Stochastic persistence of ecological communities driven by Lévy-noise

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Persistence: preliminaries

A simple stochastic LV system of prey-predator type with intraspecific competition driven by Brownian motion (here the flow of biomass is $1 \longrightarrow 2$).

$$dX_t^{(1)} = X_t^{(1)}(b_1 - a_{11}X_t^{(1)} - a_{12}X_t^{(2)}) + \sigma_1 X_t^{(1)} dW_t^{(1)}$$
(1)
$$dX_t^{(2)} = X_t^{(2)}(-b_2 + a_{21}X_t^{(1)} - a_{22}X_t^{(2)}) + \sigma_2 X_t^{(2)} dW_t^{(2)}.$$

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Initial conditions for the process $\mathbf{X} = (X^{(1)}, X^{(2)})$ in \mathbb{R}^n_+ = positive orthant $(\mathbb{R}^n_{++} = \text{strictly positive orthant}).$

Biological interpretation:

- (1) b_i 's: per-capita per-unit time birth-death rates.
- 2) a_{ij} 's: if $i \neq j$, interaction strength (increase-decrease of fitting per unit time per prey-predator encounter). If i = j: intraspecific competition (due e.g. to competition for space, soil, light, etc.)
- (3) σ_i 's: amount of the random perturbation due to environmental noise.

W = (W⁽¹⁾, W⁽²⁾): standard 2d Wiener process, carrying the environmental noise.

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Modeling issue: it is assumed that the stochastic perturbations of the fitting occur in characteristic times much shorter than the time horizon relevant to our observations

In our work we consider communities of n populations indexed by I engaged in LV dynamics driven by continuous and jump-type noise.

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• It can be proved that for initial condition on the strictly positive cone, for every t > 0, $X_t^{(1)} > 0$, $X_t^{(2)} > 0$ a.s.

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• It can be proved that for initial condition on the strictly positive cone, for every t > 0, $X_t^{(1)} > 0$, $X_t^{(2)} > 0$ a.s. \Rightarrow there is no extinction on finite time. \Rightarrow any notion of stochastic persistence involves a claim on the asymptotic behavior of the laws.

Definition

A set of species indexed by $J \subset I$ is said to be **stochastically persistent** in probability if for every $\varepsilon > 0$ there exists a compact set $K_{\varepsilon} \subset \mathbb{R}^{|J|}_{++}$ such that:

$$\liminf_{t\to\infty}\mathbb{P}_{\mathbf{x}}\left((X_t^{(i_1)},X_s^{(i_2)},\ldots,X_t^{(i_l)})\in K_{\varepsilon},(i_1,i_2,\ldots,i_l)\in J\right)>1-\varepsilon,$$

uniformly on the initial condition $\mathbf{x} \in \mathbb{R}'_{++}$.

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• Previous work has provided condition for these *weak* form of persistence for Lévy-driven ecological models: [BY11], [Mao11], [BMYY12].

Strong Stochastic Persistence

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- We ask: when started from $\mathbf{x} \in \mathbb{R}^n_{++},$
 - (1) when do the laws of X converges (in an appropriate sense) to a unique invariant probability measure, π ?
 - What is the nature of this invariant measure? Namely: does it support the whole community or just a strict subset of the community?

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 - **1** when do the laws of **X** converges (in an appropriate sense) to a unique invariant probability measure, π ?
 - What is the nature of this invariant measure? Namely: does it support the whole community or just a strict subset of the community?
- If π does not change the boundary of the positive cone, then we say that the community is **strongly stochastic persistent**.

Definition

(see [HN18a]) We say that a \mathbb{R}_{++}^n -valued, right-continuous Markov process ($\mathbf{X}_t : t \ge 0$) is strongly stochastically persistent (SSP) if there exists a unique invariant probability measure π supported on \mathbb{R}_{++}^n such that for every $\mathbf{x} \in \mathbb{R}_{++}^n$:

$$d_{TV}(\mathbb{P}_{\mathbf{x}}(\mathbf{X}_t \in \cdot), \pi(\cdot)) \to 0,$$

as $t
ightarrow \infty$.

