

Heterogeneity and chaos: Granular chains and DNA models

Haris Skokos

**Department of Mathematics and Applied Mathematics,
University of Cape Town
Cape Town, South Africa**

E-mail: haris.skokos@uct.ac.za

URL: http://math_research.uct.ac.za/~hskokos/

**Work in collaboration with
Vassos Achilleos, George Theocharis,
Adrian Schwellnus, Malcolm Hillebrand, George Kalosakas**

Outline

- Chaotic behavior of granular chains
 - ✓ Weakly nonlinear regime: Long-lived chaotic Anderson-like localization
 - ✓ Highly nonlinear regime: equilibrium chaotic state
- DNA models
 - ✓ Lyapunov exponents
 - ✓ Different dynamical regimes
 - ✓ DNA melting
 - ✓ Deviation Vector Distributions
- Summary

Energy Distributions

We consider normalized **energy distributions**

$$z_v \equiv \frac{E_v}{\sum_m E_m} \quad \text{with } E_v \text{ being the energy of particle } v.$$

Second moment: $m_2 = \sum_{v=1}^N (v - \bar{v})^2 z_v$ with $\bar{v} = \sum_{v=1}^N v z_v$

Participation number: $P = \frac{1}{\sum_{v=1}^N z_v^2}$

measures the number of stronger excited modes in z_v .

Single site excitation $P=1$. Equipartition of energy $P=N$.

Lyapunov Exponents (LEs) and Deviation Vector Distributions (DVDs)

Consider an orbit in the $2N$ -dimensional phase space with **initial condition $\mathbf{x}(0)$** and an **initial deviation vector from it $\mathbf{v}(0)$** . Then the mean exponential rate of divergence is:

$$\text{m L C E} = \lambda_1 = \lim_{t \rightarrow \infty} \frac{1}{t} \ln \frac{\|\vec{\mathbf{v}}(t)\|}{\|\vec{\mathbf{v}}(0)\|}$$

$\lambda_1=0 \rightarrow$ Regular motion $\propto (t^{-1})$

$\lambda_1 \neq 0 \rightarrow$ Chaotic motion

Deviation vector:

$$\mathbf{v}(t) = (\delta u_1(t), \delta u_2(t), \dots, \delta u_N(t), \delta p_1(t), \delta p_2(t), \dots, \delta p_N(t))$$

$$\text{DVD: } w_l = \frac{\delta u_l^2 + \delta p_l^2}{\sum_l (\delta u_l^2 + \delta p_l^2)}$$

Granular media

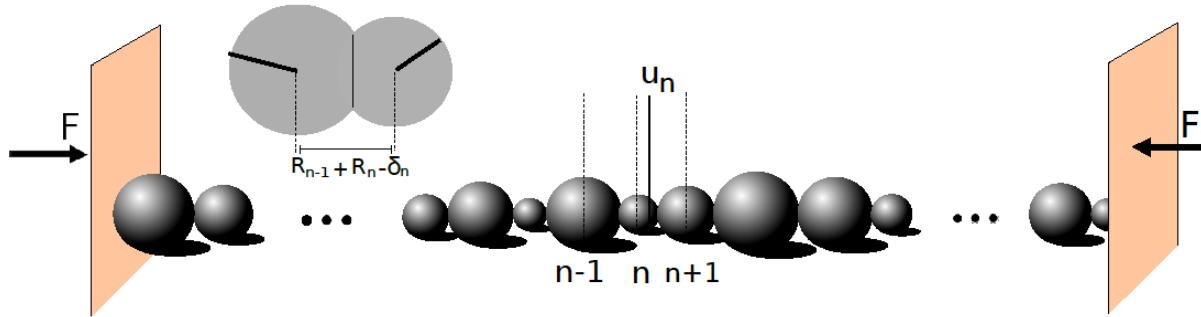


**Examples: coal, sand, rice,
nuts, coffee etc.**

1D granular chain (experimental control of nonlinearity and disorder)



Hamiltonian model



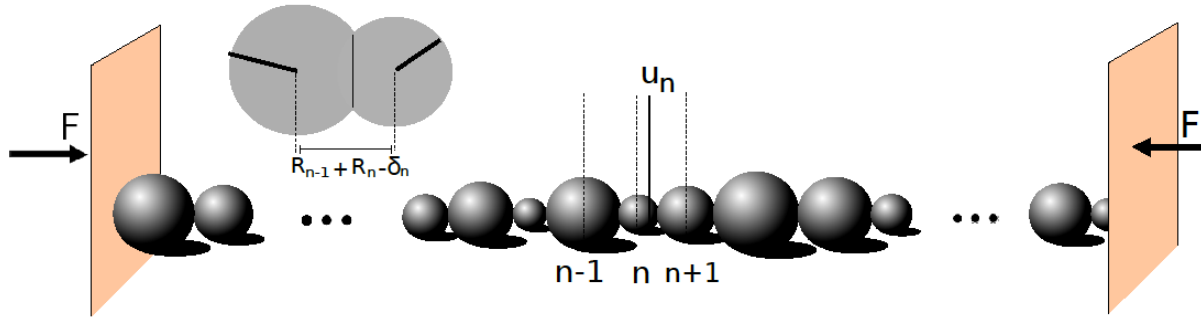
$$H = \sum_{n=1}^N \left(\frac{p_n^2}{2m_n} + \frac{2}{5} A_n [\delta_n + u_{n-1} - u_n]_+^{5/2} - \frac{2}{5} A_n \delta_n^{5/2} - A_n \delta_n^{3/2} (u_{n-1} - u_n) \right)$$

$$\delta_n = (F/A_n)^{2/3} \quad A_n = (2/3) \mathcal{E} \sqrt{(R_{n-1} R_n) / (R_{n-1} + R_n) / (1 - \nu^2)}$$

$[x]_+ = 0$ if $x < 0$: **formation of a gap**. ν : Poisson's ratio, \mathcal{E} : Elastic modulus.

Hertzian forces between spherical beads. Fixed boundary conditions.

Hamiltonian model



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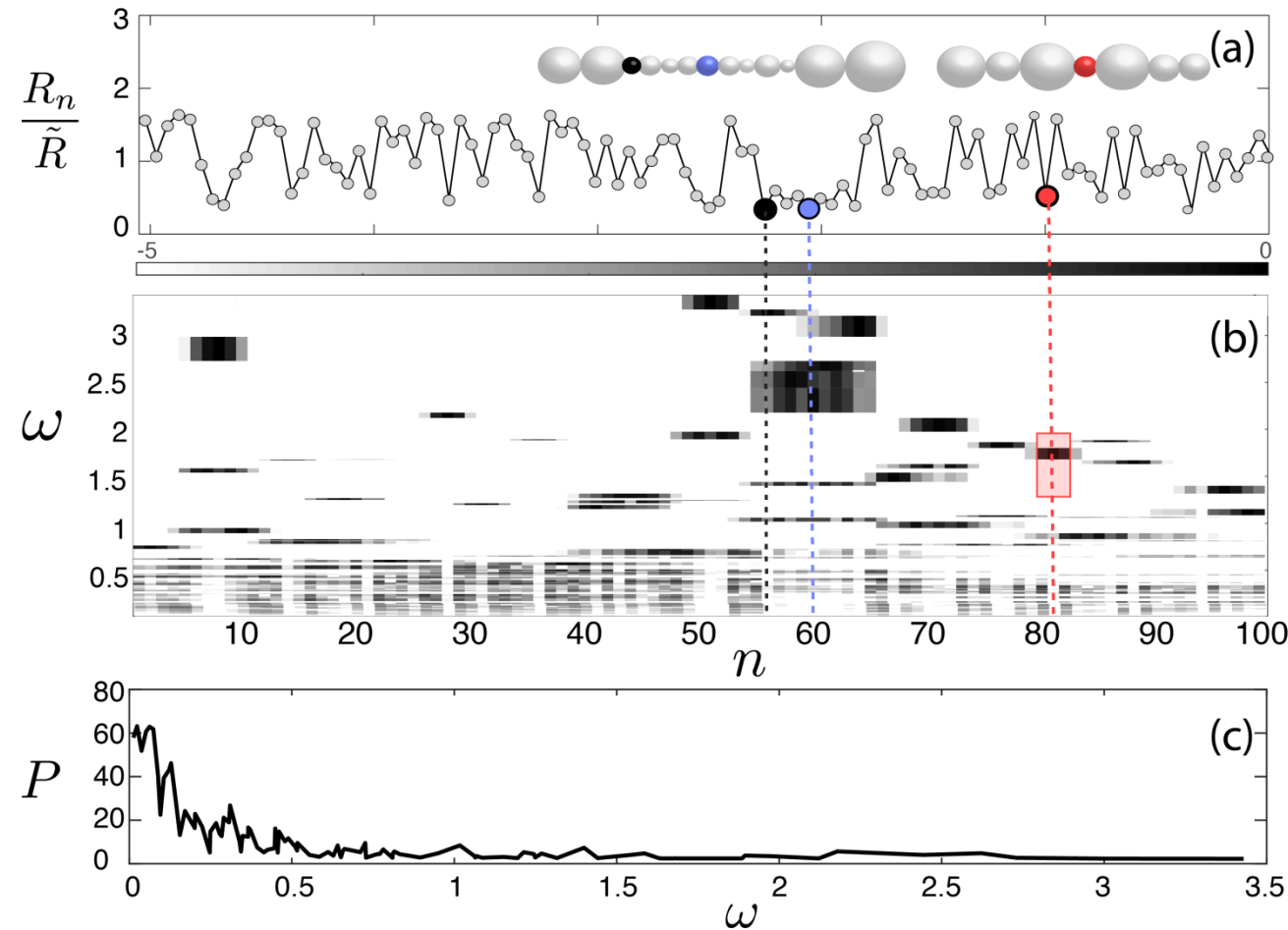
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Hertzian forces between spherical beads. Fixed boundary conditions.

