = 13C
Y_Freq         = 100.52530333[MHz]
Y_Offset       = 100[ppm]
Y_Points       = 256
Y_Prescans    = 0
Y_Resolution  = 98.64267677[Hz]
Y_Sweep        = 25.25252525[kHz]
Clipped       = FALSE
Scans          = 32
Total_Scans   = 8192

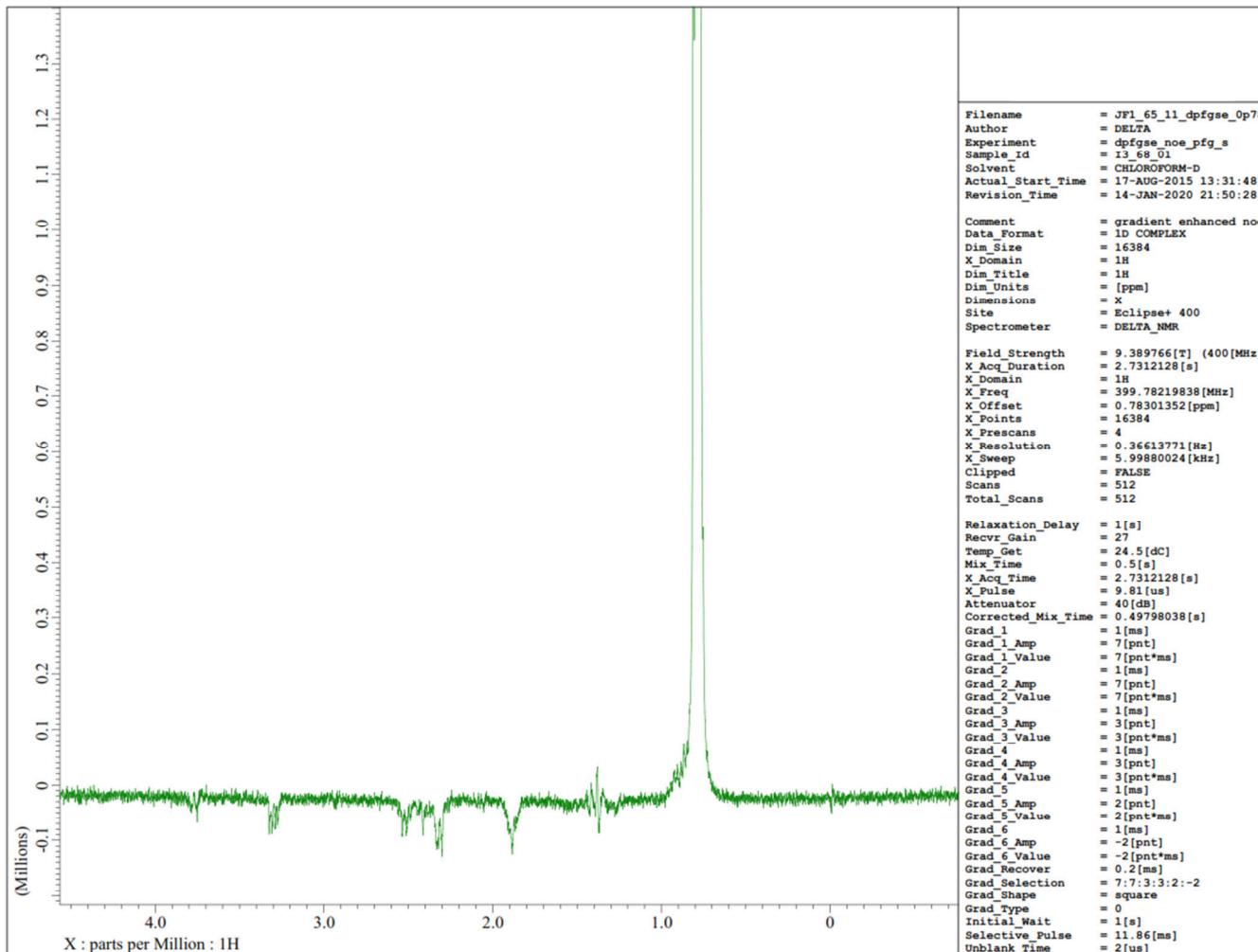
Relaxation_Delay = 2[s]
Recvr_Gain      = 30
Temp_Get         = 21.3[dC]
X_Acq_Time      = 0.2267136[s]
X_Pulse          = 9.81[us]
Y_Acq_Time      = 10.1376[ms]
Y_Pulse          = 12[us]
Grad_1           = 1[ms]
Grad_1_Amp       = 10[pnt]
Grad_1_Value     = 10[pnt*ms]
Grad_2           = 1[ms]
Grad_2_Amp       = 10[pnt]
Grad_2_Value     = 10[pnt*ms]
Grad_3           = 1[ms]
Grad_3_Amp       = 5[pnt]
Grad_3_Value     = 5[pnt*ms]
Grad_Recover    = 0.2[ms]
Grad_Selection  = 13C = 2:2:1
Grad_Shape       = square
Grad_Type        = 0
Initial_Wait    = 1[s]
J_Constant      = 140[Hz]
Long_Range_J    = 8[Hz]
T1              = 1[us]
Unblank_Time    = 2[us]

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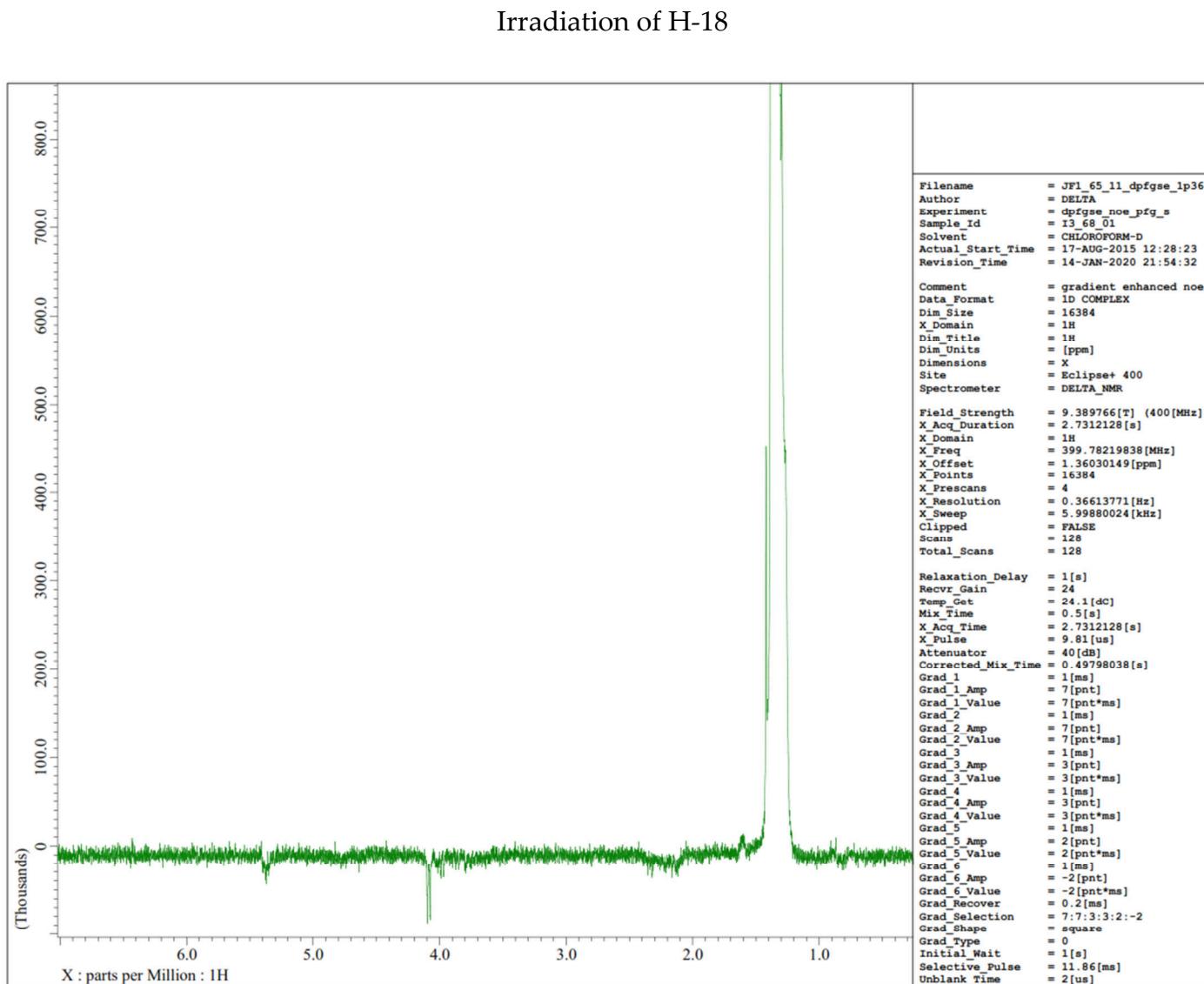


**Figure S7.** Briarellin T, NOE spectra

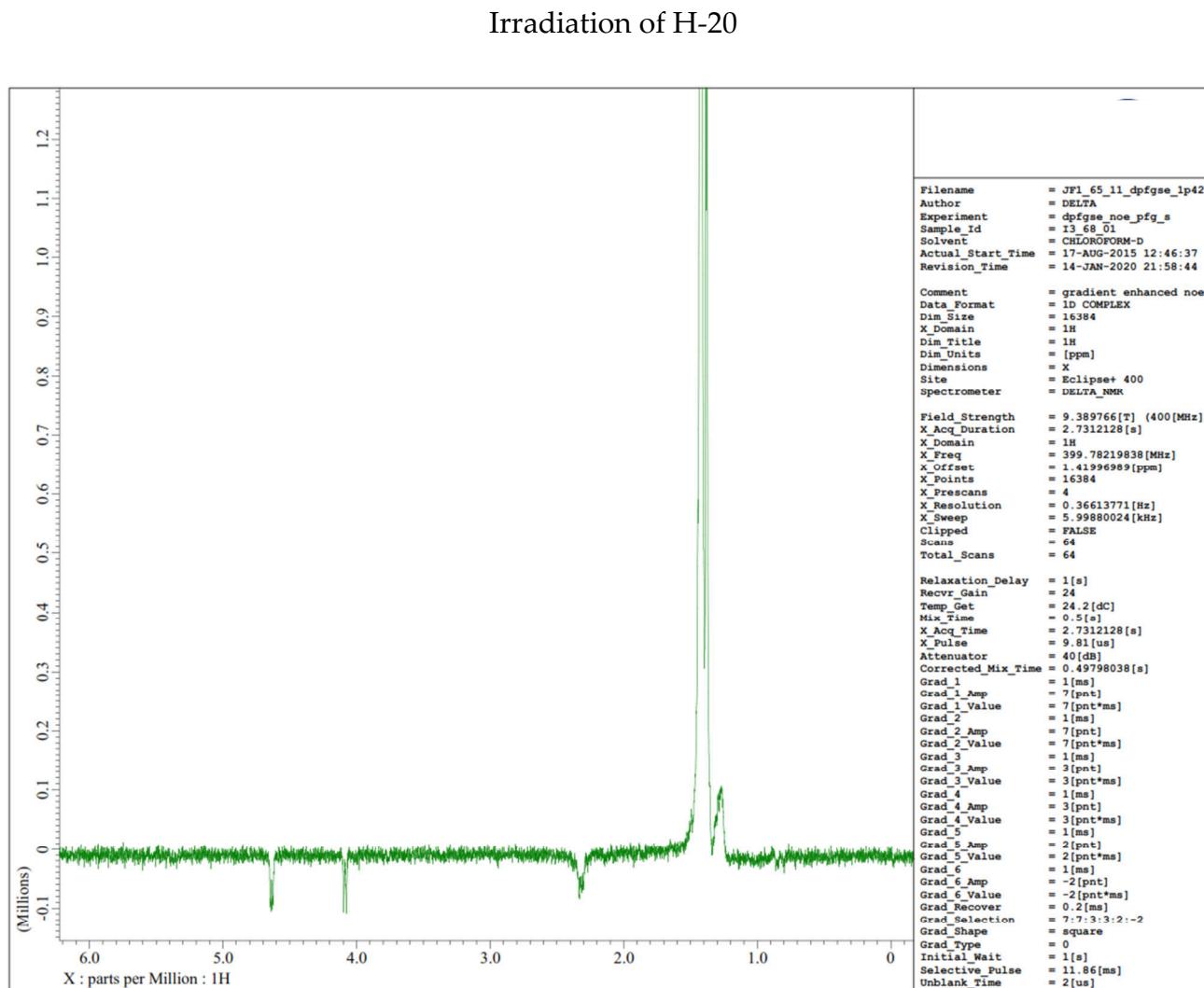
Irradiation of H-17



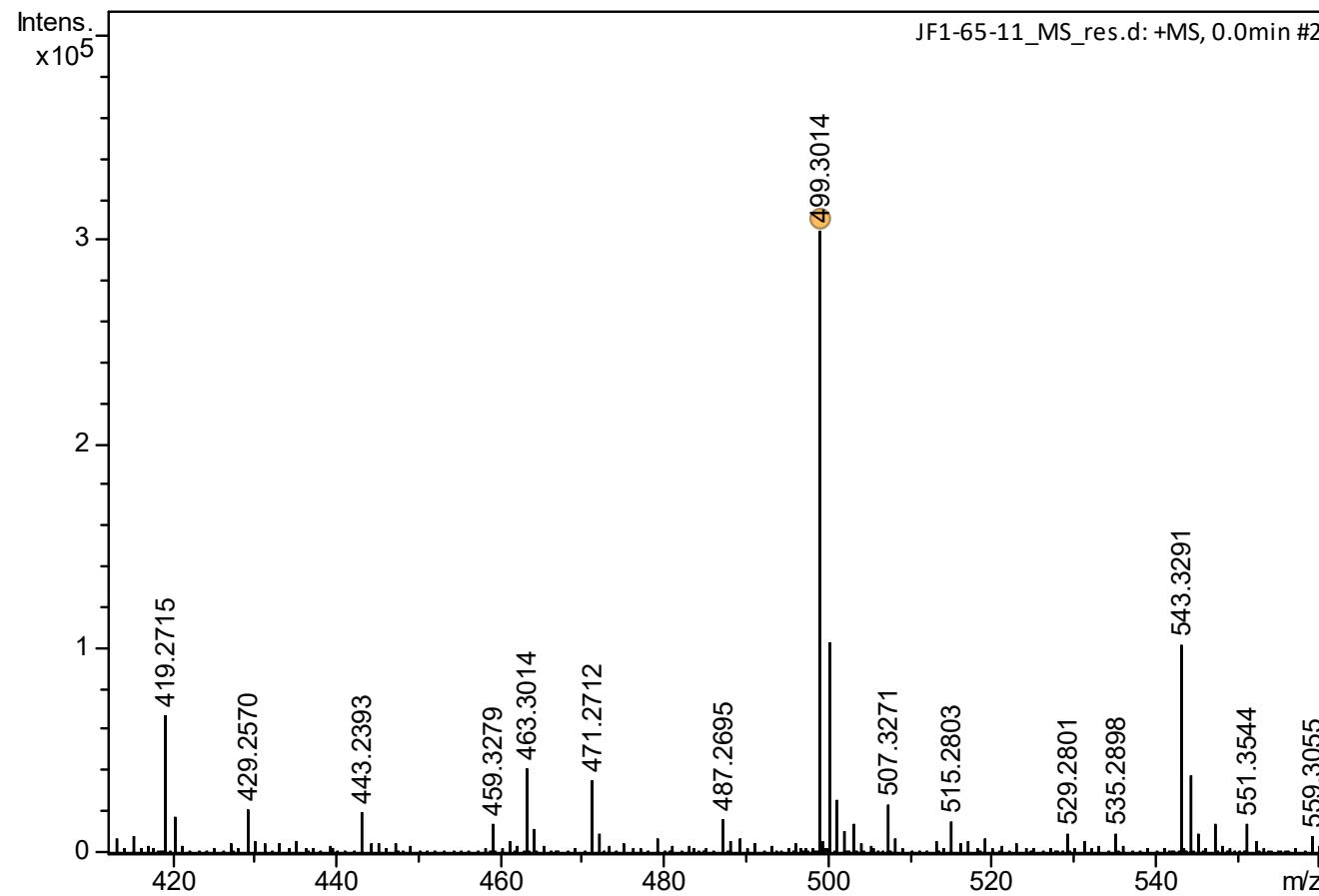
**Figure S7.** Briarellin T, NOE spectra



