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Open boundary molecular dynamics of sheared star-polymer melts

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Open Boundary Molecular Dynamics (OBMD) simulations of a sheared star polymer melt in isothermal conditions are performed to study the melt's rheology and molecular structure under a fixed normal load. Comparison is made with the standard molecular dynamics (MD) in periodic (closed) boxes with a fixed shear rate (using the SLODD dynamics). The OBMD system exchanges mass and momentum with adjacent reservoirs (buffers) where the external pressure tensor is imposed. Insertion of molecules in the buffers is made feasible by implementing there a low resolution model (blob-molecules with soft effective interactions) and then use the Adaptive Resolution Scheme (AdResS) to connect with the bulk MD. Straining with increasing shear stress induces melt expansion and a significantly different redistribution of pressure compared with the closed case. In the open sample the shear viscosity is also a bit lowered but more stable against the viscous heating. At a given Weissenberg number, molecular deformations and material properties (recoverable shear strain, normal stress ratio) are found to be similar in both setups. We also study the modelling effect of normal and tangential friction between monomers implemented in a dissipative particle dynamics (DPD) thermostat. Interestingly, the tangential friction substantially enhances the elastic response of the melt due to a reduction of the kinetic stress viscous contribution.

1 Introduction

Die swelling¹ is a well known phenomenon in polymer melts and most viscoelastic liquids which consist of the sudden expansion of the liquid after exiting out a slit or orifice. The most frequent explanation has a microscopic origin: molecules elongate in the stream direction and compress perpendicularly exerting extra elastic pressure in the normal planes. This leads to the so called *normal stress differences* which are the landmark of viscoelasticity. Despite this accepted view, the devil is in the details and although considerable effort has been carried out since the middle of the last century (see e.g. Refs.¹ for historic details) accurate modelling of polymer melts is a very difficult task. Any of the many constitutive relations¹ for continuum models cannot generally predict the rheology of a new molecular polymer design. In turn, the huge span in time scales in any standard poly-

mer melt limits the scope of molecular dynamics (MD) to simple rheological tests with extremely small samples under simple (usually steady) flows. However, a detailed account of bonded and nonbonded interactions in atomistic (AT) simulations (see e.g.²⁻⁴) is able to grasp relevant information, maybe then to feed continuum (fluid dynamics) models. In between, coarse grained (CG) molecular modelling is useful for many reasons. We give at least a couple: first, polymer science has some degree of universality which benefits the use of simplified models, quite often able to provide insight and valuable predictions^{4,5}. Second, the theory of coarse graining to extract precise coarse potential interactions pertaining to the atomistic model at hand is now advancing at relatively fast rate^{4,6,7}. More recently the community has started to recognize the relevance of the dynamic aspects of coarse graining either based on following routes such as GENERIC⁸ or the Mori-Zwanzig formalism⁹. The idea is to perform short atomistic simulations⁸ to extract the quantities determining the reversible and irreversible dynamics of the slow variables, such as the friction kernels⁹. These friction kernels (between polymer "blobs") are naturally implemented in the dissipative particle dynamics (DPD) method as, notably, Español et al.⁹ showed (under the Markovian and pairwise approximation) that DPD can be formally derived from the Mori-Zwanzig coarse graining route. Another relatively newer route is the use of hybrid models concurrently combining continuum and molecular simulations. Yasuda et al.¹⁰ and other

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(mostly Japanese) groups are exploiting a version of these hybrids to model polymer melts. We have also worked in this field using essentially the same technique used in this work for simple fluids^{11,12}. Contrary to Yasuda et al.¹⁰ and other hybrid schemes the present method is designed to open up the simulation box so as to consider melt expansions. Here we apply the idea to study the rheology of much larger, polymeric molecules. At present we restrict to the Open Boundary Molecular Dynamics (OBMD) simulation¹³, without connection to the continuum side. However, the present method naturally connects with continuum fluid dynamics via hybrid schemes^{14–20}.

Another aspect of this work is the use of DPD as a tool to check the effect of details of monomer friction in the rheological behaviour of these CG models. In particular, the tangential friction between blobs naturally arises when performing dynamic coarse graining⁹ but is seldom included in these sort of analysis (see e.g. the DPD study for melts by Fedosov et al.²¹). To conclude our comments on methodological aspects, we also note that the present simulations in open domains use a useful trick which consists of using an even lower detailed molecular model to feed the molecular reservoir close to the open system boundaries. The idea is taken from the so called “Adaptive Resolution Scheme” (AdResS)^{22–34} and permits to generalize the use of AdResS in standard periodic (closed) boxes (see e.g. Refs.^{11,12,35–37}).

The main physical question we pose here is what are the rheological consequences of imposing a fixed pressure load to a sheared sample of polymer melt, compared to the case of shearing at a fixed volume. This question was also raised by a bunch of groups spread over the last two decades^{38–43} with contradictory results. It is indeed a particularly relevant question for molecular simulations because the vast majority of numerical studies on melts have considered closed (usually periodic) systems, while many rheological experiments are carried out under a normal load: melt across slabs in Ref.⁴⁴ (cited experiments therein) or cone and plate rheometry under a fixed load¹.

From the fundamental side, this question connects with the already mentioned die swelling phenomena whose details are still not completely understood. It also connects with another interesting question which is, what consequences do boundary constraints have on flowing (far-from-equilibrium) polymer melts (see Ref.⁴⁵ for a recent study). The OBMD is a flexible tool for these questions on boundary constraints because it can be tuned to permit different ensembles⁴⁶ such as the grand canonical, isoenthalpic, isothermal, constant stress (Neumann-like) or constant shear (Dirichlet-like)¹⁴. In particular it could be useful to validate theories for non-equilibrium thermodynamics (such as Extended Thermodynamics^{47,48}), or the far from trivial fate of fluctuations of mass (related to sound modes) and momentum in sheared complex fluids, which often leads to undesirable instabilities in sheared or extruded melts, like the shark skin^{1,49}.

Even in unentangled melts the influence of boundary or global constraints on the density expansion of sheared melt is still poorly understood with studies present contradicting results. At least there is a consensus on the fact that for a given shear, the shear viscosity η is larger in the isochoric (NVT) constraint, than under either a constant pressure^{39,40} or a constant load^{43,44}. A clear

manifestation of this effect was presented in a numerical study of Thompson et al.⁴⁴: a slab of lubricating liquid (20-mers) flowing between two solid walls presented a shear thinning exponent $\beta_\eta \simeq 2/3$ under a constant normal load while just 0.5 under a constant volume (here $\eta \sim \dot{\gamma}^{-\beta_\eta}$ with $\dot{\gamma}$ the shear rate). This effect has been ascribed to the shear dilatancy manifesting in a larger hydrostatic pressure under a constant volume^{39,40}. Indeed, polymer melts viscosities often increases with the pressure⁴⁹ as also observed in Ref.⁴⁴. In Sec. 6 we offer a more precise analysis showing that the springs’ stress is reduced in the open system due to the smaller intermolecular friction in the expanded melt.

There are several kinds of “isobaric” conditions: a constant hydrostatic pressure (P_{iso}) is usually termed isobaric, while a constant normal load (here P_{22}) is closer to industrial processes like slit extrusion¹ or lubricants⁴⁴. Both can show substantially different rheological behaviour when compared with constant volume studies (see e.g. Ref.⁴⁴). Experimental works of the Couette flow under constant load are also scarce and indicate shear expansion and a measurable increase of the melt’s viscosity with increasing external pressure^{50,51}.

The most striking differences in the computational literature are found in the density variation with shear. Dlugogorski³⁸ (for a FENE dumbbell) and Daivis and Evans (modelling decane)³⁹ seem to be the first to perform a molecular simulation showing density decreasing under shear (using isobaric conditions). Daivis and Evans use the term “shear dilatancy”, following the term used by Reynolds on the same phenomena (see quotation in Ref.³⁹). Note that “shear dilatancy” has been later often used instead of “shear thinning”, but here it is not. By contrast, Xu et al.⁴⁰ (attractive linear chains up to 50-mers) report just the opposite result (compression under shear). For (purely repulsive) branched chains (under NVT), they also report a reduction in the hydrostatic pressure with shear. A subsequent study by Matin⁴³ for linear chains *at a constant load* (and chain lengths up to 50 monomers) found shear dilatancy and also a non-monotonous trend for the hydrostatic pressure (as Xu et al.⁴⁰ and others^{2,3} did). Shear dilatancy was also later also found by Bosko et al.⁴¹ when analysing dendrimer melts under isobaric conditions. Consistently, they found a pressure increase under constant volume. Shear dilatancy is the trend we also observe but, under our imposed constant normal load (which is closer to experimental setup^{1,39,44}) we find that the density (and viscosity) is controlled by the load and not by the hydrostatic pressure. However, there is a lack of studies and the question remains about universality of shear dilatancy of polymer melts under a Couette flow (consistent with die swell phenomena under Poisson flow¹).

To conclude the introduction some words should be said about the technological relevance of star molecules (see discussion in Ref.^{52–54}) which certainly arises from its unique dynamic features. Star molecules present a broad range of relaxation times associated with different molecules pulsations (rotation, elastic deformation, arm entanglements) analysed in Ref.⁵³. Each relaxation time triggers a change in rheological regime once the external perturbation (shear) exceeds the corresponding threshold rate. Present results also illustrate this phenomena. Moreover star molecules bridge the gap between linear polymers and

colloids^{54,55} and can present interesting (colloidal-like) ordering effects, sometime enhanced due to its softer character^{53,56–59}. In this context, a suggesting observation in these simulations is the onset of some ordering in the neutral direction at large shear rates whose origin (hydrodynamic or entropic) remains to be established.

In Sec. 2 we briefly present the OBMD method which was otherwise more thoroughly explained in Ref.¹³. The star molecule melt model is presented in Sec. 3 and Sec. 4 shows that the OBMD correctly reproduces thermodynamic equilibrium according to the grand canonical ensemble. In Sec. 5 we present results for sheared melts in absence of the tangential friction between monomers and analyze the results in Sec. 6 according to the pressure balance. This serves to enlighten the discussion on the effect of the tangential friction in Sec. 8. Finally Sec. 9 discusses some interesting results obtained in melts presenting severe viscous heating that depends on the characteristics of the DPD thermostats and their friction kernels. Comparison with previous results is made in Sec. 10 while conclusions and future outlook are given in Sec. 11.

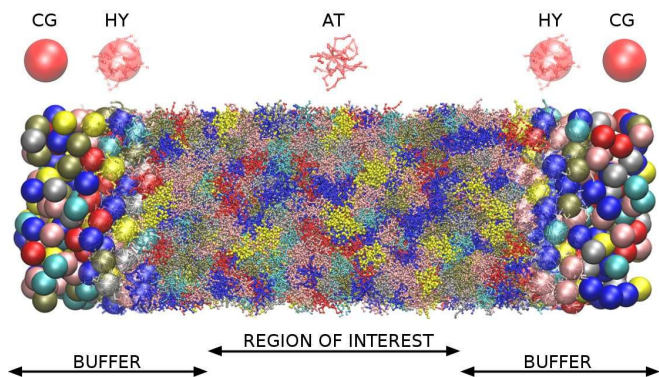


Fig. 1: Schematic representation of the open system in equilibrium along the longitudinal direction. All monomers and polymers in the system are of the same kind. Different colors are used for the sake of clarity of the picture. In the region of interest polymers are represented in the highest, i.e. atomistic (AT), resolution. Buffer regions (see text) are heterogeneous, i.e. containing regions of different resolution. The change of resolution from AT (dots), to coarse-grained, i.e. CG, (spheres) occurs in hybrid region (HY) of each buffer, carried out by AdResS. New molecules are inserted into CG part of buffers. The system is open at both ends of the box. Upper part of the figure depicts schematic representations of a molecule inside regions of different resolution.

2 Open Boundary Molecular Dynamics

We begin by briefly explaining the method for the OBMD which combines features of the open MD⁴⁶ and adaptive resolution¹¹. The reader is referred to a review on open MD¹⁴ and to Ref.¹³ for a more detailed presentation of the present OBMD implementation to star polymers. The OBMD simulation is carried out in an open rectangular box which, in the present setup, permits the MD domain to exchange mass and momentum through two of its boundaries (along the x_2 direction) with a reservoir (called buffer) which is maintained at some desired thermo-mechanic state. The buffer's two domains of finite extent embed the central part of the box (the MD domain). They allow for molecular insertion or deletion so as to keep their average molecular density

fixed (typically to a fraction between 0.5 to 0.7 of the bulk density). The OBMD is therefore not periodic in the coupling direction. Molecules are free to enter or leave the buffer from or to the MD domain, but in doing so they cross another layer where they gradually change their atomistic resolution (73 monomers for the star molecule considered hereby) to a reduced CG model, comprising one only spherical “blob” per molecule. Obviously the CG layer is placed inside the buffer domain (which here also contains a smaller atomistic part). This strategy permits to perform an otherwise impossible task: the insertion of new polymeric molecules into the melt. New molecules are inserted into the low resolution layer of the buffer, where soft CG interactions govern the dynamics of the blob-model polymers. Soft effective interactions can be obtained from the Boltzmann iteration procedure^{60,61}, although we shall see that in principle, the consistency of OBMD (in terms of pressure balance across the layers) does not depend on the CG potential chosen. The insertion of these blob molecules is carried out by the USHER scheme⁶² and the change from CG to monomer molecular resolution (usually termed atomistic resolution, AT) is carried out by the AdResS^{22–34}.

The dynamics of the monomers can be described by the following equations of motion:

$$\frac{d\mathbf{r}_i}{dt} = \mathbf{v}_i, \quad (1)$$

$$m_i \frac{d\mathbf{v}_i}{dt} = \mathbf{F}_i^{ad}(\{\mathbf{r}\}) + \mathbf{F}_i^{th}(\{\mathbf{v}\}) + \mathbf{F}_i^{ext}(\mathbf{r}_i). \quad (2)$$

Here \mathbf{r}_i denotes the position of i -th particle, \mathbf{v}_i its velocity, and m_i its mass. The total force acting on this particle has three contributions: the external force \mathbf{F}_i^{ext} acting only on the particles at the buffer (to impose the desired momentum flux); the adaptive resolution force \mathbf{F}_i^{ad} which accounts for all type of particle-particle interactions and the thermostat \mathbf{F}_i^{th} contribution (here, it is applied to the whole system).

The adaptive resolution force \mathbf{F}_i^{ad} is constructed to allow for a momentum conserving “alchemic” transformation of the molecules, which takes place gradually along the transition layer (where $0 < w < 1$, see below). The transition is achieved by the following linear combination of the AT and CG forces (respectively $\mathbf{F}_{\alpha\beta}^{AT}$ and $\mathbf{F}_{\alpha\beta}^{CG}$)²²,

$$\mathbf{F}_{\alpha\beta}^{ad} = w(x_\alpha)w(x_\beta)\mathbf{F}_{\alpha\beta}^{AT} + (1 - w(x_\alpha)w(x_\beta))\mathbf{F}_{\alpha\beta}^{CG}. \quad (3)$$

Both expressions are correspondingly weighted by a position dependent function $w(X)$, whose value equals 0 in the CG region and 1 in the AT one and gradually changes in between (transition layer). The adaptive resolution force provided by AdResS is not derived from a Hamiltonian and does not conserves the energy²³. It is however constructed to obey Newton’s third law, ensures the conservation of total linear momentum of the system and can thus be used to study fluid flows. This fact is unimportant to the present study which targets isothermal sheared systems (not-conserving energy anyhow). The OBMD model might be generalized by using the recent Hamiltonian AdResS (H-AdResS)^{63,64} which is based on a global Hamiltonian (i.e. also allows Monte Carlo simulations). In such hypothetical case, extra care should

be taken with the momentum conservation because of the presence of drift-forces in H-Adress coming from the free energy difference between the AT and CG models⁶⁵.

An essential function of the buffer region is the imposition of boundary conditions to the open MD box. This is done by adding an extra “external” force at the buffer regions, \mathbf{F}^{ext} , calculated from Eq. 4 (see e.g. Ref.⁶⁶).

$$\mathbf{F}^{ext} = (\mathbf{P}_{out} - \mathbf{P}_{in})/\delta t + A J^P \mathbf{n} \quad (4)$$

Here \mathbf{P}_{out} and \mathbf{P}_{in} represent the total linear momenta of the particles that exited and were inserted into the simulation in the last time step of integration δt , respectively. J^P is the momentum flux tensor, while \mathbf{n} is the outwards normal vector of an open-end plane of the box¹¹. In general, the pressure tensor contains normal and tangential contributions, i.e. $J^P = -p_{nn}^{ext} \mathbf{nn} - p_{nt}^{ext} \mathbf{nt}$. The external force is designed to exactly conserve the linear momentum over the whole particle system (buffer+MD) and it is distributed among the buffer particles according to $\mathbf{F}_i^{ext} = G(x_i) \mathbf{F}^{ext}$. To allow a different distribution of the normal and tangential forces, the distribution function G is chosen to be a tensor defined by Eq. 5¹³,

$$\mathbf{G}(x_i) \equiv \frac{g_{\parallel}(x_i)}{\sum_{i \in B} g_{\parallel}(x_i)} \mathbf{nn} + \frac{g_{\perp}(x_i)}{\sum_{i \in B} g_{\perp}(x_i)} \mathbf{tt}, \quad (5)$$

with g_{\parallel} determining the spatial distribution of the normal force and g_{\perp} the distribution of shear stress. Both functions are depicted in Fig. 2.

