Johari-Goldstein heterogeneous dynamics in a model polymer

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Abstract

The heterogeneous character of the Johari-Goldstein (JG) relaxation is evidenced by molecular-dynamics simulation of a model polymer system. A double-peaked evolution of dynamic heterogeneity (DH), with maxima located at JG and structural relaxation time scales, is observed and mechanistically explained. The short-time single-particle displacement during JG relaxation weakly correlates with the long-time one observed during structural relaxation.

1 Introduction

Avoiding crystallization, polymers and liquids freeze into a microscopically disordered solid-like state, a glass. On approaching the glass transition, molecular rearrangements occur via
both the primary mode, referred to as structural or $\alpha$ relaxation, and the faster secondary ($\beta$) processes as evidenced by mechanical, electrical, and thermal properties of materials.\[^{2-4}\]

There is wide interest in the $\beta$ relaxation, as attested by the large number of experiments, as well as phenomenological and theoretical studies, and simulations.\[^{4-10}\]

Here, we focus on linear polymers where there are no side groups and the secondary relaxations are usually ascribed to some movement of short lengths of the main chain, like to limited vibrational oscillation about their mean position or crankshaft motion.\[^{2,11-13}\]

The secondary relaxation in linear polymers is thought to be a genuine manifestation of the Johari Goldstein (JG) $\beta$-relaxation,\[^{5}\] a special class of secondary relaxations having strong connections to the $\alpha$-relaxation in properties, and advocated to be a universal feature of the glass transition.\[^{4,8-11}\]

It was early noted that the molecular rearrangements giving rise to JG relaxation process are similar to those that are responsible for the glass transition itself.\[^{15}\] In particular, correlation between JG and primary relaxation were reported in both molecular liquids and polymers.\[^{13,16-18}\]

It was argued that the JG relaxation, like the alpha, involve transitions between metabasins in the energy landscape\[^{19,20}\], whereas other approaches suggest that JG processes can be interpreted in structural glasses as transitions between sub-basins belonging to a same metabasin.\[^{21}\]

Furthermore, a number of experimental results indicate that the JG relaxation is sensitive to the thermodynamic variables underlying the glass transition,\[^{41}\] mimicking the $\alpha$ relaxation, being strongly pressure dependent and showing the invariance of the ratio $\tau_\alpha/\tau_\beta$ to variations of pressure and temperature, keeping $\tau_\alpha$ constant.\[^{8}\] This led to the conclusion that the JG relaxation is precursor to structural relaxation and viscous flow, having a slower dynamics due to cooperativity involving many body dynamics.\[^{6,8,22,23}\]

This has been recently substantiated by studying the invariance of the relation between $\alpha$ relaxation and $\beta$ relaxation in metallic glasses to variations of pressure and temperature by molecular dynamics simulations combined with the dynamic mechanical spectroscopy method.\[^{24}\] Further support
to the view that primary and JG relaxation are closely related also come from the evidence that the JG relaxation shows both a very broad distribution of relaxation times, and cooperative dynamics, indicated by simulations in non-polymeric liquids, experiments. The existence of spatial correlations between dynamic fluctuations - in short, dynamic heterogeneity (DH) - has been revealed by experiments and numerical studies, e.g., see the reviews in. In particular, the presence of DH in the α relaxation regime has been studied in bulk polymers, by e.g. multidimensional NMR and simulations, and tuned by nanoconfinement. Of particular interest to the present study are the findings that DHs at both α and β relaxation time scales have been reported in colloids as clusters of faster-moving particles and numerical studies found heterogeneous dynamics of JG relaxation, supporting previous suggestions. On even shorter time scales, picosecond DH - observed by incoherent quasieelastic neutron scattering - allowed the evaluation of the characteristic time scale of primary relaxation of molecular liquids. Furthermore, it has been noted that JG involves a broad distribution of processes with those occurring at longer times being characterized by a longer length scale.

A familiar tool to expose and quantify DH is the non-gaussian parameter (NGP):

\[ \alpha_2(t) = \frac{3}{5} \frac{\langle r^4(t) \rangle}{\langle r^2(t) \rangle^2} - 1 \]  

where \( r(t) \) and \( \langle \cdots \rangle \) denote the modulus of the particle displacement in a time \( t \) and the ensemble average, respectively. NGP vanishes if the displacement is accounted for by a single, i.e. spatially homogeneous, gaussian random process. Instead, if dispersion is present and the individual particles undergo distinct motions (even if with gaussian features), NGP is positive, \( \alpha_2(t) > 0 \). Notably, NGP is accessible to experiments e.g. by confocal microscopy and neutron scattering. Recently, JG β relaxation has been resolved by studying the isochronal NGP in simulations of metallic glasses.