We observe:

- **1** X is not empty (because δ_0 is trivially ergodic).
- 2 For every $J \subseteq \{1, 2\}$, the sets:

$$\mathbb{R}^{J}_{++} := \{ \mathbf{x} \in \mathbb{R}^{n}_{+} : x_{i} > 0, i \in J, x_{i} = 0, i \notin J \},\$$

are positively invariant for the dynamics. In general, thus, we can expect to have a lot of invariant probability measures on \mathbb{R}_+^n , many concentrated on $\partial \mathbb{R}_+^n$ (the boundary of the orthant, where at least one of the species is extinct).

Crucial insight

• Pick an invariant probability measure μ such that at least one of the species is extinct under μ (and thus, μ charges $\partial \mathbb{R}^n_+$). Suppose that the process X, when started from strictly positive abundances, gets close to the part of the extinction boundary that is changed by μ . To guarantee SSP we will need that, in this situation, at least one of the endangered species tend to grow rapidly, avoiding thus the extinction fate.

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• Under some regularity assumption, this will amount to ask that, under any invariant probability measure μ that charges the boundary, at least one of the absent species has positive μ -averaged Lyapunov exponent.

Back to first example

How to ensure SSP for this community?



We have many invariant (indeed, ergodic) p.m. for this system.

- δ_{0,0}.
- An ergodic p.m. μ under which species 1 persists and species 2 is extinct (thus, concentrated on {x₁ > 0, x₂ = 0}).

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We have many invariant (indeed, ergodic) p.m. for this system.

- δ_{0,0}.
- An ergodic p.m. μ under which species 1 persists and species 2 is extinct (thus, concentrated on $\{x_1 > 0, x_2 = 0\}$). This p.m. is unique, and corresponds to the product measure $m \otimes \delta_0$, where *m* is the law of a Gamma-distributed random variable (the law of the stochastic logistic equation).

Apply (formally!) Ito's formula to $\mathbf{x} \mapsto \log(x_1)$ and $\mathbf{x} \mapsto \log(x_2)$ to obtain:

$$\frac{\log(X_t^{(1)})}{t} - \frac{\log(X_0^{(1)})}{t} = \frac{1}{t} \int_0^t \left(b_1 - a_{11}X_s^{(1)} - a_{12}X_s^{(2)} - \frac{1}{2}\sigma_1^2 \right) ds + \frac{M_1(t)}{t} \frac{\log(X_t^{(2)})}{t} - \frac{\log(X_0^{(2)})}{t} = \frac{1}{t} \int_0^t \left(-b_2 + a_{21}X_s^{(1)} - a_{22}X_s^{(2)} - \frac{1}{2}\sigma_2^2 \right) ds + \frac{M_2(t)}{t}$$

where M_1 and M_2 are local martingales. Put:

$$\Xi_1(\mathbf{x}) := b_1 - a_{11}x_1 - a_{12}x_2 - \frac{1}{2}\sigma_1^2$$
$$\Xi_2(\mathbf{x}) := -b_2 + a_{21}x_1 - a_{22}x_2 - \frac{1}{2}\sigma_2^2$$

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If X expends a lot of time close to, for example, the support of μ, then for large times the first integrals should be close to μΞ₁ and μΞ₂.

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2 Since under μ the first species persists, $\mu \Xi_1 = 0$. (resident, stable species do not die or grow exponentially fast.)

- **1** $max_{i=1,2}\delta_{0,0}\Xi_i > 0$ (when both are rare, at least one of the species tends to grow exponentially fast).
- 2) $\mu \Xi_2 > 0$ (when rare, the predator tends to grow exponentially fast).

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From the first condition we obtain:

$$b_1-\frac{1}{2}\sigma_1^2>0$$

and from the second one we get:

$$-b_2+a_{21}\mathbb{E}_{Z\sim m}(Z)-rac{1}{2}\sigma_2^2>0$$

Since $\mu \Xi_1 = 0$, we have:

$$\mathbb{E}_{Z\sim m}(Z)=\frac{b_1-\frac{1}{2}\sigma_1^2}{a_{11}}$$

Thus, still intuitively, a sufficient condition for SSP is:

$$rac{b_1 - rac{1}{2}\sigma_1^2}{a_{11}} > rac{b_2 + rac{1}{2}\sigma_2^2}{a_{21}}$$

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• In the simple example of the prey-predator community, the above intuition is just right ([Ben18], [HN18b]).

Thus, still intuitively, a sufficient condition for SSP is:

$$\frac{b_1 - \frac{1}{2}\sigma_1^2}{a_{11}} > \frac{b_2 + \frac{1}{2}\sigma_2^2}{a_{21}}$$

• In the simple example of the prey-predator community, the above intuition is just right ([Ben18], [HN18b]).

