

Population dynamics and rare events

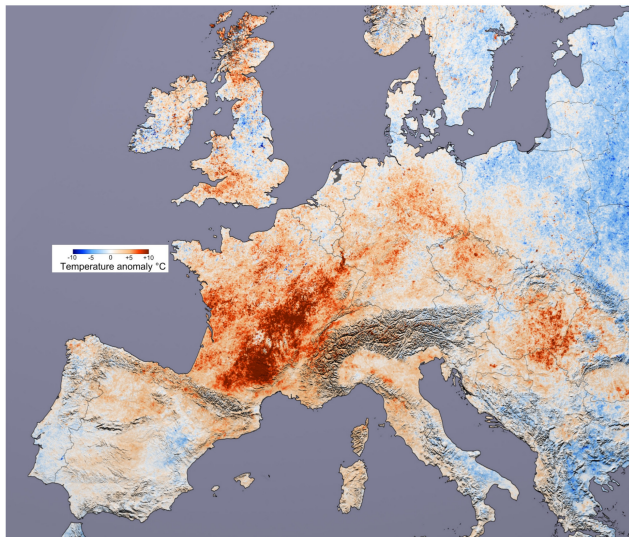
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Freddy Bouchet⁽³⁾, Rob L Jack⁽⁴⁾, Vivien Lecomte⁽⁵⁾

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(4)Bath University & Cambridge University (5) LIPhy, Grenoble

University of Oldenburg — January 24, 2019

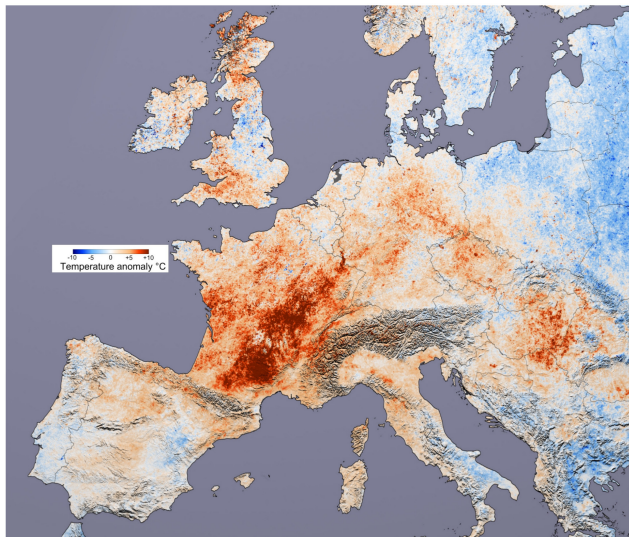


Why studying rare events?



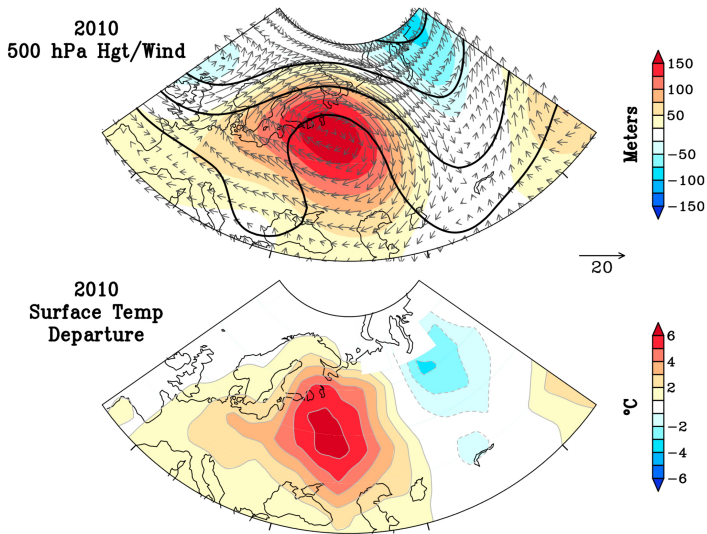
2003 heat wave, Europe [Terra MODIS]

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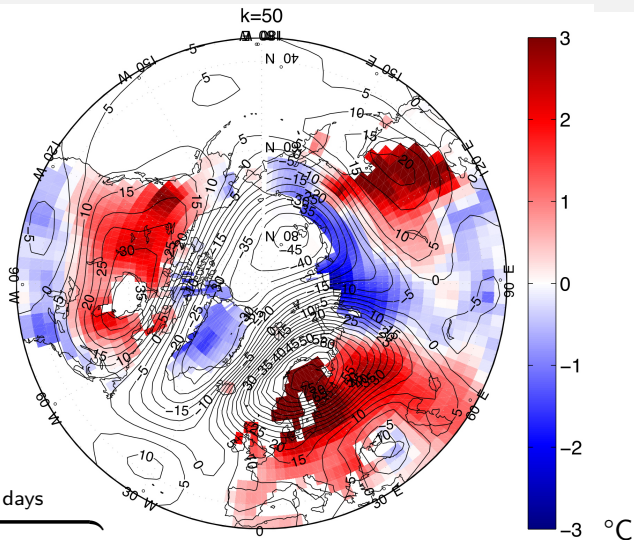
[Anomaly for **1-month** average] 2003 heat wave, Europe [Terra MODIS]

Why studying rare events?



2010 heat wave in Western Russia [Dole *et al.*, 2011]

Why studying rare events?



$$\frac{1}{t_{\max}} \int_0^{t_{\max}} dt \Delta T(t) > 2^{\circ}\text{C} \Rightarrow \text{« Teleconnection patterns » [Bouchet et al.]}$$

$t_{\max} = 40 \text{ days}$

How to study rare events?

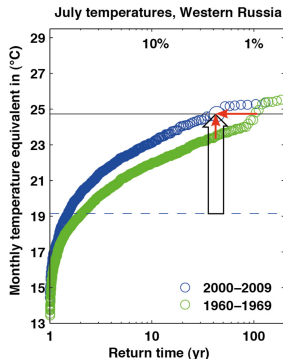
Questions for physicists and mathematicians:

- Probability and **dynamics** of rare events?
- How to **sample** these in numerical modelisation?
- Numerical **tools** and methods to understand their **formation**?

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Evolution of the return time
of the monthly averaged temperature

$$\frac{1}{t_{\max}} \int_0^{t_{\max}} dt T(t)$$

↔ anthropogenic impact on climate?

