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Large Deviations in Renewal Models of Statistical Mechanics

Marco Zamparo

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 $\{(S_i, R_i)\}_{i \ge 1}$ sequence of i.i.d. random vectors on $(\Omega, \mathcal{F}, \mathbb{P})$ so that

- $S_i \in \{1, 2, \ldots\} \cup \{\infty\}$ is a "waiting time";
- $T_i := S_1 + \ldots + S_i$ is a "renewal time" and $T_0 := 0$;
- $R_i \in \mathbb{R}^d$ is a "reward" associated with S_i .

Basics

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For each integer time $t \ge 0$

- $X_t := \mathbb{1}(t \notin \{T_i\}_{i \ge 0})$ is the (non-)renewal indicator;
- $N_t := \sup\{i \ge 0 : T_i \le t\}$ is the number of renewals by t;
- $W_t := \sum_{i=1}^{N_t} R_i$ is the total reward by t and $W_t := 0$ if $N_t = 0$.

Constrained Renewal Model (1)

Renewal equation:

$$\mathbb{P}[X_t = 0] = \begin{cases} 1 & \text{if } t = 0; \\ \sum_{s=1}^t p(s) \mathbb{P}[X_{t-s} = 0] & \text{if } t > 0, \end{cases}$$

where $p(s) := \mathbb{P}[S_1 = s]$ is the "waiting time distribution".



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Lemma

 $\mathbb{P}[X_t = 0] > 0$ for all t sufficiently large if $gcd\{s \ge 1 : p(s) > 0\} = 1$.

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If $gcd{s \ge 1 : p(s) > 0} = 1$, then conditioning on ${X_t = 0}$

• is well-defined for all *t* sufficiently large;

• yields the new model $(\Omega, \mathcal{F}, \mathbb{P}_t)$ with $\frac{d\mathbb{P}_t}{d\mathbb{P}} := \frac{1 - X_t}{\mathbb{P}[X_t = 0]}$.

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Constrained Renewal Model (2)

Distribution of waiting times:

$$\mathbb{P}_t\left[S_1=s_1,\ldots,S_n=s_n,N_t=n\right]=\frac{\mathbb{I}\left(\sum_{k=1}^n s_k=t\right)}{\mathbb{P}[X_t=0]}\prod_{k=1}^n p(s_k).$$

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Distribution of renewal indicators:

$$\mathbb{P}_t[X_1 = x_1, \dots, X_t = x_t] = \frac{1 - x_t}{\mathbb{P}[X_t = 0]} \prod_{s=1}^t \left[p(s) \right]^{\#_{s|t}(0, x_1, \dots, x_t)}$$

with $\#_{s|t}(x_0, \ldots, x_t) := \sum_{i=1}^{t-s+1} (1-x_{i-1}) \left(\prod_{k=i}^{i+s-2} x_k \right) (1-x_{i+s-1}).$

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...and they look like finite-volume Gibbs states!

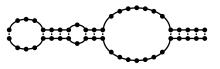
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Polymer Localization and DNA Denaturation

Pinned polymer (Fisher):



DNA molecule (Poland-Scheraga):



t monomers (per strand)

- in *n* stretches of lengths s_k with $s_1 + \cdots + s_n = t$ and only the first monomer bound;
- with binding energy ϵ ;
- with loop entropy σ_s so that $\sigma_s \leq b s$.

Statistical weight:

$$\prod_{k=1}^{n} \exp(\sigma_{s_{k}} - \epsilon + \eta \mathbf{s}_{k}).$$

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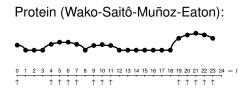
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Take η so that $p(s) := \exp(\sigma_s - \epsilon + \eta s)$ satisfies $\sum_{s=1}^{\infty} p(s) \le 1!$

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Protein Folding



t peptide bonds

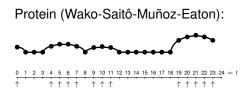
- $x_i = 1$ native, $x_i = 0$ non-native;
- interact only if belong to the same native stretch;
- with contact energy ϵ_{j-i+1} so that $u_s := \sum_{k=1}^{s} (s-k)\epsilon_k \ge b s;$
- order with entropic loss σ .

