Molecular structure of bottlebrush polymers in melts

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Bottlebrushes are fascinating macromolecules that display an intriguing combination of molecular and particulate features having vital implications in both living and synthetic systems, such as cartilage and ultrasoft elastomers. However, the progress in practical applications is impeded by the lack of knowledge about the hierarchic organization of both individual bottlebrushes and their assemblies. We delineate fundamental correlations between molecular architecture, mesoscopic conformation, and macroscopic properties of polymer melts. Numerical simulations corroborate theoretical predictions for the effect of grafting density and side-chain length on the dimensions and rigidity of bottlebrushes, which effectively behave as a melt of flexible filaments. These findings provide quantitative guidelines for the design of novel materials that allow architectural tuning of their properties in a broad range without changing chemical composition.

INTRODUCTION

Significant progress in polymerization techniques allows synthesis of hyperbranched molecules with precisely controlled architectures (1–9). Dense branching results in distinct shape of individual molecules and reduces overlap of neighboring molecules in dense systems (concentrated solutions and melts). These unique features inspire the design of new materials with physical properties that are different from properties of conventional linear polymers. Branched macromolecules were explored as molecular pressure sensors (10), pH-sensitive probes (11), supersoft elastomers (12, 13), and drug delivery agents (14–16). They have also been used as components for the construction of mesoscopic systems (6) and controlling conformations of polymer chains (17).

One of the most distinct examples of highly branched macromolecules are molecular bottlebrushes composed of many polymer side chains densely grafted to a linear chain (backbone) (Fig. 1). The high grafting density results in strong steric repulsion between the side chains, causing extension of the backbone (18–21) and, in some cases, even scission of its covalent bonds (21, 22). Because of this steric repulsion, bottlebrushes adopt a wormlike conformation controlled by side-chain length and grafting density (23). In bulk melts, this conformation promotes reduction of entanglement density of the wormlike molecules (24), resulting in unusual rheological properties (25, 26) with an ultralow plateau modulus of 10² to 10³ Pa (13, 25, 27), which is much lower than the 10⁶ to 10⁹ Pa typically observed in melts of linear polymers. Note that these fundamental changes in physical properties are achieved only through architectural control without changing the chemical composition. Varying length and grafting density of side chains allows for systematic control of conformation of individual molecules as well as overlap and entanglements with neighboring molecules in dense systems.

Given their unique physical properties, molecular bottlebrushes have been an active field for many theoretical (18, 28–32), experimental (13, 27, 33–40), and numerical investigations (20, 22, 34, 36, 41–51). Most of these studies focused on basic structural properties of bottlebrushes in solutions and in the adsorbed state. Particular attention was paid to the bending rigidity of bottlebrush macromolecules, which is characterized by the persistence length $\ell_p$ and remains a matter of debate in the scientific literature. The major difficulty is the interplay between many length scales in the bottlebrush structure and their impact on $\ell_p$. Several theoretical approaches have been proposed to address this problem using scaling analysis (28, 29, 31, 52) and the self-consistent field method (49). For bottlebrushes in dilute solutions, under good solvent conditions, the persistence length was predicted to scale as $\ell_p \propto N_{sc}^{\alpha}$, with $\alpha$ as low as $\frac{3}{4}$ (28) or as high as 1.11 (49) and $\frac{15}{8}$ (29). The exponent $\alpha$ for bottlebrushes in a θ solvent was predicted to be $\frac{3}{2}$ (28) or 1.01 (49). Significantly less attention has been paid to solvent-free systems (53). Here, we address the problem of architecture-induced increase of bottlebrush persistence length as the key feature underlying physical properties of bottlebrush melts and elastomers.

In this work, we present the results of systematic coarse-grained molecular dynamics simulations and scaling analysis of the equilibrium

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Fig. 1. Molecular architecture and conformation of a bottlebrush polymer. (A) Architecture of a bottlebrush molecule consisting of a backbone with $N_{bb}$ monomers (red beads) and $z$ side chains (blue beads) per backbone monomer. Each side chain is made of $N_{sc}$ monomers. The total number of monomers of bottlebrush macromolecule is $N = N_{bb}(1 + zN_{sc})$. All beads in the simulation are considered to be identical and interact via bonded and nonbonded potential (see Materials and Methods for details). Here, $N_{bb} = 20$, $N_{sc} = 4$, and $z = 2$. (B) The bottlebrush molecule in a melt state can be represented as a chain of effective persistence segments of length $\ell_p$ and thickness $R_{sc}$. $R$ denotes end-to-end distance of bottlebrush backbone. Here, $N_{bb} = 150$, $N_{sc} = 10$, and $z = 2$. 

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structure of bottlebrush polymer melts for a range of degrees of polymerization of the backbone \(N_{bb}\), side chains \(N_{sc}\), and backbone spacer between the neighboring side chains. The latter is inversely proportional to the grafting density \(z\), which is the number of side chains per backbone monomer. We show that the persistence length \(\xi_g\), for \(z = 1\) and \(z = 2\) bottlebrushes is on the order of the size of side chains \(\langle R^2_{sc}\rangle^{1/2}\) and scales as \(\xi_g \propto \langle R^2_{sc}\rangle^{1/2} \propto N_{sc}^{-1/2}\). This finding suggests that the entanglement plateau modulus of bottlebrush melts decreases as \(G_\infty \approx \frac{1}{zV_{per}} \approx \langle R^2_{sc}\rangle^{1/2} \propto N_{sc}^{1/2}\) (Fig. 2), where \(V_{per}\) is the volume of the effective bottlebrush Kuhn segment proportional to the pervaded volume of a side chain \(V_{per} \approx \langle R^2_{sc}\rangle^{1/2}\). The pervaded volume \(V_{per}\), of a side chain is the volume of a sphere that encompasses this side chain. Our results also indicate that the backbones of bottlebrushes for \(z = 1\) and \(z = 2\) in a melt state obey Gaussian statistics with their size \(R\) (root mean square radius of gyration and end-to-end distance) scaling as \(\langle R^2\rangle^{1/2} \propto N_{bb}^{1/2} N_{sc}^{-1/4}\) for \(N_{bb} \gg N_{sc}\). Furthermore, our molecular modeling provided vital insights into the internal organization of bottlebrush melts, including limited interpenetration of side chains of neighboring molecules, radial distribution function of backbone monomers, and the form factor of individual bottlebrushes inside melt. We conclude that bottlebrush melts behave as melts of thick and flexible filaments, with a persistence length proportional to the size of the side chains.

