

1 Multi-Chain Slip-Spring Simulations for Polyisoprene Melts

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12 Molecular simulations; rheology; entanglement;

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18 ABSTRACT

19 The multi-chain slip-spring (MCSS) model is a coarse-grained molecular model
20 developed for efficient simulations of the dynamics of entangled polymers. In this study,
21 the simulations were examined for viscoelasticity of polyisoprene (PI) melts, for which
22 the experimental data for well-characterized samples are available in the literature. The
23 conversion factor for the molecular weight was determined from the fitting of the
24 molecular weight dependence of zero-shear viscosity. According to the obtained value,
25 linear viscoelasticity of several linear PI melts was calculated, and the units of time and
26 modulus were determined. Based on the shift-factors reported experimentally for the
27 time-temperature superposition, the conversion factors for time and modulus were
28 obtained for different temperatures and used for the prediction of linear viscoelasticity
29 of 6-arm star PI melts, and viscosity growth under high shear for linear PI melts. The
30 predictions were in good agreement with the data, demonstrating the validity of the
31 method. The conversion factors determined were consistent with those reported for
32 polystyrene melt earlier, whereas the relations between the conversion factors are still
33 unknown.

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35 INTRODUCTION

36 The multi-chain slip-spring (MCSS) model is one of the coarse-grained multi-
37 chain models for simulations of entangled polymers in molecular level (Masubuchi 2014;
38 Masubuchi 2016a). Due to the slow relaxation nature, molecular dynamics simulations
39 with atomistic details are practically difficult for entangled polymers, yet attempted (Baig
40 et al. 2010b; Stephanou et al. 2010; Baig et al. 2010a). The widely used and established
41 approach is the bead-spring type simulations (Kremer and Grest 1990). Recent studies
42 with highly optimized computation codes have reported some simulation results for
43 moderately entangled polymer melts, in which the bead number per chain was increased
44 up to several hundred (Xu et al. 2015). For this approach, the computation costs are still
45 high due to the inter-bead interaction, which is short-range and repulsive like the Lennard-
46 Jones interaction. Such a short-range and strong repulsion limits the size of time step for
47 the numerical integration rather small. In this respect, the simulations with softer
48 interactions that allow interpenetration of the particles are favorable for the computation.
49 However, the simulations with such softer interactions cannot reproduce the entangled
50 polymer dynamics unless *ad-hoc* uncrossability between the chains is implemented (Pan
51 and Manke 2003). Attempts have been made for realization of uncrossability via the inter-
52 bond interaction (rather than the inter-bead interaction)(Kumar and Larson 2001) and via
53 the geometrical consideration (Padding and Briels 2001). However, such rigorous

54 treatments for uncrossability demands computation comparable to calculations with the
55 conventional bead-spring simulations. An alternative approach is a slip-spring model, in
56 which the entanglement is replaced by a virtual spring that connects bead-spring chains
57 without short-range repulsive interactions. To mimic the entanglement, the anchoring
58 points of the virtual spring slide along the connected chains. The virtual spring is
59 annihilated by a certain probability when one of the anchoring points reaches the chain
60 end, and at each chain end, a new virtual spring is created to maintain the average number
61 of springs in the system. Although the computational efficiency is less than the multi-
62 chain slip-link model (Masubuchi et al. 2001; Masubuchi et al. 2004; Masubuchi 2016b)
63 owing to the soft interactions the simulations for moderately entangled systems are quite
64 efficient.

65 Inspired by the single-chain version (Likhtman 2005; Uneyama 2011), the multi-
66 chain slip-spring models have been developed by a few groups in parallel (Chappa et al.
67 2012; Uneyama and Masubuchi 2012; Langeloth et al. 2013; Ramírez-Hernández et al.
68 2013; Vogiatzis et al. 2017). The critical issue for multi-chain modeling is the inter-bead
69 interaction. For the single chain version, because one of the ends for the virtual spring is
70 anchored at a certain point in space and is not spatially coupled with other chains, the
71 conformational distribution of the chain reduces to that of an ideal chain when we

72 integrate-out the degrees of freedom for the anchoring points. This construction ensures
73 that the ideal chain statistics is straightforwardly retained. In contrast, for the multi-chain
74 models, because both ends of the virtual spring are connected to the mobile chains, the
75 conformational distribution functions for the chains are affected by the inclusion of virtual
76 springs. Indeed, they play a role as an attractive interaction between the segments to
77 compress the chains. Because it has been established that the entanglement is purely
78 kinetic and does not affect the chain statistics, this artificial effect of the virtual spring
79 must be compensated. The rigorous treatment by the inter-bead repulsive, yet soft
80 interaction has been proposed (Chappa et al. 2012; Uneyama and Masubuchi 2012),
81 whereas the intermolecular interaction with a finite compressibility (Ramírez-Hernández
82 et al. 2013) and the weak repulsive interaction employed in dissipative particle dynamics
83 simulations (Langeloth et al. 2013; Masubuchi et al. 2016) are effective as well.