Disorder both in couplings and masses

$R_n \in [R, \alpha R]$ with $\alpha \geq 1$

Eigenmodes and single site excitations

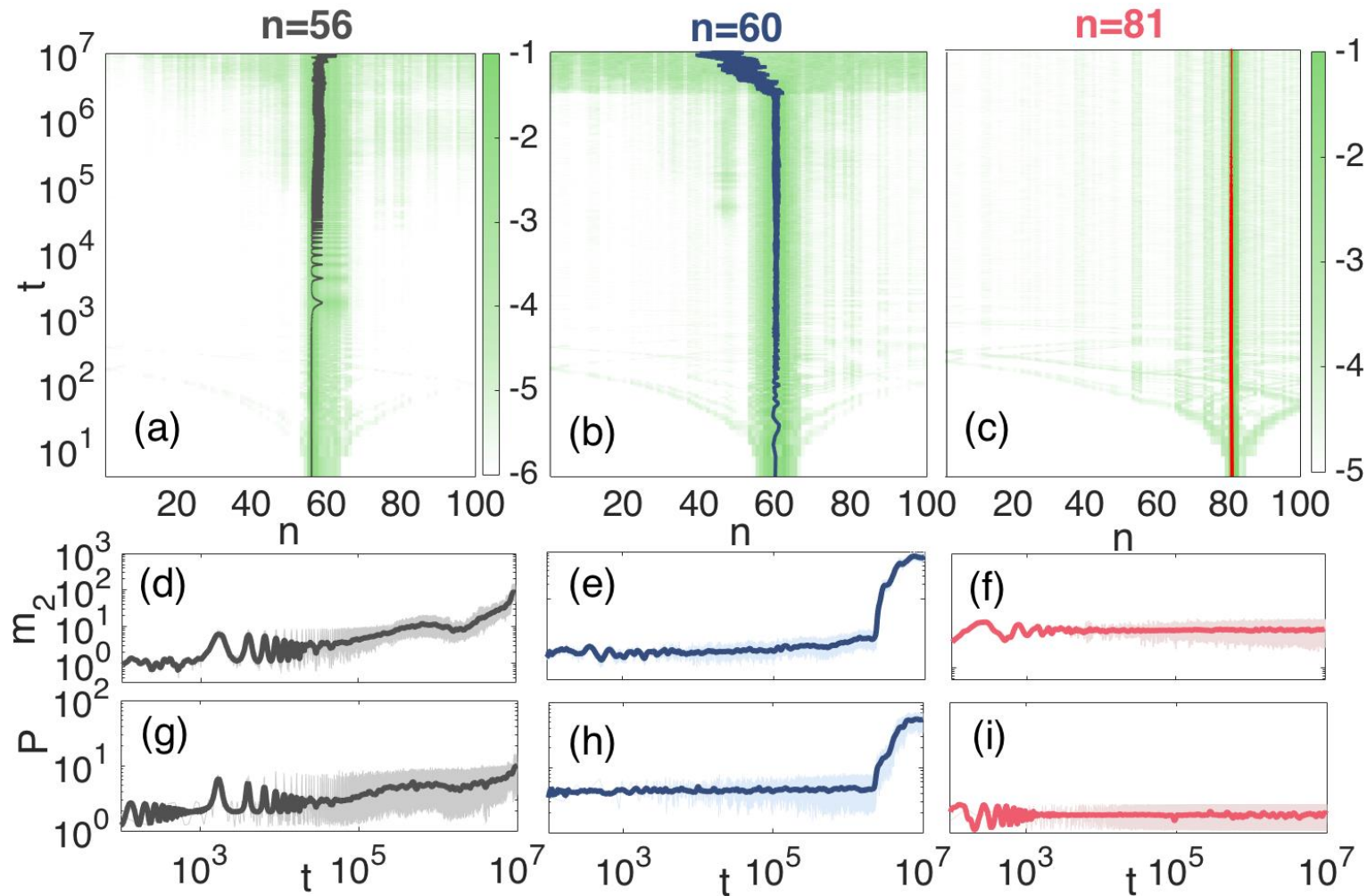


**Disorder realization
with $N=100$ beads**

**Displacement
excitation of bead n**

**Participation number
of eigenmodes.
About 10 extended
modes with $P > 40$**

Weak nonlinearity: Long time evolution

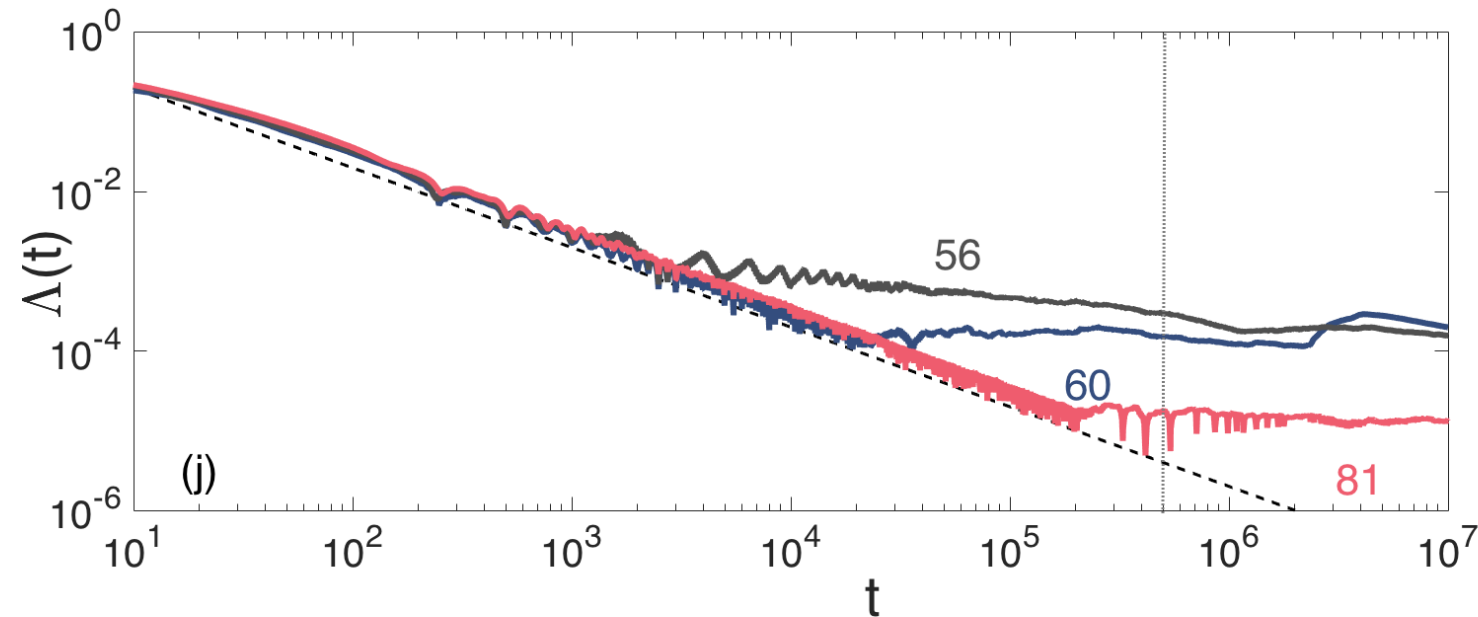


Delocalization

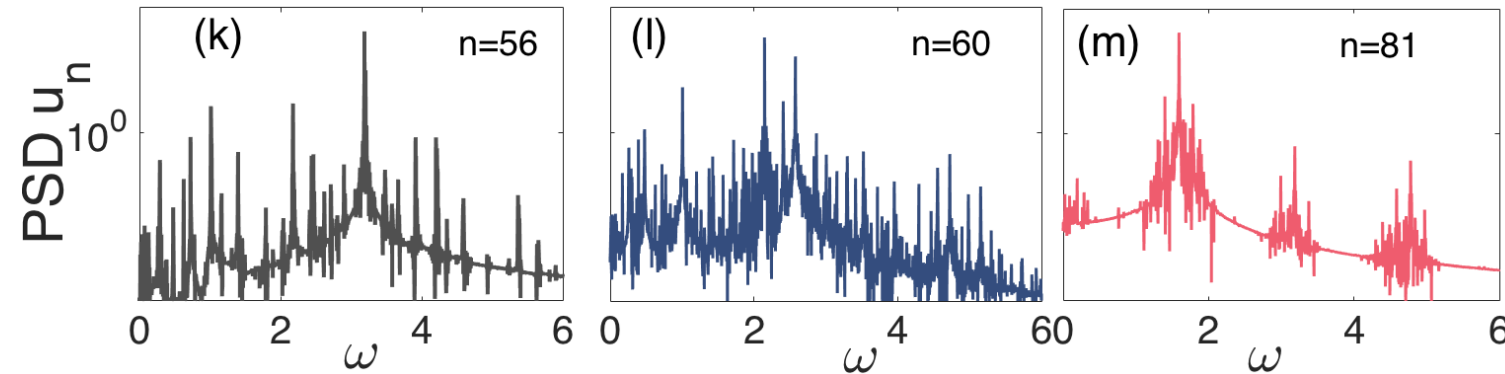
Delocalization

Localization

Weak nonlinearity: Chaoticity



mLCE

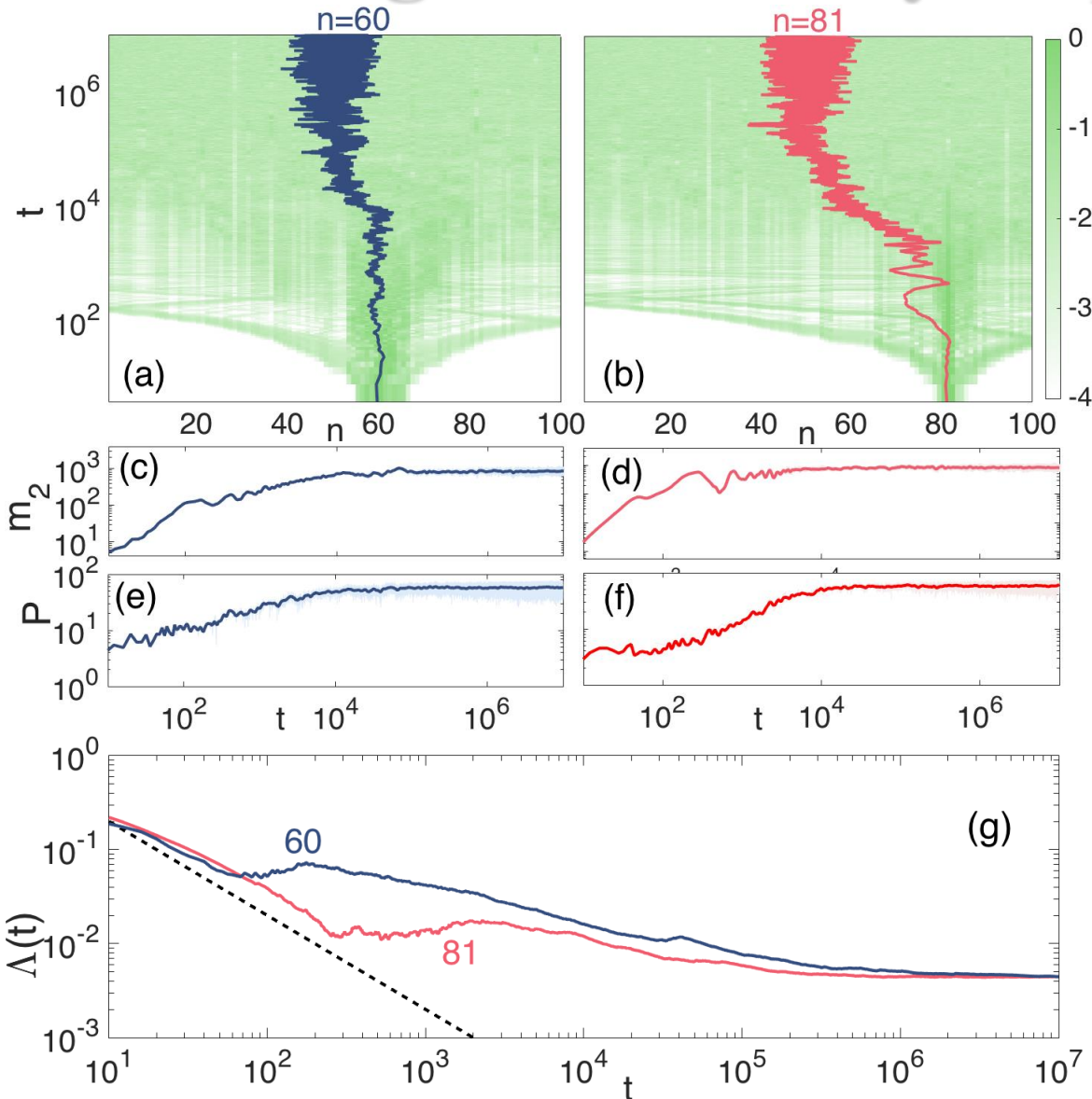


Power
Spectrum
Distribution

Weakly chaotic motion:
Delocalization

Long-lived chaotic
Anderson-like
Localization

Strong nonlinearity: Equipartition

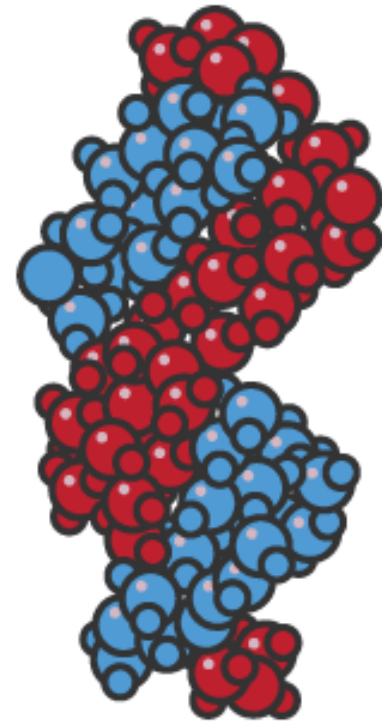
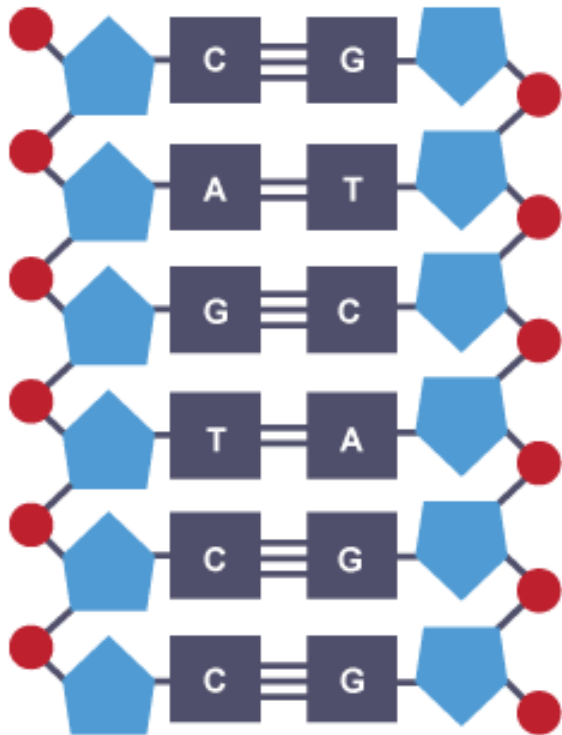


The granular chain reaches **energy equipartition** and an **equilibrium chaotic state**, independent of the initial position excitation.

