

First-Order Theory of Self-Avoiding Walks Based on Loop Exclusion

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Self-avoiding walks may be constructed through a progressive exclusion of walks with loops. A study of the process leads to critical exponents $\nu \approx (4+D)/4D$ and $\gamma \approx 8/(4+D)$ for dimension $4 \geq D \geq \frac{4}{3}$. The equations agree with the ϵ expansion to first order, fit the (known) values for $D = 3, 2$, and also those (suggested) for $D < 2$. The probability of an exclusion due to a loop of length j appears to be asymptotically equal to $(\gamma-1)j^{-2}$, for $4 > D > \frac{4}{3}$ ("strong universality").

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Self-avoiding walks (SAW's) of length $N \gg 1$ are described with the help of critical exponents ν and γ ,

$$R_N^2 = AN^{2\nu}, \quad (1)$$

and

$$W_N = A'\mu^N N^{\gamma-1}, \quad (2)$$

where R_N^2 and W_N are the mean square end-to-end distance and the number of SAW's, respectively; A , A' , and μ are lattice dependent, but ν and γ depend on the dimension D alone. For $D \geq D_c = 4$ we have the ideal values $\nu = \frac{1}{2}$ and $\gamma = 1$; at $D = 1$ SAW's constitute rigid rods, $\nu = 1$ and $\gamma = 1$. The theory of critical phenomena gives very accurate (or exact) ν and γ for $D = 4 - \epsilon$, 3, and 2.¹⁻³ However, the dependence on a continuous variable $1 < D < 4$ is also of great interest, e.g., in connection with attempts to relate random fractals, like percolating clusters and branched polymers, to SAW's at an equivalent (noninteger) dimension.^{4,5} For that purpose one may utilize an equation $\nu = 3/(D+2)$, derived by Flory⁶ with the help of a mean-field approximation. This equation disagrees with the (exact) ϵ expansion already to the first order. Still, it gives a quite accurate value for $D = 3$ and the exact values for $D = 2$ and 1, but the physical reason for its remarkably good performance is puzzling.⁷ Furthermore, its validity in the region $1 < D < 2$ is uncertain in view of recent Monte Carlo experiments.⁸ Finally, no similar equation has been derived for γ .

SAW's may be derived from random walks with the help of a progressive exclusion of walks with loops. A construction of SAW's through "dimerization"^{8,9} offers one possible way to carry out this progressive exclusion. (Alternative constructions linked to exclusion are the conventional stepwise growth, $N \rightarrow N+1$, requiring exclusion of loops formed by the newly added step, or the progressive increase of the excluded loop length in restricted

walks of constant N .¹⁰ This Letter studies in detail the loop exclusion in the course of a dimerization. On this basis it derives approximate analytical equations, correct to $O(\epsilon)$, for $\nu(D)$ and $\gamma(D)$.

Pairs of "monomer" SAW's of length $N \gg 1$ are linked together to produce "dimer" walks of length $2N$. Dimers in which one half intersects the other, forming a loop, are excluded. (We need not worry about internal intersections since each monomer is a SAW.) Upon completion of the exclusion, the remaining dimers constitute an ensemble of SAW's of length $2N$. Clearly $W_{2N} = (W_N)^2 f_{\text{suc}}$, where f_{suc} is the fraction of the ("successful") dimers in which one half does not intersect the other. In view of Eq. (2), $W_{2N} = (A'\mu^N N^{\gamma-1})^2 f_{\text{suc}} = A'\mu^{2N} (2N)^{\gamma-1}$, giving

$$f_{\text{suc}} \sim N^{-(\gamma-1)}. \quad (3)$$

The exclusion of dimers in which an m th and an n th segment, belonging to the first and to the second monomer, respectively, form a loop of length $j = m + n$ [see Fig. 1(a)] is carried out in a succession of steps. n is varied from 1 to N and, for each n , m is varied from 1 to N . $\phi_{m,n,N}$ is the loop's probability at the (mn) th step. More precisely, $\phi_{m,n,N}$ is the fraction of dimers in which the m th and the n th segments form the loop $j = m + n$ in an ensemble of dimers in which, on the second monomer, (i) segments 1 to $n-1$ are already completely self-avoiding with respect to the first monomer, (ii) the n th segment is self-avoiding for loops of length up to $j-1$ only, and (iii) segments $n+1$ to N are not yet self-avoiding with respect to the first monomer. We have [using $x = \exp(\ln x)$ and expanding $\ln x$]

$$\begin{aligned} f_{\text{suc}} &= \prod_{n=1}^N \prod_{m=1}^N (1 - \phi_{m,n,N}) \\ &= \exp \int_1^N \int_1^N (-\phi_{m,n,N} + \dots) dm dn. \end{aligned} \quad (4)$$

