Estimation of Persistence Lengths of Semiflexible Polymers: Insight from Simulations¹

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Abstract—The persistence length of macromolecules is one of their basic characteristics, describing their intrinsic local stiffness. However, it is difficult to extract this length from physical properties of the polymers, different recipes may give answers that disagree with each other. Monte Carlo simulations are used to elucidate this problem, giving a comparative discussion of two lattice models, the self-avoiding walk model extended by a bond bending energy, and bottle-brush polymers described by the bond fluctuation model. The conditions are discussed under which a description of such macromolecules by Kratky—Porod worm-like chains holds, and the question to what extent the persistence length depends on external conditions (such as solvent quality) is considered. The scattering function of semiflexible polymers is discussed in detail, a comparison to various analytic treatments is given, and an outlook to experimental work is presented.

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INTRODUCTION

Flexibility of chain molecules (or lack of flexibility, respectively) is one of their most basic general properties [1-5]. It affects the use of macromolecules as building entities of soft materials, and controls some aspects of the functions of biopolymers in a biological context. Thus, it is important to understand its origin in terms of the macromolecular chemical architecture. and the extent to which it depends on external conditions (temperature, solvent quality if the polymer is in solution, as well as polymer concentration), and one therefore needs to be able to characterize macromolecular flexibility or stiffness precisely. The quantity that is supposed to describe the local intrinsic stiffness of a polymer is termed "persistence length" and often it is introduced (e.g. [4, 5]) as a length describing the exponential decay of orientational correlations of segments with the length of the piece of the chain separating them. Thus, let us consider a linear macromolecule composed of segments vectors $\{\vec{a}_i, i = 1, ..., N\}$, all having the same bond length $\ell_b \left(\langle \bar{a}_i^2 \rangle = \ell_b^2 \right)$, if we wish to allow for thermal fluctuations of the length of these segments). Then it is assumed that the correlation of two segments *i*, *j*, that are s = |i - j| steps along the chain apart, varies as

$$\langle \cos \theta(s) \rangle = \langle \vec{a}_i \cdot \vec{a}_j \rangle / \langle a_i^2 \rangle = \exp(-s\ell_b/\ell_p),$$

$$s \to \infty,$$
(1)

where ℓ_p is the persistence length.

In fact, Eq. (1) holds for models of linear polymer chains that strictly follow Gaussian statistics (for large distances between monomeric units), however, Eq. (1) is not true for real polymers, irrespective of the considered conditions: for dilute solutions and good solvent conditions one rather finds a power law behavior [6]

$$\langle \cos \theta(s) \rangle \propto s^{-\beta},$$

$$\beta = 2(1 - \nu), \quad 1 \ll s \ll N.$$
(2)

Here v is the well-known Flory exponent, describing the scaling of the end-to-end distance $\vec{R} = \sum_{i=1}^{N} \vec{a}_i$ with the number N of segments, $\langle R^2 \rangle \propto N^{2v}$, with $v \approx 3/5$ (more precisely [7], v = 0.588) in d = 3 dimensions [1–5]. Polymer chains in dense melts do show a scaling of the end-to-end distance as predicted bv Gaussian statistics, $\langle R^2 \rangle \propto N$ (i.e., v takes the meanfield value $v_{MF} = 1/2$), and hence it was widely believed, that Eq. (1) is useful for polymer chains under melt conditions. However, recent analytical and numerical work [8, 9] has shown that this assertion is completely wrong, and there also holds a power law decay, though with a different exponent,

$$\langle \cos \theta(s) \rangle \propto s^{-3/2}, \quad 1 \ll s \ll N.$$
 (3)

More recently, it was also found by approximate analytical arguments [10], and verified in extensive simulations [11] that Eq. (3) also holds for chains in dilute solutions at the Theta point. In practice, since asymptotic power laws such as Eqs. (2), (3) hold only in the intermediate regime $1 \ll s \ll N$ and hence one must

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Fig. 1. (a) Semi-log plot and (b) Log–log plot of $\langle \cos \theta(s) \rangle$ versus *s* as obtained from Monte Carlo simulations (as described in [11]) using the pruned-enriched Rosenbluth method (PERM algorithm [12]) for a self-avoiding walk with nearest-neighbor attraction ε , under Theta point conditions. *N* = (*1*) 6400, (*2*) 3200, (*3*) 1600, (*4*) 800 and (*5*) 400. The full curve in (a) and straight line in (b) represents the relation $\langle \cos \theta(s) \rangle = 0.16s^{-3/2}$.

consider the limit $N \rightarrow \infty$, one easily could be misled if data for $\langle \cos \theta(s) \rangle$ are considered for insufficiently long chains. As an example Fig. 1 presents simulation results for the simple self-avoiding walk (SAW) model on the simple cubic (sc) lattice, where an attractive energy ε between neighboring occupied sites (representing the effective monomers of the chain) occurs and the temperature is chosen as $k_B T/\epsilon = 3.717$ which is known to reproduce Theta point conditions for this model [12]. One can see clearly that the data for $N \to \infty$ and $s \ge 10$ do approach Eq. (3), but for finite N systematic deviations from Eq. (3) clearly are visible already for s = N/10. On the semi-log plot, for rather short chains one might be tempted to apply a fit of an exponential decay proportional to $\exp(-s\ell_b/\ell_p)$ to the data for rather large s, but resulting estimates for ℓ_p/ℓ_b are not meaningful at all: for the considered model, the chain is fully flexible, any reasonable estimate for ℓ_p/ℓ_b that describes the local intrinsic stiffness of the chain should be (i) of order unity (see Fig. 1a, $\ell_p/\ell_b \approx 0.94$), and (ii) independent of N. Both conditions are dramatically violated, of course, if estimates for ℓ_p/ℓ_b were extracted from fits to an exponential decay in this way.

Since the intrinsic stiffness of a chain is a local property of a macromolecule, one might alternatively try the recipe to either fit Eq. (1) in the regime of small s to the data, or assume that Eq. (1) holds for s = 1 already and hence

$$\ell_p / \ell_b = -1 / \ln(\langle \cos \theta(1) \rangle). \tag{4}$$

This recipe works in simple cases, such as the SAW model where an energy ε_b associated with bond bending is added (every kink of the walk by $\pm 90^\circ$ on the sc lattice costs ε_b), see Fig. 2, but it fails for molecules

with more complex chemical architecture, such as bottle-brush molecules [13-16]. The dramatic failure of Eq. (4) for bottle-brush polymers is understood in terms of their multiscale structure (Fig. 3): The side chains lead to a stiffness of the backbone on a mesoscopic scale, even if on the local scale of nearest-neighbor bonds the backbone is still rather flexible. The question of understanding this stiffening of bottlebrush polymers because of their grafted linear side chains [11, 13-38] or grafted branched objects [39– 42] is an issue of longstanding debate in the literature.

Complex polymer architecture is only one out of many reasons which make the analysis of bond orientational correlations based on Eqs. (1) or (4) problematic. In dilute solutions we expect that a nontrivial crossover occurs when the solvent quality is marginal, i.e. close to the Theta point a large size ξ_T of "thermal blobs" [43] exists, such that for values of s along the backbone of the chain corresponding to distances $r(s) > \xi_T$ one expects that excluded volume effects are visible and hence Eq. (2) should hold. For semidilute solutions [43], on the other hand, in the good solvent regime the inverse effect occurs: there exists a screening length $\xi(c)$ depending on the polymer concentration c (also called size of "concentration blobs" [43]), such that excluded volume effects are pronounced for $r(s) < \xi(c)$ but are absent for $r(s) \gg \xi(c)$. Then Eq. (3) holds for the latter case and Eq. (2) for the former, for rather flexible chains. If the chains are semiflexible, in favorable cases (e.g., for simple chemical architecture of the polymers) we might observe Eq. (1) for $1 < s < s^*$ where s^* depends on the local intrinsic stiffness of the chain, which we wish to characterize by ℓ_p .

Then the question arises whether s^* is smaller than any of the other crossover chemical distances (due to marginal solvent quality, described by a Flory-Huggins



Fig. 2. Semi-log plot of $\langle \cos \theta(s) \rangle$ vs. *s* for a semiflexible version of (a) a SAW model on the sc lattice, cf. text, and (b) the bondfluctuation model of bottle-brush polymers under very good solvent conditions [11, 17]. Part (a) refers to the chains of length N =50000, and several choices of the parameter $q_b = \exp(-\varepsilon_b/k_BT)$ controlling the chain stiffness, namely $q_b = (I) 0.02$, (2) 0.05, (3) 0.1, (4) 0.2 and (5) 0.4. Using Eq. (4), the straight lines indicate the exponential decay $\exp(-s\ell_b/\ell_p)$ for the choices of q_b . Part (b) refers to the case of bottle-brush polymers where every effective monomer of the backbone has one side chain of length $N_s = 24$ grafted to it, and several choices of backbone chain length $N_b = (I)$ 259, (2) 131 and (3) 67. Here $\ell_p/\ell_b =$ $-1/\ln(\langle \cos \theta(1) \rangle)$ has been extracted from the chain backbone only.

parameter χ with $(1/2 - \chi) \ll 1$ [43, 44], or due to nonzero *c*) or not. The conclusion of this discussion is that the behavior of bond orientational correlations $\langle \vec{a}_i \cdot \vec{a}_j \rangle$ is subtle, and not always suitable to obtain straightforwardly information on the intrinsic stiffness of macromolecules; as a further caveat we mention that in general it is also not true that this correlation depends on the relative distance s = |i - j| only: it matters also, if one of the sites is close to a chain end.

