

^1H Spin-Spin Relaxation and Spin-Diffusion of Poly(vinyl phenol)/Poly(methyl acrylate) Blends

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(Received May 6, 1992)

ABSTRACT: ^1H spin-spin relaxation time (T_2) and ^1H spin-diffusion of poly(4-vinyl phenol)/poly(methyl acrylate) (PVPh/PMA) blend are measured indirectly by observing well-resolved ^{13}C CP/MAS resonance. The results indicate that at 310 K the molecular motion of the whole chain of PMA in the blends is hindered by mixing with PVPh. At 361 K, the blending hinders only the main-chain motion of PMA; the CH_3 group in the side-chain still undergoes more or less free rotation. The spin-diffusion from the CH_3 protons of PMA to the protons of PVPh occurs in the blend. It is concluded that PVPh and PMA are intimately mixed in the blend.

KEY WORDS ^1H Spin-Spin Relaxation / Spin-Diffusion / ^{13}C CP/MAS
NMR / Poly(vinyl phenol)/Poly(methyl acrylate) / Polymer Blend /

In a recent paper,¹ we reported high-resolution solid-state ^{13}C NMR studies of poly(4-vinyl phenol)/ poly(methyl acrylate) (PVPh/PMA) blend. The results indicated that the intermolecular hydrogen-bonding interaction is responsible for the two polymers to be intimately mixed on a scale of 20—30 Å. The intimate mixing is expected to bring about significant change in the molecular motion of the two polymers.

^1H spin-spin relaxation time (T_2) is a sensitive parameter for studying molecular motion of polymer.² It reflects the large-scale segmental motion related to the glass transition. It also reflects local motions of the main-chain and side-chain.³ Even a slight change in motion induced by blending would alter the relaxation time. The purpose of this paper is to obtain information about molecular motion of the PVPh/PMA blend.

We can measure the ^1H spin-spin relaxation

time by ^1H spin-echo experiment. It is, however, difficult to measure ^1H T_2 of each polymer in blend separately, because ^1H spectra of solid polymers are broad. The ^{13}C spectra observed by the magic-angle sample spinning and ^1H dipolar decoupling method are well resolved. We take this advantage to measure the ^1H T_2 indirectly by observing the high-resolution spectra of ^{13}C that directly attaches to the proton

Proton spin-diffusion provides information about the miscibility of polymer blend. Much work has been done on the spin-diffusion from theoretical and experimental viewpoints.⁴⁻¹⁴

Another purpose of this paper is to obtain information about the miscibility of the blend. We utilize the Goldman-Shen pulse sequence combined with cross polarization/magic-angle sample spinning (CP/MAS) to measure the spin diffusion between PMA and PVPh.

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EXPERIMENTAL

PVPh was obtained from Polysciences Inc. The molecular weight was 1500–7000. PMA was obtained from Aldrich Chemical Company Inc. The number- and weight-average molecular weights of PMA were 10600 and 30700, respectively.

The PVPh/PMA blend was prepared by the solution-cast method using tetrahydrofuran (THF) as a solvent. PVPh and PMA were separately dissolved in THF at a concentration of 1% (w/v). The two solutions were mixed together at room temperature at given weight ratios. After stirring the mixed solutions for 3 h, the solvent was allowed to evaporate at room temperature for several days. The resulting samples were further dried under reduced pressure at 333 K for two weeks. We prepared five blend samples; the composition of blend is expressed in weight ratio of the two polymers. Pure PVPh and PMA samples were prepared in the same way.

^{13}C CP/MAS NMR experiments were carried out on a JEOL JNM-GX270 spectrometer operating at 270 and 67.8 MHz for ^1H and ^{13}C , respectively. The radio-frequency field strength for both ^1H and ^{13}C was about 56 kHz. A double-bearing aluminum oxide rotor was used. The magic angle was adjusted by monitoring the ^{79}Br NMR spectrum of KBr incorporated in the rotor. The ^1H decoupling frequency was set 3 ppm downfield from tetramethylsilane (TMS). The ^{13}C CP/MAS spectra were measured by the normal CP/MAS pulse sequence at a magic-angle sample spinning rate of about 6 kHz. The CP contact time was 1.2 ms at 310 K and 0.6 ms at 361 K. The chemical shift of ^{13}C spectra was measured in ppm relative to the methine resonance of solid adamantane as 29.5 ppm. Variable-temperature measurements were made using a JEOL MVT temperature controller in an accuracy of 1 K. Temperature calibration was done by the conventional methanol method.

