

# Molecular Weight Dependence of Block Copolymer Micelle Fragmentation Kinetics

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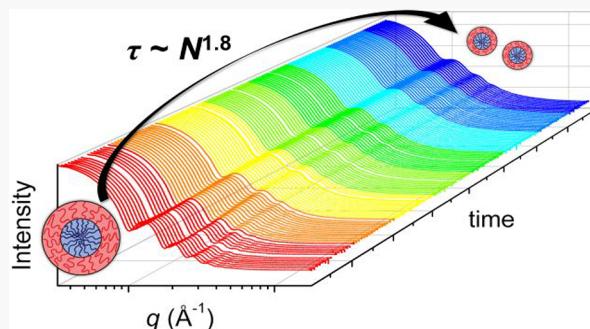
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**ABSTRACT:** The effect of molecular weight ( $M$ ) on the fragmentation kinetics of micelles formed by 1,2-polybutadiene-*block*-poly(ethylene oxide) (PB-PEO) copolymers was studied in the ionic liquid 1-ethyl-3-methylimidazolium bis(trifluoromethylsulfonyl)imide. 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 ( $\tau$ ) 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.



19 ■ INTRODUCTION

20 Diblock copolymers self-assemble into a variety of micellar 21 structures in the presence of a block-selective solvent, i.e., a 22 solvent that dissolves one block and not the other. Block 23 copolymer (BCP) micelles are utilized in a wide range of 24 applications including ion gels,<sup>1,2</sup> drug or gene delivery,<sup>3,4</sup> 25 nanolithography,<sup>5</sup> and viscosity modification.<sup>6</sup> Because of the 26 versatility of BCP micelles from an applications perspective, 27 their dynamics and equilibration kinetics in solution are of 28 interest, yet these processes remain less well understood in 29 comparison to their low molecular weight counterparts.<sup>7–10</sup> 30 The thermodynamics of micellization has been studied 31 extensively, both theoretically and experimentally.<sup>11,12</sup> The 32 morphology and size of block copolymer micelles at 33 equilibrium can be tuned by changing the molecular 34 characteristics of the block copolymer itself, such as the core 35 and corona block lengths,<sup>13–16</sup> and the solvent selectivity.<sup>17</sup> 36 There are five primary processes thought to govern the 37 formation, destruction, and equilibration of block copolymer 38 micelles in solution, namely, (i) *chain exchange*, where 39 individual copolymer chains are exchanged between micelles, 40 (ii) *fragmentation*, where a micelle much larger than the 41 equilibrium aggregation number  $Q_{\text{eq}}$  breaks up into smaller 42 micelles, (iii) *fusion*, where micelles smaller than  $Q_{\text{eq}}$  merge 43 together, (iv) *micelle creation*, where individual chains nucleate 44 a new micelle, and (v) *annihilation* of micelles, where a series 45 of chain exchange/fragmentation/fusion events eliminates one 46 micelle. The kinetics of these processes were first modeled by 47 Aniansson, Wall, and co-workers.<sup>18,19</sup> These processes of

micellization and equilibration are supported by experimental 48 studies of low molecular weight surfactants<sup>20</sup> and Pluronics.<sup>21–24</sup> 49 Halperin and Alexander developed the first detailed 50 theory of block copolymer micelle dynamics near equilibrium,<sup>51</sup> 52 where chain exchange was considered to be the only plausible 53 equilibration mechanism, as fragmentation and fusion were 54 argued to be much slower.<sup>25,26</sup> Dormidontova proposed a 55 different model that gives scaling predictions for the relaxation 56 times of chain exchange, fragmentation, and fusion.<sup>27</sup> Nyrkova 57 and Semenov focused on the rates of micelle creation and 58 annihilation in response to abrupt changes in conditions such 59 as jumps in temperature.<sup>28</sup>

The molecular weight ( $M$ ) of diblock copolymers not only 60 affects the equilibrium structure of micelles, but also the 61 dynamics and equilibration kinetics in solution.<sup>8,15,29</sup> The 62 aggregation number  $Q$  of a given micelle can vary, and micelle 63 equilibration occurs by some combination of the exchange of 64 individual chains,<sup>7,9,15,30–35</sup> fusion, or fragmentation.<sup>20,22,23,36</sup> 65 Halperin and Alexander proposed that the chain exchange 66 kinetics depends on a characteristic relaxation time,  $\tau$ , that 67 depends on the core and corona block lengths as  $\tau \sim$  68