Another popular definition is the local persistence length $\ell_p(i)$ defined as [1, 2]

$$\ell_{p}(i)/\ell_{b} = \left\langle \vec{a}_{i} \cdot \vec{R} \right\rangle / \left\langle \vec{a}_{i}^{2} \right\rangle.$$
(5)

However, it has been shown by renormalization group methods that in good solvents one has, for $N \to \infty$, $\ell_p(i) \propto [i(N-i)]^{2v-1}$, $i \ge 1$, so the behavior of $\ell_p(i)$ in the chain interior clearly is unsuitable to conclude anything about the local stiffness of a chain under good solvent conditions, and this conclusion has been corroborated by simulations [11, 17]. Sometimes it has been argued that a better choice is to take the correlation between the first bond vector and the end-to-end distance, $\ell_p(1)$ [46]. However, since in a macromolecule the chemical nature of the end monomer always differs from inner monomers, one can never expect that $\ell_p(1)$ precisely characterizes the local stiffness of a linear macromolecule in the inner parts of a chain. Moreover, since $\langle R^2 \rangle$ reflects all the crossovers (due to "thermal blobs" etc.), [43], as discussed above, it is premature to expect that $\ell_p(1)$ stays unaffected from them. We also note that for d = 2 dimensions under good solvent conditions it has been shown [47] that $\ell_p(1) \propto \ln N \to \infty$ as $N \to \infty$, so in this case $\ell_p(1)$ clearly is not a useful measure of the intrinsic stiffness of a chain at all. Since formula (5) is difficult to extract from any experiments, and inconvenient for simulation studies due to high sampling effort, we shall not discuss Eq. (5) further in the present paper.

Experimental studies try to extract the persistence length either from scattering analyses of the single chain structure factor (e.g. [28, 32, 34-38, 48]) or from analyses of extension versus force measurements of stretched chains (e.g. [49-57]). However, the interpretation of the latter experiments must rely on a theoretical model of the extension versus force curve. While this task is simple for ideal random walk models of polymers [4, 5, 58] and also for semiflexible polymers when excluded volume is neglected [59], so that the Kratky-Porod (K-P) model [60] of worm-like chains can be used, it is very difficult (due to multiple crossovers [61, 62]) if excluded volume effects are included. These excluded volume effects cause an intermediate nonlinear variation of the extension versus force curve (the chain is then a string of "Pincus blobs" [63]), making the estimation of the persistence length difficult [62], and this behavior has also been verified in recent experiments [56, 57]. Since we have given a recent extensive discussion of this problem elsewhere [62], we shall not dwell on this problem here further, and focus on the problem how the persistence length shows up in the single chain structure factor S(q). Here the key idea is that the scattering intensity

Fig. 3. Sketch of the multiple length scales that one may define for bottle-brush polymers (schematic): While on scales where one resolves the effective monomers of both the backbone and the side chains, the correlations of back-

bone vectors \vec{a}_i^b (which have length ℓ_b) and side chain bond vectors \vec{a}_i^s (which have length ℓ_s) can be studied, as well as end-to-end distances $\vec{R}_{e,s}$, s of side chains and of the backbone $\vec{R}_{e,bb}$ (and the corresponding gyration radii). In the bond-fluctuation model of Fig. 2, for simplicity no chemical difference between backbone and monomer was considered, so $\ell_b = \ell_s$ was chosen. The "microscopic" contour length of the backbone then is $L_b = N_b \ell_b$, if the backbone has N_b bonds. On a coarse-grained level the bottle-brush resembles a worm-like chain of thickness (crosssectional radius) R_{cs} and contour length $L_{cc} < L_b$, which is locally straight on the scale of the persistence length ℓ_p .

S(q) at scattering wavenumber q yields information on the structure of the macromolecule at a length scale $\lambda = 2\pi/q$. This problem also is subtle, even in the framework of simple models (see Fig. 4a, 4b, 4c, 4d) used for simulations. If λ is of the scale of the cross sectional radius R_{cs} for the models (a, b, d) or the lattice spacing in (c), local structure on the scale of effective subunits is revealed: soft (a) versus hard (b) effective cylinders, hard spheres in (d), but one could also conceive a chain where soft spheres are jointed, etc. When one considers semiflexible chains with no excluded volume, the persistence length ℓ_p would be just one half of the step length ℓ_K in cases (b), (d), where one then requires a strong bond angle potential to make these chains semiflexible rather than flexible; however, as emphasized above, such models neglecting excluded volume completely will inevitably imply Eq. (1), which is inappropriate for real polymers under all physically possible conditions. So the information on chain stiffness, as described by the persistence length, is hidden in some intermediate range of wavenumbers. E.g., for the model (c), which will be used extensively in the rest of the paper (but in d = 3 dimensions, since the case of d = 2 is rather special [62] as will be discussed below), we need wavenumbers in the range $2\pi/\sqrt{\langle R_g^2 \rangle} \ll q \ll 2\pi/a$, where *a* is the lattice spacing. The aim of the present paper is to present a discussion of how one can obtain detailed information on intrinsic chain stiffness from the gyration radius of the macromolecules and from the structure factor S(q)in the suitable intermediate range of wavenumbers q.

The outline of our paper is as follows: in the next section, we summarize some pertinent theoretical results on S(q). In the third section, our Monte Carlo simulation methods are briefly described. In the fourth section, a comparative discussion of simulation results for two models is given, the bond fluctuation model of bottle-brushes (c.f. Fig. 4a), and the self-avoiding walk model on the simple cubic lattice with variable bending energy (cf. Fig. 4c). The final section contains our conclusions.

SOME THEORETICAL RESULTS ON THE STRUCTURE FACTOR OF ISOLATED MACROMOLECULES IN SOLUTION

We consider here a single macromolecule with linear chain architecture, assuming a sequence of N + 1(effective) monomeric units at positions $\vec{r}_i, j = 1, 2, ...,$ N + 1, with effective bond vectors $\vec{a}_j = \vec{r}_{j+1} - \vec{r}_j$, j =1, ..., N. We have in mind application to standard polymers like polystyrene (disregarding here the scattering from the side groups that are attached to the backbone of the chain, see e.g. Rawiso et al. [48] for a discussion of this problem in an experimental context). We also have in mind application to the scattering from the backbone of bottle-brush polymers (this is experimentally directly accessible from neutron scattering [28] if selective deuteration only of the backbone is used, while in the case of deuteration of the whole macromolecules [32, 34] this information can be inferred only indirectly). Due to the restriction to "effective monomeric units" rather than talking about the scattering from individual atoms with the appropriate scattering lengths, we clearly disregard information on the scale of the length of an effective bond, but we then need not discuss experimental problems such as contrast factors between the scattering from the macro molecule and the solvent [48]. The effect of the cross-sectional structure of the chain (finite chain thickness D) is not explicitly considered as well (experimentally this problem often is approximated in terms of the Guinier





Fig. 4. Various models of semiflexible polymers, as discussed in the context of simulations. Case (a) shows the snapshot picture of a typical conformation of a simulated bottle-brush polymer using a backbone chain length $N_b = 1027$, side chain length $N_s = 24$, projected into the *xy*-plane (this model is discussed in more detail in Section "Monte Carlo Simulation Methods and Models"). Case (b) shows a model of freely jointed cylindrical rods of Kuhn step length ℓ_K and diameter $D = 2R_{cs}$, with R_{cs} the cross-sectional radius (if $R_{cs} = 0$ this leads to a simple off-lattice random walk configuration, while excluded volume interaction is introduced if overlap of the cylinders is forbidden). Case (c) shows the SAW model on the square lattice with lattice spacing a (D = a in this case), where 90° bends cost an energy $\varepsilon_b \gg k_B T$, so the chain consists of straight pieces where n_{str} steps go in the same lattice direction, with $n_{str} \gg 1$. Case (d) shows a model of tangent hard spheres with radius R_{cs} (and $\ell_k = 2R_{cs}$).

