#### **TEXAS A&M UNIVERSITY**



No. 394 July 1991

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#### FORTHCOMING NMR MEETINGS

33rd Rocky Mountain Conference on Analytical Chemistry, Denver, CO, July 28 - August 2, 1991; For information on the NMR Symposia, contact Dr. H. Eckert, Dept. of Chemistry, Univ. of California at Santa Barbara, Goleta, CA 93106, (805) 893-8163; For general information, contact: P. Sulik, Conference Chair, Rocky Mountain Instrumental Laboratories, 456 S. Link Lane, Fort Collins, CO 80524, (303) 530-1169.

Tenth Annual Scientific Meeting and Exhibition, Society of Magnetic Resonance in Medicine, San Francisco, August 10-16, 1991; Contact: S.M.R.M., 1918 University Ave., Suite 3C, Berkeley, CA 94704; (415) 841-1899, FAX: (415) 841-2340; See Newsletter 391, 55.

Two-Dimensional NMR Spectroscopy (ACS Short Course), New York, NY, August 23 - 25, 1991; See Newsletter 392, 33.

New Developments and Applications of Magnetic Resonance and Optical Spectroscopies (ACS Symposium), New York City, August 25-30, 1991; See Newsletter 393, 48.

International Conference on NMR Microscopy, Heidelberg, Germany, September 16-19, 1991; See Newsletter 385, 28.

1991 Joint Meeting FACSS/Pacific Conference, Anaheim, California, October 6-11, 1991; NMR/EPR Program Section Chairman: Prof. Cecil R. Dybowski, Chemistry Dept., Univ. of Delaware, Newark, DE 19716. Contact: FACSS, P.O. Box 278, Manhattan, KS 66502-0003.

Eighth Australian NMR Conference, Lorne, Victoria, Australia, February 2-6, 1992; See Newsletter 391, 38.

Eleventh Annual Scientific Meeting and Exhibition, Society of Magnetic Resonance in Medicine, Berlin, Germany, August 8-14, 1992; Contact: S.M.R.M., 1918 University Ave., Suite 3C, Berkeley, CA 94704; (415) 841-1899, FAX: (415) 841-2340.

Additional listings of meetings, etc., are invited.

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

#### DEADLINE DATES

No. 396 (September) ------ 16 August 1991 No. 397 (October) ------13 September 1991 No. 398 (November) ------11 October 1991 No. 399 (December)------8 November 1991

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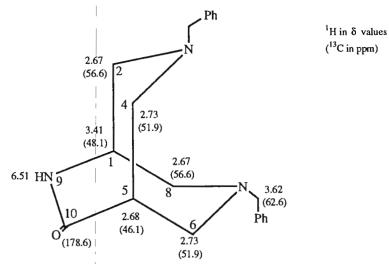
DEPARTMENT OF CHEMISTRY COLLEGE OF ARTS AND SCIENCES

Dr. B. L. Shapiro 966 Elsinore Court Palo Alto, CALIFORNIA 94303 June 17, 1991 (received 6/20/91)

TITLE: Proton and carbon analysis of 3,7-dibenzyl-3,7,9-triazabicyclo[3.3.2]decan-10-one

Dear Dr. Shapiro:

Our contribution this time concerns our studies on several novel heteroatom-substituted bicyclo[3.3.2]decane systems which are somewhat rare. We have obtained the following and 13C data on the example illustrated below. As one might expect, the nonequivalence of the protons alpha to nitrogen are observed as well as the bridgehead protons. Of course, there are sets of 13C signals in the ring system which are nonequivalent. Although we assume a chair-chair conformation, we have not been able to eliminate a chair-boat from consideration in certain



[3,7-Dibanzyl-3,7,9-triazabicyclo[3,3,2]decan-10-one]

solvents. A HETCOR 2-D plot in DCCl<sub>3</sub> confirmed the assignments, but it was not possible even at 400 MHz to resolve individual signals of pseudo axial and pseudo equatorial protons for H(2,4) and H(6,8). These materials will serve as useful synthons for certain medicinal agents in our laboratory. We trust this will be acceptable for our contribution this time.

Sincerely yours,

Derrell

K. Darrell Berlin Regents Professor 

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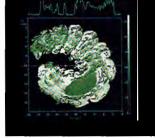
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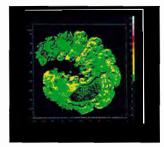
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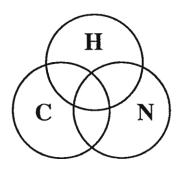
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May 17, 1991 (received 5/22/91)

Professor Bernard L. Shapiro Texas A&M University NMR Newsletter 966 Elsinore Court Palo Alto, CA 94303

#### Analysis of Spinning Sidebands for the Quadrupolar Nuclei with Half Integer Spin

Dear Barry,

NMR spectroscopy of quadrupolar nuclei in solids generally yields broad lines even when magic angle spinning is applied. Broadening of the central transition  $(-1/2 \leftrightarrow +1/2)$  of quadrupolar nuclei with half integer spins is caused by the anisotropy of the second order quadrupolar interaction. Recently we have developed a new approach for the simulation of lineshape for a quadrupolar nucleus under such a complicated variable angle sample spinning situation. It is known that an analytical expression for the central transition frequency can be obtained by average Hamiltonian theory (1). The perturbation in the modulation of the sample spinning can be written as a finite Fourier expansion in the angle  $\omega_{st}$ :

$$\omega(\beta,\theta,\phi,t) = \sum_{m=-4}^{4} W_m e^{im\omega_S t}$$

with

$$W_m(\beta,\theta,\phi) = \left(\frac{e^2 qQ}{2I(2I-1)\hbar}\right)^2 \frac{(I(I+1)-3/4)}{\omega_L} \sum_{l=0,2,4} \sum_{n=-l}^{l} A_n^l D_{n,-m}^{(l)}(\phi,\theta,0) d_{-m,0}^{(l)}(\beta),$$

where  $\omega_{\rm S}$  is sample spinning speed and the coefficient A are a function of asymmetry parameter  $\eta$ .  $\beta$  is the angle between spinning axis and external magnetic field. The Euler angles  $\theta$  and  $\phi$  specify the spinning axis in the principal frame of the electric field gradient tensor. The coefficient in the Fourier expansion  $W_0$  gives the frequency of the central line, while the  $W_m$  with  $m\neq 0$  describe the frequency modulation by the sample spinning which results in the sidebands. A method for computing the intensities of both the centerband and the sidebands for spin-1/2 nuclei has been reported by Sethi et al (2). The new method is readily extended to the case of quadrupolar nuclei and requires that only a set of banded homogeneous equations be solved to obtain the intensities for each spinning axis direction described by  $\theta$  and  $\phi$ :

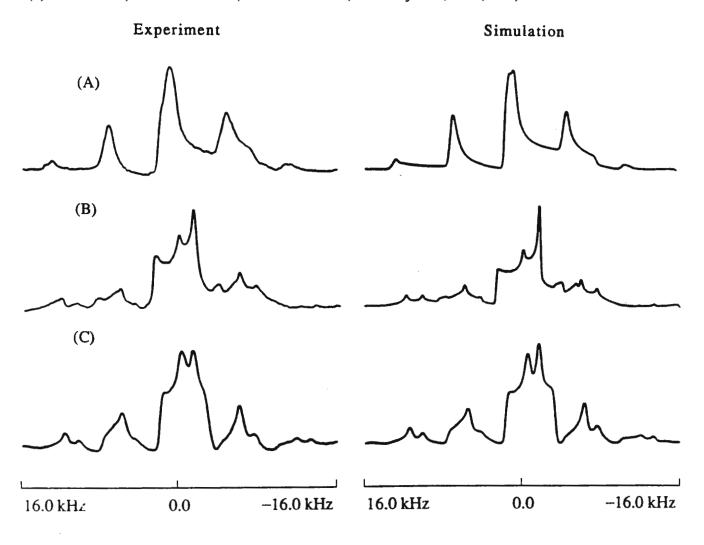
where  $w_m(\beta, \theta, \phi) = W_m(\beta, \theta, \phi)/\omega_S$ . Since terms with fourth rank Wigner rotation matrices are included in the frequency expression, the width of the banded matrix is expanded to include -4 to +4, rather than the -2 to +2 range required earlier by second rank interactions. In the powder sample for the set of all crystallites for which the spinning axis lies in a particular direction defined by  $\theta$  and  $\phi$  the magnetization is given by  $\mathbf{M}_{+}(\beta,\theta,\phi,t) = \sum_{N} \left| C_{N}(\beta,\theta,\phi) \right|^{2} \exp\{i[W_{0}(\beta,\theta,\phi) - N\omega_{S}]t\}.$ Department of Chemistry

$$\mathbf{M}_{+}(\beta, \theta, \phi, t) = \sum_{N} |C_{N}(\beta, \theta, \phi)|^{2} \exp\{i[W_{0}(\beta, \theta, \phi) - N\omega_{S}]t\}.$$
Department of Chemistry

This magnetization creates a series of peaks at frequencies  $W_0(\beta, \theta, \phi) - N\omega_S$ , each with intensity  $|C_N(\beta, \theta, \phi)|^2$ . Integrating  $\theta$  and  $\phi$  over the sphere produces the powder lineshapes including the spinning sidebands. The banded matrix method used in the simulations is computationally much more efficient than the standard method using Bessel functions. The complete details and analysis of this technique have been submitted for publication.

