Hydration of Potato Starch in Aqueous Suspensions Determined from Nuclear Magnetic Relaxation Studies by ¹⁷O, ²H, and ¹H NMR: Relaxation Mechanisms and Quantitative Analysis

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Nuclear magnetic relaxation (¹⁷O, ²H, and ¹H NMR) measurements were employed to investigate the hydration properties of potato starch in relation to functionality in food systems. A comparison of ¹⁷O, ²H, and ¹H NMR data allowed us to resolve the contributions to water relaxation made by water binding, chemical exchange, and intermolecular dipolar interaction cross-relaxation, respectively. Weakly bound water in potato starch suspensions has an average correlation time of about 20 \pm 0.6 ps (determined by ¹⁷O, NMR), compared with 5 ps for bulk water (D₂O) at 20 °C. Three hydration regions were observed between 0.02 and 0.90 g of starch/g of D₂O, respectively. The first hydration region extends from 0.02 to 0.4 g of starch/g of H₂O, whereas the second region occurred between 0.40 and 0.65 g of starch/g H₂O. Deuterium and proton NMR allowed us to resolve three water populations "bulk (or free)" water, "weakly bound" water, and trapped water in potato starch suspensions. Deuterium NMR resolved a slowly exchangeable anisotropically bound water population within the potato starch granule structure. We find, therefore, that potato starch has a potential for utilization as a stable functional ingredient in food formulations to increase stability and storage life.

INTRODUCTION

Potato starch has the potential for wide utilization as a stable functional ingredient in food formulations to increase the stability and storage/shelf life of foods (Baianu et al., 1990a). Previous deuterium nuclear magnetic resonance (NMR) studies of potato starch hydration properties showed that these are unique since they were not found in corn or wheat starches (Lechert et al., 1980; Lechert, 1981; Baianu et al., 1990b; Yakubu et al., 1990a,b). Potato starch contains about $34 \pm 1\%$ water (Schierbaum and Taufel, 1962), which we found to be trapped, or anisotropically bound, within the potato starch granule structure (Baianu et al., 1990a; Yakubu et al., 1990b,c). The X-ray diffraction studies of starches, on the other hand, classified fully hydrated potato starch as B-starch (Zobel, 1988a). It has also been recently suggested that the hydration properties of potato starch and potatoes may be due to the presence of B-type structure (Reuther et al., 1984; Zobel, 1988b), that is, the hexagonal arrangement of double helices of very large amylose molecules (Yakubu et al., 1990b). The ability of potato starch to bind, or trap, water within its granule structure makes potato starch more desirable than other (cereal) starches as a functional ingredient for improving and increasing the rigidity and firmness of gels in a number of food systems (Baianu, 1986; Lanier, 1986; Lee, 1986; Baianu et al., 1990a).

The role of water in influencing the physicochemical, hydration, and functional properties of carbohydrates and proteins has been quantitated by NMR techniques (Tait

* Address correspondence to this author at Department of Food Science, Physical Chemistry and NMR Laboratories, University of Illinois at Urbana-Champaign, 580 Bevier Hall, 905 S. Goodwin Ave., Urbana, IL 61801 [telephone (217) 244-6630, 333-4443]. et al., 1972; Lèlievre and Mitchell, 1975; Halle and Lindman, 1978; Baianu, 1982, 1985, 1989, 1992; Baianu and Förster, 1980; Kakalis and Baianu, 1988; Mora-Gutierrez and Baianu, 1989, 1991). The NMR techniques, being nondestructive, are widely utilized to monitor the molecular dynamics of water in foods, water binding, and diffusion rates (Baianu et al., 1992). Such information is useful for the prediction of stability and storage life of foods (Baianu, 1986; Baianu et al., 1990a,b).

Oxygen-17, deuterium, and proton NMR measurements were recently employed to investigate the hydration properties of starches (Baianu et al., 1990a) and proteins (Lioutas et al., 1986; Kakalis and Baianu, 1988; Baianu, 1989). Such studies are here extended to the hydration of potato starch suspensions.