**RESULTS**

**Scaling theory of combs and bottlebrush molecules**

Conformations of combs and bottlebrushes depend on the degree of polymerization of the side chains \(N_{sc}\) and their grafting density \(z\). Although most of the paper concentrates on bottlebrushes with \(z \geq 1\), in the present section, we consider a broader set of parameters, including loosely grafted bottlebrushes (LBs) and loosely grafted combs (LCs) with \(z < 1\). Depending on grafting density, we identify four conformational regimes of comb and bottlebrush melts (27), depicted in Fig. 2. At lower grafting density, we distinguish two comblike regimes characterized by Gaussian conformations of both backbone and side chains: (i) loosely grafted combs (LCs) with long backbone spacers between side chains \(z < 1/N_{sc}\), and with strongly interpenetrating neighboring molecules, and (ii) densely grafted combs (DCs) for \(1/N_{sc} < z < z^*\) with weak interpenetration between molecules, where \(z^*\) is defined in Eq. 1 below. There are also two regimes at higher grafting density: (iii) loosely grafted bottlebrushes (LBs) with extended backbones and Gaussian side chains for intermediate grafting density \(z^* < z < z^{**}\), and (iv) densely grafted bottlebrushes (DBs) with extended backbones and side chains for high grafting density of side chains \(z > z^{**}\), where \(z^{**}\) is defined in Eq. 3 below. The boundary between the comb and bottlebrush regimes can be found from the space-filling condition of \(zN_{sc}\) side chains with physical volume \(vN_{sc}\), each within their pervaded volume \((bn_{sc})\), resulting in reduced interpenetration of side chains from neighboring molecules

\[
z^* \approx \left(\frac{b}{v}\right)^{3/2} N_{sc}^{-1/2}
\]

where \(b\) is the Kuhn length, \(l\) is the monomer length, and \(v\) is the monomer volume. The present paper focuses on the melts of densely grafted bottlebrushes, whereas below we briefly review conformations of other types of molecules.

The low grafting density regime with \(z < z^*\) (combs) includes two subregimes: LC and DC. Loosely grafted combs (LC part of Fig. 2), with spacers between side chains longer than the side chains \(z < 1/N_{sc}\), and a high volume fraction of backbones (>50%), fully interpenetrate each other in melts. Densely grafted combs (DC part of Fig. 2), with spacers shorter than the side chains \(1/N_{sc} < z < z^*\), allow only partial interpenetration of the side chains because there is not enough space to accommodate side chains of neighboring molecules near the backbone of a host molecule. Both the side chains and backbones in melts of combs (LC and DC regimes) are in almost unperturbed Gaussian conformations.

Macromolecules with \(z > z^*\) correspond to the so-called bottlebrush regime, which onsets because of a lack of space for side chains emanating from the unperturbed Gaussian backbone. Interpenetration of these side chains without their significant deformation is only possible upon extension of the backbone. We can estimate \(z^*\) (Eq. 1) by considering a side chain with an unperturbed Gaussian size \((R^2_{sc})^{1/2} \approx (bn_{sc})^{1/2}\) and with pervaded volume \(V_{per} \approx \langle R^2_{sc}\rangle^{1/2} \approx (bn_{sc})^{1/2}\). This pervaded volume can only fit \(V_{per}/V_{sc} \approx (b/l)^3 N_{sc}^{1/2}/v\) side chains, each with a physical volume \(V_{sc} \approx vN_{sc}\). A section of the backbone of size \((R^2_{sc})^{1/2}\) passing through this pervaded volume contains \(N_{sc}\) monomers if it is in its unperturbed Gaussian conformation (assuming the same conformational statistics of backbone and side chains). Therefore, if grafting density is too high (\(z > z^*\)), the \(N_{sc}\) side chains grafted to the undeformed section of the backbone with combined physical volume \(vN_{sc}^2\), \(z > z^*\) can no longer fit in the pervaded volume \(V_{per}\), forcing the backbone to extend.

The backbone extension on the length scale \((R^2_{sc})^{1/2}\) assures a fixed number of grafting points along the backbone section of this size \((R^2_{sc})^{1/2} \approx (bn_{sc})^{1/2}\) equal to the number of overlapping side chains \((R^2_{sc})^{1/2}/(vN_{sc}) \approx (b/l)^3 N_{sc}^{1/2}/v\). On the small length scales, up to the size of the tension blob \((5d)\), the backbone remains unperturbed. The size of the tension blob \(\xi \approx (bl)^{1/2}\) consisting of \(g\) monomers is estimated from the condition that \(gz\) side chains emanated from this section of the backbone densely fill its pervaded volume \(\xi^3\), there are \(\xi^3/\langle V\rangle \approx gz\) such overlapping chain sections. Therefore, the tension blob size is \(\xi \approx (bl)^{1/2}/(vz)\). There is no crowding issue on length scales \(r\) smaller than the tension blob \((r < \xi)\), and bottlebrush backbones maintain the unperturbed Gaussian conformations with bare Kuhn length \(b\). On the intermediate length scales \((\xi < r < (R^2_{sc})^{1/2})\),

![Fig. 2. Diagram of states of combs and bottlebrush molecules.](image-url)
a backbone can be visualized as an extended array of tension blobs with a constant average distance between grafting points $v/(bl).$ On larger length scales ($r > (R_{sc}^2/0)^{1/2}$), backbones of bottlebrushes in a melt are represented as random walks of these extended arrays of tension blobs. The loosely grafted bottlebrush (LB part of Fig. 2) is described as a thick filament with contour length $L \approx N_{bb}vz/(bl)$, thickness equal to the end-to-end distance of its side chains $(R_{sc}^2/0)^{1/2}$, and persistence length on the same order of magnitude (see detailed derivation in the subsection “Persistence length of a bottlebrush in a melt”). Thus, bottlebrush macromolecules are considered as chains of $L/(R_{sc}^2/0)^{1/2}$ effective monomers of size $(R_{sc}^2/0)^{1/2}$. 

The filament-like bottlebrush with both thickness and persistence length on the order of $(R_{sc}^2)^{1/2}$ and bottlebrush contour length on the order of the contour of the backbone $L_{bb}$ has mean square size 

$$
\langle R^2 \rangle \approx \left( v z N_{sc} \right)^{1/2} \text{ for } z > \left( \frac{p b}{v} \right) \text{ if } v < b^2 l \text{ or }
$$

$$
\text{for } z > \left( \frac{p b}{v} \right) \sqrt{v} \text{ if } v > b^2 l \text{ (6)}
$$

The dependence of backbone and side-chain size of combs and bottlebrushes on $z$ is summarized in Fig. 3.