DNA structure

Double helix with two types of bonds:

- **Adenine-thymine (AT) – two hydrogen bonds**
- **Guanine-cytosine (GC) – three hydrogen bonds**



Hamiltonian model

Peyrard-Bishop-Dauxois (PBD) model

[Dauxois, Peyrard, Bishop, PRE (1993)]

$$H_N = \sum_{n=1}^N \left[\frac{1}{2m} p_n^2 + D_n (e^{-a_n y_n} - 1)^2 + \frac{K}{2} (1 + \rho e^{-b(y_n + y_{n-1})}) (y_n - y_{n-1})^2 \right]$$

Bond potential energy (Morse potential)

GC: $D=0.075$ eV, $a=6.9$ Å⁻¹

AT: $D=0.05$ eV, $a=4.2$ Å⁻¹

Nearest neighbors coupling potential

$K=0.025$ eV/Å², $\rho=2$, $b=0.35$ Å⁻¹

Disorder realizations

Different arrangements of **AT** and **GC** bonds.

AT AT AT AT AT AT AT AT AT AT



$P_{\text{AT}}=1$ (100% AT bonds)

Disorder realizations

Different arrangements of **AT** and **GC** bonds.

AT AT AT AT AT AT AT AT AT AT



$P_{\text{AT}}=1$ (100% AT bonds)

GC AT AT GC GC GC GC AT AT GC



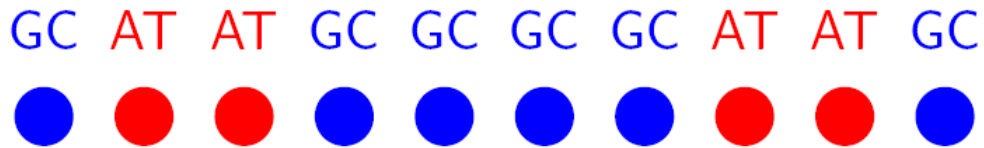
$P_{\text{AT}}=0.4$ (40% AT bonds)

Disorder realizations

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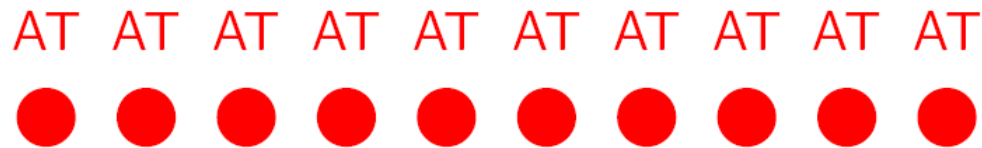


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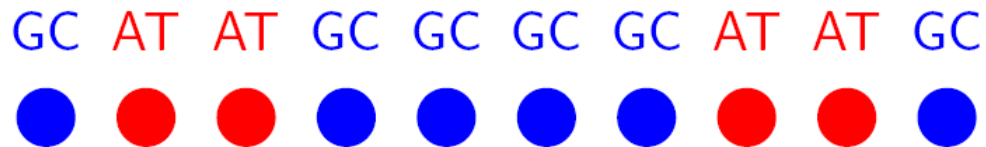


Disorder realizations

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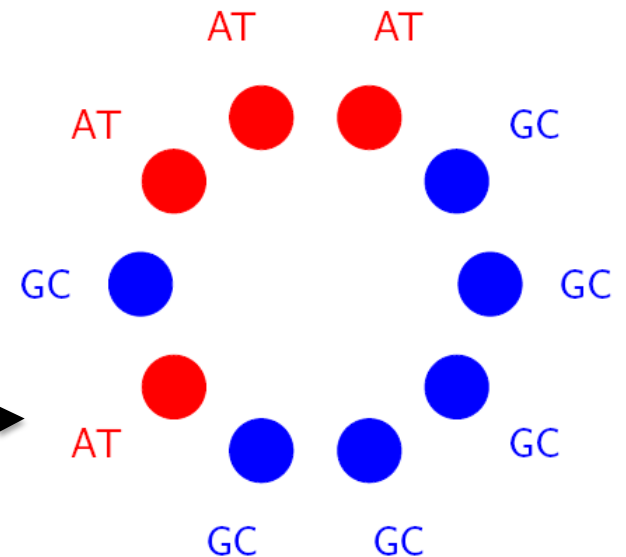
$P_{\text{AT}}=1$ (100% AT bonds)



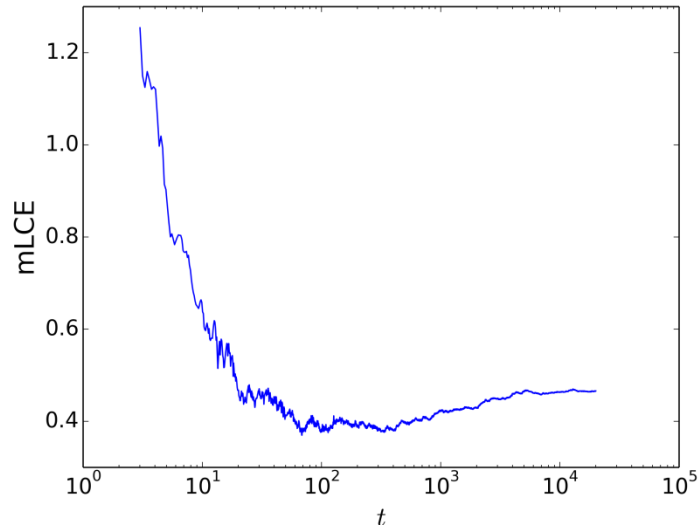
$P_{\text{AT}}=0.4$ (40% AT bonds)



Periodic boundary conditions

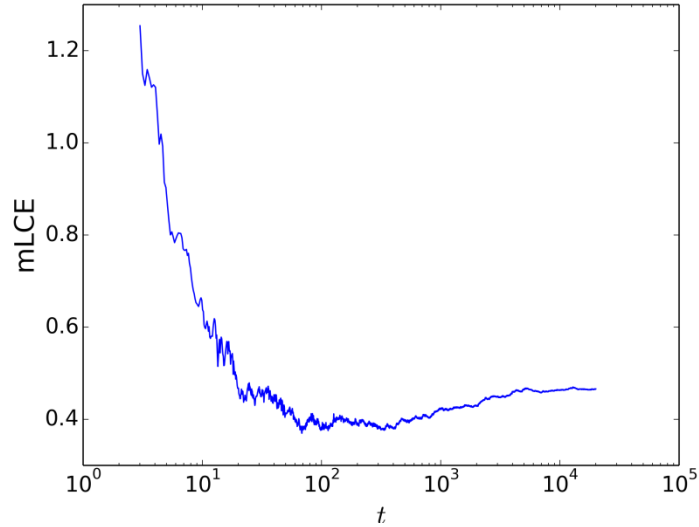


Lyapunov exponents ($E/n=0.04$, $P_{AT}=0.3$)



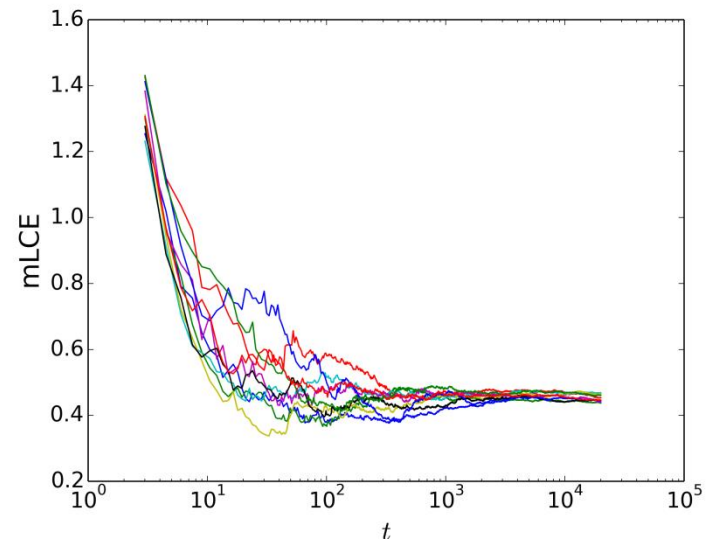
1 realization, 1 initial condition

Lyapunov exponents ($E/n=0.04$, $P_{AT}=0.3$)

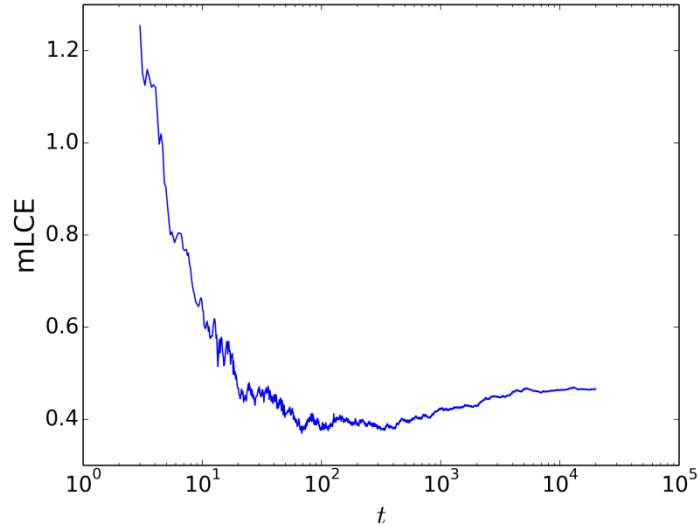


1 realization, 1 initial condition

1 realization, 10 initial conditions

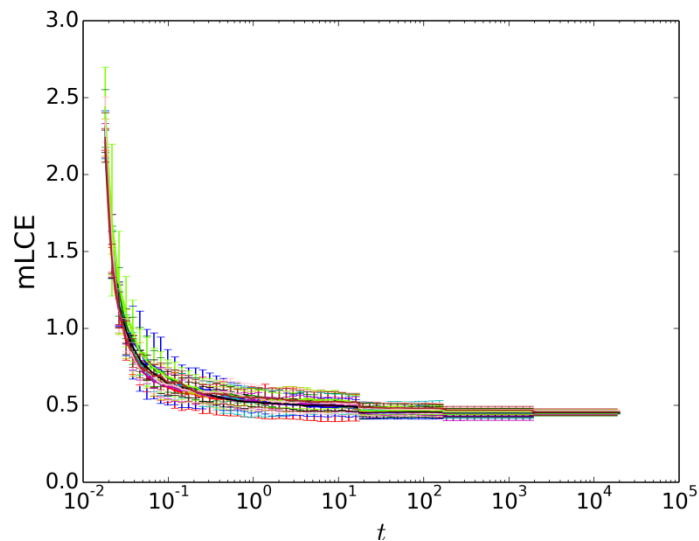
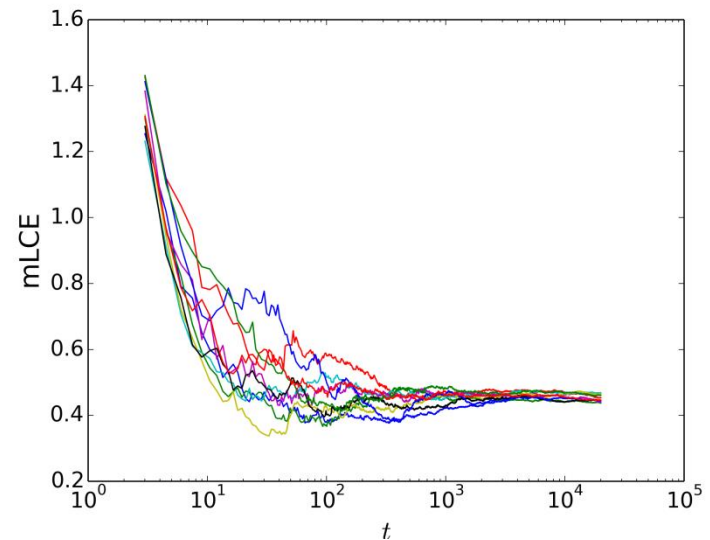


Lyapunov exponents ($E/n=0.04$, $P_{AT}=0.3$)



