

Supporting information

A Role of Electrostatic Repulsion in Gelation of Poly(vinyl alcohol)/Borax Aqueous Solutions

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I. Modified Reversible Gelation model for non-entangled semidilute solution: The complex modulus of this model is expressed as a sum of stress relaxation originating from the Rouse relaxation of the precursor chain G_X^* , the relaxation of the sol chains/gel strands formed in the mean-field and critical percolation regions, G_{MF}^* and G_{CP}^* , respectively, and the dissociation-activated relaxation of the gel network, G_N^* , as classified below for several ranges of the extent of gelation ε .^{1, 2}

for $-1 < \varepsilon < -\varepsilon_G$:

$$G^*(\omega) = G_{\text{Rouse}}^*(\omega) + G_{\text{MF}}^*(\omega)$$

$$= i\omega\nu kT \left[\sum_{q=2}^{N_\xi} \frac{\tau_X / q^2}{1 + i\omega\tau_X / q^2} + \int_{\tau_X}^{\infty} \frac{(\tau_X / \tau)^1}{i\omega + 1/\tau_{\text{char}} + 1/\tau} \frac{d\tau}{\tau} \right] \quad (\text{S1})$$

where $\tau_X = \tau_\xi N_\xi^2$ is the Rouse time of precursor chain, with τ_ξ being the relaxation time of the correlation blob of size ξ , and N_ξ is the number of blobs *per* chain. τ_{char} in eq 1 is the relaxation time of the characteristic sol chain, which scales with ε as $\tau_{\text{char}} = \tau_X |\varepsilon|^{-3}$.

for $-\varepsilon_G \leq \varepsilon < -\varepsilon_c$:

$$\begin{aligned}
G^*(\omega) &= G_{\text{Rouse}}^*(\omega) + G_{\text{MF}}^*(\omega) + G_{\text{CP}}^*(\omega) \\
&= i\omega\nu kT \left[\sum_{q=2}^{N_\xi} \frac{\tau_X / q^2}{1 + i\omega\tau_X / q^2} + \frac{(N_\xi - 1)}{N_\xi} \int_{\tau_X}^{\infty} \frac{(\tau_X / \tau)^1}{i\omega + 1/\tau_G + 1/\tau} \frac{d\tau}{\tau} + \frac{0.67}{N_\xi} \int_{\tau_G}^{\infty} \frac{(\tau_G / \tau)^{0.67}}{i\omega + 1/\tau_{\text{char}} + 1/\tau} \frac{d\tau}{\tau} \right]
\end{aligned} \tag{S2}$$

$\tau_G = \tau_X N_\xi$ is a characteristic time where a transition from the mean-field type to a critical percolation type occurs. τ_{char} denotes the relaxation time for the characteristic sol chains, which scales with ε as $\tau_{\text{char}} = \tau_G |\varepsilon / \varepsilon_G|^{-4}$, with $\tau_G = \tau_X N_\xi^{-1}$ being the characteristic time of the sol chains at $\varepsilon = -\varepsilon_G$.

for $-\varepsilon_c \leq \varepsilon < \varepsilon_G$

$$\begin{aligned}
G^*(\omega) &= G_{\text{Rouse}}^*(\omega) + G_{\text{MF}}^*(\omega) + G_{\text{CP}}^*(\omega) \\
&= i\omega\nu kT \left[\sum_{q=2}^{N_\xi} \frac{\tau_X / q^2}{1 + i\omega\tau_X / q^2} + \frac{(N_\xi - 1)}{N_\xi} \int_{\tau_X}^{\infty} \frac{(\tau_X / \tau)^1}{i\omega + 1/\tau_G + 1/\tau} \frac{d\tau}{\tau} + \frac{0.67}{N_\xi} \int_{\tau_G}^{\infty} \frac{(\tau_G / \tau)^{0.67}}{i\omega + 1/(\Lambda\tau_c) + 1/\tau} \frac{d\tau}{\tau} \right]
\end{aligned} \tag{S3}$$

In this range of ε , the system includes very large critical sol chains/gel strands of the relaxation time $\tau_c = \tau_X^{0.25} \tau_s^{0.75}$. τ_c is the Rouse time of a critical sol chain that exhibits the Rouse time identical to the effective breakup time. Λ is a fitting parameter to the order of 1. Rubinstein and Semenov³ argued that the sol chains/network strands larger than this critical chain (at $-\varepsilon_c \leq \varepsilon < \varepsilon_c$) would exhibit effective breakups continuously before the resulting pieces have the critical size, where the pieces further relax through the Rouse-type relaxation to exhibit relaxation time $\sim \tau_c$.

