

# Preparation of $C_2$ -Symmetric Allenes by Palladium-Catalyzed Double Nucleophilic Substitution on 3-Bromopenta-2,4-dienyl Acetate

*Masamichi Ogasawara,<sup>\*</sup> Mayuka Suzuki, and Tamotsu Takahashi<sup>\*</sup>*

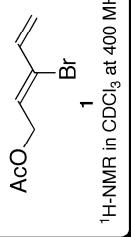
Catalysis Research Center and Graduate School of Life Science, Hokkaido University, Kita-ku, Sapporo  
001-0021, Japan

## Supporting Information

### Table of Contents

$^1\text{H}$ - and $^{13}\text{C}$ -NMR Spectra of 3-Bromo-2,4-pentadienyl Acetate <b>1</b> .....	S2-S3
$^1\text{H}$ - and $^{13}\text{C}$ -NMR Spectra of 3,3-Dibromo-2-propenyl Acetate <b>2</b> .....	S4-S5
$^1\text{H}$ - and $^{13}\text{C}$ -NMR Spectra of 1,1-Bis(phenylsulfonyl)undecane <b>5d</b> .....	S6-S7
$^1\text{H}$ - and $^{13}\text{C}$ -NMR Spectra of Functionalized $C_2$ -Symmetric Allenes <b>6a-g</b> .....	S8-S21
$^1\text{H}$ - and $^{13}\text{C}$ -NMR Spectra of 13,14-Heptacosadiene <b>9</b> .....	S22-S23
Chiral HPLC Chromatograms of ( <i>R</i> )-(-)- <b>6c</b> .....	S24

**JEOL**

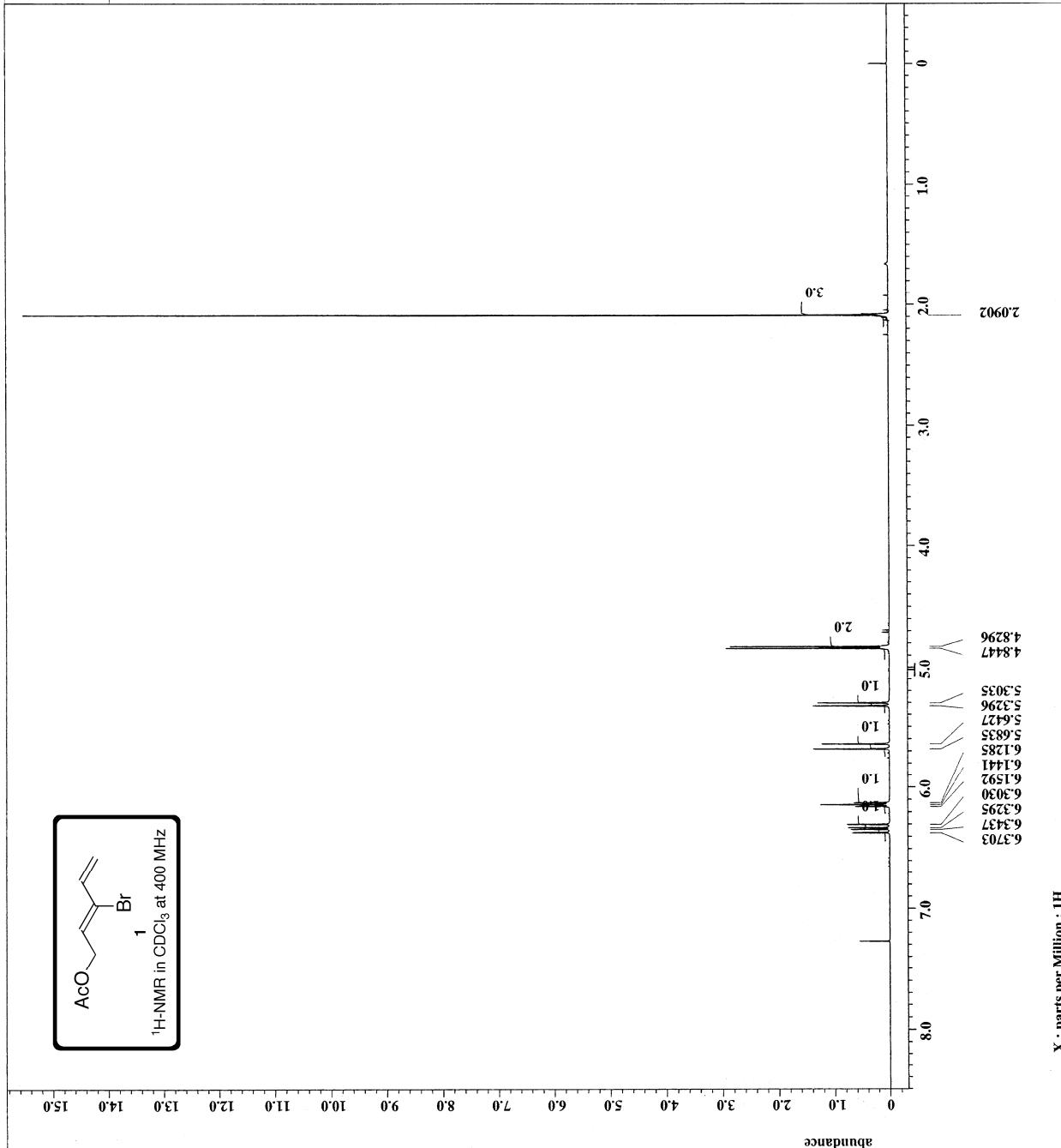


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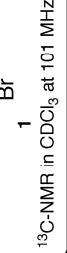
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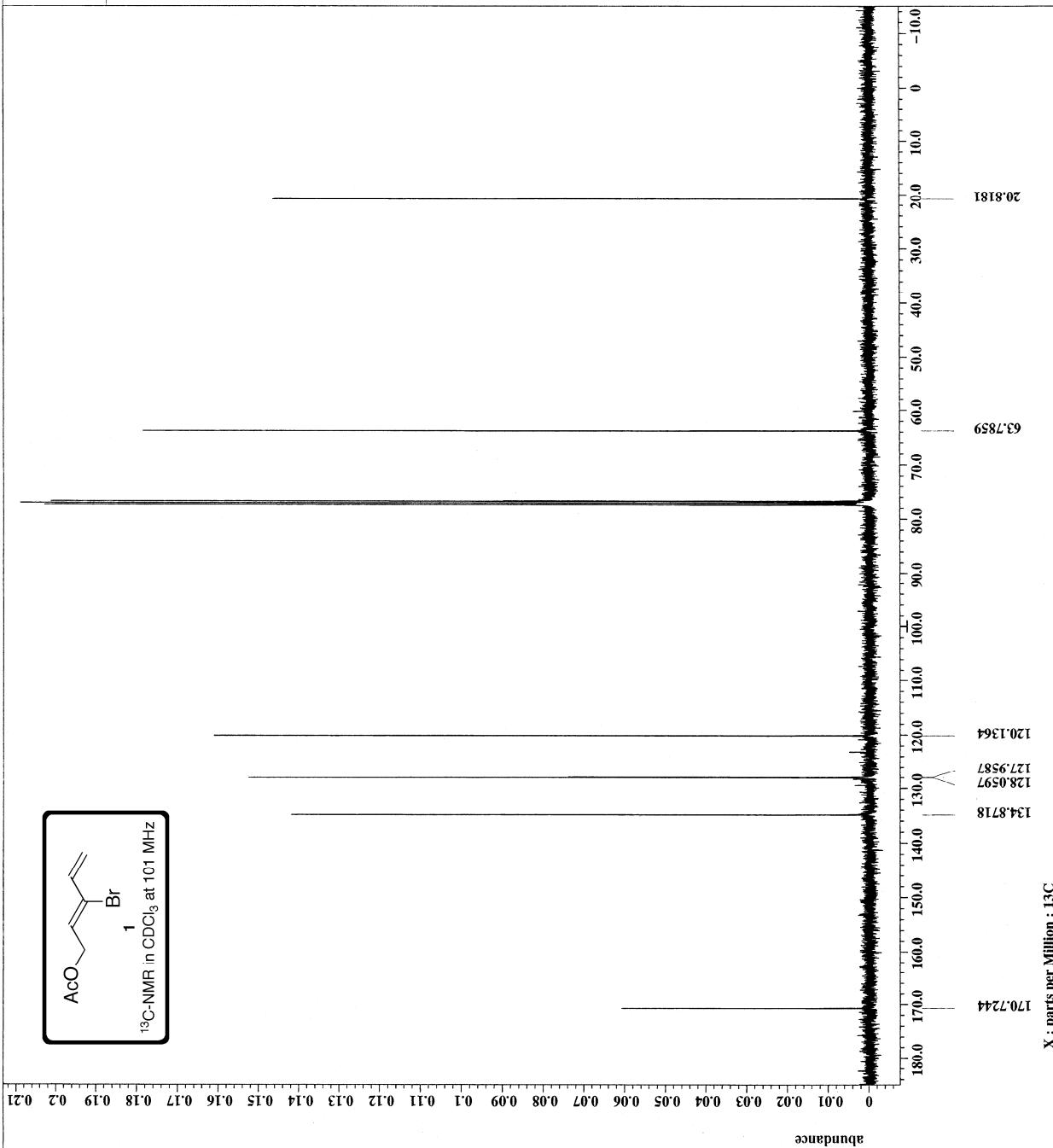
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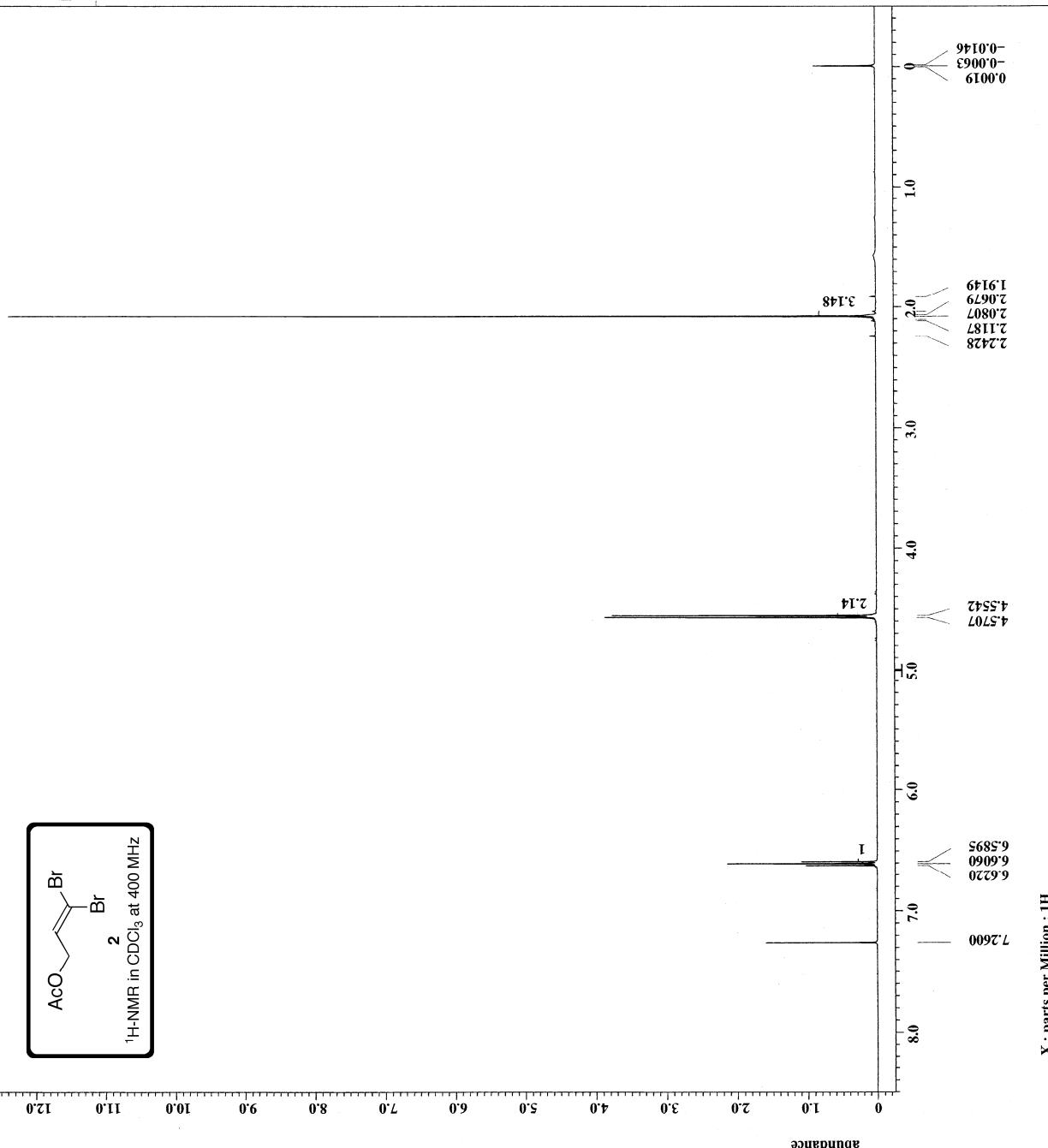
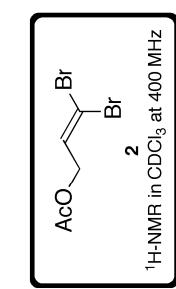


