Spatiotemporal Formation Kinetics of Polyelectrolyte Complex Micelles with Millisecond Resolution

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ABSTRACT: We have directly observed the *in situ* self-assembly kinetics of polyelectrolyte complex (PEC) micelles by synchrotron time-resolved small-angle X-ray scattering, equipped with a stopped-flow device that provides millisecond temporal resolution. A synthesized neutral-charged diblock polycation and homopolyanion that we have previously investigated as a model charge-matched, core-shell micelle system were selected for this work. The initial micellization of the oppositely charged polyelectrolytes was completed within the dead time of mixing of 100 ms, followed by micelle growth and equilibration up to several seconds. By combining the structural evolution of the radius of gyration (R_g) and aggregation number (N) with complementary molecular dynamics simulations, we develop new information on how the self-assemblies evolve incrementally in size over time through a two-step kinetic process: first, oppositely-charged polyelectrolyte chains pair to form nascent aggregates that immediately assemble into spherical micelles, and second, these PEC micelles grow into larger micellar entities. This work has determined one possible kinetic pathway for the initial formation of PEC micelles, which provides useful physical insights for increasing fundamental understanding self-assembly dynamics driven by polyelectrolyte complexation that occur on ultrafast timescales.

Under selective solution conditions, amphiphilic block polymers can be designed to self-assemble into an assortment of diverse morphological structures, depending on the relative length, chemical nature, and molecular architecture of the hydrophilic and hydrophobic polymer blocks.^{1,2} Equilibrium structures have received far more attention than the time-dependent phenomena of their formation. Theoretical works by Aniansson and Wall,³⁻⁵ Halperin and Alexander,⁶ and Dormidontova^{7,8} have proposed two possible mechanisms for the kinetics of polymeric self-assemblies toward equilibrium: (i) single-chain insertion/expulsion and (ii) micelle fusion/fission. Experimental evidence has tested these model frameworks by adjusting solvent quality to induce micelle formation. For instance, using millisecond time-resolved small-angle X-ray scattering (TR-SAXS) under stopped-flow to provide nanoscale spatial resolution and temporal resolution, Lund and coworkers reported that the initial micellization kinetics of poly(ethylenealt-propylene)-block-poly(ethylene oxide) (PEP-b-PEO) can be viewed as a fast nucleation (within 10 ms) and a slow growth process, where the elemental PEP-b-PEO aggregate growth follows a single-chain insertion mechanism akin to the Aniansson-Wall theory.9 By comparison, Kalkowski et al. investigated micellization kinetics of poly(ethylene glycol)-block-poly(caprolactone) (PEG-b-PCL) with synchrotron X-ray scattering equipped with a microfluidic device.¹⁰ In this system, they distinguished three successive stages of the micellization (nucleation, micelle fusion, and polymer insertion), evident by the temporal evolution of the radii of gyration (R_{2}). From 0-250 ms, the R_{g} increased steadily as free unimers coupled into nuclei. Immediately after 250 ms, the R_{g} abruptly doubled in size, indicating that nascent nuclei began merging together, followed by steady structural growth from free PEG-*b*-PCL chains. While distinct from the Halperin-Alexander and Lund reports, the formation kinetics are in good agreement with the particle-particle fusion model described by Dormidontova.⁸ Further advanced techniques such as *in situ* liquid-cell transmission electron microscopy and *in silico* simulations have captured such collision events in solution.¹¹ In short, understanding kinetic formation in terms of tunable fundamental variables (e.g., polymer size/composition, concentration, interfacial tension) has unlocked design principles to more easily prepare nanoparticle assemblies with greater fidelity, control, reproducibility, and stability over time for intended applications.

However, compared to the advances in understanding the assembly kinetics of micelles formed by uncharged amphiphilic polymers over the past decade, 6,8,12-18 little is known about the initial moments of self-assembly of polyelectrolyte complexation-driven systems. Polyelectrolyte complexes (PECs) are prepared from an associative phase separation (i.e., coacervation) process between oppositely charged polyelectrolytes in water; for block polyelectrolytes, the process is thought to be more complicated due to the interplay among multiple components. including the polycation/polyanion pairing (e.g., polymer polarity, charge density, sequence effect, chirality, asymmetric block length ratios, etc.), high water content and related excluded volume considerations, and the presence of salts in physiological settings.19-21 Since the formation of PEC micelles is believed to be strongly driven by the entropically favorable liberation of counterions between oppositely charged moieties, it is reasonable to infer that there are parallels to the thermodynamics of homopolyelectrolyte complexation.^{22,23} Takahashi and coworkers investigated the complexation kinetics of oppositely charged homopolyelectrolytes, sodium poly(acrylate) and poly(allylamine hydrochloride), in aqueous NaCl solutions with TR-SAXS coupled with a stopped-flow device.²⁴ They reported the evolution of the complex droplet size, structure, and molar mass in a time range from 2.5 to 8733 ms, following (*i*) an immediate (< 2.5 ms) pairing step where oppositely charged chains pair by electrostatic interactions; (*ii*) a complexation step where the ion pairs further coalesce into nearly neutral aggregates, driven by van der Waals and hydrophobic effects; (*iii*) a growth step where aggregates grew analogously to the Brownian-coagulation kinetics of spherical colloidal particles. The same group also later revealed morphological transitions of cylindrical-tospherical PEC micelles with salt by TR-SAXS,²⁵ similarly to amphiphilic micelles.²⁶