^1H T_2 measurement was made by the pulse

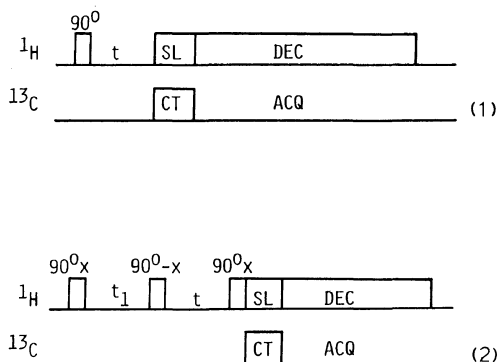


Figure 1. Pulse sequences used in the work. (1) pulse sequence for ^1H T_2 measurement with CP; (2) pulse sequence of Goldman–Shen spin-diffusion measurement with CP.

sequence shown in Figure 1(1).¹² The first ^1H 90° pulse creates a proton transversal magnetization. It decays with T_2 . The decayed magnetization at a delay time t is transferred to ^{13}C . Then we observe the ^{13}C resonance by the magic-angle spinning and broad-band ^1H decoupling technique. The delay time t was varied from $1\ \mu\text{s}$ to $100\ \mu\text{s}$.

The ^1H spin diffusion was measured using the Goldman–Shen pulse sequence⁴ combined with the CP/MAS technique shown in Figure 1(2).^{11,12} For a constant delay time t_1 ($22\ \mu\text{s}$), a shorter T_2 component of ^1H transversal magnetization ($T_2 < 22\ \mu\text{s}$) vanishes, and only a longer T_2 component remains. The second 90°_{-x} pulse, therefore, flips only the longer T_2 magnetization back in the z -direction. An inhomogeneous distribution of ^1H magnetization is created in the sample. For time t , the magnetization gradually approaches a uniform distribution through the spin-diffusion. We measure the ^1H magnetization at a time t by observing ^{13}C resonances using the CP/MAS technique.

RESULTS AND DISCUSSION

Figure 2 shows the ^{13}C CP/MAS spectra at 310 K of pure PVPh(A), PVPh/PMA = 1/1 blend(B), and pure PMA(C). The resonances

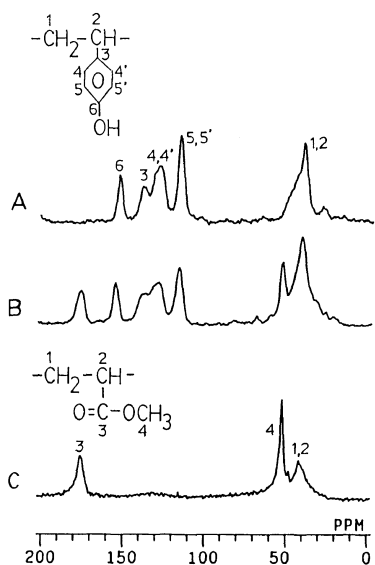


Figure 2. ^{13}C CP/MAS NMR spectra of PVPh/PMA blends at 310 K: A, pure PVPh; B, PVPh/PMA = 1/1; C, pure PMA.

of two polymers in the blend are well-resolved. We can, therefore, measure the ^1H T_2 of each polymer in the blend separately. Table I lists ^1H T_2 values at 310 K and 361 K of pure PVPh, five PVPh/PMA blends with different weight ratios, and pure PMA.

^1H T_2 decay of pure PVPh shows a fast Gaussian decay with a small T_2 value of about 13 μs for all PVPh carbon resonances at 310 and 361 K.

^1H T_2 values of the main-chain of pure PMA observed through the C-1 and C-2 resonances are also small (14 μs) at 310 K. On the other hand, neither single Gaussian decay nor single Lorentzian decay can represent the ^1H T_2 decay observed through the CH_3 (C-4) resonance of the side-chain as shown in Figure 3. The solid line represents the superposition of two types of decay: a slow Lorentzian with a T_2 value of 89 μs and a fast Gaussian with a T_2 value of 17 μs . At 361 K, the decay is a single Lorentzian with a larger T_2 value of 155–210 μs for all

