Briand



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A monthly collection of informal private letters from Laboratories of NMR. Information contained herein is solely for the use of the reader. Quotation is *not* permitted, except by direct arrangement with the author of the letter, and the material quoted *must* be referred to as a "Private Communication". Reference to the TAMU NMR Newsletter by name in the open literature is strictly forbidden.

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### FORTHCOMING NMR MEETINGS

9th EENC (European Experimental NMR Conference), May 16-20, 1988; Bad Aussee, Austria; For further information, write Professor H. Sterk, Karl-Franzens-Universitaet Graz, Institut fuer Organische Chemie, Heinrichstrasse 28, A-8010 Graz, Austria. See Newsletter 348, 15.

Teaching Course on Nuclear Magnetic Resonance, May 30 - June 3, 1988; Trondheim, Norway; Ms. I. S. Gribbestad, The MR Center, N-7034, Trondheim, Norway.

European Workshop on Nuclear Magnetic Resonance: Seminar on Relaxometry, June 6-7, 1988; Trondheim, Norway; Ms. I. S. Gribbestad, The MR Center, N-7034, Trondheim, Norway.

2nd European Congress on NMR in Medicine and Biology, June 23-25, 1988; Berlin, West Germany; contact Prof. R. Felix, Dept. of Radiology, Charlottenburg University Hospital, Spandauer Damn 130, D-1000 Berlin 19, West Germany.

XIII Intl. Conference on Magnetic Resonance in Biological Systems, Aug. 14-19, 1988; Madison, Wisconsin. See Newsletter 349, 60.

<u>NATO Summer School</u>: "A Methodological Approach to Multinuclear Magnetic Resonance in Liquids and Solids: Chemical Applications", August 22 - September 2, 1988; Maratea, Italy; See Newsletter <u>353</u>, 76.

XXIV Ampere Congress on Magnetic Resonance and Related Phenomena, August 29 - September 3, 1988; Poznan, Poland; Dr. S. Hoffmann, Instytut Fizyki Molekularnej PAN, ul. Smoluchowskiego 17/19, 60-179 Poznan, Poland.

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<u>DEADLINE DATES</u> No. 357 (June) ------ 20 May 1988 No. 358 (July) ------ 17 June 1988 No. 359 (August) ----- 22 July 1988 No. 360 (September) ---- 19 August 1988

All Newsletter Correspondence Should Be Addressed To:

Dr. Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303, U.S.A.

(415) 493-5971

### NMR Studies of Polyamine-DNA Interactions

### Michael J. Minch\*, Yong Ma and James Blankenship Departments of Chemistry and Pharmacology University of the Pacific Stockton, CA 95211

### (received 2/25/88)

Polyamines are aliphatic polycationic compounds found in all cells. A large body of data, collected over the last two decades, indicates that polyamines must play an essential role in cell growth and differentiation. Their ubiquitous distribution, their high concentration in cells, the increase in their concentration in rapidly growing tissue and the coordination of their acetylation and deacetylation with the cell cycle all point to a prominant role for such polycations. Yet their specific physiological function is not known although it is generally presumed that polyamines affect growth by interacting with DNA. There is ample experimental and theoretical evidence for strong binding between polyamine and DNA accompanied by profound DNA conformational changes. Characterization of a polyamine-nucleic acid complex involves more than an estimation of the binding forces, which was the focus of earlier studies, or even the average or lowest energy structure of the complex; the conformational complexities of complex formation and the dynamics of transitions between various conformations must also be considered.

We embarked on an NMR study of polyamines, including spermine, spermidine and N<sup>1</sup>- and N<sup>8</sup>-acetyl spermidine a few years ago and have now completely assigned all <sup>1</sup>H and <sup>13</sup>C resonances of the above compounds by a combination of pH-titration, decoupling and 2D-Hetcor experiments. We began this work with the goal to determine any physical chemical differences between the two isomeric acetyl spermidines that could account for their different physiological functions. The N<sup>8</sup>-acetyl spermidine occurs in cell nuclei associated with DNA whereas the N<sup>1</sup>-acetyl compound is an intermediate in the breakdown of spermidine and is found primarily in the cytoplasm. Even at 500 MHz the <sup>1</sup>H methylene resonances of spermidine and N<sup>1</sup>-acetyl spermidine are poorly resolved with several overlapping at pH 7.5. But the chemical shifts are pH dependent and a complete assignment of all resonance was accomplished by examining spectra at pH values above and below the pH where individual resonances begin to overlap.

The NMR spectrum of spermidine alone has narrow lines characteristic of small molecules and the addition of heterogeneous calf thymus DNA causes only minute differences in the chemical shifts and linewidths of the polyamine.<sup>1</sup> This surprising observation has since been confirmed for spermine-d(CGCGAATTCGCG) complexes by others.<sup>2</sup> Apparently the bulk of polyamine binding is to sites that do not tether the polyamine down so that the polyamine in the compex has a mobility relatively unconstrained by its strong electrostatic interaction with DNA. Does this rule out the use of NMR as a method for studying this important interaction? We would like to share with you two NMR strategies we have adopted for studying this interaction.

The binding of low concentrations of certain paramagnetic ions appears to be base-sequence specific, in that some aromatic and deoxyribose<sup>1</sup>H resonances are broadened much more than others. This affords us an opportunity to study the competitive binding of polyamines to nucleic acids.

The figure illustrates that polyamines replace these paramagnetic ions and that one can monitor the relative binding of the polyamine and paramagnetic ion by studying the linewidth of the broadened DNA resonances as a function of the ratio of these two ions. The bottom spectrum of the aromatic and H<sub>1</sub>-resonances of ds(CGCGAATTCGCG) reveals selective broadening of the C<sub>3</sub>G<sub>4</sub> resonances. In the presence of one equivalent of added polyamine (N<sup>8</sup>-acetyl spermidine) the paramagnetic ion is displaced and the top spectrum looks exactly like that observed in the absence of added ions. This is an important and heretofore unexplored approach since low levels of polyamines by themselves provoke very little changes in the NMR of dilute dsDNA samples and their association with specific bases is hard to assess.

We have also found that polyamines cause a marked broadening of the aromatic and sugar <sup>1</sup>H resonances of partially thermally denatured DNA probably by shifting the ssDNA  $\leftrightarrow$  dsDNA equilibrium. In a related variable temperature study of the Watson-Crick imino protons of ds(CGCGAATTCGCG) in 90% H<sub>2</sub>O at 500 MHz, we found that polyamines increase the proton exchange rate of the outermost GC base pair while paradoxically increasing the oligonucleotide "melting temperature."

1. M. J. Minch, J. Blankenship and H. D. Pham, "Interaction of Acetylated Polyamines with DNA," ACS 20th Western Regional Meeting, Sacramento, Oct 11-12, 1984.

2. D. E. Wemmer, K. S. Srivenugopal, Brian Reid and D. Morris, <u>J. Mol. Biol.</u>, 185, 457-459 (1985).



### 355-4

### UNIVERSITY OF CALIFORNIA, SAN FRANCISCO

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SANTA BARBARA · SANTA CRUZ

February 16, 1988 (received 2/27/88)

Dr. Bernard L. Shapiro TAMU NMR News Letter 966 Elsinore Court Palo Alto, CA 94303 University of California Service Veterans Administration Medical Center 4150 Clement Street (11D) San Francisco, California 94121 (415) 750-2146

Re: Intensity and Phasing Standards for In Vivo MRS

Dear Dr. Shapiro:

The overall goal of research at the Magnetic Resonance Unit, Veterans Administration Medical Center, the University of California at San Francisco, continues to be development and utilization of magnetic resonance spectroscopy techniques for the investigation of intact tissues in animals and human subjects. Current instrumentation includes a Philips 2.0 T whole body magnetic resonance imaging/spectroscopy system which permits MRS from isolated volumes of interest in human tissues including brain, heart, liver, kidney, muscle and various tumors. Until recently, the major localization method which has been used is the ISIS technique (1,2). We have incorporated a post-acquisition saturation sequence into ISIS to enable its application with surface coils (3).

Absolute quantitation of phosphorus metabolite levels is of obvious importance, and our initial approach in human studies has been to make use of an intensity standard consisting of a small glass vial containing hexamethylphosphorous triamide, or HMPT ( $P[N(CH_3)_2]_3$ ), fixed near the center of the loop of the probe. The chemical shift of this compound is well outside the range of tissue metabolites so it can be examined separately from the tissue.

Although the HMPT phantom is external to the subject, and thus the volume being shimmed, the field homogeneity at the site of the phantom is reasonably The phantom serves a number of purposes: first, it is visible on the good. proton image, enabling the coil position to be referenced with respect to anatomical features of the subject. Second, the phantom enables a pulse length determination to be made for the loaded coil (the length for a 180 pulse is determined on the phantom, and computer generated  $B_1$  isocontour plots of the probe are used to re-scale this value for a 90° pulse length at the center of the volume of interest (VOI)). Finally, the phantom is used as an intensity standard. Then, in a separate experiment, using a VOI identical to that used in the human study, the signal intensity of the phantom is related to that observed in a standard phosphate solution of known phosphate In addition, corrections for partial saturation of concentration (4). metabolite resonances (T1) in human studies are required for absolute In experiments with surface coils (severe  $B_1$ quantitation (5). inhomogeneity) the spatial shift of the VOI that accompanies chemically shifted resonances (2,3) results in altered resonance intensities due to the spatial sensitivity of the surface coil. Computer simulations are used to obtain an average sensitivity over the selected VOI, to facilitate quantitation of metabolites (5).

Signal intensities are obtained through integration of deconvoluted peaks as accomplished with the curve fit routine in the NMR-1 software (George Levy, Syracuse University). However, with spectra of poor signal-to-noise, the proper phasing can be ambiguous, and different individuals may choose different phase parameters, resulting in alterations in peak positions and intensities. To combat this phase uncertainty, we have begun using a second phantom, triphenoxyphosphine, which resonates 16 ppm downfield from HMPT. By positioning the carrier between the two phantom resonance positions and phasing both peaks, phase parameters can be obtained and applied to the human spectra that are virtually operator-independent. This procedure does require that the probe Q be sufficiently low so that the spectrometer response is not altered by the change in carrier frequency. The strong inductive coupling to the sample in our human spectroscopy experiments insures that this condition is well met.

We have recently implemented 3-dimensional phase encoding spectroscopic imaging techniques on the Philips system. Acceptable quality <sup>31</sup>P spectra have been obtained from human brain, liver, heart, and kidney. This approach is advantageous compared to "single point" MRS techniques in that metabolic information is obtained simultaneously from multiple voxels over the entire tissue region of interest.

A 7.0 T, 18 cm horizontal bore magnet system fully equipped with selfshielding gradient coils, and spectroscopy and imaging software will be delivered this summer. This system will be used for technique development, and studies of intact animals, perfused organs, and isolated cells.

Finally, we note that three recent additions to our laboratory have interests in proton spectroscopy: Drs. Andrew Maudsley (spectroscopic imaging), Hoby Hetherington (proton editing techniques), and Albert Thomas (multiple quantum studies). We hope to be able to report on their research activities in a future letter.

Sincerely,

MICHAEL W. WEINER

DIETER J. MEYERHOFF

DONALD B. WIEG

MATSO ROTH

HUBESCH

GREGORY S. KARCZMAR

### References:

- 1. Ordidge RJ, Connelly A, and Lohman JAB: J Magn Reson 66:283, 1986.
- 2. Segebarth C, Baleriaux D, Arnold DL, Luyten PR, and den Hollander JA: Radiology 165:215, 1987.
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- 5. Lawry TJ, Karczmar GS, Weiner MW, and Matson GB: (Submitted).

Telephone (0223) 337733

### DEPARTMENT OF PHYSICAL CHEMISTRY UNIVERSITY OF CAMBRIDGE

LENSFIELD ROAD CAMBRIDGE CB2 1EP

Dear Barry

### A BIZARRE LINESHAPE

This comes under the heading of "partial mysteries", beloved of the early contributors to your Newsletter. It could all turn out to be wrong because it has not been substantiated by experiment.

We recently had occasion to try some 3D correlation spectroscopy by an unorthodox method involving line-selective pulses (rather than evolution and Fourier transformation). The 3D frequency space was explored with two variable frequencies  $F_1$  and  $F_2$ , the third dimension (F\_3) being examined in the usual manner by transforming a free induction signal  $S(t_3)$ . The selective pulses were shaped according to the first half of a Gaussian curve (1,2). They were applied (simultaneously) to an I transition and an S transition of a scalar-coupled three-spin (ISR) system. Coherence transfer to R involves the creation of IS multiplequantum coherence - zero-quantum if the I and S transitions are regressively connected, double-quantum if they are progressively connected (3).

We used a density matrix treatment to calculate the intensity of coherence transferred to a given R transition, as a function of  $F_1$  and  $F_2$ . For simplicity we neglected relaxation and spatial inhomogeneity effects, but took into account the time-domain shaping pattern of the selective pulses. For the regressive case the predicted intensity contour map came as no particular surprise; it is basically the familiar "phase-twist" lineshape which appears in several kinds of 2D experiment. In contrast, the predicted shape for the progressive case is quite unlike anything we have seen before (Figure 1).

The diagonals  $\Delta F_1 = \Delta F_2$  and  $\Delta F_1 =$ nodes represent of zero  $-\Delta F_2$ intensity. The contour diagram is antisymmetric with respect to reflection in one or other of these There are four lobes of diagonals. intensity, alternating signal positive-negative, and they are elongated along the direction  $\Delta F_1 =$ (the condition for efficient  $-\Delta F_2$ double-quantum excitation of coherence). have observed the Although we regressive responses experimentally

the progressive responses have so far eluded detection, possibly because of mutual cancellation of the closely-spaced positive and negative regions in a spatially inhomogeneous applied magnetic (neglected in the calculation)

F2

(received 2/24/88)

field

Kindest regards

'Kay ray freehan

Friedrich, Davies and Freeman, J. Magn. Reson. <u>75</u>, 390 (1987).
Davies, Friedrich and Freeman, J. Magn. Reson. <u>75</u>, 540 (1987).