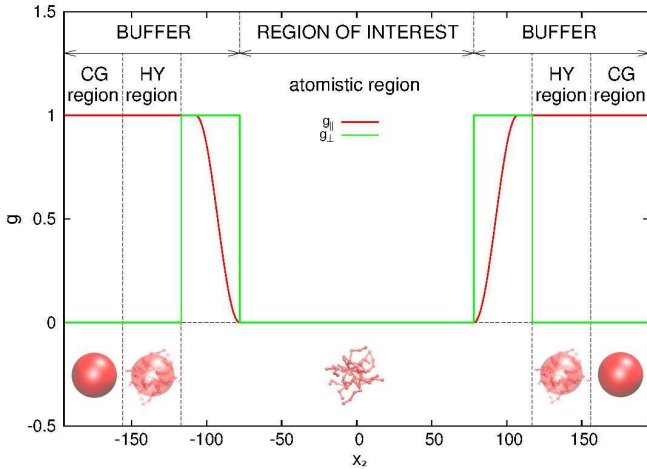


Fig. 2: Buffer distribution function. The force \mathbf{F}^{ext} , that acts on buffer regions in order to impose the boundary conditions at open ends of the box to the region of interest, is distributed among molecules inside buffers via the depicted functions, forming the distribution function tensor \mathbf{G} , given by Eq. 5. The force acting on each molecule equals $\mathbf{F}_i^{ext} = \mathbf{G}(x_i) \mathbf{F}^{ext}$.

Many OBMD applications (as those illustrated hereby) involve transfer of momentum (the pressure tensor) from outside the MD domain. This requires a momentum conserving thermostat. In production runs we used the DPD thermostats (while a strong damping Langevin for equilibration purposes). Our choice for the DPD thermostat is not only due to conserve momentum (in principle one could use the Lowe-Andersen⁶⁷) but also because of

modelling purposes. The Mori-Zwanzig formalism, under Markovian conditions leads to coarse-graining dynamics with DPD-like equations of motion⁹. The message of that solid theoretical result is that the friction kernels introduce an important modelling aspect. The friction kernels in a CG model of some real melt, should ideally be measured from force-force correlations of the detailed all-atom model⁹. Here we adopt a simpler but yet useful route, which is to study how friction affect the rheology of the model melt. The generic form the of the DPD thermostat used is

$$\mathbf{F}_i^{th} = - \sum_j \Gamma(r_{ij}) (\mathbf{v}_i - \mathbf{v}_j) + \tilde{\mathbf{R}}_{ij}, \quad (6)$$

where $\tilde{\mathbf{R}}_{ij}$ is the fluctuating force constructed to satisfy the fluctuation-dissipation under equilibrium conditions. We refer the reader to Refs.⁶⁸⁻⁷⁰ for details on the DPD implementation. As in Ref.^{9,69,70} the friction kernel has normal and tangential components,

$$\Gamma = \gamma_{\parallel} \mathbf{n}_{ij} \mathbf{n}_{ij} + \gamma_{\perp} \mathbf{t}_{ij} \mathbf{t}_{ij},$$

where \mathbf{n}_{ij} is the vector joining two monomers i and j and \mathbf{t}_{ij} determines the directions in the perpendicular plane. The kernels γ_{\parallel} and γ_{\perp} are distance dependent. Its shape (for a particular all-atom model) can be obtained from dynamic CG⁹, here we will use Heaviside functions with a certain cutoff distance (see Table 1 for details). Most thermostat require or introduce some form of friction, albeit, the great majority of simulations of polymer melts do not considered thermostating (and its added friction) as part of the molecular model, but just a way to remove the heat dissipated under shear. Also most DPD simulations, such as the relatively recent work on sheared melts²¹ do not introduce the tangential friction between blobs, but rather take the form of the most standard DPD kernel (normal friction alone). However, as also pointed out by Padding and Briels⁵, friction should be considered as a part of the CG model. Indeed, as shown by Hijon⁹ et al. (CG of a star molecule as the unit blob), tangential and normal frictions can be quite different from each other. Here we start to explore how tangential and normal frictions affect the rheology of a star molecule under far-from-equilibrium conditions.

In the following sections we present and analyse results obtained for the DPD thermostat with no tangential friction between monomers (blobs). This analysis is then used to understand the effect of the tangential friction, considered in Sec. 8. In the case of the normal friction alone, the substantial heat dissipated by the sheared melt requires from us to implement a slight modification of the DPD thermostat to keep constant (kinetic) temperature at the largest shear rates. Details of this modified DPD thermostat (we call it “adaptive”) and of friction kernels are given in Sec. 3.

Importantly, simulations in closed isothermal (periodic) boxes (NTV) are used as reference to investigate the effect of open boundaries. Shear flow in the closed system has been simulated using Lees-Edwards boundary conditions⁷¹ and the SLLOD algorithm^{72,73}.

3 Setup and melt models

The simulation setup is illustrated in Figs. 1 and 2. The polymer melt is exposed to a Couette flow in the x_1 direction being sheared

along the x_2 direction (gradient direction). The vorticity or neutral direction is x_3 . In the closed box we use the SLODD^{72,73} dynamics to impose the desired shear rate $\dot{\gamma}$ in a closed periodic box (with constant particle number and volume). In the open setup, the surfaces located at $x_2 = \pm L_2/2$ are submitted to equal normal pressures p_{22}^{ext} and (opposite sign) tangential stresses $\pm p_{12}^{ext}$, in such a way that the rotational part of the shear flow turns counter-clockwise in the flow-gradient ($x_1 - x_2$) plane. No constraint is imposed to the remaining component of the pressure tensor, resulting in $\langle p_{13} \rangle = 0$ and a self determined $\langle p_{33} \rangle$. The box is periodic in the other two directions x_1 and x_3 so this setup corresponds to an slice of polymer melt with fixed load at two of its boundaries (at $x_2 = \pm L_2/2$).

The melt is made of the star polymer model already presented in Refs.^{9,74} Each polymer consists of 73 monomers, i.e. 12 arms of 6 monomers attached to the central monomer. In what follows we use m_0 , σ_0 and ϵ_0 for mass, length and energy units and we will arbitrarily set these units to $m_0 = 1$, $\sigma_0 = 1$, and $\epsilon_0 = 1$. The resulting time unit is $\tau_0 = \sigma_0(m_0/\epsilon_0)^{1/2} = 1$. Excluded volume interactions of monomers are modelled by the repulsive Weeks-Chandler-Anderson interaction with diameter parameter $\sigma = 2.415\sigma_0$ and energy parameter $\epsilon = 1$. The interactions between two adjacent bonded monomers are harmonic with a linear spring of stiffness constant $K = 20.0\epsilon/\sigma_0^2$. The equilibrium distance between non-central monomers is $r_{ij}^{eq} = 2.77\sigma_0$ while the equilibrium distance between the central monomer and the first monomer of an arm is $r_{ij}^{eq} = 3.9\sigma_0$. The size of the simulation box is $390 \times 117 \times 117$ (in units of σ_0).

Simulations were performed at the fixed constant (monomer's kinetic) temperature ($T = 4.00 \pm 0.01$). Results presented in the following sections correspond to simulations obtained using a modified DPD thermostat which we label as "adpd". This adpd thermostat has no tangential friction $\gamma_{\perp} = 0$ while $\gamma_{\parallel} = 1$ (some results also with $\gamma_{\parallel} = 5$). The cutoff of the Heaviside friction kernel is $R_{dpd} = 2^{1/6}\sigma$. We refer to Table 1 for thermostat and kernel details. We recall that the effect of the tangential friction is analysed in Sec. 8. In Sec. 9 we illustrate the heat dissipation and temperature increase observed when using standard thermostats DPD with normal friction.

Thermostatting sheared polymer melts is a delicate issue due to the large amount of heat they dissipate. At a large enough shear rate or large shear stress, the temperature of the melt increases. The same phenomenon is also observed in experiments and industrial processes (extrusion) at high shear rates (typically above 500Hz)^{47,75,76}. Phenomenological temperature "corrections" for the melt's viscosity are often used in industry and experiments⁷⁷. Although this problem goes beyond the present manuscript, the temperature of a system under a non-equilibrium steady state is also a fundamental problem because equipartition is lost and the different temperature definitions present slight variations (kinetic versus configurational temperature⁷⁸). In the literature, few works comment on the problem of viscous heating in molecular simulation (see the exceptions in Refs.⁷⁹⁻⁸¹) and many published material elude reporting on possible temperature variation in their sheared thermostated systems. After the present exper-

Table 1: Thermostats used in simulations. Standard means a standard DPD thermostat and the adaptive DPD is explained in Eq. 7. The transverse DPD thermostat from Ref.⁷⁰ is denoted by "tdpd".

Label	Kernel cutoff (R_{cut}^{DPD})	γ_{\parallel}	γ_{\perp}
sdpdshort	$2^{1/6}\sigma$	[1.0-20.0]	0
sdpdlong	$1.5 \times 2^{1/6}\sigma$	1.0,5.0	0
adpd	$2^{1/6}\sigma$	1.0,5.0	0
tdpd	$1.5 \times 2^{1/6}\sigma$	1.0	1.0

ience, we believe that some of the data presented in previous papers might be somehow biased by temperature. We will show an indication later in Sec. 9.

We considered four different thermostats: the "standard" (sdpd)⁶⁹ and "transverse" (tdpd)⁷⁰ DPD thermostats and also a modified DPD thermostat. The latter is able to extract larger amounts of heat by self-adapting its temperature parameter T_{DPD} which controls its random force term. This "adaptive DPD thermostat" (adpd) as we call it, dynamically adjust T_{DPD} according to a sort of coupled heat equation,

$$\frac{dT_{DPD}}{dt} = -\frac{1}{\tau_{DPD}} (T_{MD} - T_{igt}), \quad (7)$$

where $T_{igt} = 4$ is the target system's temperature and T_{MD} is the kinetic temperature obtained from the variance of the monomers peculiar velocities $\mathbf{u}_i = \mathbf{v}_i - \mathbf{v}_f(\mathbf{r}_i)$. Here $\mathbf{v}_f(\mathbf{r}_i)$ is the flow velocity at the position of the monomer \mathbf{r}_i evaluated on-the-fly from (time averaged) binned x_2 coordinate. Eq. 7 resembles the characteristic equation of the Berendsen thermostat⁸², where the linear differential equation in time is solved for the current temperature of the system. In our case, on the other hand, T_{DPD} is just the temperature incorporated in the equations for the DPD thermostat and not the actual temperature of the system. The basic idea is simple: if the system under non-equilibrium sheared state is producing substantial heat due to the friction, the thermostat's noise term (or in physical terms, the hypothetical reservoir temperature T_{DPD}) should be made colder to faster extract heat. The adpd thermostat does not alter the equilibrium state (indeed we first checked this fact), but reduces the noise term under substantial shear.

The thermostat time τ_{DPD} (acting like a coefficient of heat transfer to the "reservoir") was set to $\tau_{DPD} = 100\delta t$. The thermostat nominal temperature T_{DPD} was updated using a simple explicit Euler scheme for Eq. 7 with a time step of $100\delta t$. All the thermostats used are applied to the monomers relative velocities. The friction kernels of the DPD thermostats (damping and noise terms are constructed using the same kernel⁷⁰) is chosen to be a Heaviside function with a cutoff distance R_{cut}^{DPD} , i.e. $\gamma(R) = 1$ for $R \leq R_{cut}^{DPD}$ and zero otherwise.

All results presented here correspond to a normal friction kernel $\gamma_{\parallel} = 1$ (some results for the adpd thermostat were also carried out for $\gamma_{\parallel} = 5$ to test sensitivity). To test the effect of tangential friction we also run simulations with $\gamma_{\parallel} = \gamma_{\perp} = 1$ in the tdpd thermostat. The thermostat details and labels used are given in Table 1.

The integration step ranges from $\delta t = 0.01\tau_0$ to $\delta t = 0.005\tau_0$

for the highest shear rates. Note that $\tau_0 = 1$ is smaller than the standard Lennard-Jones time (monomer-monomer interaction), $\tau = \sigma(m/\epsilon)^{1/2} = 2.415 \tau_0$ where $m = 1$ is the monomer mass.

The equilibration of the lump of melt in the OBMD simulation is conducted by a modified version of AdResS, whereby the weighting function w in Eq. 3 is gradually increased in time starting from $w = 0$ (CG model)⁸³. The weighting function is therefore switched from a position-dependent to a time-dependent one. The resolution is thus gradually sharpened from CG to AT. The procedure is in detail described in Ref.¹³. After equilibration, each simulation is run for $10000\tau_0$.

4 Melt at Equilibrium

4.1 Characteristic times

In view of the close relation between structure and dynamics that takes place in sheared melts (and complex fluids in general), it is interesting to present the range of physical times of the melt before analysing its structural transformation with shear. In star polymers one can observe three types of relaxation phenomena⁸⁴. First is elastic deformation of the overall shape of polymers, second relaxation occurs via the rotational diffusion, and the third one regards disentanglement of arms of every star polymer. Each of the relaxation processes can be estimated from the integral of the corresponding normalized autocorrelation function (ACF), via $\tau_A = \int_0^\infty C_A(t)dt$, and these are given by Eqs. 8, 9, and 10, respectively.

$$C_{elas}(t) = \frac{\sum_i \langle \mathbf{R}_i(t) \mathbf{R}_i(0) - \langle \mathbf{R}_i \rangle^2 \rangle}{\sum_i \langle \mathbf{R}_i^2 \rangle - \langle \mathbf{R}_i \rangle^2} \quad \text{Center-end correlation} \quad (8)$$

$$C_{rot}(t) = \sum_i \frac{\langle \mathbf{R}_i(t) \cdot \mathbf{R}_i(0) \rangle}{\sum_i \langle \mathbf{R}_i^2 \rangle} \quad \text{Rotational diffusion} \quad (9)$$

$$C_{arm}(t) = \frac{1}{f(f-1)} \sum_{\substack{i,j=1 \\ i \neq j}}^f \langle [\mathbf{R}_i(0) \cdot \mathbf{R}_j(0)] [\mathbf{R}_i(t) \cdot \mathbf{R}_j(t)] \rangle \quad (10)$$

Arm entanglement

\mathbf{R}_i represents the center-end vector of arm i , R_i its length, t time, f number of arms of each polymer. And i and j are indices of different arms within the same polymer. Each autocorrelation function decays with its characteristic time of the relaxation process^{84,85} and these are given in Table 2 for the different DPD thermostat friction kernels considered. We checked that under equilibrium state all thermostats produce consistent results, in terms of pressure and density, while the correlations and the characteristic times differ, as they should. The adpd and the sdpdshort correspond to the same friction kernel and produce similar relaxation times at equilibrium (within statistical uncertainty) indicating that the adpd modification does not essentially alter the dynamics. We observe that the disentanglement of the arms occurs more rapidly than rotational diffusion and that is due to the short length of the arms, as each contains only 6 monomers. The longest time is the diffusion time for the molecules' center of mass (CoM) $\tau_{dif} = R_g^2/D$, where $R_g = 7.6\sigma_0$ is the radius of gyration of

Table 2: Polymer times scales obtained at equilibrium for different setups and thermostats (see text). All times are in units of $\tau_0 = \tau/2.415$, where $\tau = \sigma(m/\epsilon)^{1/2}$ is the standard Lennard-Jones time scale for monomers. In all standard and adaptive DPD cases the thermostat damping constant is $\gamma_{DPD} = 1m_0/\tau_0$, while for the tdpd thermostat $\gamma_{||} = 1m_0/\tau_0$ and $\gamma_{\perp} = 1m_0/\tau_0 \cdot \tau_{elas}$, τ_{rot} , and τ_{arm} are defined as characteristic decay times of the ACF given by Eqs. 8, 9, and 10, respectively. Diffusion time is defined as $\tau_{dif} = R_g^2/D$, with $R_g = 7.6\sigma_0$ the average radius of gyration in equilibrium.

Simulation	τ_{arm}	τ_{elas}	τ_{dif}	τ_{rot}
open sdpdshort	33 ± 2	6 ± 1	800 ± 100	55 ± 5
closed sdpdshort	33 ± 2	6 ± 1	800 ± 100	55 ± 5
open sdpdlong	55 ± 2	10 ± 1	1200 ± 100	100 ± 5
closed sdpdlong	55 ± 2	10 ± 1	1100 ± 100	100 ± 5
open adpd	33 ± 2	3 ± 1	700 ± 100	59 ± 5
closed adpd	33 ± 2	3 ± 1	800 ± 100	59 ± 5
open tdpd	195 ± 5	35 ± 5	4800 ± 400	332 ± 5

polymers and D the diffusion constant of CoMs, which is different for every friction kernel. The differences in the relaxation times between the sdpdshort and sdpdlong thermostats are due to the larger thermostat cutoff distance R_{cut}^{DPD} leading to a larger inter-particle friction in the sdpdlong case^{70,86}. Interestingly, the ratio between relaxation times and viscosity is similar for all the thermostats in Table 1 (values coincide within error bars), regardless of the kernel and R_{cut}^{DPD} ⁸⁷. This indicates that the translational and orientational dynamics are affected in a similar way by the thermostats.