• The aim of this work is to show how these results can be extended to communities such as (2).

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Our framework, assumption and results

Our SDE framework

$$d\mathbf{X}_{t} = \mathbf{X}_{t} \circ \left((\mathbf{B} + \mathbf{A}\mathbf{X}_{t})dt + \boldsymbol{\Sigma}d\mathbf{W}_{t} + \int_{\mathbb{R}^{n} \setminus \{0\}} \mathbf{L}(\mathbf{X}, \mathbf{z})\tilde{N}(dt, dz) \right).$$
(2)

$$d\mathbf{X}_{t} = \mathbf{X}_{t} \circ \left((\mathbf{B} + \mathbf{A}\mathbf{X}_{t})dt + \Sigma d\mathbf{W}_{t} + \int_{\mathbb{R}^{n} \setminus \{0\}} \mathbf{L}(\mathbf{X}, \mathbf{z})\tilde{N}(dt, dz) \right).$$
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(Σ : constant; \tilde{N} : a compensated Poisson random measure of intensity measure $dt\nu(dz)$ for ν a Lévy measure on \mathbb{R}^n).

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(2)

(Σ : constant; \tilde{N} : a compensated Poisson random measure of intensity measure $dt\nu(dz)$ for ν a Lévy measure on \mathbb{R}^n).

• **Biological interpretation of the jump part:** Other sources of randomness on the biotic or abiotic factors of the ecosystems whose natural time-scale are much longer than those of the perturbation modeled by the Brownian part: sudden migration of resident species or incoming flow of biomass through ecosystem boundaries (see e.g. the excellent compendium on the topic of animal migration [MG11], specially chapter 9) or ENSO effects on habitat compression and deepening of nutricline (see [OMK⁺17]).

Our framework regarding the communities

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Our framework regarding the communities

• The signs of **A** determine a directed graph (with an edge from prey to predators).

Our framework regarding the communities

- The signs of **A** determine a directed graph (with an edge from prey to predators).
- We consider layered communities: food-webs with intraspecific competition with no autocalytic cycles.



basal species

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Assumptions

- The community is layered.
- (Assumption 1): $\int_{\mathbb{R}^n \setminus \{0\}} \|z\|^2 \nu(dz) < \infty$.
- (Assumption 2): For every i = 1, 2, ..., n, $\mathbf{x}, \mathbf{z} \in \mathbb{R}^n$:

$$|L_i(\mathbf{x},\mathbf{z})| \leq |z_i|\mathbf{1}_{z_i>-1}.$$

(Assumption 3): There exists a constant C such that:

$$\|\mathbf{x} \circ \mathbf{L}(\mathbf{x}, \mathbf{z})\| \leq C \|\mathbf{z}\|$$

(Assumption 4): For \mathbf{x}, \mathbf{y} in K compact, $\mathbf{z} \in \mathbb{R}^n$, there exists a constant M_k such that:

$$\|\mathbf{x} \circ \mathbf{L}(\mathbf{x}, \mathbf{z}) - \mathbf{y} \circ \mathbf{L}(\mathbf{y}, \mathbf{z})\| \leq M_k \|\mathbf{x} - \mathbf{y}\| \|\mathbf{z}\|,$$

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holds.

Just like in our example, define:

$$\Xi_i(\mathbf{x}) := (B_i + (\mathbf{A}\mathbf{x})_i) - \frac{1}{2}\sigma_i^2 - \int_{\mathbb{R}^n_+} L_i(\mathbf{x}, \mathbf{z}) - \ln(1 + L_i(\mathbf{x}, \mathbf{z}))\nu(d\mathbf{z}).$$

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More technical assumptions

Observe that any layered community has the following property: there exists c ∈ ℝⁿ₊₊ such that whenever i → j:

$$c_i a_{ij} \geq c_j a_{ji}.$$

It is proved that the conditions:

For every
$$\mu \in \mathcal{P}_{erg}(\partial \mathbb{R}^n_+) : \max_i \mu \Xi_i > 0$$

and

There exists
$$\boldsymbol{p} := (p_1, \dots, p_n) \in \mathbb{R}^n_{++}$$
 such that:

$$\inf_{\mu \in \mathcal{P}_{erg}(\partial \mathbb{R}^n_+)} \sum_i p_i \mu \Xi_i := \rho > 0, \quad (3)$$

are equivalent ([SBA11]).