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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  $\gamma$  is the interfacial tension 71 between the core block and the solvent and  $b$  is the monomer 72 size of the core block.<sup>25</sup> 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,  $N_{\text{core}}$ , in contrast to the 76 Halperin-Alexander model.<sup>8,32–34,37,38</sup> Interestingly, it has also 77 been reported that increasing the degree of polymerization of 78 the corona block,  $N_{\text{corona}}$ , of PS-PEO micelles in squalane could 79 increase the rate of chain exchange by up to 2 orders of 80 magnitude,<sup>15</sup> whereas studies of  $C_{27}\text{-PEO}_n$  in water showed 81 that increasing  $N_{\text{corona}}$  had the opposite effect.<sup>8</sup>

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.<sup>36</sup> 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.<sup>22,23,36</sup> 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.<sup>39–43</sup> 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.<sup>41</sup> They concluded 99 that the initial growth of micelles occurs via micelle fusion.<sup>41</sup> 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.<sup>36,44,45</sup>

106 A theoretical model of the  $M$  dependence, specifically the 107 dependence on  $N_{\text{corona}}$ , of micelle fusion and fragmentation has 108 been reported.<sup>27</sup> 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.<sup>27</sup> The relaxation time ( $\tau$ ) 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  $\tau \sim N_{\text{corona}}^{17/5} Q_1^{4/5} Q_2$ .<sup>27</sup> 115 The strong  $N_{\text{corona}}$  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  $V_{\text{corona}}$  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 ( $[\text{C}_2\text{mim}][\text{TFSI}]$ ) 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.<sup>40</sup> For micelles prepared by DD,  $\langle R_h \rangle$  decreased to approximately half of the original size when annealed at 170 °C.<sup>40</sup> 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).<sup>36</sup> 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 ( $[\text{C}_x\text{mim}][\text{TFSI}]$  where  $x = 1, 2, 4, 6$ , and 8).<sup>45</sup> 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.<sup>45</sup> We previously reported the direct observation of micelle fragmentation for three molecular weights of PB-PEO in  $[\text{C}_2\text{mim}][\text{TFSI}]$  using high-temperature liquid-phase transmission electron microscopy, and intermediate structures close to an apparent transition state for micelle fragmentation were observed in situ.<sup>46</sup>

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  $[\text{C}_2\text{mim}][\text{TFSI}]$  where the total molecular weight of the diblock is varied from 11 to 100 kg mol<sup>-1</sup> 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.<sup>47</sup>

## ■ EXPERIMENTAL SECTION

**Synthesis and Characterization.** Six PB-PEO copolymers were synthesized by two-step sequential anionic polymerization,<sup>48</sup> 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<sup>-1</sup>) of hydroxyl-terminated 1,2-polybutadiene were synthesized via anionic polymerization 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).<sup>176</sup> 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  $\text{BO}(x-y)$ , where  $x$  and  $y$  indicate the number-average molecular weights of the PB and PEO blocks in kg mol<sup>-1</sup>, respectively. The number-average molecular weight ( $M_n$ ), dispersity ( $D$ ), and  $f_{\text{PEO}}$  were determined by a combination of size exclusion chromatography (SEC) in THF with a multiangle laser light scattering detector (Wyatt Dawn Heleos II) and <sup>1</sup>H nuclear magnetic resonance spectroscopy in CDCl<sub>3</sub> (<sup>1</sup>H 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<sup>a</sup>**

sample	$M_{n,\text{PB}}$ (kg mol <sup>-1</sup> )	$M_{n,\text{PEO}}$ (kg mol <sup>-1</sup> )	$f_{\text{PEO}}$	$D$
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

<sup>a</sup><sup>1</sup>H NMR spectroscopy in CDCl<sub>3</sub> was used to determine the number-average molar mass ( $M_n$ ) and volume fraction of PEO ( $f_{\text{PEO}}$ ) using bulk densities  $\rho_{\text{PB}} = 0.87 \text{ g/cm}^3$  for PB, and  $\rho_{\text{PEO}} = 1.13 \text{ g/cm}^3$  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 ( $D = M_w/M_n$ ) of the diblocks.

197 index increment ( $dn/dc$ ) for a diblock was estimated as the weight-  
198 average of the refractive index increments for PB in THF ( $dn/dc =$   
199 0.119 mL/g) and PEO in THF ( $dn/dc = 0.068 \text{ mL/g}$ ).<sup>49</sup> The <sup>1</sup>H  
200 NMR spectra of the diblocks in deuterated chloroform (CDCl<sub>3</sub>) are  
201 shown in Figure S2.

