

## Supplemental Information for

# Rod-Coil Block Copolymer Aggregates via Polymerization-Induced Self-Assembly

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## CONTENT

1. The persistence length of rod blocks.....	S3
2. Relationship between $k[\mathbf{C}^*]$ and $[\mathbf{M}]$ .....	S5
3. Relationship between $k$ and $[\mathbf{C}^*]$ .....	S6

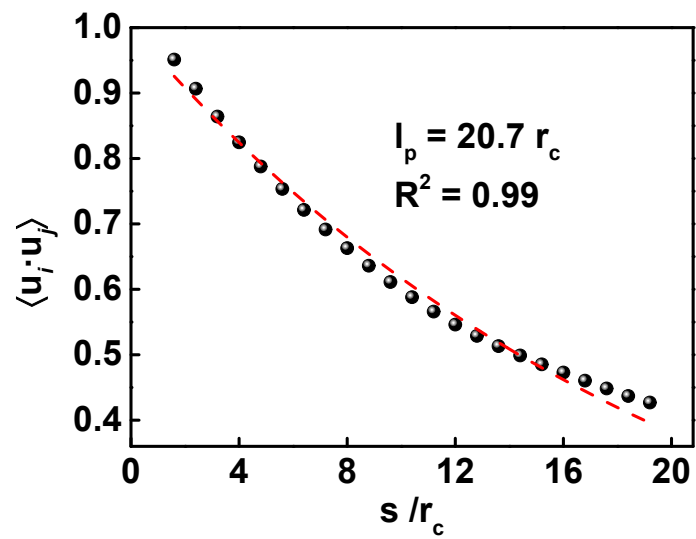
## 1. The persistence length of rod blocks

The persistence length of rod block was calculated to evaluate the rigidity of rod blocks. The persistence length  $l_p$  is given by

$$\langle \mathbf{u}_i \cdot \mathbf{u}_j \rangle = \exp(-s / l_p) \quad (\text{S-1})$$

Here,  $\mathbf{u}_i$  and  $\mathbf{u}_j$  are the unit vectors of  $i$ th and  $j$ th bond, respectively.  $s=|i-j|l_b$  is the contour distance between  $i$ th and  $j$ th bond, where  $l_b$  is the bond length. Here,  $l_b$  nearly equals to the equilibrium bond distance (*ca.*  $0.8 r_c$ ). In the simulation, we subjected a single rigid chain ( $k_\theta = 10$ ) with 25 DPD beads in a simulation box of  $L = 40r_c$  with periodic boundary conditions and full of solvent beads. The system was simulated with all  $a_{ij}$  equal to 25 for  $2.0 \times 10^6$  DPD steps.

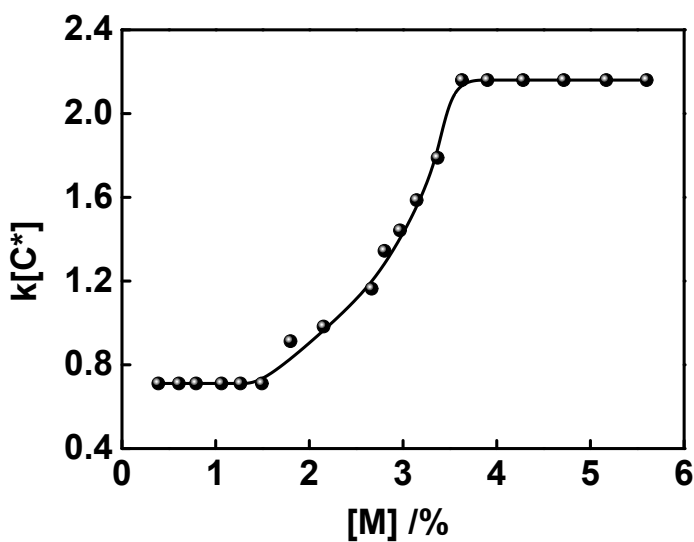
Figure S1 shows the plot of  $\langle \mathbf{u}_i \cdot \mathbf{u}_j \rangle$  as a function of  $s$ . By fitting the data with the Eq. S-1, we obtained the persistence length. The persistence length of rod blocks is *ca.*  $20.7 r_c$ . Considering that the maximum targeted DP of rod blocks studied in this work is 10, the maximum contour distance  $s$  is *ca.*  $7.2 r_c$ . The persistence length of rod blocks is significantly larger than the contour distance, indicating the sufficient rigidity of the rod blocks.