MATERIALS AND METHODS

Potato Starch. Potato starch was obtained from Sigma Chemical Co. (St. Louis, MO) and was specified to contain 25% amylose and 75% amylopectin.

Moisture and pH. Moisture content was determined according to the AOAC (1981) method and was found to be about 14%. The pH values of potato starch suspensions in H_2O and D_2O were determined with a laboratory pH meter (Hana Instruments, H18417). For pD = pH + 0.45 (Covington et al., 1968), most values were close to pD = 7.45 (pH 7.0).

Sample Preparation. Potato starch suspensions were prepared by mixing appropriate amounts of solids with deuterium oxide for ¹⁷O and ²H NMR or deionized water for ¹H NMR. Samples were transferred to 5- or 10-mm o.d. NMR tubes and were allowed to equilibrate on ice.

¹H NMR Measurements. ¹H NMR measurements were carried out at 10 MHz with a PC-10 process analyzer (Bruker/IBM Instruments, Danbury, CT). The Carr-Purcell-Meiboom-Gill (CPMG) sequence was employed for T_2 measurements. The decay of the transverse magnetization was monitored with a dualbeam Tektronix storage oscilloscope (Model 5113). Measurements were carried out in triplicate at 20 ± 1 °C.



Figure 1. Fourier transform NMR spectra of D_2O in potato starch suspensions at 20 ± 1 °C. (A) ²H NMR spectrum of D_2O in potato starch suspensions; (B) ¹⁷O NMR spectrum of D_2O in potato starch suspensions.

Oxygen-17 and Deuterium NMR Measurements. A CXP 200, high-field, multinuclear NMR spectrometer (Bruker Instruments Inc., Billerica, MA), operating at 4.7 T, was used for both ¹⁷O and ²H NMR. Measurements were carried out at resonance frequencies of 27.12 and 30.71 MHz, respectively. For oxygen-17 NMR, spectral widths were selected in the range from 50 to 200 kHz, the 90° pulse width was 16 μ s, and the recycling time was 0.16 s. The number of scans was varied from 500 in dilute suspensions to 10 000 in hydrated powders. The deuterium NMR spectral widths were selected in the range from 40 kHz to 2 MHz, the 90° pulse width was 16 μ s, and recycling time was 0.16 s. The number of scans was varied from 500 in dilute suspensions to 10 000 in hydrated powders. A signal-to-noise ratio of 1000:1 was selected to observe the deuterium powder pattern of "tightly bound" D_2O within the potato starch granules. The Fourier transforms were carried out on-line with a Bruker Aspect 2000A (24 bit) computer equipped with a CDC 90-MB drive.

Oxygen-17 and ²H NMR Theory. The oxygen-17 nucleus has a spin quantum number I = 5/2. Therefore, it has five transitions between six spin energy levels. The ¹⁷O nuclear spin relaxation is dominated by the quadrupolar interaction (Abragam, 1961; Halle, 1981; Kintzinger, 1983). The strong quadrupolar interaction causes a large relaxation effect, permitting the study of hydration even for a small number of hydration sites (Baianu et al., 1992).

Both the reorientation of the water molecules at the binding sites on a biopolymer, or a particle, and the exchange between "bound" and "free" or "bulk" water cause the narrowing of the ¹⁷O NMR absorption peak of water. When the exchange between bound and free water is fast on the time scale of the NMR measurements (that is, when the water-exchange rate is faster than the bound water nuclear spin relaxation rate), the measured relaxation rate is the weighted average of the relaxation rates of bound (R_{2B}) and free (R_{2F}) water populations (Zimmerman and Brittin, 1957; Derbyshire, 1982; Mora-Gutierrez and Baianu, 1989). Such a relationship is expressed as