In general, the self-assembly process of PEC micelles from well-defined polyelectrolyte systems has remained unclear. In this Letter, by employing in situ TR-SAXS with millisecond temporal resolution, we experimentally investigate the self-assembly of one set of well-defined PEC micelles as a starting point toward building more quantitative predictions of formation kinetics. The constituents of this model PEC micelle system are poly(ethylene oxide)-block-poly(vinyl benzyl trimethylammonium chloride) (PEO-b-PVBTMA) and sodium poly(acrylate) (PAA), shown in Figure 1-A. We have previously demonstrated the controlled fabrication of polyelectrolytes in parallel synthesis27 and interrogated aspects of the welldefined equilibrium structures of this pairing in the dilute regime.^{28,29} TR-SAXS experiments were carried out on the Biological Small Angle Scattering Beamline BL 4-2 at the Stanford Synchrotron Radiation Lighthouse (SSRL), SLAC National Accelerator Laboratory.^{30,31} The TR-SAXS facility at the SSRL BL4-2, equipped with a customized Bio-Logic 4 syringe stopped-flow mixer (SFM 400), provides access to solution kinetics on the millisecond time scale. Briefly, a typical TR-SAXS experiment proceeds in the following manner, shown schematically in Figure 1-B. First, aqueous solutions of positively-charged polymers and negatively-charged polymers were loaded in two separate syringes. Second, equal volumes of polvcation and polyanion solutions were pumped into the mixer by the motors, corresponding to the equimolar concentration of the cationic and anionic monomer units. The dead time was ~ 4 ms. Third, 30 µL of mixed solution was further dispensed into the capillary cell, where the incident X-ray radiates through. The sample solution was sealed and buffered by solvent on the two ends of the capillary cell. The exposure time for each measurement was set to 20 ms. Automated capillary cleaning assisted in checking for potential radiation damage between experiments. Additional details are provided in the Supporting Information.

The time evolution of the SAXS curves is presented in Figure 2. Three independent kinetic experiments were carried out from the 5K53-158 sample, where the diblock polycation was PEO_{5K} -*b*-PVBTMA₅₃ (molar mass of $PEO = 5000 \text{ g mol}^{-1}$; degree of polymerization of PVBTMA = 53), the polyanion was PAA₁₅₈ (degree of polymerization of PAA = 158), and the total polymer concentration was 2.5 mg mL⁻¹. To achieve high time resolution, we used a low exposure time of 20 ms, which may reduce the quality of the scattering curves due to noise. To avoid this, we independently repeated the same experiment three

times. It is shown that the scattering data are reliable and reproducible even under a very low exposure time of 20 ms. Collectively, all TR-SAXS patterns exhibited features of isolated particles with spherical morphology, well fitted by a polydisperse core-shell sphere model (see the Supporting Information). The increase in the scattering intensity over time reflects the micelle growth captured in real time. Due to the limitations of the instrument setup and experimental conditions, the lower limit of the experimentally probed time period remained 100 ms. Thus, the nucleation event for chain complexation was completed by 80 ms and was not experimentally observable.



Figure 1. (A) Polyelectrolyte complex micelles consist of a polycation, PEO-*b*-PVBTMA, and polyanion, PAA. (B) Schematic representation of the time-resolved small-angle X-ray scattering experiments equipped with a stopped-flow setup.