Table I. ^1H T_2 (μs) of PVPh/PMA blends at 310 and 361 K^a

| | | PMA C-3 | PVPh C-6 | PVPh C-4 | PVPh C-5 | PMA C-4 | PVPh,PMA C-1,2 C-2 |
|------------------|-------|------------|-------------|-------------|-------------|-----------------------|-----------------------|
| PVPh | 310 K | | 13 | 13 | 13 | | 14 |
| | 361 K | | 13 | 13 | 13 | | 13 |
| PVPh/PMA = 4/1 | 310 K | 13 | 15 | 15 | 14 | 16 | 13 |
| | 361 K | | 14 | 14 | 13 | 14 (76%) 55 (24%)L | 14 |
| PVPh/PMA = 2/1 | 310 K | 14 | 15 | 15 | 15 | 15 | 14 |
| | 361 K | 14 | 13 | 14 | 14 | 14 (65%) 51 (35%)L | 13 |
| PVPh/PMA = 1/1 | 310 K | 14 | 15 | 15 | 15 | 14 | 14 |
| | 361 K | 14 | 15 | 15 | 15 | 14 (62%) 56 (38%)L | 14 |
| PVPh/PMA = 1/1.5 | 310 K | 13 | 12 | 13 | 13 | 12 | 12 |
| | 361 K | 19 | 22 | 18 | 22 | 16 (27%) 53 (73%)L | 16 |
| PVPh/PMA = 1/3 | 310 K | 14 | 16 | 16 | 14 | 13(82%) 86(18%)L | 14 |
| | 361 K | — | — | — | — | — | — |
| PMA | 310 K | 20 | | | | 17 (52%) 89 (48%)L | 14 |
| | 361 K | 155L | | | | 210L | 180L |

^a "L" denotes Lorentzian ^1H T_2 decay and all others, Gaussian decay. The value in parenthesis represents proton-number percentage of the two components.

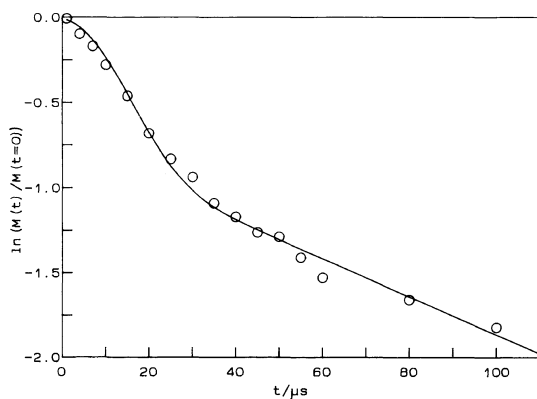


Figure 3. The normalized decay curve of ^1H magnetization of C-4 proton of pure PMA at 310 K. The solid line represents the superposition of a slow Lorentzian with a T_2 value of $89\ \mu\text{s}$ and a fast Gaussian with a T_2 value of $17\ \mu\text{s}$.

resonances of PMA.

The fast Gaussian decay with small T_2 observed for pure PVPh indicates that PVPh is in the glassy state at 310 and 361 K. This is consistent with the fact that the glass transition temperature T_g of PVPh is 393 K. The Lorentzian decay with the larger T_2 value observed for pure PMA at 361 K is in accord with the fact that PMA is in the rubbery state at this temperature.

Pure PMA is in a more complicated motional state at 310 K than PVPh. The main-chain motion of PMA is not fully excited at 310 K. The temperature of 310 K is not high enough to activate the main-chain motion on the NMR time scale. On the other hand, the CH_3 group of side-chain undergoes rotation rather freely.

Since the cross-polarization occurs through dipolar interaction, a large part of ^{13}C polarization is brought about by directly-bonded rigid protons. ^{13}C resonance of the PMA main-chain is, therefore, enhanced by the main-chain rigid protons, showing small ^1H T_2 at 310 K as seen in Table I. On the other hand, there is no proton that makes a major contribution to the ^{13}C polarization of the CH_3 group of PMA (C-4), because the directly-bonded CH_3 protons undergo rotation, and

the rigid main-chain protons are distant. It seems that the main-chain protons and CH_3 protons make a contribution to the ^{13}C polarization in a similar degree. There appear, therefore, two components: a fast Gaussian decay with T_2 of $17\ \mu\text{s}$ (52%) and a slow Lorentzian decay with T_2 of $89\ \mu\text{s}$ (48%). At 361 K, we only obtain a longer Lorentzian component with T_2 of $155\text{--}210\ \mu\text{s}$ for both the main-chain and the CH_3 carbons, indicating that rapid segmental motion occurs.

