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Generation of well relaxed all atom models of stereoregular polymers: a validation of hybrid particle-field molecular dynamics for polypropylene melts of different tacticities

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ABSTRACT

The tacticity of vinyl polymer chains strongly affects the physical properties of polymeric materials, as example the chain conformations and stiffness. In the present work we tested how the hybrid Particle-Field Molecular Dynamics (PF-MD) technique is capable to describe conformational differences of polymer chains as function of the tacticity. In particular, we focus on tacticity effect of atactic, isotactic, and syndiotactic polypropylene (PP) homopolymer melts. We found that PF-MD simulations exhibit dependence of Flory's Characteristic Ratio from the fraction of racemo diads along the PP chains in qualitative agreement with Small Angle Neutron Scattering (SANS) experiments and theoretical previsions. Finally, we calculated and compared the packing length parameter on very high stereoregular syndiotactic PP systems with rheological measurements. A qualitative agreement between the calculated and experimental packing length is found.

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Tacticity; polypropylene; hybrid Particle-Field; atomistic model; molecular dynamics; packing length; Flory's characteristic ratio

Introduction

One of the crucial aspects determining the physical properties of vinyl polymers is the tacticity of the chains. A vinyl polymer chain has along its backbone a sequence of methylene (CH₂) and pseudo asymmetric methyne groups (-CHR). The stereoregularity of a vinyl polymer can be defined based on arrangement of adjacent pseudo asymmetric centers. If the two pseudo asymmetric centers, once a given direction along the polymer chain is assigned, have the same stereochemical configuration the diad (composed of two adjacent repeating units) is called meso (m). Instead, the diad is called *racemo* (r) in case of opposite stereochemical configuration of chiral centers (Scheme 1a). The tacticity of a vinyl polymer can be usually measured in terms of diads (or sequence of diads) distributions by using the nuclear magnetic resonance spectroscopy (13 C-NMR) technique.[$^{1-3}$]

Tuning the properties of the material, by selecting appropriate tacticity, allowed the stereoregular polypropylene (and more in general vinyl stereoregular polymers) to become one of the most relevant materials of the

polymer industry.^[4,5] For semicrystalline polymeric materials, of course, tacticity is one of the key features determining most of the properties involved in several applications. Even in the case of fully amorphous materials or polymer melts, the tacticity plays an important role in influencing the final material behavior. In particular, in this case, tacticity is effective in the behavior of several properties such as: glass transition temperature (T_{σ}) ,^[6–8] self-diffusion coefficient (D),^[8] entanglement molar mass (M_e) ^[9] and other rheological properties.^[10,11] Since tacticity mainly influences the torsional angle distributions of the polymer backbone, the main molecular effects on single chain properties are connected to changes in chain stiffness. For this reason, both experimental and theoretical studies have been conducted to understand the effect of tacticity on the stiffness of polymers. The chain stiffness can be quantified by the Flory's characteristic ratio (C_{∞}) or by the Kuhn length (b). Higher values of C_{∞} correspond to stiffer chains. The stiffness of polypropylene (PP) as function of the tacticity was experimentally estimated by measuring the differences in $T_{g}^{[7]}$ intrinsic viscosity in Θ solvent,^[12,13] and chain dimension by SANS measurements.^[14,15] From the theoretical point of

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Scheme 1. (a) Chemical structure of *racemo* (r) and *meso* (m) diad of PP. (b) Assignment of coarse-grained density to lattice points for a PP segment's chain (for clarity only the backbone atoms have been used for the assignment). (c) Criterion for the assignment of a fraction of a particle of type K to mesh vertexes numbered from 1 to 4.

view, the Rotational Isomeric State model (RIS) applied to an unperturbed single chain was used to estimate the effect of tacticity on the PP chain dimension and conformational energy.^[16–18] Antoniadis *et al.* reported a Molecular Dynamics (MD) study to address the effect of tacticity on the dynamics of PP chains in melt state.^[19,20] More recently, Tzounis *et al.* studied the effect of tacticity on the conformational properties of PP homopolymer melts and poly(ethylene-propylene) block copolymers by using single-chain Monte Carlo (MC) technique.^[21] More generally, MD and MC techniques applied to atomistic and coarse-grained (CG) models were proposed to investigate the effect of tacticity on several physicochemical polymer properties.^[18–26]