Nevertheless, the structural evolution of the 5K53-158 PEC micelles provided insightful information in the time range from 100 to 5000 ms. At 100 ms, well-formed micellar aggregates with a spherical geometry were already present in solution. The scattering intensity values increased steadily over time while maintaining the overall shape until 5000 ms, when the change in scattering intensity was barely discernable and indicated that the micelle growth became exceptionally slow relative to the prior state. The final state of micelles formed after injection in the stopped-flow path was sampled with SAXS after several minutes and did not differ (see Supporting Information). We also investigated the effects of block length and polymer concentration on the formation kinetics of PEC micelles, particularly on the growth rate. This involved a longer polycation, PEO_{10K}-b-PVBTMA₁₀₀ with PAA₁₅₈ (10K100-158) at 2.5 and 3.75 mg mL⁻¹. In anticipation of slower growth kinetics, the exposure time was set to 80 ms in order to extend experimental time window for TR-SAXS characterization. The recorded time was 19,200 s and 38,400 s for the 2.5 and 3.75 mg mL^{-1} polymer concentrations, respectively. We observed that qualitatively, 10K100-158 samples formed larger micellar aggregates at a slower overall rate at both concentrations.



Figure 2. TR-SAXS curves showing the formation of PEC micelles by (A) 5K53-158 at 2.5 mg mL⁻¹ total polymer concentration, and (B) 10K100-158 at 2.5 and (C) 3.75 mg mL⁻¹ total polymer concentration, respectively. Data points represent experimental data; see Figure S-X for fittings to a polydisperse core-shell model.

To better understand the assembly mechanism, the micelle size evolution over time was quantified. The R_g of the micellar aggregates over time was extracted via the Guinier approximation using the GNOM package³² as we have previously described²⁸ The aggregation number (N) was determined from fitting to a polydisperse core-shell sphere model. Three independent measurements of 5K53-158 formation showed a similar time dependence. As seen in Figure 3, we observe that at within the initial 100 ms, PEC micelles have already formed with $R_{\rm g}$ = 10.5 nm. Afterward, these micellar aggregates continued to grow in size, steadily increasing by 14% in Rg over 5000 ms. By dynamic light scattering (see Supporting Information), preassembled micelles prepared from these polyelectrolytes generally exhibit an apparent hydrodynamic radius $(R_{\rm h})$ of ca. 30.0 nm, corresponding to a shape factor (R_g/R_h) of 0.4. This corresponds to a highly water solvated structure, well below the conventional reference of a solid sphere $(R_g/R_h = \sqrt{3/5} \sim 0.77)$.³³ For comparison, Heo et al. have reported molecular conformation measurements of multiple post-polymerization functionalized PEC micelles, which comprised roughly 80% water by neutron scattering. ³⁴ A similar trend in $R_{\rm g}$ and N versus time was found in the 10K100-158 samples in Figure 4. The kinetic equilibration process appears slower for the micelles with longer polycation block lengths. For the 5K53-158 micelle, the $R_{\rm g}$ values plateaued around 5000 ms, whereas for the 10K100-158 system, the $R_{\rm g}$ increased by 27% from 11.0 nm at 5000 ms to 14.0 nm at 38,400 ms. We attribute this delay to the higher activation energy barrier and longer time involved with polymer rearrangement in longer chains.

From the aforementioned amphiphilic block polymer literature, there are two conventional mechanisms for micelle formation: single-chain insertion/expulsion or micelle fusion/fission. Micelles formed by oppositely charged polymers have features that are distinct from uncharged polymers. According to the current theories of polyelectrolyte complexation, free individual chains are driven to form complexes with their oppositely charged counterparts due to predominately entropic contributions,³⁵⁻³⁷ even at extremely low concentrations. Under this established framework, free chains are unlikely to exist as unbound chains when micelles have already formed. In alignment with time-resolved formation kinetics using homopolyelectrolytes,³⁸ it is reasonable to infer that a similar entropically-driven process occurs with block polyelectrolytes: immediately upon mixing, oppositely charged blocks pair with each other to form effectively neutralized clusters within a few milliseconds. Chain exchange events may happen immediately after complexation, but given the short time and length scales, this is not experimentally observable, even with advanced synchrotron capabilities. However, in the hundred-millisecond time periods, the experimental data suggests that it is unlikely that micelle fusion/fission took place. Otherwise, the $R_{\rm g}$ values of the micellar aggregates would have exhibited an abrupt jump in magnitude, as Kalkowski et al. have observed in the (1) nucleation, (2) fusion, and (3) insertion stages of PEG-b-PCL micelle formation; here, the Guinier R_g correspondingly evolved from (1) \sim 2-5 nm to (2) \sim 5-9 nm, before (3) plateauing at \sim 10 nm.¹⁰



Figure 3. The evolution of the fitted Guinier R_g for 5K53-158 (red circles) in TR-SAXS experiments over the time range of 0 to 5000 ms. Error bars denote the standard deviation of each measurement.



