Dual Self-Assembly in Strongly Asymmetric A-B-A Triblock Copolymer Melts Studied by Self-Consistent Field Theory and Monte Carlo Simulations

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Abstract: Using the Self-Consistent Field Theory (SCFT) we study the dual self-assembly of ABA triblock copolymers melts and compare the numerical results with those obtained by the lattice Monte Carlo simulations. While the results are qualitatively similar for both methods, the simulation times are significantly shorter for the SCFT calculations than those for the corresponding Monte Carlo simulations.

Key words: triblock copolymer, self-consistent field theory, Monte Carlo simulation, self-assembly

I. INTRODUCTION

Self-assembly in the block copolymer melts is an active area of research [1-16]. Block copolymers self-assemble via the microphase separation which is an instant of the orderdisorder transition (ODT). This transition leads to formation of a plethora of different nanostructures, such as ordered layers, hexagonally arranged cylinder, cubically ordered spheres or a double gyroid phase. Previous studies [17-23] of molecularly asymmetric A1BA2 triblock copolymers synthesized from a parent diblock copolymer so that N_{A1} is significantly smaller than N_{A2} (where N_{A1} and N_{A2} denote the number of repeat units in the A1 and A2 blocks, respectively) have helped to elucidate the molecular and property changes accompanying the transformation from an AB diblock to a molecularly symmetric ABA triblock copolymer (with $N_{A1} = N_{A2}$). Recent Monte Carlo (MC) simulations of moderately segregated copolymers have yielded results that quantitatively agree with unexpected experimental findings, most notably a pronounced minimum in the order-disorder transition temperature as N_{A2} is progressively increased.

In the limit of superstrong segregation (SSS) [24, 25], interstitial micelles composed of the minority A2 endblock are observed to be arranged into two-dimensional hexagonal arrays along the midplane of B-rich lamellae in compositionally symmetric (50:50 A:B) copolymers. Calculations performed here establish the coupled molecular-asymmetry and incompatibility conditions under which such micelles form. Beyond an optimal length of the A2 endblock, the propensity for interstitial micelles to develop decreases, and the likelihood for colocation of both endblocks in the A 1 -rich lamellae increases. The SSS regime has also been studied experimentally [26-30] and nontraditional morphologies have been observed [27, 28, 31]. From previous studies [17-23] we know that as the A1BA2 triblock melt begins to self-organize, short A-blocks initially locate in the B-domain, and then as the χ value grows they migrate to A-domain. Moreover, for sufficiently long chains and relatively high χ values we can observe that short A-segments inside the B-domain aggregate into interstitial micelles (IM's).

This observation will be investigated by the selfconsistent field theory (SCFT) and reaffirmed by lattice Monte Carlo methods, which are described in the following section.

II. METHODS

II. 1. SCFT Method

The incompressible ABA copolymer melt is modeled as a collection of *n* diblock chains confined in volume *V*. Each chain, labeled $\alpha = 1, 2, ..., n$, can take any Gaussian configuration parameterized from *s*=0 to *s*=*f*₁ for *A*-segments, from *s*=*f*₁ to *s*=*f*₂ for *B*-segments, and from *s*=*f*₂ to *s*=1 for *A*-segments. Up to a multiplicative constant, the partition function for a *single* Gaussian chain in external fields $W_A(\mathbf{r})$ and $W_B(\mathbf{r})$ acting on segments A and B, respectively, is

$$\mathcal{Q}[W_A, W_B] \equiv \int \tilde{\mathcal{D}} \mathbf{r}_{\alpha}(\cdot) \exp\left[-\int_0^f ds W_A(\mathbf{r}_{\alpha}(s)) - \int_{f_1}^{f_2} ds W_B(\mathbf{r}_{\alpha}(s)) - \int_{f_2}^1 ds W_A(\mathbf{r}_{\alpha}(s))\right]$$
(1)

The path integral, $\int \tilde{D} \mathbf{r}_{\alpha}(\cdot)$, is taken over single-chain trajectories, $\mathbf{r}_{\alpha}(s)$, with Wiener measure expressed as $\tilde{D} \mathbf{r}_{\alpha} = \mathcal{D} \mathbf{r}_{\alpha} P[\mathbf{r}_{\alpha}; 0, 1]$, and

$$P[\mathbf{r}_{\alpha}; s_1, s_2] \propto \exp\left[-\frac{3}{2Na^2} \int_{s_1}^{s_2} ds |\frac{d}{ds} \mathbf{r}_{\alpha}(s)|^2\right] \quad (2)$$

Note that a is the segment size, and Na^2 is the mean squared end-to-end distance of a Gaussian chain. By Kac-Feynman theorem, eq 1 can be related to a Fokker-Planck partial differential equation, known also as a modified diffusion equation (MDE) and shown with appropriate details below (eqs 15 and 16).