[Otto *et al.*, 2012]

Outline

- **Introduction**
- **Tools and algorithm:**
 - Large deviation functions
 - Ingredient 1/2: population dynamics
 - Ingredient 2/2: change of ensemble
- **Use, extensions and limitations of population dynamics:**
 - Different averages
 - Feedback method
 - Finite-time and finite-population scalings
- **Open questions**

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Time-extensive observable on a time window $[0, t_f]$

- Climate dynamics:

$$\int_0^{t_f} dt \text{ temperature}(t)$$

- Fluctuating thermodynamics:

$$\text{work} = \int_0^{t_f} dt \text{ force}(t) \cdot \text{velocity}(t)$$

- Road traffic:

$$\#\{\text{cars passing through a gate}\}$$

- Molecular transport:

$$\#\{\text{steps of a motor on a filament}\}$$

- Lattice gases in 1d:

$$\text{“current”} = \#\{\text{jumps to the right}\} - \#\{\text{jumps to the left}\}$$

$$\text{“activity”} = \#\{\text{jumps to the right}\} + \#\{\text{jumps to the left}\}$$

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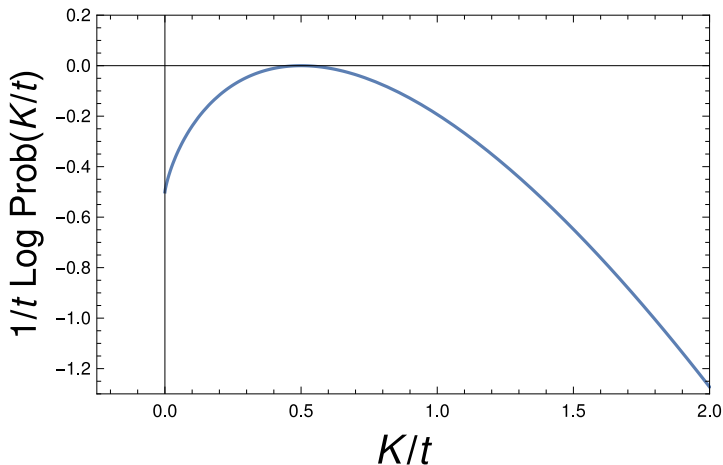
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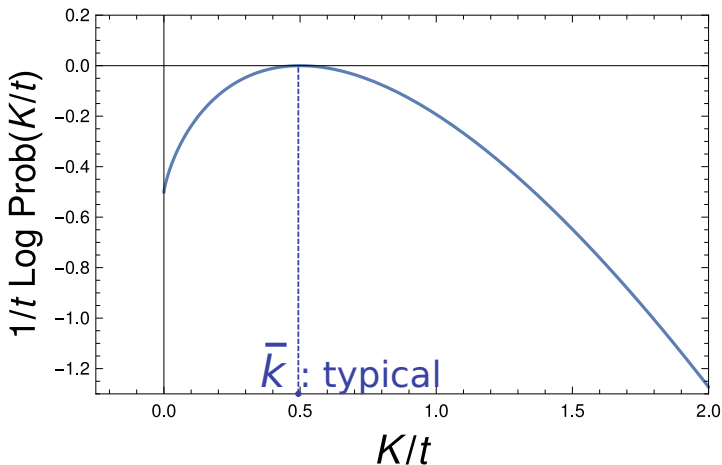
Distribution of a time-extensive observable K on $[0, t]$



$$\text{Prob}[K, t] \sim e^{t\varphi(K/t)} \text{ as } t \rightarrow \infty$$

$\varphi(k) =$ large deviation function

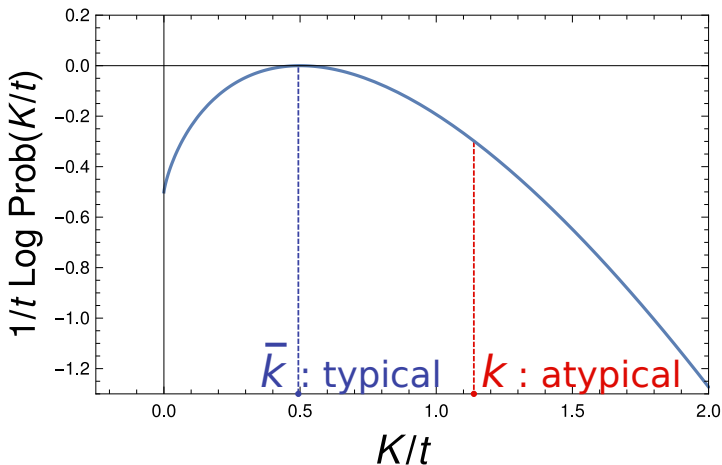
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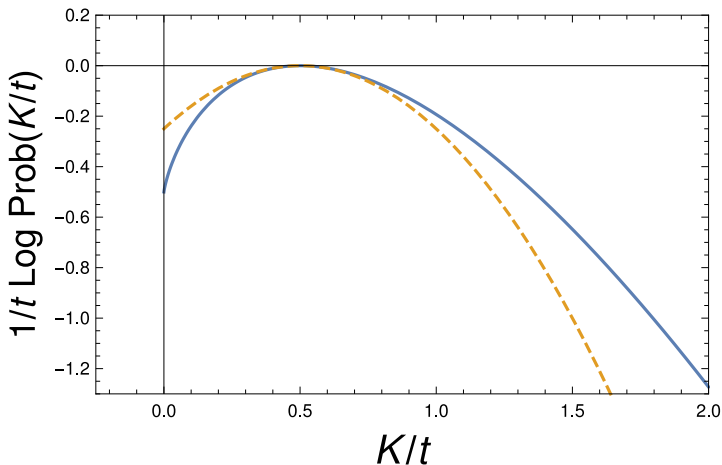
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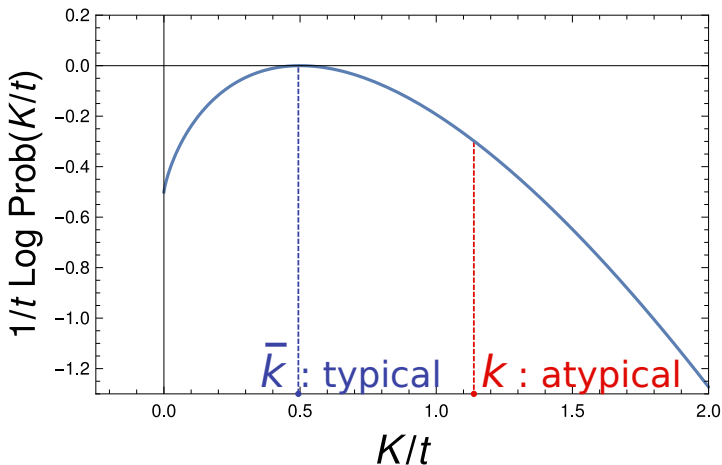
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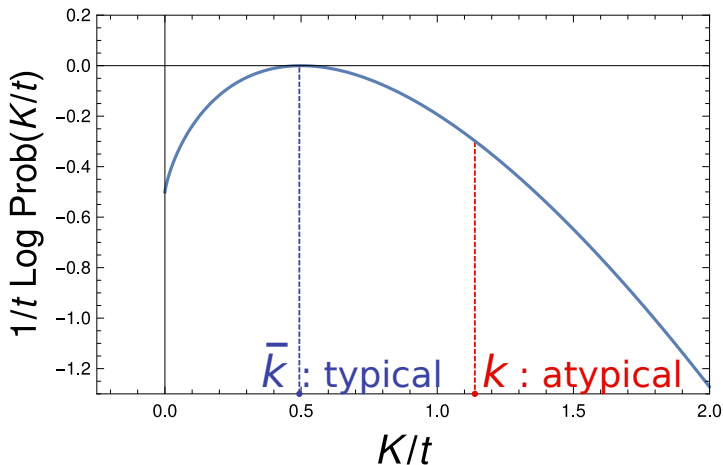
$\text{Prob}[K, t] \sim e^{t\varphi(K/t)}$ as $t \rightarrow \infty$ $\varphi(k) =$ large deviation function
 quadratic approx. $\varphi(k) = \frac{(k-\bar{k})^2}{2\sigma^2} + \dots \leftrightarrow$ Gaussian fluctuations