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### Smith Kline & French Laboratories

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**23. February 1988** (received 2/26/88)

Prof. Bernard Shapiro 966 Elsinore Court Palo Alto CA 94303

### NDS Terminals: Benefits for Hare Program Users Part II.

Dear Barry,

In my letter a few weeks back I reported how to enable gray scaling on the Northwest Digital GP-29 and GP-220 terminals, ending with a teaser comment that Dave Wemmer had told me that GP-220 terminals with 8-plane memory extensions could produce overlays of two different spectra (using Dennis Hare's FTNMR program). I have been corrected by Dave-but for the better! If I had read a little more of the NDS manual, I would have realized that both the GP-29 and GP-220 terminals, without the additional memory, can produce these plots. Since it is easy to do I thought it warranted a quickee second note.

On the GP-29: set planes to "combined" as before. The terminal is now set up to produce two graphics planes; normally these are combined to give gray scaling, but one can toggle between them using  $\langle ctr \rangle F1$  (plane 1) and  $\langle ctr \rangle F2$  (plane 2). Generate your first plot, and hit  $\langle ctr \rangle F1$ , then type  $\langle ctr \rangle F2$  and generate the second plot (with the necessary FTNMR commands). Typing  $\langle ctr \rangle$ -SHIFT-F5 OR's the planes for a nice overlay display;  $\langle ctr \rangle$ -SHIFT-F6 AND's the two planes, which yields gray scaling.

I hope NDS and FTNMR users find these suggestions helpful; perhaps some clever users can report their own nifty findings in these pages.

Sincerely Yours,

FTNMR is available from Hare Research Inc., 14810 216th NE, Woodinville WA 98072. NDS GP-29 and GP-220 terminals can be ordered from Northwest Digital Systems, Box 15288, Seattle WA 98115 (215)542-0014.



355-10



UNIVERSITY OF LEICESTER BIOLOGICAL NMR CENTRE

P.O. Box 138 Medical Sciences Building University Road LEICESTER LE1 9HN *Telephone:* Direct Line: (0533) 52 Switchboard: (0533) 522522

Prof. G. C. K. Roberts Dr. L. Y. Lian

> 18th February, 1988 (received 3/2/88)

Professor Bernard L. Shapiro, 966, Elsinore Court, Palo Alto, CA 94303, U.S.A.

Dear Professor Shapiro,

<sup>31</sup>P NMR in Phosphate Buffer?

It is often difficult to perform phosphorus NMR of biological samples in phosphate buffer, because of the strong phosphorus signal of the buffer. It is not easy to irradiate the buffer signal in the homogated mode whilst observing <sup>31</sup>P (as in presaturation for observing <sup>1</sup>H in water) because most spectrometers do not have additional frequency synthesizers or probe configurations other than the standard ones. However, a very simple solution that we have adopted is to use composite pulses to suppress the buffer <sup>31</sup>P signal, in an analagous manner to water suppression techniques.

Figure 1 is the normally-recorded spectrum  $^{31}$ P spectrum at 202 MHz of a complex of a protein (dihydrofolate reductase) and NADPH at about 3mM, in a 50mM inorganic phosphate buffer. The NADPH 2'-phosphate signal is at +2.4 ppm (w.r.t. external trimethylphosphate) and the pyrophosphates are at -14 and -16.2 ppm (a small amount of free NADPH present shows as a shoulder on the -14 ppm signal). Figure 2 is of the same sample using a 1331 pulse. The carrier was set on the buffer signal, a "1" pulse of 2 µsec and a "3" pulse of 6.2 µsec were used with gaps of 0.13 msec between them (the spectral width was 8000 Hz, a 90° pulse was 17 µsec). The 2'-phosphate signal is inverted with respect to the pyrophosphates, and its proximity to the carrier reduces its intensity. Obviously in this particular case buffer suppression was not crucial but it does illustrate how effective this technique can be. There was only a slight roll in the baseline, which could probably be eliminated by using a variation on the "binomial" pulse theme. No adjustment to the spectrometer hardware was required.

Please credit this contribution to Professor G.C.K. Roberts' subscription.

Yours sincerely, J.R.P. Arnold



FIGURE 1









### THE PENNSYLVANIA STATE UNIVERSITY

152 DAVEY LABORATORY UNIVERSITY PARK, PENNSYLVANIA 16802

College of Science Department of Chemistry

> February 23, 1988 (received 2/27/88)

Dr. B. L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

Dear Barry:

### LONG RANGE DIPOLAR COUPLING OF <sup>13</sup>C TO <sup>14</sup>N

We have been using <sup>13</sup>C CPMAS to characterize the solid state of molecules which can undergo the Cope rearrangement. The structure and two spectra for one of these molecules are shown on the next page.

Two hundred fifty microseconds of dead time inserted between the spin lock and the onset of decoupled acquisition causes the disappearance of all hydrogen-bearing carbons except the methyl carbons. The two cyano carbons are each split into asymmetric ugly doublets, with the larger downfield part of the doublet centered at about 125 ppm and the upfield part near 100 ppm. The splittings of the inequivalent cyano carbon resonances are caused by the dipolar coupling of <sup>13</sup>C to quadrupolar <sup>14</sup>N, which magic angle spinning is unable to completely average. The complicated powder lineshapes in each part of the doublet represent a superposition of powder lineshapes of the type described by Hexem et al.<sup>1</sup> The splittings, about 650 Hz, are surprisingly large for the magnetic field strength of 2.3 T, the internuclear distance of about 1.15 Å, the axial symmetry of the electric field gradient tensor, and the expected quadrupolar coupling constant of  $e^2Qq/h = 3$  MHz.<sup>1,2</sup>

What surprised us even more, however, was the <sup>13</sup>C-<sup>14</sup>N dipolar coupling apparent in the resonances of carbons 2 and 6. These are clear examples of the long-range dipolar coupling reported by Harris et al.<sup>3</sup> The magnitudes of the splittings and the lineshapes of these resonances are qualitatively consistent with their <sup>13</sup>C-<sup>14</sup>N internuclear distances.

We hope this interesting tidbit beats your ultimatum letter. [Not quite . . .BLS]

Regards,

aloyd (mm)

Alan J. Benesi and Director, NMR Facility

Lloyd M. Jackman Professor of Chemistry

AJB/jmm

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<sup>13</sup>C CPMAS Spectrum



As above, except 250  $\mu$ sec delay between spin lock and decoupled acquisition



355-14



DEPARTMENT OF HEALTH & HUMAN SERVICES 200 C St., S.W. (HFF-423)

**Public Health Service** 

Food and Drug Administration Washington DC 20204 March 11, 1988 (received 3/17/88)

Dr. Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Ct. Palo·Alto, CA 94303

Proton Homonuclear NOE Experiments: A Practical Test Compound

Dear Barry,

Despite the sophistication of today's NMR spectrometers, it is sometimes desirable to have a compound on which one can practice NOE determinations or check instruments. This is especially the case in structural elucidation work where trans-H/H orientations may have to be inferred from the absence of NOEs. We have used several compounds as reference materials and have been particularly pleased with the following: 1,5-dichloro-2,4-dimethoxybenzene (DCDMB,1). Its proton NMR spectrum consists of single lines for the observed (H-3) and irradiated (OMe) nuclei, and there is a control nucleus (H-6) which is also a singlet. The resulting NOEs are substantial, with degassed solutions giving rise to ca. 35% enhancements in CDCl<sub>3</sub> or CS<sub>2</sub> at room temperature. In addition, these values appear to be unchanged at elevated temperatures and over a freguency range of 80-400 MHz.

While DCDMB is not comercially available, it can be prepared by methylating the corresponding dichlororesorcinol with dimethyl sulfate. However, a word of <u>caution</u> is important: dimethyl sulfate is extremely toxic and a cancer-suspect agent. Please see supplier data sheets for proper handling, i.e. wearing rubber gloves and working in a hood, and disposal procedures.



Samuel W. Page

Sincerely,

Eugene P. Mazzola

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### 3737 BELLAIRE BOULEVARD HOUSTON, TEXAS

MAILING ADDRESS P. O. BOX 481 HOUSTON, TEXAS 77001

February 24, 1988 (received 2/26/88)

Dr. Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

Dear Dr. Shapiro:

INSTALLATION OF SHIELDED GRADIENT COILS ON SHELL'S GE CSI-2T NMR IMAGING SPECTROMETER

We have recently installed a prototype shielded gradient set from GE NMR Instruments on our CSI-2T imaging system. The gradients, which were designed by P. Roemer and W. Edelstein at GE Corporate R&D, have an inner bore of 6" and can accommodate an RF coil up to 4.5" in diameter. The windings produce a maximum gradient of 18 G/cm with settling time on the order of 300 usec.

The fast switching times of the shielded gradients has allowed us to decrease the minimum echo time for 2D Fourier images from 13 ms to 5 ms. Typically, our minimum echo time is limited by the finite acquisition time of the echo itself. Less than 5 ms echo times are possible at the expense of digital resolution, i.e. echo times of 2 to 3 ms have been obtained at relatively larger field of views and/or smaller data acquisition buffer sizes. Shorter echoes could be obtained with faster data acquisition, for example using a Nicolet digital oscilloscope.

Due to the relatively short T2 values of fluids in rocks, a significant improvement in image quality is obtained at the shorter echo times as shown in the figure.

Please credit this contribution to Dr. L. L. Sterna's account.

Very truly yours, Sutin

P. N. Tutunjian Research Chemist Analytical Chemistry Department

1.002921

H. J. Vinegar! Senior Staff Physicist Reservoir Mechanisms Research Dept.

SHELL DEVELOPMENT COMPANY Bellaire Research Center



2D Fourier image of a 3mm slice of Indiana limestone obtained at (a) 13 ms and (b) 5 ms echo times. The 5 ms, T2-weighted image which shows a marked improvement in image signal-to-noise clearly displays the natural bedding planes of the rock.

### ARCONNE NATIONAL LABORATORY

9700 South Cass Avenue, Argonne, Illinois 60439

Telephone: (312) 972-3524

March 8, 1988 (received 3/11/88)

Dr. B. L. Shapiro 966 Elsinor Court Palo Alto, CA 94303

RE: 27 Al NMR of "Aluminum-Free" Ceramic Rotors

Dear Barry:

We have been using <sup>27</sup>Al NMR to investigate the composition of clay minerals found in Argonne Premium Coals. In addition to the expected tetrahedral and octahedral <sup>27</sup>Al resonances of the clays, we observed a fairly intense signal from our Si<sub>3</sub>Ni<sub>4</sub> rotor (Fig. 1). The chemical shift of 108 ppm is similar to the value reported in the literature for AlN (1). We can only assume that Al<sub>2</sub>O<sub>3</sub> was used as a sintering aide during fabrication of the ceramic, even though the material was billed as being Al free. The upfield shift of 2 ppm observed from pure AlN (despite operating at the same resonance frequency) may be explained by substituting Al tetrahedra in the Si<sub>3</sub>N<sub>4</sub> lattice. <sup>29</sup>Si NMR experiments performed to resolve this issue turned out to be ambiguous (Fig. 2). While the sideband intensities were unexpectedly large for  $v_r = 10$  kHz, the <sup>29</sup>Si linewidth was significantly sharper than that expected for a Si<sub>3</sub>N<sub>4</sub> lattice substituted with Al (2). We estimate T<sub>1</sub> of the <sup>27</sup>Al species to be >50 s; consequently, the <sup>27</sup>Al resonance is not readily observed in spectra of pure clay samples obtained using fast pulse repetition times. However, <sup>27</sup>Al spectra of samples with little Al content, as in the case of coals, or spectra taken using longer recycle times may be dominated by this unwanted resonance.

N.D. Butler, R. Dupree and M.H. Lewis, J. Mater. Sci. Lett. <u>3</u> (1984) 467.
R. Dupree, M.H. Lewis, G. Leng-Ward and D.S. Williams, J. Mater. Sci. Lett. <u>4</u> (1985) 393.



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Figure 1. 78.2 MHz  $^{27}$ Al NMR spectrum of raw coal in a "Si<sub>2</sub>N<sub>4</sub>" rotor spinning at 10 KHz.

Sincerely,

Arthur R. Thompson Chemistry Division





Figure 2. 59.6 MHz <sup>29</sup>Si NMR spectrum of a "Si $_3$ N<sub>4</sub>" rotor at 10 KHz MAS.

Robert E. Botto

Ghemistry Division

U.S. DEPARTMENT OF ENERGY

The University of Chicago



Department of Pure and Applied Chemistry

Thomas Graham Building, 295 Cathedral Street Glasgow G1 1XL Tel: 041-552 4400

> 16th February, 1988 (received 3/2/88)

Dr. Bernard L. Shapiro, Editor/Publisher TAMUNMR Newletter, 966 Elsinore Court, Palo Alto, California 94303. U.S.A.

Dear Barry,

Proton and C-13 Spectra of Bicyclo-[3.3.0]-6-oxa-oct-2-ene-1-one (1).

In connection with work in this department on the use of acetylenic cobalt carbonyl complexes to form 5-membered-ring ketones (Khand reaction), it is frequently necessary to analyse in some detail the NMR spectra of intermediates. The spectra are usually nicely resolved, but sometimes there can be difficulty in assignments. The proton spectrum of the title compound forms a case in point; two of the resonances overlap and the problem was solved only by using the classical technique of selective decoupling in both the proton and carbon spectra. With regard to the carbon spectra, I feel that valuable information is gained from a proton-coupled spectrum; in this case it proved crucial in confirming the assignments.



Using the atom labels in the structure (1) the assignments are: Proton Chemical Shifts 2.895; D 3.56; E 3.67; B 7.59; С A 6.21; Н 3.88 ррп. 3.66; 4.14; G F Proton Coupling Constants CE 7.65; AB 5.65; AD -1.65; BD CD 6.00; 2.55; DH 0.55; GH -8.95 Hz CF 0.65; DG 6.95 EF -9.30; Carbon Chemical Shifts (multiplicity in proton-coupled spectrum) C1 210.65 (m?); C2 135.05 (2x3); C3 164.65 (2x6); C4 49.75 (3x2); C5 47.20 (3x2); C6 71.05 (2x4); C7 69.75 ppm. (2x2x3). Carbon-Proton Coupling Constants (absolute values) 1-Bond couplings: 2A 172.0; 3B 165.0; 4C 138.0; 5D 140.0; 148.5 Hz. (7G+7H)/2(6E+6F)/2 149.5; 2-Bond couplings: 7D 9.0 Hz 6C 9.5 2- and 3- Bond couplings (averaged): 3.5; (3A+3D+3C?+3G?+3H?)/5 3.0; (2B+2D?)/2(4E+4E+4F(/3 3.0; (5B+5C+5G+5H)/4 5.0Hz. All experiments were made using a Bruker WM 250 machine. Proton spectra were run at 250.13 MHz, using a spectrum width of 3000 Hz, and with 32K data points.

