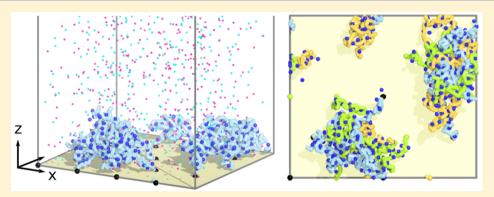


Heterogeneous Morphology and Dynamics of Polyelectrolyte Brush **Condensates in Trivalent Counterion Solution**

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Supporting Information



ABSTRACT: Recent experiments have shown that trivalent ion, spermidine³⁺, can provoke lateral microphase segregation in DNA brushes. Using molecular simulations and simple theoretical arguments, we explore the effects of trivalent counterions on polyelectrolyte brushes. At a proper range of grafting density, polymer size, and ion concentration, the brush polymers collapse heterogeneously into octopus-like surface micelles. Remarkably, the heterogeneity in brush morphology is maximized, and the relaxation dynamics of chain and condensed ion are the slowest at the 1:3 stoichiometric concentration of trivalent ions to polyelectrolyte charge. A further increase of trivalent ion concentration conducive to a charge inversion elicits modest reswelling and homogenizes the morphology of brush condensate. Our study provides a new insight into the origin of the diversity in DNA organization in cell nuclei as well as the ion-dependent morphological variation in polyelectrolyte brush layer of biological membranes.

INTRODUCTION

Polyelectrolyte brushes are ubiquitous in biological systems. As the main component of outer membrane in Gram-negative bacteria, lipopolysaccharides, consisting of charged O-polysaccharides side chains, form a brush layer and mediate the interaction of bacteria with their environment. For vertebrates, negatively charged polysaccharide hyaluronic acids play a vital role in the organization of pericellular matrix.^{2,3}

Double-stranded DNA brushes on a biochip⁴ at a cell-like density ($\sim 10^4 \text{ kb/}\mu\text{m}^3$) have been developed as a platform to study cell-free gene expression.⁵ Because of the negative charges along the backbone, this synthetic system behaves like a well-defined strong polyelectrolyte brush,5 the height of which can be varied by modulating the ionic strength of monovalent salt (NaCl) solution and the grafting density.⁶ Recently, it has been reported that trivalent counterions, spermidine³⁺ (Spd³⁺), can induce a collapse transition of DNA brush into fractal-like dendritic macroscopic domains. The heterogeneous morphology of DNA condensate points to polyamine-mediated regulation of local DNA configuration and gene expression,^{8,9} underscoring the importance of understanding the effects of multivalent counterions on polyelectrolyte brush. 10,11 Compared with uncharged polymer brushes in good or poor solvent, the electrostatic interaction and osmotic pressure, which are readily controlled by the salt concentration and pH, yield additional flexibilities in manipulating polyelectrolyte brushes and thus holding promise for its wide applications. 12,13

Conventional theoretical approaches, successful in explaining the behaviors of polyelectrolytes in monovalent salt solutions, 11,14,15 are of limited use, particularly when multivalent counterions are at work. For example, the solution to the meanfield Poisson-Boltzmann theory 16 fails to account for phenomena such as ion condensation, charge overcompensation, and collapse of like-charged polymers. 17-20 A prediction from a nonlocal density functional theory that the exponent ν in the power-law relation between brush height (H) and counterion concentration $(H \sim (C_{n+})^{-\nu})$ decreases with the valence of salt ion $(n)^{21}$ contradicts to the finding from a recent MD simulation²² that the transition is sharper for counterions with larger valence, i.e., ν increases with n.

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Polyelectrolyte brushes, which retain charged groups along the chain backbone and release ions to the solution, have been modeled explicitly with fully accounted electrostatic interactions. The dependence of brush height on counterion concentration and grafting density has been systematically investigated in mono-, di-, and trivalent ion solutions. Atomistic simulations of DNA arrays solvated with Na⁺, Mg²⁺, and spermine suggested that the transient "bridging" of multivalent cations drives the collapse of DNA. It was also shown that an excess trivalent counterions overcompensating the polyelectrolyte charges elicits re-entrance transition (decondensation) of brush.

The presence of lateral instability (or morphological heterogeneity) in polyelectrolyte brush, once noticed with AFM images, is evident in DNA (strong polyelectrolyte) brush condensates in polyamine solution. Simulations and theories have predicted the possibilities of lateral microphase segregation in homopolymer brush in *poor* solvent condition for decades, but the effects of multivalent ions on the morphological heterogeneity of polyelectrolyte brushes have not been fully investigated. Here, we carried out molecular simulations of strong polyelectrolyte brushes in trivalent counterions to study (1) the conditions for the heterogeneous collapse of brush and (2) the dynamic characteristics of polymers and multivalent ions in brush condensates.

MODEL AND METHODS

Model. A well-tested coarse-grained model of strong polyelectrolyte brush 22,23,36 was used in our study. Total M polymer chains were grafted to the uncharged surface at z=0, in a 2D triangular lattice with a lattice spacing d (Figure 1a).

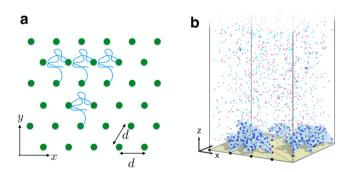


Figure 1. (a) Geometry of brush simulation. Each brush polymer is grafted in a 2D triangular lattice with spacing d. (b) A snapshot from simulation. Shown is the bottom part of simulation box where 4×4 polyelectrolyte brush (light blue) is collapsed by trivalent cations (deep blue). The monovalent counterions (magenta) are released from the polymer brush into the bulk. The monovalent anions are shown in cyan.

Each chain, consisting of N-negatively charged monomers and a neutral monomer grafted to the surface, was initially neutralized using N-monovalent counterions. The simulation box has a dimension of $L_x \times L_y \times L_z = (\sqrt{M} \, d) \times (\sqrt{3M} \, d/2) \times (2Na)$, where a is the size (diameter) of chain monomer. Periodic boundary conditions were applied in x and y dimensions, and an impenetrable neutral wall was placed at z=2Na. The location of the wall is far from the brush region, so that the density profiles of ions in the bulk region are effectively constant. 37,38

In order to model a flexible strong polyelectrolyte brush in good solvents with multivalent salts, we considered the following energy potentials. First, the finite extensible nonlinear elastic bond potential was used to constrain the bond distance between the neighboring chain monomers

$$U_{\text{bond}}(r) = -\frac{k_0 R_0^2}{2} \log \left(1 - \frac{r^2}{R_0^2} \right)$$
 (1)

with a spring constant k_0 and a maximum extensible bond length R_0 . Second, the excluded volume interaction was modeled in between ions and chain monomers by using the Weeks-Chandler-Andersen potential

$$U_{\text{excl}}(r) = 4\epsilon \left[\left(\frac{a}{r} \right)^{12} - \left(\frac{a}{r} \right)^{6} + \frac{1}{4} \right]$$
 (2)

when $r \le 2^{1/6}a$; otherwise, $U_{\text{excl}}(r) = 0$. Third, we assigned the Columbic interactions between charged particles i and j

$$U_{\text{elec}}(r) = \frac{k_{\text{B}} T \lambda_{\text{B}} q_{i} q_{j}}{r} \tag{3}$$

where $q_{i,j}$ is the corresponding charge in the unit of elementary charge e, and $\lambda_{\rm B}=e^2/(4\pi\varepsilon_0\varepsilon_{\rm r}k_{\rm B}T)$ is the Bjerrum length with the vacuum permittivity ε_0 . We used $\varepsilon_{\rm r}\approx 80$ for the relative dielectric constant of the solvent. Except for the anchored chain monomers, we considered a repulsion between the wall and any particle whose distance from the wall satisfies $z\leq 0.5a$, such that

$$U_{\text{wall}}(z) = 4\varepsilon \left[\left(\frac{a}{z+\Delta} \right)^{12} - \left(\frac{a}{z+\Delta} \right)^{6} + \frac{1}{4} \right] \tag{4}$$

with $\Delta=(2^{1/6}-0.5)$ a; otherwise, $U_{\rm wall}(z)=0$. Note that $U_{\rm wall}(z)$ and its derivative are continuous at z=0.5a. For simplicity, we assume that all the ions and chain monomers have the same values of a and $\varepsilon=k_{\rm B}T$, which are used as the basic length and energy units in this model. With $k_0=5.83~\varepsilon/a^2$, $R_0=2a$, and $\lambda_{\rm B}=4a$, MD simulations of a single grafted polyelectrolyte chain give an average bond length $\langle b \rangle=1.1a$ ($\sim a$). For double-stranded DNA this bond length maps to the effective charge separation ($\sim 1.7~{\rm \AA}$) along the chain. Since $\lambda_{\rm B}=7.1~{\rm \AA}$ in water at room temperature, one gets $\lambda_{\rm B}=7.1/1.7\times a$ ($\approx 4a$).

