

VARIED MAGNETIC FIELD, MULTIPLE-PULSE, AND MAGIC-ANGLE SPINNING PROTON NUCLEAR MAGNETIC RESONANCE STUDY OF MUSCLE WATER

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ABSTRACT The nuclear magnetic resonance linewidth of ^1H in water of frog muscle was studied as a function of magnetic field strength and angle of orientation. The results suggest that the observed spectra are dominated by demagnetization field anisotropy and dispersion, but a small static dipolar interaction of the order of a few hertz may be present. Data from line-narrowing, multiple-pulse experiments also indicate the presence of a small dipolar broadening.

INTRODUCTION

In many biological systems, water is present as a separate heterogeneous phase. In some systems in which the macromolecules are oriented, such as collagen (1, 2), oriented DNA (3, 4), silk fibroin (3), and keratin (3, 5), dipolar splitting of H_2O ($\Delta\nu_{\text{H}}$) and quadrupolar splitting of D_2O ($\Delta\nu_{\text{D}}$) have been observed in the nuclear magnetic resonance (NMR) spectra. On the other hand, the existence of dipolar or quadrupolar splitting for water in biological cells has not been unequivocally demonstrated. An observation of orientation-dependent NMR patterns for water protons in nerve fiber (6) was later attributed to the effect of demagnetization fields (7). Recently, Fung (8) reported changes in the linewidths of both proton and deuteron in muscle water as a function of the angle between the muscle fiber and the magnetic field. However, the interpretation of the results as an indication of water orientation has been questioned (9), and it was suggested that the change in the proton line shape of muscle water with orientation may also be the result of anisotropy in magnetic susceptibility (9).

In collagen (1, 2, 10), Li-DNA (4), and other nonbiological systems such as clay (11) and rayon (12), $\Delta\nu_{\text{H}}/\Delta\nu_{\text{D}} \sim 0.3$. However, the observed linewidths for H_2O in nerve (6, 7) and muscle (8) are larger than those for D_2O . $\Delta\nu_{\text{H}}$ and $\Delta\nu_{\text{D}}$ in hydrated collagen decrease as the water content increases (2, 13–16). Native collagen contains ~ 1 g H_2O /g dry protein. At this water content, the proton doublets overlap each other and are not resolvable (1). Skeletal muscle contains $\sim 75\%$ water, or 3 g H_2O /g dry weight. Since the water contents in muscle and nerve are much larger than that in native collagen, it is surprising that their proton peaks show apparent splittings (6, 8). These comparisons seem to favor the interpretation that the

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multiple proton peaks in these cell systems are due to dispersion of the demagnetization field (7, 9) rather than true dipolar splitting. It was suggested that the arguments can be settled by examining the proton linewidth as a function of the strength of the external magnetic field, H_0 (9).

Homonuclear dipolar splittings in solid or partially ordered systems can be reduced or eliminated by spinning the sample at the "magic angle" ($54^\circ 44'$) with respect to H_0 (17–19), at a frequency large compared to the dipolar interaction, H_D (s^{-1}), by using multiple pulse sequences (20–22), or by a combination of both (23, 24). The use of these techniques and a measurement of the variation of proton linewidth with H_0 might enable us to better understand certain characteristics in the NMR study of muscle water.

