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Anisotropy of transverse and longitudinal relaxations in liquids entrapped in nano- and micro-cavities of a plant stem

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Abstract

We studied the anisotropy of ¹H NMR spin–lattice and spin–spin relaxations in a fresh celery stem experimentally and modeled the sample theoretically as the water-containing nano- and micro-cavities. The angular dependence of the spin–lattice and the spin–spin relaxation times was obtained, which clearly shows the presence of water-filled nano- and micro-cavities in the celery stem, which have elongated shapes and are related to non-spherical vascular cells in the stem. To explain the experimental data, we applied the relaxation theory developed by us and used previously to interpret similar effects in liquids in nanocavities located in biological tissues such as cartilages and tendons. Good agreement between the experimental data and theoretical results was obtained by adjusting the fitting parameters. The obtained values of standard deviations (0.33 for the mean polar angle and 0.1 for the mean azimuthal angle) indicate a noticeable ordering of the water-filled nano- and micro-cavities in the celery stem. Our approach allows the use of the NMR technique to experimentally determine the order parameters of the microscopic vascular structures in plants.

Keywords

Spin–lattice relaxation; Spin-spin relaxation; Anisotropy; Nano- and microcavity; Celery stem

1. Introduction

Nuclear Magnetic Resonance (NMR) and Magnetic Resonance Imaging (MRI) have proven to be excellent tools for the study of the physical, chemical, and biological properties of matter, for diagnosing pathological tissue changes at early degradation stages and understanding organ malfunctioning due to diseases [1–6]. A wide range of biological samples has been studied by NMR, ranging from single cells to organs and tissues,

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Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

Ethical approval

All procedures performed in studies involving human participants were in accordance with the ethical standards of the institutional and/or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards.

including the study of plant metabolism impacts, a broad range of domains such as plant cultural practices, plant breeding, human or animal nutrition, phytochemistry and green biotechnologies [3,4,7]. In particular, NMR has the capability to determine the chemical composition, micro- and nano-structures and ionic transport in biological samples non-invasively and non-destructively [4,8–12].

Among the various NMR methods, a special place is occupied by the NMR relaxation technique [3,4], which is very useful for studying microstructure, internal spin interactions and the motion of molecules in solids, liquids and soft matter [1–4]. In particular, the anisotropy (angular dependence) of relaxation times of a ^1H NMR signal has been found in water-filled tissues that contains organized fibers; and whose anisotropy correlates with microscopic anatomy of the organs (their fibril structure), that can be characterized by water molecules filled between or inside fibrils of 280–300 nm in length and several tens nanometers in diameter [4]. Such anisotropy is not observed in non-fibrillar tissues. In several publications the anisotropy was explained by orientation of water molecules along the tissue fibrils [4,13–16]. Another explanation of the anisotropy is based on representation of a sample by a set of micro- and nano-cavities containing water molecules and averaging of the restricted molecular motion in such cavities [17,18]. It was shown that the relaxation times of the water contained inside the nano-cavity depend on the orientation of the cavity relative to the external magnetic field. The angular dependences of different relaxation times are different [17].

The most common relaxation mechanism is the magnetic dipole–dipole (DD) interaction between the magnetic moments of nuclei (spins). This interaction depends on the distance between nuclear spins (for example, a pair of protons in a water molecule) and on the orientation of the proton-proton vectors relative to the external magnetic field. The perturbed spin system relaxes to an equilibrium state and is characterized by two relaxation times: the spin–lattice (longitudinal relaxation) time T_1 and the spin–spin (transverse) relaxation time T_2 [1,2]. The study of the spin–spin and spin–lattice relaxations requires solving the Liouville - von Neumann equation for the spin system and considering all interactions of spins with each other and their environment. This is a very difficult task because the spin–spin and spin–lattice interactions can be time dependent. One of the effective methods for analytically describing the evolution of spin systems is to average the Hamiltonian over time; as a result of that, the systems are described by time-independent Hamiltonians [19–21]. The averaged method, based on the averaging of only a Hamiltonian *spin part*, has been successfully applied to explain the results of high-resolution NMR experiments with solids. One of the main results of the application of the averaged Hamiltonian theory was the explanation of the observed narrowing of the line in solid-state NMR experiments [19–21].

Another method, based on averaging a Hamiltonian *space part* over lattice variables [22,23] has been successfully used to explain the anisotropy of the transverse and longitudinal relaxation times in liquids entrapped in a nano-cavities [17,18,24–30]. Restriction of the Brownian motion of molecules in liquid or gas, enclosed in nano-cavities [17,18,24–30], results in the averaged Hamiltonian which describes the averaged DD interactions and can be characterized by a single averaged coupling constant. This simplification of a lattice part of the Hamiltonian by averaging over the Brownian chaotic motion has led to

analytically solvable models with non-trivial spin dynamics, which allows one to obtain the analytical expressions for the transverse and longitudinal relaxation times and the relaxation time under spin-locking [31,32]. This approach was applied to explain anisotropy obtained in the experiments with biological tissues, such as collagen fibril tissues (tendons, cartilages, nerves) [17,18,24–30]. The transverse and longitudinal relaxation times in liquid entrapped in a nano-cavity depend on the structural and dynamic parameters of samples and demonstrate noticeable anisotropy [4,17,18,33]. However, in these works various relaxations were studied in different samples; for each sample good agreement between the experimental data and the theoretical results were achieved by adjustment of several fitting parameters, which cannot be theoretically predicted. The fitting parameters can be considered as characteristics of nanocavities and fibril structure of a sample.