It is worth noting that one of the problems in studying polymer melts by MD or MC simulation techniques is to obtain well relaxed configurations at the equilibrium density.^[27-29] This target, due to the large relaxation times, is still out of reach of atomistic MD simulations for chain lengths typical of industrially relevant polymeric materials. Moreover, due to the large number of atom overlaps observed at the typical melt densities (which may cause divergence in the calculation of the potentials and forces), the generation of initial configurations, even for short oligomers, can be very difficult.^[27-30] Recently, some of us introduced a suitable and validated procedure based on the hybrid Particle-Field Molecular Dynamics based on a combination of MD with the Self Consistent Field theory (named in the following MD-SCF), to overcome these technical problems.^[30] In particular, this procedure was successfully applied to study different homopolymer melts and composite polymer materials at atomistic resolution and at CG level.^[30-34] However, the MD-SCF method has not been tested to study the effect of tacticity on the structural properties, such as Flory's characteristic ratio, of stereoregular homopolymer chains in melt state. This application represents a good test case for the proposed methodology, not only for the relevance of the systems, but also, from a more technical point of view, to understand if the effects of subtle differences in the chain microstructure can still be modeled using an approach based on combination of particle models with density fields.

The aim of the present work is to investigate if and how MD-SCF methods can reproduce structural properties of polymers with different tacticity and compare the results with reference atomistic MD simulations and experimental data. Such a study can provide a validation of the proposed approach and open application to further systems based on stereoregular polyolefins, which represent the largest class of industrial polymers produced by chemical industry. We have chosen PP melts as a test case because they have been deeply investigated by experiments and molecular dynamics simulations. In the following, a brief introduction to the hybrid MD-SCF method is given in the Computational Methods and Models Section. In the Results Section MD-SCF simulations are compared to reference MD and experimental data. Finally, in the Conclusion Section the main results are summarized.

Computational Methods and Models

Hybrid MD-SCF

A brief description of the method will be given in this section, while the full description and derivation can be found in Ref.^[30] As mentioned in the Introduction, the MD-SCF approach applied in the present work has been validated and widely employed in many previous works to get equilibrated polymer melts and models of polymer composite materials on both atomistic and coarse-grained resolution.^[32,34,35] The core feature of the method is the evaluation of non-bonded forces and potentials (the most computationally expensive terms in a MD simulation) through the calculation of an external potential depending on the local density at position $V(\mathbf{r})$.^[36] The manybody problem in the spirit of Self Consistent Field (SCF) Theory can be reduced to a problem of deriving the partition function of a single particle in an external potential $V_{\rm K}(\mathbf{r})$.^[36] The non-bonded forces acting on a particle of type K are calculated from a suitable expression of $V_{\rm K}$ (r) and of its derivatives. In the SCF scheme, a particle interacts with its own surrounding through a mean field. With this assumption, the density-dependent interaction potential *W* can be written as follows:

$$W\left[\left\{\phi_{K}(r)\right\}\right] = \int dr \left(\frac{k_{B}T}{2} \sum_{KK'} \chi_{KK'} \phi_{K}(\mathbf{r}) \phi_{K'}(\mathbf{r}) + \frac{1}{2\kappa} \left(\sum_{K} \phi_{K}(r) - 1\right)^{2}\right)$$
(1)

where $\phi_K(r)$ is the coarse-grained density of a particle of type K at position r, and $\chi_{KK'}$ are the mean field parameters describing the interaction between a particle of type K and the density field due to particles of type K'. It can be demonstrated, by using the saddle point approximation that the external potential $V_K(r)$ can be written as:

$$V_{K}(r) = \frac{\delta W[\phi_{K}(r)]}{\delta \phi_{K}(r)}$$
$$= k_{B}T \sum_{K'} \chi_{KK'} \phi_{K'}(r) + \frac{1}{\kappa \left(\sum_{K} \phi_{K}(r) - 1\right)}$$
(2)