**Figure S7.** Briarellin T, NOE spectra



**Figure S8. Briarellin T, HR-ESITOFMS spectrum**



**Figure S9. Asbestinin 27,  $^1\text{H}$  NMR spectrum**

```

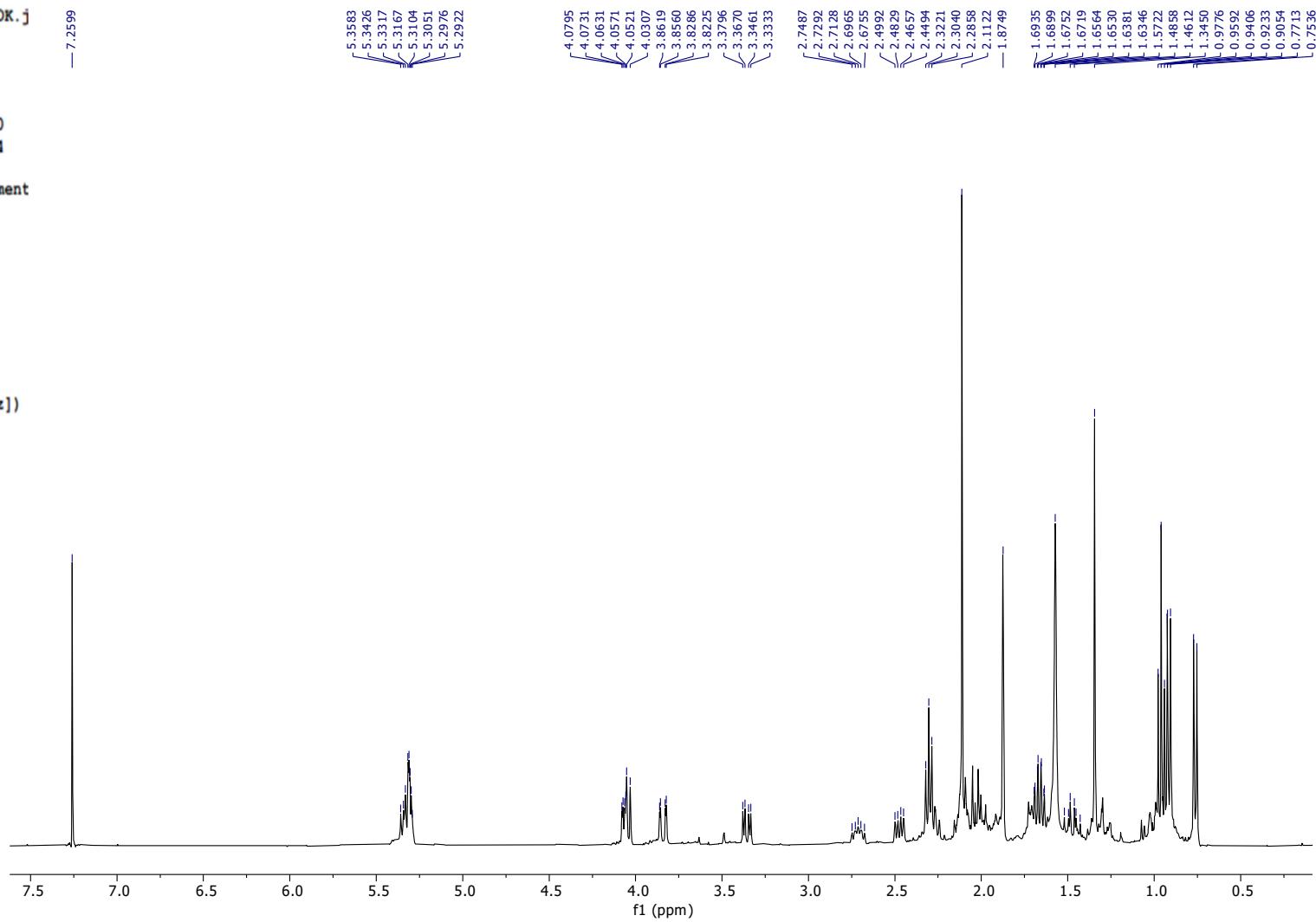
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Author        = DELTA
Experiment    = single_pulse.exp
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Solvent       = CHLOROFORM-D
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Revision_Time  = 19-AUG-2015 15:41:04

Comment       = Single Pulse Experiment
Data_Format   = 1D COMPLEX
Dim_Size      = 16384
X_Domain     = 1H
Dim_Title     = 1H
Dim_Units     = [ppm]
Dimensions    = X
Site          = Eclipse+ 400
Spectrometer  = DELTA_NMR

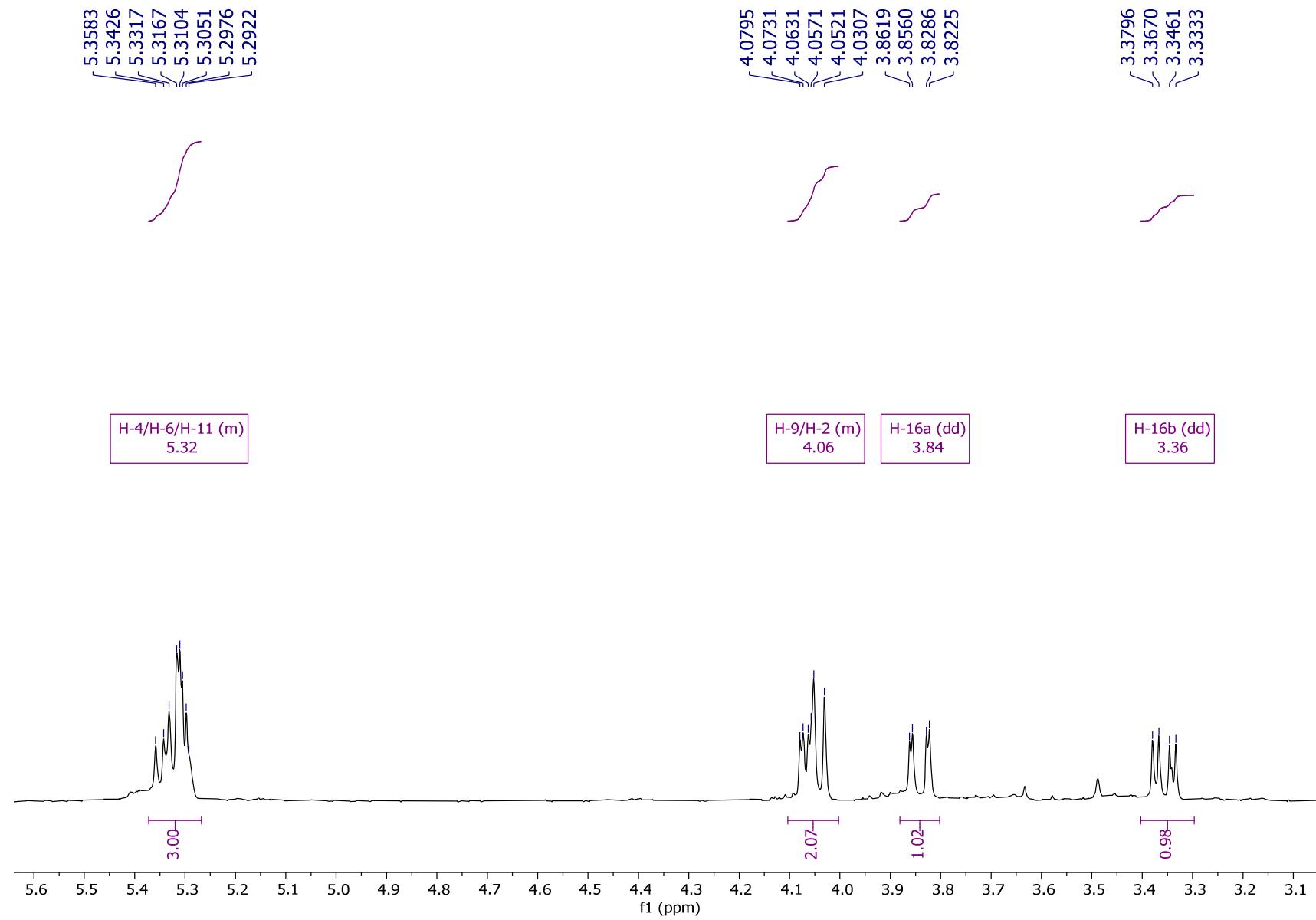
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X_Acq_Duration = 2.7312128[s]
X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 5[ppm]
X_Points       = 16384
X_Prescans    = 0
X_Resolution   = 0.36613771[Hz]
X_Sweep        = 5.99880024[kHz]
Clipped        = FALSE
Scans          = 16
Total_Scans    = 16

Relaxation_Delay = 4[s]
Recvr_Gain      = 17
Temp_Get         = 25[dC]
X_90_Width       = 9.81[us]
X_Acc_Time       = 2.7312128[s]
X_Angle          = 45[deg]
X_Pulse          = 4.905[us]
Initial_Wait     = 1[s]
Unblank_Time     = 2[us]

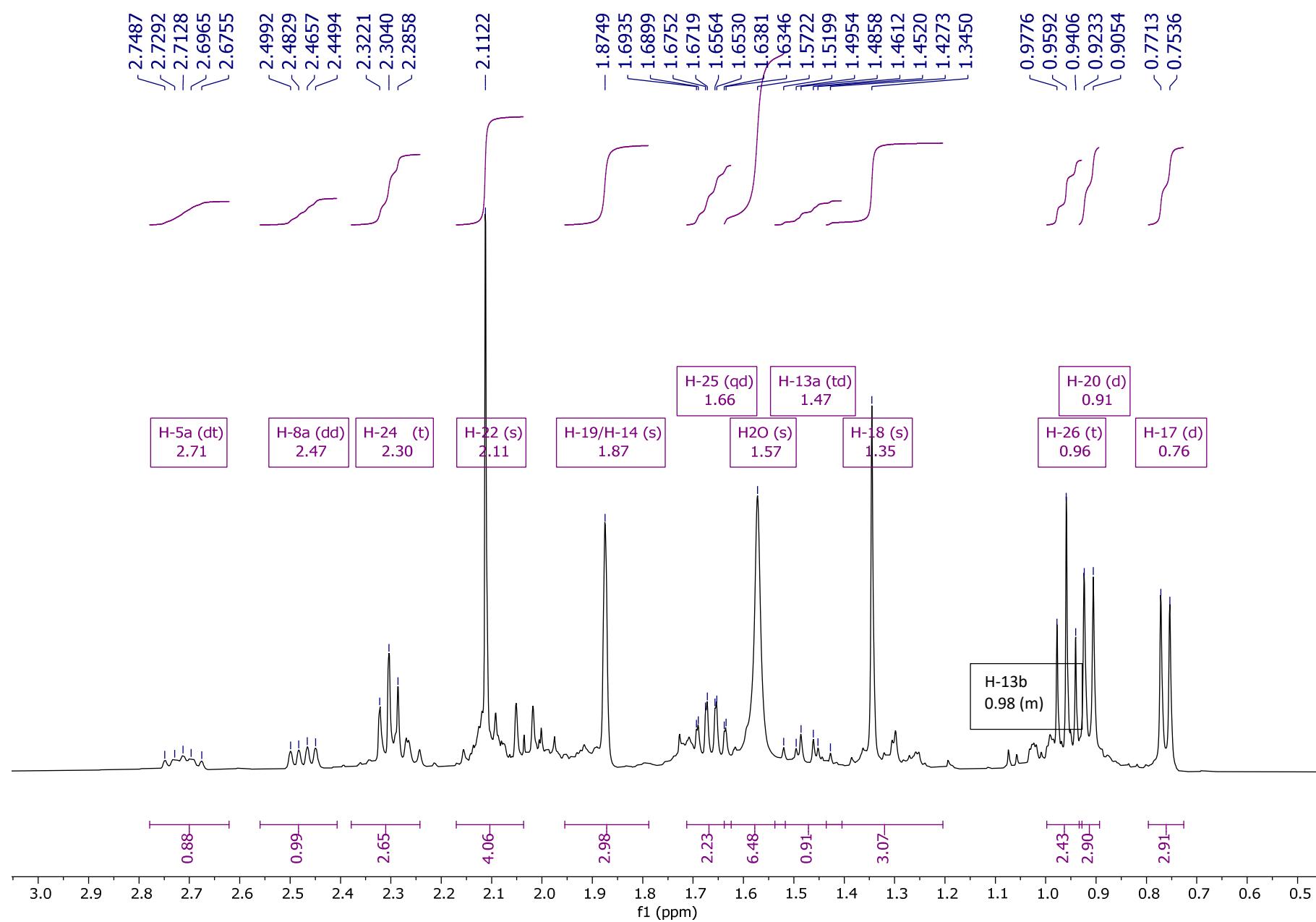
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**Figure S9. Asbestinin 27,  $^1\text{H}$  NMR spectrum (expanded)**



**Figure S9. Asbestinin 27,  $^1\text{H}$  NMR spectrum (expanded)**



**Figure S10. Asbestinin 27,  $^{13}\text{C}$  NMR spectrum**

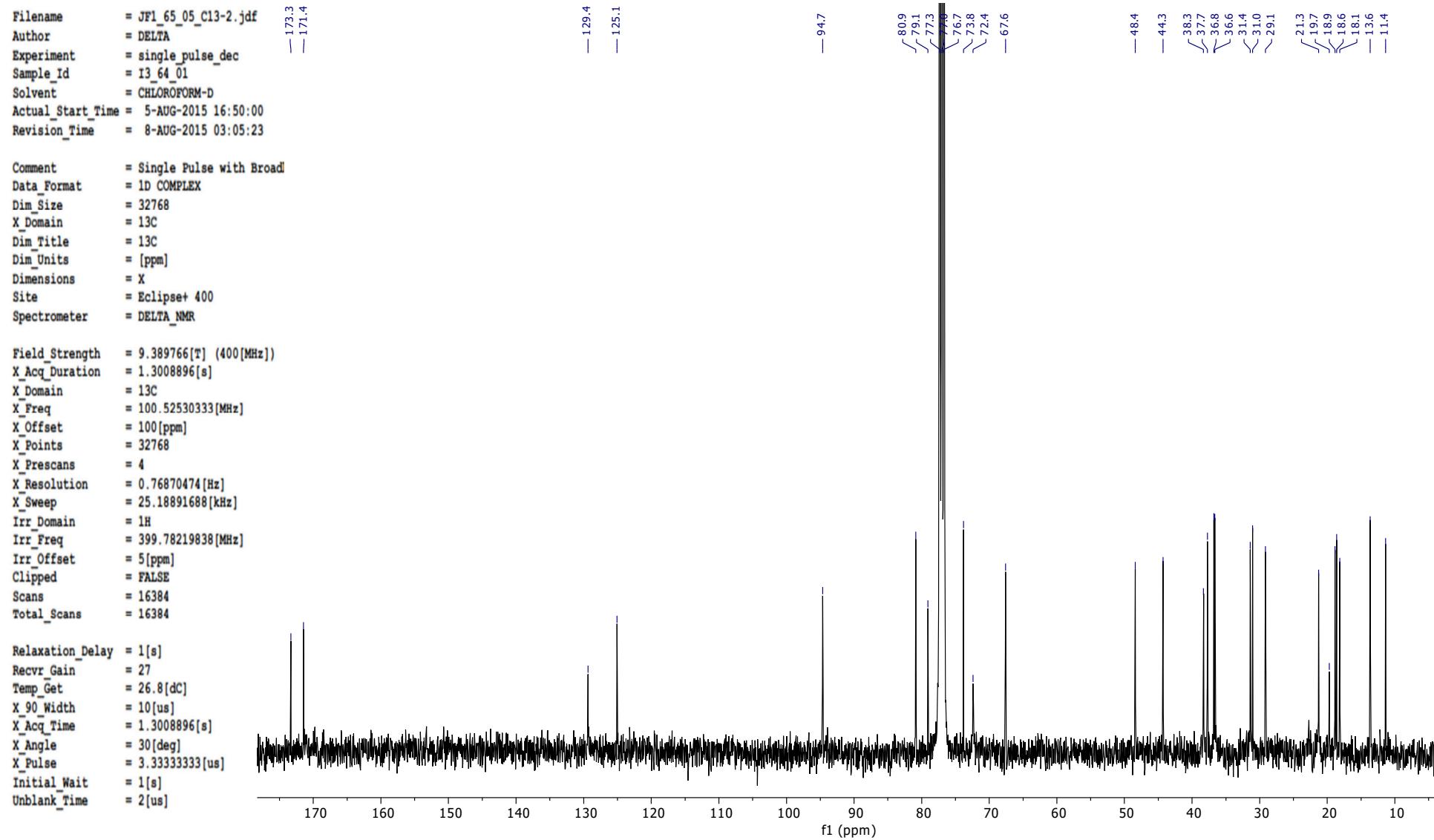


Figure S11. Asbestinin 27, DEPT-135 spectrum

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Author        = DELTA
Experiment    = dept_dec.exp
Sample_Id     = I3_68_02
Solvent       = CHLOROFORM-D
Actual_Start_Time = 17-AUG-2015 15:56:24
Revision_Time  = 14-JAN-2020 22:39:27

Comment       = DEPT with decoupling
Data_Format   = 1D COMPLEX
Dim_Size      = 32768
X_Domain     = 13C
Dim_Title     = 13C
Dim_Units     = [ppm]
Dimensions    = X
Site          = Eclipse+ 400
Spectrometer  = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 1.3008896[s]
X_Domain      = 13C
X_Freq         = 100.52530333[MHz]
X_Offset       = 100[ppm]
X_Points       = 32768
X_Prescans    = 4
X_Resolution   = 0.76870474[Hz]
X_Sweep        = 25.18891688[kHz]
Irr_Domain    = 1H
Irr_Freq       = 399.78219838[MHz]
Irr_Offset     = 5[ppm]
Clipped        = FALSE
Scans          = 1024
Total_Scans   = 1024

Relaxation_Delay = 2[s]
Recvr_Gain      = 27
Temp_Get         = 25.7[dC]
X_Acq_Time      = 1.3008896[s]
X_Pulse          = 10[us]
Irr_Pulse        = 44.5[us]
Initial_Wait    = 1[s]
J_Constant       = 140[Hz]
Selection_Angle = 135[deg]
Selection_Pulse = 66.75[us]
Unblank_Time    = 2[us]
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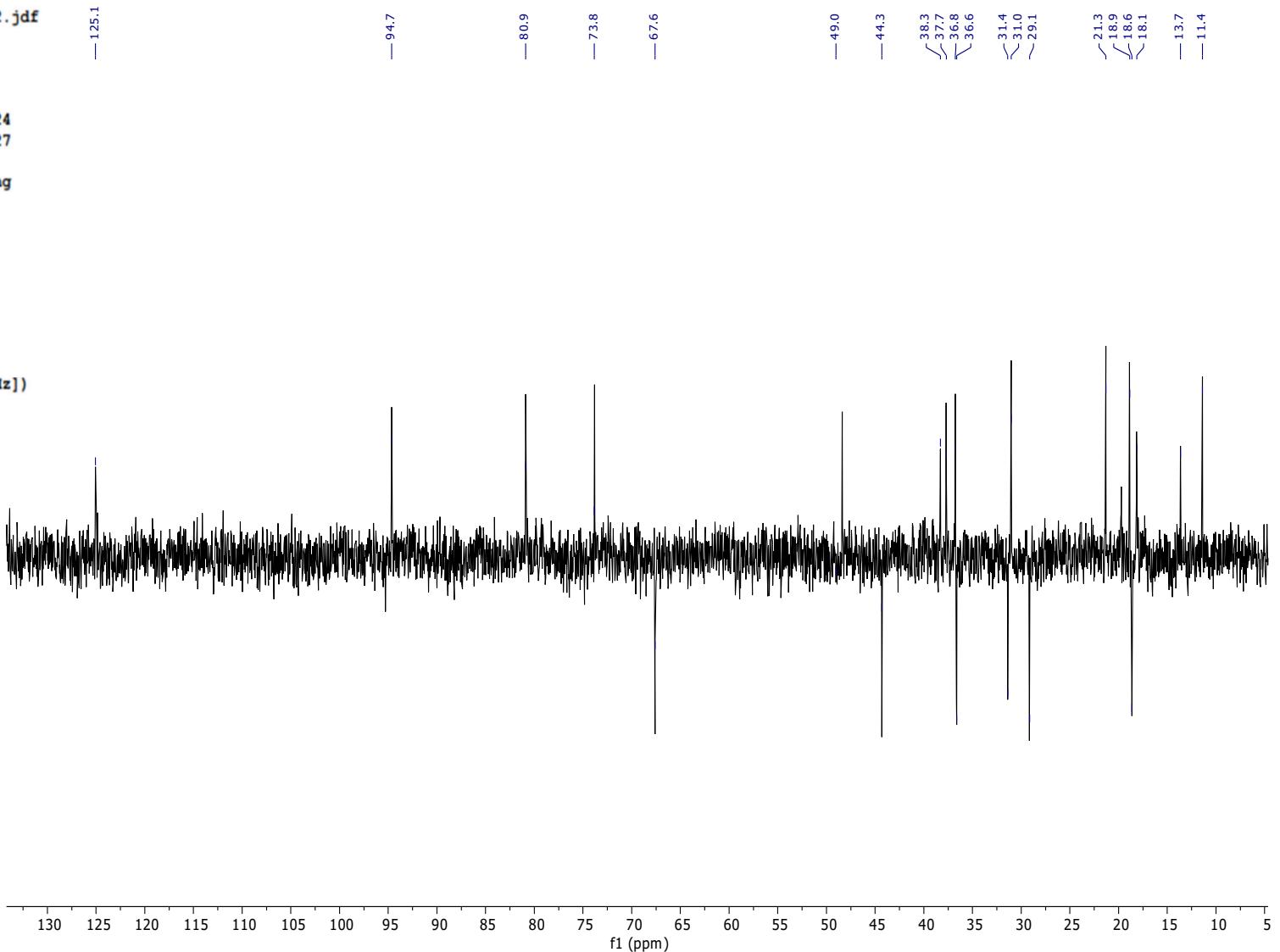


Figure S12. Asbestinin 27, COSY spectrum

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Filename      = JF1_65_05_cosy-2.jdf
Author        = DELTA
Experiment    = cosy_pfg_s.exp
Sample_Id     = I3_68_02
Solvent       = CHLOROFORM-D
Actual_Start_Time = 18-AUG-2015 04:45:39
Revision_Time = 14-JAN-2020 22:44:12

Comment       = gradient absolute value
Data_Format   = 2D REAL REAL
Dim_Size      = 1024, 1024
X_Domain     = 1H
Y_Domain     = 1H
Dim_Title    = 1H 1H
Dim_Units    = [ppm] [ppm]
Dimensions   = X Y
Site          = Eclipse+ 400
Spectrometer = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 0.2900992[s]
X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 3.62959[ppm]
X_Points       = 1024
X_Prescans    = 4
X_Resolution  = 3.44709672[Hz]
X_Sweep        = 3.52982704[kHz]
Y_Domain      = 1H
Y_Freq         = 399.78219838[MHz]
Y_Offset       = 3.62959[ppm]
Y_Points       = 256
Y_Prescans    = 0
Y_Resolution  = 13.78838687[Hz]
Y_Sweep        = 3.52982704[kHz]
Clipped       = FALSE
Scans          = 8
Total_Scans   = 2048

Relaxation_Delay = 1[s]
Recvr_Gain       = 18
Temp_Get          = 23.6[dC]
X_90_Width        = 9.81[us]
X_Acq_Time        = 0.2900992[s]
X_Pulse           = 9.81[us]
Y_Acq_Time        = 72.5248[ms]
Grad_1            = 1[ms]
Grad_1_Amp        = 5[pnt]
Grad_1_Value      = 5[pnt*ms]
Grad_2            = 1[ms]
Grad_2_Amp        = 5[pnt]
Grad_2_Value      = 5[pnt*ms]
Grad_Recover      = 1[ms]
Grad_Selection    = 1:1
Grad_Shape        = square
Grad_Type         = 0
Initial_Wait     = 0.1[s]
Pulse_1           = 9.81[us]
Pulse_2           = 9.81[us]
Pulse_Angle_1    = 90[deg]
Pulse_Angle_2    = 90[deg]
T1                = 1[us]
Unblank_Time     = 2[us]