In the present work, we investigate both experimentally and theoretically the spin–lattice and spin–spin relaxation of the spin system in the *same* sample containing a liquid-trapped in a micro-/nano-cavity - a celery stalk. We measure the spin–lattice relaxation times T_1 and the spin–spin relaxation times T_2 of protons of water molecules. These relaxation times are of particular importance since they have been shown to be sensitive to the tissue structure and responsible for the MRI visualization in samples [4].

2. Theory

2.1. Space averaged spin Hamiltonian

In a nano-cavity, in contrast to a bulk material, space where molecules of gas or liquid diffuse is restricted; but the molecules can still move randomly throughout the whole cavity with the characteristic time t_{dif} . When the characteristic time t_{dif} is much shorter than the NMR time scale t_{NMR} , $t_{dif} \ll t_{NMR}$, the spin evolution in a single nanocavity can be described by an averaged of the *space part* of the dipole–dipole interaction (DD) Hamiltonian [23,24]. The time-averaged of the secular part of the DD interaction Hamiltonian, $H_d^{(s)}$ can be presented in the following form for an axis-symmetrical nano-cavity [23,24]:

$$\bar{H}_d^{(s)} = D^{(0)} \left(3I_z^2 - \vec{I}^2 \right) \quad (1)$$

where $\vec{I}^2 = I_x^2 + I_y^2 + I_z^2$ is the square of the total nuclear spin-1/2 operator, $I_v = \sum_{\mu=1}^N I_{v\mu}$ are the operators of the projections of the total spin operator on the v - axes ($v = x, y, z$), N is the number of spins in a nano-cavity. $D^{(0)}$ is the space-averaged pair coupling constant

$$D^{(0)}(\varphi) = -\frac{\gamma^2 \hbar}{V} P_2(\cos \varphi) F, \quad (2)$$

where γ is the nuclear gyromagnetic ratio, V is the cavity volume, $P_2(\cos \varphi) = \frac{1}{2}(3\cos^2 \varphi - 1)$, φ is the angle between the main axis Z_C of the nano-cavity and the external magnetic field,

\vec{H}_0 (Fig. 1), and F is the form factor, which varies for an ellipsoidal nanocavity from $-\frac{4}{3}\pi$ to $\frac{2}{3}\pi$ and a cylindrical nano-cavity from 0 to 2π .

The non-secular part of the DD interaction Hamiltonian, $H_d^{(ns)}$ after time-averaged takes the form [34]

$$\overline{H}_d^{(ns)} = \gamma^2 \hbar \sum_{q=-2, q \neq 0}^2 D^{(q)} S^{(q)}, \quad (3)$$

where $D^{(q)}$ are the averaged dipolar coupling constants:

$$\begin{aligned} D^{(1)} &= \left(-\frac{3}{8V}\right) F \sin(2\varphi) e^{-i\psi}, D^{(2)} \\ &= \left(-\frac{3}{8V}\right) F \sin^2(\varphi) e^{-2i\psi}, D^{(-q)} = D^{(q)*}, \end{aligned} \quad (4)$$

$S^{(q)}$ are operators acting only on the spin variables:

$$S^{(1)} = I_z I_+ + I_+ I_z, S^{(2)} = I_+ I_+, S^{(-q)} = S^{(q)†}, \quad (5)$$

$I_{\pm} = \sum_{\mu=1}^N I_{\pm\mu}$ are the total raising and lowering operators, ψ is the azimuthal angle of the main Z_C -axis of the nano-cavity.

The averaged non-secular part $\overline{H}_d^{(ns)}$ of the Hamiltonian contains four effective constants for each pair of spins. Similar to the effective constant of the secular part these constants depend on the volume, form, and orientation of the nano-cavity.

2.2. Transverse relaxation time

The transverse relaxation time T_2 can be calculated using its relationship with the local dipolar field $T_2 \approx \omega_{loc}^{-1}$ [35]. Using the expression for the local dipole field calculated in the case of a liquid entrapped in a nano-cavity [19,20], the transverse relaxation rate can be obtained

$$T_2(\varphi) = \left[\frac{\gamma^2 \hbar}{4} \left| (3\cos^2\varphi - 1) F \sqrt{\frac{3C}{V}} \right| \right]^{-1} \quad (6)$$

where C is the spin density of the liquid. A remarkable result is that anisotropy of the transverse relaxation time is determined by only angle φ between the magnetic field direction and the main Z_C -axis of the nanocavity. Fig. 2a shows the angle dependence of the normalized relaxation time $\frac{T_2(\varphi)}{T_2(0)}$ calculated using Eq. (6) for a single nano-cavity or for a set of the perfectly ordered nano-cavities. One can see from Fig. 2a that the normalized transverse relaxation time has a period equaled to π and reaches the local minimum of about $2T_2(0)$ at $\varphi = \pi/2$. At the magic angles, $\varphi = 0.96$ and 2.19 the relaxation time tends to infinity.