Figure 4. The evolution of the fitted Guinier R_g for 10K100-158 (red circles) at (A) 2.5 mg mL⁻¹ and (B) 3.75 mg mL⁻¹ over the time range of 0 to 40,000 ms. Error bars denote the standard deviation of each measurement. The final R_g and N are X and X, respectively.

Furthermore, when we repeated TR-SAXS measurements at a higher 3.75 mg mL⁻¹ total polymer concentration, the slow growth of R_g remained invariant over time (Figure 4B). This implies that the formation of dilute PEO-*b*-PVBTMA/PAA micelles is largely concentration-independent. Previous work on electrostatic assemblies have shown a strong dependence on polymer concentration due to fusion-fission mechanisms.³⁹

Because the PEC micelles were highly solvated and completely hydrophilic, this result was initially puzzling. However, we have previously shown that this PEO-*b*-PVBTMA/PAA system exhibits repulsive interparticle effects at dilute conditions using SAXS.²⁸ Physically, we demonstrated that when two micelles approach, the PEO coronas may be compressed to some extent, but the depth of the compressed area is only half of the corona thickness. In other words, the micelle cores were unable to reach each other, which supports the observations that PEC micelle fusion/fission is improbable during the formation process. Similar steric arguments have been made by theories for certain amphiphilic block polymer micelles that do not exhibit fusion-based growth mechanisms.⁶ At some critical stage the cluster insertion becomes dominant when micelle coronas are too large and dense to be penetrated.

To complement the experimental TR-SAXS kinetics conditions and assess the relationship between R_g and N in this time regime, a charged Kremer-Grest model⁴⁰ was employed for the coarse-grained molecular dynamics (MD) simulation of polyelectrolyte coacervates. Polymers were simulated as beadspring chains. For simplicity in maintaining electroneutrality, the 5K53 chain was simulated as a diblock with 110 and 50 monomer beads for the neutral and positively charged blocks, respectively; the homopolyanion comprised 150 negativelycharged monomer beads. Each bead was identical in size and interacted via the shifted and truncated Lennard-Jones potential. The connectivity of polymer beads was maintained by the finitely extensible nonlinear elastic (FENE) potential. The saltfree solvent was treated implicitly with a dielectric constant ε = 78.5, and counterions were neglected. Spherical micelles at given aggregation numbers were pre-assembled and placed at the center of the simulation box. Pre-assembly was achieved by first uniformly placing the polymer chains on a spherical surface of given radius with the charged blocks pointing towards the center of the sphere. An energy minimization of the system was then performed using a conjugated gradient algorithm implemented in a large-scale atomic/molecular massively parallel simulator (LAMMPS⁴¹) with 10⁶ MD steps for equilibration. Afterward, a production run of 10⁶ timesteps was used to collect data every 10⁴ timesteps to calculate the R_g , radial density profile, and radial distribution function of the micelles. A detailed simulation methodology is provided in the Supporting Information.

The MD simulation results in Figure 5 show how the $R_{\rm g}$ and asphericity parameter (k²) of the PEC micelle correlates to the change in specified N. Given three selected aggregation numbers (N = 75, 105, and 135), it is observed that the R_g increased from 8.25 ± 0.98 nm to 9.66 ± 0.88 nm (i.e., a 17% difference), which is qualitatively consistent with experimental estimates of a 14% increase in R_g growth. Differences in the absolute R_g values from experiment can be attributed to the coarse-grained and implicit solvent nature of the simulations. Near zero values of the asphericity parameter (Figure 5) combined with the computed radial distribution functions (Figure S-5) indicated that at all molecular weights micelles exhibit spherical morphologies with similar internal structures. Further computational efforts can help capture the initial micellization kinetics that are not experimentally observable. For instance, Bos and Sprakel recently employed Langevin dynamics simulations to study fast exchange dynamics.42 In the early simulation time steps of the micellization stage, they found that initially, polyelectrolyte chains adopted a relatively stretched configuration before rapidly forming aggregates by both expulsion/insertion and fusion/fission events, revealed by the decrease of size and release of counterions. Fusion and fission events were defined by the expulsion of small neutralized clusters containing a few polyelectrolytes from one micelle into another micelle.



Figure 5. MD simulation results of the R_g (orange data with dashed arrow to the left axis) and asphericity (green data with dotted arrow to the right axis) as a function of aggregation number for the 5K53-158 system. The inset shows a snapshot of the micelle using LAMMPS analysis.