EXPERIMENTAL

Muscle of frogs (*Xenopus laevis*) was used in this study. Immediately after a frog was killed, the tibialis muscle was excised from the leg, and a striated piece was cut and aligned on a Teflon (DuPont de Nemours & Co., Inc., Wilmington, Del.) plug, which was then inserted into a NMR sample tube, placed upside down. Then, the sample was placed in the NMR probe in a magnetic field, and was rotated so that the axes of the muscle fibers could form specific angles with respect to H_0 . A proton NMR study at $H_0 = 23,487$ G was made by using a Varian XL-100 spectrometer (Varian Associates, Palo Alto, Calif.) in the continuous wave mode with external ^{19}F lock. Measurements at $H_0 = 11,744$, 5,872, and 2,936 G, respectively, were made on a home-built pulse spectrometer with a Bruker high-resolution (unshimmed) magnet (Bruker Instruments, Inc., Billerica, Mass.) at the University of Oklahoma. The total time for studying each sample at four frequencies was 2–3 h. Except at 23,487 G, the linewidth was defined as $\Delta\nu = 1/\pi T_2^*$, where T_2^* is the decay constant of the free induction decay (FID). Since at exact resonance the FID signal is sensitive to small drifts of H_0 for a system without field-frequency lock, T_2^* was determined from the envelope of the FID when the frequency was set at 1 kHz off resonance. $\Delta\nu$ determined this way was probably not exactly the same as the halfwidth of the Fourier transform of the FID. However, the error introduced was much smaller than uncertainties due to the variation of sample shape and size and the lack of perfect alignment. The values of $\Delta\nu$ reported are the average values of two measurements for the same relative θ (0° and 180° , 30° and 150° , etc.). Multiple-pulse and magic-angle spinning experiments were performed on a home-built spectrometer at Iowa State University. The spectrometer operated at 56 MHz for protons ($H_0 = 13,153$ G), with a deuterium lock. 5-mm (OD) oriented samples were used in the multiple-pulse experiments, and unoriented samples were used for the magic-angle spinning experiments. The spin-spin relaxation time, T_2 , was obtained by using the Carr-Purcell-Meiboom-Gill sequence (25, 26). The constructions of the spectrometer and the special probe are described in detail elsewhere (24, 27).

RESULTS

In a previous study of oriented frog muscle at $H_0 = 23,487$ G (8), the sample size was quite large (fitting a 12-mm OD tube). Considerable variations of proton linewidth and small changes in the peak center with the angle between the fiber axes and H_0 (θ) were observed. In the present work, samples of smaller sizes (fitting a 5-mm OD tube) were studied. The spectra differed somewhat from sample to sample, but the maximum linewidth $\Delta\nu$ (for $\theta = 0^\circ$) was generally smaller than that for the larger samples (8), and there was less variation in $\Delta\nu$ with respect to θ . The spectra of one sample are shown in Fig. 1. The variation in the position of the peak is 1.2 ppm. The maximum variation in the linewidth is 30 Hz. Changes in the center of spectrum and linewidth with orientation also observed in the proton spectra for a sample of liquid water which was placed in a container $\sim 5 \times 2 \times 7$ mm, constructed by shaping a 5-mm

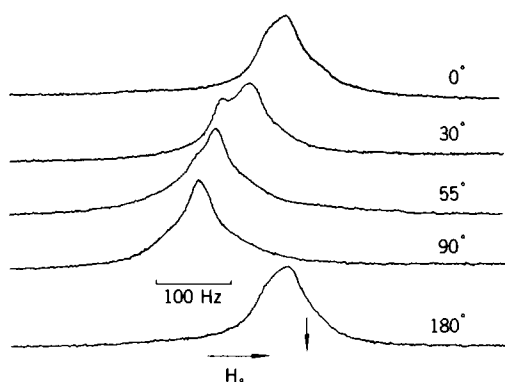


FIGURE 1 Proton NMR spectra of frog muscle oriented at different angles with respect to H_0 . The sample was placed in a 5-mm (OD) tube and the spectra were taken at 100 MHz and 25°C. The arrow indicates the peak position for liquid water.

NMR tube. The results of this experiment are given in Fig. 2, which shows the spectrum with the long side of the rectangular parallelopiped at angles 0°, 55°, and 90° to the external field. The variation in the position of the peak is 1.2 ppm. The maximum variation in the linewidth is 25 Hz.

To determine whether the line broadening of protons in muscle water is only due to dispersion of demagnetization, a study of the proton linewidth of muscle water was studied as a function of magnetic field. The results for two samples at four different fields are given in Table I.

