## Supporting Information:

## The influence of side-chain conformations on the phase behavior of bottlebrush block polymers

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## **Theoretical formalism of SCFT**



Figure S1: Schematic of the bottlebrush diblock polymer chain composed of side chains A (blue), B (red) and the backbone C (black). The grafting points along the backbone are labelled as  $\tau_i$ ,  $i = 1, 2, 3, \dots, m$  and the free ends are labelled as  $\tau_0$  and  $\tau_{m+1}$ , respectively.

We consider a system of volume V, which is comprised of n monodisperse and indistinguishable bottlebrush polymer chains. Each of chain has m side chains including  $m_A$  A-chains and  $m_B = m - m_A$  B-chains, which uniformly divide the C backbone into m + 1 equal segment, as illustrated in Fig.S1. We assume that each monomer on all blocks shares the same statistical length b and then occupies the same volume  $\rho_0^{-1}$ . The degree of polymerization of side chains A and B is  $N_A$  and  $N_B$ , while the degree of polymerization of backbone is  $N_C$ . Thus, the summed total segment number of a single bottlebrush polymer is  $N = m_A N_A + m_B N_B + N_C$ . The persistence length for the individual block is  $\lambda_{\alpha}$ ,  $\alpha = A, B, C$ . The volume fraction for each component is calculated by  $f_A = m_A N_A / N$ ,  $f_B = m_B N_B / N$ , and  $f_C = N_C / N$ . The *i*<sup>th</sup> grafting point on the backbone is located at  $\tau_i = i \times f_C / (m + 1)$ , where the grafting point index *i* ranges from 1 to *m*. Then, two remaining free ends of backbone are designated as  $\tau_0$  and  $\tau_{m+1}$ , respectively.

The Helmholtz free energy per chain for multicomponent block copolymers in the mean field approximation<sup>1</sup> can be written as

$$\frac{F}{nk_{\rm B}T} = -\ln Q + \frac{1}{V} \int d\mathbf{r} \left\{ \sum_{\alpha \neq \beta} N \chi_{\alpha\beta} \phi_{\alpha}(\mathbf{r}) \phi_{\beta}(\mathbf{r}) - \sum_{\alpha} \omega_{\alpha}(\mathbf{r}) \phi_{\alpha}(\mathbf{r}) - \xi(\mathbf{r}) \left[ 1 - \sum_{\alpha} \phi_{\alpha}(\mathbf{r}) \right] \right\}$$
(1)

The Flory-Huggins parameter  $\chi_{\alpha\beta}$  describes the interaction energy between dissimilar species. The average density for species  $\phi_{\alpha}(\mathbf{r})$  which is conjugated to the mean field  $\omega_{\alpha}(\mathbf{r})$  is enforced by the Lagrangian multiplier  $\xi(\mathbf{r})$  due to the incompressibility constraint to  $\sum_{\alpha} \phi_{\alpha}(\mathbf{r}) = 1$ .  $k_{\rm B}$  is the Boltzmann constant and *T* is the temperature of the system. For the wormlike chain model used here, the single chain partition function for bottlebrush copolymers can be computed by

$$Q = \frac{1}{4\pi V} \int d\mathbf{r} \int d\mathbf{u} q_{\rm C}(\mathbf{r}, \mathbf{u}, f_{\rm C})$$
(2)

The backbone propagator  $q_{\rm C}(\mathbf{r}, \mathbf{u}, t)$  represents the probability of finding the *t* terminal, starting from the free end  $\tau_0$ , which is located at a spatial position specified by  $\mathbf{r}$  and points to a direction specified by the unit vector  $\mathbf{u}$ .

In our model, the backbone is divided into m + 1 blocks. The propagator within the *i*<sup>th</sup> block is designated as

$$q_{\rm C}(\mathbf{r}, \mathbf{u}, t) = q_{\rm C}^{(i)}(\mathbf{r}, \mathbf{u}, t)$$
 here,  $\tau_i \le t < \tau_{i+1}$   $i = 0, 1, 2, \cdots, m$  (3)

which satisfies the modified diffusion equation<sup>2</sup>

$$\frac{\partial}{\partial t}q_{\rm C}^{(i)}(\mathbf{r},\mathbf{u},t) = \left[\frac{Nb}{2\lambda_{\rm C}}\nabla_{\mathbf{u}}^2 - Nb\mathbf{u}\cdot\nabla_{\mathbf{r}} - \omega_{\rm C}(\mathbf{r})\right]q_{\rm C}^{(i)}(\mathbf{r},\mathbf{u},t)$$
(4)