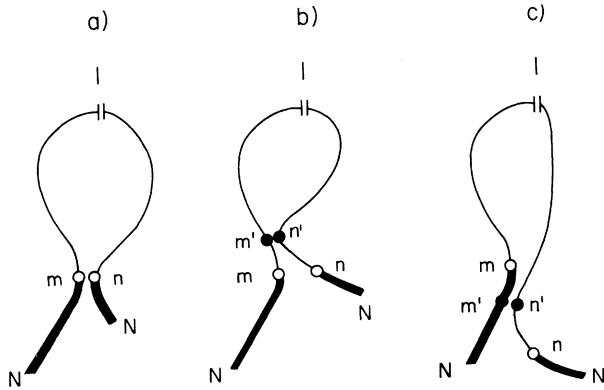


FIG. 1. Geometrical effects of the exclusion of dimers with loops: (a) Segments m and n , on the first and second monomer, respectively, form a loop $j = m + n$; the effective dimer's length is shortened from $2N$ to $(N - m) + (N - n) = 2N - j$. (b) Segments $m' < m$ and $n' < n$ form an "enclosed" loop $j' = m' + n'$, shortening the effective length between m and n from j to $j - j'$. (c) Segments $m' > m$ and $n' < n$ form a "crossed" loop $j' = m' + n'$, shortening the effective length between m and n from j to $m' - n' = j' - 2n'$ (for $m = n = j/2$).

If we assume

$$\phi_{m,n,N} \rightarrow \phi_j = (\gamma - 1)j^{-2}, \quad \text{for } j \gg 1, \quad (5)$$

then Eq. (4) at once reproduces Eq. (3) to within leading order of N . True, other expressions, not depending on j alone, might achieve as much. Yet $\phi_{m,n,N}$ should be independent of N , as well as of m and n separately, with the exception of the crossover regions $j \rightarrow N$ and $(m \text{ or } n) \rightarrow j$. Ruling out that these crossover regions provide the dominant contribution to the double integral justifies Eq. (5). Equation (5) predicts a *strong universality*, a power law independent of $D < D_c$. The coefficient equals $\gamma - 1 = O(\epsilon)$. Consequently, correlations between multiple loops are presumed to contribute terms of order higher than first in ϵ . The present estimates are limited to the first order alone.

We now consider the increase of R_N^2 upon dimerization. Prior to the exclusions, the *random linking* of pairs of monomers of (average) R_N^2 produces dimers of $R_{2N}^{2*} = 2R_N^2 = 2AN^{2\nu}$, with the asterisk denoting the stage preceding exclusions. Upon completion of the sequence of exclusions, we are left with true SAW's, having $R_{2N}^2 = A(2N)^{2\nu}$. The corresponding swelling is

$$\Delta = R_{2N}^2 / R_{2N}^{2*} = 2^{2\nu-1}. \quad (6)$$

Δ is decomposed into stepwise variation

$$\Delta = \prod_{n=1}^N \prod_{m=1}^N \delta_{n,m},$$

where $\delta_{n,m}$ is the ratio of R^2 's, after and before the exclusion at an (m,n) th step. Prior to this exclusion the dimers consist of two fractions, ϕ_j and $1 - \phi_j$, respectively, in which the n th and m th segments do, or do not, form the loop $j = m + n$. The ϕ_j fraction has its *effective* length shortened from $2N$ to $2N - j$ [see Fig. 1(a)]; the $1 - \phi_j$ fraction is unaffected. $\delta_{n,m}$ is the ratio of (the average) R^2 after and prior to the exclusion of the ϕ_j fraction. Thus, to a first approximation,

$$\delta_{n,m} \approx (2N)^{2\nu} / [\phi_j (2N - j)^{2\nu} + (1 - \phi_j) (2N)^{2\nu}].$$