84 Nevertheless, the models with adequate treatments for the artifacts of the
85 inclusion of virtual springs are capable to simulate a variety of polymeric systems
86 including block-copolymers (Ramírez-Hernández et al. 2013; Ramírez-Hernández et al.
87 2018), mixtures of long and short chains (Langeloth et al. 2014), polymer solutions
88 (Masubuchi et al. 2016), branch polymers (Masubuchi 2018), network polymers
89 (Megariotis et al. 2018; Masubuchi and Uneyama 2019), etc. The simulations under fast

90 shear flows have also been attempted (Ramírez-Hernández et al. 2013; Masubuchi 2015;
91 Ramírez-Hernández et al. 2015; Masubuchi and Uneyama 2018a). The other important
92 direction is the linkage to the atomistic models (Sgouros et al. 2017; Vogiatzis et al. 2017)
93 and the other coarse-grained models (Masubuchi and Uneyama 2018b).

94 In this specific study, we shall report the consistency of the MCSS simulations
95 with the literature data for polyisoprene (PI) melts (Auhl et al. 2008; Matsumiya et al.
96 2014). In the earlier studies, we have evaluated the model by the comparison of
97 rheological data for polystyrene (PS) (Masubuchi 2018), and by the comparison to the
98 generic bead-spring model (Masubuchi and Uneyama 2018a; Masubuchi and Uneyama
99 2018c). Because the polymer dynamics is virtually universal and not strongly dependent
100 on the chemistry (Ferry 1980), the evaluation already performed would be sufficient.
101 However, parameter determination for specific chemistries is also necessary for practical
102 use. Besides, the parameters must be evaluated for the consistency to the other materials
103 and the theories. We have performed the MCSS simulations for several linear polymers
104 and a few star-branched polymers. For the linear polymers, the simulations under fast
105 shear flows were also conducted. We have confirmed that the simulation results are in
106 semi-quantitative agreement with the experimental data. The parameters used are
107 consistent with those for PS. Details are shown below.

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109 MODEL AND SIMULATIONS

110 The MCSS model proposed earlier (Uneyama and Masubuchi 2012) is used with
111 the modifications for shear flows (Masubuchi 2015) and the branch point dynamics
112 (Masubuchi 2018). In this model, a number of Rouse chains are dispersed in a simulation
113 box with a periodic boundary condition. The Rouse chains are connected by virtual
114 springs, for which both ends hop between the Rouse beads along the chain. The virtual
115 spring is annihilated with a certain probability when one of the ends comes to the chain
116 end, and vice versa, new virtual springs are introduced at the chain ends.

117 The free energy is given by

$$\begin{aligned} 118 \quad \frac{F}{k_{\text{B}}T} = & \frac{3}{2b^2} \sum_{i,k} (\mathbf{R}_{i,k+1} - \mathbf{R}_{i,k})^2 + \frac{3}{2N_s b^2} \sum_{\alpha} (\mathbf{R}_{S_{\alpha,1}, S_{\alpha,2}} - \mathbf{R}_{S_{\alpha,3}, S_{\alpha,4}})^2 \\ 119 \quad & + e^{v/kT} \sum_{i,k,j,l} \exp \left[-\frac{3}{2N_s b^2} (\mathbf{R}_{i,k} - \mathbf{R}_{j,l})^2 \right] \end{aligned} \quad (1)$$

120 The first and second terms in the right-hand side are the contributions from the Rouse
121 springs and the virtual springs, respectively. b is the average bond length between the
122 Rouse segments, $\mathbf{R}_{i,k}$ is the position of the Rouse beads k on the chain i , and N_s is
123 the parameter that determines the intensity of the virtual spring. $S_{\alpha,j}$ is the connectivity
124 matrix for the virtual spring α . The third term is the soft-core repulsive interaction
125 between the Rouse beads to eliminate the artificial effect on the chain statistics from the

126 virtual springs. $e^{\nu/kT}$ is the activity of the virtual springs to control the density of virtual
127 springs in the system by a grand canonical manner according to the chemical potential ν .