It must be pointed out that a distribution of relaxation times, as observed in JG relax-
...does not imply necessarily non-gaussian displacements - i.e. non vanishing NGP, the customary criterion to identify DH -, e.g., see the Rouse model which predicts multiple relaxation times of an unentangled single chain with gaussian particle displacements.

In this work we report results from extensive molecular-dynamics (MD) simulations of a polymer model melt proving, when JG relaxation is present, a bimodal double peaked NGP leading to two distinct DH growths in the $\beta$ and $\alpha$ relaxation regimes. A mechanistic microscopic explanation is provided. Our findings capture a new correspondence between JG and structural relaxation in polymers.

2 Model and Numerical Methods

We adopt a variant of coarse-grained models of linear polymers having nearly fixed bond length and bond angles constrained to $120^\circ$, Fig. 1a. Details are given as Supporting Information (SI). All the data presented in this work are expressed in reduced MD units, in particular lengths are in units of the length scale $\sigma$ of the Lennard-Jones (LJ) potential. An interesting aspect of our variant is that, even if the force field does not include a torsional potential, thus saving computing times, an effective torsional barrier occurs for $l_0 < 0.5\sigma$, see Fig. 1b, due to the LJ repulsion between the farther two monomers in a chain fragment of four monomers, see Fig. 1a. We focus on $l_0 = 0.48\sigma$ and $l_0 = 0.55\sigma$ leading to considerable or missing torsional barrier, respectively, see Fig. 1b. It is worth noting that the assessment of the interplay of the intra-chain torsional barriers and other barriers, e.g. the ones arising from packing constraints, is a difficult task and goes beyond the purposes of the present work.
3 Results and discussion

3.1 Detection of JG process by bond reorientation

Experiments and simulations demonstrated that orientational correlation functions are sensitive to detect and resolve secondary motions, in particular, the reorientation of the chain bonds. Let us define the bond correlation function (BCF) $C(t)$ as:

$$C(t) = \langle \cos \theta(t) \rangle$$

where $\theta(t)$ is the angle spanned in a time $t$ by the unit vector along a generic bond of a chain. An average over all the bonds is understood. Starting from the unit value, BCF decreases in time, finally vanishing at long times when the bond orientation has spanned all the unit sphere.
Fig. 2 plots BCF for the bonds with length $l_0 = 0.48\sigma$ and $l_0 = 0.55\sigma$. In agreement with previous MD studies on similar model polymers,$^{12}$ it is seen that in the presence of shorter bonds BCF exhibits a characteristic two-step decay (in addition to the initial decay for $t \lesssim 1$), signaling the presence of two distinct relaxation processes interpreted as the JG and structural relaxation.$^{41,42,47}$ The relaxation maps of the JG and the $\alpha$ processes for the present model were reported elsewhere.$^{47}$

Figure 2: BCF with bond length $l_0 = 0.48$ (top) and $l_0 = 0.55$ (bottom) at different temperatures. If $l_0 = 0.48$, a clear two-step decay — evidencing two distinct relaxations—is observed.

3.2 Dynamic heterogeneity

Fig. 3 compares the time evolution of DH of the melts of chains with different bond length by resorting to their NGP. At an early stage ($t \lesssim 0.1$), the ballistic regime is interrupted by the first repeated collisions of the monomers with the surrounding cage and the subsequent NGP growth is largely independent of both the temperature and the bond length. The missing role played by the temperature suggests that DH is driven by static structure at short times,
Figure 3: NGP with bond length $\ell_0 = 0.48$ (top) and $\ell_0 = 0.55$ (bottom) at different temperatures. Color codes as in Fig. 2. In the presence of JG relaxation ($\ell_0 = 0.48$) two peaks increasingly grow by lowering $T$ with positions close to the knees observed in BCF associated to the $\beta$ and the $\alpha$ relaxations, see Fig. 2.
as also concluded by previous works.\textsuperscript{[24]} The small peak at $t \sim 0.1$ locates the average time needed by the monomer to hit the cage of the first neighbours.\textsuperscript{[19]} For $t > 0.1$, DH is fully developed and NGP exhibits a complex pattern, strongly dependent on both temperature and bond length.

