

Name: _____

Chem 203
December 15, 2012

Final Exam Part I
(40 points)

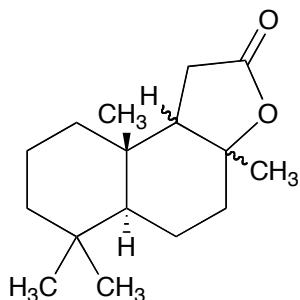
**THIS FINAL EXAM CONSISTS OF PART I
AND TWO OUT OF THE THREE PROBLEMS FROM PART II**

**IF THREE PROBLEMS FROM PART II ARE SUBMITTED,
ONLY THE FIRST TWO (Part II, PROBLEMS 1-2) WILL BE GRADED**

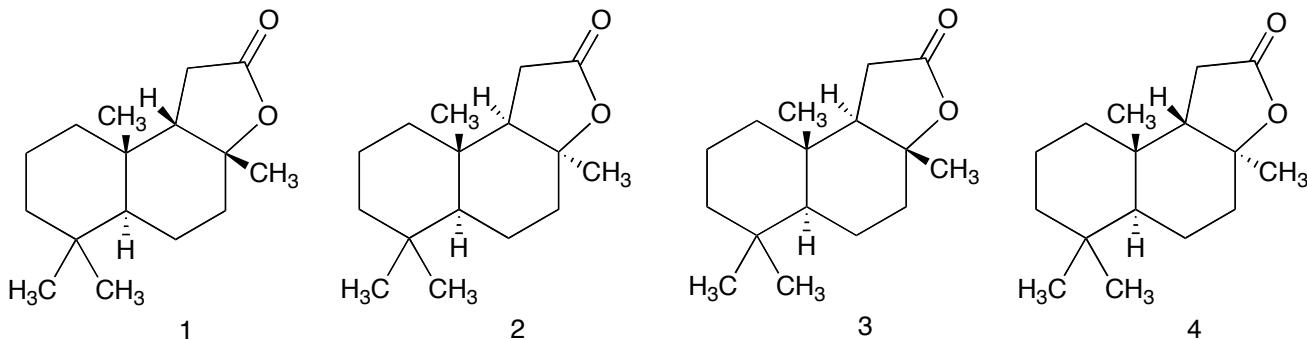
If you wish to unstaple the pages, please initial each page.

Books, notes, lecture videos, calculators, rulers, and laptop computers are permitted as is wireless (or wired) internet access and appropriate software (e.g., PyMOL, Maestro/MacroModel, Excel, ChemDoodle, Chemdraw, ElComp, MolE, etc.). Communication with other students by e-mail, text, or in person is not permitted. Catalogs of molecular structures (e.g., the Aldrich catalog, the Merck Index, etc.) or databases of molecular structures (such as wireless access to SciFinder Scholar, the Sigma-Aldrich website, etc.) are NOT PERMITTED. INAPPROPRIATE COMMUNICATION OR USE OF SUCH ITEMS CONSTITUTES ACADEMIC DISHONESTY, WILL RESULT IN A FAILING GRADE (F) IN THE CLASS, AND MAY RESULT IN EXPULSION FROM THE Ph.D. PROGRAM.

The following spectral data are provided for a tricyclic compound: 500.22 MHz ^1H NMR, 125.79 MHz ^{13}C NMR, DEPT, COSY, TOCSY, HMQC, HMBC, NOESY, and HSQC-TOCSY spectra with 5-, 10-, 20-, and 100-ms mixing times. All NMR spectra were measured in C_6D_6 solution.



Using these data, determine the stereochemistry and assign all of the ^1H and ^{13}C resonances to their respective atoms in the structure. Specifically, assign the stereochemistry of the two stereocenters shown with squiggly lines, and hence which of the four possible diastereomers (**1**, **2**, **3**, or **4**) is consistent with these data.



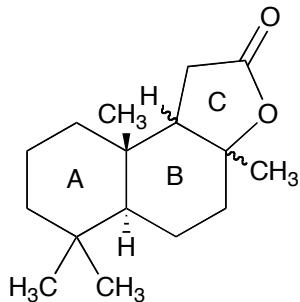
MAKE SURE TO COMPLETELY ANSWER THE QUESTIONS **a-g** ON PAGES 2-6.

a. Build energy-minimized molecular models of diastereomers **1**, **2**, **3**, and **4** using PyMOL and the "clean" function. Rotate each molecule into an appropriate orientation. Use the side menu (S) to show each molecule as sticks (Show-sticks). Use the pulldown menu to display each molecule in maximum quality with (Display-Quality-Maximum Quality). Save the .pse files as diastereomer1.pse, diastereomer2.pse, diastereomer3.pse, and diastereomer4.pse.

Feel free to use the *trans*-decalin template .pse file or .pdb file on the course web page "Simple Conformational Analysis of Cyclic and Bicyclic Compounds" which is linked to the "Assignments" and "Class Materials" web pages. You are also welcome to just make your own *trans*-decalin.

NOTE: For diastereomer **4** you will have to build the B ring in a boat conformation, because the B–C ring junction does not allow the B ring to adopt a chair conformation. After minimization, the B ring should be in a twist-boat conformation. The A–B ring junction will look like that of the other diasteromers, but the B–C ring junction will be enforcing a twist-boat conformation of the B ring.

If you would like to review the conformations of cyclohexane, please see my Chem H52A course web page "Cyclohexanes" at: <https://eee.uci.edu/10f/40600/cyclohexane.html>. The page contains links to .pdb files of boat and twist-boat cyclohexane, which can be opened in PyMOL. If you would like to see an example of a related exercise in which ring fusion induces a twist boat conformation in a cyclohexane ring, please see my course web page "Perhydroanthracenes" and the associated .pdb files at: <https://eee.uci.edu/10f/40600/perhydroanthracenes.html>.



E-mail the .pse files to me (jsnowick@uci.edu).

b. Examine the ^1H NMR spectra and familiarize yourself with the resonances, which have been lettered *a*–*l* for you. Examine the ^{13}C NMR spectra and number the sixteen unique resonances associated with the molecule 1–16.

Examine the DEPT spectra and identify the quaternary (C), methine (CH), methylene (CH₂), and methyl (CH₃) peaks. NOTE: Although the DEPT 90 contains small resonances associated with the methyl and methylene groups, it is easy to identify the methine groups.

Examine the HMQC spectra and correlate the numbers of ^{13}C resonances with the letters of the ^1H resonances.

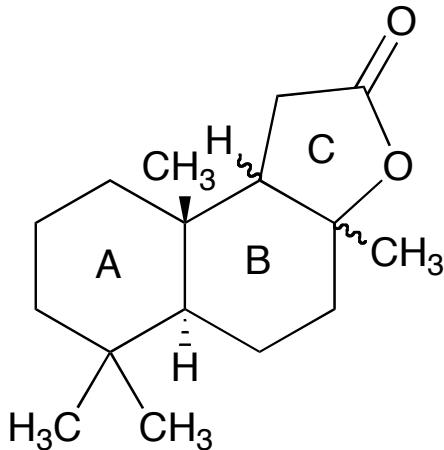
Finally, examine the HSQC-TOCSY spectrum with 100-ms mixing time and identify which methine and methylene resonances are associated with the three main spin systems in the molecule, which we will call the A-ring spin system, the B-ring spin system, and the C-ring spin system:

Numbers associated with the ^{13}C resonances associated with the A-ring spin system: ___, ___, ___.

Numbers associated with the ^{13}C resonances associated with the B-ring spin system: ___, ___, ___.

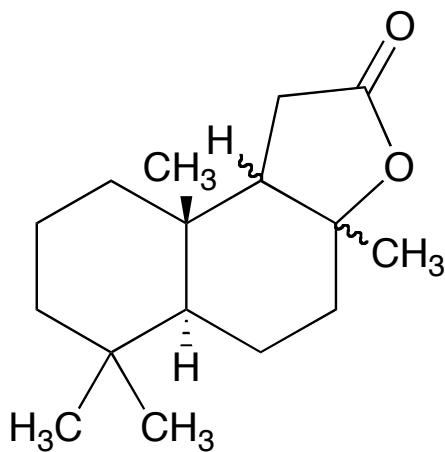
Numbers associated with the ^{13}C resonances associated with the C-ring spin system: ___, ___.

c. Examine the HSQC-TOCSY spectra with 5, 10, 20, and 100-ms mixing times. Assign the *five* methine and methylene ^{13}C resonances from the B- and C-ring spin systems, to the corresponding atoms in the structure, below. That is, write the number next to the atom in the structure below.



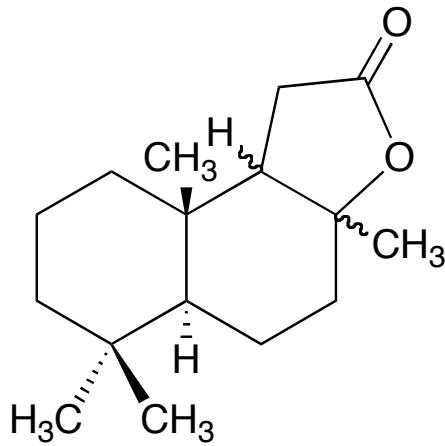
NOTE: Work the problem the way we did in class and is described in the assigned reading in "Two-Dimensional NMR Spectroscopy: Applications for Chemists and Biochemists." Start with the track that grows in under ^1H resonances *a* and *b* and assign one spin system. Then work with the track that grows in under ^1H resonance *c* and assign another spin system.

d. Use the HMBC spectrum and particularly the expansion that gives the higher-level contours of the isolated methyl groups to assign the remaining ^{13}C resonances to the corresponding atoms in the structure, below. That is, write the number next to the atom in the structure below.



e. Use the NOESY spectrum to stereospecifically assign the diastereotopic geminal dimethyl ^{13}C resonances to the corresponding atoms in the structure, below. That is, write the number next to the atom in the structure below. HINT: Consider the proximity of the methyl groups to each other; use the models that you have built in Part **a** for insights.