128 The governing equations and algorithms for the dynamics were derived from the
129 total free energy of the system to fulfill the detailed balance. The time development of
130 $\mathbf{R}_{i,k}$ obeys the Langevin equation of motion, and the SLLOD method was implemented
131 for the application of shear flows (Masubuchi 2015). The Glauber dynamics manage the
132 hopping of the virtual springs. For the branch polymers, the hopping across the branch
133 point (the so-called SHAB) is also considered, yet such a mechanism is not essential for
134 symmetric star polymers (Masubuchi 2018). The annihilation and creation of the virtual
135 springs are in balance under equilibrium by the master equation. For these dynamical
136 equations, in addition to the parameters that appear in the free-energy ($e^{\nu/kT}$ and N_s),
137 the friction coefficient for the bead motion ζ and that for the kinetics of virtual spring
138 ζ_s are introduced. For simplicity, we assume $\zeta = \zeta_s$. See the previous publication for
139 further details, the numerical implementation (Uneyama and Masubuchi 2012).

140 Simulations were performed for linear, and star-branched polymer melts with
141 dimensionless units, for which the units of length, energy, and time are the bond length
142 b , the thermal energy $k_B T$, and the diffusion time of a single bead $\tau_0 \equiv \zeta b^2 / k_B T$. For
143 convenience, to compare the simulated results with experimental data, we employ the unit

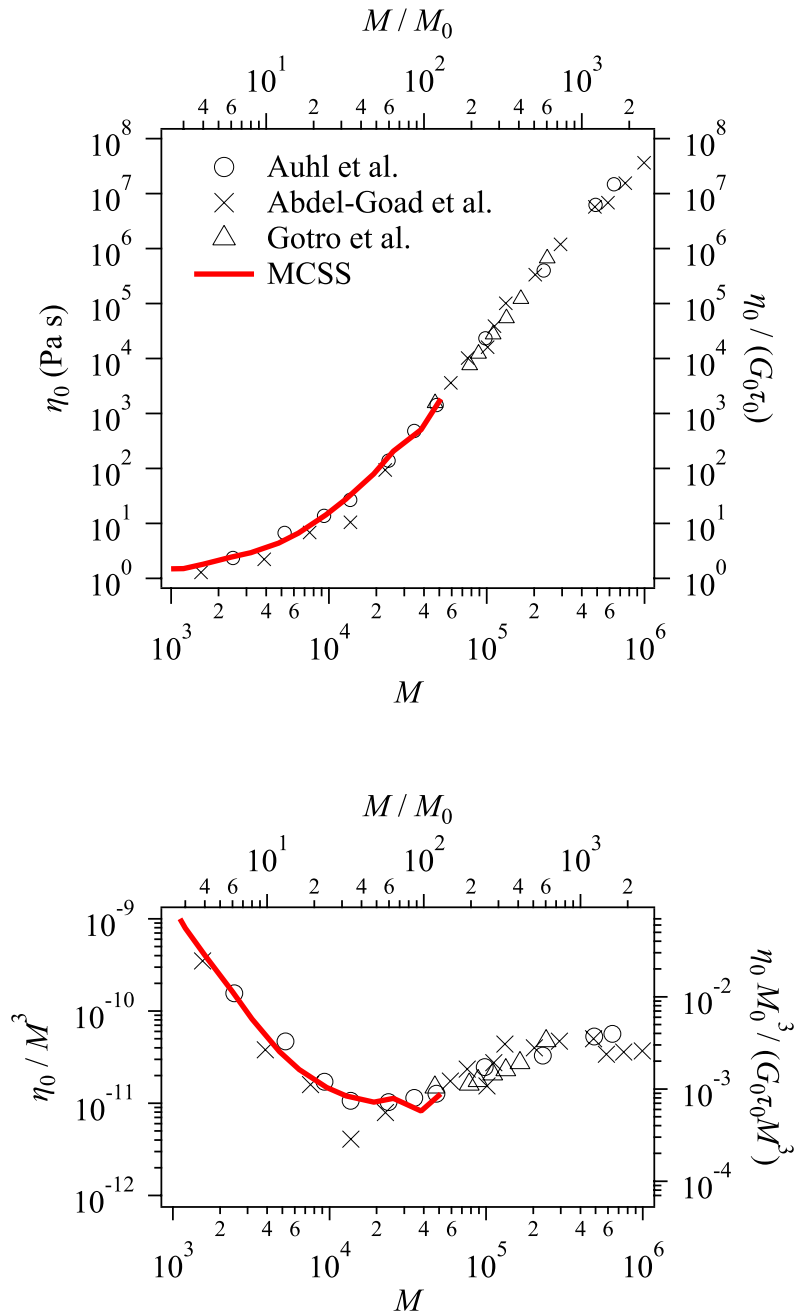
144 of molecular weight (i.e., corresponding molecular weight for the single bead) M_0 and
145 the unit of modulus $G_0 \sim b^3/k_B T$, instead of the units of length and energy. For each case,
146 200 molecules are placed in the simulation box. Owing to the previous studies, $e^{v/kT}$
147 and N_s were fixed at 0.036 and 0.5, respectively. The bead number density ρ_b was 4.
148 These values attain a particular density of slip-springs, which are distributed along the
149 chain with the average number of Rouse beads between consecutive anchoring points of
150 slip-springs, $N_e^{SS} = 3.5$.