Simulation Methods. Simulations were carried out using ESPResSo 3.3.1 package.^{39,40} Depending on the property of interest (static or dynamic), we used two different algorithms for simulations.^{41–44} In order to get a well-sampled ensemble of configurations and obtain static (equilibrium) property of brush system for a given condition, we set the friction coefficient to a low value and integrated the following underdamped Langevin equation of motion using the velocity Verlet algorithm.

$$m\frac{d^{2}\vec{r_{i}}}{dt^{2}} = -\zeta_{\text{MD}}\frac{d\vec{r_{i}}}{dt} - \vec{\nabla}_{\vec{r_{i}}}U(\vec{r_{1}}, \vec{r_{2}}, ...) + \vec{\xi}(t)$$
(5)

where the random force noise, satisfying $\langle \dot{\xi}(t) \rangle = 0$ and $\langle \dot{\xi}(t) \cdot \dot{\xi}(t') \rangle = 6\zeta_{\text{MD}}k_{\text{B}}T\delta_{ij}\delta(t-t')$, was used to couple the simulated system to Langevin thermostat. The characteristic time scale of this dynamics is given to be $\tau_{\text{MD}} = (ma^2/\varepsilon)^{1/2}.^{45}$ We chose the time step of integration $\delta t = 0.01\tau_{\text{MD}}$ and a small friction coefficient $\zeta_{\text{MD}} = 0.1m\tau_{\text{MD}}^{-1}$. A salt-free brush was first equilibrated for $1 \times 10^6 \ \delta t$, corresponding to $\mathcal{T}_{\text{eq}} = 10^4 \tau_{\text{MD}}$,

Table 1. Parameters for the Simulated Brush Systems

group	N	χ	q_{3+}	σ	C_{3+}	group	N	χ	q_{3+}	σ	C_{3+}	group	N	χ	q_{3+}	σ	C_{3+}
A	80	2.0	0.05	8.30	0.07	В	80	0.5	2.0	2.08	0.69	С	120	1.0	1.0	2.09	0.35
A	80	2.0	0.1	8.30	0.14	В	80	0.5	4.0	2.08	1.38	C	120	1.8	1.0	3.76	0.63
A	80	2.0	0.2	8.30	0.28	C	40	1.0	1.0	12.5	2.08	C	120	2.0	1.0	4.18	0.70
A	80	2.0	0.4	8.30	0.55	С	40	1.8	1.0	22.4	3.74	C	120	2.2	1.0	4.59	0.77
A	80	2.0	0.8	8.30	1.11	C	40	2.0	1.0	24.9	4.15	C	120	3.0	1.0	6.27	1.04
A	80	2.0	1.0	8.30	1.38	C	40	2.2	1.0	27.4	4.57	C	120	5.0	1.0	10.4	1.74
A	80	2.0	2.0	8.30	2.77	C	40	3.0	1.0	37.4	6.23	C	120	9.0	1.0	18.8	3.13
A	80	2.0	4.0	8.30	5.54	C	80	1.0	1.0	4.15	0.69	C	160	1.0	1.0	1.22	0.20
В	80	0.5	0.05	2.08	0.02	C	80	1.8	1.0	7.47	1.25	C	160	1.8	1.0	2.20	0.37
В	80	0.5	0.1	2.08	0.03	C	80	2.0	1.0	8.30	1.38	C	160	2.0	1.0	2.44	0.41
В	80	0.5	0.2	2.08	0.07	С	80	2.2	1.0	9.13	1.52	C	160	2.2	1.0	2.68	0.45
В	80	0.5	0.4	2.08	0.14	C	80	3.0	1.0	12.5	2.08	C	160	3.0	1.0	3.66	0.61
В	80	0.5	0.8	2.08	0.28	С	80	5.0	1.0	20.8	3.46	C	160	5.0	1.0	6.10	1.02
В	80	0.5	1.0	2.08	0.35	C	80	9.0	1.0	37.4	6.23	C	160	9.0	1.0	11.0	1.83

^aValues of polymer chain length N, dimensionless overlap parameter χ , dimensionless trivalent counterion concentration q_{3+} , grafting density σ in units of $10^{-3}a^{-2}$, and trivalent ion concentration C_{3+} in units of $10^{-3}a^{-3}$.

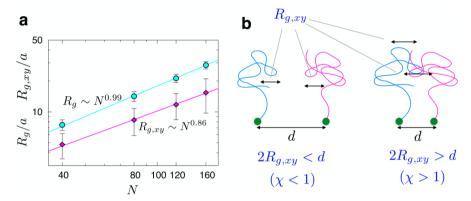


Figure 2. (a) Radius of gyration $R_{\rm g}$ and its in-plane component $R_{\rm g,xy}$ obtained from simulations of a single grafted polyelectrolyte chain with varying lengths N=40, 80, 120, and 160, grafted on surfaces. $R_{\rm g}^2=\frac{1}{2(N+1)^2}\sum_{i=0}^N\sum_{j=0}^N\sum_{\alpha=1}^N(r_{i\alpha}-r_{j\alpha})^2$ and $R_{\rm g,xy}^2=\frac{1}{2(N+1)^2}\sum_{i=0}^N\sum_{j=0}^N\sum_{\alpha=1}^N(r_{i\alpha}-r_{j\alpha})^2$, where $r_{i\alpha}$ is the α -th Cartesian component of monomer i. The scaling relationship of $R_{\rm g,xy}^2\sim N^{1.7}$ acquired from simulations is used to discuss the scaling arguments made in eqs 10 and 14. (b) The chain overlap parameter χ is illustrated.

from an initial configuration where polymer chains were vertically stretched with monovalent counterions homogeneously distributed in the brush region. To simulate the effect of trivalent cations on the brush systems, we randomly added cations and its monovalent co-ions (anions) into the brush-free zone. The production runs were generated for $10^5 \times \tau_{\rm MD}$. A snapshot from simulations was saved every $25 \times \tau_{\rm MD}$ for the analysis.

When examining dynamic behaviors of system (such as configurational relaxation of brush) and evaluating the associated time scale of dynamics, which we report in Figure 8 by using eq 15, simulations under overdamped condition are more appropriate. 44,46 In order to probe the dynamic behavior of brush, we switch our simulation algorithm to Brownian dynamics (BD). In BD simulations the following equation of motion is integrated:

$$\frac{d\vec{r}_{i}}{dt} = -\frac{D_{i0}}{k_{B}T} \vec{\nabla}_{\vec{r}_{i}} U(\vec{r}_{1}, ..., \vec{r}_{N}) + \vec{R}_{i}(t)$$
(6)

where D_{i0} is the bare diffusion coefficient of the *i*th particle, and $\vec{R}_i(t)$ is the Gaussian random noise satisfying the fluctuation—dissipation theorem $\langle \vec{R}_i(t) \cdot \vec{R}_j(t') \rangle = 6D_{i0}\delta_{ij}\delta(t-t')$. For the diffusion constant of ions and chain monomers, we used $D_{i0} = 1$

1.33, 2.03, 0.48, and $0.24 \times 10^3 \ \mu m^2/s$ for Na⁺, Cl⁻, Spd³⁺, and DNA monomer, respectively. ^{47–49} For the integration time step of simulation, we chose $\delta t_{\rm BD} = 5 \times 10^{-3} \tau_{\rm BD}$, which is much smaller than the Brownian time $\tau_{\rm BD} = a^2/D$. The last configuration of brush obtained from MD simulation was used as the starting configuration of BD simulation. The total simulation time for BD in each trajectory is $7.5 \times 10^4 \tau_{\rm BD}$.