```

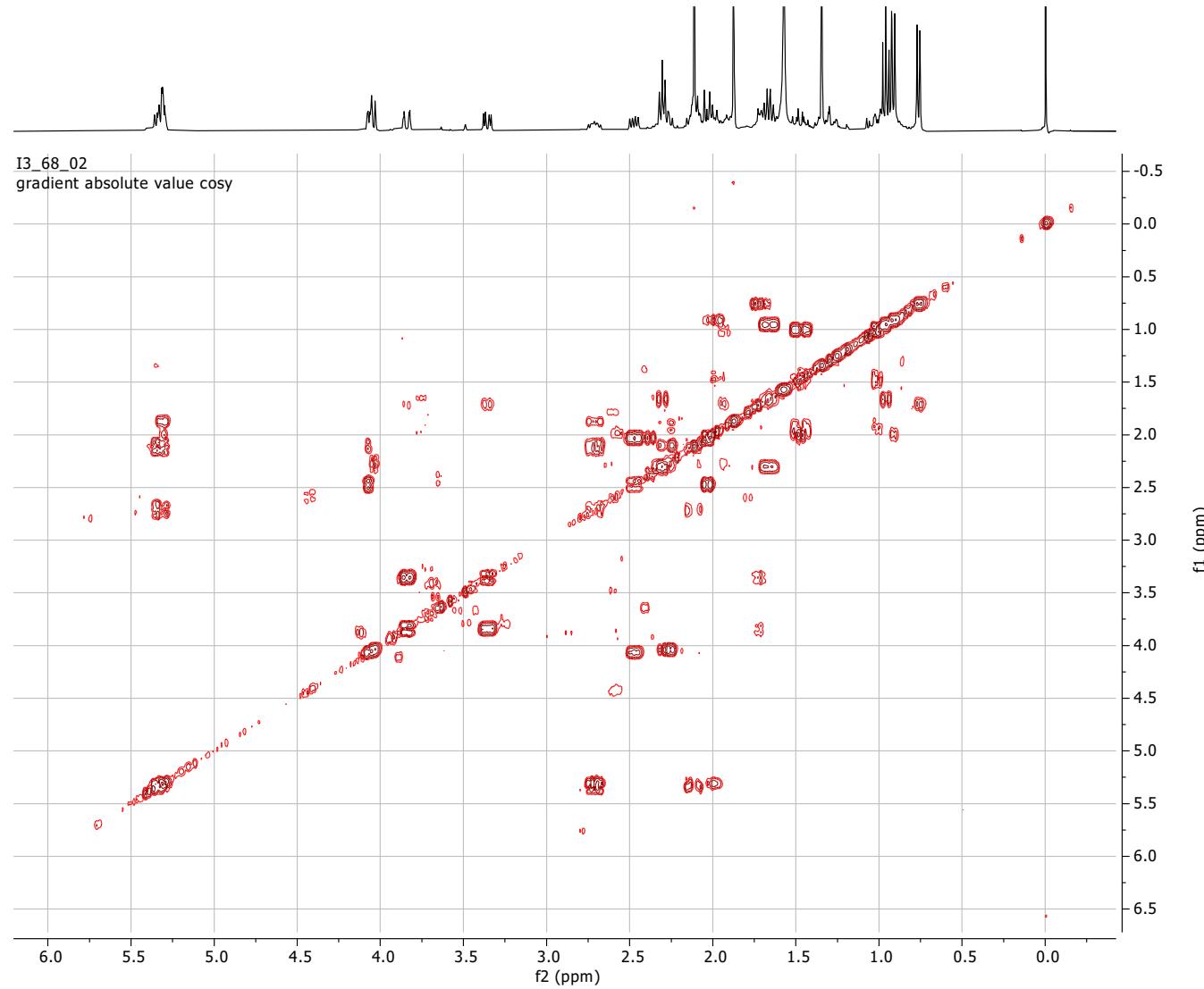


Figure S13. Asbestinin 27, HSQC spectrum

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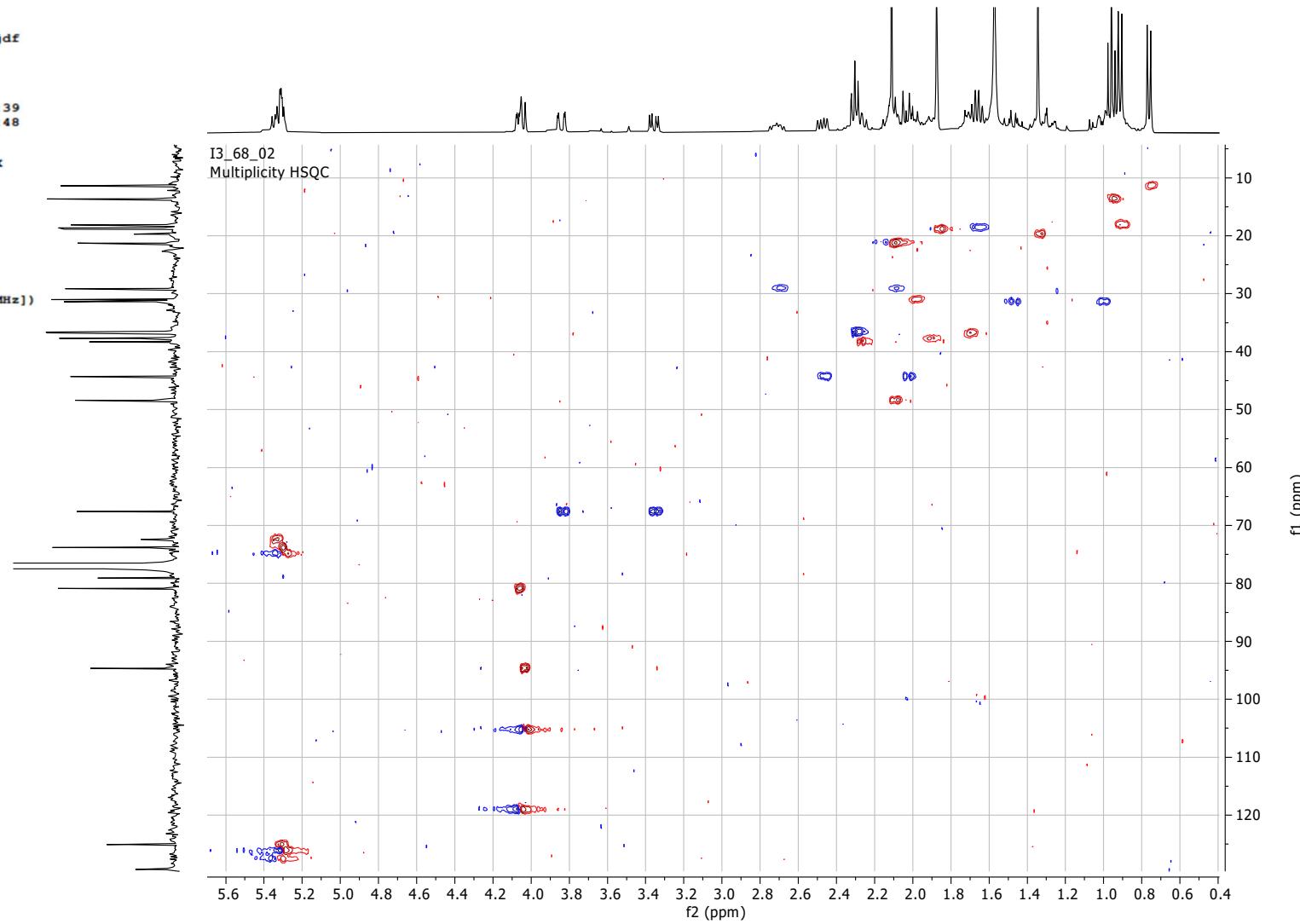
Filename      = JF1_65_05_hsqc-2.jdf
Author        = DELTA
Experiment    = multiplicity_hsq
Sample_Id     = I3_68_02
Solvent       = CHLOROFORM-D
Actual_Start_Time = 17-AUG-2015 17:02:39
Revision_Time  = 14-JAN-2020 22:50:48

Comment       = Multiplicity HSQC
Data_Format   = 2D COMPLEX COMPLEX
Dim_Size      = 2048, 512
X_Domain     = 1H
Y_Domain     = 13C
Dim_Title     = 1H 13C
Dim_Units     = [ppm] [ppm]
Dimensions    = X Y
Site          = Eclipsed+ 400
Spectrometer  = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 0.2900992[s]
X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 3.62959[ppm]
X_Points       = 1024
X_Prescans    = 4
X_Resolution  = 3.44709672[Hz]
X_Sweep        = 3.52982704[kHz]
Y_Domain      = 13C
Y_Freq         = 100.52530333[MHz]
Y_Offset       = 100[ppm]
Y_Points       = 256
Y_Prescans    = 0
Y_Resolution  = 98.64267677[Hz]
Y_Sweep        = 25.25252525[kHz]
Clipped        = FALSE
Scans          = 16
Total_Scans   = 4096

Relaxation_Delay = 2[s]
Recvr_Gain      = 30
Temp_Get         = 24.7[dC]
X_Acq_Time      = 0.2900992[s]
X_Pulse          = 9.81[us]
Y_Acq_Time      = 10.1376[ms]
Y_Pulse          = 12[us]
Enhance_Temp    = 12
Enhancement     = 1/6J
Grad_1           = 1[ms]
Grad_1_Amp       = 4[pnt]
Grad_1_Value     = 4[pnt*ms]
Grad_2           = 1[ms]
Grad_2_Amp       = -1[pnt]
Grad_2_Value     = -1[pnt*ms]
Grad_3           = 1[ms]
Grad_3_Amp       = 1[pnt]
Grad_Recover    = 0.1[ms]
Grad_Selection  = 13C = 4:1
Grad_Shape       = square
Grad_Type        = 0
Initial_Wait    = 1[s]
J_Constant      = 140[Hz]
T1              = 1[us]
Tau              = 0.5952381[ms]

```



**Figure S14. Asbestinin 27, HMBC spectrum**

```

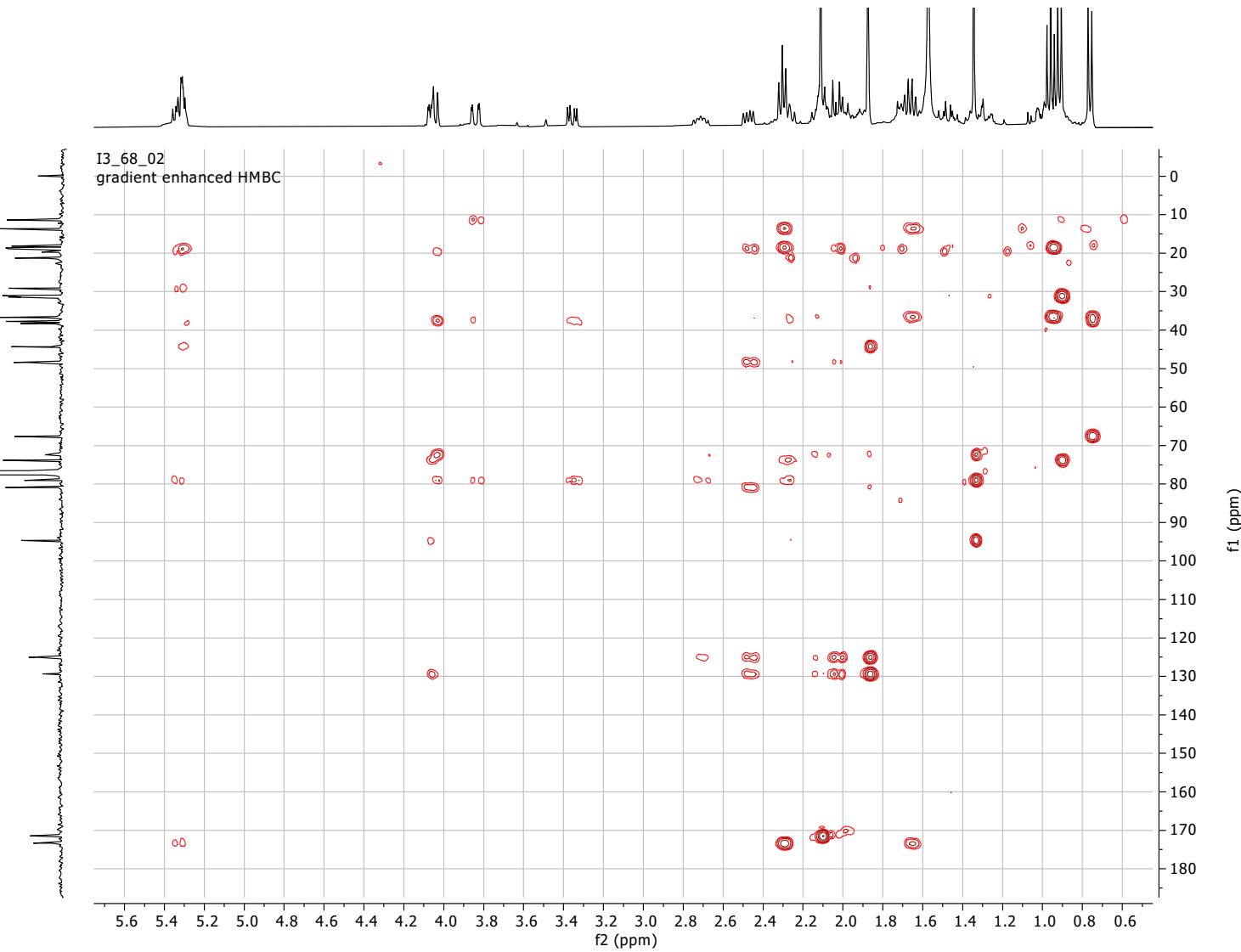
Filename      = JF1_65_05_hmboKOKOK-2.
Author        = DELTA
Experiment    = hmbo_pfg_s.exp
Sample_Id     = I3_68_02
Solvent       = CHLOROFORM-D
Actual_Start_Time = 17-AUG-2015 22:19:20
Revision_Time = 5-JUL-2019 17:15:31

Comment       = gradient enhanced HMBC
Data_Format   = 2D REAL REAL
Dim_Size      = 1024, 512
X_Domain     = 1H
Y_Domain     = 13C
Dim_Title    = 1H 13C
Dim_Units    = [ppm] [ppm]
Dimensions   = X Y
Site          = Eclipse+ 400
Spectrometer = DELTA_NMR

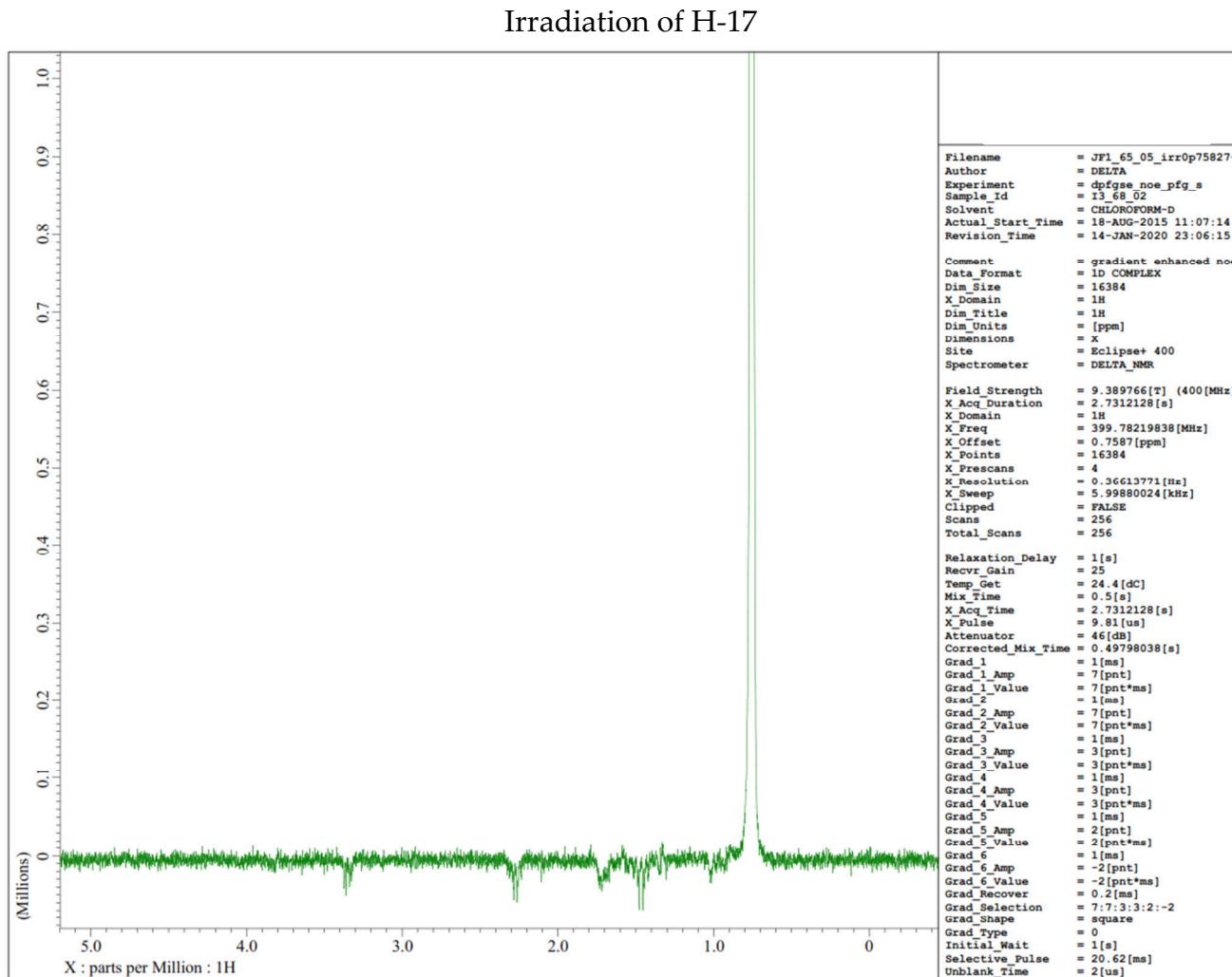
Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 0.2900992[s]
X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 3.62959[ppm]
X_Points       = 1024
X_Prescans    = 4
X_Resolution  = 3.44709672[Hz]
X_Sweep        = 3.52982704[kHz]
Y_Domain      = 13C
Y_Freq         = 100.52530333[MHz]
Y_Offset       = 100[ppm]
Y_Points       = 256
Y_Prescans    = 0
Y_Resolution  = 98.64267677[Hz]
Y_Sweep        = 25.25252525[kHz]
Clipped       = FALSE
Scans          = 32
Total_Scans   = 8192