This equation is subject to the following initial conditions

$$q_{\rm C}^{(0)}(\mathbf{r}, \mathbf{u}, 0) = 1 \tag{5}$$

and

$$q_{\rm C}^{(i)}(\mathbf{r},\mathbf{u},\tau_i) = q_{\rm C}^{(i-1)}(\mathbf{r},\mathbf{u},\tau_i) \times \frac{1}{4\pi} \int \,\mathrm{d}\mathbf{u} q_\alpha(\mathbf{r},\mathbf{u},t=f_\alpha/m_\alpha) \quad i=1,2,\cdots,m$$
(6)

Herein, we assume that the side chain is grafted on the backbone in an orientation-independent manner. It means that side chains are grafted to the backbone with the equal probability for all directions. The species index  $\alpha$  = A,B is determined by the component of the side chain grafted at the specific point along the backbone. The propagator for the side chain satisfies

$$\frac{\partial}{\partial t}q_{\alpha}(\mathbf{r},\mathbf{u},t) = \left[\frac{Nb}{2\lambda_{\alpha}}\nabla_{\mathbf{u}}^{2} - Nb\mathbf{u}\cdot\nabla_{\mathbf{r}} - \omega_{\alpha}(\mathbf{r})\right]q_{\alpha}(\mathbf{r},\mathbf{u},t)$$
(7)

which is subject to the initial condition  $q_{\alpha}(\mathbf{r}, \mathbf{u}, 0) = 1$  for the free end.

Because the side chains grafted at two sides of the backbone are distinct, a complementary segment distribution function for the backbone  $\tilde{q}_{C}^{(i)}(\mathbf{r}, \mathbf{u}, t)$  is also required. It represents the probability of finding the *t* terminal belonging to the *i*<sup>th</sup> block along the backbone, starting from the other free end  $\tau_{m+1}$ , which is at the spatial position **r** and points in a direction specified by the unit vector  $-\mathbf{u}$ . The propagator within the  $(j-1)^{\text{th}}$  block is designated as

$$\tilde{q}_{C}(\mathbf{r}, \mathbf{u}, t) = \tilde{q}_{C}^{(j-1)}(\mathbf{r}, \mathbf{u}, t)$$
 here,  $\tau_{j-1} < t \le \tau_{j}$   $j = m+1, m, m-1, \cdots, 1$  (8)

which satisfies the modified diffusion equation

$$\frac{\partial}{\partial t}\tilde{q}_{\rm C}^{(j-1)}(\mathbf{r},\mathbf{u},t) = \left[-\frac{Nb}{2\lambda_{\rm C}}\nabla_{\mathbf{u}}^2 - Nb\mathbf{u}\cdot\nabla_{\mathbf{r}} + \omega_{\rm C}(\mathbf{r})\right]\tilde{q}_{\rm C}^{(j-1)}(\mathbf{r},\mathbf{u},t)$$
(9)

This equation is subject to the following initial conditions

$$\tilde{q}_C^{(m)}(\mathbf{r}, \mathbf{u}, t = f_C) = 1 \tag{10}$$

and

$$\tilde{q}_{\mathrm{C}}^{(j-1)}(\mathbf{r},\mathbf{u},\tau_j) = \tilde{q}_{\mathrm{C}}^{(j)}(\mathbf{r},\mathbf{u},\tau_j) \times \frac{1}{4\pi} \int \mathrm{d}\mathbf{u} q_\alpha(\mathbf{r},\mathbf{u},t=f_\alpha/m_\alpha) \quad j=m,m-1,m-2,\cdots,1$$
(11)

where the species index  $\alpha = A, B$  is determined by the component of the side chain grafted at the junction point. Similarly, the other segment distribution function for the side chain propagating

from the  $i^{th}$  junction point satisfies

$$\frac{\partial}{\partial t}\tilde{q}_{\alpha}^{(i)}(\mathbf{r},\mathbf{u},t) = \left[-\frac{Nb}{2\lambda_{\alpha}}\nabla_{\mathbf{u}}^{2} - Nb\mathbf{u}\cdot\nabla_{\mathbf{r}} + \omega_{\alpha}(\mathbf{r})\right]\tilde{q}_{\alpha}^{(i)}(\mathbf{r},\mathbf{u},t)$$
(12)

which is subject to the initial condition

$$\tilde{q}_{\alpha}^{(i)}(\mathbf{r},\mathbf{u},t=f_{\alpha}/m_{\alpha}) = \frac{\frac{1}{4\pi}\int d\mathbf{u}q_{C}^{(i-1)}(\mathbf{r},\mathbf{u},t=\tau_{i})\tilde{q}_{C}^{(i)}(\mathbf{r},\mathbf{u},t=\tau_{i})}{\left[\frac{1}{4\pi}\int d\mathbf{u}q_{\alpha}(\mathbf{r},\mathbf{u},t=f_{\alpha}/m_{\alpha})\right]^{2}}$$
(13)

