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The role of intramolecular barriers on the glass transition of polymers: Computer simulations versus mode coupling theory

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We present computer simulations of a simple bead-spring model for polymer melts with intramolecular barriers. By systematically tuning the strength of the barriers, we investigate their role on the glass transition. Dynamic observables are analyzed within the framework of the mode coupling theory (MCT). Critical nonergodicity parameters, critical temperatures, and dynamic exponents are obtained from consistent fits of simulation data to MCT asymptotic laws. The so-obtained MCT λ -exponent increases from standard values for fully flexible chains to values close to the upper limit for stiff chains. In analogy with systems exhibiting higher-order MCT transitions, we suggest that the observed large λ -values arise form the interplay between two distinct mechanisms for dynamic arrest: general packing effects and polymer-specific intramolecular barriers. We compare simulation results with numerical solutions of the MCT equations for polymer systems, within the polymer reference interaction site model (PRISM) for static correlations. We verify that the approximations introduced by the PRISM are fulfilled by simulations, with the same quality for all the range of investigated barrier strength. The numerical solutions reproduce the qualitative trends of simulations for the dependence of the nonergodicity parameters and critical temperatures on the barrier strength. In particular, the increase in the barrier strength at fixed density increases the localization length and the critical temperature. However the qualitative agreement between theory and simulation breaks in the limit of stiff chains. We discuss the possible origin of this feature. © 2009 American Institute of Physics. [doi:10.1063/1.3266852]

I. INTRODUCTION

Since they do not easily crystallize, polymers are probably the most extensively studied systems in relation with the glass transition phenomenon. Having said this, their macromolecular character, and in particular chain connectivity, must not be forgotten. The most evident effect of chain connectivity is the sublinear increase in the mean squared displacement (MSD) (Rouse-like)¹ arising after the decaging process, in contrast to the linear regime found in nonpolymeric glass-formers. Moreover, in the case of strongly entangled polymer chains, the reptation model predicts other two sublinear regimes between the Rouse and linear regimes.^{1–3}

Another particular ingredient of polymers is that, apart from fast librations or methyl group rotations,⁴ every motion involves jumps over carbon-carbon rotational barriers and/or chain conformational changes. Intramolecular barriers play a decisive role in the physical properties of polymer systems. Thus, they are responsible of partial or total crystallization.^{5,6} They also enhance dynamic features which are usually associated to reptation,^{7,8} which controls rheological properties.³

Models for semiflexible polymers are of great interest, since they can be applied to many important biopolymers such as proteins, DNA, rodlike viruses, or actin filaments.^{9–11} Moreover, chain stiffness seems to play an important role in the absorption behavior of polymers at interfaces.^{12,13} Thus, an understandig of the role of intramolecular barriers on structural, dynamic, and rheological properties of polymers is of practical as well as of fundamental interest.

In this work we investigate, by means of MD simulations, the role of intramolecular barriers on the glass transition of polymer melts, by systematically tuning barrier strength in a simple bead-spring model. We discuss the obtained results within the framework of the mode coupling theory (MCT) of the glass transition.^{14–19} We extend preliminary results reported by us in Ref. 20 by testing a large set of predictions, including the factorization theorem and timetemperature superposition principle (TTSP). A consistent set of dynamic exponents associated to asymptotic scaling laws is obtained. By increasing the barrier strength a crossover is observed for the values of the so-called λ -exponent. In the limit of fully flexible chains λ takes values ~0.7, characteristic of simple fluids dominated by packing effects. On the contrary, for strong intramolecular barriers the λ -values approach the upper limit $\lambda = 1$ characteristic of higher-order MCT transitions. The latter arise in systems with different

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coexisting mechanisms for dynamic arrest.^{21–23} In the system investigated here, the obtained results suggest an interplay between general packing effects and polymer-specific intramolecular barriers.

Chong and co-workers^{24,25} have recently presented an extension of the MCT to simple fully flexible bead-spring models of polymer systems, in the framework of the polymer reference interaction site model (PRISM).²⁶⁻²⁸ In this formalism each molecule is divided into interaction sites corresponding to monomers. A key assumption of the PRISM is the replacement of the site-specific intermolecular surroundings of a monomer by an averaged one (equivalent-site approximation), while keeping the fully intramolecular dependence. We have tested the PRISM approximations used by MCT in the polymer model here investigated, which incorporates intramolecular barriers. Likewise, we have solved the MCT equations for the location of the MCT "glass transition" temperatures (MCT critical temperatures) and for the nonergodicity parameters, which quantify the stability of density fluctuations in the reciprocal space. We compare solutions of the MCT equations with the results obtained from the phenomenological analysis of the simulation data. We observe that the theory reproduces qualitative trends in the nonergodicity parameters and critical temperatures. However, the agreement breaks as the limit of stiff chains is approached. We discuss the possible origins of this feature.

The article is organized as follows. In Sec. II we describe the model and give simulation details. Static correlators are shown in Sec. III. Moreover the PRISM approximations are tested for representative values of the barrier strength. Section IV presents qualitative dynamic trends as a function of the barrier strength. In Sec. V we summarize the universal predictions of the MCT and the equations of motion of the version for polymer melts introduced by Chong and co-workers.^{24,25} In Sec. VI we perform a phenomenological analysis of simulation data within the MCT, by testing universal scaling laws, and deriving their associated dynamic exponents. In Sec. VII we compare the results of the former analysis with numerical solutions of the MCT equations. We discuss the observed differences for stiff chains in Sec. VIII. Conclusions are given in Sec. IX.

II. MODEL AND SIMULATION DETAILS

We have performed molecular dynamics (MD) simulations of a bead-spring model for which we have implemented bending and torsional intramolecular barriers. The monomermonomer interaction is given by a corrected soft-sphere potential

$$V(r) = 4\epsilon [(\sigma/r)^{12} - C_0 + C_2(r/\sigma)^2],$$
(1)

where $\epsilon = 1$ and $\sigma = 1$. The potential V(r) is set to zero beyond the cutoff distance $r \ge c\sigma$, with c=1.15. The values C_0 $=7c^{-12}$ and $C_2=6c^{-14}$ guarantee continuity of potential and forces at $r=c\sigma$. The potential V(r) is purely repulsive. It does not show local minima within the interaction range $r < c\sigma$. Thus, it drives dynamic arrest only through packing effects. Along the chain backbone, of *N* monomers, an additional finitely extensible nonlinear elastic (FENE) potential^{29,30} is used to introduce bonds between consecutive monomers:

$$V_{\text{FENE}}(r) = -\epsilon K_F R_0^2 \ln[1 - (R_0 \sigma)^{-2} r^2], \qquad (2)$$

where $K_F=15$ and $R_0=1.5$. The superposition of potentials (1) and (2) provides an effective bond potential for consecutive monomers with a sharp minimum at $r \approx 0.985$, which makes bond crossing impossible.

Intramolecular barriers are implemented by means of a combined bending V_B , and torsional potential V_T . We have used the potentials proposed by Bulacu and van der Giessen in Refs. 8 and 31. The bending potential acts on three consecutive monomers along the chain. The angle between adjacent pairs of bonds is mantained close to the equilibrium value θ_0 =109.5° by the cosine harmonic bending potential

$$V_{\rm B}(\theta_i) = (\epsilon K_{\rm B}/2)(\cos \theta_i - \cos \theta_0)^2, \qquad (3)$$

where θ_i is the bending angle between consecutive monomers i-1, i and i+1 (with $2 \le i \le N-1$).

The torsional potential constrains the dihedral angle $\phi_{i,i+1}$, which is defined for the consecutive monomers i-1, i, i+1, and i+2 (with $2 \le i \le N-2$), as the angle between the two planes defined by the sets (i-1,i,i+1) and (i,i+1,i+2). The form of this potential is

$$V_{\mathrm{T}}(\theta_i, \theta_{i+1}, \phi_{i,i+1}) = \epsilon K_{\mathrm{T}} \sin^3 \theta_i \sin^3 \theta_{i+1} \sum_{n=0}^3 a_n \cos^n \phi_{i,i+1}.$$
(4)

The values of the coefficients a_n are $a_0=3.00$, $a_1=-5.90$, $a_2=2.06$, and $a_3=10.95$.^{8,31} The torsional potential depends both on the dihedral angle $\phi_{i,i+1}$ and on the bending angles θ_i and θ_{i+1} . As noted in Refs. 8 and 31, numerical instabilities arising when two consecutive bonds align are naturally eliminated by choosing the torsional potential (4), without the need of imposing rigid constraints on the bending angles.