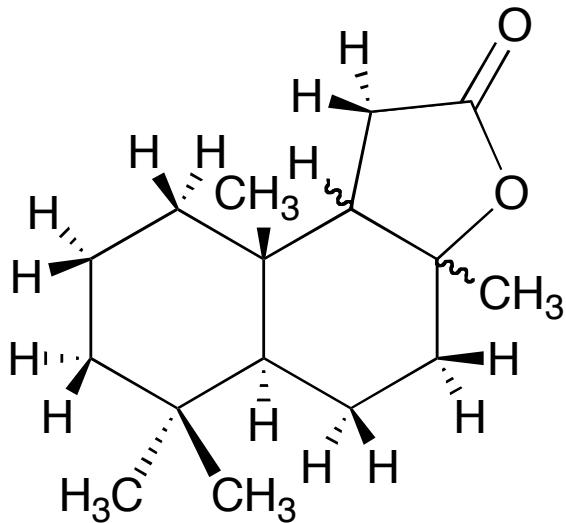
Transcribe your other numbers from Parts **c** and **d**, so that every carbon is numbered. Write the letters of the corresponding ^1H resonances next to the numbers.



f. Determine the stereochemistry of the molecule. Which diastereomer is it? _____

Explain how you determined the stereochemistry of the molecule. Make a conformationally realistic drawing of the molecule to help aid in your explanation. If there are any aspects of the stereochemistry you are uncertain about, please include this in your explanation.

g. Insofar as possible, assign the ^1H resonances to the corresponding atoms in the structure, below. That is, write the letter (a–l) next to the atom in the structure below. You will not likely be able to assign them all. Explain briefly about those that you are uncertain.



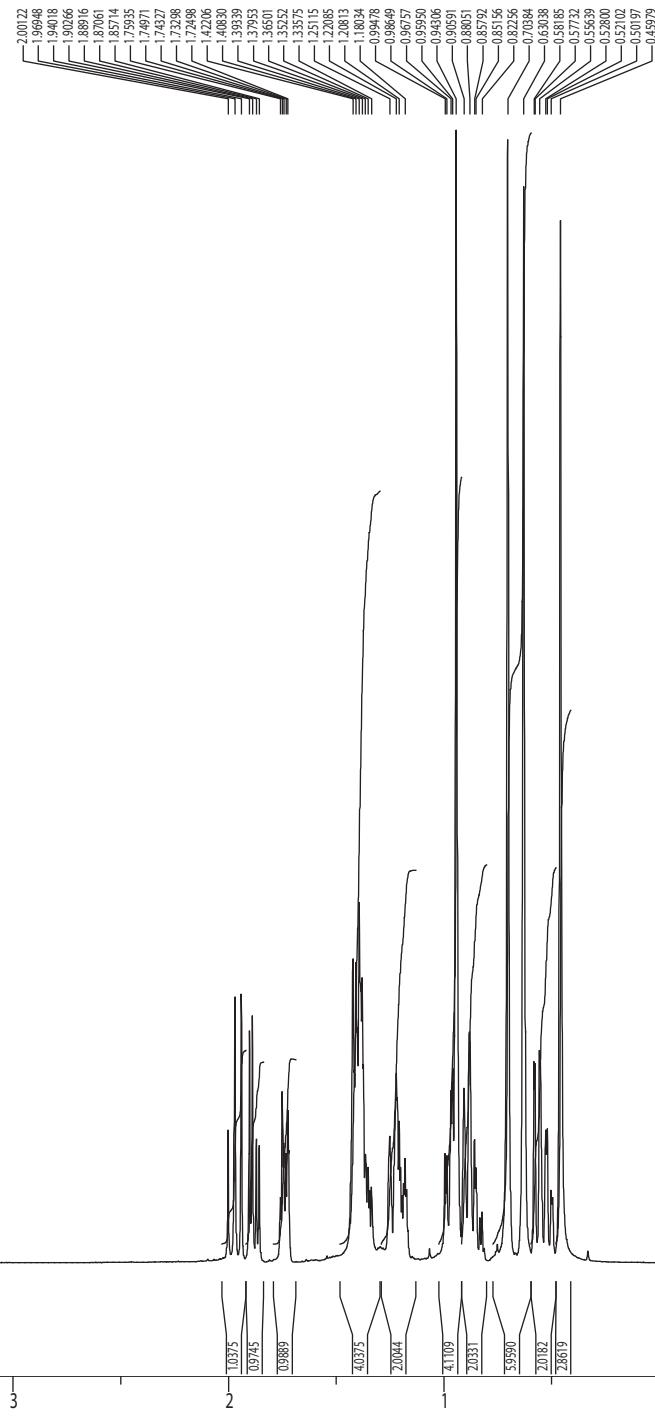
¹H spectrum

ppm

7.1600

Integral

ppm



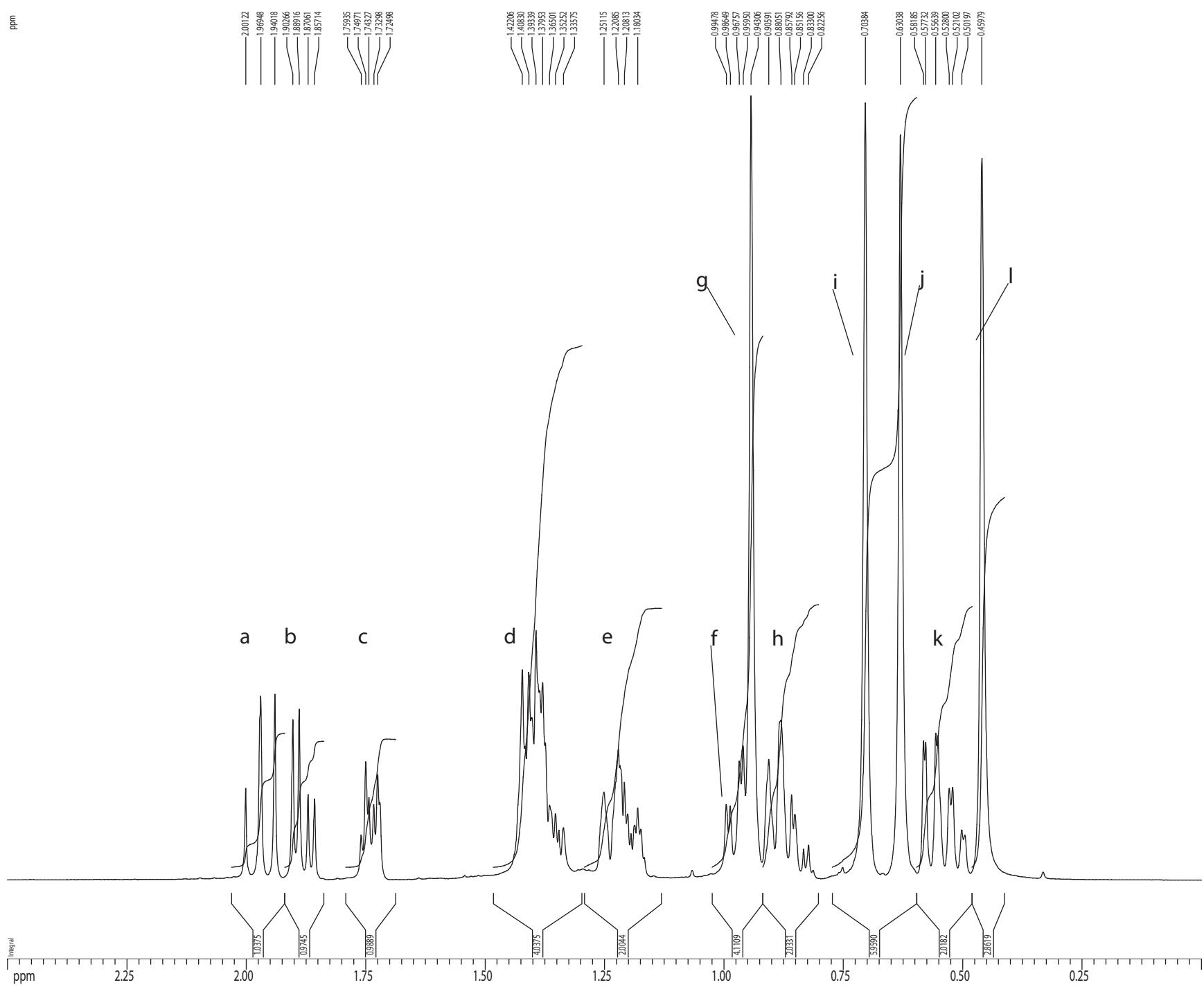
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4	53467.1	970.516	1.9402	3.56
5	53620.6	951.747	1.9027	3.08
6	53675.8	944.998	1.8892	3.27
7	53751.7	935.718	1.8706	1.64
8	53806.8	928.978	1.8571	1.56
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11	54272.7	872.018	1.7433	1.59
12	54314.8	866.869	1.7330	1.45
13	54347.5	862.871	1.7250	2.02
14	55586.9	711.343	1.4221	4.02
15	55643.1	704.461	1.4083	3.98
16	55704.1	697.002	1.3934	4.78
17	55760.8	690.071	1.3795	3.78
18	55820.3	682.803	1.3650	1.44
19	55871.4	676.555	1.3525	1.27
20	55940.0	668.169	1.3357	1.01
21	56286.1	625.849	1.2511	1.68
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23	56462.1	604.332	1.2081	1.88
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26	57368.9	493.461	0.9865	1.42
27	57446.3	483.997	0.9676	2.28
28	57479.3	479.960	0.9595	2.58
29	57546.6	471.736	0.9431	15.00
30	57698.5	453.156	0.9059	2.32
31	57802.5	440.450	0.8805	3.06
32	57894.9	429.147	0.8579	1.63
33	57920.9	425.965	0.8516	1.26
34	57996.8	416.683	0.8330	0.60
35	58039.6	411.462	0.8226	0.68
36	58525.3	352.074	0.7038	14.86
37	58825.8	315.331	0.6304	14.24
38	59024.4	291.052	0.5818	2.67
39	59042.9	288.788	0.5773	2.64
40	59128.5	278.316	0.5564	2.82
41	59244.7	264.115	0.5280	1.76
42	59273.2	260.624	0.5210	1.78
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¹H spectrum