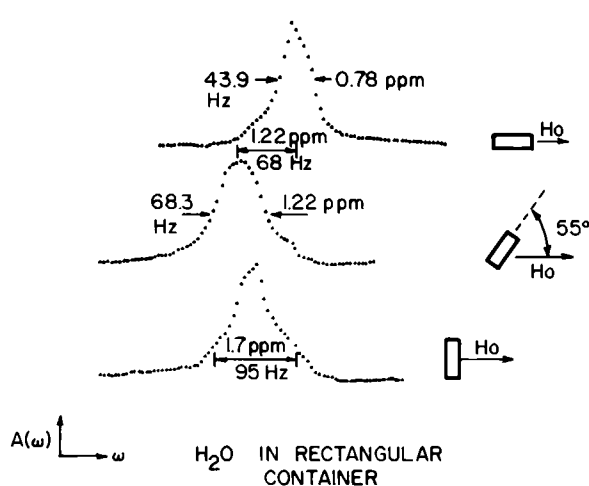


FIGURE 2

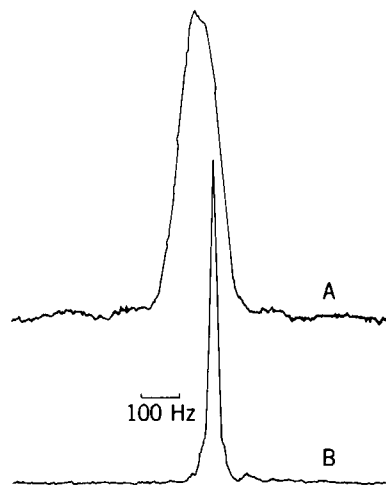


FIGURE 3

FIGURE 2 Line shape of a rectangular water sample as a function of orientation to static field at 56 MHz. Note change in linewidth due to dispersion of demagnetization field, and change in center of spectrum due to change in average demagnetization field.

FIGURE 3 Proton NMR spectra of frog muscle at 56 MHz and 25°C. (A) Without spinning. (B) With magic-angle spinning at a rate of 2 kHz. The linewidth was reduced only by a factor of 2 when the sample was spun at the same rate about an axis perpendicular to H_0 .

DISCUSSION

Spectra at High Field

To discuss the results of the proton linewidth study, it is necessary to realize that the effective magnetic field at a given proton in sample within an external field consists of a number of contributions:

$$\mathbf{H}_{\text{eff}} = \mathbf{H}_0 + \mathbf{H}_{\text{demag}} + \mathbf{H}_{\text{shield}} + \mathbf{H}_{\text{dipole}}. \quad (1)$$

\mathbf{H}_0 is the external field of the magnet. $\mathbf{H}_{\text{demag}}$ is the demagnetization field associated with the surface geometry of the sample. $\mathbf{H}_{\text{shield}}$ is the field associated with the currents induced in the local electronic environment by the external field \mathbf{H}_0 . $\mathbf{H}_{\text{dipole}}$ is the field due to the through space interaction with all other nuclear magnetic moments in the sample.

In terms of the demagnetization factor, n , the volume susceptibility tensor $\bar{\chi}_v$, the shielding tensor $\bar{\sigma}/s^{-1}$, the nuclear gyromagnetic ratio γ , and the internuclear vector \mathbf{r}_{ik} between the i th moment and the k th moment, the field at the i th nucleus is given by:

$$\mathbf{H}_{\text{eff}}^i = \mathbf{H}_0(\bar{I} - n\rho\bar{\chi}_v - \bar{\sigma}_i) - \sum_k [3r_{ik}(\boldsymbol{\mu}_k \cdot \mathbf{r}_{ik})/r_{ik}^5 - \boldsymbol{\mu}_k/r_{ik}^3]. \quad (2)$$

For a sample with appropriate geometry, e.g., an ellipsoid of revolution, n is exactly calculable. For a needle with \mathbf{H}_0 parallel to the needle axis, $n = 0$; for a needle with \mathbf{H}_0 perpendicular to the needle axis, $n = 2\pi$ (24). A sample with nontractable boundary conditions, e.g., a cube, will have a demagnetization field dispersion in the sample. This dispersion will have the effect of an external field inhomogeneity, and will result in a corresponding broadening of the NMR spectrum associated with dephasing of the transverse components of spin throughout the sample. For a particular sample geometry and orientation with respect to the external field, the dispersion of the demagnetization field within the sample may be calculated by, e.g., relaxation techniques applied to solutions of the appropriate Maxwell equations. Alternatively, in the absence of exact knowledge of the boundary conditions for the sample in question, one may choose to estimate this dispersion via an experiment on an appropriately chosen model system.

The example of the rectangular cell (Fig. 2) shows the effect of sample geometry on the spectrum. In general, the average value of the demagnetization field determines the center of the spectrum, and dispersion of the field determines the line shape and width. Muscle cells are elongated and highly anisotropic. By comparing the spectra in Figs. 1 and 2, we conclude that the main contribution to the linewidth and the change in the peak position for muscle water is the demagnetization field. The larger linewidth and apparent splitting previously observed (8) are most likely due to the larger size of samples used.