Carbon spectra were run at 62.9 MHz, using a spectrum width of 15000 Hz, and with 32 K data points.

Regards and Best Wishes,

Yours sincerely,

Dr. Peter Bladon

### FORTHCOMING NMR MEETINGS, continued

Teaching Course on Nuclear Magnetic Resonance, September 5 - 9, 1988; Trondheim, Norway; Ms. I. S. Gribbestad, The MR Center, N-7034, Trondheim, Norway.

European Workshop on Nuclear Magnetic Resonance: Seminar on Contrast in MRI and MRS, September 12 - 13, 1988; Trondheim, Norway; Ms. I. S. Gribbestad, The MR Center, N-7034, Trondheim, Norway.

27th Annual Eastern Analytical Symposium and Exposition, October 2 - 7, 1988; New York Hilton Hotel, New York; General Chairman - Dr. Harvey S. Gold, Polymer Products Dept., E256/308, E. I. du Pont de Nemours & Co., Wilmington, DE 19898; (302) 695-3669.

International Post-Graduate Course 'NMR in Agriculture, Plants and Products', October 3 - 15, 1988; Wageningen, The Netherlands; Dr. Ir. J. H. de Ru, Foundation for Post-Graduate Courses, Agricultural University, Hollanseweg 1, NL-6706 KN Wageningen, The Netherlands.

### CALIFORNIA INSTITUTE OF TECHNOLOGY Chemical Engineering 206-41 Pasadena, CA 91125

March 9, 1988 (received 3/11/88)

Dr. Bernard L. Shapiro Editor/Publisher TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

### Protein Hydration by <sup>17</sup>O NMR

Dear Barry:

In order to obtain a better understanding of the role of water in affecting protein stability in nonaqueous solvents, we have studied the hydration of a hydrophobic protein, crambin, as a function of the quantity of water added to the solvent (N,N dimethylformamide). The fraction of water that is bound to the protein surface is determined by first integrating the NMR water signal at ambient temperatures (where water exists in both "bound" and "free" states), and then comparing it with the integrated intensity when the temperature is lowered to well below the freezing point of the solvent (1).

The results of our NMR studies on <sup>17</sup>O labelled  $H_2O$  (12 atom %) are shown by the solid curve in Fig. 1 for four concentrations corresponding to 1, 2, 4 and 6 percent v/v water content. The dashed curve represents the water that would be associated with the crambin if there were no selective interactions between the protein and the solvent components. It is clear that crambin exhibits a clear tendency to "extract" water to its surface in a solution that consists primarily of the nonaqueous solvent.

Further studies on protein hydration in other solvents are presently underway; it is expected that the nature of the protein hydration will provide a model that will complement current ideas concerning the role of water in enzymatic activity (2,3).

Sincerely, Akbar Mayreen Frances Arnold

### References

- 1. Kuntz, I. D., Brassfields, T. S., Law, G. D. and Purcell, G. V., Science 163, 1329 (1969).
- 2. Rupley, J. A., Yang, P. H. and Tollin, G., in "Water in Biopolymers", ed. S. P. Rowland (Am. Chem. Soc., Washington, D.C., 1980).
- 3. Finney, J. L. in "Water in Aqueous Solutions", ed. G. W. Neilson and J. E. Enderby (Bristol, 1986).

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# How to bring true computer power to your NMR research





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Amsterdam, 2nd March 1988

KONINKLIJKE/SHELL-LABORATORIUM, AMSTERDAM BADHUISWEG 3 1031 CM AMSTERDAM-N. 020-

> Professor Bernard L. Shapiro - <del>Texas A&M University</del> -

-Department-of-Ghomistry-

- College Station - TEXAS-77843-

Verenigde Staten

### 0352

Dear Prof. Shapiro,

To my opinion NMR is one of the most fascinating fields to work in. Nevertheless it is sometimes good to change and therefore I very recently switched to a job in a completely different field. Consequently, this contribution to TAMU Newsletters will my last one. Fortunately, the NMR research at our laboratory will continue and Dr. Alex de Groot, which is also one of the authors of the present contribution, will be my successor.

I kindly request to continue our subscription to TAMU Newsletters under his name.

Finally we would like to acknowledge the contribution of Jos van Boxtel to the work presented below. Jos spent 6 months on our laboratory as a trainee of the Landbouw Universiteit Wageningen and did most of the data acquisition and processing.

### A correction procedure for non-uniformities in NMR images

It is a well-known fact that NMR images may show relatively large intensity variations over the imaging plane due to non-uniform RF excitation. This non-uniformity is fully determined by the probe design and, therefore, a phenomenon we sometimes have to live with. As a consequence, quantitative comparison of the intensities of different features in an image is hindered.

In order to circumvent this problem we developed the following procedure to <u>correct for this non-uniformity</u>.

First, the non-uniformity itself is recorded by measuring a so-called <u>reference</u> image of a tube filled with water (note the variation in the intensity profile due to non-uniform excitation in Figure 1). In the second step the image manipulation software is used to "invert" the shape of the non-uniformity in order to obtain the so-called <u>correction</u> <u>image (Figure 2)</u>. The procedure is illustrated in Figure 3, note that since no division of images is required, division by zero is avoided. The removal of the non-uniformity effect is demonstrated in Figure 4 where the original reference image of Figure 1 is multiplied with the correction image of Figure 3 resulting in the desired flat intensity profile.

It will be clear that once the correction image has been obtained this image can be used to correct for the non-uniformity effects in any other image by simply multiplying the image to be corrected with this correction image.

Yours sincerely, Alen de Groot

N.C.M. Alma-Zeestraten and A. de Groot



Figure 1. Image of a 30 mm diameter (maximum) tube filled with water doped with 20 mM NiCl<sub>2</sub>. Clearly the intensity is non-uniform (high intensity is white, low is black). A typical intensity profile is shown (from left to right along the line across the image).



Figure 2. The correction image derived from the image in Fig. 1



Figure 3. Schematic outline of the procedure used to derive the correction image from the reference image applied to an arbitrary intensity profile. At each step the minimum and maximum (relative) values of the intensity profile are indicated.

- a. A normalized arbitrary intensity profile of the reference image (R in text)
- b. Image a. multiplied by -1
- c. First order correction profile
- d. First order corrected profile (note that the non-uniformity is already partially removed)
- e. First step for second-order correction. as b.
- f. Second-order correction profile
- g. Combination of first- and second-order corrections yielding the final correction profile (C in text)
- h. The corrected reference profile R' (see text) obtained by multiplication of a. by g. The non-uniformity is removed.



Figure 4. The corrected image.

355–28	University of Illinois at Urbana-Champaign	<b>School of Chemical Sciences</b> 505 South Mathews Avenue
		Urbana, IL 61801
		February 12, 1988 (received 3/4/88 [sic])
	Dr. Bernard Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303	
	Dear Barry:	

Solid-State 170 NMR of a Picket-Fence Porphyrin and  $YBa_2Cu_3O_{7-x}$ 

We have recently been using  $^{17}$ O NMR to investigate oxygen-transport proteins and model systems, both in solution and in the crystalline solid-state. The results we have obtained on  $^{17}$ O<sub>2</sub>-picket fence porphyrin are shown below:



and suggest that both static and dynamic structural information can be obtained via such studies. For example, we find the isotropic chemical shifts of the bridging and terminal oxygens are at 1200 and 2000 ppm (in moderate accord with the solution results on a single-face hindered iron porphyrin-dioxygen complex of 1755 and 2488 ppm, JACS 1987, 109, 6944), that the shielding tensors are  $\sigma_{11} = 100$ ,  $\sigma_{22} = 1200$  and  $\sigma_{33} = 2400$  ppm for the bridging oxygen, and  $\sigma_{11} = 850$ ,  $\sigma_{22} = 850$  and  $\sigma_{33} = 4200$  ppm for the terminal oxygen, at 77 k, and that the Fe-O-O bond angle must be ~130°. Attempts of observing solid hemoglobin are underway.

We have also been investigating with Tom Rauchfuss in this Department, the <sup>17</sup>O NMR spectra of <sup>17</sup>O-labelled  $YBa_2Cu_3O_{7-x}$ , a high T<sub>c</sub> superconductor. As shown in the following Figure, there appear to be two main types of (narrow) oxygen signal in the room temperature (metallic) state:

Oldfield *et allios* [sic] have herewith been declared the winners and permanent titleholders in the undeclared *Most Authors* competition, which is now closed. Forever. Please.



with isotropic shifts of  $\approx 1800$  and  $\approx 200$  ppm (from H<sub>2</sub>O). In the superconducting state (at 77°K) there still appear to be two main sites. While these results are very preliminary, they do seem promising. T<sub>1</sub> measurements suggest a sizable contribution of Korringa relaxation due to the apparently large Knight shift, in the metallic state.

Yours sincerely,

Hecher Lee K. D. Park E.

C. Coretsopoulos T. Rauchfuss B. Montes

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### Department of Chemistry

LOUISIANA STATE UNIVERSITY AND AGRICULTURAL AND MECHANICAL COLLEGE BATON ROUGE · LOUISIANA · 70803-1804

504/388-3361

March 2, 1988 (received 3/5/88)

Dr. Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

### A LabView-Controlled ADLF Spectrometer

Dear Dr. Shapiro:

We use field cycling methods, developed by Hahn and Redfield in the 1960's and later expanded by Edmonds and Brown, to acquire deuterium quadrupole coupling constants in solid samples. In the present instrument version, the sample is physically transported from high to zero magnetic fields. While the sample is in zero magnetic field, deuterium transitions are saturated with an rf field; the extent of saturation is measured through a double resonance technique that terminates with an Ostroff Waugh experiment on the proton spin system. Then, the zero-field rf transmitter is retuned and the cycle repeated.

Because of all of the functions that must be controlled, computer programs written in traditional languages have tended to be long and complex (ca. 5000 lines of Pascal), such that only the instrument guru can make any changes. However, we recently switched to LabView, a data flow programming language that uses a block diagram approach with a graphical user interface rather than the typical sequential line coding. The results have been most satisfactory. It took THREE WEEKS for two students to reprogram the instrument in LabView. Table 1 lists the main steps in the experiment. Figure 1 shows the instrument layout; figure 2 shows part of the capacitor tuning subroutine.

Table 1, Major Experiment Steps in ADLF (Mechanical Sample Motion)

- based on the 90° and 70° pulse lengths and probe ringdown times, create and load pulse program; signal averagers set to digitize 4 points at each of 128 spin echoes
- get starting parameters from front panel (starting frequency, stopping frequency, frequency increment, H<sub>1</sub> level, low magnetic field setting (ranges from 0 to 7 Gauss))
- initialize IEEE-488 bus and CAMAC crate controller, set zero or low magnetic field
- LOOP: frequency synthesizer, capacitors (series and parallel set according to formula and look-up table); these use BCD interfaces as does the zero magnetic field coil power supply.
- put sample in zero field coil (even at low frequency, it changes coil tuning)
- increment DAC driving variable attenuator until H<sub>1</sub> correct; H<sub>1</sub> monitored with Fluke DMM that senses current in zero field coil.
- set timing generator, signal averagers, Look-At-Me status register in CAMAC
- monitor CAMAC crate until signal averages' Look-At-Me bit set
- DMA signal averager data, calculate integral of spin echoes.
- increment frequency; if not at final frequency, then LOOP (not a GOTO but a WHILE control "statement"). Finally, I note that FORTRAN, Pascal, and Basic are older, or nearly so, than some of my graduate students. This

is a weird situation for research that is, I hope, state-of-the-art.

Sincerely.

Les Butler Assistant Professor of Inorganic Chemistry

<sup>1</sup>National Instruments, 12109 Technology Blvd., Austin, TX 78727-9989 (800) 531-4742



Figure 1. Layout of adiabatic demagnetization in the laboratory frame (ADLF) spectrometer; sample is mechanically shuttled between high and low magnetic fields.



Figure 2. A portion of the  $H_1$  tune program. In this "filmstrip", a WHILE loop will operate until the DAC driving the attenuator is set to give the desired  $H_1$  level. The unattached input "Freq, kHz", was used in an early "filmstrip" to set the frequency synthesizer and series/parallel capacitors. The returned variable, "new", is the new DAC setting in Volts and is displayed in the front panel window in a stripchart format. This format is nice for detecting oscillations that would indicate that the gain in the "circuit", now set for 20, is too large.

355-32

(303)491-6480 Colorado State University Fort Collins, Colorado 80523

Department of Chemistry Dr. Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Ct. Palo Alto, CA 94303 March 7, 1988 (received 3/10/88)

### Preliminary Solid-Sample NMR Spectra on the AM-600.

Dear Barry:

Installation of the Bruker AM-600 spectrometer in the Colorado State University Regional NMR Center has proceeded extremely smoothly. Within a few days of completion of the magnet shimming by the Bruker engineers, our laboratory (primarily Bruce Hawkins, Steve Dec and Bob Zeigler) had obtained some very promising results on solid samples. The spectra below show preliminary Al MAS and H spectra obtained with home-built probes.



We are at present engaged in a vigorous and systematic program to explore what a 14 T spectrometer can do in solid-state NMR. Special emphasis is being directed to comparisons with spectra obtained on 500 MHz and 360 MHz spectrometers.

