

## ELECTRONIC SUPPORTING INFORMATION

### A methodology for radiolabeling of the endocannabinoid 2-arachidonoylglycerol (2-AG)

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(5'*Z*,8'*Z*,11'*Z*,14'*Z*)-lcosa-5',8',11',14'-tetraenoate (**4**, 2-AG borate ester)

1,3-dibutyrate glycerol  
after re-purification  
chloroform

Pulse Sequence: s2pul

Solvent: CDCl<sub>3</sub>

Ambient temperature

INOVA-500 "waters500"

Pulse 42.2 degrees

Acq. time 2.048 sec

Width 8000.0 Hz

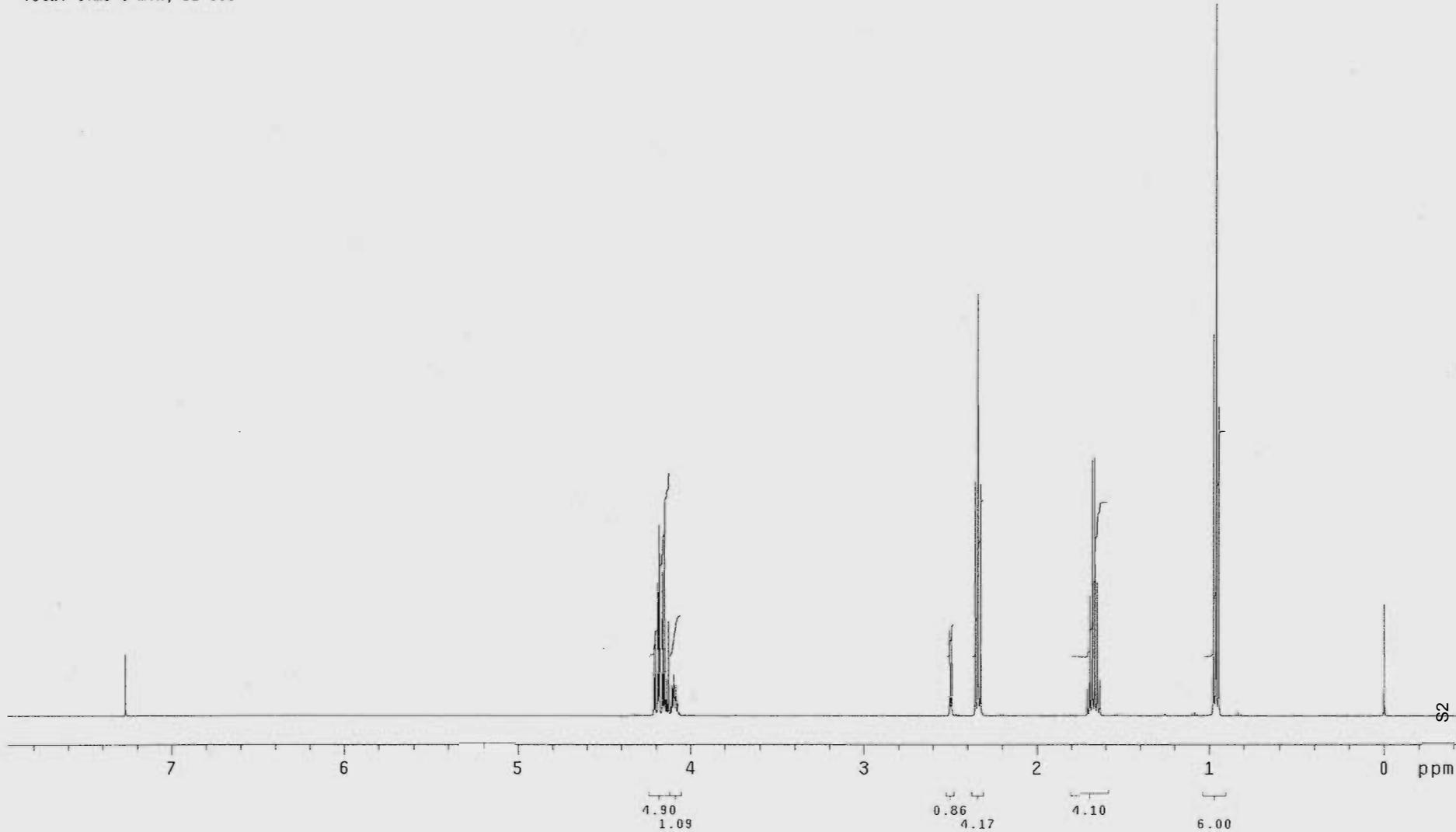
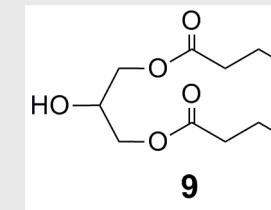
16 repetitions

