

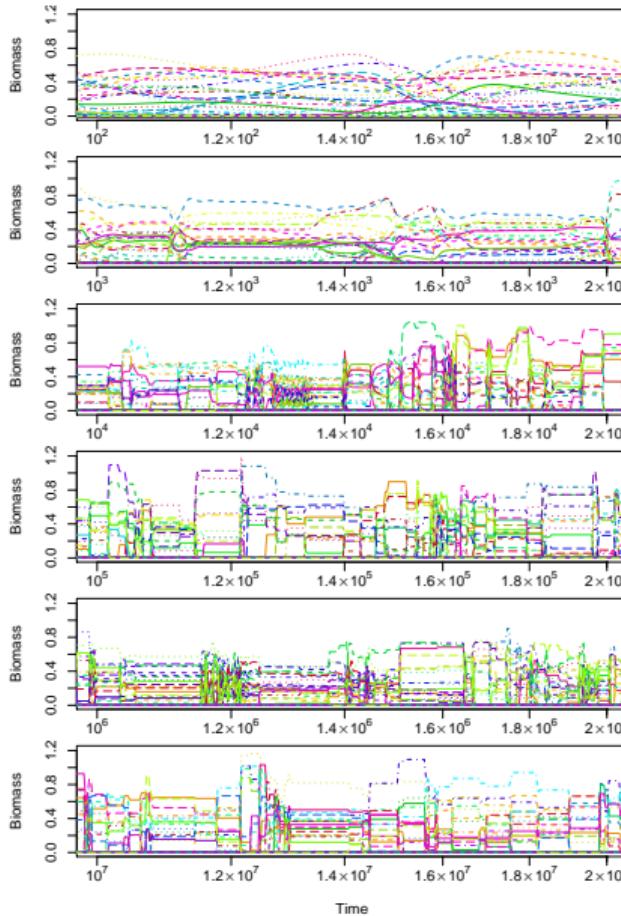
More Recent Evidence for Self-Organised Ecological Structural Instability in Nature



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Queen Mary University of London

Paris, 29 October 2024



Simulation of generic behaviour of random competitive
LV models deeply inside the MA phase:
heteroclinic networks
(Hofbauer 1994, *Tatra Mountains Math. Publ.*).

$$\frac{dB_i}{dt} = \left[r_i - \sum_j^S A_{ij} B_j \right] B_i \quad (1 \leq i \leq S)$$

$$S = 300$$

$$r_i = \mathcal{N}(1, 0.1)$$

$$A_{ij} = 0.5 \cdot \text{Bernoulli}(0.5) \quad (i \neq j)$$

$$A_{ii} = 1$$

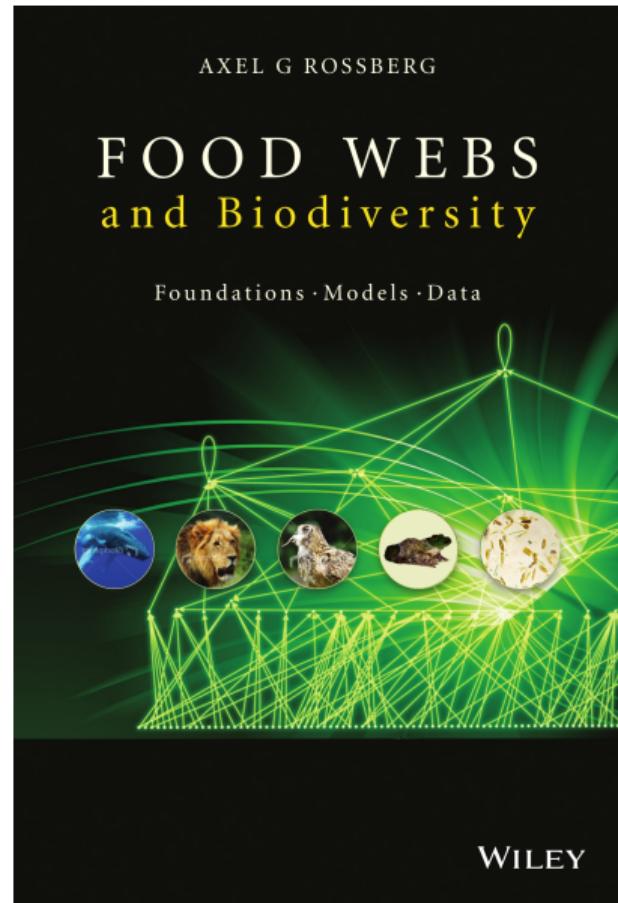
$$B_i = 0.001 \text{ at } t = 0$$

O'Sullivan, Terry, and Rossberg 2021, *Nat Commun*

First evidence that
(ecological) structural instability
controls natural community structure.

Wiley, 2013

PROSE award Biological Sciences 2013



Ecological Structural Instability

Consider the Lotka-Volterra competition model:

$$\frac{dB_j}{dt} = \left(1 - \sum_k^S G_{jk} B_k\right) B_j$$

G_{jk} : Competition (overlap) matrix

S : Species richness

- $G_{jj} = 1$
- G_{jk} = random i.i.d., mean μ , variance σ^2 ($j \neq k$).

What is the equilibrium distribution of biomasses B_j ?

