

Molecular Weight Dependence of Block Copolymer Micelle Fragmentation Kinetics

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Cite This: <https://doi.org/10.1021/jacs.1c02147>



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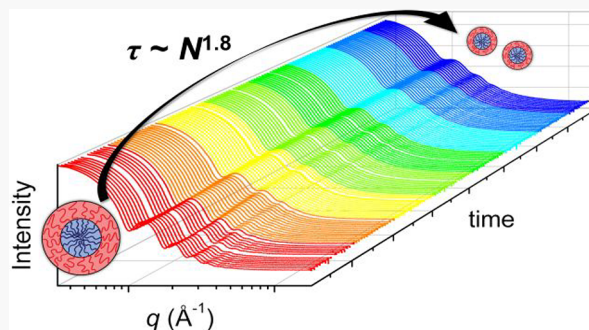


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ABSTRACT: The effect of molecular weight (M) on the fragmentation kinetics of micelles formed by 1,2-polybutadiene-*block*-poly(ethylene oxide) (PB-PEO) copolymers was studied in the ionic liquid 1-ethyl-3-methylimidazolium bis(trifluoromethylsulfonyl)imide. A series of six samples, with total M ranging from 10^4 to 10^5 g mol $^{-1}$ and nearly constant composition ($f_{\text{PEO}} \approx 0.4$), were examined; all six formed spherical micelles with PEO coronas. Nonequilibrium PB-PEO micelles were prepared by direct dissolution, a process that systematically produces nanoparticles with mean aggregation numbers more than twice the equilibrium values. When subjected to high temperature annealing (170 °C), the average micelle radius was found to decrease substantially, as determined by temperature-jump dynamic light scattering (T -jump DLS) and time-resolved small-angle X-ray scattering (TR-SAXS). The characteristic fragmentation times (τ) were found to increase strongly with increasing degree of polymerization N , as $\tau \sim N^{1.8}$. This result compares favorably with the prediction of a previously untested model.



INTRODUCTION

Diblock copolymers self-assemble into a variety of micellar structures in the presence of a block-selective solvent, i.e., a solvent that dissolves one block and not the other. Block copolymer (BCP) micelles are utilized in a wide range of applications including ion gels,^{1,2} drug or gene delivery,^{3,4} nanolithography,⁵ and viscosity modification.⁶ Because of the versatility of BCP micelles from an applications perspective, their dynamics and equilibration kinetics in solution are of interest, yet these processes remain less well understood in comparison to their low molecular weight counterparts.^{7–10} The thermodynamics of micellization has been studied extensively, both theoretically and experimentally.^{11,12} The morphology and size of block copolymer micelles at equilibrium can be tuned by changing the molecular characteristics of the block copolymer itself, such as the core and corona block lengths,^{13–16} and the solvent selectivity.¹⁷ There are five primary processes thought to govern the formation, destruction, and equilibration of block copolymer micelles in solution, namely, (i) *chain exchange*, where individual copolymer chains are exchanged between micelles, (ii) *fragmentation*, where a micelle much larger than the equilibrium aggregation number Q_{eq} breaks up into smaller micelles, (iii) *fusion*, where micelles smaller than Q_{eq} merge together, (iv) *micelle creation*, where individual chains nucleate a new micelle, and (v) *annihilation* of micelles, where a series of chain exchange/fragmentation/fusion events eliminates one micelle. The kinetics of these processes were first modeled by Aniansson, Wall, and co-workers.^{18,19} These processes of

micellization and equilibration are supported by experimental studies of low molecular weight surfactants²⁰ and Pluronics.^{21–24} Halperin and Alexander developed the first detailed theory of block copolymer micelle dynamics near equilibrium, where chain exchange was considered to be the only plausible equilibration mechanism, as fragmentation and fusion were argued to be much slower.^{25,26} Dormidontova proposed a different model that gives scaling predictions for the relaxation times of chain exchange, fragmentation, and fusion.²⁷ Nyrkova and Semenov focused on the rates of micelle creation and annihilation in response to abrupt changes in conditions such as jumps in temperature.²⁸

The molecular weight (M) of diblock copolymers not only affects the equilibrium structure of micelles, but also the dynamics and equilibration kinetics in solution.^{8,15,29} The aggregation number Q of a given micelle can vary, and micelle equilibration occurs by some combination of the exchange of individual chains,^{7,9,15,30–35} fusion, or fragmentation.^{20,22,23,36} Halperin and Alexander proposed that the chain exchange kinetics depends on a characteristic relaxation time, τ , that depends on the core and corona block lengths as $\tau \sim$

Received: February 24, 2021



$69 N_{\text{core}}^{2/25} N_{\text{corona}}^{9/5} \exp(E_a/k_B T)$, where E_a is the activation
70 barrier given by $\gamma N_{\text{core}}^{2/3} b^2$, where γ is the interfacial tension
71 between the core block and the solvent and b is the monomer
72 size of the core block.²⁵ Experimentally, the effect of M on
73 micelle chain exchange kinetics has been found to be very
74 strong, with the barrier to exchange increasing linearly with
75 core block degree of polymerization, N_{core} , in contrast to the
76 Halperin-Alexander model.^{8,32–34,37,38} Interestingly, it has also
77 been reported that increasing the degree of polymerization of
78 the corona block, N_{corona} , of PS-PEP micelles in squalane could
79 increase the rate of chain exchange by up to 2 orders of
80 magnitude,¹⁵ whereas studies of C_{27} -PEO_{*n*} in water showed
81 that increasing N_{corona} had the opposite effect.⁸

82 Despite the dramatic effect of M on chain exchange,
83 essentially nothing is known experimentally about the
84 influence of M on the kinetics of fragmentation.³⁶ In general,
85 fragmentation is much slower than chain exchange for micelles
86 that are not too far from equilibrium, thus increasing molecular
87 weight should lead to longer fragmentation time scales.^{22,23,36}
88 Techniques including dynamic light scattering (DLS) and
89 time-resolved small-angle X-ray scattering (TR-SAXS) are
90 useful for monitoring changes in micelle size over a wide range
91 of time scales.^{39–43} This is primarily due to the relatively short
92 acquisition times required for DLS (i.e., minutes), and
93 particularly for synchrotron SAXS (seconds). Previous work
94 by Kelley et al. showed that micellization of 1,2-polybutadiene-
95 block-poly(ethylene oxide) (PB-PEO) in mixtures of tetrahy-
96 drofuran and water occurs via a distinct bimodal pathway, and
97 the increase of the average hydrodynamic radius, $\langle R_h \rangle$, was
98 monitored using DLS for as long as 90 days.⁴¹ They concluded
99 that the initial growth of micelles occurs via micelle fusion.⁴¹
100 Additionally, the combination of DLS and TR-SAXS proves
101 invaluable for determining micelle fragmentation kinetics, as
102 the total micelle radius, i.e., $\langle R_h \rangle$, and the average micelle core
103 radius, $\langle R_{\text{core}} \rangle$, are readily determined. In general, chain
104 exchange by itself is not expected to change $\langle R_{\text{core}} \rangle$ to a
105 significant extent.^{36,44,45}

106 A theoretical model of the M dependence, specifically the
107 dependence on N_{corona} , of micelle fusion and fragmentation has
108 been reported.²⁷ Dormidontova proposed a scaling model to
109 account for micellization in systems far from equilibrium, and
110 assumes that micelle fragmentation proceeds by the reverse
111 mechanism as micelle fusion.²⁷ The relaxation time (τ) for
112 fusion of two micelles with similar aggregation numbers ($Q_1 \leq$
113 Q_2) scales with aggregation number and the degree of
114 polymerization of the corona block as $\tau \sim N_{\text{corona}}^{17/5} Q_1^{4/5} Q_2$.²⁷
115 The strong N_{corona} dependence on the characteristic time for
116 fusion is attributed to the corona chain deformation during the
117 merging process, where the corona chains of a smaller micelle
118 penetrate into the corona region of a larger micelle. On the
119 basis of this observation for micelle fusion, the fragmentation
120 times for micelles with small aggregation numbers were
121 estimated to scale with $N_{\text{corona}}^{17/5}$ as well. The fragmentation
122 time was predicted to scale as $\tau \sim N_{\text{corona}}^{9/5} Q^{13/5}$ for micelles
123 with very large aggregation numbers $Q > (cV_{\text{corona}})^{5/2} N_{\text{corona}}^2$,
124 where c is the micelle concentration and V_{corona} is the molar
125 volume of a corona-forming chain.