Because of the slab geometry of our system, an electrostatic layer correction to the particle–particle particle–mesh algorithm (P³M+ELC) was used to calculate the long-range Coulomb interactions, 50 which required an extension of the simulation box along the +z direction so that there is a vacuum slab above the top impenetrable wall. The height of the slab was chosen to be Na/4, and the error tolerance was set to $10^{-5}\varepsilon$. Differences between the electrostatic potentials of some exemplary configurations, calculated by P³M+ELC and by MMM2D⁵¹ algorithm, were negligible $\sim 0.01\varepsilon$, indicating that Coulomb interactions have been fairly accounted.

Brushes of varying chain length (N) and grafting density (σ) were studied at varying trivalent counterion concentration C_{3+} . The parameters explored for the brush system are listed in Table 1. We introduce two dimensionless parameters to describe the system. The extent of lateral overlap between

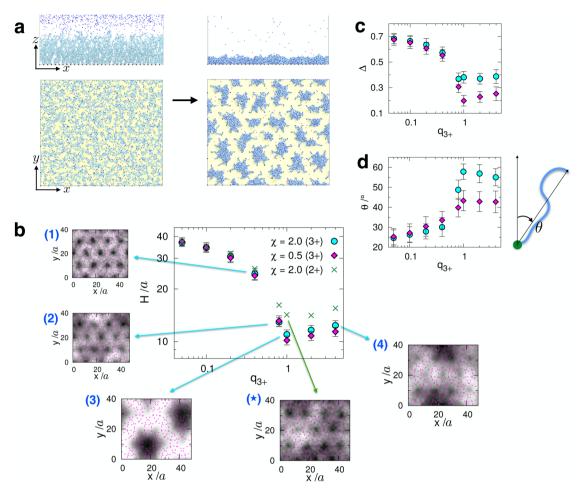


Figure 3. (a) Configuration of brush (side and top views) before and after trivalent counterion-induced collapse to condensates $(M=16\times 16, N=80, \chi=2.0, \text{ and } q_{3+}=1.0)$. (b) Brush height $H=2\sum_{i=0}^N z_i/(N+1)$. Lateral monomer density maps of brushes $(\chi=2.0)$ averaged over time, $(\rho(x,y))$, are shown in the subpanels (1)-(4) at $q_{3+}=0.4$, 0.8, 1.0, and 4.0. (the first monomers anchored to the surface (z<2a) are excluded from the visualization using $(\rho(x,y))$). The dashed magenta circles on the profiles depicts the radius $R_{g,xy}$ with respect to the anchoring point of chain. The brush height with increasing divalent counterion concentration is depicted with the symbols \times and monomer density map for $q_{2+} \left(= \frac{2C_2+V}{M\times N} \right) = 1.0$ is shown. (c) Asphericity $\Delta = \sum_{i=1}^3 (\lambda_i - \overline{\lambda})^2/6\overline{\lambda}^2$ where λ_i are the eigenvalues of inertia tensor, $\overline{\lambda} = \sum_{i=1}^3 \lambda_i/3.5^{25,53}$ $\Delta = 0$ for a perfect sphere. (d) Polar angle $\theta = \angle(\overline{z}, \overline{r}_e)$, where \overline{r}_e is the end-to-end vector of individual chains, as a function of trivalent ion concentration q_{3+} .

neighboring polymer chains is formulated using the chain overlap parameter $\chi \equiv (2R_{\rm g,xy}/d)^2$, where $R_{\rm g,xy}$, which scales as $R_{\rm g,xy}^2 \sim N^{1.7}$ (Figure 2), is the in-plane component of the gyration radius of a *single* grafted polyelectrolyte chain. The parameter χ can be related to the grafting density σ via $\chi = 2\sqrt{3\sigma}R_{\rm g,xy}^2 \sim \sigma N^{1.7}$. Collective features of brushes are expected to become significant only for $\chi \geq 1$ (see Figure 2).

Next, we used another dimensionless parameter q_{3+} which defines the ratio of the total positive charges on the trivalent counterions in the simulation box of volume $V\left(Q_{3+}=3C_{3+}V\right)$ to the total negative charges on the chain monomers $\left(Q_{\text{mono}}=M\times N\right)$, i.e., $q_{3+}=Q_{3+}/Q_{\text{mono}}$. Thus, q_{3+} becomes 1 $\left(q_{3+}=1\right)$ when the concentration of trivalent counterions is set to the 1:3 stoichiometric ratio at which a trivalent ion (+3) neutralizes three monomers in the brush.

RESULTS

Effect of Trivalent Ion on Polymer Brush. Upon addition of trivalent counterions to the bulk phase, the trivalent ions are spontaneously pulled into the brush phase, replacing the monovalent counterions neutralizing the bare charges on

the brush monomers (Figure 3a). The morphology of brush condensate varies depending on the size of brush chain (N), grafting density (σ) or equivalently χ), and trivalent ion concentration (q_{3+}) .

For N=80, $\chi=2.0$ with an increasing trivalent ion concentration (C_{3+}) but with $q_{3+}<1$ (group A in Table 1), the brush height drops gradually owing to the decrease of the osmotic pressure,³ and the lateral distribution of the brush becomes more inhomogeneous. Our study finds that at $q_{3+} = 1$ the height of brush (H) is minimized (Figure 3b) and that the grain boundary between chain-free and chain-rich phase projected on the xy plane becomes clear in the same condition. The density profiles of chain monomers in xy plane $\langle \rho(x,y) \rangle$ (Figure 3b) clearly visualize the morphology of brush condensate at various q_{3+} , highlighting that the development of lateral heterogeneity in the brush condensate is maximized at $q_{3+} = 1$. At this ideal charge compensation point $(q_{3+} = 1)$, the six chains stretched from the corners of a hexagon are organized into aggregates entangled at the center, forming octopus-like surface micelles (Figure 3b-(3), see also a snapshot in Figure 4b), demarcated by the grain boundary whose width is about $d = 2R_{g,xy}/\sqrt{\chi}$. A further increase of q_{3+} beyond q_{3+}

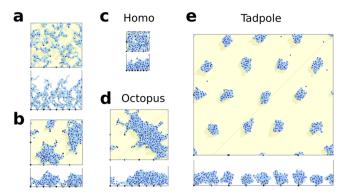


Figure 4. Top and side views of snapshots from simulations of polyeletrolyte brush condensates at various conditions. (a) N=80, $\chi=2.0$, $q_{3+}=0.4$. (b) N=80, $\chi=2.0$, $q_{3+}=1.0$. (c) N=40, $\chi=3.0$, $q_{3+}=1.0$. (d) N=120, $\chi=3.0$, $q_{3+}=1.0$. (e) N=160, $\chi=1.0$, $q_{3+}=1.0$. (b), (d), and (e) correspond to the brush morphologies in (6), (4), and (3) of the structural diagram (Figure 7), respectively.

 $1 (q_{3+} > 1)$ elicits reswelling of the brush along the *z*-direction, obscuring the boundaries between the surface micelles (Figure 3b-(4)).