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(Assumption 5): The condition (3) holds.

Define for $\mathbf{x} \in \mathbb{R}^{n}_{++}$

$$\hat{\mathcal{V}}(\mathbf{x}) = rac{1+\mathbf{c}^T\mathbf{x}}{\prod_{i=1}^n x_j^{p_j}},$$

and for $\alpha > 0$ define:

$$ilde{\mathcal{W}}(\mathsf{x}) := \hat{\mathcal{V}}^lpha(\mathsf{x}).$$

(the c_i 's are those ensured by the layered hypothesis; the p_i 's are *small* and satisfy the condition (3); ; $\alpha > 0$ will be fixed appropriately).

Observe that for p_i small (namely, $\sum_i p_i < 1$), the function \hat{V} satisfy:

$$\lim_{\|\mathbf{x}\|\to\infty} \hat{V}(\mathbf{x}) = \liminf_{\mathbf{x}:\min_i x_i\to 0} \hat{V}(\mathbf{x}) = +\infty$$

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For $\mathbf{x} \in \mathbb{R}^{n}_{++}$, set:

$$I(\mathbf{x}) := \int_{\mathbb{R}^n} \left\{ \tilde{W}((x_i(1+L_i(\mathbf{x},\mathbf{z}))_i) - \tilde{W}(\mathbf{x}) -\alpha \tilde{W}(\mathbf{x}) \sum_{i=1}^n \left(L_i(\mathbf{x},\mathbf{z})(\frac{c_i x_i}{1+c^T \mathbf{x}} - p_i) \right) \right\} \nu(d\mathbf{z}).$$

Assumption 6:

 $I(\mathbf{x}) \leq C \tilde{W}(\mathbf{x})$

for some positive constant C. Furthermore, for some small positive α_0 , the function $\mathbf{z} \mapsto \exp(\alpha_0 \|\mathbf{z}\|)$ is ν -integrable.

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Theorem

Suppose Assumptions 1 through 6 hold. Then $(X_t : t \ge 0)$ is a \mathbb{R}^n_+ -valued, right-continuous, C_b -Feller Markov process with the SSP property.

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Define:

$$\mathsf{Safe}(\varepsilon, R) := \{ \mathbf{x} \in \mathbb{R}_{++}^n : \min x_i > \varepsilon, \|\mathbf{x}\| \le R \},\$$

and let $\mathcal{I}(\mathbf{x})$ be a positive Lipschitz function, bounded by 1, that approximates $\mathbf{1}_{\mathsf{Safe}(\varepsilon,R)}(\mathbf{x})$ from below. Assume that the community is layered and $\nu(d\mathbf{z}) = \lambda \mu(d\mathbf{z})$, with μ the law of a \mathbb{R}^n -valued random variable with compact support contained in $\{z_i > -\gamma, i = 1, \ldots, n\}$, where $\gamma \in (0, 1)$. Assume also that $L_i(\mathbf{x}, \mathbf{z}) = \beta \mathcal{I}(\mathbf{x}) z_i$, where $0 < \beta < 1$, $\varepsilon > 0$ and R > 0. On this setting, the hypotheses 1, 2, 3, 4 and 6 required by Theorem 3 are satisfied.

Example: Lévy-driven, Lotka-Volterra food-chains

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For i = 1, ..., n, let $(N_t^{(i)} : t \ge 0)$ be a family of independent compound Poisson processes on \mathbb{R} with intensity $\tilde{\lambda}_i > 0$ and jump increment distributed according to a law m_i with support contained in $\{z > -1\}$. As before, let \tilde{N}_i be the compensated process. Let $\beta \in (0, 1)$ and $0 < \varepsilon_i < M_i$, i = 1, ..., n be some constants. Let $\ell_i : \mathbb{R}_+ \to [0, 1]$ be a Lipschitz function that equals 1 on $[\varepsilon_i, M_i]$ and vanishing outside the interval $[\varepsilon_i/2, 2M_i]$. Set $L_i(\mathbf{x}, \mathbf{z}) = \ell_i(x_i)\beta z_i$.