ppm

Integral



¹H spectrum

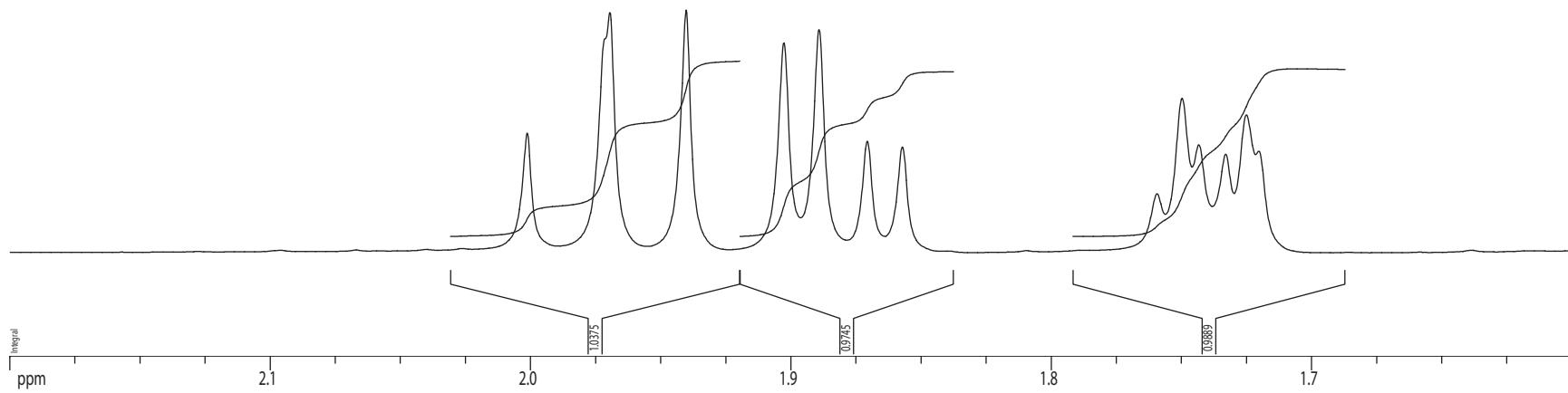
ppm



a

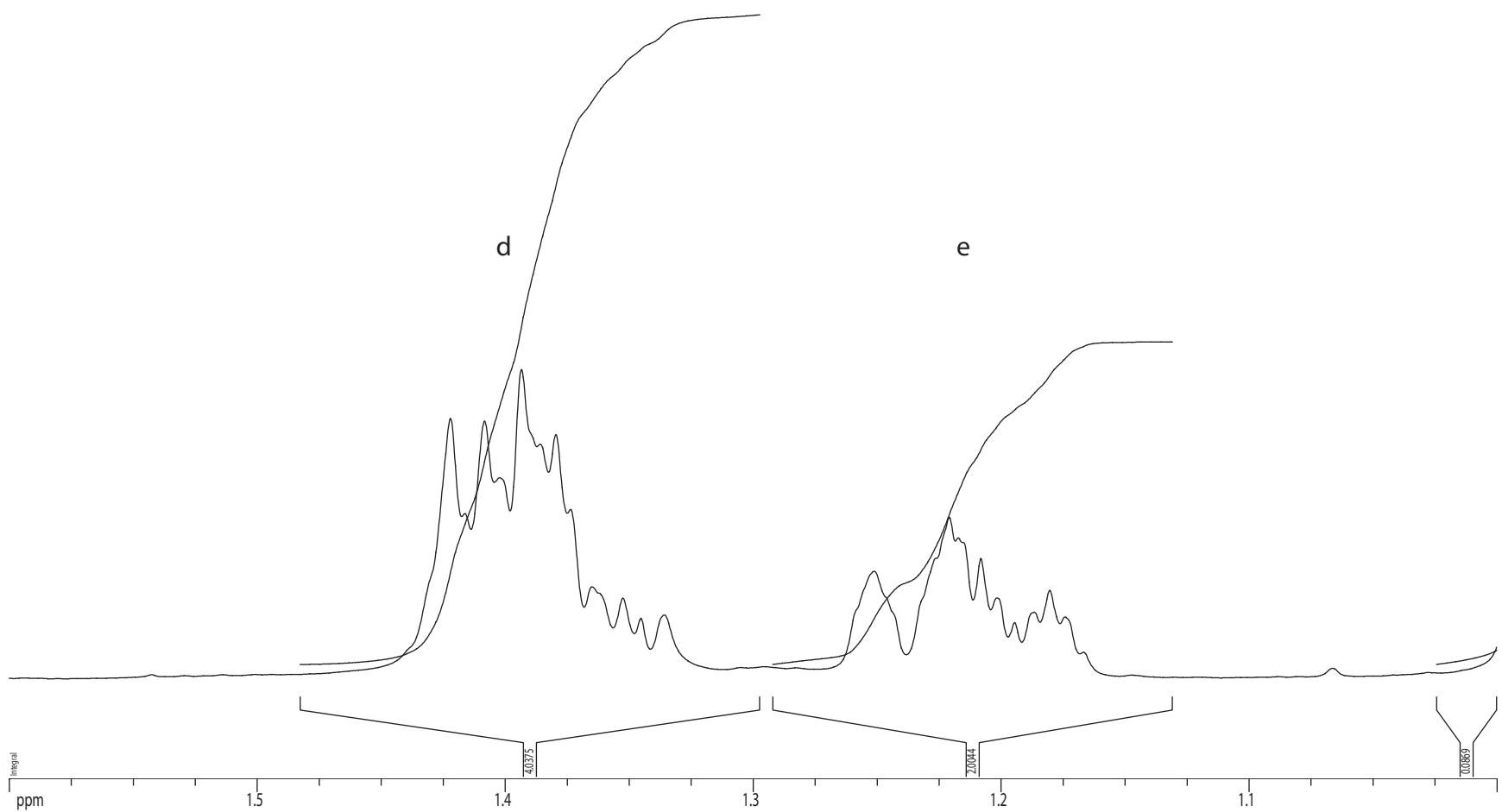
b

c



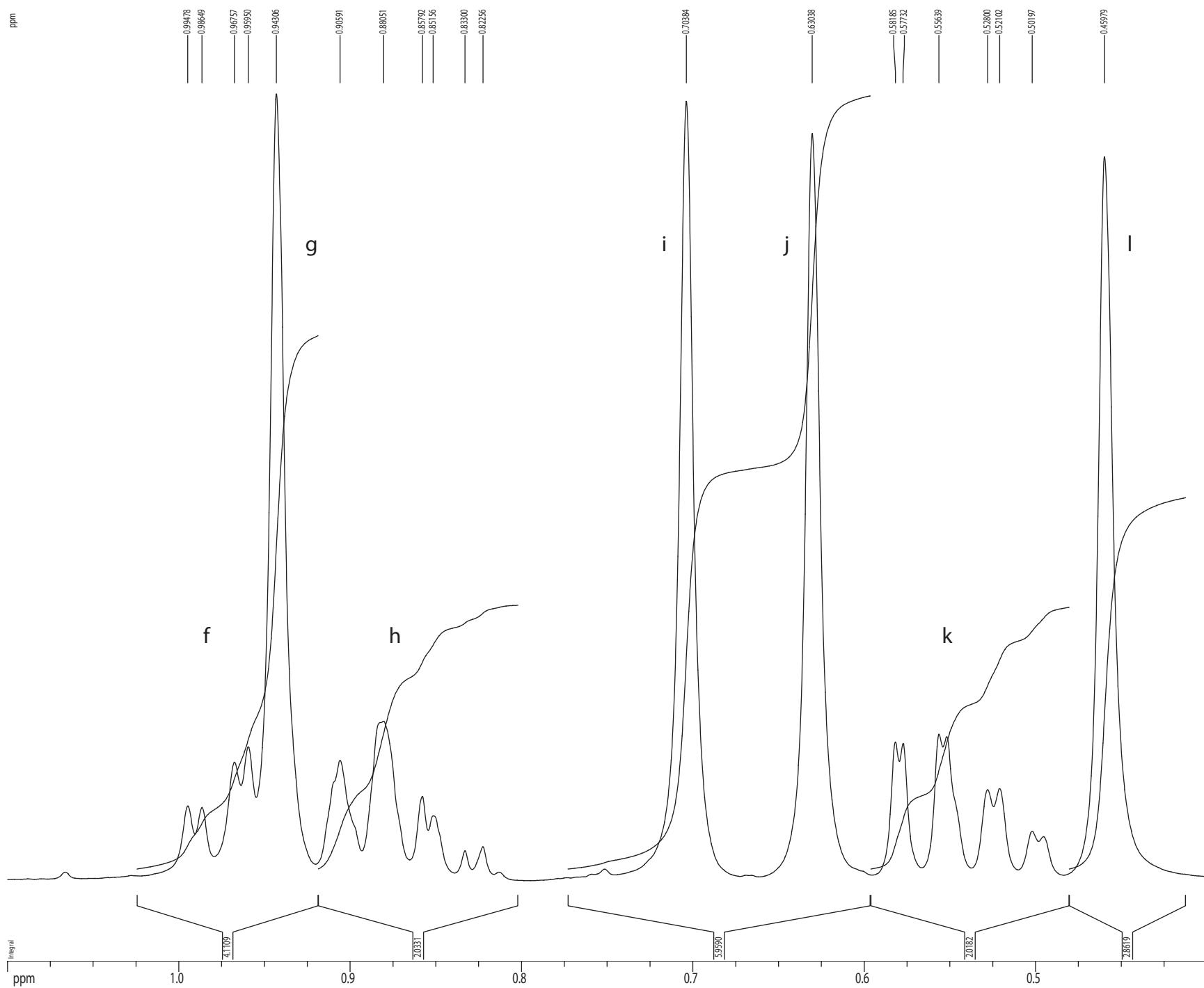
¹H spectrum

ppm

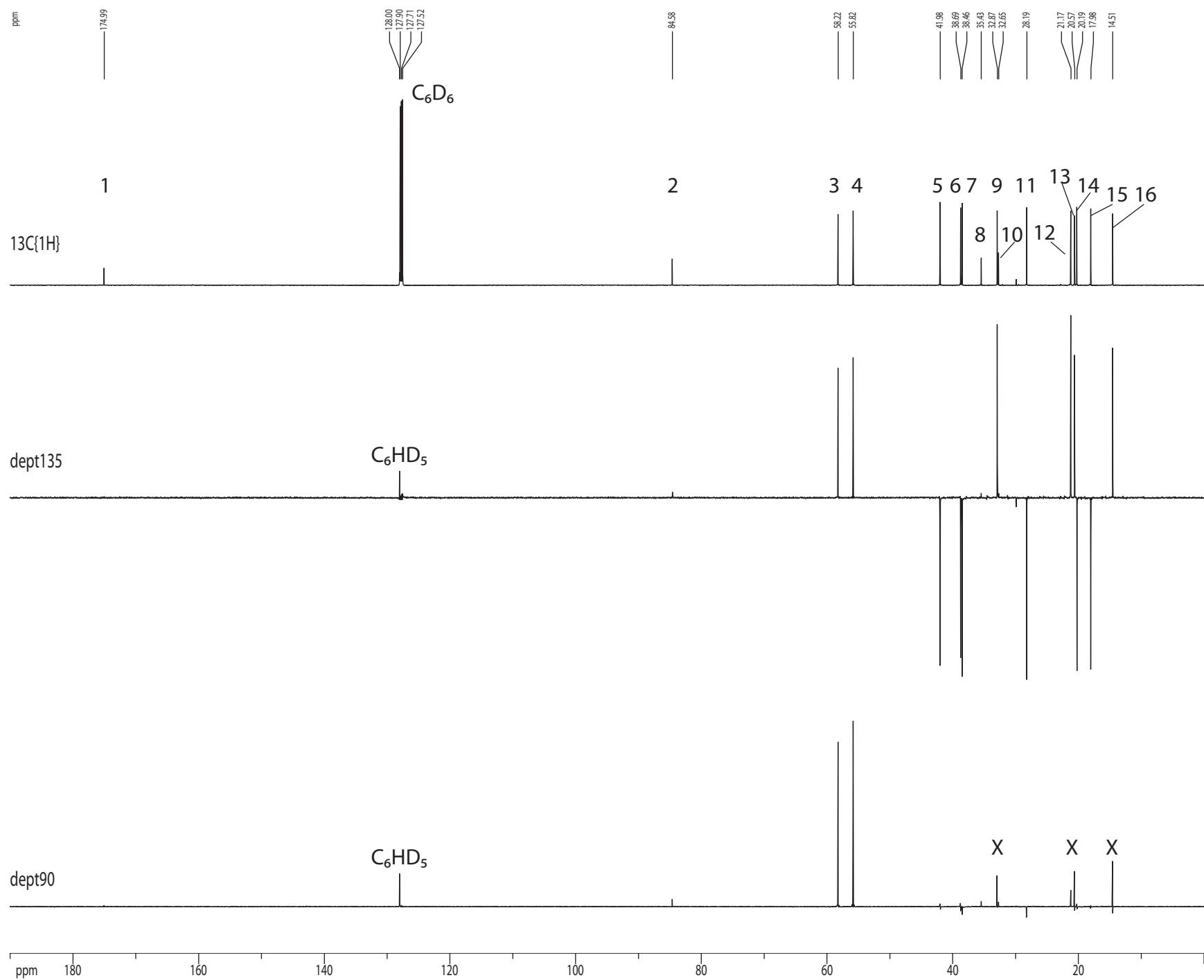


¹H spectrum

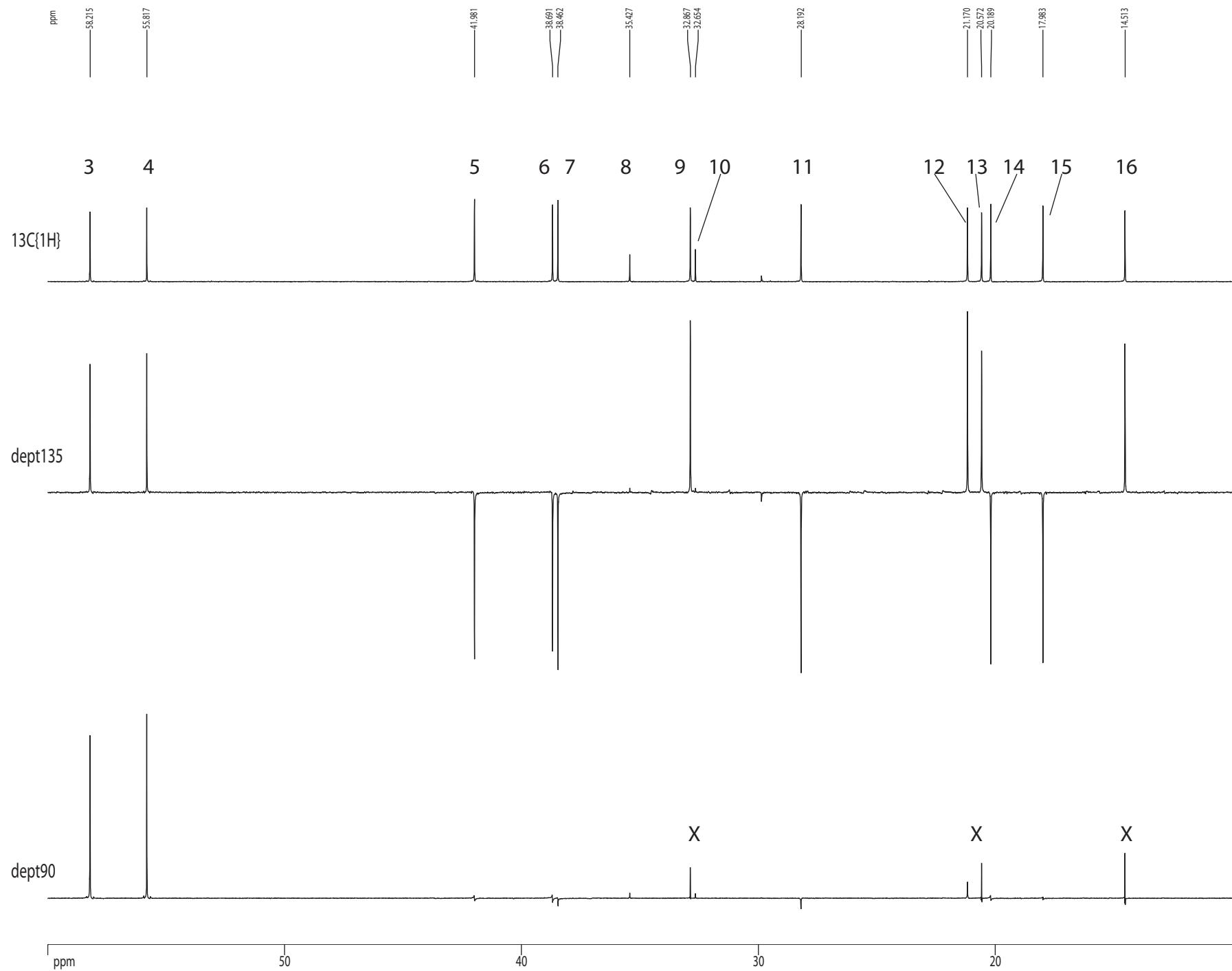
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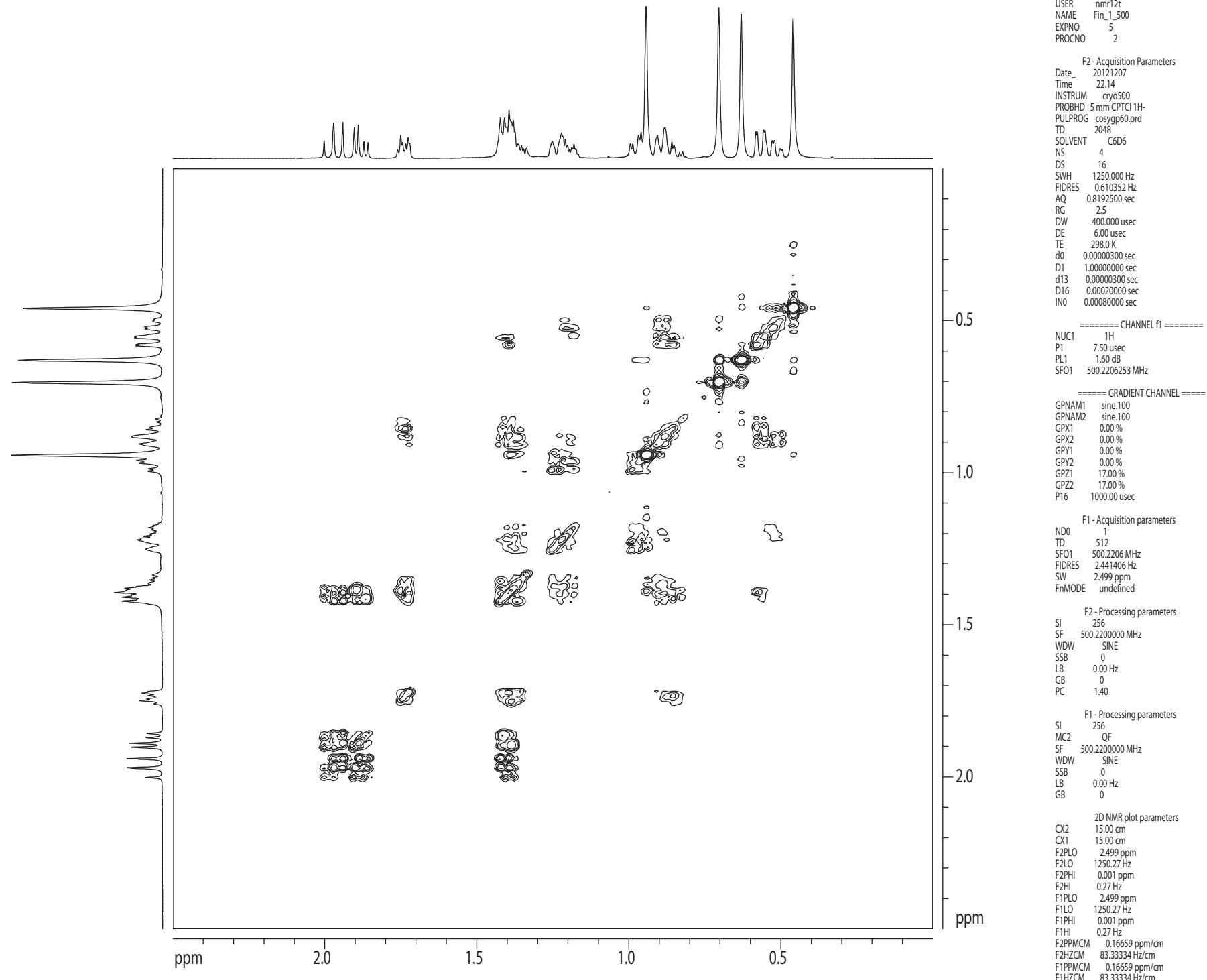
Z-restored spin-echo ^{13}C spectrum with ^1H decoupling



