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Unzipping of DNA with correlated base sequence

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We consider force-induced unzipping transition for a heterogeneous DNA model with a correlated base sequence. Both finite-range and long-range correlated situations are considered. It is shown that finite-range correlations increase stability of DNA with respect to the external unzipping force. Due to long-range correlations the number of unzipped base pairs displays two widely different scenarios depending on the details of the base sequence: either there is no unzipping phase transition at all, or the transition is realized via a sequence of jumps with magnitude comparable to the size of the system. Both scenarios are different from the behavior of the average number of unzipped base pairs (non-self-averaging). The results can be relevant for explaining the biological purpose of correlated structures in DNA.

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I. INTRODUCTION

Structural transformations of DNA under changing of external conditions are of importance for molecular biology [1] and biophysics [2]. They take place in transcription of genetic information from DNA and in duplication of DNA during cell division [1]. The common scenario of these processes is unwinding of the double-stranded structure of DNA under influence of external forces. Recall that DNA consists of two strands with one wound around the other. These two strands interact via hydrogen bonds due to which the double-helix structure is formed. The individual strand is constructed by covalent bonds whose strength is thus much larger than the interstrand coupling. Each strand is a polymer based on nucleotides. A nucleotide is a deoxyribose sugar molecule bearing on one side purine or pyrimidine group (the base) and on the other a phosphate group. The purines can be of two types: adenine (A) and guanine (G), whereas pyrimidines are cytosine (C) and thymine (T) [an additional purine uracil (U) is found in RNA]. A, G, C, and T groups differentiate the nucleotides and constitute the genetic code carried by a DNA molecule. The bounds between neighboring nucleotides within one strand are formed via the corresponding phosphate groups. Hydrogen bonds between opposite strands are formed either by A-T bases or by G-C bases. Since the bases A, G, C, and T are hydrophobic, they are located at the core of the double helix. In contrast, the sugar molecules and the phosphate groups are hydrophilic and they are located in the outside part of the DNA molecule. Thus in a regular DNA molecule the letters of the genetic code are hidden from the molecular environment. This appears as a problem for the polymerase enzymes whose role is to read the genetic code. The polymerase may function if only they unzip the needed part of the DNA molecule, so that the bases are exposed to the environment. This is the main reason why DNA unzipping, in particular, under an external force is important for functioning of all living organisms. Force-induced unzipping has been actively investigated only recently [3-7] motivated by the new generation of micromanipulation experiments [7,8].

It is expected that features of the unzipping process depend on the base sequence of DNA, because AT and GC base pairs do have different formation energies. It is more difficult to break a single GC base pair, since it is made of three hydrogen bonds, while a single AT base pair is made of two hydrogen bonds only. Thus, the formation energy difference between AT and GC base pairs is of the order of one hydrogen bond energy, that is, 0.1–0.2 eV. This is comparable with the average formation energy itself. We note in addition that for a given DNA molecule the overall concentrations of AT and GC base pairs are approximately equal [1]. This is especially true for higher organisms, e.g., the concentration of GC base pairs for primates is between 49% and 51% [1].

The above energy difference may not be relevant for certain bulk properties of DNA. Therefore, the latter is frequently modeled assuming a homogeneous base sequence. However, in natural conditions the energy supplied for unzipping can be comparable to the average formation energy, and then the heterogeneous character of the base sequence becomes relevant. One of the first steps in this direction was made in Ref. [6], where it was shown that short-range heterogeneity does influence the unzipping process in the region where the energy supplied by an external unzipping force is comparable to the average formation energy of a DNA base pair.

Our main purpose is to make the next step towards real DNAs and to analyze force-induced unzipping for a DNA model, where the structural features of the base sequence are taken into account. One of the known features of DNA is that its base sequence displays substantial correlations which, in particular, can be of long-range characters [9-12]: two base
pairs separated from each other by thousands of pairs appear to be statistically correlated. Initial studies reported long-range correlations for noncoding regions of DNA (introns). For higher organisms, e.g., humans, these regions constitute more than 90% of DNA [1]. It was believed for some time that coding regions, which carry the majority of genetic information, can have only short-range correlations. However, more recent results indicate on the existence of weak long-range correlations in coding regions as well [11] (this point was controversial for a while, but the general consensus on its validity emerged gradually). Moreover, systematic changes were found in the structure of correlations depending on the evolutionary category of the DNA carrier [10]. In spite of ubiquity of long-range correlations, their biological reason remains largely unexplored. Some attempts in this direction were made in Ref. [13], where it was studied why long-range correlations are absent in certain biologically active proteins.

Our basic purpose in the present paper will be to determine how statistical correlations, in particular long-range correlations, influence on the unzipping process. Due to the biological relevance of unzipping, indications of such influences can provide useful information for explaining the presence of long-range correlations in DNA.

This paper is organized as follows. The basic model we work with is described in Sec. II. The situation with finite-range correlated base sequence is investigated in Sec. III A. The next three sections study various aspects of the long-range correlated situation. We conclude with a summary of our results. Several technical points are outlined in appendices.

II. THE MODEL

There are three basic mechanisms which determine the physics of the unzipping process: An external force tending to unbind the double-helix structure of a DNA molecule,

thermal noise generated by an equilibrium environment into which the molecule is embedded, and finally structural features of the molecule itself. Among various structural features which may be of relevance, the most important ones are connected with the base sequence of the molecule.

We shall work with a model which takes into account these three physical ingredients in the most minimal way. It was recently proposed in Ref. [6] for studying DNA unzipping.

(i) A DNA molecule is lying along the $x$ axis between the points $x=a$ and $x=L$.

(ii) Among all degrees of freedom of the molecule we consider only base pairs; they are located at points $x_i$, $a<x_i<L$, $i=1,...,M$. Indeed, for that range of external force where the molecule is close to be unbound completely, those degrees of freedom which are related to hydrogen bonds have much shorter characteristic times as compared to other degrees of freedom. The latter ones can therefore be considered as adiabatically frozen, and excluded from the effective description we are developing.

(iii) Any base pair can be in one of two states: bound or disconnected (broken). We choose the overall energy scale in such a way that the latter case contributes to the Hamiltonian a binding energy $\phi(x_i)$, whereas the former case brings nothing. As we stressed in the Introduction, different types of base pairs do have different binding energies: even when considering the ideal situation, where there are no “wrong base pairs” such as AC and GT, the “correct” base pairs AT and GC are different with respect to energy needed to unbind them. Thus $\phi(x_i)$ is a random quantity with an average $\langle \phi \rangle$:

$$\phi(x_i) = \langle \phi \rangle + \eta(x_i).$$

(iv) An external force is acting on the left end $x=a$ of the molecule pulling apart the two strands. Thus, if a bond $x_i$ is broken, all the base pairs $x_j$ with $j<i$ are broken as well. Each broken bond brings additionally to the Hamiltonian a term $-F$, where $F$ is proportional to the acting force.

(v) Summarizing all of these, one comes to the Hamiltonian

$$H(x) = -F x + \sum_{i=1}^{x} \phi(x_i) = (\langle \phi \rangle - F)x + \sum_{i=1}^{x} \eta(x_i),$$

where $x$ is the number of broken base pairs.

In the thermodynamical limit, where $L,M \gg 1$, one applies the continuum description with $x$ being a real number, $a<x<L$, and ends up with the following Hamiltonian:

$$H(x) = (x-a)f + \int_{a}^{x} ds \eta(s),$$

where $f=\langle \phi \rangle - F$ and $\beta=1/T$ is the inverse temperature ($k_B = 1$).

(vi) For characteristic time scales of unzipping experiments we can certainly neglect any changes of the base sequence for a single DNA molecule. Thus, once it is modeled via the random noise $\eta$, it is legitimate to assume that this noise is frozen, i.e., its single realization corresponds to a single molecule. It is assumed that the DNA molecule is embedded into a thermal bath with temperature $T$, and had sufficient time to reach equilibrium. Thus, the partition function and the free energy corresponding to the Hamiltonian (3) read

$$Z = \int_{a}^{L} \text{d}x \ e^{-\beta H(x)}, \quad F = -T \ln Z.$$

These quantities are still random together with $\eta$. Average results of many experiments with various realizations of $\eta$ can be described with help of the average free energy $\langle F \rangle$.