Segments A and B interact via the χ parameter which provides an effective measure of incompatibility between them. Evaluation of the full partition function of n interacting diblock chains, shown below (eq 3), is a highly challenging task, involving many-body interactions, both intermolecular and intramolecular.

$$Z = \int \prod_{\alpha=1}^{n} \tilde{\mathcal{D}} \mathbf{r}_{\alpha} \, \delta[1 - \hat{\phi}_{A} - \hat{\phi}_{B}] \, \exp\left[-\chi \rho_{0} \hat{\phi}_{A} \hat{\phi}_{B}\right], \, (3)$$

where δ -function enforces incompressibility (the melt is assumed to be incompressible), and

$$\hat{\phi}_{A}(\mathbf{r}) = \frac{N}{\rho_{0}} \sum_{\alpha=1}^{n} \left[\int_{0}^{f_{1}} ds \, \delta(\mathbf{r} - \mathbf{r}_{\alpha}(s)) + \int_{f_{2}}^{1} ds \, \delta(\mathbf{r} - \mathbf{r}_{\alpha}(s)) \right]$$
(4)

$$\hat{\phi}_B(\mathbf{r}) = \frac{N}{\rho_0} \sum_{\alpha=1}^n \int_{f_1}^{f_2} ds \,\delta(\mathbf{r} - \mathbf{r}_\alpha(s)) \tag{5}$$

are the microscopic segments densities of A and B, respectively; $\rho_0 = nN/V$ is the segment number density. After replacing microscopic segment (or particle) densities with a variety of fields, by inserting and spectrally decomposing the appropriate δ -functionals, the partition function of an incompressible diblock melt is

$$Z = \mathcal{N} \int \mathcal{D}\phi_A(\cdot) \mathcal{D}W_A(\cdot) \mathcal{D}\phi_B(\cdot) \mathcal{D}W_B(\cdot) \mathcal{D}\Psi(\cdot)$$
$$\exp\left[-\frac{F\left[\phi_A, W_A, \phi_B, W_B, \Psi\right]}{k_B T}\right],$$
(6)

where \mathcal{N} is a normalization factor. The functional integral is taken over the relevant fields $\phi_A(\mathbf{r}), W_A(\mathbf{r}), \phi_B(\mathbf{r}), W_B(\mathbf{r}), \text{ and } \Psi(\mathbf{r})$, with the free energy functional, $F[\phi_A, W_A, \phi_B, W_B, \Psi]$, including the single chain partition function (in external fields $W_A(\mathbf{r})$ and $W_B(\mathbf{r})$), as shown below

$$\frac{F\left[\phi_{A}, W_{A}, \phi_{B}, W_{B}, \Psi\right]}{nk_{B}T} \equiv$$

$$\equiv -\ln\frac{Q}{V} + V^{-1} \int d\mathbf{r} [N\chi\phi_{A}\left(\mathbf{r}\right)\phi_{B}\left(\mathbf{r}\right) \qquad (7)$$

$$-W_{A}\left(\mathbf{r}\right)\phi_{A}\left(\mathbf{r}\right) - W_{B}\left(\mathbf{r}\right)\phi_{B}\left(\mathbf{r}\right)$$

$$-\Psi\left(\mathbf{r}\right)\left(1 - \phi_{A}\left(\mathbf{r}\right) - \phi_{B}\left(\mathbf{r}\right)\right)\right]$$

Fields $\phi_A(\mathbf{r})$ and $\phi_B(\mathbf{r})$ are associated with normalized concentration profiles of A and B, and fields $W_A(\mathbf{r})$ and $W_B(\mathbf{r})$ with chemical potential fields acting on A and B, respectively; field $\Psi(\mathbf{r})$ enforces incompressibility. Evaluating functional integrals in eq 6 is a challenging task which, in principle, can be performed by field theoretic simulations. A simpler, but approximate, approach is based on the mean-field idea, where the dominant and, in fact, only contribution to the functional integral in eq 6 comes from the fields satisfying the saddle point condition expressed as the following set of equations:

$$\frac{\delta F}{\delta \phi_A} = \frac{\delta F}{\delta \phi_B} = \frac{\delta F}{\delta W_A} = \frac{\delta F}{\delta W_B} = \frac{\delta F}{\delta \Psi} = 0 \qquad (8)$$