126 Previous work on the micellization of PB-PEO in the ionic
127 liquid (IL) 1-ethyl-3-methylimidazolium bis-
128 (trifluoromethylsulfonyl)imide ($[C_2\text{mim}][\text{TFSI}]$) showed
129 that the solution preparation method influences $\langle R_h \rangle$. For
130 micelles prepared by direct dissolution (DD) of the bulk
131 copolymer in the IL, the $\langle R_h \rangle$ of the as-prepared micelles is

quite large compared to micelles prepared by the cosolvent
method, in which a good solvent is introduced to the solution
and slowly evaporated away.⁴⁰ For micelles prepared by DD,
 $\langle R_h \rangle$ decreased to approximately half of the original size when
annealed at 170 °C.⁴⁰ Further work on this system showed that
the time-dependent decrease in $\langle R_h \rangle$ was consistently well-
described by a compressed exponential function $\exp(-(t/\tau)^n)$
with an exponent $n \approx 2$, even at lower annealing temperatures
($T = 120$ °C).³⁶ To further understand the mechanism, the
fragmentation kinetics for one molecular weight of PB-PEO
was studied in five 1-alkyl-3-methylimidazolium bis-
(trifluoromethylsulfonyl)imide-based ILs ($[C_x\text{mim}][\text{TFSI}]$)
where $x = 1, 2, 4, 6$, and 8).⁴⁵ The solvent quality with
respect to the PB core-forming block was improved by
increasing the length of the alkyl chain on the IL cation, but it
was found that the improvement in solvent quality did not
affect the fragmentation kinetics of PB-PEO micelles to any
significant extent.⁴⁵ We previously reported the direct
observation of micelle fragmentation for three molecular
weights of PB-PEO in $[C_2\text{mim}][\text{TFSI}]$ using high-temperature
liquid-phase transmission electron microscopy, and intermedi-
ate structures close to an apparent transition state for micelle
fragmentation were observed in situ.⁴⁶

The absence of experimental results on the molecular weight
dependence of fragmentation kinetics motivates this work.
Here we monitor the fragmentation of PB-PEO in the IL
 $[C_2\text{mim}][\text{TFSI}]$ where the total molecular weight of the
diblock is varied from 11 to 100 kg mol^{−1} while maintaining a
constant volume composition $f_{\text{PEO}} \approx 0.4$. The thermal stability
and nonvolatility of ILs is exploited here as the solutions can
be heated to much higher temperatures than organic solvents,
allowing the kinetics of fragmentation to be studied over more
accessible time scales.⁴⁷

EXPERIMENTAL SECTION

Synthesis and Characterization. Six PB-PEO copolymers were
synthesized by two-step sequential anionic polymerization,⁴⁸ as briefly
described here. 1,3-Butadiene (Sigma-Aldrich, $\geq 99\%$) and ethylene
oxide (Sigma-Aldrich, $\geq 99.5\%$) were purified by stirring with
vacuum-dried *n*-butyllithium (Sigma-Aldrich, 2.5 M in hexanes) twice
for 30 min. Six molecular weights (6–53 kg mol^{−1}) of hydroxyl-
terminated 1,3-polybutadiene were synthesized via anionic polymer-
ization of 1,3-butadiene with *sec*-butyllithium in tetrahydrofuran
(THF) at −75 °C for 8 h; the polymer chains were end-capped with a
single ethylene oxide unit at 27 °C, and the polymerization was
terminated by the addition of acidic methanol (1:10 HCl:methanol).
A concentrated solution of PB-OH in dichloromethane was
precipitated into an excess of cold methanol, isolated via vacuum
filtration, and dried under vacuum (<100 mTorr) at 40 °C for 72 h
prior to use. PB-PEO diblocks with a nearly constant volume fraction
of PEO ($f_{\text{PEO}} \approx 0.4$) were prepared by subsequent anionic
polymerization of ethylene oxide. The polymerization was performed
in THF at 40 °C for 24 h in the presence of PB-OH with potassium
naphthalenide. The polymerization was terminated by the addition of
acidic methanol (1:10 HCl:methanol). PB-PEO diblocks were freeze-
dried in benzene with 0.1 wt % BHT as an antioxidant under vacuum
(<100 mTorr) at 27 °C for 24 h prior to use. The samples are
denoted BO(*x*–*y*), where *x* and *y* indicate the number-average
molecular weights of the PB and PEO blocks in kg mol^{−1}, respectively.
The number-average molecular weight (M_n), dispersity (\bar{D}), and f_{PEO}
were determined by a combination of size exclusion chromatography
(SEC) in THF with a multiangle laser light scattering detector (Wyatt
Dawn Heleos II) and ¹H nuclear magnetic resonance spectroscopy in
CDCl₃ (¹H NMR, Varian Inova 500) as shown in Table 1. The
refractive index detector traces from SEC in THF of all diblocks are
shown in the Supporting Information (SI Figure S1). The refractive

Table 1. Characteristics of BO(x - y) Diblock Copolymers^a

sample	$M_{n,PB}$ (kg mol ⁻¹)	$M_{n,PEO}$ (kg mol ⁻¹)	f_{PEO}	\bar{D}
BO(6-5)	6.5	5.2	0.38	1.14
BO(8-7)	9.4	7.6	0.38	1.07
BO(10-9)	10	9.4	0.43	1.05
BO(25-22)	25	22	0.41	1.09
BO(27-27)	27	27	0.43	1.06
BO(53-46)	53	46	0.40	1.05

^a¹H NMR spectroscopy in CDCl₃ was used to determine the number-average molar mass (M_n) and volume fraction of PEO (f_{PEO}) using bulk densities $\rho_{PB} = 0.87$ g/cm³ for PB, and $\rho_{PEO} = 1.13$ g/cm³ for PEO. SEC with a multiangle light scattering detector in THF was used to determine the weight-average molar mass (M_w) and molar mass dispersity ($\bar{D} = M_w/M_n$) of the diblocks.

index increment (dn/dc) for a diblock was estimated as the weight-average of the refractive index increments for PB in THF ($dn/dc = 0.119$ mL/g) and PEO in THF ($dn/dc = 0.068$ mL/g).⁴⁹ The ¹H NMR spectra of the diblocks in deuterated chloroform (CDCl₃) are shown in Figure S2.

The ionic liquid (IL) 1-ethyl-3-methylimidazolium bis(trifluoromethylsulfonyl)imide ([C₂mim][TFSI]) was synthesized by an anion exchange reaction between 1-ethyl-3-methylimidazolium bromide and lithium bis(trifluoromethylsulfonyl)imide, carried out in water at 70 °C for 24 h.⁵⁰ The reaction mixture was diluted with dichloromethane, the aqueous layer was removed, and the organic layer was washed three times with deionized water. The organic layer was stirred on activated charcoal for 24 h. The solution was passed through an alumina column, dichloromethane was removed in vacuo, and [C₂mim][TFSI] was dried under vacuum (<100 mTorr) at 60 °C for 72 h. The dried IL was characterized by ¹H, ¹³C, and ¹⁹F NMR spectroscopy in DMSO-*d*₆. The ¹H NMR spectrum for [C₂mim][TFSI] is shown in Figure S3.