The derivation of Equation 2 and the details of the implementation of the MD-SCF approach are reported in the Refs.^[36,37] To connect the particles and the field it is crucial to obtain a smooth *coarse-grained* density function $\phi_K(r)$ from the position of the particles. To this aim, and to obtain both the coarse-grained density function and its density derivatives (needed to calculate the forces acting on the particles) a mesh-based approach was used. In a symbolic form this procedure can be written as follows:

$$\overline{S}\left\{\widehat{\phi}(r;\Gamma)\right\} = \phi(r) \tag{3}$$

where \overline{S} denotes the mapping from particle position to the coarse-grained density $\phi(r)$ and Γ represents the positions of all particles in the system. The density field is obtained by mapping particle positions on a density mesh. In particular, the simulation box is divided into sub-cells. Then, according to the particle positions inside a cell, a fraction of the particle is assigned to each vertex of the sub-cell. A simplified two-dimensional case showing this procedure is reported in panels B and C of Scheme 1. A fraction of the particle, proportional to the rectangle areas (panel C of Scheme 1), is assigned to each lattice point. The coarse-grained level of the density is tuned by the mesh size l. As example, for a large *l* value several particles are included in each cell, thus determining a coarser description of the density. Once the coarse-grained density is calculated from the particle coordinates and assigned to all mesh points, the spatial derivatives of the density fields are calculated by numerical interpolation. Then, forces acting on particles can be calculated. In the present work we used the so-called "stencil" method to calculate the density gradient. Such a method is based on the Taylor series developed by Alfaraj et al.-^[38] Recently, the stencil method has been implemented in the MD-SCF by Sevink.^[39] Further details about the derivation of the potential in the Equation 2 are reported in the Supporting Information, while a complete description of the algorithm used for the interpolation of the density gradient can be found at Ref.^[39]

Depending on the coarse-grained level of the density, a frequency Δt_{update} is set to update the density field from a new coordinate set at time $t + \Delta t$. For all MD-SCF simulations reported in this work, the frequency update Δt_{update} was set to 0.1 ps, similarly to the value used for equivalent systems.^[30]

Atomistic Model

The OPLS-AA^[40] force-field was used for the traditional MD atomistic simulation, which serves as a reference simulation for the different PP melts. The functional form and parameters of the harmonic bond stretching, angle bending, and torsions can be found in Ref.^[40] The intramolecular interactions, including bond, angle, and torsional terms of the MD-SCF model are coincident with those of the OPLS-AA.^[40] All bonded and non-bonded parameters are reported in the Supporting Information.

Computational Details

All hybrid MD-SCF simulations were performed in the *NVT* ensemble at the constant temperature of 473 K, which was controlled by the Andersen thermostat^[41] with a collision frequency of 7 ps⁻¹. Three different mesh sizes (l = 0.2, 0.4, and 0.8 nm) were employed. The time step for particle displacement using velocity Verlet algorithms has been set to 1 fs. The OCCAM code^[42] was used for all hybrid MD-SCF simulations.

Reference atomistic simulations performed with standard MD technique were carried out in the *NPT* ensemble by using the GROMACS package (ver. 5.0.1).^[43] The temperature was set equal to that of MD-SCF simulations (T = 473 K) and controlled by the Berendsen thermostat^[44] ($\tau_T = 0.1 \text{ ps}$). The target pressure was held constant at 1 atm with the Berendsen barostat^[44] ($\tau_P = 1$ ps). A cutoff distance of 1.2 nm was used for both van der Waals and Coulomb interactions. The long-range correction to the electrostatic interactions was calculated using the Ewald summation algorithm.^[45] A time step of 2 fs was used for all reference simulations. The bonds involving hydrogen atoms were constrained by using the LINCS algorithm.^[46] System compositions and details are reported in Table 1.

Multi-Stage Relaxation Procedure

The strategy to relax PP melts by MD-SCF approach is based on an increase of the model resolution (tuning the mesh size *l*) to equilibrate stepwise by successive equilibrations starting from low (large value of *l*) to finer resolutions (small values of *l*). The procedure is intended to recover an increasing number of structural correlations from each stage. The advantage of this method is that the model resolution can be tuned simply by changing the mesh size and does not involve the parameterization or reconstruction of the degree of freedom as it is standardly done in coarse-grained model based on particle reduction.^[30] The steps of the procedure applied in the

 Table 1. System composition of reference MD (50 ns long) and

 MD-SCF simulations.