Our order parameter is the number of broken base pairs $X$. Along with its average it is defined for $t=0$ as

$$X = \partial_x F, \quad \langle X \rangle = \partial_x \langle F \rangle.$$

A. Finite-range and long-range correlated situations

It remains to specify the properties of the noise $\eta$. Within the adopted description we assume it is a Gaussian stationary process with an autocorrelation function

$$K(t-t') = \langle \eta(t) \eta(t') \rangle, \quad K(t) = K(-t).$$
Two major classes can now be distinguished depending on the behavior of $K(t)$ for large $t$. The finite-range correlated situation is defined by requiring that the integral

$$D = \int_0^\infty ds \ K(s)$$  \hspace{1cm} (7)

determining the total intensity of the noise is finite. There are three particular cases of the finite-range correlated situation. The white-noise case

$$K(t) = D \delta(t)$$ \hspace{1cm} (8)
describes completely uncorrelated noise. The physical situation given by Eqs. (3) and (8) is well known, and was used to describe interfaces, random walks in a disordered media, and population dynamics [14]. It was recently applied for the unzipping transition in DNA [6]. Similar models were considered in Refs. [5,15].

The second case corresponds to the noise having some finite—though possibly large—correlation length $\tau$. The simplest and most widely used model for this case is provided by Ornstein-Uhlenbeck (OU) noise

$$K(t) = \frac{D}{\tau} e^{-|t|/\tau},$$ \hspace{1cm} (9)

where $D$ is the total intensity of the noise and $\tau$ is the correlation time; $\tau \to 0$ corresponds to the white noise. The third case is when $K(t)$ has a power-law dependence for large $t$, but still decays sufficiently quickly so that the integral in Eq. (7) is finite: $K(t) \propto |t|^{-\alpha}$ with $\delta > 1$.

The second major class is the long-range correlated situation, where the integral in Eq. (7) is infinite, that is, when $K(t)$ for sufficiently large $t$ behaves according to a power law [10]:

$$K(t) = \langle \eta(t) \eta(0) \rangle = \sigma |t|^{-\alpha},$$ \hspace{1cm} (10)

where

$$0 < \alpha < 1$$ \hspace{1cm} (11)

is the exponent characterizing the long-range correlation, and where $\sigma$ is the (local) intensity. Note that $K(t)$ has to be regular and finite for small $t$ [12], as one would expect from physical reasons.

The OU noise (9), as the typical representative of the finite-range correlated situation, and the long-range correlated noise (10) are relevant for modeling correlations in base sequence of DNA [9–12,16]. Note, however, that the real noise distributions in DNA can be much more complicated [10,11]. In particular, this concerns the Gaussian property we assume (see in this context Sec. V A, where we study a model of a non-Gaussian noise to show that its predictions in the thermodynamic limit do not differ from those given by the corresponding Gaussian noise). For the long-range correlated situation there can exist several characteristic exponents for different ranges of $t$. Nevertheless, Eqs. (9) and (10) are certainly the minimal models of noise which are sufficiently simple and which allow to study both finite- and long-range correlations.

### B. Reduction to Langevin equation

The basic method of solving the present model will be to reduce it to the physics of a Brownian particle whose dynamics is described by a stochastic differential equation. In Eq. (4) one fixes $L$, and views $a$ as a parameter varying from the highest possible value $L$, where $Z=0$, to the lowest possible value which we define to be $a=0$. The quantity $t=\tau-a$ will thus monotonically increase and can be interpreted as a time variable. Differentiating $Z$ in Eq. (4) over $a$ and changing the variable as $t=\tau-a$, one gets

$$\frac{dZ}{dt} = 1 - \beta fZ - \beta \eta(t)Z, \quad -L < t < 0,$$ \hspace{1cm} (12)

where we used $\eta(t) = \eta(-t)$, as follows from the Gaussian stationary property of the noise. This is a Langevin equation with a multiplicative noise. From Eq. (12) one can obtain a stochastic equation for $F=-T \ln Z:$

$$\frac{dF}{dt} + V'(F) = \eta(t), \quad V(F) = T^2 e^{2F} - f F.$$ \hspace{1cm} (13)

This is the basic stochastic equation we will work with.

### III. FINITE-RANGE CORRELATED NOISE

#### A. Ornstein-Uhlenbeck noise

Our main purpose here is to study the process of unzipping in the presence of the finite-range correlated noise given by Eq. (9). We wish to understand how the magnitude of $\tau$ influences unzipping.

Note that the OU noise (9) can itself be modeled via a white noise:

$$\tau \dot{\eta} = - \eta + \sqrt{D} \xi(t),$$ \hspace{1cm} (14)

where $\xi(t)$ is a Gaussian noise with $\delta$-correlated spectrum:

$$\langle \xi(t) \xi(t') \rangle = 2 \delta(t - t').$$ \hspace{1cm} (15)

Indeed, Eq. (9) is recovered directly from Eqs. (14) and (15), since their exact solution is

$$\langle \eta(t) \eta(t') \rangle = e^{-\gamma(t-t')/\tau} \left( \langle \eta^2(0) \rangle - \frac{D}{\tau} \right) + \frac{D}{\tau} e^{-|t-t'|/\tau}.$$ \hspace{1cm} (16)

We get back from here to Eq. (9) under an additional consistency condition $\langle \eta^2(0) \rangle = D/\tau$. Moreover, $\eta(t)$ is a Gaussian random process, because $\xi(t)$ is Gaussian and Eq. (14) is linear [17].

To handle Eq. (13) one differentiates it over $t$ and uses Eqs. (14) and (15). Changing the variable as $s = t/\sqrt{\tau}$ one gets [18]

$$\frac{d^2 F}{ds^2} + \gamma(F) \frac{dF}{ds} = -V'(F) + \frac{\sqrt{D}}{\tau^{1/2}} \xi(s),$$ \hspace{1cm} (17)

where

$$\gamma(F) = \tau^{-1/2} + V''(F) \tau^{1/2}.$$ \hspace{1cm} (18)

Equation (17) has the same form as a Langevin equation for a particle with unit mass in the potential $V(F)$ and subjected...
to a white noise and a $F$-dependent friction with a coefficient $\gamma(F)$. Note that the potential $V(F)$ is confining only for $f > 0$: $V(F) \to \infty$ for $F \to \pm \infty$.

We can rewrite Eq. (17) introducing an additional variable $\bar{F}(s) = dF(s)/ds$, which in the above language of the Brownian motion corresponds to the velocity:

$$\frac{dF}{ds} = \bar{F},$$

$$\frac{d\bar{F}}{ds} = -\gamma(F)\bar{F} - V'(F) + \sqrt{\frac{4\pi}{\sqrt{D}}} \xi(s).$$

As $\xi(t)$ is a Gaussian white noise, one uses the standard tools, see, e.g., Ref. [17], and writes down from Eqs. (19) and (20) a Fokker-Planck-Klein-Kramers equation for the common probability distribution:

$$P(F, \bar{F}, s) = \langle \delta(F - F(s)) \, \delta(\bar{F} - \bar{F}(s)) \rangle,$$

where $F(s)$ and $\bar{F}(s)$ are particular noise-dependent solutions of Eqs. (19) and (20), and where the average is taken over the white noise $\xi(t)$ given by Eq. (15):

$$\frac{\partial P(F, \bar{F}, s)}{\partial s} = -\bar{F} \frac{\partial P(F, \bar{F}, s)}{\partial \bar{F}} + \gamma(F) \frac{\partial \left[ \bar{F}P(F, \bar{F}, s) \right]}{\partial \bar{F}} + V'(F) \frac{\partial P(F, \bar{F}, s)}{\partial \bar{F}} + \frac{D}{\tau^{1/2}} \frac{\partial P(F, \bar{F}, s)}{\partial \bar{F}^2}.$$ 