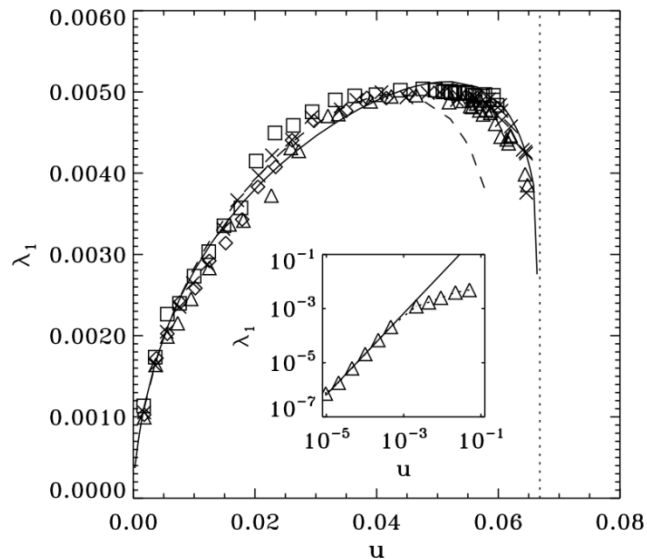
1 realization, 1 initial condition

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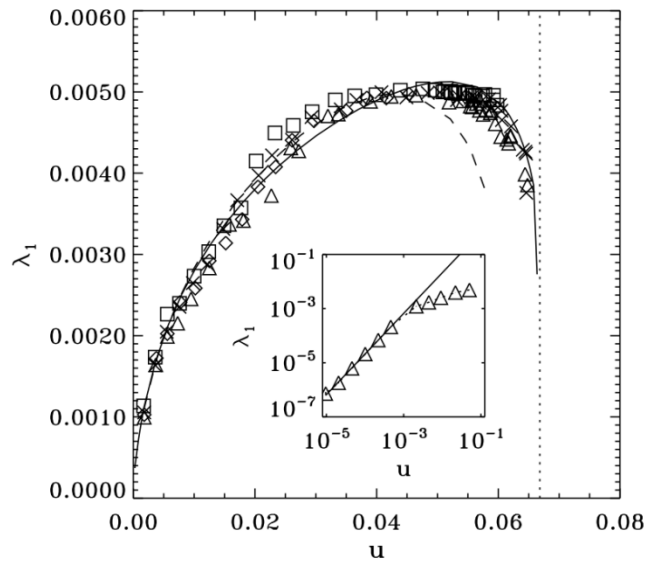
10 realizations, 10 initial conditions

Lyapunov exponent vs. energy per particle

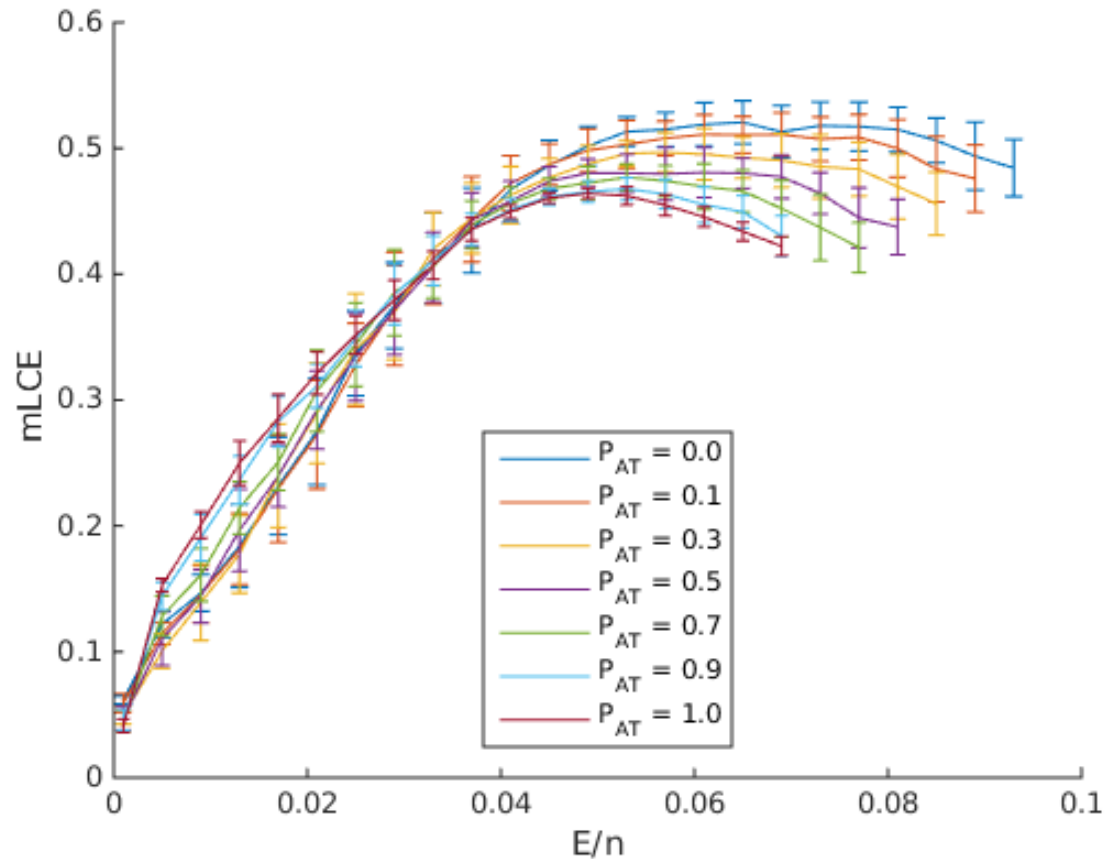


Homogeneous chain
[Barré & Dauxois,
EPL (2001)]

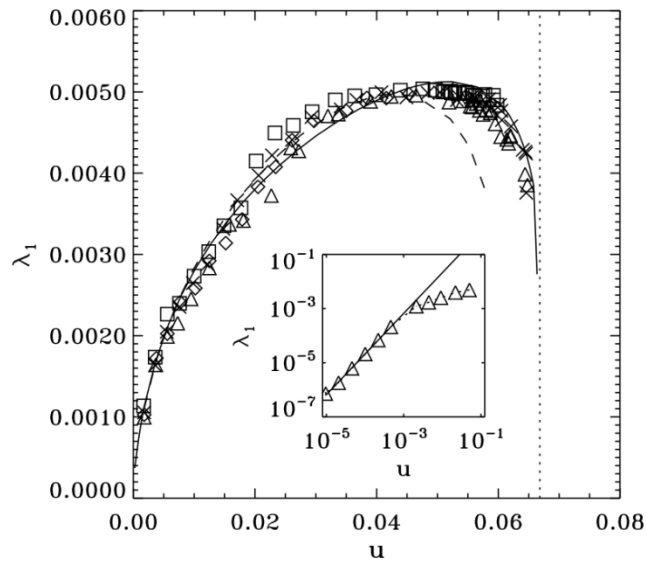
Lyapunov exponent vs. energy per particle



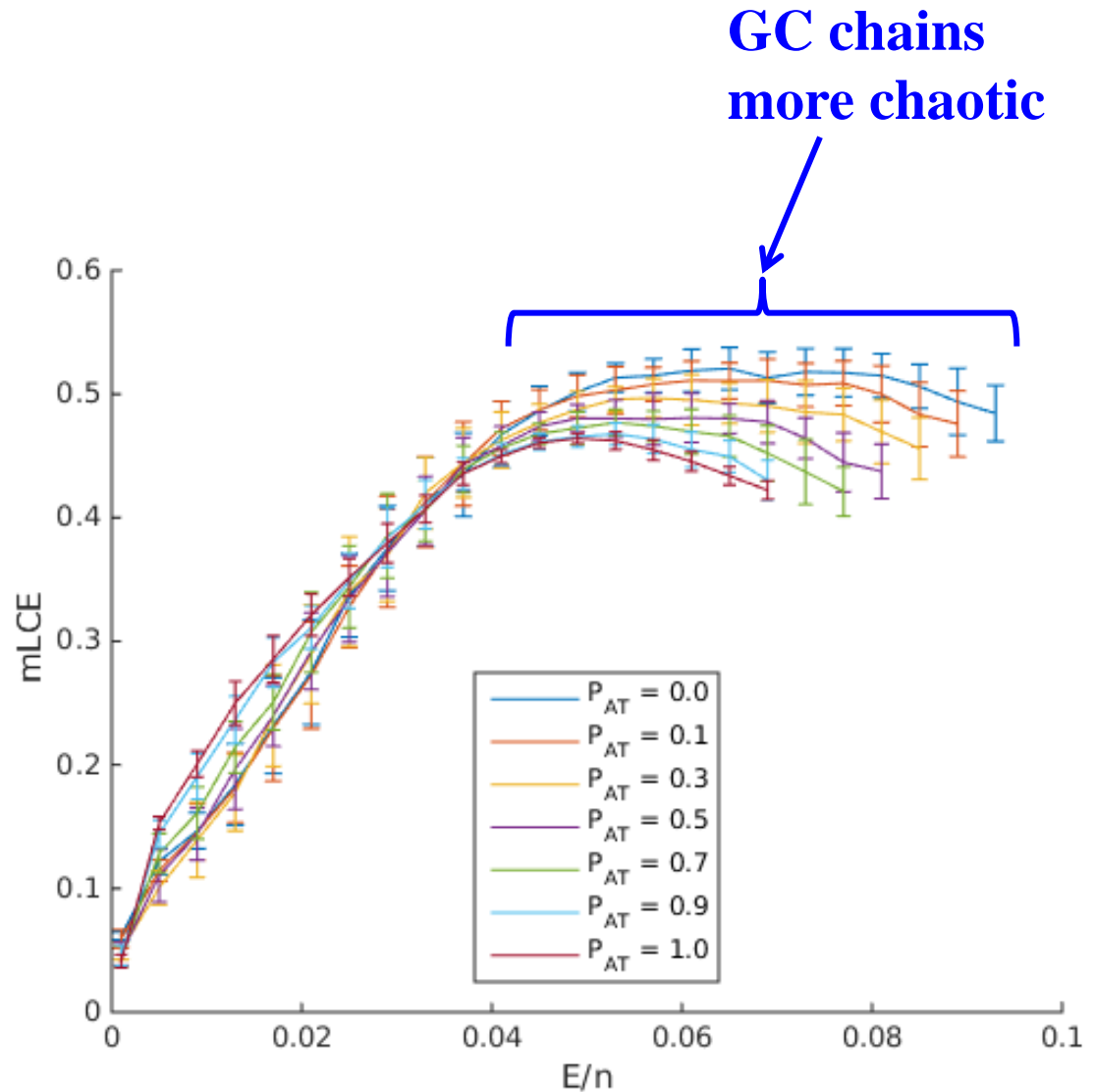
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Lyapunov exponent vs. energy per particle