Using the adpd thermostat (normal friction alone) we get $\tau_{elas} = 3 \pm 1$, $\tau_{arm} = 33 \pm 2$, $\tau_{rot} = 59 \pm 5$ and $\tau_{dif} = 700 \pm 100$, which illustrates the wide range in time scales involved in these sort of simulations. These times are similar in the open and closed systems (in equilibrium) and are compared with those obtained with other thermostats and kernels in Table 2. However, as shown below the rheology of the melt is not determined by the molecular diffusion, but rather by the molecular relaxation times. For the star molecule under study with $f = 12$ arms of length $l_a = 6\sigma$, molecular rotation is slower than the arms elastic relaxation, the ratio being $\tau_{rot}/\tau_{arm} > 10$ in all cases considered (see Table 2). This ratio determines the type of rheological behaviour of the melt according to a theoretical approach based on solving the Fokker-Planck equation for the bonds distribution^{88,89}. We will come back to this issue in Sec. 5.3.

4.2 Equation of state

As explained in Ref.¹³, the OBMD simulations of the melt at equilibrium provide the correct average thermodynamic variables. The pressure equation of state obtained in OBMD agrees with that obtained from a standard NVT simulation for all volume fractions studied¹³. More precisely, the average equilibrium density $\langle \rho \rangle (p_{22}^{ext})$ obtained with OBMD at a fixed normal pressure p_{22}^{ext} are consistent with the equilibrium pressure p calculated in a closed MD simulation at fixed density $\rho = \langle \rho \rangle (p_{22}^{ext})$. In passing,

it should be highlighted that the same CG potential was used for all the melt densities considered, indicating that the pressure consistency at the MD domain is independent of the coarse-grained potential used in the buffer.

In the following we work with the polymer volume fraction, defined as $\Phi = \rho_N \pi \sigma^3 / 6$, where $\rho_N = N/V$ is the number density (N and V represent the number of monomers and the volume of the region of interest and the mass density is $\rho = m\rho_N$ with monomer mass $m = 1$, thus $\rho = 0.136\Phi$). For the NVT ensemble, the study conducted here corresponds to $p = (0.093 \pm 0.001)$ and $\Phi = 0.20$, fixed for any shear rate. In the open box we fix $p_{22}^{ext} = 0.093$ and for zero shear rate (equilibrium) we get $\langle \Phi \rangle = (0.20 \pm 0.01)$. Fig. 3 shows the normalised density profile of polymers ($\rho_M^{slab} / \rho_M^{avg}$). Here ρ_M^{slab} denotes mass density of polymers in each slab, where it is measured, and ρ_M^{avg} its average value. The latter is constant in closed simulations, i.e. $\rho_M^{avg} = 0.0271$, while its calculated value in open cases equals $\rho_M^{avg} = 0.0271 \pm 0.0001$. We observe that the obtained density profile is flat in the region of interest with some minor artefacts, which are due to the lack of statistics. Along the buffer zones the density gradually decreases as a consequence of the application of the external pressure p_{22}^{ext} . As it has been explained in previous related works on hybrid atomistic-continuum schemes^{14,66}, this inhomogeneity does not affect the transfer of pressure and stress from the exterior, provided that the density profile is flat around the hybrid interface. This is indeed the case, as it can be seen in Fig. 3 (see the interface between “region of interest” and “buffer”). The radial distribution function (RDF) of CoMs of molecules is in perfect agreement with NVT simulations.

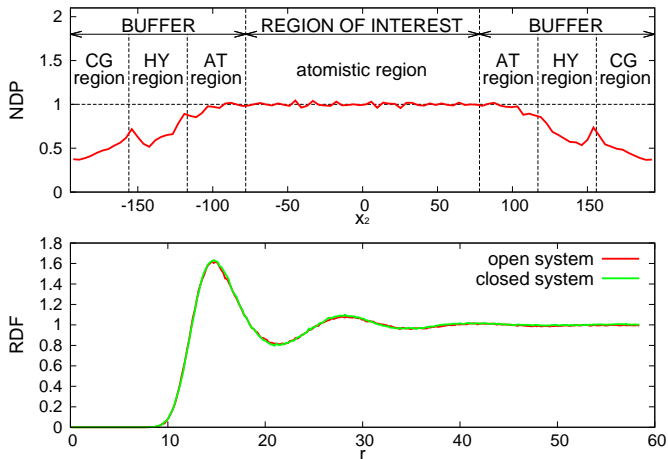


Fig. 3: Normalised density profile (NDP - top panel) and comparison of RDFs (bottom). NDP is depicted in the direction, in which the box is open. It is normalised to the desired value of the density, which corresponds to occupational factor $\Phi = 0.2$. RDFs are calculated from molecules inside the region of interest in open simulation (red) and in closed system (green).

4.3 Mass fluctuations

In the grand canonical ensemble, mass fluctuations are related to the integral of the RDF⁹⁰, so the excellent agreement between the RDF's from NVT and open boxes suggest that the fluctuations in the number of molecules inside the interest domain should be

thermodynamically consistent with the grand canonical prescription: namely, the relative fluctuation in polymer mass $M = \rho_M V$ should be $\text{Std}[M]/M = [k_B T / (M c_T^2)]^{1/2}$, where c_T is the isothermal sound velocity $c_T^2 = (\partial p / \partial \rho)_T$. c_T , which is evaluated using the pressure equation of state from Ref.¹³, is shown in Fig. 4. The agreement between OBMD and the NVT ensemble is excellent. In the same figure we plot the results for the relative mass fluctuations at different pressures (average densities) and compare with the grand canonical prediction. We find that the agreement is very good, notably because of the tiny relative mass fluctuations in the (not small $V = 3504384\sigma_0^3 = 248805\sigma_0^3$) volume considered for our open box: which range between 3.6% and 0.26% at the largest density considered ($\Phi = 0.2$). In terms of mass density variance $\sigma_\rho^2 = \rho k_B T / (V c_T^2)$, for the state we considered hereafter under shear ($p = 0.093\epsilon_0 / \sigma_0^3$, $\Phi = 0.2$ and $\rho = 0.0271 m_0 / \sigma_0^3$) the OBMD result is $\sigma_\rho^2 = (5.0 \pm 0.3) \times 10^{-9} m_0 / \sigma_0^3$ in excellent agreement with the variance predicted for the μVT ensemble 4.79×10^{-9} . The conclusion of this study strongly supports our claim that the OBMD equilibrium simulation is sampling the grand canonical ensemble without any (or negligible) bias. It has to be said that the value of the external chemical potential $\mu^{ext} = \mu(p_{22}^{ext}, T)$ can however not be imposed in OBMD, although it could be reconstructed following the standard Gibbs-Duhem route with varying external pressure. New implementations of AdResS^{29,65} might also be used to evaluate μ .

A typical outcome for the time evolution of the *total* mass of polymer in the MD domain is shown in Fig. 4. It presents oscillations, suggesting that it might contain some information about the sound velocity of the system.

Sound propagation can indeed be studied in our simulations because we work with momentum preserving (DPD) dynamics (by contrast, sound is damped in the standard Langevin dynamics). It is noted that an “ideal” open system should be transparent to all waves, meaning that all waves, either created by inner mass fluctuations of any wavelength or by external waves travelling across, should leave the system and do not reflect back. This implies, in particular, that in the absence of external longitudinal forces, fluctuations of the *total* mass should have no memory, being a white noise (or at least a broad-band signal)⁹¹. The presence of correlations in the total mass of our “region of interest” is in fact due to partial reflections of waves at the rarefied buffers, where density fluctuations are reduced (recall we fix the average total mass of these reservoirs, see e.g.⁶⁶). In the autocorrelation (ACF) of the total mass fluctuation, the dominant wavelength should obviously be the largest possible one compatible with this condition and the *total* simulation box length. A density mode with wavenumber $k = 2\pi/\lambda$ decays like $\langle \rho(\mathbf{k}, t) \rho(\mathbf{k}, 0) \rangle = \langle \delta \rho^2 \rangle \exp[\nu_L k^2 t] \cos[\omega t]$ with ν_L the sound attenuation coefficient⁹² and $\omega = c_T k$ the oscillation frequency. Inspection of Fig. 3 indicates that the density profile of the *whole* system (MD+ buffer) roughly conforms to Dirichlet boundary conditions with fixed density at the end of the buffer domains $\rho(x_2 = \pm L_2/2) < \rho_0$. In such case, the longest wavelength available to the system's mass fluctuations would be $\lambda_{eff} \simeq 2L_2$. We fitted the time autocorrelation function of the MD mass to extract ω and compare it with the ansatz $\omega = c_T k_{eff}$ using $\omega = c_T k_{eff}$

with $k_{eff} = 2\pi/\lambda_{eff}$. The best fit to the simulation results corresponds to $\lambda_{eff} = 760\sigma_0$, while $2L_2 = 780\sigma_0$ and it is shown in Fig. 4 in terms of ω/k_{eff} and compared with the melt's sound velocity c_T . The excellent match confirms that the mass in the MD domain has memory induced by reflections of sound waves against the low density domains (buffers). Such reflections could be reduced by coupling the MD with a continuum hydrodynamic field outside⁶⁶, or by imposing a non-reflecting boundary condition⁹¹ (still to be generalized to MD, see also⁹²). However, in the present scenario we find that this result is quite interesting because it suggests the possibility of measuring the sound velocity c_T from the fluctuations of the total MD mass. In particular, it might allow to measure how the sound velocity is modified in a sheared melt, $c_T(\text{Wi})$. Although a detailed study would be certainly required; just as an indication, we find that $c_T(\text{Wi})$ (estimated in this way, i.e. from mass fluctuations), does decrease.

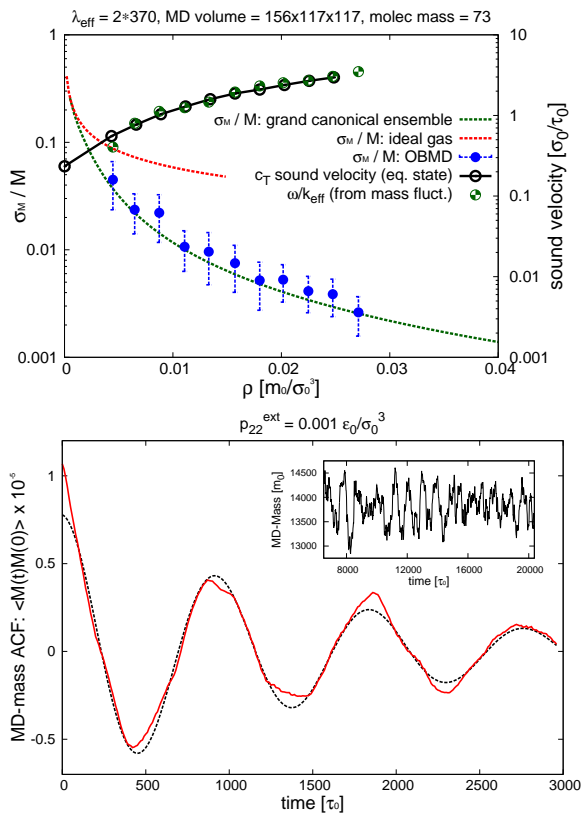


Fig. 4: Top panel: The relative fluctuation of the polymer mass (standard deviation over average polymer mass M) in the interest MD-domain of the open setup (see Fig. 2). Comparison is made with the grand canonical theoretical result (green dashed line) $\text{Std}[M]/M = [k_B T / (M c_T^2)]^{1/2}$ and, also included, the ideal gas limit (dot dashed) obtained with the isothermal sound velocity $c_T^{id} = (k_B T / M_m)^{1/2}$ with $M_m = 73m_0$ the molecular mass. The volume of the interest domain is $V = 156 \times 117 \times 117 \sigma_0^3$. Recall that the monomer LJ diameter is $\sigma = 2.415\sigma_0$ and its mass is $m_0 = 1$. The right ordinate axis shows the values of the isothermal sound velocity $c_T = (\partial p / \partial \rho)_T$ (from the pressure equation of state) that is compared with ω/k_{eff} obtained from the oscillation frequency ω of the total mass autocorrelation function in the MD domain (see bottom panel). The effective wavenumber is $k_{eff} = (\pi/370)\sigma_0^{-1}$ and the total open box is $L_2 = 390\sigma_0$. Bottom panel: the time autocorrelation function of the mass in the MD domain at equilibrium with imposed external pressure $p_{22}^{ext} = 0.001\epsilon_0/\sigma_0^3$ (dashed line is the fit to extract ω , see text); the inset illustrates the evolution of the total mass in the MD domain for this case.

5 Sheared melt with normal friction

This section presents results for the star molecule model with zero tangential friction between monomers. Kernel and thermostat details (adpd) are given in Secs. 3 and 4. We decided to first focus on this model to avoid embarrassing the discussions with details of different cases (open versus closed, normal versus tangential friction) and also because this model presents a richer dynamical behaviour, whose analysis will be useful to understand the effect of tangential friction in Sec. 8.

5.1 Weissenberg number

The Weissenberg number Wi is a useful number to compare and dissect different regimes in polymer rheology. It is defined as

$$\text{Wi} = \tau_{rel} \dot{\gamma}, \quad (11)$$

where τ_{rel} is the longest molecular relaxation time and $\dot{\gamma}^{-1}$ is the “shear flow time” needed to affinely deform a square box of sheared fluid into a parallelepiped with an angle of 45° between adjacent planes. As stated, around Eq. 9 and in Table 2, the longest relaxation of our star polymer is related to the molecular rotation so $\tau_{rel} = \tau_{rot}$ in Eq. 11. In fact, the diffusion of the CoM of the molecules is much slower (see Table 2). The CoM diffusion does not directly sample the molecule’s structure whose modification under flow is related to the non-Newtonian character of polymers. The Peclet number determines the ratio between CoM diffusion and flow advection $\text{Pe} = \tau_{dif} \dot{\gamma}$ and for our setup is about 10 times larger than Wi . In colloidal suspensions, the shear thinning typically starts for $\text{Pe} > 1$ due to shear banding. Interestingly, star polymers constitute a sort of a bridge between the open structures of linear polymers and the compactness of colloids. Thus, one might elucidate that the onset of shear thinning in compact stars could well be due to shear banding (collective molecular ordering in lanes), rather than by (individual) polymer elongations. We shall see later on that both (collective and individual) effects take place in our sheared system, although we advance that the transition to the non-Newtonian regime takes place at $\text{Wi} = \tau_{rot} \dot{\gamma} > 1$. Hence at least for the (not so compact) star polymer studied here, shear thinning is not determined by collective ordering at straining rates faster than the CoM diffusion. The hybrid character of star molecules (between colloids and polymers) is the subject of current research^{56–58}.

5.2 Density and Hydrostatic pressure

Denoting $P_{\alpha\beta}$ the symmetric pressure tensor (exerted by the melt), the hydrostatic pressure is defined as

$$P_{iso} = \frac{P_{11} + P_{22} + P_{33}}{3}. \quad (12)$$

We find that our model for a star polymer melt expands when sheared under the normal load (see Fig. 17 which collects results from different cases). This behaviour is consistent with the die swell phenomena of polymer melts¹ and with the increase in the normal pressure P_{22} observed in all cases. By contrast, the hydrostatic pressure decreases with Wi in the open domain, while

in the NVT box presents a non-monotonous trend (with not large variations). As explained in Sec. 6 P_{iso} is influenced by several molecular mechanisms, whose relevance changes with $\dot{\gamma}$ along with the molecular structure. The density (in the open domain) and the pressures (P_{22} in NVT and also P_{iso} in both cases) are quite sensible to temperature changes, as shown in Sec. 9.