In the following, temperature *T*, time *t*, distance, wave vector *q*, and monomer density ρ are given, respectively, in units of ϵ/k_B (with k_B the Boltzmann constant), $\sigma(m/\epsilon)^{1/2}$ (with *m* the monomer mass), σ , σ^{-1} , and σ^{-3} . We investigate, at fixed monomer density ρ =1.0, the temperature dependence of the dynamics for different values of the bending and torsion strength, (K_B, K_T)=(0,0), (4,0.1), (8,0.2), (15,0.5), (25,1), (25,4), and (35,4). In the following, all the data presented in the figures and discussed in the main text will correspond to ρ =1.0. This value will not be, in general, explicitly mentioned there. We have also studied the case (K_B, K_T)=(35,4) at density ρ =0.93. The specific information of this case is given in Table I (see below). We investigate typically 8–10 different temperatures for each set of values (K_B, K_T).

We simulate 300 chains, each chain consisting of N = 10 monomers of mass m=1, placed in a cubic simulation box of lenght $L_{\text{box}} = 14.4225$ for $\rho = 1.0$, or $L_{\text{box}} = 14.7756$ for $\rho = 0.93$, with periodic boundary conditions. Equations of motion are integrated by using the velocity Verlet scheme.³² Computational expense is reduced by implementing a linkedcell method.³² We use a time step ranging from 10^{-4} to 5 $\times 10^{-3}$. We take shorter and longer steps for, respectively, higher and lower values of temperatures, bending, and tor-

TABLE I. Values of the MCT λ -exponents and critical temperatures T_c for different ρ and barrier strength. Also included are the mean chain end-to-end radii R_{ee}^c at T_c .

ρ	K _B	K _T	$R_{\rm ee}^{\rm c}$	$T_{\rm c}$	λ
1	0	0	3.6	0.48	0.761
1	4	0.1	4.4	0.54	0.767
1	8	0.2	4.7	0.62	0.773
1	15	0.5	5.2	0.75	0.785
1	25	1	5.5	0.92	0.827
1	25	4	6.4	1.02	0.845
1	35	4	6.5	1.23	0.862
0.93	35	4	6.9	1.02	0.885

sional constants. The system is prepared by placing and growing the chains randomly in the simulation box, with a constraint avoiding monomer core overlap. The initial monomer density is ρ =0.375. Equilibration consists of a first run where the box is rescaled periodically by a factor 0.99 < f<1 until the target density ρ is reached, and a second isochoric run at that ρ . Thermalization at the target T is achieved by periodic velocity rescaling. After reaching equilibrium, energy, pressure, chain radii of gyration, and end-to-end distances show no drift. Likewise, dynamic correlators show no aging effects. Once the system is equilibrated, a microcanonical run is performed for production of configurations, from which static and dynamic correlators are computed. Static correlators presented here are averaged over typically 300 equispaced configurations. Dynamic correlators are averaged over typically 40 equispaced time origins. The typical duration of a production run is of 40-200 million time steps for, respectively, high and low temperatures.

III. STATIC PROPERTIES

Simulation results presented in this work correspond to isotropic phases. We do not observe signatures of global orientational order induced by chain stiffness for the investigated state points. Thus, we obtain negligible values for the quantity $P_2(\Theta) = (3\langle\cos^2 \Theta\rangle - 1)/2$, where Θ is the angle between the end-to-end vectors of two chains, and average is performed over all pairs of distinct chains. Local orientational order is also negligible. This is evidenced by computing a similar correlator $P_2(\Theta; r_{cm})$. In this case the average is performed only over pairs of distinct chains for which the distance between their respective centers-of-mass is less than r_{cm} . Negligible values of $P_2(\Theta; r_{cm})$ are obtained for r_{cm} > 2.0. Weak local orientational order $|\langle P_2(\Theta; r_{cm}) \rangle| \leq 0.1$ is observed only for very small interchain distances, without any systematic dependence on the barrier strength.

A. Static structure factors and chain form factors

Now we present results for static structure factors and chain form factors, both for fully flexible chains and for a representative case of stiff chains. Let us consider an isotropic homogeneous system of volume V containing n identical chains of N monomers. The densities of chains and monomers are, respectively, denoted by $\rho_c = n/V$ and $\rho = nN/V$. Let

us denote the location of a monomer along its chain by the index $1 \le a \le N$. The site-site static structure factor for monomers of indices *a* and *b* is defined as:

$$S_{ab}(q) = \frac{1}{n} \langle \rho_a(-\mathbf{q}, 0) \rho_b(\mathbf{q}, 0) \rangle.$$
(5)

Brackets denote both ensemble and orientational average (the latter performed over all the orientations of the wave vector \mathbf{q}). The monomer density distribution for wave vector \mathbf{q} is given by

$$\rho_a(\mathbf{q}) = \sum_{j=1}^n \exp[i\mathbf{q} \cdot \mathbf{r}_j^a].$$
(6)

In this expression \mathbf{r}_{j}^{a} is the position vector of the *a*th monomer in the *j*th chain $(1 \le j \le n)$. The quantity $S_{ab}(q)$ can be splitted into intrachain and interchain *a-b* correlations:

$$S_{ab}(q) = \omega_{ab}(q) + \rho_c h_{ab}(q), \tag{7}$$

or in matrix form, $\mathbf{S}(q) = \mathbf{w}(q) + \rho_c \mathbf{h}(q)$. In Eq. (7) $\omega_{ab}(q)$ and $h_{ab}(q)$, respectively, denote the intrachain and interchain correlations between monomers of type *a* and *b*. By averaging over all the possible pairs (a, b) we obtain the static correlators S(q), $\omega(q)$ and h(q), which are related through:

$$S(q) = \omega(q) + \rho h(q). \tag{8}$$

In this expression S(q) is the total static structure factor, which equivalently can be obtained as $S(q) = (nN)^{-1} \langle \rho(-\mathbf{q}, 0)\rho(\mathbf{q}, 0) \rangle$, where $\rho(\mathbf{q}) = \sum_{a=1}^{N} \rho_a(\mathbf{q})$ is the total monomer density distribution. In Eq. (8) the chain form factor, $\omega(q)$, accounts for all the static intrachain correlations, while h(q) accounts for all the static interchain correlations.

Figure 1 (top panel) shows simulation results for S(q) as a function of temperature for fully flexible chains, $(K_{\rm B}, K_{\rm T})$ =(0,0). Data for representative stiff chains, $(K_{\rm B}, K_{\rm T})$ =(25,1), are shown in the bottom panel. In both cases, no signature of crystallization is present. Indeed no sharp Bragg peaks are observed. In both cases S(q) shows a maximum at $q_{\rm max} \approx 7.0$. Since $S(q_{\rm max})$ comes from the packing in the first shell around a monomer, the latter corresponds to a typical distance $2\pi/7.0\approx 0.90$ in the real space between neighboring monomers. On cooling, the peak at $q_{\rm max} \approx 7.0$ increases in intensity, which is a signature of increasing short-range order.

In Fig. 2 we show, for the former values of $(K_{\rm B}, K_{\rm T})$, the corresponding results for the form factors $\omega(q)$. We note that in the case of fully flexible chains the form factor is nearly independent on temperature. The form factor for stiff chains exhibits a certain *T*-dependence, which is however rather weak in comparison with that of S(q). The *T*-dependence of $\omega(q)$ becomes more clear at low *q*-values. The way the form factor behaves on lowering the temperature is directly connected with the values of the mean chain end-to-end radius $R_{\rm ee}$. Thus, by decreasing temperature from T=2.0 to T=0.96, the computed $R_{\rm ee}$ increases from 4.8 to 5.5 for the selected stiff chains. This leads, for lower *T*, to a stronger decay in $\omega(q)$ at low-*q*. On the other hand, the value $R_{\rm ee} = 3.6$ for the fully flexible chains is almost *T*-independent, leading to a negligible *T*-dependence of $\omega(q)$.



FIG. 1. Temperature dependence of the static structure factor S(q) for fully flexible chains (top panel) and for chains with barrier strength $(K_{\rm B}, K_{\rm T}) = (25, 1)$ (bottom panel).

B. Test of the PRISM approximations

The MCT for polymer melts developed by Chong and co-workers^{24,25} invokes several approximations of the PRISM theory.²⁷ In this subsection we summarize such approximations and test their validity for all the investigated range of barrier strength. The site-site direct correlation function, $c_{ab}(q)$, is introduced via the generalized Ornstein-Zernike relation for polyatomic molecules, or "reference interaction site model,"³³

$$h_{ab}(q) = \sum_{x,y=1}^{N} \omega_{ax}(q) c_{xy}(q) [\omega_{yb}(q) + \rho_c h_{yb}(q)], \qquad (9)$$

in which intramolecular contributions are accounted by the form factor terms $\omega_{ab}(q)$. By inserting Eq. (7) in Eq. (9), $c_{ab}(q)$ is related to $S_{ab}(q)$ and $\omega_{ab}(q)$ as:



FIG. 2. Temperature dependence of the form factor $\omega(q)$ for fully flexible chains (lines) and for chains with barrier strength ($K_{\rm B}, K_{\rm T}$)=(25,1) (symbols). For clarity, the inset shows results in the range of low-q. Different colors correspond to different temperatures, following the legends of Fig. 1.