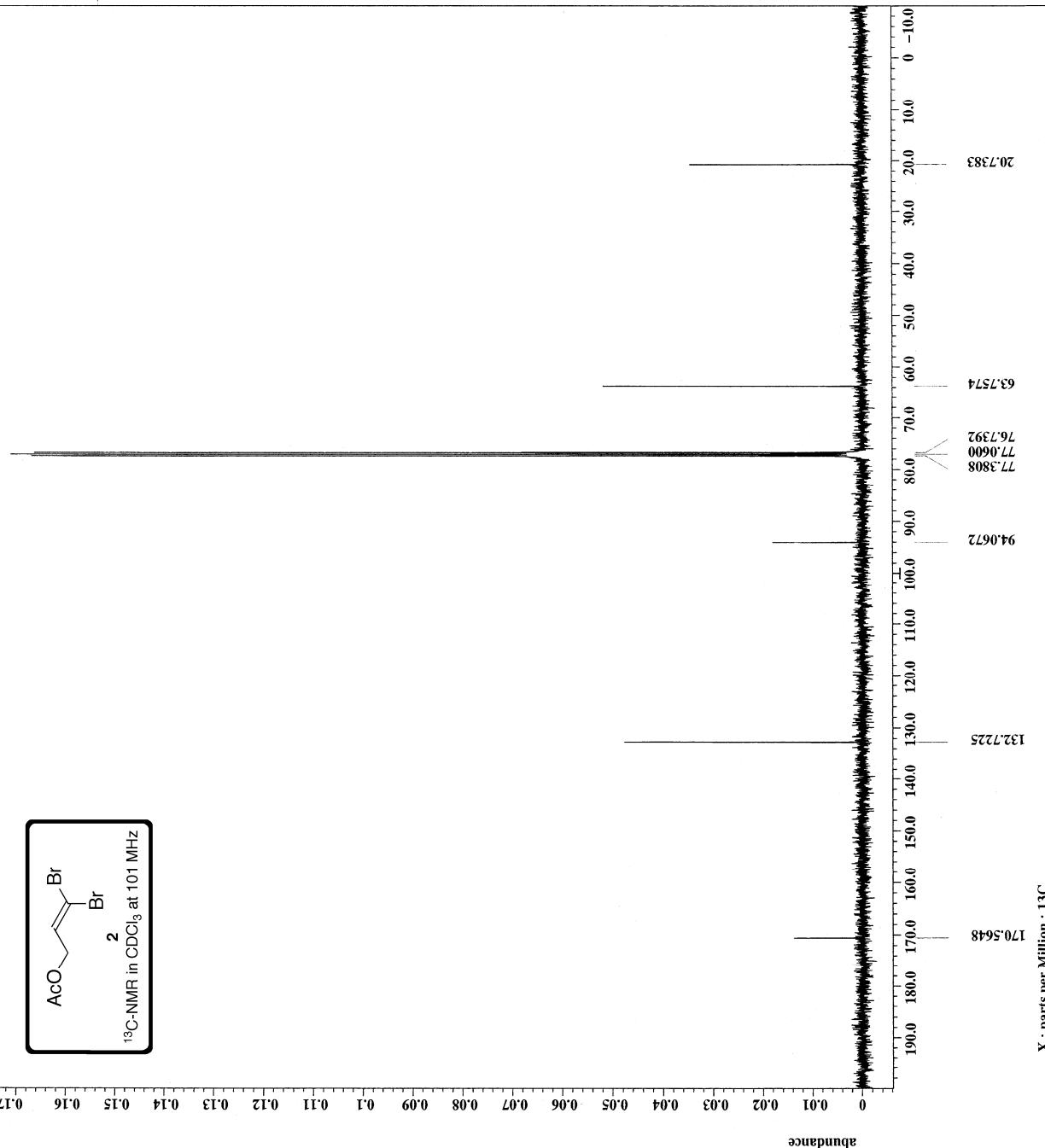
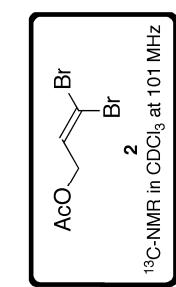
**AJEOL**



<sup>13</sup>C-NMR in CDCl<sub>3</sub> at 101 MHz





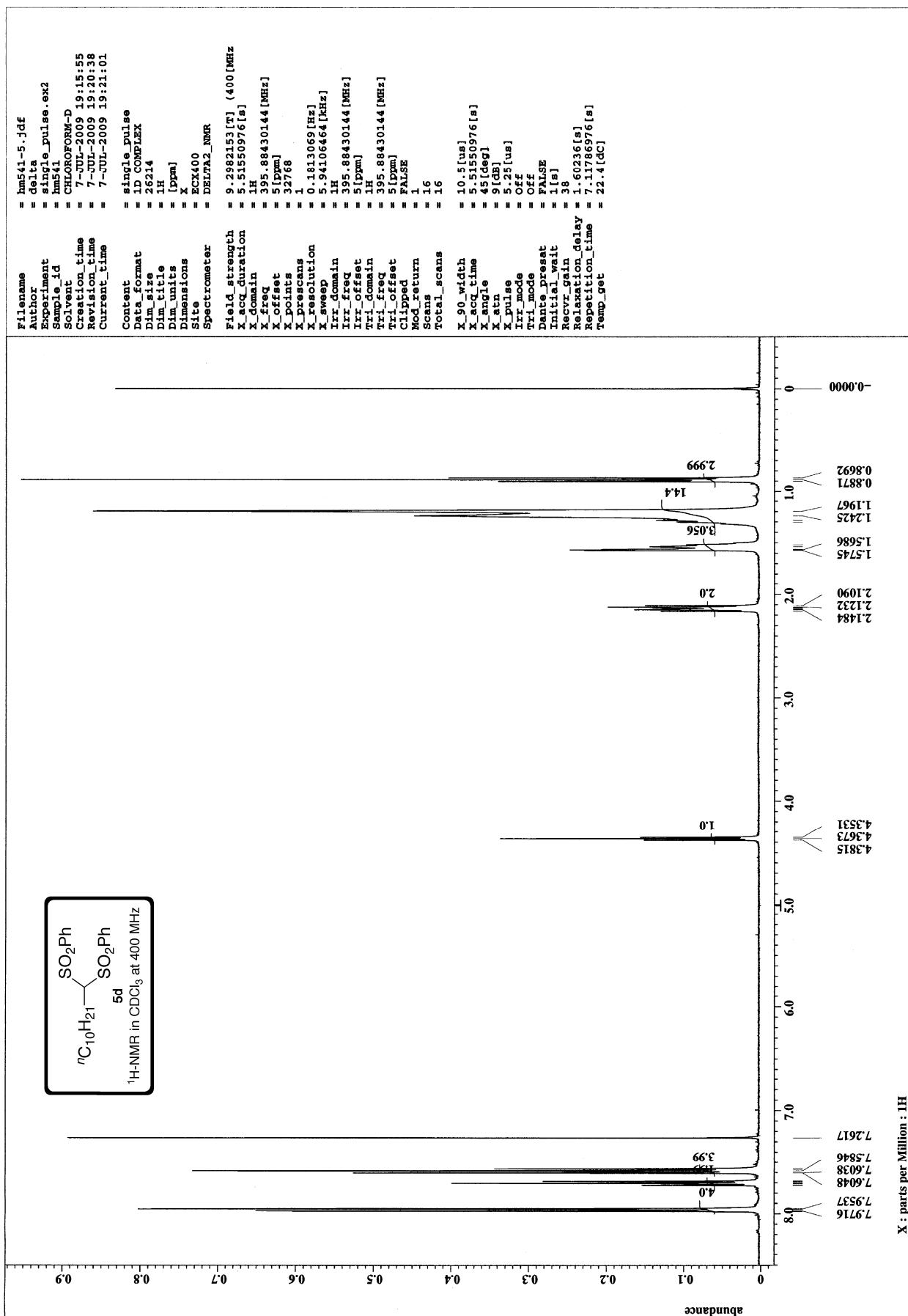


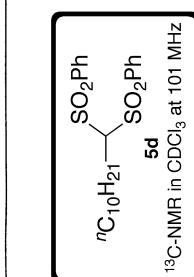
**AJEOL**

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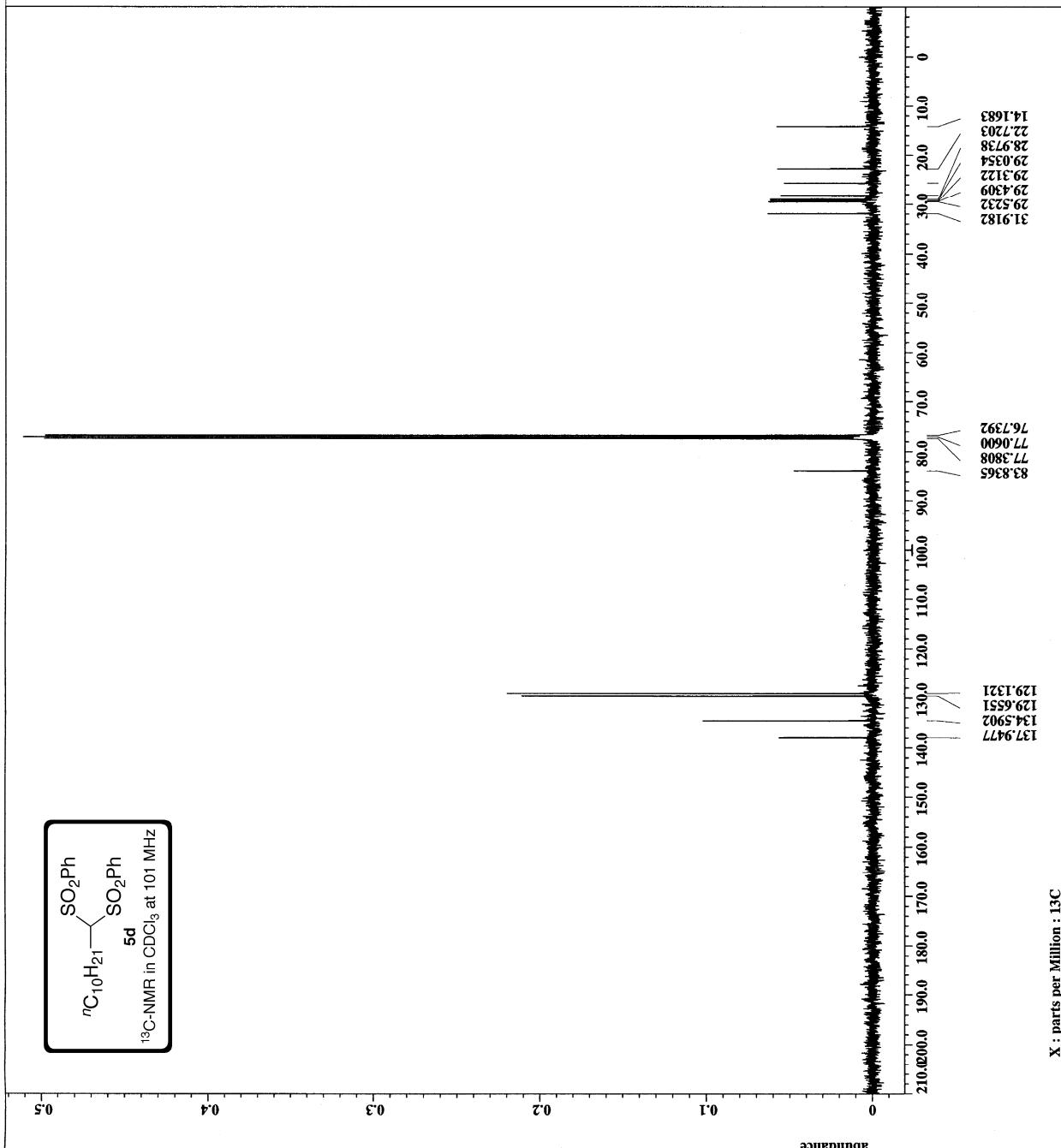
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X_domain      = 13C
X_freq        = 99.54517646 [MHz]
X_offset      = 100 [ppm]
X_points      = 65536
X_precs      = 1
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X_sweep       = 28 - 569.2477 [kHz]
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Irr_freq      = 395.88430144 [MHz]
Irr_offset    = 5 [ppm]
Clipped      = FALSE
Mod_return   = 1
Scans         = 256
Total_scans  = 256
X_90_width   = 9.2 [us]
X_acq_time   = 2.24584564 [s]
X_angle      = 30 [deg]
X_stn        = 8.4 [dB]
X_pulse      = 3.06665667 [us]
Irr_attn_dec = 29.2 [dB]
Irr_attn_noe = 29.2 [dB]
Irr_noise    = 0.0172
Decoupling   = TRUE
Initial_wait = 1 [s]
Noe          = TRUE
Noe_time     = 1.13759 [s]
Revr_gain    = 50
Relaxation_delay = 1.13759 [s]
Repetition_time = 3.4234856 [s]
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<sup>13</sup>C-NMR in CDCl<sub>3</sub> at 101 MHz