202 The ionic liquid (IL) 1-ethyl-3-methylimidazolium bis-  
203 (trifluoromethylsulfonyl)imide ([C<sub>2</sub>mim][TFSI]) was synthesized  
204 by an anion exchange reaction between 1-ethyl-3-methylimidazolium  
205 bromide and lithium bis(trifluoromethylsulfonyl)imide, carried out in  
206 water at 70 °C for 24 h.<sup>50</sup> The reaction mixture was diluted with  
207 dichloromethane, the aqueous layer was removed, and the organic  
208 layer was washed three times with deionized water. The organic layer  
209 was stirred on activated charcoal for 24 h. The solution was passed  
210 through an alumina column, dichloromethane was removed in vacuo,  
211 and [C<sub>2</sub>mim][TFSI] was dried under vacuum (<100 mTorr) at 60 °C  
212 for 72 h. The dried IL was characterized by <sup>1</sup>H, <sup>13</sup>C, and <sup>19</sup>F NMR  
213 spectroscopy in DMSO-*d*<sub>6</sub>. The <sup>1</sup>H NMR spectrum for [C<sub>2</sub>mim]-  
214 [TFSI] is shown in Figure S3.

215 **Solution Preparation by Direct Dissolution (DD).** All  
216 solutions were prepared by the direct dissolution (DD) method  
217 described here. The desired amounts of PB-PEO and [C<sub>2</sub>mim][TFSI]  
218 were combined by weight in a 20 mL scintillation vial equipped with a  
219 stir bar to obtain a 0.25 wt % solution. The vial was placed in an oil  
220 bath and stirred vigorously at 70 °C for 48 h. The resulting DD  
221 solutions were slightly blue-tinged for BO(6–5), BO(8–7), and  
222 BO(10–9) micelles, whereas those for BO(25–22), BO(27–27), and  
223 BO(53–46) were white-tinged.

224 **Dynamic Light Scattering (DLS).** Dynamic light scattering  
225 measurements were performed on a home-built light scattering setup  
226 equipped with a Brookhaven BI-DS photomultiplier mounted to an  
227 adjustable goniometer, a Lexel Ar<sup>+</sup> laser ( $\lambda = 488 \text{ nm}$ ), and a  
228 Brookhaven BI-9000 correlator. During a typical annealing experi-  
229 ment, the temperature was controlled with an index-matching high-  
230 temperature silicone oil bath to within  $\pm 0.1$  °C. To verify the size of  
231 the micelles before and after annealing, multiangle light scattering  
232 experiments were performed at  $T = 27$  °C with a range of scattering  
233 angles  $\theta$  from 60° to 120°.

234 Each micelle solution was passed through an 0.45 μm PTFE  
235 syringe filter to remove any dust, and subsequently flame-sealed under  
236 vacuum ( $\leq 70 \text{ mTorr}$ ) in a dust-free glass tube (I.D. 0.51 cm) to  
237 prevent degradation of the block copolymer and to avoid contact with  
238 moisture and dust. In an annealing experiment, the oil bath  
239 temperature was allowed to equilibrate for 30 min prior to  
240 introducing the sample. The normalized intensity autocorrelation  
241 function,  $g_2(t)$ , was measured at  $\theta = 90^\circ$  as a function of time. For  
242 each time point,  $g_2(t)$  was acquired for 90 s. The intensity  
243 autocorrelation function was converted to the electric field  
244 autocorrelation function,  $g_1(t)$ , via the Siegert relation,<sup>51</sup> which was  
245 then fit to a second-order cumulant expansion for single populations  
246 to obtain the average decay rate,  $\bar{\Gamma}$ , and the variance,  $\mu_2/\bar{\Gamma}^2$ . The  
247 mutual diffusion coefficient  $D_m$  was calculated from  $D_m = \bar{\Gamma}/q^2$ , where  
248  $q$  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 [C<sub>2</sub>mim][TFSI], and  $\lambda_0$  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;  $\eta_s(170 \text{ }^\circ\text{C}) = 2.4 \text{ kPa}\cdot\text{s}$  for [C<sub>2</sub>mim][TFSI].<sup>36</sup> The temperature dependence of the viscosity is well described by the Vogel–Fulcher–Tamman equation for [C<sub>2</sub>mim][TFSI] and many other ILs.<sup>52–55</sup> Additional analysis of the micelle size distribution was performed by applying the regularized positive exponential sum (REPES) Laplace inversion to the intensity correlation function.<sup>56</sup> 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 [C<sub>2</sub>mim][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 ( $\lambda = 0.918 \text{ \AA}$ ) 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.<sup>57</sup> The background-corrected intensity traces were fit using an established block copolymer micelle model.<sup>58,59</sup> In this model, the scattering intensity is defined as follows:

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

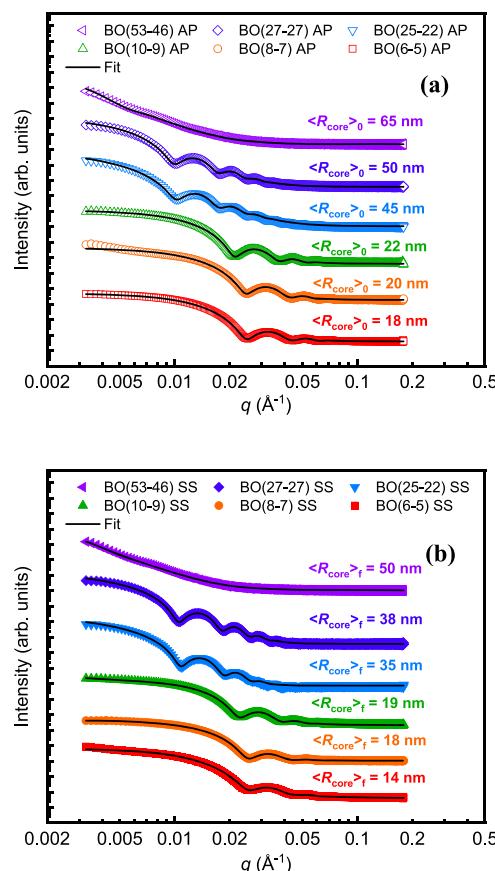
where  $R_{\text{core}}$  is the micelle core radius with distribution  $D(R_{\text{core}})$ ,  $P_{\text{mic}}(q)$  is the spherical form factor for the micelle core,<sup>58</sup>  $A_{\text{mic}}(q)$  is the form factor amplitude, and  $S(q)$  is the hard-sphere structure factor with the Percus–Yevick closure.<sup>59</sup>

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  $\lambda = 0.729 \text{ \AA}$  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.

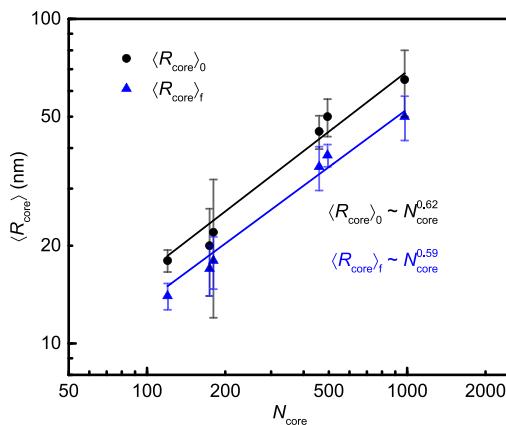
To determine the effect of  $M$  on  $R_{core}$ , SAXS was performed at room temperature before and after annealing at 170 °C. The background corrected scattering intensity traces versus  $q$  for the as-prepared and steady-state micelles are shown in Figure 1a and 1b, respectively. The as-prepared micelles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



**Figure 1.** SAXS intensity versus  $q$ , on logarithmic scales, for 0.25 wt % PB-PEO in  $[C_2mim][TFSI]$  ( $T = 27$  °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.

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.<sup>58,60</sup> 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.

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  $R_{core} \sim N_{core}^{3/5}$ .<sup>61–63</sup> 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.<sup>36,45</sup> Thus,

**Table 2. Micelle Dimensions for 0.25 wt % Solutions in [C<sub>2</sub>mim][TFSI] at T = 27 °C, Before and After Annealing at 170 °C**

sample	as prepared by direct dissolution						
	$\langle R_h \rangle_0$ (nm)	$\langle \mu_2/\Gamma^2 \rangle_0$	$\langle R_{\text{core}} \rangle_0$ (nm)	$\sigma_{\text{core},0}$ (nm)	$Q_0^a$	$s_{\text{core},0}^b$	$s_{\text{corona},0}^c$
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
sample	steady-state after annealing at 170 °C						
	$\langle R_h \rangle_f$ (nm)	$\langle \mu_2/\Gamma^2 \rangle_f$	$\langle R_{\text{core}} \rangle_f$ (nm)	$\sigma_{\text{core},f}$ (nm)	$Q_f^a$	$s_{\text{core},f}^b$	$s_{\text{corona},f}^c$
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

<sup>a</sup>The aggregation number ( $Q$ ), where the subscripts 0 and  $f$  denote initial and final values, respectively, was calculated as  $4\pi\langle R_{\text{core}} \rangle^3/(3V_{\text{PB}})$  assuming the core is devoid of solvent, and  $V_{\text{PB}} = M_{\text{n,PB}}/\rho_{\text{PB}}N_{\text{Av}}$  is the volume per core chain,  $\rho_{\text{PB}} = 0.89 \text{ g/cm}^3$ ,  $N_{\text{Av}}$  is Avogadro's number, and  $M_{\text{n,PB}}$  is the molecular weight of PB, as reported in Table 1. The degree of core chain stretching ( $s_{\text{core}}$ ), was calculated as  $\langle R_{\text{core}} \rangle$  divided by the root-mean-square end-to-end distance of the core block using the statistical segment length of PB,  $b = 5.9 \text{ \AA}$ .<sup>64</sup> <sup>b</sup>The degree of corona chain stretching ( $s_{\text{corona}}$ ) calculated as the corona thickness ( $L_{\text{corona}} = \langle R_h \rangle - \langle R_{\text{core}} \rangle$ ) divided by the root-mean-square end-to-end distance of the corona block using the statistical segment length of PEO,  $b = 6.0 \text{ \AA}$ .<sup>65</sup>