OBSERVE H1, 499.7029680 MHz

DATA PROCESSING

FT size 32768

Total time 0 min, 32 sec



MJ0668 good

chloroform

Pulse Sequence: s2pul

Solvent: CDCl<sub>3</sub>

Ambient temperature

File: MJ0668good

INOVA-500 "waters500"

Pulse 42.2 degrees

Acq. time 2.048 sec

Width 8000.0 Hz

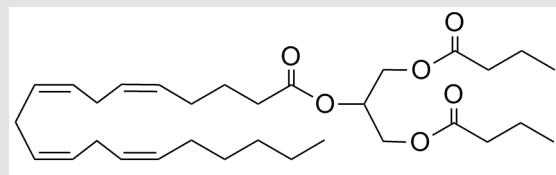
16 repetitions

OBSERVE H1, 499.7029680 MHz

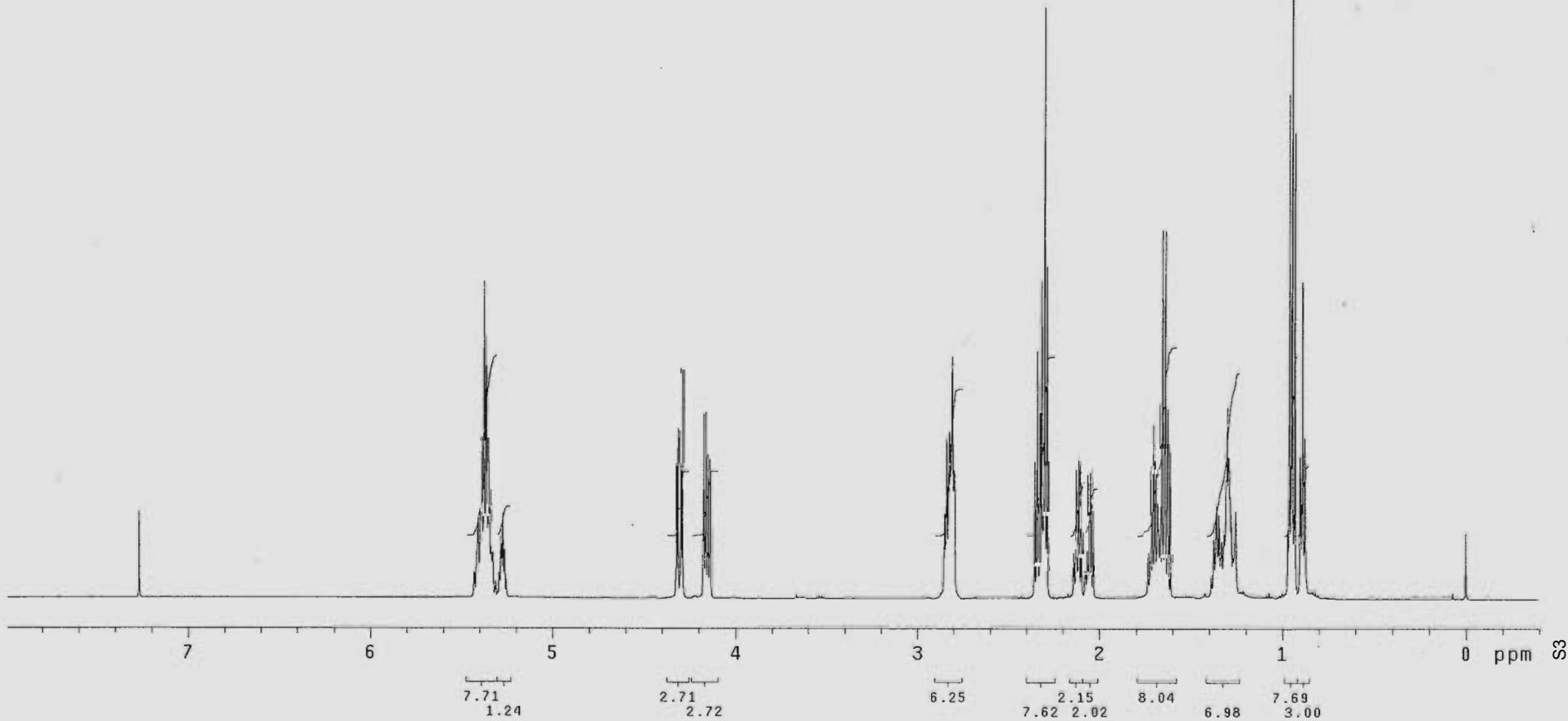
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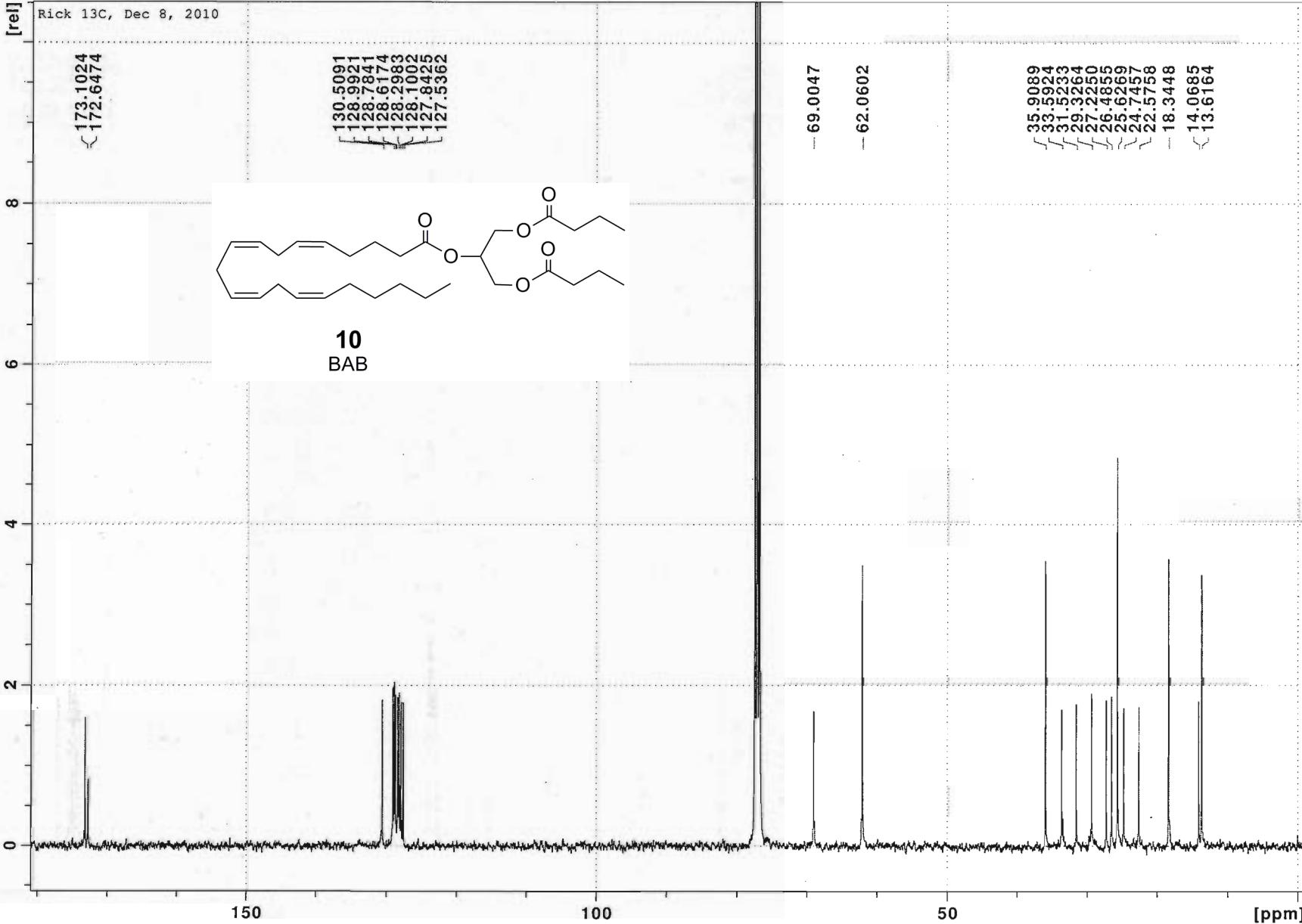