Estimation for a nanocavity with characteristic volume $V \approx 10^6 \text{ nm}^3$, the spin density for water $C = 66 \frac{\text{spin}}{\text{nm}^3}$, $|(3\cos^2\varphi - 1)F| = 2.3$, and for a proton $\gamma^2\hbar = 2\pi \times 120\text{Hz} \cdot \text{nm}^3$ gives the spin–spin relaxation time $T_2 \approx 170 \text{ ms}$, which agrees with the experimental data (Fig. 3a).

2.3. Longitudinal relaxation time

Method for the classical description of a lattice is well adapted to study of relaxation in liquids, where the Brownian motion of molecules is responsible for random variation in the lattice parameters [1,2]. The lattice parameters included in the non-secular part of the DD interaction Hamiltonian are then given by functions of time.

However, in the case of a liquid enclosed in a nano-cavity, the Hamiltonian after averaging does not contain terms describing random Brownian motion and is independent of time. The averaged Hamiltonian contains parameters characterizing a nano-cavity (volume V , form factor F), its orientation relative to the magnetic field, and liquid in the nano-cavity. Random oscillations of these parameters cause spin–lattice relaxation in a liquid enclosed in a nano-cavity. Assume that the reduced correlation functions $g^{(1)}(\tau)$ and $g^{(2)}(\tau)$ are represented by [34]

$$g^{(q)}(\tau) = \frac{|\tau|}{\tau_c} e^{-\frac{|\tau|}{\tau_c}}, q = 1, 2, \quad (7)$$

where τ_c is the correlation time of random fluctuation of a nano-cavity wall and using the standard procedure (see e.g. [1]) we obtain the spin–lattice relaxation rate:

$$T_1 = \left\{ 3 \left(\frac{3}{8} \right)^3 \frac{C}{V} (\gamma^2\hbar)^2 \tau_c B(\varphi) \right\}^{-1} \quad (8)$$

where

$$B(\varphi) = F^2 \left[\frac{(\omega_0^2 \tau_c^2 - 1)}{(\omega_0^2 \tau_c^2 + 1)} \sin^2 2\varphi + \frac{(4\omega_0^2 \tau_c^2 - 1)}{(4\omega_0^2 \tau_c^2 + 1)} \sin^4 \varphi \right] \quad (9)$$

Eq. (8) gives the dependence of the spin–lattice relaxation time on the Larmor frequency ω_0 , volume V , density of nuclear spins C , shape (form factor F) and nano-cavity orientation relative to the magnetic field given by the angle φ . Estimation of the spin–lattice relaxation time T_1 at the field strength of 8.0 T according to Eq. (8) with $C = 66 \frac{\text{spin}}{\text{nm}^3}$, $B \approx 10$,

characteristic size $V \approx 10^6 \text{ nm}^3$ and $\tau_c \approx 10^{-2} \text{ s}$ gives $T_1 \approx 1.7 \text{ s}$ which agrees with the experimental data (Fig. 3b). Fig. 2b shows the angle dependence of normalized relaxation time $\frac{T_1(\varphi)}{T_1(0.96)}$ calculated using Eq. (8) for a single cavity or for a set of the perfectly ordered nanocavities.

3. Experimental

3.1. Sample

The stem of celery consists of four main types of structural components: (i) the vascular bundles that contain water-transporting xylems and (ii) nutrient-transporting phloems, (iii) collenchyma which provides support for the plant and (iv) parenchyma (ground tissue). Xylem and phloem vessels are created by overlapping cells, which form hollow cavities interconnected into one long micrometer-diameter tube. Collenchyma is made up of elongated living cells filled with water and is located along the outer edges of the stem and the vascular bundle tissue, being in a longitudinal orientation. The cell walls possess a fibril structure and consist of nano-fibers with their diameters in the range of 6–25 nm [36]. Parenchyma is made up of non-oriented living cells filled by water and cellular liquid, which are bigger in diameter and often shorter when comparing with collenchyma [37]. Water is also a part of the fibril structure and is located between the microfibrils, presumably serving to stabilize them through hydrogen bonding.

Celery studied was purchased from a local supermarket. For the NMR measurements, we cut out from the outer (lateral) part of the celery stalk a small piece of ~ 1 cm long and ~ 2 mm wide, which mainly consists of collenchyma, xylem and phloem, avoiding the parenchyma as much as possible. Thus, we worked in the most part with oriented fibril tissues. The specimen did not contain the skin (epidermis). We note that celery is highly hydrated, having the water content up to 95% w.t., thus the NMR signal comes mainly from water molecules.