$$R_{2\rm obs} = P_{\rm B}R_{2\rm B} + P_{\rm F}R_{2\rm F} \tag{1}$$

where $P_{\rm B}$ and $P_{\rm F}$ are fractions of bound and free water in the system, respectively, and $R_{\rm 2obs}$ is the measured transverse relaxation rate of the samples at various hydration levels. A single Lorentzian ¹⁷O NMR peak was observed in potato search suspensions in D₂O from 0.02 to 0.6 g of starch/g of D₂O. Our observations are consistent with a fast-exchange, two-state model between bound and free water populations, as shown in Figure 1 (spectra A and B). The two-state model was previously employed to analyze ¹⁷O NMR data for water (D₂O) in dilute lysozyme solutions (Lioutas et al., 1986; Kakalis and Baianu, 1988), corn maltodextrins (Mora-Gutierrez and Baianu, 1990), and wheat starch (Baianu and Mora-Gutierrez, 1985; Baianu, 1986).

In the case of ${}^{2}H$ NMR studies of potato starch suspensions in D₂O, the analysis with the fast-exchange model applies only to the water populations responsible for the central Lorentzian peak. The relevant deuterium NMR theory for water (D_2O in potato starch suspensions and hydrated powders) was outlined in several recent papers (Baianu et al., 1990a, 1992; Yakubu et al., 1990a-c). Deuterium NMR quadrupole splittings of about 1 kHz were resolved in potato starch suspensions ranging from 12% starch (88% water) to hydrated powders (down to $\leq 2\%$ water).

In the case of water (D_2O) molecules that undergo anisotropic reorientation, there is a partial ordering of the water molecules (Mantsch et al., 1977; Smith, 1983). This is characterized by an order parameter, S, defined by the expression

$$\Delta \nu_{\Omega} = (3e^2 q Q/4\hbar)S \tag{2}$$

where $S_D = [(3 \cos^2 \theta - 1)/2]$ is the order parameter, eq is the electric field gradient at the deuterium nucleus, eQ is the deuterium quadrupole moment, θ is the angle made by the direction of the oxygen-deuterium bond with the applied magnetic field, H_o , and \hbar is Planck's constant devided by 2π . The order parameter, S_D , is equal to 1.0 for a perfectly ordered system, whereas it is zero for a completely disordered system. This property, S_D , makes deuterium NMR an excellent technique for the study of the degree of water ordering in foods and other biological systems (Baianu et al., 1984). The order parameter, S_O , in the case of the oxygen-17 NMR of water has twice the value of S_D measured for D_2O by deuterium NMR. This is mainly because of the different orientations of the oxygen-17 and deuterium quadrupole tensors with respect to the magnetic field (Abragam, 1961).

The transverse relaxation rates, R_2 (s⁻¹), for both oxygen-17 and deuterium NMR were calculated at each concentration with (Dwek, 1973; Mora-Gutierrez and Baianu, 1989)

$$T_2^{-1} = R_{2\text{Corr}} \,(\text{s}^{-1}) \propto \pi \Delta \nu_{\text{obs}}^{\text{Corr}} \tag{3}$$

where $\Delta \nu_{obs}^{Corr}$ is the line width at half-height for each spectrum, after correcting for a small inhomogeneity broadening of the order of 2% or smaller.

The ¹⁷O NMR relaxation data were fitted with a Systat nonlinear regression program on a 32-bit Macintosh II microcomputer equipped with a 68020 CPU, a 68882 mathematical coprocessor, 5-Mb RAM, and an 80-Mb hard drive. Both Simplex and quasi-Newton algorithms were employed.

For calculations of correlation times, the "excess", or differential transverse relaxation rates, ΔR_2 (s⁻¹), were employed in

$$\Delta R_2 (s^{-1}) = (\Delta \nu_{\rm obs}^{\rm Corr} - \Delta \nu_{\rm free})$$
⁽⁴⁾

where Δv_{free} is the line width at half-height for water (D₂O).