Sincerely. Garry aciel Professo/

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Deuterium oxide, 99.8 atom % D	30,875-7	\$5.00	26,979-4	.5 pkg \$8.70; 1 pkg \$13.10
<b>Deuterium oxide</b> , 99.8 atom % D (contains 0.75% 3-(trimethylsilyl)propionic- 2,2,3,3-d, acid, sodium salt)	30,876-5	\$6.30	29,838-7	.5 pkg \$10.90; 1 pkg \$15.20
<b>Dichloromethane-</b> <i>d</i> <sub>2</sub> , 99.6 + atom % D	30,877-3	\$28.00	23,694-2	.5 pkg \$50.70; 1 pkg \$84.50
<i>N</i> , <i>N</i> - <b>Dimethylformamide</b> - $d_7$ , 99 atom % D	30,878-1	\$56.60	26,980-8	.5 pkg \$107.00; 1 pkg \$179.00 5 pkg \$595.00
<b>1,4-Dioxane-</b> <i>d</i> <sub>8</sub> , 98.5 atom % D	30,880-3	\$37.50	26,981-6	.5 pkg \$70.90; 1 pkg \$117.90
Methyl-d <sub>3</sub> alcohol-d, 99.5 atom % D	30,881-1	\$18.90	26,982-4	.5 pkg \$32.70; 1 pkg \$54.60
(Methyl sulfoxide)- $d_6$ , 99.9 atom % D	30,883-8	\$6.25	23,692-6	.5 pkg \$10.40; 1 pkg \$16.60
Nitrobenzene-d <sub>5</sub> , 99 atom % D	30,884-6	\$12.00	26,975-1	.5 pkg \$34.00; 1 pkg \$40.00
Nitromethane-d <sub>3</sub> , 99 atom % D	30,885-4	\$13.20	26,983-2	.5 pkg \$22.00; 1 pkg \$36.70
<b>Pyridine-</b> d <sub>s</sub> , 99 atom % D	30,886-2	\$16.70	23,695-0	.5 pkg \$29.00; 1 pkg \$48.50
Tetrahydrofuran-d <sub>8</sub> , 99.5 atom % D	30,887-0	\$63.40	26,984-0	.5 pkg \$110.20; 1 pkg \$183.70
			-	5 pkg \$609.00
<b>Toluene-</b> $d_8$ , 99 + atom % D	30,888-9	\$16.25	26,985-9	.5 pkg \$29.40; 1 pkg \$49.00
Trifluoroacetic acid-d, 99 atom % D	30,889-7	\$7.25	26,977-8	.5 pkg \$12.60; 1 pkg \$19.90

The following 100.0 atom % D\* solvents are available in 0.5-ml ampules:

	1  pkg = 10  x  0.5 ml			
Description	Catalog No.	Price		
Acetone-d <sub>6</sub>	23,696-9	.5 pkg \$26.50; 1 pkg \$44.50		
Acetonitrile-d <sub>3</sub>	23,701-9	.5 pkg \$35.70; 1 pkg \$57.50		
Benzene-d6	23,697-7	.5 pkg \$29.30; 1 pkg \$48.90		
Chloroform-d	23,691-8	.5 pkg \$11.00; 1 pkg \$18.00		
Deuterium oxide	26,978-6	.5 pkg \$10.90; 1 pkg \$16.30		
Dichloromethane-d <sub>2</sub>	23,702-7	.5 pkg \$68.20; 1 pkg \$113.40		
(Methyl sulfoxide)-d <sub>6</sub>	23,693-4	.5 pkg \$32.60; 1 pkg \$54.30		
Toluene-d <sub>8</sub>	23,703-5	.5 pkg \$57.50; 1 pkg \$95.90		
*Minimum isotopic purity 09.96	tom % D			

\*Minimum isotopic purity 99.96 atom % D



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SANTA BARBARA • SANTA CRUZ

DEPARTMENT OF CHEMISTRY

DAVIS, CALIFORNIA 95616

Manipulating NOEs in Paramagnetic Proteins

March 3, 1988 (received 3/8/88)

Professor B. L. Shapiro Editor/Publisher TAMU NMR Newsletter 966 Elsinore Court Palo Alto, California 94303

Dear Barry:

We have been interested in exploring the scope and limitations of the utility of NOEs and 2D NOESY in paramagnetic molecules for both resonance assignment and structure determination. Our emphasis has been to implement such studies on ferric enzymes such as heme peroxidases.

The major problem associated with detecting such NOEs is the strong relaxation caused by the unpaired spins of the metal which markedly decrease the magnitude of steady-state NOEs and makes it difficult to implement time-dependent NOEs.

One approach with which we have had some success is to take advantage of the fact that the correlation time for paramagnetic relaxation is usually some Orbach process which is independent of the overall motion of the system, and hence paramagnetic and diamagnetic contributions to relaxation are completely uncoupled.

This dictates that  $\rho$ , the intrinsic relaxation rate in  $\eta = \sigma/\rho$ , is independent of  $\tau_c$ , the molecular reorientation time, while  $\sigma$ , the cross relaxation rate, is linearly dependent on  $\tau_c$ . Thus doubling the solvent viscosity (with ethylene-glycol as solvent or addition of sucrose) should lead to a doubling of an NOE to a resonance experiencing predominantly paramagnetic relaxation. As shown in the accompanying figure for the NOEs between a pair of geminal 6-propionate  $\alpha$ -CH<sub>2</sub> in high-spin ferric myoglobin, such an increase is indeed observed. In fact, the method seems to work surprisingly well.

A manuscript on the full details on the quantitative discription of such methods, as well as their limitations, is in preparation.

Best regards,

Sert

Gerd N. La Mar Professor of Chemistry

Research Associate

GNLM:des


#### FIGURE CAPTION

Downfield region of the 360 MHz <sup>1</sup>H NMR spectrum of met-aquo myoglobin at pH 6.0,  $20^{\circ}$ C. A) Reference spectrum in D<sub>2</sub>O. B) Difference spectrum showing irradiation of 6  $\alpha$  proton and NOE to 6  $\alpha$ ' in D<sub>2</sub>O. C) Same as B but in 30% (w/w) ethylene glycol-d<sub>6</sub>-D<sub>2</sub>O solvent mixture. \* indicates the off resonance irradiation.



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0304-616672 February 29, 1988 (received 3/10/88)

Professor B. Shapiro, TAMU NMR Newsletter, 966 Elsinore Court, Palo Alto, California 94303, U.S.A.

#### **Trace Analysis of Process Streams by Carbon-13 NMR**

Dear Professor Shapiro,

We needed to measure low levels of two organic molecules in an aqueous wash stream. They were:  $(CH_3)_3 S^+O \Gamma$  {Trimethyl Sulphoxonium Iodide (TMSOI)} and  $CH_3 SO_3^- Na^+$  {Sodium Methane Sulphonate (MESNA)}.

The approximate concentrations had to be mesured quickly in order to determine whether a routine quality control assay needed to be developed. Carbon-13 NMR was selected as the analytical technique for the following reasons:

- Other techniques such as ion chromatography need longer development times.
- (2) Little pre-treatment of aqueous samples is needed.
- (3) Each compound gives a single peak in the broad-band decoupled carbon spectrum.
- (4) Acquisition of spectra could be done overnight with our GE QE300 spectrometer, equipped with an autochanger.
- (5) Calibration solutions are readily available.
- (6) The analysis of results is quickly performed.

The carbon-13 spectra of calibration solutions of TMSOI in water  $/D_2O$  (90:10 v/v) are shown in Figures 1A to 1D. The carbon-13 spectrum of the test sample with the same concentration of  $D_2O$  added, is shown in Figure 1E. All the spectra were processed under identical conditions with the same scaling factor.

The level of TMSOI in the test sample had to be determined by extrapolating the calibration curve, because TMSOI started to crystallise from solution at a concentration of 50 mg/ml. The concentration of TMSOI in the test sample was determined as 66 mg/ml and the increased solubility is probably due to the presence of other ions in the test sample which increase the ionic strength and hence the solubility by " the salting-in" effect. The detection limit for TMSOI is less than 5 mg/ml.

The level of MESNA in the test sample was very low and no signal could be observed in the aqueous wash sample. The detection limit of MESNA was determined by a spiking experiment, which also showed that the MESNA and TMSOI peaks did not overlap at the pH used for the experiment. Figure 2A shows the carbon-13 spectrum of the test sample diluted 1:1 with water/D<sub>2</sub>O. Figure 2B shows the test sample diluted 1:1 with a 5 mg/ml MESNA solution. The concentration of water/D<sub>2</sub>O was 90:10 v/v in both cases. A peak at 38.5 ppm due to MESNA is separated from the TMSOI peak by 1.1 ppm in Figure 2B. The detection limit for MESNA is less than 5mg/ml.

Please credit this contribution to David V. Bowen's account.

Yours sincerely,

Pfizer Central Research

Mike Kinns Mike Newros M. Kinns M.J.Newman.

Anaytical Chemistry Department



# DOW CHEMICAL U.S.A.

Dow

March 7, 1988 (received 3/11/88)

MIDLAND, MICHIGAN 48667

Dr. Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

MAXIMUM PULSE WIDTH CRITERIA AND ENHANCEMENT RATE

Dear Barry,

It's common to try to get a semi-quantitative high resolution NMR spectrum from weak signals and as-yet-unknown components, using ordinary multiple scan single pulse experiments. I approach this from a signal/noise (S/N) "enhancement rate" viewpoint, with the goal of getting a signal fast <u>and</u> being confident that detected signals do not differ in degree of saturation by more than some chosen factor, such as two-fold.

One concern is reduced signal size at large frequency offset  $\Delta\nu$  Hz from the pulse radiofrequency. A basic relation we are taught for 90 degree pulses in a quantitative experiment (repeat time > 3 x T<sub>1</sub>) is PW(90) x  $\Delta\nu \leq 0.25$  to assure 99% of the zero offset signal size, where PW(90) is the 90 degree pulsewidth time in seconds. In the day-to-day world where T<sub>1</sub> has not been determined and we already accept partial saturation in order to raise S/N rapidly, we use pulses less than 90 degrees and would be satisfied to know that the offset is not causing more than 5 or 10 percent signal loss relative to the center position.

I have modelled the offset effect for the rapid-pulse experiment as a function of on-resonance flip angle from 0 to 90 degrees, and find as safe overall guide:

> 95% of maximum for (PW x  $\Delta \nu$ ) < 0.3 > 90% of maximum for (PW x  $\Delta \nu$ ) < 0.4.

The table below of reduced FID ENHANCEMENT RATES

= (steady-state magnetization) x sin(flip angle) x  $(T_1/T_r)^{1/2}$ 

illustrates this. The rate ratios and effective flip angle also are shown. The repeat time  $T_{r}$  is chosen to produce optimum (maximum) signals using flip angle  $\omega(0)$  for on-resonance nuclei with longitudinal relaxation time  $T_{1}$ , according to the Ernst relation,

 $\cos(w(0)) = \exp(-T_{repeat}/T_1),$ 

except for 90 degree flip angle where  $T_r/T_1 = 1.25$  (the optimum value when  $T_2 << T_1$ ). The corresponding off-resonance flip angles  $\omega(\Delta\nu)$  were derived from published relations (e.g. Martin, Martin, and Delpuech) and used to calculate the respective steady-state signals for the same values of  $T_r/T_1$ . The effective flip angles determined for PW x  $\Delta\nu = 1$  are less accurate because computer round-off error may be present.



REDUCED FID ENHANCEMENT RATE, S/N PER  $(T_r/T_1)^{1/2}$ RAPID-SCAN CONDITIONS  $T_r/T_1$  is optimum for w = w(0) at offset,  $\Delta \nu = 0$ 

<u>ω(0</u> )	$P W x \Delta \nu =$	<u>o</u>	<u>.1</u>	.25	<u>.4</u>	.5	<u>1</u>
10°	Rate	.707	.707	.703	.680	.640	.000
Ι	$(\Delta \nu)/I(0)$	1.00	1.00	.99	.96	.91	.00
ω	$(\Delta \nu)$	10	9.9	9.0	7.6	6.4	0
45°	Rate	.704	.704	.698	.670	.622	.010
I	$(\Delta \nu) / I(0)$	1.00	1.00	.99	.95	.88	.01
ພ	$(\Delta \nu)$	45	44	40	34	28	0
70°	Rate	.676	.676	.668	.629	.572	.022
Ι	$(\Delta \nu)/I(0)$	1.00	.96	.95	.89	.81	.03
ω	$(\Delta \nu)$	70	69	62	51	42	1
90°*	* Rate	.638	.644	.663	.655	.611	.042
I	$(\Delta \nu) / I(0)$	1.00	1.01	1.04	1.03	.96	.07
w	$(\Delta \nu)$	90	88	79	64	52	3

\*\*  $T_r/T_1 = 1.25$  to maximize the net FT response for  $T_2^* << T_1$ .

The point is that <u>pulse width</u>, <u>not flip angle</u> per se, is the critical factor in pulse offset considerations. This point has been made many times in development of multi-pulse compensation schemes for spin inversion, etc., but is not necessarily appreciated by users at a lower level of sophistication.

Implications? Pulsewidth is a problem for rapid scanning of larger full-range spectra of nuclei such as F-19 and P-31. Pulsewidth is not a problem for C-13 on modern spectrometers. For example, at 100 MHz and maximum shift range  $\pm$  150 ppm,  $\Delta \nu = \pm$  15,000 Hz, a pulse width of 27 microseconds satisfies the relation, PW x  $\Delta \nu = 0.4$ ; this is more than PW(90) for many spectrometers.

The run time to achieve a given FID S/N ratio is inversely proportional to the squared enhancement rate and actual time is proportional to  $T_1$ . Nuclei with different  $T_1$  values would behave according to their own  $T_1/T_1$  ratios and net flip angles. Remember that the minimum possible value for  $T_1$  is the data acquisition time, and that the S/N obtained after Fourier transform is maximum and proportional to  $T_2^*$  (=  $1/(\pi \times \text{observed linewidth})$ ) only when data acquisition time is greater than 1.5 x  $T_2^*$  (preferably 3 x  $T_2^*$ ).

len

Jerry P. Heeschen Analytical Sciences Instrumental Methods 1897 Building (517) 636-5330 335-39

355-40



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Tacoma, Washington 98477 (206) 924-2345

March 10, 1988 (received 3/17/88) Dr. Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto CA 94303

SUBJECT: One User's Cautions on PSZ Rotors for CP/MAS NMR

Dear Dr. Shapiro:

Several months ago I switched from boron nitride to phase-stabilized zirconia as CP/MAS sample rotor body material. There are few more aggravating events than sudden decomposition of a rotor at high speed. This always seems to occur during a crucial experiment and requires a couple hours downtime to clean the probe and start over.

Over the years I have destroyed about a dozen BN rotors and except for the time loss and the need to repeat some experiments, no real damage has occurred. This spinner system is of the double-air-bearing design, with polyimide bearing surfaces at the top and bottom of the ceramic body which float in brass stator journals. When BN rotors come apart under stress, they normally break into very small pieces (powder); even though these microchunks fly everywhere inside the spinner/coil compartment, they cause no physical damage to the static components - rather like throwing handfuls of powdered sugar against the kitchen wall.

A different situation pertains to PSZ rotor bodies, however. A few months ago, I had one of these self-destruct at 3 krps and I was startled to find that PSZ breaks into larger pieces than BN and that, owing to PSZ's greater hardness and density, these pieces hit things with sufficient momentum to damage them. Besides putting dents and gouges in the rf coil, some of these particles became wedged between the rotor bearings and stator journals, digging divots in the brass journal surfaces and chewing up the bearings.