151 Periodic boundary conditions were employed with the Lees-Edwards boundary
152 under shear. The systems were sufficiently equilibrated before data acquisition. The linear
153 relaxation modulus $G(t)$ was calculated from the stress auto-correlation obtained from
154 equilibrium simulations for an extended period that is at least ten times larger than the
155 longest relaxation time. From $G(t)$, storage and loss moduli, $G'(\omega)$ and $G''(\omega)$, were
156 obtained by the fitting of $G(t)$ to a multi-mode Maxwell function. The zero-shear
157 viscosity η_0 was obtained from $G(t)$. For statistics, eight independent simulation runs
158 starting from different initial configurations were performed for each condition, and the
159 results reported below are the averaged values.

160

161 RESULTS

162 Figure 1 shows the molecular weight M dependence of η_0 for linear polymers
163 in comparison to the data for PI melts reported experimentally (Gotro and Graessley
164 1984; Abdel-Goad et al. 2004; Auhl et al. 2008). The simulation result indicated by the
165 solid red curve reasonably captures the data, including the onset of entanglement. Namely,
166 the viscosity increases with an increase of M with a power-law manner, for which the
167 exponent is unity in the low- M regime, whereas it is around 3.5 in the high- M regime.
168 The transition between unentangled to entangled regimes is shown in a further clarity in
169 the bottom panel, in which the viscosity is reduced by M^3 , which is the scaling prediction
170 given by the classical tube model (Doi and Edwards 1986). Indeed, the critical molecular
171 weight for the onset of entanglement is indicated by the minimum of η_0/M^3 . From this
172 comparison, the conversion factor for the molecular weight of PI melt is determined as
173 $M_0 = 400$ (g/mol).



174

175 **Figure 1** Molecular weight dependence of zero-shear viscosity for linear polymers (top)
 176 and the viscosity normalized by the classical reptation prediction (bottom). The data for
 177 PI melts at 25°C (Gotro and Graessley 1984; Abdel-Goad et al. 2004; Auhl et al. 2008)
 178 are shown for comparison by symbols, and the simulation result is indicated by the solid
 179 curve.