Performing the above functional derivatives yields

$$W_A(\mathbf{r}) = N\chi\phi_B(\mathbf{r}) + \Psi(\mathbf{r})$$
(9)

$$W_B(\mathbf{r}) = N\chi\phi_A(\mathbf{r}) + \Psi(\mathbf{r})$$
 (10)

$$1 = \phi_A(\mathbf{r}) + \phi_B(\mathbf{r}) \tag{11}$$

$$\phi_A(\mathbf{r}) = \frac{V}{Q} [\int_0^{J^1} ds \ q(\mathbf{r}, s) q^{\dagger}(\mathbf{r}, s)$$
(12)

$$+ \int_{f_2}^{1} ds \ q(\mathbf{r}, s) q^{\dagger}(\mathbf{r}, s)]$$

$$\phi_B(\mathbf{r}) = \frac{V}{Q} \int_{f_1}^{f_2} ds \ q(\mathbf{r}, s) q^{\dagger}(\mathbf{r}, s), \qquad (13)$$

where Q/V can be calculated as

$$\frac{\mathcal{Q}}{V} = \frac{1}{V} \int d\mathbf{r} \, q(\mathbf{r}, 1) \tag{14}$$

and $q(\mathbf{r}, s)$ is the forward chain propagator which is the solution of the following modified diffusion equation

$$\frac{\partial q}{\partial s} = \frac{1}{6} N a^2 \nabla^2 q - W_A(\mathbf{r}) q, \quad 0 \le s \le f_1$$

$$\frac{\partial q}{\partial s} = \frac{1}{6} N a^2 \nabla^2 q - W_B(\mathbf{r}) q, \quad f_1 \le s \le f_2$$

$$\frac{\partial q}{\partial s} = \frac{1}{6} N a^2 \nabla^2 q - W_A(\mathbf{r}) q, \quad f_2 \le s \le 1 \quad (15)$$

with the initial condition $q(\mathbf{r}, 0) = 1$. Similarly $q^{\dagger}(\mathbf{r}, s)$ is the backward chain propagator which is the solution of the conjugate modified diffusion equation:

$$-\frac{\partial q^{\dagger}}{\partial s} = \frac{1}{6}Na^{2}\nabla^{2}q^{\dagger} - W_{A}(\mathbf{r})q^{\dagger}, \quad 0 \le s \le f_{1}$$
$$-\frac{\partial q^{\dagger}}{\partial s} = \frac{1}{6}Na^{2}\nabla^{2}q^{\dagger} - W_{B}(\mathbf{r})q^{\dagger}, \quad f_{1} \le s \le f_{2}$$
$$-\frac{\partial q^{\dagger}}{\partial s} = \frac{1}{6}Na^{2}\nabla^{2}q^{\dagger} - W_{A}(\mathbf{r})q^{\dagger}, \quad f_{2} \le s \le 1$$
(16)

with the initial condition $q^{\dagger}(\mathbf{r}, 1) = 1$.

While the set of equations 9, 10, 11, 12, and 13 can be solved, in principle, in a self-consistent manner, it is difficult to solve this set without some additional assumptions. First, we assume that the melt forms a spatially ordered nanophase. Second, we use the UCA which is a considerable simplification, limiting our attention to a single D-dimensional spherical cell of radius R, and volume V. All fields within this cell have radial symmetry, which reduces this problem computationally to a single radial coordinate, r. Thus eq 14 can be rewritten as

$$\frac{\mathcal{Q}}{V} = D \frac{\int_0^R r^{D-1} q(r,1) dr}{R^D}$$
(17)

Note that the factor, D, in front of the above integral originates from the ratio of the area of a sphere with radius 1 to

the volume of a spherical cell with the same radius, both in D dimensions.

While in integrals (eqs 12, 13 and 14) we replace \mathbf{r} with r, and $d\mathbf{r}/V$ with $Dr^{D-1}dr/R^D$, in the modified diffusion equations, 15 and 16, we replace \mathbf{r} with r and use the spherically symmetric form of the Laplacian

$$\nabla^2 f = \frac{\partial^2 f}{\partial r^2} + \frac{D-1}{r} \frac{\partial f}{\partial r}$$
(18)