**Figure S1.** The plot of  $\langle \mathbf{u}_i \cdot \mathbf{u}_j \rangle$  as a function of  $s$ . The red dashed line is the fitting line.

## 2. Relationship between $k[C^*]$ and $[M]$

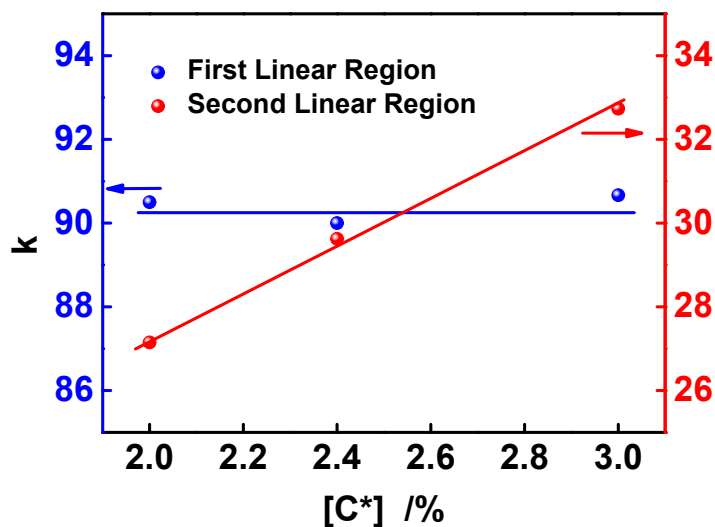
We calculated pseudo-first-order reaction rate constant  $k[C^*]$  at different monomer concentrations  $[M]$  to figure out whether or not the  $k[C^*]$  depends upon the concentration of monomers in the PISA. The results are shown in Figure S2. As shown,  $k[C^*]$  depends significantly on  $[M]$ . However,  $k[C^*]$  is unchanged at lower and higher  $[M]$ , which correspond to the two linear regions in Figure 2. The lower and higher  $[M]$  correspond to the aggregate-dominated and free-chain dominated states, respectively.  $k[C^*]$  decreases with the decrease of  $[M]$  in the range from  $[M]=1.5\%$  to  $[M]=3.6\%$  because of the heterogeneous growths of the free chain and the aggregated chain. As such,  $k[C^*]$  is mainly governed by the state of copolymer chains (free chains and aggregated chains) rather than the concentration of monomers during the PISA.



**Figure S2.** Plots of  $k[C^*]$  as a function of monomer concentrations  $[M]$  during the PISA. The targeted DP of the rod blocks, the macromolecular initiator, and the copolymer concentrations are 7,  $C_3^*$ , and 8%, respectively.

### 3. Relationship between $k$ and $[C^*]$

Figure S3 shows the rate coefficient  $k$  against the initiator concentrations  $[C^*]$ . The  $k$  values for the first linear (free-chain dominated) regions and the second linear (aggregated dominated) region are respectively given in the Figure. The values of  $k$  in the first linear region are almost unchanged with the increasing of  $[C^*]$ , denoting that  $k$  is independent of  $[C^*]$ . This fact indicates that the polymerization in the first linear (free-chain dominated) region is similar to the free radical polymerization. In the second linear region,  $k$  increases with the increase of  $[C^*]$ . In this region, the growth of the chains is limited by the aggregate. At higher  $[C^*]$  (hydrophobic chain), smaller aggregates are formed, and therefore, the chains are less confined and can grow more rapidly.



**Figure S3.** Plots of the rate coefficient  $k$  against initiator concentrations  $[C^*]$  in the first and second linear regions. The macromolecular initiator and the copolymer concentrations are  $C_3^*$  and 8%, respectively.