[73] approximation, writing the observed scattering intensity $S_{obs}(q) = S(q) \exp(-q^2 R_c^2/2)$, with R_c some "effective" cross-sectional radius of the chain [48]). Thus only wavenumbers $qD \ll 2\pi$ are physically meaningful: in the case of the lattice model, Fig. 4c, D = a, of course. The structure factor then is defined as

$$S(q) = \frac{1}{(N+1)^2} \left\langle \sum_{j=1}^{N+1} \sum_{k=1}^{N+1} \exp[i\vec{q} \cdot (\vec{r}_j - \vec{r}_k)] \right\rangle, \quad (6)$$

and does not depend on the direction of the scattering wavevector q. In d = 3 dimensions, it has the small q expansion

$$S(q) = 1 - \left\langle R_g^2 \right\rangle q^2 / 3 + \dots, \quad q \to 0, \tag{7}$$

where the mean square gyration radius $\langle R_g^2 \rangle$ enters

$$\left\langle R_g^2 \right\rangle = \frac{1}{\left(N+1\right)^2} \left\langle \sum_{j=1}^{N+1} \sum_{k=j+1}^{N+1} \left(\vec{r}_j - \vec{r}_k\right)^2 \right\rangle.$$
 (8)

POLYMER SCIENCE Series C Vol. 55 No. 1 2013

Other characteristic lengths of the chain molecule are the mean square end-to-end distance

$$\left\langle \boldsymbol{R}^{2}\right\rangle = \left\langle \left(\sum_{i=1}^{N} \vec{a}_{i}\right)^{2}\right\rangle$$
 (9)

and the contour length

$$L = N\ell_b, \tag{10}$$

but neither of these lengths can be inferred directly from the scattering. For chains in dense melts or in dilute solutions under Theta conditions one typically uses an ideal chain approximation (disregarding, e.g., logarithmic corrections at the Theta point [3, 43, 64])

$$\left\langle R^{2}\right\rangle = 6\left\langle R_{g}^{2}\right\rangle = C_{\infty}\ell_{b}^{2}N, \quad N \to \infty,$$
 (11)

with C_{∞} a characteristic constant [1–5]. In this case one introduces an equivalent freely jointed Kuhn chain with the same contour length, $\langle R^2 \rangle = n_k \ell_K^2$, where n_K is the number of equivalent Kuhn segments and ℓ_K their length,

$$\ell_K = C_{\infty} \ell_b, \quad n_K = N/C_{\infty}, \quad N \to \infty.$$
 (12)

For a semiflexible worm-like chain with $C_{\infty} \ge 1$ Eq. (1) holds. However, since under Theta conditions (and melts) formula (12) is approximately true, one finds

$$\ell_p = 3 \langle R_g^2 \rangle / (N \ell_b), \quad \langle R^2 \rangle = 2 \ell_p \ell_b N, \quad N \to \infty, (13)$$

if the relation $\ell_p = \ell_k/2 = C_\infty \ell_b/2$ then simply is taken as an alternative definition of a persistence length. For the simple SAW model of Fig. 1 this gives $\ell_p = 0.94$ lattice spacings: but as expected, using this value in the simple exponential $\exp(-s\ell_b/\ell_p)$ one does not obtain a description of the actual data in Fig. 1 on the basis of this description, because the actual behavior of bond orientational correlations is a power law decay, Eq. (3). Note, however, that the relation $\ell_p = \ell_K/2$ makes only sense for semiflexible chains for which $C_\infty \ge 1$ at the Theta point, which is not the case for the model of Fig. 1.

In the case of good solvent conditions excluded volume interactions invalidate formula (11) and one finds instead [3, 7, 43, 64]

$$\langle R^2 \rangle = 2\ell_p^R \ell_b N^{2\nu}, \quad \langle R_g^2 \rangle = \frac{1}{3} \ell_p^{R_g} \ell_b N^{2\nu}, \quad N \to \infty$$
(14)

with [7] $v \approx 0.588$ instead of the mean field value $v_{MF} = 1/2$ that appears in formula (13). Note that we have defined the prefactors of the relations $\langle R^2 \rangle \propto N^{2v}$, $\langle R_g^2 \rangle \propto N^{2v}$ in formula (14) in complete analogy with formula (13) [11], but we shall see shortly that the lengths ℓ_p^R , $\ell_p^{R_g}$ do not play the role of a persistence length that describes the local intrinsic stiffness of the chains.

For a better understanding of this problem, in particular when ℓ_p is very large, it is of interest to consider the crossover from the rod limit (that occurs for $L < \ell_K$, i.e. $n_K < 1$) to the Gaussian coil limit. This problem can be worked out easily for various models of discrete chains [1–5] as well as for the Kratky–Porod model. Describing the chain by a continuous curve $\vec{r}(s)$, *s* being the curvilinear coordinate along the chain contour, the potential energy of a particular conformation of the chain is given by

$$\mathcal{H} = \frac{\kappa}{2} \int_{0}^{L} \left(\frac{\partial^2 \vec{r}(s)}{\partial s^2} \right)^2 ds, \quad \kappa = k_B T \ell_p \quad (d = 3).$$
(15)

In Eq. (15) it is clearly assumed that κ is a constant, independent of the contour length L (or chain length N, respectively), and the same holds for ℓ_p . The physical interpretation of κ is in terms of the local bending stiffness of the chain. Eq. (15) can be used for arbitrary values of the ratio $L/\ell_p = n_p$, and one can show [60, 65]

$$\frac{\langle R^2 \rangle}{2\ell_p L} = 1 - \frac{1}{n_p} [1 - \exp(-n_p)], \qquad (16)$$

and

2

$$\frac{3\langle R_g^2 \rangle}{\ell_p L} = 1 - \frac{3}{n_p} + \frac{6}{n_p^2} - \frac{6}{n_p^3} [1 - \exp(-n_p)].$$
(17)

One immediately recognizes that for $n_p = 2n_K \rightarrow \infty$ one recovers Eq. (13), while in the opposite limit the results for rigid rods of length *L* are obtained,

$$\left\langle R^2 \right\rangle = 12 \left\langle R_g^2 \right\rangle = L^2, \quad n_p \ll 1.$$
 (18)

However, the generalization of these results to the good solvent case, where excluded volume matters, is not straightforward. Of course, for $n_p \ll 1$ excluded volume is irrelevant, Eq. (18) remains valid. It turns out, however, that in d = 2 Eqs. (16), (17) are not valid at all, one has no regime of Gaussian chain behavior as described in Eq. (13), and rather near $n_p = 1$ a cross-over from rigid rod behavior to the behavior of two-dimensional self-avoiding walks occurs [62, 66] (v = 3/4)

$$\langle R^2 \rangle \propto \langle R_g^2 \rangle \propto \ell_p^{1/2} L^{3/2}, \quad L > \ell_p.$$
 (19)

For d = 3, however, Eqs. (16), (17) for semiflexible chains remain valid for $n_p < n_p^*(\ell_p)$ where $n_p^*(\ell_p \to \infty) \to \infty$. This crossover contour length $L^* = n_p^* \ell_p$ has first been estimated by a Flory argument as [62, 67, 68]

$$L^* \propto \ell_p^3 / D^2, \quad n_p^* \propto (\ell_p / D)^2. \tag{20}$$

Note, however, that Flory arguments imply v = 3/5 in d = 3 (rather than the precise value $v \approx 0.588$ [7]) and cannot predict any prefactors in Eq. (20); they are based on a crude balancing of the elastic energy of chain stretching (taken as Gaussian) and a mean field estimate of binary interactions: having in mind a model description as in Fig. 4b, one takes the second virial coefficient proportional to the rod volume on the scale of the persistence length, $v_2 \propto \ell_p^2 D$, and in this way the effective chain diameter D enters the estimated Eq. (20) [62, 67, 68]. Numerical results, however, seem to suggest that rather [62] $n_p^* \propto (\ell_p/D)^{\zeta}$ with an exponent $\zeta \approx 1.5$.

In any case, the conclusion of this discussion is that for semiflexible chains in d = 3 the mean square radii as a function of the reduced contour length $n_p = L/\ell_p$ exhibit two successive crossovers, from rods to Gaussian coils near $n_p = 1$ and from Gaussian coils to swollen chains (described by Eq. (14)) near $n_p = n_p^*$. These two crossovers have in fact been seen nicely in both experiment [69] and computer simulation of the model of Fig. 4c [62, 70, 71].

We now turn to a discussion how these behaviors show up in the scattering function S(q) at larger wavenumbers, when Eq. (7) does not hold. In the regime $n_p < 1$, when the chain behaves like a rigid rod, one can work out the scattering function in the continuum limit as [72]

$$S_{\rm rod}(q) = \frac{2}{qL} \left[\int_{0}^{qL} dx \frac{\sin x}{x} - \frac{1 - \cos(qL)}{qL} \right]$$
(21)

while for a discrete chain of N + 1 scatterers along a rod of length $L = N\ell_b$ one has

$$S_{\rm rod}(q) = \frac{1}{N+1} \left[-1 + \frac{2}{N+1} \sum_{k=0}^{N} (N+1-k) \frac{\sin(q\ell_{b}k)}{q\ell_{b}k} \right],$$
$$q\ell_{b} < 2\pi.$$
(22)

It is noteworthy to recall that the large q-limit of Eq. (21) contains information on the contour length L and shows a 1/q decay,

$$S_{\rm rod}(q \to \infty) = \pi/(qL).$$
 (23)

In the Gaussian regime, that applies for chain lengths that correspond to $1 \ll n_p < n_p^*(\ell_p)$ in d = 3, the structure factor S(q) is described by the well-known Debye function,

$$S_{\text{Debye}}(q) = \frac{2}{X} \left\{ 1 - \frac{1}{X} [1 - \exp(-X)] \right\}, \quad X \equiv q^2 \left\langle R_g^2 \right\rangle. (24)$$