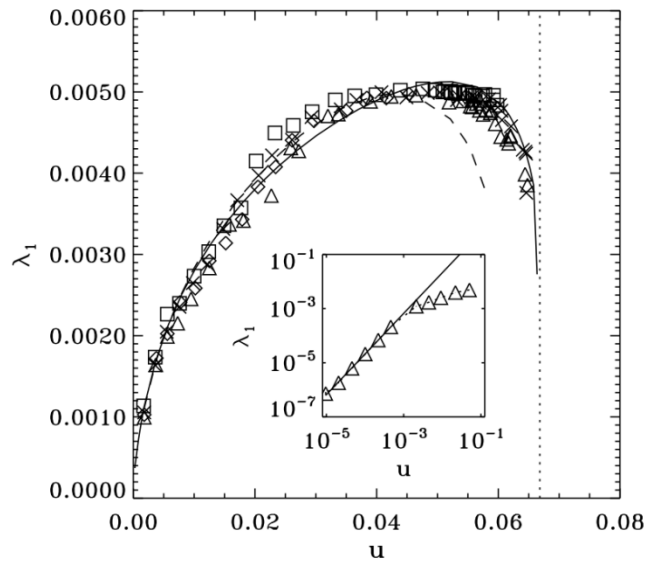
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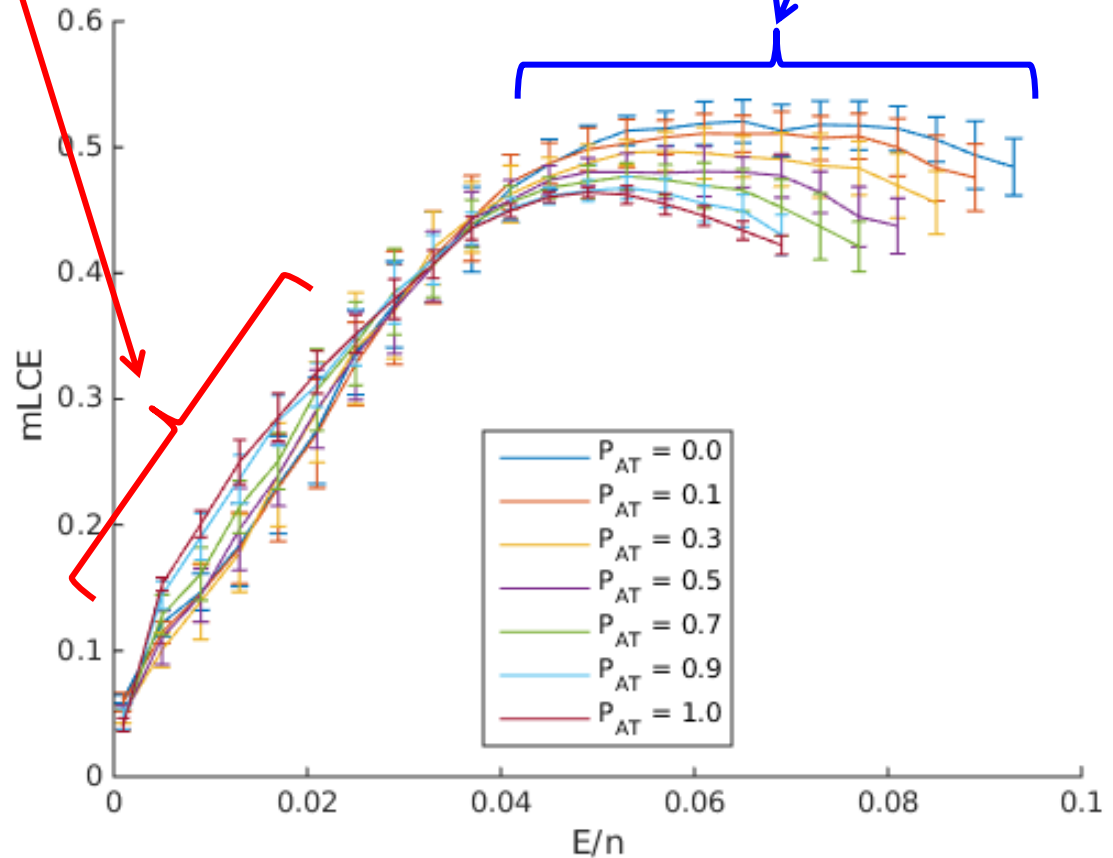
Lyapunov exponent vs. energy per particle

**AT chains
more chaotic**

**GC chains
more chaotic**



**Homogeneous chain
[Barré & Dauxois,
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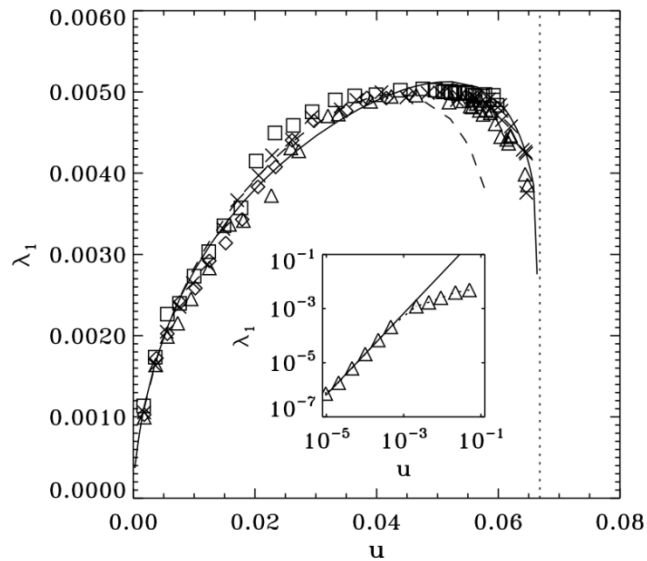


Lyapunov exponent vs. energy per particle

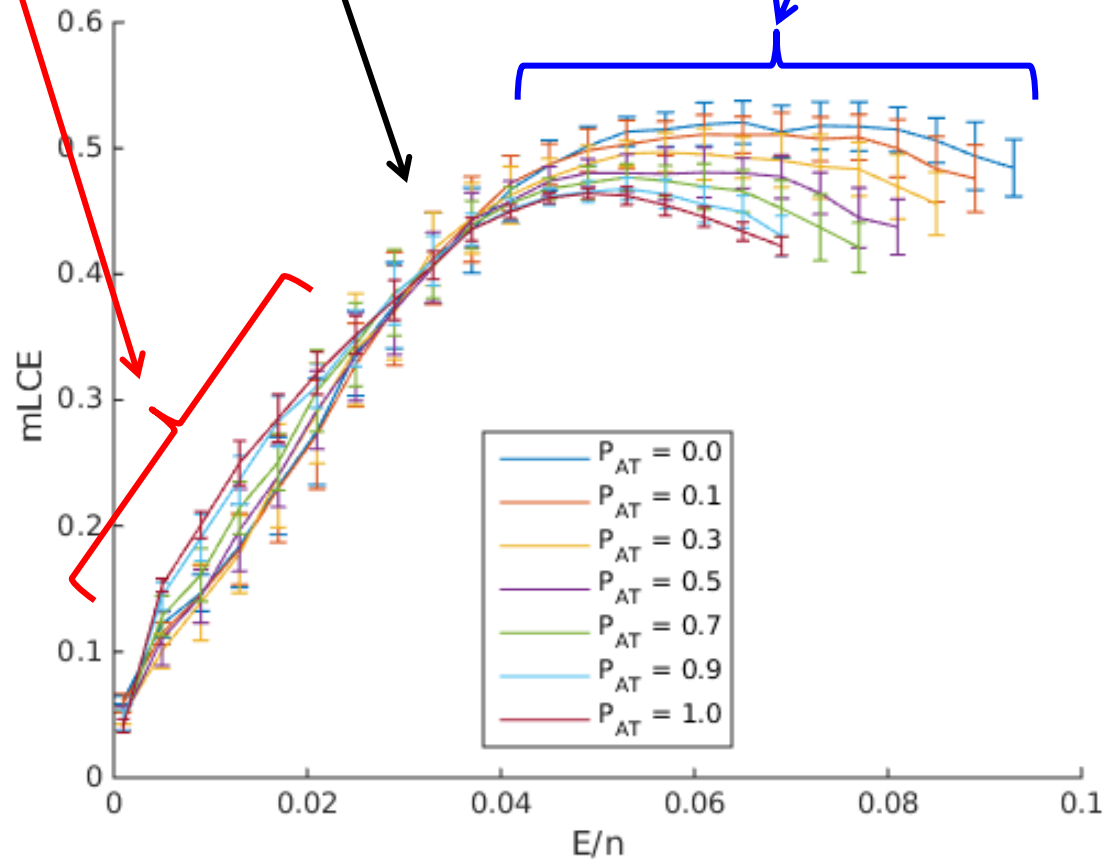
**AT chains
more chaotic**

**Type of chain
does not play
a role**

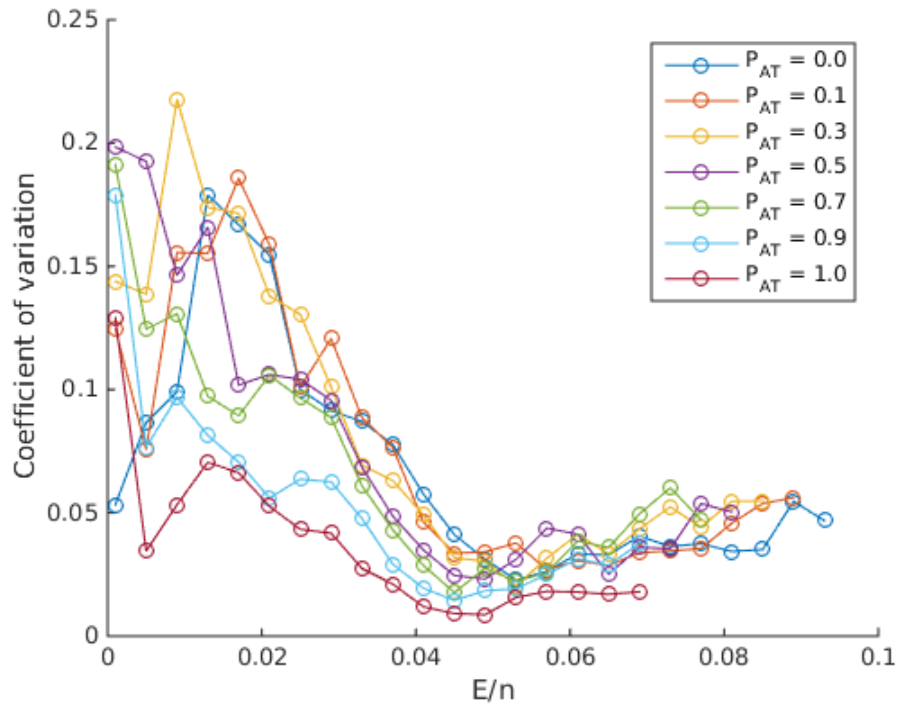
**GC chains
more chaotic**



**Homogeneous chain
[Barré & Dauxois,
EPL (2001)]**

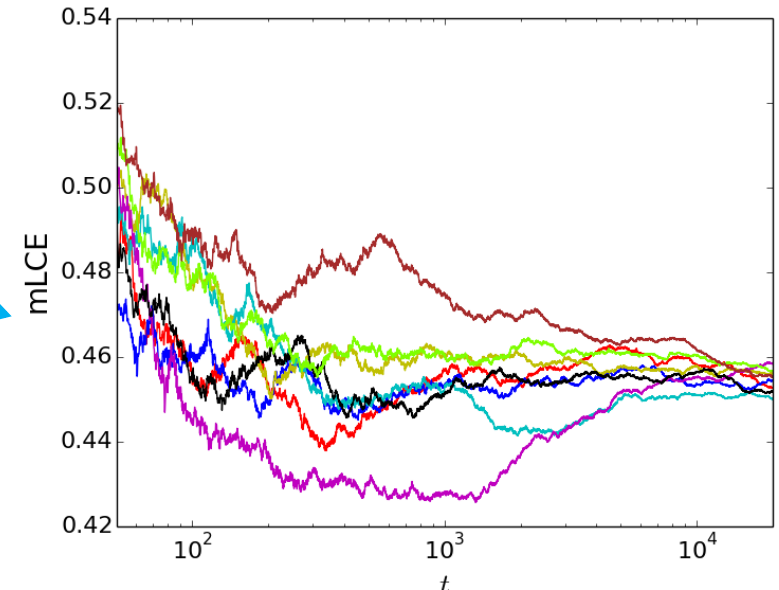
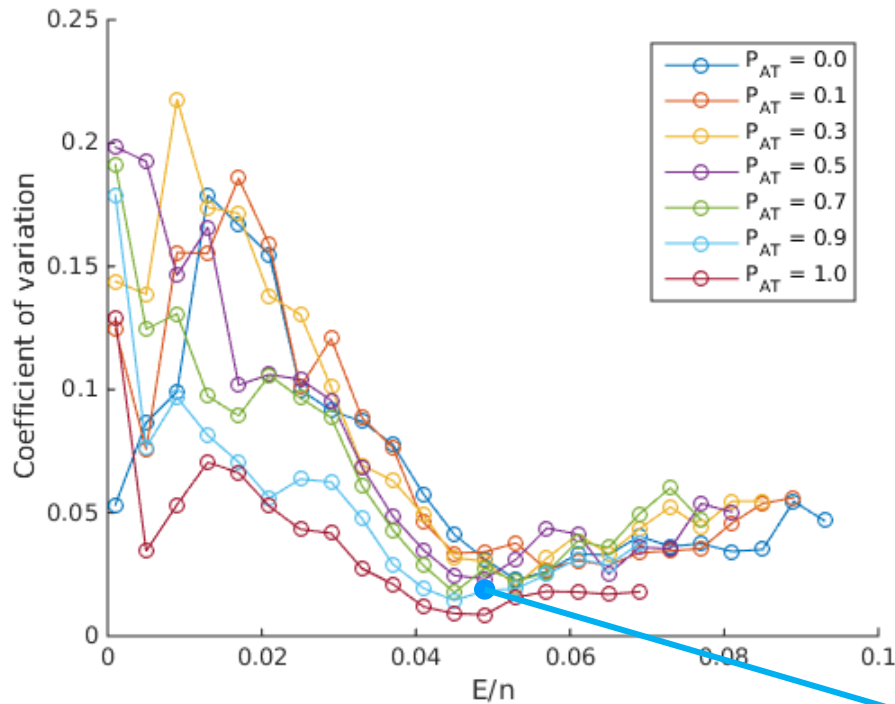