Shown in the following Figure are the experimental (left) and theoretical (right)  $^{23}$ Na NMR spectra of Na<sub>2</sub>SO<sub>4</sub> operating at 52.921 MHz. The angles of the spinning axis  $\beta$  are 45°(A), 54.7°(B) and 65°(C), respectively. The spinning speed is 6 kHz. Previously determined values of quadrupolar coupling constant  $e^2qQ/\hbar = 2.6$  MHz and asymmetry  $\eta = 0.6$  were used in the simulation. Good agreement between the experiment and theory is obtained. Na<sub>2</sub>SO<sub>4</sub> lineshapes are quite sensitive to  $\beta$ .

<sup>(2).</sup> N. K. Sethi, D. W. Alderman, and D. M. Grant, Mol. Phys. 71, 217 (1990).



Best wishes

Zhiwen Zheng

Zhiwen Zheng D. W. Alderman

David M. Grant

<sup>(1).</sup> A. Samoson, E. Kundla, and E. Lippmaa, J. Magn. Reson. 49, 350 (1982).

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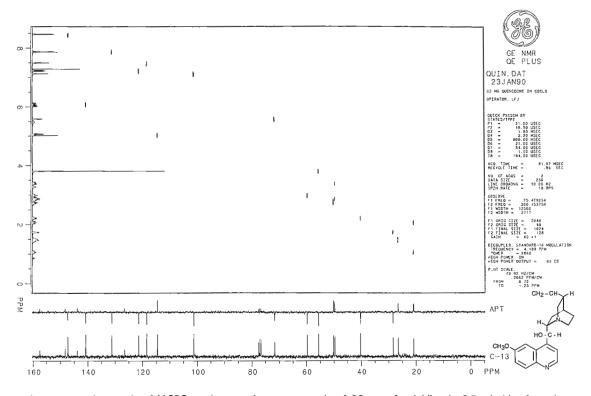
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Joseph R. Duke Department of Chemistry 142 Schrenk Hall Rolla, Missouri 65401-0249 (314)-341-6453 (or 4420) jd@blumiris.chem.umr.edu

May 30, 1991 (received 6/3/91) Dr. B. L. Shapiro 968 Elsinore Court Palo Alto, CA 94303

Dear Barry,

#### LOCAL ORDERING POTENTIALS IN <sup>2</sup>H POWDER PATTERN SIMULATIONS

SHBS forms lamellar liquid crystals with water. In these liquid crystals, the mobility of the phenyl ring is increased with respect to the dry surfactant. We have done lineshape simulations using two different models for phenyl group reorientation, free rotational diffusion and anisotropic viscosity. These are based on the method of Freed et al. (1-2) as adopted in our lab for <sup>2</sup>H NMR (3). For free rotational diffusion the motion is simulated by the inclusion of a fast reorientation about the molecular symmetry axis (the 1'-4' axis) (10 MHz), and a slower motion perpendicular to this axis (1 kHz). For anisotropic viscosity the motion is simulated by reorientation about a space fixed axis, the director axis. There are two rotational diffusion rates, one about the director axis (10 MHz) and one perpendicular to it (1 MHz). In this model, a restoring potential is used to mimic the intermolecular interactions which produce long range order. Spectra from the anisotropic viscosity model (bottom of Figure) seem to fit the experimental spectra best suggesting the headgroup motion is restricted with respect to reorientation away from the liquid crystalline director axis, but allowed to rotate freely about this axis. The local ordering, therefore, dominates to large extent the spectral features. Thus, these experiments provide evidence of the local environment in terms of structure as well as dynamics. We have submitted a paper on this topic.

Incidentally, I am planning to graduate soon with a Ph.D. in the area of NMR spectroscopy. I would be pleased to know of any job openings in this area.

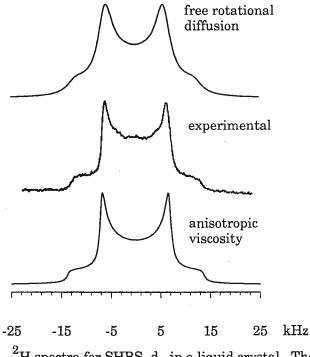
- 1. Freed, J. H. Spin Labeling: Theory and Applications, Vol. 1; Academic Press, Inc., New York, 1976.
- Schneider, D. J.; Freed, J. H. <u>Spin Labeling: Theory and Applications</u>, Vol. 8; Academic Press, Inc., New York, 1989.
- 3. Jagannathan, S.; Polnaszek, C.F.; Blum, F.D. J. Chem. Inf. Comput. Sci. 1987, 27, 167.

Sincerely.

Joe R. Duke

\*Please credit this contribution to F. D. Blum.

Sodium 4-(1'-heptylnonyl)benzenesulfonate (SHBS)



 $^2\mathrm{H}$  spectra for SHBS-  $\mathrm{d_4}$  in a liquid crystal. The top and bottom spectra are simulations.

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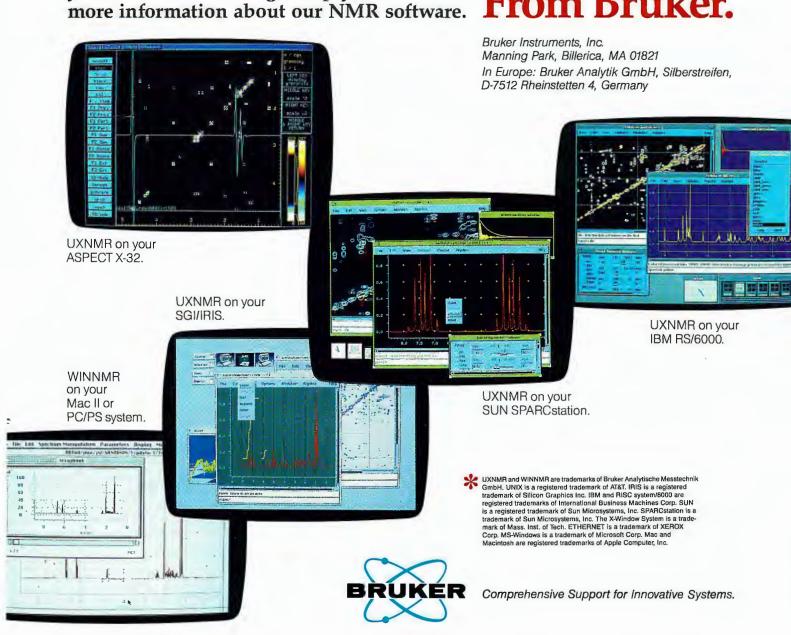
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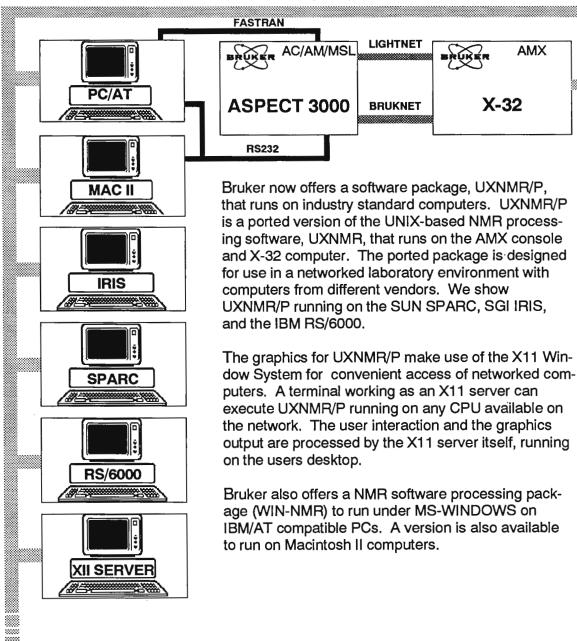
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June 3, 1991 (received 6/10/91)

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

Dear Dr. Shapiro:

Observation of  $^{15}N$ -labeled metabolites in isolated rat brain by  $^{1}H$ - $^{15}N$  HMQC.