			Box length		
	No. PP	No. rep. units/	(x = y = z)	Fraction	Fraction
System	Chains	chain	[nm]	of r	of rrrr
a-PP-	27	50	5.2	0.5	-
50					
<i>i-</i> PP	27	50	5.2	0	-
s-PP	27	50	5.2	0.4	-
sPP	45	200	9.455	0.74	0.47
	45	200	9.455	0.83	0.60
	45	200	9.455	0.87	0.71
	45	200	9.455	0.95	0.91
	45	200	9.455	0.98	0.93

present work are listed in the following: (i) The initial configurations of the systems are built by randomly placing the center of mass (c.o.m.) of each PP chain in the simulation box. The box size and the number of chains is chosen according to the experimental density of the PP melt at 473K. (ii) Three subsequent MD-SCF simulations at increasing density field resolution (reducing the mesh size *l* from 0.8 to 0.2 nm) are performed. In the first stage of the procedure the chains are described by a very coarse density (l = 0.8 nm, almost 5.3 times larger than the C–C bond distance in the backbone) which allows to equilibrate the chains c.o.m. positions and to pack them in a proper manner (further details in the Supporting Information). The use of coarser density field allows for faster simulations, in terms of both the intrinsic dynamics and the computational effort.^[30] (iii) Configurations obtained from the highest density field resolution (l = $0.2 \text{ nm} \sim 1.3$ times the backbone length) are used in the final stage of the procedure where non bonded Lennard-Jones potentials and Coulomb interactions are employed for a short relaxation.

To show in a pictorial way how the mesh size *l* changes the polymer chain model resolution, the density field, reported as isosurfaces that describe regions with the density (ρ) > 3.5 atoms/nm³, is reported in Figure 1. From the figure it is clear that, with a resolution of *l* = 0.8 nm, the bulkiness of the chain backbone is described, while, at higher resolutions (from *l* ≤ 0.4 nm) the bulkiness of the small CH₃ lateral group emerges. In the inset of Figure 1 the details of the density field calculated at *l* = 0.2 nm are compared with the atomistic structure. As can be seen, the shape and the details of PP chain reproduced by density field are extremely detailed in comparison with the fullatom resolution.

It is important to underline that no parametrization is needed to generate the well relaxed structures of different polymers by using the described procedure (for homopolymer melt $\chi = 0$) as reported in the Ref.^[30]

Results and Discussion

Short PP Chains

The first set of systems investigated is composed of 27 PP chains (N = 50 monomers, Table 1). Each system is composed of chains characterized by a Bernoullian distribution of diads with the following fractions of *racemo* r diads: $P_r = 0$ (isotactic *i*-PP), 0.5 (atactic *a*-PP) and 1.0 (syndiotactic *s*-PP). For each of these systems we performed MD-SCF simulations at constant temperature of 473 K. According to the multi-stage relaxation procedure reported previously, 10 million steps/stage have been



Figure 1. Density isosurfaces calculated from density field generated from a single PP chain (N = 50) at different mesh size (*l*) lengths. The inset showing atom details is reported together with the density isosurface calculate for l = 0.2 nm. All isosurfaces correspond to a density of $\rho > 3.5$ atoms/nm³.

performed for each system and the trajectory produced was used for the analysis.

The stiffness of a polymer chain can be expressed by the Flory's characteristic ratio C_n (Equation 4).

$$C_n = \frac{R^2}{nl_b^2} \tag{4}$$

where R^2 is the mean-square end-to-end vector distance of PP chains, *n* is the number of backbone bonds and l_b is the bond length (0.1529 nm). The C_n values calculated at each stage of relaxation procedure (different mesh size *l*) by using the MD-SCF technique are reported in Table 2. In Figure 2, the calculated C_n values from MD-SCF and reference MD simulations are compared with experimental values measured with SANS reported by Fetters *et al.*^[12,14,15], RIS calculations from Suter,^[16,18] Alfonso^[47] and Allegra,^[48] and MC simulations of a single unperturbed chain reported by Tzounis.^[21]

As a general behavior, the stiffness of PP chains calculated as Flory's characteristic ratio C_n is well

Table 2. C _n	, values	for	each	P_m	distribution.
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		Mesh size /		
System	N (monomer)	(nm)	Pr	Cn
РР	50	0.8	0.0	3.7 (0.2)
			0.5	4.8 (0.2)
			1.0	4.1 (0.2)
PP	50	0.4	0.0	4.2 (0.2)
			0.5	4.8 (0.2)
			1.0	4.7 (0.2)
PP	50	0.2	0.0	5.3 (0.1)
			0.5	4.3 (0.2)
			1.0	4.5 (0.2)
PP (Reference MD)	50	-	0.0	5.8 (0.4)
			0.5	6.9 (0.2)
			1.0	8.6 (0.3)

reproduced by MD simulations after the relaxation procedure, based on MD-SCF. This is apparent from the comparison of the results from the last stage of procedure (black square in the figure) with SANS measurements, RIS calculation^[16,18,47,48] and MC simulation of single unperturbed chain.^[21] The final stage of the relaxation procedure requires approximatively 350 ps