(Error of mLCE)/mLCE



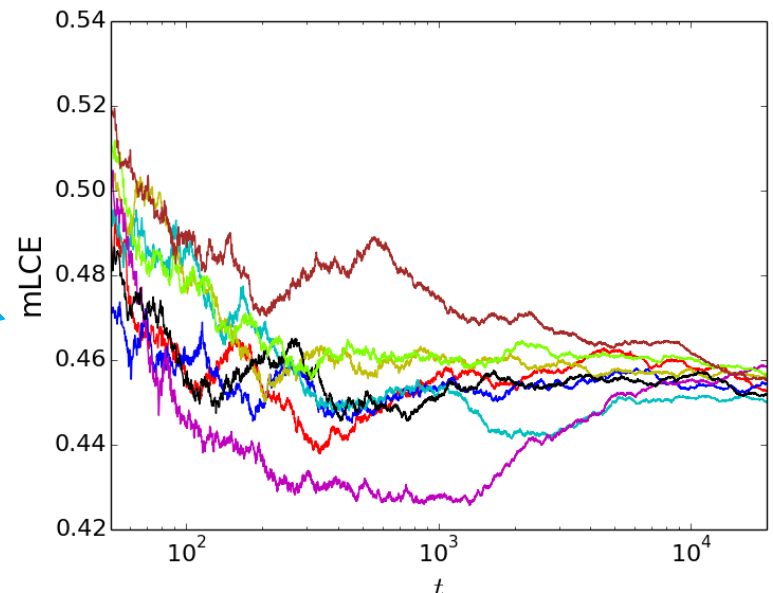
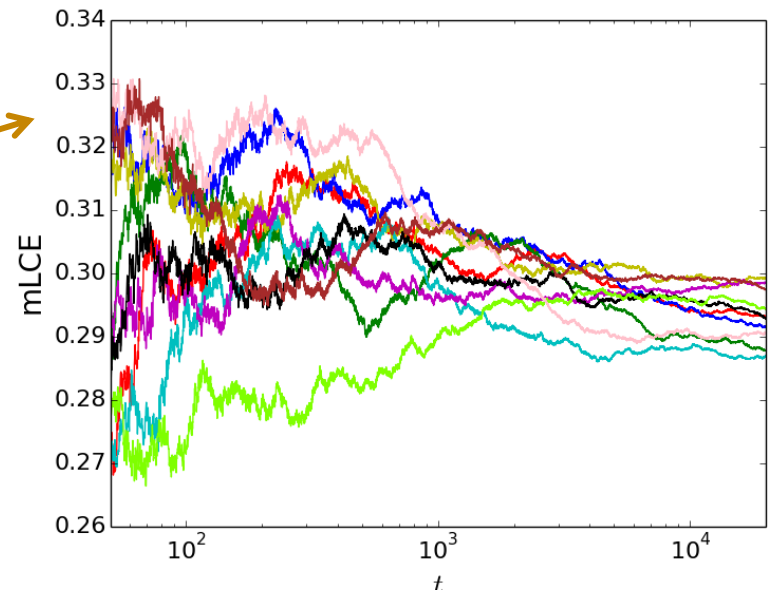
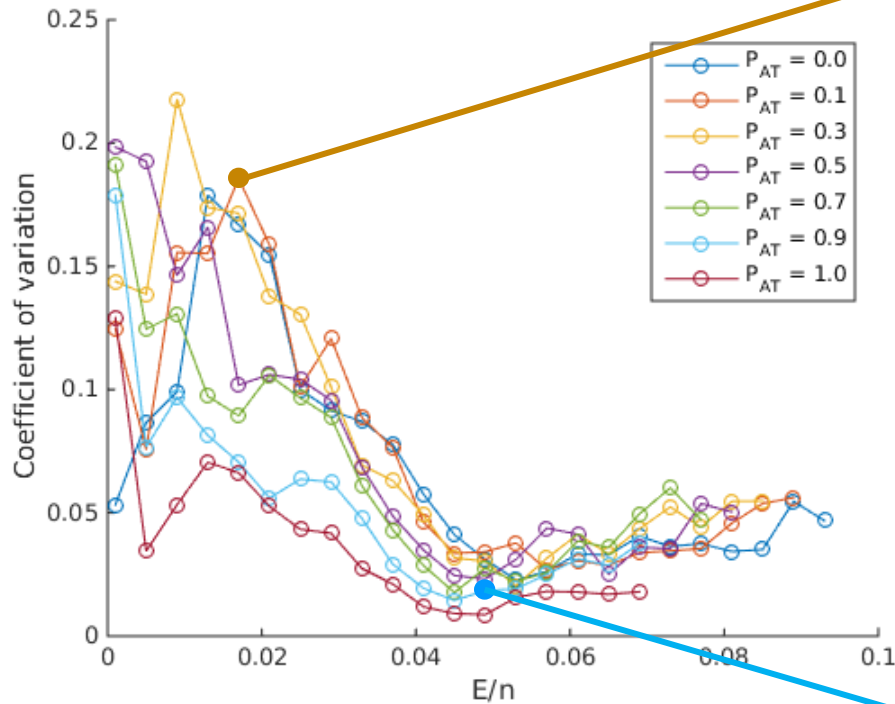
Values of Lyapunov exponents

(Error of mLCE)/mLCE



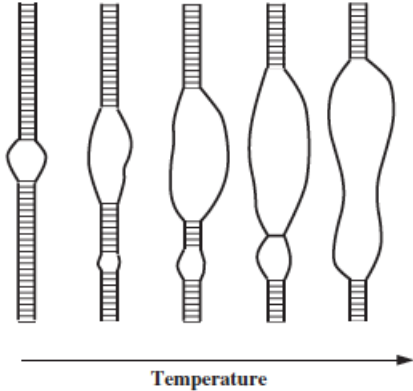
Values of Lyapunov exponents

(Error of mLCE)/mLCE



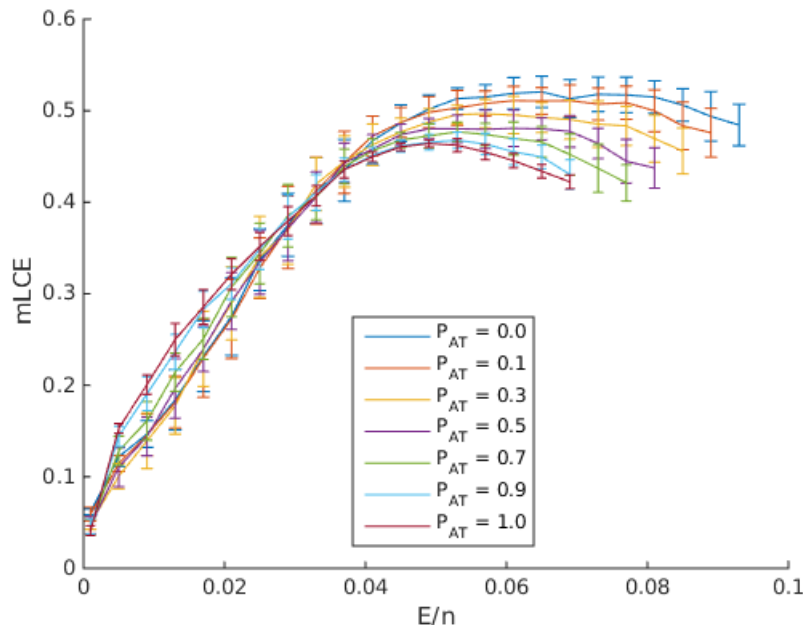
DNA denaturation (melting)

Melting: large **bubbles forming** in the DNA chain as bonds break



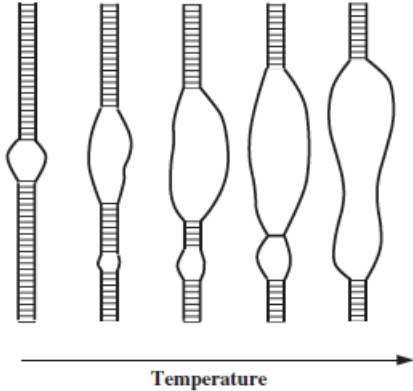
As y_n increases the exponentials in

$$D_n(e^{-a_n y_n} - 1)^2 + \frac{K}{2}(1 + \rho e^{-b(y_n + y_{n-1})})(y_n - y_{n-1})^2$$
 tend to 0, the system becomes effectively linear and the mLCE $\rightarrow 0$.