The lateral heterogeneity in brush condensate is unique to the counterions whose valence is greater than two. It is worth noting that as depicted in Figure 3b-(\bigstar), addition of divalent counterions even at $q_{2+} = 1$, where the divalent ions can fully neutralize the total bare charge of chain monomers, does not elicit the heterogeneous brush collapse as in Figure 3b-(3).

The morphology of a sparsely grafted polyelectrolyte brush condensate (χ < 1) differs greatly from that of a dense brush condensate (group B in Table 1). Even at high C_{3+} (or high q_{3+}), ion-mediated interchain attractions are not strong enough to sustain a fused micelle overcoming the penalty of stretching individual polymer from the grafting point. As a consequence, each grafted chain collapses into a separate tadpole-like/mushroom structure (Figures 4e).

Changes of the in-plane morphology of the brush are also mirrored in the configuration of individual polyelectrolyte chain. Of immediate notice are the nonmonotonic dependences of brush height (H), chain asphericity (Δ) , and tilt angle (θ) on q_{3+} (Figure 3). The chains in dense brush at $q_{3+}=1$ display the greatest compression along the z-axis and tilt angle, which can be explained by laterally stretched configurations of the polyelectrolyte chain in the octopus-like condensate. Both Δ and θ are smaller in a sparse brush $(\chi=0.5)$ than in a dense one $(\chi=2.0)$, pointing to their tadpole-like configuration.

Next, of great interest are the distribution of counterions in the brush and bulk regions. For $q_{3+} \ll 1$, when there is only a small amount of trivalent counterions in the system, the monovalent cations are still dominant in the brush region $(\rho_{3+}(z) \ll \rho_{+}(z))$, compare the red solid lines in Figure 5a,b). But, as $q_{3+}(<1)$ increases, more trivalent counterions are condensed into the brush region, compressing the brush into a more compact condensate, and when the 1:3 stoichiometric condition of $q_{3+} = 1$ is reached, the distribution of trivalent counterions in the brush region exceeds that of monovalent counterion $(\rho_{3+}(z) \gg \rho_{+}(z))$, compare the green lines in Figures 5a and 5b). For $q_{3+} > 1$, the excess trivalent counterions are condensed on polyelectrolyte brush region together with monovalent anions (see the black solid/dashed lines in Figure 5a,b and symbols in Figure 5c). When all the charge ratios of ions are summed up $(\varphi_{tot} = \varphi_{3+} + \varphi_{+1} - \varphi_{-1})$ for $q_{3+} > 1$, the

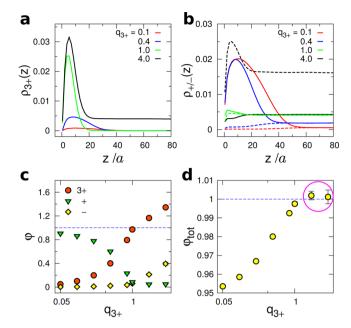


Figure 5. Distribution of ions. (a) Distributions (along *z*-axis) of trivalent counterions, (b) monovalent counterions (solid lines) and co-ions (dashed lines), in units of a^{-3} , calculated from a collapsed brush system with N=80 and $\chi=2.0$ for varying $q_{3+}=0.1, 0.4, 0.8, 1.0$, and 4.0. (c) Ratio of total charge of trivalent counterions (φ_{3+}) , monovalent counterions (φ_+) , and monovalent co-ions (φ_-) , trapped within the range of Bjerrum length $\lambda_B \sim 4a$ from any chain monomer, to the amount of bare charges in the brush. (d) $\varphi_{\rm tot}$ (= $\varphi_{+3} + \varphi_{+1} - \varphi_{-1}$) as a function of q_{3+} . The condition of brush charge neutralization is marked with a blue dashed line, and the charge inversions by counterions and co-ions at $q_{3+} > 1$ are highlighted with a magenta circle enclosing the two points greater than $\varphi_{\rm tot} = 1$.

total charge ratio (φ_{tot}) is slightly greater than the unity, $\varphi_{tot} \gtrsim 1$ (Figure 5d). In fact, this charge inversion is responsible for the reswelling of brush suggested in Figures 3b–d. Here, it is noteworthy that the density profile of trivalent ions is uniform in the polymer-free zone, which indicates that the simulation box size is sufficiently large and that the effect of box size in the z-direction on brush structure is negligible.

In order to further quantify the lateral configuration of the brush, we calculated the in-plane static structure factor

$$S_{xy}(k) = \left\langle \left\langle \frac{1}{N_m^2} \left| \sum_{i,j=1}^{N_m} e^{i\vec{k} \cdot (\vec{r}_i - \vec{r}_j)} \right| \right\rangle_{|\vec{k}|} \right\rangle$$
(7)

where $N_m = M \times N$ is the total number of nongrafted chain monomers in the brush, \vec{r}_i is the position of the ith monomer, and \vec{k} is 2D wave vector in the xy plane. $S_{xy}(k)$ is evaluated by first integrating over the space of $|\vec{k}| = k$ and then averaging over the simulation trajectory. As shown in Figure 6a, $S_{xy}(k)$, calculated for N = 80 and $\chi = 2.0$, in the range of $k > 2\pi/L_y$ is maximized at $ka \approx 0.21$. This indicates that at a characteristic length scale of $R \sim 2\pi/k \approx 30a$ there is a repeated pattern in the system, which corresponds to the octopus-like surface micelle domains of size $\sim 30a$ (see the monomer density map of $\langle \rho(x,y) \rangle$ in Figure 3b-(3)).

In accord with our observation that the lateral heterogeneity becomes most evident at $q_{3+}=1$ in Figure 3, nonmonotonic dependence of $S_{xy}(k=0.21a^{-1})$ on q_{3+} is again maximized sharply at $q_{3+}=1.0$ (Figure 6a, inset). At $q_{3+}=0.1$, which

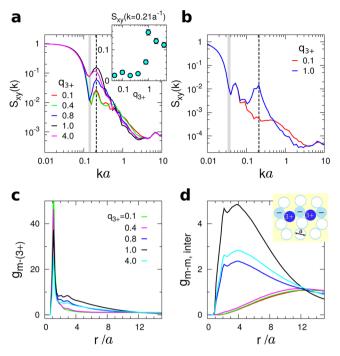


Figure 6. Lateral static structure factor $S_{xy}(k)$ as a function of wavenumber k calculated at N=80 and $\chi=2.0$ for (a) 4×4 and (b) 16×16 brush systems with various trivalent ion concentrations $q_{3+}=0.1$, 0.4, 0.8, 1.0, and 4.0. Dashed vertical lines indicate the peaks around ka=0.21, whose amplitude is plotted with varying q_{3+} (inset). The gray solid lines demarcate the range $2\pi/L_x \le k \le 2\pi/L_y$, i.e., the periodic boundary of the simulation box. Pair correlation functions between (c) monomer and trapped trivalent ion (3+) and (d) monomers from different chains as a function of the interparticle distance r. The insert of (d) is a cartoon of the chain "bridging" mechanism of trivalent ions, which explains the peak positions of $g_{m-m,inter}$ around r=2a and 4a.

supposedly corresponds to a homogeneous brush phase, a peak is still identified in $S_{xy}(k)$, but this is due to an artifact of periodic boundaries imposed on the 4× 4 brush system (Figure 6a). In $S_{xy}(k)$ calculated from the 16 × 16 brush system (Figure 6b), this peak at $q_{3+}=0.1$ vanishes, while the peak of $S_{xy}(k)$ at ka=0.21 in the salted brush ($q_{3+}=1.0$) still remains. It is worth noting that the domain size of surface micelle ($R=2\pi/k\approx 30a$, the value of k which maximizes $S_{xy}(k)$) is identical for 4 × 4 and 16 × 16 brush systems.