The size of side chains of densely grafted bottlebrushes with almost fully stretched backbones

The size of side chains increases with degree of polymerization $N_{sc}$ (see fig. S1 and table S1). Their size also increases with grafting density $z$ along the backbone. This effect is illustrated in Fig. 4A, which exhibits the variation of the ratio of the mean square distance $(R_{sc}^2(s))$ of side-chain monomer $s$ from the grafting point and the corresponding Gaussian size $sv^2$ as a function of the bond index $s$ for different grafting densities $z$. Different colors and symbols correspond to bottlebrushes with different values of $N_{bb}$, $N_{sc}$, and $z$, as shown in Fig. 4B and table S2. To understand the bond index $s$ dependence of the mean square distance $(R_{sc}^2(s))$, we consider the average of the square of the size $R_{sc}(s) = \langle R_{sc}(s) \rangle + \delta R_{sc}(s)$ of these side-chain segments containing $s$ monomers

$$
\langle R_{sc}^2(s) \rangle = \langle R_{sc}(s) \rangle^2 + \langle \delta R_{sc}^2(s) \rangle
$$

We assume that the nontrivial $s$ dependence of $\langle R_{sc}^2(s) \rangle$ observed in Fig. 4A is due to chain extension $\langle R_{sc}(s) \rangle$, whereas the fluctuations $\langle \delta R_{sc}^2(s) \rangle$ of the size of these $s$-segments can be described by the mean square size of chain sections containing $s$ monomers of a free linear 16-mer ($z = 0$, red crosses).

For monomers near the free ends of side chains, the mean distance $\langle R_{sc}(s) \rangle$ can be expanded in the Taylor series of the variable $1 - s/N_{sc}$:

$$
\langle R_{sc}(s) \rangle = \langle R_{sc} \rangle \left[ 1 + \sum_{n>0} a_n \left( 1 - \frac{s}{N_{sc}} \right)^n \right]
$$

The backbone is almost fully stretched in the case of lower monomer volume $v < b^2 l$ if $z > z^{**} = (p b)^3/v^2$, or, for higher grafting density, $z > (p b^2/v)^{1/2}$ in the case of higher monomer volume $v > b^2 l$. In this case, the dense packing of side chains forces them to extend to the mean square size

$$
\langle R_{sc}^2 \rangle \approx \frac{v z}{l} N_{sc} \text{ for } z > \left( \frac{p b}{v} \right) \text{ if } v < b^2 l \text{ or }
$$

$$
\text{for } z > \left( \frac{p b}{v} \right) \sqrt{v} \text{ if } v > b^2 l \text{ (5)}
$$

The filament-like bottlebrush with both thickness and persistence length on the order of $(R_{sc}^2)^{1/2}$ and bottlebrush contour length on the order of the contour of the backbone $L_{bb}$ has mean square size
where \( \langle R_{sc}(s) \rangle = \langle R_{sc}(N_{sc}) \rangle \) is the average size of a side chain. The first coefficient is \( a_1 = 0 \) due to the boundary condition \( dR_{sc}/ds = 0 \) at the free end \( s = N_{sc} \). The condition \( \langle R_{sc}(s) \rangle \ll \langle R_{sc} \rangle \) for small \( s \ll N_{sc} \) leads to the constraint for the sum of all coefficients \( \Sigma a_n = -1 \). Note that the asymptotic expressions for the higher-order coefficients \( a_n \) in Eq. 8 can be found by expanding the dependence \( \langle R_{sc}(s) \rangle \ll \langle R_{sc} \rangle \) for small \( s \ll N_{sc} \) in the power series of \( (1 - s/N_{sc}) \). Comparing this expansion with expansion in Eq. 8 term by term, we estimate \( a_1 \approx -1/16 \) and coefficients \( a_n \) decay with \( n \) as \( n^{-3/2} \). The small values of these coefficients justify omission of the higher-order terms in the expansion in Eq. 8. Thus, we take all \( a_{s,2} = 0 \) and \( a_2 = -1 \) and obtain

\[
\langle R_{sc}(s) \rangle \approx \frac{s}{N_{sc}} \left( 2 - \frac{s}{N_{sc}} \right) \langle R_{sc} \rangle \quad \text{for} \quad \gamma N_{sc} \leq s \leq N_{sc} \tag{9}
\]

The parameter \( \gamma \sim 0.3 \) to 0.5 defines the lower boundary of the interval of validity of the above approximation. The mean square size of linear chain segments containing \( s \) monomers \( \langle \delta R_{sc}^2(s) \rangle \) obtained from molecular dynamics simulations is presented by the lowest set of points denoted by \( \times \) symbols in Fig. 4A. This dependence can be approximated by

\[
\langle \delta R_{sc}^2(s) \rangle = s \sigma^2 C_{lin} \left( 1 + s/N_{sc} \right) \tag{10}
\]

with the data obtained from simulations of bottlebrushes with grafting density \( z = 1, 2, \) and 4 using single fitting parameter \( \langle R_{sc} \rangle \) and the value of \( C_{lin} = 1.55 \) and \( \tilde{z} = 0.61 \) from the fit to linear chain data \( (z = 0) \). This simple estimate (Eq. 10) demonstrates excellent agreement with the simulation data.

The average side-chain size \( \langle R_{sc} \rangle \) can be estimated from the monomer dense packing condition. The transverse slice of a bottlebrush can be approximated by a disc of volume \( dR_{sc}^2 \) and thickness \( d \approx \sigma \) of the backbone bond projection onto the contour of the molecule. Assuming that there is no (or limited) overlap between the side chains of neighboring bottlebrushes, the disc volume is occupied by \( \nu \) side chains of volume \( v N_{sc} \) each, where \( \nu \approx 3^{3/2} \) is the volume of one monomer. Therefore, the square of the average size of side chains can be estimated as

\[
\langle R_{sc}(s) \rangle^2 \approx \frac{v N_{sc} \sigma^2}{d} = C_{sc} N_{sc} \tilde{z} \sigma^2 \tag{11}
\]

where \( C_{sc} \) is the numerical coefficient accounting for the scaling form of this expression. The inset in Fig. 4A shows good agreement with Eq. 11, with the value of the fitting parameter \( C_{sc} = 0.17 \).

Combining Eqs. 9 and 11, we can write

\[
\langle R_{sc}(s) \rangle = C_{sc}^{1/2} \left( 2 - \frac{s}{N_{sc}} \right) \frac{\sigma \tilde{z}^{1/2}}{N_{sc}^{1/2}} \quad \text{for} \quad \gamma N_{sc} \leq s \leq N_{sc} \tag{12}
\]