Our interest is in the large-$s$ limit of this equation (thermodynamic limit), and we want to have the reduced probability distribution $P(F, s)$ of $F$ only:

$$P(F, s) = \langle \delta(F - F(s)) \rangle = \int d\bar{F} P(F, \bar{F}, s).$$

To this end let us introduce

$$Q_n(F, s) = \int d\bar{F} \bar{F}^n P(F, \bar{F}, s), \quad n = 0, 1, 2, 3 \ldots,$$

where $Q_0(s) = P(F, s)$. From Eq. (22) one gets an infinite set of coupled equations for $Q_n(F, s)$:

$$\frac{\partial Q_0(F, s)}{\partial s} = -\frac{\partial Q_1(F, s)}{\partial F}.$$ 

$$\frac{\partial Q_1(F, s)}{\partial s} = -\frac{\partial Q_2(F, s)}{\partial F} - \gamma(F)Q_1(F, s) - V'(F)Q_0(F, s),$$

$$\frac{\partial Q_2(F, s)}{\partial s} = -\frac{\partial Q_3(F, s)}{\partial F} - 2V'(F)Q_1(F, s) - 2\gamma(F)Q_2(F, s) + \frac{2D}{\tau^{1/2}} Q_0(F, s),$$

$$\frac{\partial Q_3(F, s)}{\partial s} = -\frac{\partial Q_4(F, s)}{\partial F} - 2V'(F)Q_2(F, s) - 2\gamma(F)Q_3(F, s) - \lambda(F)Q_4(F, s).$$

When deriving Eqs. (25) and (26) we used integration by parts, and the following standard boundary conditions:

$$P(F, \bar{F}, s) \to 0 \quad \text{if} \quad F \to \pm \infty \quad \text{or} \quad \bar{F} \to \pm \infty.$$ 

These conditions are physically meaningful if the potential $V(F)$ is confining, and thus the motion of the corresponding Brownian particle takes place in a finite domain. According to the above discussion on the confining character of the potential $V(F) = T^2 e^{BF} - F$, the boundary conditions (29) are reliable only for $f > 0$.

Recall that the “time variable” $t$ moves between $-|L|$ and 0. For large lengths, i.e., for $L \gg 1$ (thermodynamic limit) and as the consequence $t \ll s \to 0$, any solution of Eq. (22) relaxes towards the unique stationary distribution $P_{st}(F, \bar{F})$. A rather general proof of this fact is presented in Ref. [17].

We shall now use Eqs. (25) and (26) to get explicitly the stationary distribution function $P_{st}(F)$ of $F$. Putting to zero the left-hand side (LHS) of Eq. (25) one gets that $Q_{1,st}(F)$ does not depend on $F$. Taking into account the boundary condition (29) one concludes that it is equal to zero:

$$Q_{1,st}(F) = 0.$$ 

Putting to zero the LHS of Eq. (26) and using Eq. (30) we get

$$\frac{\partial Q_{2,st}(F)}{\partial F} = -V'(F)Q_{0,st}(F).$$

It remains to determine $Q_{2,st}(F)$ putting to zero the LHS of Eq. (27). One can conjecture that the stationary state $P_{st}(F, \bar{F})$ is symmetric with respect to $\bar{F} \to -\bar{F}$, and then $Q_{1,st}(F) = 0$ in the same way as for $Q_{1,st}(F)$ in Eq. (30). Alternatively, one can assume that $\gamma(F)$ and $D$ are sufficiently large so that the term $\partial Q_{2,st}(F)/\partial F$ can be simply dropped in the right-hand side (RHS) of Eq. (27). If $V'(F)$ is of order 1, then a large $\gamma(F)$ is realized both for large and small $\tau$ [18].

Thus we conclude from Eq. (27):

$$\gamma(F)Q_{2,st}(F) = \frac{D}{\tau^{1/2}} Q_{0,st}(F).$$

In view of Eqs. (31) and (32) one has a single differential equation

$$\frac{D}{\sqrt{\tau}} \frac{\partial}{\partial F} \left( \frac{Q_{nst}(F)}{\gamma(F)} \right) = -V'(F)Q_{0,st}(F),$$

and gets for $Q_{0,st}(F) = P_{st}(F)$:

$$P_{st}(F) \propto \gamma(F) \exp \left[ -\frac{\tau}{2D} [V'(F)]^2 - \frac{1}{D} V(F) \right].$$
\[
P_\sigma(F) = \mathcal{N}(1 + \tau e^{\beta F}) \exp \left[ \frac{fE}{D} - \frac{T(T - \tau f)e^{\beta F} - \tau T^2}{2D} \right].
\]

(34)

where \(\mathcal{N}\) is the normalization factor. The white-noise, \(\tau \rightarrow 0\), limit of \(P_\sigma(F)\) was obtained in Refs. [6,14].

According to Eqs. (5) and (34) the average free energy reads

\[
\langle F \rangle = T f + \int_{0}^{\infty} d\tau \frac{1}{u} \Gamma(u, 0) \left[ \left( \mu - \frac{T^2}{D} \right) u - \frac{\tau T^2}{2D} u^2 \right] - \frac{T^2}{2D} \tau^2 u^2,
\]

\[
\int_{0}^{\infty} d\tau \frac{1}{u} \Gamma(u, 0) \left[ \left( \mu - \frac{T^2}{D} \right) u - \frac{\tau T^2}{2D} u^2 \right] - \frac{T^2}{2D} \tau^2 u^2,
\]

(35)

where

\[
\mu = \frac{Tf}{D}.
\]

(36)

Note that both integrals in Eq. (35) can be expressed through the Gamma function \(\Gamma(x)\) and the confluent hypergeometric (Kummer) function \(F_1(a, b; x)\), since

\[
\int_{0}^{\infty} d\tau \frac{1}{u} \Gamma(u, 0) \left[ \left( \mu - \frac{T^2}{D} \right) u - \frac{\tau T^2}{2D} u^2 \right] = \frac{T^2}{2D} \mu^2
\]

\[
\int_{0}^{\infty} d\tau \frac{1}{u} \Gamma(u, 0) \left[ \left( \mu - \frac{T^2}{D} \right) u - \frac{\tau T^2}{2D} u^2 \right] = \frac{T^2}{2D} \mu^2
\]

(37)

Similar formulas can be written for \(\int_{0}^{\infty} d\tau \frac{1}{u} \Gamma(u, 0) \left[ \left( \mu - \frac{T^2}{D} \right) u - \frac{\tau T^2}{2D} u^2 \right] - \frac{T^2}{2D} \tau^2 u^2\) for \(n=1,2\). These representations facilitate numerical calculations.

The average number of broken base pairs \(\langle X \rangle\) can be calculated from Eqs. (5) and (34). Note that for the white-noise situation \(\tau \rightarrow 0\) a simple formula is obtained:

\[
\langle X \rangle = \frac{T^2}{D} \frac{d\psi(\mu)}{d\mu},
\]

(38)

where \(\psi(\mu) = \Gamma(\mu) / \Gamma'(\mu)\). For \(\mu \rightarrow 0\), \(\langle X \rangle\) does not depend on temperature and on \(\tau\) and becomes very large:

\[
\langle X \rangle = DF^{-2},
\]

(39)

for \(f \rightarrow 0\). When the external force reaches its critical value, the average number of broken base pairs diverges in the thermodynamic limit.