3.2. NMR measurements

We measured the spin–lattice (T_1) and spin–spin (T_2) relaxation times of water molecules in the celery stem at different angles between the sample’s long axis and the direction of the applied magnetic field. The ^1H NMR measurements were carried out at ambient temperature (293 ± 0.5 K) using a Tecmag pulse solid state NMR spectrometer and an Oxford superconducting magnet with the magnetic field $B_0 = 8.0$ T, corresponding to the ^1H resonance frequency $\omega_0 = 340.54$ MHz, and a home-built NMR probe. The spin–spin relaxation times T_2 were measured using the Carr-Purcell-Meiboom-Gill (CPMG) pulse sequence [38]; in addition (for comparison), the Hahn echo measurements were also carried out. The duration of the $\pi/2$ RF pulse in the sequences was 1.6 μs . To determine the relaxation time T_2 we processed the decays of echo amplitudes as $M(t) \sim \exp(-2t/T_2)$ and $M(2n\tau) \sim \exp(-2n\tau/T_2)$ for the Hahn echo and CPMG method, respectively (here t is the pulse spacing and τ is the echo spacing, respectively). The CPMG pulse train consisted of 2048 pulses separated by a time interval of 2 msec. Spin-lattice relaxation time T_1 was measured using the inversion recovery pulse sequence [1]. Magnetization recovery in the T_1 measurements of a sample was well described by a stretched exponential function $M(t) = M_\infty \left\{ 1 - \exp\left(-\frac{t}{T_1}\right)^\alpha \right\}$, where M_∞ is the equilibrium magnetization and α is the stretched exponential parameter. Such approach is usually used in the case of a distribution of the relaxation times in a system with a number of different relaxation environments, resulting in a multi-component relaxation behavior with the impossibility of separating into

individual components. Repetition time was chosen as $5 T_1$. To get an intense signal, 16 acquisitions were usually required.

All measurements were carried out with non-spinning sample to preserve the contributions of the proton DD interactions to the spin–lattice and spin–spin relaxations, which are studied in the present article. To obtain the orientation dependences of the relaxation times, the sample was placed at different angles between the stem axis and the direction of the applied magnetic field with an accuracy of $\sim 1.5^\circ$.

4. Experimental results and discussion

The relaxation times of bulk distilled water are $T_1 = 3.8$ s and $T_2 = 2.4$ s [39] and are known to be independent on the sample orientation relative to the applied magnetic field. Herewith our measurements of the celery stem demonstrate noticeable dependences of both the transverse and longitudinal relaxations of the water protons on the angle θ between the stem axis and the direction of the external magnetic field (Fig. 3a and b). According to the above theory, such anisotropy of relaxation times should be observed in a liquid located in a restricted space/volume formed by nano- and micro-cavities in a material [17,18]. In our case, the observed anisotropy of relaxation times in celery should be associated with nano- and micro-sized water-filled cavities in the studied tissues. The stretched exponential parameter when fitting of the T_1 data was found to be around 0.9.

We note that the CPMG measurements show longer spin–spin relaxation times compared with the Hahn echo measurements, since the latter are affected by diffusion of the water molecules [26] and will not be considered further. The CPMG curves are well fit by a single exponential, corresponding to the water-filled elongated nano/micro-cavities in the tissues relaxing with short T_2 . As it was mentioned above, our sample is almost free of the parenchyma tissue with non-oriented water-filled cavities. Even if such tissue is present in the sample, it represents only a small fraction of water and, therefore, corresponds to a minor component of the relaxation signal, relaxing with long T_2 and located at the tail of the CPMG signal. Therefore, its contribution can be practically neglected. Thus, the observed angular dependence of the relaxation times (Fig. 3) is associated with the elongated micro- and nano-sized water-filled cavities.

Comparison of the experimental results shown in Fig. 3 with the calculations presented for a set of perfectly ordered cavities (Fig. 2) shows a significant difference. The reason is that the above-mentioned calculations were done for a single cavity or for a number of cavities perfectly oriented in the same direction. However, the celery stem contains a set of the cavities which are somewhat differently oriented and have different shapes and volumes. To explain the difference between the angular dependences of the relaxation times T_2 and T_1 given by Eqs. (6) and (8) and the experimentally observed dependences, we consider a plant tissue as a set of imperfectly orientated water-containing cavities. For further analysis, it is convenient to use expressions for the normalized relaxation rates

$$R_k(\theta) = \frac{\max_{\theta}(T_k(\theta))}{T_k(\theta)} \quad (10)$$

where $R_k(\theta)$ is the normalized relaxation rates for a single nanocavity, $k = 1$ for the spin–lattice relaxation time T_1 and $k = 2$ for the spin–spin relaxation time T_2 , respectively. In Eq. (10) we assume that the spin–lattice relaxation is described by an exponential function of time with ($\alpha = 1$). Calculations according to Eq. (8) with $\alpha = 1$ and with $\alpha = 0.9$ give almost the same result.