Relaxation_Delay = 2[s]
Recvr_Gain      = 30
Temp_Get         = 23.9[dC]
X_Acq_Time      = 0.2900992[s]
X_Pulse          = 9.81[us]
Y_Acq_Time      = 10.1376[ms]
Y_Pulse          = 12[us]
Grad_1           = 1[ms]
Grad_1_Amp       = 10[pnt]
Grad_1_Value     = 10[pnt*ms]
Grad_2           = 1[ms]
Grad_2_Amp       = 10[pnt]
Grad_2_Value     = 10[pnt*ms]
Grad_3           = 1[ms]
Grad_3_Amp       = 5[pnt]
Grad_3_Value     = 5[pnt*ms]
Grad_Recover    = 0.2[ms]
Grad_Recover    = 13C = 2:2:1
Grad_Selection   = square
Grad_Type        = 0
Initial_Wait    = 1[s]
J_Constant      = 140[Hz]
Long_Range_J    = 8[Hz]
T1              = 1[us]
Unblank_Time    = 2[us]

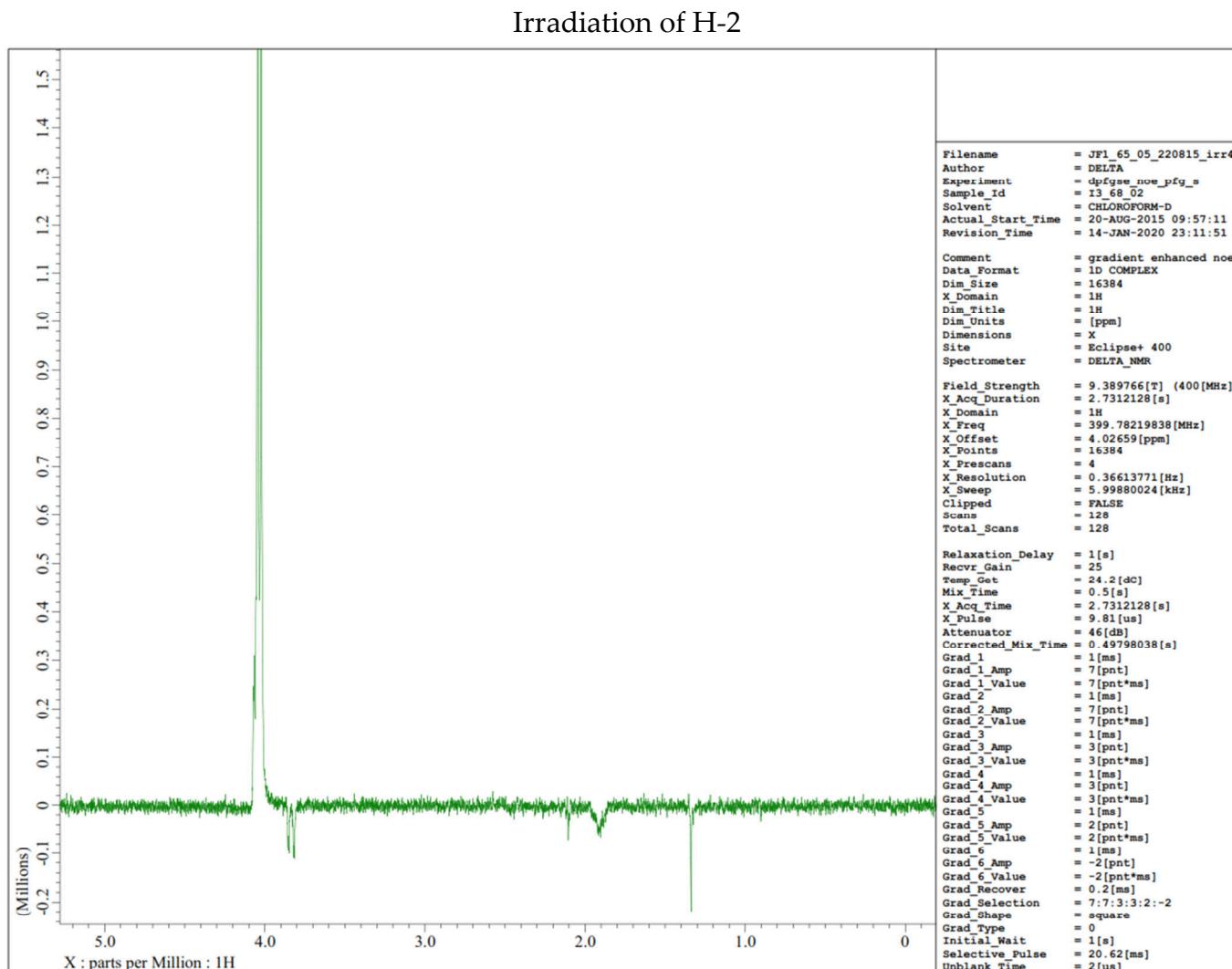
```



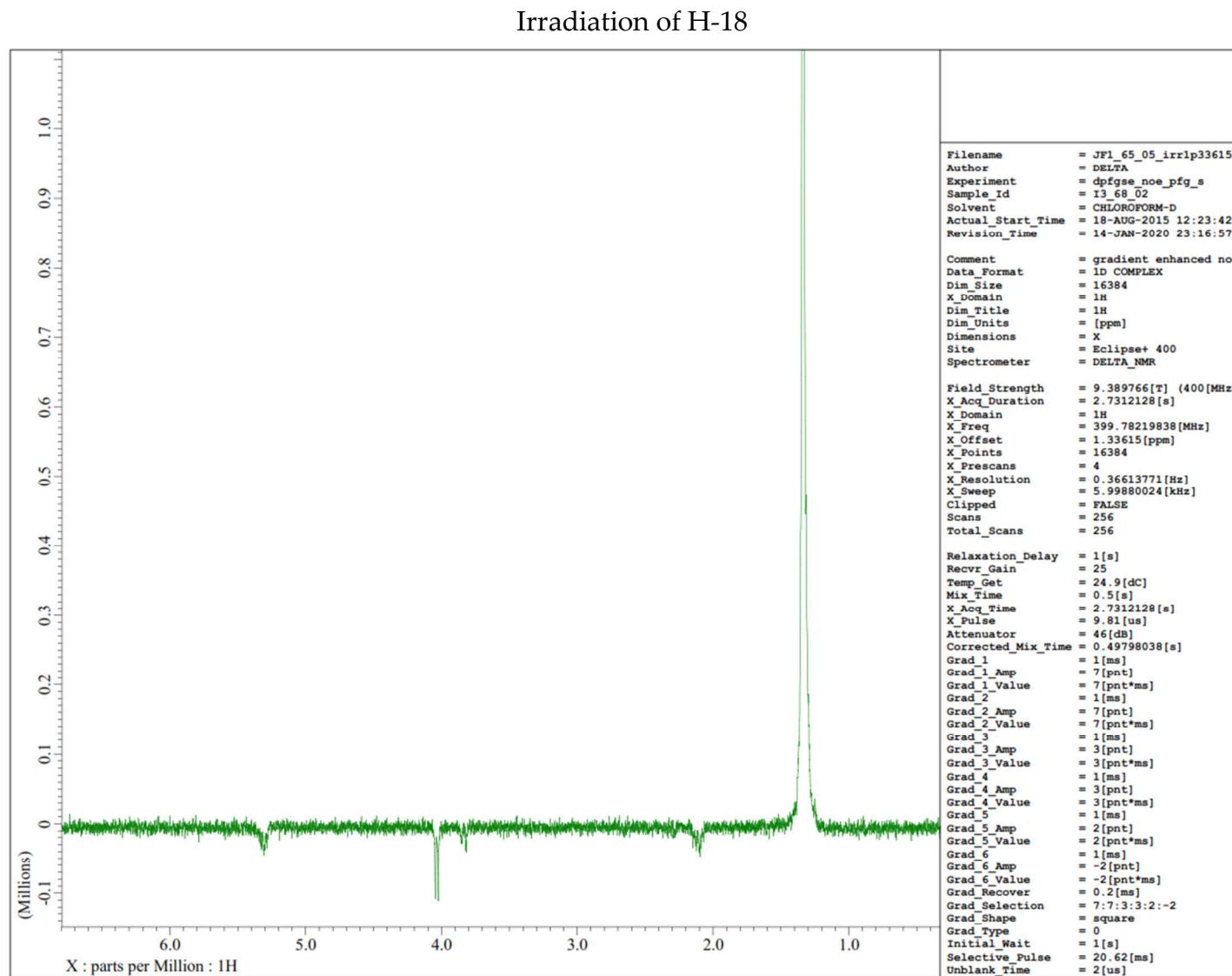
**Figure S15. Asbestinin 27, NOE spectra**



**Figure S15. Asbestinin 27, NOE spectra**

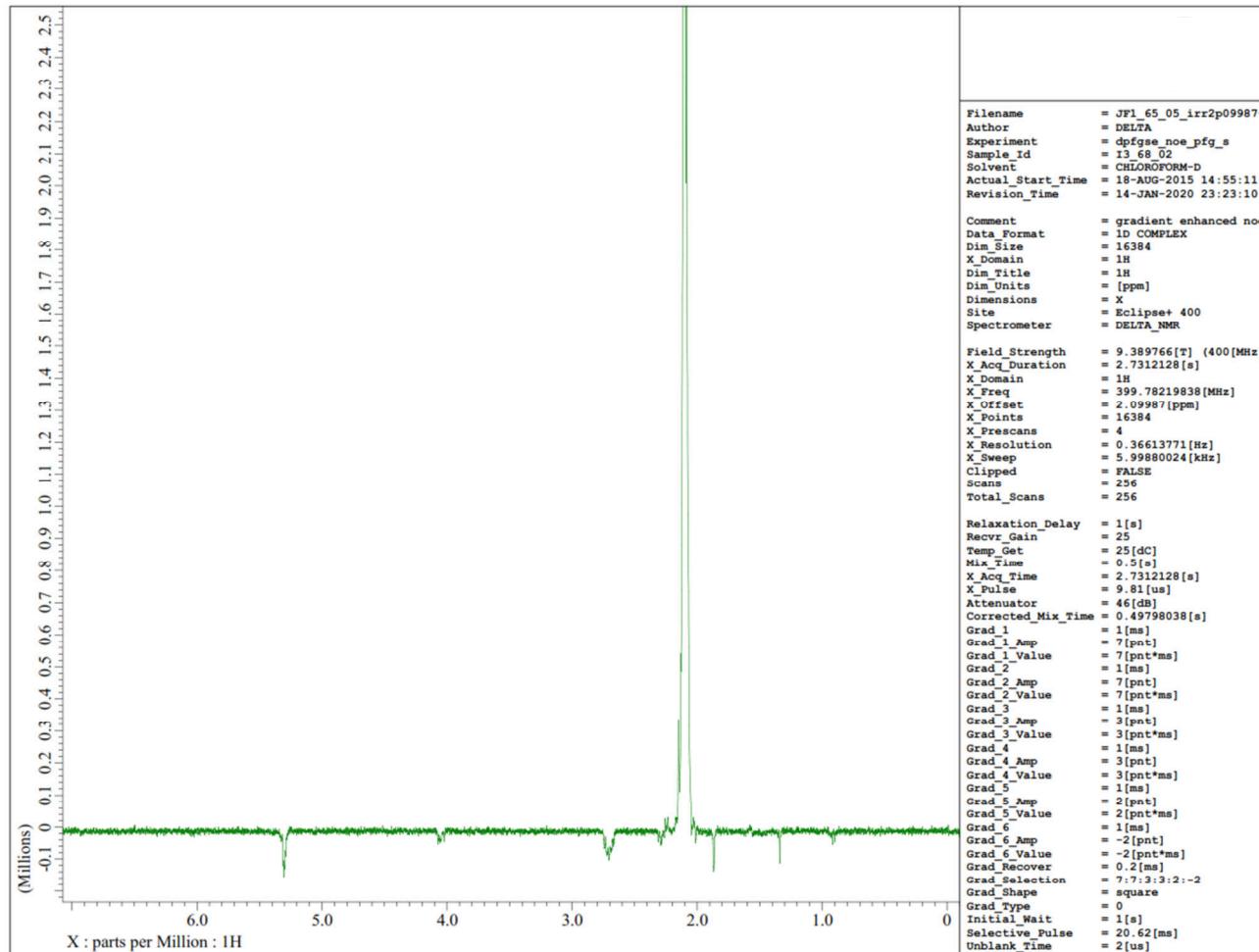


**Figure S15. Asbestinin 27, NOE spectra**

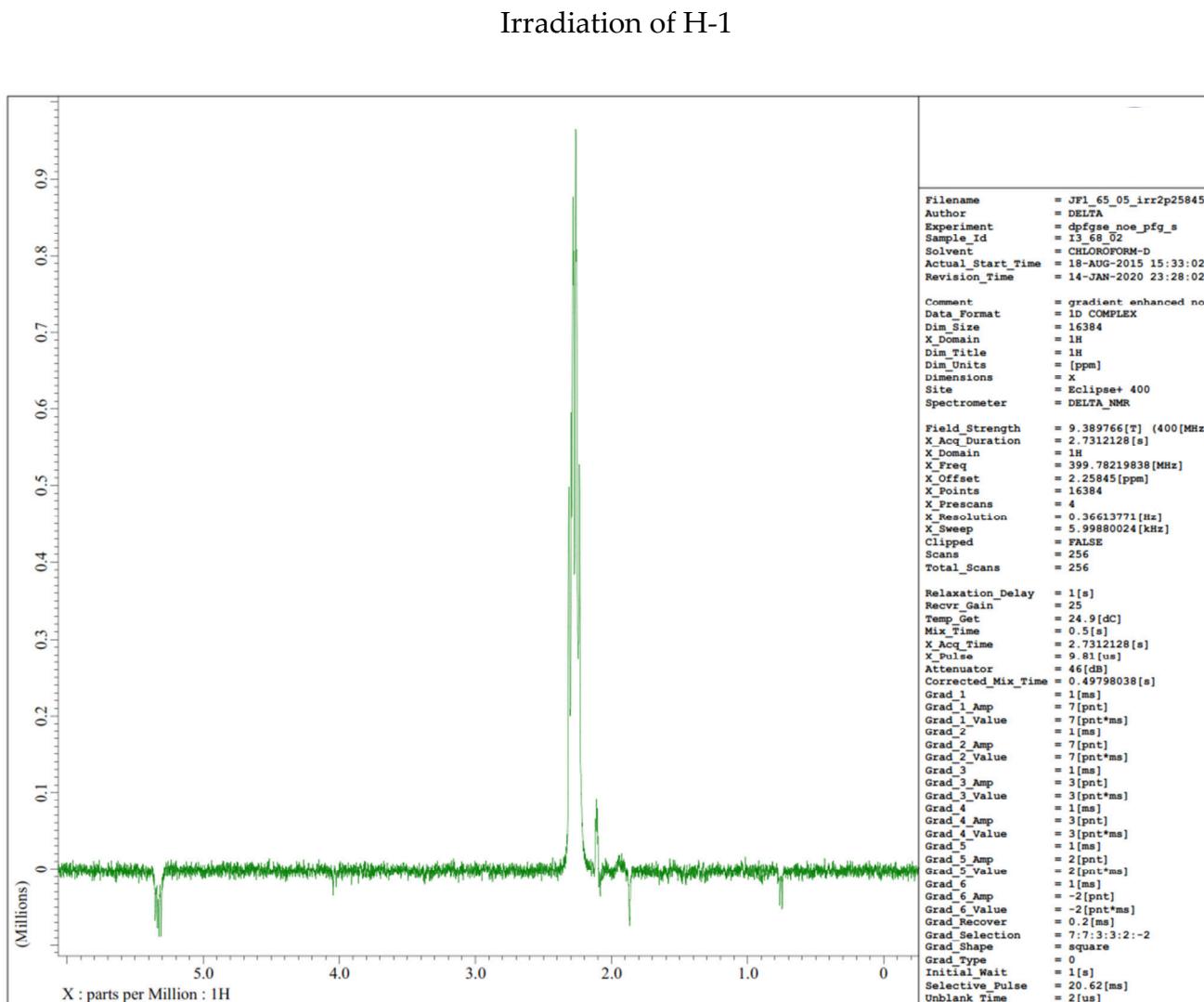


**Figure S15. Asbestinin 27, NOE spectra**

Irradiation of H-10

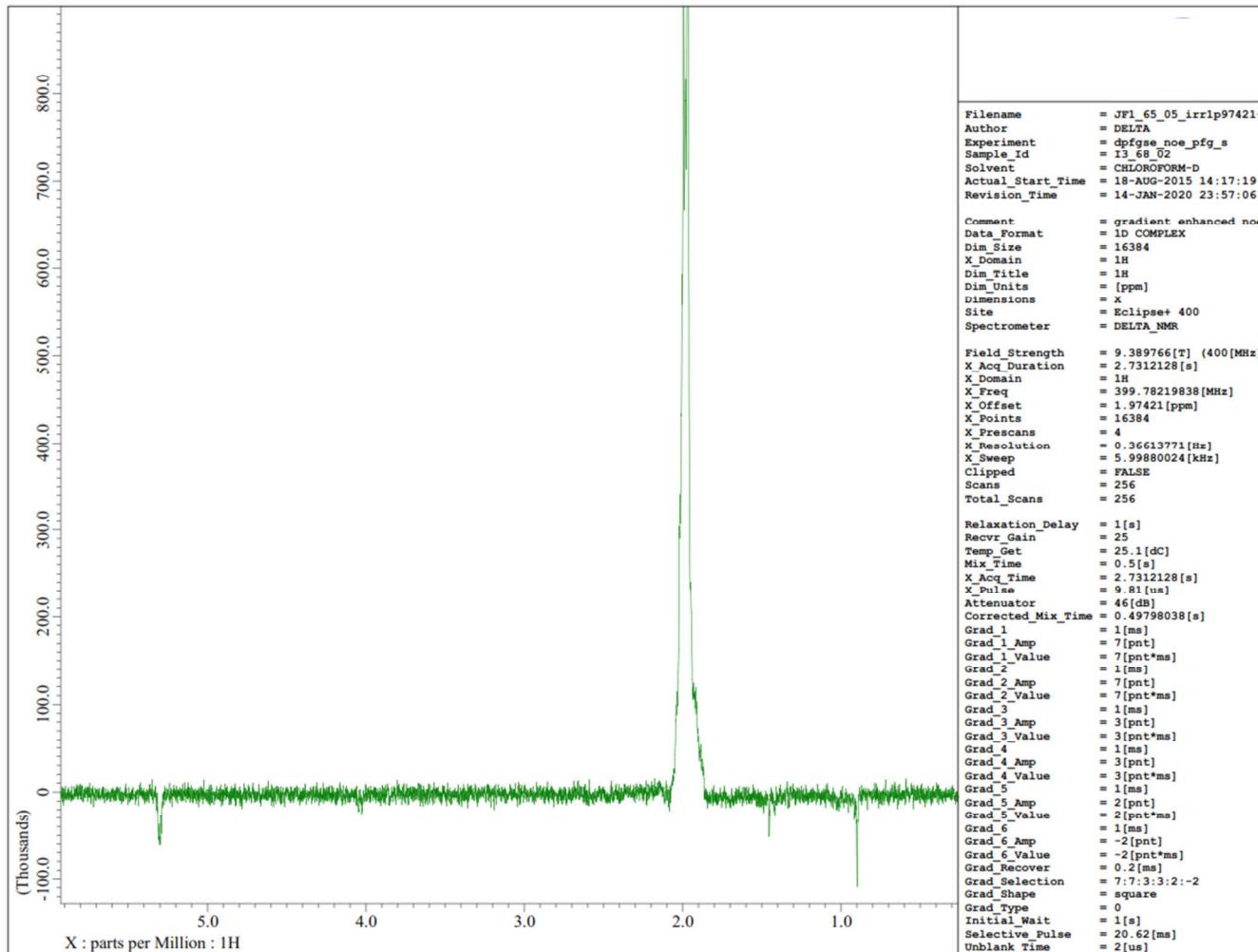


**Figure S15. Asbestinin 27, NOE spectra**

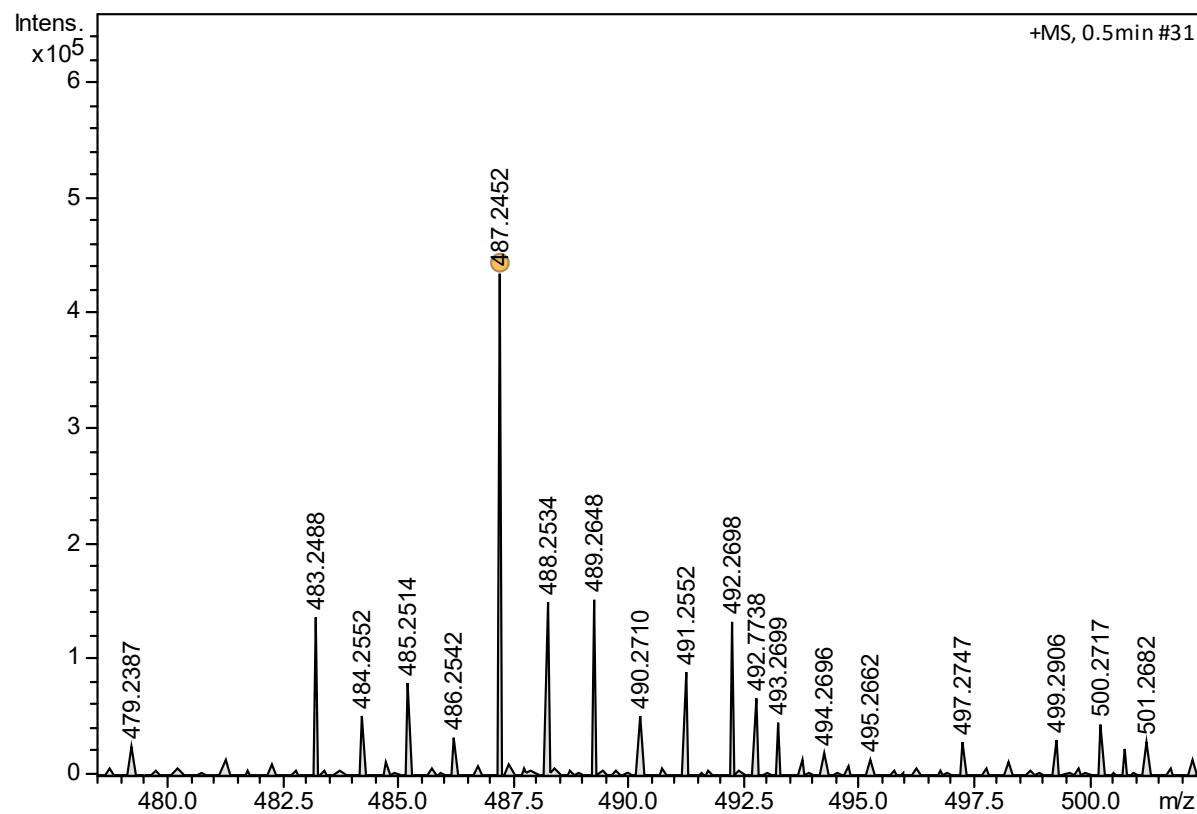


**Figure S15. Asbestinin 27, NOE spectra**

Irradiation of H-12



**Figure S16. Asbestinin 27, HR-ESITOFMS spectra**



**Figure S17. Asbestinin 28,  $^1\text{H}$  NMR spectrum**