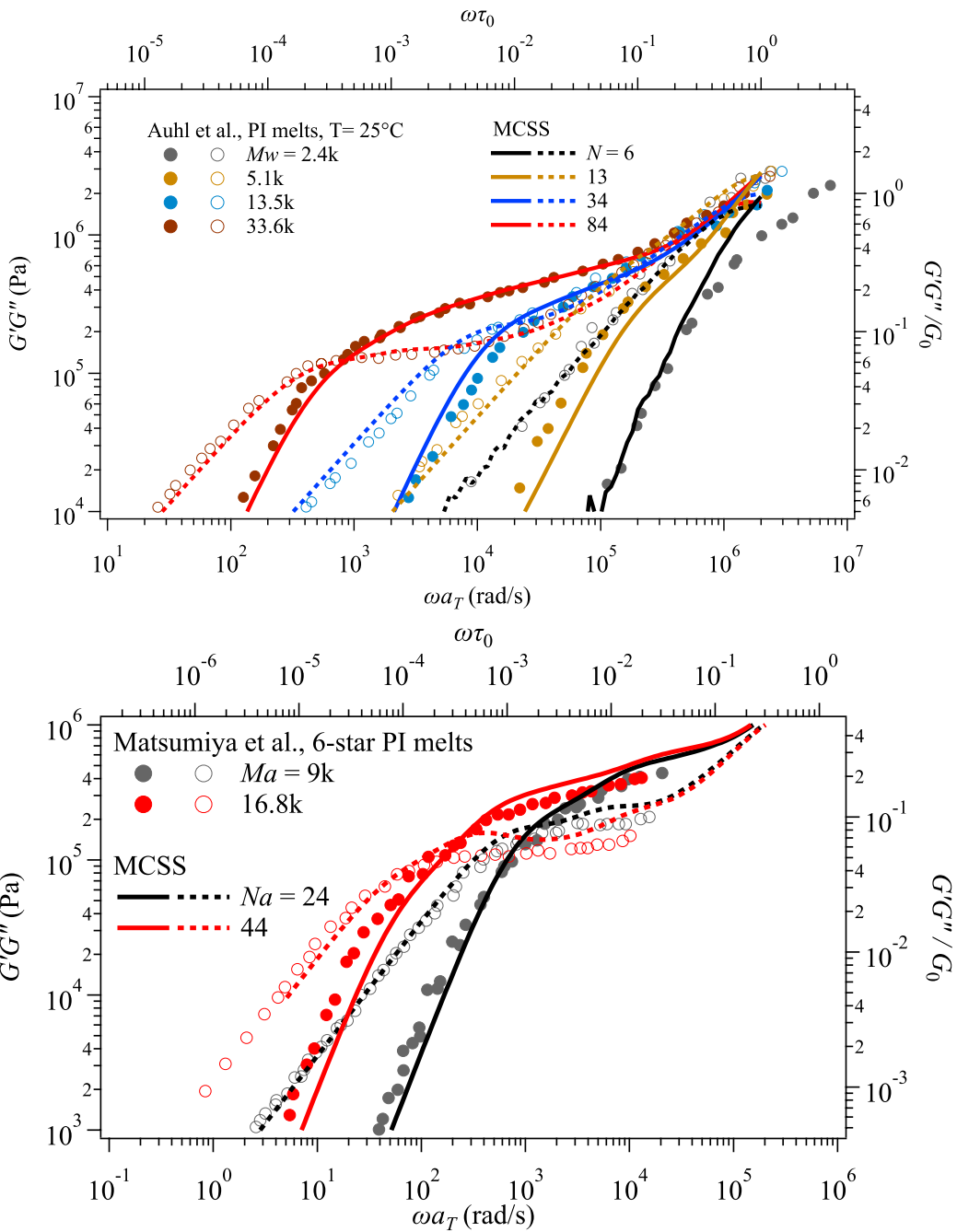
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Figure 2 shows the linear viscoelastic response for linear and star polymers. The

182 literature data of PI melts (Auhl et al. 2008; Matsumiya et al. 2014) are shown for
183 comparison for the samples that have the molecular weights comparable to the simulated
184 chains according to the conversion factor M_0 obtained above. For linear PI (seen in the
185 top panel), the conversion factors for time and modulus are chosen as $\tau_0 = 5 \times 10^{-7}$ sec
186 and $G_0 = 2 \times 10^6$ Pa at 25°C. These values give the unit viscosity $\tau_0 G_0 = 1$ Pa s, which
187 is consistent with the result shown in Figure 1. With the same set of the conversion factors,
188 the data for several different molecular weights are reasonably reproduced, except the
189 high- ω behavior for the shortest chain, due to an unavoidable cut-off of the model.

190 For branch polymers, the conversion factors must be shared with those for linear
191 polymers. Because the experimental data for star PI employed here (Matsumiya et al.
192 2014) were taken at 40°C, which is different from that for the linear polymers, we utilized
193 the shift factors for PI melts reported experimentally (Auhl et al. 2008) to obtain the
194 conversion factors. The determined values are $\tau_0 = 1.5 \times 10^{-6}$ sec and $G_0 = 2.1 \times$
195 10^6 Pa (at 40°C), and the comparison based on the conversion is shown in the bottom
196 panel. The prediction is not excellent, although it works to some extent. We wish to note
197 that better reproducibility has been reported for linear and branch polystyrenes
198 (Masubuchi 2018), for which the reference temperature is identical and the data are
199 reported from the same group.