The food-chain equations

$$dX_{t}^{(1)} = X_{t}^{(1)} \left((b_{1} - a_{11}X_{t}^{(1)} - a_{12}X_{t}^{(2)})dt + \sigma_{1}dW_{t}^{(1)} + \int_{\{z_{1} > -1\}} L_{1}(\mathbf{X}_{t}, \mathbf{z})\tilde{N}_{1}(dz_{1}, dt) \right),$$

$$dX_{t}^{(i)} = X_{t}^{(i)} \left((-b_{i} + a_{i,i-1}X_{t}^{(i-1)} - a_{i,i}X_{t}^{(i)} - a_{i,i+1}X_{t}^{(i+1)})dt + \sigma_{i}dW_{t}^{(i)} + \int_{\{z_{i} > -1\}} L_{i}(\mathbf{X}_{t}, \mathbf{z})\tilde{N}_{i}(dz_{i}, dt) \right), \quad i = 2, \dots, n-1,$$

$$dX_{t}^{(n)} = X_{t}^{(n)} \left((-b_{n} + a_{n,n-1}X_{t}^{(n-1)} - a_{nn}X_{t}^{(n)})dt + \sigma_{n}dW_{t}^{(n)} + \int_{\{z_{n} > -1\}} L_{n}(\mathbf{X}_{t}, \mathbf{z})\tilde{N}_{n}(dz_{n}, dt) \right).$$

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For $n \ge 2$, set:

$$d_{1} = b_{1} - \frac{1}{2}\sigma_{1}^{2}, \quad d_{i} = b_{i} + \frac{1}{2}\sigma_{i}^{2}; \quad i = 2..., n$$

$$\Delta_{i} = \mathbb{E}_{Z \sim m_{i}}(\beta Z - \ln(1 + \beta Z)) \ge 0; \quad \lambda_{i} = \tilde{\lambda}_{i}\Delta_{i}$$

$$r_{1} := d_{1} - \lambda_{1}, \quad r_{i} := d_{i} + \lambda_{i}, i = 2, ..., n;$$

$$\mathbf{r}_{m} = (-r_{1}, r_{2}, ..., r_{m}), 1 \le m \le n.$$

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Theorem

For $2 \le m \le n$, let \mathbf{A}_m be the leading m-dimensional submatrix of \mathbf{A} . Consider the linear systems:

$$\mathbf{A}_m \mathbf{s} = \mathbf{r}_m, \quad 2 \le m \le n$$

and let $s^{(*,m)} \in \mathbb{R}^m$ be the unique solution of the m-th system . Assume that:

$$a_{n,n-1}s_{n-1}^{(*,n-1)} > r_n.$$
 (5)

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holds. Then the SDE (4) is SSP.

Remarks:

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The above result extends a Theorem of Hening and Nguyen ([HN18c]), which in turn extends a result of Gard ([GH79]) on deterministic LV food-chains.

Appendix: sketch of the proof

Steps toward the proof

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• To prove that there exists an **embedded chain** with tight laws in \mathbb{R}_{++}^n (uniformly for initial conditions on compact sets). A sufficient condition is:

Proposition

For some continuous log-Lyapunov function \tilde{V} , for some $T^* > 0$, $m \in (0, 1)$ and C > 0, the inequality:

$$\mathbb{E}_{\mathbf{x}}(\tilde{V}(\mathbf{X}_{T^*})) \leq m\tilde{V}(\mathbf{x}) + C,$$

holds for every $\mathbf{x} \in \mathbb{R}^n_{++}$

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• This is the hard part.

We prove:

Lemma

For every T > 0 and $\mathbf{X}_0 \in \mathbb{R}^n_{++}$, under $\mathbb{P}_{\mathbf{X}_0}$ the Markov chain $\mathbf{X}^T := (\mathbf{X}_{mT} : m \ge 0)$ is irreducible and aperiodic. Furthermore, every compact set is a petite set for the chain.

By Lemma 5 and Theorem 6.3 of [MT92], there exists a probability measure π such that:

$$\lim_{n\to\infty} \|\mathbb{P}_{\mathbf{x}}(\mathbf{X}_{nT^*}\in\cdot)-\pi(\cdot))\|_{TV}=0,$$

and the convergence is geometric.

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and the convergence is geometric. Then, \mathbf{X}^{T*} is positive Harris recurrent, and thus this chain has hitting times (of open sets) with finite expectation; this in turn also implies that \mathbf{X} is positive Harris recurrent (see theorem 1 of [KM94]), and thus has a unique invariant probability measure in \mathbb{R}^{n}_{++} . Of course, this probability measure is just π .

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