For small X, Eq. (24) reduces to Eq. (7), as it must be, while for large X Eq. (24) yields $S_{\text{Debye}}(q) \approx 2/X =$ $2/[q^2 \langle R_g^2 \rangle]$. While for flexible chains at the Theta point Eq. (24) is expected to hold for large q, up to $q\ell_{h}$ of order unity where effects due to the local structure of monomeric units comes into play, the validity of Eq. (24) for semiflexible chains is much more restricted, since then the rod to coil crossover matters also with respect to the intrinsic structure of these polymers, as it is probed by S(q). In oder to discuss this problem, it is useful to cast S(q) in the representation of the so-called Kratky plot [73], qLS(q) is plotted as a function of qL = Y. For rigid rods, one simply would have a linear increase of qLS(q) with Y for small Y, which smoothly crosses over near Y = 1 to a flat plateau (which has the value π , cf. Eq. (23)). For chains where intermonomer distances follow Gaussian distributions, at all scales, the Kratky plot exhibits a maximum at Y_{max} , and then a crossover to a decay proportional to Y^{-1} occurs. To locate this maximum, it is convenient to write qLS(q) as $\sqrt{X}\left(L/\sqrt{\langle R_g^2 \rangle}\right)S_{\text{Debye}}(q)$ as a function of X, noting that the maximum occurs at $X_{\text{max}} \approx 2.13$, i.e. the Kratky plot has its maximum at $Y_{\text{max}} \approx \sqrt{2.13} L / \sqrt{\langle R_g^2 \rangle}$, and the height of this maximum also is of order $L/\sqrt{\langle R_g^2 \rangle}$. Using now Eq. (13) in the

POLYMER SCIENCE Series C Vol. 55 No. 1 2013

form $\langle R_g^2 \rangle = L\ell_p/3$, we recognize that the maximum of the Kratky plot occurs at

 $(qL)_{\text{max}} \approx \sqrt{6.4} (L/\ell_p)^{1/2}, \quad 1 \ll L/\ell_p < n_p^*(\ell_p)$ (25) and also the height of this maximum scales proportional to $\sqrt{L/\ell_p}$. However, while for flexible chains under Theta conditions (for which ℓ_p and ℓ_b are of the same order), one observes on the Kratky plot for $Y \gg Y_{\text{max}}$ a decay $qLS(q) \propto q^{-1}$, for semiflexible chains a crossover from this decay to the plateau value π (given by Eq. (23)) is expected. This is also true for semiflexible chains under good solvent conditions, if the persistence length ℓ_p is large enough so that

 $n_p(=L/\ell_p) < n_p^*(\ell_p)$, and hence excluded volume effects still can be ignored. The description of this decay of the structure factor from its peak towards this so-called "Holtzer plateau" [74] has been a long-standing problem in the literature [75–96]. Only in the limit $N \to \infty$ a simple explicit result derived from Eq. (15) is available [80],

$$LqS(q) = \pi + \frac{2}{3}(q\ell_p)^{-1}, \quad L \to \infty, \quad q \to \infty, \quad (26)$$

but we should keep in mind that the limit $q \to \infty$ is well-defined for a simple mathematical continuum model such as Eq. (15), while for real chains (and for simulations) the regime $q\ell_b > 2\pi$ is not at all meaningful. Although the decay $S(q) \propto q^{-2}$, that Eq. (26) predicts for $q\ell_p \ll 1$, is compatible with the power law decay of the Debye function, Eq. (24), for large q

$$S(q) \approx \frac{2}{q^2 \langle R_g^2 \rangle} \rightarrow qLS(q) \approx 6(q\ell_p)^{-1}, \quad q \rightarrow \infty, \quad (27)$$

the prefactor in Eq. (27) is by a factor of 9 larger than the prefactor of the q^{-1} term in Eq. (26), so both Eqs. (24) and (26) are inconsistent with each other. This inconsistency is due to the fact that Eq. (26) is only accurate for $q\ell_p > 3$, it should not be used for small $q\ell_p$. After many less successful attempts, Kholodenko [86-89] achieved a description which interpolates between the limiting cases of rigid rods and of Gaussian coils, capturing the scattering law of both limits exactly, but deviating from the exact result ("exact" refers to the Hamiltonian Eq. (15), so no excluded volume effects are being accounted for) in the intermediate regime; this exact behavior is known from systematic expansions [94-96] whose use requires heavy numerical work, and will not be considered here. Recently we have shown [97] that the exact method of Stepanow [95, 96] deviates only very little from the approximation of Kholodenko [89], which can be cast in the form

$$S(q) = \frac{2}{x} \left[I_1(x) - \frac{1}{x} I_2(x) \right], \quad x = 3L/2\ell_p,$$
(28)

where

$$f(z) = \begin{cases} \frac{1}{E} \frac{\sinh(Ez)}{\sinh z}, & q \le 3/2\ell_p, \\ \frac{1}{E'} \frac{\sin(E'z)}{\sinh z}, & q > 3/2\ell_p, \end{cases}$$
(29)

with

 $E = [1 - (2q\ell_p/3)^2]^{1/2}, \quad E' = [(2q\ell_p/3)^2 - 1]^{1/2}. \quad (30)$ We stress that all these analytical results Eqs. (11)– (13), (16), (17), (24)–(30) are only applicable if excluded volume effects are negligible. When we consider very long semiflexible chains, such that $n_p = L/\ell_p > n_p^*(\ell_p)$, we expect that the Gaussian results $\langle R^2 \rangle = 2\ell_p L = 2\ell_p^2 n_p$ and $\langle R_g^2 \rangle = (1/3)\ell_p L = (1/3)\ell_p^2 n_p$ hold roughly up to $n_p^*(\ell_p)$, and there a smooth crossover to the excluded volume power laws, Eq. (14), occurs. We first note that hence $n_p^*(\ell_p)$ corresponds to a crossover radius R^* of the chains as well, $R^{*2} = 2\ell_p L^* = 2\ell_p^2 n_p^*$. Omitting factors of order unity, we conclude

$$R^* = \sqrt{\ell_p L^*} = \ell_p \sqrt{n^*} \propto \ell_p^2 / D, \qquad (31)$$

where in the last step Eq. (20) was used. For $n_p > n_p^*(\ell_p)$ we hence expect, invoking the fact that the crossover in the linear dimensions for $n_p = n_p^*$ should be smooth,

$$\left\langle R^2 \right\rangle = R^{*2} (n_p/n_p^*)^{2\nu}$$

$$\propto \ell_p^4 / D^2 (D/\ell_p)^{4\nu} n_p^{2\nu} \approx \ell_p^2 \left(\frac{D}{\ell_p}\right)^{2/5} n_p^{6/5},$$
(32)

where in the last step the Flory estimate $v \approx 3/5$ was used (recall that in Eq. (31) the exponent ζ defined above has also been put to its Flory value, $\zeta = 2$). In terms of *N* and ℓ_p , Eq. (32) becomes $\langle R^2 \rangle \propto \ell_b^{6/5} (\ell_p D)^{2/5} N^{6/5}$. In terms of the constant ℓ_p^R defined in Eq. (14), we would have $\ell_p^R \propto \ell_b^{1/5} (\ell_p D)^{2/5}$.

The consequences for the scattering function S(q) are now clear, since the gyration radius shows the same scaling behavior as $\langle R^2 \rangle$, apart from prefactors of order unity. Hence we have

$$\sqrt{\langle R_g^2 \rangle} \propto \ell_b^{3/5} (\ell_p D)^{1/5} N^{3/5}$$
(33)

and only for $1/\sqrt{\langle R_g^2 \rangle} < q < 1/R^*$ we can expect to see the nontrivial power law

$$S(q) \propto q^{-1/\nu},$$
 (34)

while at q^* defined from $q^*R^* = 1$ we have a smooth crossover to the standard Debye law, $S(q) \propto q^{-2}$. Near $q\ell_p = 1$ then a smooth crossover to the rod-like scattering law $S(q) \propto q^{-1}$ occurs. So the three power laws for the radii as a function of chain length $(\sqrt{\langle R^2 \rangle} \propto N)$ in the rod regime, $\propto N^{1/2}$ in the regime of Gaussian coils, and $\propto N^{\vee}$ in the regime of swollen coils) find their counterpart in the scattering function, if N is large enough. The schematic Fig. 5 illustrates these crossover behaviors. The three regimes of the $\langle R^2 \rangle$ versus N (or n_p , respectively) curve, namely rods, Gaussian coils, and swollen coils (Fig. 5a) appear in the S(q)vs. q curve (or qLS(s) vs. qL-curve, in the Kratky representation) in inverse order: the rods occur for large q_{1} , then occurs a first crossover to Gaussian coils, and a second crossover to swollen coils. Of course, if the chains are very stiff but not extremely long, it may be that the regime $n_p > n_p^*$ is not reached: then in part (a) the swollen coil regime is absent, and in part (b) as well: then the K–P model can describe S(q) fully, including the regime of the maximum of the Kratky plot. Since the crossovers are smooth, it may be difficult to identify the different power laws in Fig. 5b in practice, however.

We also note that the different regimes are also only well separated if both ℓ_p is very large (in comparison to ℓ_b) and also ℓ_p/D needs to be very large. If ℓ_p is very large, but *D* also (as in the case of bottle-brush polymers [11, 17, 70, 71]) then the regime of Gaussian coils disappears from both Fig. 5a and 5b, and the K–P model loses its applicability.

MONTE CARLO SIMULATION METHODS AND MODELS

In the present work, we focus on lattice models exclusively, because for them particularly efficient simulation methods exist; pertinent work on coarsegrained off lattice models of bottle-brush polymers studied in Molecular Dynamics methods for variable solvent quality [98] will be mentioned in the conclusions section.

The archetypical lattice model of a polymer is the self-avoiding walk on the simple cubic lattice [99]. Each effective monomer takes a single lattice site, the length of an effective bond is the lattice spacing, so adjacent monomers along the chain are nearest neighbors on the lattice. Double occupancy of lattice sites being forbidden, excluded volume interactions under very good solvent conditions are modelled.