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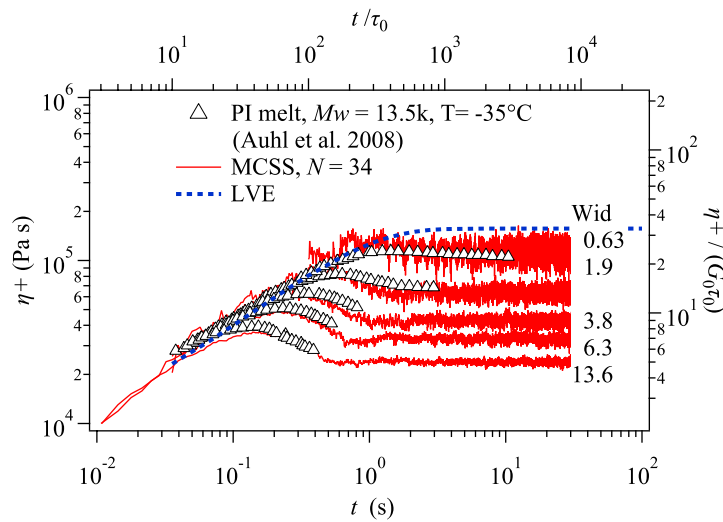
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Figure 2 Linear viscoelasticity of linear (top) and 6-arm star-branched (bottom) polymer melts compared to the experimental data for PI with corresponding molecular weights indicated in the figures (Auhl et al. 2008; Matsumiya et al. 2014). For the star polymers, the molecular weight and the bead number of each arm are indicated. The experimental data were taken at 25°C for linear PI and 40°C for star PI, respectively.

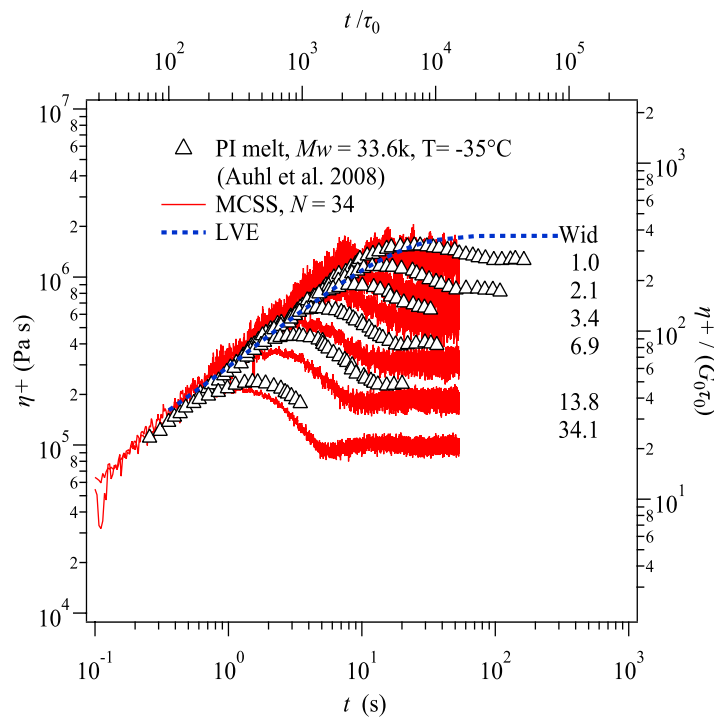
Figure 3 shows the viscosity growth for a few linear polymers, for which the

209 linear viscoelasticity is shown in Figure 2. For the prediction of nonlinear viscoelasticity,
210 the conversion factors must be shared with those for linear viscoelasticity. We
211 accommodated the temperature difference employing the shift factors reported (Auhl et
212 al. 2008) and obtained the conversion factors at -35°C as $\tau_0 = 2.8 \times 10^{-3}\text{sec}$ and $G_0 =$
213 $1.7 \times 10^6\text{Pa}$. The simulation results according to these conversion factors are in excellent
214 agreement for the short PI ($M_w = 13.5\text{k}$ shown in the top panel), whereas the simulations
215 are not entirely consistent with the data for the long PI ($M_w = 33.6\text{k}$ shown in the bottom
216 panel). In particular, the predictions for high shear rates are underestimated. This
217 inconsistency might be due to the model construction, in which we assume that the system
218 is not far from equilibrium. Meanwhile, we wish to note that the MCSS prediction under
219 high shear is in good agreement with the results obtained from the bead-spring
220 simulations (Masubuchi and Uneyama 2018c). Note also that better agreement can be
221 attained if the conversion factors could be tuned.