Figure 2. Comparison of C_n , as function of *racemo* diad content P_r, between: MD-SCF simulations (l = 0.2 nm blue open circle, l = 0.4 nm light blue circle, l = 0.8 nm purple blue circle), SANS measurements from Fetters^[12,14,15] (open red square), RIS calculation from Alfonso^[47] (blue line), Suter^[16,18] (orange line), Allegra^[48] (dark green line) and MC simulation of single unperturbed chain from Tzounis^[21] (green diamonds). The final stage of the relaxation procedure consisting in a short relaxation (350 ps) by MD simulation is reported with filled black square.

to fully allow the C_n values to become indistinguishable from long independent MD simulations. In particular, the C_n for *a*-PP is slightly stiffer than the experimental value, while the *s*-PP stiffness is slightly lower than SANS data.^[12,14,15]

By considering the results obtained from the stages of the procedure involving the MD-SCF simulations, no difference between C_n values have been found for syndiotactic (*s*-PP), atactic (*a*-PP), and isotactic (*i*-PP) in the whole explored tacticity range (see Figure 2).

In addition to the C_n evaluation, the structural correlation reproduced by the MD-SCF model at different resolutions (from coarser l = 0.8 nm to finer l = 0.2 nm) can be evaluated by comparing the backbone-backbone intramolecular radial distribution function (RDF) with the reference MD simulations (see Figure 3).

As can be seen from Figure 3, the intermolecular RDF of backbone-backbone with mesh size l = 0.8 nm does not match reference MD simulations at low *r* values. Instead, with a mesh size $l \le 0.4$ nm (see panel B, C, D, of Figure 3) the RDFs gain more details approaching to the correlation obtained in standard MD simulations, as found also in several previous works.^[28,30–32,34,49] However, the small

difference between the RDF calculated for different tacticity, which is kept by reference MD simulation and in agreement with previous studies,^[50] is not completely reproduced by MD-SCF simulations (see panel A of Figure 3). The larger deviation from MD is observed for *s*-PP (panel B of Figure 3). In particular, with the mesh size l = 0.2 nm, the position of first peak of RDF is shifted at higher *r* values of ~0.2 nm with respect to the MD reference simulation. In case of CH₃ substituent group, the RDFs calculated from MD-SCF simulations with a mesh size $l \leq 0.4$ nm show similar correlations for different tacticity (Figure 4b–d), as also found in the case of reference MD simulations (panel A of Figure 4).

In both cases, i.e., backbone-backbone and CH_3-CH_3 , the first and most intense peak in the RDF distributions has a very local nature and can be recovered only using a pair potential. We demonstrated that a short relaxation is required to fully recover this local pair correlation.^[28] In fact, as reported in Figure 5, a relaxation time of about 70 ps is enough to obtain indistinguishable RDF distributions for the backbone-backbone pair for all investigated tacticity range. In particular, starting from the last configuration of MD-SCF with mesh size l = 0.2 nm, the longest relaxation



Figure 3. Comparison of backbone–backbone intermolecular RDF distribution: (a) Reference MD simulations for *s*-PP (red line), *a*-PP (black line) and *i*-PP (light blue line). (b) Reference MD of *s*-PP versus MD-SCF simulations. (c) Reference MD of *i*-PP versus MD-SCF simulations. (d) Reference MD of *a*-PP versus MD-SCF simulations.



Figure 4. Comparison of CH₃-CH₃ intermolecular RDF distribution: (a) Reference MD simulations for *s*-PP (red line), *a*-PP (black line), and *i*-PP (light blue line). (b) Reference MD of *s*-PP versus MD-SCF simulations. (c) Reference MD of *i*-PP versus MD-SCF simulations. (d) Reference MD of *a*-PP versus MD-SCF simulations.