```

Filename      = JF1_65_09-4.jdf
Author        = DELTA
Experiment    = single_pulse.exp
Sample_Id     = I3_62_05
Solvent       = CHLOROFORM-D
Actual_Start_Time = 3-AUG-2015 18:05:47
Revision_Time = 22-AUG-2015 07:42:29

Comment       = Single Pulse Experiment,
Data_Format   = 1D COMPLEX
Dim_Size      = 16384
X_Domain     = 1H
Dim_Title     = 1H
Dim_Units     = [ppm]
Dimensions    = X
Site          = Eclipse+ 400
Spectrometer  = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 2.048[s]
X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 7[ppm]
X_Points       = 16384
X_Prescans    = 0
X_Resolution   = 0.48828125[Hz]
X_Sweep        = 8[kHz]
Clipped        = FALSE
Scans          = 8
Total_Scans    = 8

Relaxation_Delay = 4[s]
Recvr_Gain      = 13
Temp_Get         = 24.7[dC]
X_90_Width      = 9.81[us]
X_Acq_Time      = 2.048[s]
X_Angle          = 45[deg]
X_Pulse          = 4.905[us]
Initial_Wait    = 1[s]
Unblank_Time    = 2[us]

```

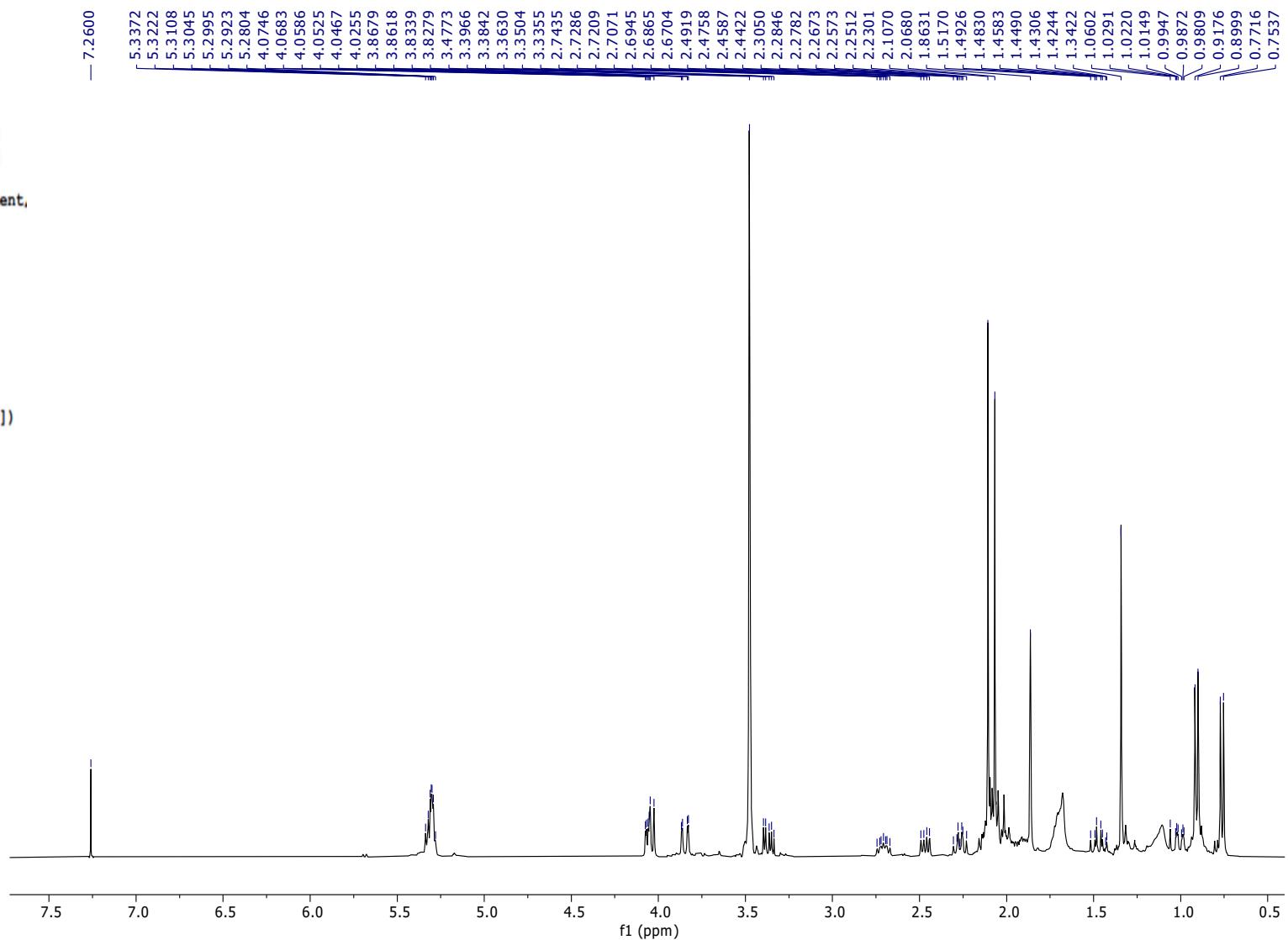
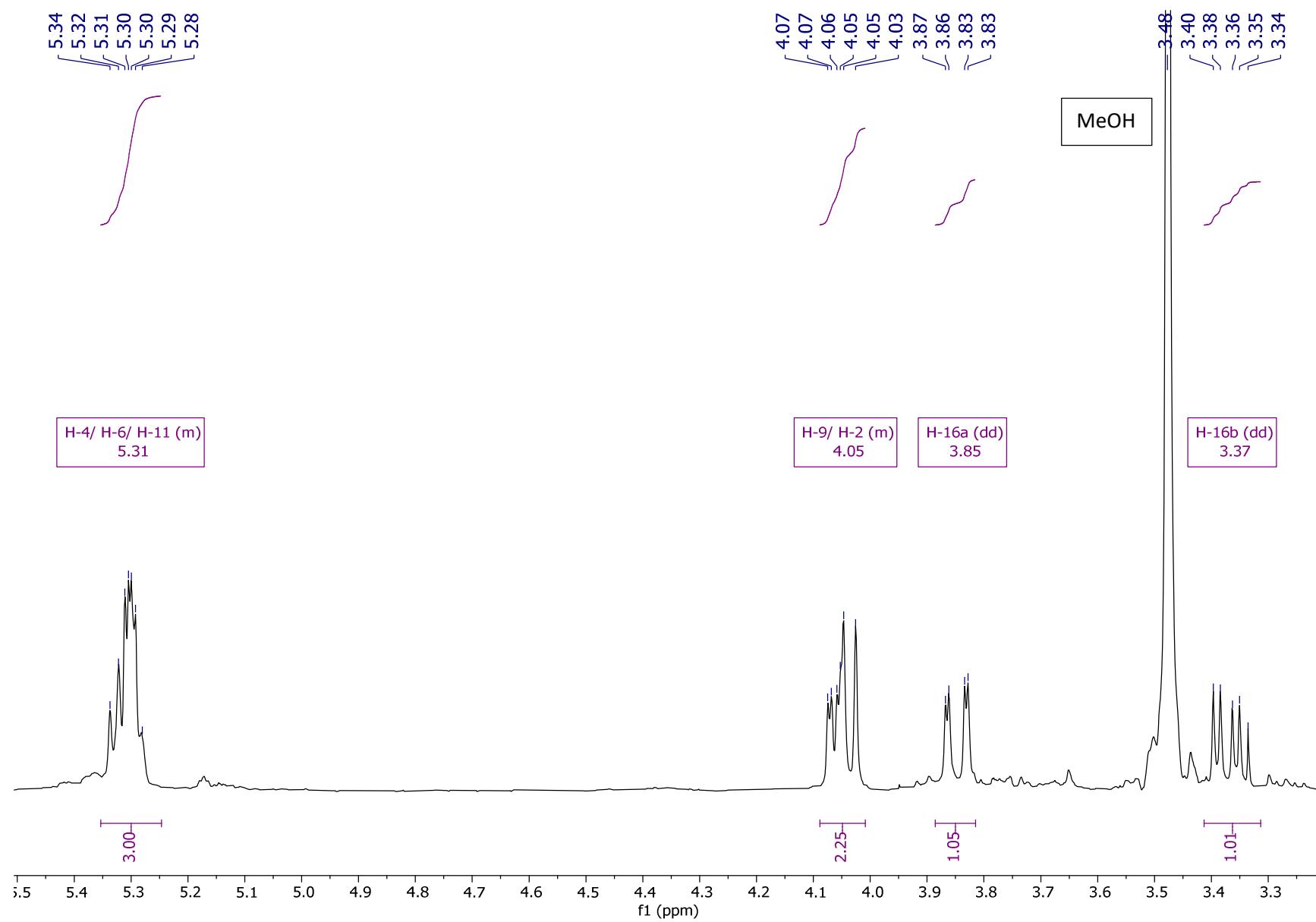
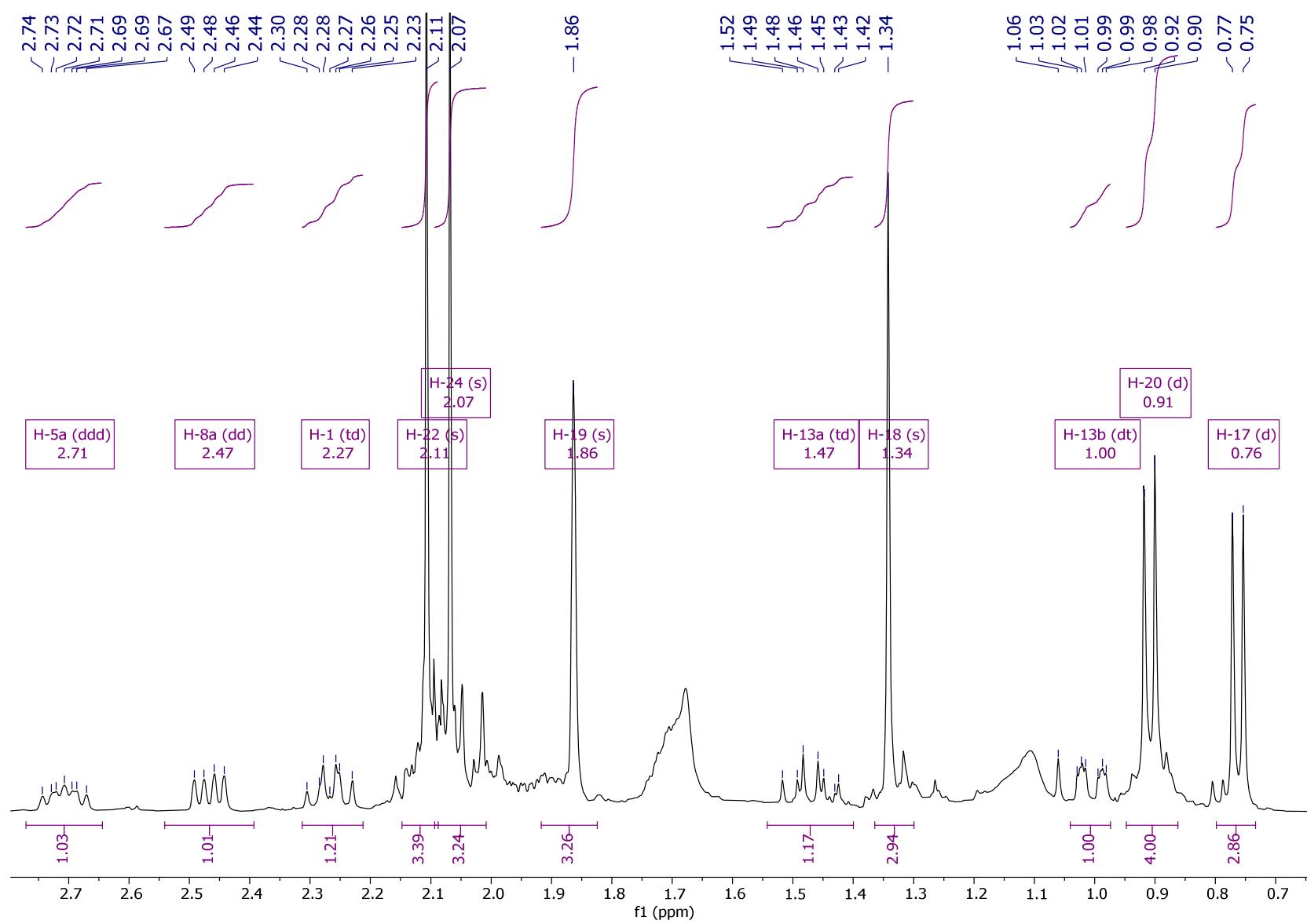


Figure S17. Asbestinin 28,  $^1\text{H}$  NMR spectrum (expanded)



**Figure S17. Asbestinin 28,  $^1\text{H}$  NMR spectrum (expanded)**



**Figure S18. Asbestinin 28,  $^{13}\text{C}$  NMR spectrum**

```

Filename      = JF1_65_09 C13-3.jdf
Author        = DELTA
Experiment    = single_pulse_dec
Sample_Id     = I3_62_05
Solvent       = CHLOROFORM-D
Actual_Start_Time = 3-AUG-2015 18:13:56
Revision_Time = 5-AUG-2015 22:35:58