for $\varepsilon_c \leq \varepsilon < \varepsilon_G$

$$\begin{aligned}
G^*(\omega) &= G_{\text{Rouse}}^*(\omega) + G_{\text{MF}}^*(\omega) + G_{\text{CP}}^*(\omega) + G_{\text{N}}^*(\omega) \\
&= i\omega\nu kT \left[\sum_{q=2}^{N_\xi} \frac{\tau_X / q^2}{1 + i\omega\tau_X / q^2} + \frac{(N_\xi - 1)}{N_\xi} \int_{\tau_X}^{\infty} \frac{(\tau_X / \tau)^1}{i\omega + 1/\tau_G + 1/\tau} \frac{d\tau}{\tau} + \frac{1}{0.67 N_\xi} \int_{\tau_G}^{\infty} \frac{(\tau_G / \tau)^{0.67}}{i\omega + 1/\tau_{\text{strand}} + 1/\tau} \frac{d\tau}{\tau} + \frac{(\varepsilon / \varepsilon_G)^{2.7}}{N_\xi} \frac{\tau_{\text{life}}}{1 + i\omega\tau_{\text{life}}} \right]
\end{aligned} \tag{S4}$$

$\tau_{\text{strand}} = \tau_G |\varepsilon / \varepsilon_G|^{-4}$ is an average Rouse time of a network strand, and $\tau_{\text{life}} = \tau_s N_\xi^{-1/3} (\varepsilon / \varepsilon_G)^{1.33}$ is the lifetime of the strand controlled by ionic dissociation time τ_s .

for $\varepsilon_G \leq \varepsilon < 1$

$$\begin{aligned}
G^*(\omega) &= G_{\text{Rouse}}^*(\omega) + G_{\text{MF}}^*(\omega) + G_{\text{N}}^*(\omega) \\
&= i\omega\nu kT \left[\sum_{q=2}^{N_\xi} \frac{\tau_X / q^2}{1 + i\omega\tau_X / q^2} + \frac{(N_\xi - 1)}{N_\xi} \int_{\tau_X}^{\infty} \frac{(\tau_X / \tau)^l}{i\omega + 1/\tau_{\text{strand}} + 1/\tau} \frac{d\tau}{\tau} + \varepsilon^3 \frac{\tau_{\text{life}}}{1 + i\omega\tau_{\text{life}}} \right] \quad (\text{S5})
\end{aligned}$$

where $\tau_{\text{strand}} = \tau_X \varepsilon^{-3}$ and $\tau_{\text{life}} = \tau_s \varepsilon$.

Equations S1-S5 give the modulus for the ionomers at $-1 \leq \varepsilon \leq 1$, i.e., those still containing sol chains. At $\varepsilon > 1$, all the precursor chains in the system are incorporated in the gel, and the modulus becomes sticky Rouse type:^{1, 2, 4}

for $1 \leq \varepsilon$

$$\begin{aligned}
G^*(\omega) &= G_{\text{Rouse}}^*(\omega) + G_{\text{SR}}^*(\omega) \\
&= i\omega\nu kT \left[\sum_{q=\varepsilon+1}^{N_\xi} \frac{\tau_X / q^2}{1 + i\omega\tau_X / q^2} + \sum_{q=1}^{\varepsilon} \frac{\tau_s \varepsilon^2 / q^2}{1 + i\omega\tau_s \varepsilon^2 / q^2} \right] \quad (\text{S6})
\end{aligned}$$

Here, G_{Rouse}^* and G_{SR}^* are the *intrinsic* Rouse and sticky Rouse parts of the moduli, respectively. When the sticky Rouse relaxation manifests (at $\varepsilon \geq 1$), the intrinsic Rouse relaxation of the chain is limited for a portion of chain between stickers and thus to the mode indices $q > \varepsilon$. The terminal relaxation of the chain becomes sticky-Rouse type at larger length scales, as represented by the second summation for $q \leq \varepsilon$.^{1, 2, 4}

II. Shear rate dependences of viscosity: As an example, Figure 1 plots the viscosity η against shear rate $\dot{\gamma}$ for the PVA135 and PVA92 aqueous solutions of different weight fraction w as indicated. At low concentration ($w \leq 6\text{wt}\%$ for PVA135 and $w \leq 10\text{wt}\%$ for PVA92), the solutions behave as Newtonian fluid for shear rate $\dot{\gamma} < 10^3 \text{s}^{-1}$, for which the zero-shear viscosity η_0 can be determined as an average value of all data points (lines attached). In contrast, for high concentration ($w \geq 8\text{wt}\%$ for PVA135 and $w \geq 12\text{wt}\%$ for PVA92), shear thinning is observed at high shear rate, for which the data are fit to the Carreau model (see curves),⁵

$$\eta(\dot{\gamma}) = \frac{\eta_0}{\left[1 + (\dot{\gamma}\tau)^2 \right]^m} \quad (\text{S7})$$

where η_0 is the zero-shear viscosity, τ is a characteristic relaxation time for shear thinning to occur, and m characterizes power law behavior at high shear.