Nuclear Magnetic Relaxation Mechanisms of Water in Potato Starch Suspensions. Oxygen-17 nuclear spin relaxation is essentially quadrupolar, and in the case of "heavy" water (D_2O), it is not altered to any significant degree by either cross-relaxation (Kakalis and Baianu, 1988) or chemical exchange (Dwek, 1973; Kintzinger, 1983).

Deuterium nuclear spin relaxation mechanism is also quadrupolar in solutions, but the ²H NMR relaxation rate is affected by the chemical exchange between bound and free water protons. The ²H NMR is, therefore, quite useful for separating, or resolving, "liquidlike" from "solidlike" behaviors in biological systems where mixed phases are present (Baianu et al., 1984); other major uses of NMR, in general, and deuterium NMR in particular, are concerned with the conformational changes induced by heat treatments of biopolymers (Lechert et al., 1980; Baianu et al., 1982; Baianu, 1983, 1992; Mora-Gutierrez and Baianu, 1989; Yukubu et al., 1989).

The proton with a spin 1/2 has a large magnetic moment, and in the absence of paramagnetic species, its nuclear spin relaxation is dominated by *dipolar interactions* (Abragam, 1961). In protonated or partially deuterated biopolymers, the dipolarcoupled spin group behavior in solids becomes important (Baianu et al., 1981). Major concerns in ¹H NMR spin-lattice relaxation studies are the contributions of cross-relaxation and chemicalexchange processes to the relaxation of water protons. Crossrelaxation is essentially a spin diffusion process (Edges and Samulski, 1978) involving a magnetic energy transfer across a "solid" particle-liquid water interface. This process can only occur if the relaxing water molecules are closer than about 5 Å,



Figure 2. ¹⁷O NMR transverse relaxation rates, R_2^* (s⁻¹), of water in potato starch suspensions. (A) Dependence of the transverse relaxation rates, R_2^* (s⁻¹), on potato starch-to-water ratios from 0.02 to 0.4 g of starch/g of D₂O; (B) dependence of the transverse relaxation rates, R_2^* (s⁻¹), on potato starch-towater ratios from 0.4 to 0.6 g of starch/g of D₂O; (C) residual plots of ¹⁷O NMR transverse relaxation rates, R_2^* (s⁻¹), for potato starch-to-water ratios [(A) 0.02–0.4 g of starch/g of D₂O, SD = 17.31; (B) 0.4–0.6 g of starch/g of D₂O, SD = 26.13].

or less, to the particle surface. Because of the marked effect of the protons attached to the biopolymer on the rate of water proton cross-relaxation, the ¹H nucleus is useful for the investigation of aggregation and denaturation of proteins (Baianu, 1989; Myers-Betts and Baianu, 1990a,b) and gelatinization and retrogradation of starch (Baianu and Mora-Gutierrez, 1985; Baianu, 1986; Mora-Gutierrez and Baianu, 1989, 1990).

RESULTS

Oxygen-17 and deuterium NMR spectra of potato starch suspensions in D₂O are presented in Figure 1A,B. In both spectra, a single Lorentzian peak is observed for a highly mobile water population in fast exchange with bulk water, observed in the hydration range from 0.02 to ~ 0.5 g of potato starch/g of D₂O. This is consistent with the twostate, fast-exchange model between bound and free water populations (eq 1).

The dependence of the ¹⁷O NMR transverse relaxation rate, R_2^{+Corr} , potato starch suspensions on the starch-to-D₂O ratio is shown in Figure 2A; the ¹⁷O NMR line width increases linearly with the starch-to-water ratio for low to intermediate values, as predicted by eq 1. Deviations from linearity were observed above 0.4 g of starch/g of D₂O (Figure 2B). Previous studies on other systems have attributed such deviations from linearity to either (1) a change in the hydration of macromolecules (Baianu, 1989), (2) a change in the relaxation rate of bound water (Baianu et al., 1990b), or (3) restricted diffusion of water molecules between tightly packed starch granules in hydrated powders (Baianu, 1986; Mora-Gutierrez and Baianu, 1989, 1990; Yakubu et al., 1990b,c). Note also the marked increase in the relaxation rate gradient above 0.4 g of starch/g of D₂O (Figure 2B).