Comment       = Single Pulse with Broad
Data_Format   = 1D COMPLEX
Dim_Size      = 32768
X_Domain     =  $^{13}\text{C}$ 
Dim_Title    =  $^{13}\text{C}$ 
Dim_Units     = [ppm]
Dimensions    = X
Site          = Eclipsit 400
Spectrometer  = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 1.3008896[s]
X_Domain      =  $^{13}\text{C}$ 
X_Freq         = 100.52530333[MHz]
X_Offset       = 100[ppm]
X_Points       = 32768
X_Prescans    = 4
X_Resolution   = 0.76870474[Hz]
X_Sweep        = 25.18891688[kHz]
Irr_Domain    = 1H
Irr_Freq       = 399.78219838[MHz]
Irr_Offset     = 5[ppm]
Clipped        = FALSE
Scans          = 7168
Total_Scans    = 7168

Relaxation_Delay = 1[s]
Recvr_Gain      = 26
Temp_Get         = 25.9[dC]
X_90_Width      = 10[us]
X_Acq_Time      = 1.3008896[s]
X_Angle          = 30[deg]
X_Pulse          = 3.333333333[us]
Initial_Wait    = 1[s]
Unblank_Time    = 2[us]

```

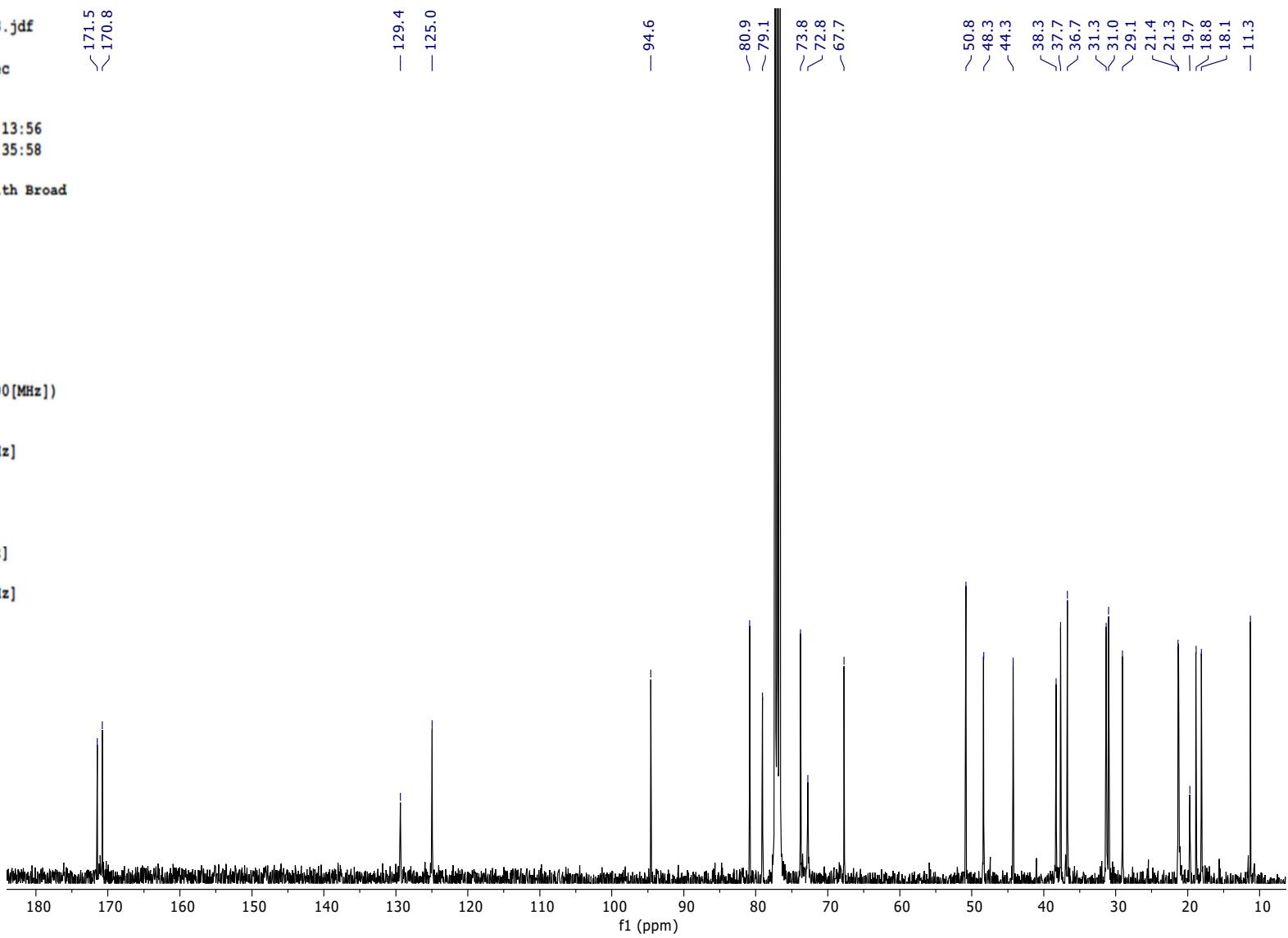


Figure S19. Asbestinin 28, DEPT-135 spectrum

```

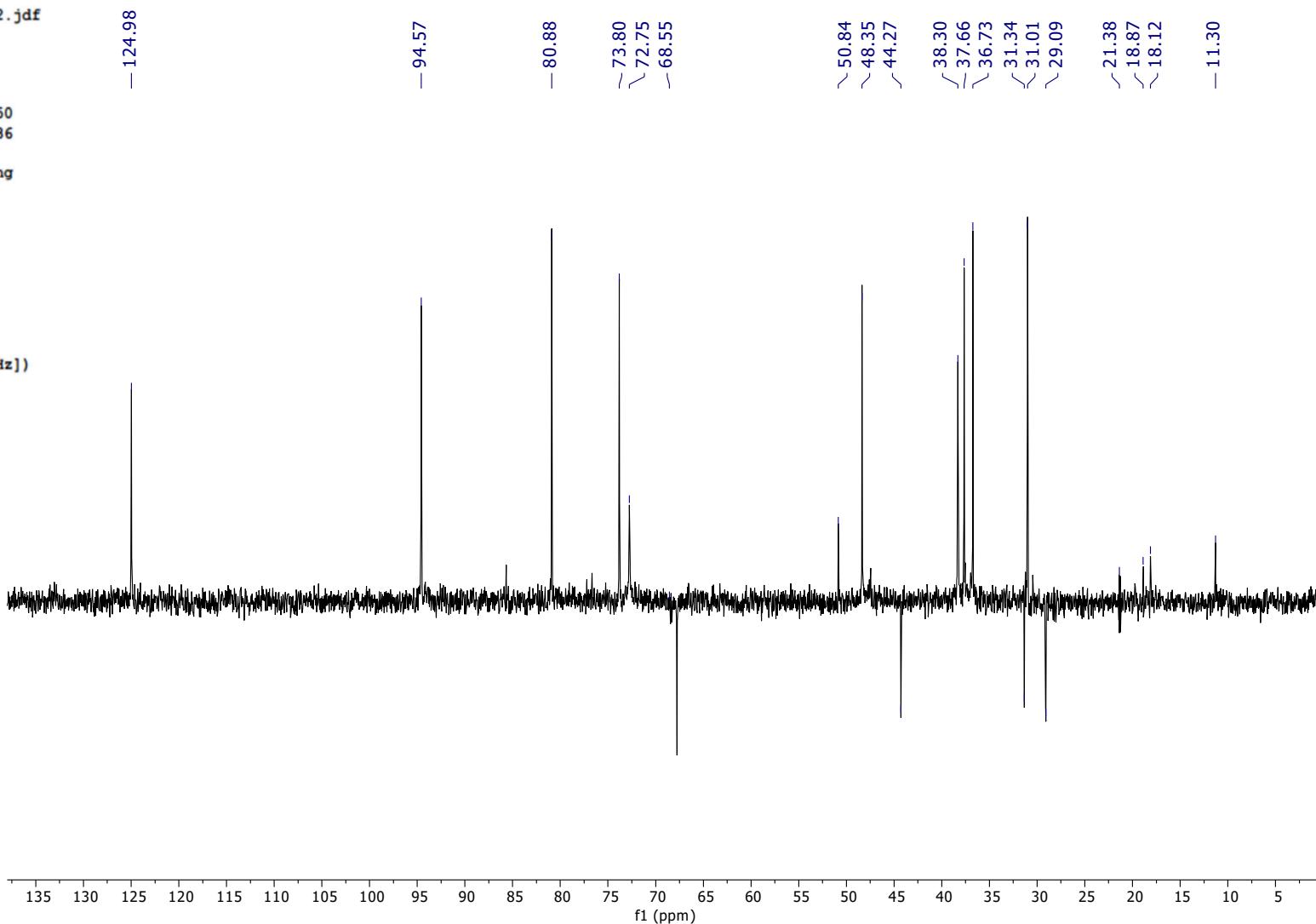
Filename      = JF1_65_09 dept135-2.jdf
Author        = DELTA
Experiment    = dept_dec.exp
Sample_Id     = I3_62_05
Solvent       = CHLOROFORM-D
Actual_Start_Time = 3-AUG-2015 22:49:50
Revision_Time  = 6-AUG-2015 01:26:36

Comment       = DEPT with decoupling
Data_Format   = 1D COMPLEX
Dim_Size      = 32768
X_Domain     = 13C
Dim_Title     = 13C
Dim_Units     = [ppm]
Dimensions    = X
Site          = Eclipse+ 400
Spectrometer  = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acc_Duration = 1.3008896[s]
X_Domain     = 13C
X_Freq        = 100.52530333[MHz]
X_Offset      = 100[ppm]
X_Points      = 32768
X_Prescans   = 4
X_Resolution  = 0.76870474[Hz]
X_Sweep       = 25.18891688[kHz]
Irr_Domain   = 1H
Irr_Freq      = 399.78219838[MHz]
Irr_Offset    = 5[ppm]
Clipped       = FALSE
Scans         = 3072
Total_Scans   = 3072

Relaxation_Delay = 2[s]
Recvr_Gain      = 26
Temp_Get         = 24.9[dC]
X_Acc_Time      = 1.3008896[s]
X_Pulse         = 10[us]
Irr_Pulse       = 44.5[us]
Initial_Wait    = 1[s]
J_Constant      = 140[Hz]
Selection_Angle = 135[deg]
Selection_Pulse = 66.75[us]
Unblank_Time    = 2[us]

```



**Figure S20. Asbestinin 28, COSY spectrum**

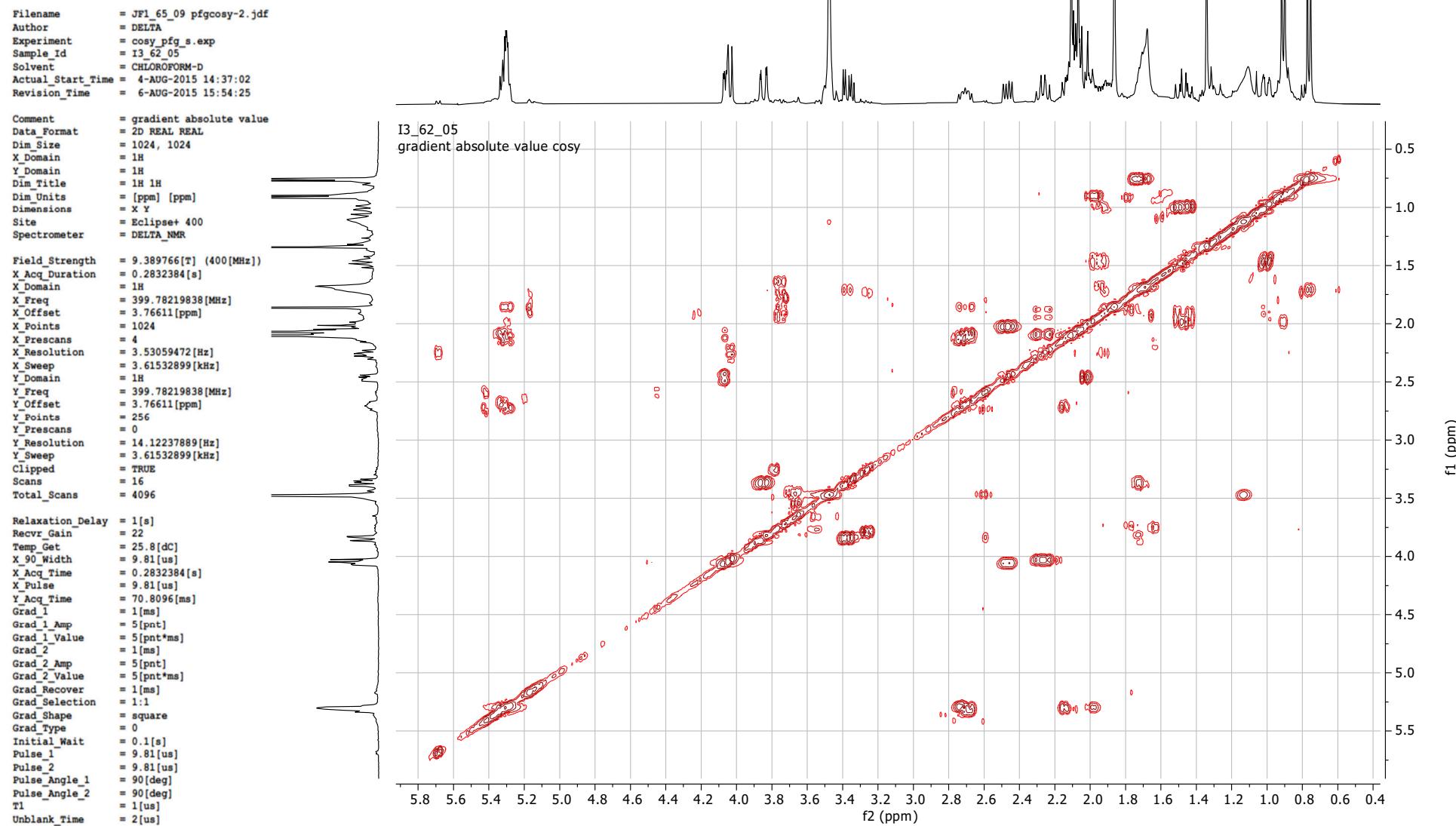


Figure S21. Asbestinin 28, HSQC spectrum

```

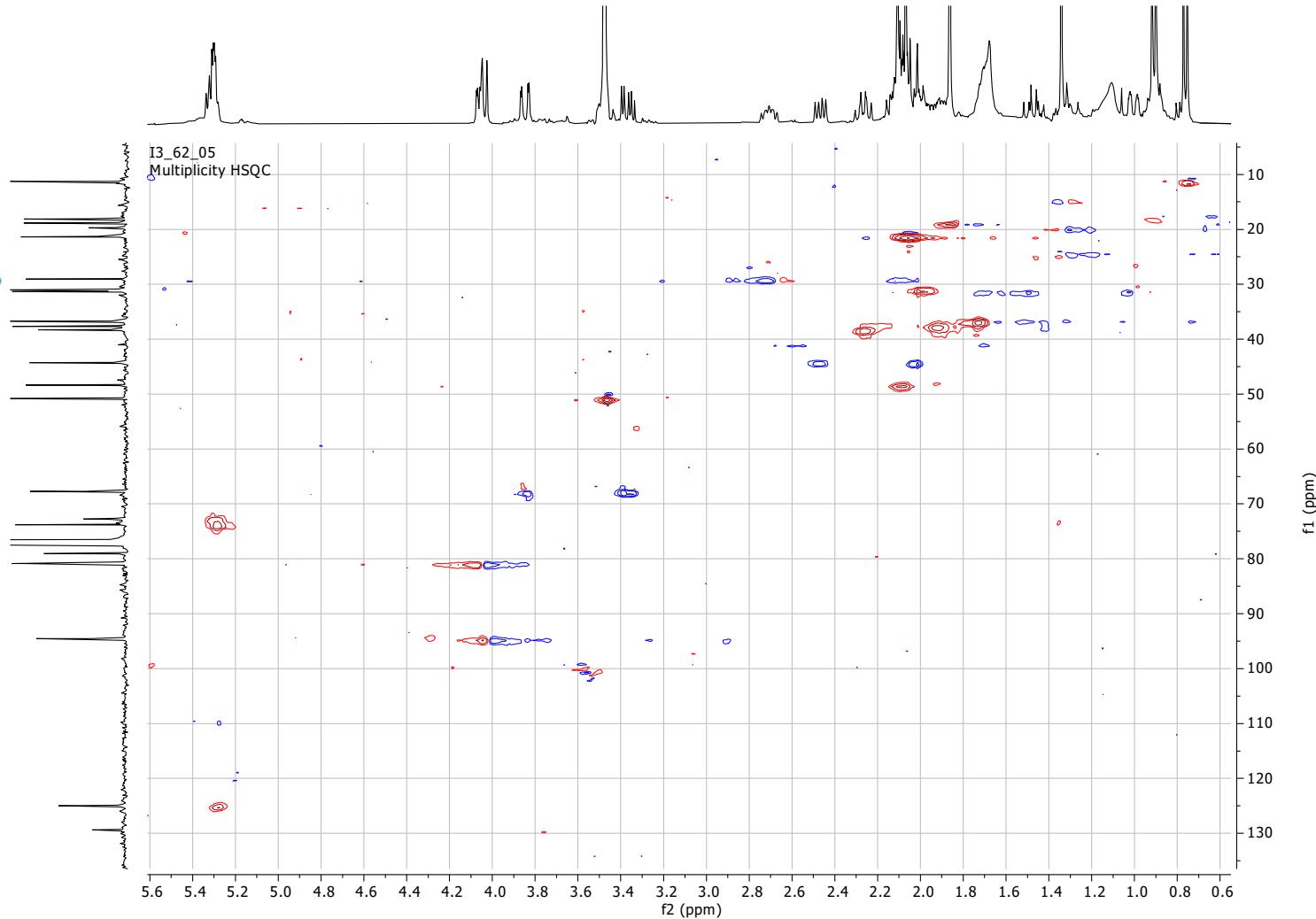
Filename      = JF1_65_09 hsqc-4.jdf
Author        = DELTA
Experiment    = multiplicity_hsq
Sample_Id     = I3_62_05
Solvent       = CHLOROFORM-D
Actual_Start_Time = 4-AUG-2015 04:31:29
Revision_Time = 21-AUG-2015 06:18:42

Comment       = Multiplicity HSQC
Format        = 2D COMPLEX COMPLEX
Dim_Size      = 1024, 512
X_Domain     = 1H
Y_Domain     = 13C
Dim_Title    = 1H 13C
Dim_Units    = [ppm] [ppm]
Dimensions   = X Y
Site          = Ecliptet 400
Spectrometer = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 0.2832384[s]
X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 3.76611[ppm]
X_Points       = 1024
X_Prescans    = 4
X_Resolution  = 3.53059472[Hz]
X_Sweep        = 3.61532899[kHz]
Y_Domain      = 13C
Y_Freq         = 100.52530333[MHz]
Y_Offset       = 100[ppm]
Y_Points       = 256
Y_Prescans    = 0
Y_Resolution  = 98.64267677[Hz]
Y_Sweep        = 25.25252525[kHz]
Clipped       = FALSE
Scans          = 8
Total_Scans   = 2048