FIG. 3. Test of the equivalent-site approximation, Eq. (11), for fully flexible chains at T=0.50. Top and bottom panels compare c(q) with, respectively, matrix elements $c_{aa}(q)$ and $c_{a5}(q)$. The insets enhance the region around the wave vector q_{max} for the maximum of the static structure factor S(q).

$$\rho_{\rm c} c_{ab}(q) = \omega_{ab}^{-1}(q) - S_{ab}^{-1}(q). \tag{10}$$

Here $\omega_{ab}^{-1}(q)$ and $S_{ab}^{-1}(q)$ are the elements of, respectively, the matrices $\mathbf{w}^{-1}(q)$ and $\mathbf{S}^{-1}(q)$, which are defined as the inverses of $\mathbf{w}(q)$ and $\mathbf{S}(q)$.

In the *equivalent-site approximation* (which is exact for polymer rings) of the PRISM, chain end effects are neglected and all sites are treated equivalently for interchain correlations. Thus, c_{ab} is replaced by the average over all (a,b)-pairs:

$$c_{ab}(q) \approx c(q). \tag{11}$$

By introducing this approximation in Eq. (9) and averaging over all (a,b) pairs we find $h(q) = \omega(q)c(q)[\omega(q) + \rho h(q)]$. By introducing Eq. (8) in the latter expression we arrive to the scalar equation

$$\rho c(q) = 1/\omega(q) - 1/S(q),$$
 (12)

also known as PRISM equation.

In Figs. 3 and 4 we test the validity of the equivalent-site approximation $c_{ab}(q) \approx c(q)$. We calculate $c_{ab}(q)$ and c(q), respectively, through Eqs. (10) and (12), by using the quantities $\omega_{ab}^{-1}(q)$, $S_{ab}^{-1}(q)$, $\omega(q)$, and S(q) as computed from the simulations. Figure 3 shows results for the fully flexible case.



FIG. 4. Test of the equivalent-site approximation, Eq. (11), for stiff chains with $(K_{\rm B}, K_{\rm T})=(25, 1)$, at T=0.96. Top and bottom panels compare c(q) with, respectively, matrix elements $c_{aa}(q)$, and $c_{a5}(q)$. The insets enhance the region around the wave vector $q_{\rm max}$ for the maximum of the static structure factor S(q).

Data for the case $(K_{\rm B}, K_{\rm T}) = (25, 1)$ are displayed in Fig. 4. Both data sets correspond to the respective lowest investigated temperatures. We use a representation analogous to that of Ref. 34. Thus, top and bottom panels in both figures show the comparison of the averaged c(q) with, respectively, the matrix elements $c_{aa}(q)$ and $c_{a5}(q)$. The data of Fig. 3 are consistent with results of Ref. 34 for a similar fully flexible bead-spring model. Data in Fig. 4 constitute new results for the case of implemented intramolecular barriers. By looking at both figures we conclude that the quality of the equivalentsite approximation is not altered by the introduction of strong intramolecular barriers. Data in Fig. 4 display the same trends as in the fully flexible case. Thus, $c_{ab}(q) \approx c(q)$ is an excellent approximation except for correlations involving chain end monomers a=1 (and a=N by symmetry). The latter show deviations from c(q) which are moderate around $q_{\rm max}$, this q-range being the dominating one in the MCT kernel.

An additional approximation of the PRISM is the *ring* approximation (which is again exact for polymer rings). First we define the quantities $\tilde{S}_a(q) = \sum_{b=1}^N S_{ab}(q)$ and $\tilde{S}_a^{-1}(q) = \sum_{b=1}^N S_{ab}^{-1}(q)$. By exploiting the fact that for a ring polymer $\tilde{S}_a(q)$ is *a*-independent, i.e., $\tilde{S}_a(q) \approx N^{-1} \sum_{a=1}^N \tilde{S}_a(q)$, we find



FIG. 5. Test of the ring approximation, Eqs. (13) and (14). Top panel: fully flexible case at T=0.50. Bottom panel: barrier strength ($K_{\rm B}, K_{\rm T}$)=(25,1), at T=0.96. Main panels and insets compare S(q) (symbols) with, respectively, $\tilde{S}_a(q)$ and $1/\tilde{S}_a^{-1}(q)$, for the sites a=1, 2 and 5 (lines).

$$\widetilde{S}_a(q) \approx S(q).$$
 (13)

From the definition of $S_{ab}^{-1}(q)$ and $\tilde{S}_b(q)$ the relation $\Sigma_{b=1}^N S_{ab}^{-1}(q) \tilde{S}_b(q) = 1$ is exact. By introducing the ring approximation $\tilde{S}_b(q) \approx S(q)$ the former relation is transformed into:

$$\frac{1}{\tilde{S}_a^{-1}(q)} \approx S(q). \tag{14}$$

Figure 5 shows a test of the ring approximation of Eq. (13) (main panels) and Eq. (14) (insets). This is done both for fully flexible chains (top panel) and for stiff chains with $(K_{\rm B}, K_{\rm T}) = (25, 1)$ (bottom panel). The comparison between S(q), $\tilde{S}_a(q)$, and $1/\tilde{S}_a^{-1}(q)$ as computed from simulations is in general excellent, with the same quality for fully flexible and stiff chains. Only for the end monomers a=1 (and a=N by symmetry) significant differences between S(q) and $1/\tilde{S}_a^{-1}(q)$ are observed around the wave vector $q_{\rm max}$.

With all these results we conclude that the approximations assumed by the PRISM theory and introduced in the MCT equations for polymer melts (see below) are fulfilled, with the same quality for all the investigated range of barrier strength.

IV. DYNAMIC PROPERTIES

In this section we show some phenomenological dynamic features induced by the introduction of intramolecular barriers in our model. Panels in Fig. 6 show the T-dependence of the monomer MSD for fully flexible and

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FIG. 6. Temperature dependence of the monomer MSD for fully flexible (top) and stiff chains with $(K_{\rm B}, K_{\rm T})$ =(25,1) (bottom). The solid and dashed lines indicate, respectively, sublinear ($\sim t^{0.65}$) and linear behavior.

representative stiff chains with $(K_{\rm B}, K_{\rm T}) = (25, 1)$. We observe similar features in both cases, but also some differences. After the initial ballistic regime, a plateau extends over longer times with decreasing temperature. This plateau corresponds to the caging regime—i.e., the temporary trapping of each monomer in the shell of neighboring monomers around itwhich is usually observed when approaching a liquid-glass transition. At longer times, leaving the plateau, a crossover to a Rouse-like sublinear regime $\langle (\Delta r)^2 \rangle \propto t^{0.65}$ (Refs. 30 and 35) is observed for the fully flexible case. The final crossover to linear diffusion $\langle (\Delta r)^2 \rangle \propto t$ is reached at long times only for the highest investigated temperatures. However, for the case of stiff chains it is difficult to discriminate power law behavior over significant time windows. Apparently, the linear diffusive regime is not reached within the simulation time window.

Figure 7 shows the monomer MSD, for fixed values of density $\rho = 1.0$ and temperature T = 1.5, as a function of the barrier strength. Consistently with results in Ref. 31, we observe that increasing the strength of the internal barriers at fixed ρ and T leads to slower dynamics.



FIG. 8. Temperature dependence of f(q,t) for fully flexible (top) and stiff chains with $(K_{\rm B}, K_{\rm T}) = (25, 1)$ (bottom). The wave vector is $q_{\rm max} \approx 7$, in both cases corresponding to the maximum of the static structure factor S(q).

Figure 8 shows simulation results at several temperatures, both for fully flexible and stiff chains, for the normalized density-density correlator f(q,t). The latter is defined as $f(q,t) = \langle \rho(-\mathbf{q},0)\rho(\mathbf{q},t) \rangle / \langle \rho(-\mathbf{q},0)\rho(\mathbf{q},0) \rangle$. In both cases the correlator is evaluated at the maximum, $q_{\text{max}} \approx 7$, of the static structure factor S(q). As in the case of the MSD, both the fully flexible and stiff cases exhibit the standard behavior in the proximity of a glass transition.^{30,35} After the initial transient regime, f(q,t) shows a first decay to a plateau connected with the caging regime. On lowering the temperature this plateau extends over longer time intervals. At long times, a second decay is observed from the plateau to zero. This second decay corresponds to the structural α -relaxation.