Aim: modify dynamics to make atypical values k typical



Original dynamics: \bar{k} is typical and k atypical

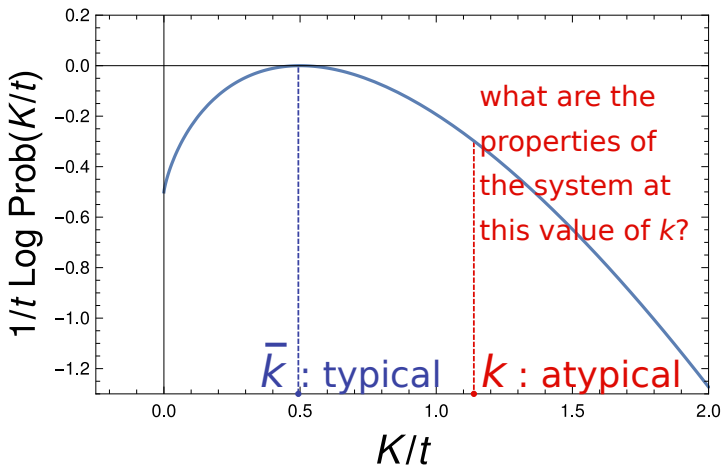
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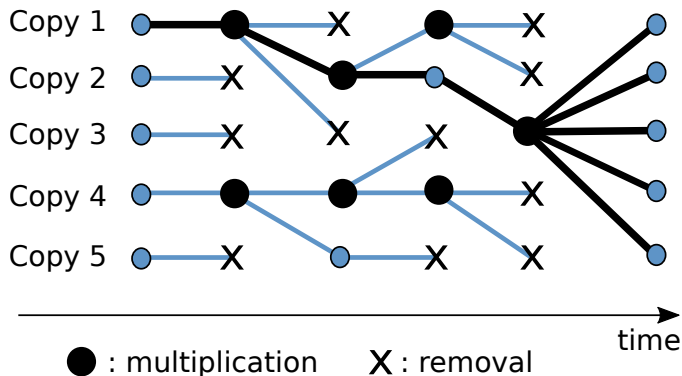
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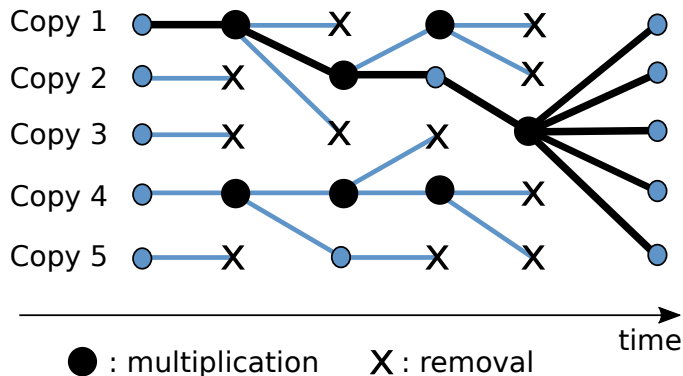
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Ingredient 1/2: population dynamics



Many copies of the system of interest evolve in parallel.

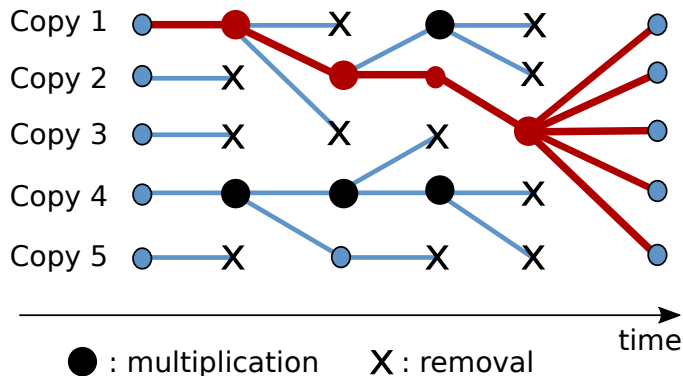
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Selection rules favor the normally **atypical value of k**

Ingredient 1/2: population dynamics



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Selection rules favor the normally **atypical value of k**

→ **typical population trajectories** sample the original system at **atypical k**

Ingredient 2/2: dynamical change of ensemble @ $t \rightarrow \infty$

Consider an observable \mathcal{O} [trajectory].

$$\underbrace{\frac{\langle \mathcal{O}[\text{traj.}] \delta\left(\frac{1}{t}K[\text{traj.}] - k\right) \rangle}{\langle \delta\left(\frac{1}{t}K[\text{traj.}] - k\right) \rangle}}_{\text{average of } \mathcal{O} \text{ for trajectories with atypical } k = K/t}$$

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Consider an observable \mathcal{O} [trajectory]. One can show:

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For s and k suitably “conjugated”.

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Next goal: show that $\dots = \underbrace{\langle \mathcal{O}[\text{traj.}] \rangle}_{\text{population dynamics}}$
 average of \mathcal{O} for trajectories in fixed-size population dynamics

For s and k suitably “conjugated”.

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For s and k suitably “conjugated”.

Analogy: $k \equiv \text{energy/volume}$; $s \equiv \text{inverse temperature } \beta$

Ingredient 2/2: dynamical change of ensemble @ $t \rightarrow \infty$

Main message:

Fixed $k = K/t \iff$ bias by e^{-sK}

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Fixed $k = K/t$ \iff bias by e^{-sK} \iff Selection rules in population dynamics

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Relation between s and k ; Cumulant Generating Function (CGF):

$$\text{Prob}[K/t = k] \sim e^{t\varphi(k)} \iff \langle e^{-sK} \rangle \sim e^{t \overbrace{\psi(s)}^{\text{CGF}}}$$

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Saddle-point at large t :

$$\max_k \{ \varphi(k) - sk \} = \psi(s)$$

Maximum reached for k conjugated to s

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Remaining question: how to represent e^{-sK} by pop. dynamics?

s-modified dynamics (for discrete stochastic processes)

- Markov processes: Configs. \mathcal{C} , jump rates $W(\mathcal{C} \rightarrow \mathcal{C}')$

$$\partial_t P(\mathcal{C}, t) = \sum_{\mathcal{C}'} \left\{ \underbrace{W(\mathcal{C}' \rightarrow \mathcal{C}) P(\mathcal{C}', t)}_{\text{gain term}} - \underbrace{W(\mathcal{C} \rightarrow \mathcal{C}') P(\mathcal{C}, t)}_{\text{loss term}} \right\}$$

s-modified dynamics

 $K = \text{activity} = \# \text{events}$

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- More detailed dynamics for $P(\mathcal{C}, K, t)$:

$$\partial_t P(\mathcal{C}, K, t) = \sum_{\mathcal{C}'} \left\{ W(\mathcal{C}' \rightarrow \mathcal{C}) P(\mathcal{C}', K-1, t) - W(\mathcal{C} \rightarrow \mathcal{C}') P(\mathcal{C}, K, t) \right\}$$

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- Biased ensemble: s conjugated to K (canonical description)

$$\hat{P}(\mathcal{C}, s, t) = \sum_K e^{-sK} P(\mathcal{C}, K, t)$$

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s-modified dynamics

$$K = k_{c_0 c_1} + k_{c_1 c_2} + \dots$$

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Numerical method [JB Anderson; D Aldous; P Grassberger; P Del Moral; ...]