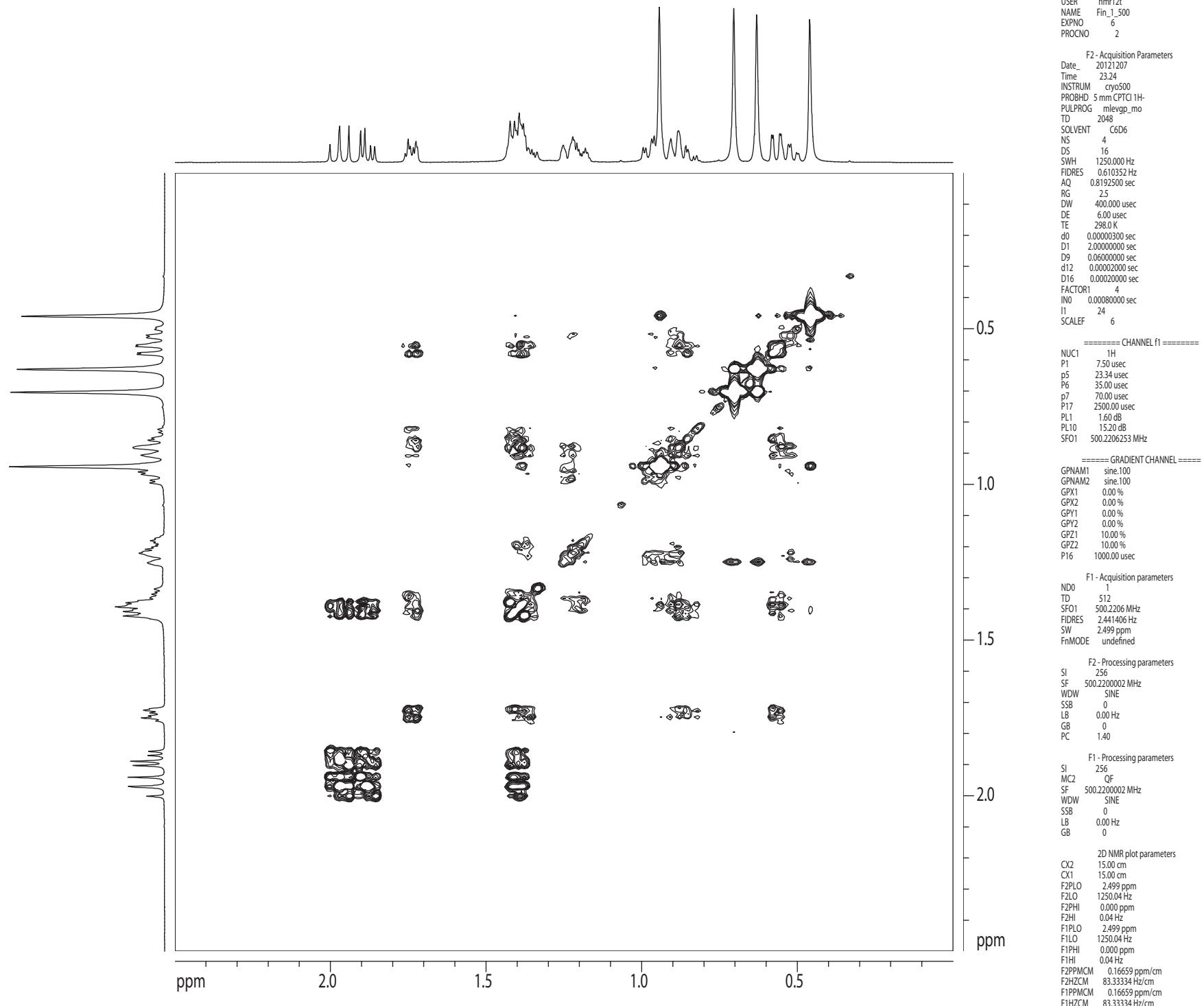
Z-restored spin-echo ^{13}C spectrum with ^1H decoupling