Figure 5. Comparison of intermolecular RDF for the backbonebackbone pair between reference MD (black line), MD-SCF (blue circle), and after: (a) 65 ps for *s*-PP, (b) 10 ps for *i*-PP, (c) 10 ps for *a*-PP.

time of 65 ps is required in case of *s*-PP (see panel A of Figure 5), while, 10 ps are sufficient for *a*-PP and *i*-PP (see panel B and C of Figure 5). For CH_3 – CH_3 , a relaxation time of 10 ps is sufficient to fully recover the short-range correlations (more details in the Supporting Information).

According to the results discussed in this section, it is possible to obtain for the proposed procedure results practically equivalent to models based on traditional MD simulations without the limitations related to the slow relaxation times.

Long PP Chains

In order to study and to properly include the effects of tacticity in our models it is necessary to consider high molecular weight (M.W.) models for different reasons.

The main reason is that, to model highly stereoregular (but not fully stereoregular) polymer chains, as for example a high fraction of diads [r] and pentads [rrrr] close to one but not fully syndiotactic, a long pseudo asymmetric center sequence (and hence a high molecular weight) is needed. The MD-SCF method well fits these requirements due to the fact that the mean-field approximations are more accurate for polymer systems with large molecular weights.^[30] In particular, in addition to the mesh size l and density update frequency, the fluctuation of the collective density depends on the invariant degree of polymerization $\tilde{N} = (\rho_0 b^3)^2 N$, where ρ_0 is the average number density of repeating units, b is the Kuhn length and N is the number of monomers. In the limit of $\tilde{N} \to \infty$ the collective density fluctuations are suppressed leading to an increase in the accuracy of the approximations in the SCF theory.^[30,51,52] This means that upon increasing the monomer number N, the accuracy of the MD-SCF simulations increases in turn. Last but not least, MD-SCF allows to efficiently relax polymer melts with high molecular weights, which is practically not feasible or extremely hard by using atomistic MD simulations. For example, it is extremely difficult to move the c.o.m. of polymer chains (N = 200) at least of their radius of gyration by using traditional MD simulations (see Supporting Information).

For all these reasons, the second set of systems are composed of long *s*-PP chains (45 chains of N = 200 monomers, Table 1) with high tacticity. In particular, five increasing *racemo* diads fraction (P_r) have been chosen: 0.74, 0.83, 0.87, 0.95, and 0.98, respectively. For each P_r fraction, five different distributions (according to the Bernoulli probability distribution) were generated to build the PP chains. In total, each system contains 45 *s*-PP chains with 5 independent *racemo* diads distributions according to the fixed values of P_r. A Bernoullian probability distribution of the diads was generated with the algorithm described by Ladd.^[53]

The MD-SCF simulations of the second set of systems were performed at constant temperature of 473 K. As in the short chain case, the resolution of the coarse grain density was increased step by step (from a mesh size of l = 0.8 to 0. nm and from 0.4 to 0.2 nm). In Figure 6, the stiffness of the *s*-PP chains has been calculated through the C_n parameter and compared with MD-SCF simulations results with N = 50, experimental and theoretical behaviors.

As it can be seen from Figure 6, the MD-SCF simulations of higher molecular weight reproduce, better than those with N = 50, the main trend of C_n which increases with increasing P_r , as predicted by theoretical RIS model of Allegra,^[48] Suter^[16,18] and Alfonso^[47] and observed experimentally by Fetters.^[12,14,15] In particular, the MD-SCF



Figure 6. Comparison of Flory's Characteristic ratio C_n as function of *racemo* diad content P_r , between: MD-SCF simulations of *s*-PP with I = 0.2 nm (N = 200 blue open circle, N = 50 blue circle), I = 0.4 nm (N = 200 light blue open circle, N = 50 light blue circle), I = 0.4 nm (N = 200 light blue open circle, N = 50 light blue circle), I = 0.8 nm (N = 200 yellow open circle, N = 50 yellow circle). SANS measurements from Fetters^[12,14,15] (open red square). RIS calculation from Alfonso^[47] (blue line), Suter^[16,18] (orange line), Allegra^[48] (dark green line). MC simulation of single unperturbed chain from Tzounis^[21] (green diamonds). Last stage of relaxation procedure for N = 50 (black square) and N = 200 (black triangle).

