#### Population dynamics and rare events



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#### Centre Henri Lebesgue, Rennes - 7 June 2018







Motivations

#### Why studying rare events?



#### 2003 heat wave, Europe [Terra MODIS]

Vivien Lecomte (LIPhy)

Motivations

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

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Motivations

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

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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  $\frac{1}{t_{max}} \int_{0}^{t_{max}} dt T(t)$ 

 $\longleftrightarrow$  anthropogenic impact on climate?

[Otto et al., 2012]

#### Outline

- Introduction
- Settings
- Different averages
- Feedback method
- Finite-time and finite-population scalings
- Open questions

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

Tools



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Tools



#### Changing ensemble

#### s-modified dynamics

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

#### s-modified dynamics

#### K =activity= #events

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Configs.  $\mathcal{C}\text{, jump rates }\textit{W}(\mathcal{C}\rightarrow\mathcal{C}')$ 

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Tools

• More detailed dynamics for P(C, K, t):

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

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

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#### s-modified dynamics

$$\mathbf{K} = \mathbf{k}_{\mathcal{C}_0 \mathcal{C}_1} + \mathbf{k}_{\mathcal{C}_1 \mathcal{C}_2} + \dots$$

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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"]

$$\sum_{s} \hat{P}(\mathcal{C}, s, t) = \left\langle e^{-s K} \right\rangle \sim e^{t \psi(s)} \qquad (\psi(s) = \mathsf{CGF})$$

• discrete time: Giardinà, Kurchan, Peliti [PRL 96, 120603 (2006)]

continuous time: VL, Tailleur [JSTAT P03004 (2007)]



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# Cloning dynamics $\partial_{t}\hat{P}(\mathcal{C},s) = \underbrace{\sum_{\mathcal{C}'} W_{s}(\mathcal{C}' \to \mathcal{C})\hat{P}(\mathcal{C}',s) - r_{s}(\mathcal{C})\hat{P}(\mathcal{C},s)}_{\text{modified dynamics}} + \underbrace{\delta r_{s}(\mathcal{C})\hat{P}(\mathcal{C},s)}_{\text{cloning term}}$ • $W_{s}(\mathcal{C}' \to \mathcal{C}) = e^{-s}W(\mathcal{C}' \to \mathcal{C})$ • $r_{s}(\mathcal{C}) = \sum_{\mathcal{C}'} W_{s}(\mathcal{C} \to \mathcal{C}')$ • $\delta r_{s}(\mathcal{C}) = r_{s}(\mathcal{C}) - r(\mathcal{C})$

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

- handle a large number  $N_{\rm c}$  of copies of the system
- implement a selection rule: on a time interval Δt a copy in config C is replaced by Y = e<sup>Δt δr<sub>s</sub>(C)</sup> copies
- $\psi(s) =$  the rate of exponential growth/decay of the total population
- optionally: keep population constant by non-biased pruning/cloning

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How to take into account loss/gain of probability?

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CGF estimator: 
$$\psi(s) = \mathbb{E}\Psi(s)$$
 with  $\Psi(s) = \log \prod_{t} \frac{N_c + Y_{t-1}}{N_c}$ 

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#### Biological interpretation

- $\bullet$  copy in configuration  $\mathcal{C}\equiv$  organism of  $genome \ \mathcal{C}$
- dynamics of rates  $W_s \equiv$  mutations
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- the CGF  $\psi(s)$  is a measure of the **fitness** of the population

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



## How to perform averages? (i)

# [spectral analysis]

#### $\star$ Final-time distribution: *proportion* of copies in ${\cal C}$ at t

$$\begin{split} \langle N_{\rm nc}(t) \rangle_s \\ \langle N_{\rm nc}(\mathcal{C},t) \rangle_s \\ p_{\rm end}(\mathcal{C},t) &= \frac{\langle N_{\rm nc}(\mathcal{C},t) \rangle_s}{\langle N_{\rm nc}(t) \rangle_s} \end{split}$$

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

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Final-time distribution governed by right eigenvector.

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Mid-time distribution governed by left and right eigenvectors.

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



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#### How to perform averages?