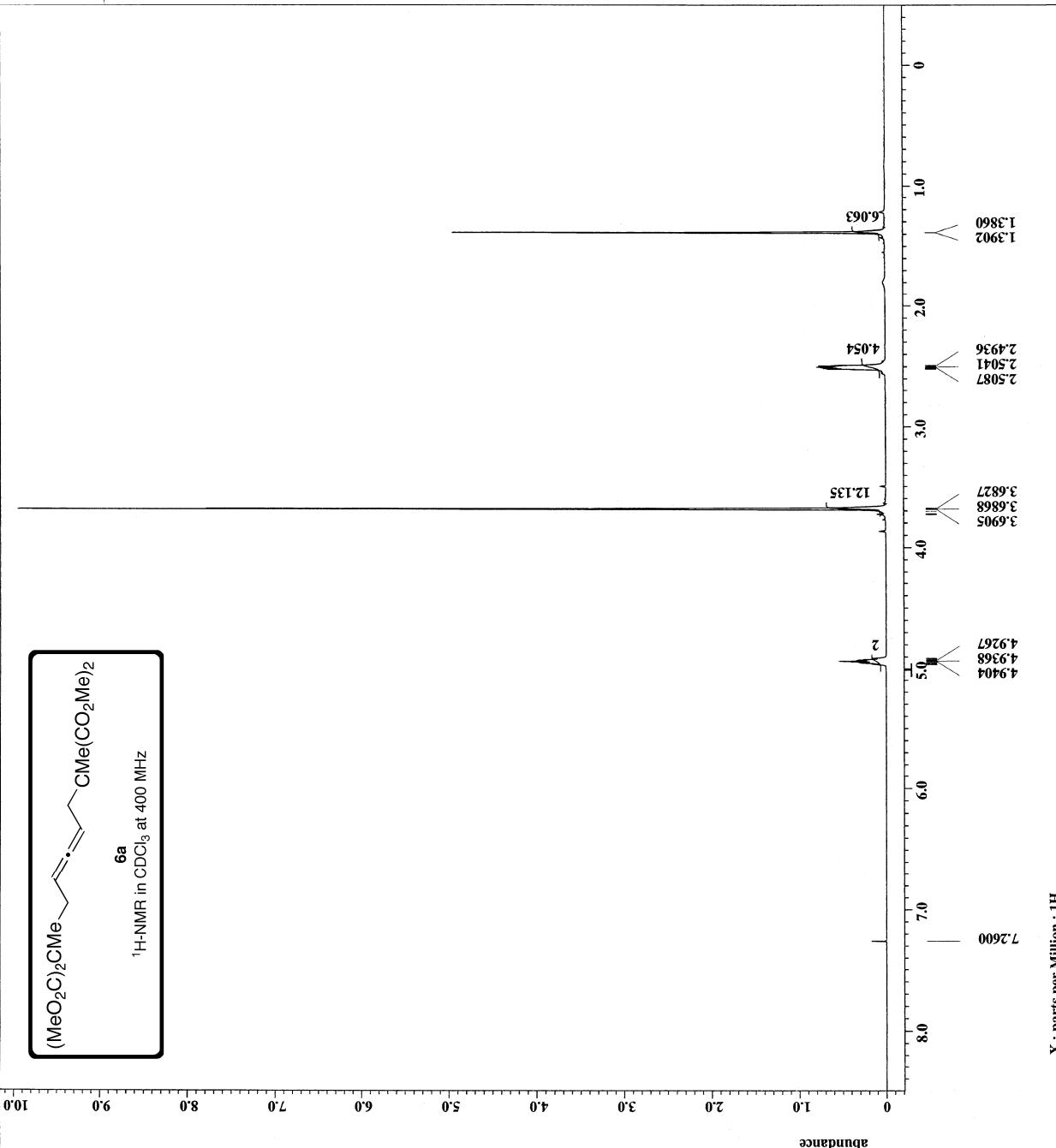
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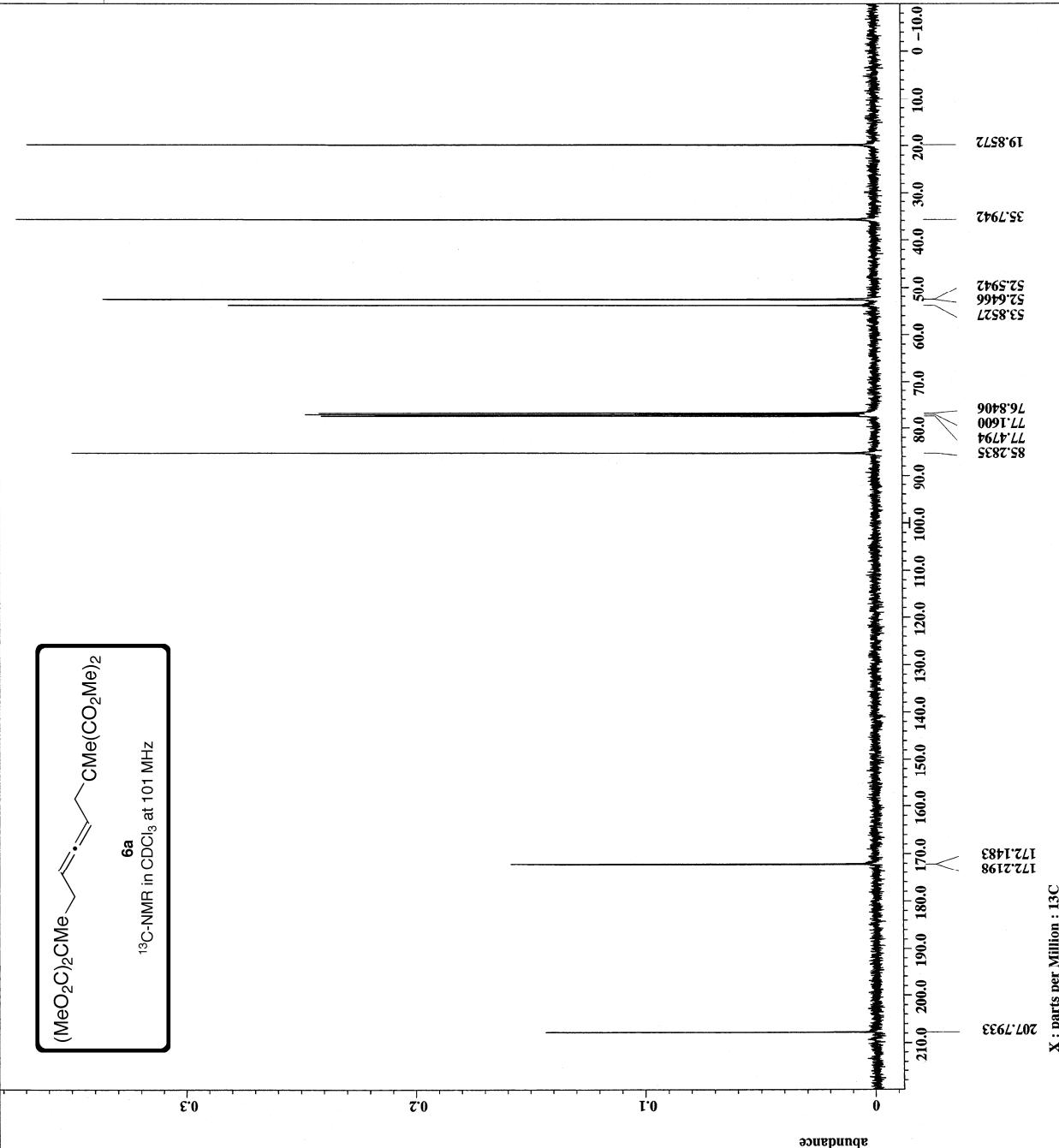
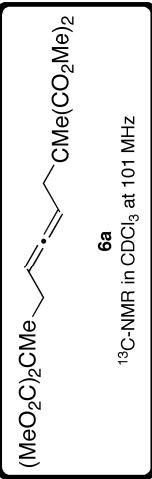
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Dimensions      = X
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X_sweep          = 28.66972477[Hz]
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Irr_freq         = 395.88430144[MHz]
Irr_offset       = 5[ppm]
Clipped          = FALSE
Mod_return       = 256
Scans            = 256
Total_scans     = 256
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X_90c_time      = 2.28589568[s]
X_angle          = 30[deg]
X_atten          = 8.4[db]
X_pulse          = 3.16666667[us]
Irr_att_dec     = 29.2[db]
Irr_att_noe     = 29.2[db]
Irr_noise        = WALTZ
Decoupling       = TRUE
Initial_wait     = 1[s]
Noe              = TRUE
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Revr_Gain        = 60
Relaxation_delay = 0.89[s]
Repetition_time  = 3.17589568[s]
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**6a**