Ecological structural instability as amplification of indirect interactions

Equilibrium of population B_1 :

$$\text{population growth rate} = 0 = 1 - \sum_k G_{jk} B_k \quad (1 \leq j \leq S)$$

For $j = 1$:

$$(1 - \mu)^2 \text{var } B_1 \approx S\sigma^2 \left[(\text{EB}_{\text{other}})^2 + \text{var } B_{\text{other}} \right]$$

$$(1 - \mu)^2 \text{var } B_1 \approx S\sigma^2 \left[(\text{EB}_1)^2 + \text{var } B_1 \right]$$

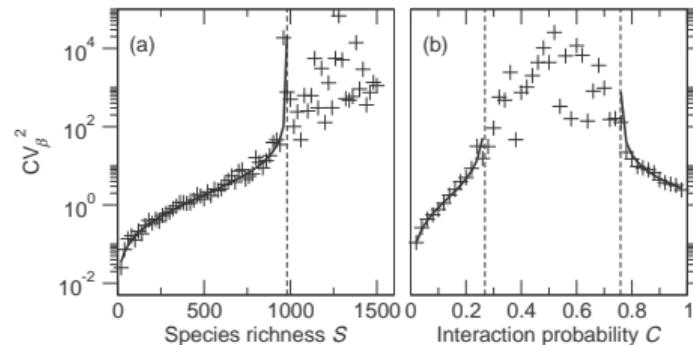
Solve for $\text{var } B_1$:

$$\text{var } B_1 = \dots,$$

$$CV_B^2 = \frac{S\sigma^2}{(1 - \mu)^2 - S\sigma^2}.$$

Rossberg 2013, *Food Webs and Biodiversity*

→ Feedback amplification of $\text{var } B_i$ through $S\sigma^2 \implies$ singularity at $S = S_{\text{ESI}} = \frac{(1-\mu)^2}{\sigma^2}$!



History of self-consistency/mean-field/disordered-systems method

Diederich and Opper 1989, <i>Phys. Rev. A</i>	replicator model
Rieger 1989, <i>J. Phys. A: Math. Gen.</i>	various models
Tokita 2004, <i>Phys. Rev. Lett.</i>	replicator, various interaction patterns
Tokita 2006, <i>Ecol. Inform.</i>	symmetric replicator
Yoshino, Galla, and Tokita 2007, <i>J. Stat. Mech.</i> , Yoshino, Galla, and Tokita 2008, <i>Phys. Rev. E</i>	general replicator
Rossberg 2013, <i>Food Webs and Biodiversity</i>	LV assembly model, it's real!
Bunin 2017, <i>Phys. Rev. E</i>	general LV model
Tikhonov and Monasson 2017, <i>Phys. Rev. Lett.</i>	resource competition
Advani, Bunin, and Mehta 2018, <i>J. Stat. Mech.</i>	consumer-resource model
Dougoud et al. 2018, <i>PLoS Comput. Biol.</i>	strict proof
Pettersson, Savage, and Jacobi 2020, <i>Phys. Rev. E</i>	using von Neumann expansion

Assembly model!

The Lotka-Volterra competition model:

$$\frac{dB_j}{dt} = \left(1 - \sum_k^S G_{jk} B_k\right) B_j$$

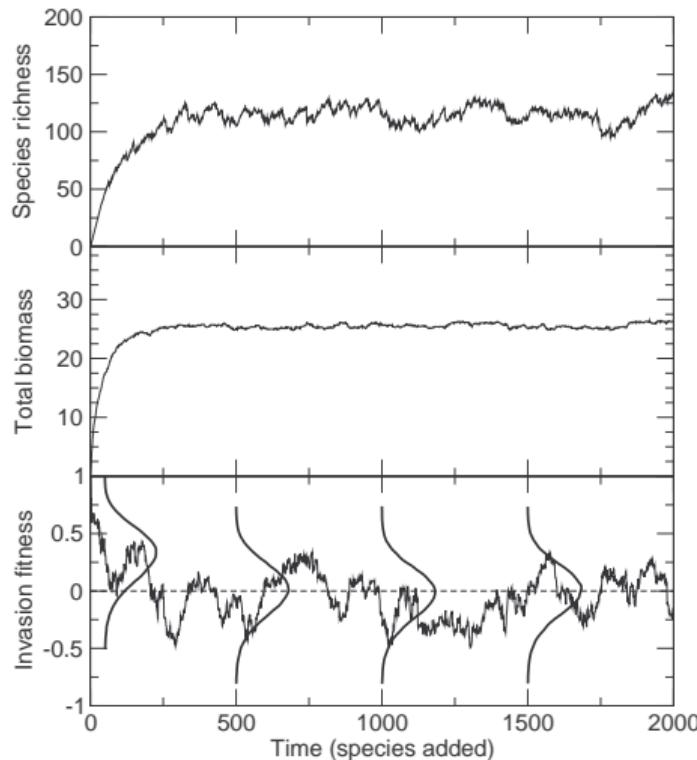
G_{jk} : Competition (overlap) matrix
 S : Species richness

Here

- $G_{jj} = 1$
- G_{jk} = random i.i.d., mean μ , variance σ^2 ($j \neq k$).
- Add species one-by-one, simulate in between, remove those going extinct.