The properties of this basic model are very well established [100]. Solvent quality can be included as a variable into this model implicitly, by allowing for an (attractive) energy ε that is won if two monomers (that



Fig. 5. (a) Schematic plot of the normalized mean square radius $\langle R^2 \rangle / (2\ell_p L)$ versus $n_p = L/\ell_p$ (apart from a factor of 2 this is the number of Kuhn segments), on log–log scales. The Kratky–Porod (K–P) model describes the crossover from rods $(\langle R^2 \rangle = L^2)$ to Gaussian coils $(\langle R^2 \rangle = 2\ell_p L)$. At $n_p^* = (\ell_p/D)^2$, according to the Flory theory a crossover to swollen coils occurs, where $\langle R^2 \rangle \propto n_p^{2\nu}$ with $\nu = 3/5$ (according to the Flory theory). (b) Schematic Kratky plot of the structure factor of a semiflexible polymer, qLS(q) plotted vs. qL, on log–log scales. Four regimes occur: in the Guinier-regime, $S(q) \approx 1 - q^2 \langle R^2 \rangle / 3$; it ends at the maximum of the Kratky plot, which occurs roughly at $q_{\max} \sqrt{\langle R^2 \rangle} \approx 1$ (constants of order unity being ignored throughout). For very large L then a regime of swollen coils with $S(q) \propto q^{-1/\nu}$ is observed, until near $qR^* \approx 1$ a crossover to Gaussian coil behavior occurs ($R^* \approx \ell_p^2/D$). In the Gaussian coil regime $S(q) \propto q^{-2}$, until at $q\ell_p$ of order unity the crossover to the rod-like regime occurs ($qLS(q) = \pi$). Only the latter two regimes are captured by the Kratky–Porod model.

are not nearest neighbors along the chemical sequence of the chain) are nearest neighbors on the lattice. One then finds that the Theta point, at which (apart from logarithmic corrections [3, 64]) the mean square radius $\langle R^2 \rangle$ scales like a Gaussian chain, $\langle R^2 \rangle \propto N$, occurs for $q \equiv \exp(\varepsilon/k_{\rm B}T) = q_{\theta}(\approx 1.3087)$ [12]. On the other hand, if one introduces an energy cost ε_b whenever the walk makes a turn by $\pm 90^{\circ}$ (of course, reversals by 180° are forbidden, because of the excluded volume constraint), one can vary the local intrinsic stiffness of the chain (cf. Fig. 4c, which illustrates this model for d = 2 dimensions). For $q_b = \exp(-\varepsilon_b/k_B T) = 1$ one recovers the standard SAW, while the limit $\varepsilon_b \rightarrow \infty$ corresponds to rigid straight rods. Following up on our previous work [11, 62, 70, 71, 97], we shall focus on this model in the present paper, applying the prunedenriched Rosenbluth method (PERM) [12, 101]. PERM is a biased chain growth algorithm with resampling and allows to get accurate data up to N = 50000for this model [70, 71], PERM yields a direct estimate of the partition function of a self-avoiding walk with N steps and N_{bend} 90°-bends

$$Z_N(q_b) = \sum_{\text{config}} C_{N,N_{\text{bend}}} q_b^{N_{\text{bend}}}$$
(35)

where $C_{N,N_{\text{bend}}}$ is the number of configurations of SAW's with N bonds and a number N_{bend} of $\pm 90^{\circ}$ turns. It would be interesting to extend the approach

POLYMER SCIENCE Series C Vol. 55 No. 1 2013

from athermal semiflexible chains ($q = \exp(\varepsilon/k_B T) = 1$) to semiflexible chains in solvents of variable quality (q > 1), which would mean an estimation of

$$Z_N(q,q_b) = \sum_{\text{config}} C_{N,N_{\text{bend}},N_{\text{pair}}} q_b^{N_{\text{bend}}} q^{N_{\text{pair}}}, \qquad (36)$$

with N_{pair} the number of nonbonded nearest neighbor pairs of monomers in the considered configuration. However, we are not aware of any study of the full problem, Eq. (36), yet.

Sampling suitable data on the monomer coordinates of the configurations that contribute to the partition function Eq. (35), one can obtain reasonably accurate estimates of the radii and of S(q), as defined in previous section.

The second model that is studied here is the bond fluctuation model of bottle-brush polymers. In the bond fluctuation model [102–104], each effective monomer blocks all eight corners of the elementary cube of the simple cubic lattice from further occupancy. Two successive monomers along a chain are connected by a bond vector $\vec{\ell}_b$, chosen from the set {(±2, 0, 0), (±2,±1,0), (±2,±1,±1), (±2,±2,±1), (±3,0,0), (±3,±1,0)}, including also all permutations. Originally configurations were relaxed by an algorithm where a monomer of the chain is chosen at random, and one also randomly chooses one of the six directions (±x, ±y, or ±z), respectively, and attempts to move the monomers by one lattice unit in the chosen direction. Of course, the move is accepted only if it does not violate excluded volume or bond length constraints. This move is called the "L6" move. Recently Wittmer et al. [9] provided evidence that a much faster algorithm results if one allows monomers to move to one of the 26 nearest and next nearest neighbor sites surrounding a monomer. With this "L26" move bonds can cross one another, and while such moves do not correspond to a real dynamics of macromolecules, it leads to a much faster exploration of phase space and hence a faster equilibration [105].

This model for linear polymers is generalized to the bottle-brush architecture by adding side chains at regular spacings $1/\sigma$ (which must be integer, e.g. for $\sigma = 1/2$ a side chain is attached to every second monomer of the backbone; the densest packing that is studied here is $\sigma = 1$). The side chains have chain length N_s , and are described by the bond fluctuation model as well. Furthermore, one more monomer is added to each chain end, to clearly identify the latter. The number N_b of monomers that constitute the backbone then is related to the number of side chains n_c via $N_b = (n_c - 1)/\sigma + 3$ and the total number of monomers of the bottle-brush polymer is $N_{tot} = N_b + n_c N_s$. For the sake of computational efficiency, the L26 move is combined with Pivot moves [100]. We refer to [17, 105] for implementation details.

As an example for well-equilibrated bottle-brush polymers as studied in [11, 17, 70, 71] and in the present paper, Fig. 6a shows selected snapshot pictures for side chain length $N_s = 24$ and various backbone chain lengths N_b . According to the visual impression, it seems rather natural to describe these bottle-brush polymers by the worm-like chain model, but as we shall see below, this conclusion would be totally misleading. Experimentalists often are led to a similar conclusion from microscope images of semiflexible polymers (e.g. DNA) adsorbed at a substrate (see e.g. [106–108]). However, such a conclusion is misleading for several reasons: (i) depending on the speed of adsorption of the polymer on the substrate, the conformation of the adsorbed polymer may be a frozen "projection" of the three-dimensional coil, which did not have enough time to relax to the two-dimensional equilibrium. (ii) In d = 2 dimensions, excluded volume forces render the Kratky–Porod (K–P) model of wormlike chains inapplicable [62, 66], one encounters a direct crossover from the rod regime to two-dimensional self-avoiding walk behavior (cf. Eq. (19)) when the contour length L exceeds the persistence length ℓ_n . One also should note that the persistence length of a polymer in d = 2 dimensions is not at all identical to the persistence length of the same polymer in d = 3dimensions [62, 66]. The experimental work (see, e.g., [106-108]) seems to be unaware of these problems and the resulting conclusions from this work need to be considered with care. It is also interesting to note that the snapshot pictures of the semiflexible SAW model (Fig. 6c) do not yield an immediate visual impression that the chains can be described by the K-P model, because of the 90° kinks; however, as we shall see, despite this difference of the local structure the statistical properties on the mesoscopic length scales are well described by the K-P model, for $q_b \leq 10^{-2}$ and n_p less than $n_p^*(\ell_p)$. Thus, we argue that on the basis of the inspection of AFM images of semi-flexible polymers one should be very careful on drawing conclusions which model is appropriate to describe these polymers.