<sup>1</sup>H-NMR in CDCl<sub>3</sub> at 400 MHz

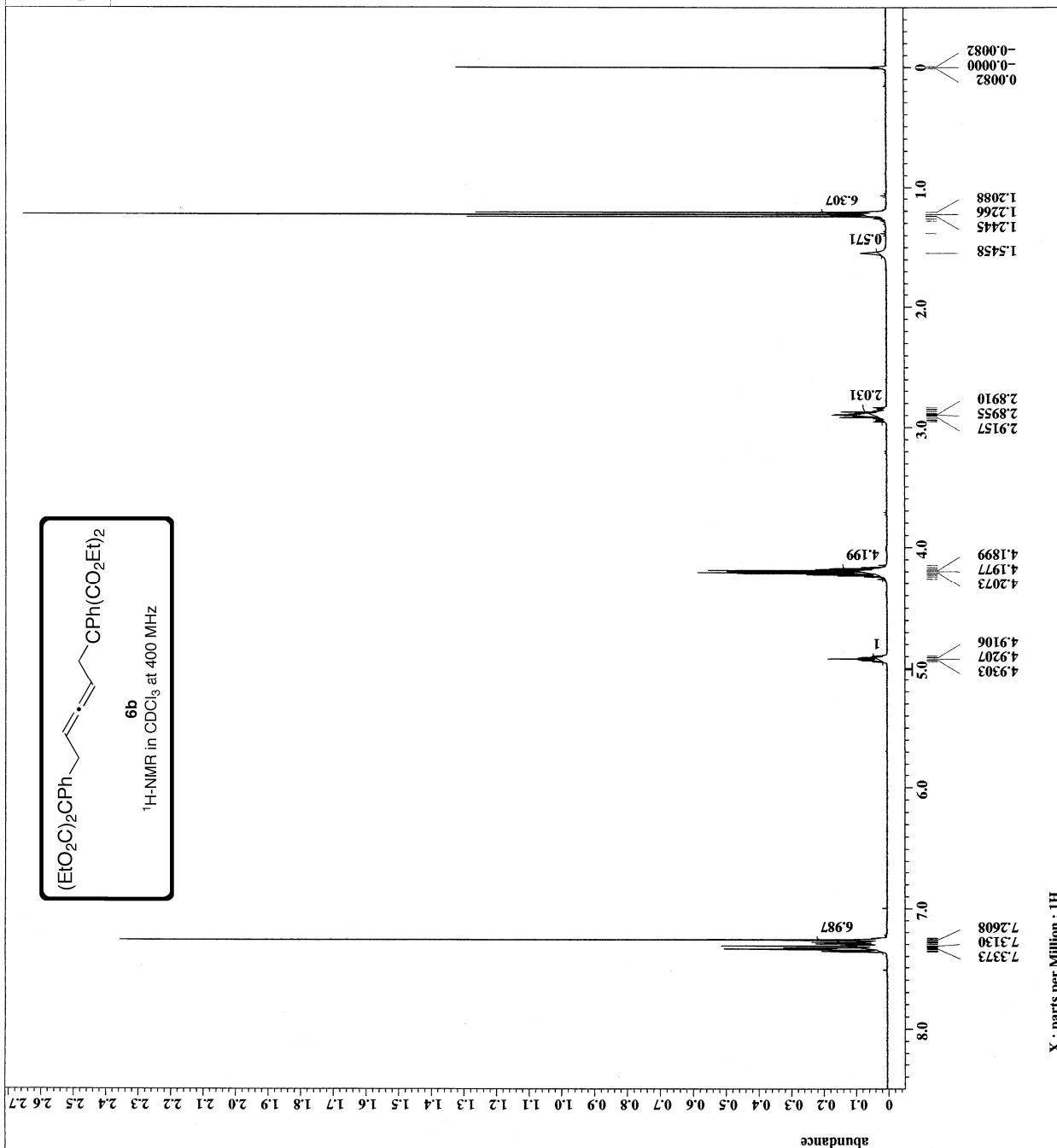




**JEOL**

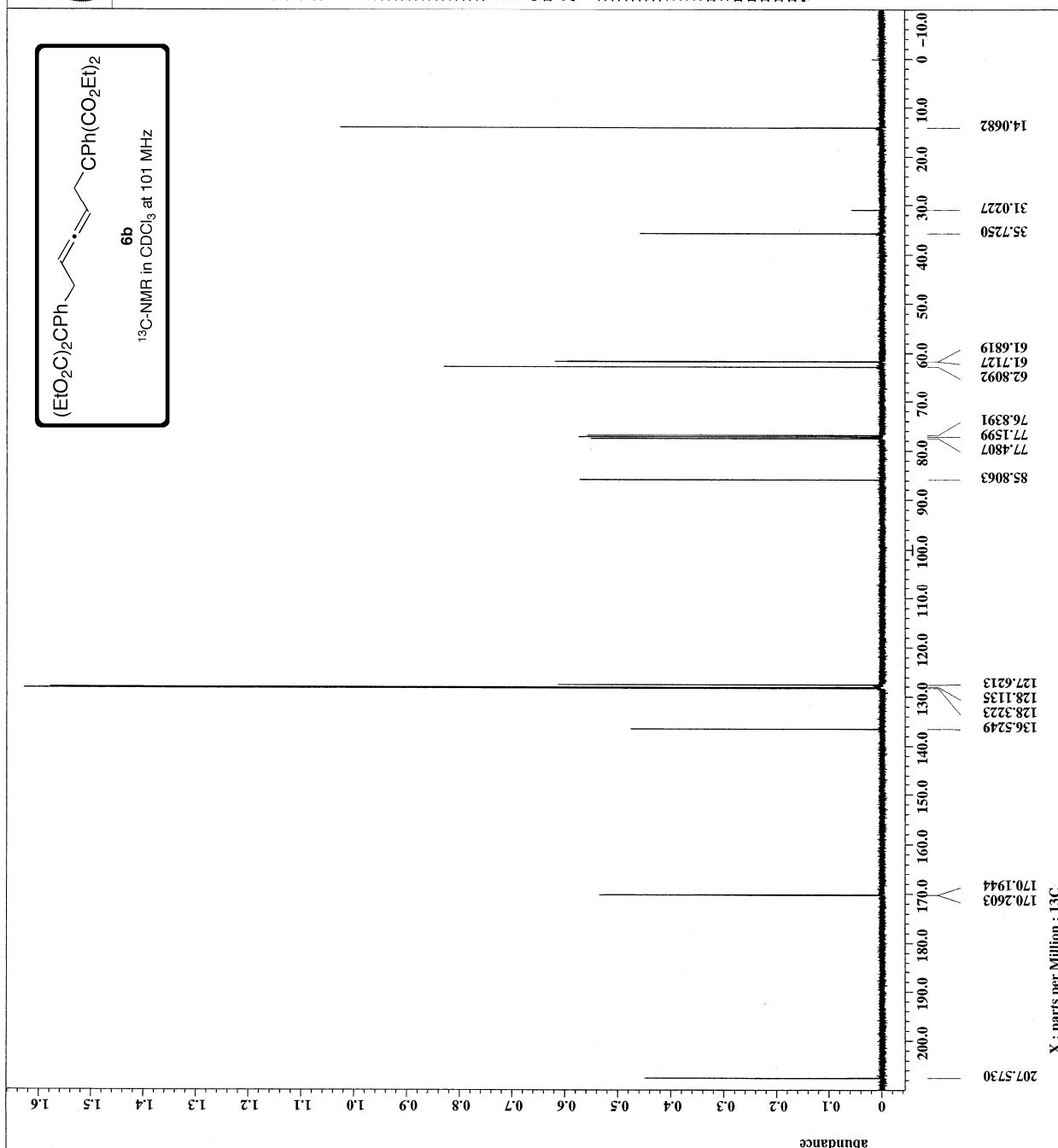


**6b**  
<sup>1</sup>H-NMR in CDCl<sub>3</sub> at 400 MHz



X: parts per Million : 1H

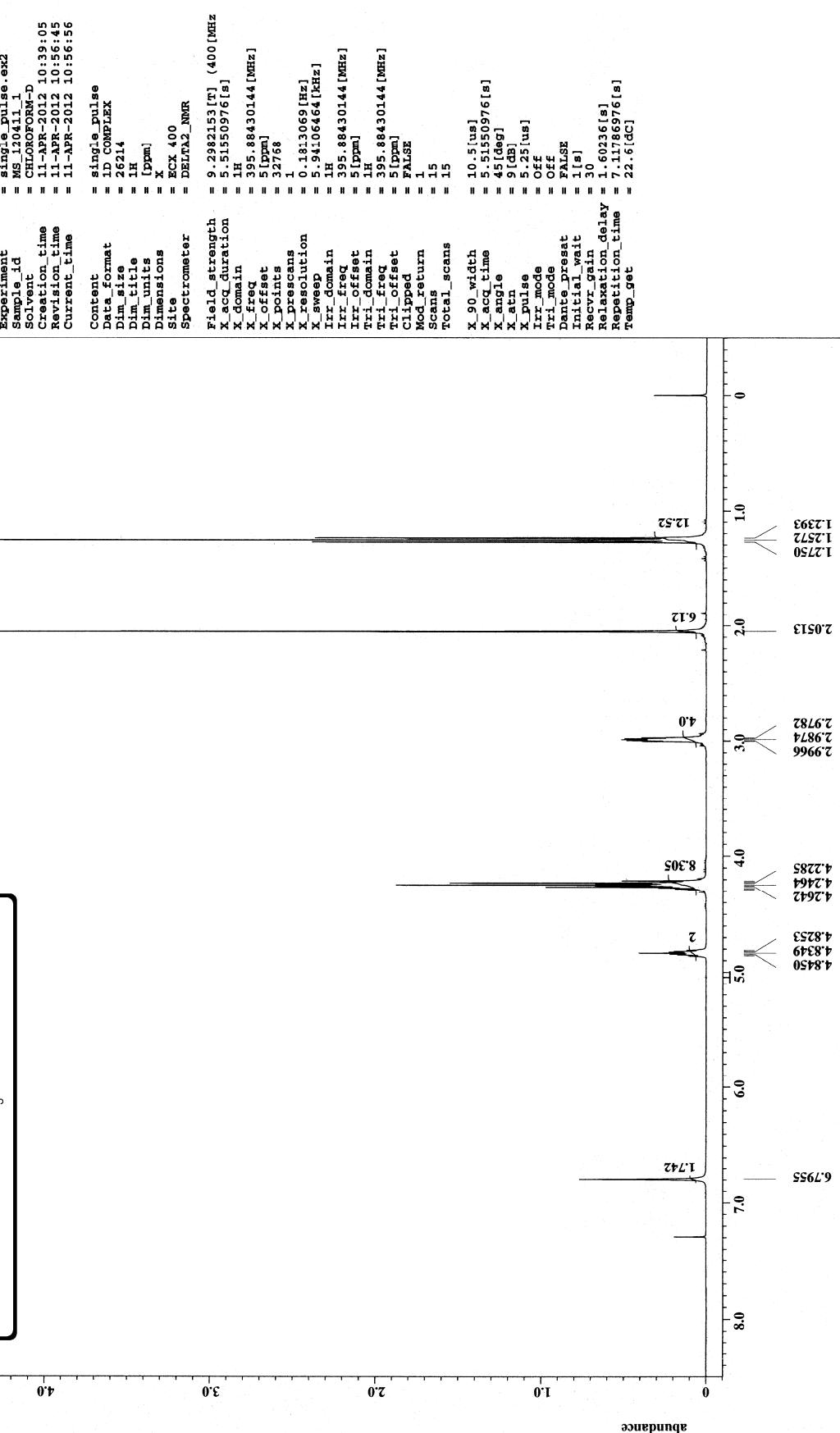