Gamarra et al. 2005, *Biological Invasions*

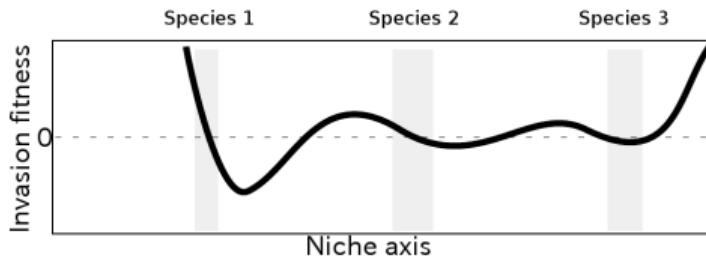
Community saturation in assembly models



- Separate saturation of richness and biomass
- Community remains open to invasion
- Invasion fitness of 'test species' fluctuates

Solve this model?

Solving the model I (harvesting resistance)



Harvesting resistance of a species:

$$h \stackrel{\text{def}}{=} - \left(\frac{d \ln B(H)}{dH} \Big|_{H=0} \right)^{-1}$$

where

H : Harvesting rate [1/Time]

$B(H)$: Equilibrium biomass of harvested species

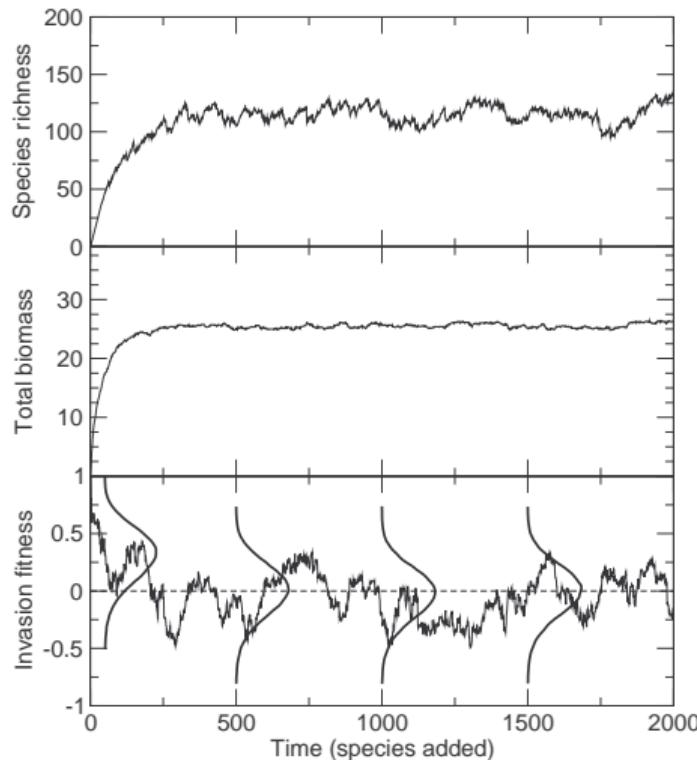
Important property:

$$(h \text{ after invasion}) \approx (\text{invasion fitness before invasion})$$

with '=' for Lotka-Volterra systems and close to extinction.

Furthermore, $h_j \approx \text{const. } B_j$ and $h_j \rightarrow 0$ at extinction.

Community saturation in assembly models



Now, consider interactions between:

- Fluctuations of invasion fitness
- Fluctuations of harvesting resistance
- Species turnover

Rossberg 2013, *Food Webs and Biodiversity*

Solving the model II (self-consistent assembly equilibrium)

Premises:

- Ecological community is in invasion-extinction quasi-equilibrium (Pawar 2009, *J. Theor. Biol.*) with S species on average.'
- Invasions and extinctions occur at rate 1 (defines time unit).
- As the community turns over, the invasion fitness r_i of a given potential invader i follows an Ornstein-Uhlenbeck process, i.e. a mean-reverting random walk, with identical parameters for all i .
- Harvesting resistance of residents j follows the same process, but with $h_j = r_i > 0$ at time of invasion, and extinction if $h_j = 0$.
- ~~Reversion rate of r_i is $1/S$.~~

Solve this self-consistency condition: the reversion rate (ρ) of r_i for potential invaders equals that of h_j for residents, **while taking species turnover into account**.

Some of the maths...

We now evaluate the relaxation rate of R_i using moment equations, denoting expectation values by brackets $\langle \cdot \rangle$. With our assumption of negligible correlations, suppressing the index i and abbreviating $Z = |\mathcal{A}|$, we can write $\langle R \rangle = \langle Z \rangle \langle y \rangle$, where the second factor is the expectation of y_i for a randomly chosen extant species. According to above considerations, $\langle R \rangle$ value changes over a small time interval $\delta T \geq 0$ to:

$$\begin{aligned} \langle R(T + \delta T) \rangle &= \left\langle \sum_{j \in \mathcal{A}} y_j - \delta T \rho(y_j - \bar{y}) + \delta T C y^{\text{inv}} \right\rangle \quad (9) \\ &= \langle Z \rangle \langle y \rangle + [-\rho \langle Z \rangle (\langle y \rangle - \bar{y}) + C y^{\text{inv}}] \delta T, \end{aligned}$$

where y^{inv} denotes the mean associated with $P^{\text{inv}}(y)$. The sum above is over the extant species at time T . Extinctions of species can be disregarded at lowest order, because shortly before extinction their contribution to R is small. In the steady state the time-dependent term in Eq. (9) vanishes:

$$-\rho(\langle y \rangle - \bar{y}) \langle Z \rangle + C y^{\text{inv}} = 0. \quad (10)$$