COSY

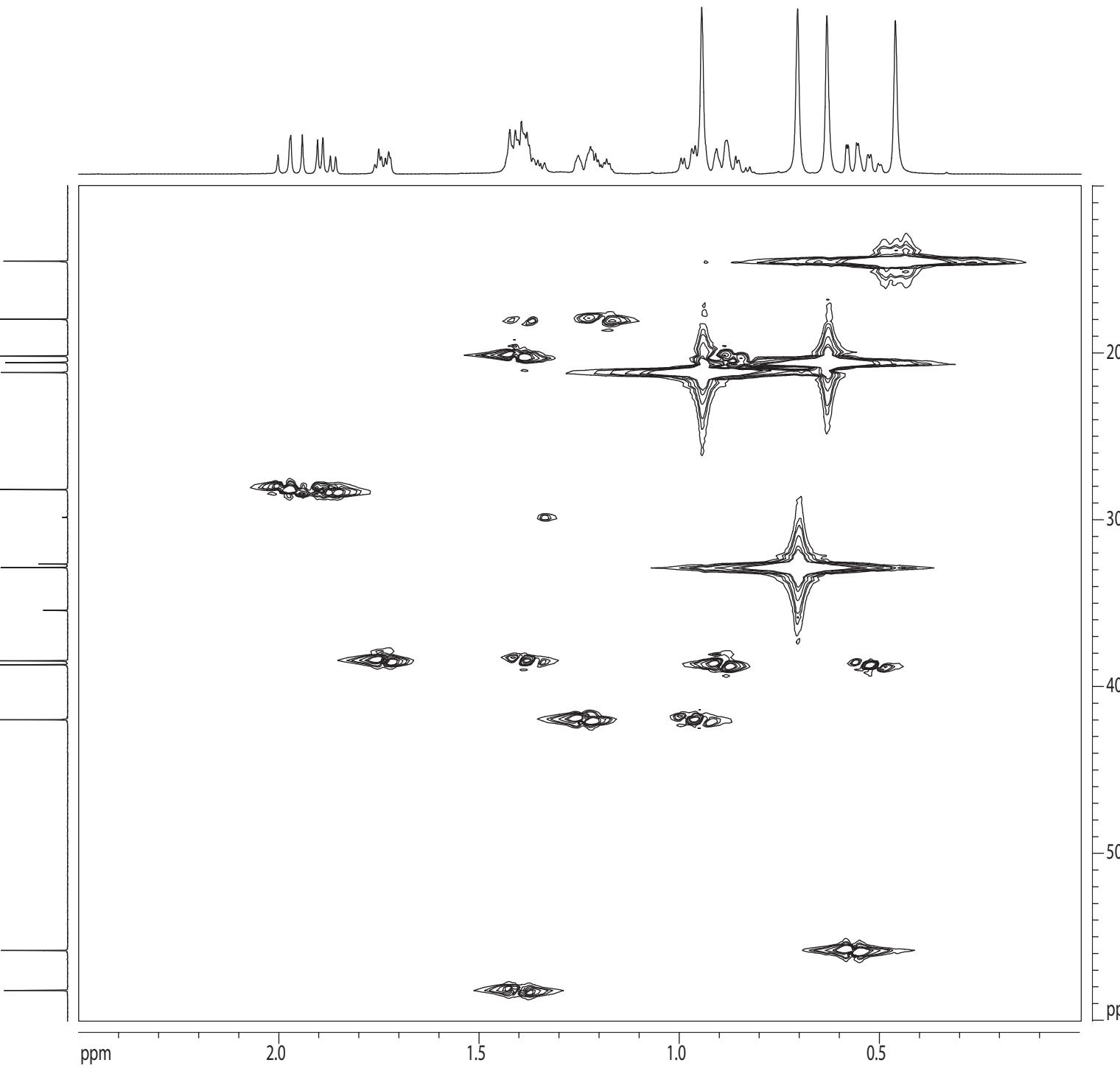


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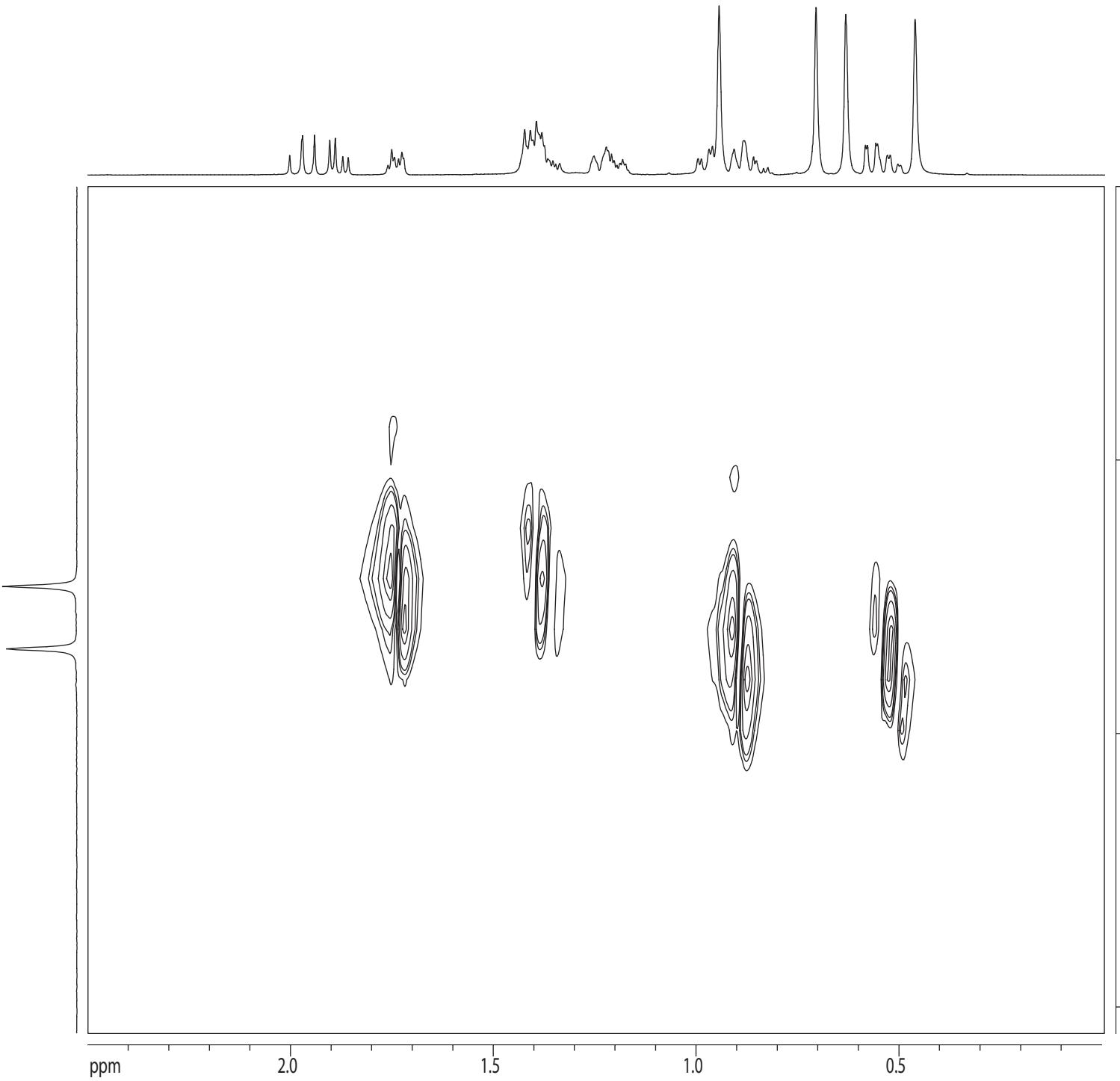
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ghmqc



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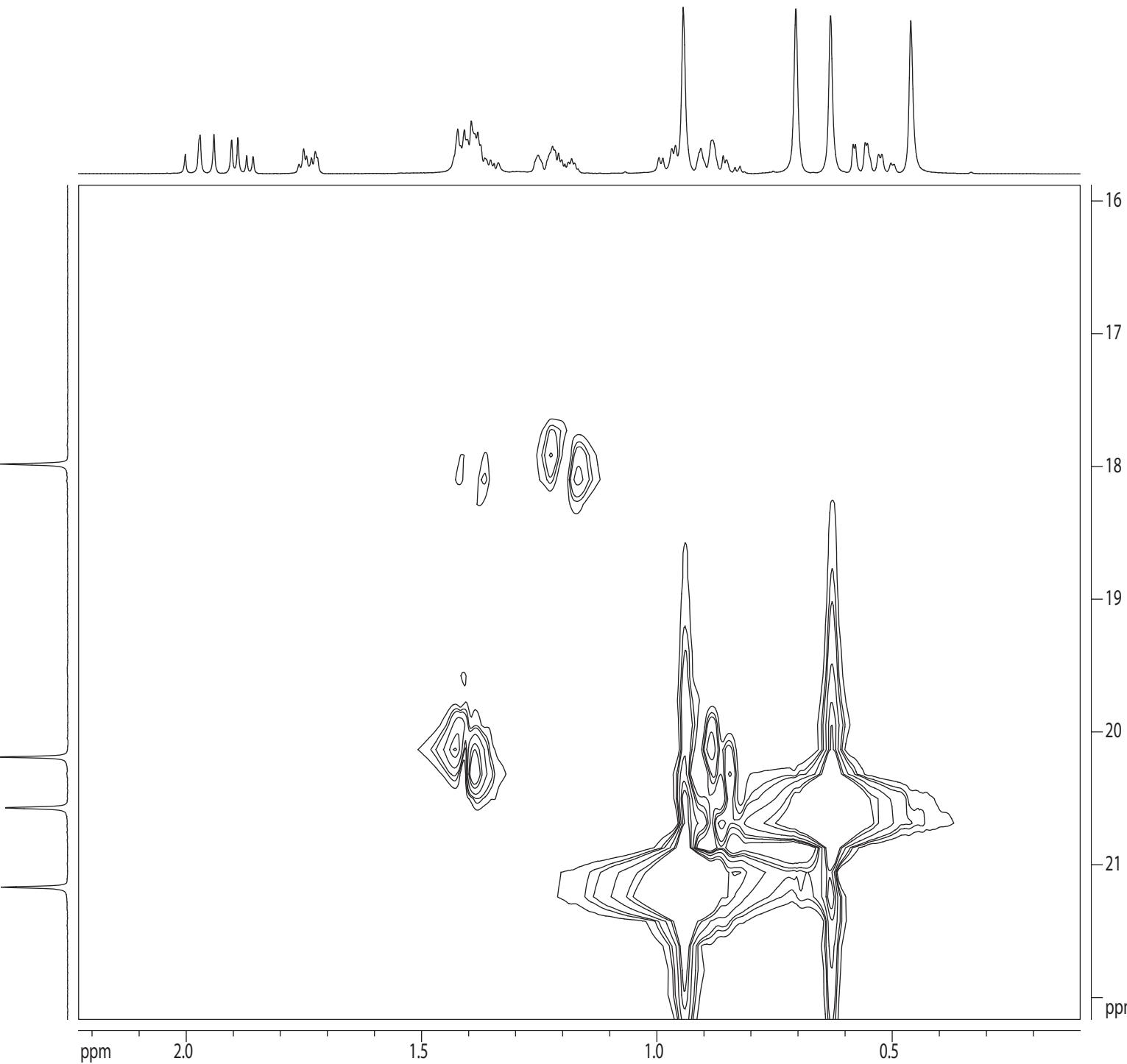