**JEOL**



**JEOL**

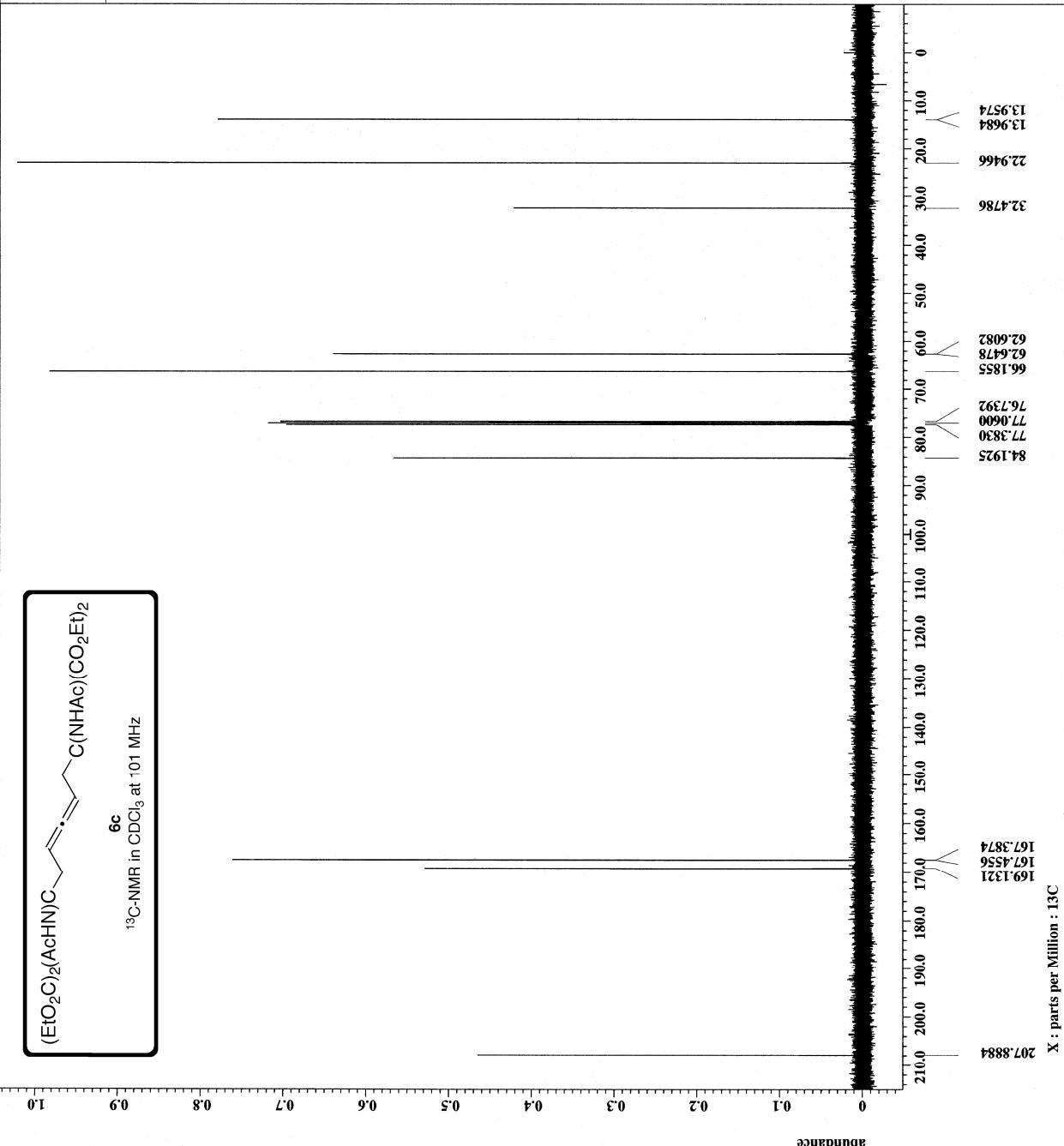
**6c**

<sup>1</sup>H-NMR in CDCl<sub>3</sub> at 400 MHz

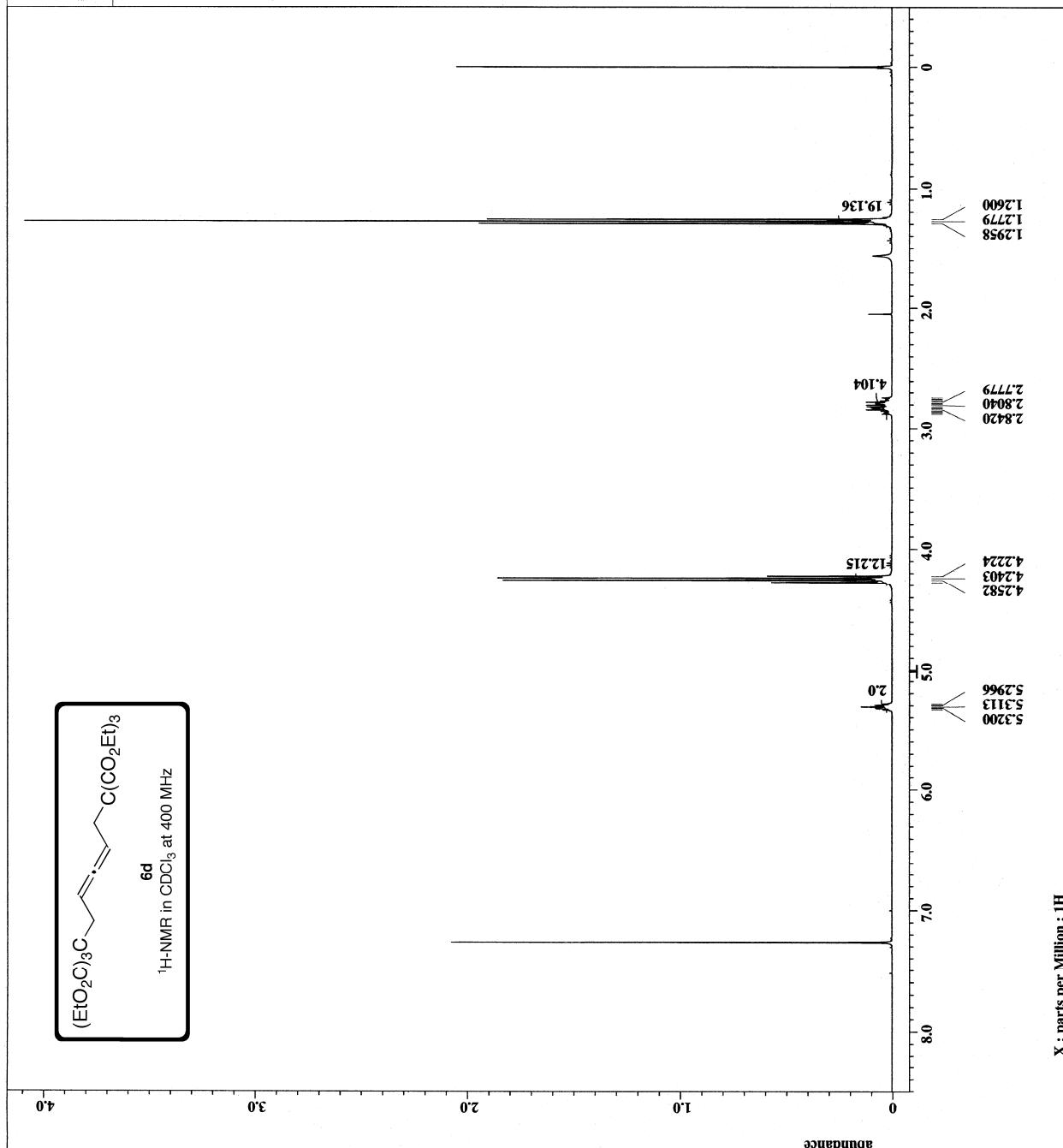
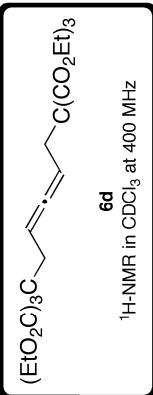




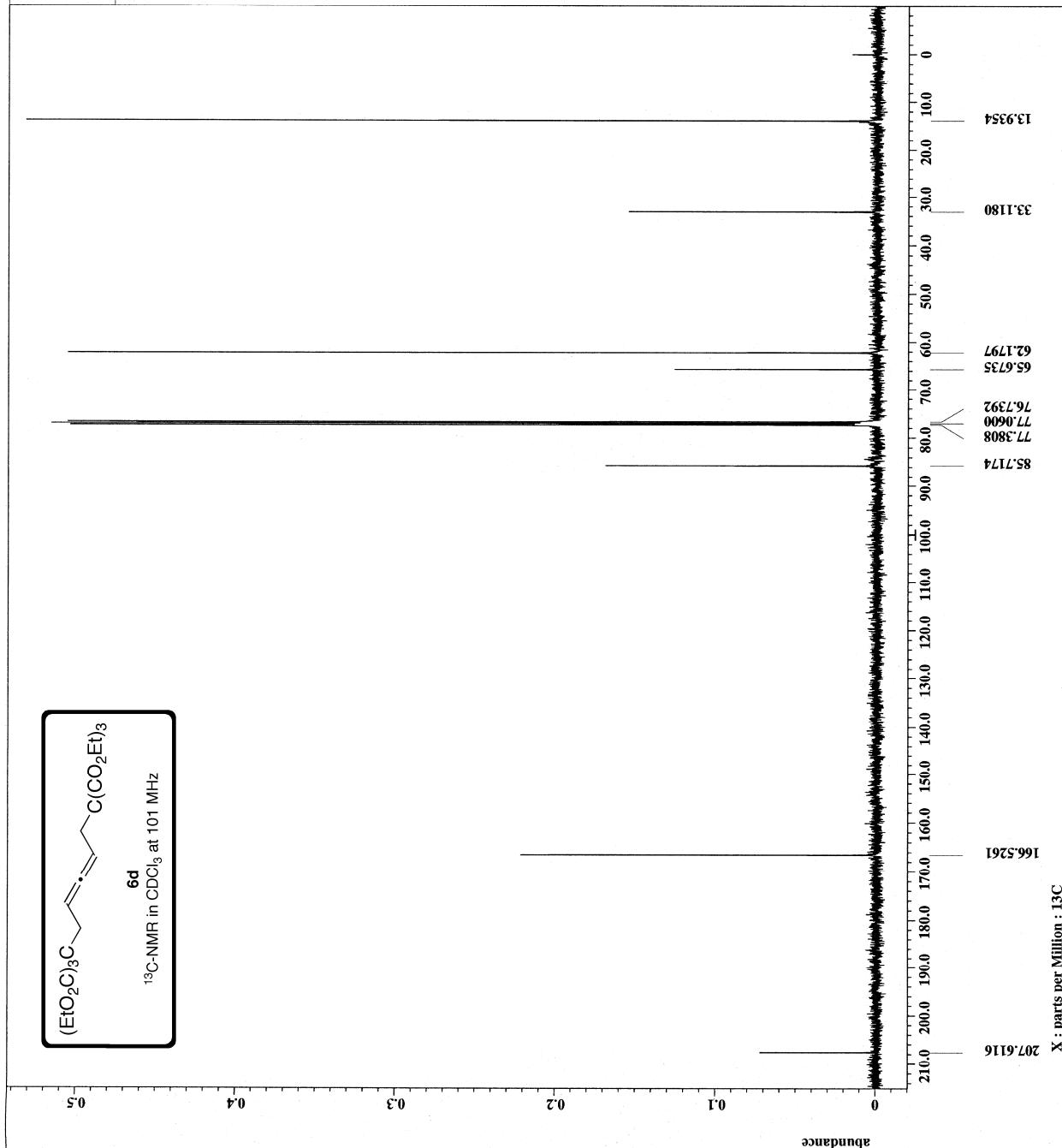
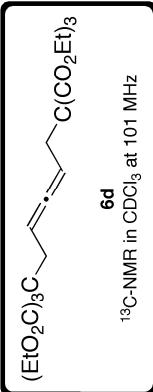
13C-NMR in CDCl<sub>3</sub> at 101 MHz