To study the influence of \(\tau\) on this unzipping phase transition, one should keep in mind the realistic situation, where DNA molecules belonging to different evolutionary classes have different correlation properties of their base sequences [10]. At the same time the concentration (fraction) of AT and GC base pairs is known to be (approximately) equal for sufficiently long DNA molecules in natural conditions [1,19]. Therefore, in comparing two situations having different correlation characteristics, it is legitimate to keep fixed the intensity of the noise defined by Eq. (7) —this corresponds to fixed concentration of various base pairs — and to study how the average number of broken base pairs \(\langle X \rangle\) depends on \(\tau\) for some fixed value of \(f\). This dependence is displayed in Fig. 1 following Eqs. (5) and (35). It is seen that the behavior of \(\langle X \rangle\) for very small \(f\) depends on \(\tau\) rather weakly. Indeed, as follows from Eq. (34), for \(f \rightarrow 0\) the relevant domain of \(F\) contributing into \(\langle F \rangle\) is \(F \sim D/f\). As it does not depend on \(\tau\), we get back Eq. (39). However, a nontrivial dependence on \(\tau\) does exist for moderately small values of \(f\), where as seen in Fig. 1, \(\langle X \rangle\) is a decreasing function of \(\tau\) for a fixed \(f\): longer correlations present in the base sequence increase the stability of the DNA molecule, since larger external forces \(\tau\) needed to achieve the same average amount of broken base pairs. This is our main qualitative conclusion on how a finite correlation length influences the unzipping process.

B. Arbitrary finite-range correlated noise at low temperatures

In the preceding section we reduced the nonlinear equation (13) with the finite-range correlated noise (9) to a Fokker-Planck equation, and solved the latter exactly in the thermodynamic limit. The essential feature that made this analytical solution possible is that the OU noise has a single and well-defined characteristic time and due to this allows representations (14) and (15).

In general it is impossible to solve Eq. (13) for an arbitrary Gaussian noise, and, in particular, for the situation given by Eq. (10): there is no exact Fokker-Planck equation for this case. There is, however, a particular case which allows analytical treatment. For very low temperatures, \(T \rightarrow 0\), one can approximately substitute \(V(F)\) in Eq. (13) by \(-f F\) for \(F < 0\) and by an infinite potential wall standing at \(F=0\). Thus, all values \(F > 0\) become prohibited. For this particular form of potential one can get a Fokker-Planck equation for

\[
P(F,t) = \langle \delta(F - F[t, \eta,t]) \rangle
\]

(40)

with an arbitrary Gaussian noise in the RHS of Eq. (13). The derivation goes as follows. Write Eq. (13) as

\[
\frac{dF}{dt} = f + \eta(t),
\]

(41)

where the stochastic variable \(F\) is restricted to be negative due to the above infinite wall. Differentiating \(P(F,t)\) in Eq. (40) over \(t\) one gets
\[
\frac{\partial P(F,t)}{\partial t} = -f \frac{\partial P(F,t)}{\partial F} - \frac{\partial}{\partial F} (\delta(t) \delta(F - F[\eta,t])).
\] (42)

It remains to handle the last term in this equation. One uses Novikov's theorem [20]

\[
(\delta(t) \delta(F - F[\eta,t])) = -\frac{\partial}{\partial F} \int_{-L}^{t} ds K(t-s)
\]

\[
\times \left( \delta(F - F[\eta,t]) \frac{\delta F[t]}{\delta \eta(s)} \right),
\] (43)

where \(\partial F[\eta]/\partial \eta(s)\) is the variational derivative and \(\delta F[t]/\delta \eta(s)\) is obtained from Eq. (41):

\[
\frac{d}{dt} \frac{\delta F[t]}{\delta \eta(s)} = \delta(t-s), \quad \frac{\delta F[t]}{\delta \eta(s)} = \theta(t-s),
\] (44)

where \(\theta(t-s)\) is the step function. Combining Eqs. (42)–(44) we get finally

\[
\frac{\partial P(F,t)}{\partial t} = -f \frac{\partial P(F,t)}{\partial F} + D \frac{\partial^2 P(F,t)}{\partial F^2},
\] (45)

\[
D = \int_{0}^{+L} ds K(s).
\] (46)

Equation (45) should additionally be supplemented by a boundary condition which reflects the presence of the infinite wall at \(F=0\). Equation (45) can be written as the continuity equation

\[
\frac{\partial P(F,t)}{\partial t} + \frac{\partial J(F,t)}{\partial F} = 0, \quad J(F,t) = f - \mu_{\tau} \frac{\partial P(F,t)}{\partial F},
\] (47)

where \(J(F,t)\) is the probability current. The infinite wall at \(F=0\) is now implemented by requiring

\[
\int_{-\infty}^{0} dF P(F,t) = 1,
\] (48)

\[
J(0,t) = 0,
\] (49)

for all \(t\). Conditions (48) and (49) are imposed on any solution of Eq. (45).

In the thermodynamic limit \(L \to \infty\) and \(t=0\) one gets from the stationarity condition \(\partial P(F,t)/\partial t = 0\)

\[
P(F) = \frac{f}{D} \exp \left[ \frac{f F}{D} \right] \quad \text{for} \quad F < 0
\] (50)

\[
= 0 \quad \text{for} \quad F \gg 0,
\] (51)

where the total intensity, as given by Eq. (7), is finite for the considered short-range correlated situation. Note that in the thermodynamical limit conditions (48) and (49) are satisfied automatically as seen from Eqs. (50) and (51). It is now seen from Eq. (5) that

\[
\langle X \rangle = \frac{D f^2}{\tau},
\] (52)

which has the same \(f\) dependence as the white-noise case for small \(f\); see Eq. (39). We conclude that, not unexpectedly, for low temperatures the behavior of \(X\) is determined only by the total intensity of the noise. All other details of \(K(t)\) do not matter. It remains to stress that the present analysis certainly does not apply to the long-range correlated situation (10), since the total intensity \(D\) diverges in the thermodynamical limit.

In closing this section, let us note that Eq. (52) can be applied to finite-range correlated noise that for \(t \ll L\) has the same autocorrelation function as Eq. (10). As an example take

\[
K_{\alpha}(t) = \sigma |t|^{-\alpha} \quad \text{for} \quad |t| \leq l
\] (53)

\[
= 0 \quad \text{for} \quad |t| > l,
\] (54)

where \(l\) is some parameter that is finite in the thermodynamical limit \(L \to \infty\). Therefore, the noise given by Eq. (53) is obviously finite-range correlated. Equation (52) now reads

\[
\langle X \rangle = \frac{\sigma}{1 - \alpha} \int |t|^{-\alpha} f^2 dt.
\] (55)

If one chooses to take \(l \sim (X)\) then \(\langle X \rangle \sim f^{2/\alpha}\) as predicted in Ref. [6]. However, there is no any a priori reason for this choice, and at any rate this result refers to the finite-range correlated noise \(K_{\alpha}\). The real long-range correlated situation, where \(l \sim L\), is still not described by it.

IV. LONG-RANGE CORRELATED SITUATION: THE FROZEN NOISE LIMIT

The present and the following section are devoted to the long-range correlated situation, where according to Eq. (10) the autocorrelation function \(K(t)\) of the noise has a power-law behavior with the single characteristic exponent \(1 > \alpha > 0\).

To start with, let us consider the case with \(\alpha \to 0\). The noise is now completely frozen: \(\eta(s)\) in Eq. (3) does not depend on \(s\). This situation is less physical as compared to that with \(\alpha > 0\). However, it is exactly solvable, and one can hope it catches at least some features of the realistic situation where \(\alpha\) is larger than zero, but certainly smaller than 1. This intuitive expectation will be confirmed later on.

The problem with \(\alpha = 0\) is easily solved from Eq. (3). Moreover, the exact solution can be obtained for an arbitrary value of \(L\):

\[
\frac{X}{L} = g[\beta L (f + \eta)],
\] (56)

\[
g[x] = \frac{1}{x} - \frac{1}{e^x - 1}.
\] (57)

It is seen that in the thermodynamical limit \(L \to \infty\), \(g[\beta L (f + \eta)]\) behaves as roughly the step function \(g[\beta L (f + \eta)] = \theta(-\eta - f)\); for any single realization of the noise there is a sharp
phase transition with a jump at the realization-dependent point \( f = -\eta \). Exactly at this point \( f = -\eta \) one has \( g(0) = 1/2 \) and \( \langle X \rangle = L/2 \).