In Fig. 4C, we test this prediction by plotting the \( s \)-dependence of the ratio of the average distance \( \langle R_{sc}(s) \rangle = \langle R_{sc}(s) \rangle^2 - \langle \delta R_{sc}^2(s) \rangle \) and \( s \sigma \tilde{z}^{1/2} \) using the simulation data presented in Fig. 4A. For larger values of \( s \) for \( \gamma N_{sc} \leq s \leq N_{sc} \), this rescaled function is \( z \)-independent and exhibits linear dependence on \( s \langle R_{sc}(s) \rangle / (s \sigma \tilde{z}^{1/2}) = C_{sc}^{1/2} (2 - s/N_{sc}) / N_{sc}^{1/2} \), with \( N_{sc} \)-dependent negative slope predicted by Eq. 12. The red and black dashed lines in Fig. 4C (for \( s > 6 \)) have slopes of \(-0.012\) and \(-0.0064\) for \( N_{sc} = 10 \) and \( N_{sc} = 16 \), respectively, which are consistent with the predicted negative slopes \(-C_{sc}^{1/2} N_{sc}^{-1/2} = -0.013 \) and \(-0.0063\) from Eq. 12.
The mean square fluctuations of the size of chain segments containing large number $s \gg 1$ of monomers are Gaussian. Therefore, the normalized mean square size of side-chain segments (Eq. 10) can be approximated for large $s$ by

$$\frac{\langle R_{sc}^2(s) \rangle}{s^2 \sigma^2} = C_{sc} \frac{2s}{N_{sc}} (2 - \frac{s}{N_{sc}})^2 + C_{lin} \frac{1 + s}{1 + 3/s}$$

$$\approx C_{sc} \frac{2s}{N_{sc}} (2 - \frac{s}{N_{sc}})^2 + C_{lin} \text{ for } \gamma N_{sc} \leq s \leq N_{sc}$$

This equation predicts a maximum at $s_{max} = 2N_{sc}/3$. This prediction is in good agreement with the simulations (see Fig. 4A). Note that the position of the maximum (for both points and lines) has a slightly higher value of $s$ than $2N_{sc}/3$ because of the residual $s$ dependence of the ratio $\langle \delta R_{sc}^2(s) \rangle/(s \sigma^2)$ for short side chains. The physical explanation of this peak is that not all of the chains extend all the way to $R_{sc}$. There is a wide distribution of the positions of side-chain ends around their average value ($R_{sc}$), because fewer side chains tend to extend to larger radial distances from the backbone, they provide an additional contribution to the ratio $\langle R_{sc}^2(s) \rangle/(s \sigma^2)$ in Eq. 13 for $\gamma N_{sc} < s < s_{max}$ and a relatively smaller contribution for larger values of $s > s_{max}$. As a result, side chains that do not extend to large radial distances from the backbone, the crowding of remaining side chains at these large radial distances decreases. This decrease in crowding weakens the stretching of the remote side-chain sections, resulting in a relatively smaller average extension of chain sections with $s > 2N_{sc}/3$. The stretching decreases with $s$ and vanishes at the free side-chain ends in the overlapping zone of neighboring bottlebrushes.

Conformations of side-chain segments with $s \leq \gamma N_{sc}$ near the grafting point are determined by the monomer packing condition due to the limited penetration of monomers with index $s' > s$ into this zone near the backbone, similar to packing restrictions for the entire side chain (see Eq. 11)

$$\langle R_{sc}(s) \rangle^2 = \frac{zv}{d} \approx z \sigma^2 \text{ for } s \leq \gamma N_{sc}$$

Therefore, the ratio $\langle R_{sc}(s) \rangle/(s \sigma^2)^{1/2}$ should be independent of $N_{sc}$ for small $s \leq \gamma N_{sc}$, as observed in Fig. 4C. However, note that the $s$ dependence of $\langle R_{sc}(s) \rangle$ for $s \leq 6$ differs from our prediction (Eq. 14) because of strong crowding of side-chains near the backbone and the non-Gaussian behavior of these short chain segments. The $s$ dependence of $\langle \delta R_{sc}^2(s) \rangle/(s \sigma^2)$ significantly deviates from a constant for $s \leq 6$ (see red crosses in Fig. 4A), as described by the crossover expression $C_{lin} \approx (1 + 3/s)$.

**Persistence length of a bottlebrush in a melt**

The rigidity of bottlebrush is only due to the mutual repulsion of the crowded side chains. The excluded volume interactions in a melt state are highly screened. In the unrealistic case of complete screening of steric interactions, the resulting persistence length of a bottlebrush is on the order of its monomer size $\approx \sigma$. To estimate the persistence length of "real" bottlebrushes, we have to account for partially screened excluded volume interactions between side chains. The physical volume of $s \sigma$ side chains grafted to a persistent bottlebrush section is $v_{sp} z N_{sc}$ and the radius of this section is $R_{sc}$. Therefore, the length of this cylindrical-like section is $v_{sp} zN_{sc}/(R_{sc}^2)$ and its pervaded volume is $\approx (v_{sp} z N_{sc}/(R_{sc}^2))^3$. The pervaded volume of a persistent bottlebrush section is the volume of a sphere that encompasses this cylindrical-like persistent segment. The excluded volume interactions between polymer sections in the melt are reduced by the degree of polymerization $P_w = s_p z N_{sc}$ of these sections (54) (see Fig. 5). Thus, the free energy of the excluded volume interactions between these persistent bottlebrush sections within their pervaded volume is

$$E_{sc} = k_B T \langle v \rangle \langle z v N_{sc} \rangle^2$$

$$P_w (v_{sp} z N_{sc} / (R_{sc}^2))^3$$

The persistent segment $s_p$ is determined by the condition that the excluded volume interaction energy $E_{sc}$ is on the order of thermal energy $k_B T$, resulting in

$$s_p \approx \langle R_{sc}^2 \rangle_{\gamma N_{sc}}^{1/2} \approx \left( \frac{z v N_{sc}}{d} \right)^{1/2}$$

where $\langle R_{sc}^2 \rangle_{\gamma N_{sc}}^{1/2} \approx \left( \frac{z v N_{sc}}{d} \right)^{1/2}$ was used. In this case, the size of the persistence segment $\ell_p$ is

$$\ell_p \approx s_p d \approx \left( \frac{z v N_{sc}}{d} \right)^{1/2} \approx \langle R_{sc}^2 \rangle_{\gamma N_{sc}}^{1/2}$$

The conformations of bottlebrush backbones at small length scales are similar to those of flexible polyelectrolytes that are almost undeformed on scales up to electrostatic blob size but extended into a linear array of electrostatic blobs on larger length scales with persistence length determined by the screening length (55). By analogy with the polyelectrolytes, bottlebrushes are flexible on small length scales and have large persistence length, induced by side-chain repulsion, on intermediate length scales.

Our simulations confirm the scaling prediction that the persistence length of bottlebrush backbones in a melt state is comparable to the size of side chains. To determine the length of persistence segments $s_p$.