**AUJEOL**



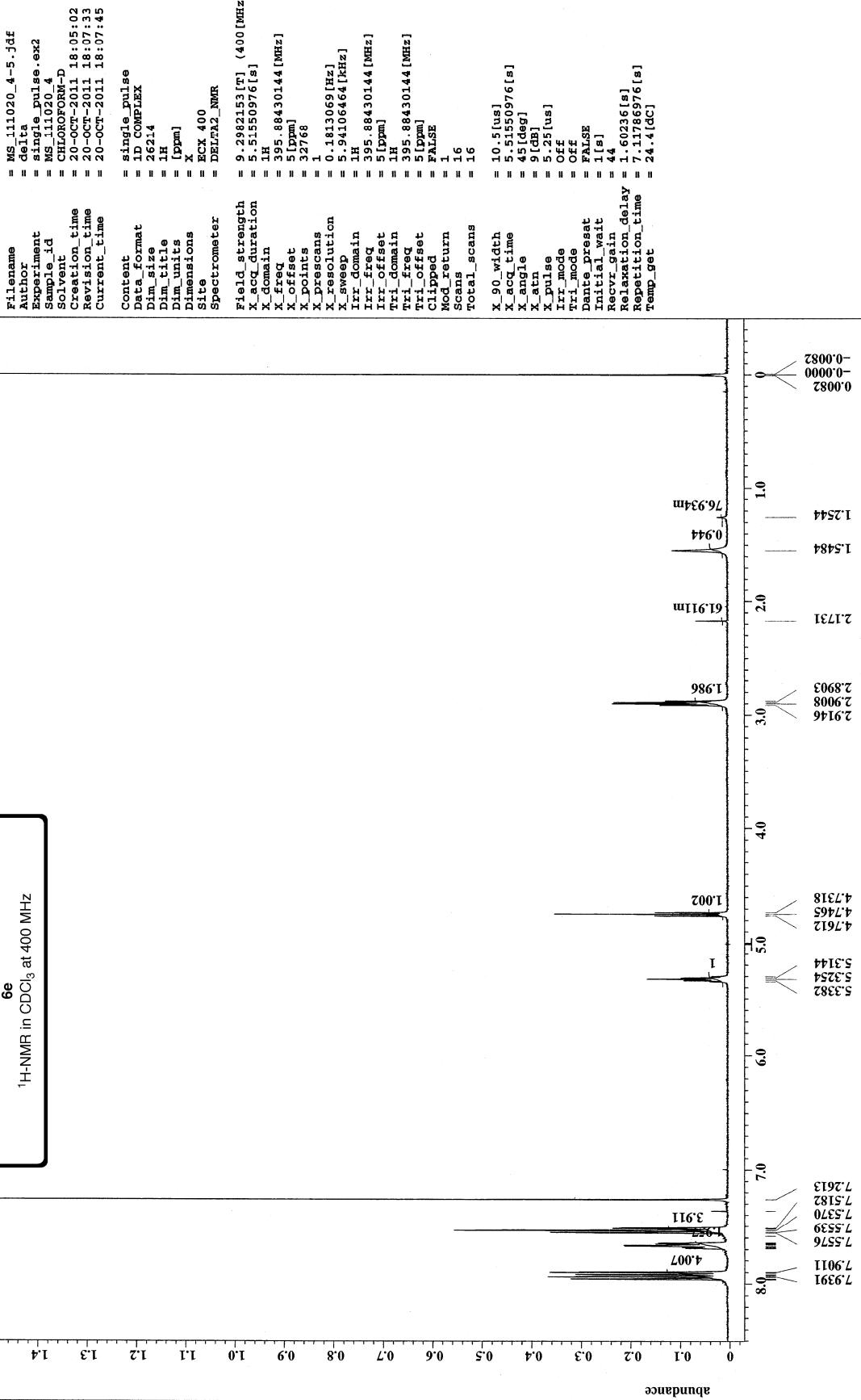
**JEOL**



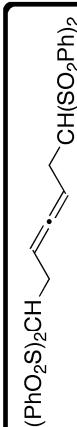
**JEOL**



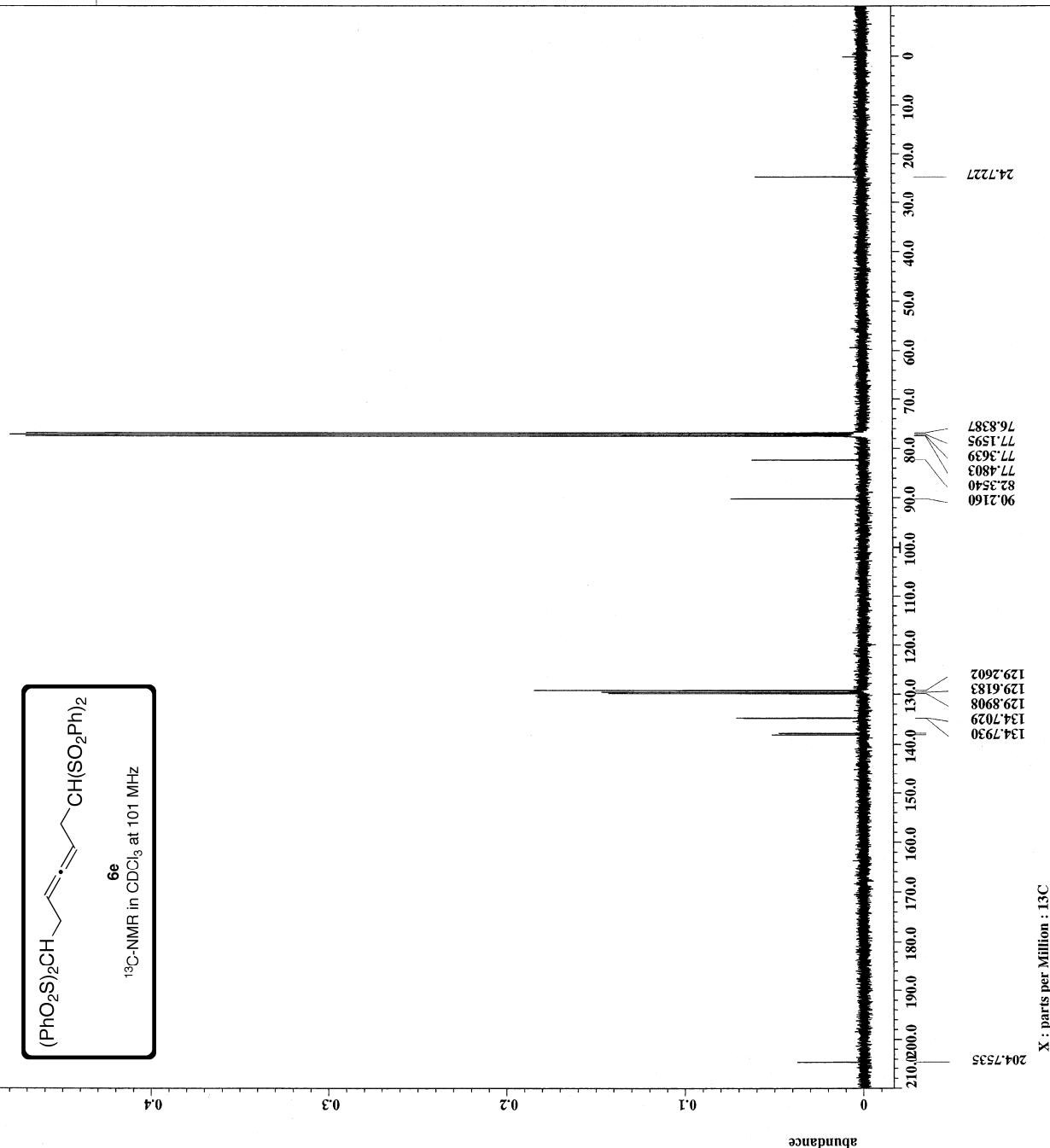
1H-NMR in CDCl<sub>3</sub> at 400 MHz



**JEOL**



**6e**  
 $^{13}\text{C}$ -NMR in  $\text{CDCl}_3$  at 101 MHz



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Author        = delta
Experiment   = single_pulse_dec
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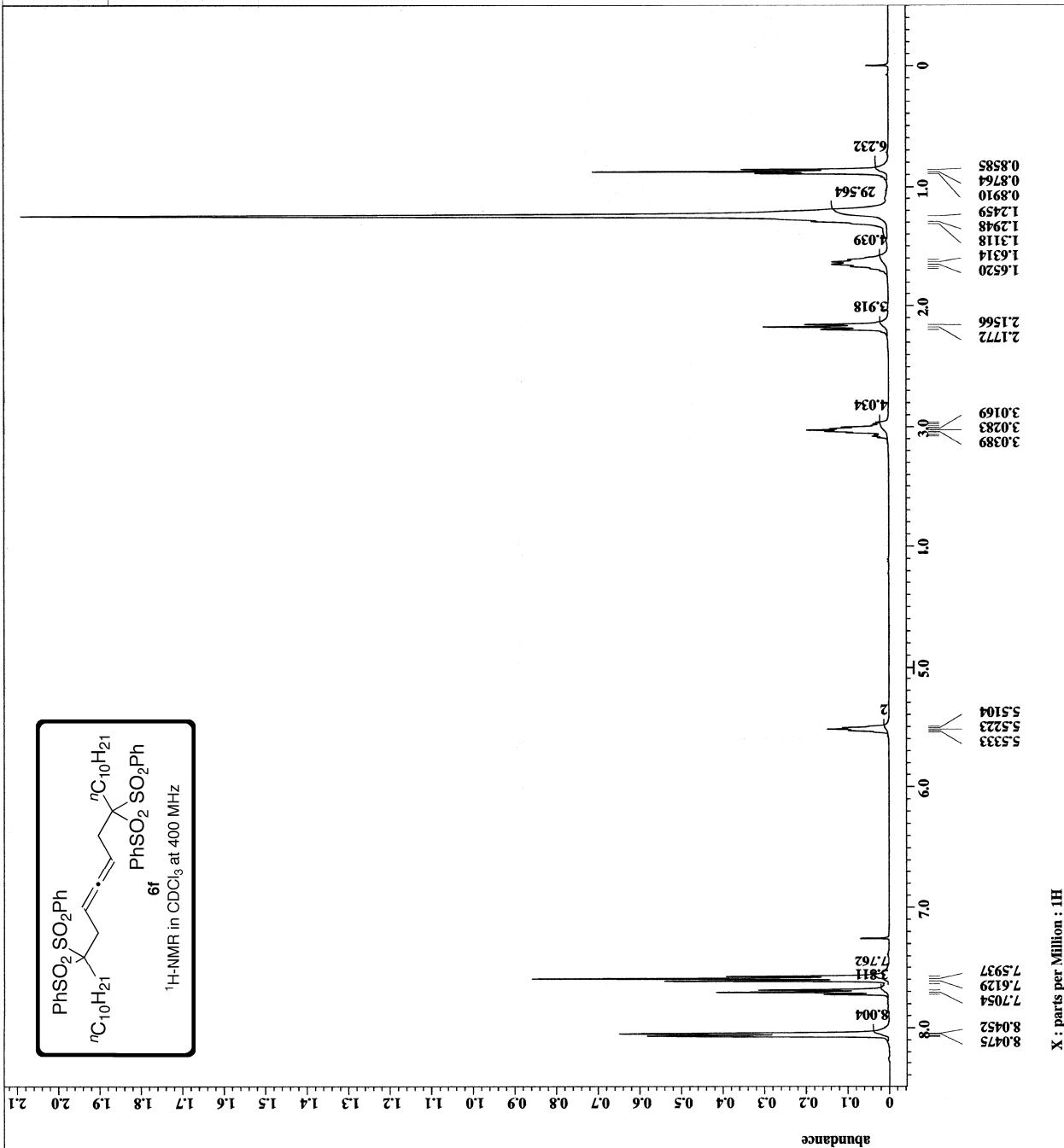
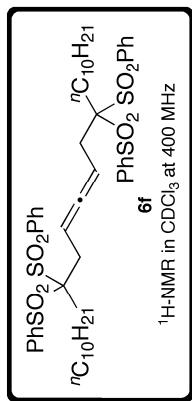
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X.domain      = 13C
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X.offset       = 100 [ppm]
X.jpoints      = 131072
X.precs        = 1
X.resolution  = 0.21873264 [Hz]
X.sweep        = 28.66972477 [kHz]
Irr_domain    = 1H
Irr_freq       = 395.88430144 [MHz]
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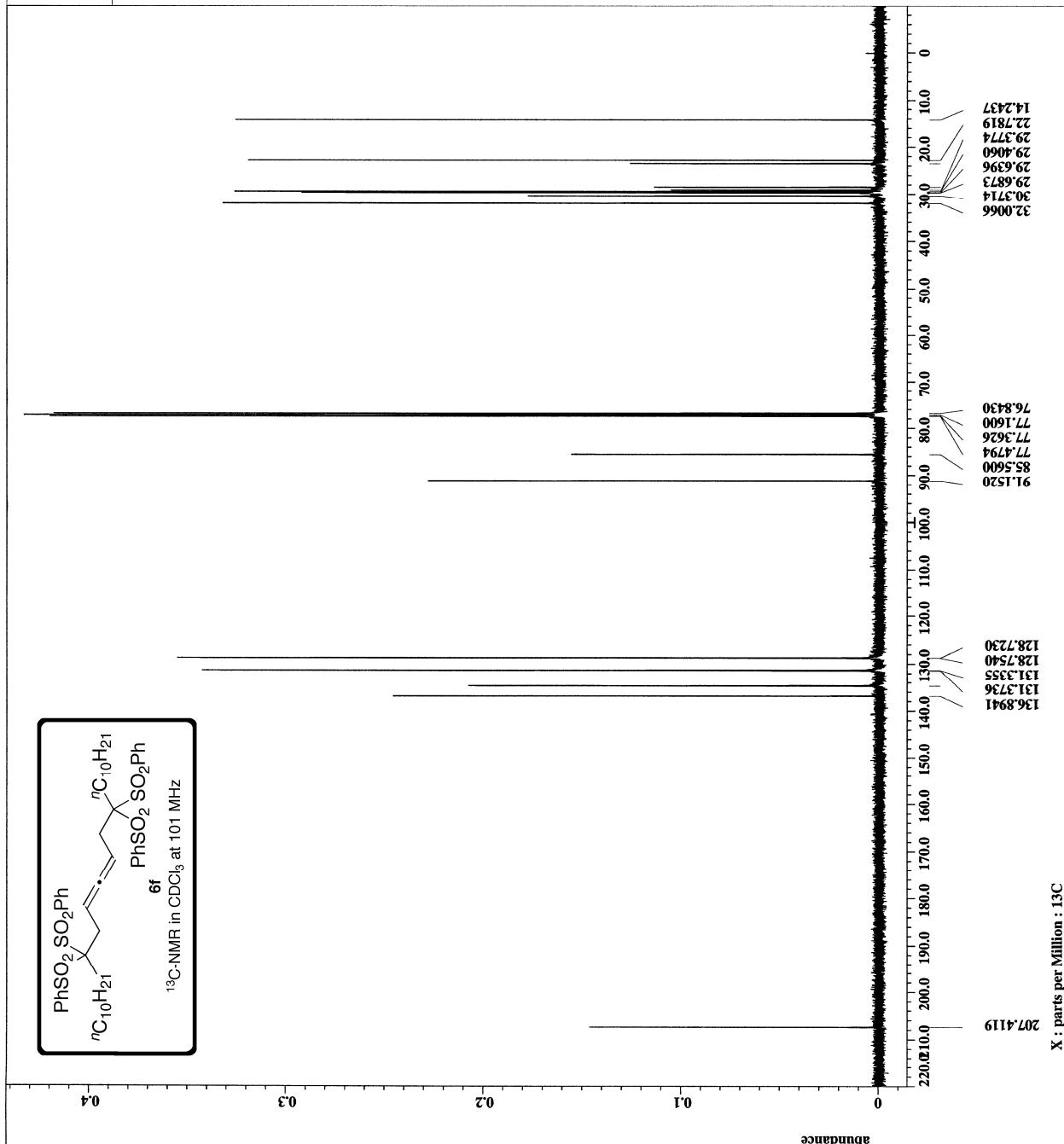
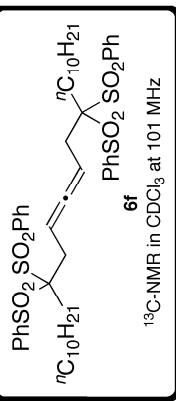
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X_pulse        = 3.16666667 [us]
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Irr_actn_noe  = 29.79 [dB]
Irr_noise      = WALTZ
Decoupling    = TRUE
Initial_wait   = 1 [s]
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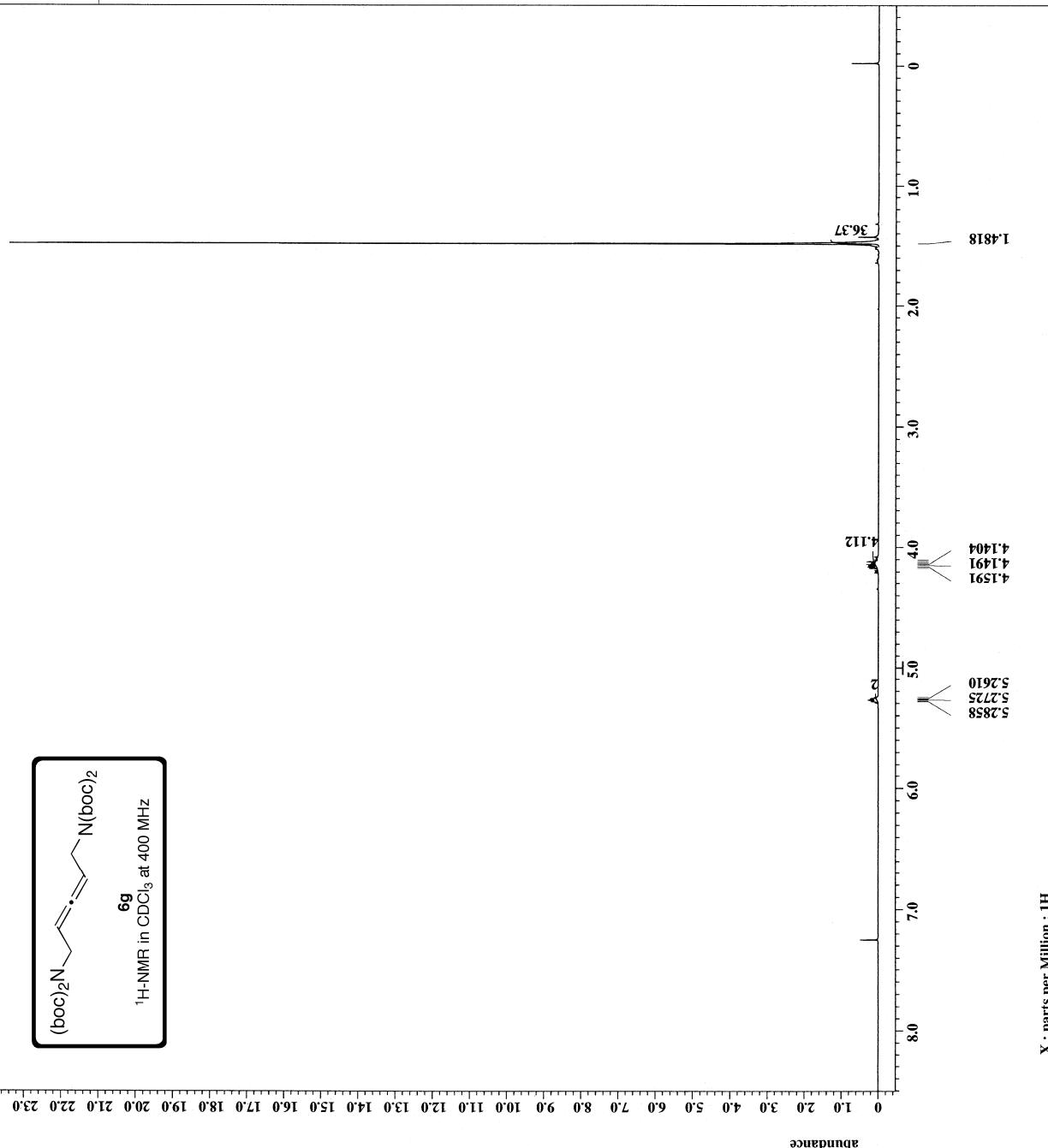
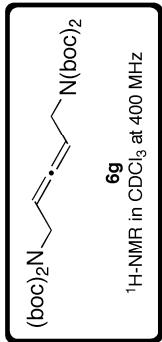
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**JEOL**