Evaluation of large deviation functions

[à la “Diffusion Monte-Carlo”]

$$\sum_{\mathcal{C}} \hat{P}(\mathcal{C}, \mathbf{s}, t) = \langle e^{-\mathbf{s}K} \rangle \sim e^{t\psi(\mathbf{s})} \quad (\psi(\mathbf{s}) = \text{CGF} = \text{max eigenv. } \mathbb{W}_{\mathbf{s}})$$

- discrete time: Giardinà, Kurchan, Peliti [PRL **96**, 120603 (2006)]
- continuous time: VL, Tailleur [JSTAT P03004 (2007)]

Cloning dynamics

$$\partial_t \hat{P}(\mathcal{C}, \mathbf{s}) = \underbrace{\sum_{\mathcal{C}'} W_{\mathbf{s}}(\mathcal{C}' \rightarrow \mathcal{C}) \hat{P}(\mathcal{C}', \mathbf{s}) - r_{\mathbf{s}}(\mathcal{C}) \hat{P}(\mathcal{C}, \mathbf{s})}_{\text{modified dynamics}} + \underbrace{\delta r_{\mathbf{s}}(\mathcal{C}) \hat{P}(\mathcal{C}, \mathbf{s})}_{\text{cloning term}}$$

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Explicit construction

$$\partial_t \hat{P}(\mathcal{C}, s) = \underbrace{\sum_{\mathcal{C}'} W_s(\mathcal{C}' \rightarrow \mathcal{C}) \hat{P}(\mathcal{C}', s) - r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{modified dynamics}} + \underbrace{\delta r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{cloning term}}$$

How to take into account loss/gain of probability?

- handle a large number N_c of copies of the system
- implement a **selection** rule: on a time interval Δt a copy in config \mathcal{C} is replaced by $Y = e^{\Delta t \delta r_s(\mathcal{C})}$ copies
- $\psi(s)$ = the rate of exponential growth/decay of the total population
- optionally: keep population constant by non-biased pruning/cloning

Explicit construction

$$\partial_t \hat{P}(\mathcal{C}, s) = \underbrace{\sum_{\mathcal{C}'} W_s(\mathcal{C}' \rightarrow \mathcal{C}) \hat{P}(\mathcal{C}', s) - r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{modified dynamics}} + \underbrace{\delta r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{cloning term}}$$

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- implement a **selection** rule: on a time interval Δt a copy in config \mathcal{C} is replaced by $Y = \lfloor e^{\Delta t \delta r_s(\mathcal{C})} + \varepsilon \rfloor$ copies, $\varepsilon \sim [0, 1]$
- $\psi(s)$ = the rate of exponential growth/decay of the total population
- optionally: keep population constant by non-biased pruning/cloning

Explicit construction

$$\partial_t \hat{P}(\mathcal{C}, s) = \underbrace{\sum_{\mathcal{C}'} W_s(\mathcal{C}' \rightarrow \mathcal{C}) \hat{P}(\mathcal{C}', s) - r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{modified dynamics}} + \underbrace{\delta r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{cloning term}}$$

How to take into account loss/gain of probability?

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CGF estimator: $\psi(s) = \langle \Psi(s) \rangle$ with $\Psi(s) = \log \underbrace{\prod_t \frac{N_c + Y_t - 1}{N_c}}_{\text{reconstituted population size}}$

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Biological interpretation

- copy in configuration $\mathcal{C} \equiv$ organism of **genome** \mathcal{C}
- dynamics of rates $W_s \equiv$ **mutations**
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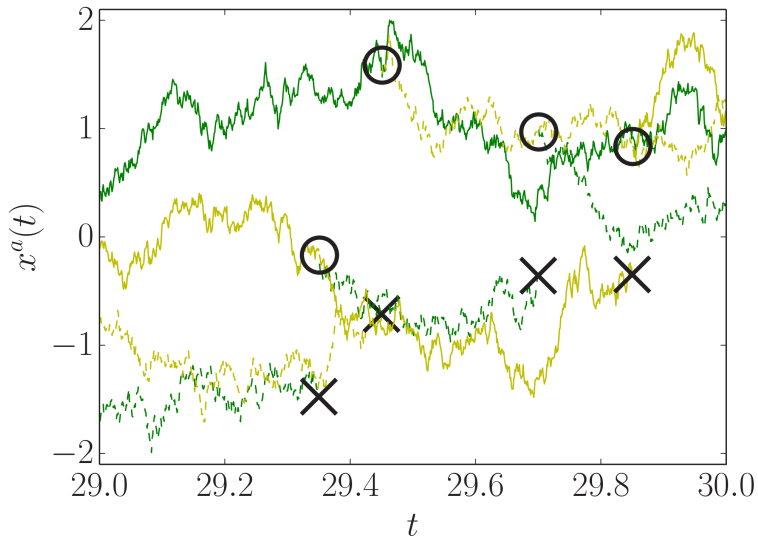
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Generic idea

- **Different dynamics can share equivalent statistical properties.**
- Constrained trajectories (fixed atypical $k = K/t$) \equiv pop. dynamics

An example: 4 copies, 1 degree of freedom $\mathcal{C} = x \in \mathbb{R}$



How to perform averages? (i)

[spectral analysis]

- ★ Final-time distribution $p_{\text{end}}(\mathcal{C})$: *proportion of copies in \mathcal{C} at t*

$$\langle N_{\text{nc}}(t) \rangle_s$$

$$\langle N_{\text{nc}}(\mathcal{C}, t) \rangle_s$$

$$p_{\text{end}}(\mathcal{C}, t) = \frac{\langle N_{\text{nc}}(\mathcal{C}, t) \rangle_s}{\langle N_{\text{nc}}(t) \rangle_s}$$

[N_{nc} = number of copies in non-constant population dynamics]

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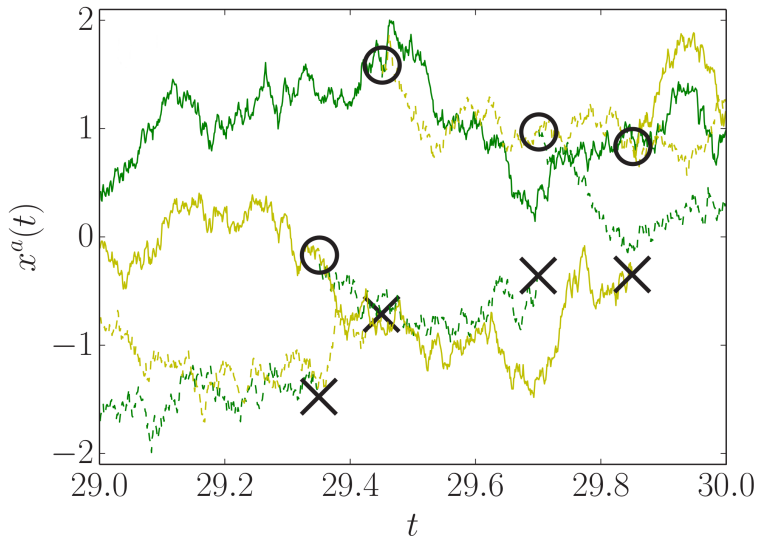
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[N_{nc} = number of copies in non-constant population dynamics]

Final-time distribution $p_{\text{end}}(\mathcal{C})$ governed by **right** eigenvector.