Selective observation of <sup>1</sup>H coupled to <sup>15</sup>N by heteronuclear multiple quantum coherence (HMQC) spectroscopy permits indirect detection of <sup>15</sup>N-labeled compounds with a substantially higher sensitivity than direct <sup>15</sup>N NMR (1). Using this technique in Bruker AM-500 spectrometer, we have recently observed the amide protons of biologically <sup>15</sup>N-enriched [γ-<sup>15</sup>N]glutamine in isolated rat brain in 2 min of acquisition time (Fig.A) (2). Comparison with the conventional <sup>1</sup>H spectrum of the same brain (Fig.B) shows that proton signals from all other cerebral metabolites not spin-coupled to <sup>15</sup>N were suppressed. Sensitivity enhancement over direct <sup>15</sup>N detection was 50-fold. In an isolated liver of a similarly <sup>15</sup>N-enriched rat, [<sup>15</sup>N]urea protons were observed in 16 min (Fig.C). The HMQC method is likely to be effective for in vivo study of cerebral and hepatic nitrogen metabolism.

References.

1. A. Bax, R.H. Griffey, B.L. Hawkins, J Magn Reson. 55, 301 (1983).

2. K. Kanamori, B.D. Ross, and F. Parivar, J. Magn. Reson. 93, 319 (1991).

Yours sincerely,

Keiko Kanamori,

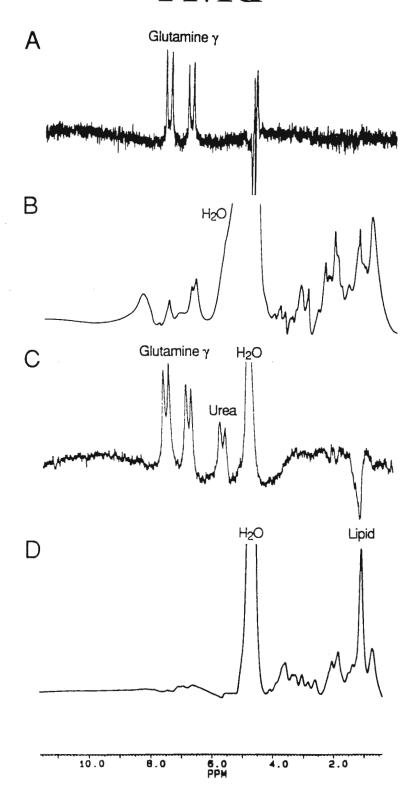
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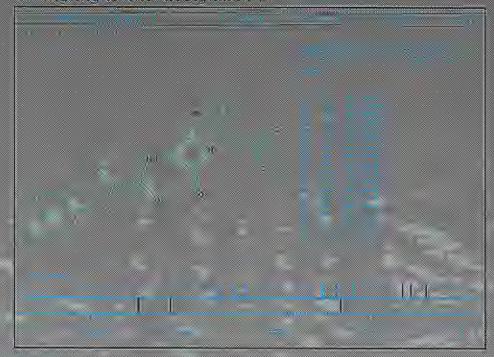
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Dr. B. L. Shapiro TAMU NMR Newsletter 966 Elsinore Court Palo Alto, California 94303 U.S.A.

(received 6/7/91) 31 May 1991

Hypophosphite Two-Spin Order Relaxation and Biological Applications

Dear Barry,

We have previously used  $^{31}P$  spin—transfer,  $^{31}P$  longitudinal relaxation measurements and  $^{31}P$  pulsed field gradient NMR measurements to study the molecular dynamics and transport of the hypophoshite ion ( $H_2PO_2^-$ ; HP) in human erythrocytes [1–3]. Later experiments revealed differential longitudinal relaxation in solutions of low viscosity (i.e., the rate of molecular reorientation is within the extreme motional narrowing limit). This differential relaxation is caused by chemical shift anisotropy — dipole—dipole cross—relaxation. The presence of this cross—relaxation gives rise to new methods with which to study biological systems. Selective study of cross—relaxation, which is extremely sensitive to reorientational correlation—time, has been shown to be a useful tool for elucidating relaxation mechanisms and molecular dynamics [4–9].

The first step in this series of experiments was to properly characterize the  $^{31}P$  relaxation mechanism, including terms for cross—relaxation, in  $D_2O$  solution [10]. The relative orientation of the chemical shift shielding tensor with respect to the molecular axes was determined from two—spin order relaxation measurements (e.g., [7–9]). The chemical shift anisotropy—dipole—dipole cross—interaction gives rise to the two—spin order signal. Next, we studied HP  $^{31}P$  two—spin order relaxation in  $H_2O$ , glycerol solutions and in suspensions of human erythrocytes with different cell volumes [11]. Using two—spin order relaxation to study the molecular dynamics of the hypophosphite ion has the great advantage of allowing the separation of the intra— and intermolecular (i.e., random field) contributions to the total relaxation rate. The results revealed that random field effects can have significant effects on the relaxation of species in biological systems, and must be accounted for if accurate estimates of the reorientational correlation time are to be obtained.

Differential line broadening [4–6], which occurs when the rate of molecular reorientation is outside the extreme narrowing limit, is observed for the proton—coupled <sup>31</sup>P spectrum of HP (a triplet) in a solution of albumin (see Fig. 1). We are currently investigating the use of two—spin order relaxation and differential line broadening measurements to study binding processes.

Yours sincerely,

Lian-Pin Hwang William S. Price Ching-Lung Tsai Baw-Ching Perng

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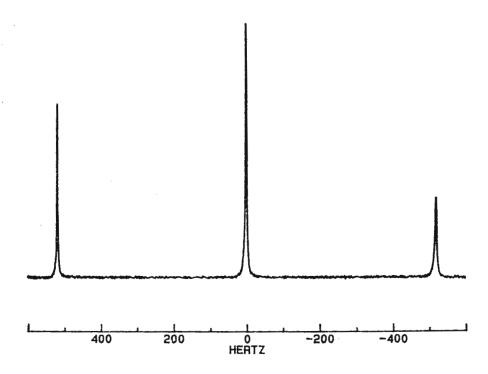


Figure 1. Differential linebroadening of the proton-coupled <sup>31</sup>P spectrum of HP (80 mM; pH 7.4) in bovine serum albumin (0.1 g/ml) at 121.4 MHz and 280 K. The effects of differential linebroadening are clearly shown by the difference in intensity of the satellite peaks. The line widths (from left to right) are 2.0, 3.3 and 5.5 Hz, respectively.

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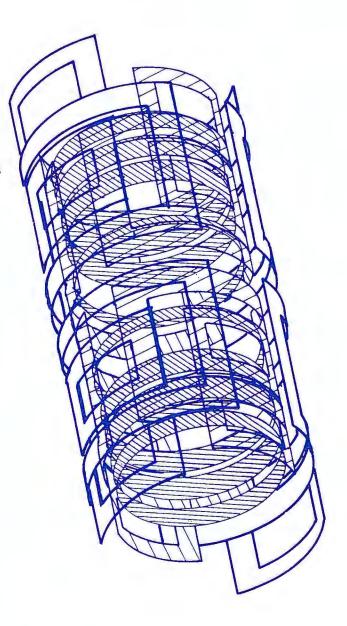
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FAX # 608-262-0381

5 June 1991 (received 6/8/91)

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

RE: Deuterium Quadrupole Coupling Constants (Theoretical and Experimental).

Greetings, Barry!

We have recently been carrying out deuterium NMR relaxation time measurements and ab initio calculations of deuterium quadrupole coupling constants,  $\chi_D$ , in small, garden variety molecules and ions, such as the phosphite ion,  $PHO_3^-$ . At a pH of 9 or higher there is no exchange (at least not over the period of five years that we have been monitoring a sealed sample) of the deuterium with the protons in the  $H_2O$  solvent. By measuring the  $^2H$  longitudinal relaxation time as a function of temperature, one can obtain accurate experimental values for  $\chi_D$ ; the value we obtain is  $94.7 \pm 0.5 \text{ kHz}$ . This result is in excellent agreement with Gaussian 86 ab initio calculations, if the proper bond length is used for the P-D bond. Because of vibrational motion the average P-D bond length is about three percent longer than the  $r_e$  value obtained from an ab initio calculation. The values for  $\chi_D$  depend on the internuclear distance. For the phosphite anion they change as shown in the figure below. Using an  $r_{PD}$  value of 150.5 pm obtained from NMR measurements, the ab initio value obtained for  $\chi_D$  in  $PDO_3^-$  is 95.0 kHz, in excellent agreement with the experimental value. We would be most interested in hearing from other readers who have similar results on other molecules. Are we again a mailee in good standing?  $\nu$ 

Golf Ho!

Thomas C. Farrar Professor of Chemistry

Tan.

TCF/mm

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\* Yeach! Ats Mobil Res. & Dev. Corporation Dallas Research Laboratory 13777 Midway Road Dallas, Texas 75244

June 14, 1991 (received 6/21/91)

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

Re: NMR Imaging of Pore Structures and Flow

Dear Barry:

Here at Mobil we have been getting very interesting NMR images of liquids in porous rocks from 1.0 to 100 mm diameter using our Bruker system with micro-, mini-, and macro-imaging probes together with a 15 cm bore vertical magnet and a 40 cm bore horizontal magnet. I wish to briefly mention some of our work performed with the mini-imaging probe.