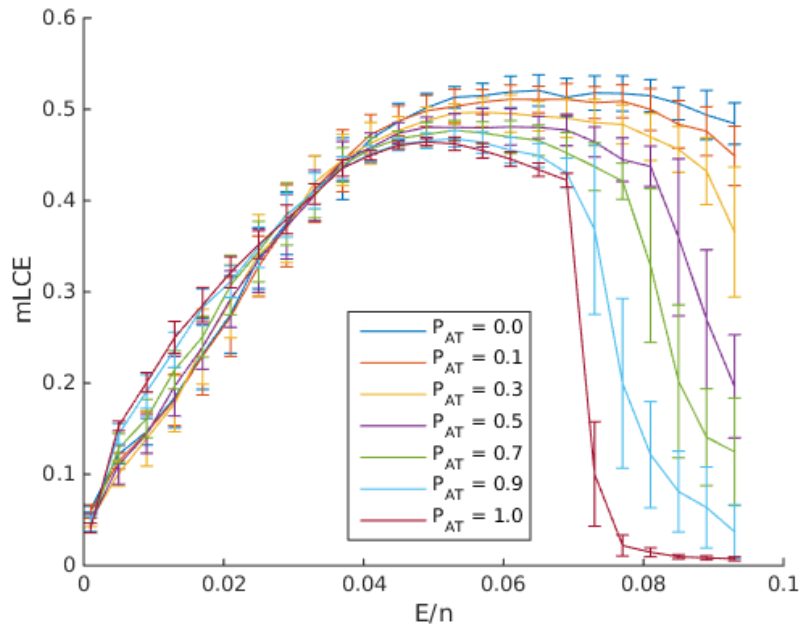
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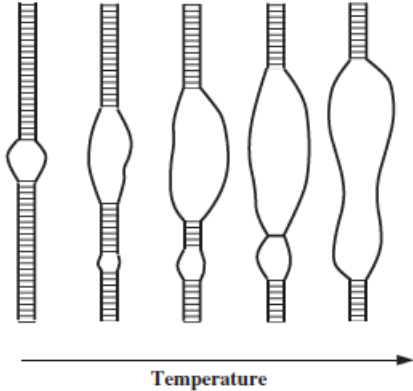
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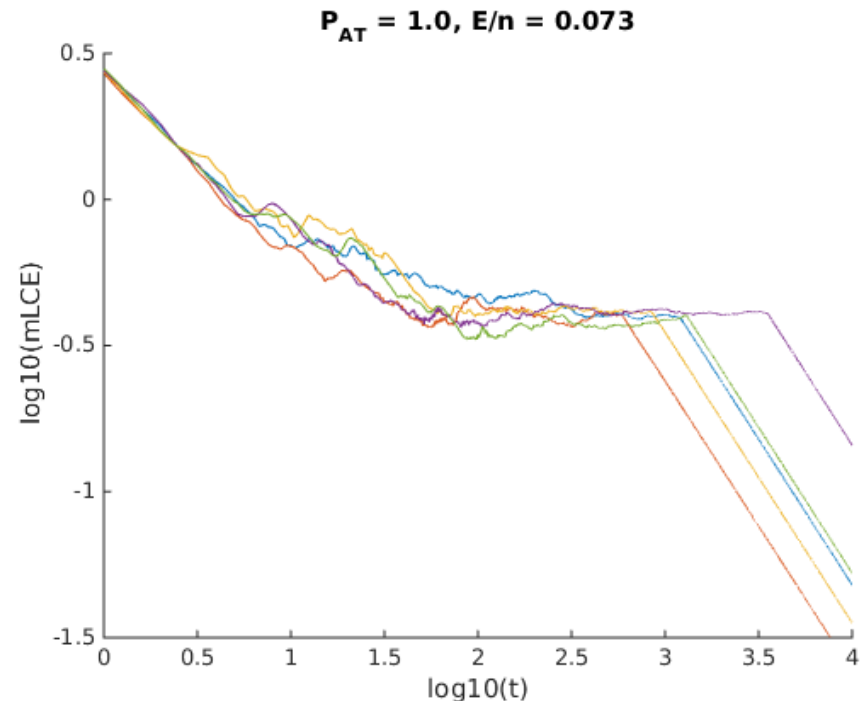
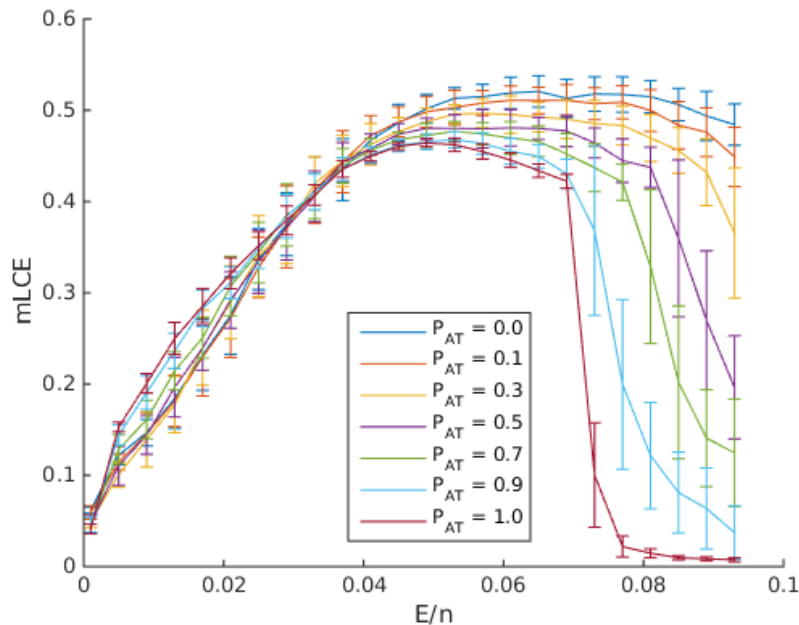
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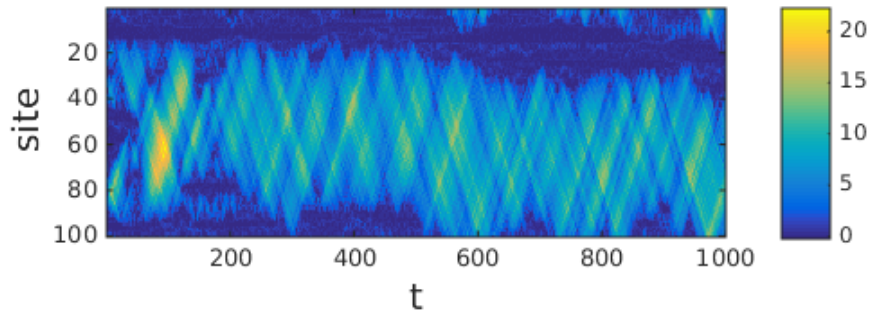
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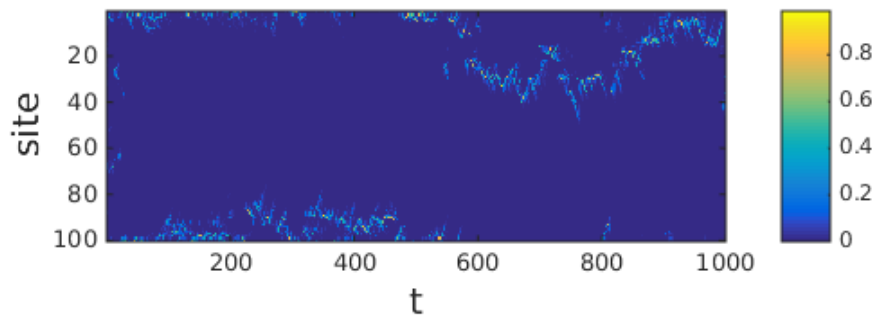


DVD and the formation of bubbles

$E/n = 0.071$ - displacement

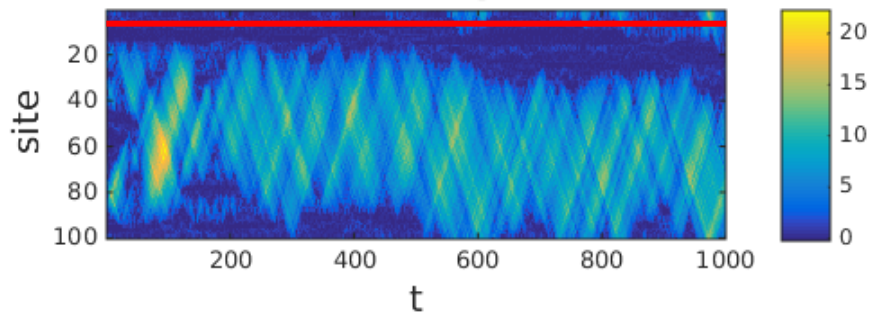


$E/n = 0.071$ - DVD

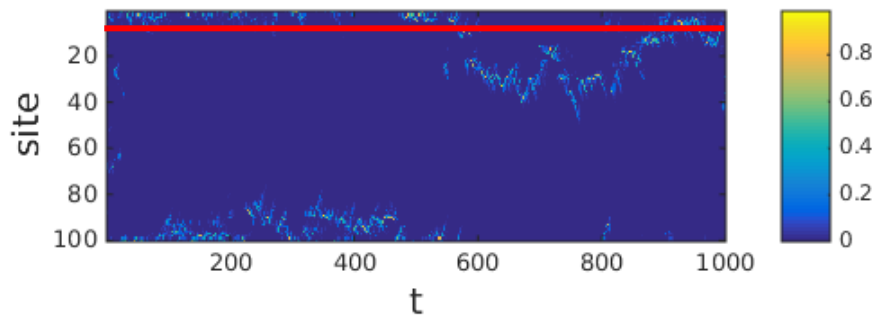


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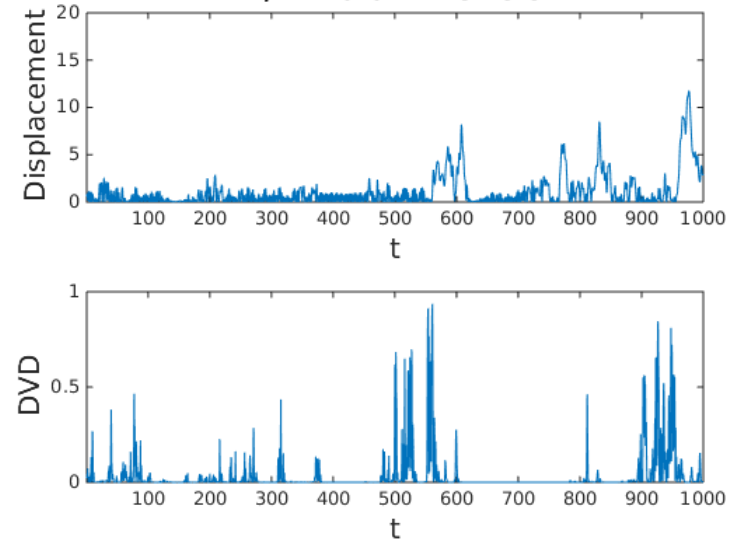
$E/n = 0.071$ - displacement



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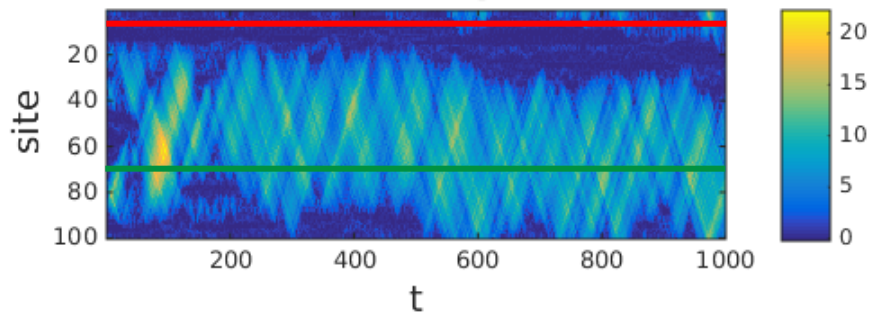


$E/n = 0.071$ - site 5

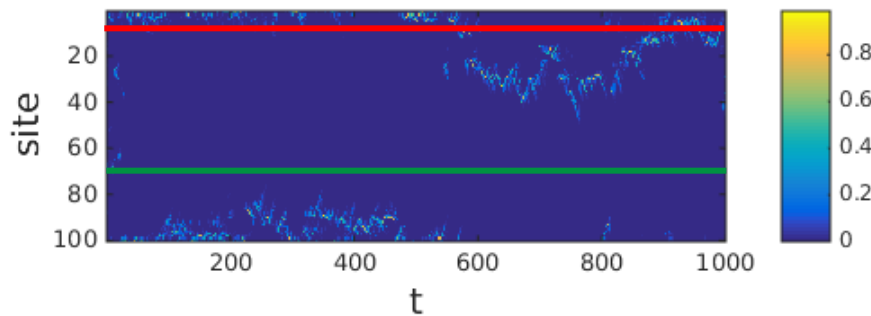


DVD and the formation of bubbles

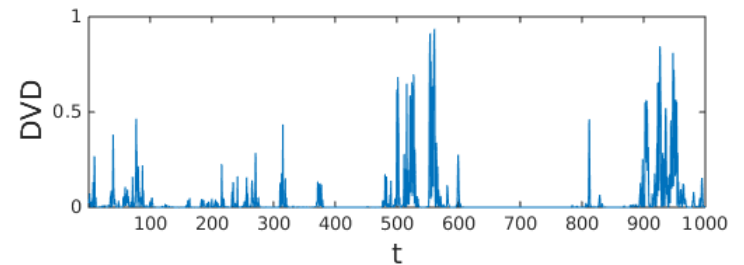
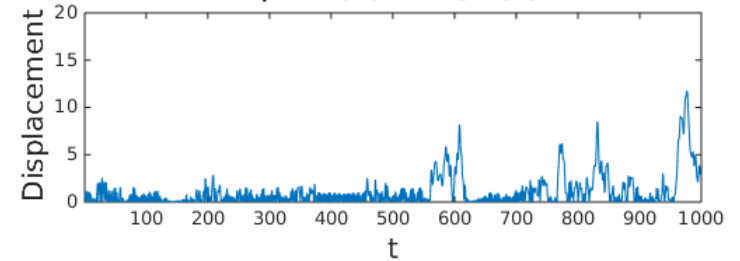
$E/n = 0.071$ - displacement



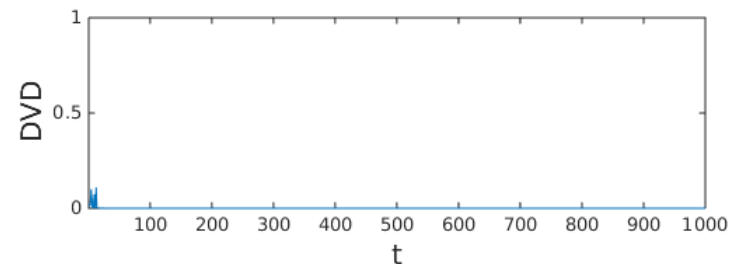
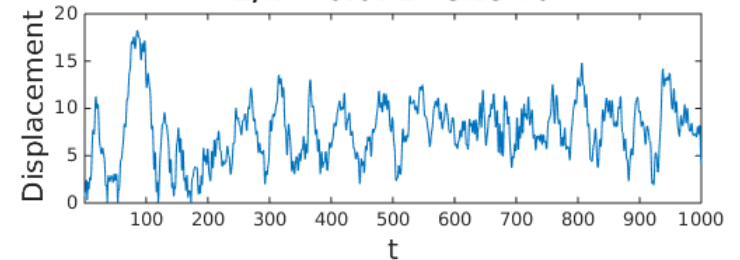
$E/n = 0.071$ - DVD



$E/n = 0.071$ - site 5



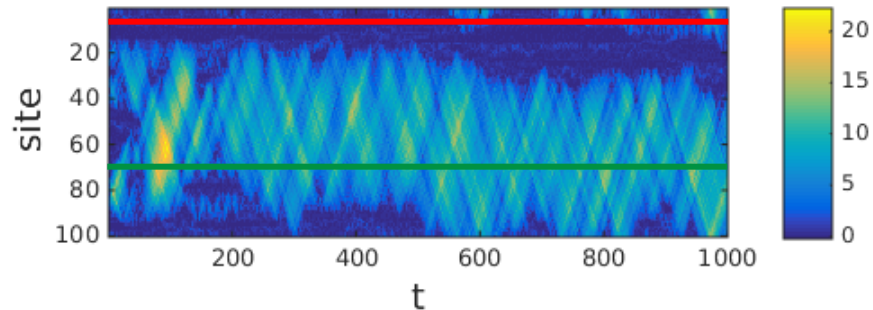
$E/n = 0.071$ - site 70



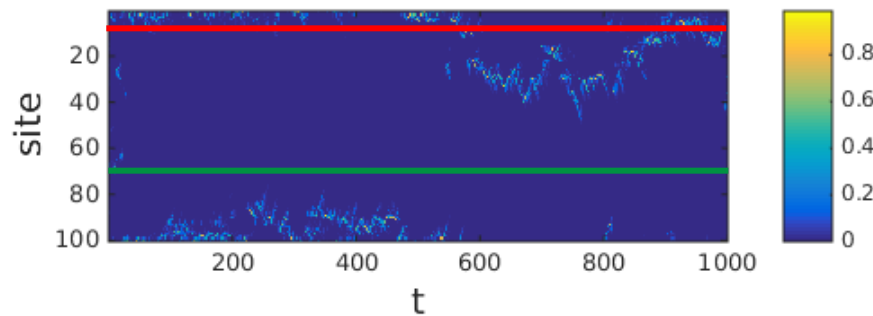
DVD and the formation of bubbles

Relation between the concentration of the deviation vector at a site and the formation of a bubble at that site.