Solution Preparation by Direct Dissolution (DD). All solutions were prepared by the direct dissolution (DD) method described here. The desired amounts of PB-PEO and [C₂mim][TFSI] were combined by weight in a 20 mL scintillation vial equipped with a stir bar to obtain a 0.25 wt % solution. The vial was placed in an oil bath and stirred vigorously at 70 °C for 48 h. The resulting DD solutions were slightly blue-tinged for BO(6-5), BO(8-7), and BO(10-9) micelles, whereas those for BO(25-22), BO(27-27), and BO(53-46) were white-tinged.

Dynamic Light Scattering (DLS). Dynamic light scattering measurements were performed on a home-built light scattering setup equipped with a Brookhaven BI-DS photomultiplier mounted to an adjustable goniometer, a Lexel Ar⁺ laser ($\lambda = 488$ nm), and a Brookhaven BI-9000 correlator. During a typical annealing experiment, the temperature was controlled with an index-matching high-temperature silicone oil bath to within ± 0.1 °C. To verify the size of the micelles before and after annealing, multiangle light scattering experiments were performed at $T = 27$ °C with a range of scattering angles θ from 60° to 120°.

Each micelle solution was passed through an 0.45 μ m PTFE syringe filter to remove any dust, and subsequently flame-sealed under vacuum (≤ 70 mTorr) in a dust-free glass tube (I.D. 0.51 cm) to prevent degradation of the block copolymer and to avoid contact with moisture and dust. In an annealing experiment, the oil bath temperature was allowed to equilibrate for 30 min prior to introducing the sample. The normalized intensity autocorrelation function, $g_2(t)$, was measured at $\theta = 90^\circ$ as a function of time. For each time point, $g_2(t)$ was acquired for 90 s. The intensity autocorrelation function was converted to the electric field autocorrelation function, $g_1(t)$, via the Siegert relation,⁵¹ which was then fit to a second-order cumulant expansion for single populations to obtain the average decay rate, $\bar{\Gamma}$, and the variance, $\mu_2/\bar{\Gamma}^2$. The mutual diffusion coefficient D_m was calculated from $D_m = \bar{\Gamma}/q^2$, where q is the magnitude of the scattering vector defined by $q = (4\pi n/\lambda_0)\sin(\theta/2)$, where n is the refractive index of [C₂mim][TFSI], and λ_0 is the wavelength of light in a vacuum. For multiangle light scattering experiments, D_m was estimated by taking the slope of the line in plots of $\bar{\Gamma}$ versus q^2 with zero intercept. For dilute solutions, D_m can be used as an approximation of the tracer diffusion coefficient, D_t . This is used to calculate the hydrodynamic radius using the Stokes–Einstein equation:

$$R_h = \frac{k_B T}{6\pi\eta_s D_t} \quad (1)$$

where k_B is the Boltzmann constant, T is the temperature, and $\eta_s(T)$ is the solvent viscosity; $\eta_s(170$ °C) = 2.4 kPa·s for [C₂mim][TFSI].³⁶ The temperature dependence of the viscosity is well described by the Vogel–Fulcher–Tamman equation for [C₂mim][TFSI] and many other ILs.^{52–55} Additional analysis of the micelle size distribution was performed by applying the regularized positive exponential sum (REPES) Laplace inversion to the intensity correlation function.⁵⁶ Application of this algorithm results in the decay rate distribution, $G(\Gamma)$, which can be expressed in terms of the hydrodynamic radius from eq 1. When the REPES routine showed a sample with a bimodal distribution, $g_1(t)$ was fit to a double-exponential function, shown in eq 2, to obtain the decay rates of each population.

$$g_1(t) = A_1 \exp(-\bar{\Gamma}_1 t) + A_2 \exp(-\bar{\Gamma}_2 t) \quad (2)$$

Time-Resolved Small Angle X-ray Scattering (TR-SAXS). SAXS experiments were conducted at the 11-BM Complex materials Scattering (CMS) beamline at the National Synchrotron Light Source II (NSLS-II), Brookhaven National Laboratory. Samples of 0.25 wt % PB-PEO directly dissolved in [C₂mim][TFSI] were syringe-filtered into 1.5 mm diameter borosilicate capillaries (Charles Supper Co.) and then placed into a custom 15-capillary temperature-controlled stage. SAXS measurements were conducted with samples under dynamic vacuum at room temperature and 170 °C. Two-dimensional scattering patterns were obtained using a Dectris Pilatus 300k pixel-array detector (pixel size $172 \times 172 \mu\text{m}^2$) using a 30 s exposure time to 13.5 keV X-rays ($\lambda = 0.918$ Å) and a sample-to-detector distance of 2 m. The position where the incident beam hits the capillary was changed throughout a time-resolved experiment to mitigate beam damage effects. The scattering vector q was calibrated using silver behenate. The 2D data were azimuthally averaged to yield 1D scattering patterns as intensity versus q . Due to the presence of higher q upturns in ionic liquid scattering, which has been observed previously, and significant capillary-to-capillary variations, the background (i.e., ionic liquid and capillary scattering) was fit to a power law ($I(q) = A + Bq^{-m}$, where $2 \leq m \leq 4$) and subtracted from the solution scattering data.⁵⁷ The background-corrected intensity traces were fit using an established block copolymer micelle model.^{58,59} In this model, the scattering intensity is defined as follows:

$$I(q) = \int D(R_{\text{core}}) [P_{\text{mic}}(q) + A_{\text{mic}}^2(q)(S(q) - 1)] dR_{\text{core}} \quad (3)$$

where R_{core} is the micelle core radius with distribution $D(R_{\text{core}})$, $P_{\text{mic}}(q)$ is the spherical form factor for the micelle core,⁵⁸ $A_{\text{mic}}(q)$ is the form factor amplitude, and $S(q)$ is the hard-sphere structure factor with the Percus–Yevick closure.

SAXS experiments for the bulk BO copolymers were conducted at the 5-ID-D beamline of the Dupont-Northwestern-Dow Collaborative Access Team (DND-CAT) at the Advanced Photon Source, Argonne National Laboratory. All samples were hermetically sealed in aluminum DSC pans under argon with <0.1 wt % BHT as an antioxidant. The samples were annealed at 70 °C for 10 min prior to measurement. SAXS measurements of bulk polymers were conducted at 70 °C. Two-dimensional scattering patterns were obtained using a Rayonix MX170-HS CCD area detector using a 0.5 s exposure time to X-rays with a wavelength of $\lambda = 0.729$ Å and a sample-to-detector distance of 8.5 m. The 2D data were azimuthally averaged to yield 1D scattering patterns as intensity versus q .

RESULTS AND DISCUSSION

Initial and Final Micelle Dimensions. SAXS and DLS were used to determine the effect of molecular weight on the size of micelles, both as prepared by direct dissolution and also at steady-state after annealing at 170 °C. The initial size of micelles prepared by DD and after a T -jump to 170 °C were determined by multiangle DLS and SAXS, in terms of the overall micelle radius R_h and the micelle core radius R_{core} , respectively. The overall radii were determined by fitting the intensity autocorrelation function from DLS to a second order cumulant expansion. The radii increased monotonically with increasing M , as seen in SI Figures S4 and S5. REPES analysis of DLS data, shown in Figure S4, for the as-prepared and steady-state $\langle R_h \rangle$ of 0.25 wt % PB-PEO in $[C_2mim][TFSI]$ corroborates the fitting results in that the $\langle R_h \rangle$ decreases after annealing at 170 °C for all molecular weights. The percent decrease in $\langle R_h \rangle$ after long-time annealing at 170 °C was 31%, 36%, 33%, 17%, 25%, and 29% for BO(6–5), BO(8–7), BO(10–9), BO(25–22), BO(27–27), and BO(53–46), respectively.