Relaxation_Delay = 2[s]
Recvr_Gain       = 30
Temp_Get         = 24.1[dC]
X_Acq_Time       = 0.2832384[s]
X_Pulse          = 9.81[us]
Y_Acq_Time       = 10.1376[ms]
Y_Pulse          = 12[us]
Enhance_Temp    = 12
Enhancement     = 1/6J
Grad_1           = 1[ms]
Grad_1_Amp       = 4[pnt]
Grad_1_Value     = 4[pnt*ms]
Grad_2           = 1[ms]
Grad_2_Amp       = -1[pnt]
Grad_2_Value     = -1[pnt*ms]
Grad_3           = 1[ms]
Grad_3_Amp       = 1[pnt]
Grad_Recover    = 0.1[ms]
Grad_Selection  = 13C = 4:1
Grad_Shape       = square
Grad_Type        = 0
Initial_Wait    = 1[s]
J_Constant      = 140[Hz]
Tau              = 1[us]
Tau              = 0.5952381[ms]

```



**Figure S22. Asbestinin 28, HMBC spectrum**

```

Filename      = JF1_65_09_210815_hmhc-3
Author        = DELTA
Experiment    = hmhc_pfg_s.exp
Sample_Id     = I3_68_06
Solvent       = CHLOROFORM-D
Actual_Start_Time = 19-AUG-2015 15:04:21
Revision_Time = 22-AUG-2015 01:38:21

Comment       = gradient enhanced HMBC
Data_Format   = 2D REAL REAL
Dim_Size      = 1024, 512
X_Domain     = 1H
Y_Domain     = 13C
Dim_Title     = 1H 13C
Dim_Units     = [ppm] [ppm]
Dimensions    = X Y
Site          = Eclipse+ 400
Spectrometer  = DELTA_NMR

Field_Strength = 9.389766[T] (400[MHz])
X_Acq_Duration = 0.2961408[s]
X_Domain      = 1H
X_Freq         = 399.78219838[MHz]
X_Offset       = 3.59946[ppm]
X_Points       = 1024
X_Prescans    = 4
X_Resolution  = 3.37677213[Hz]
X_Sweep        = 3.45781466[kHz]
Y_Domain      = 13C
Y_Freq         = 100.52530333[MHz]
Y_Offset       = 100[ppm]
Y_Points       = 256
Y_Prescans    = 0
Y_Resolution  = 98.64267677[Hz]
Y_Sweep        = 25.25252525[kHz]
Clipped        = FALSE
Scans          = 64
Total_Scans   = 16384