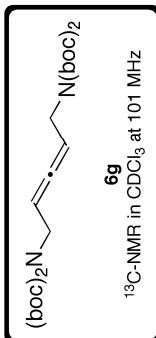


**JEOL**





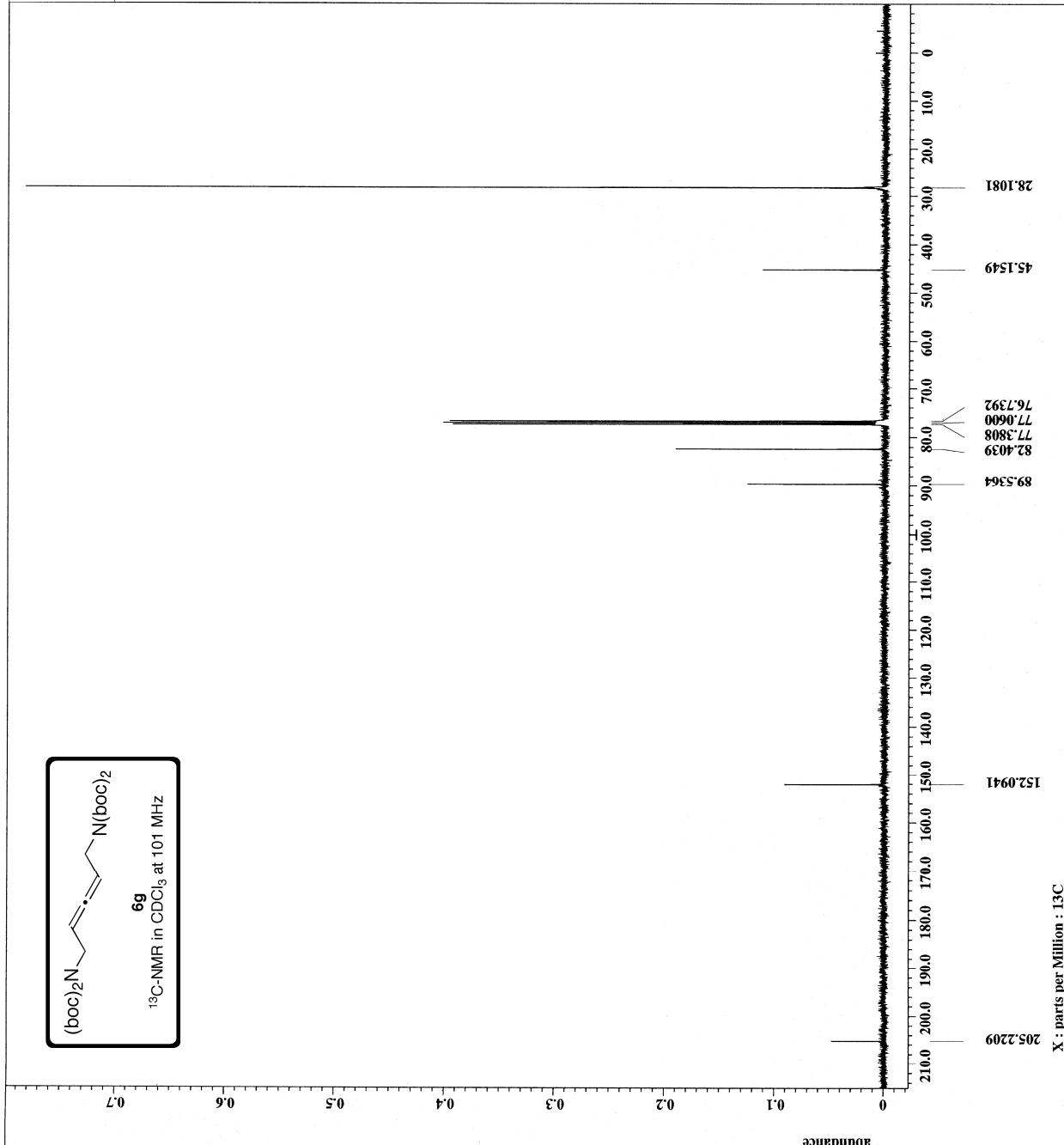
**JEOL**



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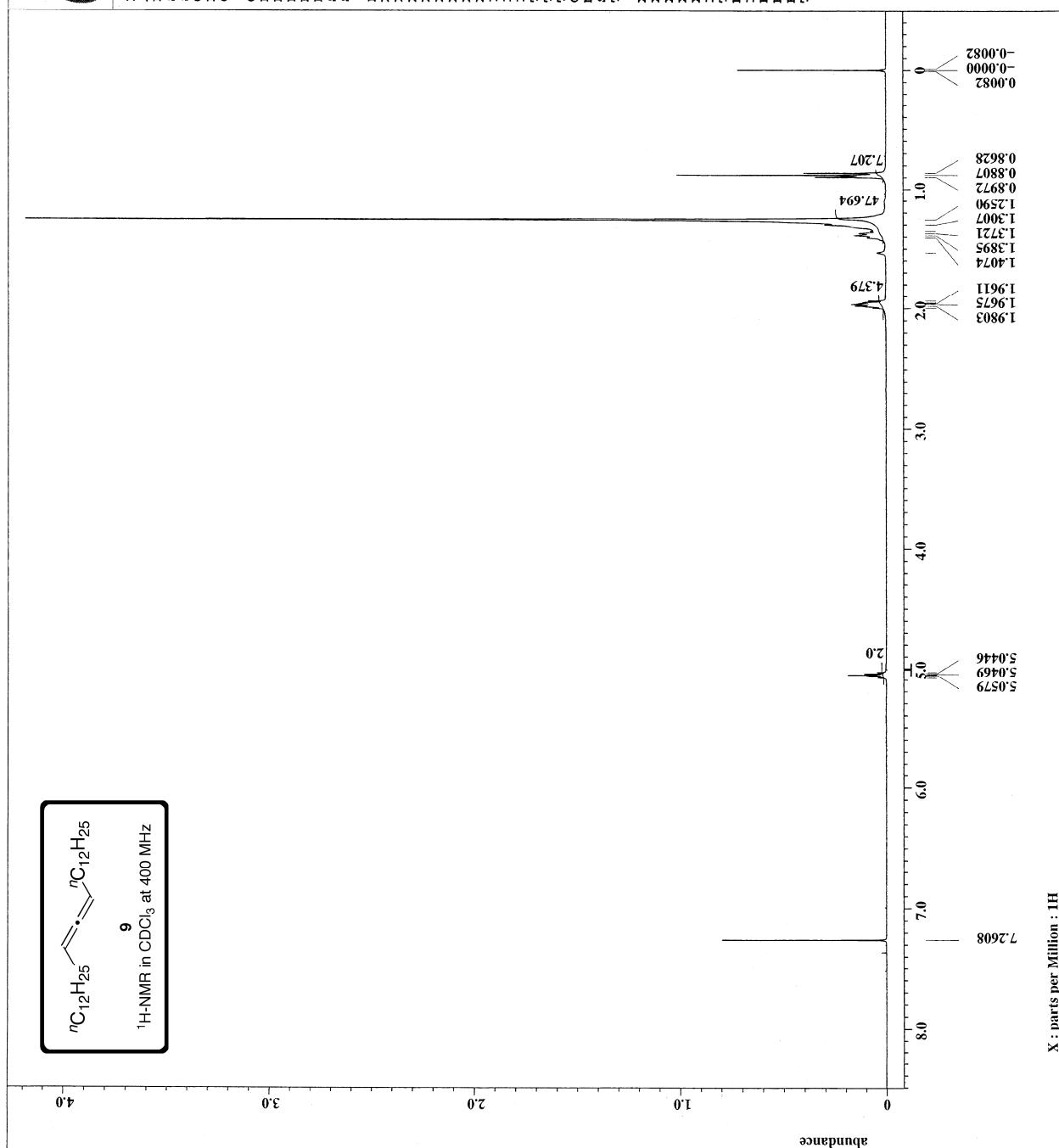
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X_f0eq             = 99.54517646 [MHz]
X_offset           = 100 [ppm]
X_points           = 65516
X_precans          = 1
X_resolution       = 0.43746528 [Hz]
X_sweep            = 28.66572477 [kHz]
Xr_domain         = 1H
Irr_domain        = 395.88430144 [MHz]
Irr_freq           = 5 [ppm]
Irr_offset         = FALSE
Mod_return         = 1
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Total_scans        = 321
X_90_width         = 9.5 [us]
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X_angle            = 30 [deg]
X_dtn              = 8.4 [deg]
X_pulse             = 3.16666667 [us]
Irr_atn_dec        = 29.79 [dB]
Irr_atn_noe        = 29.79 [dB]
Irr_noise           = WALTZ
Decoupling          = TRUE
Initial_wait        = 1 [s]
Noe                = TRUE
Noe_time            = 1.13759 [s]
Revr_Gain           = 58
Relaxation_delay    = 1.13759 [s]
Repetition_time     = 3.42348558 [s]
Temp_get            = 22.9 [dc]

```





**9**  
 $^1\text{H-NMR}$  in  $\text{CDCl}_3$  at 400 MHz



AUJEOL