Table I. Hydration Parameters of Sigma Potato Starch for the Weakly Bound Water Population, Derived from ¹⁷O NMR Measurements

calcd parameters	correl coeff	suspension range, g of starch/g of D_2O
$n_{\rm H}^{\rm w} = 0.31 \pm 0.04$ $R_{2\rm B}^{\rm w} = 1000 \pm 20 {\rm s}^{-1}$ $\tau_{\rm c}^{\rm wb} = 20 \pm 0.6 {\rm ps}$	0 .999	0.00-0.35
$\vec{R}_{2F} = 236 \pm 9 \mathrm{s}^{-1}$	 Λ 	(60%)
(A)	\sum	(05%)
(B)		(60%)

FREQUENCY, Hz

Figure 3. Fourier transform ²H NMR spectra of D_2O in potato starch suspensions and powders at 20 °C. (A) Anisotropic slowly exchangeable water population within the potato granule structure at 0.65 g of starch/g of D_2O ; (B) spectrum of a 0.6 g of starch/g of D_2O suspension showing a Lorentzian peak on top of the doublet. The water population within the potato starch granule structure does not exchange rapidly with the other water populations.

The residual plot of $(R_{2obs} - R_{2calc})$ vs potato starchto-water ratio for values below 0.4 g of starch/g of D₂O and above this value, are presented in parts C and D, respectively, of Figure 2. In such a plot, a "run" is defined as a series of consecutive points with a residual of the same sign, positive or negative (Motulsky and Ransnas, 1987). At lower starch-to-water ratios from 0.02 to 0.4 g of starch/g of D₂O, there are seven runs for a total of 11 points, whereas in the range of higher starch-to-water rations from 0.4 to 0.60 g of starch/g of D₂O, there are four runs for a total of 5 points.

Nonlinear regression analysis of ¹⁷O NMR transverse relaxation data for potato starch in the low (0.2–0.4 g of starch/g of D₂O) and high (0.4–0.6 g of starch/g of D₂O) solid ranges are presented in Table I. The value of the correlation time of weakly bound water to potato starch granule surface in a low concentration range (≤ 0.4 g of starch/g of D₂O) is also given in the same table.

In addition to the single Lorentzian peak observed by both ^{17}O and ^{2}H NMR, deuterium NMR resolved a bound water population with anisotropic motions within the potato starch granule structure (Figure 3A,B). The "tightly" bound water is the only population remaining in potato starch powders above 0.6 g of starch/g of D₂O, as shown in Figure 3A. Quadrupole splittings of about 1 kHz were resolved in potato starch even in dilute starch suspensions (88% water). Deuterium NMR resolved a (35%) slowly exchangeable water population (D₂O) which is trapped within the potato starch granule and has strongly anisotropic motions.

A comparison of the normalized, or scaled, ¹⁷O and ²H NMR transverse relaxation rates, $R_{20be}^{Corr}/R_{2free}$, is presented in Figure 4A as a function of potato starch-to-water ratio. Oxygen-17 and ²H NMR relaxation rates have different, unscaled values because of the difference(s) between the magnitudes of the nuclear quadrupole moments of the two nuclei (Kakalis and Baianu, 1988; Mora-Gutierrez and Gaianu, 1990). The difference in the normalized ¹⁷O and ²H NMR transverse relaxation rates as a function of potato starch-to-water ratio is presented



Figure 4. (A) Dependence of the normalized ¹⁷O and ²H NMR transverse relaxation rates, R_2° (s⁻¹), on the starch-to-water ratios. Normalized rates were calculated with the following Rd_2° values of D₂O: $R_2^{\circ}({}^{17}\text{O}) = 235 \text{ s}^{-1}$ and $R_2^{\circ}({}^{2}\text{H}) = 1.86 \text{ s}^{-1}$. (B) Dependence of the difference between normalized ¹⁷O and ²H NMR transverse relaxation rates, R_2° (s⁻¹), on potato starch-to-water ratios. Normalized rates were calculated with the same R_2° values as in (A).