The outcome of this disaster was replacement of the entire stator assembly. Naturally, the manufacturer does not stock supplies of this item, so the spectrometer collected dust while waiting six weeks for custom engineering and manufacturing to be completed. Eventually, I succeeded in getting new journals fitted in the original stator, leaving me with a handy back-up for the next rotor explosion.

All in all, this was a costly and time-consuming chain of events. I recommend that those who use PSZ rotors keep spare stator assemblies on hand; sooner or later, this will happen to them.

Respectfully,

Lany Amos

Larry W. Amos Analytical Laboratories Weyerhaeuser Analytical and Testing Services



The PTS 300 joins our growing line of direct synthesizers that cover 0.1 to 500 MHz. A new development that answers the demand for compact devices (31/2")rack space), it still benefits from a decade of experience and manufacturing expertise that have made PTS synthesizers the number one choice in OEM applications. With thousands in use they serve many diverse applications, including communications, ECM, mode locking, ATE and MR Imaging.

The PTS 300 is a generator of precision frequencies. It transfers the accuracy and stability of a frequency standard (built-in or external) to any output frequency between 0.1 and 300 MHz. Resolution is 1 Hz throughout the range, remote-only versions are available. With its low spurious outputs, fast switching, low phase noise, choice of signal purity and digital phase modulation, it sets new standards in performance/ price for low noise synthesized sources.

Both versions of the PTS 300 switch phase-continuously for all steps 1 Hz through 100 KHz.

The calculated MTBF of PTS synthesizers is 7,000 hours. Evaluation of service records for many thousands of units show a figure of 25,000 hours. A yearly failure rate of 3-4% is typical. Warranty is two years from date of shipment. For ease of service the PTS 300 uses plug-in modular design throughout.

FREQUENCY	Range:	0.1 MHz to 300 MHz (299.999 999 MHz)						
	Resolution:	1 Hz						
	Control:	Local by 10-position switches. Remote by TTL-BCD, 1248, buffered par. entry						
	Switching Time:	me: 20 micro-sec (10 MHz, 100 MHz steps)						
		5 micro-sec (1 MHz steps)						
		for all steps 1 Hz –100 KHz); (pha	se continuous)					
OUTPUT	Level:	+ 3 to + 13dBm, (1V) into 50 ohms, me	etered in dBm and volt					
	Flatness:	± 0.5dB						
	Impedance:	i0 ohms						
	Control:	Manual by F/P control, remote by voltage $(+0.63 \text{ to } +2.00\text{V})$						
		Туре 1	Type 2					
SPURIOUS	Discrete	– 70/65dB (typ./spec)	– 60/55dB					
OUTPUT	Harmonics	– 30dB						
	Phase Noise:	– 68dBc (0.5Hz-15KHz)	– 63dBc (0.5Hz–15KHz)					
	Noise Floor:	– 135dB/Hz						
	Phase Setting:	0°, 90°, 180°, 270°	0-360° in .225° steps					
	(digital)	standard; 5° resolution	optional					
		optional 0-360°						
EBEQUENCY	Internal <sup>.</sup>	$3 \times 10^{-9}$ /day or	$1 \times 10^{-8}/dav$					
STANDARD	(OPTIONAL)	$+1 \times 10^{-8}/0-50^{\circ}C$	$\pm 1 \times 10^{-6}/0-50^{\circ}C$					
Int. Std. or ext. Drive	•	$1 \times 10^{-6}$ /year	$2 \times 10^{-6}$ /year					
required for operation		(OVEN)	(TCXO)					
GENERAL	External Drive:	10 MHz, 0.4Vrms into 300 ohms; 5 MH	z, 0.5Vrms into 300 ohms					
	Aux. Output:	10 000 MHz, 0.4V into 50 ohms						
	Oper. Ambient:	0 to 55°C, 95% R.H.						
	Power:	105–125V, 50–400 Hz, 40 Watts (100V, 220V, 240V available)						
	Dimensions/Weight:	t: $19 \times 3\frac{1}{2} \times 17^{\prime\prime}$ (relay rack or bench cabinet) 25 lbs.						
PRICES (domestic)	Price: (Manual & Remote Controls, TCXO)	\$5,550.00	\$5,050.00					

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0430

Professor B.L. Shapiro TAMU NMR Newsletter 966 Elsinore Court PALO ALTO California 94303 USA

Zürich, March 10, 1988 (received 3/14/88)

#### COMPUTER FITTING, A LAST RESORT REMEDY IN 2D SPECTROSCOPY

Dear Barry,

It is wellknown that spectra which exuberate with information are notoriously difficult to analyze. Two-dimensional correlation spectra of medium-size molecules are typical in this respect. As soon as strong coupling is present, a logical straight forward analysis by a skilled spectroscopist or by a trained computer is difficult or impossible and accurate chemical shifts and coupling values cannot be determined.

In this situation, least squares fitting is usually a last resort remedy to reach the goal. We have explored the feasibility of iterative fitting of cross-peak multiplets in 2D correlation spectra of the E.COSY-type in order to determine coupling constants and chemical shifts. The approach is similar to fitting 1D spectra. A reasonable set of initial values is chosen, and selected regions of a 2D spectrum are simulated and compared with the experimental spectrum. The parameters are then iteratively refined following one of the wellknown strategies of optimization. The error surface can also in 355-44

this situation be very treacherous with countless crevasses and local minima. Temporary smoothing of the error surface by peak broadening is mormally indispensible.

A novice in computer fitting of 2D spectra will rapidly become disappointed by the fact that his recently acquired pocket calculator is hardly of any use and that even most powerful computers are kept busy for appreciable time before a satisfactory fit is reached. As a demsonstration example, the strongly coupled proline-2 seven-spin system in a 300 MHz E.COSY spectrum of the cyclic decapeptide antamanide has been analyzed. Two chemical shifts and six coupling constants were determined in a first step from the  $C_{\delta}H_1 - C_{\delta}H_2$  cross peak. remaining five chemical shifts and 15 coupling constants, together with two linewidth parameters and the overall intensity were obtained from a 23 parameter fit in 14 iteration steps of the cross peaks between  $C_{\delta}H_1$  and  $C_{\beta}H_1$ ,  $C_{\beta}H_2$ ,  $C_{\gamma}H_1$  and  $C_{\gamma}H_2$ . The Modula-2 program was running for 55h 43min (!) on a DEC µVAXII-GPX computer with an ULTRIX-32w Operating The fit shown for the overlapping cross peaks system.  $C_{\delta}H_1 - C_{\beta}H_1$  and  $C_{\delta}H_1 - C_{\gamma}H_1$  is certainly as good as it can be expected and the coupling constants obtained are accurate to about 0.1 Hz. The full story is presently in print in the Journal of Magnetic Resonance.

Sincerely yours,

Lollan K. Urddi

Zoltán L. Mádi

Michaul

Richard R. Ernst



Cross-peak region from a 300 MHz E.COSY spectrum of the cyclic decapeptide antamanide containing two cross peaks of proline-2. A: experimental spectrum, B: computer fit.

355-46

# UNIVERSITY of PENNSYLVANIA

School of Medicine Department of Biochemistry and Biophysics Philadelphia, PA 19104-6089

February 26, 1988

Dr. Bernard L. Shapiro Editor - TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

Dear Dr. Shapiro:

We would like to report on a resonator based on the "cosine coil" design built for NMR studies on human heads and limbs. Basically, the coil consists of linear wire segments distributed around a cylinder in a way that there is a "cosine" distribution of current leading to a very large volume of B1 homogeneity. (A detailed paper describing this coil design has just been accepted for publication: L. Bolinger et al., J. Magn. Res.)

Our head coil has been built around a 32cm polypropylene cylinder as the former (a standard distilled water carboy with its neck and bottom sawn off). The coil segments are made from 1 mil thick adhesive backed copper strips (0.7cm wide & 32cm long). On either end of the coil we have used 4cm wide copper strips as one plate of the tuning capacitor. We have fabricated two end rings also made from polypropylene to slide over the formeer. Two more 4cm wide copper strips have been glued to the end rings to form a high voltage variable capacitor. Tuning is achieved simply by sliding the end rings over the former. The self resonance frequency of this coil is over 100 MHz and with our capacitor we can tune it from 50 MHz to 90 MHz. This tuned circuit is coupled to the T/R switch inductively.

The unloaded Q of this coil is over 250 while loaded Q (with a human head inside) drops to about 100. The 90 degree pulse width on a human head at 78 MHz proton frequency is about 250 microsec with a 400W transmitter power. The B1 inhomogeneity over a human head seems to be less than 10%. A simplified diagram of this coil is given in the figure.

The major features of this coil design are:

- 1. It is extremely easy and relatively inexpensive to build.
- 2. It has a very large volume of B1 homogeneity.
- 3. Contrary to other designs such as a birdcage etc., the whole coil can be treated electrically as a simple inductor and hence one can readily build multiple tuned circuits with this coil.

Thanking you,

Sincerely yours,

Hari Subramanian

Lizann Bolinger

n S. Leigh



#### POSTDOCTORAL POSITIONS AVAILABLE

Opportunities for Postdoctoral research in biomedical NMR. Instrumentation available includes 500 MHz, 360 MHz and 200 MHz Bruker spectrometers, and 40 cm bore, 2.0 T and 5.0 T in vivo spectrometers. Applicant should be skilled in modern NMR techniques and instrumentation, or highly enthusiastic about learning these techniques. A keen interest in either metabolic studies in vivo (spectroscopy and imaging of brain, heart, liver, muscle), or high resolution spectroscopy of extracts is a plus, since research opportunities exist in these areas.

If interested please send Curriculum Vitae, biblography and 3 letters of reference to: Dr. John S. Leigh, Professor, Biochemistry and Biophysics, University of Pennsylvania, D501 Richards Building, Philadelphia, PA 19104-6089.

# A Poor Man's "Ethernet" Link to the DEC VAX from Nicolet 1280 Computers in the NT Series NMR Spectrometers

C. Allen Bush

Department of Chemistry, Illinois Institute of Technology, Chicago, IL 60616 U.S.A. Bitnet address: "GLYCOBUSH @ IITVAX"

The inadequacies of the computers attached to the older NMR instruments such as the Nicolet 1280 supplied with the NT series spectrometers become readily apparent when one attempts some of the currently available 2-dimensional experiments such as phased COSY (DQF COSY). Although the electronics can generate pulse sequences for acquisition of suitable data sets, the computer lacks adequate storage for processing of the large matrices needed to avoid cancellation of antiphase cross peaks multiplets. Similarly the software provided with these older spectrometers lacks many desirable features such as the ability to plot negative and positive contour levels or to integrate cross peak intensities in phased NOESY spectra. An unreasonable effort would be required to modify the assembler coded software to incorporate these features and it would not be possible to explore methods of linear algebra or MEM for extraction of the spectral information from the FID. It has been suggested that these new methods could be very useful in improving the resolution in the  $f_1$  dimension of 2-d NMR by avoiding the severe apodization which accompanies zero-filling necessary in the  $t_1$  dimension.

The obvious solution to the shortcomings of an obsolete computer is to transfer the 2-d NMR data from the instrument computer to a modern general purpose computer for data processing thus freeing the spectrometer to do what it does best, namely to acquire NMR data. As a byproduct of this measure, one gains a simple and efficient method for back-up of data on the tape drive of the general purpose computer as well as faster processing of the 2-d data without the purchase of an array processor for the proprietary instrument computer. Among the several available software packages for processing NMR data on general purpose computers, we have selected the FTNMR package of Dennis Hare. It is especially well suited for analysis of the 2-d NMR data used in our research on the structure and conformation of carbohydrates which exploits technology similar to that used in the biophysical chemistry of proteins and nucleic acids. A major obstacle to the use of a general purpose computer for 2-d NMR data processing is the need for transmitting large data sets. Typically, the raw data in a 256 x 2k x 2 complex data set with phase encoding may involve as much as 6 megabytes of data. Although it is expensive, the

generally lack this capability and their serial data transfer rates are slow. For older NMR computers with limited disk storage, high speed data transfer on a parallel data port is desirable.

The Nicolet 1280 computer supplied with the NT series spectrometers has a parallel 8-bit port intended for use with a Centronics printer which is capable of high speed data transfers. Support is supplied with the original software for data output on this port. In our laboratory we have connected the output from this port to a serializer intended for remote use of a Centronics printer. Ordinary twisted pair wires lead to the VAX computer site where the data are converted back to parallel form by a second converter. The two converters necessary were purchased from Black Box Corp. of Pittsburgh, Pa. for a total cost of about \$600. The parallel output of the Black Box was connected to the VAX Unibus through a DR-11-W interface. The equivalent Q-bus interface is the DRV-11-B which would be suitable for a MicroVax installation. The VAX running VMS provides software support via QIO calls for DMA transfer of data in large blocks to reduce software overhead to negligible levels. Thus the data transfer rate is limited only by the speed of the Black Box serializer which we have chosen for this implementation. If the VAX were in the same room as the NMR spectrometer, one might be able to eliminate the Black Box and do the transfers directly over parallel ribbon cable. The data blocks are transferred from the Nicolet memory directly in native format and written to the VAX disk by a VAX FORTRAN program. A second formatting program is required to convert the native Nicolet character codes and floating point numbers in the data header into standard formats and to convert the NMR data from Nicolet format (20 or 40 bit integers) into 32 bit floating point suitable for processing by the Dennis Hare software.

In our hands the scheme described above is capable of transferring a typical 2-d data set from the CDC Hawk disks on the Nicolet to the VAX in about 10 to 15 minutes. Although a serializer superior to the Black Box would result in improved transfer times, the data transfer is much shorter than the time for acquisition of a typical 2-d data set (4 to 24 hrs.). It will be recognized that since the Centronics port on the 1280 is for output only, handshaking is provided in this scheme only at the byte level. We had anticipated that it might be necessary to use the serial port on the 1280 to receive parity check information on entire data blocks to ensure integrity of the data and to request retransmission of any faulty data blocks. We have not implemented the block parity checking since data errors have not been a problem in more than a year of routine use of this system in our laboratory.