Introducing Eq. (5) and expanding $(2N - j)^{2\nu}$ in a series leads to

$$\delta_{n,m} \approx 1 + \nu(\gamma - 1)/Nj. \quad (7)$$

Higher-order terms of the expansion are proportional to $(2\nu - 1)(\gamma - 1) = O(\epsilon^2)$ and are therefore neglected. The same applies to the neglected progressive variation of R_x^2 ($x = 2N$ or $2N - j$) by a factor $\approx 2^{2\nu-1}$, as n varies from 1 to N [cf. Eq. (6)]. Furthermore, R^2 of the length $2N - j$ is affected by the presence of extra "dangling" lengths m and n . However, in view of the remark following Eq. (5), this should contribute terms that are of order higher than first in ϵ . Transforming $\prod \prod \delta_{n,m}$ into $\exp[\int \int (\delta_{n,m} - 1) \dots]$ [similarly to Eq. (4)] gives $\Delta \approx 2^{2\nu(\gamma-1)}$. Together with Eq. (6) this leads to

$$(2\nu - 1)/2\nu \approx \gamma - 1. \quad (8)$$

The above estimate of the effect of exclusions on R_{2N}^2 may be reexpressed more formally as

$$R_{2N}^2 = D^*(N_{\text{eff}}) = 2AN_{\text{eff}}^{2\nu}. \quad (9)$$

The notation requires explanation: The "effective" length $N_{\text{eff}} = N \times [\Delta(N^{2\nu})]^{1/2\nu}$, where $\Delta(N^{2\nu})$ denotes the swelling $\Delta = 2^{2\nu(\gamma-1)}$ estimated in Eqs. (7) and (8). The present notation stresses that Δ has been computed with respect to N raised to the power 2ν , which is the quantity associated with R_{2N}^2 ; in the expression for N_{eff} the $1/2\nu$ root of $\Delta(N^{2\nu})$ is taken to express it as a length. Finally, D^* denotes an operator corresponding to a *random linking of effective lengths*, saying "Take N_{eff} , raise it to power 2ν , multiply by A [cf. Eq. (1)], and double the result, to obtain R_{2N}^2 of the dimer SAW's." [One easily verifies that the recipe reproduces Eq. (8).]

Equation (9) helps us to estimate a second effect of exclusions, notably upon the dependence of ϕ_j on j . We consider again the pair of segments m and n on the two monomers. Prior to the exclusions,

the monomers are mutually uncorrelated. Hence the probability that m and n intersect one another, forming the loop $j = m + n$, scales as the segments' density in a hypersphere in D space, viz., $\phi_j^* \sim R_j^{-D} \sim j^{-\nu D}$. (Here we take for the sake of simplicity $m \simeq n \simeq j/2$.) This initial dependence applies to dimers in which other segment pairs, n' and m' , may form loops $j' = m' + n'$. If $1 \leq n' < n$ (while $1 \leq m' \leq N$), such dimers are excluded in the sequence preceding the (m, n) th step. This causes the drop from the initial ϕ_j^* to $\phi_j \sim j^{-2}$ [cf. Eq. (5)], by the time the process attains the n th step. In analogy with Eq. (9), this drop is estimated by assuming

$$\phi_j \sim \phi^*(j_{\text{eff}}) \sim (j_{\text{eff}})^{-\nu D}. \quad (10)$$

Here, however, $j_{\text{eff}} = j \times [\Delta(j^{-2})]^{-1/2}$. Or, the effect of exclusions is estimated with respect to the reciprocal square length, j^{-2} , since this quantity is associated with ϕ_j . Except for that difference, the estimation of $\Delta(j^{-2})$ parallels entirely that of $\Delta(N^{2\nu})$ before. At an (m', n') th step the dimers consist of two fractions, $\phi_{j'}$ and $1 - \phi_{j'}$, respectively, with or without the loop $j' = m' + n'$. We first consider the case $m' < m$ [illustrated in Fig. 1(b)], where an "enclosed" loop j' shortens the effective length between m and n from j to $j - j'$. Similarly to $\delta_{n,m}$ the ratio of the reciprocal square lengths, after and prior to the exclusion of the $\phi_{j'}$ fraction, is estimated as $\delta(j^{-2})_{n', m' < m} \simeq j^{-2} / [\phi_{j'}(j - j')^{-2} + (1 - \phi_{j'})j^{-2}]$. For a "crossed" loop, $m' > m$ (see Fig. 1(c)), the length is shortened from j to $j - 2n'$, for $j' < j + 2n'$ (for longer j' there is no effect). $\delta(j^{-2})_{n', m' > m}$ is therefore estimated correspondingly, with $(j' - 2n')^{-2}$ replacing $(j - j')^{-2}$. The total variation, $\Delta(j^{-2})$, is obtained by taking products of $\delta(j^{-2})_{n', m'}$ over m' and n' . Passing to integrals [again, as in Eq. (4)] gives