To determine the effect of M on R_{core} , SAXS was performed at room temperature before and after annealing at 170 °C. The background corrected scattering intensity traces versus q for the as-prepared and steady-state micelles are shown in Figure 1a and 1b, respectively. The as-prepared micelles for all

polymers, except for BO(53–46), are well-defined spheres. This is confirmed by the distinct first minimum in the form factor and oscillations at higher q values, which generally reflect a narrow size distribution and a modest interfacial width at the core–corona interface.^{58,60} The increased size dispersity of BO(53–46) micelles may be due to the fixed time used to prepare the solutions, and longer dissolution times may be required to obtain more narrowly distributed initial micelles. Similarly, well-defined smaller spherical micelles are obtained after annealing at 170 °C, as shown in Figure 1b, for all but the largest polymer.

As with the total micelle radius, $\langle R_{core} \rangle$ from SAXS was found to increase monotonically with M for the as-prepared and steady-state micelles, as shown in Figure 2. The changes in

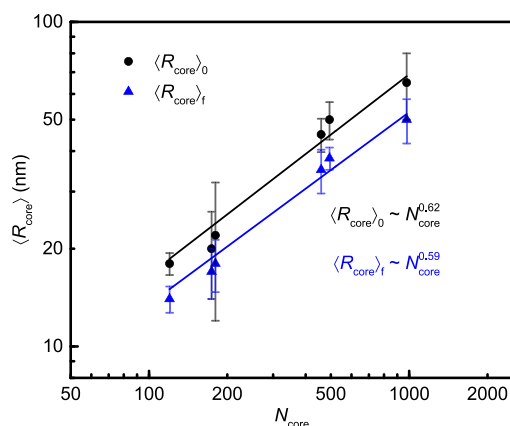


Figure 2. Scaling of $\langle R_{core} \rangle$ versus N_{core} for 0.25 wt % PB-PEO in $[C_2mim][TFSI]$ as prepared by DD (black circles) and steady-state after annealing at 170 °C (blue triangles). The solid lines for the as prepared and steady-state values represent the scaling obtained for $\langle R_{core} \rangle_0 \sim N_{core}^{0.62}$ (black line) and $\langle R_{core} \rangle_f \sim N_{core}^{0.59}$ (blue line). The error bars represent $\pm \sigma_{core}$ shown in Table 2.

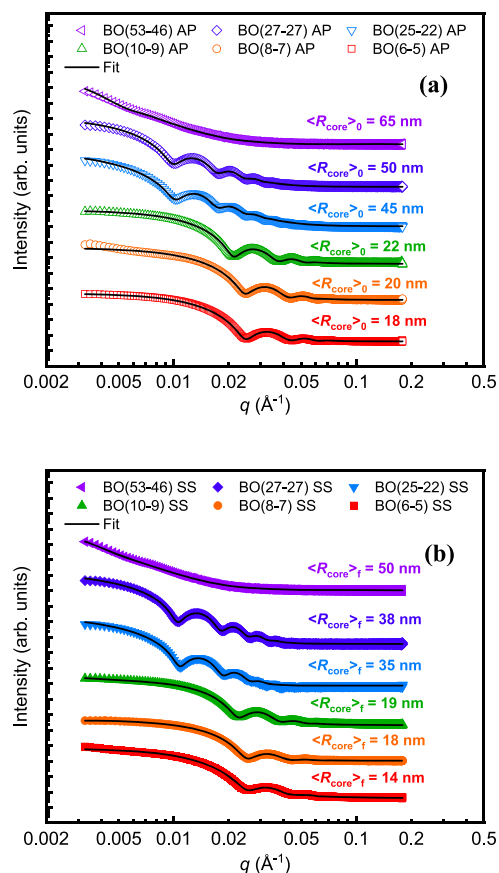


Figure 1. SAXS intensity versus q , on logarithmic scales, for 0.25 wt % PB-PEO in $[C_2mim][TFSI]$ ($T = 27^\circ C$) (a) as prepared by DD and (b) steady-state after annealing at 170 °C. The black lines represent the best fits of the scattering data to eq 4 with the Pedersen model for block copolymer micelles. The data sets are shifted vertically for clarity.

micelle size before and after annealing at 170 °C are summarized in Table 2. The percent decrease in $\langle R_{core} \rangle$ exhibits a similar trend to that observed in $\langle R_h \rangle$; the percent decrease in $\langle R_{core} \rangle$ with increasing M was 22%, 20%, 14%, 22%, 24%, and 23%. From $\langle R_{core} \rangle$, the average aggregation number Q was calculated assuming that the micelle core is devoid of solvent (i.e., $Q = (4\pi\langle R_{core} \rangle^3)/3V_{core}$, where $V_{core} = M_{n,PB}/\rho_{PB}N_{Av}$ is the molar volume of one core block). This is a reasonable approximation for such a strongly segregated system. From these values, the fragmentation process results in a population of micelles with an average aggregation number that is approximately half that of the initial aggregation number. The core size of strongly segregated block copolymer micelles should scale as $R_{core} \sim N_{core}^{3/5}$.^{61–63} As shown in Figure 2, $\langle R_{core} \rangle$ increases with N_{core} essentially as predicted for both the as-prepared and steady-state samples. This former result is somewhat unexpected, in that micelles prepared by DD apparently adhere to an equilibrium scaling relationship of $\langle R_{core} \rangle$ with N_{core} , even though they are clearly far from equilibrium. This result highlights another difficulty in assessing the true equilibrium state in block copolymer micelles—apparent adherence to an equilibrium scaling law is far from a sufficient criterion.

Fragmentation Kinetics. The substantial decrease in micelle size after annealing has previously been established as being due to equilibration via micelle fragmentation.^{36,45} Thus,

Table 2. Micelle Dimensions for 0.25 wt % Solutions in [C₂mim][TFSI] at T = 27 °C, Before and After Annealing at 170 °C

as prepared by direct dissolution							
sample	$\langle R_h \rangle_0$ (nm)	$\langle \mu_2 / \Gamma^2 \rangle_0$	$\langle R_{core} \rangle_0$ (nm)	$\sigma_{core,0}$ (nm)	Q_0^a	$s_{core,0}^b$	$s_{corona,0}^c$
BO(6–5)	29	0.084	18	1.4	1260	2.05	2.22
BO(8–7)	42	0.251	20	6.0	1900	2.45	2.50
BO(10–9)	51	0.229	22	10.0	2400	2.38	3.31
BO(25–22)	70	0.086	45	5.3	8060	2.93	1.86
BO(27–27)	83	0.166	50	6.6	11000	3.38	2.20
BO(53–46)	108	0.177	65	15.0	11400	3.15	2.07
steady-state after annealing at 170 °C							
sample	$\langle R_h \rangle_f$ (nm)	$\langle \mu_2 / \Gamma^2 \rangle_f$	$\langle R_{core} \rangle_f$ (nm)	$\sigma_{core,f}$ (nm)	Q_f^a	$s_{core,f}^b$	$s_{corona,f}^c$
BO(6–5)	20	0.050	14	1.3	930	1.97	0.78
BO(8–7)	27	0.050	16	3.0	950	1.60	1.51
BO(10–9)	34	0.042	19	3.3	1500	2.05	1.71
BO(25–22)	58	0.072	35	5.4	3800	2.28	1.72
BO(27–27)	62	0.085	38	3.0	4900	2.57	1.60
BO(53–46)	77	0.085	50	7.8	5200	1.96	1.80