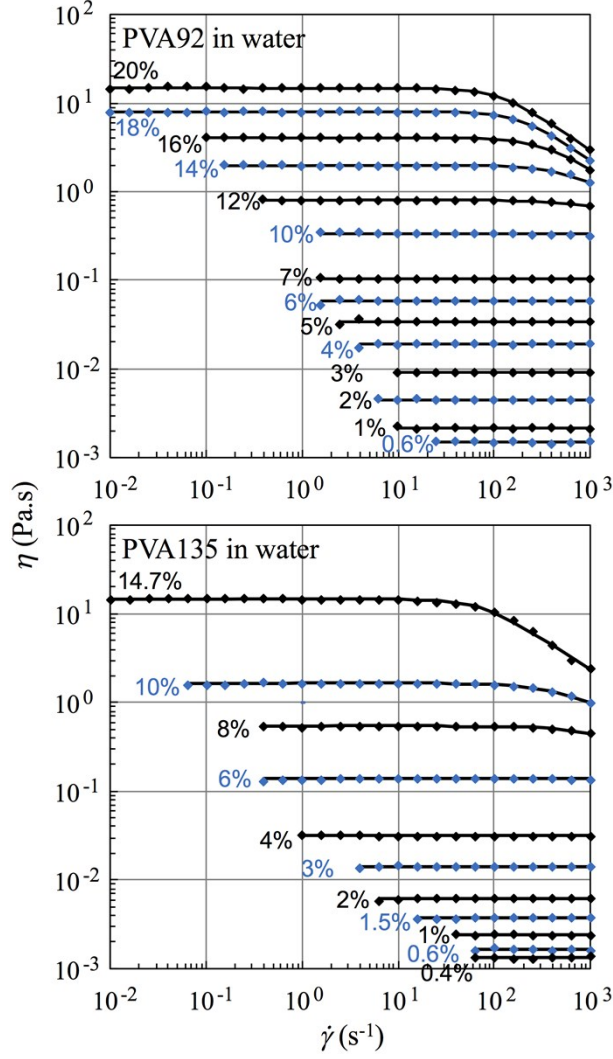


Figure S1: plots of shear viscosity η as a function of shear rate $\dot{\gamma}$ for the PVA92 and PVA135 aqueous representative solutions with concentrations of PVA as indicated.

III. Analysis of LVE reported by Narita and Indei: Since the concentration of PVA is similar to present study, we used $M_\xi \sim C_{PVA}^{-1/(3\nu-1)}$ and $\tau_0 \sim M_\xi \zeta^2 \sim M_\xi^{1+2\nu}$ to estimate $M_\xi = 14.1\text{kDa}$ and $\tau_0 = 0.13\mu\text{s}$ for the 5.5wt% PVA89 ($M_w = 89\text{kDa}$) solution on a basis of $M_\xi = 16.0\text{kDa}$ and $\tau_0 = 0.16\mu\text{s}$ of the 5wt% PVA92 solution. τ_s is assumed to be the same as that of the PVA92 sample. By taking ε as the only fitting parameter, we can fit LVE as shown in Figure S2. During the fitting, we tried to fit both G'' and G' for the samples of $[\text{borax}] \geq 1.7\text{mM}$. Nevertheless, for samples having $[\text{borax}] \leq 1.2\text{mM}$, the similar agreement cannot be achieved for G'' and G' , where we try to fit G'' that is less vulnerable to the inertia effect than G' .

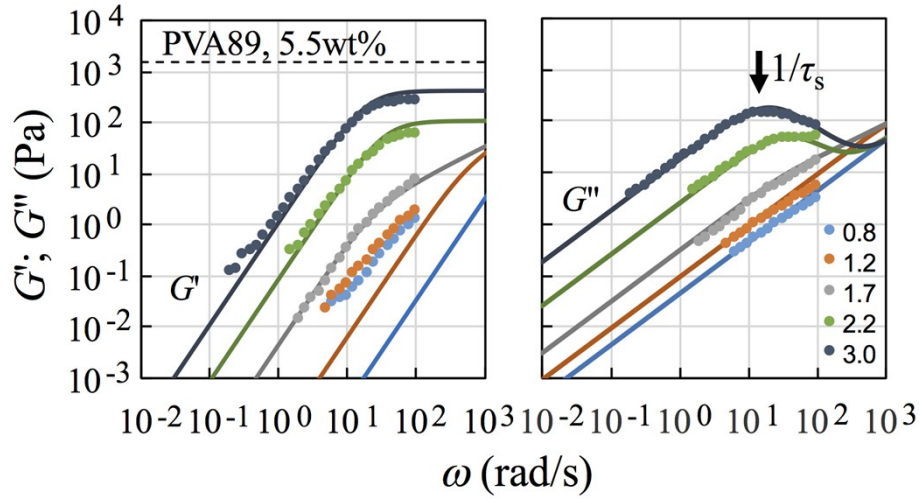


Figure S2: Storage and loss moduli, G' and G'' as function of angular frequency ω reported by Narita and Indei. The symbols are experimental results and the curves are fitting curve of the reversible gelation model, with degree of gelation ε as an adjustable parameter. For the two samples having 0.8mM and 1.2mM borax, the fitting can only be achieved simultaneously for G' and G'' , the fitting is focused on G'' that is less vulnerable to instrument inertia.

References

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