**Fig. 5. Geometry of a bottlebrush polymer.** A bottlebrush is composed of $z$ side chains with $N_{sc}$ monomers each grafted to every backbone monomer ($x = 2$ in this figure). $R_{sc} = R_{sc}(N_{sc})$ and $R_{sc}(s)$ denote instantaneous values of size of side chains (bottlebrush thickness) and distance of a side-chain monomer $s$ from the grafting point, respectively. The number of monomers per persistence segment is $s_p$ and persistence length is $\ell_p$. $P_w = s_p z N_{sc}$ is the total degree of polymerization of cylindrical-like section composed of $s_p$ backbone monomers and $s_p z N_{sc}$ side-chain monomers. $d$ is average projection of a backbone bond onto the direction of the backbone contour.
The data obtained from simulations (symbols) have been fitted to the expression for the decay of bond orientational correlations plotted as a function of the side-chain and number of side chains grafted per backbone monomer (see Fig. 4B and the corresponding caption for the definition of symbols). Dashed lines represent best fits to the expression for g(s) given by Eq. 18. (B) Persistence segments obtained from the decay of bond orientational correlations plotted as a function of the side-chain polymerization degree N_{sc} for various backbones N_{bb} and grafting densities z of side chains, as indicated. The dashed lines represent the best power-law fit for data sets with N_{bb} = 100: s_{p} = 1.91 N_{sc}^{0.53±0.04} for z = 1 and s_{p} = 2.75 N_{sc}^{0.48±0.03} for z = 2.

we calculated the decay rate of the correlations between bond orientations g(s) ≈ (cos θ)^\(s^{3/2}\) (see simulation data in Fig. 6A and fig. S2). The cosine of the angle is cos \(θ = (r_i/|r_i|) \cdot (r_{i+1}/|r_{i+1}|)\), where \(r_i = R_{i+1} - R_i\) is the bond vector between monomers \(i\) and \(i + 1\), and \(r_{i+1} = R_{i+2} - R_i\) is the bond vector between monomers \(i + 1\) and \(i + 2\). The data obtained from simulations (symbols) have been fitted to the function (dashed lines) of the following form

\[
g(s) = (1 - A)e^{-s/\xi} + Ae^{-s/z} + \xi s^{-3/2} \tanh [(s/3)^2]
\]

where A, s_{0}, s_{p}, and \(ξ\) are fitting parameters (see table S3) and denote the following: (1 - A) and A are the magnitudes of short- and long-range correlations, respectively, between bond vectors of the backbone; \(s_{0}\) is the characteristic number of beads in the backbone, which undergo "local" bond vector correlations; \(s_{p}\) is the number of beads per persistence segment; and \(ξ\) is the magnitude of long-range inter-actions induced by the connectivity of backbone bonds (see the section on persistence length in the Supplementary Materials for details). The power-law decay \(s^{1/2}\) at \(s \gg 1\) was reported for polymers melts (56) and for \(θ\) solutions (57). The origin of these interactions is explained either by effective compression of polymer coils due to the correlation hole effect (56) or by the shift of the monomeric Mayer f-function due to the finite interaction range and chain connectivity (57). The function \(\tanh[(s/3)^2]\) presents in Eq. 18 describes a cutoff at minimal loop size \(s = 3\).

In Fig. 6B, the number of monomers along the bottlebrush backbone in the persistent segment \(s_{p}\) is presented as a function of the degree of polymerization of side chains \(N_{sc}\) for grafted densities \(z = 1\) and \(2\) and various backbone degrees of polymerization \(N_{bb}\). These data demonstrate the effect of backbone straightening with the increasing \(N_{sc}\) and \(z\). The exponents of the observed power laws agree with the exponent \(1/2\) predicted from the scaling arguments, as presented in the beginning of this subsection (see Eq. 16), and indicate that the number of monomers \(s_{p}\) in the persistence segments of the bottlebrush backbones with \(z = 1\) or \(2\) chain per backbone monomer and \(N_{bb} \gg 50\) is proportional to the brush radius (side-chain size \(R_{sc}^{z}\)) (see fig. S3).

**Size of a bottlebrush in a melt**

The size of a bottlebrush in a melt state can be estimated from a simple physical picture of nonoverlapping flexible filaments with \(ξ_{p} \approx (R_{sc}^{z})^{1/2}\). Consider a bottlebrush backbone that is much longer than persistence segment \(N_{bb} \gg s_{p}\) (see Fig. 1B). The bottlebrush conformation can be represented as a chain of effective monomers consisting of \(s_{p}\) backbone monomers and \(s_{c}\) side chains of size \(〈R_{sc}^{z}〉^{1/2}\). The mean square end-to-end distance of the backbone can be estimated as the number of these effective monomers \(N_{bb}/s_{p}\) times the square of their size, which is proportional to the mean square size of side chains \(〈R_{sc}^{z}〉\), resulting in \(〈R^{2}〉 \approx (N_{sc}/s_{p}) 〈R_{sc}^{z}〉\). Because the number of monomers in a persistence segment \(s_{p}\) is proportional to the size of a side chain \(〈R_{sc}^{z}〉\) (see Eqs. 17, 19, and 20), we predict that the mean square size of the backbone is also proportional to the size of a side chain \(〈R^{2}〉 \approx \sigma (N_{sc}/s_{p}) 〈R_{sc}^{z}〉\). Our simulation results corroborate this prediction (cf. Eq. 6), as discussed below.

The mean square end-to-end distance of the backbone \(〈R^{2}〉\) and the mean square radius of gyration \(〈R_{g}^{2}〉\) of the whole bottlebrush are plotted as functions of the degree of polymerization of side chains \(N_{sc}\) and backbone \(N_{bb}\) in Fig. 7 and fig. S4 (for the definition of symbols, see Fig. 4B and the corresponding caption as well as table S2). The mean square size increases with \(N_{bb}\) and \(N_{sc}\) and obeys the power law

\[
\frac{〈R^{2}〉}{N_{bb}} = 1.33 N_{sc}^{0.47±0.01} + 1\quad \text{for}\quad z = 1
\]

\[
\frac{〈R^{2}〉}{N_{bb}} = 2.76 N_{sc}^{0.47±0.01} + 1\quad \text{for}\quad z = 2
\]

and

\[
\frac{〈R_{g}^{2}〉}{N_{bb}} = 0.23 N_{sc}^{0.49±0.01} + 1\quad \text{for}\quad z = 1\quad \text{a nd}\quad \frac{〈R_{g}^{2}〉}{N_{bb}} = 0.46 N_{sc}^{0.47±0.01} + 1\quad \text{for}\quad z = 2.
\]

Similar to linear chains in melts, the
conformations of bottlebrushes with long backbones are well described by the ideal chain statistics, that is, \( \langle R^2 \rangle / \langle s^2 \rangle \approx 6 \), as verified by the distributions of the end-to-end distances \( R \) and the radii of gyration \( R_g \) (cf. fig. S5). From Fig. 7A, one can observe that the mean square end-to-end distance of molecules with backbones \( N_{bb} = 10 \) and 20 becomes insensitive to further increase in \( N_{sc} \). This is attributed to the crossover from the crew-cut bottlebrushes to the starlike configurations.