SIMULATION RESULTS

Mean Square Gyration Radii and Their Analysis

We start with a description of our results for the mean square gyration radius $\langle R_g^2 \rangle$ of the semiflexible SAW model (similar data for the mean square end-toend distance $\langle R^2 \rangle$ of this model have already been presented elsewhere [70, 71]), Fig. 7. We clearly see that there are three regimes (Fig. 7a): in the chosen normalization where we divide out the asymptotic power law $\langle R_g^2 \rangle \propto N^{2\nu}$, we first have a regime where $3 \langle R_g^2 \rangle / \ell_b N^{2\nu}$ increases with $N^{2-2\nu}$. For small q_b this is interpreted as a rod-like regime; for $q_b \ge 0.2$ the chains are still too flexible, however, so a strictly rodlike behavior cannot yet be seen. Then a maximum occurs, and the ratio $3\langle R_g^2 \rangle / \ell_b N^{2\nu}$ decreases, before it settles down, after a second smooth crossover at a horizontal plateau (which according to Eq. (14) defines the value $\ell_p^{R_g}$). While this plateau for $q_b \ge 0.05$ is (pre-sumably) actually reached for N = 50000, the data also indicate that for $q_b \le 0.03$ even chains of length N =50000 are at least an order of magnitude too short to allow a direct convincing estimation of the amplitude value $\ell_p^{R_g}$. On the other hand, even for $q_b = 0.005$ (where we estimate from Eq. (4) that the persistence length ℓ_p is as large as $\ell_p \approx 52$ [62]) the slope of the data in the intermediate regime has not fully reached the theoretical value 1 - 2v, the slope of the data in Fig. 7a is still affected by crossover effects: the gradual crossover away from the Gaussian plateau towards the excluded volume-dominated behavior already starts when the gradual crossover from the rod-like regime to the Gaussian regime ends. Thus, even stiffer chains would be required to have a fully developed Gaussian behavior of the gyration radius. Figure 7b now attempts a scaling plot, where the persistence length estimates extracted from Eq. (4) were used to rescale $\langle R_g^2 \rangle$ in the K–P model representation (cf. Eq. (17)). It is evident that the rod-like regime and the onset of



Fig. 6. (a) Snapshot pictures of bottle-brush polymers as described by the bond-fluctuation model, for side chain length $N_s = 24$, and backbone chain length $N_b = (I)$ 99, (2) 195, (3) 387, and (4) 643. (b) Same as (a), but displaying the backbone of these bottle-brush polymers (c) Snapshot pictures of the SAW model with N = 25600 and two choices of q_b , $q_b = (I)$ 0.05 and (2) 0.005.

the crossover towards the K–P plateau are very well described by Eq. (17). For $q_b \ge 0.2$, of course, there is basically a direct crossover from the rod-like regime to the excluded volume dominated regime, but even then it is evident that the curves do not

superimpose on a master curve, as they do in d = 2 dimensions [62, 66], but rather splay out systematically, and the smaller q_b becomes (and hence the larger ℓ_p becomes) the more the data still are slightly above the K–P plateau.

Using the estimates for $n_p^*(q_b)$ extracted from the analysis of $\langle R^2 \rangle$ for this model in our previous work [70, 71], the data for $n_p \ge n_p^*$ do collapse on a simple straight line on the log–log plot, however (Fig. 7c).



For n_p near n_p^* the curves splay out, the master curve describing this second crossover from the K–P plateau to the excluded volume power law emerges as an envelope of the curves for individual values of q_b (which fall increasingly below the master curve in the crossover region the larger q_b is). Of course, $n_p^* \propto (\ell_p/\ell_b)^{\zeta}$ cannot produce a scaling of the crossover towards the rodbehavior, there the curves must splay out, irrespective of how small q_b is, but the deviation of the data from the horizontal K–P plateau moves more and more to the left of the plot the smaller q_b becomes.

Recalling that for the semiflexible SAW model the effective chain thickness D simply is $D = \ell_b = 1$, the Flory theory, Eq. (20), simply predicts $n_p^* \propto \ell_p^{\zeta}$; $\zeta = 2$, while the rod to Gaussian coil behavior occurs around $n_p = 1$, of course. Qualitatively, our data are in good agreement with these predictions, but not quantitatively: This is illustrated in Fig. 8, where n_p^* and $\ell_p^{R_g}$ are plotted in log–log form versus ℓ_p/ℓ_b . It is seen that instead of the theoretical value $\zeta = 2$ an exponent $\zeta = 1.5$ is observed. Now it is clear that Flory arguments imply also v = 3/5 instead of $v \approx 0.588$ [7], but this small difference cannot account for the large discrepancy encountered here. It would be desirable to study much larger values of N to confirm whether this discrepancy is a real effect (or our estimation of the crossover master curves in Fig. 7c for $\langle R_g^2 \rangle$ (and for in [71]) are systematically off). Thus, more work is still needed to fully clarify the situation.

It turns out that the behavior of our model for the bottle-brush polymers (which can describe actual scattering data for bottle-brush polymers very well, as demonstrated by Hsu et al. [17]) is much simpler: a plot of the mean square gyration radius $\langle R_{g,b}^2 \rangle$ of the

Fig. 7. (a) Log–Log plot of the relaxed mean square gyration radius $3\langle R_g^2 \rangle / \ell_b N^{2\nu}$ versus *N*, for chain lengths *N* up to N = 50000, and the stiffness parameter $q_b = (1) 0.005$, (2) 0.01, (3) 0.02, (4) 0.03, (5) 0.05, (6) 0.1, (7) 0.2, (8) 0.4, and (9) 1.0. The straight line with slope $2 - 2\nu$ shows the slope reached for small *N* in the rod-like regime, where $\langle R_g^2 \rangle = \ell_b^2 N^2 / 12$; the straight dotted line with slope $1 - 2\nu$ (for intermediate values of *N*) indicates the behavior expected for Gaussian chains, $\langle R_g^2 \rangle = \ell_b \ell_p N/3$. (b) Log–log plot of $3\langle R_g^2 \rangle / \ell_p L$ versus $n_p = N \ell_b / \ell_p$, using the same data as in (a) to test the K–P model (full curve). The different choices of q_b are shown by different symbols, as indicated. The slope that one expects for all q_b for $N \to \infty$, (2 ν – 1), is indicated by a broken straight line. (c) Same as (b), but choosing n_p/n_p^* as an abscissa variable



Fig. 8. Log–log plot of (1) n_p^* and (2) $\ell_p^{R_g}/\ell_b$, as indicated in the figure, versus ℓ_p/ℓ_b , using data for $q_b = 0.2$ to 0.005 (left to right). Dotted lines indicate the observed exponents while full straight lines show the Flory predictions for the exponents.

backbone versus backbone chain length N_b , for different side chain lengths N_s , Fig. 9a, normalized by $N_b^{2\nu}$ reveals a monotonic increase towards a plateau, there is not the slightest indication of a regime where the ratio $\langle R_{g,b}^2 \rangle / N_b^{2\nu}$ decreases, unlike the behavior of the semiflexible SAW (Fig. 7a). Thus, there is no evidence whatsoever for a Gaussian K–P plateau for this model. But the increase of the plateau value $\ell_p^{R_g}$ with increasing side chain length N_s does indicate that the chain

considerably stiffens, as N_s increases. However, this stiffening goes along with an increase in the effective chain thickness D. The latter can be estimated from the radial density profile (Fig. 10) by identifying the diameter D of the bottle-brush as D = $2R_{cs}(N_s) \equiv 2\sqrt{\langle R_{cs}^2 \rangle}$. Hsu et al. [70, 71] suggested to coarse-grain the bottle-brush along the backbone, dividing it into "blobs" along the chemical sequence of the backbone. The chemical distance s_{blob} along the backbone between its exit and entry points into a blob is found from a simple construction which assumes that the blobs are essentially spherical, so the geometrical distance $\Delta r(s)$ between exit and entry points of the backbone should be equal to D. Recording $\Delta r(s)$ for arbitrary s, Fig. 10a, using the equation $\Delta r(s_{\text{blob}}) =$ *D* allows us to simply read off the numbers s_{blob} for the choices of N_s , as illustrated in Fig. 10. The success of the rescaling shown in Fig. 9b shows that the persistence length ℓ_p of bottle-brushes simply is proportional to D. We also recognize that the asymptotic SAW-like behavior (where the horizontal plateau in Fig. 9 is reached) only occurs for about $N_b/s_{\text{blob}} \approx 60$. Comparing this behavior to Fig. 7b, we see that there the power law (for $q_b = 0.4$) or K–P plateau (for small q_b) is reached for $n_p = N\ell_b/\ell_p \approx 40$, Roughly, these successive blobs then are equivalent to one persistence length. This comparison suggests that for the bottlebrushes we should identify $\frac{2}{3}(N_b/s_{blob})$ with n_p , i.e. the number of monomers along the backbone corre-



Fig. 9. (a) Plot of the normalized mean square gyration radius of the backbone of the model for bottle-brush polymers, $3\langle R_{g,b}^2 \rangle/\bar{\ell}_b N^{2\nu}$, versus the backbone chain length N_b , for four different side chain lengths $N_s = (I) 6$, (2) 12, (3) 18 and (4) 24. Here $\bar{\ell}_b \approx 2.7$ is the average bond length of the single bond in the bond fluctuation model under good solvent conditions. The horizontal plateaus allow to extract the estimates for $\ell_p^{R_g}$ defined in Eq. (14). (b) Same data as in (a), but ordinate is rescaled with $\ell_p^{R_g}$ so for $N_b \to \infty$ all data converge to one, and abscissa is rescaled by the effective blob-size s_{blob} (see text), on a log–log plot. Straight line indicates the rigid-rod behavior, with slope $2 - 2\nu$ in this representation.



