Population dynamics and rare events



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Motivations

Why studying rare events?



2003 heat wave, Europe [Terra MODIS]

Vivien Lecomte (LPMA & LIPhy)

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

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Motivations

Why studying rare events?



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

Vivien Lecomte (LPMA & LIPhy)

Why studying rare events?



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Questions for physicists:

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

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Evolution of the return time of the monthly averaged temperature $\frac{1}{t_{max}} \int_{0}^{t_{max}} dt T(t)$

Due anthropogenic impact on climate?

[Otto et al., 2012]

Distribution of a time-extensive observable K

Tools



Distribution of a time-extensive observable K

Tools



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

• Markov processes:

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

K = activity = #events

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

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Numerical method

[with J. Tailleur]

Evaluation of large deviation functions

[à la Monte-Carlo diffusion]

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

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

Tools

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

Cloning dynamics

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

odified dynamics

cloning term

•
$$W_{s}(\mathcal{C}' \to \mathcal{C}) = e^{-s}W(\mathcal{C}' \to \mathcal{C})$$

•
$$r_s(\mathcal{C}) = \sum_{\mathcal{C}'} W_s(\mathcal{C} \to \mathcal{C}')$$

•
$$\delta r_{s}(\mathcal{C}) = r_{s}(\mathcal{C}) - r(\mathcal{C})$$

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Which configurations will be visited?

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

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

where

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

Vivien Lecomte (LPMA & LIPhy)

Explicit construction

Explicit construction (2/3)



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

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

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



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

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

 $e^{-(t-t_K)r_s(\mathcal{C}_K)}$

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

- 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^{Δt δr_s(C)} 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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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

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



How to perform averages? (i)

[with R Jack, F Bouchet, T Nemoto]

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

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

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

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

Final-time distribution governed by **right** eigenvector.

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



How to perform averages? (ii) Intermediate times

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Mid-time distribution governed by left and right eigenvectors.

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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



final-time: $p_{end}(x)$

intermediate-time: $p_{ave}(x)$

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



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

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

- Probability preserving
- No mismatch between pave and pend
- Constructed as

 $\mathbb{W}^{\mathsf{aux}}_{s} = L \mathbb{W}_{s} L^{-1} - \psi(s) \mathbf{1}$

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

$$\mathbb{W}_{s}^{\mathsf{test}} = L_{\mathsf{test}} \mathbb{W}_{s} L_{\mathsf{test}}^{-1}$$

• Whichever *L*_{test}, the simulation is still correct. **Iterate**

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$$\mathbb{W}_{s}^{\mathsf{test}} = \mathcal{L}_{\mathsf{test}} \mathbb{W}_{s} \mathcal{L}_{\mathsf{test}}^{-1}$$

• Whichever *L*_{test}, the simulation is still correct. **Iterate** Similar in spirit to **multi-canonical** (e.g. Wang-Landau) approach in static thermodynamics

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



Physical insight: probability loss transformed into effective forces

Vivien Lecomte (LPMA & LIPhy)

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.].

Vivien Lecomte (LPMA & LIPhy)

Summary and questions (1)

Multicanonical approach

[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 questions (1)

Multicanonical approach [with F Bouchet, R Jack, T Nemoto]

- Sampling problem (depletion of ancestors)
- On-the-fly evaluated auxiliary dynamics
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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 ← selection?

Questions (2): why is it working?

Improvement of the depletion-of-ancestors problem:



Vivien Lecomte (LPMA & LIPhy)



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





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





time \longrightarrow

[Merrolle, Garrahan and Chandler, 2005]

Conclusion



Exponential divergence of the susceptibility