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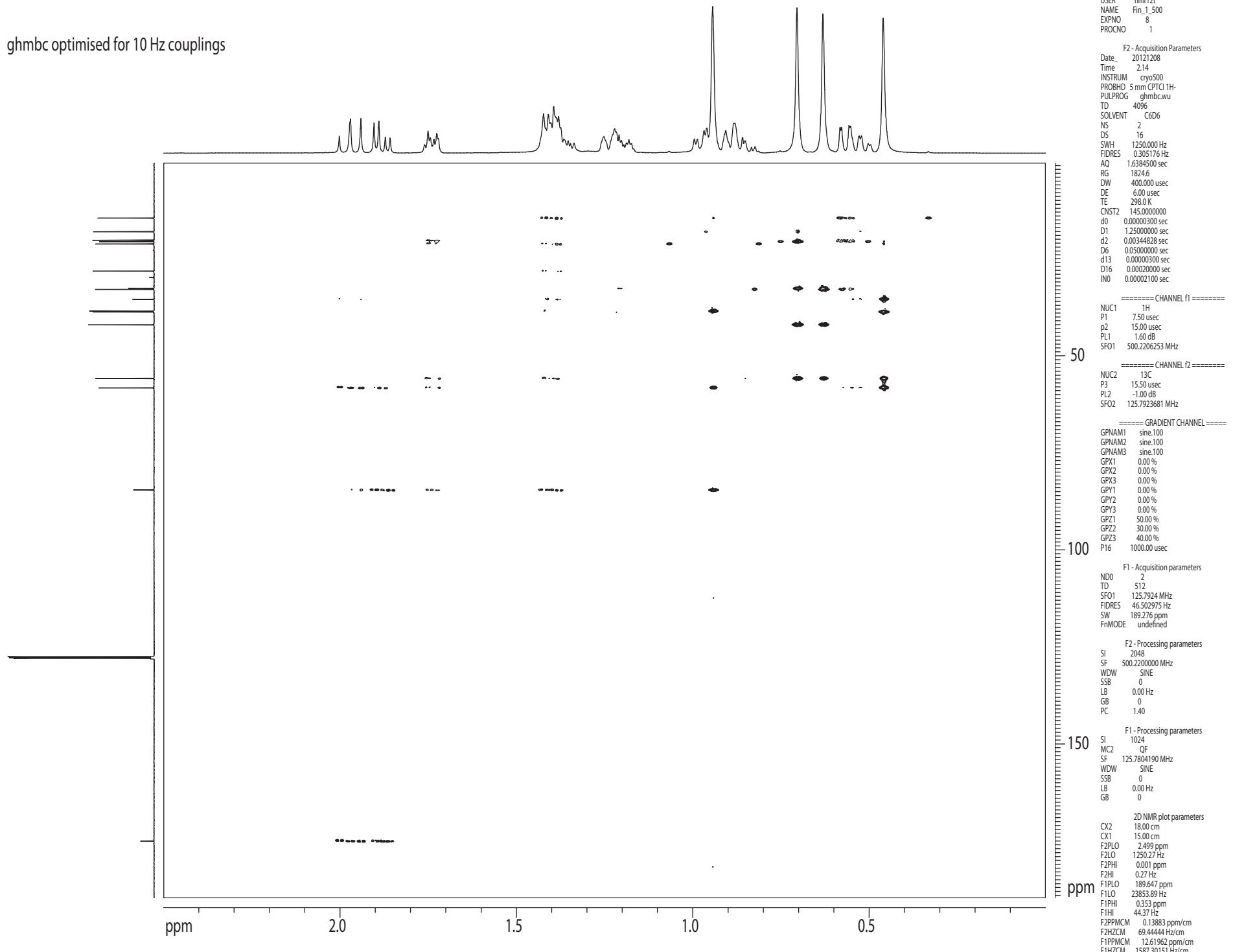
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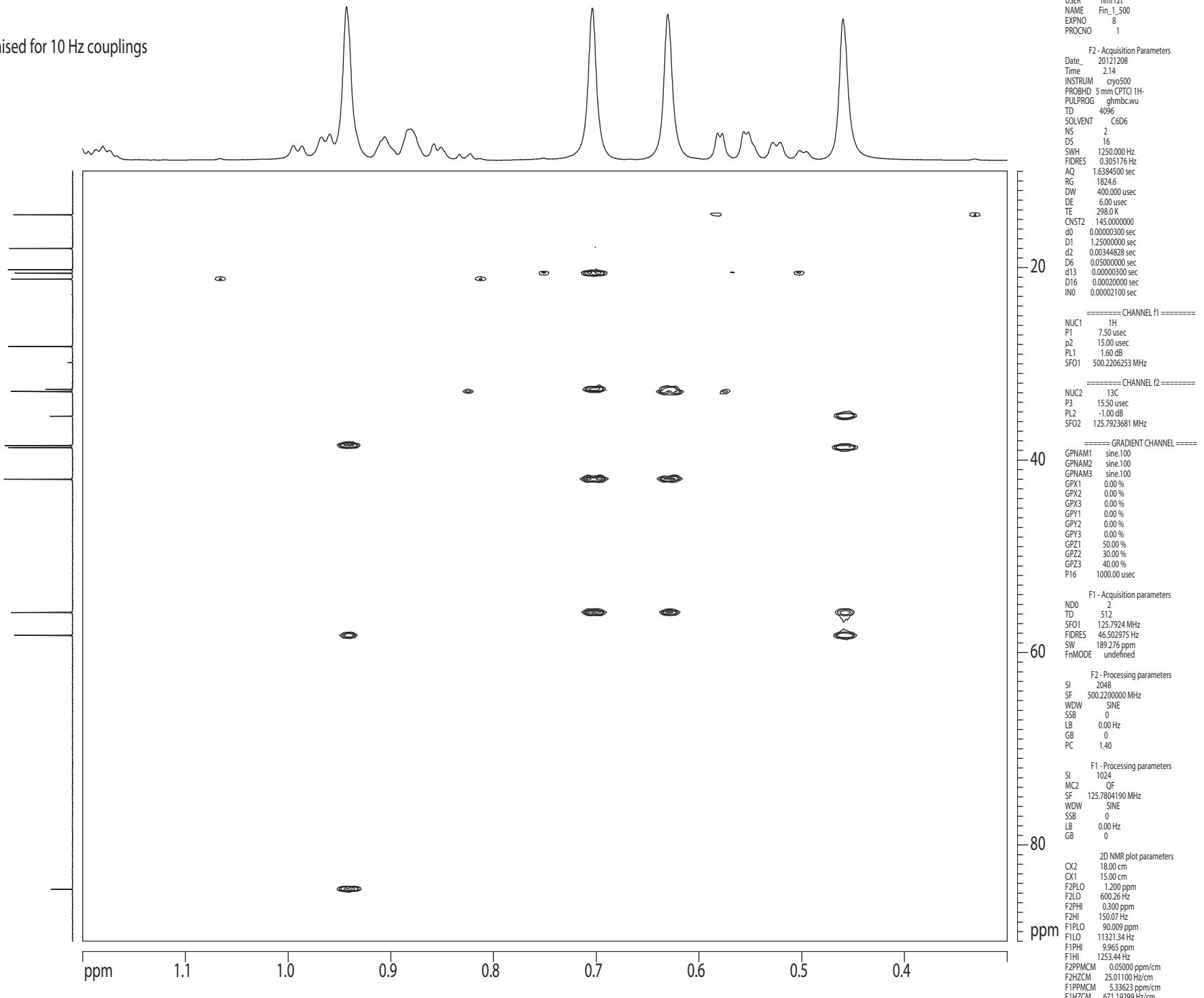
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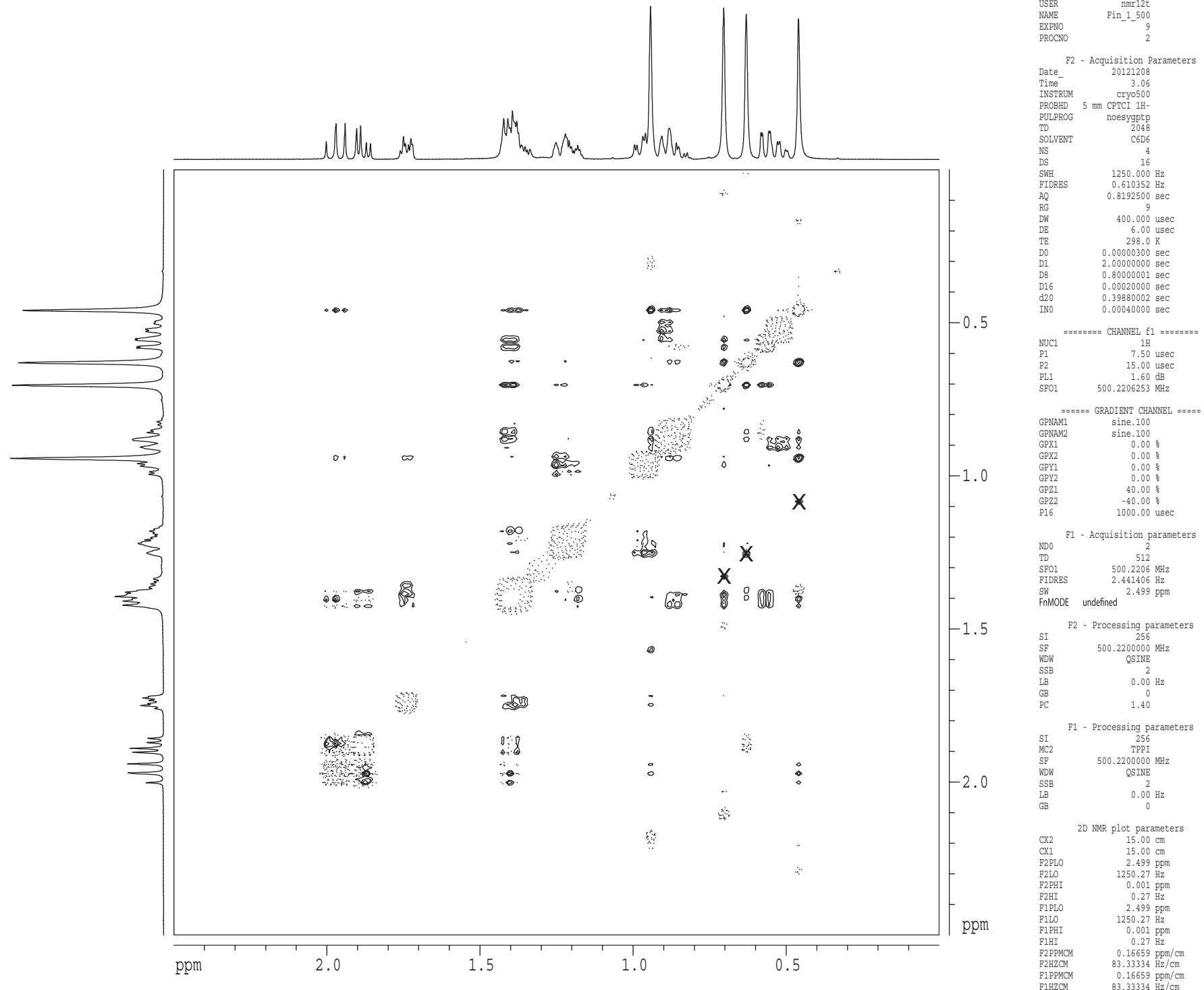
ghmbc optimised for 10 Hz couplings