Fig. 10. (a) End-to-End distance $\Delta r(s)$ of subchains containing *s* successive backbone monomers for $N_s = 6$, 12, 18, and 24. The horizontal solid lines and the numbers shown on the ordinate indicate the choices $\Delta r(s) = 2R_{cs}(N_s)$, from which the corresponding values of s_{blob} can be read off (vertical straight lines), namely, $s_{blob} = 6$, 10, 12, and 14 for $N_s = (1) 6$, (2) 12, (3) 18, and (4) 24, respectively. $R_{cs}(N_s)$ is the cross-sectional radius, which is extracted from the radial monomer density profiles $\rho(r)$ as shown in (b). (b) Radial monomer density profiles $\rho(r)$ in planes locally perpendicular to the backbone of bottle-brush polymers with backbone length $N_b = 1027$ plotted versus radial distance *r* for the same side as in (a) chain lengths N_s . The cross-sectional radius then follows as $\langle R_{cs}^2 \rangle = 2\pi \int_0^\infty r dr \rho(r) r^2$ with the density profile being normalized as $2\pi \int_0^\infty r dr \rho(r) = N_s$. The values of R_{cs} are pointed out by arrows.

sponding to one persistence length is $\frac{3}{2}s_{blob} = 9, 15, 18$ and 21 for $N_s = 6$, 12, 18 and 24, respectively. Noting that the average bond length ℓ_b in the bond fluctuation model is $\overline{\ell}_b = 2.7$, we would obtain persistence lengths $\ell_p(N_s) = \frac{3}{2}D \approx 17, 27, 34 \text{ and } 41 \text{ for } N_s = 6, 12, 18$ and 24, respectively. The result that $\ell_p(N_s)$ is of the same order as D irrespective of the side chain length agrees with early theoretical predictions [18, 19] but is at variance with the result of Fredrickson [20] who predicted a much faster increase of ℓ_p with N_s . However, Feuz et al. [30] pointed out that the result of Fredrickson [20] can only be expected to hold for extremely long side chains, such as $N_s = 1000$. Such long side chains are neither relevant for simulations nor for experiment, however. We stress that the range of N_s accessible to simulations (Figs. 9, 10) nicely corresponds to the range of N_s studied experimentally [28, 32-38].

The mapping performed in Fig. 9 means that we have coarse-grained the bottle-brush polymers (Fig. 4a) into an effective bead-spring model (Fig. 4d). If this mapping is taken literally, it can also be used to obtain the resulting coarse-grained contour length L_{cc} (Fig. 3) as

$$L_{cc} = 2R_{cs}(N_s)N_h/s_{\text{blob}}(N_s). \tag{37}$$

Instead of the "chemical" contour length $L = N_b \ell_b \approx 2773$ a reduced length is found, namely $L_{cc} \approx$

1989, 1824, 1963 and 2016, for $N_s = 6$, 12, 18 and 24, respectively. This means that the coarse-grained contour length L_{cc} is about 30% smaller than the "chemical" contour length in this model.

Analysis of the Structure Factor

We now turn to the structure factor of the semiflexible SAW model presenting Kratky plots for two contour lengths, L = 400 and L = 25600, in Fig. 11. As expected, cf. Fig. 5, one first has a linear increase with qL, then a round maximum followed by a decrease which then gradually settles down at a horizontal plateau, that again is compatible with the theoretical prediction, π .

While for the short chain length (L = 400) the agreement with the theoretical prediction (due to Kholodenko [89], Eqs. (28)–(30), which were found [97] to be numerically almost indistinguishable from the exact result provided by Stepanow [95, 96]) is almost perfect, for the very long chains (L = 25600) we note systematic deviations between Kholodenko's prediction [89] and the data for relatively large q_b near the maximum of the Kratky plot. This must be expected, since the input in the Kholodenko formula is just the persistence length ℓ_p {which we have extracted from Eq. (4)} and implicit in the theory is the Gaussian prediction for $\langle R_g^2 \rangle$, namely $\langle R_g^2 \rangle = L\ell_p/3$ {Eq. (13)}. As seen in Fig. 7, for $q_b = 0.2$ already rather clear deviations from this result occur for N = 25600,

while for small N such as N = 400 such deviations still are rather small. In contrast, in the Debye formula the correct (as observed) value of $\langle R_g^2 \rangle$ was used as an input: then deviations from the Debye function are only seen near the region where the crossover to $qLS(q) = \pi$ starts to set in at large q (the Debye function does not describe this crossover at all). Since the shape of the Kholodenko function always is rather similar to the actual function, it is obvious that one always can fit the data to the Kholodenko function, if ℓ_p is not known: however, the resulting fitted persistence length will be systematically too large, if excluded volume effects are present as shown in Fig. 11c.

To elucidate the significance of excluded volume on the structure factor further, we show a magnification of the region near the maximum for $q_b = 1, 0.2$, and 0.005 in Fig. 12. It is seen that the identification of the two power laws suggested for the decay of qS(q) in the region beyond the maximum of the Kratky plot is rather subtle. In particular, for rather stiff chains the crossover to the rod-like scattering sets in rather early, so for the clear identification of power laws the available range of q simply is not large enough. This very gradual crossover between the three different regimes (rods to Gaussian coils to coils swollen by the excluded volume interaction) complicates the data analysis, if only a restricted range of parameters (such as the chain length N and the wavenumber q) can be investigated.

The smoothness of the crossover also becomes evident when one studies the dependence of the position of the peak in the Kratky-plot (and its height) on the persistence length (Fig. 13). Typically, the data fall neither in the regime where strict Gaussian behavior occurs, nor in the regime where excluded volume scaling is fully developed.

Despite all these difficulties due to the gradual crossovers, the semiflexible SAW nevertheless is a rel-

atively simple case, since one knows that here $D = \ell_b(=1)$, and ℓ_p can be varied over a wide range by variation of q_b , keeping all other parameters constant, and moreover ℓ_p can be estimated precisely from the initial decay of the bond vector autocorrelation function (or, equivalently, from Eq. (4)). For the second model studied here, bottle-brush polymers under good



Fig. 11. Kratky plot of the structure factor of the semiflexible self-avoiding walk model, $\log -\log \operatorname{plot} \operatorname{of} qLS(q)$ versus qL, for two chain lengths, namely (a) L = 400 and (b) L = 25600. (c) Persistence length ℓ_p plotted vs. q_b for $0.005 \le q_b \le 0.4$. In (a)(b) several choices of q_b are included, namely $q_b = (1) 0.2$, (2) 0.05, (3) 0.02, (4) 0.01, and (5) 0.005. The scattering functions of a rigid rod and the Debye function are included, as well as the prediction of Kholodenko {Eqs. (28)-(30)}. The predicted large qlimit of π is indicated. In the Debye function the observed value of $\langle R_g^2 \rangle$ was used as an input, while for the Kholodenko formula the persistence length (estimated from Eq. (4)) was used as an input. In (c) the persistence length is taken as the best fitting parameter such that the prediction of Kholodenko formulas describes the correct maximum in the Kratky plot for our simulation data of chain lengths (1) L = 400 and (2) 25600. The estimates using (3) Eq. (1) and (4) Eq. (4) are also shown in (c) for comparison.



Fig. 12. Kratky plot for $q_b = (1) 0.005$, (2) 0.2, and (3) 1.0 plotted in a rescaled form, $qS(q)/(qS(q))_{max}$ vs. q/q_{max} , restricting the ordinate range to the decade from 1 to 0.1 and the abscissa to the range from 1 to 10. The theoretical power laws (slope = $1 - 1/\nu \approx 2/3$ and slope = -1, respectively, as predicted in Fig. 5) are included.



Fig. 13. Log–log plot of (1) q_{max} and (2) $(qS(q))_{\text{max}}$ versus ℓ_p (as estimated using Eq. (4)) using data for $q_b = 0.2$ to 0.005. Straight lines indicated the exponents $q_{\text{max}} \propto \ell_p^{-1/5}$ and $q_{\text{max}} \propto \ell_p^{-1/2}$ that one expects according to the Flory treatment in the excluded volume region and Gaussian region, respectively. All data were taken for N = 50000.



Fig. 14. Kratky plot of $qLS_b(q)$ versus qL for bottle-brush polymers with (a) $N_b = 131$ and (b) 259, where " $L = N_b \overline{\ell}_b$ " is the "chemical" contour length and $S_b(q)$ is the scattering function of the backbone only. Four side chain lengths $N_s = (1) 6$, (2) 12, (3) 24 and (4) 48 are included, as indicated. The horizontal straight line shows the Holtzer plateau ($=\pi$) if the coarse-grained contour length could be identified with the chemical contour length.

solvent conditions, we have seen that varying the side chain length N_s we change D and ℓ_p together, and also the coarse-grained contour length L_{cc} is significantly smaller than the chemical contour length $N_b \overline{\ell}_b$, and it is nontrivial to estimate L_{cc} accurately.

In previous work [11, 17] we have already considered the decomposition of the total scattering function of bottle-brush polymers into the scattering from the backbone and from the side chains. This analysis which has the advantage that it provides a direct link to corresponding experiments [32, 33] will not be addressed here, but we rather focus on the scattering function of the backbone only. Figure 14 shows Kratky plots for relatively short backbone chain lengths $(N_b = 131 \text{ and } 259, \text{ respectively})$. One recognizes that for short side chain lengths $(N_s = 6, 12) qLS_b(q)$ does not settle down to a well-defined "Holtzer Plateau", at least not within the available window of wavenumbers. Clearly, also the range over which $qLS_b(q)$ decays from the maximum to the horizontal part that appears for



Fig. 15. Log–log plot of $qS_b(q)$ vs. q for bottle-brush polymers with side chain length $N_s = 24$ and various backbone chain lengths $N_b = (1)$ 131, (2) 195, (3) 259, (4) 387, (5) 515, (6) 771, and (7) 1027. The point of intersection between two broken straight lines illustrates the estimation of $q^*(N_b)$, the wavenumber where the onset at the Holtzer plateau occurs.