$$\Delta(j^{-2}) \sim j^{-2(\gamma-1)}, \quad (11)$$

where the decrease is shared equally by the enclosed and crossed loops. [It is important to note that $\Delta(j^{-2})$ diverges with j , whereas $\Delta(N^{2\nu})$ is merely equal to a factor; this difference seems to be related to the additional divergence of small intersegmental distances.¹¹] From Eq. (11) it follows that $j_{\text{eff}} = j \times [\Delta(j^{-2})]^{-1/2} \sim j^\gamma$. Introducing this into Eq. (10) and comparing with Eq. (5) leads to

$$\phi_j \sim j^{-\gamma\nu D} \sim j^{-2}. \quad (12)$$

Therefore $\gamma\nu D = 2$. Combining this result with Eq. (8) gives

$$\nu = (4 + D)/4D, \quad (13a)$$

and

$$\gamma = 8/(4 + D). \quad (13b)$$

Expansion at $D = 4 - \epsilon$ gives $\nu = \frac{1}{2}(1 + \epsilon/8 + \epsilon^2/32)$ and $\gamma = 1 + \epsilon/8 + \epsilon^2/64$ which agrees with the correct expansion¹ to first order, but gives a much too small second-order term. At $D = 3$, present results are $\nu = 0.583$ and $\gamma = 1.142$ to be compared with the accurate estimate² $\nu = 0.588$ and $\gamma = 1.162$. At $D = 2$ the result reproduces the presumably exact value³ $\nu = \frac{3}{4}$ and gives $\gamma = \frac{4}{3}$, believed until recently to be exact, but slightly smaller than the very recent³ $\gamma = 1.344$. Throughout $2 \leq D \leq 3$, Eq. (13a) and the Flory equation differ very slightly. For $D < 2$, however, Eq. (13a) predicts a much steeper increase, comparable to that of Ref. 8. Thus Ref. 8, Eq. (13a), and the Flory equation give, respectively, $\nu(1.5) = 0.94, 0.92,$ and 0.86 ; $\nu(\frac{4}{3}) = 1, 1,$ and 0.90 . Below $D = \frac{4}{3}$, Eq. (13a) gives a nonphysical $\nu > 1$. This is interpreted as a transition to rigid rods, $\nu = 1$, already at $D = \frac{4}{3}$; at that point γ reaches $\frac{3}{2}$.

The strong universality of progressively excluded loops, $\phi_j \sim j^{-2}$ for $D < D_c$ (above D_c one has the ideal $\phi^0 \sim j^{-D/2}$), plays a major role here. A similar power law has been proposed^{12,13} for somewhat differently defined "tadpole" loops; see, however,, Ref. 14. (The present loops are relevant to a construction of SAW's through dimerization, $N \rightarrow 2N$; the tadpole loops are relevant to the stepwise construction $N \rightarrow N + 1$.) Finally, one should note that the basic assumptions of the present and the Flory theory are entirely different. In the Flory theory, the (mean-field) binary repulsions are balanced by the (ideal) entropy of stretching, both terms greatly overestimated.⁷ Here ν (of swelling) and γ (of exclusion) become interrelated; first, because exclusion causes swelling, and second, because swelling decreases exclusion, keeping it at the marginal level $\phi_j \sim j^{-2}$. A more negative power would make the exclusion negligibly small; a less negative power would lead to a tearing exponential increase of R_N^2 with N [as can be verified by following Eqs. (7) and (8)].

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