★ Mid-time ancestor distribution:

fraction of copies (at time  $t_1$ ) which were in configuration C, knowing that there are in configuration  $C_f$  at final time  $t_f$ :

$$p_{\rm anc}(\mathcal{C}, t_1; \mathcal{C}_{\rm f}, t_{\rm f}) = \frac{\langle N_{\rm nc}(\mathcal{C}_{\rm f}, t_{\rm f} | \mathcal{C}, t_1) \rangle_s}{\sum_{\mathcal{C}'} \langle N_{\rm nc}(\mathcal{C}_{\rm f}, t_{\rm f} | \mathcal{C}', t_1) \rangle_s} \underset{t_{\rm f,1} \to \infty}{\sim} \langle L | \mathcal{C} \rangle \langle \mathcal{C} | \mathcal{R} \rangle = p_{\rm ave}(\mathcal{C})$$

The "ancestor statistics" of a configuration  $C_f$  is thus independent (far enough in the past) of the configuration  $C_f$ .

#### Example distributions for a simple Langevin dynamics



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

CGF as a function of the noise amplitude  $\epsilon$ :



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#### The feedback method

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

- Probability preserving
- No mismatch between pave and pend
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Issue: determining L is difficult

• Solution: evaluate L as L<sub>test</sub> on the fly [feedback] and simulate

 $\mathbb{W}_{s}^{\text{test}} = L_{\text{test}} \mathbb{W}_{s} L_{\text{test}}^{-1}$  (induces *effective forces*)

• Iterate. [For any L<sub>test</sub>, the simulation is in principle correct.]

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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  $\mathbb{W}_{s}^{\text{test}}$ .]

#### Results

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

CGF as a function of the noise amplitude  $\epsilon:$ 



Physical insight: probability loss transformed into effective forces.

Vivien Lecomte (LIPhy)

Results

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



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.].

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#### 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





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)

# 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

#### 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 ← selection?
- Question: relation to recent work of Ferré and Touchette?

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

Improvement of the depletion-of-ancestors problem:



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# Open questions (3)

#### Finite-population and -time scalings

- Anomalous fluctuations (invalid 1/N<sub>c</sub> asymptotics)
- Correct description of the meta-dynamics?
- Finite-N<sub>c</sub> and -t scaling with feedback
- Phase transition in the distribution of the CGF estimator?

# Thanks 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



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



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





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Population dynamics & rare events

07/06/2018 31/29





time  $\longrightarrow$ 

[Merolle, Garrahan and Chandler, 2005]



Exponential divergence of the susceptibility

# Explicit construction (1/3)







#### Which configurations will be visited?

Configurational part of the trajectory:  $\mathcal{C}_0 \to \ldots \to \mathcal{C}_K$ 

$$\mathsf{Prob}\{\mathsf{hist}\} = \prod_{n=0}^{K-1} \frac{W_{\mathsf{s}}(\mathcal{C}_n \to \mathcal{C}_{n+1})}{r_{\mathsf{s}}(\mathcal{C}_n)}$$

where

$$r_{s}(\mathcal{C}) = \sum_{\mathcal{C}'} W_{s}(\mathcal{C} \to \mathcal{C}')$$

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

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

 $r_{s}(\mathcal{C}_{n-1})e^{-(t_{n}-t_{n-1})r_{s}(\mathcal{C}_{n-1})}$ 



When shall the system jump from one configuration to the next one? • probability density for the time interval  $t_n - t_{n-1}$ 

$$r_{s}(\mathcal{C}_{n-1})e^{-(t_{n}-t_{n-1})r_{s}(\mathcal{C}_{n-1})}$$

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

$$e^{-(t-t_{\mathcal{K}})r_{s}(\mathcal{C}_{\mathcal{K}})}$$

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

- handle a large number of copies of the system
- implement a selection rule: on a time interval Δt a copy in config C is replaced by e<sup>Δt δr<sub>s</sub>(C)</sup> copies
- $\psi(s) =$  the rate of exponential growth/decay of the total population
- optionally: keep population constant by non-biased pruning/cloning

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

- handle a large number of copies of the system
- implement a selection rule: on a time interval  $\Delta t$ a copy in config C is replaced by  $\lfloor e^{\Delta t \, \delta r_{\rm s}(C)} + \varepsilon \rfloor$  copies,  $\epsilon \sim [0, 1]$
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#### Biological interpretation

- $\bullet$  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