All these nonmonotonic dependences on q_{3+} arise from the effective interchain monomer attractions mediated by the condensed trivalent counterions. The pair correlations between the charged monomers and trivalent ions are strongest at q_{3+} = 1.0 (Figure 6c). Furthermore, the interchain monomer correlation has the greatest values at r = 2a and r = 4a (Figure 6d). It is worth noting that the pair correlation function between trivalent ions condensed to the brush is not as sharp or structured as that of Wigner crystal which was observed in flexible polyelectrolyte chain in free space at relatively good solvent condition collapsed by multivalent ions.⁵⁴ In fact, the diffusivity of the trivalent ions is finite even when they are trapped in the polyelectrolyte brush (see below). Still, the pronounced correlation at r = 2a lends support to the "bridging" mechanism by which charged monomers in polyelectrolyte are in effective attraction mediated by trivalent

Structural Diagram. At $q_{3+}=1.0$, the stoichiometric condition at which the most heterogeneous brush structures emerge, the effects of chain size (N) and chain overlap (χ) , which is related to the grafting density σ as $\chi \sim \sigma R_{g,xy}^{2}$, on the morphology of brush are further studied (group C in Table 1). As depicted in Figure 7 (see also Figure 4), three distinct phases are present. As a trivial case, tadpole-like condensates are expected if σ is very low (see the brush of N=160, $\chi=1.0$, Figure 7-(3)). When the polyelectrolyte chains are either short

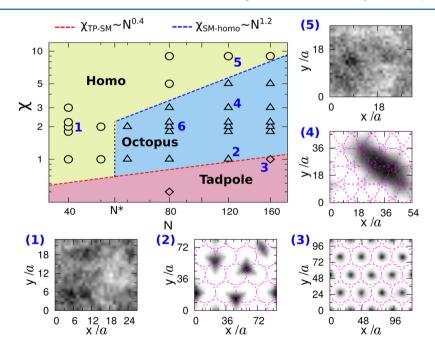


Figure 7. Structural diagram of polyelectrolyte brush simulated at $q_{3+}=1.0$ as a function of chain length N and overlap parameter χ . Typical lateral density profiles of brushes are shown together. Dashed lines are phase boundaries $\chi \sim N^{0.4}$ and $\chi \sim N^{1.2}$, which are derived from a scaling argument. The vertical phase boundary formed due to the competition between chain fluctuations and trivalent ion-mediated interaction is marked at $N^* \approx 50-60$ between homogeneous and octopus phases. The density profile at state point (6) is shown in Figure 3b-(3).

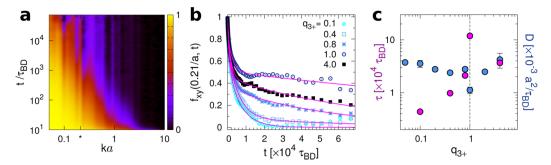


Figure 8. Dynamics of brush condensates. (a) Heat map of normalized intermediate scattering function $f_{xy}(k,t)$ for a polyelectrolyte brush with N=80 and $\chi=2.0$ and at $q_{3+}=1.0$. (b) $f_{xy}(k=0.21a^{-1},t)$ for bushes at $q_{3+}=0.1$, 0.4, 0.8, 1.0, and 4.0. Solid lines are fittings of $f_{xy}(k,t)$ to a multiexponential function. (c) Relaxation time τ and diffusion coefficient of trapped trivalent ions as a function of q_{3+} .

(N=40, Figure 7-(1)) or grafted with a very high density $\chi > 5.0$ (Figure 7-(5)), the morphology of brush condensates is homogeneous. With $1 \le \chi \le 3$, octopus-like surface micelle domains are formed at an intermediate range of χ (Figure 7-(2) or (4)). Assuming that the collapsed phase of polyelectrolyte brush with trivalent ions is reasonably mapped onto the collapsed polymer brush in a poor solvent, we treated the counterion-mediated interactions as an effective attraction and used the scaling argument³⁴ to describe the phase boundaries in Figure 7.

Boundary between Tadpole and Surface Micelle. The free energy of individual tadpole composed of a single chain, like the case of N=160 and $\chi=1.0$ (Figure 7-(3)), is dominated by its surface energy, $F_{\rm TP}\sim\gamma R^2$, where γ is an effective surface tension mediated by the salt concentration, and R is the size of a tadpole-like structure. Assuming that the surface energy is around $k_{\rm B}T$ per monomer, $\gamma\sim\xi k_{\rm B}T/a^2$, where the parameter ξ , which is maximized at $q_{3+}=1$, denotes a scale of counterion mediated-effective attraction between chain monomers, and $R\sim aN^{\nu}$ for a collapsed globule of polyelectrolyte chain, $F_{\rm TP}$ is then given by

$$\frac{F_{\rm TP}}{k_{\rm B}T} \sim \xi N^{2\nu} \tag{8}$$

The free energy of a bigger fused micelle composed of three neighboring chains, like the case of N=120 and $\chi=1.0$ (Figure 7-(2)), can be expressed as a summation of a surface energy term γ ($(3N)^{\nu}a)^2$, and a elastic energy term $3k_{\rm B}Td^2/(N_{\rm s}a^2)$ as a result of stretching $N_{\rm s}=d/a$ monomers in each chain to reach the surface micelle. Taken together, it yields

$$\frac{F_{\rm FM}}{k_{\rm B}T} \sim \xi (3N)^{2\nu} + 3\frac{d}{a}$$
 (9)

Finally, the balance between $F_{\rm FM}$ and $F_{\rm TP}$ leads to a critical grafting distance $d_{\rm c}$ (or grafting density, $\sigma_{\rm c}$), $\sigma_{\rm c} \sim d_{\rm c}^{-2} \sim \xi^{-2} N^{-4\nu}$. Therefore, the phase boundary between tadpole and fused surface micelle has the following scaling behavior in terms of the chain overlap parameter ($\chi \sim \sigma R_{\rm g,xy}^2 \sim \sigma N^{1.7}$) and N.

$$\chi \sim \xi^{-2} N^{1.7-4\nu} \tag{10}$$

Boundary between Surface Micelle and Homogeneously Collapsed Layer. In a similar way, the free energy per unit area of a homogeneously collapsed layer, like the case of N=120 and $\chi=9.0$ (Figure 7-(5)), is $f_{\text{homo}}=2\gamma$

$$\frac{f_{\text{homo}}}{k_{\text{B}}T} \sim \xi \frac{2}{a^2} \tag{11}$$

where the factor 2 arises because there are two interfaces with the collapsed polymer: one at the grafting points and another at the interface with the bulk.

Meanwhile, for an octopus-like domain of size R_n , containing n chains within a surface area defined by $\sim R_c^2$, the surface energy is $F_{n,\text{surf}} \sim \gamma R_n^2 \sim \xi k_\text{B} T (nN)^{2\nu}$, and the elastic energy is $F_{n,\text{elastic}} \sim n k_\text{B} T R_\text{c}^2 / (N_\text{s} + a^2) = n k_\text{B} T R_\text{c} / a = k_\text{B} T \sigma R_\text{c}^3 / a$ where we used the relationships of $R_c = N_\text{s} \times a$ and $\sigma = n / R_c^2$. Hence, the total free energy, per unit area, in octopus-like condensate with n arms, is

$$\frac{f_{\text{octo}}}{k_{\text{B}}T} = \frac{1}{k_{\text{B}}T} \frac{F_{n,\text{surf}} + F_{n,\text{elastic}}}{R_{\text{c}}^2}$$

$$= \frac{\xi(nN)^{2\nu} + \sigma R_{\text{c}}^3/a}{R_{\text{c}}^2}$$

$$= \frac{\xi(\sigma N)^{2\nu}}{R_{\text{c}}^{2-4\nu}} + \frac{\sigma R_{\text{c}}}{a}$$
(12)