ghmbc optimised for 10 Hz couplings

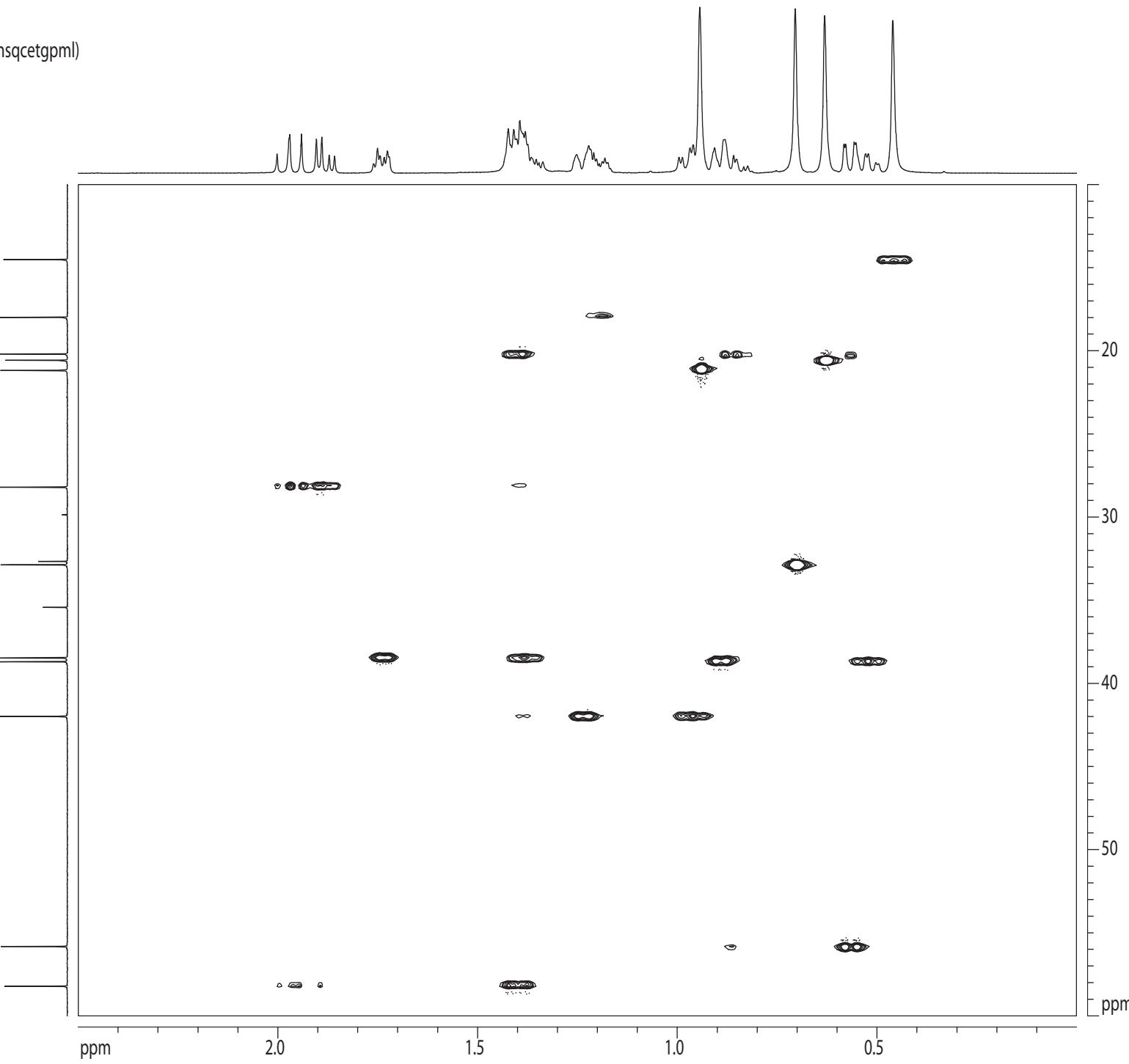


NOESY



5ms

HSQC-TOCSY (hsqcetgpm)