Figure 5. (A) Dependence of the normalized ²H and ¹H NMR transverse relaxation rates, R_2 (s⁻¹), on the potato starch-to-water ratios. Normalized rates were calculated with the following R_2^* values of D₂O (²H) and H₂O (¹H): $R_2^*(^2H) = 1.86 \text{ s}^{-1}$ and $R_2^*(^1H) = 0.33 \text{ s}^{-1}$. (B) Dependence of the difference between the normalized ²H and ¹H NMR transverse relaxation rates, (R_2) , on potato starch-to-water ratios. Normalized rates were calculated with the same R_2^* values in (A).

in Figure 4B. This plot reflects the effect of *chemical* exchange on the ²H NMR transverse relaxation rates of potato starch hydration.

The dependences of the normalized, or scaled, ²H and ¹H NMR transverse relaxation rates on the potato starchto-water ratio are shown in Figure 5A. ¹H NMR transverse relaxation rates vary more rapidly than the scaled ²H NMR relaxation rates (Figure 5B), as a result of proton intermolecular dipolar interactions (PIDI), which contributes only to ¹H. The difference between the scaled ²H and ¹H NMR transverse relaxation rates is shown in Figure 5B as a function of the starch-to-water ratio and reflects the effects of PIDI or cross-relaxation; note that the effect of PIDI cross-relaxation on the ¹H NMR relaxation rates is more marked above 0.2 g of potato starch/g of D_2O (Figure 5B). Comparing the relaxation contributions to 2 H and ¹H NMR relaxation rates presented in Figures 5 and 6, one notes that PIDI cross-relaxation contributes significantly more than proton/deuterium chemical exchange to the transverse relaxation rates of H_2O/D_2O in potato starch suspensions with less than 45% water (w/w).

DISCUSSION

Oxygen-17 NMR observations help resolve two hydration regions and three water populations in potato starch. The first region, which is linear, extends from 0.02 to 0.4g of starch/g of D₂O (Figure 2A). In this region, the free,



Figure 6. Dependence of the difference between normalized ${}^{17}\text{O}/{}^{2}\text{H}$ and ${}^{2}\text{H}/{}^{1}\text{H}$ NMR transverse relaxation rates, R_{2}^{*} (s⁻¹), on potato starch-to-water ratios. (A) Effect of PIDI cross-relaxation; (B) effects of chemical exchange.

or bulk, water and the surface-bound water populations make the major contributions to the observed transverse relaxation rates of water in potato starch suspensions. The second hydration region extends from 0.4 to 0.6 g of starch/g of D_2O (Figure 2B); this is also the beginning of aggregation and trapped water region in hydrated potato starch powders. A complete loss ("disappearance") of the oxygen-17 NMR peak was observed above 0.6 g of starch/g of D_2O . The water present within this region could only be observed by ²H and ¹H NMR.

The percentage of weakly bound water to potato starch is about 31.1 ± 4.3%, whereas the value of $R_2^{\rm wb}$ for ¹⁷O NMR of weakly bound water to potato starch is about $1000 \pm 29 \text{ s}^{-1}$. Previous ²H NMR hydration studies of hydrated potato starch granules resolved the slowly exchangeable water population trapped within the potato starch granule, which is about 35% w/w (Baianu et al., 1990a,b; Yakubu et al., 1990a,b). The percentage of weakly bound water to potato starch suspensions is less than that of the water population within the potato starch granule (35%). The weakly bound water and the bulk water populations contribute to the linear relaxation rate within the first hydration region (from 0.02 to 0.04 g of starch/g of D_2O). The percentage of trapped water between the potato starch granules is about $9.1 \pm 0.07\%$, whereas the ¹⁷O NMR transverse relaxation rate, R_2 , of the trapped water is about $2000 \pm 48 \text{ s}^{-1}$.