Statistical weight:

$$\exp\left[-\sum_{i=0}^{t-1}\sum_{j=i}^{t-1}\epsilon_{j-i+1}\prod_{k=i}^{j}x_{k}+\sigma\sum_{i=0}^{t-1}(1-x_{i})+\eta t\right]$$

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Statistical weight:

$$\exp\left[\sum_{s=1}^{t} \left(\sigma - \boldsymbol{u}_{s} + \eta \boldsymbol{s}\right) \#_{s|t}(\boldsymbol{x}_{0}, \dots, \boldsymbol{x}_{t-1}, \boldsymbol{0})\right]$$

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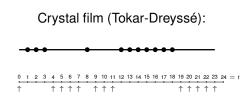
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Strained Epitaxy



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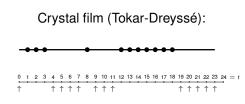
t lattice sites

- $x_i = 1$ occupied, $x_i = 0$ empty;
- particles interact only if belong to the same cluster;
- with energetic gain u_s so that u_s ≥ bs;
- with chemical potential μ .

 $\exp\left[-\sum_{s=1}^{t} u_{s} \#_{s|t}(\mathbf{x}_{0}, \dots, \mathbf{x}_{t-1}, 0) + \mu \sum_{i=0}^{t-1} x_{i} + (\eta - \mu)t\right]$

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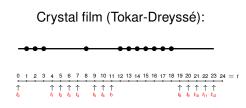
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Total Reward $W_t := \sum_{i=1}^{N_t} R_i$ in Statistical Mechanics

In pinned-polymer model and Poland-Scheraga model

- $R_i := 1 \implies W_t$ counts bound monomers;
- $R_i := \sigma_{S_i} \implies W_t$ is the total loop entropy;
- $R_i := (1, \sigma_{S_i}) \implies W_t$ combines both of them;
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In Wako-Saitô-Muñoz-Eaton model and Tokar-Dreyssé model

- $R_i := S_i 1 \implies W_t$ counts native bonds or particles;
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In all cases

R_i := (1(*S_i* = 1),..., 1(*S_i* = *d*)) ⇒ *W_t* counts waiting times;
 ...

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Quantifying Rare Events

Problem: study $\mathbb{P}_t \left[\frac{W_t}{t} \in A \right]$ for large *t* and some $A \subseteq \mathbb{R}^d$ measurable.

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Definition

 W_t satisfies a large deviations principle (LDP) with rate function I if

(a)
$$I: \mathbb{R}^d \to [0,\infty) \cup \{\infty\}$$
 is lower semicontinuous;

(b)
$$\limsup_{t\uparrow\infty} \frac{1}{t} \ln \mathbb{P}_t \left[\frac{W_t}{t} \in F \right] \le -\inf_{w\in F} \{I(w)\} \text{ for each } F \subseteq \mathbb{R}^d \text{ closed};$$

(c)
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Weak large deviations principle if (b) valid only for F compact.

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Main Results

Let $\zeta(k)$ be the extended real number defined for each $k \in \mathbb{R}^d$ by

$$\zeta(k) := \inf \left\{ z \in \mathbb{R} \, : \, \mathbb{E} \Big[\exp \left(k \cdot \boldsymbol{R}_1 - z \, \boldsymbol{S}_1 \right) \mathbb{1}(\boldsymbol{S}_1 < \infty) \Big] \le 1 \right\} > -\infty.$$

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$$\zeta(k) := \inf \left\{ z \in \mathbb{R} \, : \, \mathbb{E} \Big[\exp \left(k \cdot \boldsymbol{R}_1 - z \, \boldsymbol{S}_1 \right) \mathbb{1}(\boldsymbol{S}_1 < \infty) \Big] \le 1 \right\} > -\infty.$$

Theorem

$$\lim_{t\uparrow\infty}\frac{1}{t}\ln\mathbb{E}_t\left[\exp(k\cdot W_t)\right] =: c(k) \text{ exists and } c(k) = \zeta(k) - \zeta(0).$$

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Theorem

(a) W_t satisfies a weak LDP with rate function I.

(b) If $0 \in int\{k \in \mathbb{R}^d : c(k) < \infty\}$, then W_t satisfies an LDP with I.



c non-differentiable on $int \{ k \in \mathbb{R}^d : c(k) < \infty \}$ in general

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Gärtner-Ellis Theorem does not apply

(based on a change of measure)

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an original proof is needed!

(based on subadditivity arguments like for Cramér's Theorem)

Final Remarks

On the mathematical side

• an LDP established for the constrained renewal model $(\Omega, \mathcal{F}, \mathbb{P}_t)$;

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On the physical side

• rate function / with singularities.