The mean square internal distances \( \langle R^2(s) \rangle \) between backbone monomers are plotted in Fig. 8 as a function of the number of bonds \( s \) in a backbone section for molecules with various side-chain degrees of polymerization \( N_{sc} \) and different number \( z \) of side chains grafted per backbone monomer (the definition of symbols is displayed in Fig. 4B and described in the corresponding caption). The simulation data for all values were fitted to the crossover expression

\[
\langle R^2(s) \rangle = s \sigma^2 \frac{C_{bb}}{1 + \tilde{s}/s}
\]

with fitting parameters \( C_{bb} \) and \( \tilde{s} \) plotted in Fig. 8B and reported in table S4. The overall good agreement between the crossover expression and simulation data is observed by the overlap between points (simulation data) and lines (Eq. 22) in Fig. 8A. The backbone stretching (characterized by both parameters \( C_{bb} \) and \( \tilde{s} \)) increases significantly with the grafting density of side chains \( N_{sc} \) and with the degree of polymerization of side chains \( N_{sc} \) because of steric repulsion between densely grafted side chains. The Flory characteristic ratio \( C_{bb} \) and parameter \( \tilde{s} \) increase proportionally to the power of the degree of polymerization of side chains (Eq. 22) and \( \tilde{s} = 2.1 N_{sc}^{0.51 \pm 0.01} \) for \( z = 1 \) and \( C_{bb}^{0.53 \pm 0.01} \) and \( \tilde{s} = 3.4 N_{sc}^{0.55 \pm 0.01} \) for \( z = 2 \) (see Fig. 8B and table S4). The scaling exponents are close to \( 1/2 \), indicating that \( s \) dependence of distances between backbone bonds separated by \( s \) monomers for \( s > \tilde{s} \) is consistent with the backbone size scaling (Eq. 21). We observe that \( \tilde{s} \) is 60% and 25% larger than corresponding \( C_{bb} \) values (see

**Fig. 7. Size of bottlebrushes in a melt.** Mean square end-to-end distance \( \langle R^2 \rangle \) (A) and mean square radius of gyration \( \langle R_g^2 \rangle \) (including side chains) (B) of bottlebrushes in a melt normalized by the ideal mean square size of backbones \( N_{bb} \sigma^2 \) as functions of the degree of polymerization of side chains \( N_{sc} \). (C) Mean square radius of gyration \( \langle R_g^2 \rangle \) of bottlebrushes normalized by the ideal mean square size of side chains \( N_{sc} \sigma^2 \) as a function of the degree of polymerization of backbones \( N_{bb} \). See Fig. 4B and the corresponding caption for the definition of symbols. In (A) and (B), the number of side chains grafted per backbone monomer is \( z = 1 \) and 2. (C) displays data for \( z = 2 \). Dashed lines represent fitted scaling laws: (A) \( 1.33 N_{sc}^{0.47 \pm 0.01} \) for \( z = 1 \) and \( N_{bb} = 100 \) and \( 2.76 N_{sc}^{0.47 \pm 0.01} \) for \( z = 2 \) and \( N_{bb} = 100 \). (B) \( 0.23 N_{bb}^{0.49 \pm 0.01} \) for \( z = 1 \) and \( N_{bb} = 100 \) and 0.46\( N_{sc}^{0.42 \pm 0.01} \) for \( z = 2 \) and \( N_{bb} = 100 \), and (C) 0.39\( N_{sc}^{0.41 \pm 0.04} \) for \( z = 2 \) and \( N_{sc} = 10 \). The error bars for all data points are smaller than the size of symbols.

**Fig. 8. Extension of a bottlebrush in a melt.** (A) Dependence of the mean square internal distances \( \langle R^2(s) \rangle / (s \sigma^2) \) between bottlebrush backbone monomers normalized by their ideal mean square size \( s \sigma^2 \) on the number of monomers \( s \) in the backbone sections plotted for various side-chain polymerization degrees \( N_{sc} \) and grafting densities \( z \) (see Fig. 4B and its caption for the definition of symbols). Solid lines represent best fits to the crossover expression \( \langle R^2(s) \rangle / (s \sigma^2) = C_{bb}^{0.55 \pm 0.01} \), with fitting parameters \( C_{bb}^{0.55 \pm 0.01} \) and \( \tilde{s} \) listed in table S4. (B) Fitting parameters \( C_{bb}^{0.55 \pm 0.01} \) and \( \tilde{s} \) plotted as a function of side-chain polymerization degree \( N_{sc} \) for \( z = 1 \) (full symbols) and \( z = 2 \) (open symbols). Dashed lines represent fitted scaling laws: \( C_{bb}^{0.55 \pm 0.01} = 1.3 N_{sc}^{0.52 \pm 0.01} \) and \( \tilde{s} = 2.1 N_{sc}^{0.51 \pm 0.01} \) for \( z = 1 \), whereas \( C_{bb}^{0.55 \pm 0.01} = 2.8 N_{sc}^{0.53 \pm 0.01} \) and \( \tilde{s} = 3.4 N_{sc}^{0.55 \pm 0.01} \) for \( z = 2 \).
Fig. 8B and table S4) for \( z = 1 \) and 2, respectively, pointing out the wiggling of the backbone \( \langle R^2(s) \rangle^{1/2} \approx s^{1/2} \delta \) on length scales smaller than the tension blob. The bottlebrushes undergo a conformational transformation from a random coil to a rod as the grafting density \( z \) increases from 1 to 4, which is evidenced by the increase of both the Flory characteristic ratio \( C_{bb}^{0} \) and the crossover value \( \delta \) by more than an order of magnitude. Because the backbone degree of polymerization \( N_{bb} \ll \delta \), the simulated bottlebrush with side-chain grafting density \( z = 4 \) is effectively rodlike.

Note that it is hardly possible to determine persistence segment \( s_p \) from correlations of bond orientations (see Eq. 18) for simulated bottlebrushes with short backbones \( N_{bb} \leq 50 \) and \( z = 4 \) because the corresponding \( g(s) \) functions do not decay sufficiently.

Interpenetration of neighboring bottlebrushes in a melt

As discussed below, reduced interpenetration of side chains from neighboring macromolecules is a distinct feature that distinguishes segregated filaments (bottlebrushes) from overlapped molecules (linear chains and combs). This interpenetration is crucial for the understanding of the friction between these molecules and for the explanation of the complex rheological response of bottlebrush-based materials observed in recent experiments (12, 26, 27, 58, 59). We have analyzed the number of intramolecular \( h_s \) and intermolecular \( g_s \) contacts between side-chain monomers. Contacts were defined between pairs of monomers within distance \( r_{shell} = 1.5\sigma \) from each other. This separation corresponds to the position of the minimum in the interbead correlation function \( g(r) \), as discussed below. We denote by \( \langle h_s \rangle \) the average number of intramolecular nonbonded contacts between \( s \)th monomer of a side chain and monomers within the same host molecule, whereas \( \langle g_s \rangle \) stands for the average number of intermolecular contacts between the \( s \)th monomer of a side chain of a given molecule with monomers belonging to all other molecules.