Making use of Eq. (10), we similarly derive in Appendix the second moment

$$\begin{aligned} \langle R(T)R(T + \delta T) \rangle &= \\ &= \langle Z \rangle^2 \langle y \rangle^2 + \langle Z \rangle \langle y^2 \rangle + \langle Z \rangle \rho (\langle y \rangle \bar{y} - \langle y^2 \rangle) \delta T \quad (11) \end{aligned}$$

Derivation of Eq. 11

We derive Eq. (11), making first use of Eq. (10) and then of the fact that $\text{var } Z = \langle Z^2 \rangle - \langle Z \rangle^2 = \langle Z \rangle$ on account of the Poisson distribution of Z :

$$\begin{aligned} &\langle R(T)R(T + \delta T) \rangle \\ &= \left\langle \sum_{j,l \in \mathcal{A}} y_j(T)y_l(T + \delta T) + \sum_{j \in \mathcal{A}} y_j C y^{\text{inv}} \delta T \right\rangle \\ &= \left\langle \sum_{j,l \in \mathcal{A}} y_j [y_l - \rho(y_l - \bar{y})\delta T] \right\rangle + \langle y \rangle \langle Z \rangle C y^{\text{inv}} \delta T \\ &= \left\langle \sum_{\substack{j,l \in \mathcal{A} \\ j \neq l}} y_j [y_l - \rho(y_l - \bar{y})\delta T] \right\rangle \\ &\quad + \left\langle \sum_{j \in \mathcal{A}} y_j [y_j - \rho(y_j - \bar{y})\delta T] \right\rangle \\ &\quad + \langle y \rangle \langle Z \rangle C y^{\text{inv}} \delta T \\ &= \langle Z(Z-1) \rangle [\langle y \rangle^2 - \rho(\langle y \rangle^2 - \langle y \rangle \bar{y}) \delta T] \\ &\quad + \langle Z \rangle [\langle y^2 \rangle - \rho(\langle y^2 \rangle - \langle y \rangle \bar{y}) \delta T] \\ &\quad + \langle y \rangle \langle Z \rangle \rho(\langle y \rangle - \bar{y}) \langle Z \rangle \delta T \\ &= \langle Z \rangle^2 \langle y \rangle^2 + \langle Z \rangle \langle y^2 \rangle \\ &\quad + (\langle Z \rangle^2 - \langle Z^2 \rangle) \rho(\langle y \rangle^2 - \langle y \rangle \bar{y}) \delta T \\ &\quad + \langle Z \rangle \rho(\langle y \rangle^2 - \langle y^2 \rangle) \delta T \\ &= \langle Z \rangle^2 \langle y \rangle^2 + \langle Z \rangle \langle y^2 \rangle \\ &\quad - \langle Z \rangle \rho(\langle y \rangle^2 - \langle y \rangle \bar{y}) \delta T \\ &\quad + \langle Z \rangle \rho(\langle y \rangle^2 - \langle y^2 \rangle) \delta T \\ &= \langle Z \rangle^2 \langle y \rangle^2 + \langle Z \rangle \langle y^2 \rangle \\ &\quad + \langle Z \rangle \rho(\langle y \rangle \bar{y} - \langle y^2 \rangle) \delta T. \end{aligned}$$

Combining the moment equations, we first evaluate

$$\text{var } R = \langle R^2 \rangle - \langle R \rangle^2 = \langle Z \rangle \langle y^2 \rangle. \quad (12)$$

Then we calculate the short-term autocorrelation function

$$\text{cor}[R(T), R(T + \delta T)] = \frac{\text{cov}[R(T), R(T + \delta T)]}{\text{var } R} \quad (13)$$

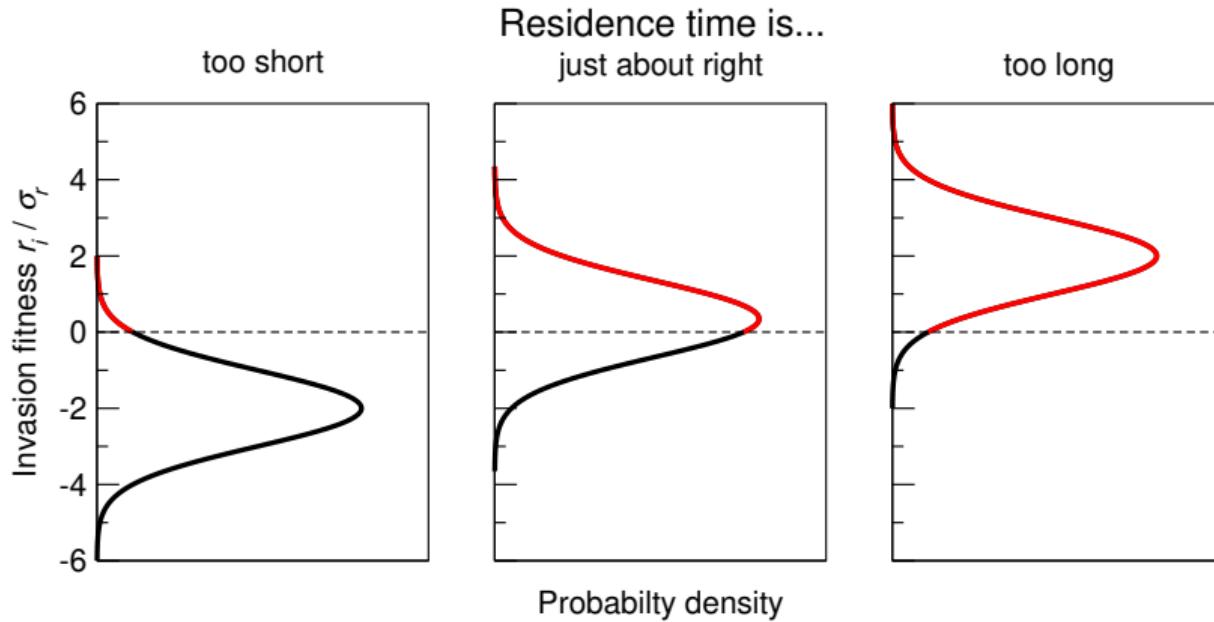
and from this, considering that for an Ornstein Uhlenbeck process $\text{cor}[R(T), R(T + \delta T)] = \exp(-\tilde{\rho}|\delta T|)$, the relaxation rate of R as:

$$\begin{aligned} \tilde{\rho} &= -\lim_{\delta T \rightarrow 0^+} \frac{d \text{cor}[R(T), R(T + \delta T)]}{d \delta T} \\ &= -\frac{\langle Z \rangle \rho (\langle y \rangle \bar{y} - \langle y^2 \rangle)}{\langle Z \rangle \langle y^2 \rangle} = -\frac{\rho (\langle y \rangle \bar{y} - \langle y^2 \rangle)}{\langle y^2 \rangle}. \end{aligned} \quad (14)$$

Equating $\tilde{\rho}$ and ρ in Eq. (14) yields our self-consistency condition. It simplifies to

$$\bar{y} = 0, \quad \text{and so} \quad p_{\text{inv}} = \Phi(\bar{y}) = \frac{1}{2}, \quad (15)$$

Invasion probability (intuition)



- Previously: $\bar{r}/\sigma_r \approx 0.347$, invasion probability is 0.635.

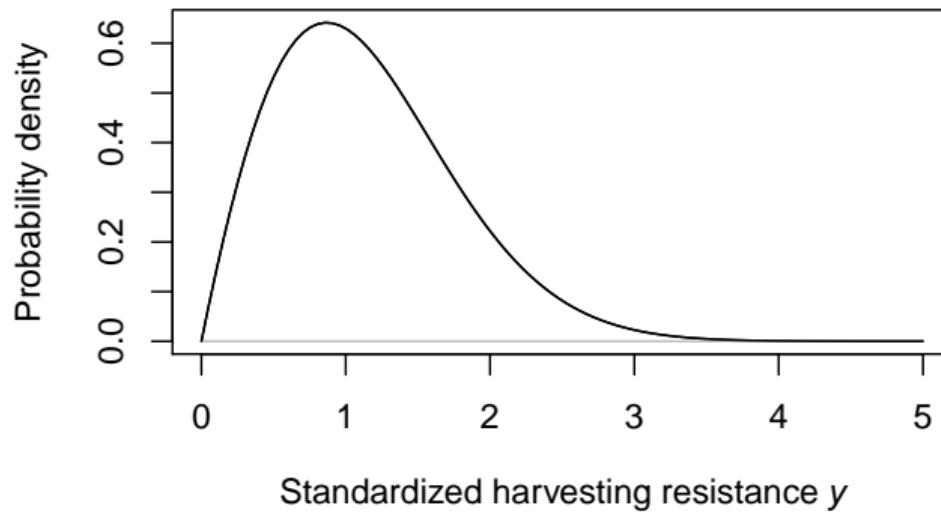
Rossberg 2013, *Food Webs and Biodiversity*

- New: $\bar{r} = 0$, invasion probability is 1/2.

Cockrell et al. 2024, *Phys. Rev. Res.*

Solving the model III (distribution of scaled harvesting resistance)

$$0 = -\rho \underbrace{\frac{d[(0-y)P(y)]}{dy}}_{\text{mean reversion}} + \rho \underbrace{\frac{d^2P(y)}{dy^2}}_{\text{diffusion}} + \sqrt{\frac{2}{\pi}} \exp\left(-\frac{y^2}{2}\right).$$



Solving the model IV (self-consistent community response)

To get S and the reversion rate, match parameters of Ornstein-Uhlenbeck process with elementary description of community-response to invasion. (Rossberg 2013, *Food Webs and Biodiversity*)

Compute the resident's response ΔB_k to an invader with biomass B_{inv} and interaction strengths γ_j from:

$$0 = \gamma_j B_{\text{inv}} + \sum_k G_{jk} \Delta B_k \quad (1 \leq j \leq S).$$

Problem is very similar to finding the equilibrium:

$$0 = 1 - \sum_k G_{jk} B_k \quad (1 \leq j \leq S).$$

This is where ESI enters the calculation. Negative ΔB_k are natural!

Analytic solution of species-rich LV assembly model

Let as above

μ = mean interspecific interaction,
 σ^2 = variance of interspecific interactions.