Let us define the relaxation time as a time scale probing the α -structural relaxation. This can be done by introducing the time τ_x for which the correlator for q_{max} takes the value $f(q_{\text{max}}, \tau_x)=x$, provided x is small in comparison with the plateau height. Here we use x=0.2. Figure 9 shows $\tau_{0.2}$ as a function of T, for different values of the bending and torsional constants. As observed in the analysis of the MSDs, increasing the chain stiffness slows down the dynamics. At fixed



FIG. 7. Monomer MSD, for several values of the barrier strength, at fixed density $\rho = 1.0$ and temperature T = 1.5.



FIG. 9. Temperature dependence of the relaxation times $\tau_{0.2}$ for several values of the barrier strength.

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temperature, the relaxation time for the stiffest investigated chains increases up to three decades with respect to the fully flexible case.

In this section we have demonstrated a main dynamic feature: the slowing down of the dynamics, at fixed density and temperature, by progressively increasing the strength of the intramolecular barriers. This feature strongly suggests that intramolecular barriers constitute and additional mechanism for dynamic arrest, coexisting with the general packing effects induced by density and temperature. In the following we summarize the main predictions of the MCT and discuss simulation dynamic features within this theoretical framework.

V. MCT: SUMMARY

In this section we briefly summarize universal dynamic scaling laws concerning the MCT liquid-glass dynamics, and test them in the simulated polymer melt for all the investigated range of barrier strength. Extensive reviews on MCT can be found, e.g., in Refs. 14-19, 36, and 37. Though initially derived for simple hard-sphere systems, these predictions follow as consequences of the mathematical structure of the MCT equations. More specifically, they are associated to the bilinear dependence of the memory kernel on the density correlators (see below). Thus, MCT predicts the same dynamic scaling laws of the monoatomic case if such a mathematical structure is retained in systems of polyatomic molecules. This is indeed the case of the MCT for polymer melts developed by Chong and co-workers^{24,25} (see below). Therefore, the phenomenological analysis of our simulation results in terms of MCT dynamic scaling laws is justified within the theory.

By starting from the fundamental Liouville equation of motion and using the Mori–Zwanzig projection operator formalism one arrives to an integrodifferential equation for the normalized density-density correlator:

$$\ddot{f}(q,t) + \frac{q^2 k_B T}{mS(q)} f(q,t) + \frac{q^2 k_B T}{mS(q)} \int_0^t dt' m(q,t-t') \dot{f}(q,t') = 0.$$
(15)

This equation is obtained by using projectors over the subspace spanned by the densities and the longitudinal currents. The memory kernel $m(q,t-t') \propto \langle R_q^f(0) R_q^f(t-t') \rangle$, where the quantities R_q^f are, within the Mori–Zwanzig formalism, the associated fluctuating forces. Since the kernel cannot be exactly expressed in terms of f(q,t) and/or its time derivatives, Eq. (15) is not solvable. MCT introduces several approximations for the memory kernel, in order to provide a closed solvable form of Eq. (15). These approximations are:

(i) It is assumed that the long-time, slow dynamic regime of any observable coupled to density fluctuations can be expressed as a linear combination of "mode pairs," $\rho_k \rho_{q-k}$. Since the *exact* expression of the correlator of the fluctuating forces contains a slow contribution which is a linear combination of mode pairs (see, e.g., Ref. 17 for details), the former assumption is equivalent to neglecting the fast contribution of the fluctuating forces. In other words, it is equivalent to assuming a large separation between the time scales of the former contributions.

 (ii) Convolution approximation: three-point static correlations are approximated as products of static structure factors,

$$\langle \rho_{-\mathbf{q}}(0)\rho_{\mathbf{k}}(0)\rho_{\mathbf{q}-\mathbf{k}}(0)\rangle \approx nNS(q)S(k)S(|\mathbf{q}-\mathbf{k}|).$$
 (16)

(iii) Kawasaki approximation: dynamic four-point correlations are factorized in terms of products of dynamic two-point correlations (see, e.g., Ref. 17 for details).

By making use of these three approximations, the memory kernel m(q,t) becomes a bilinear form in f(q,t),

$$m(q,t) = \int \frac{d^3 \mathbf{k}}{(2\pi)^3} \mathcal{V}(\mathbf{q} - \mathbf{k}, \mathbf{k}) f(k, t) f(|\mathbf{q} - \mathbf{k}|, t), \qquad (17)$$

where the vertex $\mathcal{V}(\mathbf{q}-\mathbf{k},\mathbf{k})$ is given by:

$$\mathcal{V}(\mathbf{q} - \mathbf{k}, \mathbf{k}) = \frac{\rho}{2q^4} S(q) S(k) S(|\mathbf{q} - \mathbf{k}|)$$
$$\times [\mathbf{q} \cdot \mathbf{k} c(\mathbf{k}) + \mathbf{q} \cdot (\mathbf{q} - \mathbf{k}) c(|\mathbf{q} - \mathbf{k}|)]^2. \quad (18)$$

In a monoatomic fluid the direct correlation function c(q) is related to the static structure factor via the exact Ornstein– Zernike relation³⁸ $\rho c(q)=1-S^{-1}(q)$. With all this, Eq. (15) has been reduced to a closed set of coupled equations which can be solved self-consistently, provided S(q) and c(q) are known (the latter are *external* inputs in the MCT equations).

For the case of systems with molecular architecture, Chong and Hirata³⁹ have obtained, by using projectors over site-densities and site-currents, generalized MCT equations of motion for site-site correlators. The general mathematical structure of the kernel (bilinear in site-site correlators), and of the MCT equations of motion is retained. Except for very small values of N, numerical solution of the MCT equations for site-site correlators is extremely expensive, and further simplifications are needed in order to obtain a tractable set of equations. For the case of simple bead-spring chains, Chong and co-workers^{24,25} have reduced such equations to a scalar form for f(q,t). This is achieved by introducing the equivalent site, Eq. (11), and ring, Eqs. (13) and (14), approximations of the PRISM theory. The so-obtained scalar MCT equations of motion, memory kernel, and vertex for polymer chains are formally *identical* to Eqs. (15), (17), and (18). The polymer character of the system only enters implicitly through the PRISM relation $\rho c(q) = 1/\omega(q) - 1/S(q)$, which differs from the Ornstein-Zernike equation, $\rho c(q)$ $=1-S^{-1}(q)$, for monoatomic systems. With this, general MCT predictions which originate from the mathematical structure of Eqs. (15), (17), and (18) will be, due to the mentioned formal equivalence, analogous both for monoatomic systems and for polymer chains. Now we summarize such general predictions.

In MCT, nonergodic arrested states (glasses) are defined as those for which density correlators do not exhibit full relaxation. More specifically, if we introduce the nonergodicity parameters, defined as $f_q = \lim_{t\to\infty} f(q, t)$, MCT discriminates between fluid states $(f_q=0)$ and glassy states $(f_q>0)$. At the MCT critical temperature T_c , the nonergodicity parameters jump from zero to nonzero values.⁴⁰ In the following we use the notation f_q^c for referring to the critical nonergodicity parameters, i.e., the values of f_q at $T=T_c$.

By Laplace transform $(t \rightarrow z)$ of Eqs. (15) and (17) and taking the limit $z \rightarrow 0$, one finds a coupled set of equations for the nonergodicity parameters:

$$\frac{f_q}{1-f_q} = \mathcal{F}_q(\lbrace f \rbrace) = \int \frac{d^3 \mathbf{k}}{(2\pi)^3} \mathcal{V}(\mathbf{q} - \mathbf{k}, \mathbf{k}) f_{|\mathbf{q} - \mathbf{k}|} f_k,$$
(19)

where $\mathcal{F}_q(\{f\})$ denotes a functional, whose explicit expression is given in the right-hand side of the equation. Note that Eq. (19) always has the trivial solution $f_q=0$. Thus, glassy states take place when solutions $f_q>0$ also exist.