Let us now study the behavior of \( \langle X \rangle \). Since the noise is completely frozen, the calculation of \( \langle X \rangle \) reduces to the averaging over a Gaussian variable with dispersion \( \sigma \). We have

\[
\frac{\langle X \rangle}{L} = \frac{\langle g[\beta L(f + \eta)] \rangle}{L} = \int \frac{d\eta}{\sqrt{2\pi \sigma}} \exp \left[ -\frac{\eta^2}{2\sigma^2} \right] g[\beta L(f + \eta)]
\]

where we changed the integration variable as \( \xi = \beta(\eta + f) \). In the thermodynamical limit \( L \to \infty \), we shall obtain for \( \langle X \rangle / L \) the main term of order \( O(L^0) \), and the first correction to it which will appear to be of order \( O(1/L) \). To this end, let us divide the integration in the RHS of Eq. (58) into three pieces:

\[
\int = \int_{-\infty}^{-2L} + \int_{-2L}^{2L} + \int_{2L}^{\infty}.
\]

For each piece we shall use the following approximate expressions obtained from Eq. (57):

\[
g[L\xi] = \frac{1}{L\xi} \quad \text{for} \quad L\xi \geq 2,
\]

\[
g[L\xi] = \frac{1}{2} \quad \text{for} \quad -2 \leq L\xi \leq 2,
\]

\[
g[L\xi] = 1 + \frac{1}{L\xi} \quad \text{for} \quad L\xi \leq -2.
\]

To obtain Eqs. (60) and (62) we neglected terms of order \( O(e^{-\beta L^0}) \). For Eq. (61) which corresponds to the second integration piece in Eq. (59), we have taken the value of \( g[L\xi] \) at \( \xi = 0 \). The boundary points of \( L\xi \) were chosen such as to ensure a continuous matching. However, neither the precise value of \( g[L\xi] \) within the second piece of integration in Eq. (59) nor the precise values of the points separating this piece from the remaining ones are important, since as we show below the contribution coming from this second piece as well as the contributions from the boundary points of the two other integration pieces produce factors of order \( O(1/L^2) \) at best.

The same concerns factors of order \( O(e^{-\beta L^0}) \) that were neglected in Eqs. (60)–(62). Combining Eqs. (60)–(62) with Eq. (58) one gets

\[
\frac{\langle X \rangle}{L} = \int_{-\infty}^{-2L} \frac{d\xi}{\sqrt{2\pi \sigma \beta^2}} \exp \left[ -\frac{(\xi - \beta f)^2}{2\sigma^2} \right] \left( 1 + \frac{1}{L\xi} \right)
\]

\[
+ \frac{1}{2} \int_{-2L}^{2L} \frac{d\xi}{\sqrt{2\pi \sigma \beta^2}} \exp \left[ -\frac{(\xi - \beta f)^2}{2\sigma^2} \right]
\]

\[
+ \int_{2L}^{\infty} \frac{d\xi}{\sqrt{2\pi \sigma \beta^2}} \exp \left[ -\frac{(\xi - \beta f)^2}{2\sigma^2} \right] \frac{1}{L\xi}.
\]

One notes that both Eqs. (65) and (66) are of order \( O(1/L^2) \). This can be verified by directly expanding integrals in Eqs. (65) and (66) for small \( 2/L \). Skipping these terms, one gets

\[
\frac{\langle X \rangle}{L} = \int_{-\infty}^{\infty} \frac{d\xi}{\sqrt{2\pi \sigma \beta^2}} \exp \left[ -\frac{\xi^2}{2\sigma^2} \right] \frac{1}{L\xi} \int_{0}^{\infty} \frac{d\xi}{\xi} \exp \left[ -\frac{\xi^2}{2\sigma^2} \right]
\]

\[
\times \left( \exp \left[ -\frac{(\xi - \beta f)^2}{2\sigma^2} \right] - \exp \left[ -\frac{(\xi + \beta f)^2}{2\sigma^2} \right] \right)
\]

\[
\times \int_{0}^{\beta f} \frac{d\xi}{\sqrt{2\pi \sigma \beta^2}} \exp \left[ -\frac{\xi^2}{2\sigma^2} \right] \frac{1}{\sigma \beta^2 L} \exp \left[ -\frac{\xi^2}{2\sigma^2} \right]
\]

\[
× \int_{0}^{\beta f} \frac{d\xi}{\sqrt{2\pi \sigma \beta^2}} \exp \left[ -\frac{\xi^2}{2\sigma^2} \right].
\]

When obtaining the last term in the RHS of Eq. (67), we used a tabulared identity for the error function.

For \( f \) not very large as compared to \( \sqrt{\sigma} \), the first term in the LHS of Eq. (67) is dominating: \( \langle X \rangle / L \) is of order \( 1/2 \). In particular, it is exactly equal to \( 1/2 \) for \( f = 0 \) as can be deduced directly from Eq. (58). The dependence of \( \langle X \rangle \) on \( f \) becomes thus very weak for \( f \to 0 \). The second, subdominant term becomes non-negligible for \( f \gg \sqrt{\sigma} \), where using asymptotic identities (see Appendix B)

\[
\int_{-\infty}^{\infty} \frac{d\xi}{\sqrt{2\pi}} \exp \left[ -\frac{\xi^2}{2} \right] = \frac{1}{\sqrt{2\pi}} \left( 1 - \frac{1}{a^2} + \cdots \right), \quad a \gg 1,
\]

\[
\int_{0}^{a} \frac{d\xi}{\sqrt{2\pi}} \exp \left[ -\frac{\xi^2}{2} \right] = \frac{a}{\sqrt{2\pi}} \left( 1 + \frac{1}{a^2} + \cdots \right), \quad a \gg 1,
\]

one gets from Eq. (67) noting \( a = f \sqrt{\sigma} \)

\[
\langle X \rangle = \frac{1}{\beta f} \left( 1 + \frac{\sigma}{f} + \frac{\sqrt{\sigma}}{\sqrt{2\pi} L \beta \exp \left[ -\frac{f^2}{2\sigma} \right]} \right).
\]

Note that for \( f \gg \sqrt{\sigma} \), \( \langle X \rangle \) has — within the leading order — the same \( 1/f \) dependence as it will be in the completely homogeneous situation without noise (\( \sigma = 0 \)). In the considered regime, the noise only renormalizes this behavior modifying the subdominant terms.
The quantity \( \langle X^2 \rangle \) can be calculated in the same way as in Eqs. (63) and (67). We shall bring the result only for \( f \) not very large as compared to \( \sqrt{\sigma} \), that is, when \( \langle X^2 \rangle \sim L^2 \):

\[
\frac{\langle X^2 \rangle}{L^2} = \int_{\beta \ell}^{\infty} d\xi \frac{d\xi}{\sqrt{2\pi\alpha\beta}} \exp \left[ -\frac{\xi^2}{2\sigma^2} \right] + \frac{2}{L} \ln \left( \frac{L}{2} \right) \exp \left[ -\frac{1}{2\ell^2} \right] \sqrt{2\pi\beta^2\sigma^2} + O\left( \frac{1}{L} \right).
\]

Substituting this into Eq. (71), we see that \( \frac{\langle X^2 \rangle - \langle X \rangle^2}{\langle X \rangle^2} \) remains finite in the thermodynamical limit:

\[
\frac{\langle X^2 \rangle - \langle X \rangle^2}{\langle X \rangle^2} = \frac{\int_{\beta \ell}^{\infty} d\xi \frac{d\xi}{\sqrt{2\pi\alpha\beta}} \exp \left[ -\frac{\xi^2}{2\sigma^2} \right]}{\left( \int_{\beta \ell}^{\infty} d\xi \frac{d\xi}{\sqrt{2\pi\alpha\beta}} \exp \left[ -\frac{\xi^2}{2\sigma^2} \right] \right)^2} - 1.
\]

In particular, for \( f \to 0 \)

\[
\frac{\langle X^2 \rangle - \langle X \rangle^2}{\langle X \rangle^2} \to 1,
\]

indicating essential non-self-averaging.