An example: 4 copies, 1 degree of freedom $\mathcal{C} = x \in \mathbb{R}$



How to perform averages? (ii) Intermediate times

$$\partial_t |\hat{P}\rangle = \mathbb{W}_s |\hat{P}\rangle$$

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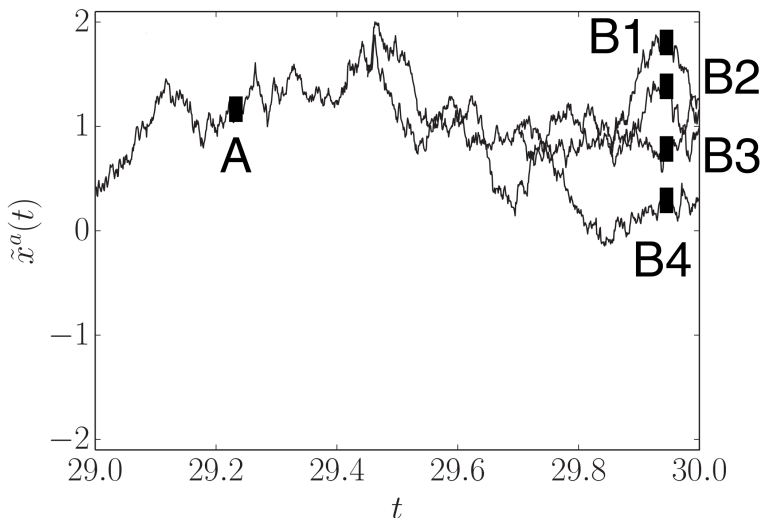
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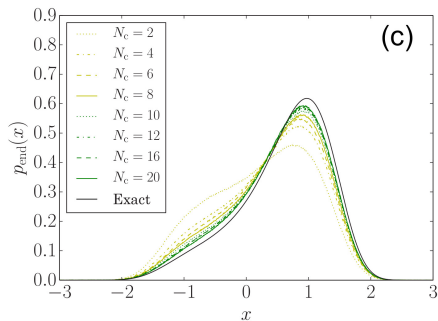
Mid-time distribution $p_{\text{ave}}(\mathcal{C})$ governed by **left** and **right** eigenvecs.

An example: 4 copies, 1 degree of freedom $\mathcal{C} = x \in \mathbb{R}$

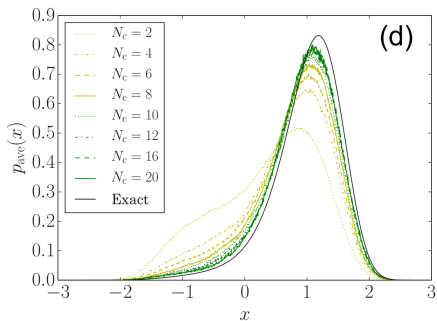


Huge sampling issue

Example distributions for a simple Langevin dynamics



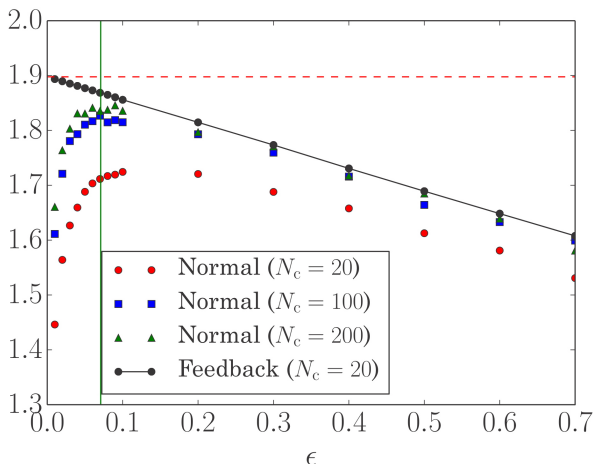
final-time: $p_{\text{end}}(x)$
 (= $R(x)$)



intermediate-time: $p_{\text{ave}}(x)$
 (= $R(x)L(x)$)

The small-noise crisis: systematic errors grow as $\epsilon \rightarrow 0$

CGF as a function of the noise amplitude ϵ :



Cause: as $\epsilon \rightarrow 0$, $p_{\text{ave}}(x)$ & $p_{\text{end}}(x) \rightarrow$ sharply peaked at *different points*
i.e. the clones do not sample correctly the phase space

The feedback method

How to make mid- and final-time distribution closer?

Driven/auxiliary dynamics: [Maes, Jack&Sollich, Touchette&Chetrite]

- Probability preserving
- No mismatch between p_{ave} and p_{end}
- Constructed as [Different dynamics can share \equiv statistical properties.]

$$W_s^{\text{aux}} = L W_s L^{-1} - \psi(s) \mathbf{1}$$

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- Issue: determining L is difficult
- Solution: evaluate L as L_{test} on the fly [feedback] and simulate

$$\mathbb{W}_s^{test} = L_{test}\mathbb{W}_sL_{test}^{-1} \quad (\text{induces } \textit{effective forces})$$

- **Iterate.** [For any L_{test} , the simulation is in principle correct.]

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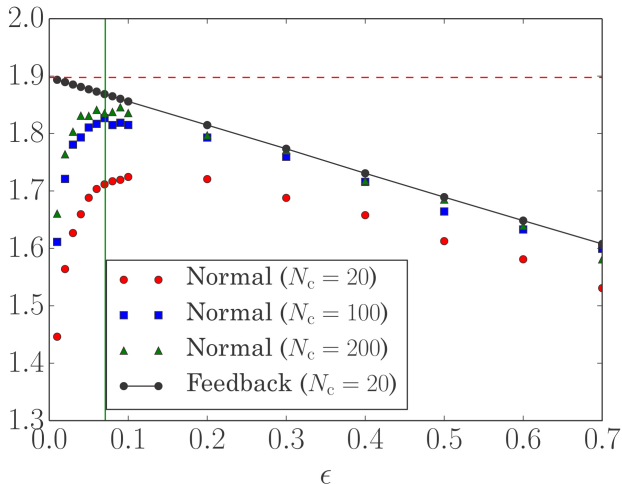
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Similar in spirit to **multi-canonical** (e.g. Wang–Landau) approaches in static thermodynamics. [Here, one flattens the left-eigenvector of \mathbb{W}_s^{test} .]