Minimization of the free energy, f_{octo} , with respect to R_c gives a minimum value of $f_{\text{octo}}(R_c^*) \sim N^{2\nu/(3-4\nu)} \sigma^{(2-2\nu)/(3-4\nu)}$, at the following size R_c^* :

$$R_c^* \sim N^{2\nu/(3-4\nu)} \sigma^{(2\nu-1)/(3-4\nu)}$$
 (13)

By balancing f_{octo} with f_{homo} , we also obtain the critical grafting density between these two phases, and hence the chain overlap parameter χ which scales with N and ξ as follows:

$$\chi \sim \xi^{(1-2\nu)/(1-\nu)} N^{1.7+(\nu/(\nu-1))}$$
 (14)

To recapitulate, for $\nu=1/3$, which is the scaling exponent for collapsed polymer, the phase boundary between tadpole and fused micelle (surface micelle) is $\chi_{\text{TP-SM}} \sim \xi^{-2} N^{0.4}$, and the phase boundary between homogeneously collapsed brush and octopus-like surface micelle is $\chi_{\text{SM-homo}} \sim \xi^{1/2} N^{1.2}$. Therefore, the surface micelle phase is enclosed by these two boundaries, and its area in Figure 7 is maximized when $q_{3+}=1$, where the strength of effective attraction between monomers reaches its maximum value ($\xi \leq \xi_{\text{max}}$). The vertical boundary (the dotted line in Figure 7) between homogeneous and octopus phases at $N^* \approx 50$ –60 is due to large fluctuations of short chains which predominate over the counterion-induced attraction between neighboring chains and prevent them from aggregating into a metastable octopus-like surface micelle.

Dynamics of Brush Condensate. The intermediate scattering function for chain monomers⁵⁵

$$F_{xy}(k, t) = \left\langle \left\langle \frac{1}{N_m^2} \sum_{m=1}^{N_m} e^{i\vec{k} \cdot \vec{r}_m(t+t_0)} \sum_{n=1}^{N_m} e^{-i\vec{k} \cdot \vec{r}_n(t_0)} \right\rangle_{|\vec{k}|} \right\rangle_{t_0}$$
(15)

is evaluated using the trajectories generated from Brownian dynamics simulations. $F_{xy}(k,t)$ describes the conformational relaxation of brush chains on different spatial scales. The heat map of $f_{xy}(k,t)$ [= $F_{xy}(k,t)/F_{xy}(k,0)$], for the brush with N=80, $\chi=2.0$ is plotted in Figure 8a.

The fast decaying behavior of $f_{xy}(k,t)$, i.e., on small length scales (ka > 1), indicates that the monomers inside the collapsed surface micelles are fluidic and that the trivalent ionmediated polymer-polymer attractions are still transient. In contrast, the relaxation dynamics of polymer brush in a larger length scale (ka < 1) is rather slow. The scattering function $f_{xy}(k = 0.21a^{-1}, t)$, which probes the dynamics of brush on the length scale of surface micelle size $(R = 2\pi/k \approx 30a)$, exhibits slow decaying behavior, which is fitted to a multiexponential function (Figure 8b) or to a stretch-exponential function $\sim \exp(-(t/\tau)^{\beta})$ with $\beta < 1$ (Figure S1). Together with the exponent of stretch-exponential β less than 1, which is a quintessential marker of glassy dynamics, 56 the nonmonotonic dependence of τ on q should be noted (Figure 8c). The relaxation dynamics of octopus-like surface micelle at $q_{3+} \approx 1$ is ~30-fold slower than that at $q_{3+} = 0.1$. Given that the size of polyelectrolyte chain we simulated is finite (N = 80), it is expected that the relaxation time of octopus-like surface micelle would effectively be divergent at a mesoscopic chain size $N \gg 1$ as in ref 7. Once trapped in a dynamically metastable state, the heterogeneous configurations of surface micelle are maintained for an extended amount of time.

At $q_{3+}=1.0$, when the relaxation dynamics of polyelectrolyte chain is the slowest, the trivalent ions trapped in the brush region exhibit the slowest dynamics as well; the diffusion coefficient of trivalent ions trapped in the brush region, which can be calculated from an ensemble averaged mean square displacement, is minimal $(D_{\rm ion}^{\rm brush} \sim 10^{-3} a^2/\tau_{\rm BD})$, Figure 8c) and is only $\sim 1/100$ of that in the bulk $(D_{\rm ion}^{\rm bulk} \sim 10^{-1} a^2/\tau_{\rm BD})$. When $q_{3+}>1.0$, the effective correlation between trapped ions and monomers becomes smaller than the value at $q_{3+}=1.0$ (see Figure 6c), giving rise to a greater diffusivity (Figure 8c).

DISCUSSION

At high trivalent counterion concentration ($q_{3+} \gtrsim 1.0$), dense polyelectrolyte brushes composed of long chains ($\chi > 1.0$ and N > 40) collapse to spatially inhomogeneous brush condensates. The octopus-like surface micelle domains, demarcated by chain-depleted grain boundaries, become most evident at the ionic condition of q_{3+} = 1. Our structural diagram (Figure 7), the accompanying scaling arguments, and dynamics of brush condensates assessed by intermediate scattering functions (eq 15 and Figure 8) qualitatively capture the essence of the experimental observations by Bracha et al. In their experiment, DNA brushes underwent a transition into macroscopic dendritic domains, only if the spermidine concentration C_{Spd} was greater than the critical value C_{Spd}^* (\approx 61 μ M). Just like the vertical phase boundary at $N^* \approx 50$ – 60 demonstrated in the structural diagram of our study, the minimal chain length of ~1 kb was required for the growth of dendritic domains. It was also observed that for $C_{\text{Spd}} > C_{\text{Spd}}^*$ the collapsed domains were smaller in a denser brush of shorter chains, which is consistent with the prediction from our scaling

argument $R_c \sim N^{0.4}\sigma^{-0.2}$. The scaling relationship of $R_c \sim N^{0.4}\sigma^{-0.2}$ further explains the various size of octopus-like micelle domain depicted in (2), (4), and (6) in Figure 7. According to Figure 7, the heterogeneous morphology of collapsed brush is more likely to be acquired for large N; however, when the grafting density is extremely high $(\chi \gg 1)$ crossing the phase boundary of $\chi_{\rm SM-homo} \sim N^{1.2}$, the morphology of brush condensate becomes homogeneous regardless of N.

It is noteworthy that there have been a large amount of studies on a collection of free polyelectrolytes in solution. S7-59 Yet, the geometric constraint of the brush system with one end being grafted makes substantial difference. While the two systems share similarity in that the attraction between two polymer chains is initiated and mediated by counterions, which in a certain condition for free polyelectrolytes leads to a macrophase segregation through infinite bundling, S7-59 the very constraint on the one termini in brush system limits the size of aggregates leading to lateral microphase segregation, the domain size of which is determined by N and σ (or χ), i.e., $R_{\rm c} \sim N^{0.4} \sigma^{-0.2}$.

A few cautionary remarks are in place in comparing our studies with the existing dsDNA biochip experiments. Compared with the DNA brush experiment, the simulated brush is more than an order of magnitude shorter, but with a higher grafting density. Importantly, we have not taken into account the finite chain stiffness of double-stranded DNA⁶⁰ by reasoning that the ratio of contour (L) and persistence lengths (l_p) of dsDNA used in biochip experiment $(L/l_p \sim O(10^2))$ is large enough to ignore the effect of chain stiffness on overall conclusion of our study. However, in reality the local chain stiffness does affect the local coalignment of brush condensate, which is propagated and amplified into a macroscopic scale. As a consequence, the chain organization inside octopus-like domain differ from that inside the dendritic domain in ref 7, where tens to hundreds of chains are coaligned in a width of 20-100 nm and tightly packed in the hexagonal columnar order. 7,61 In addition, different models for ions and polyamine compounds such as their size and charge distribution 62,63 and spatial inhomogeneity in grafting density could modulate the critical salt concentration for maximal heterogeneity.