In closing this section, let us repeat that the character of the thermodynamical for the considered case \( \alpha = 0 \) is different from that of the finite-range correlation situation, where — for \( L \to \infty \) — the behavior of \( X \) becomes independent of \( L \) at least in the physical range of other parameters (e.g., \( 1 > f > 0, T > 0, \) etc.). For the \( \alpha = 0 \) case, as seen from Eqs. (67) and (70), there is an explicit dependence on \( L \) in the whole range of physical range of the involved parameters. According to Eq. (70), if \( L \) is kept large but finite, then this dependence is very weak for external forces far form their critical value \( f = 0 \), that is, for \( f \gg \sqrt{\sigma} \). There are no reasons for taking this explicit dependence on \( L \) as something unphysical. In contrast, the actual size of physically relevant examples of DNA is never more than \( L \sim 10^3 \sim 10^6 \); see Ref. [1]. This is certainly much smaller than the number \( 10^{23} \) which in the standard statistical physics is taken as the typical size. Therefore, it is rather natural to study the physics of unzipping for a large but fixed \( L \).

For the considered frozen situation, we could solve the problem analytically for a given realization. However, for \( \alpha > 0 \) this is not possible, and one has to rely on numerical methods. This is what we intend to do in the following section.

V. NUMERICAL RESULTS

As we have seen in the preceding section, there are reasons to expect that for the long-range correlated situation, especially for sufficiently small index \( \alpha \), the typical — that is, frequently met among many independent realizations of the noise — behavior of \( X(f) \) in the thermodynamic limit is not described adequately by the average quantity \( \langle X \rangle \) (non-self-averaging). We note in this context that the correlator
((X^2)−<X>^2)/<X>^2 studied in Sec. V can indicate on non-self-averaging, but by itself does not provide any direct information on typical realizations. It is perhaps needless to stress that once we expect the effect of non-self-averaging, the attention should be shifted towards typical realizations, since they do have a direct physical meaning for single-molecule experiments.

In the present section we study numerically the behavior of the number of broken base pairs X as a function of f both for the long-range correlated situation and for the uncorrelated noise. For the discrete version of the model the partition function reads

\[ Z = \sum_{k=1}^{L} \exp \left[ -\beta \left( f k + \sum_{i=1}^{k} \eta_i \right) \right] , \tag{75} \]

where for the long-range correlated situation \( \eta_i \) are Gaussian random variables with the autocorrelation function given by Eq. (10). Note that for the purposes of numerical computations the behavior of \( K(i) \) in Eq. (10) was regularized at short distances so as to avoid superfluous short-range singularities; see Appendix A for details. The generation of \( \eta_i \), \( i = 1, \ldots, L \) is described in Appendix A following optimized recipes proposed in Ref. [23]. For numerical computations we have chosen \( T=1 \) and \( L=10^4 \) or \( L=5 \times 10^4 \).

As \( L \) is now explicitly finite, one should be careful with the selection of the thermodynamical domain, since due to the very statement of the problem the limit \( L \rightarrow \infty \) is taken before \( f \rightarrow 0 \). As a plausible estimate of this domain, one can use a condition \( f \sqrt{L} \gg 1 \). We confirmed it in several ways, reproducing predictions which were made in the thermodynamical limit \( L \rightarrow \infty \).

### A. Uncorrelated noise

Let us start with the uncorrelated-noise case, where \( \eta_i \)'s are independent Gaussian variables with zero average, \( \langle \eta_i \rangle = 0 \), and variance \( \eta_i^2 = D=1 \) (white noise), and where \( X \) is given by Eqs. (5) and (75). For comparison we also studied a case, where \( \eta_i \) are independent random variables assuming values \( \eta_i = \pm 1 \) with equal probability (dichotomic noise).

The results are illustrated by Figs. 3 and 4, where we display \( \langle X \rangle \) and \( X \) for several typical realizations. It is seen that \( \langle X \rangle \) and \( X \) do not coincide exactly, as it is in general expected due to the finite magnitude of \( L \) if not by any other reason. However, in the considered thermodynamical domain of \( f \) the behavior of various typical realizations qualitatively resembles each other, and, therefore, resembles that of \( \langle X(f) \rangle \). In particular, for all typical realizations \( X(f) \) grows for \( f \rightarrow 0 \). In that sense \( f = 0 \) is a special point for both typical \( X \) and \( \langle X \rangle \). It should be mentioned that for \( f \leq 0.05 \) we have seen realizations containing relatively sudden jumps at realization-dependent values of \( f \). This differs from the behavior of \( \langle X \rangle \) and is in agreement with the results of Ref. [6].

However, such small values of \( f \) are not in the thermodynamical domain. Acknowledging reservations connected with the numerical character of our study, we, nevertheless, conclude that the uncorrelated-noise situation is self-averaging at least for not very small, \( f \sqrt{L} \gg 1 \), values of \( f \).

### B. Long-range correlated noise

#### 1. Typical realizations

The situation for the long-range correlated noise for \( \alpha =0.5, T=s=1 \) is illustrated by Figs. 6–9. The first point to

![Figure 3](https://via.placeholder.com/150)

**FIG. 3.** (Color online) Solid curves: \( X(f) \) for several realizations of the white uncorrelated noise. Dotted curve: \( \langle X(f) \rangle \) obtained by averaging over \( 10^3 \) realizations. \( T=D=1, L=5 \times 10^4 \).

It is seen from Fig. 4 that the white and dichotomic noise produce very similar results. This is to be expected for the considered large values of \( L \) (law of large numbers). Figure 5 shows that the power law \( \langle X \rangle \propto f^{-2} \) for the white-noise case is recovered by direct averaging over various realizations. Indeed, it is seen from this figure that one recovers

\[ \langle X \rangle \propto f^{-1.84} \tag{76} \]

after averaging over \( 10^3 \) realizations in the domain \( 0.05 < f < 0.25 \). This result is stable upon increasing the number of realizations, e.g., from \( 10^3 \) to \( 2 \times 10^3 \).
note is that now there are typical realizations with radically different properties. The first type of realizations is presented by Figs. 6 and 7: \(X(f)\) increases by several sudden jumps followed by flat regions. It is seen that \(X_u=0\) is either equal to its maximal possible value \(L\) or is close to it. Points where \(X_s\) has jumps vary from one realization to another. However, the overall number of jumps when varying \(f\) between zero and one is typically two or three.

In contrast to this, Figs. 8 and 9 present a strictly different situation: It is characterized by very smooth behavior of \(X(f)\) for \(f \neq 0\). In particular, \(X|_{f=0}\) is much smaller than \(L\) (typically by few orders of magnitude). \(X(f)\) is still a monotonic function of \(f\), but the point \(f=0\) — where the energy supplied by the external unzipping force is equal to the average binding energy of a base pair — is by no means special.

To estimate the frequency by which each scenario is met among all possible realizations, we have taken the following criteria for deciding whether a given realization belongs to one of the above classes: for \(L=5 \times 10^4\) we prescribe the given realization to the first class if \(X_s(f=0) > 4.8 \times 10^4\), while it is prescribed to the second class if \(X(f=0) < 10^2\). These criteria appeared to be sufficiently adequate, as they are consistent with the fact of presence (for the first class) of absence (for the second class) of sudden jumps for \(X(f)\).

In this way the frequencies of each class were estimated in a sample of \(10^3\) realizations. It appeared for \(L=5 \times 10^4\) and \(T=\sigma=2\alpha=1\) that the first scenario is met in \(84\%\) of all cases (839 in \(10^3\) realizations), while the second scenario is present in \(12\%\) of all cases (118 in \(10^3\) realizations). These fractions are stable upon increasing the size of the sample on which the above estimations were carried out. Interestingly enough, realizations where \(X(f)\) as a function of \(f\) fall into neither of the above two classes amount only to \(4\%\) of all possible cases.

It is relevant to note that the fractions of the two classes show tendency to move towards each other upon decreasing the size of the system. For instance, the fractions of the first and the second class amount to \(18\%\) and \(76\%\), respectively, for \(L=10^4 (T=\sigma=2\alpha=1)\). These fractions were estimated by criteria \(X(f=0) > 0.8 \times 10^4\) and \(X(f=0) < 10^2\), respectively.
Recall in this context that the chosen values for $L$ are sensible, since the typical DNA samples used in experiment have $L \sim 10^4 - 10^5$; see, e.g., Refs. [1,2] and also the following section.