Given a tagged chain (labeled s), the density distribution for the *a*th monomer of the tagged chain is defined as $\rho_a^{s}(q) = \exp[i\mathbf{q}\cdot\mathbf{r}_{s}^{a}]$. The site-site intrachain correlator is defined as $F_{ab}^{s}(q,t) = \langle \rho_a^{s}(-\mathbf{q},0)\rho_b^{s}(\mathbf{q},t) \rangle$. Note that $F_{ab}^{s}(q,0) = \omega_{ab}(q)$. For the derivation of the MCT equations for $F_{ab}^{s}(q,t)$ we refer to Ref. 25. In this case the reduction to a scalar form is not possible. The corresponding nonergodicity parameters $f_{ab}^{s}(q) = \lim_{t\to\infty} F_{ab}^{s}(q,t)$ are obtained by solving the $N \times N$ -matrix equation⁴¹

$$f_{ab}^{s}(q) = \sum_{x,y=1}^{N} \mathcal{F}_{ax}^{s}(q) [\mathbf{I} + \mathcal{F}_{q}^{s}]_{xy}^{-1} \omega_{yb}(q), \qquad (20)$$

with **I** the identity matrix. The corresponding functional $\mathcal{F}^s_{ab}(q)$ is given by

$$\mathcal{F}_{ab}^{s}(q) = \sum_{x=1}^{N} \omega_{ax}(q) \int \frac{d^{3}\mathbf{k}}{(2\pi)^{3}} \mathcal{V}^{s}(\mathbf{q}-\mathbf{k},\mathbf{k}) f_{xb}^{s}(k) f_{|\mathbf{q}-\mathbf{k}|}, \quad (21)$$

with the vertex

$$\mathcal{V}^{s}(\mathbf{q}-\mathbf{k},\mathbf{k}) = \frac{\rho}{q^{4}} S(|\mathbf{q}-\mathbf{k}|) [\mathbf{q} \cdot (\mathbf{q}-\mathbf{k})]^{2} c^{2} (|\mathbf{q}-\mathbf{k}|). \quad (22)$$

The normalized self-correlator, usually introduced as $f^{s}(q,t) = (nN)^{-1} \sum_{j=1}^{n} \sum_{a=1}^{N} \langle \exp[i\mathbf{q} \cdot (\mathbf{r}_{j}^{a}(t) - \mathbf{r}_{a}^{a}(0))] \rangle$, can be equivalently obtained as $f^{s}(q,t) = N^{-1} \sum_{a=1}^{N} F_{aa}^{s}(q,t)$. Likewise, the corresponding nonergodicity parameters, defined as the long-time limit of $f^{s}(q,t)$, can be obtained as $f_{q}^{s} = N^{-1} \sum_{a=1}^{N} f_{aa}^{s}(q)$. Thus, the solution of Eq. (20) also provides trivially the nonergodicity parameters for the self-correlator.

The separation parameter, $\epsilon = (T - T_c)/T_c$, is introduced to quantify the relative distance to the critical temperature T_c . We are interested in the behavior of f(q,t) in the ergodic fluid by approaching T_c from above. Thus we express the long-time behavior of the density-density correlators as:

$$f(q,t) = f_{q}^{c} + g_{q}(t),$$
(23)

where $g_q(t)$ quantifies (small) deviations around f_q^c for $|\epsilon| \rightarrow 0$. By introducing Eq. (23) in Eqs. (15) and (17), expanding the functional \mathcal{F}_q of Eq. (19) in a power series of $|\epsilon|$, comparing the so-obtained resulting expressions and retaining the lower-order terms (see, e.g., Ref. 36 for a detailed exposition), one finds that $g_q(t)=h_qG(t)$, where h_q only depends on q, and G(t) is a q-independent term which contains

the full time dependence of the deviations of f(q,t) around f_q^c . Thus, we rewrite Eq. (23) as:

$$f(q,t) = f_q^c + h_q G(t).$$
 (24)

This expression is known as the *first universality* of the MCT or *factorization theorem*. It predicts a scaling function G(t) (known as the β -correlator) that is *common* for all the density correlators (since it is *q*-independent). Following the procedure mentioned in the previous paragraph,³⁶ the function G(t) is found to obey the equation:

$$\sigma - z^2 \tilde{G}^2(z) = \lambda z L[G^2(t)], \qquad (25)$$

where $\tilde{G}(z)$ and $L[G^2(t)]$ are the Laplace transform of, respectively, G(t) and $G^2(t)$. In this equation $\sigma = c|\epsilon|$, with *c* a constant (see Ref. 36 for its explicit expression), and λ is another constant given by

$$\lambda = \sum_{qk} e_q^{\mathrm{T}} C^{\mathrm{c}}(q, k, |\mathbf{q} - \mathbf{k}|) e_k e_{|\mathbf{q} - \mathbf{k}|}.$$
(26)

The quantities e_q and $e_q^{\rm T}$ are, respectively, the eigenvectors of the so-called stability matrix $\mathbf{C}^{\rm c}$ (see below) and its traspose, with the normalization conditions $\Sigma_q e_q^{\rm T} e_q = 1$ and $\Sigma_q e_q^{\rm T} (1 - f_q^{\rm c}) e_q^2 = 1$. The elements of the stability matrix are given by

$$C^{\rm c}(q,k) = (1 - f_k^{\rm c})^2 \left(\frac{\partial \mathcal{F}_q}{\partial f_k}\right)_{\{f=f^{\rm c}\}}.$$
(27)

The terms $C^{c}(q,k,|\mathbf{q}-\mathbf{k}|)$ in Eq. (26) are given by:

$$C^{c}(q,k,|\mathbf{q}-\mathbf{k}|) = \frac{1}{2}(1-f_{k}^{c})^{2}(1-f_{|\mathbf{q}-\mathbf{k}|}^{c})^{2} \left(\frac{\partial^{2}\mathcal{F}_{q}}{\partial f_{k} \partial f_{|\mathbf{q}-\mathbf{k}|}}\right)_{\{f=f^{c}\}}.$$
(28)

Equation (25) for the β -correlator does not have an analytical solution. Still, asymptotic expressions can be obtained for different time windows. With this idea in mind the β -time scale is first defined as

$$\tau_{\beta} = t_0 |\sigma|^{-1/(2a)} \tag{29}$$

with t_0 a microscopic time scale and a an exponent. The β -correlator is then rewritten as $G(t) = |\sigma|^{1/2} g_{\sigma}(t/\tau_{\beta})$. By introducing this expression in Eq. (25) and taking the limits $t \ll \tau_{\beta}$ and $t \gg \tau_{\beta}$ one finds¹⁶ the asymptotic solutions $g_{\sigma}(t/\tau_{\beta}) = (t/\tau_{\beta})^{-a}$ for $t \ll \tau_{\beta}$, and $g_{\sigma}(t/\tau_{\beta}) = -B(t/\tau_{\beta})^{b}$ for $t \gg \tau_{\beta}$, where *B* is a constant.³⁶ The exponents *a* and *b* follow the constraint

$$\lambda = \frac{\Gamma^2(1-a)}{\Gamma(1-2a)} = \frac{\Gamma^2(1+b)}{\Gamma(1+2b)},$$
(30)

where Γ denotes the Euler's Gamma function. According to the former equations for $g_{\sigma}(t/\tau_{\beta})$, one finds for Eq. (24) the asymptotic expressions $f(q,t)=f_q^c+h_q(t/t_0)^{-a}$ for $t \ll \tau_{\beta}$ and $f(q,t)=f_q^c-h_q(t/\tau_{\alpha})^b$ for $t \gg \tau_{\beta}$. Inclusion of higher-order corrections to the latter equations yield:³⁶

$$f(q,t) = f_q^{c} + h_q(t/t_0)^{-a} + \tilde{h}_q^{(2)}(t/t_0)^{-2a} \quad t \ll \tau_\beta,$$
(31)

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$$f(q,t) = f_q^{\rm c} - h_q (t/\tau_{\alpha})^b + h_q^{(2)} (t/\tau_{\alpha})^{2b} \quad t \ge \tau_{\beta}.$$
 (32)

As h_q , the prefactors $\tilde{h}_q^{(2)}$ and $h_q^{(2)}$ only depend on q (see also below). Equation (32) is also known as the von Schweidler expansion and is often used in the analysis of the long-time decay (see below). In the former equations τ_{α} is the α -time scale, defined as:

$$\tau_{\alpha} = B^{-1/b} t_0 |\sigma|^{-\gamma}. \tag{33}$$

The exponent γ follows the constraint:

$$\gamma = \frac{1}{2a} + \frac{1}{2b}.\tag{34}$$

Another important prediction of the MCT for states approaching T_c from above, is the *second universality* or *TTSP*. This prediction arises as a long-time scaling property of the MCT equations of motion.¹⁶ According to the TTSP, the long-time decay of any correlator f(q,t) (i.e., the final part of the α -relaxation) is invariant under scaling by the α -relaxation time τ_{α} . In other words, for two temperatures T_1 and T_2 above T_c one finds

$$f(q,t/\tau_{\alpha}(T_1);T_1) = f(q,t/\tau_{\alpha}(T_2);T_2) = \bar{f}(q,\hat{t}),$$
(35)

where $\tilde{f}(q, \hat{t})$ is a *T*-independent master function of the normalized time \hat{t} . While G(t) is common to all correlators, the master function $\tilde{f}(q, \hat{t})$ associated to the TTSP is different for each correlator f(q, t). The superposition principle implies that the estimated α -relaxation time, defined in this work as the time τ_x where $f(q_{\text{max}}, t)$ takes a value *x* well below the plateau, is proportional to τ_{α} . Thus, it also follows the asymptotic power law

$$\tau_x(T) \propto (T - T_c)^{-\gamma}.$$
(36)

The α -decay from the plateau to zero is often well described by an empirical Kohlrausch–Williams–Watts (KWW) function,

$$f(q,t) = A_q \exp\left[-\left(t/\tau_q^K\right)^{\beta_q}\right],\tag{37}$$

with A_q , $\beta_q < 1$. Note that the latter does not come out as an analytical solution of the MCT equations. However in the limit $q \rightarrow \infty$ of the KWW time τ_a^K , MCT predicts that⁴²

$$\tau_q^K \propto q^{-1/b} \quad q \to \infty, \tag{38}$$

where b is the von Schweidler exponent introduced above.