Combining the experimental and simulation results with literature evidence, we propose that a possible kinetic pathway of micellization can be described in a two-stage process, as illustrated in Scheme 1. First, immediately upon mixing in the stopped-flow apparatus, oppositely charged polyelectrolytes form charge-neutralized clusters. From these cluster aggregates, although the ultrafast time scale of the PEC micellization process was not commensurate with TR-SAXS, well-defined spherical micelles with $R_g \sim 10-11$ nm. Second, the micelle products equilibrate by exchanging either unimer chains or ionpaired clusters, as evident by incremental growth in $R_{\rm g}$ and N. During this process, we believe that fusion events are highly unlikely due to repulsive interparticle interactions that this system has been shown to exhibit. The universality of this kinetic pathway is the subject of ongoing investigations. In the former step, the mixing method of the stopped-flow apparatus and the ionic strength of the solution may play key roles in adjusting the timescale of cluster formation proceeding the PEO-b-PVBTMA and PAA micellization. For the latter step, greater scrutinization of the polyelectrolytes' physical attributes, e.g., varying the ratio and the total chain lengths of blocks, may reveal differences in the likelihood of micelle fusion/fission, and therefore, alter the spatiotemporal evolution of micelles upon equilibration.

Scheme 1. Illustrated kinetic pathways of the investigated PEC micelles.



Furthermore, the formation kinetics may be strongly dependent on the chemical nature of the polymer constituents themselves. Charged heteropolymers and sequence-defined macromolecules give rise to distinct hierarchical assemblies at equilibrium,43 depending on the local interactions and sequence variation of the primary structure. By extension, the dynamics of PEC micelle formation, chain exchange, and equilibration are also likely highly influenced by how oppositely-charged polyelectrolytes associate at the molecular level. With this vast parameter space, greater systematic exploration of different polyelectrolytes is vital for more quantitative prediction capabilities and theories of complex coacervation. Recently, Amann et al. reported kinetic pathways for PEC micelle formation using TR-SAXS using PEO-b-PVBTMA (2K6, 2K8, 2K12, and 2K19) and poly(styrene sulfonate) (PSS₁₉).⁴⁴ Their data shows the formation of metastable aggregates, followed by rearrangement and pinch-off into smaller micellar-like entities. We hypothesize that this additional kinetic step arises because of the choice of the strongly-charged PSS polyanion. We have previously shown that upon complexation, PEO-b-PVBTMA forms micron-scale, kinetically frozen aggregates with PSS, in contrast to the spherical micelles with PAA.45 Furthermore, direct comparison of the homopolymer PVBTMA with PAA and PSS reveal liquid and solid complexes, respectively, by optical microscopy and SAXS (Figure S-1), which may account for subtle differences in the initial pairing step. Amann and coworkers also provide evidence that during the growth and equilibration step, fusion/fission events were not observed, in agreement with the subsequent behavior of PEO-*b*-PVBTMA/PAA. A direct comparison of formation pathways suggests that the early-stage micellization kinetics might be challenging to fully generalize, with the exact pathway likely to be highly dependent on the environmental and polymer parameters.

Altogether, this Letter provides an early study in understanding the ultrafast kinetics of ionic polymer assemblies. Further effort is underway to compare the kinetic pathway dependency of molecular attributes that are known to affect complex coacervation, such as the neutral-to-charged block lengths, the strength and type of the charged moieties, counterion valency, solvent quality, and other orthogonal association effects. Uncovering the underlying polymer physics from a fundamental molecular engineering standpoint can lead to new insights in an array of bioapplications. Such endeavors can potentially improve the loading capacity, robustness, and stability of non-viral vector formulations for proteins,⁴⁶ peptides,⁴⁷ and nucleic acids⁴⁸ in the future.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge on the ACS Publications website.

Materials, Methods (file type, i.e., PDF)

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Notes

The authors declare no competing financial interest. #These authors contributed equally to this work.

Author Contributions

All authors contributed to proofreading the manuscript. H.W. conceived and implemented the project, carried out the micelle formation and scattering experiments, drafted the initial manuscript, and analyzed the scattering data. J.M.T. performed the polymer synthesis, carried out polymer characterization, and conducted the scattering experiments. B.Y. conducted the molecular dynamics simulations and analysis. N.E.J. assisted with the development and interpretation of the simulations. S.M. assisted with the scattering experiments and analyzed the bulk complex materials. J.J.d.P. was the principle investigator and contributed to the data analysis and interpretation. M.V.T. was the principal investigator of the project and directed the research, managed the project, and contributed to the data analysis and interpretation of the experiments.

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