Relaxation_Delay = 2[s]
Recvr_Gain       = 30
Temp_Get         = 23.5[dC]
X_Acq_Time       = 0.2961408[s]
X_Pulse          = 9.81[us]
Y_Acq_Time       = 10.1376[ms]
Y_Pulse          = 12[us]
Grad_1           = 1[ms]
Grad_1_Amp       = 10[pnt]
Grad_1_Value     = 10[pnt*ms]
Grad_2           = 1[ms]
Grad_2_Amp       = 10[pnt]
Grad_2_Value     = 10[pnt*ms]
Grad_3           = 1[ms]
Grad_3_Amp       = 5[pnt]
Grad_3_Value     = 5[pnt*ms]
Grad_Recover     = 0.2[ms]
Grad_Selection   = 13C = 2:2:1
Grad_Shape       = square
Grad_Type        = 0
Initial_Wait    = 1[s]
J_Constant       = 140[Hz]
Long_Range_J     = 8[Hz]
T1               = 1[us]
Unblank_Time    = 2[us]

```

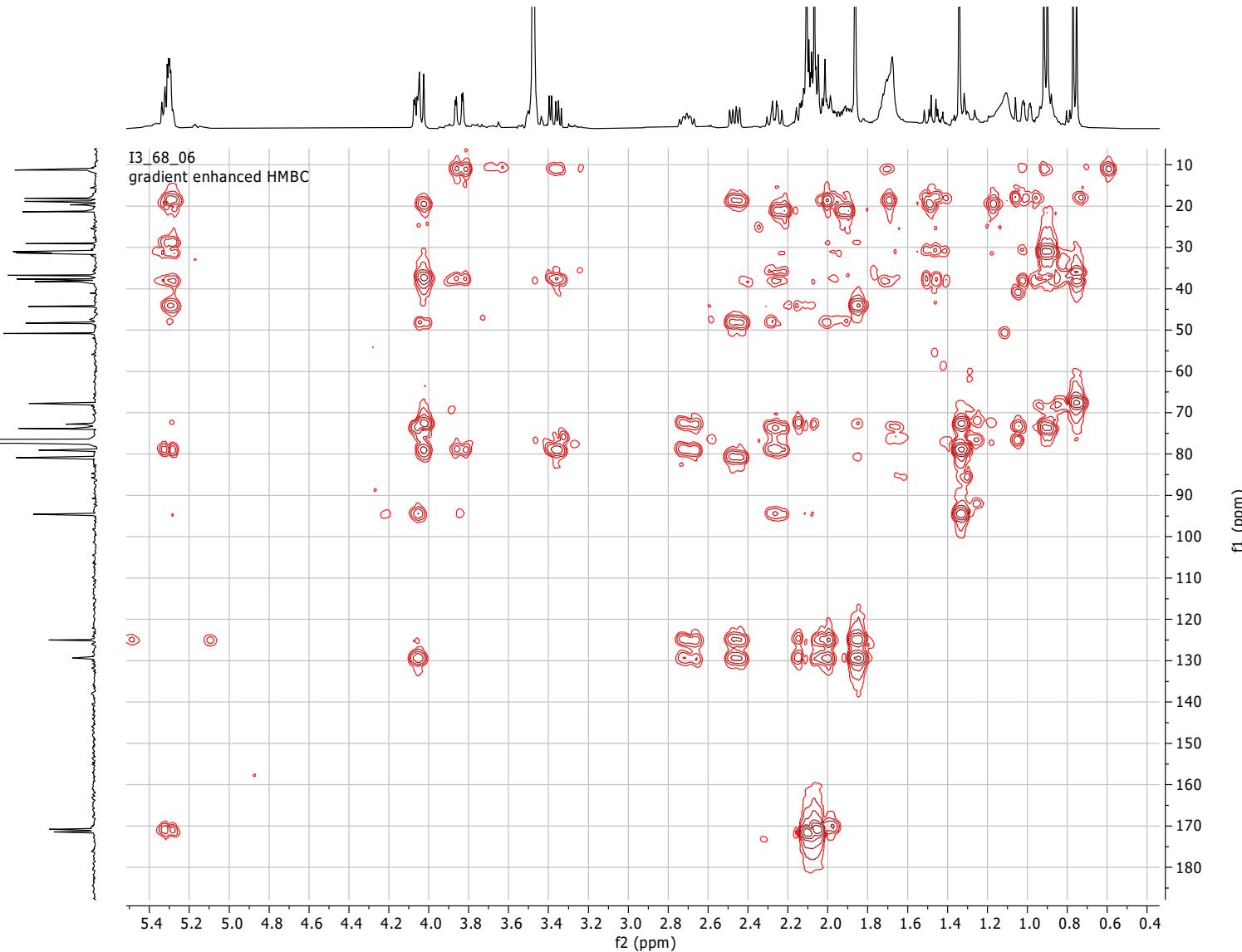


Figure S23. Asbestinin 28, NOE spectra

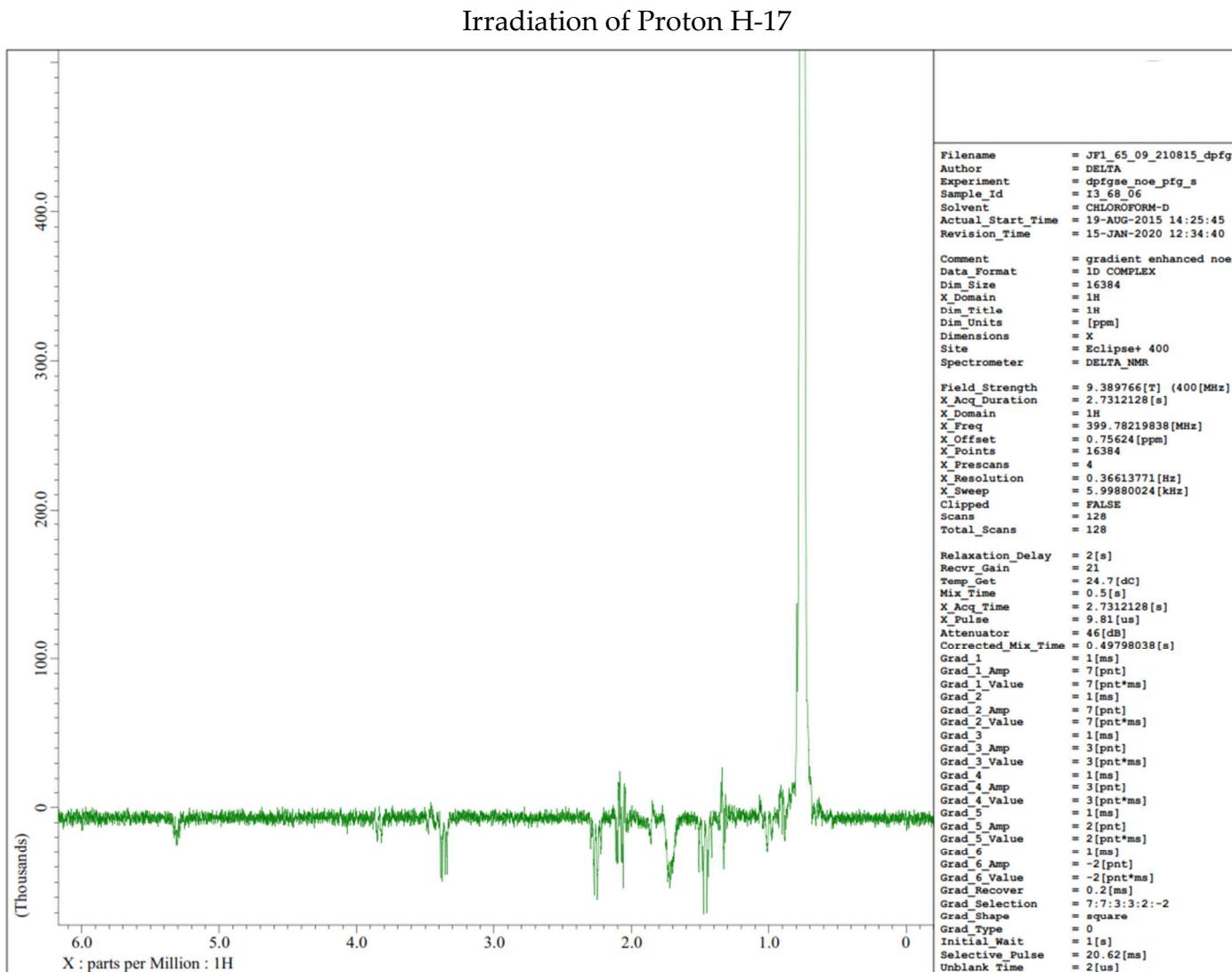


Figure S23. Asbestinin 28, NOE spectra

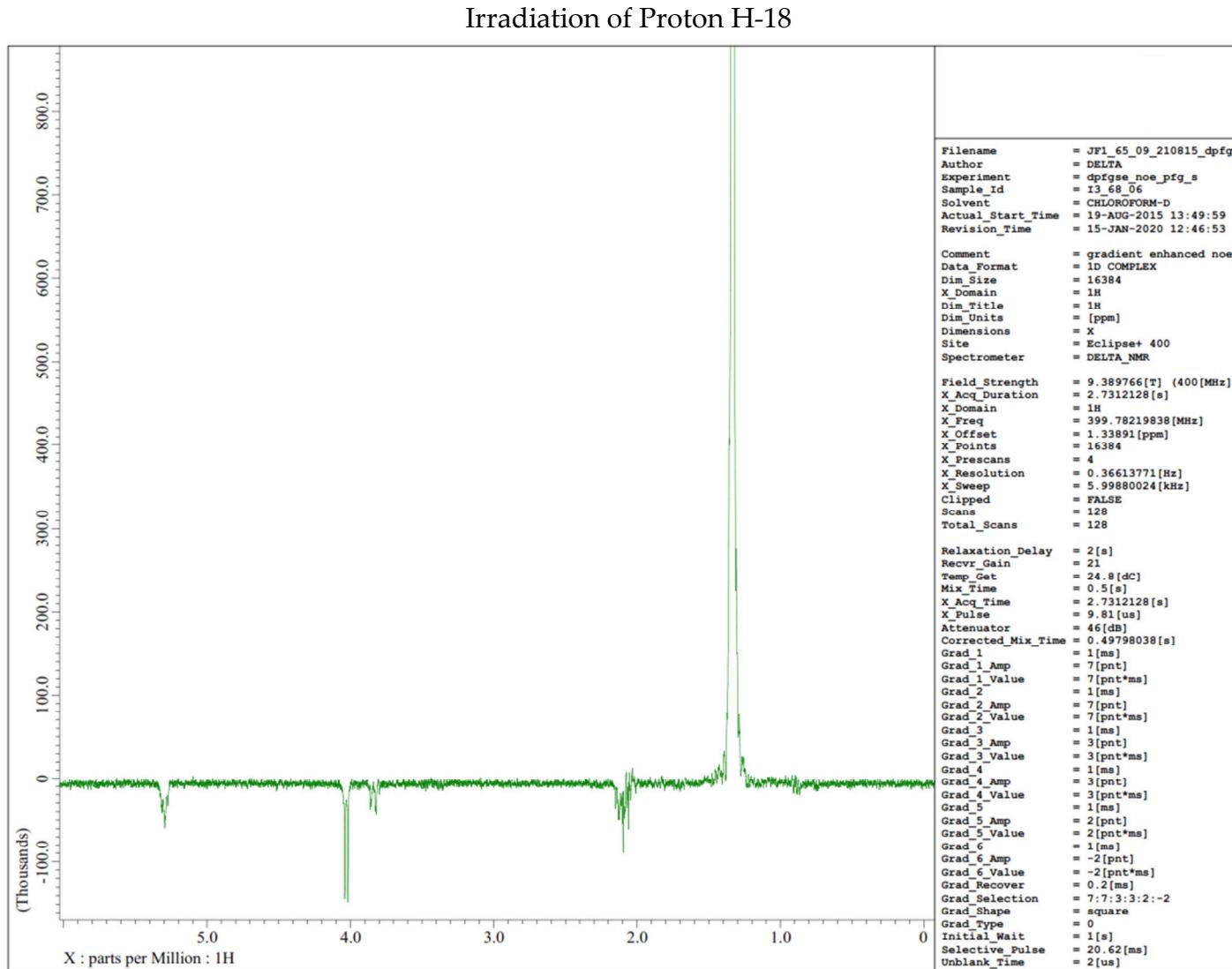


Figure S23. Asbestinin 28, NOE spectra

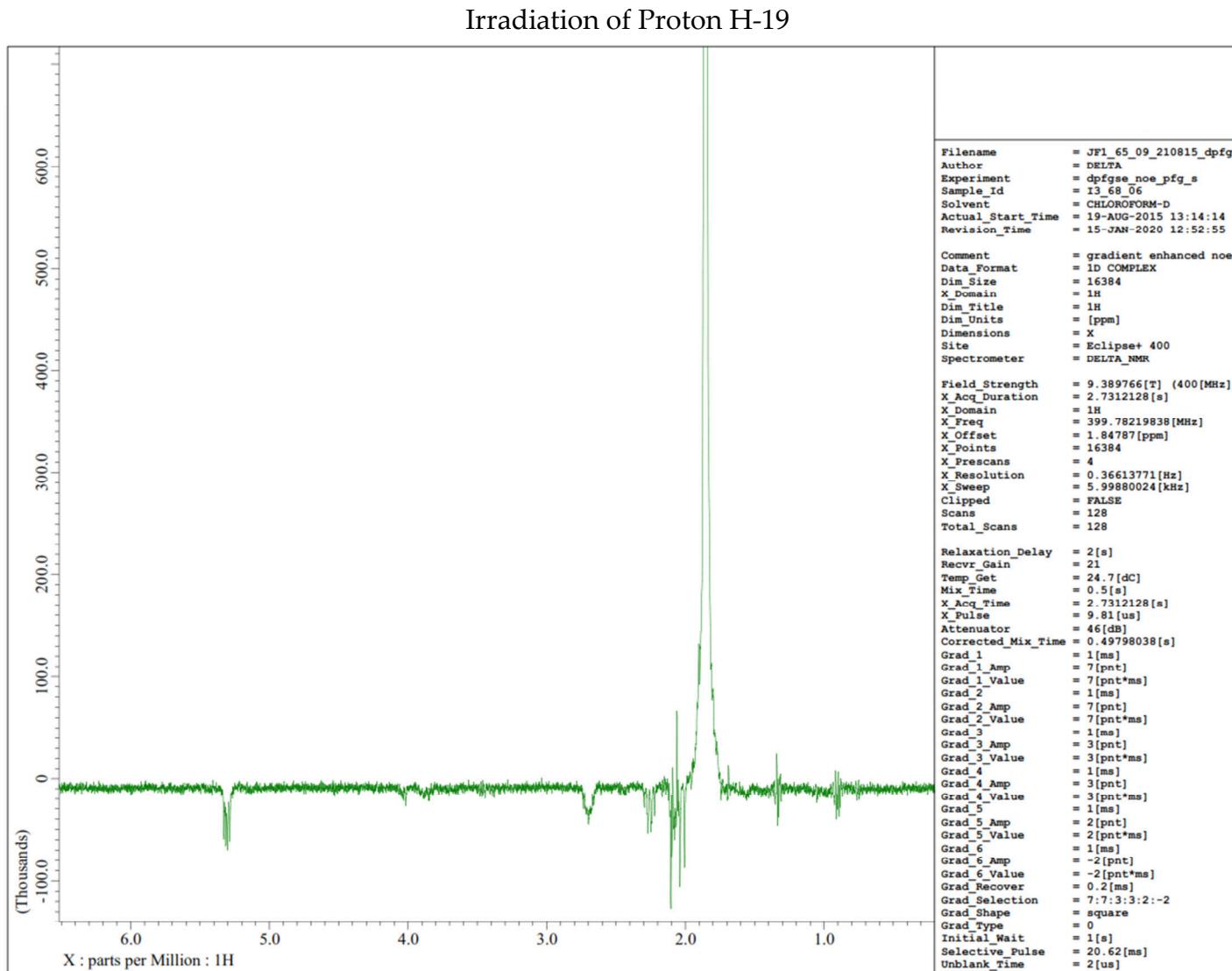
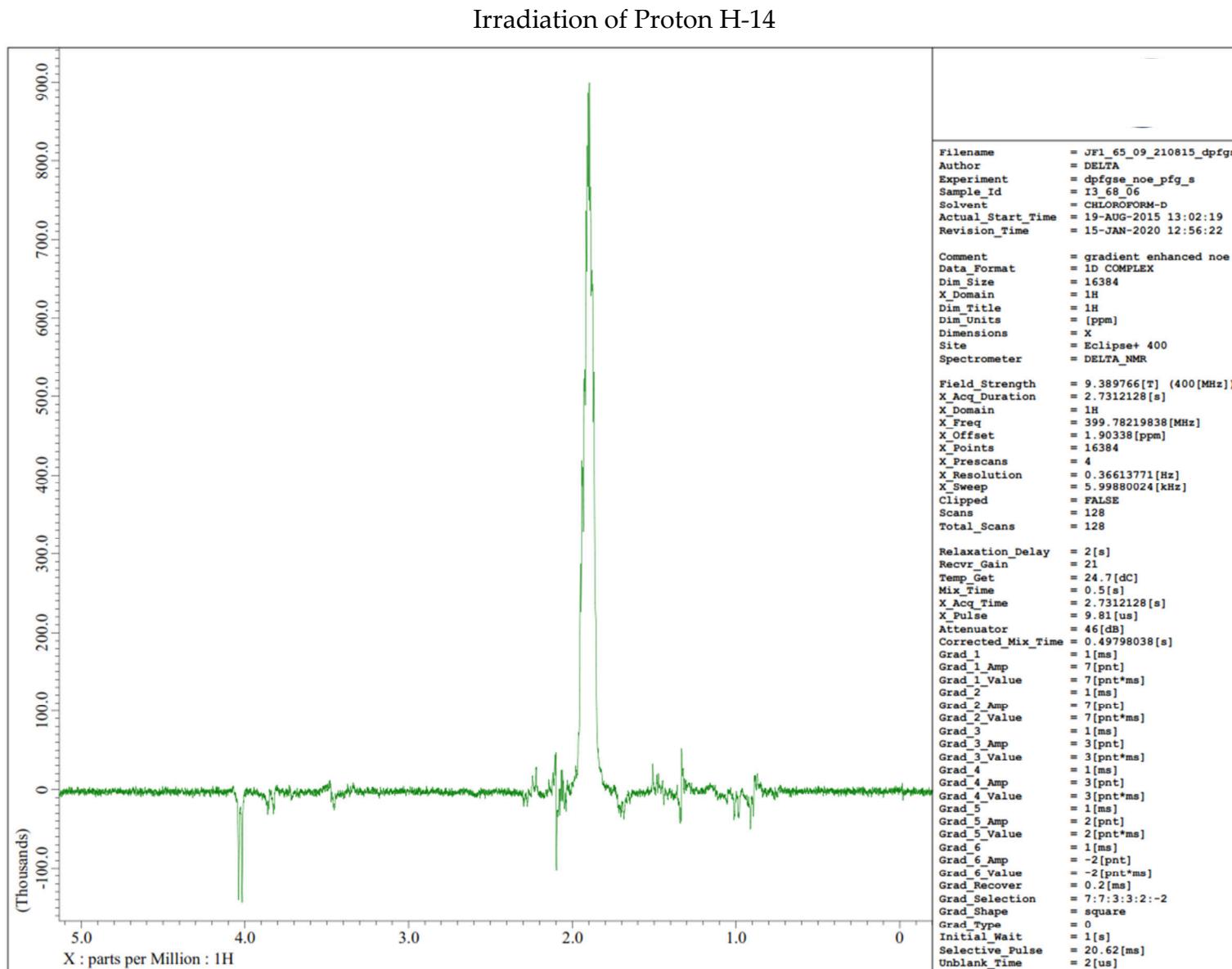
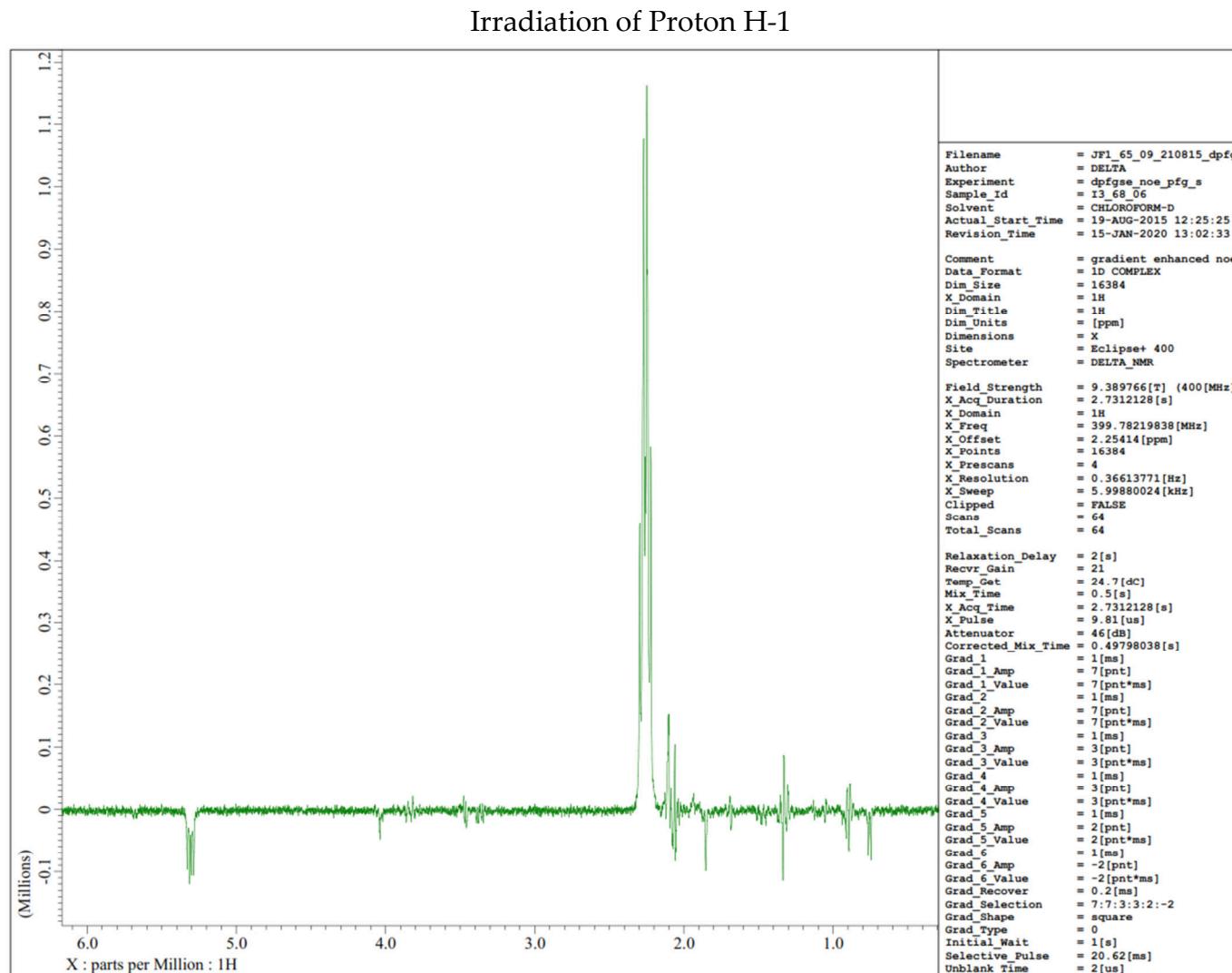


Figure S23. Asbestinin 28, NOE spectra

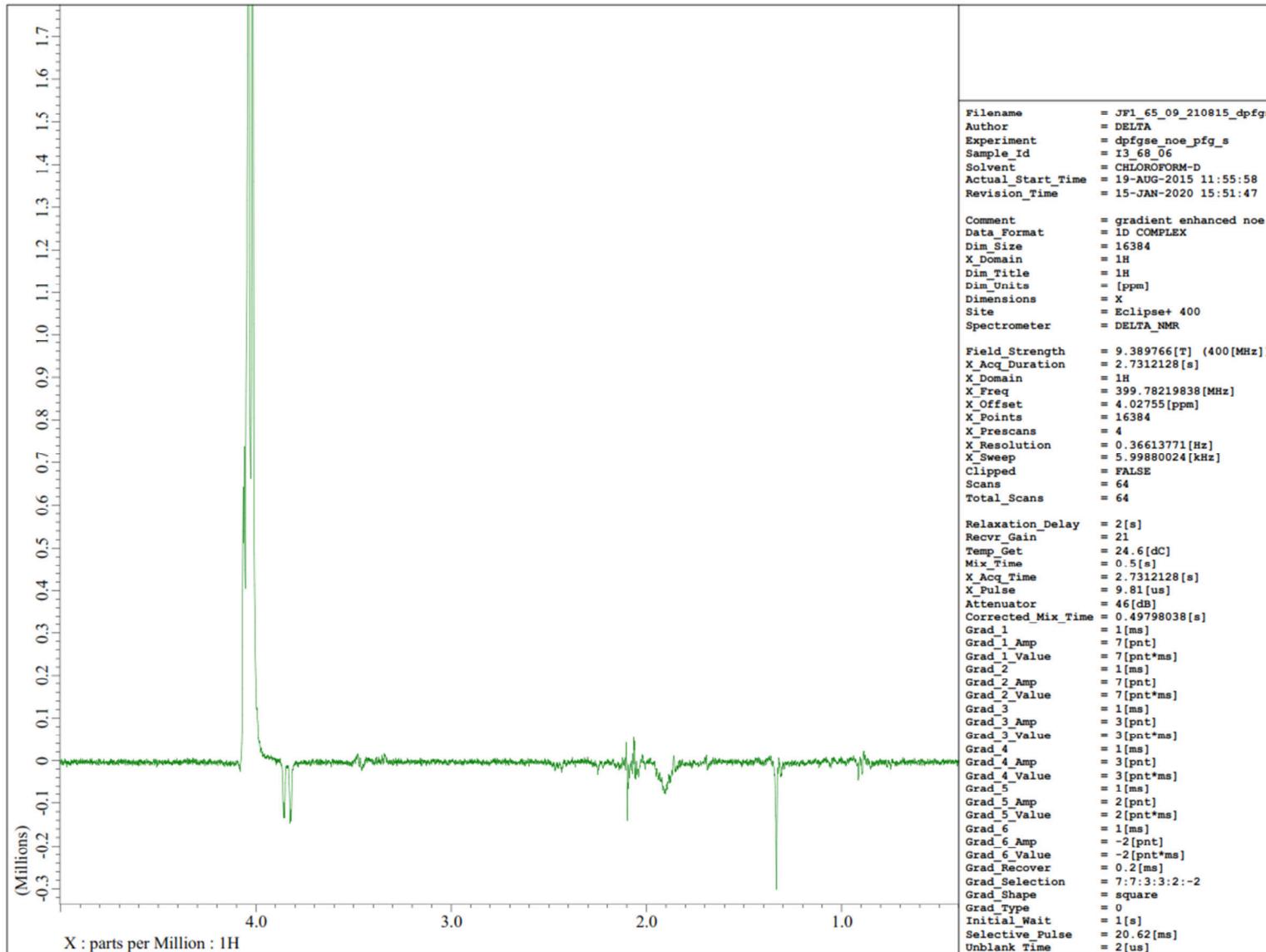


**Figure S23. Asbestinin 28, NOE spectra**



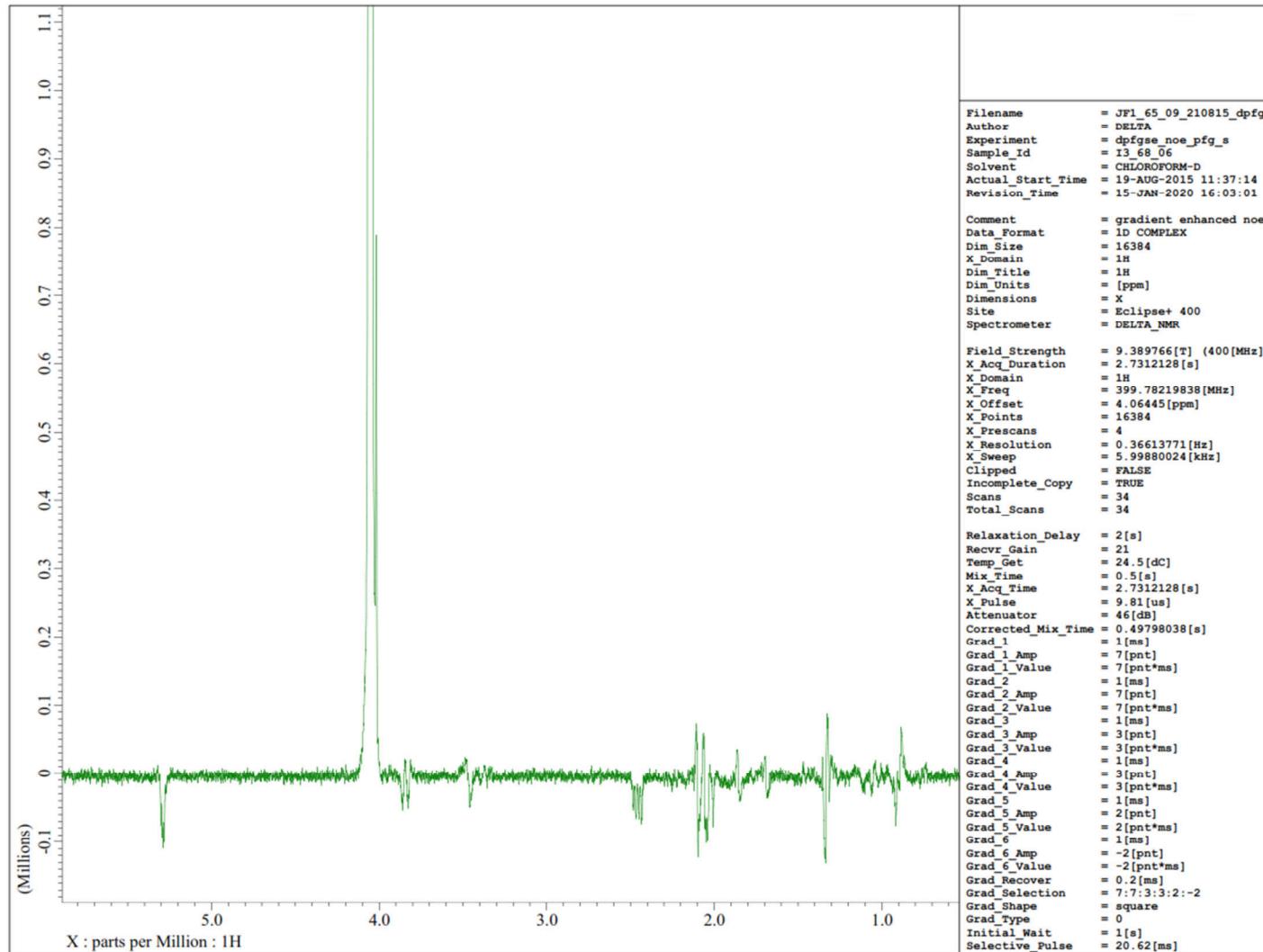
**Figure S23. Asbestinin 28, NOE spectra**

Irradiation of Proton H-2

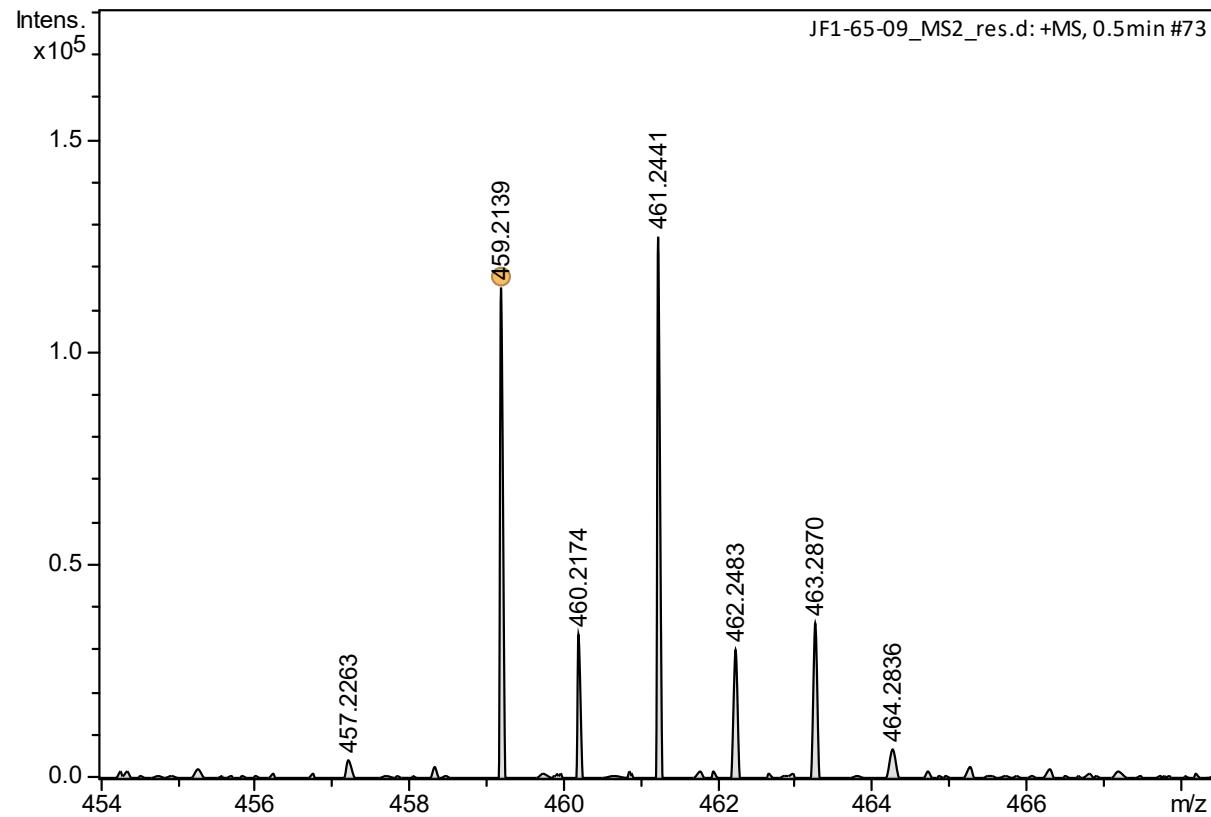


**Figure S23. Asbestinin 28, NOE spectra**

Irradiation of Proton H-9



**Figure S24. Asbestinin 28, HR-ESITOFMS spectrum**



**Table S1.** Viability of THP-1 human macrophages treated with different concentrations of diterpenes. Values are mean  $\pm$  SEM (%) of three independent experiments in duplicate ( $n = 3$ ).

	% Viability THP-1 macrophages (24 h)				
( $\mu$ M)	10	20	50	100	$IC_{50}$
Briarellin T ( <b>1</b> )	99.5 $\pm$ 2.0	98.5 $\pm$ 1.2	101.4 $\pm$ 0.5	98.5 $\pm$ 1.0	> 100
Asbestinin 27 ( <b>2</b> )	100.5 $\pm$ 1.3	99.5 $\pm$ 0.9	98.6 $\pm$ 2.1	97.4 $\pm$ 1.1	> 100
Asbestinin 28 ( <b>3</b> )	98.5 $\pm$ 3.1	99.1 $\pm$ 1.4	97.0 $\pm$ 1.0	99.5 $\pm$ 2.0	> 100
Asbestinin 17 ( <b>4</b> )	102.2 $\pm$ 1.5	100.5 $\pm$ 0.8	98.0 $\pm$ 2.0	99.5 $\pm$ 1.2	> 100