Volume and form factor are various for different nanocavities. Assuming that the angular distributions of all cavity types are the same, we obtain the normalized relaxation rate averaged over whole sample [34]

$$\langle R_k(\theta) \rangle = \frac{\langle \max_{\theta}(T_k(\theta)) \rangle}{\langle T_k(\theta) \rangle} \quad (11)$$

In the NMR experiments, the relaxation times were measured as functions of the angle θ between the sample Z_0 -axis and magnetic field. Herewith the main axes of the cavities are distributed about some “preferred direction”, an averaged orientation of the elongated cavities. Let us determine the deviation of the main axis of each cavity from the Z_0 -direction by the polar angle ζ and azimuthal angle ξ . Suggesting the Gaussian distribution of the elongated cavities directions over both the angles, the polar ζ and azimuthal ξ angles with the bivariate normal distribution function

$$\langle R_k(\theta) \rangle = \frac{1}{\Gamma} \int_0^{2\pi} d\xi \int_0^{\pi} d\zeta \sin(\zeta) \Psi_k(\zeta, \xi, \rho) \frac{1}{T_k(\theta, \zeta, \xi)}, \quad (12)$$

where $\Psi_k(\zeta, \xi, \rho)$ is the probability density function

$$\Psi_k(\zeta, \xi, \rho) = \frac{1}{2\pi\sigma_{k\zeta}\sigma_{k\xi}\sqrt{1-\rho^2}} \exp\left\{-\frac{1}{2\sqrt{1-\rho^2}}\left[\frac{(\zeta-\zeta_{0k})^2}{\sigma_{k\zeta}^2} - 2\rho\frac{(\zeta-\zeta_{0k})(\xi-\xi_{0k})}{\sigma_{k\zeta}\sigma_{k\xi}} + \frac{(\xi-\xi_{0k})^2}{\sigma_{k\xi}^2}\right]\right\}, \quad (13)$$

ρ is the correlation parameter of the polar ζ and azimuthal ξ angles, $\sigma_{k\zeta}$ and $\sigma_{k\xi}$ are the standard deviations, ζ_{0k} and ξ_{0k} are the means of the distributions of the angles and give distinction of the “preferred direction” of the elongated cavities from the sample axis,

$$\Gamma = \max_{\theta} \int_0^{2\pi} d\xi \int_0^{\pi} d\zeta \sin(\zeta) \Psi(\zeta, \xi, \rho) \frac{1}{T_k(\theta, \zeta, \xi)}. \quad (14)$$

The angular dependences of the normalized relaxation rate $\langle R_k(\theta) \rangle$ are given by Eq. (12) in which there are unknown parameters: the correlation parameter (ρ), standard deviations ($\sigma_{k\zeta}$ and $\sigma_{k\xi}$) and distribution means (ζ_{0k} and ξ_{0k}). These parameters can be considered as parameters which characterize the fine structure of celery.

Our numerical experiments showed that varying the correlation parameter ρ from 0 to 1 has practically no effect on the fitting parameters, which indicates that there is no correlation between the polar and azimuthal angles.

The best matching of the calculations for two data sets is achieved simultaneously for both relaxation rates at the standard deviations of $\sigma_{k\zeta} = 0.33$ and $\sigma_{k\xi} = 0.1$ for the polar and azimuthal angles, respectively (here $k = 1, 2$). The averaged transverse relaxation $\langle R_2(\theta) \rangle$ and longitudinal relaxation $\langle R_1(\theta) \rangle$ rates calculated according to Eq. (12) together with the experimental data are presented in Fig. 4. Since the standard deviation describes the distribution of nano-cavities within the sample and measurements of the relaxation rates were carried out in the same sample, this result is expected.

We consider the agreement between the theoretical values and the experimental data to be quite satisfactory. Some discrepancies between the calculations and experimental data must come from the heterogeneities of the specimen containing several types of tissues (collenchyma, xylem, phloem). We found that for the mean values of the distributions, the best agreement of calculations with experimental data is achieved with the following values: $\zeta_{01} = 0.43$ and $\zeta_{02} = 2.3$ and $\xi_{01} = \xi_{02} = 0.1$. Note that the relaxation rates in a single nanocavity possess a period of π relatively to the angle φ (Fig. 3). For the sample we observed 3D space symmetry of the relaxation rates and change of the averaged angle by π : $\tilde{\zeta}_{0k} \rightarrow \pi - \zeta_{0k}$ and $\tilde{\xi}_{0k} \rightarrow \pi - \xi_{0k}$ that gives the same dependence of the relaxation rates on angle θ . For example, using the averaged values: $\zeta_{01} = 0.43$ and $\zeta_{02} = 0.814$ and $\xi_{01} = 0.1$ and $\xi_{02} = 3.04$ we obtained the relaxation rate angle dependences presented in Fig. 4. The standard deviations and averaged angles can be considered as parameters which characterize the fine structure of tissue components of celery.

5. Conclusion

We have studied both experimentally and theoretically the proton spin–spin and spin–lattice relaxations of water molecules entrapped in the celery stem. The angular dependence of both relaxation times was obtained. This behavior was attributed to the elongated water-filled nano-/micro-cavities. The obtained findings were interpreted in the frameworks of the theory, previously developed by us [17,18,26,32], which was successfully applied to explain similar effects in liquids located in nanocavities of biological tissues. We averaged the normalized relaxation rates over orientations of the cavities relative to the applied magnetic field. Good agreement between the calculations and experimental data was obtained by adjustment of a few fitting parameters - the standard deviations and averaged cavities direction. The approach allows using NMR data to non-invasively and non-destructively determine the order parameters of plant tissues.