^aThe aggregation number (Q), where the subscripts 0 and f denote initial and final values, respectively, was calculated as $4\pi\langle R_{core} \rangle^3 / (3V_{PB})$ assuming the core is devoid of solvent, and $V_{PB} = M_{n,PB} / \rho_{PB} N_{Av}$ is the volume per core chain, $\rho_{PB} = 0.89 \text{ g/cm}^3$, N_{Av} is Avogadro's number, and $M_{n,PB}$ is the molecular weight of PB, as reported in Table 1. ^bThe degree of core chain stretching (s_{core}), was calculated as $\langle R_{core} \rangle$ divided by the root-mean-square end-to-end distance of the core block using the statistical segment length of PB, $b = 5.9 \text{ Å}$. ^cThe degree of corona chain stretching (s_{corona}) calculated as the corona thickness ($L_{corona} = \langle R_h \rangle - \langle R_{core} \rangle$) divided by the root-mean-square end-to-end distance of the corona block using the statistical segment length of PEO, $b = 6.0 \text{ Å}$.

the fragmentation kinetics were studied for six molecular weights of PB-PEO in [C₂mim][TFSI] using T -jump DLS and TR-SAXS to determine the dependence of the fragmentation time τ on the total degree of polymerization of the copolymer N_{total} . The change in $\langle R_{core} \rangle$ was monitored with high temporal resolution by heating micelle solutions directly on the beamline to 170 °C under vacuum, and the scattering was measured during annealing for 48 h. Representative TR-SAXS data for 0.25 wt % BO(27–27) in [C₂mim][TFSI] are shown in Figure 3, where the background-subtracted intensity traces versus q are plotted as a function of annealing time. Analogous TR-SAXS data for BO(8–7), BO(10–9), and BO(25–22) are provided in the SI, Figures S6–S8.

As shown in Figure 3b, the form factor for BO(27–27) shows a first minimum at $q_{min} = 0.009 \text{ Å}^{-1}$ after annealing at 170 °C for approximately 26 min, which corresponds to $\langle R_{core} \rangle = 50 \text{ nm}$ from the hard sphere approximation where $qR_{core} = 4.493$. Throughout the course of an annealing experiment, the q -position of the first minimum shifts progressively to higher values of q , indicative of a smaller $\langle R_{core} \rangle$. After annealing at 170 °C for approximately 2500 min, q_{min} shifts to 0.012 Å^{-1} and $\langle R_{core} \rangle = 37.4 \text{ nm}$. Fitting the scattering data to the block copolymer micelle model before and after annealing at 170 °C shows good agreement with the hard sphere approach used to determine $\langle R_{core} \rangle$ from TR-SAXS.

To quantify the fragmentation kinetics, the change in micelle size, either $\langle R_h \rangle$ or $\langle R_{core} \rangle$, is normalized according to the following: eq 4:

$$R(t) = \frac{\langle R_i \rangle_t - \langle R_i \rangle_f}{\langle R_i \rangle_0 - \langle R_i \rangle_f} = \exp[-(t/\tau)^n] \quad (4)$$

where $R(t)$ is the normalized relaxation function, and $i = h$ or c denotes either the average hydrodynamic radius from DLS or the average micelle core radius from SAXS. The normalized change in size was fit to the Avrami equation or “compressed” exponential, shown in eq 4, where τ is the fragmentation time constant and n is the Avrami exponent.

The normalized change in $\langle R_h \rangle$ from DLS is shown in Figure 4, and that for $\langle R_{core} \rangle$ in Figure 5.

From DLS, the change in $\langle R_h \rangle$ while annealing at 170 °C is well-described by eq 4, with an exponent of 2.0–2.3. The fitting was performed using n as an adjustable parameter, resulting in an average value of $n = 2.2 \pm 0.3$. Refitting the data with a fixed exponent of $n = 2.0$ resulted in fits of similar quality with only modest variations in τ , as shown in Table S1. Attempts to fit the change in normalized $\langle R_h \rangle$ to a single or a double exponential resulted in poor fit quality for the former, and physically unreasonable values of τ for the latter. This is consistent with previous reports on a single, low molecular weight of PB-PEO in various ionic liquids.^{36,40,45} From Figure 4, the fragmentation time τ increases strongly with increasing M ; the 11 kg mol^{−1} diblock has a τ on the order of 100 min, whereas the 100 kg mol^{−1} diblock has $\tau \approx 5000 \text{ min}$. To corroborate the results from T -jump DLS, the time-dependent change in R_{core} from SAXS at 170 °C was normalized according to eq 4. The results for all polymers except BO(53–46) are shown in Figure 5. The dispersity in R_{core} for BO(53–46) made estimating the radius using the hard sphere approximation based on the first minimum of the form factor unreliable. Although the intensity traces for BO(53–46) could be fit to the Pedersen model, the error in $\langle R_{core} \rangle$ was quite large. The large micelle size dispersity and exponent by DLS shown in Table 3 for BO(53–46) are likely due to the fact that this sample is prepared using the same dissolution time as the lower M samples. Due to this uncertainty, the fragmentation kinetics were determined only for the other five M . In general, values of τ obtained from DLS and SAXS agree remarkably well. The fitting results for both experiments are summarized in Table 3.

Molecular Weight Dependence of Fragmentation Time. As noted above, the fragmentation kinetics for PB-PEO in [C₂mim][TFSI] depend strongly on M . Figure 6 shows the values of τ obtained from fitting the normalized change in micelle size to eq 4 as a function of the total degree of polymerization; the results from DLS and SAXS are very

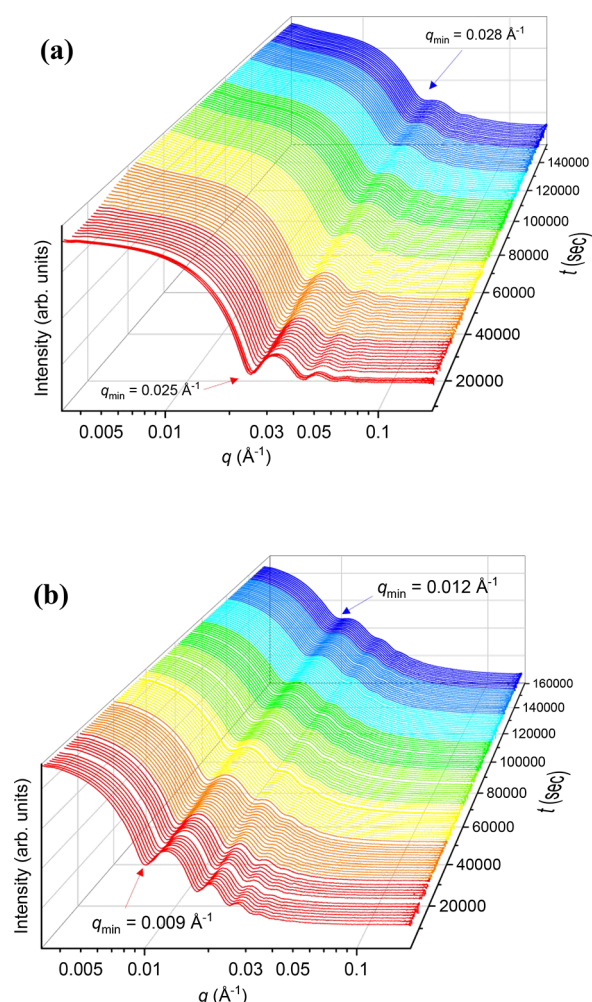


Figure 3. TR-SAXS intensity (logarithmic scale) as a function of q showing the evolution in the micelle core radius while annealing at 170 °C for 0.25 wt % (a) BO(6-5) in $[C_2mim][TFSI]$. The position of the first minimum in the form factor increases from $q_{min} = 0.025 \text{ \AA}^{-1}$ for short annealing times to $q_{min} = 0.028 \text{ \AA}^{-1}$ at longer annealing times. (b) BO(27-27) in $[C_2mim][TFSI]$. The first minimum in the form factor increases from $q_{min} = 0.009 \text{ \AA}^{-1}$ at short times to $q_{min} = 0.012 \text{ \AA}^{-1}$ at longer annealing times. The PB core radius can be estimated assuming a hard sphere ($R_{core} \cong 4.493/q_{min}$), so the radius of the micelle core decreases with increasing annealing time.