the fragmentation kinetics were studied for six molecular weights of PB-PEO in [C<sub>2</sub>mim][TFSI] using *T*-jump DLS and TR-SAXS to determine the dependence of the fragmentation time  $\tau$  on the total degree of polymerization of the copolymer  $N_{\text{total}}$ . The change in  $\langle R_{\text{core}} \rangle$  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 [C<sub>2</sub>mim][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  $q_{\text{min}} = 0.009 \text{ \AA}^{-1}$  after annealing at 170 °C for approximately 26 min, which corresponds to  $\langle R_{\text{core}} \rangle = 50 \text{ nm}$  from the hard sphere approximation where  $qR_{\text{core}} = 4.493$ . 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  $\langle R_{\text{core}} \rangle$ . After annealing at 170 °C for approximately 2500 min,  $q_{\text{min}}$  shifts to  $0.012 \text{ \AA}^{-1}$  and  $\langle R_{\text{core}} \rangle = 37.4 \text{ nm}$ . 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  $\langle R_{\text{core}} \rangle$  from TR-SAXS.

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

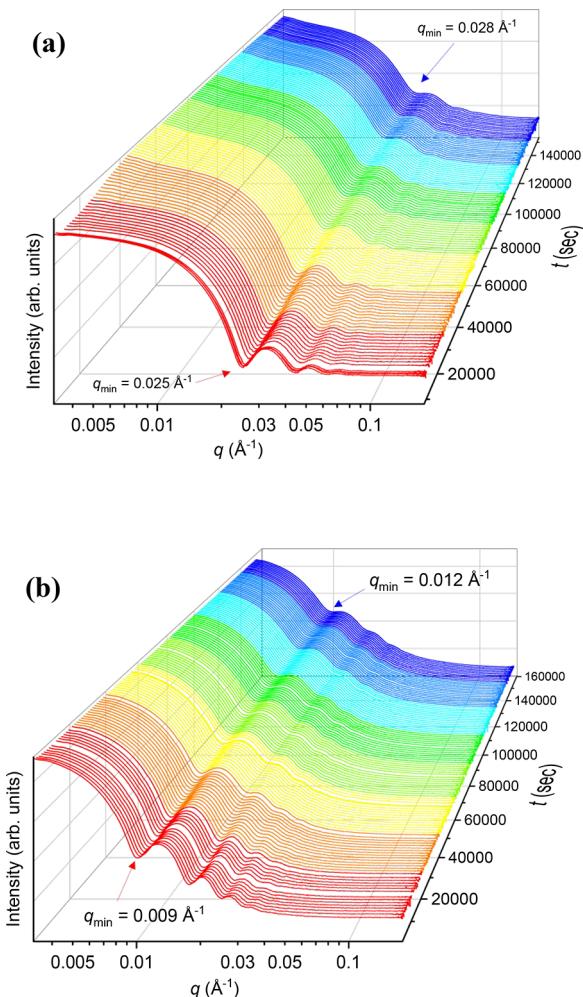
$$R(t) = \frac{\langle R_i \rangle_t - \langle R_i \rangle_f}{\langle R_i \rangle_0 - \langle R_i \rangle_f} = \exp[-(t/\tau)^n] \quad (4)$$

where  $R(t)$  is the normalized relaxation function, and  $i = h$  or core 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  $\tau$  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  $\langle R_{\text{core}} \rangle$  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  $\tau$ , 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  $\tau$  for the latter. This is consistent with previous reports on a single, low molecular weight of PB-PEO in various ionic liquids.<sup>36,40,45</sup> From Figure 4, the fragmentation time  $\tau$  increases strongly with increasing  $M$ ; the 11 kg mol<sup>-1</sup> diblock has a  $\tau$  on the order of 100 min, whereas the 100 kg mol<sup>-1</sup> diblock has  $\tau \approx 5000$  min. To corroborate the results from *T*-jump DLS, the time-dependent change in  $R_{\text{core}}$  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  $R_{\text{core}}$  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  $\langle R_{\text{core}} \rangle$  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  $\tau$  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 [C<sub>2</sub>mim][TFSI] depend strongly on  $M$ . Figure 6 shows the values of  $\tau$  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