The set of equations exposed in this section constitute a series of universal results which originate from the structure of the MCT equations of motion, Eqs. (15), (17), and (18). As mentioned above, the latter were initially derived for simple hard-sphere systems, but the corresponding ones for polymer melts become formally identical following the derivation by Chong and co-workers.^{24,25} With this, the scaling laws exposed in this section will also hold in the MCT for polymer melts. Thus, the phenomenological analysis of our simulation data in terms of such scaling laws is justified within the framework of MCT. This analysis is presented in the next section.



FIG. 10. Test of the factorization theorem, Eq. (39), for density-density (top) and self-correlators (bottom) at T=0.80 and $(K_{\rm B}, K_{\rm T})=(15, 0.5)$. The different curves correspond to equispaced wave vectors in the range 3.8 $\leq q \leq 16.35$. The fixed times (indicated by filled squares) are t'=300 and t''=3.

VI. MCT ANALYSIS OF SIMULATIONS

In order to test the factorization theorem, Eq. (24), we compute the ratio:

$$R_q(t) = \frac{f(q,t) - f(q,t')}{f(q,t') - f(q,t')} = \frac{G(t) - G(t')}{G(t') - G(t')}$$
(39)

where t' and t'' are arbitrary times in the β -regime. The ratio for the self-correlators, $R_a^{s}(t)$, is defined analogously. If the factorization theorem, and then also the right-hand side of Eq. (39), is fulfilled, the ratios $R_q(t)$ and $R_a^s(t)$ do not depend on the specific correlator. Figure 10 shows $R_q(t)$ and $R_q^s(t)$ over a broad range of wave vectors $3.8 \le q \le 16.5$. The data correspond to barrier strength $(K_{\rm B}, K_{\rm T}) = (15, 0.5)$ at T = 0.80. The fixed times t''=3 and t'=300 roughly correspond to the beginning and the end of the plateau regime. There is an intermediate time window of about two decades where the data for density-density and self-correlators collapse onto a q-independent master curve, while they split at both short and late times. Figure 11 demonstrates that the master curve is, moreover, the same for both density-density and selfcorrelators. Thus, Figs. 10 and 11 demonstrate the validity of the MCT first universality.

The inset in Fig. 11 shows a test of MCT predictions for the deviations from the β -scaling behavior. This is done following the procedure proposed by Gleim and Kob in Ref. 43. Thus, we take a large set of *M* correlators. In the present case we take density-density and self-correlators over the former *q*-range, up to a total of M=32. Then we draw two vertical lines (see Fig. 11) at a time t'_d a bit longer than t' and at a time t''_d a bit shorter than t''. The correlators are labeled as i''=1,2,...M according to the order they intersect, from top to bottom, the line at t''_d . Then a second label i'=1,2,...M is assigned to each i''-curve according to the orden it intersects, again from top to bottom, the line at t'_d . This procedure pro-



FIG. 11. Main panel: common representation of $R_q(t)$ (empty symbols) and $R_q^s(t)$ (lines), for selected wave vectors (common colors correspond to common *q*-values). As in Fig. 10, data correspond to T=0.80 and ($K_{\rm B}, K_{\rm T}$) = (15,0.5), and the selected fixed times (indicated by filled squares) are t' = 300 and t''=3. Inset: following Ref. 43, test for the deviations of the β -scaling behavior. The vertical dashed lines in the main panel indicate the selected times t'_d and t''_d (see text for explanation).

vides a function i'(i''). The prefactors of the leading corrections to the β -scaling in Eqs. (31) and (32) are given by³⁶ $\tilde{h}_q^{(2)} = h_q(K_q + \kappa)$ and $h_q^{(2)} = h_q K_q$, where κ is a constant and K_q only depends on q. According to this, MCT predicts that the former function i'(i'') will be just i' = i''. Data in the inset of Fig. 11 show some scattering, which is comparable to that in Fig. 5 of Ref. 43 for an analogous test in a Lennard-Jones system. Still, data are in reasonable agreement with MCT predictions, since they clearly follow the trend i' = i'' (straight line in the inset).

Figure 12 shows a test of the TTSP, Eq. (35), for the density-density correlator evaluated at q_{max} [maximun of the static structure factor S(q)]. The data correspond to the case $(K_{\text{B}}, K_{\text{T}}) = (15, 0.5)$ and cover a broad temperature range $0.80 \le T \le 1.5$. Data collapse onto a master curve after rescaling the absolute time by the relaxation time $\tau_{0.2}$. Thus, the MCT second universality also holds for chains with strong intramolecular barriers.

Solving numerically the MCT equations and determining the dynamic exponents (a, b, γ, λ) is in general a difficult task. When numerical solutions are not available, nonergodicity parameters, prefactors and exponents in Eqs. (31), (32), (36), and (38) can be obtained as fit parameters from simulation or experimental data (see, e.g., Refs. 15, 35, and 44–



FIG. 12. Test of the TTSP, Eq. (35), for the density-density correlators at $q_{\text{max}} \approx 7$, for the case ($K_{\text{B}}, K_{\text{T}}$)=(15,0.5).



FIG. 13. Symbols: simulations results for density correlators. Top panel: f(q,t) for $(K_{\rm B},K_{\rm T})=(15,0.5)$, at T=0.80. Bottom panel: $f^{\rm S}(q,t)$ for $(K_{\rm B},K_{\rm T})=(35,4)$, at T=1.33. Identical symbols in both panels correspond to identical wave vectors q [values are given in panel (a)]. Lines are fits to the von Schweidler expansion, Eq. (32) (up to second-order terms), with b = 0.50 (top) and 0.37 (bottom).

46). Consistency of the analysis requires that dynamic correlators and relaxation times are described by a common set of exponents, all of them related to a single λ -parameter through Eqs. (30) and (34).

We have performed this consistency test for all the investigated range of barrier strength. The following figures in this section illustrate, for some representative cases, the analysis of simulation data in terms of MCT asymptotic laws. Figure 13 shows for a broad *q*-range $(2.0 \le q \le 14.4)$, fits to the von Schweidler expansion, Eq. (32) (up to second-order terms). Data correspond to density-density correlators f(q,t) for the state point $(K_B, K_T) = (15, 0.5)$, T = 0.80 (labeled S1), and to self-correlators $f^s(q,t)$ for $(K_B, K_T) = (35, 4)$, T = 1.33 (labeled S2). A good description of the simulation data is achieved, for all the range of *q*-values and over almost four time decades, with a fixed *b*-exponent (*b*=0.50 and 0.37 for, respectively, S1 and S2).

Figure 14 displays, for the former values of the barrier strength, the *q*-dependence of the so-obtained critical nonergodicity parameters $[f_q^c \text{ for } f(q,t)]$ and $f_q^{\text{sc}} \text{ for } f^{\text{s}}(q,t)]$. For comparison, we also include the fully flexible case $(K_{\text{B}}, K_{\text{T}}) = (0, 0)$. As deduced from the stronger decay of f_q^c and f_q^{sc} for stronger barriers, the introduction of chain stiffness yields a weaker stability of density fluctuations. It also induces a weaker localization for self-motions at fixed density. Thus, by making an approximate fit of f_q^{sc} to Gaussian behavior, $f_q^{\text{sc}} \approx \exp(-q^2 l_c^2/6)$, we estimate, at fixed $\rho=1.0$, a localization length $l_c=0.19$, 0.21, and 0.23 for, respectively, $(K_{\text{B}}, K_{\text{T}}) = (0,0)$, (15,0.5), and (35,4).