NMR imaging can be used to map the pore space distribution in liquid-saturated porous media. When the resolution is sufficient, the actual pore structure can be imaged. dimensional imaging (both spin-warp and back-projection) is especially powerful for obtaining the structure of a whole system of pores. We found that imaging carbonate rocks, with their very complex pore systems, gives very interesting results. connectivity algorithms, we can select networks of connected pores and estimate liquid flow paths. Alternatively, we can set up the imaging protocol to greatly attenuate the imaging signal from the stationary liquid when liquid is flowing through the The relatively unattenuated signal indicates the major Because of dead-end pore space, the latter method is flow paths. the more accurate. In some cases, we found that only a fraction of the connected pore space contains the liquid flow path. have used NMR imaging to visualize the various types of pore structures and flow paths in limestones with two-dimensional slices and three-dimensional rendering.

Sincerely,

Don

D. E. Woessner

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May 22, 1991 (received 5/28/91)

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

<sup>1</sup>H-Detected C-C TOCSY Experiments: Tempting the Probe Gods

Dear Barry;

Over the last year, we have been employing isotope enrichment techniques and the associated NMR techniques to our ongoing studies in polysaccharide structure determinations (Byrd, Freese, & Vann, 32<sup>nd</sup> ENC, St. Louis, MO). One of the problems which is particularly troublesome in these systems, namely small <sup>3</sup>J<sub>HH</sub> coupling constants by which H-H correlations may be generated, is efficiently overcome by making <sup>13</sup>C an abundant spin and using the techniques developed by Fesik & Zuiderweg and by Bax, Kay, Clore, Gronenborn and coworkers.

We have generated several isotopically enriched polysaccharide systems; however, for clarity, we would like to show here the efficient principle of generating H-H correlations via C-C and H-C coupling interactions with glucose. Figure 1 shows calculations, done in collaboration with Mark Rance of Scripps Institute, for the TOCSY transfer functions amongst <sup>13</sup>C spins in a uniformly labeled glucose sample. The pertinent <sup>1</sup>JCC couplings were measured in a conventional 2D J spectrum, and the plot shows transfer from C1 to sequential C2, C3, C4, C5, and C6 as a function of idealized isotropic mixing time. The dashed vertical lines correspond to mixing times of 6 msec, 12 msec, and 24 msec. It is clear that at 6 msec, one would expect transfer to both the 2 and 3 positions, while at 12 msec and 24 msec there should be transfers to C5 and C6 respectively. Experimental verification of this is provided in Figure 2, which shows f2 traces through the α-anomeric proton for the corresponding mixing times of Figure 1.

This experiment was performed using a 2D sequence in order to demonstrate the mixing. The actual sequence consists of a  $^1H^{-13}C$  coherence transfer (INEPT) step with frequency encoding of the  $^1H$  shift in  $t_1$  followed by  $^{13}C^{-13}C$  TOCSY mixing via a DIPSI-2 sequence and finally a coherence transfer back to  $^1H$  via an INEPT process for acquisition (Bax, et al., JMR 88 425-431, 1990). The real benefit, and application in our lab, comes from utilizing the experiment as a 3D sequence where the resultant spectrum contains  $^1H^{-1}H$  "TOCSY" planes separated by the  $^{13}C$  chemical shift of the site where the first coherence transfer occurred.

This data was collected on a GN300 spectrometer using a home-built X-decoupler channel, ENI 550L amplifier ( $\gamma B_2$  was 5 kHz), and a Cryomagnet inverse probe. Our 3D studies using these sequences have been performed on a JEOL GSX-500 with basically standard inverse probe and components except for the use of the ENI 550L amplifier for the  $^{13}$ C channel ( $\gamma B_2$  was 7.5 kHz). These experiments really do *tempt the probe gods*, since most manufacturers never intended to see spin lock RF pulses of up to 30 msec on the X-channels of liquid probes...but just be careful!

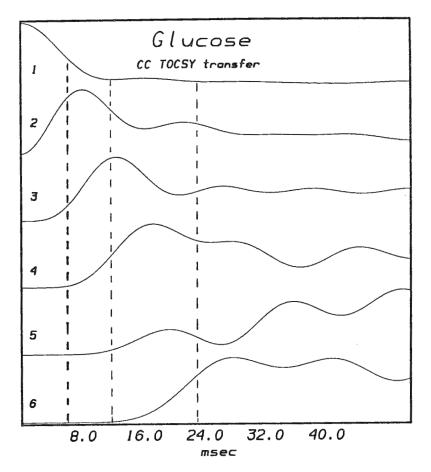


Figure 1: TOCSY transfer functions for Glucose. The calculation ignores relaxation and utilizes measured <sup>1</sup>J<sub>CC</sub> values.

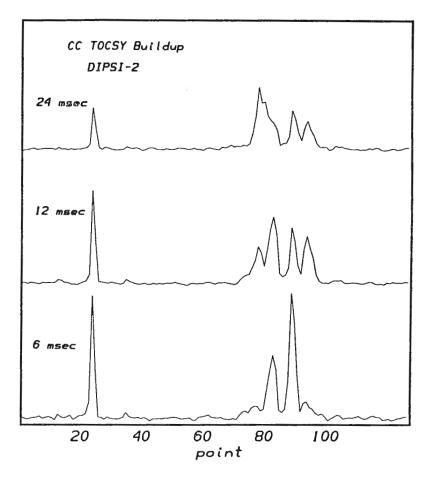
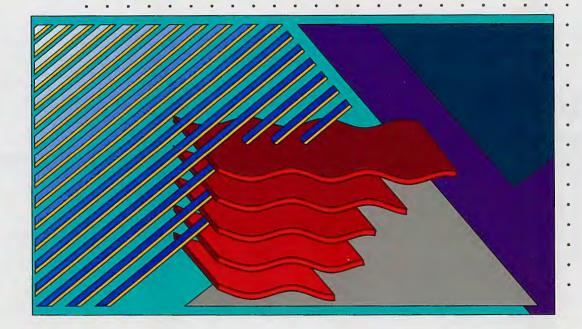


Figure 2. Traces through a 2D HCCH TOCSY experiment. Note the agreement with intensities both of the origin site and the sites of transfer.

Best regards;

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

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#### Dear NMR Colleagues:

I'd like to invite you to an NMR workshop that you just shouldn't miss--especially if you're attending the Rocky Mountain Conference. Each year following the conference in Denver, Chemagnetics invites NMR scholars to give seminars on advanced solids topics and organizes a 2-day opportunity for NMR scholars to visit and get up-to-date on the latest in NMR. This year's workshop will be held August 2 & 3 at the new Chemagnetics facility in Ft. Collins, Colorado.

We are privileged to have luminaries from both academia and industry. Below is a list of seminars that will be presented on Friday. Saturday will feature hands-on demonstrations of advanced solids NMR techniques. Registrants may attend one or both days.

Those presenting Friday and their topics are:

■ Dr. Gary Drobney, University of Washington	"Solids NMR Study of DNA Structures."
■ Dr. Bob Griffin, MIT	"Distance Measurements with MAS."
■ Dr. Mark Conradi, Washington University	"High Temperature NMR, Technique & Application to the Anomaly in Metal Hydrides."
■ Dr. Bernie Gerstein, Iowa State University	"Technology and Gimmicks for Low Temperature CPMAS Without Tears."
■ Dr. Naresh Sethi, AMOCO Corporation	"New Methods to Simulate NMR Powder Spectra."
■ Dr. Jim Haw, Texas A&M	"Isotropic Reconstruction: The Second Chapter."
■ Dr. Yue Wu , U. of California, Berkeley	"DOR, Technique and Applications."
■ Dr. Dan Weitekamp , California Institute of Technology,	"MRI of Solids with RF Field Gradients and Multiple-Pulse Line Narrowing."

If you'd like to get up-to-date on the latest in solids NMR, I urge you to register today. Registrations are limited to 60 people on the first day and 15 for day two. There is no fee to attend the seminar, but registrants are responsible for their own transportation from Denver to Ft. Collins (approx. \$13), for the hotel, and meals. To help defray your costs, rooms at Chemagnetics' discount rate of \$33.50 single/\$37.50 double have been booked at the nearby Ramada Inn. Please call today to register at 1-800-4-OTSUKA and ask for Sheila or Patti.

"Solid-State Imaging: Review and Prospects."

I look forward to seeing you.
Robert Wind

Dr. Robert Wind

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May 31, 1991 (received 6/21/91)

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

IN VIVO DETERMINATION OF INTRACELLULAR pH AND OTHER PHYSIOLOGICAL PARAMETERS BY <sup>31</sup>P NMR SPECTROSCOPY, IN THE STUDY OF PLANT RESPONSE TO ANOXIA. POSITION WANTED FOR A RUSSIAN COLLEAGUE.