Since there are typical realizations which are so much different from each other, we conclude that this long-range situation is essentially non-self-averaging in the whole physical domain $0 < f < 1$ and, in particular, in the thermodynamical domain of $f$. This fact distinguishes between the uncorrelated (white noise) and long-range correlated situations. It should be noted that due to the law of large numbers any non-self-averaging present in the whole domain $0 < f < 1$ is certainly impossible for the uncorrelated (or weakly correlated) noise [21,22]. For the long-range correlated case the very law of large numbers does not apply, and the above effect becomes possible.

Our discussion of the frozen noise presented in Sec. IV allows to provide a qualitative explanation for features of the above two classes of typical realizations. One notes that a sizable portion of long-range correlated noise realizations can be seen as several pieces of the frozen noise with different $\eta$’s put next to each other. Now recall from Eq. (56) that every sufficiently long piece of that type has a single first-order phase transition with a jump proportional to its length.

The same reasoning can be applied for the understanding of the existence of the second class, where $X(f)$ is a smooth function of $f$ and $X(f=0) \ll L$. Here one should note that — within the above qualitative image of a long-range correlated random sequence — there are realizations of the noise where all $\eta$’s are positive, and thus all jumps of $X(f)$ can occur only for negative $f < 0$, that is, beyond the domain of our interest.

2. Inferring phase transitions

Let us finally discuss on whether we can infer phase transitions by studying the typical realizations. First of all, it is obvious that once we do not have self-averaging, phase transitions should be studied on typical scenarios of behavior for $X$ and not on the behavior of its average $\langle X \rangle$. There is another aspect which is certainly more subtle: phase transitions are typically defined in the thermodynamical limit and one needs special tools of finite-size scaling for their identification in results of numerical computations which are necessarily done on finite systems. The idea of the finite-size scaling is thus to extrapolate these results to the thermodynamical limit. However, there is another, somewhat different line of thought [24] which identifies the proper thermodynamical quantities (such as entropy, free energy, order parameter, etc.) directly for finite systems, and then searches in the space of parameters some points having a special character for these quantities. This approach well recommended itself for studying phase transitions in atomic and nuclear physics, and in systems with long-range interactions (e.g., a gas of self-gravitating particles). For the present study of DNA there is a related aspect that should be taken into account: in natural conditions the number of base pairs is large, but finite. Here $L$ is of order of $10^4 - 10^5$, see, e.g., Refs. [1,2], as we have mentioned already. It is, therefore, clear that the considered finite size aspect of DNA is something generic, and not only connected with natural limitations of numerical methods.

Let us now return to the situation presented in Figs. 7–9. We are going to use the analogy with the case of the totally correlated noise described in Sec. IV. It was seen already that this analogy helped us to draw useful qualitative conclusions on the numerical data. For the totally correlated noise the point of the phase transition is unambiguously identified with the realization-dependent value $f = -\eta$. At this point the order parameter $X$ has a jump of order $L$; see Eqs. (56) and (57). It may be useful to repeat that the most unusual aspect of this phase transition is that its point is strictly realization dependent. The same philosophy can now be applied to Figs. 6 and 7: there are realization-dependent values of $f$, where $X$ has jumps of order of $L/2$ (recall that for the figures we have taken $L=10^4$ or $L=5 \times 10^4$). It is seen as well that there can be several such phase transitions for a single system (single realization of noise). The latter fact can by itself appear to be rather surprising. However, it is known that some disordered [22], or deterministic but strongly frustrated [25], systems can experience several phase transitions; there can even exist quasicontinuous domains of criticality [25]. With the same logic one sees that the typical realizations presented in Figs. 8 and 9 do not have phase transitions in the domain $0 < f < 1$ at least for the considered values of $L$.

C. The behavior of $\langle X \rangle$

As we already noted, once the effect of non-self-averaging is present, the basic physical quantities are the typical realizations, since it is these features that are directly observed in experiments. It is, however, still of relevance to know the behavior of the average number of unzipped bonds $\langle X \rangle$, since it illustrates what are the precise differences as compared to typical realizations.

Here we report on two features of $\langle X \rangle$ as a function of $f$. The first one is how does $\langle X \rangle$ depend on $f$ for small values of $f$. In particular, is there any power-law dependence similar to $\langle X(f) \rangle \propto f^2$ present in the uncorrelated-noise situation, and verified by us numerically in Sec. VA? Note that for the
The values of $\delta(\alpha)$ for several $\alpha$’s are shown by Fig. 14. For $\alpha=0$, we get $\delta=0.01$ which is in a good agreement with exact value $\delta(\alpha=0)=0$ obtained in Sec. IV.
different correlation properties of their base sequences. Molecules belonging to different evolutionary classes have inclusions. The results will be useful especially for drawing qualitative conclusions. In the present paper we have studied how statistical correlations present in the base sequence of a DNA molecule influence the process of unzipping. There were two related motivations for our study. On the one hand, the existence of these correlations — that can have both finite-range and long-range character — is by now a well-established fact [9–12]. It is, therefore, legitimate to study how they influence on the DNA physics. On the other hand, general qualitative predictions drawn on the above influence can be used for explaining the reason of rich correlated structures found in the base sequence of DNA. Recall that various segments of a DNA molecule can have different — finite-range or long-range [11,12] — correlation structures. Moreover, DNA molecules belonging to different evolutionary classes have different correlation properties of their base sequences [10].

The model we studied contains only the most minimal number of ingredients needed to describe unzipping, and to account for correlations in the base sequence of DNA. Therefore, many realistic features of the unzipping process remain beyond our study. We, nevertheless, believe that the obtained results will be useful especially for drawing qualitative conclusions.

Let us now summarize our results starting from the finite-range correlated situation. In Sec. III A we have shown that the presence of a finite correlation length $\tau$ plays a stabilizing role for the unzipping process: for a fixed external force $f$, the average number of broken base pairs decreases under increasing $\tau$. If only finite-range correlations are present, the process of unzipping does not depend much on the detailed structure of the base sequence: all typical — i.e., frequently met among all possible base sequences — scenarios of unzipping have the same qualitative pattern of behavior, that is, the number of the broken base pairs diverges as the external force approaches its critical value: $f \to 0$. This divergence can be adequately understood by studying the average — over all possible base sequences — number of broken base pairs. All by one, one can say that the basic influence of finite-range correlations is in stabilizing the DNA molecule with respect to the external unzipping force.

The influence of long-range correlations is certainly more drastic. Possibly the most important aspect is that the situation is essentially non-self-averaging: there are two radically different scenarios of typical unzipping which depend on the detailed structure of the base sequence and which do not coincide with the behavior averaged over all possible base sequences. Within the first scenario, the number of broken base pairs $X(f)$ shows as a function of the external force $f$ a sequence of sharp jumps at sequence-dependent values of $f$. The overall number of jumps is nearly constant within the class. Each jump has the magnitude comparable with $L$, that is, under small change of $f$ a large number of base pairs can be opened. The point $f=0$ is special, since $X(0)$ either coincides with $L$ or at least is very close to it. We argued in Sec. V that it is sensible to describe this scenario as a sequence of phase transitions. Such an effect is known from other disordered or strongly frustrated systems [22,25].

The second typical scenario is crucially different. Now $X$ is a smooth, slowly changing function of the external force $f$ in the whole relevant domain $0 < f < 1$. There is no sign of phase transition, and the value $f=0$ is not distinguished from $f>0$ as far as $X$ is concerned. DNA molecules which due to the structure of their base sequence fall into this class are thus rather stable with respect to the external unzipping force.

It appears, interestingly enough, that the qualitative and even some quantitative features of the long-range correlated situation can be understood via the analytical solution of the model with the totally correlated (frozen) noise, which we presented in Sec. IV. In particular, this allows us to explain why there exist two typical scenarios with widely different behavior of the number of unzipped base pairs, and provides rather robust analytical indications for the phenomenon of non-self-averaging.