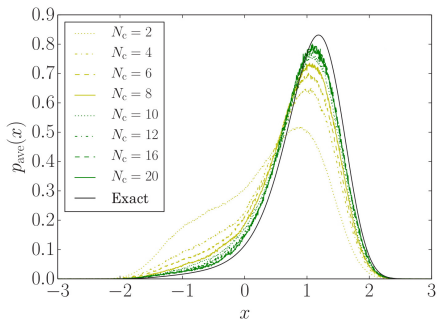
Improvement of the small-noise crisis (i.i)

CGF as a function of the noise amplitude ϵ :

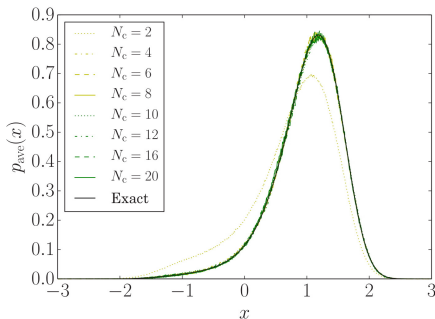


Physical insight: probability loss transformed into *effective forces*.

Improvement of the small-noise crisis (i.ii)



Without feedback

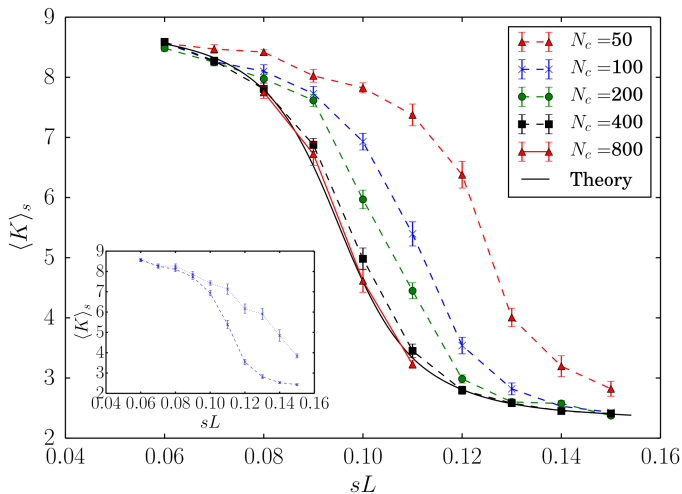


With feedback

Much more efficient evaluation of the biased distribution.

Even for a very crude (polynomial) approximation of the effective force.

Improvement of the small-noise crisis (ii)

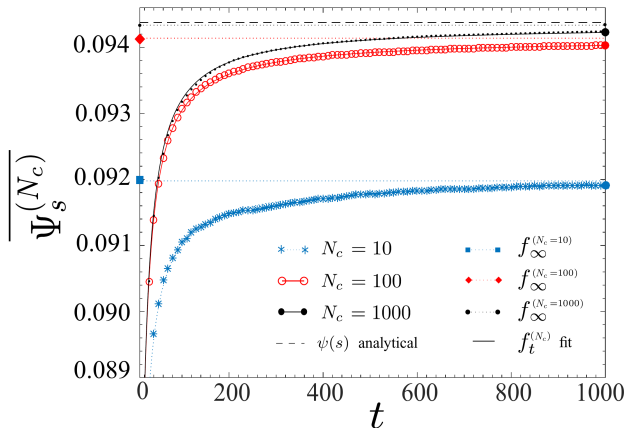


Interacting system in 1D.

Effective force: 1-, 2-, 3- body interactions only [also crude approx.].

Finite-time and -population effects

Finite-time scaling

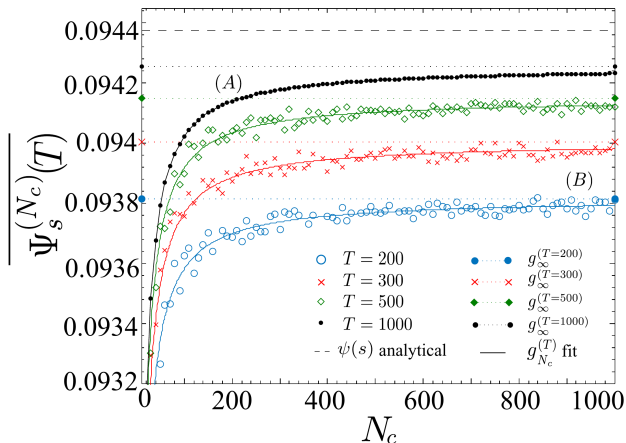
[fixed population N_c]

Estimator converges in $1/t$ to its infinite-time limit

Understanding: the estimator is an additive observable of the pop. dyn.

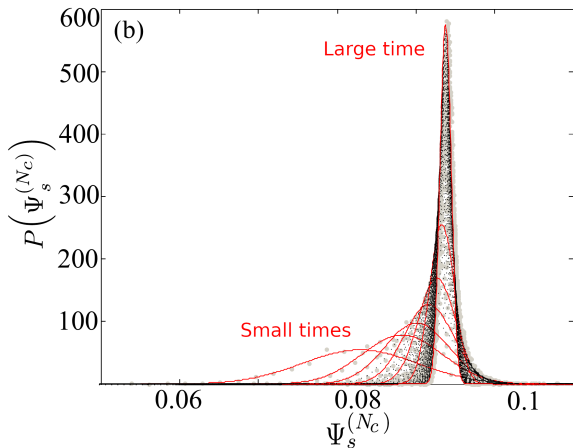
Finite- N_c scaling

[fixed time]



Estimator converges in $1/N_c$ to its infinite-population limit
 Understanding: large N_c expansion, small-noise description

Distribution of the CGF estimator [fixed population N_c]



In the numerics: \approx Gaussian when finite- N_c scaling is $O(1/N_c)$
 A way to check why one is / is not in that regime

Summary and open questions (1)

Feedback method

[with F Bouchet, R Jack, T Nemoto]

- Sampling problem (depletion of ancestors)
- On-the-fly evaluated auxiliary dynamics
- Solution to the small-noise crisis
- Systems with large number of degrees of freedom

Summary and open questions (1)

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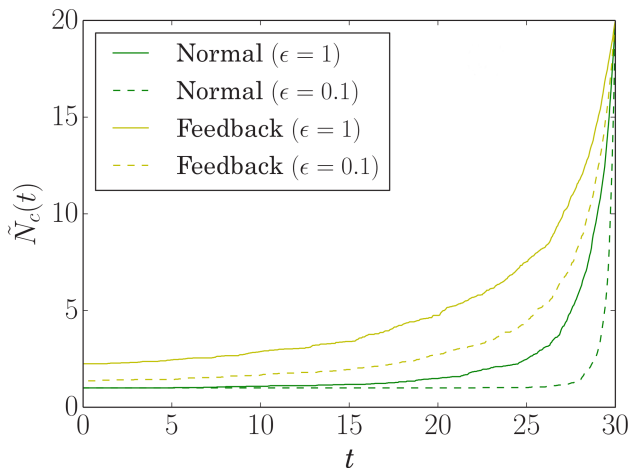
Finite-population effects

[with E Guevara, T Nemoto]

- Quantitative finite- N_{clones} scaling \rightarrow interpolation method
- Initial transient regime due to small population
- Analogy with biology: many small islands vs. few large islands?
- Question: effective forces \leftarrow selection?

Open questions (2): why is the feedback working?

Improvement of the depletion-of-ancestors problem:



Dashed line: lower noise

Continuous line: higher noise

Open questions (3)

Finite-population and -time scalings

- Anomalous fluctuations (invalid $1/N_c$ asymptotics)
- Correct description of the meta-dynamics?
- Finite- N_c and $-t$ scaling **with feedback**
- Phase transition in the distribution of the CGF estimator?