Dear Barry,

as probably I have been already dropped from your mailing list, I apologize for the trouble that my delay induces. It is a pleasure for me to send you a note from my young colleague, Dr. Enzio Ragg, as a contribution of our laboratory. At the same time, I would ask you a favor for a Russian colleague, who wants to find a position in a western country, to work in NMR field. He has 25 years experience in NMR application to inorganic, analytical, organometallic chemistry and asked me not to mention at present his name. If somebody of the NMR Newsletter readers is interested, would he please contact me.

Thank you very much, with kindest regards

Ofelli Rosanna Mondelli

In vivo  $^{31}$ P NMR spectroscopy was used to study plant response to  $^{0}$ 2 deficit, comparing the highly tolerant rice seedlings with the intolerant wheat seedlings. NMR spectroscopy indicated a strong acidification of wheat cytoplasm (pH = -0.8 pH units) and a milder one in rice cytoplasm (pH= -0.4 pH units) in agreement with the higher tolerance to anoxia of rice. The previously reported (MENEGUS F. et al., Plant Physiol. 90,29,1989) cell sap alkalinization during anoxia in rice, as compared to acidification in wheat, was reflected in the alkalinization of vacuolar content in rice but not in wheat. The requirement of cytoplasm acidification, as predicted by the widely accepted model of plant resistance to anoxia (ROBERTS JKM et al., Proc. Natl. Acad. Sci. USA 81, 3379,1984) is therefore satisfied also in rice. Details of the above results are reported elsewhere.

Here we focus on some other events, concomitant with the changes of intracellular pH, that have received less attention. Oxygen deficit reduces ATP level in rice tissues to 50% of that found in air (Fig. 1). In wheat tissues ATP signals practically disappear under anoxia. The same is true for UDPG level which is hardly affected in rice, whereas it is strongly depressed in wheat. This last finding seems to be connected to the ability of rice coleoptile, but not of wheat, to elongate during anoxia.

Concomitant with the decrease of ATP and other phosphorylated compounds, there is an increase in cytoplasmic content of the inorganic phosphate (Pc). Cytoplasmic acidification and cytoplasmic phosphate build-up are simultaneous events.

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The increase of cytoplasmic phosphate content may be relevant in determing plant resistance to anoxia, because it negatively affects the phosphorylation potential: P = [ATP]/[ADP][Pc].

During anoxia, therefore, P will be lower in wheat than in rice, not only because ATP level is lower, but also because the inorganic phosphate level is higher. Changes of ADP level induced by anoxia in the two species (<sup>31</sup>P NMR spectra of HClO<sub>4</sub> tissue extracts) are not much different, so their influence on P is less important.

The qualitative changes of the phosphorylation potential, outlined here from the in vivo NMR measurements, help to explain the higher tolerance of rice with respect to wheat to anoxia.

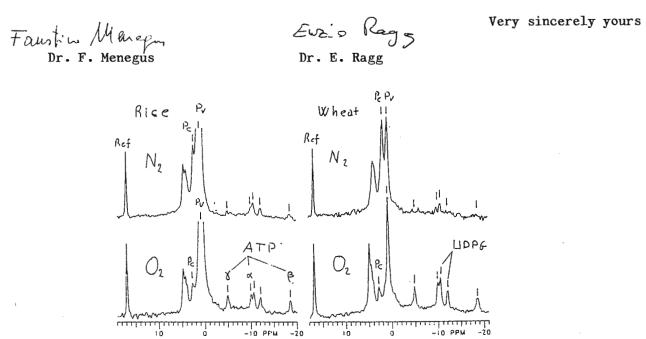
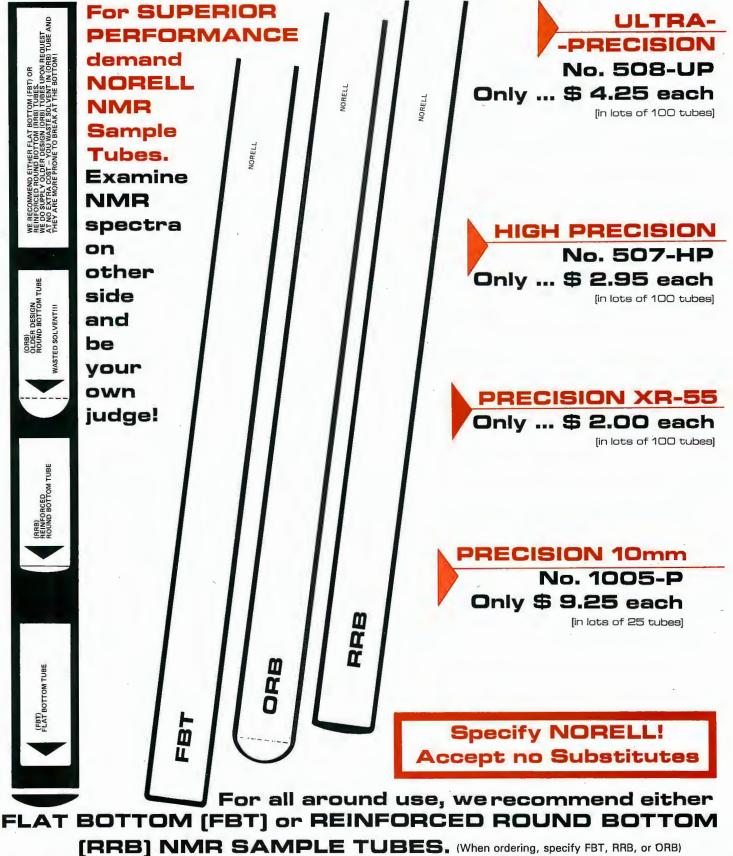


Fig. 1 - In vivo  $^{31}$ P NMR spectra (121.4 MHz) of aerobic (0<sub>2</sub>) and anoxic (N<sub>2</sub>) rice and wheat tissues. Pc and Pv refer to the signals of inorganic phosphate pools of cytoplasm and vacuole respectively.



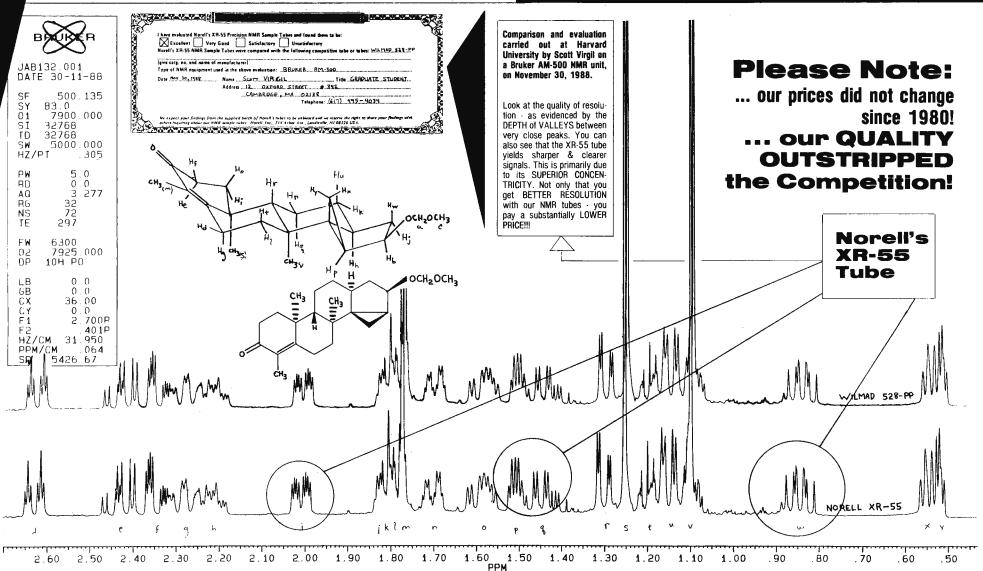
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Department of Physics Tel: 523523 Ext. 2403 DR. RAY DUPREE

28th May 1991 (received 6/3/91)

#### SLOW PROTONATION IN ISOPOLYTUNGSTATES

Dear Barry,

We have been having a lot of fun recently trying to sort out the aqueous chemistry of tungsten (VI) using <sup>183</sup>W and <sup>17</sup>O NMR. It is proving possible to recognise the relationships between many anions which have up till now been either unknown, or only known through one-off methods of preparation. The figures below show two of the better known isopolytungstate anions. Both of them normally hold two protons in their central cavity, in order to achieve a favourable distribution of charge. We have shown that metatungstate can be made with just one internal proton. It pulls in a second one within about one day, in aqueous solution. The H/D exchange rate in the diprotonated

The cavity in paratungstate B is less obviously closed off to the solvent, and sure enough one cannot now observe any separate resonance from the internal protons. However, we were initially surprised to observe significant line-broadening in the tungsten resonances. The broadening increases from nil to a maximum as the anion protonates. This was a surprise to us, because the tungsten-oxygen framework shows every sign of being stable on this timescale. We therefore tried moving from H<sub>2</sub>O to D<sub>2</sub>O solvent, and sure enough, all the linewidths more than doubled. It is not easy to get accurate linewidths in this system because the protonated paratungstate B anion undergoes further chemical changes, over a day or so, which alter the pH. (It goes to a new 11-tungsten species, but that's another story). Nevertheless, we are convinced that the widths shown in the Table below are reasonably reliable. As the broadening only occurs in the (tri)protonated anion, we reckon that it arises from protonation isomerism between a species with two internal and one external protons and one with three internal protons. Our oxygen data support this.