Minimization of free energy functional Eq. (1) with respective of functions  $\phi_{\alpha}(\mathbf{r})$ ,  $\omega_{\alpha}(\mathbf{r})$ ,  $\xi(\mathbf{r})$ , we directly arrive at a set of saddle-point approximation equations

$$\omega_{\rm A}(\mathbf{r}) = N\chi_{\rm AB}\phi_{\rm B}(\mathbf{r}) + N\chi_{\rm AC}\phi_{\rm C}(\mathbf{r}) + \xi(\mathbf{r}) \tag{14}$$

$$\omega_{\rm B}(\mathbf{r}) = N\chi_{\rm AB}\phi_{\rm A}(\mathbf{r}) + N\chi_{\rm BC}\phi_{\rm C}(\mathbf{r}) + \xi(\mathbf{r})$$
(15)

$$\omega_{\rm C}(\mathbf{r}) = N \chi_{\rm AC} \phi_{\rm A}(\mathbf{r}) + N \chi_{\rm BC} \phi_{\rm B}(\mathbf{r}) + \xi(\mathbf{r})$$
(16)

$$\phi_{\mathrm{A}}(\mathbf{r}) + \phi_{\mathrm{B}}(\mathbf{r}) + \phi_{\mathrm{C}}(\mathbf{r}) = 1 \tag{17}$$

$$\phi_{\mathrm{A}}(\mathbf{r}) = \frac{1}{4\pi Q} \sum_{i=1}^{m_{\mathrm{A}}} \int \mathrm{d}\mathbf{u} \int_{0}^{f_{\mathrm{A}}/m_{\mathrm{A}}} \mathrm{d}t q_{\mathrm{A}}(\mathbf{r},\mathbf{u},t) \tilde{q}_{\mathrm{A}}^{(i)}(\mathbf{r},\mathbf{u},t)$$
(18)

$$\phi_{\rm B}(\mathbf{r}) = \frac{1}{4\pi Q} \sum_{i=m_{\rm A}+1}^{m} \int d\mathbf{u} \int_{0}^{f_{\rm B}/m_{\rm B}} dt q_{\rm B}(\mathbf{r},\mathbf{u},t) \tilde{q}_{\rm B}^{(i)}(\mathbf{r},\mathbf{u},t)$$
(19)

$$\phi_{\mathrm{C}}(\mathbf{r}) = \frac{1}{4\pi Q} \sum_{i=0}^{m} \int \mathrm{d}\mathbf{u} \int_{\tau_{i}}^{\tau_{i+1}} \mathrm{d}t q_{\mathrm{C}}^{(i)}(\mathbf{r},\mathbf{u},t) \tilde{q}_{\mathrm{C}}^{(i)}(\mathbf{r},\mathbf{u},t)$$
(20)

In the present work, we are mainly interested on the lamellar phase self-assembled by the symmetric bottlebrush copolymers. The system can be simplified further that only one spatial variable *z* and one orientational variable  $\theta$  are concerned. Then, Eqs.(4), (7), (9) and (12) can be numerically solved by the pseudo-spectral methods.<sup>1,3</sup> Once the propagators are calculated, we are ready to revise the external fields  $\omega_{\alpha}$  according to the newly obtained component density. In order

to accelerate the convergence, we use the Anderson mixing scheme,  $^{4,5}$  which updates the original field by incorporating the results from several preceding iterations. The free energy in Eq. (1) is used to analyze the thermodynamical stability of a specific morphology.

The degree of microphase segregation for bottlebrush copolymers strongly depends on the number of side chains *m*. For a fixed Flory-Huggins parameter  $\chi$ , a larger *m* commonly results in a stronger phase segregation. In order to estimate the degree of segregation varying with the increase of *m*, we set  $N\chi_{AB} = (m/2)(N_A + N_B)\tilde{\chi}_{AB}$ , where  $(N_A + N_B)\tilde{\chi}_{AB} = 15$  corresponding to the intermediate degree of segregation is fixed. On account of the radial stretching of side chains, we assume that the backbone is neutral to side chains, i.e.  $\chi_{AC} = \chi_{BC} = 0$ . Then, the distribution of the backbone is affected by the steric congestion of side chains and the incompressibility constraint, as a consequence of A-B segregation. The present model facilitates us to understand the impact of polymer conformations of the backbone and side chains on the self-assembly behaviors for bottlebrush block polymers.