Thank you for your attention!

References:

- ★ *Population dynamics method with a multi-canonical feedback control*
Takahiro Nemoto, Freddy Bouchet, Robert L. Jack and Vivien Lecomte
PRE **93** 062123 (2016)
- ★ *Finite-time and finite-size scalings in the evaluation of large deviation functions: Analytical study using a birth-death process*
Takahiro Nemoto, Esteban Guevara Hidalgo and Vivien Lecomte
PRE **95** 012102 (2017)
- ★ *Finite-size scaling of a first-order dynamical phase transition: adaptive population dynamics and effective model*
Takahiro Nemoto, Robert L. Jack and Vivien Lecomte
PRL **118** 115702 (2017)
- ★ *Finite-time and finite-size scalings in the evaluation of large deviation functions: Numerical approach in continuous time*
Esteban Guevara Hidalgo, Takahiro Nemoto and Vivien Lecomte
PRE **95** 062134 (2017)

Supplementary material

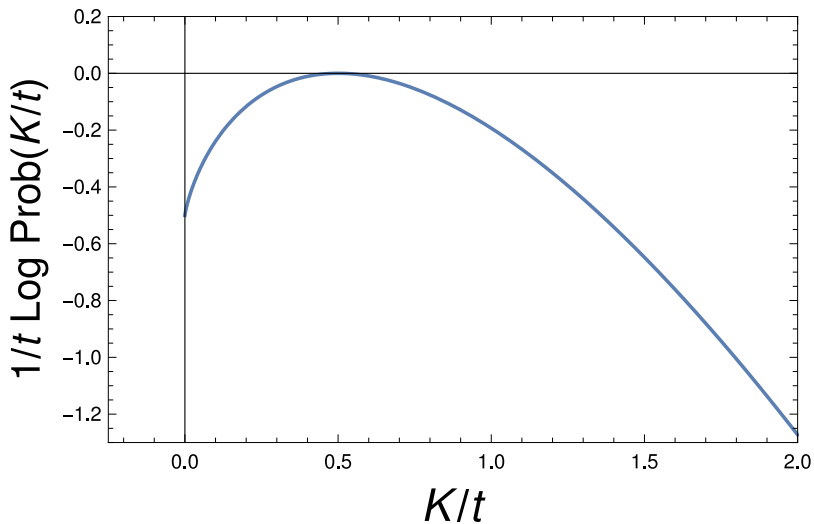
How to perform averages?

★ Mid-time ancestor distribution:

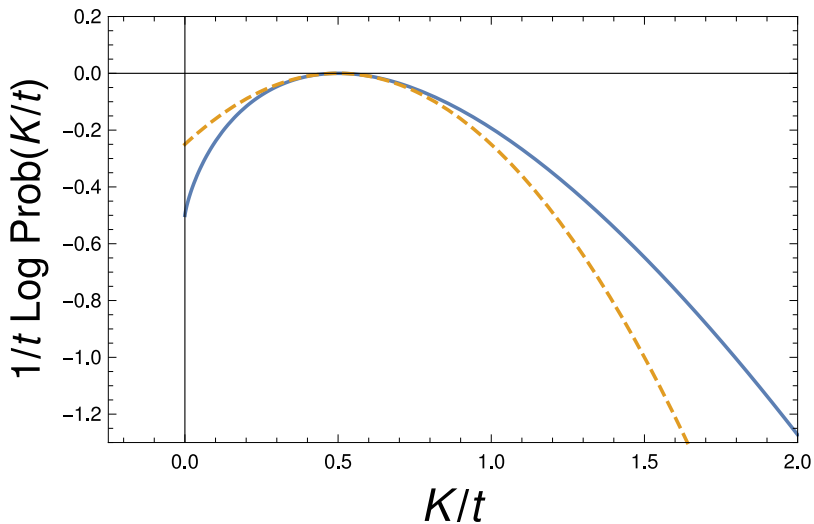
fraction of copies (at time t_1) which were in configuration \mathcal{C} , knowing that there are in configuration \mathcal{C}_f at final time t_f :

$$p_{\text{anc}}(\mathcal{C}, t_1; \mathcal{C}_f, t_f) = \frac{\langle N_{\text{nc}}(\mathcal{C}_f, t_f | \mathcal{C}, t_1) \rangle_s}{\sum_{\mathcal{C}'} \langle N_{\text{nc}}(\mathcal{C}_f, t_f | \mathcal{C}', t_1) \rangle_s} \underset{t_f, 1 \rightarrow \infty}{\sim} \langle L | \mathcal{C} \rangle \langle \mathcal{C} | R \rangle = p_{\text{ave}}(\mathcal{C})$$

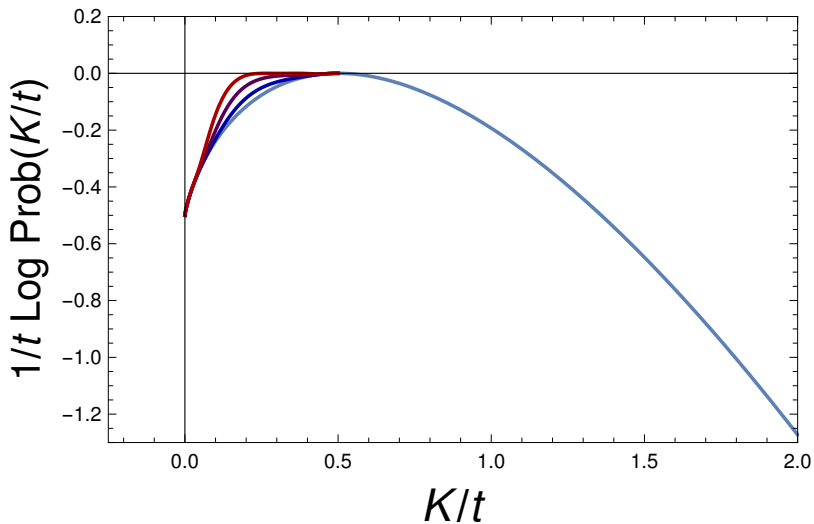
The “ancestor statistics” of a configuration \mathcal{C}_f is thus independent (far enough in the past) of the configuration \mathcal{C}_f .