simulation with l = 0.8 nm (yellow open circle in the figure) follows the same monotonic increase of C_n versus P_p , while, both MD-SCF simulations with $l \le 0.4$ nm show a deviation from the monotonic behavior at $P_r = 0.87$. More in general, the stiffness of s-PP chains we calculated is close to the results of Suter (RIS at 413 K, yellow line in Figure 6) and experimental SANS measurements at 465 K^[12,14,15] and slightly lower than the estimation of Tzounis et al.^[21] calculated from single unperturbed chain MC simulations at 450 K. The final stage of the relaxation procedure, which requires a short relaxation with standard MD simulation, in case of N = 200 gives results (black triangle) very close to those ones obtained from the different stage involving the MD-SCF simulation. More in general, the stiffness of the s-PP chains is better reproduced from MD-SCF simulations with larger N. As mentioned before, this can be attributed to the different value of the invariant degree of polymerization *N* which is 1.1×10^4 for *N* = 50 and 5.6×10^4 for *N* = 200.

To analyze in more detail the *s*-PP chain conformations we computed the distribution of torsional angles (ϕ) belonging to the backbone of polymer chains for each P_r (See the Supporting Information). Then, we identified the *trans* (*t*) and *gauche* (*g*) torsional states at 180° and 78°, respectively.

In panel A of Figure 7 we report a comparison of the average fractions of *trans* (P_t) and *gauche* (P_g) state at increasing P_r content. In particular, a qualitative agreement between our calculations and the results of Tzounis^[21] calculated from MC simulations of a single unperturbed chain, emerges from the comparison. In addition, we reported in panel B of the same figure the maximum length of trans sequence defined as the number of consecutive torsional ϕ angles in the backbone corresponding to *trans* state. As expected, moving from relatively lower to higher P_r fraction the maximum length of trans sequence becomes higher (from 17 to 22). In Figure 7c, the X-ray scattering calculated from simulations of PP (N = 50 and 200) systems, relaxed by using the described procedure, are compared with the experimental data.^[50] Likely, due to the larger molecular weight of s-PP (N = 200), the region at low q (from 2 to 5 AA^{-1}) is slightly better reproduced.

As mentioned in the Introduction, the relation between the microstructure of a polymer (tacticity) and the macroscopic parameters of a polymer melt have been experimentally well studied. To this aim, SANS and plateau modulus (G_N^0) measurements have been used.^[10,14,15] Recently, Ahmad *et al.* reported a detailed



Figure 7. (a) Average of *trans* (P_t) and *gauche* (P_g) probability as function of P_r fraction. (b) Maximum length of trans sequence calculated from MD-SCF simulations as function of P_r . All data referred to MD-SCF simulation have been calculated on systems with a mesh size I = 0.2 nm. (c) Scattering function s(q) calculated for different chain length *s*-PP are compared with Experimental behavior ^[50] (red line). The Scattering functions have been calculated on the relaxed MD simulation by averaging configurations over the last 2 ns. (d) longest trans sequence, for each tacticity of *s*-PP, is reported as function of the packing length *p* (red open circle). The segmented black line is the best linear fitting (y = a + bx, a = 4.6 Å, and b = -0.094 Å/transseq.).

study on the relationship between high stereoregular *s*-PP melts and the viscoelasticity of the polymer.^[10] Using single-center metallocene and half-metallocene catalysts they synthetized *s*-PP samples characterized by different content of [*rrrr*] pentads and investigated their viscoelastic properties. In addition, in the work of Ahmad *et al.* the packing length (*p*) was estimated by using the plateau modulus G_N^0 . They found that the higher is the content of *racemo* pentads [*rrrr*] the smaller is the packing length *p*. It is worth noting that a low value of the packing length value corresponds to a chain segment which occupies less volume.^[54]

As a test case, we applied our method to calculate, and compare with experiments, the values of the packing length calculated on polymer melts having the same experimental pentad fraction reported by Ahmad *et al.* (see the pentad fraction in Table 1).

The packing length (p) is defined as the ratio between the effective volume of the polymer chain and its mean square end-to-end distance.^[55] From rheological measurements, p can be calculated through the Equation 5:

$$p = \sqrt[3]{\left(\frac{A^2kT}{864G_N^0}\right)} \tag{5}$$

where A^2 is a constant independent on temperature (A = 1.545), k is the Boltzmann constant, T is the absolute temperature, and G_N^0 is the plateau modulus of the polymer.