In our model the singularity of Ecological Structural Instability occurs at richness

$$S_{\text{ESI}} \stackrel{\text{def}}{=} \frac{(1 - \mu)^2}{\sigma^2}.$$

Then, with

$$\langle y \rangle = 2^{1/2} \pi^{-1/2} \ln(2)^{-1} = 1.1511$$

and

$$\langle y^2 \rangle = 1 + \ln(4)^{-1} = 1.7213,$$

in the saturated LV assembly model,
species richness is

$$\langle S \rangle = \frac{S_{\text{ESI}}}{\langle y^2 \rangle} \approx 0.58 S_{\text{ESI}},$$

standard deviation of invasion fitness

$$\text{SD}_r = \frac{\langle y^2 \rangle}{\langle y \rangle} \cdot \frac{\sigma^2}{\mu(1 - \mu)},$$

and **relaxation rate** of a species' invasion fitness

$$\rho = \frac{\ln 2}{\langle S \rangle} \text{ invasions.}$$

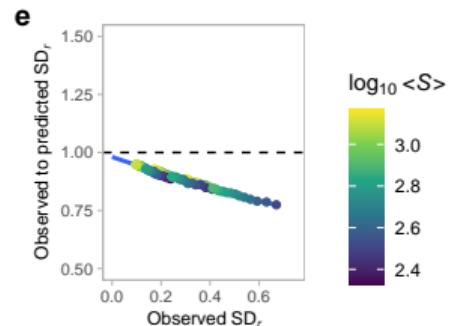
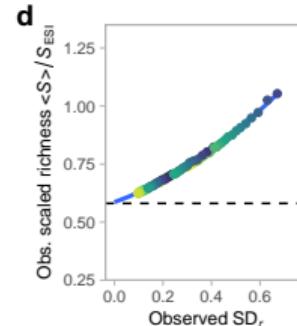
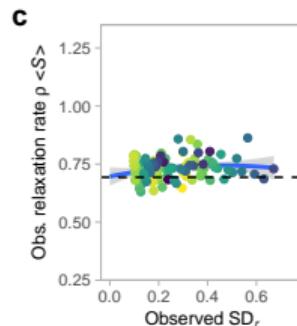
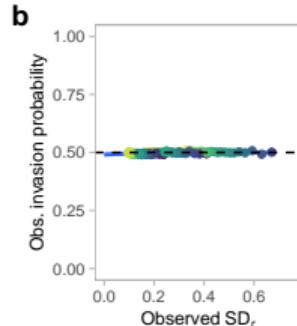
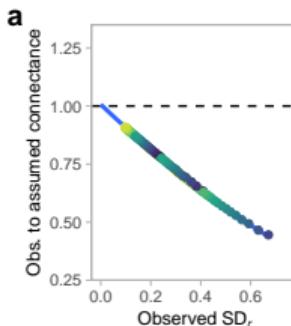
Invasion probability is $1/2$.

Cockrell et al. 2024, *Phys. Rev. Res.*

Numerical test of analytic solution

— — —: universal analytic results

Coloured: simulations



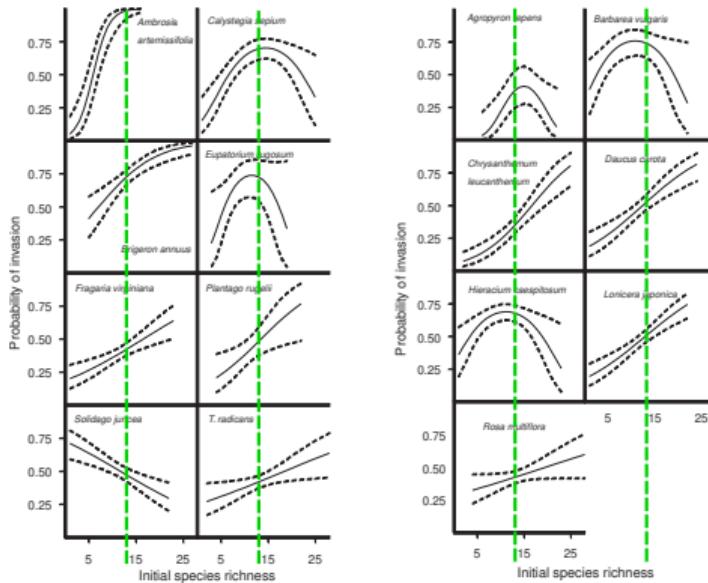
← Limit of large and complex communities

Cockrell et al. 2024, *Phys. Rev. Res.*

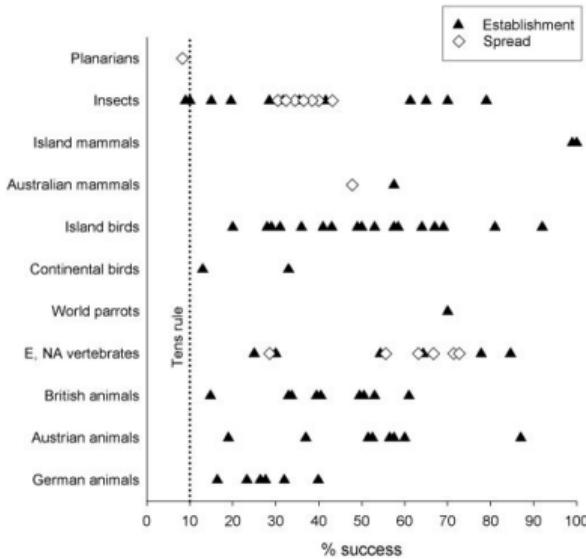
Empirical support for model predictions

Empirical invasion probabilities (1)

Plants in old fields



Europe & N.-America



"Successional communities are model systems for the regulation of community invasibility because they are characterized by continuous species invasions, [...] While the proportion of exotic species has decreased over time, and field scale richness increased, species richness per plot has remained near 13 species per plot [...]." (Meiners, Cadenasso, and Pickett 2004, *Ecol. Lett.*)
"...we extract a median of 51% and an average of 56 ± 5%" (Cockrell et al. 2024, *Phys. Rev. Res.*)