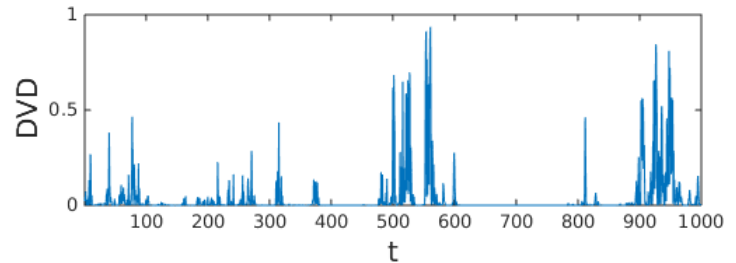
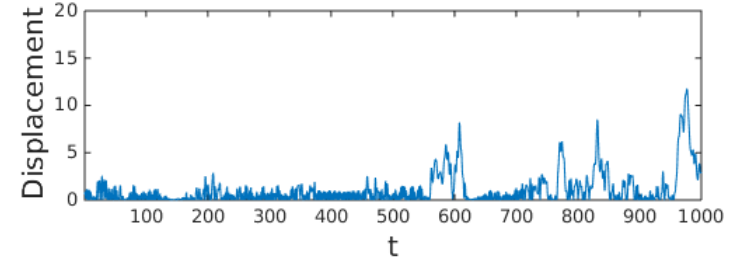
$E/n = 0.071$ - displacement



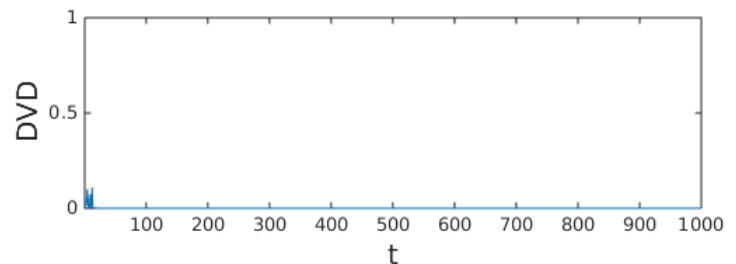
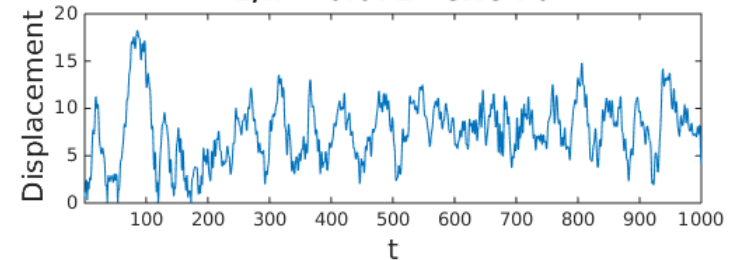
$E/n = 0.071$ - DVD



$E/n = 0.071$ - site 5

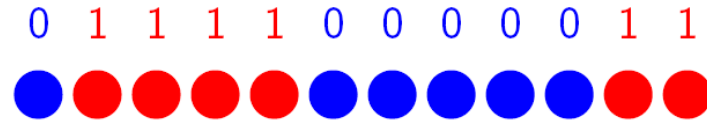


$E/n = 0.071$ - site 70



Mixing of the DNA chain

Mixing parameter α = Number of alternations in the chain (AT and GC).



$\alpha=4$

(1) $\bar{0}$ 1 1 1 $\bar{1}$ 0 0 0 0 $\bar{0}$ 1 $\bar{1}$ (0)

Mixing of the DNA chain

Mixing parameter α = Number of alternations in the chain (**AT** and **GC**).

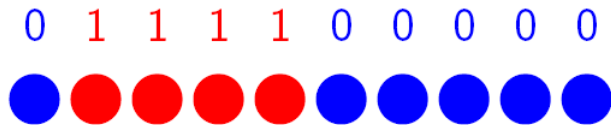
$\alpha=4$

0	1	1	1	1	0	0	0	0	0	1	1
●	●	●	●	●	●	●	●	●	●	●	●

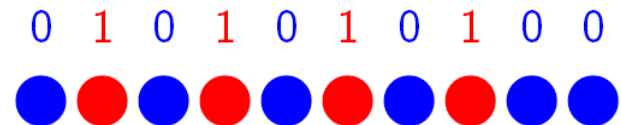
$$(1)\bar{0}111\bar{1}0000\bar{0}1\bar{1}(0)$$

Example case: $N=10$, $N_{AT}=4$, $N_{GC}=6$.

Extreme cases: $\alpha=2$ and $\alpha=8$



$$\alpha = 2$$



$$\alpha = 8$$

Mixing of the DNA chain

Mixing parameter α = Number of alternations in the chain (**AT** and **GC**).

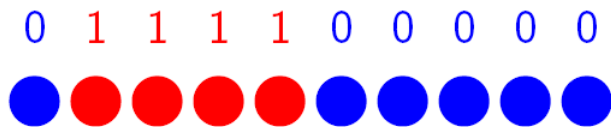
$\alpha=4$

$$\begin{array}{cccccccccccc} 0 & 1 & 1 & 1 & 1 & 0 & 0 & 0 & 0 & 0 & 1 & 1 \\ \bullet & \bullet & \bullet & \bullet & \bullet & \bullet & \bullet & \bullet & \bullet & \bullet & \bullet & \bullet \end{array}$$

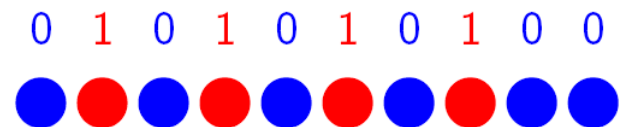
$$(1)\bar{0}111\bar{1}0000\bar{0}1\bar{1}(0)$$

Example case: $N=10$, $N_{AT}=4$, $N_{GC}=6$.

Extreme cases: $\alpha=2$ and $\alpha=8$



$$\alpha = 2$$

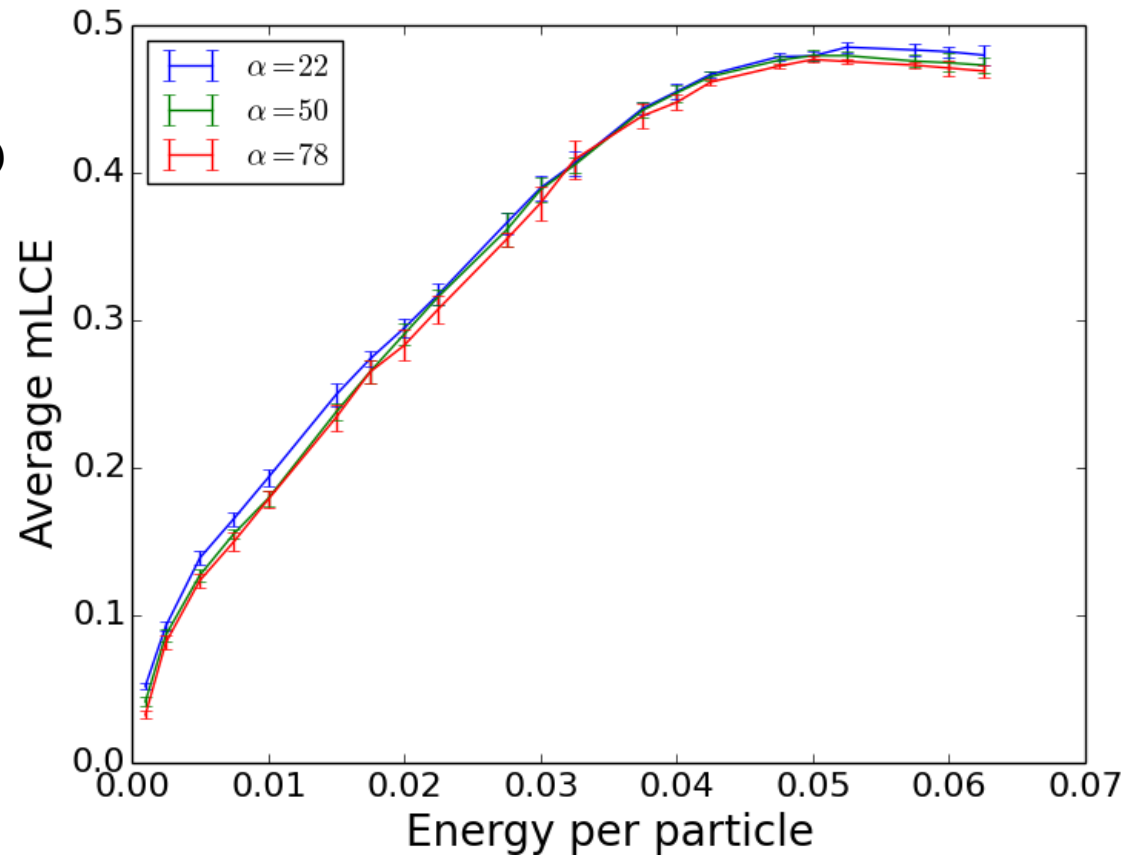
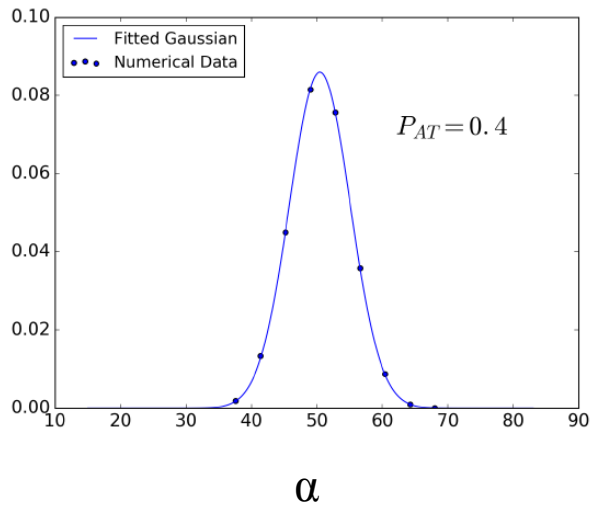


$$\alpha = 8$$

$$2 \leq \alpha \leq \min\{2N_{AT}, 2N_{GC}\}, \quad \alpha \text{ even}$$

Effect of mixing

Probability distribution function $P(\alpha)$



The more heterogeneous chains are slightly less chaotic

Summary

- **Granular chain model**

- ✓ **Moderate nonlinearities:** although the overall system behaves chaotically, it can exhibit **long lasting energy localization for particular single particle excitations.**
- ✓ **Sufficiently strong nonlinearities:** the granular chain reaches **energy equipartition and an equilibrium chaotic state,** independent of the initial position excitation.

- **DNA model**

- ✓ **Heterogeneity influences the behavior of the mLE and the system's chaotic behavior.**
- ✓ **There seems to be a relation between the concentration of the DVD at a site and the formation of a bubble.**
- ✓ **Mixing does not influence significantly the system's chaoticity.**
- ✓ **The behavior of DVDs can provide important information about the chaotic behavior of a dynamical system.**