Both quantities were calculated as functions of monomer index \( s \) along a side chain and normalized by the average number of nonbonded neighbors per \( s \)th monomer \( Z_s = \langle h_s \rangle + \langle g_s \rangle \). The average value of \( Z_s \) is independent of the monomer index \( s \), \( Z_s \approx \langle Z \rangle \approx 4.7, \) except for terminal monomers \( Z_1 \approx 5.0 \) and \( Z_{N_{sc}} \approx 5.3 \). The results of this analysis are presented in Fig. 9A (for the definition of symbols, see Fig. 4B and the corresponding caption as well as table S2). The average fraction of intermolecular contacts for the first side-chain monomers \( (s = 1) \) is low and decreases with increasing grafting density \( z \) of \( \langle g_s \rangle / \langle Z_s \rangle \approx 0.2 \) for \( z = 1 \), \( \langle g_s \rangle / \langle Z_s \rangle = 0.1 \) for \( z = 2 \), and \( \langle g_s \rangle / \langle Z_s \rangle < 0.01 \) for \( z = 4 \). The limited interpenetration is ascribed to dense crowding of the side-chain monomers belonging to the same bottlebrush in the vicinity of its backbone, which hinders penetration of guest monomers to the central region of the host bottlebrush. The probability of encountering guest monomers \( \langle g_s \rangle / \langle Z_s \rangle \) increases with \( s \) and reaches the maximum value for terminal monomers of the side chains. This maximum value was found to be \( \langle g_{N_{sc}} \rangle / \langle Z_{N_{sc}} \rangle \approx 1/2 \) independent of grafting density \( z \) and degree of polymerization of side chains \( N_{sc} \). The terminal monomers are in the interpenetration zone between two neighboring bottlebrushes. This zone contains 50:50 composition of monomers from both molecules. The low interpenetration of bottlebrushes in a melt state is demonstrated by the "territorial map" (60–62) of the simulation box snapshot in Fig. 9B.

For a reference system, we have calculated the number of contacts between monomers of neighboring linear chains in a melt (cf. the inset of Fig. 9A). The interchain contacts for linear polymers are almost uniformly distributed along the backbone except for the chain ends. The estimated fraction of guest monomers for linear chains was found to be \( \langle g_s \rangle / \langle Z_s \rangle \approx 0.65 \) (independent of the degree of polymerization) and is higher than the maximum fraction of 0.5 (encountered by end monomers of bottlebrush side chains). The increase in the fraction of intermolecular contacts of up to \( \approx 0.78 \) is observed at the ends of linear chains. The intrachain nonbonded contacts in linear chains are due to the formation of self-loops. For inner monomers of a linear chain, one can have self-loops at both sides, whereas for end monomers, self-loops can only be formed from one side. This explains the higher number of interchain contacts for end monomers of linear chains in a melt. The main conclusion of this analysis is that the overlap of bottlebrush molecules in the melt is qualitatively different on a monomer level from that of linear chains. Below, we show that melts of bottlebrushes become qualitatively similar to those of linear chains if we describe bottlebrushes as thick flexible filaments, which are chains of "effective monomers" of size \( \langle R^2_s \rangle^{1/2} \).
As the degree of polymerization of side chains monomers in a melt various degrees of polymerization of side chains results for bone monomers. As indicated in the legend, various colors are used to distinguish between the lines with different values of grafting density ρg. In melts of bottlebrushes with short side chains: g intra(r) ≈ r for ρg = 1, which is the optimal distance between pairs of nearest-neighbor monomers. The second peak at r ≈ 1.9 σ displays the influence from the shell of the second-nearest neighbors. In contrast to melts of linear chains, the large-scale structure is observed in bottlebrush melts. Bottlebrush backbones are shielded by densely grafted side chains and do not approach each other. This fact is clearly demonstrated in the case of bottlebrushes with the long side chains, that is, Nsc = 10 (solid red line for z = 2 and dashed red line for z = 4). At distances comparable to bottlebrush thickness, equal to the average side-chain size ⟨Rsc2⟩ ≈ 4.18 σ for z = 2 and at ⟨Rsc2⟩ ≈ 4.58 σ for z = 4, the corresponding values of g intra(r) are low g intra(r)/ρbb ≈ 0.1 because of the "coat" of side chains around a given backbone prevents neighboring backbones from approaching it. Note that the highest probability to find monomers of neighboring backbones does not exactly match the brush diameter 2⟨Rsc2⟩1/2 but is rather at a shorter distance because of partial interpenetration of side chains. The first maxima are observed at rmax ≈ 7.25 σ for z = 2 and ≈ 8.35 σ for z = 4, respectively. The smaller value of rmax/(2⟨Rsc2⟩1/2) for z = 2 (≈ 0.87) with respect to z = 4 (≈0.91) implies a stronger overlap of grafted side chains in bottlebrush melts with lower z (see Fig. 9). The inset of Fig. 10B shows the correlation functions g intra(r), with the abscissa rescaled by the position of the first maximum rmax. The positions of the peaks for bottlebrush systems correlate very well with the peaks observed for linear chain, indicating a similar "liquid-like" origin of melt structure (63, 64) but with stronger correlations at correspondingly larger distances. This result justifies representing bottlebrushes in melts by chains of "effective" monomers of size ≈ (⟨Rsc2⟩)1/2, which are thick flexible filamentous objects.

The form factor of a bottlebrush backbone in a melt
The form factor S(q) of the backbones of bottlebrushes with different grafting densities z of side chains is presented as the Holzer plot in Fig. 11. The form factor of linear chains (z = 0) in a melt state (solid black line) is well represented by the Debye function (black dashed line) and scales as S(q) ∝ q−2 for large wave vectors q. The backbone form factors of bottlebrushes exhibit different behavior because of the backbone extension induced by side chains. For bottlebrushes with z = 2 (solid red line), the backbone S(q) at intermediate values of wave vector q < 0.5 σ−1 is similar to the form factor of a semiflexible chain. The simulation data for z = 2 (solid red line) were fitted to the theoretical prediction of the structure factor of a semiflexible chain (blue
a bottlebrush is proportional to the product of the backbone degree of polymerization $N_{bb}$ and the persistence length of the molecule $\propto N_{sc}^{1/2}$, that is, $\langle R^2 \rangle \propto N_{bb} N_{sc}^{1/2}$. For high grafting densities $z > 3$, bottlebrushes with short backbones $N_{bb} \lesssim 50$ adopt rodlike conformations.

**MATERIALS AND METHODS**

Simulations of bottlebrush melts were performed using three-dimensional coarse-grained bead-spring model (63). An individual bottle molecule is composed of $N_{bb}$ backbone monomers (beads) connected by bonds and $z$ side chains of $N_{sc}$ monomers grafted to every backbone monomer (see Fig. 1A). Thus, the total number of beads in a bottlebrush is $N = N_{bb} + z N_{bb} N_{sc}$, where $z$ is the grafting density. The case of $N_{sc} = 0$ (or $z = 0$) corresponds to a linear chain.