5.3 Rheology

We start by determining the zero shear viscosity, $\eta_0 = \eta(\dot{\gamma} = 0)$. A standard way is the Green-Kubo equilibrium route (via the integral of the time autocorrelation of shear stress)⁹³. To keep consistency with the non-equilibrium route we follow the experimental approach which fits the trend for $\eta(\dot{\gamma})$ for a range of values of the shear rate with some suitable expression such as the popular Carreau fit^{50,94}. We also checked that Green-Kubo viscosities⁹⁵ are similar to the Carreau-fitted ones within statistical uncertainty (about 10%). The Carreau fit has the following expression,

$$\eta = \eta_0 \left[1 + (\tau_\eta \dot{\gamma})^2 \right]^{-\beta_\eta/2}, \quad (13)$$

shown in Fig. 5 along with simulation data. Fig. 5 shows the shear viscosity η obtained under open and closed setups for several models with different thermostats and friction kernels (see Table 1). $\eta(\dot{\gamma})$ was calculated from the off-diagonal pressure tensor component $P_{12} = -\eta\dot{\gamma}$, which was measured in simulations. From Eq. 13, the viscosity shear thinning exponent β_η , is such that,

$$\eta \rightarrow \dot{\gamma}^{-\beta_\eta} \text{ for large } \dot{\gamma}$$

and for polymer melts β_η ranges between 0.4 and 1^{1,88}. This fit of Eq. 13 also provides an estimation of the zero-shear viscosity η_0 and a characteristic time τ_η related to the onset of the shear thinning regime. η_0 differs for different friction kernels of DPD thermostat (see Table 1). The adpd and sdpdshort thermostats (identical friction kernels) consistently provide the same zero shear rate viscosity $\eta_0 = 0.5$ (that does not increase largely for $\gamma_\parallel \leq 10$). The sdpdlong model with increased kernel cutoff $R_{cut}^{DPD} = 1.5 \times 2^{1/6} \sigma$ presents $\eta_0 = (0.60 \pm 0.1) + 0.29\gamma_\parallel$ an increase consistent with an increasing relaxation time with friction* as deduced in the analysis of Kindt and Briels⁹⁶. Tangential friction (tdpd) leads to $\eta_0 = 2.6$ ($\gamma_\parallel = \gamma_\perp = 1$). The meaning of τ_η becomes clear when it is compared with the estimated star rotational relaxation time τ_{rot} . Here are their values for different thermostats: $\tau_\eta = 61$ and $\tau_{rot} = 59$ for the adpd; $\tau_\eta = 58$ and $\tau_{rot} = 55$ for the sdpdshort; $\tau_\eta = 125$ and $\tau_{rot} = 100$ for the sdpdlong; and $\tau_\eta = 287$, $\tau_{rot} = 332$ for the tdpd. The error bar of the given values is approximately ± 5 . We observe $\tau_\eta \simeq \tau_{rot}$ indicating that the onset of shear thinning, which takes place at $\dot{\gamma}\tau_{rot} > 1$, coincides with the molecular deformation altering the equilibrium rotational diffusion.

We now focus on the adpd model and defer the discussion on the tangential friction to Sec. 8. The system temperature is

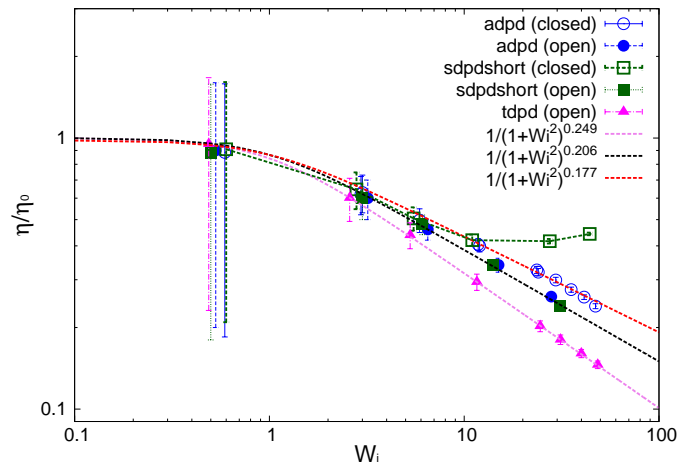


Fig. 5: Normalized shear viscosity obtained for several models under open and closed conditions. Details of the model (varying in thermostat and kernels) are given in Tables 1 and 2. Lines are best Carreau fits.

fixed to $T = 4.0 \pm 0.01$. In this case we get a zero shear viscosity $\eta_0 = 0.50 \pm 0.05$. At larger Wi , the shear thinning exponents β obtained from Carreau fits (Eq. 13) are found to be slightly steeper in the open domain $\beta_\eta = 0.41(2)$ than in the NVT box $\beta_\eta = 0.35(4)$. Thus at a fixed shear rate, the open system is slightly less viscous than the closed sample. This is in agreement with previous studies for linear and branched melts carried out at constant and unconstrained density (see Sec. 10).

The first and second normal stress coefficients $\Psi_1 = N_1/\dot{\gamma}^2$ and $\Psi_2 = N_2/\dot{\gamma}^2$ (for the adpd model, discussed in this section) are shown in Fig. 6. For $Wi < 20$ we find a decrease in Ψ_1 consistent with the Carreau standard behaviour (Eq. 13), providing $\Psi_1(\dot{\gamma} = 0) = 21 \pm 1$ and an exponent $\beta_{\Psi_1} \simeq 1.30 \pm 0.04$ (i.e. $N_1 \sim \dot{\gamma}^{0.7}$) quite similar for both ensembles. The relaxation time for Ψ_1 obtained from the fit is also consistent with $\tau_\eta = \tau_{rot}$ within error bars. At larger $Wi > 20$ we find a measurable decrease of Ψ_1 with respect the Carreau trend (see Fig. 6), which takes place at slightly smaller Wi in the open case. This corresponds to a loss in the elastic component of the melt at large shear rates. The second normal stress coefficient $\Psi_2 = N_2/\dot{\gamma}^2$, shown in Fig. 6 also takes quite similar values in both environments and at large $\dot{\gamma}$ scales like $\Psi_2 \sim -\dot{\gamma}^{-1}$. The similar behaviour for N_2 under open and closed boxed might be due to the fact that we just fix the normal load (in the x_2 direction) and not the hydrostatic pressure as in some other studies^{39,40,42}, presenting different trends for N_2 in NVT and $NP_{iso}T$ constraints.

The normal stress ratio $VR \equiv -\Psi_2/\Psi_1$ and the recoverable shear strain⁸⁷

$$SR \equiv \Psi_1/(2\eta\dot{\gamma}) = (P_{22} - P_{11})/(2P_{12}) \quad (14)$$

are standard indicators of viscoelasticity⁸⁷ (e.g. SR vanishes for a Newtonian fluid). As shown in Fig. 7 in our melt model, SR increases with the shear rate, as expected; however, both indicators (SR and VR) clearly show that a change in the elastic component of the model takes place for $Wi > 20$. Notably, SR decreases, so the melt becomes less compliant to shear strain and stores less

*An interesting remark is that the recoverable shear compliance⁸⁷ obtained from $J_e = \tau_{rot}/\eta_0$ results to be $J_e \simeq 110$ independently on the friction kernel.

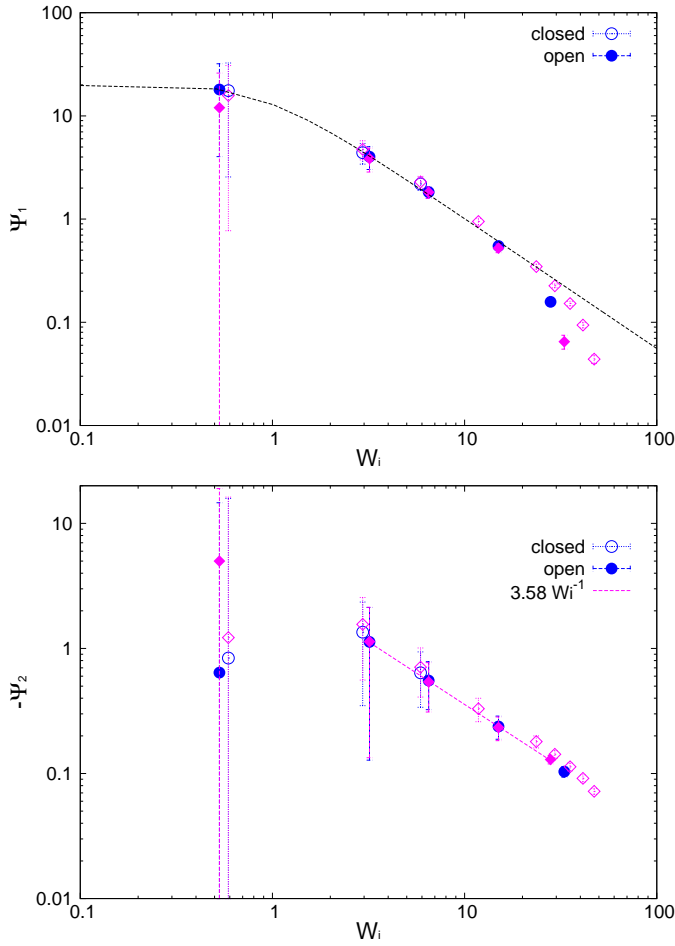


Fig. 6: First and Second normal stress coefficients of the star polymer melt under open and closed conditions and the adpd thermostat for $\gamma_{||} = 1$ (blue) and $\gamma_{||} = 5$ (purple). Dashed line for Ψ_1 is the Carreau fit (see text).

elastic energy across the flow-gradient plane, with a jump in the VR. We advance that the amount of the elastic energy loss at large Wi depends on friction forces (notably on the presence of tangential friction as shown in Sec. 8). This is a clear indication that calibration of friction from detailed all-atom models⁹ is crucial to represent or simulate some particular real melt.

5.4 Gyration tensor

Thermodynamic and rheological properties of any polymeric system are intimately related to the deformation of the molecules induced by flow. It is therefore convenient to start by presenting the results for the average gyration tensor of our star polymer model under shear, whose components are shown in Fig. 8. For $Wi > 1$ the flow induces the alignment of the star molecules in the flow direction and at the same time a compression in the gradient direction. The width of the stars in the neutral direction slightly increases up to $Wi \sim 20$ and for larger shear rate, they also start to contract in this direction. It is also instructive to observe the molecule's shape in their principal deformation axes, obtained from the diagonalization of the gyration tensor. The molecular axes are $s_1 = \cos(\theta_G)x_1 - \sin(\theta_G)x_2$, $s_2 = -\sin(\theta_G)x_1 + \cos(\theta_G)x_2$, and $s_3 = x_3$ (since $G_{13} = G_{23} = 0$). The i^{th} eigenvalue is noted

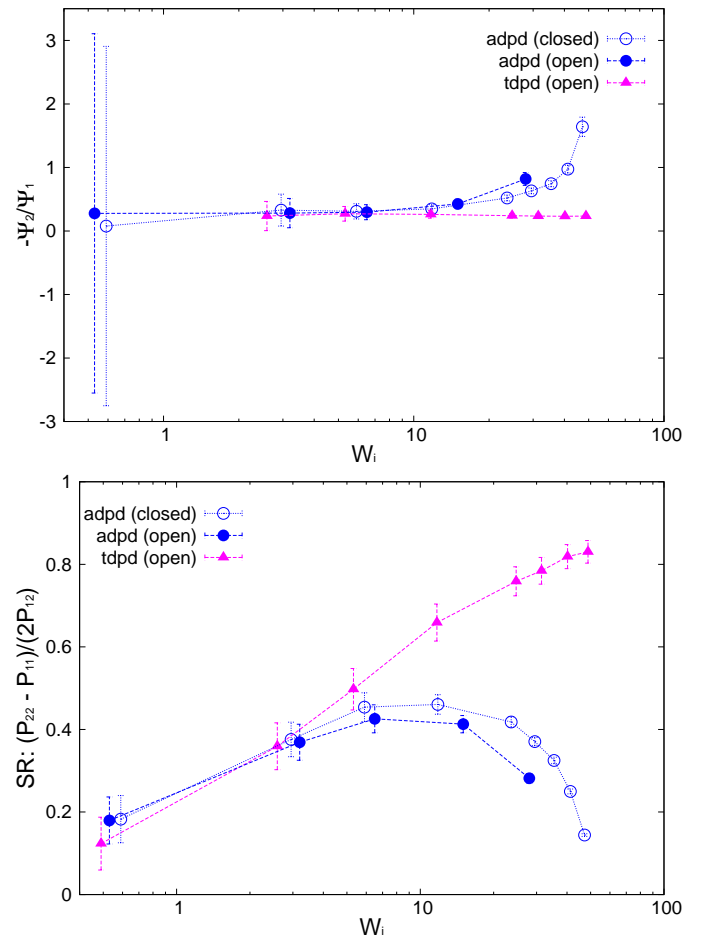


Fig. 7: (Top panel) Normal stress ratio and (Bottom panel) Recoverable shear strain index for some model melts under open and closed environments. Results for the case with tangential friction (tdpd) are also included.

as \tilde{G}_i . The angle θ_G is the average molecular tilt over the flow direction that satisfies:

$$\tan(2\theta_G) = \frac{2G_{12}}{G_{11} - G_{22}}. \quad (15)$$

Results for the eigenvalues of \mathbf{G} are shown in Fig. 8 and reveal the strong contraction of the molecule in the s_2 direction. In particular for $Wi \sim 50$ we observe that $\tilde{G}_2^{1/2} \sim 1.12\sigma$ so the chain width in the contracted axis reaches the monomer (or blob) diameter. In a real polymer at this shear rate, the flow starts altering the structure of the chain “blobs” smaller than σ . But this is precisely our modelling unit length so we will take this as the maximum Wi to be explored here (roughly $Wi < 100$). Above this (size dependent) shear rate, one should also expect a shear rate dependence of the blob-blob friction kernel (see Ref.⁹⁷ for a related study on this issue). It is also interesting to note that the total volume of the molecule (evaluated from the product of the eigenvalues of the gyration tensor) first increases (molecules expand) up to $Wi \sim 10$ and due to contraction in the neutral direction, it tends to contract (in the form of a highly extended ellipsoid) for larger Wi . In this sense, the behaviour of the arms in the neutral direction of the star have some similarities with what was found in

tethered polymers^{98,99}, which also presents a maximum volume at intermediate Wi .

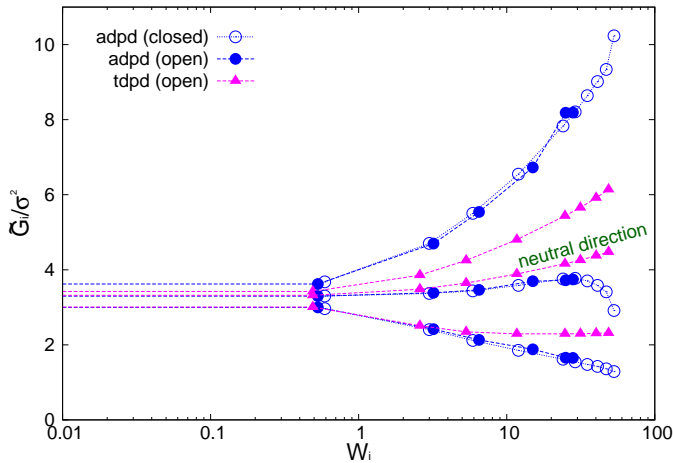


Fig. 8: Eigenvalues of the gyration tensor scaled with the monomer diameter σ^2 . Results include cases with adpd friction kernel $\gamma_{\parallel} = 1$ (no tangential friction) and the tdpd model $\gamma_{\parallel} = \gamma_{\perp} = 1$ discussed in Sec. 8. Estimated error bars of the data are approximately 5%.

6 Analysis of stress-structure and dynamics relationship

The present analysis provides relations between stress components and the molecular structure and dynamics. It focuses on the results presented in the previous section, which corresponds to the adpd star model, with zero tangential friction. The conclusions will be useful for the comparison done in Sec. 8.

6.1 Pressure balance in closed system

Pressure is the leading mechanical variable in rheology as it directly connects the microscopic world with material properties. We now elaborate on the pressure balance in an attempt to connect the molecular structure with the observed rheology.

The pressure exerted by the melt[†] is calculated using the Irving-Kirkwood method and the method of planes¹⁰⁰, which is particularly suited to the open setup as it was designed for non periodic boundaries. The pressure can be first decomposed in virial and kinetic parts. The virial pressure includes contributions from spring forces and intramolecular forces (both acting amongst pairs of monomers of the same molecule) and intermolecular forces (between monomers of different molecules). The pressure balance allows to analyze the different molecular contributions,

$$P_{\alpha,\beta} = P_{\alpha,\beta}^{kin} + P_{\alpha,\beta}^{spring} + P_{\alpha,\beta}^{intra} + P_{\alpha,\beta}^{inter}. \quad (16)$$

The pressure tensor is symmetric for the type of molecules consid-

ered hereby (in fact for most polymers)¹. The “average” pressure on the system is given by the hydrostatic pressure which is just the third part of the trace of \mathbf{P} (see Eq. 12). The hydrostatic pressure is not involved in (although indirectly affects) the main rheological quantities such as the shear stress P_{12} and the first and second normal stress differences, defined as

$$N_1 = P_{22} - P_{11}, \quad (17)$$

$$N_2 = P_{33} - P_{22}. \quad (18)$$

It is customary to introduce the traceless pressure tensor

$$\hat{\mathbf{P}} \equiv \mathbf{P} - P_{iso}\mathbf{I},$$

with $\mathbf{I} = \delta_{\alpha,\beta}$ the unit second rank tensor. Indeed, the hydrostatic pressure can be decomposed in different molecular contributions $P_{iso} = \sum_A (1/3)Tr[\mathbf{P}^A]$ with $A = \text{kinetic, springs, etc.}$ Also, due to the linearity of the trace operator we can decompose the traceless stress tensor $\hat{\mathbf{P}}$ in a sum of contributions of traceless tensors, i.e., $\hat{\mathbf{P}} = \sum_A \hat{\mathbf{P}}^A$ with $\hat{\mathbf{P}}^A = \mathbf{P}^A - (1/3)Tr[\mathbf{P}^A]$. This is useful to simplify the analysis. In fact, before discussing the differences between open and closed ensembles, we inspect some generalities of the pressure balance focusing on the results obtained for closed (NVT) ensemble. To that end we present in Fig. 9 the contributions to the melt’s pressure components under shear: to the hydrostatic pressure and to the different traceless stress tensors (component-wise: gradient \hat{P}_{22} , flow \hat{P}_{11} , neutral \hat{P}_{33} and shear P_{12} obviously $P_{13} = P_{23} = 0$).