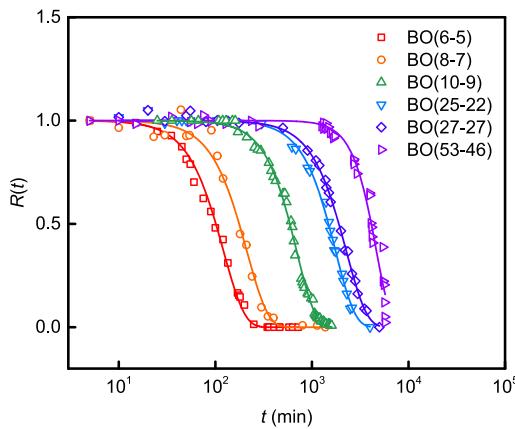


**Figure 3.** TR-SAXS intensity (logarithmic scale) as a function of  $q$  showing the evolution in the micelle core radius while annealing at 170 °C for 0.25 wt % (a) BO(6-5) in  $[\text{C}_2\text{mim}][\text{TFSI}]$ . The position of the first minimum in the form factor increases from  $q_{\min} = 0.025 \text{ \AA}^{-1}$  for short annealing times to  $q_{\min} = 0.028 \text{ \AA}^{-1}$  at longer annealing times. (b) BO(27-27) in  $[\text{C}_2\text{mim}][\text{TFSI}]$ . 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 ( $R_{\text{core}} \approx 4.493/q_{\min}$ ), so the radius of the micelle core decreases with increasing annealing time.

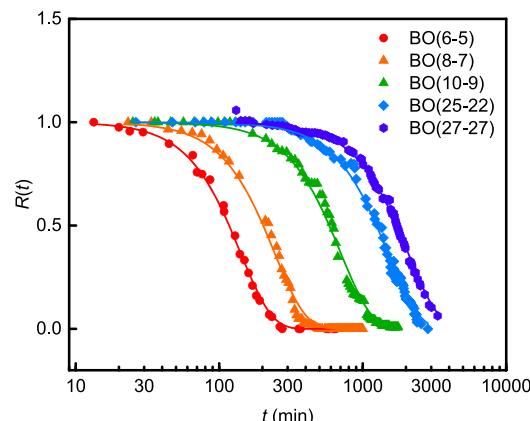
consistent. The fit to a power law gives the scaling  $\tau \sim N_{\text{total}}^{1.8 \pm 0.1}$ . Dormidontova proposed a scaling model for micelle fusion kinetics, and further assumed that fragmentation would proceed as the reverse of micelle fusion, and thus the  $N$  scaling would be the same for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

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

Assuming the fragmenting micelle is large, Dormidontova's theory proposes that  $\tau \sim N_{\text{corona}}^{1.8}$  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  $[\text{C}_2\text{mim}][\text{TFSI}]$ . 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  $n = 2.2 \pm 0.3$ .



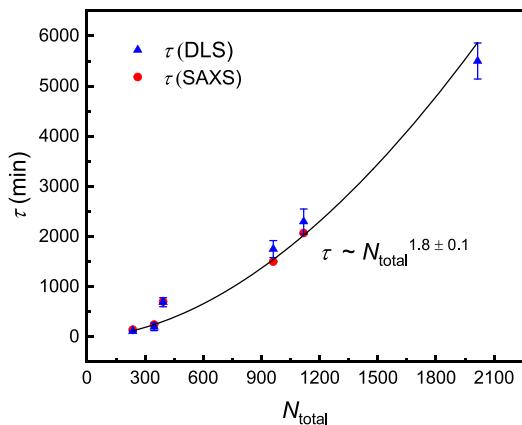
**Figure 5.** Time dependence of normalized  $\langle R_{\text{core}} \rangle$  for 0.25 wt % solutions of PB-PEO in  $[\text{C}_2\text{mim}][\text{TFSI}]$ . TR-SAXS measurements were performed at a relaxation temperature of 170 °C. Solid lines represent best fits to eq 4 with  $n = 2.0 \pm 0.1$ .

**Table 3. Molecular Weight Dependence of  $\tau$  Determined by T-Jump DLS and TR-SAXS at 170 °C for 0.25 wt % BO in  $[\text{C}_2\text{mim}][\text{TFSI}]$**

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

dependence to  $Q$ . The  $Q$  dependence of  $\tau$  is shown in the SI,<sup>465</sup> where  $\tau \sim Q_0^{1.7 \pm 0.1}$ . The agreement between the scaling of  $\tau$  with  $N_{\text{corona}}$  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  $N_{\text{total}}$  for very large  $Q$ .<sup>466</sup>

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,<sup>473</sup> so there is clearly an aspect of the fragmentation mechanism