An equivalent method to calculate p which is more suitable for the quantities computed from MD simulations is reported in the Equation 6:

$$p = \frac{M}{R^2 \rho N_A} \tag{6}$$

where *M* is the mass of the polymer chain, *R* is the endto-end vector distance, ρ is the mass density of the system, and N_A is the Avogadro number.

In order to have a direct comparison between the experimental and calculated data, the same experimental samples with high *racemo* pentad fraction of 0.93, 91.5, 70.6, 60.1, 46.9 of [*rrrr*] were generated for *s*-PP with N = 200 (see the Table 1). Starting from the last configuration of each MD-SCF simulation (l = 0.2 nm) performed in

NVT ensemble, a short relaxation (1 ns in *NVT* ensemble) with standard MD has been performed. Then, *NPT* simulations long enough to have equilibrated densities were performed at same experimental temperature T = 473 K. From those simulations, the equilibrium end-toend distance and density values needed to compute the packing length have been calculated. In Table 3 experimental and calculated *p* (with Equation 6) are reported.

We found that *p* values calculated from the MD systems, equilibrated by applying the procedure described previously, are in good agreement with the experimental values measured through plateau modulus $G_N^{0,[10]}$ As a general behavior, the increase of the pentad content *rrrr* causes a decrease of the packing length. In particular, going from [rrrr] = 0.47 to 0.93 *p* decreases from 3 to 2.5, similarly to the experimental trend reported by Ahmad et al.^[10], which shows a decrease of p from 3.4 to 2.8. This result confirms that the MD-SCF method is suitable to model fine structural differences due to the polymer microstructure. Moreover, similar trends have been measured on high stereoregular polypropylene, with different racemo diad fraction, by Liu^[56] and Fetters.^[57] We argue that the decrease of packing length upon increasing the rrrr content is a direct consequences of the more compact average set of accessible chain conformations as indicated by the simultaneous increase of the maximum length of trans sequences (Figure 7b). Moreover, we observe a linear correlation between the increase of the longest trans sequence and the packing length p as reported in panel D of Figure 7.

Conclusions

The MD-SCF approach has been tested in the reproduction of the effect of tacticity on the structural properties of PP melts by applying a multi-stage relaxation procedure validated in a previous work.^[30] We propose a systematic study, by using MD-SCF simulations, investigating two set of systems. In the first set atactic, isotactic, and syndiotactic PP chains of low M.W. have been considered. We applied a relaxation procedure which successfully reproduces the stiffness of the PP chains, in terms of C_n , with equal and indistinguishable accuracy with respect to independent MD simulations.

Table 3. Experimental^[10] and calculated p values.</sup>

System	[rrrr] fraction	p (calc.) (Å)	p (exp.) (Å)
s-PP (N = 200)	0.93	2.51 (0.2)	2.83
	0.91	2.78 (0.3)	3.04
	0.71	3.01 (0.3)	3.26
	0.60	3.02 (0.3)	3.34
	0.47	3.03 (0.3)	3.39

In the second set of simulations, a higher M. W. of s-PP has been investigated. In particular, syndiotactic PP chains (N = 200) were generated with an increased fraction of racemo diads (according to Bernoullian distribution) going from 0.74 to 0.98. From MD-SCF simulations we found that the stiffness of the polymer is well reproduced with respect to SANS measurements, RIS predictions and MC simulations of a single unperturbed chain. In addition, the results from MD-SCF simulations indicate that even with a lower resolution of the coarse grain density (which can be easily controlled by changing the mesh size l) the stiffness of s-PP chains, and hence the excluded volume of polymer chains, shows a good match with experiments. This result is not surprising since the approximations of the MD-SCF approach gain accuracy at high M.W., as proved in a previous work.^[30]

Since the MD-SCF approach is capable to reproduce fine structural differences as a function of the tacticity, we reported a first application studying the packing length of very highly stereoregular syndiotactic PP. To this aim, we generated chains with a Bernoullian distribution of pentads (*rrrr*) equal to experimental samples reported by Ahmad *et al.*^[10] Applying the relaxation procedure on the systems with high stereoregularity, and performing a short relaxation by standard MD simulations, we computed and compared the packing length for all different pentad fractions. We found a very good agreement between the calculated packing length and those estimated from rheological measurements.

The present study expands the field of applicability of MD-SCF method at atomistic resolution to the study of stereoregular polymeric materials, in particular to the description and understanding of the effects of fine microstructural features of vinyl polymers on structural properties.

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