![Graph showing NGP time evolution](image)

Figure 4: Time evolution of NGP at $T = 0.85$ for the two bond lengths.

To reach better insight, Fig.4 compares the NGP time evolution of the chains with the two bond lengths at the same temperature (the lowest investigated). Three regions are seen.

- Region I: at short times ($t \lesssim 2$) the NGPs are nearly coincident.

- Region II: at intermediate times ($2 \lesssim t \lesssim 7 \cdot 10^2$) in the presence of shorter bonds, switching the $\beta$ relaxation on in this region, NGP is lower, i.e. DHs are weaker, and, after a maximum, decreases to a local minimum. Chains with longer bonds, with no apparent JG relaxation, do not show the minimum, as already known.\textsuperscript{[20]}

- Region III: at long times ($t \gtrsim 7 \cdot 10^2$), where the $\alpha$ relaxation takes place, in the presence of shorter bonds, NGP reaches a second maximum and finally decays, whereas the NGP of chains with longer bonds decreases monotonously.
3.3 Correlation between relaxation and dynamic heterogeneity

There is a well-defined correlation between the DH evolution and relaxation. To deepen this aspect, in addition to BCF, dealing with bond reorientation, we also consider the correlation loss of the torsional angle (TACF),\cite{11,12} see SI for rigorous definition. We also inspect quantities concerning the monomer dynamics: (i) the mean square displacement (MSD) $\langle \delta r^2(t) \rangle$ where $\delta r^2(t)$ is the square modulus of the monomer displacement, $\delta r(t)$, in a time $t$, (ii) the self-part of the intermediate scattering function (ISF) $F_s(q,t) = \langle \exp[iq \cdot \delta r(t)] \rangle$ where $i^2 = -1$, $q$ is the modulus of the wavevector $q$. ISF is negligibly small if the displacement exceeds the length scale $2\pi/q$. We choose $q = q_{\text{max}}$, where the static structure factor is maximum, so that $2\pi/q_{\text{max}} \sim \sigma^*$, i.e. about the monomer diameter. We stress that, differently from TACF and BCF, both MSD and ISF are single-particle observables.

Figure 5: from top left to bottom right: BCF, TACF, ISF and MSD at $T = 0.85$ for the two bond lengths. Panels are partitioned in the same three regions of Fig. 4. The dashed curves in the TACF panel are best fits with the stretched function $A \exp\left[-(t/\tau_0)\beta\right]$ with $A = 0.93$, $\beta = 0.4$, $\tau_{0.48} = 190$ (red dashed), $\tau_{0.55} = 797$ (blue dashed).

Fig. 5 presents the angular relaxation functions (BCF and TACF) together with monomer MSD and ISF, for the two bond lengths at the lowest temperature. Both BCF and TACF
clearly show a larger decay in region I and II if the bond is shorter, i.e. the latter undergoes faster small-angle reorientation at short times, especially in the β region. In spite of the higher angular mobility at short times, both BCF and TACF signal that shorter bonds slow down in the α region (region III). While the reduced decay rate is apparent in BCF, to better visualise the effect in TACF, we first fitted the decay of TACF of chains with longer bonds with a stretched decay in regions II and III which, then, was log-shifted and superimposed to the TACF of chains with shorter bonds. We argue that the slowing down is due to the torsional hindrance due to shorter bonds, see Fig.4. The accelerated reorientation of shorter bonds at short times and their slowing down at long times have strong influence on the pattern of NGP. In fact, the crossover between regions I and II of NGP, see Fig.4 occurs when the reorientation of shorter bonds becomes apparent, see BCF in Fig.5. This suggests that the reduction of NGP observed in polymer systems with shorter bonds follows by a partial averaging of DH. The latter effect is due to the fast bond reorientation which inhibits the DH increase, finally resulting in the bump observed in NGP at \( t \sim 10 \). Naturally, the complete DH erasure needs wide changes in the chain conformation and then full dihedral torsions. This explains why NGP, like BCF, decays more slowly in region III in the presence of shorter bonds. In particular, the peak of NGP in region III for shorter bonds is interpreted as due to the large scale structural relaxation triggered by the wide-angle torsions. Finally, by inspecting both ISF and MSD in Fig.5 one sees that the chains with shorter bonds exhibit smaller monomer MSD in regions I and III, and slower structural relaxation in region III, as previously reported. The role of ISF to model the memory kernel in a generalized Langevin equation theory dealing with both primary and JG relaxations has been emphasized.