Summarizing features of these two scenarios, one can say that long-range correlations increase the adaptability of the corresponding DNA molecule, since in some typical scenarios it becomes more stable with respect to the force (any sharp transition is absent), while in others the unzipping is realized via a sequence of sharp phase transitions. The actual scenario for a single molecule will crucially depend on the detailed structure of the base sequence.

We also studied how the average number $\langle X \rangle$ of unzipped base pairs depends on the applied force $f$. In contrast to...
white-noise situation, where the behavior of \( \langle X \rangle \) for small (that is, critical) forces \( f \to 0 \) is governed by a power law \( \langle X \rangle \sim f^2 \), we found numerically no indications of a power law for small forces in the long-range correlated situation. In contrast, the dependence of \( \langle X(f) \rangle \) on \( f \) for small \( f \)'s is very weak and to a large extent is governed by \( \langle X(f=0) \rangle \). The latter quantity displays a power-law behavior (78) as a function of \( L \). The region of validity of this power law appeared to be unexpectedly wide.

We hope that these results will contribute to the understanding of the role and the purpose of correlation structures in DNA.

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**APPENDIX A: GENERATION OF THE LONG-RANGE CORRELATED NOISE**

Using ideas of Ref. [23] we shall here describe a method for numerical generation of a Gaussian random noise \( \eta(t) \) with zero average and an arbitrary symmetric autocorrelation function:

\[
K(t-t') = \langle \eta(t) \eta(t') \rangle, \quad (A1)
\]

\[
K(t) = K(-t). \quad (A2)
\]

Assume that the noise is periodic with period \( M \):

\[
\eta(t) = \eta(t+M). \quad (A3)
\]

Therefore \( K(t) \) is also periodic with the same period and can be expanded as

\[
K(t) = \sum_{n=-\infty}^{\infty} k_n e^{-\text{i} \omega_0 t}, \quad \omega_0 = \frac{2 \pi}{M}, \quad (A4)
\]

where \( k_n \) is given by Fourier formula

\[
k_n = \frac{1}{M} \int_{-M/2}^{M/2} dt K(t) e^{\text{i} \omega_0 t}. \quad (A5)
\]

Since \( K(t) \) is a real and symmetric function, \( k_n = k_n^* = k_{-n} \), and thus

\[
k_n = \frac{2}{M} \int_0^{M/2} dt K(t) \cos(n \omega_0 t). \quad (A6)
\]

It is now straightforward to see that the noise \( \eta \) we are looking for is represented as:

\[
\eta = \sum_{n=-\infty}^{\infty} \sqrt{k_n} \eta_n e^{-\text{i} \omega_0 t}, \quad (A7)
\]

where \( \eta_n \) are complex Gaussian random variables with

\[
\langle \eta_n \eta_m \rangle = \delta(n+m), \quad (A8)
\]

where \( \delta(0)=1 \) and \( \delta(k)=0 \) for \( k \neq 0 \). Indeed, once \( \eta_n \) are assumed to be Gaussian, \( \eta(t) \) is Gaussian as well; it is seen as well that Eq. (A1) is valid. Complex random variables \( \eta_n \) can be conveniently expressed via real random variables:

\[
\eta_n = \frac{1}{\sqrt{2}} (a_n + ib_n) \quad \text{for} \quad n \geq 1, \quad (A9)
\]

\[
\eta_n = \frac{1}{\sqrt{2}} (a_n - ib_n) \quad \text{for} \quad n \leq -1, \quad (A10)
\]

\[
\eta_0 = a_0. \quad (A11)
\]

where quantities \( a_n \) and \( b_n \) are independent, zero-average Gaussian random variables normalized to one:

\[
\langle a_n a_{n'} \rangle = \delta_{n0}, \quad \langle b_n b_{n'} \rangle = \delta_{n0}, \quad \langle a_n b_{n'} \rangle = 0. \quad (A12)
\]

Using this one writes

\[
\eta = \sum_{n=1}^{\infty} \sqrt{2k_n} [a_n \cos(n \omega_0 t) + b_n \sin(n \omega_0 t)] + \sqrt{k_0} a_0. \quad (A13)
\]

Let us consider an example:

\[
K(t) = \sigma \quad \text{for} \quad t < 1
\]

\[
= \frac{\sigma}{\sqrt{t}} \quad \text{for} \quad t \geq 1. \quad (A14)
\]

This represents a long-range correlated noise regularized for small \( t \). For this autocorrelation function the coefficients \( k_n \) read from Eq. (A6):

\[
\frac{k_n}{k_0} = 2 \left( \frac{\sqrt{2}}{\sqrt{M}} - \frac{1}{M} \right), \quad (A15)
\]

where \( F_n(x) \) is Fresnel’s C function:

\[
F_n(x) = \int_0^x dt \cos \left( \frac{\pi t^2}{2} \right). \quad (A16)
\]

**Numerical implementation**

Equations (A13) and (A15) are sufficient for generating long-range correlated, periodic Gaussian random noise. However, for numerical implementations this noise has to be
discretized. First we note that the above method produces periodic random noise (with period $M$), while our problem does not have any periodicity. Therefore, we have chosen $M=2L$, and took discrete values of $t=1,2,\ldots,L$ in Eq. (A13), thereby generating $L$ long-range correlated random numbers without any periodicity. Note that Eq. (A13) contains infinity as the upper limit of the summation in its RHS. For numerics this infinity should obviously be substituted by some number larger than $L$, and additionally one should check that the situation is stable with respect to varying this number. As for concrete calculations we have used, e.g., $L=10^4$, we found sufficient to take for this upper summation number. As for concrete calculations we have used, e.g., $L=10^4$, we found sufficient to take for this upper summation limit $10^4$.

Numerical simulations in Sec. V were performed by using the Gaussian independent random variables generated by the “gasdev” algorithm of Ref. [26]. The long-range correlated noise was generated following the scheme proposed in this appendix.

APPENDIX B: DERIVATION OF TWO ASYMPTOTIC RELATIONS

Here we derive the following asymptotic identities used in the main text:

$$\int_0^\infty d\xi \frac{e^{-a\xi^2}}{\sqrt{2\pi}} = \frac{e^{-a/2}}{a^{1/2} \pi^{1/2}} \left(1 - \frac{1}{a^2} + \cdots\right), \quad a \gg 1, \quad (B1)$$

$$\int_0^a d\xi e^{a\xi^2} = \int_0^a d\xi e^{a\xi^2} = \frac{e^{a/2}}{a} \left( 1 + \frac{1}{a^2} + \cdots \right), \quad a \gg 1. \quad (B2)$$

The first one is easily done via integration by parts:

$$\int_0^\infty \frac{d\xi}{\sqrt{2\pi}} e^{-\xi^2} = -\int_0^\infty \frac{d[ae^{-\xi^2}]}{\sqrt{2\pi} \xi} = \frac{e^{-a/2}}{a \sqrt{2\pi}} + \int_0^\infty \frac{d[ae^{-\xi^2}]}{\sqrt{2\pi} \xi}. \quad (B3)$$

For the second relation one notes that for $a \gg 1$ the relevant domain of integration is $\xi \sim a$. In more details,

$$\int_0^a d\xi e^{a\xi^2} = e^{a/2} \int_0^\infty d\xi e^{-a\xi^2} = \frac{e^{a/2}}{a} \int_0^a dy \ e^{-y^2/(2a^2)}. \quad (B4)$$

Now one can expand inside of the second exponent in the RHS of Eq. (B4), since the main contribution to the integral comes from $y \sim 0$ (the other side, that is, $y \sim a^2$, is strongly suppressed as seen):

$$\frac{e^{a/2}}{a} \int_0^a dy e^{-y^2/(2a^2)} = \frac{e^{a/2}}{a} \int_0^a dy \ e^{-\left(1 + \frac{y^2}{2a^2} + \cdots \right)}. \quad (B5)$$

Neglecting exponentially small terms, one gets get finally Eq. (B2).

[19] Note, however, that there are exclusions from this rule for certain bacteria [1] for them the concentration of GC pairs can differ substantially from that of AT pairs. These situations are, nevertheless, fairly rare.