consistent. The fit to a power law gives the scaling $\tau \sim N_{total}^{1.8 \pm 0.1}$. Dormidontova proposed a scaling model for micelle fusion kinetics, and further assumed that fragmentation would proceed as the reverse of micelle fusion, and thus the N scaling would be the same for both processes.²⁷ For micelles with large aggregation numbers (Q) with respect to their equilibrium size, Dormidontova defined a characteristic time for corona deformation (τ_{def}) as the inverse translational diffusion constant for chains, where V_B is the molar volume per monomer unit in the corona.²⁷

$$\tau_{def} \cong \frac{\eta_s V_B}{k_B T} N_{corona}^{9/5} Q^{13/5} \quad (5)$$

Assuming the fragmenting micelle is large, Dormidontova's theory proposes that $\tau \sim N_{corona}^{1.8}$ at constant Q , as shown in eq 5. It should be noted that for very large aggregation numbers, such as the ones obtained for the as-prepared micelles, that there is at most a small molecular weight

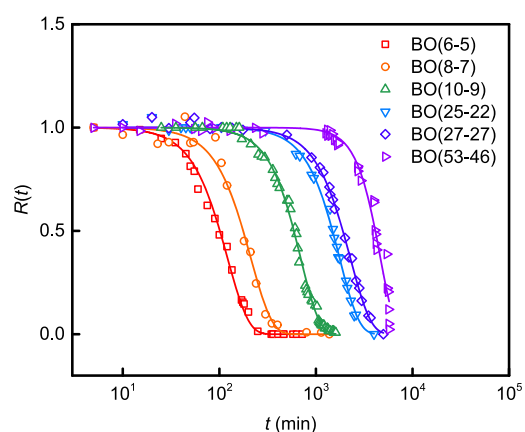


Figure 4. Time dependence of normalized $\langle R_h \rangle$ for 0.25 wt % solutions of BO molecular weight series in $[C_2mim][TFSI]$. DLS measurements were performed at a scattering angle of 90° and a relaxation temperature of 170 °C. Solid lines represent best fits to eq 4 with $n = 2.2 \pm 0.3$.

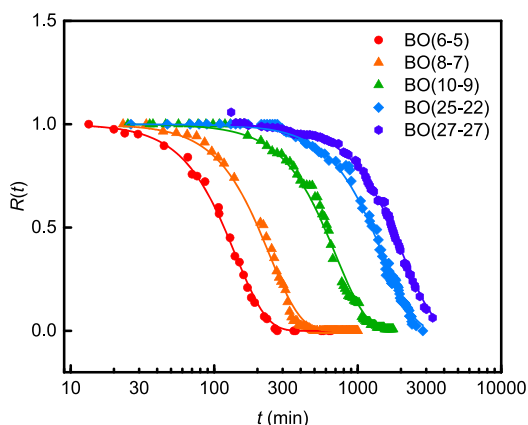


Figure 5. Time dependence of normalized $\langle R_{core} \rangle$ for 0.25 wt % solutions of PB-PEO in $[C_2mim][TFSI]$. TR-SAXS measurements were performed at a relaxation temperature of 170 °C. Solid lines represent best fits to eq 4 with $n = 2.0 \pm 0.1$.

Table 3. Molecular Weight Dependence of τ Determined by T-jump DLS and TR-SAXS at 170 °C for 0.25 wt % BO in $[C_2mim][TFSI]$

sample	T-jump DLS		TR-SAXS	
	τ (min)	n	τ (min)	n
BO(6-5)	120 ± 10	2.0	140	2.1
BO(8-7)	200 ± 75	2.0	250	2.2
BO(10-9)	690 ± 90	2.2	710	2.2
BO(25-22)	1750 ± 170	2.2	1500	2.1
BO(27-27)	2300 ± 250	2.1	2070	2.3
BO(53-46)	5000 ± 360	2.7		

dependence to Q . The Q dependence of τ is shown in the SI, where $\tau \sim Q_0^{1.7 \pm 0.1}$. The agreement between the scaling of τ with N_{corona} and Q_0 implies that the strong Q dependence from eq 5 does not reflect our experimental results, and this is consistent with the assumption that Q does not depend strongly on N_{total} for very large Q .

The results are apparently in excellent agreement with the theory. However, some caution is warranted. First, the theory does not anticipate the compressed exponential form in eq 4, so there is clearly an aspect of the fragmentation mechanism

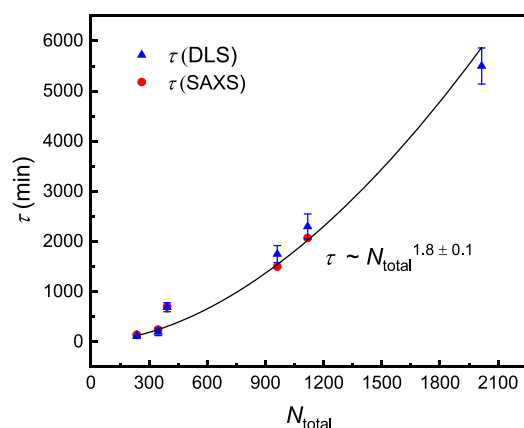


Figure 6. Dependence of the fragmentation time constants (τ) determined by T -jump DLS (red circles) and TR-SAXS (blue triangles) on the degree of polymerization of PEO (N_{PEO}).

that is not yet fully appreciated. Second, the separate effects of N_{core} and N_{corona} on the experimental fragmentation times cannot be assessed individually because the volume fraction of PEO in this work has been constant. Thus, the experimental scaling represents the dependence of the fragmentation time on the total degree of polymerization, which may not coincide with the dependence on N_{corona} . However, it does seem physically reasonable that the corona plays a much larger role in the process than the core. Further discussion of the time constants from the model and from estimates of corona dynamics are provided in the SI.

To rationalize the dependence of the fragmentation kinetics on molecular weight, one must consider the transition state and activation barrier of this process. Previous work on this system revealed that the fragmentation kinetics were unaffected by changes in the solvent selectivity toward the core-forming block,⁴⁵ thus indicating independence of interfacial tension. Accordingly, we propose that severe corona crowding in the transition state is the primary barrier to fragmentation. When considering the free energy of polymer chains in a block copolymer micelle at equilibrium, the interfacial free energy and the free energy of chains in the corona contribute more to the total free energy of the micelle compared to the free energy of chains in the core.^{12,61,66} We calculate the free energy of PB-PEO micelles as prepared by DD, after annealing at 170 °C, and for the proposed “peanut-shaped” transition state (Figure S10) observed in previous work on PB-PEO in $[\text{C}_{2}\text{mim}][\text{TFSI}]$ by high-temperature liquid phase transmission electron microscopy.⁴⁶ A schematic illustration of the transition state is shown in Figure S11. The total free energy of a spherical block copolymer micelle in solution (F_{mic}) was well described by Zhulina et al.,¹² and is given by the sum of the free energy contribution from the core chains (F_{core}), the free energy of the corona chains (F_{corona}), and the interfacial energy (F_{int}). Therefore, the free energy can be estimated for the as-prepared and steady state micelles. Some details on this calculation are provided in the SI, and were previously reported by Zhulina et al.¹² Because the transition state morphology is “peanut-shaped” the transition state free energy cannot be treated in the same manner as the spherical micelles before and after fragmentation. The calculated free energies are reported in Table S4.