The finding that, in the presence of JG relaxation, DH built up in the JG regime persists up to the \( \alpha \) relaxation, see Fig.4 conveys the impression of the possible interplay of DHs in the JG and the long-time regimes, also on the basis that long-time DH is sensed at short times, including JG time scale for non-polymeric systems, and even at vibrational time scales.
3.4 Negligible memory between particle displacements occurring in JG and $\alpha$ time scales

The sound assessment of the previous hinted correlation must consider if it holds not only on the whole system but on subsets too. In non-polymeric liquids NMR experiments answered affirmatively by selecting subensembles of particles with given mobility, whereas simulations on diatomic liquids concluded that there is no connection between properties associated to beta and alpha regimes of a single molecule. On the other hand, the invariance of the ratio $\tau_{JG}/\tau_{\alpha}$ to $P$ and $T$ variations was reported in linear polymer melts represented by a simple bead-necklace model, a variant of which is adopted in the present paper. We investigate this aspect by considering the system with JG relaxation ($l_0 = 0.48$)

![Figure 6: Fraction of monomers of polymer system with JG relaxation ($l_0 = 0.48$) retaining memory of their slow (or fast) mobility at JG time scale up to primary relaxation, $f_{\text{slow} \rightarrow \text{slow}}$ and $f_{\text{fast} \rightarrow \text{fast}}$ respectively. In the case of full memory $f_{\text{slow} \rightarrow \text{slow}} = f_{\text{fast} \rightarrow \text{fast}} = 1$. Dashed line sets the level $f_{\text{random}} = 0.05$ corresponding to the absence of memory. Cross-conversion fractions are also plotted. The results point to very weak memory. See text for details. Inset: same analysis with $l_0 = 0.55$.](image)

and focusing on the DH developed at $t_\beta$, the time of the first peak of NGP (located in the $\beta$ region) and DH still surviving at $t_\alpha$, the time of the second peak (located in the $\alpha$ region), see fig. We search for "memory" effects between $t_\beta$ and $t_\alpha$ by: i) selecting a subset of $n$ monomers ($n = 0.05N$, $N$ being the total monomers) with smallest (or largest) displacement in a time lapse $t_\beta$, and ii) evaluating the fraction $f_{\text{slow} \rightarrow \text{slow}}$ (or $f_{\text{fast} \rightarrow \text{fast}}$) of these monomers still belonging to the same mobility subset of $n$ monomers with lowest (or highest) mobility after displacement in a time $t_\alpha$ (in the case of full memory $f_{\text{slow} \rightarrow \text{slow}} = f_{\text{fast} \rightarrow \text{fast}} = 1$,
whereas in the *absence* of memory \( f_{\text{slow} \rightarrow \text{slow}} = f_{\text{fast} \rightarrow \text{fast}} = f_{\text{random}} \) with \( f_{\text{random}} = n/N \), with the latter result following if the initial subset in step i) is assembled by picking-up the \( n \) monomers *randomly*. Fig.6 shows the results. It is seen that both \( f_{\text{slow} \rightarrow \text{slow}} \) and \( f_{\text{fast} \rightarrow \text{fast}} \) are only slightly larger than \( f_{\text{random}} \), i.e. the correlation between the two subsets is quite low. The same conclusion is reached by considering the fractions \( f_{\text{slow} \rightarrow \text{fast}} \) and \( f_{\text{fast} \rightarrow \text{slow}} \) accounting for the cross-conversion between the two mobility subsets (note that in the case of full memory both quantities *vanish*). The pattern does not change by considering *longer* bonds, i.e. no JG relaxation, and different ratios \( n/N \) (\( n/N = 0.03, 0.1 \), not shown).

4 Conclusions

In conclusion, we evidenced the heterogeneous character of the JG relaxation by MD simulations of a model polymer with secondary relaxation tuned by an effective adjustable torsional barrier. The DH evolution exhibits two distinct maxima at JG and structural relaxation time scales. We find that subsets of monomers lose memory of their mobility acquired in JG time scale before the occurrence of structural relaxation.