"Mean establishment success [was] $59.6 \pm 11.6\%$ for introductions from Europe to North America and $52.4 \pm 11.9\%$ for the opposite direction [...]" meta-analysis by Jeschke and Strayer 2005, *PNAS*

Empirical invasion probabilities (2)

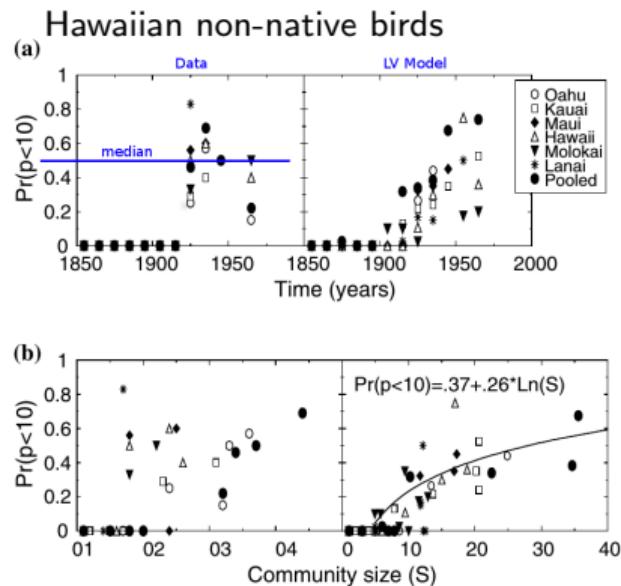
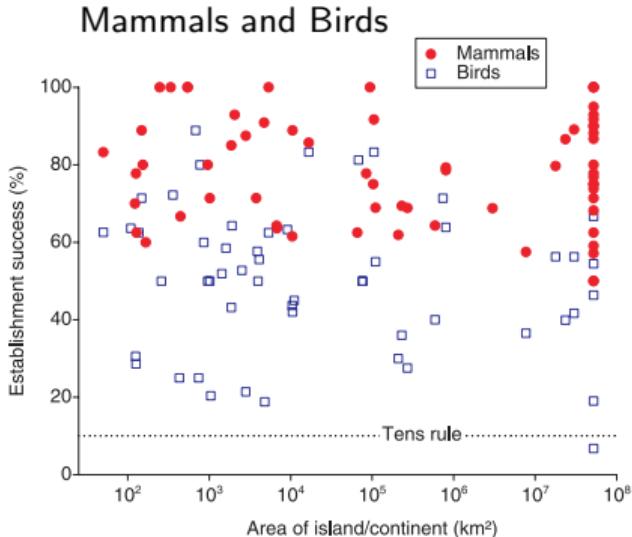


Figure 4. Species-specific probabilities of persisting less than 10 years in the islands as a function of (a) time, and (b) community size. Left: original field data. Right: LVM simulations. Observe the presence of marked thresholds in both analysis and the asymptotic behavior of these probabilities in islands with higher number of species.

(Gamarra et al. 2005, *Biological Invasions*)

Rossberg et al. (Queen Mary U London)



"Establishment success was generally higher for mammals ($79 \pm 1.7\%$, mean \pm standard error (SE), $n = 65$) than for birds ($50 \pm 2.6\%$, $n = 53$) [...]"

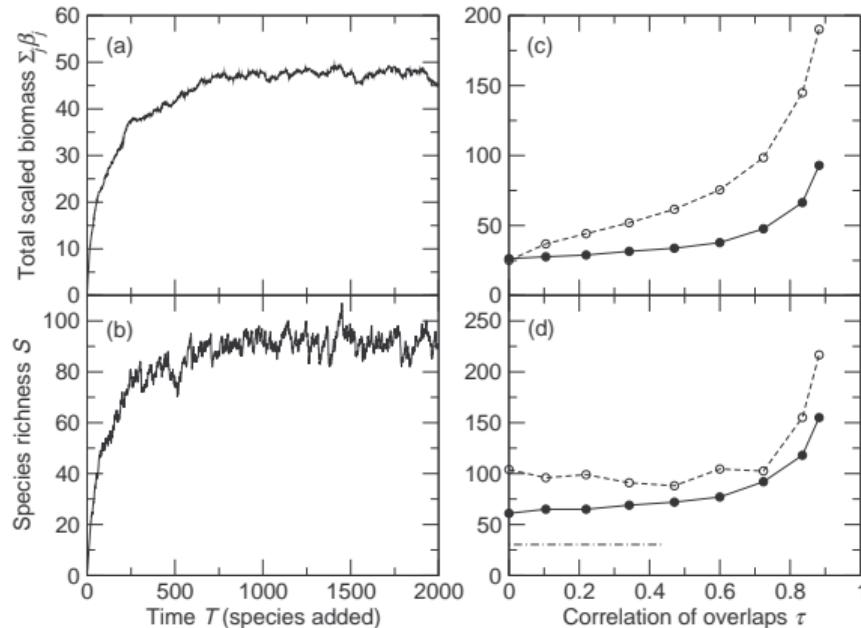
(Jeschke 2008, *Divers. Distrib.*)