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 GPZ2 20.10 %
 P16 100.00 usec

F1 - Acquisition parameters

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 SF01 500.2200000 MHz
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 SW 189.276 ppm
 FnMODE Echo-Antiecho

F2 - Processing parameters

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 GB 0
 PC 1.40

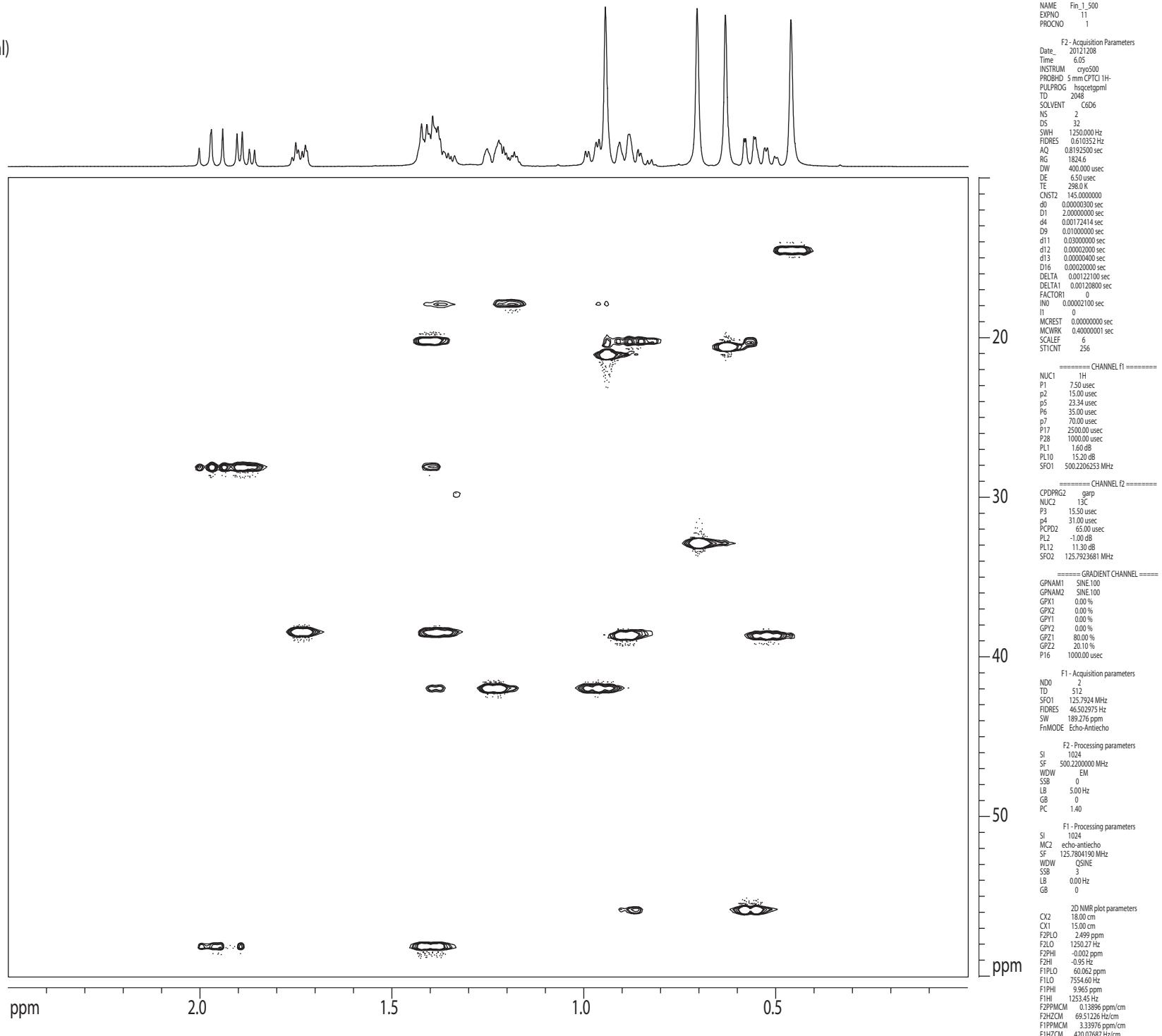
F1 - Processing parameters

SI 1024
 MC2 echo-antiecho
 SF 125.7804190 MHz
 WDW QSINE
 SSB 3
 LB 0.00 Hz
 GB 0

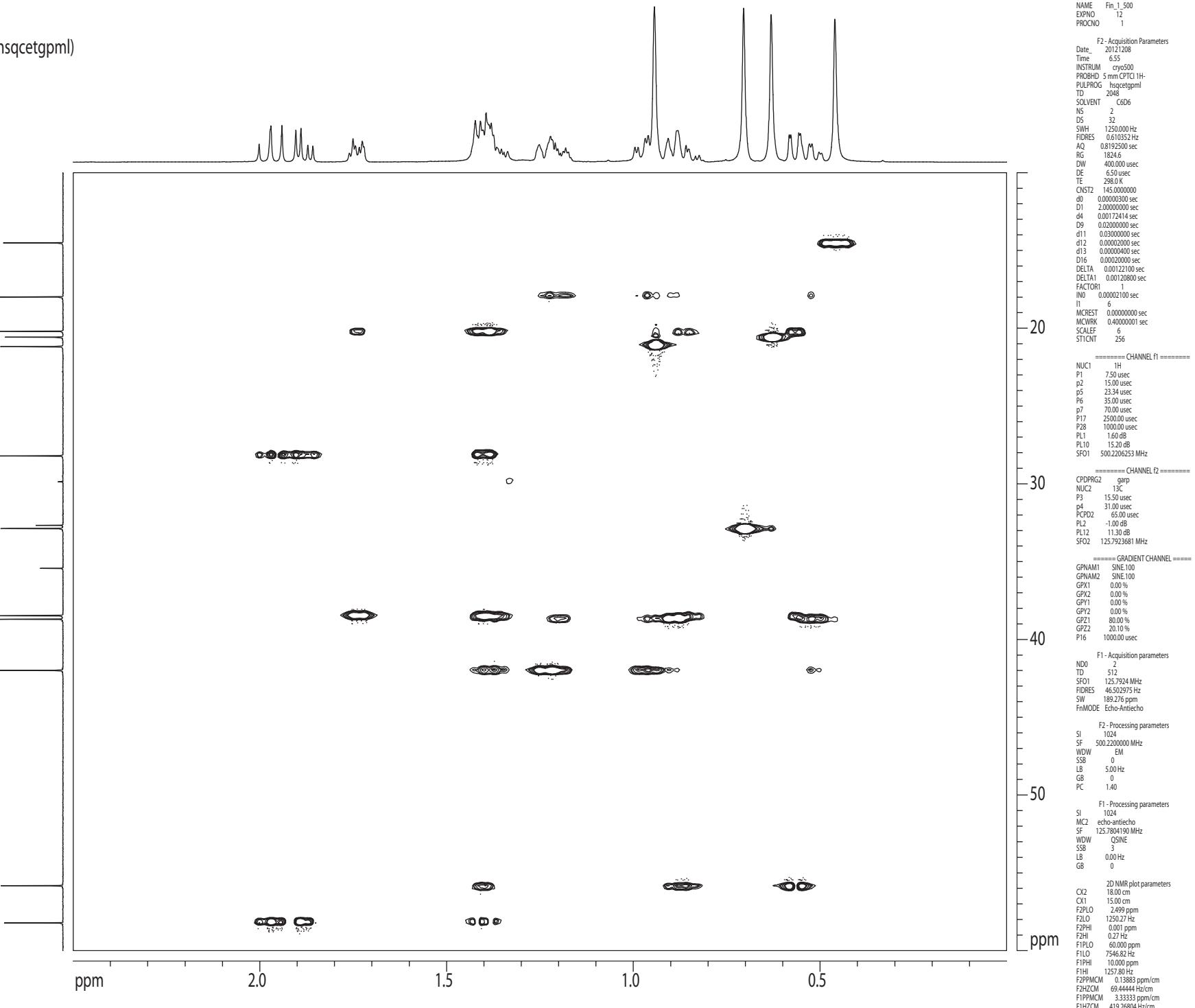
2D NMR plot parameters

CX2 18.00 cm
 CX1 15.00 cm
 F2PLO 2.499 ppm
 F2LO 1250.27 Hz
 F2PHI 0.001 ppm
 F2HI 0.27 Hz
 F1PLO 60.00 ppm
 F1PHI 754.002 Hz
 F1HI 10.00 ppm
 F1HL 1257.880 Hz
 F2PPCM 0.13884 ppm/cm
 F2HZCM 69.4444 Hz/cm
 F1PPCM 3.33333 ppm/cm
 F1HZCM 419.26804 Hz/cm

10ms
HSQC-TOCSY (hsqcetgpm1)



20ms
HSQC-TOCSY (hsqcetgpm1)



100ms
HSQC-TOCSY (hsqcetgpm1)