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Supporting Information Available

Simulation Details

We study a melt of coarse-grained linear polymer chains with $N_c = 512$ linear chains made of $M = 25$ monomers each, resulting in a total number of monomers $N = 12800$. Adjacent bonded monomers belonging to the same chain interact via the harmonic potential $U_{bond}(r) = k_{bond} (r - l_0)^2$, where the constant $k_{bond}$ is set to $2000\epsilon/\sigma^2$ to ensure high stiffness. A bending potential $U_{bend}(\alpha) = k_{bend} (\cos \alpha - \cos \alpha_0)^2$, with $k_{bend} = 2000\epsilon$ and $\alpha_0 = 120^\circ$, is introduced to maintain the angle $\alpha$ between two consecutive bonds nearly constant. Non-adjacent monomers in the same chain or monomers belonging to different chains are defined as "non-bonded" monomers. Non-bonded monomers, when placed at mutual distance $r$, interact via a shifted Lennard-Jones (LJ) potential:

$$U_{LJ}(r) = \epsilon \left[ \left( \frac{\sigma^*}{r} \right)^{12} - 2 \left( \frac{\sigma^*}{r} \right)^6 \right] + U_{cut},$$

where $\sigma^* = 2^{1/6}\sigma$ is the minimum of the potential, $U_{LJ}(r = \sigma^*) = -\epsilon + U_{cut}$. The potential is truncated at $r = r_c = 2.5\sigma$ for computational convenience and the constant $U_{cut}$ adjusted to ensure that $U_{LJ}(r)$ is continuous at $r = r_c$ with $U_{LJ}(r) = 0$ for $r \geq r_c$.

It is important to note that the above model allows LJ interactions between all non-bonded monomers. This is the feature to build the torsional barrier up when $l_0 < 0.5\sigma$ discussed in the paper. Alternatives reported in the literature exclude the LJ interactions between atoms separated by three bonds or less.$^{12}$

All the data presented in the work are expressed in reduced MD units: length in units of $\sigma$, temperature in units of $\epsilon/k_B$, where $k_B$ is the Boltzmann constant, and time in units of $\tau_{MD} = (m\sigma^2/\epsilon)^{1/2}$. We set $\sigma = 1$, $\epsilon = 1$, $m = 1$ and $k_B = 1$.

Simulations were carried out with the open-source software LAMMPS.$^{58,59}$ Equilibration runs were performed at constant number of monomers $N$, constant vanishing pressure $P = 0$.
and constant temperature $T$ ($NPT$ ensemble). For each state the equilibration lasted at least for $3\tau_{ee}$, being $\tau_{ee}$ the relaxation time of the end-to-end vector autocorrelation function. Production runs have been performed within the $NVT$ ensemble (constant number of monomers $N$, constant volume $V$ and constant temperature $T$). Additional short equilibration runs were performed when switching from $NPT$ to $NVT$ ensemble. No signatures of crystallization were observed in all the investigated states.

**Torsional correlation function**

Alternatively to the BCF, the relaxation of the chain backbone arrangement has been characterized by the torsional autocorrelation function (TACF):

$$TACF(t) = \frac{1}{N_c M - 3} \sum_{n=1}^{N_c} \sum_{m=1}^{M-3} \frac{\langle |\varphi_{m,n}(t)| |\varphi_{m,n}(0)| \rangle - \langle |\varphi_{m,n}(0)| \rangle^2}{\langle |\varphi_{m,n}(0)|^2 \rangle - \langle |\varphi_{m,n}(0)| \rangle^2}, \quad (4)$$

where $|\varphi_{m,n}(t)|$ is the modulus of the $m$-th dihedral angle of the $n$-th chain at a given time $t$. $\varphi$ is trivially related to the angle $\phi$ considered in Fig. 1 of the manuscript by the relation $\varphi = \phi - 180^\circ$. The dihedral angle features the torsion of a given bond. Terminal bonds are not subject to torsion, therefore, in a chain of length $M$, there are $M-3$ dihedral angles. A graphical representation of this angle is reported in Fig. $\varphi$ is given from the intersection of the two planes defined by four consecutive monomers in a chain: the first plane is defined considering the first set of three monomers while the second one is defined by the last set of three.

The temperature dependence of the TACF curves is reported in Fig. 8.
Figure 7: Representation of the dihedral angle, $\varphi$, characterizing the torsion of the central bond.

Figure 8: Temperature dependence of TACF in the two studied systems.

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Graphical TOC Entry

Dynamic heterogeneity

\( \alpha \) relaxation

\( \beta \) relaxation

Time

Johari-Goldstein ON

Johari-Goldstein OFF

Evolution of mobility spatial distribution