Although the cavity in paratungstate B is plenty large enough for protons, it is too small for any water molecule. Evidently protons do not move all that rapidly on an oxide-type surface without assistance from the solvent.

Table. Linewidth variations in acidified paratungstate B

Peak label	<sup>2</sup> H <sub>2</sub> O, 283K	<sup>1</sup> H <sub>2</sub> O, 283K	<sup>1</sup> H <sub>2</sub> O, 283K	<sup>1</sup> H <sub>2</sub> O, 268K	<sup>1</sup> H/ <sup>2</sup> H
	pH=2.8a	pH=3.1	$pH = 4.2^{b}$	pH=3.1	pH ≈ 3
I	7.6 <sup>c</sup>	3.4	3.0	6.0	$2.8^{d}$
11	15.2	7.2	3.9	16.2	2.3
III	16.2	6.8	3.2	11.5	2.6
lV	5.2	3.0	2.3	4.6	2.8

<sup>a</sup> Almost fully protonated (i.e. H<sub>3</sub>) <sup>b</sup> Just over half-protonated (i.e. H<sub>2 6</sub>)

<sup>c</sup> Width at half height in Hz, after removal of broadening due to window function

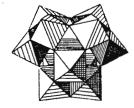
d Natural linewidth = 1Hz, as observed in the unacidified anion (i.e. H2)

With best wishes

Oliver Howarth and Jeremy Hastings

metatungstate, [H<sub>2</sub>W<sub>12</sub>O<sub>40</sub>]<sup>6-</sup>

paratungstate B,  $[H_2W_{12}O_{42}]^{10}$ 





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(617) 726-5819

Phone: 617-726-3083

Internet: jerry@nmr-r.mgh.harvard.edu

June 18, 1991 (received 6/20/91)

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

Modifying Standard Bruker CPMAS Probes for <sup>19</sup>F

#### Dear Barry:

We have two standard Bruker 7 mm CPMAS probes for our MSL400. In both of these probes, the high frequency channel is tuned for protons. We recently had occasion to measure some <sup>19</sup>F MAS spectra, and were dismayed to find that the proton channel of neither probe could be tuned to the fluorine frequency, only 6 percent away. Bruker quickly confirmed that, indeed, they themselves had tried it and failed, and that our only recourse was to purchase a probe built for <sup>19</sup>F.

Owing to my inherently cheap nature and to our desire to get the experiment done that same day, we decided to take a closer look at the Bruker tuning circuitry.

The high frequency channel of these probes employs a resonant line or coupling loop, the length of which is adjusted by the <sup>1</sup>H tune knob at the probe base. Although this adjustment reaches its low frequency stop well before the vicinity of the <sup>19</sup>F frequency, it turns out to be a simple matter to loosen the clamp which supports the moving part of the line and allows it to slide with respect to a stationary member. By pushing the clamp away from the probe head by about 0.5 cm and tightening it again (it's got to remain loose enough to allow sliding) it becomes possible to reach the <sup>19</sup>F resonance. The matching capacitor has the range to match perfectly at the new frequency. The limit stop on the tuning rod at the probe base may be backed up to a new position suitable for the full <sup>1</sup>H to <sup>19</sup>F range.

Placing marks on the resonant line to indicate the clamp positions for each frequency allows the nucleus switching operation to be completed in under five minutes. The more ambitious among your readers might consider machining a new clamp with a longer slot so that the retuning is possible without removing the probe cover.

Best regards,

Jerry Ackerman



Fremont Magnetic Resonance 3315 Seldon Court Fremont, CA 94539 415/623-0722 FAX 415/623-0851

#### File Transfers via Ethernet

**GOSSIP** provides transparent file transfers between a Nicolet 1280 and any Ethernet system supporting the tcp/ip protocol. You can configure your 1280 to upload every file as it is aquired, or you can manually select the files to upload in background.

At 19.2 KBaud, file transfers occur at about 21 K words/min. So a typical 512 x 4K 2D file can be transferred in about 100 minutes. Since the transfer occurs in parallel with the data aquisition, if the experiment takes more than 100 minutes, data transfer takes practically no additional time, just the time required to transfer the last block. Multi-block files larger than the available disk space on the host spectrometer can also be created.

GOSSIP version 1.12 offers automatic configuration and file conversion (conversion requires purchase of the appropriate data translation software from FMR). Configuration is achieved via a downloaded text file, therefore no user input at the PC is required. A keyboard and screen at the PC might be desired for greater flexibility.

Template macros for the Nicolet 1280 are provided that demonstrate: 1) data aquisition, then simultaneous data processing and file transfer, and 2) 2D data aquisition with concurrent file transfer.

#### **Features**

- No-hassle installation
- Automatic, customizable operation
- 20K data words/minute transfers
- Background transfers for 1D data
- Concurrent background transfers for 2D or kinetics data
- 2D files can be larger than the spectrometer disk space

FMR offers flexible packaging of GOSSIP to meet your requirements. You may purchase a turnkey system, or aquire the parts independently and buy only the software from FMR. Since there need be no user interaction at the PC, you may choose a 'black box' PC, without keyboard or moniter.

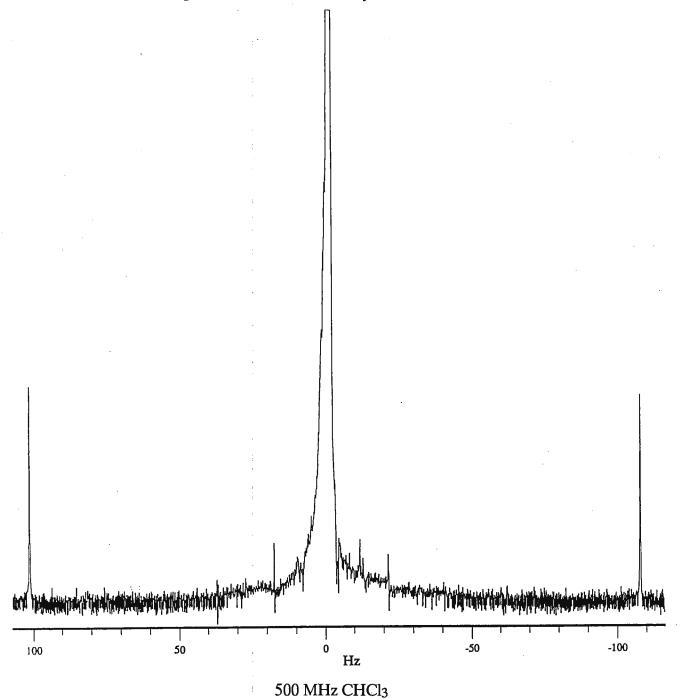
GOSSIP is very reasonably priced, and is available from stock. It can be a very cost-effective way to add your older spectrometers to the lab network. Why not order a copy and see for yourself?



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# **Probes and Probe Upgrades**

FMR offers probes for most systems. Upgrading an old, unused probe to a modern technology reverse detect or 1H can bring new usefulness to older systems.



#### The University of Texas Medical Branch at Galveston

School of Medicine Graduate School of Biomedical Sciences School of Allied Health Sciences School of Nursing Marine Biomedical Institute Institute for the Medical Humanities UTMB Hospitals



Marine Biomedical Institute

(received 6/24/91)

Barry Shapiro, Ph.D. TAMU NMR NEWSLETTER 966 Elsimore Ct. Palo Alto, CA 94303

Measurement of Water Content in Bone Graft Samples

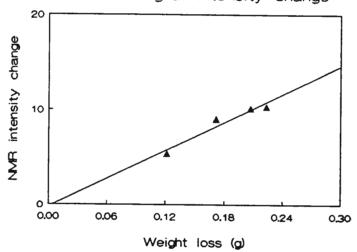
Hey Barry,

The tissue bank at Shrine Burns Institute here on our campus asked us to assess moisture content in bone samples using NMR spectroscopy. According to their technical manual, NMR is a reliable way to quantify the percentage water content. In order for their samples to be useful they must contain less than 5% water by weight. Their air drying procedure is carried out at room temperature according to standard procedures. To perform quality control of moisture content we have used the following procedure.