The nonbonded interactions between monomers separated by distance $r$ were modeled by the truncated and shifted LJ potential

$$
V_{\text{LJ}}(r) = \begin{cases} 
4\varepsilon[(\sigma/r)^{12} - (\sigma/r)^6] + (\sigma/r_c)^6 - (\sigma/r_c)^{12} & r \leq r_c \\
0 & r > r_c 
\end{cases}
$$

where the interaction strength $\varepsilon$ is measured in units of thermal energy $k_B T$, $\sigma$ is the monomer diameter, and $r_c$ is the cutoff. In the NVT ensemble, we have used $\varepsilon = k_B T$ and $r_c = 2^{1/6}\sigma$. This choice of LJ potential results in purely repulsive interactions between monomers. The bonded interactions in a molecule were described by the Kremer-Grest potential (63), $V_{\text{FENE}}(r) = V_{\text{FENE}}(r) + V_{\text{LJ}}(r)$, with the "finitely extensible nonlinear elastic" (FENE) potential

$$
V_{\text{FENE}} = -\frac{1}{2} k r_0^2 \ln \left[ 1 - \left( \frac{r}{r_0} \right)^2 \right]
$$

where the bond stiffness $k = 30 \varepsilon / \sigma^2$ and the maximum bond length $r_0 = 1.5 \sigma$ (63). All simulations were performed in a cubic box with periodic boundary conditions imposed in all spatial dimensions. In the NVT ensemble, the simulations were carried out at the overall monomer density $\rho = 0.85 \sigma^{-3}$ corresponding to the intermolecular pressure $\langle P \rangle \approx 4.75 \varepsilon / \sigma^3$. In a separate set of simulations, we have also investigated melts of bottlebrushes with attractive LJ potential with the interaction strength $\varepsilon = 0.84 k_B T$ and the cutoff $r_c = 2.5 \sigma$ using NPT ensemble with $P = 0$, ensuring that the average density $\langle \rho \rangle \approx 0.85 \sigma^{-3}$ is the same as in the NVT runs. The static properties obtained from both NVT and NPT simulations, for example, the average bottlebrush size, demonstrate good agreement with each other within the error bars after rescaling by the corresponding average bond length $l$ (bonds in NPT simulations are $1\%$ shorter than those in NVT simulations).

The molecular dynamics simulations were performed by solving the Langevin equation of motion for the position $r = [x_i, y_i, z_i]$ of each bead (66)

$$
m \ddot{r}_i = F_{i}\text{LJ} + F_{i}\text{FENE} - \zeta \dot{r}_i + F_i^B, \quad i = 1, \ldots, N
$$

which describes the motion of a set of interacting monomers. Forces $F_{i}\text{LJ}$ and $F_{i}\text{FENE}$ in Eq. 25 above are obtained from the LJ (Eq. 23) and FENE
(Eq. 24) interaction potentials between the i th monomer and surrounding monomers. The third and fourth terms on the right-hand side of Eq. 25 are a slowly evolving viscous force \(-\mathbf{\zeta}\mathbf{\dot{\mathbf{r}}}_i\) and a rapidly fluctuating stochastic force \(\mathbf{F}_i^\varphi\), respectively. This random force \(\mathbf{F}_i^\varphi\) is related to the friction coefficient \(\zeta\) by the fluctuation-dissipation theorem \(\langle \mathbf{F}_i^\varphi(t)\mathbf{F}_j^\varphi(t') \rangle = k_B T \mathbf{\zeta}_{ij} \delta(t - t')\). The friction coefficient used in simulations was \(\zeta = 0.5\, m\, t^{-1}\), where \(m\) is the monomer mass and \(\tau = \sqrt{m\zeta/k_B T}\) is the LJ time. The velocity Verlet scheme (67) was used to correct for the temperature constant. All simulations were carried out using Large-scale Atomic/Molecular Massively Parallel Simulator (LAMMPS) (68). Simulation snapshots were rendered using Visual Molecular Dynamics (VMD) (69). Initially, molecules were grown using a self-avoiding random walk technique and placed randomly in the simulation cell. The initial density of all systems was small (\(\approx 0.03\, \text{Å}^{-3}\)). Overlapping monomers in the initial configuration were pushed off using soft potential with slowly ramped interaction strength. To obtain the desired melt density (\(\rho = 0.85\, \text{Å}^{-3}\)), the simulation box was gradually decreased in size (see fig. S6) at constant velocity \(10^{-3}\, \text{Å}/\text{t}\). Equivalently, a short (\(\approx 10^5\, \text{t}\)) NPT simulation was performed at pressure \(P = 0.016\, \text{e}/\text{Å}^3\). We have verified that the results do not depend on the sample preparation method. Once the target density was reached, simulations were continued for up to at least three relaxation times of the corresponding system. During the equilibrium stage, the molecules diffused, on average, at least the root mean square end-to-end distance of their backbones.

Simulations of bottlebrush and linear chain melts were carried out for the following number of backbone monomers \(N_{bb} = 10, 16, 20, 50, 100\). The number of side-chain monomers \(N_m\) was varied between 0 and 32 for bottlebrushes with \(z = 1\) and between 0 and 16 for bottlebrushes with \(z = 2\) side chains attached to each backbone monomer. We assign unique symbols to denote data for each particular system. The convention of symbols used throughout the article is displayed in Fig. 4B. The complete list of symbols can be found in table S2. In addition, for molecules with \(N_{bb} = 50\), the number of side chains per backbone monomer was varied (\(z = 0, 1, 2\), and 4). To avoid the finite size effects, the number of molecules \(M\) in a simulation box was changed; thus, the box size \(a\) was at least \(\approx 2.5\) times larger than the root mean square end-to-end distance \((R^2)^{1/2}\) of bottlebrush backbones. Table S2 summarizes all parameters used in our computer simulations.

**SUPPLEMENTARY MATERIALS**

Supplementary material for this article is available at http://advances.sciencemag.org/cgi/content/full/2/11/e1601478/DC1

The size of a bottlebrush side chains in a melt

Persistence length of a bottlebrush in a melt

The size of a bottlebrush in a melt

Bottlebrush melt preparation

table S1. Summary of the adjustable parameters \(C_i^{bb}\) and \(N_m\) describing the mean square size of side chains.
table S2. Summary of system parameters for simulations of bottlebrush melts and linear chain melts.
table S3. Summary of the adjustable parameters \(A\), \(s_i\), \(s_p\), and \(\zeta\) for the bond angle correlation function.
table S4. Parameters \(C_i^{bb}\) and \(s\) describing the sizes of backbone sections.
fig. S1. The size of side chains of a bottlebrush in a melt.
fig. S2. The bond angle correlation functions \(g(s)\).
fig. S3. Persistent segments of bottlebrushes in a melt.
fig. S4. The size of backbones for bottlebrushes in a melt.
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