Furthermore, although our study was carried out through a specific protocol of preparation, that is, by adding trivalent salt to the chain-free zone (bulk) and waiting for the system to be equilibrated, it may well be that the result of this exercise changes with the protocol of how the trivalent ions are added. As reported by the experiments, 7,13 different modes of incorporating spermidine into brush (e.g., the rate by which trivalent counterions are added) lead to slightly different critical concentrations, different collapse dynamics, and different morphologies. All these kinetic effects of spermidine addition are closely linked to the ultraslow dynamics of brush condensates at $q_{3+} \approx 1$, underscored in Figure 8, and are of great interest for the future study.

To recapitulate, our study indicates that when a condition is met, lateral heterogeneity develops in pericellular matrices as well as in DNA biochip. The results of our study also imply that the local compactness of DNA in interphase chromosome is manipulated through dynamic controls over the DNA density and the biogenic polyamine concentration, which in turn regulates DNA transcription activities.

ASSOCIATED CONTENT

S Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.macromol.6b02685.

Figure S1 (PDF)

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Notes

The authors declare no competing financial interest.

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REFERENCES

- (1) Schneck, E.; Papp-Szabo, E.; Quinn, B. E.; Konovalov, O. V.; Beveridge, T. J.; Pink, D. A.; Tanaka, M. Calcium ions induce collapse of charged O-side chains of lipopolysaccharides from Pseudomonas aeruginosa. *J. R. Soc., Interface* **2009**, *6* (Suppl 5), S671–S678.
- (2) Attili, S.; Borisov, O. V.; Richter, R. P. Films of End-Grafted Hyaluronan Are a Prototype of a Brush of a Strongly Charged, Semiflexible Polyelectrolyte with Intrinsic Excluded Volume. *Biomacromolecules* **2012**, *13*, 1466–1477.
- (3) Cohen, M.; Joester, D.; Geiger, B.; Addadi, L. Spatial and Temporal Sequence of Events in Cell Adhesion: From Molecular Recognition to Focal Adhesion Assembly. *ChemBioChem* **2004**, *5*, 1393–1399.
- (4) Karzbrun, E.; Tayar, A. M.; Noireaux, V.; Bar-Ziv, R. H. Programmable on-chip DNA compartments as artificial cells. *Science* **2014**, 345, 829–832.
- (5) Bracha, D.; Karzbrun, E.; Daube, S. S.; Bar-Ziv, R. H. Emergent Properties of Dense DNA Phases toward Artificial Biosystems on a Surface. *Acc. Chem. Res.* **2014**, *47*, 1912–1921.
- (6) Bracha, D.; Karzbrun, E.; Shemer, G.; Pincus, P. A.; Bar-Ziv, R. H. Entropy-driven collective interactions in DNA brushes on a biochip. *Proc. Natl. Acad. Sci. U. S. A.* **2013**, *110*, 4534–4538.
- (7) Bracha, D.; Bar-Ziv, R. H. Dendritic and Nanowire Assemblies of Condensed DNA Polymer Brushes. *J. Am. Chem. Soc.* **2014**, *136*, 4945–4953.
- (8) Cherstvy, A. G. Electrostatic interactions in biological DNA-related systems. *Phys. Chem. Chem. Phys.* **2011**, 13, 9942–9968.
- (9) Carrivain, P.; Cournac, A.; Lavelle, C.; Lesne, A.; Mozziconacci, J.; Paillusson, F.; Signon, L.; Victor, J.-M.; Barbi, M. Electrostatics of DNA compaction in viruses, bacteria and eukaryotes: functional insights and evolutionary perspective. *Soft Matter* **2012**, *8*, 9285–9301.
- (10) Brettmann, B. K.; Laugel, N.; Hoffmann, N.; Pincus, P.; Tirrell, M. Bridging contributions to polyelectrolyte brush collapse in multivalent salt solutions. *J. Polym. Sci., Part A: Polym. Chem.* **2016**, 54, 284–291.
- (11) Ballauff, M.; Borisov, O. V. Phase transitions in brushes of homopolymers. *Polymer* **2016**, *98*, 402–408. Special Issue: Polymer Brushes.
- (12) Szilagyi, I.; Trefalt, G.; Tiraferri, A.; Maroni, P.; Borkovec, M. Polyelectrolyte adsorption, interparticle forces, and colloidal aggregation. *Soft Matter* **2014**, *10*, 2479–2502.
- (13) Pardatscher, G.; Bracha, D.; Daube, S. S.; Vonshak, O.; Simmel, F. C.; Bar-Ziv, R. H. DNA condensation in one dimension. *Nat. Nanotechnol.* **2016**, *11*, 1076–1081.

(14) Pincus, P. Colloid stabilization with grafted polyelectrolytes. *Macromolecules* **1991**, 24, 2912–2919.

- (15) Israels, R.; Leermakers, F. A. M.; Fleer, G. J.; Zhulina, E. B. Charged Polymeric Brushes: Structure and Scaling Relations. *Macromolecules* **1994**, *27*, 3249–3261.
- (16) Netz, R. R.; Orland, H. Beyond Poisson-Boltzmann: Fluctuation effects and correlation functions. *Eur. Phys. J. E: Soft Matter Biol. Phys.* **2000**, *1*, 203–214.
- (17) Deserno, M.; Holm, C.; May, S. Fraction of Condensed Counterions around a Charged Rod: Comparison of Poisson-Boltzmann Theory and Computer Simulations. *Macromolecules* **2000**, 33, 199–206.
- (18) Hsiao, P.-Y. Overcharging, Charge Inversion, and Reentrant Condensation: Using Highly Charged Polyelectrolytes in Tetravalent Salt Solutions as an Example of Study. *J. Phys. Chem. B* **2008**, *112*, 7347–7350.
- (19) Grosberg, A. Y.; Nguyen, T. T.; Shklovskii, B. I. Colloquium: The physics of charge inversion in chemical and biological systems. *Rev. Mod. Phys.* **2002**, *74*, 329–345.
- (20) Angelini, T. E.; Liang, H.; Wriggers, W.; Wong, G. C. L. Likecharge attraction between polyelectrolytes induced by counterion charge density waves. *Proc. Natl. Acad. Sci. U. S. A.* **2003**, *100*, 8634–8637
- (21) Jiang, T.; Wu, J. Ionic Effects in Collapse of Polyelectrolyte Brushes. J. Phys. Chem. B 2008, 112, 7713-7720.
- (22) Guptha, V. S.; Hsiao, P.-Y. Polyelectrolyte brushes in monovalent and multivalent salt solutions. *Polymer* **2014**, *55*, 2900–2912.
- (23) Csajka, F. S.; Seidel, C. Strongly Charged Polyelectrolyte Brushes: A Molecular Dynamics Study. *Macromolecules* **2000**, 33, 2728–2739.
- (24) Yoo, J.; Aksimentiev, A. The structure and intermolecular forces of DNA condensates. *Nucleic Acids Res.* **2016**, 44, 2036–2046.
- (25) Hsiao, P.-Y.; Luijten, E. Salt-Induced Collapse and Reexpansion of Highly Charged Flexible Polyelectrolytes. *Phys. Rev. Lett.* **2006**, *97*, 148301.
- (26) Feng, L.; Laderman, B.; Sacanna, S.; Chaikin, P. Re-entrant solidification in polymer-colloid mixtures as a consequence of competing entropic and enthalpic attractions. *Nat. Mater.* **2014**, *14*, 61–65.
- (27) Hur, J.; Witte, K. N.; Sun, W.; Won, Y.-Y. On the Origins of the Salt-Concentration-Dependent Instability and Lateral Nanoscale Heterogeneities of Weak Polyelectrolyte Brushes: Gradient Brush Experiment and Flory-Type Theoretical Analysis. *Langmuir* **2010**, *26*, 2021–2034.
- (28) Lai, P.-Y.; Binder, K. Structure and dynamics of polymer brushes near the Θ point: a Monte Carlo simulation. *J. Chem. Phys.* **1992**, *97*, 586–595.
- (29) Grest, G. S.; Murat, M. Structure of grafted polymeric brushes in solvents of varying quality: a molecular dynamics study. *Macromolecules* **1993**, *26*, 3108–3117.
- (30) Williams, D. R. M. Grafted polymers in bad solvents: octopus surface micelles. *J. Phys. II* **1993**, *3*, 1313–1318.
- (31) Zhulina, E.; Singh, C.; Balazs, A. C. Behavior of tethered polyelectrolytes in poor solvents. *J. Chem. Phys.* **1998**, *108*, 1175–1183.
- (32) Carrillo, J.-M. Y.; Dobrynin, A. V. Morphologies of Planar Polyelectrolyte Brushes in a Poor Solvent: Molecular Dynamics Simulations and Scaling Analysis. *Langmuir* **2009**, *25*, 13158–13168.
- (33) He, G.-L.; Merlitz, H.; Sommer, J.-U. Molecular dynamics simulations of polyelectrolyte brushes under poor solvent conditions: Origins of bundle formation. *J. Chem. Phys.* **2014**, *140*, 104911.
- (34) Pattanayek, S. K.; Pham, T. T.; Pereira, G. G. Morphological structures formed by grafted polymers in poor solvents. *J. Chem. Phys.* **2005**, *122*, 214908.
- (35) Tagliazucchi, M.; de la Cruz, M. O.; Szleifer, I. Self-organization of grafted polyelectrolyte layers via the coupling of chemical equilibrium and physical interactions. *Proc. Natl. Acad. Sci. U. S. A.* **2010**, *107*, 5300–5305.