An important feature of this study is that both transverse and longitudinal relaxation times were measured with the same sample and under the same conditions. The results obtained are well described by theoretical expressions with the same parameters characterizing the ordering of the micro-/nano-cavities in the sample. The obtained values of the standard deviations ($\sigma_{\zeta} = 0.33$ and $\sigma_{\xi} = 0.1$) indicate a noticeable ordering of the cavities by both the transverse and the longitudinal relaxation times .

Our straightforward approach (measuring different relaxation times) showed that NMR methods can be used to study the microscopic vascular structure of plants. Variations in these vascular structures in plants can be associated with the economic characteristics of

plants, for example, high/low yields of food crops or long/short growth times of vegetables [40,41]. Since NMR and MRI methods are totally non-invasive and non-destructive, we can conclude that the considered experimental NMR approach [29,32] can be used to study the fine vascular structures of *living* organisms, as well as to determine deviations of the fine anatomical structure of tissues from the norm in *living* organisms.

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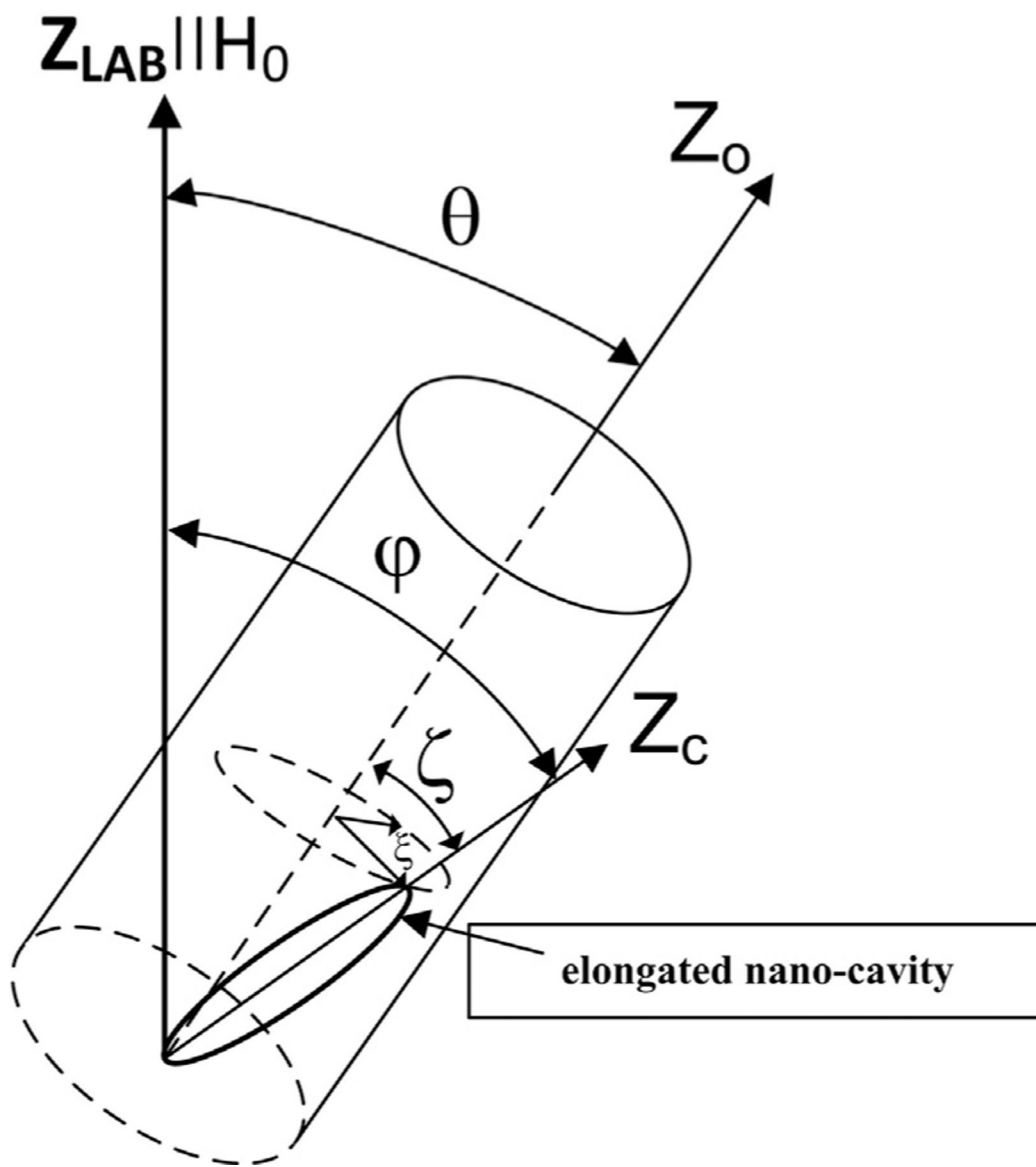


Fig. 1. The coordinate systems and angles used to describe the model of plant tissue. Z_{LAB} denotes the z-axis of the laboratory frame. θ is the angle between the external magnetic field \vec{H}_0 and the Z_0 -axis chosen as an axis of the sample in the experiment. ζ and ξ are the polar and azimuthal angles characterizing the deviation of the main Z_C -axis of an elongated nano-cavity from the Z_0 -axis of the sample.