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225 **Figure 3** Viscosity growth under high shear for linear polymers compared with the
 226 experimental data for PI melts taken at -35°C (Auhl et al. 2008). The bead number per
 227 chain and the corresponding molecular weight of PI are $N = 34$ and $M_w = 13.5\text{k}$ (top),
 228 and $N = 84$ and $M_w = 33.6\text{k}$ (bottom). The simulation results and the experimental data
 229 are shown by the red curve and triangle, respectively. The linear viscoelastic envelope is
 230 shown by the blue dotted curve. The shear rate is indicated in terms of the Weissenberg
 231 number for the longest relaxation time.

232

233

234 DISCUSSION

235 Let us discuss the unit of molecular weight M_0 , which has been determined as
236 400 for PI in this study. This value is consistent with the previous study for PS, for which
237 $M_0 = 1250$ (Masubuchi 2018) if we consider the entanglement molecular weight.
238 Because M_0 is the molecular weight being carried by the single bead, and $N_e^{SS} = 3.5$,
239 the molecular weight between two anchoring points of slip-springs M_e^{SS} is 1400 for PI
240 and 4375 for PS. Both of these values are around four times smaller than the tabulated
241 values for the entanglement molecular weight M_e . For instance, M_e values for PI and
242 PS have been reported as 4820 (Auhl et al. 2008) and 14470 (Likhtman and McLeish
243 2002) based on the tube model. One of the reasons for the difference between M_e^{SS} and
244 M_e is the model dependence of the relation between the plateau modulus and the density
245 of entanglement. As we have discussed in the previous publications (Masubuchi et al.
246 2003; Masubuchi and Uneyama 2018b), the relation between the plateau modulus and the
247 number density of entanglements is model dependent as written by $G_N = Ank_B T$. Here,
248 A is the model-dependent factor that depends on the fluctuations considered, and n is
249 the number density of entanglement segments. For most of the cases, the value of M_e
250 experimentally reported is obtained for $A = 1$, which is for the affine network theory

251 without any fluctuation at entanglement. Meanwhile, $A = 0.8$ for the tube model due to
252 the chain sliding at the entanglement, and $A = 0.5$ for the cross-link network models
253 with Brownian motion of the entanglement. We have reported that our MCSS simulation
254 gives $A = 0.17$, which seems reasonable because of the various fluctuations including
255 the chain sliding (i.e., the hopping of the slip-spring, which causes the fluctuation of
256 numbers of beads between entanglements), and the Brownian motion of the beads and
257 the conformational fluctuations of the slip-spring (which causes the fluctuation of
258 transient link positions) (Masubuchi and Uneyama 2018b). Based on $A = 0.17$, M_e^{SS}
259 can be calculated from M_e mentioned above as 1024 and 3074 for PI and PS,
260 respectively. These values are even smaller than the M_e^{SS} determined from the fitting,
261 possibly due to the difference of the segment density.

262 The unit of modulus G_0 obtained for PI is also consistent with that obtained for
263 PS earlier (Masubuchi 2018). For PI, $G_0 = 2 \times 10^6 \text{ Pa}$ (at 25°C), whereas for PS, $G_0 =$
264 $9.1 \times 10^5 \text{ Pa}$ (at 169.5°C). Both of these values are around three times larger than the
265 plateau modulus reported, implying the consistency between the simulations for PI and
266 PS. However, the relation between G_0 and M_0 seems non-trivial. From dimensional
267 analysis, we obtain a relationship $G_0 = B\rho RT/M_0$. Here, B is a constant, ρ is the
268 density, and R is the gas constant. Note that this relation is similar to, yet different from

269 that discussed above for G_N and M_e . For the tube model proposed by Likhtman and
270 McLeish (Likhtman and McLeish 2002), it has been reported that $M_0^{\text{LM}} = 4820$ and
271 $G_0^{\text{LM}} = 6 \times 10^5 \text{ Pa}$ for the PI melt examined in this study. With the density of $\rho =$
272 0.913 g/cm^3 , these values give $B^{\text{LM}} = 1.3$. For our case, $B = 0.35$. We have no
273 explanation for this value of B .