To calculate the corona free energy in the transition state, the fraction of corona chains overlapped in the neck region

must be estimated. The transition state is represented schematically in Figure S11. We propose that the increase in the local concentration of corona chains near the neck point, i.e., the increase in the number of blobs per corona chain n_{blob} in the formation of the transition state is the primary barrier to fragmentation. To estimate the degree of corona crowding in the transition state first, we estimate the fraction of corona overlap in the transition state, and then relate the overlap fraction to the concentration of blobs per PEO chain in the neck region using the theory from Zhulina et al.¹² Using a simplified model for the transition state, illustrated in Figure S12, to determine the area of corona overlap in the transition state results in approximately 1/6 of the corona chains overlapping for all molecular weights. Details on this calculation are provided in the SI. Then, the concentration of PEO chains in the neck region of the transition state is determined by estimating the number of blobs per corona chain. In the neck region, the number of blobs per chain is expected to be approximately twice the number of blobs per chain in the as-prepared micelles ($n_{\text{blob,neck}} = 2n_{\text{blob,sphere}} = 2F_{\text{corona,0}}/k_{\text{B}}T$), whereas the outer spherical regions would have $n_{\text{blob,sphere}} = F_{\text{corona,0}}/k_{\text{B}}T$, where the free energy is calculated using the as-prepared size of the micelle as shown in eq S11.

The transition state free energy for the core and the interface was calculated using the steady state micelle dimensions. The transition state corona free energy was calculated assuming 5/6 of the chains are described by $F_{\text{corona,0}}/k_{\text{B}}T$, and for the 1/6 of corona chains overlapped in the neck region, the corona free energy is $2F_{\text{corona,0}}$. Therefore, the total corona free energy in the transition state is calculated as $F_{\text{corona,TS}}/k_{\text{B}}T = (5/6)F_{\text{corona,0}}/k_{\text{B}}T + (1/3)F_{\text{corona,0}}/k_{\text{B}}T$. Details on this calculation are provided in the SI. From Table S4, the average n_{blob} is 16 for BO(53–46) before a T -jump to 170 °C. Assuming that the number of blobs in the neck point is double the calculated $n_{\text{blob,0}}$ shown in Table S4, approximately 1/6 of the corona chains experience corona crowding where there are approximately 32 blobs per chain near the neck, whereas n_{blob} is 16 for chains in the spherical region of the micelle indicated in Figure S11. The fraction of corona chain overlap was determined to be 1/6 for all molecular weights by eq S8, and thus n_{blob} at the neck point is 8, 9, 10, 21, 24, and 32 for BO(6–5), BO(8–7), BO(10–9), BO(25–22), BO(27–27), and BO(53–46). This large increase in the local concentration of PEO chains in the neck region of the transition state supports the strong molecular weight dependence of micelle fragmentation kinetics. Because the confinement free energy of corona chains is directly proportional to n_{blob} , the chains near the neck point become much more confined and the entropic penalty associated with this chain confinement contributes to this being the primary activation barrier to fragmentation.

Although these estimates are encouraging, we emphasize that the uncertainties in estimating the transition state free energy are quite large, and the differences in the calculated free energies for the spherical and “peanut” shaped micelles are only $\sim 5\text{--}10k_{\text{B}}T$. Given that the aggregation numbers are huge and the barrier is known to be of order 30–40 kT ,^{36,45} the estimated barriers per chain are small fractions of kT , and beyond the resolution of this calculation.

Molecular Weight Dependence of Initial Micelle Size by Direct Dissolution. To understand the N dependence of the initial core size for micelles prepared by DD, SAXS measurements on the pure PB-PEO diblock copolymers were conducted. The 1D scattering intensity traces for these

582 samples are shown in Figure S9. It has been suggested that the
 583 morphology of micelles prepared by direct dissolution depends
 584 on the morphology of the bulk diblock copolymer.^{36,67} The
 585 scattering data confirms that BO(6–5), BO(8–7), BO(10–9),
 586 BO(25–22), and BO(27–27) exhibit a lamellar morphology at
 587 70 °C, which is the temperature used for preparing micelle
 588 solutions by direct dissolution. The domain size of the PB
 589 lamellae was estimated from the q -position of the primary
 590 scattering peak ($d = 2\pi/q^*$) and the known volume fraction of
 591 PB. The domain sizes were found to be approximately 16, 17,
 592 19, 47, and 54 nm for BO(6–5), BO(8–7), BO(10–9),
 593 BO(25–22), and BO(27–27), respectively. Interestingly, the
 594 domain sizes obtained from SAXS of the bulk polymers are
 595 comparable to the initial micelle core size. This dependence is
 596 represented in Figure 7 as a plot of $\langle R_{\text{core}} \rangle_0$ vs d_{PB} . From this

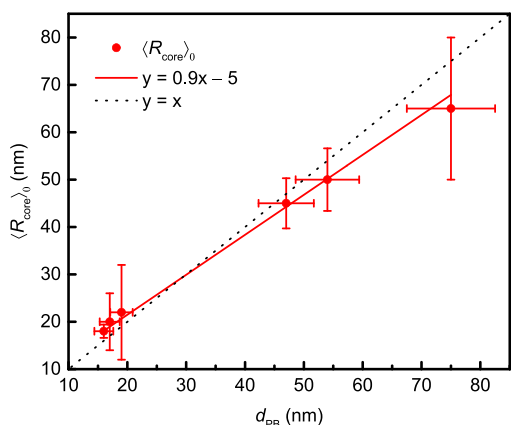


Figure 7. Dependence of $\langle R_{\text{core}} \rangle_0$ (nm) on the PB domain size (d_{PB}) indicating a direct relationship between $\langle R_{\text{core}} \rangle_0$ for micelles prepared by DD and the PB domain size of the bulk diblock copolymer. The solid red line represents the best linear fit to $\langle R_{\text{core}} \rangle_0$ as a function of d_{PB} , and the dotted black line represents $y = x$. The error bars in $\langle R_{\text{core}} \rangle_0$ represent $\pm \sigma_{\text{core}}$ from Table 2, and 10% error in d_{PB} is assumed.

597 figure, the direct relationship between the domain size of the
 598 bulk copolymer and the initial micelle core radius is apparent.
 599 This result is also consistent with the previous reports on
 600 BO(8–7) in 1-alkyl-3-methylimidazolium bis-
 601 (trifluoromethylsulfonyl)imide-based ionic liquids.^{36,45}
 602 It should be noted that a linear dependence should not
 603 necessarily be expected. The dependence begins to weaken
 604 slightly at higher molecular weights, as evidenced by the slight
 605 downward curvature of $\langle R_{\text{core}} \rangle$ for the two largest polymers.
 606 The dependence of $\langle R_{\text{core}} \rangle_0$ on d_{PB} follows the strong
 607 segregation limit scaling of $2/3$,⁶⁸ as shown in Figure S13,
 608 while the apparent equilibrium scaling of $\langle R_{\text{core}} \rangle$ with N_{core} for
 609 the initial micelles has an exponent closer to $3/5$, as shown in
 610 Figure 2. These slightly different dependences give rise to the
 611 curvature evident in Figure 7. Figure S13 shows the
 612 relationship of d_{PB} as a function of N_{core} for the domain size
 613 determined experimentally by SAXS, and the estimated
 614 domain spacing calculated from self-consistent field theory
 615 (SCFT); the experimental domain size of the PB lamellae
 616 agrees well with SCFT. We therefore speculate that during
 617 direct dissolution the solvent penetrates the PEO domains,
 618 peeling apart layers of PEO-decorated PB layers. Some surface
 619 instabilities in these separate layers drive a pinching-off process
 620 to give large, disperse spherical aggregates in solution. This