I would like to acknowledge Tom Lew of Toronto for originally suggesting the use of the Centronics port for data output and for substantial help with the software. HUNTINGTON MEDICAL RESEARCH INSTITUTES

Professor B.L. Shapiro, TAMU NMR Newsletter, 966 Elsinore Court, Palo Alto, CA 94303

March 11th 1988 (received 3/17/88)

File transfers from Nicolet 1280 to IBM PC

Dear Professor Shapiro,

Readers of the TAMU NMR newsletter who have Nicolet 1280 computers may be interested in a program we have developed to transfer files to an IBM PC (or equivalent) microcomputer. There were several reasons for the development of the program, the most important one being the need for off-line data analysis. We also wished to be able to transfer data to a more "friendly" programming environment, one which is readily available to a large number of our users. Indeed, with the advent of portable computers (and personal computers at home), our spectroscopists may now also examine their spectra at home, if they so desire!

We have primarily been using this program to download image files (".DAT") recorded on General Electric "CSI" spectrometer; however, it should be equally applicable to other types of data files (and possibly other Nicolet computers if they use the "SECS" file transfer protocol). Use is made of the Nicolet utility program "FILTRN" to transmit data via serial RS-232 port at rates of up to 9600 baud. 19.2 kbaud should be possible over very short distances, although we haven't tried this. The receiving program on the PC converts the Nicolet 20-bit word into a 3 byte/sample format before storing it on disk; if the dynamic range of the data permits (virtually always the case for image files from our system which has a 12-bit ADC) a separate routine can be used to pack this into a 2 byte integer format. This program also discards the file header plus some other garbage. Finally, a routine is available to graphically inspect the downloaded file in either format.

Anybody who is interested in obtaining a copy of these programs (and the source code if desired, written in "Turbo Pascal") should send a blank 5 1/4" floppy disk to me at the address below,

yours sincerely,

Barker Peter

Peter B. Barker

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200/400	4.7T	400 mm	324 mm	1.8 G/cm	140 mm DSV ± 4 ppm	80 mm DSV 0.1 ppm	8.50 m	6.75 m
85/310	2.0T	310 mm	225 mm	3.0 G/cm	100 mm DSV ±5 ppm	70 mm DSV 0.1 ppm	4.50 m	3.63 m

DSV=Oiameter Spherical Volume HHLW = Half-Height Line Width PPM = Parts Per Million



Spectroscopy Imaging Systems 1120 Auburn Road Fremont, California 94538 (415) 659-2600 200/400, 200/330, and 300/180 magnets for the NMR Imaging Spectrometer System. (Photos courtesy Oxford Instruments.)

Note: Equipment described is intended for investigational purposes, and is not approved by the FDA for clinical use.



March 10, 1988. (received 3/16/88)

Dr. B.L. Shapiro 966 Elsinore Court Palo Alto CA 94303 U.S.A.

Title: "Solid Phase <sup>13</sup>C NMR of Dibenzo-12-Crown-4"

Recently we have become interested in the applications of  $^{13}$ C CPMAS to stereochemical problems in crown ether chemistry. As an example of spectral obtained I enclose that for DB-12-C-4. In solution at room temperature one observes the expected 4 resonances. The solid phase spectrum is much more interesting, with a doubling of resonances due to a stereochemically rigid macrocyclic conformation in the crystal. We have found that the crystal possesses a true molecular centre of symmetry, so that indeed 8 resonances are expected.

The observed chemical shift difference of 7.3 ppm at the <u>ortho</u>-aromatic carbons is worthy of note. The x-ray structure shows a torsional angle of  $24.64^{\circ}$  for the C3-C2-O-C7 network, while the value for the C6-C1-O-C8 network is  $61.64^{\circ}$ . We attribute the more shielded resonance to C3, where the steric  $\gamma$ -effect is presumably more pronounced than for C6.

the steric  $\gamma$ -effect is presumably more pronounced than for C6. At present we are in the process of calculating (INDO MO) the angular dependence of  $^{13}$ C shifts for O-C-C-O and C-O-C-C torsional networks. Such data have only been calculated previously for the C-C-C-C case in butane.



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DEPARTMENT OF CHEMISTRY AND BIOCHEMISTRY 405 HILGARD AVENUE LOS ANGELES, CALIFORNIA 90024-1569 March 11, 1988 (received 3/14/88)

Dr. Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

Dear Barry,

USES OF NMR SAMPLE CHANGERS IN AN ACADEMIC ENVIRONMENT

Since postdoctoral and graduate student labor is cheap, the purchase of equipment for automatic sample changing for NMR spectrometers in academic labs is often regarded as frivolous by granting agencies, administrators, and even end users who are accustomed to running all spectra manually. From a purely economic point of view, it should be pointed out that NMR spectrometer capital costs and operating costs are anything but cheap and thus any method of increasing the productivity of the spectrometer in an environment where the demand for spectrometer time is heavy should be considered. Sample changers don't eat, sleep, or even answer the telephone. From a more academic point of view, high throughput with minimal intervention by busy human beings will allow some experiments to be carried possible due to lack of time.

Our departmental NMR facility has made use of automatic sample changers in a variety of ways for almost three years now. The following spectrometers in our laboratory are fitted with sample changers: 1) 500 MHz (Bruker AM500) with 7 different probes and a large variety of accessories to maximize flexibility, 2) 360 MHz (Bruker AM360) with 2 probes and a smaller variety of accessories, and 3) 200 MHz (IBM Instruments/Bruker AF200) with 2 probes and no other accessories. Additionally, a Bruker data station running the same software is linked via optical fiber to these spectrometers. The availability of some type of off-line processing is essential to the operation of this facility using sample changers.

There are three different modes in which the sample changers are used. The first is operation by the NMR laboratory staff to run service samples submitted by the users. The proton probe is used for maximum sensitivity two mornings per week and the 5mm broadband probe is used for one overnight run where both <sup>1</sup>H and <sup>13</sup>C spectra are obtained, sometimes with extended signal averaging. This mode is used most heavily for the 500 MHz spectrometer where the user must have adequate training to ensure that the hardware is set up properly for observation of the chosen nucleus. Advantages of this mode are 1) users with no training at all can obtain high field spectra every few days and 2) the operator runs the spectrometer in exactly the same mode on a regular basis including standard samples. This latter advantage provides a form of quality control that may be overlooked in academic laboratories where most samples are run by the users.

The second mode is operation with the sample changer by a particular user

in a time slot scheduled in advance. This mode allows for a great deal of flexibility in obtaining routine spectra provided the user is trained in setting up the hardware properly to observe the desired nucleus. In a regularly scheduled time slot every week, one student sets up and runs up to forty <sup>11</sup>B samples in about three hours including <sup>1</sup>H decoupled and coupled spectra on each sample. Another group runs several <sup>15</sup>N samples including extended signal averaging in one overnight run per week. There have been other examples of use of the sample changer for a large number of spectra on other "X-nuclei" that would not have been done manually for lack of operator time. NMR staff personnel rapidly set up the sample changer for a novice user who then ran the samples. A few groups have such a heavy need for <sup>1</sup>H spectra that they designate an operator within their group to gather up the samples and run them using the sample changer in a regularly scheduled time.

The third mode is first-come first-served on the 200 MHz spectrometer using a 5mm probe that is switched under computer control between  ${}^{1}$ H and  ${}^{13}$ C observation. The hardware is kept constant on this spectrometer to assure easy operation and fast turn-around times for novice users. This mode is in operation a few hours per day Monday through Friday. Other times the spectrometer is available for experienced users, but the hardware must be returned to the state required for novice users.

The 360 MHz spectrometer is just now being installed. Its sample changer has a bar code reader for labels made for each sample. A 5mm probe that can be switched between  ${}^{1}$ H,  ${}^{13}$ C, and  ${}^{31}$ P under computer control will probably be used the most with the sample changer.

The actual use of the sample changers in our laboratory has evolved in the past as the users needs and equipment availablity have changed. It is clear that the availablity of the automatic sample changers has been extremely important to those experiments that are amenable to automation and has aided those that are not by minimizing the amount of spectometer time devoted to routine work.

Sincerely,

Jane Frouse

Mane Strouse Director, Instrumentation Facility

GROUPE DE DYNAMIQUE DES PHASES CONDENSEES

Laboratoire Associé au C.N.R.S. Nº 233

Montpellier, le March 4, 1988 (received 3/15/88)

"NMR <sup>13</sup>C signal enhancement by cross polarization in graphitic powders dispersed in a polymer matrix"

P.BERNIER and R.DELTOUR

Dear Pr. Shapiro,

The <sup>13</sup>C resonance observation in graphitic powder at room and lower temperature is hindered by the weak natural abundance of these nuclei (1.1%), the wide line powder spectrum ( 4-5 gauss) and the rather long spin lattice

relaxation time  $(T_1(T)$  between 2 and 90 minutes). As a consequence various methods have been used to investigate these nuclei, including electronsnuclei spin polarization transfer by Overhauser effect<sup>(1)</sup>, fast passage techniques<sup>(2)</sup>, high resolution pulse methods<sup>(3)</sup> and the use of lamellae oriented graphite samples<sup>(4)</sup>.

The techniques used for the powder samples are very much time consuming ranging from hours up to weeks of data accumulation<sup>(3)</sup>.

In order to reduce the observation time, we have dispersed high conductivity carbon black powder particles (Ketjenblack EC) in a polyethylene polymer matrix.

The carbon particles, with a high surface-volume ratio (1200  $m^2/gr$ ) have an onion skin structure consisting of fine pellicular graphite lamellae forming connected hollow partly spherical bundles<sup>(5)</sup>.

By cross polarization of the  $^{13}$ C nuclei with the protons of the polyethylene at the surface of the carbon black particles and using magic angle spinning, we have obtained an appreciable increase in sensitivity. This is illustrated in fig. 1, showing the  $^{13}$ C carbon black signal at room temperature after 14 hours data signal accumulation (- 10.000 signal sweeps) using a Bruker CXP Fourier Transform pulse spectrometer at 50.3 MHz. The line shift of 160 ppm (from TMS) for the graphitic  $^{13}$ C is somewhat smaller than the value (180 ppm) reported for bulk graphite by Suganuma et al.<sup>(6)</sup> and in the measurements on oriented graphite lamellae of Kume<sup>(4)</sup>;

A tentative explanation which will need more experimental support would be to suggest a redistribution of the electronic wave functions near the surface increasing locally  $n(E_{\rm F})$  and inducing a Knight shift displacement of about 40 ppm as is the case with intercalated graphite<sup>(7)</sup>. The possibility of the observation of two carbon sites at the surface as may be apparent in the doublet structure is not excluded.

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R.DELTOUR Physique des Glides Université Libre Bruxelles 7050 BRUXELLES BELGIUM

P. BERNIER

#### UNIVERSITE D'AIX - MARSEILLE - FACULTE DE MEDECINE CENTRE DE RESONANCE MAGNETIQUE BIOLOGIQUE ET MEDICALE

Unité de Recherche 1186 Associée au Centre National de la Recherche Scientifique

(received 3/17/88)

Professeur Patrick J. COZZONE

Prof. B.L. SHAPIRO TAMU NMR Newsletter 966 Elsinore Court PALO ALTO CA 94303

#### Title : PHANTOM FOR LOCALIZED SPECTROSCOPY

Dear Barry,

The CRMBM is actively involved in research on muscular diseases in animals and humans using a Bruker Biospec NMR system equipped with a 4.7T - 30 cm horizontal magnet. Optimizing pulsing conditions with a surface coil and shimming the magnet can be rather lengthy procedures. They cannot be conveniently performed when the patient - often affected with a form of muscular dystrophy - is already uncomfortably installed with a limb in the magnet. We have found it very practical to conduct most of the adjustments prior to real clinical examination using a pork roast as a phantom. Usually, a 2-pound roast is adequate, especially if you have asked your butcher to shape the roast so that it approximates the section of a human limb. For P-31 spectroscopy, pulsing conditions are calibrated on the Pi peak which dominates the phosphorus spectrum recorded on the roast (Figure 1). The peak of water protons and its separation from the lipid resonance are used to adjust the field homogeneity. Fine adjustments are subsequently completed in a few minutes on the patient or animal limbs. Results are pretty satisfactory as shown on the attached spectrum (Figure 2) which was obtained in 2 min. on the leg muscle of a 50-pound pig, using a home-built 3 cm-diameter surface coil (collaborative work with Dr G. MONIN and Dr G. KOZAK-REISS).

Sylviane Confort-Gouny Patrick Cozzone Figure 1 Figure 2 И.И 8.8 -18.8 PPH PPH

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March 9, 1988 (received 3/10/88)

Dr. Barry Shapiro Editor/Publisher TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

#### Frankenstein-ian NMR: An Old World Solution to a New World Problem

Dear Barry;

Finding myself in danger of the pink slip after a stormy return to the TAMU fold, I can only ask for forgiveness and wish you a warm spring season in your new surroundings, and may the single traffic light between you and Pebble Beach be always GREEN.

On a more technical note, I would like to describe our recent solution to what must be becoming a more common problem: Old consoles die, but superconducting magnets live on!! Our old Bruker WM-300 spectrometer, originally purchased in 1980, had reached near expiration in the New World of NMR. It suffered from a small disk drive, no pulse programmer nor decoupler phase shifts to permit sophisticated pulse sequences of 1D, let alone 2D. After many years of fruitful service, we felt drastic measures were in order.

We shopped for a new 300 MHz console among the vendors, and we ended up with a GN-300 system. Our reasons were several and clearly of an individual nature, but I only want to describe how one weds Bruker and GE into a single functional spectrometer.

We retained the magnet, magnet power supply, room temperature shims, variable temperature unit, and probes from the Bruker system. Since both systems operate in a 50 ohm environment, probe interfacing and compatibility was not a problem; however, the major obstacle was to match the lock circuitry of the GN-300 console to the field control system of the Bruker power supply/shim system and to interface the shim gradient controls to the Bruker power supply. The shim gradient interface was accomplished with minimal patchwork via a box adapting voltage levels and connector types. The lock system proved somewhat more difficult!

The response of the GE lock system is much faster than the old Bruker WM-era. This resulted in dramatic oscillations of the lock/field control system-on even the strongest lock signals. Not only was this bad for resolution; it is impossible to shim! The problem was rectified by addition of capacitance to the integrator circuit in the GE lock system (Sorry to reduce performance of a nice circuit, guys!) to match the time response of the field control circuitry of the Bruker power supply. This led to a very happy and operationally transparent system.