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

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

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

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

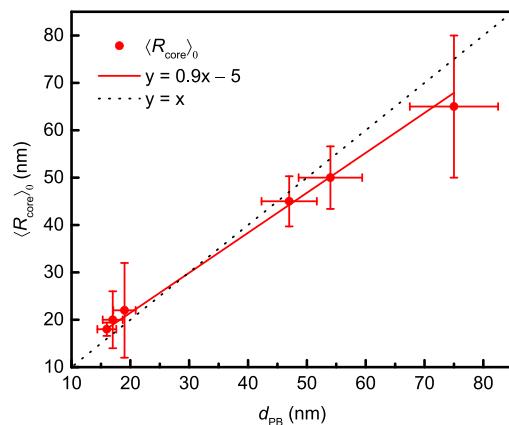
519 must be estimated. The transition state is represented  
520 schematically in Figure S11. We propose that the increase in  
521 the local concentration of corona chains near the neck point,  
522 i.e., the increase in the number of blobs per corona chain  $n_{\text{blob}}$ ,  
523 in the formation of the transition state is the primary barrier to  
524 fragmentation. To estimate the degree of corona crowding in  
525 the transition state first, we estimate the fraction of corona  
526 overlap in the transition state, and then relate the overlap  
527 fraction to the concentration of blobs per PEO chain in the  
528 neck region using the theory from Zhulina et al.<sup>12</sup> Using a  
529 simplified model for the transition state, illustrated in Figure  
530 S12, to determine the area of corona overlap in the transition  
531 state results in approximately 1/6 of the corona chains  
532 overlapping for all molecular weights. Details on this  
533 calculation are provided in the SI. Then, the concentration  
534 of PEO chains in the neck region of the transition state is  
535 determined by estimating the number of blobs per corona  
536 chain. In the neck region, the number of blobs per chain is  
537 expected to be approximately twice the number of blobs per  
538 chain in the as-prepared micelles ( $n_{\text{blob,neck}} = 2n_{\text{blob,sphere}} =$   
539  $2F_{\text{corona,0}}/k_{\text{B}}T$ ), whereas the outer spherical regions would have  
540  $n_{\text{blob,sphere}} = F_{\text{corona,0}}/k_{\text{B}}T$ , where the free energy is calculated  
541 using the as-prepared size of the micelle as shown in eq S11.  
542

The transition state free energy for the core and the interface  
542 was calculated using the steady state micelle dimensions. The  
543 transition state corona free energy was calculated assuming  
544 of the chains are described by  $F_{\text{corona,0}}/k_{\text{B}}T$ , and for the 1/6 of  
545 corona chains overlapped in the neck region, the corona free  
546 energy is  $2F_{\text{corona,0}}$ . Therefore, the total corona free energy in  
547 the transition state is calculated as  $F_{\text{corona,TS}}/k_{\text{B}}T = (5/6)$   
548  $F_{\text{corona,0}}/k_{\text{B}}T + (1/3)F_{\text{corona,0}}/k_{\text{B}}T$ . Details on this calculation  
549 are provided in the SI. From Table S4, the average  $n_{\text{blob}}$  is 16  
550 for BO(53–46) before a  $T$ -jump to 170 °C. Assuming that the  
551 number of blobs in the neck point is double the calculated  
552  $n_{\text{blob,0}}$  shown in Table S4, approximately 1/6 of the corona  
553 chains experience corona crowding where there are approx-  
554 imately 32 blobs per chain near the neck, whereas  $n_{\text{blob}}$  is 16 for  
555 chains in the spherical region of the micelle indicated in Figure  
556 S11. The fraction of corona chain overlap was determined to  
557 be 1/6 for all molecular weights by eq S8, and thus  $n_{\text{blob}}$  at the  
558 neck point is 8, 9, 10, 21, 24, and 32 for BO(6–5), BO(8–7),  
559 BO(10–9), BO(25–22), BO(27–27), and BO(53–46). This  
560 large increase in the local concentration of PEO chains in the  
561 neck region of the transition state supports the strong  
562 molecular weight dependence of micelle fragmentation  
563 kinetics. Because the confinement free energy of corona chains  
564 is directly proportional to  $n_{\text{blob}}$ , the chains near the neck point  
565 become much more confined and the entropic penalty  
566 associated with this chain confinement contributes to this  
567 being the primary activation barrier to fragmentation.  
568

569 Although these estimates are encouraging, we emphasize  
570 that the uncertainties in estimating the transition state free  
571 energy are quite large, and the differences in the calculated free  
572 energies for the spherical and “peanut” shaped micelles are  
573 only  $\sim 5$ –10  $k_{\text{B}}T$ . Given that the aggregation numbers are huge  
574 and the barrier is known to be of order 30–40  $kT$ ,<sup>36,45</sup> the  
575 estimated barriers per chain are small fractions of  $kT$ , and  
576 beyond the resolution of this calculation.  
577