 $N_s = 24$ and 48 is rather small, and does not warrant any analysis in terms of the power laws suggested in Fig. 5b. This mismatch between the actual plateau values (for $N_s = 24$ and 48), which are close to 4, and the theoretical value π can be attributed to reduction of L_{cc} in comparison to $N_b \overline{\ell}_b$ since the actual orientations of the backbone vectors are not strictly aligned with the coarse-grained backbone (Fig. 4), as is also evident from the fact that $\langle \vec{a}_i \cdot \vec{a}_{i+1} \rangle / \langle \vec{a}_i^2 \rangle$ (Fig. 2b) is already reduced to about 0.7, but the further decrease of $\langle \vec{a}_i \cdot \vec{a}_{i+s} \rangle / \langle \vec{a}_i^2 \rangle$ is rather slow, due to the side chain induced stiffening of the backbone on mesoscopic scales. The ratio at about $4/\pi$ is compatible with the reduction of L_{cc} by about 30% relative to L noted previously, so gratifyingly our analysis is internally consistent.

Figure 15 shows plots of S(q) vs. q for fixed side chain length $N_s = 24$ but different backbone chain lengths. In this plot, an attempt is made to locate an onset wavenumber q^* for the Holtzer plateau, in terms of a fit of two intersection straight lines. Of course, the data are smooth and the onset of the Holtzer plateau does not occur sharply but rather gradual; thus q^* can be estimated only with considerable error (for large N_b we estimate $q^* \approx 0.06 \pm 0.01$, while for $N_b = 131$ the estimate rather is $q^* \approx 0.075 \pm 0.020$ [11]). Now the question is, how can one relate q^* explicitly to the persistence length? Should one take $\ell_p = 2\pi/q^*$, or $\ell_p =$



Fig. 16. (a) Plot of qLS(q) vs. $(q\ell_p)^{-1}$ for the semiflexible SAW and L = 25600, including 4 choices of q_b , as indicated. (b) Same as (a) for bottle-brush polymers with fixed side chain length $N_s = 24$ and varying backbone length N_b as indicated. Broken straight line is the des Cloizeaux [80] prediction, Eq. (26), full straight line an empirical fit to the data.

 $1/q^*$? Lecommandoux et al. [28] who were the first to try such a method suggested the relation $\ell_p \approx 3.5/q^*$, but we see little theoretical support for this choice either.

It would be advantageous if one could rely on the des Cloizeaux relation, Eq. (26), which suggests to plot qS(q) vs. 1/q for $q\ell_p \ge 1$: one should find a straight line, the intercept at the ordinate should yield π , the slope of the straight line should yield $2/(3\ell_p)$.

However, when one tests this method for the semiflexible SAW, one finds that the data that can be fitted to a straight line are at $q\ell_p \approx 1$ rather than at $q\ell_p \gg 1$, and the slope of the straight line disagrees with the prediction (Fig. 16a). Thus, it is not really a big surprise



Fig. 17. (a) Plot of $qL_{cc}S_b(q)$ (q) versus $L_{cc}q$ for bottlebrush polymer with side chain length $N_s = 24$ and several values of $N_b = (I)$ 131, (2) 259, (3) 387, (4) 515, (5) 771, and (6) 1027. For each choice of N_b both L_{cc} and ℓ_p were individually adjusted. (7) The fit of Kholodenko formulas (Eqs. (28)–(30)). (b) Plot of L_{cc} vs. N_b , showing that $L_{cc} \approx 2.03N_b$ rather than being equal to the "chemical" contour length $L = \overline{\ell}_b N_b \approx 2.7N_b$. (c) Plot of the persistence length ℓ_p (from the fit in (a)) versus N_b . In the loglog plot (inset), the theoretical power law $\ell_p \propto N_b^{2\nu-1}$ is shown by the straight line.

that this does not work well for our bottle-brush model either (Fig. 16b).

An interesting alternative of data analysis is, however, a fit of the Kholodenko Eqs. {formulas (28)– (30)} to the structure factor, using both L_{cc} and ℓ_p as individual adjustable parameters for each value of N_b (Fig. 17). First of all, one sees that the Kholodenko structure factor provides a good fit in all cases, and the result for the coarse-grained contour length, $L_{cc} =$ $2.03N_b$, even is physically very reasonable: we have obtained that there is a 30% reduction of the L_{cc} in comparison to the "chemical" contour length L = $N_t \tilde{\ell}_t = 2.7N_t$ in the previous subsection.

 $N_b \ell_b = 2.7 N_b$ in the previous subsection. However, the problem of this fit is the unphysical behavior of the persistence length ℓ_p : since we know that the Kholodenko [89] approach involves necessarily the Gaussian result $\langle R_g^2 \rangle = \frac{1}{3} \ell_p L_{cc}$ but we know that for our model $\langle R_g^2 \rangle \propto N_b^{2\nu}$ and $L_{cc} \approx 2.03 N_b$, the only way to reconcile these results is a persistence length scaling as $\ell_p \propto N_b^{2\nu-1}$, and this is what we see in Fig. 17c. Thus, despite the seemingly good fit (Fig. 17a) and good results for L_{cc} (Fig. 17b), the results for the persistence lengths are completely unreliable!

In order to apply this approach, one must make sure that one works with data in the Gaussian regime, and this is not at all the case for bottle-brush polymers under good solvent conditions.

CONCLUSIONS

In this paper, we have focused on the behavior of single semiflexible polymers under very good solvent conditions, considering how the chain stiffness affects polymer properties such as the mean square gyration radius, the structure factor, etc. Our analysis focused on the question how the variation of chain stiffness affects these properties, and hence one can infer from these properties a characterization of the "intrinsic stiffness" of the polymer chain in terms of the socalled "persistence length".

We have contrasted two models, the self-avoiding walk on the simple cubic lattice where a bending energy ε_b causes pronounced stiffening of the polymer when $\varepsilon_b \ge k_B T$, and a lattice model for bottle-brush polymers, where backbone stiffening is caused by increasing the length of side chains. These two models constitute two quite distinct limiting cases: in the SAW model, increase of $\varepsilon_b/k_B T$ causes stiffening without any effect on the local thickness of the chain, which strictly remains the lattice spacing. For the bond fluctuation model of polymer brushes, however, we have found that backbone stiffening is caused by the thickness of the (coarse-grained) worm-like chain, the per-

sistence length increases proportional to the crosssectional diameter of the bottle-brush.

Since snapshot pictures (Fig. 6) suggest that the bottle-brush polymers (or their backbones, respectively) resemble worm-like chains (and the same conclusion is often drawn from AFM pictures or electron micrographs of actual polymers), the use of the Kratky-Porod worm-like chain model has become very popular. However, we demonstrate here that for bottle-brush polymers this model yields very misleading results: since the mean square gyration radii of bottle-brushes are found to scale with their contour length L as $\langle R_g^2 \rangle \propto L^{2\nu}$, the Kratky–Porod (K–P) result $\langle R_g^2 \rangle = \ell_p L/3$ invariably causes a spurious contour length dependence of the persistence length when fit to the data, namely $\ell_p(L) \propto L^{2\nu-1} \to \infty$ as $L \to \infty$. Although the fits of the K–P model look almost perfect (Fig. 17a) and numbers for L resulting for the contour length from the fit are rather reasonable, the result for "the" persistence length simply is meaningless!

Already in our earlier papers we have shown that similar ambiguous results for the persistence length are gotten when orientational correlations along the chain backbone are analyzed, or the projection of bond vectors on the end-to-end distance are studied (although the resulting numbers for $\ell_p(N_b)$ seem to be somewhat smaller than those shown in Fig. 17c). The large qbehavior of the structure factor S(q) yields a qualitatively more reasonable behavior, but a unique choice for a well-defined persistence length as a measure for intrinsic chain stiffness does not emerge. All these difficulties in understanding the stiffness of bottle-brush polymers in good solvents are intimately linked to the fact that one can coarse-grain into some effective selfavoiding walk model (Figs. 9, 10), and no regime exists where the polymers resemble Gaussian chains. Of course, this fact is different if we would consider bottle-brush polymers in Theta-solvents (as done by Theodorakis et al. [98]), since then $\langle R_g^2 \rangle \propto L$ and the use of the K–P model is qualitatively reasonable. Another interesting possibility to extract a persistence length of bottle brushes would be an attempt to estimate an effective bending modulus. One would have to estimate the coarse-grained free energy of bent versus non-bent configurations of suitable pieces of bottlebrush polymers, which in principle can be deduced from sampling suitable angular distribution functions for such sub-chains. However, the implementation of such an approach is not straightforward and has not been attempted.

We have found that the situation in some respects is simpler if one considers polymers where the stiffness can be enhanced while keeping their thickness constant, as modeled by a semiflexible extension of the standard SAW model. Then an intermediate Gaussian-like behavior of the mean square radii and the structure factor emerges, and this can be understood theoretically (Fig. 5), at least in qualitative terms. While still the asymptotic decay of bond vector autocorrelation functions is unsuitable to infer anything about the intrinsic stiffness (due to the fact that the asymptotic decay is not exponential but rather described by a power law), in favorable cases the initial decay of these autocorrelation functions provided useful estimates of the persistence length, which then can be used as input in the K–P model. While still some problems occur to understand for very long chains the crossover between the K-P model and the ultimate SAW behavior quantitatively, in qualitative terms the situation is understood. We emphasize, however, that all these comments only address the three-dimensional case: in d = 2 dimensions, the K–P model does not work at all, and one has a direct crossover from rod-like polymers to SAW's.

It is hoped that our analysis will help experimentalists with a proper interpretation of their data on semiflexible polymers.

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