274 Let us turn our attention to the unit of time that was determined as $\tau_0 =$
275 $5 \times 10^{-7} \text{ sec}$. According to the study by Auhl et al. (Auhl et al. 2008), the values for the
276 tube model are $\tau_0^{\text{LM}} = 1.3 \times 10^{-5}$. The difference for the unit of time can be explained
277 by the unit of molecular weight. For the tube model, $M_0^{\text{LM}} = 4820$ has been reported,
278 whereas our simulation gives $M_0 = 400$. Because the unit of time corresponds to the
279 Rouse relaxation time of the segment, the ratio for the unit of time would be the square
280 of the ratio for the unit of molecular weight. This estimation is consistent with the values
281 mentioned above. Similar consistency with the tube model can be confirmed for PS.

282 It is fair to note that Ramírez-Hernández et al. (Ramírez-Hernández et al. 2015)
283 have reported a similar simulation study for the rheology of linear PI melts examined
284 above. Their model, so-called TIEPOS (theoretically informed entangled polymer
285 simulations), is the same with our MCSS model in the fundamental construction, in which
286 bead-spring chains with weakly repulsive interactions are connected by virtual springs.

287 The main difference is that they consider the inter-bead interaction based on the
288 compressibility, whereas we introduce the interaction to retain the ideal chain statistics.
289 The other difference is that they allow only one slip-spring connected to a single bead to
290 avoid passing between slip-springs along the chain. Probably due to these differences, the
291 effect of slip-spring on the chain dynamics seems not the same with each other. To
292 reproduce the viscoelasticity of PI melt with $M_w = 33.6\text{k}$, we need the bead number per
293 chain of $N = 84$, whereas they only have $N = 32$. The average number of slip-springs
294 on one chain Z is 25 for our case, whereas 8.6 for them. This comparison demonstrates
295 that their model has an advantage in the level of coarse-graining and for a reduction of
296 computation costs. However, the efficiency of their entanglement is inexplicable from the
297 discussion for M_e^{SS} mentioned above. According to the difference of Z , a naïve
298 estimation of A value for TIEPOS would be ca. 0.5, which is close to the cross-link
299 networks. This estimation implies that the incompressibility for the chains, and the
300 uncrossability between slip-springs significantly suppress the fluctuations around their
301 entanglement. Besides, they compared the statistical distributions of their model with
302 predictions by a mean-field single-chain model, and reported that the simulation data
303 agree well with the theoretical predictions. Their result implies that the fluctuations in
304 TIEPOS model is close to those in the mean-field single-chain model. However, in multi-

305 chain models, in general, the environment around a chain fluctuates (because the
306 environment itself consists of fluctuating chains) and the statistical distributions deviate
307 from the mean-field statistics without fluctuations. The friction for the slip-spring kinetics
308 is the other important parameter, but the setting in TIEPOS seems the same with our
309 simulations. Nevertheless, further assessment is necessary for the nature of the multi-
310 chain slip-spring models.

311

312 CONCLUSIONS

313 We performed MCSS simulations with the model parameters fixed at $e^{v/kT} =$
314 0.036, $N_s = 0.5$, $\zeta = \zeta_s$, and $\rho_b = 4$ to compare the results with the experimental data
315 for PI melts. The conversion factor for the molecular weight (i.e., the molecular weight
316 carried by the single Rouse bead) was obtained as $M_0 = 400$ (g/mol) from the molecular
317 weight dependence of the zero-shear viscosity for linear polymers. According to the M_0
318 value, we calculated the linear viscoelasticity for several linear and 6-arm star-branch
319 polymers, for which the experimental data are available for PI with the corresponding
320 molecular weight. For the linear polymers, the simulation results agree with the data for
321 the conversion factors obtained by the fitting as $\tau_0 = 5 \times 10^{-7}$ sec and $G_0 = 2 \times 10^6$ Pa
322 at 25°C. Based on these values, the conversion factors for different temperatures were

323 obtained according to the shift factors reported experimentally. With these conversion
324 factors, the simulation results for linear viscoelasticity of star PI taken at 40°C, and for
325 viscosity growth under high shear of linear PI obtained at -35°C are in reasonable
326 agreement with the experimental data. The conversion factors thus determined are
327 consistent with those for PS. However, the relations between the conversion factors are
328 still unknown, and further evaluation and systematic studies are necessary. For example,
329 the dependence of conversion factors on the model parameters is worth investigating.
330 Studies in such directions are ongoing, and the results will be published elsewhere.

331

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