```

File_name      = MS_111202_1-5.jdf
Author        = delta
Experiment    = single_pulse_ex2
Sample_id     = MS_111202_1
Solvent       = CHLOROFORM-D
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Revision_time = 2-DEC-2011 19:22:16
Current_time  = 2-DEC-2011 19:22:26

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Dim_title     = 1H
Dim_units     = [ppm]
Dimensions    = X
Site          = ECX 400
Spectrometer  = DELTA2_NMR

Field_strength = 9.2982153 [T] (400 [MHz])
X_acq_duration = 5.51550.976 [s]
X_domain      = 1H
X_freq         = 395.88430144 [MHz]
X_offset       = 5 [ppm]
X_points       = 32768
X_prescans    = 1
X_resolution  = 0.181069 [Hz]
X_sweep        = 5.9410646 [Hz]
Irr_domain    = 1H
Irr_freq       = 395.88430144 [MHz]
Irr_offset     = 5 [ppm]
Trl_domain    = 1H
Trl_freq       = 395.88430144 [MHz]
Trl_offset     = 5 [ppm]
Clipped       = FALSE
Mod_return    = 1
Scans          = 16
Total_scans   = 16

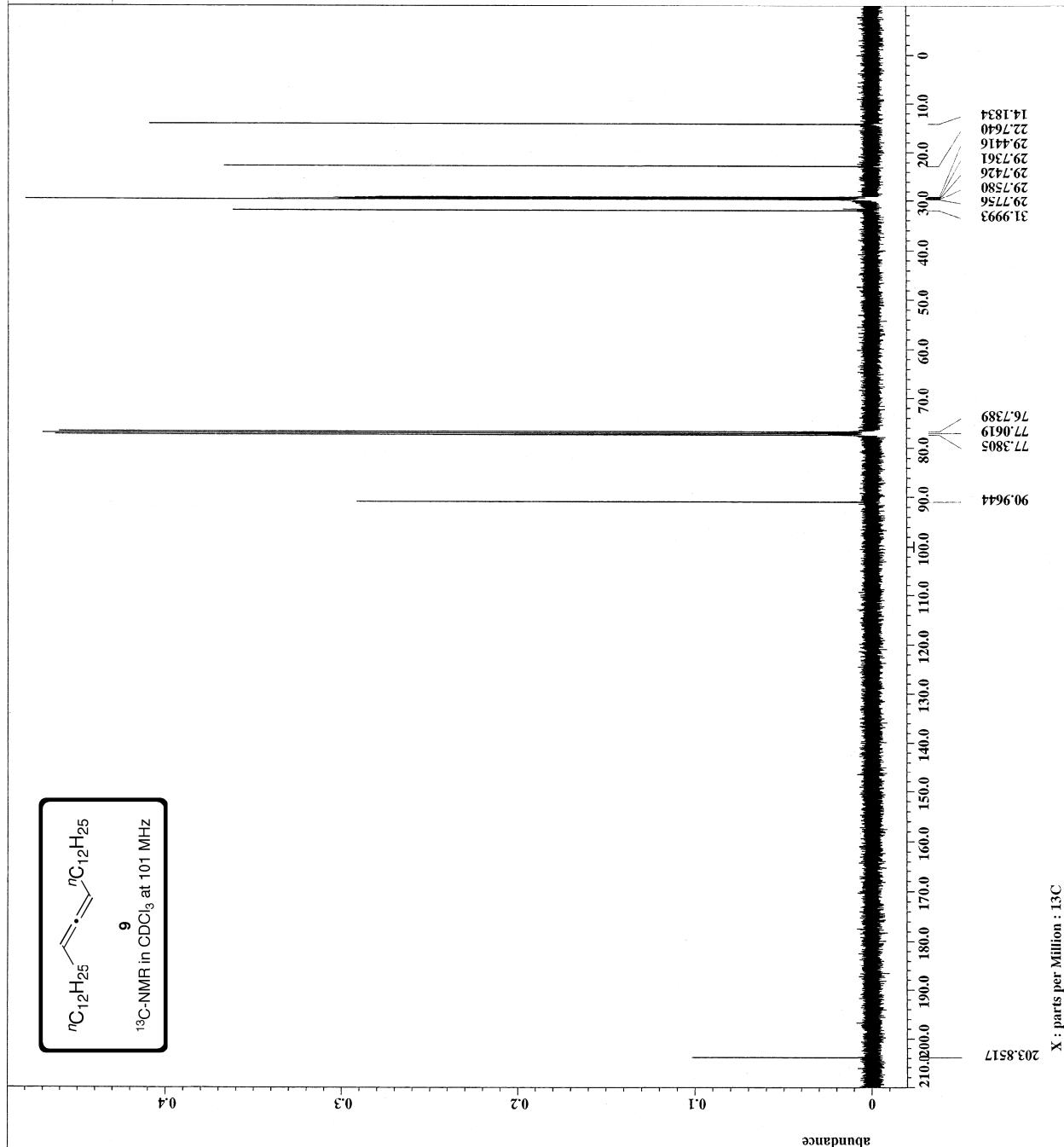
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X_pulse        = 5.25 [us]
Irr_mode       = Off
Trl_mode       = Off
Date_presat   = FALSE
Initial_wait   = 1 [s]
Revr_gain      = 38
Relaxation_delay = 1.60235 [s]
Repetition_time = 7.11769.76 [s]
Temp_get       = 23.9 [dc]

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**JEOL**



**9**  
<sup>13</sup>C-NMR in CDCl<sub>3</sub> at 101 MHz



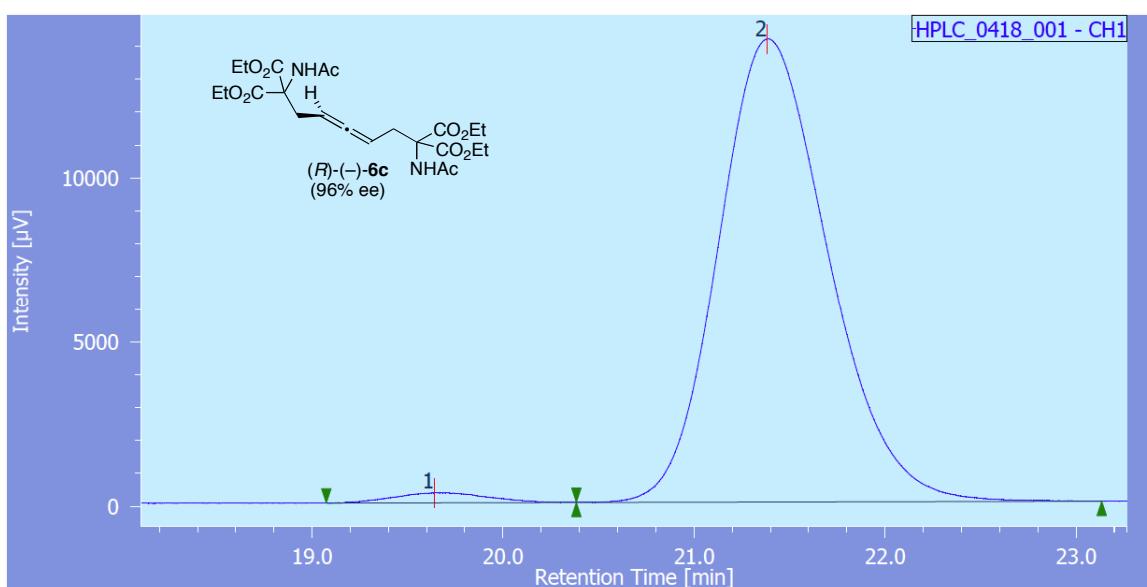
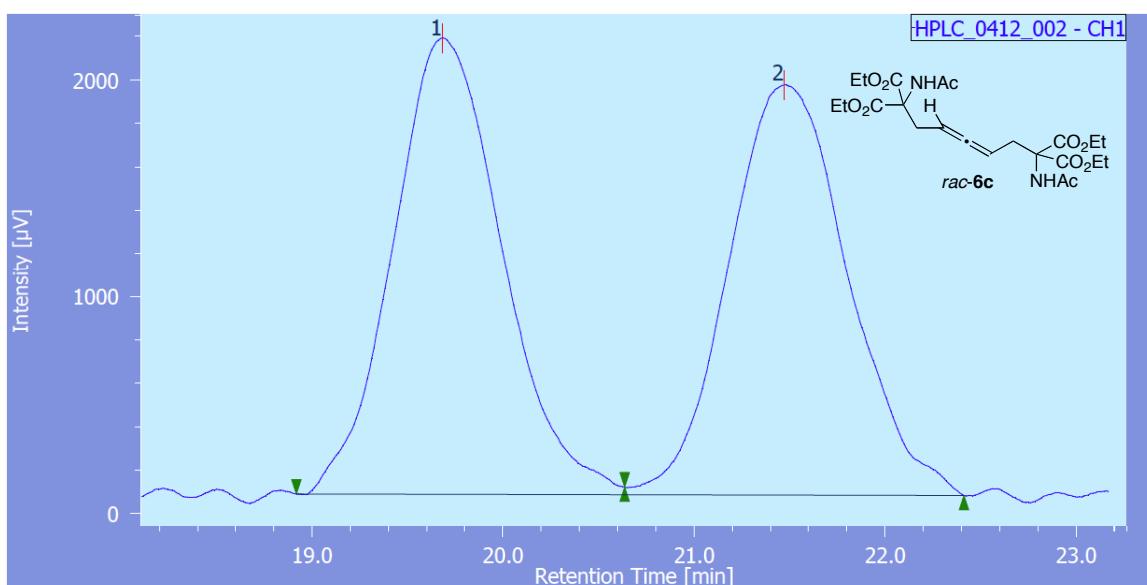
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Current_Time = 2-DEC-2011 20:24:32
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Data_format = 1D COMPLEX
Dim_size = 104857
Dim_title = 13C
Dim_units = [ppm]
Dimensions = X
Site = ECX 400
Spectrometer = DELTA2_NMR
Field_strength = 9.2982453 [T] (400 [MHz])
X_acq_duration = 4.51717136 [s]
X_domain = 13C
X_freq = 99.451766 [MHz]
X_offset = 100 [ppm]
X_points = 131072
X_prescans = 1
X_resolution = 0.21873264 [Hz]
X_sweep = 28.66972477 [Hz]
Irr_domain = 1H
Irr_freq = 395.88430144 [MHz]
Irr_offset = 5 [ppm]
Clipiped = TRUE
Mod_return = 1
Scans = 564
Total_scans = 564
X_90_width = 9.5 [us]
X_acq_time = 4.517136 [s]
X_angle = 30 [deg]
X_att = 8.4 [dB]
X_pulse = 3.1666667 [us]
Irr_att_dec = 29.79 [dB]
Irr_att_noe = 29.79 [dB]
Irr_noise = WALTZ
Decoupling = TRUE
Initial_wait = 1 [s]
No_time = TRUE
No_time = 0 [s]
Recv_gain = 60
Relaxation_delay = 0 [s]
Repetition_time = 4.517136 [s]
Temp_get = 24.1 [deg]

```

### Chiral HPLC Analysis of (*R*)-(-)-6c (Scheme 4).

chiral column: Chiralpak AD-H; eluent: hexane/<sup>i</sup>PrOH = 4/1; flow rate: 0.8 mL/min.



#	peak name	CH	tR [min]	area [µV·sec]	height [µV]	area %	height %	NTP	resolution	symmetry coefficient
1	(S)-isomer	1	19.642	10431	304	1.821	2.113	7068	1.763	1.124
2	(R)-isomer	1	21.383	562317	14083	98.179	97.887	6679	N/A	1.191