(36) Crozier, P. S.; Stevens, M. J. Simulations of single grafted polyelectrolyte chains: ssDNA and dsDNA. *J. Chem. Phys.* **2003**, *118*, 3855–3860.

- (37) Ahrens, H.; Förster, S.; Helm, C. A.; Kumar, N. A.; Naji, A.; Netz, R. R.; Seidel, C. Nonlinear Osmotic Brush Regime: Experiments, Simulations and Scaling Theory. *J. Phys. Chem. B* **2004**, *108*, 16870–16876.
- (38) Kumar, N. A.; Seidel, C. Polyelectrolyte Brushes with Added Salt. *Macromolecules* **2005**, *38*, 9341–9350.
- (39) Limbach, H. J.; Arnold, A.; Mann, B. A.; Holm, C. ESPResSo An Extensible Simulation Package for Research on Soft Matter Systems. *Comput. Phys. Commun.* **2006**, *174*, 704–727.
- (40) Arnold, A.; Lenz, O.; Kesselheim, S.; Weeber, R.; Fahrenberger, F.; Roehm, D.; Košovan, P.; Holm, C. ESPResSo 3.1 Molecular Dynamics Software for Coarse-Grained Models. Meshfree Methods for Partial Differential Equations VI. 2013; pp 1–23.
- (41) Veitshans, T.; Klimov, D.; Thirumalai, D. Protein folding kinetics: timescales, pathways and energy landscapes in terms of sequence-dependent properties. *Folding Des.* **1997**, *2*, 1–22.
- (42) Hyeon, C.; Thirumalai, D. Mechanical unfolding of RNA hairpins. *Proc. Natl. Acad. Sci. U. S. A.* **2005**, *102*, 6789–6794.
- (43) Hyeon, C.; Thirumalai, D. Forced-unfolding and force-quench refolding of RNA hairpins. *Biophys. J.* **2006**, *90*, 3410–3427.
- (44) Hyeon, C.; Thirumalai, D. Multiple probes are required to explore and control the rugged energy landscape of RNA hairpins. *J. Am. Chem. Soc.* **2008**, *130*, 1538–1539.
- (45) Honeycutt, J. D.; Thirumalai, D. The nature of folded states of globular proteins. *Biopolymers* **1992**, *32*, *6*95–709.
- (46) Hinczewski, M.; von Hansen, Y.; Dzubiella, J.; Netz, R. R. How the diffusivity profile reduces the arbitrariness of protein folding free energies. *J. Chem. Phys.* **2010**, *132*, 245103.
- (47) Lefebvre, X.; Palmeri, J.; David, P. Nanofiltration Theory: An Analytic Approach for Single Salts. *J. Phys. Chem. B* **2004**, *108*, 16811–16824
- (48) Andreasson, B.; Nordenskiöld, L.; Schultz, J. Interactions of spermidine and methylspermidine with DNA studied by nuclear magnetic resonance self-diffusion measurements. *Biophys. J.* **1996**, *70*, 2847–2856.
- (49) Fernandes, M. X.; Ortega, A.; López Martínez, M. C.; García de la Torre, J. Calculation of hydrodynamic properties of small nucleic acids from their atomic structure. *Nucleic Acids Res.* **2002**, *30*, 1782–1788.
- (50) de Joannis, J.; Arnold, A.; Holm, C. Electrostatics in periodic slab geometries. I+II. *J. Chem. Phys.* **2002**, *117*, 2496–2502.
- (51) Arnold, A.; Holm, C. MMM2D: A fast and accurate summation method for electrostatic interactions in 2D slab geometries. *Comput. Phys. Commun.* **2002**, *148*, 327–348.
- (52) Aronovitz, J. A.; Nelson, D. R. Universal features of polymer shapes. J. Phys. (Paris) 1986, 47, 1445–1456.
- (53) Hyeon, C.; Dima, R. I.; Thirumalai, D. Size, shape, and flexibility of RNA structures. *J. Chem. Phys.* **2006**, *125*, 194905.
- (54) Lee, N. K.; Thirumalai, D. Dynamics of collapse of flexible polyelectrolytes in poor solvents. *Macromolecules* **2001**, *34*, 3446–3457.
- (55) Hansen, J.-P.; McDonald, I. R. *Theory of Simple Liquids*, 3rd ed.; Academic Press: 2006.
- (56) Kang, H.; Yoon, Y.-G.; Thirumalai, D.; Hyeon, C. Confinement-induced glassy dynamics in a model for chromosome organization. *Phys. Rev. Lett.* **2015**, *115*, 198102.
- (57) Stevens, M. J. Bundle binding in polyelectrolyte solutions. *Phys. Rev. Lett.* **1999**, 82, 101.
- (58) Sayar, M.; Holm, C. Finite-size polyelectrolyte bundles at thermodynamic equilibrium. *Europhys. Lett.* **2007**, *77*, 16001.
- (59) Fazli, H.; Golestanian, R. Aggregation kinetics of stiff polyelectrolytes in the presence of multivalent salt. *Phys. Rev. E* **2007**, *76*, 041801.
- (60) Stevens, M. J. Simple Simulations of DNA Condensation. *Biophys. J.* **2001**, *80*, 130–139.

(61) Hud, N. V.; Downing, K. H. Cryoelectron microscopy of λ phage DNA condensates in vitreous ice: The fine structure of DNA toroids. *Proc. Natl. Acad. Sci. U. S. A.* **2001**, 98, 14925–14930.

- (62) Lyubartsev, A. P.; Nordenskiöld, L. Monte Carlo Simulation Study of DNA Polyelectrolyte Properties in the Presence of Multivalent Polyamine Ions. *J. Phys. Chem. B* **1997**, *101*, 4335–4342.
- (63) Yu, J.; Mao, J.; Yuan, G.; Satija, S.; Jiang, Z.; Chen, W.; Tirrell, M. Structure of Polyelectrolyte Brushes in the Presence of Multivalent Counterions. *Macromolecules* **2016**, *49*, 5609–5617.
- (64) Yoo, J.; Kim, H.; Aksimentiev, A.; Ha, T. Direct evidence for sequence-dependent attraction between double-stranded DNA controlled by methylation. *Nat. Commun.* **2016**, *7*, 11045.