6.1.1 Springs stress

Bonded interactions are hereby modelled by linear springs with non zero equilibrium distance. The contribution of the star molecules’ springs to the traceless stress tensor are illustrated in Fig. 9 (again, for the case of closed systems). As expected, as the shear rate is increased, molecules tend to be stretched in the flow direction and compressed in the gradient direction. Springs tend to restore the equilibrium (spherical) shape of the molecules by producing a negative stress (compressing) in the x_1 direction and a positive (expanding) in the gradient direction. As in any material with elastic properties, both effects contribute to enlarge the first normal stress difference $N_1 = P_{22} - P_{11}$. In turn, we find that elastic contribution to the second normal stress difference $N_2 = P_{33} - P_{22}$ is negative, as it happens in real polymeric fluids¹. Fig. 9 shows that the springs crossed tension P_{12}^{spring} is the most important contribution to the shear stress, although as discussed below, the kinetic pressure becomes significant at large Wi (above $Wi > 20$ for the adpd model discussed in this section).

6.1.2 Kinetic pressure

The kinetic pressure tensor $P_{\alpha,\beta}^{kin} = \frac{1}{V} \sum_i \langle \delta v_i^\alpha \delta v_i^\beta \rangle$ was obtained from the peculiar velocities $\delta \mathbf{v}_i = \mathbf{v}_i - \mathbf{u}(\mathbf{r}_i)$, where $\mathbf{u}(\mathbf{r}) = u_1(x_2)\mathbf{e}_1$ is the average velocity field obtained by binning the velocity gradient direction x_2 . Indeed, the ideal part of the hydrostatic pressure is one third of the trace of the kinetic pressure tensor, $(1/3)Tr[\mathbf{P}^{kin}] = \rho T$. However, its traceless part, $\hat{\mathbf{P}}^{kin} \equiv \mathbf{P}^{kin} - \rho T\mathbf{I}$, contains relevant information about the correlations in velocity

[†] Traditionally, following the experiment standpoint, rheological magnitudes are expressed in terms of the pressure on the system (here given by $-\mathbf{P}$). For instance, the normal stress difference is then $N_1 = -(P_{11} - P_{22})$ and this explains the opposite sign used in this (and others) simulation studies.

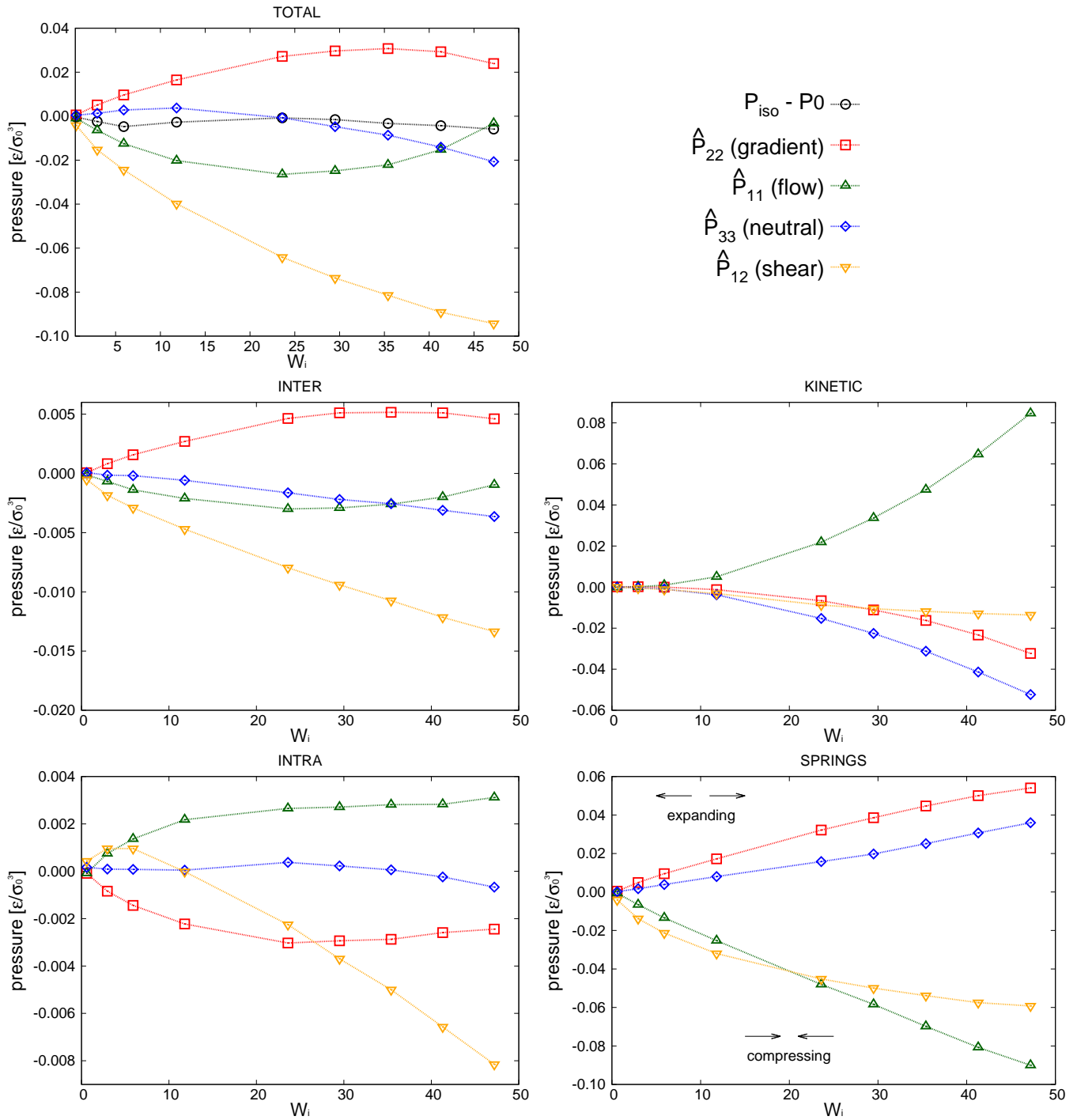


Fig. 9: Components of the different contributions to the traceless tensor ($\hat{\mathbf{P}} = \sum_A \hat{\mathbf{P}}^A$, with $A = \text{kinetic, springs, etc}$) of the traceless stress tensors $\hat{\mathbf{P}}^A = \mathbf{P}^A - (1/3)\text{Tr}[\mathbf{P}^A]$. Results correspond to the closed setup using adpd thermostat with $\gamma_{DPD} = 5$. P_0 is the (isotropic) pressure at $\dot{\gamma} = 0$, the rest of terms are explained in the text. The error bars of the data are around $0.002 - 0.005\epsilon/\sigma_0^3$.

fluctuations over different axes. In particular, monomer's velocity fluctuations in the x_1 and x_2 directions become correlated for $\dot{\gamma} > 0$ and contribute to the shear stress and viscosity. In our setup this correlation becomes negative (owing to the imposed counter-clockwise shear) and leads to a negative contribution to P_{12}^{kin} with a non-negligible viscous contribution to the shear viscosity $\eta = -P_{12}/\dot{\gamma}$ which tends to reduce the shear thinning exponent (see the comparison with the case with tangential friction in Sec. 8). Although seldom pointed out in the literature (see Ref. 4), at high enough shear rates the monomers kinetic pressure becomes significant in polymer melts. This is in fact what we clearly observe in Fig. 9 for the star model with normal friction (adpd). The anisotropy in the kinetic pressure observed in the covariance of monomer's peculiar velocities is certainly much larger than what would be observed in a simple fluid^{47,78}. It is also of a different nature because in the melt, the monomers' motions are constraint and they fluctuate and rotate being tethered to the center of their star molecule. This leads to different monomer peculiar velocities in the flow, gradient, and neutral directions which are reflected in the kinetic pressure. In Fig. 9 we show $\hat{\mathbf{P}}^{kin}$ versus Wi (recall that $\mathbf{P}^{kin} = \hat{\mathbf{P}}^{kin} + \rho T$ and diagonal components of \mathbf{P}^{kin} are positive). For $Wi > 1$, the kinetic pressure contributes to the first and second normal stress N_1^{kin} and N_2^{kin} . As soon as the molecule become stretched in the flow direction (for $Wi > 1$), velocity fluctuations in the flow direction are enhanced with respect to those along the gradient and neutral directions: $\hat{P}_{11}^{kin} > 0$ and $\hat{P}_{22}^{kin} < 0$, $\hat{P}_{33}^{kin} < 0$. We find that $N_2^{kin} < 0$ (see Fig. 10 below), so the kinetic stress acts in the same way as the elastic (springs) components. However, the kinetic contribution to the first normal stress differences is negative $N_1^{kin} < 0$ so it goes just opposite of the elastic contribution of the chain. Our conclusion is that the kinetic pressure of monomers tends to reduce the elasticity of the melt in the flow-gradient direction at large shear rates.

6.1.3 Intramolecular pressure

Intramolecular pressure gives an indication of excluded volume effects and molecular collisions within one molecule. For our moderate-size molecules it has a minor contribution to the total stress. Intramolecular collisions induce a viscous (not-restoring) stress in the melt which slightly contributes to the shear stress and tends to counterbalance the elastic first normal stress difference $N_1^{intra} < 0$ (just like the kinetic pressure - see Fig. 10). It has the same effect in the gradient-neutral plane ($N_2^{intra} > 0$) where due to the molecule stretching, monomers tend to collide less in the gradient direction. However, above $Wi \sim 20$ a shallow maximum is observed in P_{33}^{intra} (and a minimum for P_{22}^{intra} - see Fig. 9) suggesting that the reduction of the arm extension in the neutral direction has the consequence of increasing monomer collisions in the gradient direction (thus N_2^{intra} diminishes - see Fig. 10). Evidence of arm retraction in the neutral direction is shown in Fig. 8 as a clear the reduction of $\bar{G}_3 = G_{33}$ for $Wi > 20$ (also observed in "sdpdshort" and "sdpdlong" thermostats). Somewhat similar rebound in the intramolecular pressure at a large shear rate has been observed in simulations of entangled and disentangled linear polymer melts (see Ref. 3 and reference therein). However, in these works the monomer-monomer interaction is attractive,

and this fact modifies (reverses) the intramolecular contributions (polymer compression then reduces the intramolecular pressure, unlike for our purely repulsive potential).

6.1.4 Intermolecular pressure

According to Fig. 9, the contribution of the intermolecular pressure to the melt is minor. The seemingly irrelevant contribution of intermolecular stress is assumed in many theoretical models⁸⁷ to explain viscoelasticity (stress-optic rule). However, this is a simplistic view because intermolecular forces are responsible to spread the external momentum introduced through the system's boundaries (note that, in this respect, the open boundary setup behaves like a real experiment). In fact, internal forces (between monomers of the same molecule) sum up to zero so they cannot modify the CoM's velocity of the molecule. For the present star molecule (with relatively short arms) momentum in the flow direction is transferred and maintained across x_2 by intermolecular collisions⁹. These friction forces gradually build up the elastic stress in the molecules, until it finally collapses to a stationary value in the steady state. The central role of intermolecular forces can be also seen by considering the alternative molecular formulation of the pressure tensor¹⁰¹ (based on molecules (μ) CoM $R_\mu = (1/M_\mu) \sum_i m_i \mathbf{r}_{i,\mu}$ and virial pressure proportional to $\sum_\mu \mathbf{R}_\mu \mathbf{F}_\mu$). This "molecular pressure" formulation is however less informative because the effect of all internal forces, like the springs, are hidden in the spatial CoM distribution $g(\{\mathbf{R}\})$. Nevertheless, it serves to illustrate the central role of intermolecular forces and to shed some light on the apparently striking similarity between the intermolecular pressure and the total pressure dependence with Wi , which can be seen by comparison between the corresponding (inter and total) panels of Fig. 9 (note the difference in values). This similarity between intermolecular (IM) and global magnitudes (such as IM energy and hydrostatic pressure, see Fig. 9) has also been reported as "striking" in previous sheared melt simulations^{2,3}. It is interesting to note that the dominant velocity gradient component of the IM force P_{22}^{inter} reaches a plateau around $Wi \sim 20$ and slightly decreases at larger Wi (as the total P_{22} does - see Fig. 9). This is indicative of a change in the dynamics of polymers, which according to the concomitant increase observed in P_{11}^{inter} , most probably start to rotate and collide more often in the flow direction.

6.1.5 Hydrostatic pressure

The hydrostatic pressure is key in shear induced polymeric phenomena, such as shear induced crystallization⁸⁷ or separation of blends⁴⁷. Its dependence with the shear rate is not well understood. P_{iso} depends on the molecules size^{40,43} and on its architecture^{3,40,42,88}. However, there have been reports of both increase and decrease in different (sometimes contradicting) studies. Fig. 9 indicates that P_{iso} can present a non-monotonous trend with Wi due to changes in the molecular structure under shear. Non-monotonous trends for $P_{iso}(\dot{\gamma})$ have also been observed for different polymers in Refs. 2,3,40,43. The present analysis (see Fig. 9) reveals in fact that P_{iso} depends on a competition of several mechanisms. Below $Wi \simeq 10$ the hydrostatic pressure varies little but presents a slight descend, probably due to the chain expan-

sion (revealed by the analysis of the gyration tensor) which decreases the intramolecular pressure. At larger shear rates $Wi > 10$, two opposite mechanisms enter in play: First, an increase in P_{11} due to the strong increase of kinetic pressure (and to a lesser extent to intermolecular collisions). This is probably due to molecular rotations similar to the tank-thread motion reported for a star molecule solution¹⁰² and also pointed out in Ref.^{2,3}. And second, a decrease in P_{33} due to the contraction of the stars in the neutral direction (see Fig. 8) and consequent reduction of the kinetic pressure P_{33}^{kin} . Both effects nearly counterbalance each other (see Fig. 9) leaving small variations in P_{iso} . The relevance of these effects depends on the boundary conditions (the open case is analysed in next section) and type of friction (Sec. 8). More generally variations of P_{iso} depend on the presence of attractive monomer interactions and on the molecular structure.

It is interesting to note that the decrease of neutral kinetic pressure in our adpd star model starts to take place around $Wi \simeq 20$, which is precisely the ratio $\tau_{rot}/\tau_{elas} \simeq 20$. Above this shear rate the flow strains faster than the elastic relaxation of the molecule thus reducing the fluctuations of the arms in the neutral direction. This effect finally induces a net decrease P_{iso} above $Wi > 30$. We shall see (Sec. 8) that this effect is absent (or at least delayed) when introducing the tangential friction, again indicating that friction should be essential part of any CG model⁹.

6.1.6 Intermolecular forces and the Hookean limit

To summarize, at low shear rates the elastic energy stored by the melt grows in response to intramolecular (nonbonded) interactions (here mainly friction forces). At large enough shear rates viscous (Newtonian) effects coming from the kinetic and intramolecular pressures, tend to modify (normally reduce) the elastic response of the melt (notably first normal stress differences). Our findings are in agreement with the conclusions of Kroger et al.⁴ clearly and succinctly summarized in his book (page 144) in relation with the breakdown of the linear stress-optic-response (SOR) due to the Newtonian viscous transport at large Wi . In our simulations we find a linear relation between $G_{22} - G_{11}$ and $P_{22} - P_{11}$ (essentially similar in open and closed setup), indicating the validity of the linear SOR up to $Wi \simeq 10$. In Kroger's approach (applied to linear multibead FENE chains), both, kinetic and intramolecular pressures, are collected in what he calls the "simple fluid" stress contribution^{4,89}. We find here that both contributions can have different roles, which more generally should probably depend on the molecular shape (kinetic pressure) and monomers (intramolecular) interactions (attraction/repulsion).

To highlight the relevancy of the intermolecular stress in the linear SOR regime, we plot in Fig. 11 (top panel) the total and elastic contributions (springs) to the normal and shear stress against the intermolecular counterparts. At low shear rates, about $Wi < 10$, the same linear relation is found for the first normal stress difference and shear stress, $N_1 \simeq 8N_1^{inter}$ and $P_{12} \simeq 8P_{12}^{inter}$, while we find $N_2 \simeq 4N_1^{inter}$. Approximately the same linear relation holds for the elastic stress and also for the normal molecular strains evaluated with the gyration tensor ($G_{11} - G_{22}$ and $G_{22} - G_{33}$, scaled in Fig. 11(top panel)). This provides the Hookean limit of the melt $N_1 \simeq C(G_{11} - G_{22})$ and $N_2 \simeq (C/2)(G_{33} - G_{22})$

with $C = 62.5$ (we find $P_{12} \sim CG_{12}$ holds only for smaller $Wi < 5$). In the open (adpd) setup, the linear regime for N_1 and P_{12} perfectly agree with closed simulations; however we found deviation from linearity in the case of N_2 , an issue which deserves further investigation. The bottom panel of Fig. 11 presents results for stars with the tangential friction, analysed in Sec. 8.