**10**  
BAB



Meghan 1001 1 /opt/topspin tian

Rick 13C, Dec 8, 2010



MJ0660 top spot  
chloroform

Pulse Sequence: s2pul

Solvent: CDCl<sub>3</sub>

Ambient temperature  
File: M10660temp.txt

File: MJ0660topspot  
TNOVA-500 "waters500"

INUVA-500 "waters500"

Pulse 42.2 degrees  
Acc time 3.048 sec

Acq. time 2.048 sec  
Width 8000.0 Hz

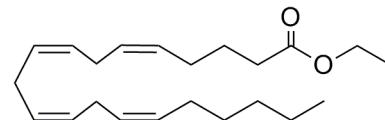
width 8000.0 Hz  
16 repetitions

OBSERVE H1, 499.7029690 MHz  
DATA PROCESSING

DATA PROCESSING  
ET size 32768

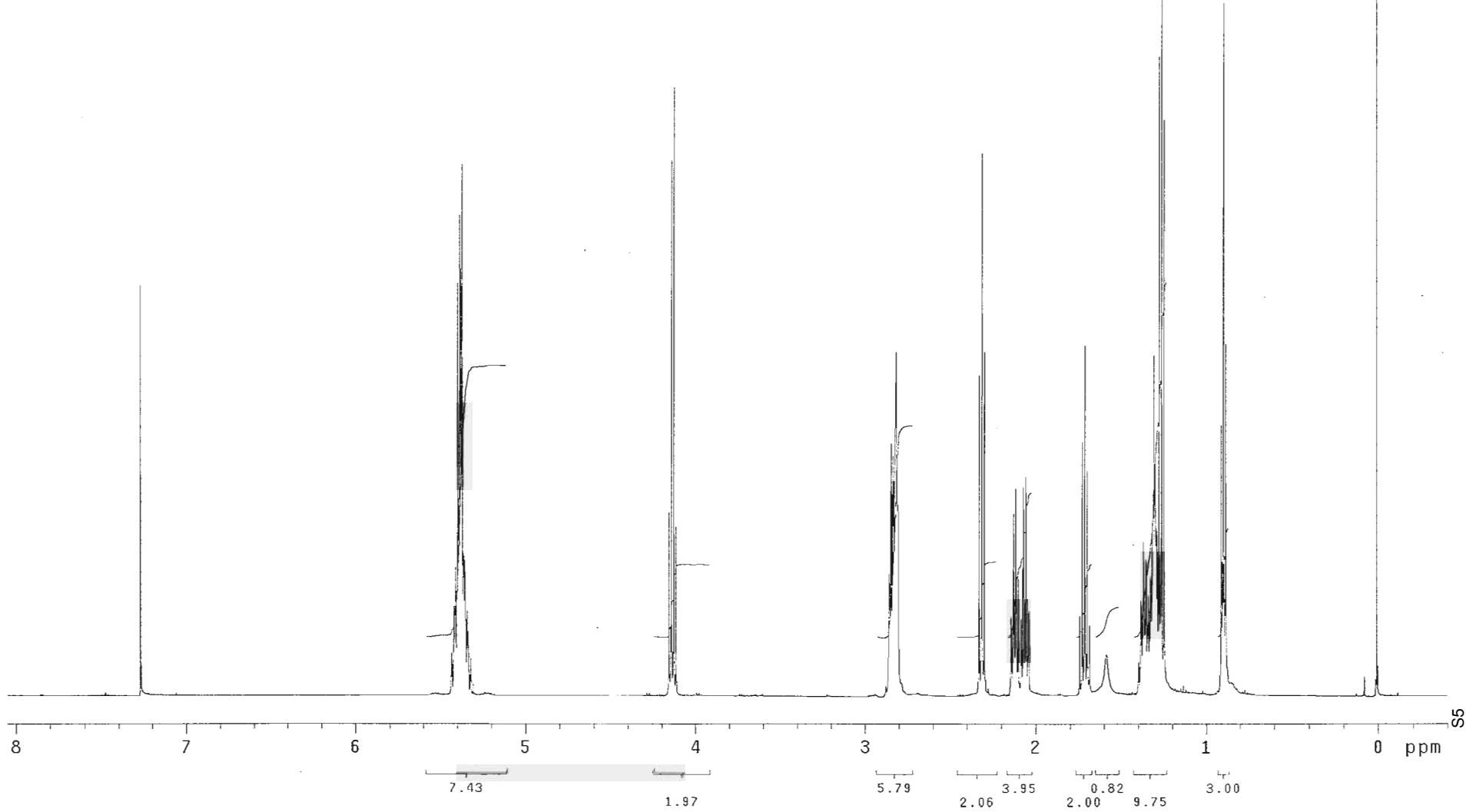
Total time 0 m

Total time 0 min, 32 sec



12

ethyl arachidonate



MJ0684  
fraction 4

chloroform

Pulse Sequence: s2pul

Solvent: CDCl<sub>3</sub>

Ambient temperature

File: MJ0684fraction4

INOVA-500 "waters500"