The final shot in the magnet has been the addition of new probes from Cryomagnet Systems, Inc. We purchased 5 mm probes for normal broadband operation and a <sup>1</sup>H probe equipped with a broadband decoupling coil for indirect detection, etc. The performance of the system (<sup>1</sup>H S/N 137:1 and <sup>13</sup>C ASTM S/N 85:1, both measured in standard thin-wall tubes) is basically the same as a new instrument of a single maker. The performance of the broadband decoupler coil in the <sup>1</sup>H probe is especially exciting for <sup>1</sup>H{<sup>13</sup>C} applications to polysaccharides (c.f. Carb. Res, 166, 47-58(1987) and Tsui, et al. Carb. Res, in press). The broadband decoupler is of my own making. Performance of this unit is not available just yet. Hardware details of the lock circuit modifications are available for anyone who finds themselves in a similar situation. This summer we should be able to provide your readers with some performance reviews of the JEOL GSX-500, since we are expecting replacement of our JEOL GX 400 with a GSX-500 within a month or so.

Best regards,

R. Andrew Byrd Biophysics Laboratory, FDA 8800 Rockville Pike Bethesda, MD 20892

P.S. Also, I have an opening for a Postdoctoral position available Fall 1988. The applicant would be involved in solid-state studies of model and biological membranes.

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474 Medical Sciences Building, Telephone (403) 432-3006

Dr. Bernard Shapiro 966 Elsinore Court Palo Alto, CA 94303 U.S.A.

March 17, 1988 (received 3/21/88)

BACKGROUND RESONANCE IN 5 MM <sup>1</sup>H PROBE

Dear Dr. Shapiro,

Thank you for your pink reminder. We would like to report on some interesting background resonance results that we have obtained with our H probe. The probe, which is a 5mm H photo-CIDNP, will be used on our Nicolet 300 WB spectrometer. When we were building it, two critical questions arose: (a) what epoxy could we use for securing the coil to the insert; and (b) what materials should be used for the internal and optical supports that would produce the minimum background? This prompted us to investigate a selection of compounds. Difference spectra of the compounds, substracting the spectrum of the empty probe, show the varying background resonances that were obtained. The conditions were sw = +/- 50,000 Hz, 32 transients, 5 µsec pulse width, and line broadening 25 Hz.

We have since designed a self supporting coil which totally eliminates the use of epoxy.

Please credit this to Brian Sykes' subscription.



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#### UNIVERSITÄT TÜBINGEN PHYSIKALISCHES INSTITUT Prof. Dr. O. Lutz

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(received 2/27/88)

Prof. Dr. Bernhard L. Shapiro Editor/Publisher TAMU NMR Newsletter 966 Elsinore Court Palo Alto, California 94303 U.S.A.

Heteronuclear Imaging: <sup>23</sup>Na, <sup>27</sup>Al, <sup>51</sup>V

Dear Barry,

in the last time we try to transform our experience in the field of the usual heteronuclear spectroscopy to tomographic and spectroscopic investigations on a whole body Siemens Magnetom working at 1.5 Tesla. In fig. 1 a spectrum of the nucleus <sup>51</sup>V is presented which has been obtained with the standard <sup>23</sup>Na head coil of the imager. Some further results on imaging with hetero-nuclei are given in the paper: <sup>23</sup>Na, <sup>27</sup>Al, and <sup>51</sup>V Multinuclear NMR - Imaging and NMR - Spectroscopy with a 1.5 T Imager: M. Braun, O. Lutz, W.I. Jung, C.S. Kischkel, R. Oeschey, M. Pfeffer, Z. Naturforsch. <u>42a</u>, 1037 (1987).

Another well known spectroscopic technique, the <u>selective non excitation</u> of special spectral lines has also been transformed to the requirements of whole body imagers. With the five pulse SENEX - sequence images of the water or the fat distribution can be obtained with high signal-to-noise ratio and independently of the homogeneity of the RF - field. A paper appeared recently in Z. Naturforsch. 42 a, 1391 (1987): Selective Non - Excitation of Water or Fat Protons in Magnetic Resonance Imaging by M. Braun, W.I. Jung, O. Lutz, R. Oeschey. Applications are running in cooperation with the colleagues of the Radiologische Universitätsklinik Tübingen.

Sincerely

(Otto Lutz)



Fig. 1: <sup>51</sup>V spectrum of a 0.14 molal alkaline aqueous solution of NaVO<sub>3</sub>, using the <sup>23</sup>Na head coil of the 1.5 T imager. The signal arise from the VO4<sup>3-</sup> (large peak) and the V<sub>2</sub>O7<sup>4-</sup> species; their chemical shift difference amounts to 23,8 ppm. 32 acquisitions.

#### UNIVERSITÉ D'AIX - MARSHILE - FACULTÉ DE MÉDICINE CENTRE DE RÉSONANCE MAGNÉTIQUE BIOLOGIQUE ET MÉDICALE

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Professeur Patrick J. COZZONE

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CRNEM would welcome postdoctoral fellows or colleagues on sabbatical. Pinancial assistance can be arranged. Please contact P. COZZONE at 33 (Country code) 91 79 91 10 extension 1506.

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355-66

#### DOW CORNING



February 23, 1988 (received 3/17/88)

Professor Bernard L. Shapiro 966 Elsinore Court Palo Alto, CA 94303

SUBJECT: 2-D APPLICATION OF INAPT

Dear Professor Shapiro:

Greetings from Dow Corning.

Assignment of the structure in Figure 1 was straightforward from a C-C autocorrelated double quantum coherence experiment except for the sterochemistry about the ene. It was much too crowded for NOE to be useful so we had to resort to analysis of the long range C-H coupling constants from the vinyl proton to carbons across the double bond.

Bax's INAPT experiment<sup>2</sup> is ideal for transferring polarization from a single resolved proton to carbons 2, 3 or 4 bonds distant. When the  $\Delta$ , delay is incemented, Figure 2, a selective long range heteronuclear <sup>2D</sup>J resolved experiment is created, Figure 3. The long range H-C coupling constants now available are very useful for sterochemical assignments. This modification is similar to Nagayama's except that here all the proton pulses are soft. This is necessary on Nicolet NT series spectrometers that only allow rapid switching of the decoupler power between two levels.

Figure 1 shows the aliphatic correlations and coupling constants obtained from the 2-D INAPT experiment with the vinyl proton selected. The large 10Hz coupling to the methylene carbon at 36.9 ppm must be the three bond coupling constant trans to the vinyl proton and thus the stereochemistry of the ene is established.

This experiment should be applicable to the solution of many such stereochemical questions.

Best Regards,

Dick

Norman Rabjohn, Professor Michael S. Tempest, Assistant Professor Department of Chemistry University of Missouri-Columbia Richard B. Taylor Project Chemist Analytical Research Dow Corning Corporation

DOW CORNING CORPORATION, MIDLAND, MICHIGAN 48686-0995 TELEPHONE 517 496-4000



A C<sub>23</sub>H<sub>46</sub> OLEFIN

ALIPHATIC REGION OF THE 2D-INAPT SPECTRUM



FIGURE 2: Basic pulse sequence of 2-D INAPT. All proton pulses are soft and applied to a single proton resonance. This affords a selective long range heteronuclear 2DJ resolved spectrum, Figure 3.

- <sup>1</sup> Turner, D.L. <u>Mol. Phys.</u> 1981, <u>44</u>, 1051-1058; Turner, D.L. <u>J. Magn. Reson.</u> 1982, <u>49</u>, 175-178; Turner, D.L. <u>J. Magn. Reson.</u> 1983, <u>53</u>, 259-271.
- <sup>2</sup> Bax, A. J. Magn. Res. 1984, 57, 314-318; Bax, A.; Ferretti, J.A.; Nashed, N.; Jerina, D.M. J. Org. Chem. 1985, 50, 3029-3034; Taylor, R.B.; Corley, D.G.; Tempesta, M.S. J. Nat. Prod. 1986, 49, 670-673.
- <sup>3</sup> Jippo, T.; Kamo, O.; Nagayama, K. J. Magn. Reson. 1986, <u>66</u>, 344-348.

355-68

#### DEPARTMENT OF HEALTH & HUMAN SERVICES



#### **Public Health Service**

National Institutes of Health National Institute of Environmental Health Sciences P.O. Box 12233 Research Triangle Park, N.C. 2770

March 8, 1988 (received 3/14/88)

Dr. B.L. Shapiro Editor/Publisher TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

Title: Minimization of Baseline Artifacts in Phased 2D NMR

Dear Barry:

As anyone who has had experience with phased, absorption mode 2D-NMR probably knows, baseline curvature in the  $f_2$ -dimension can play havoc with the presentation and interpretation of the data set. As described in detail by David Hoult <u>et al.</u> (JMR <u>51</u>, 110, 1983), a major source of the curvature comes from the transient response of the audio filters to the incoming signal. By adjusting the time after the observe pulse one begins to sample the FID, one can minimize this distortion substantially. For some 2D experiments, however, this not enough.

The problem is that, to a first aproximation, each point in the f<sub>2</sub>-domain is modulated by a term proportional to  $\Sigma_i M_i$  cos $2\pi\delta_i t_1$ , where  $\delta_i$  is the chemical shift of the i-th spin (In phased 2D, a separate data set modulated by sin $2\pi\delta_i t$ terms is also generated). After now appears, when viewed in the f<sub>1</sub>-dimension, as an image of the entire spectra on each f<sub>2</sub>-file. Aside from the fact that these image spectra may show up as ugly ridges parallel to the f<sub>2</sub>-axis in a 2D contour plot, they can also obscure the interpretation when spectra with true crosspeaks and false image peaks superimpose.

To illustrate our point, we show in Figure 1A, spectra from the initial files of 2D NOESY experiments on 0.05M Gramicidin S in DMSO-d<sub>6</sub> using a mixing time for cross-relaxation of 0.1 sec. The problem created by the curved baselines can be seen in 1B, which show  $f_1$ -slices taken through the Leu  $C_{\alpha}$ H peak at 4.55 ppm. Although the major cross peak with the PheNH at 9.12 ppm is readily observed in all three spectra, the presence of smaller cross peaks signifying weaker or long range dipolar cross relaxation with the LeuNH, (8.36 ppm);  $C_{\beta}$ H<sub>2</sub> (1.35 ppm) and  $C_{\delta}$ H<sub>3</sub> (0.82 ppm) groups is quite ambiguous in the top and bottem spectra of 1B. The source of the ambiguity can be seen in 1C, which shows slices taken at 4.60 ppm, two files to the left, where there are no lines in the f<sub>2</sub>-domain (except for a little bit of the low field wing of the LeuC<sub>a</sub>H resonance).

The spectrum with the flat baseline was generated by inserting a delay and a 180° refocusing pulse between the observe pulse and the receiver 'on' flag, so that the FID reaches a maximum sometime after the receiver and the A/D converter have been turned on. Data points sampled before the maximum were discarded by left shifting prior to apodization and Fourier transformation.

## **A Few Useful Compounds**



٠	CAT. NO.	COMPOUND <sup>§</sup>	QUANT. <sup>†</sup>	PRICE <sup>4</sup> U.S.\$	CAT. NO.	COMPOUND <sup>§</sup>	QUANT. <sup>†</sup>	PRICE
-	84-70001	Deuterium-depleted Water (<0.5 ppm D)	25g 4 x 25g 10 x 25g	25. 90. 200.	83-84003	Toluene-α- <sup>13</sup> C	0.5g 1g	210. 360.
	82-70001	Deuterium Oxide	1kg	500. 45.	83-84002	1,2-Dimethyl- <sup>13</sup> C <sub>2</sub> -benzene (o-Xylene)	0.25g 0.5g 1g	160. 275. 475.
		(99.9 atom % D)	5 x 100g 10 x 100g 1kg	215. 425. 400.	83-84019	1,3-Dimethyl- <sup>13</sup> C <sub>2</sub> -benzene ( <i>m</i> -Xylene)	0.25g 0.5g	160. 275.
	82-79041	Deuterium (gas) (99.8 atom % D)	25L 50L 100L 500L	50.* 75.* 140.* 650.*	83-84009	1,4-Dimethyl- <sup>13</sup> C <sub>2</sub> -benzene (p-Xylene)	1g 0.25g 0.5g 1g	475. 160. 275. 475.
	83-00005	Methanol- <sup>13</sup> C	1g 5g 10g	105. 400. 775.	83-42001	Octanoic Acid-1- <sup>13</sup> C	1g 5g	105. 375.
	83-00508	Iodomethane- <sup>13</sup> C	1g 5g	35. 165.	83-42015	Decanoic Acid-1- <sup>13</sup> C ( <i>n</i> -Capric Acid)	1g 5g	100. 375.
			10g 25g	310. 725.	83-42004	Dodecanoic Acid-1- <sup>13</sup> C (Lauric Acid)	1g 5g	95. 350.
	83-70005	Potassium Cyanide- <sup>13</sup> C	1g 5g 10g	100. 415. 775.	83-02030	Hexadecanoic Acid-1- <sup>13</sup> C (Palmitic Acid)	1g 5g	90. 350.
	83-70006	Sodium Bicarbonate- $^{13}$ C (>98 atom $\%$ $^{13}$ C)	1g 5g	50. 200.	83-42011	Octadecanoic Acid-1- <sup>13</sup> C (Stearic Acid)	1g 5g	105. 375.
			10g 25g	300. 750.	83-02030	Hexanedioic Acid-1,6- <sup>13</sup> C <sub>2</sub> (Adipic Acid)	0.5g 1g	245. 410.
	83-02014	Sodium Formate- <sup>13</sup> C	1g 5g 10g	120. 500. 900.	83-12205	L-Alanine-3- <sup>13</sup> C	0.5g 1g	200. 355.
	83-02015	Sodium Acetate-1- <sup>13</sup> C	1g 5g 10g	40. 175. 340.	83-12301	L-Aspartic-3- <sup>13</sup> C Acid	0.05g 0.1g 0.25g	200. 375. 750.
	83-02018	Sodium Acetate-2- <sup>13</sup> C	25g 1g 5g	800. 85. 375.	83-12302	L-Aspartic Acid-4- <sup>13</sup> C	0.1g 0.25g 0.5g	150. 300. 475,
1 1	83-02020	Sodium Acetate-1,2- <sup>13</sup> C <sub>2</sub>	10g 1g	725. 175.	83-12308	L-4-Hydroxyphenylalanine-3 (Tyrosine)	- <sup>13</sup> C 0.1g 0.25g	400. 900.
			5g	825.	83-12202	L-Phenylalanine-3- <sup>13</sup> C	0.1g	175.