Data of self-correlators from the plateau to the limit of the simulation window have been fitted to KWW functions, Eq. (37) (not shown). Figure 15 shows the q-dependence of the so-obtained KWW relaxation times τ_q^{K} for the former values of the barrier strength, at their respective lowest in-

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FIG. 14. Critical nonergodicity parameters, as determined from fits to Eq. (32), for different barrier strength. Top and bottom panels show data for, respectively, f_a^c and f_a^{sc} .

vestigated temperatures. The lines represent tests of the MCT prediction $\tau_q^{K} \propto q^{-1/b}$ for large q. A good description of the data is obtained with the same *b*-exponents used for the independently obtained von Schweidler fits of Fig. 13.

Figure 16 shows, for the same values of (K_B, K_T) in Fig. 15, a test of the power law $\tau_{0.2} \propto (T-T_c)^{-\gamma}$ for the temperature dependence of the estimated α -relaxation times. The fit covers about three time decades. By representing the data in terms of the separation parameter T/T_c-1 , clearly different γ -exponents are evidenced for different barrier strength. A good description of the data is obtained with the γ -values derived, through Eqs. (30) and (34), from the *b*-values used in Figs. 13 and 15. This result demonstrates the consistency of the MCT analysis for the representative examples showed here, which cover all the range of investigated barrier strength between fully flexible and stiff chains.

Similar consistent tests (not shown) have been performed for the rest of investigated systems. Table I displays



FIG. 15. Symbols: *q*-dependence of KWW relaxation times for different barrier strength. Lines are fits to $\propto q^{-1/b}$ (see text). From top to bottom *b* = 0.54, 0.50 and 0.37.



FIG. 16. Symbols: *T*-dependence of relaxation times $\tau_{0.2}$ for different barrier strength. Lines are fits to $\propto (T-T_c)^{-\gamma}$ (see text). From top to bottom $\gamma = 2.60, 2.74$ and 3.43.

the results for the so-obtained λ -exponents and critical temperatures T_c as a function of (K_B, K_T) . We also include the corresponding value of the mean end-to-end radius (computed at $T_{\rm c}$), which provides a qualitative characterization of chain stiffness. From the numerical values in Table I a clear correlation between the strength of the internal barriers and the values of T_c and λ is unambiguously demonstrated. The interplay between packing effects and intramolecular barriers induces a progressive increase in T_c at fixed density. A similar effect is observed for the λ -exponent, which increases from $\lambda = 0.761$ for fully flexible chains to $\lambda = 0.885$ for the stiffest investigated chains. The smallest λ -values in Table I are typical of simple glass-formers as the archetype hardsphere fluid (λ =0.74) (Ref. 36) where dynamic arrest is driven by packing effects. The largest ones, $\lambda \leq 0.9$, are similar to those observed in realistic models of polymer melts which incorporate the chemical structure of the chains. Some examples include poly(vinyl methylether),⁴⁷ polybutadiene,⁴⁸ or poly(vinyl ethylene),⁴⁹ with respective values of $\lambda = 0.87$, 0.93. and 0.93.

Thus, the analysis presented here rationalizes the difference in the MCT exponents between fully flexible beadspring models and real polymers. The systematic study performed by tuning the barrier strength suggests that large λ -exponents in real polymers arise from the interplay between two distinct mechanisms for dynamic arrest. These are general packing effects and polymer-specific intramolecular barriers. Large λ -values arising from the interplay between distinct arrest mechanisms have been observed in systems of very different nature. as short-ranged attractive colloids^{22,50,51} (competition between hard-sphere repulsion and short-ranged reversible bonding), polymer blends^{52,53} and colloidal mixtures with strong dynamic asymmetry^{54,55} (bulklike caging and matrix-induced confinement), or densified silica⁵⁶ (presumably bonding and packing). Dynamic features revealed by recent simulations on a globular native protein are also consistent with large λ -values.⁵⁷ Numerical solutions of the MCT equations in short-ranged attractive colloids^{22,50} and quenched-annealed mixtures²³ have revealed the existence of higher-order MCT transitions, which are characterized by the upper limit $\lambda = 1$. Whether higher-

his article is copyrighted as indicated in the article. Reuse of AIP content is subject to the terms at: http://scitation.aip.org/termsconditions. Downloaded to IF 161.116.80.50 On: Mon. 03 Feb 2014 14:12:29 order MCT transitions are present at some region of the control parameter space of the investigated model is an open question.

In this section we have performed a phenomenological analysis of the simulation data within the framework of MCT. In the next section the observed trends are compared with numerical solutions of the MCT equations.

VII. SOLUTION OF THE MCT EQUATIONS

We have solved Eqs. (19) and (20) for the nonergodicity parameters, for all the investigated range of barrier strengths. In analogy with the procedure exposed in, e.g., Refs. 36 and 37, the integrals over the reciprocal space in the corresponding MCT functionals of Eqs. (19) and (21) are discretised to a grid of M=600 equispaced points, with *q*-spacing Δq =0.1, leading to the expressions:

$$\frac{f_q}{1-f_q} = \frac{\rho(\Delta q)^3}{32\pi^2} \sum_{x_k} \sum_{x_p}' \frac{x_k x_p}{x_q^5} S(q) S(k) S(p) \\ \times [(x_q^2 + x_k^2 - x_p^2)c(k) + (x_q^2 + x_p^2 - x_k^2)c(p)]^2 f_k f_p$$
(40)

and

$$\mathcal{F}_{ab}^{s}(q) = \frac{\rho(\Delta q)^{3}}{16\pi^{2}} \sum_{x=1}^{N} \omega_{ax}(q) \sum_{x_{k}} \sum_{x_{p}}' \frac{x_{k}x_{p}}{x_{q}^{5}} S(p) \\ \times [(x_{q}^{2} + x_{p}^{2} - x_{k}^{2})c(p)]^{2} f_{xb}^{s}(k) f_{p}.$$
(41)

In these expressions the wave vectors are defined as $q = x_q \Delta q$, $k = x_k \Delta q$, and $p = x_p \Delta q$, with $x_q, x_k, x_p = 1/2, 3/2, \dots 1199/2$. The prime at the sums over x_p means that the latter are restricted to x_p -values following the condition $|x_q - x_k| + 1/2 \le x_p \le x_q + x_k - 1/2$.

The solutions of Eq. (40) are found by a standard iterative procedure $f_q^{j+1}/[1-f_q^{j+1}] = \mathcal{F}_q(\{f^j\})$, with *j* the iteration step, and with the initial condition $f_q^0 = 1$. It can be demonstrated that the stability matrix in Eq. (27) has always a maximum nondegenerate eigenvalue $E \le 1$, which takes the upper value $E_c = 1$ at the critical point.³⁶ Thus, by following the drift of *E* with changing temperature it is possible to bracket the values of the critical nonergodicity parameters f_q^c , and the critical temperature T_c , with very high precision. Once the values of f_q^c are obtained, they are fixed in the functional of Eq. (41), and a small number of iterations is needed to find the corresponding critical values $f_{ab}^{sc}(q)$. Finally, the critical nonergodicity parameters for selfcorrelations are obtained as $f_q^{sc} = N^{-1} \sum_{a=1}^{N} f_{aa}^{sc}(q)$.

Following the procedure exposed above, we solved Eq. (40) by inserting as external inputs the structural quantities, S(q) and c(q), as directly computed from the simulations. However, as previously reported in Ref. 25 for fully flexible chains, a MCT transition was not observed for any of the investigated barrier strength. This means that the theoretical critical temperature T_c is below the lowest simulation temperature for which equilibration was possible. This result is different from the usual observation in nonpolymeric systems, for which the theoretical critical point is accessible in simulation time scales. The reason of this difference is, in



FIG. 17. Comparison of the critical nonergodicity parameters f_q^c (top) and f_q^{sc} (bottom) obtained from MCT calculations (solid lines) with those obtained from the phenomenological analysis of the simulation data (symbols, dashed lines are guides for the eyes). Results are shown both for fully flexible chains and for representative stiff chains with (K_B, K_T)=(15,0.5).

some way, related with the unability to crystallize of beadspring models, which avoids a fast growing of peaks under cooling in the static structure factor S(q), leading to MCT kernels which are not sufficiently strong to provide nonzero solutions of f_q .