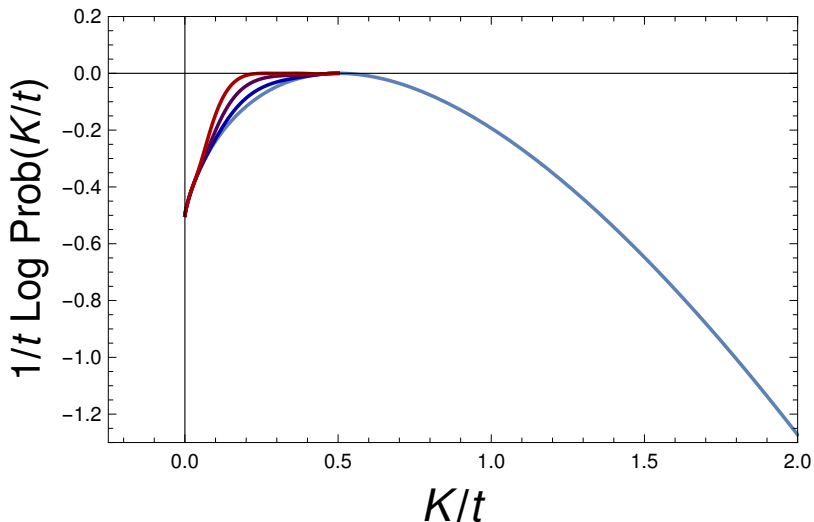
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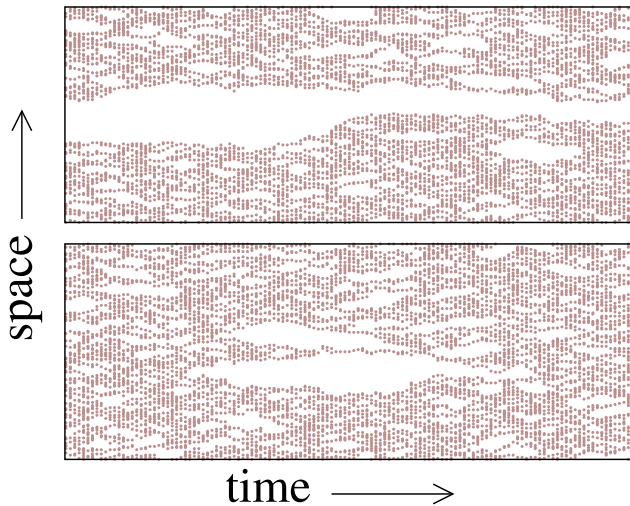


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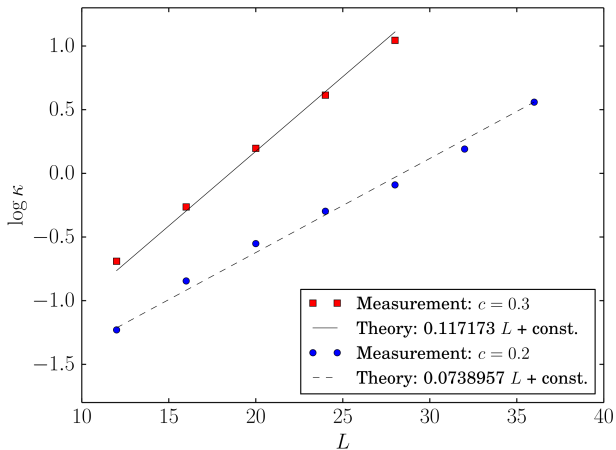


$$\text{Prob}[K] \sim e^{t\varphi(K/t)}$$

Finite-time & -size scalings matter.

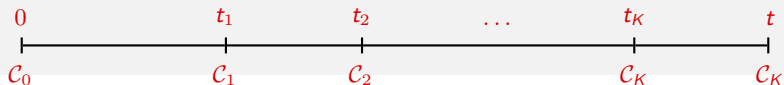


[Merolle, Garrahan and Chandler, 2005]



Exponential divergence of the susceptibility

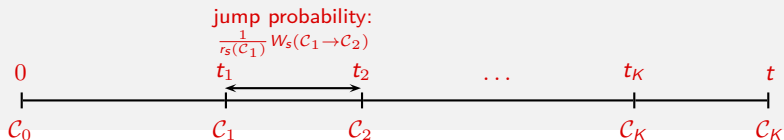
Explicit construction (1/3)



Probability-preserving contribution

$$\partial_t \hat{P}(C, t) = \sum_{C'} \left\{ \underbrace{W_S(C' \rightarrow C) \hat{P}(C', t)}_{\text{gain term}} - \underbrace{W_S(C \rightarrow C') \hat{P}(C, t)}_{\text{loss term}} \right\}$$

Explicit construction (1/3)



Which configurations will be visited?

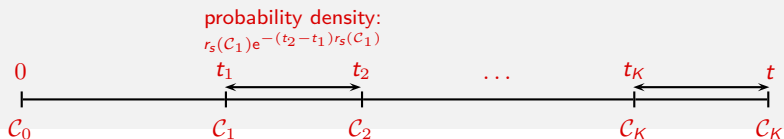
Configurational part of the trajectory: $C_0 \rightarrow \dots \rightarrow C_K$

$$\text{Prob}\{\text{hist}\} = \prod_{n=0}^{K-1} \frac{W_s(C_n \rightarrow C_{n+1})}{r_s(C_n)}$$

where

$$r_s(C) = \sum_{C'} W_s(C \rightarrow C')$$

Explicit construction (2/3)

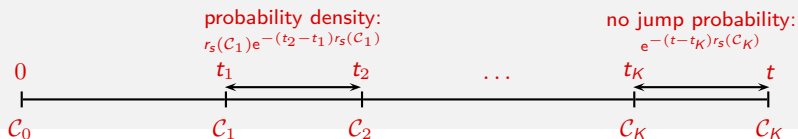


When shall the system jump from one configuration to the next one?

- probability density for the time interval $t_n - t_{n-1}$

$$r_s(C_{n-1})e^{-(t_n-t_{n-1})r_s(C_{n-1})}$$

Explicit construction (2/3)



When shall the system jump from one configuration to the next one?

- probability density for the time interval $t_n - t_{n-1}$

$$r_s(C_{n-1})e^{-(t_n-t_{n-1})r_s(C_{n-1})}$$

- probability not to leave C_K during the time interval $t - t_K$

$$e^{-(t-t_K)r_s(C_K)}$$

Explicit construction (3/3)

$$\partial_t \hat{P}(\mathcal{C}, s) = \underbrace{\sum_{\mathcal{C}'} W_s(\mathcal{C}' \rightarrow \mathcal{C}) \hat{P}(\mathcal{C}', s) - r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{modified dynamics}} + \underbrace{\delta r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{cloning term}}$$

How to take into account loss/gain of probability?

- handle a large number of copies of the system
- implement a **selection** rule: on a time interval Δt a copy in config \mathcal{C} is replaced by $e^{\Delta t \delta r_s(\mathcal{C})}$ copies
- $\psi(s) =$ the rate of exponential growth/decay of the total population
- optionally: keep population constant by non-biased pruning/cloning

Explicit construction (3/3)

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How to take into account loss/gain of probability?

- handle a large number of copies of the system
- implement a **selection** rule: on a time interval Δt a copy in config \mathcal{C} is replaced by $\lfloor e^{\Delta t \delta r_s(\mathcal{C})} + \epsilon \rfloor$ copies, $\epsilon \sim [0, 1]$
- $\psi(s)$ = the rate of exponential growth/decay of the total population
- optionally: keep population constant by non-biased pruning/cloning

Explicit construction (3/3)

$$\partial_t \hat{P}(\mathcal{C}, s) = \underbrace{\sum_{\mathcal{C}'} W_s(\mathcal{C}' \rightarrow \mathcal{C}) \hat{P}(\mathcal{C}', s) - r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{modified dynamics}} + \underbrace{\delta r_s(\mathcal{C}) \hat{P}(\mathcal{C}, s)}_{\text{cloning term}}$$

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Biological interpretation

- copy in configuration $\mathcal{C} \equiv$ organism of **genome** \mathcal{C}
- dynamics of rates $W_s \equiv$ **mutations**
- cloning at rates $\delta r_s \equiv$ **selection** rendering atypical histories typical