<sup>§</sup>All <sup>13</sup>C-labelled compounds are a minimum 99 atom % <sup>13</sup>C unless otherwise stated.
<sup>†</sup>Please request prices for quantities that you do not see listed.
<sup>‡</sup>Prices are FOB Miamisburg, Ohio for delivery in North America; please request prices for delivery to the other continents.
\*Cylinder charge, please request prices.

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All spectra were accumulated using a 200  $\mu$ sec homospoil pulse during the mixing period and a 16-step phase cycling scheme per t<sub>1</sub><sup>-</sup> interval (256 intervals in all). In the experiment with the extra refocusing pulse, the 16 steps included an EXORCYCLE rotation of the phases of that pulse and the receiver.

This contribution is to be credited to Barry Selinsky's subscription.

Best regards,

lon

Dr. Donald G. Davis Laboratory of Molecular Biophysics



## UNIVERSITY OF CALIFORNIA, RIVERSIDE

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DEPARTMENT OF CHEMISTRY

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#### March 15, 1988

Professor Barry Shapiro Editor/Publisher TAMU NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

#### Title: One Dimensional and Two Dimensional Multiple Quantum Filtered <sup>1</sup>H NMR of Monomeric Insulin

Dear Barry:

The recently successful installation of a GN-500 spectrometer in our laboratory finally enabled us to apply some of the more sophisticated NMR experiments to our research.

As our title suggests, we want to report some triple quantum filtered 1D and 2D data on monomeric insulin (M.W. ~7000) and show how they simplify the aromatic region and help us in the assignments of tyrosine, phenylalanine and histidine residues.

Figure 1A-B compares the one pulse <sup>1</sup>H spectrum of insulin to the 3QF 1D spectrum of insulin using a pulse sequence as follows:

Since spins in tyrosine and histidine residues normally do not possess triple quantum transitions, they are effectively edited out in the 3QF 1D spectrum.

Figure 2A and B show the contour plots of COSY and 3 QF-COSY experiments of the aromatic region of monomeric insulin. In the 3 QF-COSY only spins from phenylalanine residues are present and their connectivities are established by the cross peaks.

The pulse sequence is as follows:

 $90^{\circ}-t_{1}-\Delta_{1}-180^{\circ}-\Delta_{1}-90^{\circ}-\Delta_{2}-90^{\circ}-Acq(t)$ 

where  $\Delta_1$  in both sequences is approximately  $\frac{1}{4J_{HH}}$  (i.e. 30 MS) and  $\Delta_2$  is 5-10 µs.

We shall discuss the complete assignments of the aromatic region and the pulse sequence including the phase cycling scheme in our forthcoming publication. This work was carried out in collaboration with the Novo Research Institute (Bagsvaerd, Denmark), and the monomeric insulin sample was provided by Novo. Please credit this to Dr. Robert W.-K. Lee's subscription.

Robert W Los

Robert W.-K. Lee Academic Coordinator

RWK:nc

Enclosure

Sincerely,

Michael F. Dunn Professor of Biochemistry

Melinda Roy M. Roy



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Dr. Bernard L. SHAPIRO

Editor/Publisher

TAMU NMR Newsletter

966 Elsinore Court

### TRANSITION METAL INDIRECT DETECTION

Dear Barry,

Although the very first indirect detection experiments were accomplished on "non common" nuclei (N14, Hg199, ...), everybody is now applying these sequences on C13 or N15 isotopes.

I think that inorganic/organometallic chemists should use this detection scheme more often as it allows very comfortable experimental conditions (5 mm tubes, no enrichment,  $10^{-2}$  M solutions), for very unsensitive but important nuclei (Fe 57, 0s187, Rh103, ...).

Of course, indirect detection of these transition metal MR requires a slightly different approach when compared to C13 or N15 work, because :

- the H1 (F19, P31) X coupling may be unknown,

- the X chemical shift range can be spread over 20.000 ppm.

For this purpose, we developped a microprogramme, based on a 2Q heteronuclear sequence, which automatically steps through a wide X frequency range, recording the H1 (F19, P31) response of the X satellites for an array of JH1-X coupling constant values.

The attached figure examplifies such an automated procedure, run on an AM 400.

The knowledge of those two parameters (J value and  $\delta$  Fe range) makes very easy the start of the 2D version of the double quantum experiment to precisely measure the X chemical shift.

Best regards,

BREVARD

Compound with courtesy of Prof. R. BENN (Uni. Mulheim)



- Figure A : Frequency/Coupling search (X = Fe). Note the clearcut/maximum (arrow) at one precise frequency value in the Fe57 frequency space. Note also the explored Fe57 chemical shift range : 1540 ppm.
- Figure B : is an expansion of the arrowed region which again shows clearly a maximum response around the 1JH1-Fe57 value (9 Hz).

## 國立清華大學

# NATIONAL TSING HUA UNIVERSITY

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REPUBLIC OF CHINA

DEPARTMENT OF CHEMISTRY

March 17, 1988

CHEMISTRY BUILDING Professor Bernard L. Shapiro 966 Elsinore Court Palo Alto, CA 94303 U. S. A.

"2QF-COSY Spectrum of Cobrotoxin in Aromatic reagion"

Dear Professor Shapiro:

Cobrotoxin, contains 62 amino acid residues (Mr6949) with four disulfide bridges, is a neurotoxic protein isolated from the venom of Taiwan cobra (<u>Naja</u> naja <u>atra</u>). This protein blocks the neuromuscular transmission at the post-synaptic membrane by the specific binding to acetylcholine receptors.

Shown in the Figure is a 2QF-COSY spectrum of aromatic region for a 20 mM cobrotoxin in  $D_2O$  (pH=3.6). There are two tyrosines, two histidines, and one tryptophan residues that cause resonances in the plot. From the coupling pattern in 2QF-COSY spectrum, we can assign tryptophan and tyrosines resonances unambiguously and they are labelled on the top of convensional 1D spectrum. As to those two isolated siglets, labelled with "\*", are due to C4 protons of His-32( $\delta$ =7.36) and His-4( $\delta$ =7.05).

The 2QF-COSY experiment were recorded on our Bruker AM-400 spectrometer equipped with an Aspect-3000 computer. A shifted sine bell window function was applied on both dimensions, and spectrum were not symmetrized.

Sincerely,

Chi-Ying Wang

Chi-Ying Wang Research Assistant

rin //w

Chin Yu / Associate Professor of Chemistry



region

355-78



Professor B L Shapiro 966 Elsinore Court Palo Alto California 94303 USA

# Wilton Materials Research Centre

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8 March 1988 (received 3/19/88)

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Dear Professor Shapiro

WHY TWO-DIMENSIONAL NMR?

An attractive feature of two-dimensional NMR plots is their ability to convey an impression of sophistication in an NMR experiment. Gone are one-dimensional spin decoupling and nuclear Overhauser experiments to be replaced by the COSY and NOESY experiments with the seemingly complex contour or stacked plots. Of course two-dimensional experiments can be more efficient methods of obtaining spin and spatial connectivities but how often are the data fully analysed to realise this advantage?

None of us are above the desire to produce an attractive data set, particularly for the consumption of managers and supervisors. Take as an example determining the rate of a chemical exchange process. Doesn't the contour plot of the chemical exchange caused by ring flipping in 1,4 diphenoxybenzene, Figure 1, look better than the equivalent spectrum using a one-dimensional analogue? But which is a more efficient use of spectrometer time, why the one-dimensional experiment. In the time required to acquire the 2D data for one mixing time, eight one-dimensional experiments can be performed with a better signal-to-noise ratio allowing the rate constant of  $27s^{-1}$  to be derived.

I raise this issue merely to suggest some thought be given to the purpose of the two-dimensional experiment. Or is this simply envy, since my two-dimensional plots rarely have off-diagonal peaks?

Nigel Clayden

N J CLAYDEN Spectroscopy Group Room D115 Wilton Centre



FIGURE 1: 2D Exchange <sup>13</sup>C CP MAS HOR spectrum of 1,4 diphenoxybenzene at 294K with a mixing time of 50ms.

#### FIGURE 2

.

One dimensional exchange <sup>13</sup>C CP MAS NMR spectrum of 1,4 diphenoxybenzene at 294K with a 50 ms mixing time



Dr. V. Wray GBF Mascheroder Weg 1 D-3300 Braunschweig

Dr. B. L. Shapiro Editor/Publisher TAMU NMR Newsletter 966 Elsinore Court

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#### Transferred nOe's in flavonoid systems

Dear Dr. Shapiro,

We have been collaborating recently with Dr. Proksch of the Department of Pharmaceutical Biology of the Technical University of Braunschweig on the structure elucidation of a multitude of methoxyl substituted flavonoids. Although these compounds appear simple the actual determination of the substitution pattern is often no trivial matter. An observation that has proved helpful is that transferred nOe's can be observed when spectra are run in dry DMSO-d<sub>6</sub>. Although this phenomenon has been described in a standard text (J.K.M. Sanders and B. Hunter, "Modern NMR Spectroscopy" OUP 1987, p. 229) I cannot remember seeing any useful examples. I, therefore, proffer an example here.

Shown is the <sup>1</sup>H nOe difference spectrum of 6-methoxyluteolin obtained upon low power irradiation of the signal at 3.4 ppm. Transfer of saturation from this signal to the other slowly exchanging OH signals occurs, which in turn lead to positive nOe's for protons adjacent to these groups. Thus null signals are observed for H-6' and H-3, which allows a ready distinction between the latter and H-8. A similar result is obtained if the low field hydroxyl signals are irradiated.

Yours sincerely,

itor Wray

(Dr.) Victor Wray



Gesellschaft für Biotechnologische Forschung mbH

### Abteilung

Molekulare Strukturforschung

355-81





14/5/4/11 IS:WS

7 March 1988

Dr Bernard L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto California 94303 UNITED STATES OF AMERICA

Dear Dr Shapiro

## WOOD EXTRACTIVES ANALYSIS BY QUANTITATIVE <sup>13</sup>C NMR SPECTROSCOPY

Apologies for the late payment of our subscription. Our newsletters seem to have been taking some time to reach us, coming over here by seamail, back to the States and eventually back to us again. We thought this was all sorted out, but no, our overdue subscription reminder arrived via Jakarta, Indonesia.

Due to our interest in wood extractives, materials which can be extracted from wood with organic solvents, and their effects on the production and properties of mechanical pulps we required a quick method for analysing softwood extractives. The existing method, while giving good accurate results, is very time consuming to carry out taking of the order of 2 days per analysis.

Wood extractives contain a large number of compounds but may be conveniently grouped into four classes: free fatty acids; diterpene resin acids; fatty acid triglycerides; and unsaponifiables, a diverse group of compounds containing diterpene alcohols, hydrocarbons and sterols. We have found that in deuterochloroform the carbonyl carbons of fatty acids, resin acids and fatty acid esters appear at different places in the carbon spectrum. Hence by running the <sup>13</sup>C NMR spectrum quantitatively in the presence of a suitable internal standard (vanillin) it was possible to determine the amounts of fatty acids, resin acids and fatty acid esters in an extractives sample with an accuracy comparable to that of the classical procedure.

The T1's of the three carbonyl carbons were measured and found to range from 1 - 3s. The spectra were acquired with a repetition time of 15 s, a flip angle of close to 90° and broad band proton decoupling only during carbon signal acquisition.

Using this method we have found that wood extracts often contain significant amounts of fatty acid esters other than triglycerides so it is better to use the integral over the glycerol part of the triglyceride to determine these esters - this assumes that no di- and monoglycerides are present in the sample. An estimate may also be made from the spectrum of the amounts of other fatty acid esters present in the sample.

## WOOD TECHNOLOGY DIVISION

# FOREST RESEARCH INSTITUTE

0

Postal Address: Private Bag, Rotorua Telegraphic Address: 'Frestra' Rotorua, N.Z. Telex: NZ21080 Telephone: (073) 475-899 Fax: 479-380 This method is now routinely used at our Institute. Sample sizes range from 1.5 g (2 h/analysis) to 100 mg (overnight), and smaller samples if we could be persuaded to use the spectrometer to run them.

Yours sincerely

1/ Meders

R. Meder

Ian D. Suckling

Huckline



Fig. 1 Quantitative spectrum of a <u>Pinus radiata</u> dichloromethane wood extract.

Fig. 2 Expansion showing the carbonyl and glyceride regions from the spectrum of the same sample rerun with the addition of vanillin, the internal standard.

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No. 355 April 1988

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A <sup>1</sup>H spectrum, <sup>13</sup>C spectrum, an attached proton test (APT), and a <sup>1</sup>H-<sup>13</sup>C chemical shift correlation map (CSCM). All these analyses can be performed in as little as ½ hour, on as little as 50 mg. of sample, for most organic compounds. And the QE-300 does them all – automatically.

# With the NMR industry's most advanced automation.

This performance is made possible by the QE-300's automated software, hardware, and powerful MACRO programming capability.

Set-up starts with Autolock. Lock on as little as 10% CDCl3 in a 5 mm tube.

Use *Compushim* for touchingup spinning shims or complete shimming with both spinning and non-spinning gradients using the lock signal or observe FID. Autogain optimizes the receiver gain independently for

sequential <sup>1</sup>H and <sup>13</sup>C acquisition. After data acquisition, *Autophase* accurately phases <sup>1</sup>H and <sup>13</sup>C spectra.

And finally, the analysis is completed with Autointegrate.

All these routines can be called up from QE-300 MACROs. In fact, any QE-300 operation, including pulse programs, can be implemented via MACROs for automatic, unattended sample analysis.

#### And the most complete package of hardware accessories.

The QE-300 is available with the industry's most reliable, highest capacity (100 positions!) *Automatic Sample Changer*. Plus, you can add an array processor, a variety of hard disks, and switchable probes for even higher sample throughput and performance.

# Structural elucidation simplified.

For many organic molecules, the four experiments presented above will be all you need to determine or confirm molecular structure. For more complex applications, GE/NMR offers an extensive <sup>13</sup>C library with outstanding search capability. This library contains data from over 10,000 compounds and is currently being expanded using a QE-300 in operation at the Aldrich Chemical Company.

# High throughput and performance demonstrated.

Get all the facts on the GE/NMR QE-300. Better yet, arrange for a demonstration. Call the GE/NMR group at (415) 490-8310. Or write General Electric Company, NMR Instruments, 255 Fourier Avenue, Fremont, CA 94539.



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Linalool 1.5 6.0 5.5 4.5 3.5 5.0 4.0 3.0 2.5 2.0 6.0 5.5 5 0 0 5.5 10 **8** 8 8 6.0

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