7 Effect of open environment

7.1 Density and hydrostatic pressure

In our open domain we fix the load of the melt in the gradient direction ($P_{22}^{ext} = P_{22}$) and this produces a redistribution of the pressure tensor, reducing its component in the flow and neutral directions and also, indirectly, its shear stress. This is deduced from Fig. 12 where we compare the traceless stress tensor and the hydrostatic pressure in the open and closed setups at the fixed $T = 4$ temperature. Note that $P_{22}^{ext} = P_{22} = \hat{P}_{22} + (1/3)P_{iso}$ is constant in the open setup (within statistical uncertainties). In the open domain, the sheared melt expands in the gradient direction; a phenomenon similar to the die swell observed in polymer extrusion at the pipe's orifice and related to other viscoelastic phenomena¹⁰³. In the open domain this corresponds to a decreasing melt's density (at faster shear rates) and brings about a smaller hydrostatic pressure than in the closed environment at similar Wi (see Fig. 12). However, the relative decrease of P_{iso} is larger than the density jump. This fact is due to several related effects we now analyze from the inspection of Fig. 10. Indeed, at the fixed temperature, a lower density brings the lower kinetic pressure $P_{iso}^{kin} = \rho T$ found in the open domain (this trend also applies here to the intramolecular pressure P_{iso}^{intra} because our model considers purely repulsive nonbonded interactions, the opposite effect could arise for attracting chains^{2,3}) However, an even larger reduction in P_{iso} with respect to the closed box comes out from the smaller intermolecular pressure in the open box (see Fig. 10 for $Wi > 10$). Indeed, at a high shear, a less dense melt presents less molecular collisions, less intermolecular friction and thus less elastic load. As stated, at the fixed temperature, the elastic strain is essentially activated by intermolecular friction in the melt. Notably, for $Wi \sim 20$, these intertwined effects induce a reduction of about 25% of the open domain's hydrostatic pressure mainly arising from the decrease of elastic stress. At that Wi , density has only decreased about 10% (see Fig. 17). In agreement with this comment, we note this depressurizing effect is doubled when the tangential friction is added as commented in Sec. 8.

7.2 Normal stress differences and shear stress

Fig. 10 compares the different contributions to the normal stress differences and the shear rate in the open and closed systems. Much of what has been already said in the previous section applies here. The results for the shear stress nicely corroborates what we pointed out before about the close relation between intermolecular and elastic stresses. Fig. 10 (right bottom) clearly shows that the kinetic and intramolecular shear stresses are essentially equal in the open and closed domains. The decrease in elastic shear stress found in the open case is due to the reduction in the intermolecular friction at a lower density (although

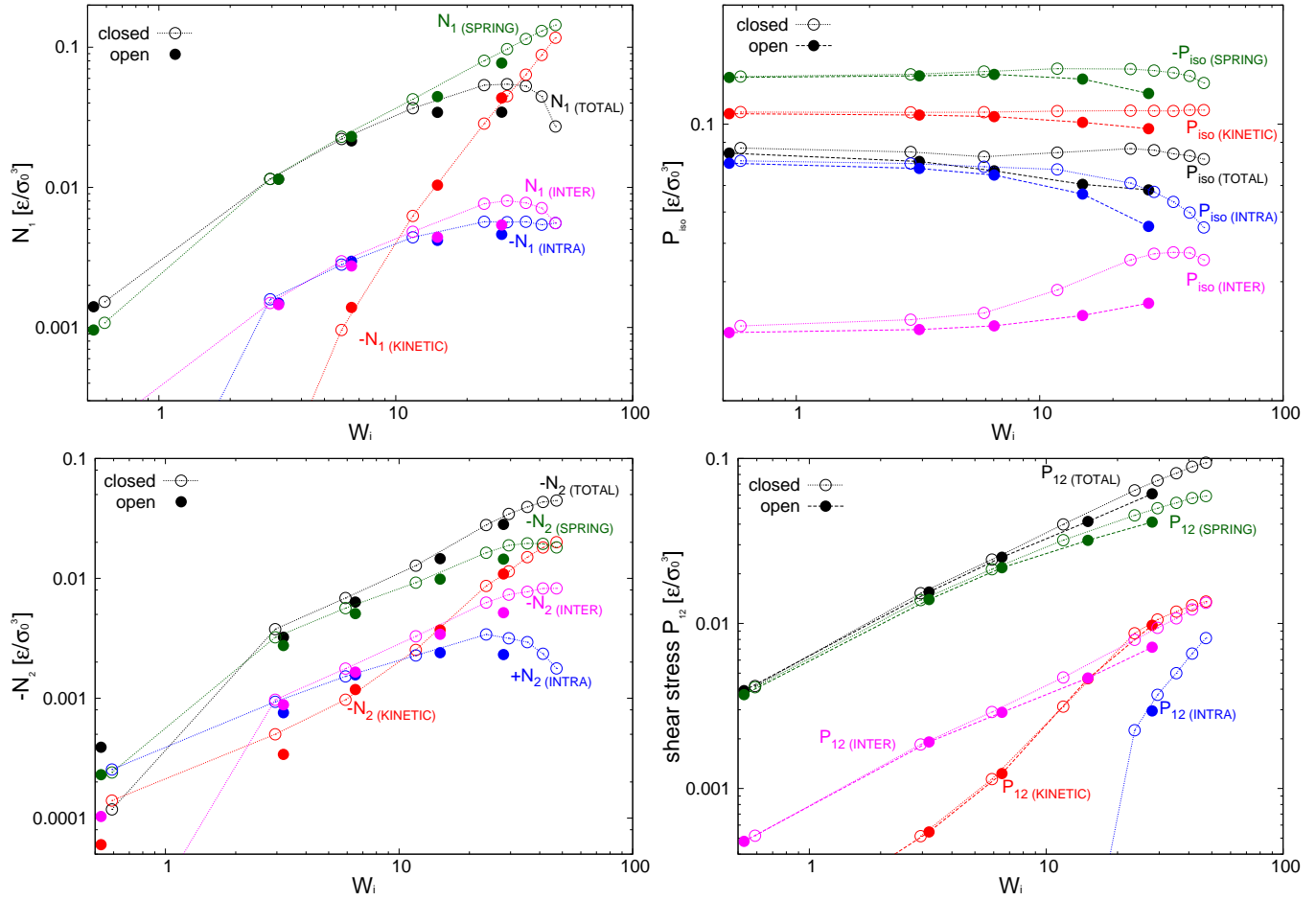


Fig. 10: Contributions of different mechanisms to the summing up the total normal stress differences, shear stress, and hydrostatic pressure. Results for adp thermostat under open and closed conditions. The error bars of the data in graphs are around $0.002 - 0.005\epsilon/\sigma_0^3$.

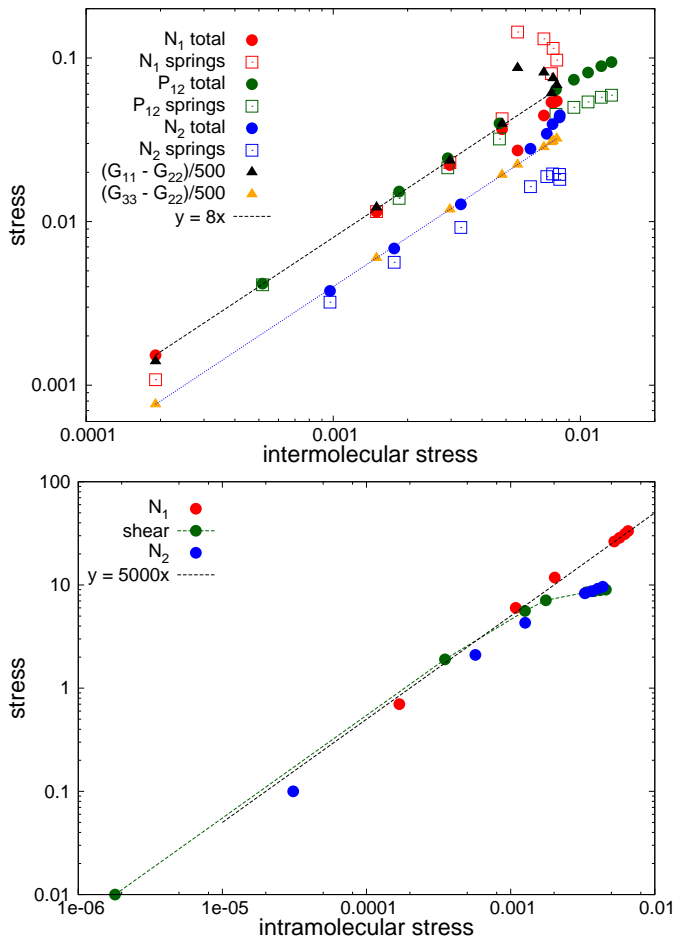


Fig. 11: Total stress, elastic stress components, and molecular strain differences in normal planes measured from the average gyration tensor ($G_{11} - G_{22}$ and $G_{33} - G_{22}$) plotted against the intermolecular stress component corresponding to each case (i.e., N_1^{intra} , N_2^{intra} , etc). Results for adpd (top panel) and for the tdpd model under open conditions (bottom panel). Estimated error bars of the depicted quantities are approximately 5%.

seemingly paradoxical, the “nominal” contribution to P_{12}^{intra} in the monomer pressure balance is minor). In all instances, at low enough shear ($Wi < 10$) the elastic stress is close to the total stress (and proportional to the intramolecular stress). This is the regime of validity of the stress optic rule which is broken at larger shear due to viscous (and compressible) effects related to the “simple fluid” of monomers. Remarkably, for the present model of star polymer melt, the kinetic normal stress becomes the dominant “viscous” contribution and at large $Wi > 20$ it even induces a decrease in the first normal stress difference N_1 .

7.3 Molecular ordering under shear

Fig. 13 presents the angle of the largest eigenvector of different contributions to the pressure tensor with the flow direction (measured according to Eq. 15). We also include the molecular orientation, measured from the angle associated to the gyration tensor (Eq. 15). This plot condenses what has been already mentioned in previous sections. Recall that a spherical molecular structure provides $\theta_G = 45^\circ$ and similarly from Eq. 15 a Newtonian fluid

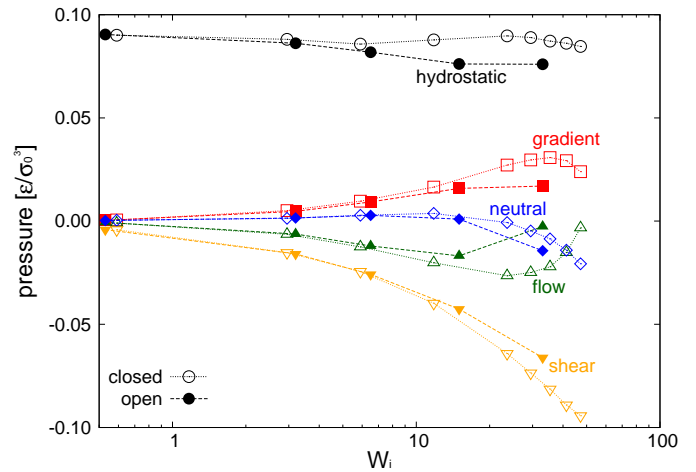


Fig. 12: Hydrostatic pressure and components of the traceless stress obtained in closed and open domains at fixed $T = 4$ and under the adpd thermostat (normal friction). Error bars of the measured quantities are approximately $0.005\epsilon/\sigma_0^3$.

without elastic component, necessarily presents $\theta = 45^\circ$. This is what is observed for the total pressure tensor angle θ_p and the molecular orientation θ_G at $Wi \rightarrow 0$ in Fig. 13. As the straining rate is made faster both angles decrease in a similar fashion, however, for $Wi > 10$, the pressure tensor angle θ_p presents a minimum and starts increasing towards 45° . By contrast, the molecular orientation θ_G keeps aligning with the flow direction. This is an indication of the loss of the Hookean behaviour of the melt which here is mainly due to the kinetic pressure (see its principal direction in Fig. 13). The springs stress direction also aligns with the flow, although its angle is larger than the molecular orientation (a similar outcome was observed in Ref.³). Finally, note the close match between the direction of intermolecular forces and the total pressure tensor. As stated, intermolecular forces are the driving mechanism of transformation between the viscous flow and elastic energy. Lastly, as observed by other authors^{39,42,43}, the open boundary does not modify the molecular structure or orientation with Wi , when compared with the closed case. Here, Fig. 13 show another remarkable result: the significant redistribution of pressure in the open case (see Fig. 12) does not alter the orientation of the different contributions of the pressure tensor at increasing shear rate. The orientation of pressure eigenvectors is a function of its eigenvalues which in turn determine the material properties of the polymer (its viscoelasticity⁸⁷). A relevant example is the the recoverable shear strain (SR) given by Eq. 14. *Material* properties should not depend on the constraints used to perturb the polymer and this precisely what our analysis provide.

To observe the collective order of the star molecular, we calculated the CoM pair distribution function $g(\mathbf{R}_{ij})$. Fig. 14 illustrates the marginal distributions $g_{2D}(X_\alpha, X_\beta) = \int g(\mathbf{R})dX_\gamma$ for different planes and at increasing shear rates. It is illustrative to draw the directions of the principal components of the pressure and the gyration tensor to observe the departure from the Hookean (linear SOR) regime. Above $Wi > 20$ molecules start to orient in lanes in the flow-gradient plane, as indicated by the elongated shape of the CoM distribution in Fig. 14. In linear polymers at

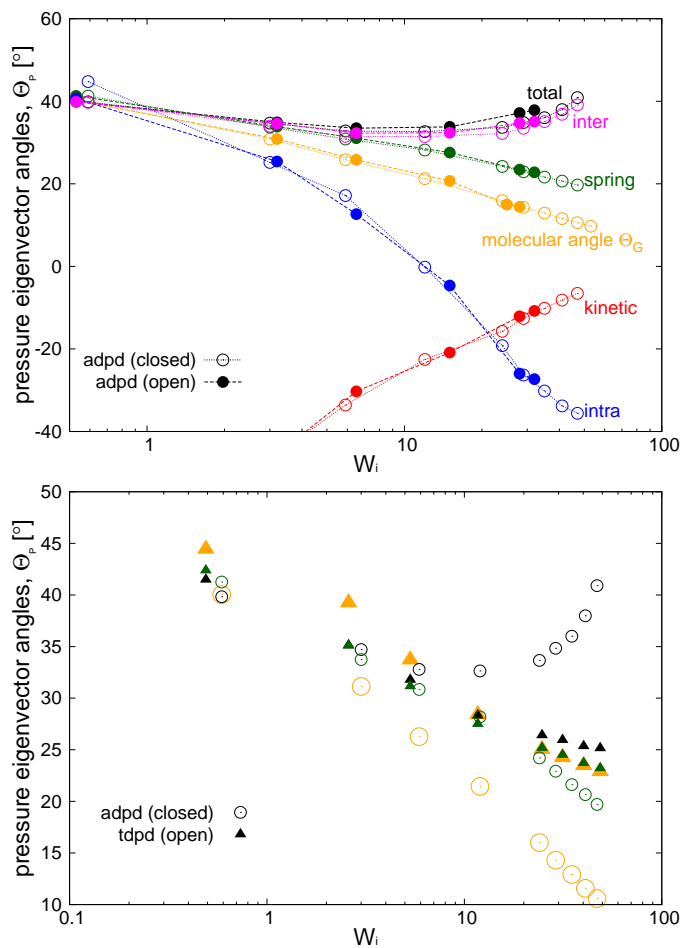


Fig. 13: Top panel: Angle between the principal eigenvectors of the contributions to the stress tensor and the principal eigenvector of the average gyration tensor. Bottom panel: Comparison with the model including transversal friction (tdpd model). Angles are with the same color coding.

large shear rate this effect creates order and even crystallization (Ref.⁸⁷). This collective order is also clearly visible in one snapshot of the system, in Fig. 15. Molecules with different relative velocities (closeby in the gradient direction) slide over creating a stress which is larger along a direction differing from the individual molecule's orientation. This direction of maximal stress is correlated with the CoM distribution (see Fig. 14) which shows bright spots where molecules slide over and depleted regions in the “wake” of each molecule. At the largest shear rates considered we also observe some collective ordering in the neutral (vorticity) direction. In sheared colloids, lanes of particles in the neutral direction appear due to hydrodynamic interactions¹⁰⁴. This could be a plausible hypothesis, in view of the hybrid colloid-polymer nature of star molecules. direction.