In order to calibrate the NMR method, we compared proton NMR signal intensity change to gravimetric weight loss in sections of fibulas weighing between 1 and 3 grams. Our gravimetric water determination was done by heating the sample in a vacuum dessicator containing DRIERITE (CaSO<sub>4</sub>) at 65 degrees C until a constant weight was achieved (about 3 days). Other labs have successfully used higher temperatures for drying bone samples. The NMR measurement was made by 12 signal averages using a 90 degree pulse and a 10 second interpulse delay. Samples were run on a horizontal bore system operating at 200 MHz. The broad-line background signal originating from protons in the plexiglass coil body was subtracted out of each spectrum. Bone water  $T_1$  relaxation times, measured by inversion recovery, were around 1.5 seconds. Line widths ranged from 500 to 1800 Hz, and were found to be inversly related to water As shown in the graph, weight loss in bone samples varied linearly with change in NMR signal intensity. The constant relating NMR intensity change and water content was 0.02094 +/-0.00157 (mean +/- SD). This corresponds to a 7.5% uncertainty in the NMR measurement vs. gravimetric. On the initial samples we tested, the average water content was 6.83% +/- 0.566, indicating that their standard drying procedure did not produce a useful sample for bone grafting. The simple expedient of using pre-dried air in the drying procedure instead of ambient air (typically 90% humidity) is expected to achieve optimal water content.

Please credit this to L.L. Smith's account.

Gravimetric Weight Loss vs. NMR signal intensity change



#### References

- American Association of Tissue Banks Technical Manual, Section 2, B2.155.
- 2. Joe Harris, AO Labs & Research, Inc., Indianapolis, IN, Personal Communication.

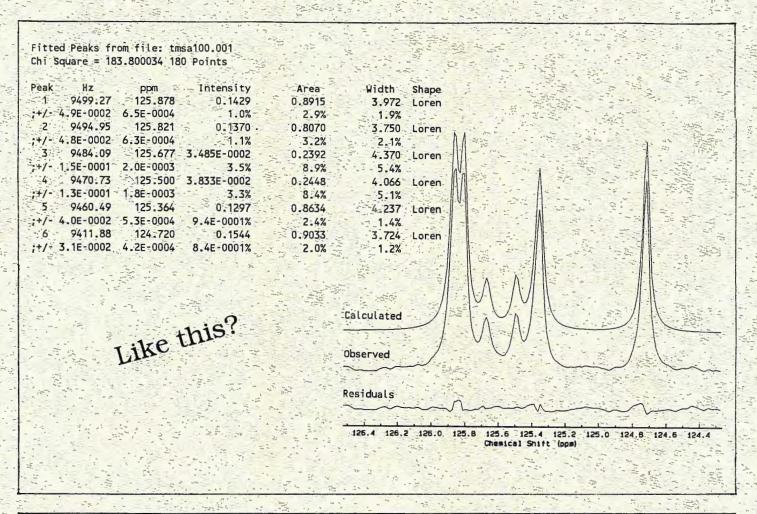
Yours in the double bond,

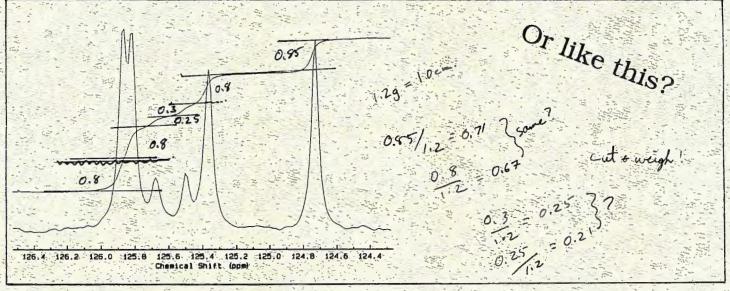
Mike Quast

Jose Gonzales

Stacy Klinke

# How does your present NMR software let you measure integrals?





# Quantitative NMR with NMR-286

NMR is inherently quantitative but our software usually leaves us guessing at the peak areas. NMR-286 uses an interactive Marquardt driven peak fitting routine to help you get the most out of your spectra. NMR-286 is a **must** for researchers doing serious quantification of their spectra. Look at these features:

- Parameters can be frozen at will to incorporate prior knowledge of the spectrum.
- The output contains the best-fit parameters as well as their standard deviations and the full variance-covariance matrix.
- You can compare the calculated and observed spectra and see the residuals in Dual Display.
- Clusters of up to 30 overlapping peaks can be pulled apart, even if the component peaks appear only as shoulders.

# NMR-286 is not just a line fitting program.

It is a powerful and complete PC based NMR processing package for one dimensional NMR. Here are some of its features:

- Lightning Fast FTs the fastest that you will find on a PC. Sizes up to 32k supported.
- Window Functions. Select from eight different window functions.
- Plotting. Plotting is extremely flexible. Send the HPGL output to a plotter, a file or a laser printer.
- Automated Processing. Almost anything that you can do from the keyboard, you can write a program to do. A powerful editor is a part of NMR-286.
- Manual. A comprehensive reference manual comes with NMR-286. It explains not only

the commands used but also the ideas behind them.

- Public File Formats. NMR-286 uses a file formats which are well described in the manual. You can import and export spectra as ASCII text files.
- · And much much more!

#### Computer Requirements

A Personal Computer which conforms entirely to the IBM standard. CPU: 80286/287 or better. OS: MS-DOS 3.0 or greater. Memory: 640k required. Graphics: EGA. VGA, Hercules or AT&T400.

#### **Spectrometers**

NMR-286 is in itself independent of the spectrometer. It requires a translation program to convert the data files to NMR-286 format. Currently such programs exist for Bruker DISNMR, DISCXP, DISMSL, TOMIKON and UXNMR files as well as for Lybrics format

files. Have another type of machine? Call!

#### Upgrade to 3.2

NMR-286 V3.1 users can receive an upgrade to Release 2 for a shipping and handling fee of \$50.00. Contact SoftPulse for more information.

#### Software Support

Each purchaser will receive a year of free software support by facsimile.

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Ī	1500.00
2-5	1250.00
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it, you'll like it. You can even try it **Risk Free!** Buy NMR-286 and if within 30 days you decide that NMR-286 is not for you, your purchase price will be refunded. Questions? Call, fax or write.

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NMR Processing Software for the Personal Computer





### Acorn NMR

46560 Fremont Blvd., #418 Fremont CA 94538-6482 Telephone: (415) 683-8595 FAX: (415) 683-6784

June 22, 1991

Dear Barry,

We have just established Acorn NMR as a spectroscopy service. As part of our preparations, we acquired a 10 year old NT360-WB. Shipping an old instrument across the country and getting it running is always fun. One nice feature of the NT console is the extensive operator feedback provided by all the lights and switches. These make debugging consoles much easier. The instrument is now up and running and we are in business. In the process of setting up we discovered a few modifications which have helped to make this older instrument more versatile. In the belief that they are of general utility, we would like to share some of them with the TAMU Newsletter readers.

To get rid of the old "clunky" TTY-43 keyboard, we substituted a 386 PC running the terminal emulator program within Windows 3.0. This configuration adds several new features to spectrometer operation. One very convenient feature is the ability to define a single mouse-activated "button" to execute a series of frequently used commands. This supplements the NT software limitation of having only one "link" list. For example, this gives us single button activation of various auto-shim sequences, another button for the ever useful ZGBCEMFTPS command string, and several others which are always available with a mouse click. The terminal emulation program also provides the ability to capture text output by the 1280, such as parameter lists, peak lists and pulse programs, into files on the PC.

Another useful feature of this configuration is that Windows lets you have multiple programs active simultaneously. It also permits rapid toggling between them which eliminates delays for each program to be started. Using this feature allows several programs to be simultaneously running in separate windows. A common configuration for us is 1) the terminal emulator for controlling the NT console, 2) GOSSIP for serial data transfers to and from the PC and 1280 at Baud rates up to 38,400, and 3) Felix/PC for data processing. All programs are available for immediate use.

Once data has been transferred to the NMR's PC, it is immediately accessible from one of several PCs at anyone's desk in the facility via a Local Area Network. Plots are output from Felix to our HP Series II laser printer, equipped with a Pacific Data Products "Plotter in a Cartridge" for HPGL emulation, obviating the need to purchase a dedicated plotter. Plots can also be output to a file in HPGL format, which can be imported into a variety of desktop publishing programs for customizing or incorporating into reports. For example, the sucrose plot included in this communication was imported into PageMaker. Using this capability in conjunction with the "captured" peak lists and other information from the 1280 allows us to prepare complete reports for our customers in both magnetic and hard copy form. They can then reprocess it later at their facilities. We can even E-mail the data and report for faster turn-around.