and similarly, in equations for both propagators q(r, s) and $q^{\dagger}(r, s)$, we replace **r** with r. Obviously the solution depends on radius, R, and dimensionality, D = 1, 2 and 3, corresponding to 3 different nanophases. We use the Crank-Nicholson scheme to solve iteratively the modified diffusion equations (eqs 15 and 16) in their radial form, until the self-consistency condition is met, obtaining the saddle point fields, $\overline{\phi_A}(r), \overline{\phi_B}(r), \overline{W_A}(r)$ and $\overline{W_B}(r)$ for a given R and D. In the MF approximation, the free energy functional becomes the free energy, and therefore we calculate the reduced free energy (per chain in k_BT units) by substituting the saddle point fields into eq 7:

$$\frac{F(R,D)}{nk_BT} \equiv -\ln\frac{Q}{V} + \frac{D}{R^D} \int_0^R r^{D-1} [N\chi\overline{\phi_A}(r)\overline{\phi_B}(r) - \overline{W_A}(r)\overline{\phi_A}(r) - \overline{W_B}(r)\overline{\phi_B}(r)] dr$$
(19)

Since in the MF theory, the stability of a nanophase depends on the product χN and composition, f, we start, at a given point of the phase diagram, $(f, \chi N)$, with numerical calculation of F(R, D) (eq 19) for various D's (1, 2, and 3) and R's. In order to solve the MDE's (eqs 15 and 16) we use up to $N_T = 1000$ and up to $N_R = 2000$ steps for the "time", s, and space, r, variables, respectively.

Numerically, we find R and D which minimize F(R, D), and this allows us to determine the dimensionality, D, of the most stable nanophase, and therefore the most favorable nanophase itself. But the free energy of this nanophase has to be compared to that of the disordered phase. Therefore, we calculate the difference

$$\frac{\Delta F}{nk_BT} \equiv \frac{F}{nk_BT} - \frac{F_{dis}}{nk_BT} \tag{20}$$

where F_{dis} is the free energy of the disordered phase:

$$\frac{F_{dis}}{nk_BT} = N\chi f(1-f) \tag{21}$$

If ΔF is negative then the appropriate nanophase is thermodynamically stable for the point considered, $(f, \chi N)$; otherwise the system is the disordered phase.

II. 2. MC Method

The cooperative motion algorithm (CMA) [32-36], based on FCC lattice, is used to simulate the triblock solutions. We apply standard Monte Carlo (MC) simulations with the Metropolis algorithm [37] as well as parallel tempering (PT) method [38-40]. In the PT case, M replicas of system are simulated in parallel, each in different temperature T_i , with i ranging from 1 to M. After 3000 MC steps we try to exchange replicas with neighboring T_i in random order with probability:

$$p(T_i \leftrightarrow T_{i+1}) = \\= \min[1, \exp(-(\beta_i - \beta_{i+1})(U_{i+1} - U_i))],$$
(22)

where $\beta_i = 1/kT_i$ and U_i is potential energy of replica at T_i . This method offers efficient equilibration at low temperatures.

We repeat the experiment at least 3 times starting with different initial configurations, in which the polymer chains assume statistical conformations and random orientations, and are uniformly distributed within the simulation box. A single MC step is defined as an attempt to move a given segment. Usually, the first half of run is used to equilibrate the system and the second one to collect the data. The results are averaged over all simulation runs. While the morphologies obtained in the simulations were quantitatively dependent on the box size, this dependence was weak and did not change the main conclusions qualitatively.

We use the following set of interaction energies which are limited to the nearest neighbors:

$$\epsilon_{AB} = \epsilon \tag{23}$$

$$\epsilon_{AA} = 0 \tag{24}$$

$$\epsilon_{BB} = 0, \qquad (25)$$

where ϵ is an energy unit, and we define the reduced energy per lattice site and the reduced temperature as:

$$\frac{E^*}{n_a} = \frac{E/\epsilon}{n_a} \tag{26}$$

$$T^* = \frac{kT}{\epsilon} \tag{27}$$

On the basis of considerations presented in reference [41] we can relate T^* used in this paper to the Flory χ parameter employed in the self-consistent field theory by the following approximate equation:

$$\chi = \frac{7.5}{T^*} \tag{28}$$

The above equation can also be used in order to relate theoretical T^* 's to experimental χ 's.

III. RESULTS

Monte Carlo simulations were carried out using the cooperative motion algorithm (CMA) for the following chain lengths: N = 32, 64, 80, 96 and 112. In each case, the ratio of segments A and B is 1 : 1. For each chain length, the number of monomers in the short A1-block is $N_{A1} = 1, 2, 3$ and 4, respectively. The SCFT calculations were performed for the corresponding parameters.