8 Effect of tangential friction

To investigate how the tangential friction between monomer blobs alters the rheology of the melt model we use a standard (i.e. not adaptive) DPD thermostat⁷⁰ with $\gamma_{\parallel} = 1$ and $\gamma_{\perp} = 1$ (see Secs. 3, 4, and 5.3 for details). As stated the friction kernels are Heaviside functions, in this case with cutoff distance $R_{dpd} = 1.5 \times 2^{1/6} \sigma$.

In the following we label this tangential friction thermostat as “tdpd”. This tdpd thermostat was found to be strong enough to keep the system's temperature relative increase smaller than 5% at the largest shear rates considered.

In a real system friction acts by reducing the relative velocities of interacting monomers, generally in the normal and also in the tangential directions. Under Markovian and pairwise interaction assumptions, this form of friction leads to the DPD equations as shown by the Mori-Zwanzing dynamic coarse graining applied to the microscopic Liouvillian dynamics⁹. The same effect is properly captured by the tdpd thermostat, although here at a qualitatively level. Reducing the monomers relative velocities immediately leads to a reduction in kinetic pressure which has large consequences in the system's rheology. In particular, the behaviour of the melt is essentially ruled by its elastic component, activated by the more effective intermolecular friction. Just to illustrate this point, we plot in Fig. 16 the contributions of the first normal stress difference (a direct measure of viscoelasticity) for the tdpd case. Compared with Fig. 10 (for the $\gamma_{\perp} = 0$ adpd case) the tdpd model has a much smaller kinetic pressure and N_1 is essentially determined by the elastic stress (particularly as Wi increases). The same conclusion applies to P_{12} in Fig. 16.

An interesting difference related to the presence of tangential friction concerns shear dilatancy. Fig. 17(a) presents the relative density expansion $\delta\rho/\rho_0 = 1 - \rho/\rho_0$ for different models. Let us now focus on the adpd and tdpd models which are kept isothermal (non-isothermal cases are discussed in Sec. 9). The density expansion of the adpd model (without tangential friction) scales like $\delta\rho/\rho_0 \sim Wi$, while tangential friction (tdpd) leads to a much softer trend $\delta\rho/\rho_0 \sim Wi^{0.5}$ (although it expands relatively more at moderate shear rates). Under a constant normal load, shear dilatancy is a consequence of the growth of pressure in the velocity-gradient direction. In the case of small kinetic effects (tdpd) this growth is controlled by the expansion force arising from the compressed springs. This elastic pressure appears as soon as molecules start to align with the flow and to compress in the gradient direction. Under enough tangential friction, the elastic stress is dominant and also controls the hydrostatic pressure, which in absence of kinetic pressure effects, presents faster decrease at large Wi compared to the adpd case (see Fig. 18). Of course, this decrease is also related to the fact that the tdpd simulations were done in the open system; notably for the tdpd we get about 50% reduction in hydrostatic pressure for less than 10% reduction in density (see Fig. 17).

The eigenvalues of the gyration tensor shown in Fig. 8 also indicate that adding tangential friction makes star molecules “stiffer”, in the sense that one needs larger values of the Weissenberg number to deform them. This observation is however somewhat misleading because for a fixed Wi , the real (physical) shear rate $\dot{\gamma} = Wi/\tau_{rot}$ is now smaller due to the increase in τ_{rot} with the friction. In any case, the tangential friction is expected to alter the stress-strain relations in the Hookean regime (related to the linear stress-optic rule coefficient). This is (indirectly) seen in Fig. 11 where we plot the normal strain differences $G_{11} - G_{22}$ and $G_{22} - G_{33}$ (also G_{12}) against the corresponding intermolecular stress differences (against P_{12}^{intra}). We choose this plot to il-

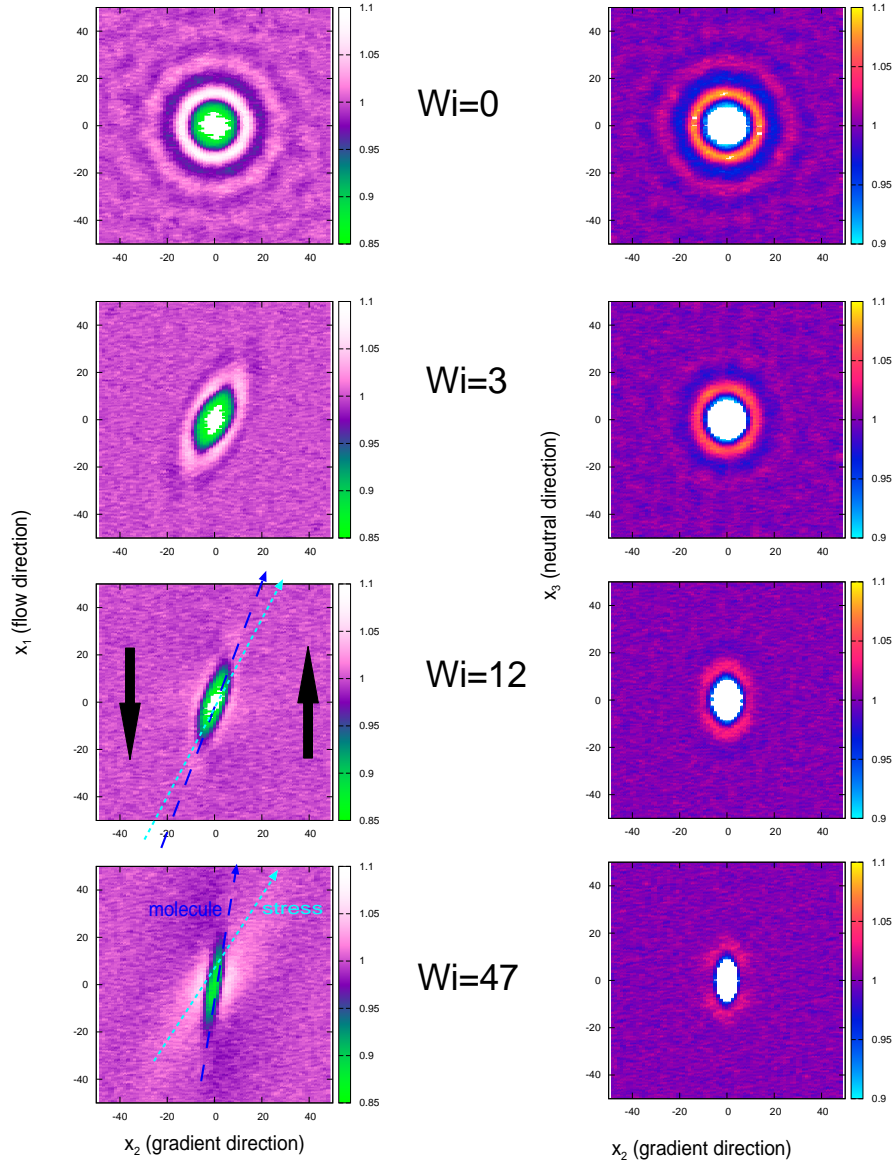


Fig. 14: The pair distribution of molecules, $g(\mathbf{R})$ providing the probability of finding the CoM of a molecule at a distance (vector) \mathbf{R} from the target molecule's CoM. Left: Marginal probability $g_{12}(X_1, X_2) = \int dX_3 g(\mathbf{R})$ in the flow-gradient plane and (right) in the gradient-neutral. Results for increasing Wi in a closed domain (open simulations at similar Wi are visually indistinguishable). Green line denotes the direction of the deviatoric stress and the blue line the molecular orientation obtained from the gyration tensor Eq. 15.

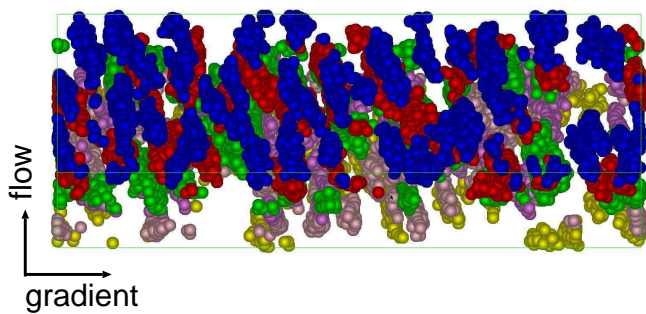


Fig. 15: Snapshot of a closed box simulation under shear rate ($Wi = 42$). For the sake clarity only 10 % of randomly chosen molecules are shown. Colors indicate position in the neutral direction. The snapshot clearly shows the formation of lanes tilted in the flow-gradient plane created due to sliding over rows of oriented molecules. Compared to Fig. 14, the snapshot is rotated for 180° with respect to the gradient axis.

illustrate two facts: first, if the kinetic pressure is minor, the intermolecular friction is the leading mechanism driving the molecular deformation (and its elastic response). Second, molecular strains (and elastic stresses, not shown in the figure) in the flow-gradient and gradient-neutral planes (corresponding to first and second normal stress differences) present a quite similar linear scaling (the shear deformation also) with the intermolecular stress. This is to be contrasted with the top pannel of Fig. 11 (model in absence of tangential friction), where the second normal stress (and $G_{22} - G_{33}$) is half of the first N_1 counterpart (also $G_{22} - G_{11}$). A perfect alignment between intermolecular forces, elastic stress and molecular strain was also found by Kroger⁸⁹ in linear FENE chains. It thus seems that tangential friction (tdpd) helps to reduce the second normal stress in such way that N_1 and N_2 present similar scaling laws $N_1 \sim \dot{\gamma}^{0.68 \pm 0.02}$ with $N_2 \sim -0.3N_1$. This is to be compared with the adpd case in Fig. 6 ($N_1 \sim \dot{\gamma}^{0.70}$ and $N_2 \sim \dot{\gamma}^{1.0}$). As shown in Fig. 7 the tdpd model yields $-N_2/N_1 \simeq 0.3$ for any $Wi < 100$. This value is characteristic of disentangled melts (being $-N_2/N_1 = 2/7$ the theoretical prediction for small shear rates^{1,88}).

The monotonous increase of elastic storage with $\dot{\gamma}$ found in the tdpd model is reflected in the recoverable shear strain (SR) shown in Fig. 7. Somewhat paradoxically, adding tangential friction increases the melt's elasticity. Albeit, this reinforces the conclusions in Sec. 6: the intermolecular friction is the principal mechanism loading elastic stress into an disentangled melt. In passing we note that in the tdpd model the orientational resistance parameter $m_G = Wi \tan(2\theta_G)$ grows like $m_G = 3.7Wi^{0.65}$ (at least for $Wi < 100$), a scaling which agrees with that reported for stars in solution¹⁰⁵ (the prefactor being however about twice larger in our melt.)

Not unexpectedly, the zero shear viscosity for the tdpd star model is larger $\eta_0 = 2.6$ than the adpd case $\eta_0 = 0.5$. The relaxation time is also larger $\tau_\eta = 287$ (compared with 60). However, the tdpd viscosity shear thins faster we find $\beta_\eta \simeq 0.5$ compared with 0.4 for the adpd case (recall $\eta \sim \dot{\gamma}^{-\beta_\eta}$). Again, this is also a consequence of a much smaller contribution of the viscous stress coming from kinetic effects. If the tangential friction is absent, the kinetic (and intramolecular) contribution increases the shear stress and the viscosity at any Wi leading to a softer shear thin-

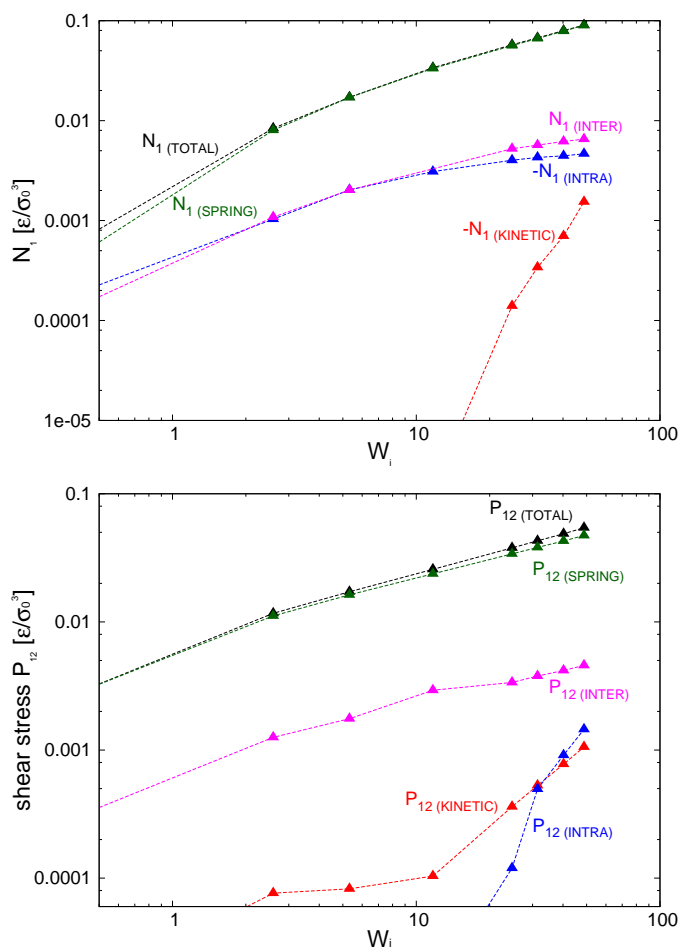


Fig. 16: Balances for the first normal stress difference and the shear stress for the tdpd model including tangential friction. Compare with Fig. 10 for the zero tangential friction case. Error bars of the measured quantities are approximately $0.002 - 0.005\epsilon/\sigma_0^3$.

ning exponent.

9 Thermostats and heat dissipation

9.1 Density and temperature

We now briefly analyze the effect of the temperature increase due to heat dissipation in the sample. In all cases, the temperature reaches a steady state, but as shown in Fig. 17 (bottom pannel) plotting $\delta T/T_0 = T/T_0 - 1$, we face severe viscous heating when performing the first row of simulations with the sdpdshort and the (stronger) sdpdlong thermostat (see Table 1). Heating is observed for $Wi > 10$ and irrespective of the damping parameter (we tried up to $\gamma_{DPD} = 50m_0\tau_0^{-1}$). Tangential friction drastically reduces heating ($\delta T/T_0 < 0.045$ for $Wi < 70$) however the adpd thermostat enabled us to simulate zero tangential friction at fixed temperature, providing $\delta T/T_0 < 0.01$.

Let us focus on the “heated” runs at increasing shear rate to illustrate the effect of an uncontrolled temperature. Fig. 17 shows that heating introduces further melt expansion under shear and this kinetic energy induces larger hydrostatic pressure (which it is seen to increase with shear in the sdpdshort case). Pressurization due to viscous heating can also alter the rheology response. This

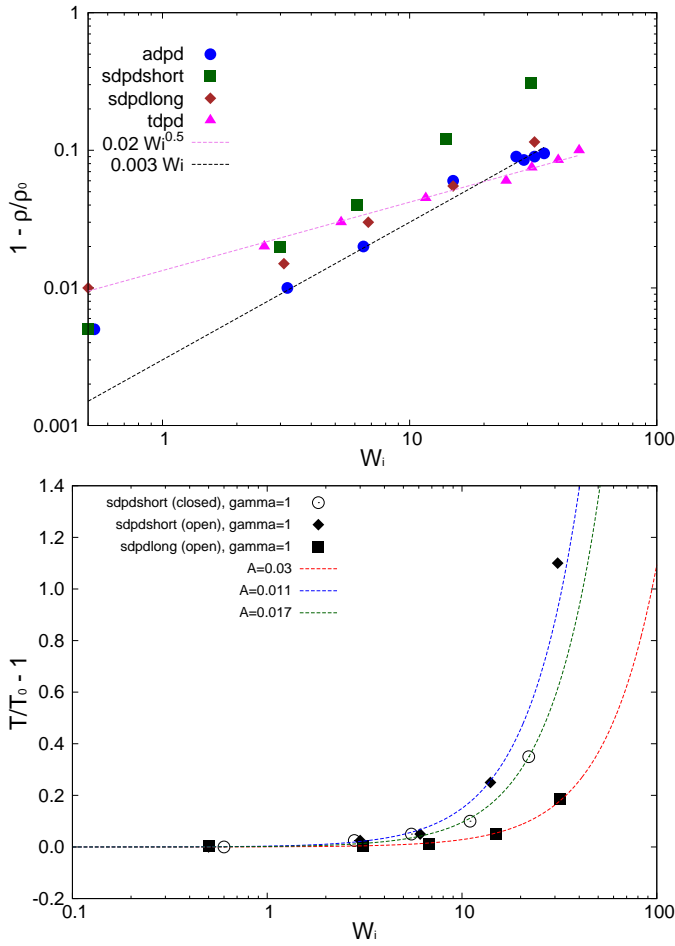


Fig. 17: Top panel: Relative density variation for the different cases studied (open and closed, and different models and thermostats, see Table 1). Bottom panel: Relative temperature increase observed in simulations with *standard* DPD thermostats with vanishing tangential friction. The relative variation of T for the adpd thermostat is smaller than 0.01 and less than 0.045 for the tdpd case (both of them not shown in the graph). Details of the thermostats are given in Table 1. The lines correspond to Eq. 21, with the characteristic constant A defined there. Error bar estimations of the data are approximately 0.005 and 0.01 for the upper and lower graph, respectively.