Variation of Linewidth with Magnetic Field

To determine whether there are other contributions to the proton linewidth of muscle water, we studied the linewidth at different magnetic field strengths (Table I). If the line broadening is due to the dispersion of demagnetization only, $\Delta\nu$ would be directly proportional to \mathbf{H}_0 for each sample. The data in Table I show that $\Delta\nu$ does decrease with the decrease of \mathbf{H}_0 , but the change in $\Delta\nu$ is somewhat slower. It is interesting to notice that, at low values of \mathbf{H}_0 , $\Delta\nu$ was

smallest for $\theta = 55^\circ$, at which $(3 \cos^2\theta - 1) \approx 0$. This is difficult to explain by the effect of dispersion in the demagnetization field, because the factor $(3 \cos^2\theta - 1)$ appears in the dipolar term but not in the demagnetization term of the spin Hamiltonian (Eq. 2). At this angle, the proton linewidth of muscle water approached that of a spherical sample of liquid water of comparable size when H_0 was lowered. The variation in $\Delta\nu$ from one sample to another was also much less at low values of H_0 . These facts suggest that, whereas the dispersion of demagnetization is the main factor of the increased proton linewidth and its variation with θ for muscle water, there may also be a small contribution to line broadening from nonzero static dipolar interaction. The data in Table I indicate that the broadening for $\theta = 0^\circ$ would be of the order of only a few hertz. This is much smaller than that suggested by the apparent splitting pattern shown by Fung (8). Packer has suggested (30) that there is a small residual quadrupole splitting of several hertz for a fraction of D_2O ($\sim 20\%$) in muscle. Since neither of these estimations is accurate, it is not clear whether the ratio $\Delta\nu_H/\Delta\nu_D$ for muscle water is comparable to that in other oriented heterogeneous systems (1, 2, 4, 10–12) or not.

Multiple Pulse Experiments

The possible existence of a small static dipolar interaction in muscle water is further demonstrated in the results of the eight-pulse experiment (Table II). If the line broadening of the proton signal of muscle water is due to dispersion in the demagnetization field only, the ratio $\Delta\nu$ for the signal obtained under REV-8 sequence and that for a single-pulse experiment would be $3/\sqrt{2} = 2.12$ or less. If the proton signal under a single pulse is further broadened by nonzero static dipolar interaction, which vanishes in the REV-8 sequence, this ratio would be larger than 2.12. The data in Table II show that this was indeed the case, indicating the existence of dipolar broadening in the single-pulse linewidth. It is to be noted that, for a perfectly aligned sample, there would be no dipolar broadening at $\theta = 54.7^\circ$ for both the single-pulse and the REV-8 results. The fact that the linewidth ratio is larger than 2.12 at this angle (Table II) is probably due to the lack of perfect alignment for the muscle sample. Considering that the uncertainties in the linewidths shown in Table II are $\sim 5\%$, the dipolar broadening for the single pulse data would be of the order of a few hertz, consistent with the result of the varied H_0 experiments.

Magic-Angle Spinning

The magic-angle spinning spectrum (Fig. 3) also demonstrated the large contribution of anisotropic magnetic susceptibility to the proton linewidth of muscle water at high fields. On the other hand, proton T_2 did not change substantially with magic-angle spinning at a rate of 2 kHz. This implies that the longest correlation time of the motion of part (or all) of water molecules is $< 8 \times 10^{-5}$ s, but tells nothing about other details of the molecular motion. In a previous study by one of us, it was suggested that nonzero static dipolar interaction in muscle water may have a large contribution to $1/T_2$ of protons (8). However, since we have demonstrated that the nonzeror dipolar interaction for muscle water is likely to be very small, this contribution would not be substantial. Other explanations of the large ratio of T_1/T_2 for proton in muscle water have been offered recently (31, 32).

In summary, we conclude that the proton spectra of muscle water in high magnetic fields are mainly determined by demagnetization field anisotropy and dispersion. The varied magnetic field and multiple-pulse experiments demonstrated that a small static dipolar

interaction of the order of a few hertz may be present and would have a small contribution to the proton linewidth.

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