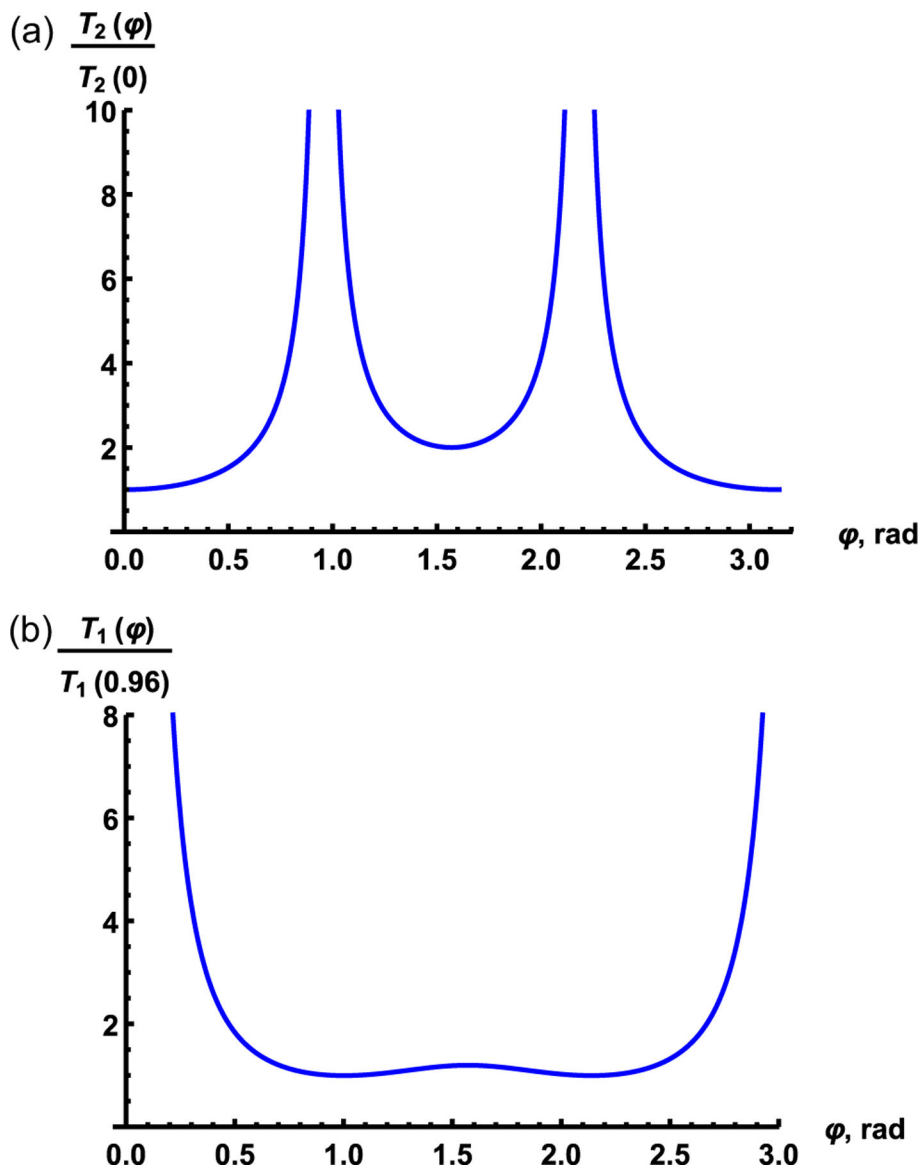


Fig. 2. Angular dependence of the normalized transverse $\frac{T_2(\varphi)}{T_2(0)}$ (a) and longitudinal $\frac{T_1(\varphi)}{T_1(0.96)}$ (b) relaxation times in a single nano-cavity, calculated using Eq. (6) and Eq. (8), respectively.