Three hydration regions and four water populations were resolved by ²H NMR in conjunction with ¹⁷O and ¹H NMR data. The first hydration region extended from 0.02 to 0.4 g of starch/g of D₂O, comprising a bulk water population, a weakly bound water fraction, and an anisotropic (slowly exchangeable) water population. In the third hydration region, above 0.6 g of starch/g of D₂O, only the anisotropic water (D₂O) population in hydrated potato starch remains consistent with previous continuous wave ²H NMR studies on potato starch powders (Lechert et al., 1980; Lechert, 1981). The latter, continuous wave, ²H NMR studies were also carried out on other starches



Figure 7. Two-component fittings of ¹H transverse magnetization decays in a 0.235-T magnetic field for water protons in potato starch suspensions at ~80% water content, 25 °C, and pH ~7. (A) Starch samples prepared from LaSoda (cultivars) potatoes (1991 and 1990 harvests; raw, 20%); (B) starch samples prepared from Norchip (cultivars) potatoes (1991 and 1990 harvests; raw, 20%); (C) data for commercial potato starch samples from Sigma (raw, 20% w/w). (\Box) M_{obs} ; (\diamond) residuals; (\bullet) M_{est} .

and revealed that bound water in potato starch powders had a resolved ²H NMR "powder pattern". On the other hand, water (D_2O) in corn, wheat, and pea starch powders gave essentially only a single Lorentzian isotropic water peak.

Determination of the Average Correlation Time, $\bar{\tau}_c^w$, of Weakly Bound Water. The average correlation time of weakly bound water at the potato starch granule surface can be calculated in the extreme narrowing limit $(\omega_0 \tau_c \ll 1)$ with

$$\bar{\tau}_{\rm c} = R_2^{\rm wb} / [K(e^2 q Q/\hbar)^2 (1 + \eta^2/3 - S^2)]$$
(5)

where $K(^{17}\text{O}) = 12 \pi^2/125$ and $K(^2\text{H}) = 3\pi/2$; e^2qQ/\hbar is the quadrupole coupling constant expressed in hertz (Abragam, 1961); for oxygen (^{17}O), $e^2qQ/\hbar = 6.67$ MHz (Halle et al., 1981), and for deuterium (^2H), $e^2qQ/\hbar = 0.214$ MHz (Boden and Mortimer, 1978) or 0.216 MHz (Pessen and Kumosinski, 1985); and η is the asymmetry parameter. R_2^{wb} was found to be 1000.2 s⁻¹ for ^{17}O and 300 s⁻¹ for ^{2}H NMR relaxation, respectively. The correlation time, $\bar{\tau}_c^{\text{w}}$, for weakly bound water at the potato starch granule surface was calculated with eq 5 and with the above R_2^{wb} value from ^{17}O NMR measurements. The value of the correlation time, $\bar{\tau}_c^{\text{w}}$, was thus found to be about 20 \pm 0.6 ps.

Conclusions. Oxygen-17, deuterium, and proton transverse, nuclear spin relaxation measurements were employed to quantitate the hydration properties of potato starch from dilute starch suspensions to hydrated powders. The weakly bound water at the potato starch granule surface has an average correlation time, τ_c^w , of about 20 ps in the range from 0.02 to 0.4 g of starch/g of D_2O . Deuterium NMR, on the other hand, resolved a slowly exchangeable anisotropic water population in the range from 0.12 g of starch/g of D_2O to 2% (g of D_2O/g of total) hydrated powders. Proton NMR was successful in monitoring the beginning of aggregation of potato starch suspensions at a low concentration (0.2 g of starch/g of) H_2O), because of the effects of proton intermolecular dipolar interactions cross-relaxation on the ¹H NMR relaxation (Figures 6A and 7). The presence of a slowly exchangeable, anisotropically bound water population with highly restricted motions within the starch granules (even in dilute potato starch suspensions) makes potato starch a suitable functional ingredient in food formulations for increasing their stability and storage life.

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