**Molecular Weight Dependence of Initial Micelle Size by Direct Dissolution.** To understand the  $N$  dependence of the  
578 initial core size for micelles prepared by DD, SAXS  
579 measurements on the pure PB-PEO diblock copolymers  
580 were conducted. The 1D scattering intensity traces for these  
581

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



**Figure 7.** Dependence of  $\langle R_{\text{core}} \rangle_0$  (nm) on the PB domain size ( $d_{\text{PB}}$ ) 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  $d_{\text{PB}}$ , 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  $d_{\text{PB}}$  is assumed.

figure, the direct relationship between the domain size of the bulk copolymer and the initial micelle core radius is apparent. This result is also consistent with the previous reports on BO(8–7) in 1-alkyl-3-methylimidazolium bis-(trifluoromethylsulfonyl)imide-based ionic liquids.<sup>36,45</sup> It should be noted that a linear dependence should not necessarily be expected. The dependence begins to weaken slightly at higher molecular weights, as evidenced by the slight downward curvature of  $\langle R_{\text{core}} \rangle$  for the two largest polymers. The dependence of  $\langle R_{\text{core}} \rangle_0$  on  $d_{\text{PB}}$  follows the strong segregation limit scaling of  $2/3$ ,<sup>68</sup> as shown in Figure S13, while the apparent equilibrium scaling of  $\langle R_{\text{core}} \rangle$  with  $N_{\text{core}}$  for the initial micelles has an exponent closer to  $3/5$ , as shown in Figure 2. These slightly different dependences give rise to the curvature evident in Figure 7. Figure S13 shows the relationship of  $d_{\text{PB}}$  as a function of  $N_{\text{core}}$  for the domain size determined experimentally by SAXS, and the estimated domain spacing calculated from self-consistent field theory (SCFT); the experimental domain size of the PB lamellae agrees well with SCFT. We therefore speculate that during direct dissolution the solvent penetrates the PEO domains, peeling apart layers of PEO-decorated PB layers. Some surface instabilities in these separate layers drive a pinching-off process 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 micelles,<sup>69</sup> and experimentally in cylinder-to-sphere transitions<sup>70–75</sup> 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,<sup>62</sup> followed by rapid chain exchange.<sup>74</sup> On the basis of the lack of chain exchange in BO/[C<sub>2</sub>mim][TFSI] systems reported previously,<sup>36</sup> 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  $d_{\text{PB}}$  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 ( $s_{\text{PB}}$ ) 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 [C<sub>2</sub>mim]-[TFSI]. Disperse micelles prepared by direct dissolution in [C<sub>2</sub>mim][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 fragmentation. 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 characteristic 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  $N_{\text{corona}}$ . Future measurements on a series of polymers with constant  $N_{\text{core}}$  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.

## 683 ■ ASSOCIATED CONTENT

## 684 ■ Supporting Information

685 The Supporting Information is available free of charge at  
686 <https://pubs.acs.org/doi/10.1021/jacs.1c02147>.687 SEC-RI traces of PB-PEO diblocks,  $^1\text{H}$  NMR spectra of  
688 PB-PEO diblocks and  $[\text{C}_2\text{mim}][\text{TFSI}]$ , REPES results  
689 for PB-PEO in  $[\text{C}_2\text{mim}][\text{TFSI}]$  as-prepared by DD and  
690 steady-state after annealing, dependence of  $\langle R_h \rangle$  on the  
691 total degree of polymerization  $N_{\text{total}}$  for 0.25 wt % PB-  
692 PEO in  $[\text{C}_2\text{mim}][\text{TFSI}]$ , fragmentation times with fixed  
693  $n$ , TR-SAXS traces for BO(8–7), BO(10–9), and  
694 BO(25–22), calculations of corona chain disentanglement  
695 time, SAXS of bulk PB-PEO, high temperature  
696 liquid-phase transmission electron microscopy of  
697 BO(53–46) in  $[\text{C}_2\text{mim}][\text{TFSI}]$ , calculations of the  
698 fraction of corona overlap in the transition state, high  
699 temperature LP-TEM micrograph of BO(53–46) in  
700  $[\text{C}_2\text{mim}][\text{TFSI}]$  at 170 °C, schematic illustration of  
701 fragmentation transition state, illustration of simplified  
702 model transition state for calculating corona chain  
703 overlap, calculation of micelle free energy, domain  
704 spacing from SAXS of of BO( $x-y$ ) diblock copolymers  
705 in the bulk, Comparison of the domain size of PB  
706 determined by SAXS and the estimated domain size for  
707 a lamellar diblock copolymer using self-consistent field  
708 theory, estimated fragmentation lag time as a function of  
709 degree of polymerization, and dependence of fragmentation  
710 time on as-prepared aggregation number ([PDF](#))

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