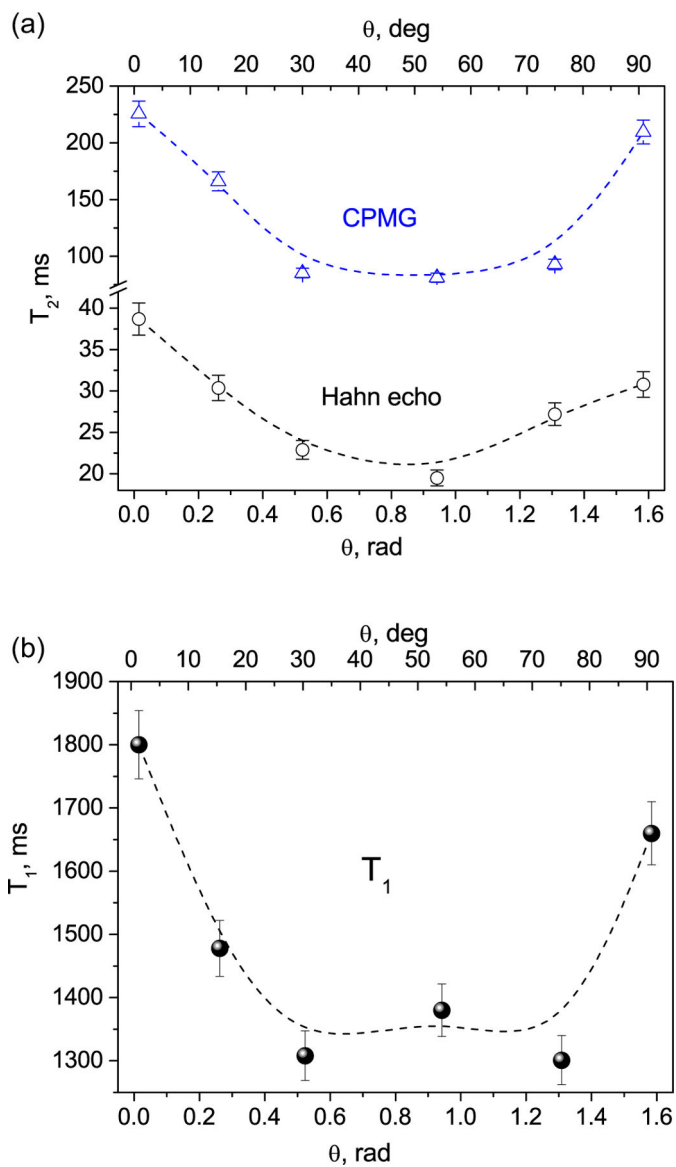


Fig. 3. Experimental dependence of the transverse relaxation T_2 (a) and longitudinal relaxation T_1 (b) times of water molecules in the celery stem on the angle θ between the sample axis and direction of the applied magnetic field. Dash lines are B-Spline fits. Error bars include both experimental and simulation errors.

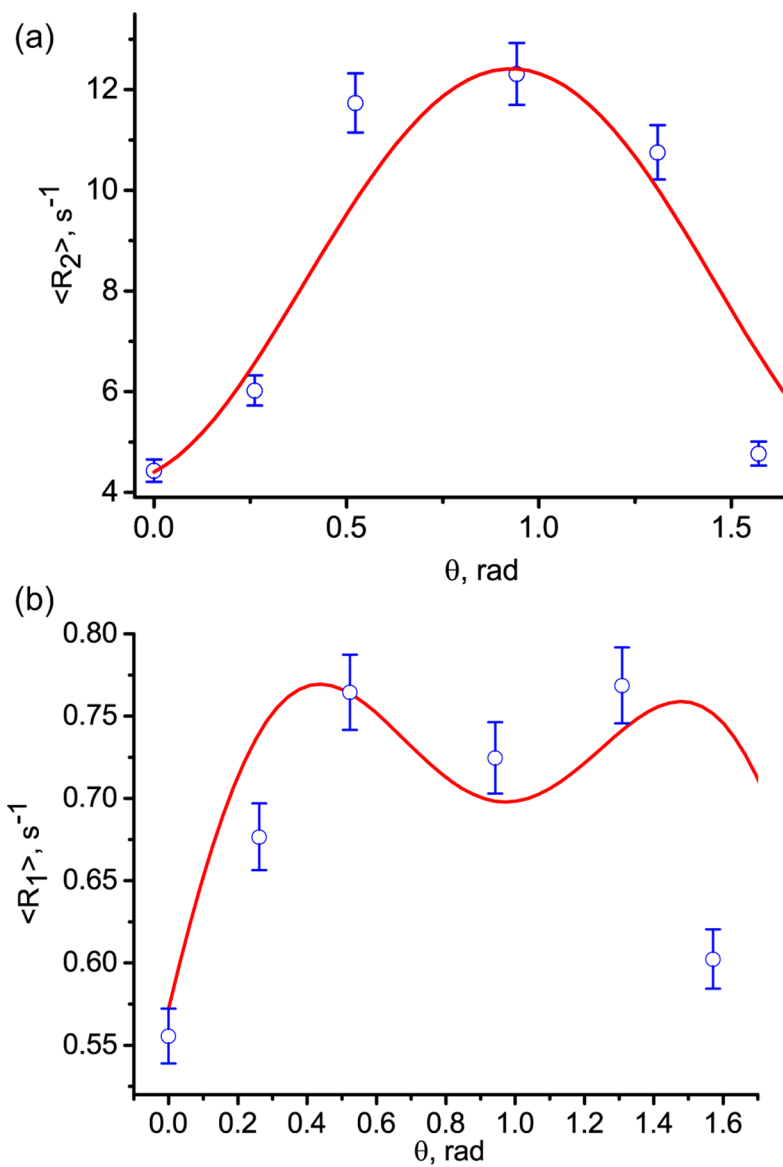


Fig. 4. Angular dependence of the transverse $\langle R_2(\theta) \rangle$ (a) and longitudinal $\langle R_1(\theta) \rangle$ (b) relaxation rates in the celery stem. The blue circles are the experimental rate $\langle R_2(\theta) \rangle$ measured with the CPMG pulse sequence and $\langle R_1(\theta) \rangle$ measured with inversion recovery pulse sequence. The solid red curves are according to Eq. (12) for the standard deviations of 0.33 and 0.1 for the polar and azimuthal angles, respectively, for both relaxation rates.