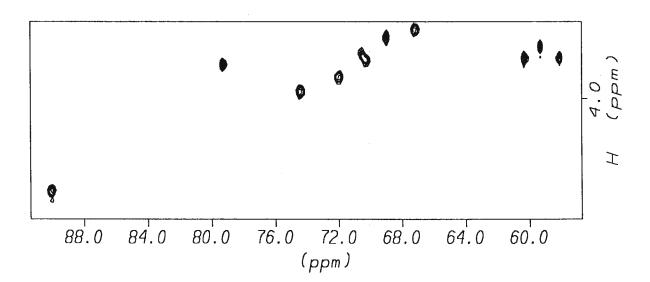
As we began working with 2D data, the advantages of Felix became more apparent. The 10 year old NT software is limited in its capability for user-interactive viewing and manipulation of 2D data. 2D FTs are also <u>faster</u> on our PCs: the sucrose spectrum shown (1K x 64) took 45 seconds for complete processing on a 486. Readers may also benefit from our experience with Felix 2D pro-

cessing, as it took a bit of trial-and-error to determine the correct combination of commands. An essential point is understanding how the data is actually stored in the computer. Complex NMR data is stored by different computers and software in different ways. Nicolet software stores complex data in two parts. The first half of the file contains the real half of the complex data set and the second half of the file contains the imaginary half. On the other hand, Felix stores the data intermingled, as alternating real and imaginary points. Listed below is a macro for transforming a magnitude (not phase sensitive) data set. The "trick" is in keeping the matrix type as real, even though we keep the imaginary part of the data until the last step of processing. The real and imaginary data are written to the matrix after the first FT as separate real and imaginary halves, with odd numbered rows being real and even numbered rows being the corresponding imaginary half. (This necessitates "lying" to Felix about the size of the data set and its real/complex nature, which can make the process rather confusing.) When a column is loaded for the 2nd dimension FT, it consists of alternating real and imaginary points, as Felix expects, and a complex FT is performed. (Thanks to Dennis Hare for explaining this.)

Gina Miner

Woody Conover

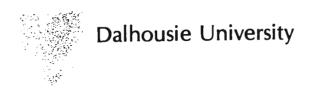
The plot below is the result of a 1K x 64 C-H correlation experiment on sucrose processed in Felix/PC with the mag2d.mac macro described.



#### Macro for 2D data processing in Felix/PC.

#### "mag2d.mac"

```
;close file & reset pointer
cl
                                             ;time-domain data file
get 'Enter data file name: ' name
get 'No. of t1 slices?' nrow
                                             ; number of spectra acquired
get 'Size of each?' ncol
                                             ; size (in COMPLEX points)
get '1st dimension Line broadening:' linbr1
get '2nd dimension Line broadening:' linbr2
                                             ; name for matrix file
get 'Enter name for matrix file:' mname
eva nrowx2 (2*&nrow)
eva ncolx2 (&ncol*2)
bld &mname 2 &ncol &nrowx2 0
                                             ;build matrix (real)
                                             ; open matrix, write-enabled
mat &mname write
                                             ; counter for matrix writing
def rrow 1
                                             ;loop for 1st dimension FT
for row 1 &nrow
re &name
                                             ;data type = complex
dat 1
em &linbr1
ft
                                             ;separate real/imag parts to
sep
                                             ; left/right sides
dat 0
                                             ;data type = real
                                             ;size
si &ncol
                                             ; write real half to matrix
sto 0 &rrow
eva rrow (&rrow+1)
                                             ; increment counter
                                             ;double size
si &ncolx2
shl &ncol
                                             get rid of 1st half
si &ncol
                                             ;reset size
                                             ; write imag. half to matrix
sto 0 &rrow
eva rrow (&rrow+1)
                                             ;increment counter
ty row=&row $
                                             ; type row number (no scroll)
next
                                             ;end 1st loop
for col 1 &ncol
                                             ;loop for 2nd dimension FT
loa &col 0
                                             ;load column from matrix
dat 1
                                             ;data type = complex
si &nrow
                                             ;size
zf &nrowx2
                                             ;zero-fill 2x
em &linbr2
ft
                                             ; magnitude spectrum
ms
                                             ; reverse spectrum
rev
                                             ; reduce to real
red
sto &col 0
                                             ;write to matrix
ty col=&col $
                                             ; type column number
next
                                             ;end 2nd loop
end
                                             ; end of macro
```



Department of Chemistry Halifax, Nova Scotia Canada B3H 4J3

> Tel: (902) 494-3305 Fax: (902) 494-1310

May 22, 1991 (received 5/30/91)

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

Dear Barry:

<sup>31</sup>P nmr study of solid mercury phosphonates

We have recently examined the  $^{31}P$  nmr spectra of several mercury phosphonates, (EtO) $_2P(0)\,\mathrm{HgX}$ , in the solid state. These compounds are characterized by very large  $^1J(^{31}P,^{199}\mathrm{Hg})$  values ranging from 12.6 kHz (X=I) to 13.8 kHz (X=Cl) and values of  $\delta_{11}$  -  $\delta_{33}$  which range from 135 ppm (X=Cl) to 170 ppm (X=SCN). For some of the derivatives it was possible to analyze the MAS and static  $^{31}P$  nmr powder line shapes to obtain estimates of the anisotropy in the  $^{31}P,^{199}\mathrm{Hg}$  spin-spin coupling constant. Values of  $\Delta J(^{31}P,^{199}\mathrm{Hg})$  were 2700 Hz (X=CH $_3$ COO), 1600 Hz (X=SCN) and 1500 Hz (X=I). The error in these values is estimated to be  $\pm$  250 Hz. Interestingly, the value of  $\Delta J/J$  in the mercury phosphonates (0.1 to 0.2) is significantly less than the corresponding values in a series of mercury phosphines (0.4 to 0.7).

The most important conclusion from this work is that mechanisms other than the Fermi-contact mechanism make substantial contributions to  ${}^{1}J({}^{31}P, {}^{199}Hg)$ , particularly in the mercury phosphines. Further details will appear in the ACS publication, Inorganic Chemistry.

Best regards,

Yours sincerely,

Rod Wasylishen

Rol Wary lishen

Bill Power

Michael Lumsden

RW: jw

<sup>1</sup>G.H. Penner, W.P. Power, and R.E. Wasylishen, Can. J. Chem., <u>66</u>, 1821 (1988).

P.S. We are preparing to write a review which will tentatively be titled, "Phosphorus NMR Studies of Solids." I would appreciate copies of any recent reprints and preprints that readers of the Newsletter would like us to include in the review. Thanks, Rod

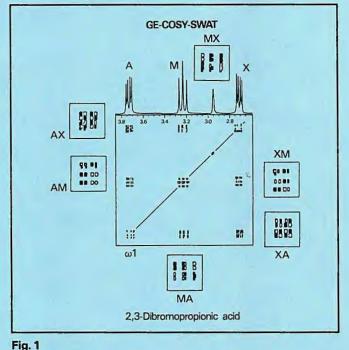
# Gradient Enhanced Spectroscopy SWAT

GE introduces the use of Switched Acquisition Time (SWAT) gradients to achieve pure phase 2D spectra with quadrature detection in both the acquisition ( $\omega_2$ ) and evolution ( $\omega_1$ ) dimensions without any phase cycling and without an additional set of  $t_1$  data.

One example of a pure phase gradient enhanced COSY spectrum of a solution of 2,3-dibromopropionic acid in benzene-d6 is shown in Fig. 1. SWAT gradients and a single acquisition per block were used. Data was collected on an Omega 300WB with Microstar actively-shielded gradients. A 5mm inverse probe was built for use within the gradient coils.

Digital resolution of 1.2 Hz in  $\omega_1$  and 2.4 Hz in  $\omega_2$  was achieved by collection of a 512 x 512 matrix with  $t_1$  evolution time of 840ms and a  $t_2$  acquisition time of 420ms. A single acquisition per  $t_1$  evolution data block and an average recycle time of 1.84s resulted in a 15 minute total collection period.

Since the SWAT gradient method encodes the necessary information in a single t<sub>2</sub> acquisition time, it avoids the collection of additional data blocks required by traditional pure phase methods. This time efficiency is especially important for collection of large multidimensional data sets.



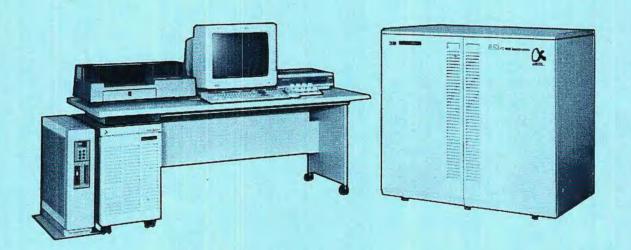
Contour plot of a 300 MHz pure-phase COSY spectrum of a solution of 2,3-dibromopropionic acid in benzene-d6 acquired with only a single acquisition per t<sub>1</sub> evolution time increment using the GE-COSY-SWAT method. Cross peaks are shown in expanded insets with positive peaks as darkened contours and negative peaks as open contours. A one dimensional spectrum is plotted across the top of the 2D spectrum.



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