For the N = 32 chain we used the $32 \times 32 \times 32$ and $64 \times 32 \times 32$ simulation boxes, for the N = 64 chain the $64 \times 32 \times 32$ and $64 \times 64 \times 64$ boxes, for the N = 80 chain the $80 \times 40 \times 40$ box, for the N = 96 chain the $96 \times 48 \times 48$ box, and for the N = 112 chain the $112 \times 56 \times 56$ box.

We determined the order-disorder (ODT) χ 's (and thus temperatures). We also introduce interface-domain transition (IDT) χ 's as those in which half of the dangles are within B-domain and half in A-domain (interface). The χN_{IDT} 's describes quantitatively the process of migration of short A1's from B-domain to the native A-domain. χN_{IDT} in SCFT calculations are determined from short A1's density profiles in the direction normal to the layers, whereas in Monte Carlo simulations they are determined directly from the configurations. In addition we determine the χ range in which aggregation into interstitial micelles (IM) within B-domain occurs.

In Fig. 1 triangles refer to SCFT calculations, while squares and diamonds to MC results. Open symbols denote ODT, solid triangles and squares refer to IDT, while solid diamonds and dashed lines indicate the area, where the IM's are observed.

In Fig. 1(a) we can see χN_{ODT} in the function of f_{A1} (open symbols) and χN_{IDT} (closed symbols). Above χN_{ODT} , most of the short A1's are in the B-domain. As χ grows A1's share in the B-domain and A-domain gets even (IDT line). Then, for the larger χ values, the share of A1's in domain A is dominant.

In Fig. 1(b) we add the Monte Carlo simulation results, where open squares mean χN_{ODT} while solid squares refer to χN_{IDT} . We can see clearly that both qualitatively and quantitatively SCFT calculations and Monte Carlo simulations give consistent results when it comes to determining order-disorder transition and interface-domain transitions.

In Fig. 1(c), we additionally apply the information about the area in which we observe the aggregation of short A1's into micelles within the B-domain (solid diamonds and dashed lines).

We can see clearly that as f_{A1} grows (e.g. shorter chains) the temperature range of the occurrence of IM's becomes narrow but moves towards higher (more easily accessible) temperatures (lower χ 's). On the other hand, when the value of f_{A1} decreases and we deal with this by maintaining the length of A1-block while increasing the length of the chain, N, the χ range in which the aggregation occurs is wider, but shifted towards higher χ values, i.e. lower temperatures.



Fig. 1. Diagram containing information on the order-disorder transition (ODT), location of shorter A1-blocks within individual domains, and aggregation of A1-blocks into micelles for molten A1-B-A2 triblock copolymer in the function Flory parameter, χ : (a) ODT (open triangles) and interface-domain transition, IDT (solid triangles), from SCFT calculations; (b) SCFT results with Monte Carlo ODT (open squares) and IDT (solid squares) lines; (c) the area in which interstitial micelles (IM) has been observed (solid diamonds and dashed lines) against the background of results shown in (a) and (b).

In Fig. 2 we present an MC snapshot of molten 2(A1)-48(B)-46(A2) copolymer. The A-domain is marked in blue and the short A1's in yellow. B-domain is not shown for clarity purposes. Red color denotes B-segments of these chains whose A1's form one of the micelles within the B-domain. We use a recursive algorithm which allows us to determine whether a given chain forms a bridge, a loop, or is part of an IM.

Finally, it is also worthwhile underlining that the differences in results obtained by these two methods are relatively small. This can be attributed to the fact that the mean-field approach, such as the SCFT, is known to work very well in the dense polymer systems. These small differences are usually explained in terms a large coordination number, which scales \sqrt{N} , where N is large for polymers, suppressing fluctuations as argued by Fredrickson [42].



Fig. 2. Monte Carlo snapshot of 2(A1)-48(B)-46(A2) triblock copolymer melt. Domain A is marked in blue. Short A-segments (A1) marked in yellow. B-domain is not shown for clarity. The red segments are the B-segments of these chains, which form a single micelle inside the B domain.

IV. CONCLUSIONS

Using the Self-Consistent Field Theory (SCFT) we study the dual self-assembly of ABA triblock copolymers melts and compare the numerical results with those obtained by the lattice Monte Carlo simulations. While the results are qualitatively similar for both methods, the simulation times are significantly shorter for the SCFT calculations than those for the corresponding Monte Carlo simulations. It is also worthwhile underlining that the differences in results obtained by these two methods are relatively small. This can be attributed to the fact that the mean-field approach, such as the SCFT, is known to work very well in the dense polymer systems. These small differences are usually explained in terms of a large coordination number, which scales as \sqrt{N} , where N is large for polymers, suppressing fluctuations.

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