Pulse 42.2 degrees

Acq. time 2.048 sec

Width 8000.0 Hz

16 repetitions

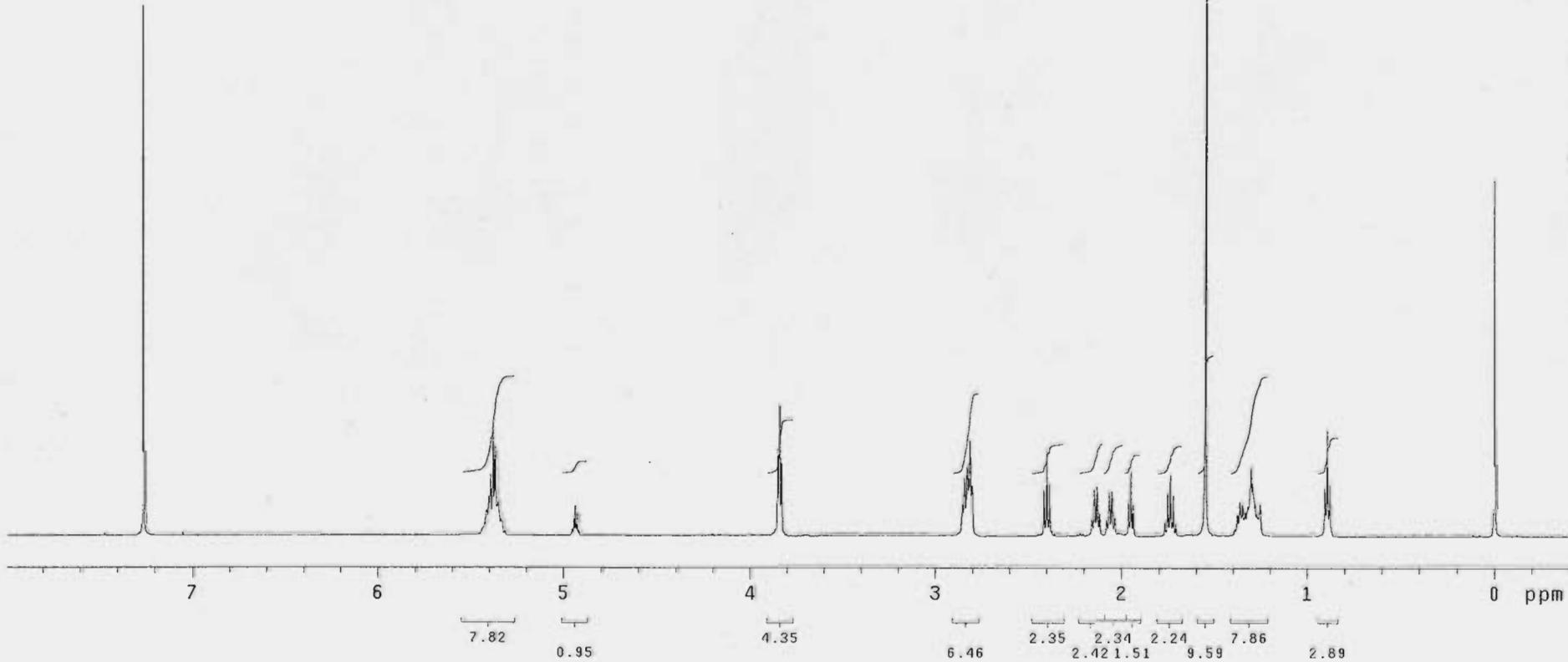
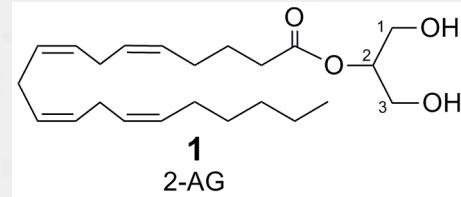
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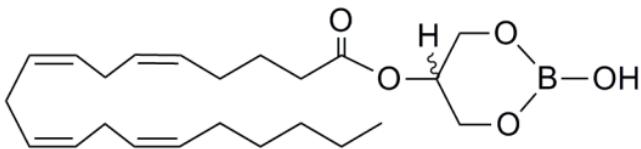
DATA PROCESSING