is seen in Fig. 5 where the sdpsdshort case present shear thickening for $Wi > 10$, but only *under closed conditions*. The shear thickening reported in some of the published works on polymer melts (closed box simulations) might in fact have been due to viscous heating. (see e.g. Ref.³⁹). More interesting than this elucubration is the result of Fig. 5 for the sdpsdshort-open. The viscosity obtained for the same thermostat in an *open* environment is *not* affected by the temperature increase with $\dot{\gamma}$. In fact it presents the very same trend as the adpd case. This observation indicates that shear viscosity is dominated by the normal load. In fact, the same outcome was also observed for the sdpsdlong model (with shear exponent $\beta_\eta = 0.39$) and for all $\gamma_{||} \leq 10$ considered. Thus, this insensibility of viscosity to temperature found only under normal load is not probably due to a cancellation of effects sometimes observed in experiments⁵⁰ (viscosity decreasing with T and increasing with P_{22}). Rather it should be due to the viscosity dominated by pressure as happens in highly pressurized melts⁵⁰. Here monomers

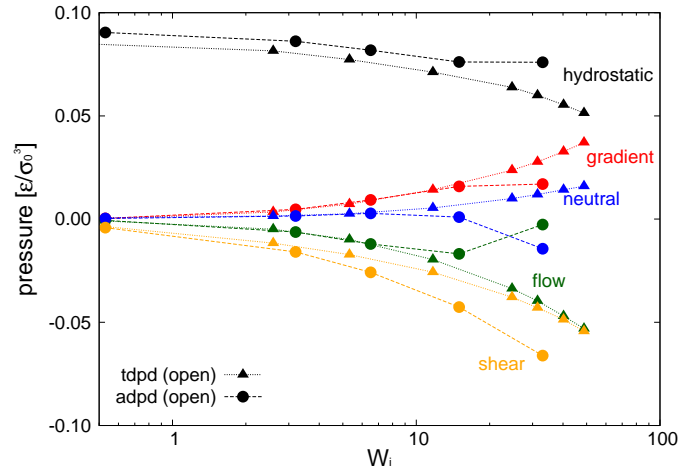


Fig. 18: Hydrostatic pressure and components of the traceless stress for the adpd model (normal friction between blobs) compared with the tdpd model (including tangential friction). Both cases in open boxes at fixed $T = 4$. Error bars of the measured quantities are approximately $0.002 - 0.005\epsilon/\sigma_0^3$

interact via purely repulsive forces (WCA) which might map a sample under large pressure, in fact, adding attractive interactions would probably trigger temperature effects on viscosity.

9.2 Viscous heating

The rate of heat production per unit volume due to viscous dissipation is $\dot{Q}_\eta = \eta \dot{\gamma}^2$ leading to a larger steady temperature whose value depends on the heat extraction rate. The onset of temperature increase is usually determined by a non-dimensional parameter which depends on Wi and on the rate of cooling \dot{Q}_c (see the recent computational study in Ref.¹⁰). Although in this work we shall not focus on heat and entropy productions, we believe it is interesting to share our observations on this phenomena, partly because of the relative small simulation literature accurately reporting heating effects in sheared, thermostatted melts. A simple equation for the heat produced in the sheared melt includes frictional gain and cooling,

$$\dot{Q} = \dot{Q}_\eta + \dot{Q}_c = \frac{\eta Wi^2}{\tau_{rot}^2} - c_X \alpha (T - T_0), \quad (19)$$

where c_X the specific heat capacity (molar) at constant pressure ($X = p$) or volume ($X = V$) and $dQ_c = \rho_n c_X dT$ (here ρ_n is the monomer number density). The DPD thermostat extracts (kinetic) energy upon pair collisions, at a rate which is proportional to the temperature difference $T - T_0$, where T is the system's (kinetic) temperature and T_0 the thermostat's nominal temperature[‡]. The value of cooling rate α (which has units of number

[‡] This can be proved from the equation for the time dependence of the covariance $\langle dv_i/dt(t) dv_i/dt(t') \rangle$ of the DPD Langevin's equation and can be easily checked (and the cooling rate α measured) upon observation of an exponential convergence of T towards T_0 after an instantaneous change (increase or decrease) of T_0 .

of colliding pairs divided by volume and time) should scale as

$$\alpha \propto \frac{1}{2} \rho_n^2 \int 4\pi r^2 g(r) w(r) \gamma_{DPD} dr, \quad (20)$$

where we note that $w(r)$ is the DPD kernel, which is simply a Heaviside function in our thermostat. The heat cooling rate of the thermostat increases with its damping coefficient γ_{DPD} , the kernel cutoff R_{cut}^{DPD} , and with the square of the monomer local density, which determines the number of thermalizing monomer collisions. In the steady state $\dot{Q} = 0$ we thus have,

$$\frac{T - T_0}{T_0} = \frac{\eta}{\alpha c_X T_0 \tau_{rot}^2} Wi^2. \quad (21)$$

Using the viscosity trend $\eta = \eta(Wi)$ obtained from simulations (see Sec. 5.3 above), we plot the prediction Eq. 21 in Fig. 17 (lines) for different cases considered. The agreement is quite reasonable, indicating that the temperature increase in the melt can be forecast using a simple thermodynamic argument. Best fits to Eq. 21 provide $A \equiv \alpha c_P = 0.011$ in the open-sdpdshort while somewhat larger $A = 0.017$ for the closed-sdpdshort. For the closed sdpdlong thermostat, which has about twice as much colliding partners within the dpd kernel, we consistently get $A = 0.03$. A preliminary calculation of c_X from the variance of the system's energy in a closed (NVT) equilibrium simulation $c_V = \langle \delta U^2 \rangle_{NVT} / (NT^2)$ provides $c_V \simeq 1.7$. From Eq. 21 we get the same order of magnitude ($A \simeq 0.05$ for sdpdshort and $A = 0.09$ for sdpdlong). A better agreement is found when comparing with the tdpd (sdpdlong) thermostat whose best fit provides $A = 0.06(5)$ against the prediction 0.09. This indicates that Eq. 20 should also depend on the number of degrees of freedom the thermostat acts upon (3 in the case of tdpd, 1 otherwise). A more refined calculation would also require including the dependence of c_X and $g(r)$ with the shear rate.

10 Comparison with previous studies

It is interesting to compare our results with previous rheological studies, some of them carried out at isobaric (constant P_{iso}) or constant load (P_{22}) constraints. As stated in the introduction, the number of studies of flowing melts under the constant pressure (either hydrostatic pressure or normal load) is not large. However, they present significant discrepancies on the density and pressure variations with shear. For instance, Ref.⁴² presents results for dendrimer melts under the isobaric condition (constant hydrostatic pressure) revealing a decrease in the melt's density under shear. For linear chains, Ref.⁴⁰ presents just the opposite effect (contraction under shear) while Ref.⁴³ (constant load) reports shear expansion (density increase).

The shear thinning exponents found here for a star polymer melt are consistent with those found in other simulations for somewhat similar systems such as hyperbranched and dendrimer polymers^{41,42}. Recall that the shear thinning exponent of any quantity Φ is β_Φ with $\Phi \sim \dot{\gamma}^{-\beta_\Phi}$ at large $\dot{\gamma}$. For viscosity we find $\beta_\eta \simeq 0.4$ (adpd) and $\beta_\eta \simeq 0.5$ (tdpd), for the first normal stress coefficient $\beta_{\Psi_1} \sim 1.30$ (adpd) and 1.31 (tdpd), while for the second one the values are $\beta_{\Psi_2} \sim 1.0$ (adpd) and 1.31 (tdpd). For dendrimers, Bosko et al.⁴¹ reports shear thinning exponents in-

creasing with M under NVT ($\beta_\eta \in [0.28 - 0.36]$) while, for $NP_{iso}T$ the were roughly independent on M ($\beta_\eta \in [0.37 - 0.39]$). The same work reports $\beta_{\Psi_1} \simeq 1.27$ and $\beta_{\Psi_2} \simeq 1.23$ under $NP_{iso}T$ while $\beta_{\Psi_1} \simeq 1.1$ $\beta_{\Psi_2} \simeq 1.0$ under NVT. Closed simulations for hyperbranched polymer melts⁴² predict $\beta_\eta \simeq 0.3$ (slightly increasing with molecular mass M) while $\beta_{\Psi_2} \simeq 0.95$ and $\beta_{\Psi_1} \simeq 1$ (both roughly independent on M). The scaling of (first and second) normal coefficients are probably sensible to the type of external constraint (either isobaric P_{iso} or constant load P_{22}). As an indication, a numerical study at constant load for linear chains⁴³ reports the same exponents found in this work ($\beta_{\Psi_1} = 1.35$ and $\beta_{\Psi_2} \simeq 1$ respectively). Concerning shear viscosity, if the density is allowed to relax (under isobaric or constant load conditions) the shear thinning exponents reported in the literature show much less variation with the molecular weight than under a fixed density. This has been shown for linear chains under constant load⁴³ and also for dendrimers⁴². In both cases, the shear thinning exponent in the variable density case was found to be close to 0.4, while it ranged from 0.25 to 0.4 in the closed system (increasing with molecular weight in both cases). This larger insensitivity of shear thinning exponents under fixed load conditions agrees with our observations in simulations presenting viscous heating mentioned in Sec. 9.

More recently, very similar papers^{106,107} studied the rheology and dynamics¹⁰⁸ of star polymers with different functionalities in solution. These groups find that the contribution of the stars to the sheared solution viscosity scales like $Wi^{-0.4}$ which is quite close to the shear thinning exponents we find here for the melt. According to these simulations^{106,107}, the first normal stress coefficient scales like $\Psi_1 \sim Wi^{-1}$ in solution, although the authors claim an exponent of $-4/3$ at very high shear rate $Wi > 100$. In melt, we observe the $-4/3$ power law at smaller shear rates. The second normal coefficient seems also to scale slightly differently in solution $\Psi_2 \sim Wi^{-4/3}$ than in melt $\Psi_2 \sim Wi^{-1}$. The strong similarities in solution and melt indicate that rheological properties are mainly ruled by conformational changes of the chains and maybe also that the hydrodynamic effects in melt are somewhat similar to those in a liquid solvent. Finally, it is interesting to note that the range of values for β_η and β_{Ψ_1} for stars, dendrimers and hyperbranched molecules are consistent with the theoretical calculation of Kroger⁸⁸ based on the Fokker-Planck equation for the bond vector distribution of multibead linear chains having slow rotational diffusion (compared with the entanglement relaxation) and a finite deformation energy⁸⁸. These two features are indeed consistent with the nature of short-armed stars, dendrimers and hyperbranched molecules with internal excluded volume interactions.

The number of experimental work on rheology of sheared star polymers is not abundant either, but it is growing fast due to the foreseen technological applications^{52,53}. Most of these works consider long armed stars which present entanglements. However, the shear thinning exponent we found for the tangential friction case $\beta_\eta = 0.5$ is quite close to the star case of the experiments by Tezel et al.¹⁰⁹ (4 arms stars with $M_a = 140\text{kg/mol}$; the lowest molecular weight considered in these experiments). Although entanglements in star polymers are not yet fully understood in

the non-Newtonian regime⁵², they should be responsible for the substantially increase in shear thinning (exponents close to 0.8) observed in the recent experiments with star molecule melts of Snijkers et al.⁵². Shear thinning exponents found in melts of linear chains are around $\beta_\eta = 1$ (predicted by reptation theory). An open question is why star molecules significantly reduce the shear thinning⁵².

The analysis presented in Sec. 6 based on the (exact) balance of the pressure coming from the different contributions (kinetic, intramolecular, intermolecular and bonds) provides some clues which can be useful in a broader study. These type of analysis were performed by Baig et al.^{2,3}, Matin⁴³ and Tschopp⁴⁵. However these balances were based on the energy budget which has less direct rheological consequences than the stress. Also, surprisingly, the kinetic contribution was not considered (leading to unbalanced analyses). The sole exception we found is the work of Kroger⁸⁹ for linear FENE chains which, like ours, is based on the exact stress budget. In elongational flow of linear chain melts, Kroger⁸⁹ found that viscous effects came mainly from intramolecular collisions (he called it “simple fluid” stress offset). Here, we find that in star molecules the kinetic pressure can be the dominant viscous contribution to the melt rheology at large shear rates (a possibility in fact recognized by Kroger in his book⁴). The kinetic pressure is however reduced with the tangential friction and this warns about the need of dynamic coarse-graining⁹ to represent a realistic model. While linear chains tumble in shear flow¹¹⁰, star molecules perform a quite different motion called tank-threading^{102,105} (whereby arms rotate around the molecule’s center). One can speculate that the reduction of shear thinning observed in star melts is due to an enhanced kinetic pressure related to the tank-threading. Above $Wi > 10$ we do observe the tank-threading motion in our melt which, with monomer angular momentum growing much faster ($\omega \sim \dot{\gamma}^{0.6}$ from preliminary results) than it does in stars in solution (where ω is constant^{102,111}). Concerning intermolecular interactions, quite often neglected in the literature (see e.g. Ref.⁸⁷), we find that these are key to transfer the externally imposed stress into molecular strain and elastic stress. This observation agrees with Kroger’s picture⁸⁹. Intermolecular *friction* is the sole possible mechanism if entanglements are not relevant (like in our short arm star molecules).

11 Conclusions

We have conducted OBMD simulations of the melt of star polymers (73 monomers, 12 arms of 6 monomers per arm) using a relatively new modelling technique which combines the adaptive resolution and open-MD (respectively, introduced in Refs.²² and⁴⁶ and used in many other studies). From a technical point of view, this work substantially enlarges the size of molecules exchanged between the open system and the reservoir¹³. In this work, we however focus on what happens if a melt is sheared under a constant normal load instead of a constant volume. The constant load is in fact the condition in many experiments (see Refs.^{40,44}) although the simulation literature on the subject is not abundant. We have also presented some conclusions on how the tangential and normal monomer-monomer frictions affect these

CG molecular models.

The OBMD permits to perform several new features in a molecular simulation. At equilibrium, the OBMD correctly represents the mass fluctuations of the grand canonical ensemble and interestingly, it could permit to study how fluctuations and sound velocity are altered under non-equilibrium (e.g. sheared) states. OBMD also allows to impose the external shear stress P_{12}^{ext} as required for (flux based) hybrid continuum-MD simulations^{15,66}. It could also enable the validation of theories like Extended Thermodynamics⁴⁷ predicting different outcomes for the “conjugate” constant-stress and constant shear rate $\dot{\gamma}$ non-equilibrium ensembles.

Concerning the present study, we observe that under a constant normal pressure, the melt expands when sheared (shear dilatancy) leading to substantial depressurization and slightly decreasing the shear viscosity. This behaviour was observed in most previous simulations on sheared melts, but surprisingly it is still unclear if it is the general (universal) trend (e.g. see⁴⁰). This study provides new information on the rheology of sheared melts: notably, we see that the type of monomer’s friction is a key aspect for the system’s rheology. From a theoretical standpoint⁹, friction between monomer (or rather “polymer blobs”) arises as soon as one consider a CG view of the complex molecule (which is the standard case in polymer science). The pressure balance analysis reveals that in absence of tangential friction, the monomer’s kinetic stress becomes significant at large shear rates, increasing the system viscosity (reducing the shear thinning exponent) and diminishing its elastic response (e.g. normal stress difference, stress recovery). We also observed viscous heating in some simulations (e.g. using the sdpdshort and sdpdlong thermostat) revealing a viscosity jump (shear thickening) in closed systems. By contrast, the viscosity of the heated (and less dense) open samples did not changed in trend (shear thinning) under a constant load. This indicates that the melt’s viscosity is controlled by the normal pressure, at least for the present type of molecules with purely repulsive interactions and no significant entanglements. We expect the OBMD to become useful in other studies, such as adding the energy transfer⁶³ or extending the incompressible coupling in Ref.¹⁰ to perform hybrid simulations of compressible melts including the transfer of dissipated heat through/across system’s boundaries. Thus the properties related to heat conduction could be investigated¹¹².

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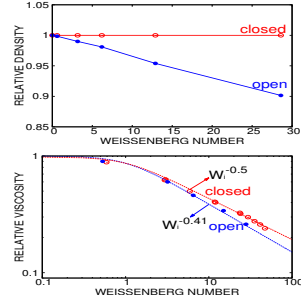
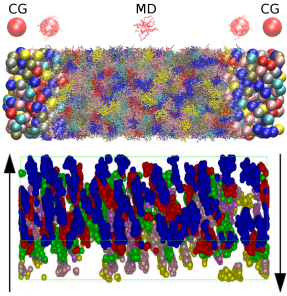
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Shearing polymer melts at constant normal pressure produces different rheology than shearing at constant volume, as revealed by AdResS-enabled open boundary MD simulations of star polymer melts.