Since static correlations computed from our equilibrium simulations do not induce a MCT transition, we are forced to use a structural theory for estimating S(q) and c(q) at lower temperatures, which will allow us to insert them in the MCT equations and to search for the critical temperature. Thus, we solve numerically the PRISM equation

$$\rho c(q) = 1/\omega(q) - 1/S(q), \qquad (42)$$

with the Percus–Yevick (PY) closure relation³⁸ for the nonbonded potential V(r) of Eq. (1). The PY relation is given by:

$$c(r) = [1 - \exp(V(r)/k_B T)](h(r) + 1),$$
(43)

where c(r) and h(r) are the Fourier transforms in the real space of c(q) and h(q). The coupled set of nonlinear Eqs. (42) and (43) is solved by a standard Picard iteration method⁵⁸ for the quantity $\Gamma(r)=h(r)-c(r)$, which is a smooth function over all the range of r. The form factor $\omega(q)$ is an *external* input in this procedure. We observed (see above) that $\omega(q)$ exhibits a very weak temperature dependence in comparison to the total static structure factor S(q). Thus we just use for each barrier strength the $\omega(q)$, as computed from the simulations, at the lowest temperature for which equilibration was possible.

In Fig. 17 we show a comparison of the critical nonergodicity parameters f_q^c (top panel) and f_q^{sc} (bottom panel) as obtained from numerical solution of the MCT equations, with the results of the fitting procedure of simulation data



FIG. 18. Critical temperature T_c as a function of the end-to-end radius R_{ce}^c . The theoretical T_c^{MCT} , as obtained by solving the MCT equations, is compared with the T_c^{MD} from the phenomenological analysis of the simulation data.

(see above). The theoretical results qualitatively reproduce the simulation trends, and in particular the observation that at fixed density the intramolecular barriers induce a weaker localization length. Quantitatively, the MCT solutions overstimate the amplitude of the nonergodicity parameters, except in the low-q region of f_q^c , for which MCT clearly understimates the results.

In Fig. 18 we show a representation of the critical temperature T_c as a function of the end-to-end radius R_{ee}^c , which quantifies chain stiffness. Values of T_c obtained from the phenomenological analysis of the simulations (T_c^{MD}) and from the numerical solutions of the MCT equations (T_c^{MCT}) are compared. We note that T_c^{MD} seems to grow monotonously with chain stiffness. This trend is well reproduced by the theory for low and moderate values of the internal barriers. Thus, for values of bending and torsional constants $K_B < 15$ and $K_T < 0.5$, the dependence of T_c^{MCT} on R_{ee}^c roughly displays the same slope as for T_c^{MD} , with a shift factor $T_c^{MCT}/T_c^{MCT} \approx 1.25$. Similar shifts between simulation and theory, which have their origin in the mean-field character of the MCT, are observed in other systems. 44,59,60 The range of barrier strength for which T_c^{MCT} and T_c^{MD} are roughly parallel is significant. Note that for $(K_B, K_T) = (8, 0.2)$ the end-to-end radius R_{ee}^c is a factor 1.3 longer than for fully flexible chains.

By further increasing chain stiffness the differences between T_c^{MD} and T_c^{MCT} progressively increase. We observe a saturation of the theoretical T_c^{MCT} around ≈ 0.55 , while the simulation T_c^{MD} grows up to a value of 1.23 for the stiffest investigated chains. Thus, the agreement between theory and simulation clearly breaks for stiff chains.

Finally, we have computed the corresponding theoretical λ -exponents according to the definitions of Eqs. (26)–(28). We find an almost constant value of $\lambda \approx 0.72$ for all the investigated range of barrier strength. This result is clearly different from the observations in the phenomenological

MCT analysis of simulation data (see data in Table I), which provides a strong dependence of λ on the barrier strength.

VIII. DISCUSSION

The results reported in the previous section show that, though reproducing some qualitative simulation trends for low and moderate barriers, numerical solutions of the MCT equations exhibit important differences with simulation values as the limit of stiff chains is approached. Another result to be understood is the clear disagreement between the almost constant value of the theoretical λ -exponent, and the observed strong dependence of simulation values on the barrier strength.

The observed disagreement between theory and simulation for strong barriers does not seem to be related with the failure of the PRISM approximations for stiff chains, which have been introduced in the derivation of the MCT equations. Indeed we have shown that the quality of the used PRISM approximations is the same for fully flexible and stiff chains (Figs. 3–5). Having said this, it might be argued that the theory is simply wrong: the phenomenological MCT analysis is apparently successful, but one finds that it has little to do with the theory, for stiff chains, when solving the MCT equations. However, we remind that the phenomenological analysis has shown, for all the investigated range of barrier strength: (i) the validity of the two MCT universalities, i.e., the factorization theorem (Figs. 10 and 11) and the TTSP (Fig. 12) and (ii) the possibility of a good description of different dynamic observables (Figs. 13, 15, and 16) with a set of dynamic exponents which are consistently transformed, through Eqs. (30) and (34), to a single λ -exponent.

We believe that all these observations, for all the investigated cases, are not fortuitous. At this point it must be noted that the predictions referred to in points (i) and (ii) arise, within MCT, as a consequence of the mathematical structure of the equations of motion, more precisely they originate from the bilinear form of the memory kernel. The specific values of the numerical solutions clearly depend on the coefficients of the bilinear products (which enter through the vertices of the kernel), but the factorization theorem, the TTSP, and the asymptotic scaling laws are universal properties provided the kernel is bilinear. Thus, the results of the phenomenological analysis suggest that the underlying physics may be connected to a bilinear memory kernel, though for high barriers the actual coefficients strongly differ from those introduced by MCT through the vertices, thus leading to theoretical results which strongly differ from simulations.

In other words, the present results suggest that there may be relevant static contributions for the case of stiff chains which are missing in the MCT vertices. Thus, the inclusion of such contributions will increase the strength of the kernel and will induce the theoretical transition at higher values of T_c , which might improve the comparison between T_c^{MCT} and T_c^{MD} of Fig. 18. Recalling the three main approximations of MCT, we suggest that the convolution approximation, Eq. (16), might break for stiff chains. Though possibly it is not the case for intermolecular contributions, its breakdown for *intramolecular* contributions in stiff chains is plausible. It is

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known that the convolution approximation fails when static correlations show a strong directionality at near-neighbor distances, as for, e.g., network-forming liquids as silica.⁶¹ This directionality is clearly enhanced for intrachain correlations by increasing the barrier strength, as evidenced by the progressively larger values of the end-to-end radius (see Table I). For the case of silica, it has been shown that the explicit inclusion of three-point static correlations in the MCT vertex improves significantly the quality of the comparison between theory and simulations.⁶¹ A similar improvement might be achieved in the present case by similarly incorporating the intrachain three-point static contributions. Work in this direction is in progress.

IX. CONCLUSIONS

We have performed simulations on a simple bead-spring model for polymer melts with intramolecular barriers. The role of such barriers on the glass transition has been investigated by systematically tuning the barrier strength. Dynamic correlators probing the structural relaxation have been analyzed in the framework of the MCT. We have obtained critical nonergodicity parameters, critical temperatures, and dynamic exponents of the theory from consistent fits of simulation data to MCT asymptotic laws. From the analysis of the critical nonergodicity parameters we deduce that the presence of the barriers induces a weaker localization length in the system at fixed density. The increase in the barrier strength at fixed density also induces a higher critical temperature $T_{\rm c}$. The values of the dynamic exponents, as obtained from the phenomenological analysis of the simulation data, exhibit significant differences between the limit of fully flexible and stiff chains. In particular the so-called λ -exponent takes standard values $\lambda \sim 0.7$ for the fully flexible case and values approaching the upper limit $\lambda = 1$ for strong intramolecular barriers. While the former λ -values are characteristic of simple systems dominated by packing effects, transitions with $\lambda \approx 1$ arise in systems with different competing mechanisms for dynamic arrest. In our systems these large λ -values suggest a competition between two distinct mechanisms: general packing effects and polymerspecific intramolecular barriers.

For a comparison between simulation and theory, we have numerically solved the MCT equations, following a recent extension of the MCT by Chong and co-workers^{24,25} for polymer melts. The approximations assumed by the structural PRISM theory, which are introduced in the MCT equations, are fulfilled for all the investigated values of the barrier strength. We have compared the critical nonergodicity parameters and critical temperatures $T_{\rm c}$, as obtained by solving the MCT equations, with the corresponding values from the phenomenological analysis of the simulation data. The theoretical calculations qualitatively reproduce the trends observed in the simulations for low and moderate barriers. However strong discrepancies are observed as the limit of high barriers is approached. The reason for such a disagreement possibly lies in the nature of the approximations made in the derivation of the MCT equations. In particular, the convolution approximation for three-point static correlations

might be unadequate for stiff chains. We suggest that a reformulation of MCT equations for polymer melts, explicitly including intrachain three-point static correlations, might lead to a better agreement between simulations and theory. Work in this direction is in progress.

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