Line broadening 1.0 Hz

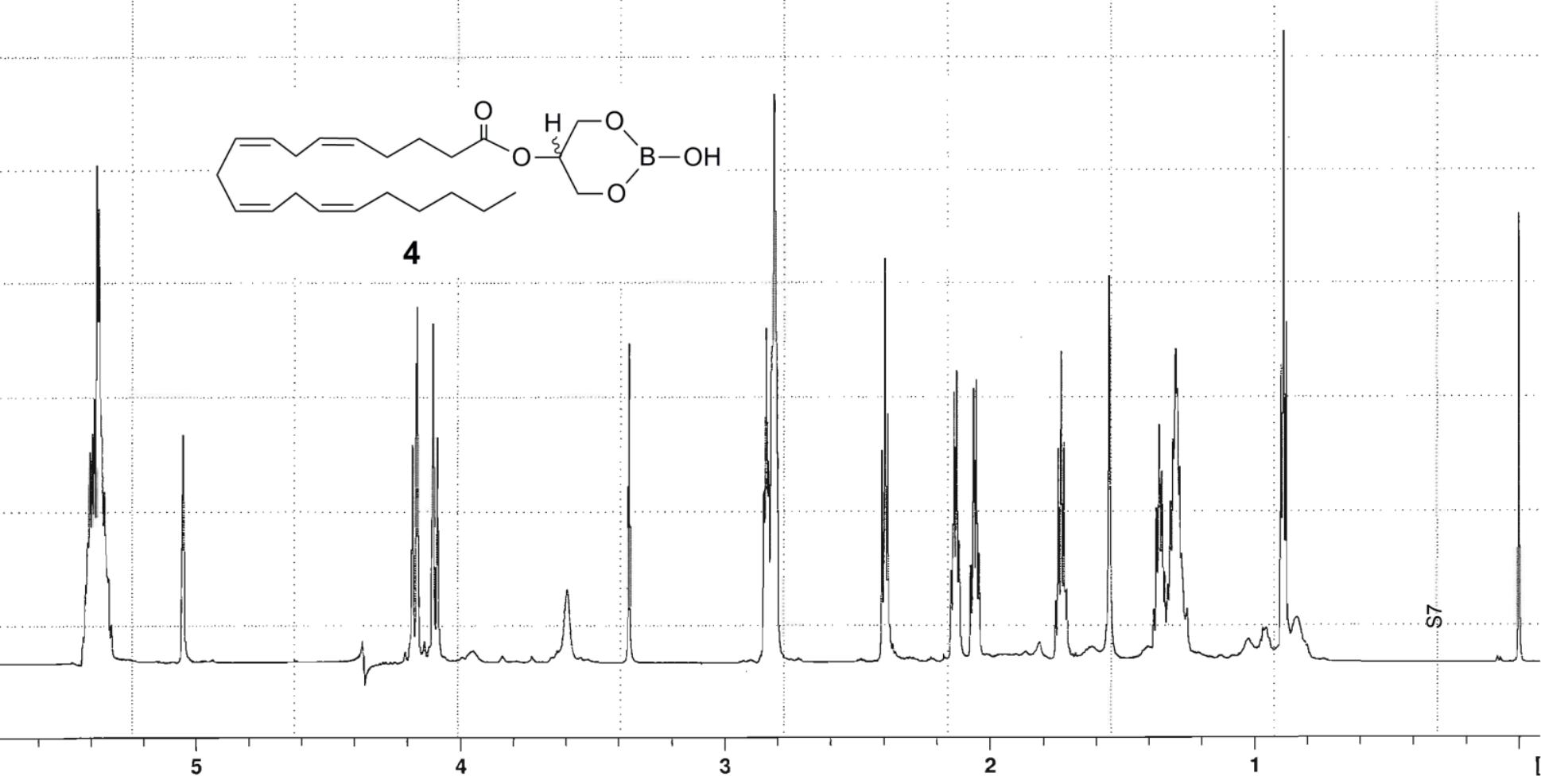
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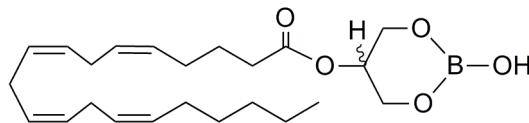


4



57

130.7515  
129.2888  
128.9568  
128.8475  
128.5146  
128.3561  
128.0927  
127.7691



150

100

50

[ppm]

77.4131  
77.3912  
77.0505

67.5347  
65.0307

33.8554  
31.7493  
29.5522  
27.4477  
26.6974  
25.8684  
25.8288  
24.8951  
22.8032

14.3012

1.2 [ref]

- 0.0

0.4

0.6

0.8

1.0

1.2 [ref]

S8