behavior is broadly analogous to a Rayleigh instability, and has
 been referenced in computations of surfactant-based mi-
 celles,⁶⁹ and experimentally in cylinder-to-sphere transi-
 tions^{70–75} in block copolymer micelles. In experimental
 works on the cylinder-to-sphere transitions in block copolymer
 micelles, it is concluded that the surface-instability-mediated
 transitions induce pinching of immature spherical micelles,
 followed by rapid chain exchange.⁷⁴ On the basis of the lack of
 chain exchange in BO/[C₂mim][TFSI] systems reported
 previously,³⁶ the direct dissolution mechanism is related to
 this process, but not identical. Furthermore, even accounting
 for the molar mass difference, our system undergoes
 fragmentation about 3 orders of magnitude slower than the
 PEP-PEO cylindrical micelles, implying a very small barrier in
 the latter case.

Another possibly surprising result of the dependence of
 $\langle R_{\text{core}} \rangle_0$ on d_{PB} is that the micelles prepared by direct
 dissolution have a core radius that is approximately equal to
 the full PB domain size, rather than $d_{\text{PB}}/2$. The interfacial
 tension in the swollen state is clearly greater than in the dry
 copolymer, so an increase in core chain stretching is certainly
 expected. Moreover, the degree of interdigitation of the PB
 chains in the dry lamellae can affect this picture of the
 dissolution mechanism. In the dry state, the degree of
 stretching of PB (s_{PB}) can be estimated from two limits: the
 first assumes no interdigitation of chains, where $s_{\text{PB},1} = d_{\text{PB}}/$
 $\langle h^2 \rangle_0^{1/2}$; the second method assumes full interdigitation of
 chains, where $s_{\text{PB},2} = d_{\text{PB}}/2\langle h^2 \rangle_0^{1/2}$. Calculating the degree of
 stretching for BO(8–7) by these two methods gives $s_{\text{PB},1} = 1.1$
 and $s_{\text{PB},2} = 2.2$, respectively. These may be to the degree of
 stretching of chains in the micelle core from Table 2, $s_{\text{core}} = 2.5$.
 This result implies some significant degree of interdigitation in
 the bulk state.

CONCLUSIONS

In this work, the effect of polymer molecular weight on micelle
 fragmentation kinetics was studied for the first time, using six
 molecular weights of PB–PEO with a near constant volume
 fraction of PEO ($f_{\text{PEO}} \approx 0.40$) in the ionic liquid [C₂mim]-
 [TFSI]. Disperse micelles prepared by direct dissolution in
 [C₂mim][TFSI] were found to be larger than the equilibrium
 size, and all molecular weight micelles decreased in size, both
 $\langle R_h \rangle$ and $\langle R_{\text{core}} \rangle$, when subjected to high temperature
 annealing. The decrease in micelle size while annealing at
 170 °C was monitored by T -jump DLS and TR-SAXS, and it
 was concluded that PB–PEO micelles equilibrate by fragmen-
 tation. The final aggregation number was approximately half of
 the original aggregation number prior to a T -jump. The decay
 in $\langle R_h \rangle$ and $\langle R_{\text{core}} \rangle$ was consistently well described by a
 compressed exponential with an exponent of 2; the origin of
 this functional form remains to be elucidated. The character-
 istic fragmentation times determined by fitting the normalized
 change in micelle size to this equation was found to depend
 strongly on the molecular weight of the block copolymer,
 where $\tau \sim N_{\text{total}}^{1.8 \pm 0.1}$. A previous model of micelle fragmentation
 by Dormidontova predicts a similar scaling, albeit in terms of
 N_{corona} . Future measurements on a series of polymers with
 constant N_{core} will be required to assess whether this apparent
 agreement is robust. The core size of the initial micelles was
 shown to correlate closely with the PB domain dimensions in
 the precursor dry, lamellar copolymer, suggesting a formation
 mechanism involving peeling apart of layers, followed by
 pinching off into spherical domains.

■ ASSOCIATED CONTENT

■ Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/jacs.1c02147>.

SEC-RI traces of PB-PEO diblocks, ^1H NMR spectra of PB-PEO diblocks and $[\text{C}_2\text{mim}][\text{TFSI}]$, REPS results for PB-PEO in $[\text{C}_2\text{mim}][\text{TFSI}]$ as-prepared by DD and steady-state after annealing, dependence of $\langle R_h \rangle$ on the total degree of polymerization N_{total} for 0.25 wt % PB-PEO in $[\text{C}_2\text{mim}][\text{TFSI}]$, fragmentation times with fixed n , TR-SAXS traces for BO(8–7), BO(10–9), and BO(25–22), calculations of corona chain disentanglement time, SAXS of bulk PB-PEO, high temperature liquid-phase transmission electron microscopy of BO(53–46) in $[\text{C}_2\text{mim}][\text{TFSI}]$, calculations of the fraction of corona overlap in the transition state, high temperature LP-TEM micrograph of BO(53–46) in $[\text{C}_2\text{mim}][\text{TFSI}]$ at 170 °C, schematic illustration of fragmentation transition state, illustration of simplified model transition state for calculating corona chain overlap, calculation of micelle free energy, domain spacing from SAXS of $\text{BO}(x-y)$ diblock copolymers in the bulk, Comparison of the domain size of PB determined by SAXS and the estimated domain size for a lamellar diblock copolymer using self-consistent field theory, estimated fragmentation lag time as a function of degree of polymerization, and dependence of fragmentation time on as-prepared aggregation number (PDF)

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Notes

The authors declare no competing financial interest.

■ ACKNOWLEDGMENTS

This work was supported primarily by the National Science Foundation (DMR-1707578). Additional support came from the University of Minnesota Doctoral Dissertation Fellowship (JTE), the University of Minnesota Robert and Jill DeMaster Excellence Fellowship (J.T.E.), the Research Experiences for Undergraduates (REU) Program of the National Science

Foundation under Award Number DMR-1559833 (through the University of Minnesota MRSEC under Award Number DMR-2011401, A.B.), and from the U.S. Department of Energy (DOE), Office of Science, Office of Workforce Development for Teachers and Scientists, Office of Science Graduate Student Research (SCGSR) program (J.T.E.). The SCGSR program is administered by the Oak Ridge Institute for Science and Education (ORISE) for the DOE. ORISE is managed by ORAU under contract number DE-SC0014664. Portions of this work were performed at the Center for Functional Nanomaterials, and the National Synchrotron Light Source II, Brookhaven National Laboratory, which are supported by the U.S. DOE Office of Science under Contract DE-SC0012704. Portions of this work were performed at the DuPont-Northwestern-Dow Collaborative Access Team (DND-CAT) located at Sector 5 of the Advanced Photon Source (APS). DND-CAT is supported by Northwestern University, E.I. DuPont de Nemours & Co., and The Dow Chemical Company. This research used resources of the Advanced Photon Source, a U.S. Department of Energy (DOE) Office of Science User Facility operated for the DOE Office of Science by Argonne National Laboratory under Contract No. DE-AC02-06CH11357. We thank Dr. Ruipeng Li and Dr. Esther Tsai for assistance with TR-SAXS measurements at the 11-BM CMS beamline at the NSLS-II and Dr. Lihua Zhang for insightful discussions on high temperature LP-TEM. We thank Claire Seitzinger for assisting with SAXS measurements of bulk PB-PEO diblocks.

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