## References

- Fredrickson, G. H. *The Equilibrium Theory of Inhomogeneous Polymers*; Oxford University Press, 2006.
- (2) Jiang, Y.; Li, S.; Chen, J. Z. The European Physical Journal E 2016, 39, 91.
- (3) Jiang, Y.; Zhang, W.-Y.; Chen, J. Z. Y. Phys. Rev. E 2011, 84, 041803.
- (4) Thompson, R. B.; Rasmussen, K. O.; Lookman, T. *The Journal of Chemical Physics* 2004, 120, 31–34.
- (5) Matsen, M. W. The European Physical Journal E 2009, 30, 361.



Figure S2: The density profiles for (a) the side chain A and (b) the backbone C in the relative length unit over one domain spacing of lamellar phase with the various grafting density  $\sigma$ . The chain flexibility parameters  $L_A/2\lambda_A = 40$  and  $L_C/2\lambda_C = 7.5$  are used for the block A and the block C, respectively.



Figure S3: The probability distribution of the backbone segments in lamellae at the given grafting density  $\sigma = 0.53$ . (a) The schematic chain conformation of bottlebrush polymers which are specifically investigated in plots (b) and (c). (b) The orientational probability distribution  $P_{\rm C}(u,t_{\rm C})$  of the backbone monomer labelled by the contour variable  $t_{\rm C}$ . (c) The positional probability distribution  $P_{\rm C}(u,t_{\rm C})$  of the backbone monomer labelled by the contour variable  $t_{\rm C}$ . (c) The positional probability distribution  $P_{\rm C}(z,t_{\rm C})$  of backbone monomers labelled by the contour variable  $t_{\rm C}$ . The chain flexibility parameters  $L_{\rm A}/2\lambda_{\rm A} = 40$  and  $L_{\rm C}/2\lambda_{\rm C} = 7.5$  are used for the block A and the block C, respectively. Note, we calculate the probability distribution for the bottlebrush molecule whose junction point is located at one specifically selected interface by imposing a restriction of a the interface-location dependence. Thus, herein, the orientation perpendicular to the interface is identified by u = -1, which is distinguishable from u = 1.



Figure S4: The interfacial width W in the relative length unit over b as the function of the chain flexibility parameter  $L_{SC}/2\lambda_{SC}$  of side chains for the given grafting density as labelled in the plot.



Figure S5: Positional and orientational probability distributions of the backbone segments specifically for three different side-chain conformations parameterized by the flexibility parameter (a)  $L_{\rm SC}/2\lambda_{\rm SC} = 50$ , (b)  $L_{\rm SC}/2\lambda_{\rm SC} = 6.67$ , and (c)  $L_{\rm SC}/2\lambda_{\rm SC} = 1.25$  at the grafting density  $\sigma = 0.47$ . The probability functions follows the same definitions in Fig. S3. The chain flexibility parameter  $L_{\rm C}/2\lambda_{\rm C} = 17$  for the backbone is used here.



Figure S6: The orientational probability distributions for different segments along the sidechain A with the various side-chain flexibility (a)  $L_{SC}/2\lambda_{SC} = 20$ , (b)  $L_{SC}/2\lambda_{SC} = 5.0$ , and (c)  $L_{SC}/2\lambda_{SC} = 3.33$  at a given grafting density  $\sigma = 0.47$ . The probability function  $P_{A,\gamma}(u;t_C)$  represents the orientational probability of finding the specific segment  $\gamma$  along the side-chain A attached to the backbone segment labelled by the contour variable  $t_C$ . The definition on  $t_C$  is same as the one in Fig. S3a. The side-chain segment index  $\gamma$  denotes the grafting point (1), middle segment (2), and free end (3) of the specified side chain. The orientational variable is defined as  $u = \cos \theta$ , where the angle  $\theta$  is graphically defined in Fig. S3a. Note, herein, the orientation perpendicular to the interface is identified by u = -1, which is distinguishable from u = 1, because the restriction of one specifically selected interface is imposed on the computation of  $P_{A,\gamma}(u;t_C)$ .



Figure S7: The positional probability distributions for two different ends of one side-chain A with the various side-chain flexibility (a)  $L_{SC}/2\lambda_{SC} = 20$ , (b)  $L_{SC}/2\lambda_{SC} = 5.0$ , and (c)  $L_{SC}/2\lambda_{SC} = 3.33$ at a given grafting density  $\sigma = 0.47$ . The probability function  $P_{A,\gamma}(z;t_C)$  represents the positional probability of finding the specific segment  $\gamma$  along the side-chain A attached to the backbone segment labelled by the contour variable  $t_C$ . The definition on  $t_C$  is same as the one in Fig. S3a. The side-chain segment index  $\gamma$  denotes the free end (solid lines) and the grafting point (dotted lines) of the specified side chain. Note, we calculate  $P_{A,\gamma}(z;t_C)$  for the bottlebrush molecule whose junction point is located at one specifically selected interface as indicated, by imposing a restriction of the interface-location dependence.