All PVPh/PMA blends studied here show a fast Gaussian component with a ^1H T_2 value of about $13\text{--}16\ \mu\text{s}$ at 310 K. The longer T_2 component of the CH_3 resonance of PMA is detected in a small fraction (18%) only for PVPh/PMA = 1/3 blend. At 361 K, the CH_3 resonance of PMA in the blends shows the fast Gaussian component, though the CH_3 of pure PMA does not. The T_2 value of the slow Lorentzian component of the CH_3 resonance of PMA in the blend is $51\text{--}56\ \mu\text{s}$, smaller than that of pure PMA ($210\ \mu\text{s}$).

These results suggest that PMA and PVPh are intimately mixed and that neighboring "hard" PVPh chains hinder the main-chain motion of PMA in the blend. The rotation of CH_3 group of PMA is not so greatly influenced by blending. This is comparable to polystyrene/poly(vinyl methyl ether) blend.¹⁵

Goldman-Shen experiments were performed to examine spin-diffusion in PVPh/PMA = 1/1.5 blend at 361 K. The CH_3 protons of PMA have long T_2 as compared to other protons. Immediately after the second 90°_x pulse, an inhomogeneous distribution of magnetization is created. For delay time t , the magnetization of CH_3 protons declines and those of other protons recover by spin diffusion.

On the basis of the method suggested by VanderHart,¹³ $P(t) = (M(t) - M_{\text{eq}}) / M_{\text{eq}}$ was plotted against $t^{1/2}$ as shown in Figure 4, where $M(t)$ is the magnetization after diffusion time t and M_{eq} is the magnetization in internal equilibrium. We see a linear relationship between $P(t)$ and $t^{1/2}$ at least at the initial stage.

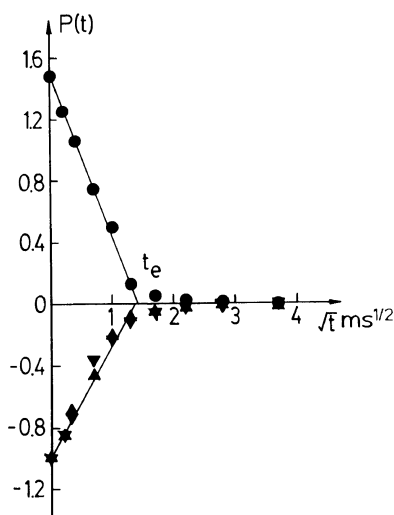


Figure 4. Plot of $P(t)$ vs. square root of spin-diffusion time $t^{1/2}$ for PVPh/PMA = 1/1.5 blends at 361 K. (●), (▼), and (▲) represent, respectively, data observed through the C-4 resonance of PMA, the C-5,5' resonances of PVPh, and the C-1, 2 resonance of both PVPh and PMA in the blend. Straight lines are drawn through the linear portion of the data at early stage to determine the intercept time t_e .

The decay rate of the CH_3 protons is comparable to the recovery rate of the C-5,5' protons of PVPh and those of the main-chain protons of PVPh and PMA. The results indicate that the spin-diffusion occurs from the CH_3 protons of PMA to other protons including the protons of PVPh in the blend in a similar way. This is other evidence for the good miscibility of the blend.

We define an effective spin-diffusion time t_e as the intercept of the straight line with the abscissa: time required for magnetization to reach the internal equilibrium.¹³ We obtain t_e of 2ms ($t_e^{1/2} = 1.4$). Because ^1H T_1 of the blend is longer than t_e , we can neglect the T_1 effect in the spin-diffusion process.

In summary, the present study indicates that at 310 K molecular motion of PMA in the blends is considerably hindered by the mixing with PVPh. At 361 K, the blending hinders only

the main-chain motion of PMA. The side-chain still undergoes more or less free rotation. The spin-diffusion between the CH_3 protons of the PMA side-chain and the other protons of PVPh occurs. The results obtained here provide additional evidence that PVPh and PMA are intimately mixed in the blend.¹ The two polymers associate each other so closely that they act as a single polymer.

Acknowledgment. This work was supported by Grants-in-Aid from the Ministry of Education, Science, and Culture of Japan (Grant No. 02554014 and 02750623). Xiaoqing Zhang was a recipient of the scholarship for foreign students from the Japanese Government.

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