Evidence for ESI in Nature

Paris, 29 October 2024

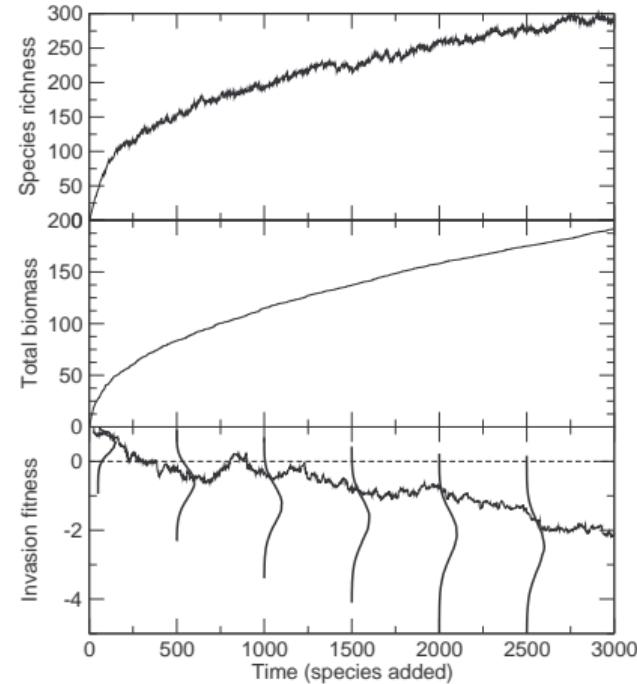
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Extension to partially symmetric interactions remains incomplete:



- : Analytic upper bounds
- : Numerical result

Increasing “competition avoidance” with increasing symmetry.
Singularity for perfect symmetry:



Dimensionless constants of nature in Physics:

For example the *dimensionless magnetic moment* of the electron:

$$g_e = 2 \quad (1928, \text{ experiment})$$

$$g_e = 2 \quad (1928, \text{ theory: Dirac equation})$$

$$g_e = 2.00236 \quad (1948, \text{ experiment})$$

$$g_e = 2.00232 \quad (1948, \text{ theory: quantum electrodynamics})$$

$$g_e = 2.0023318416(13) \quad (2006, \text{ experiment})$$

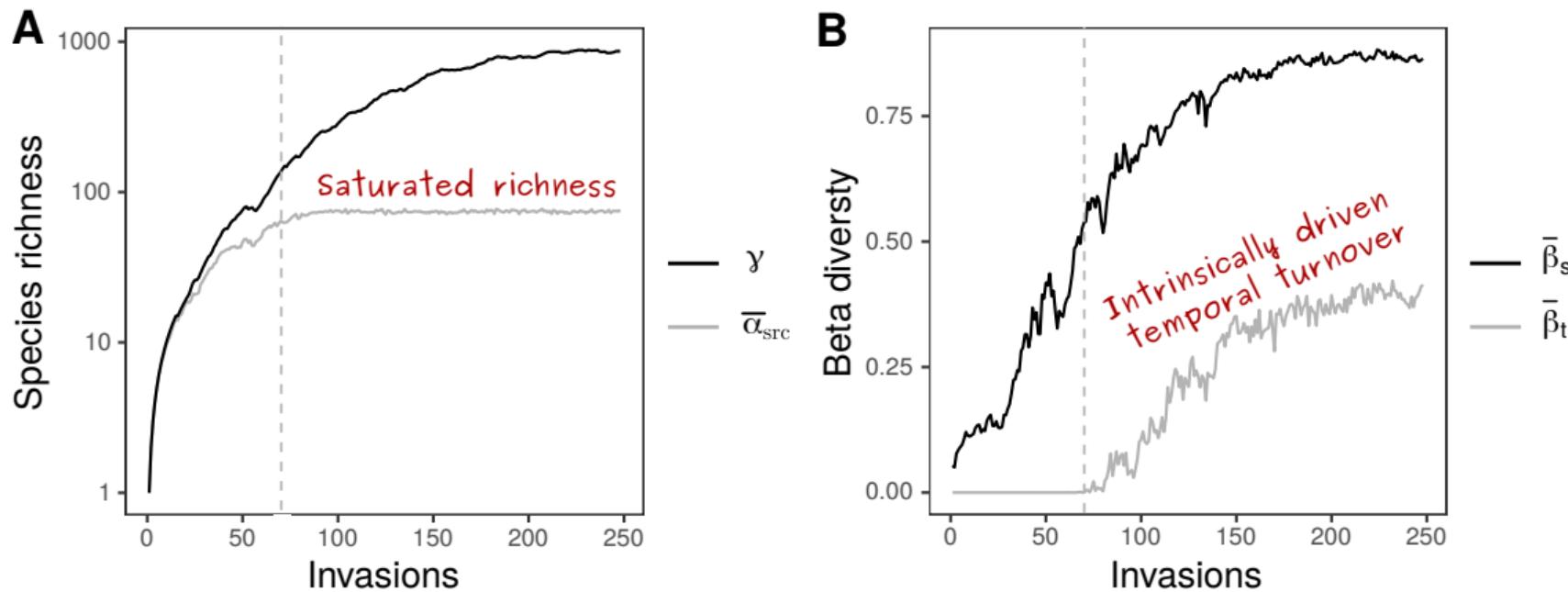
$$g_e = 2.0023318361(10) \quad (2007, \text{ theory: standard model})$$

→ New physics?

Ecological Structural Instability in Metacommunities

Species turnover caused by ecological structural instability

Simulations of Lotka-Volterra metacommunity model



O'Sullivan, Terry, and Rossberg 2021, *Nat Commun*

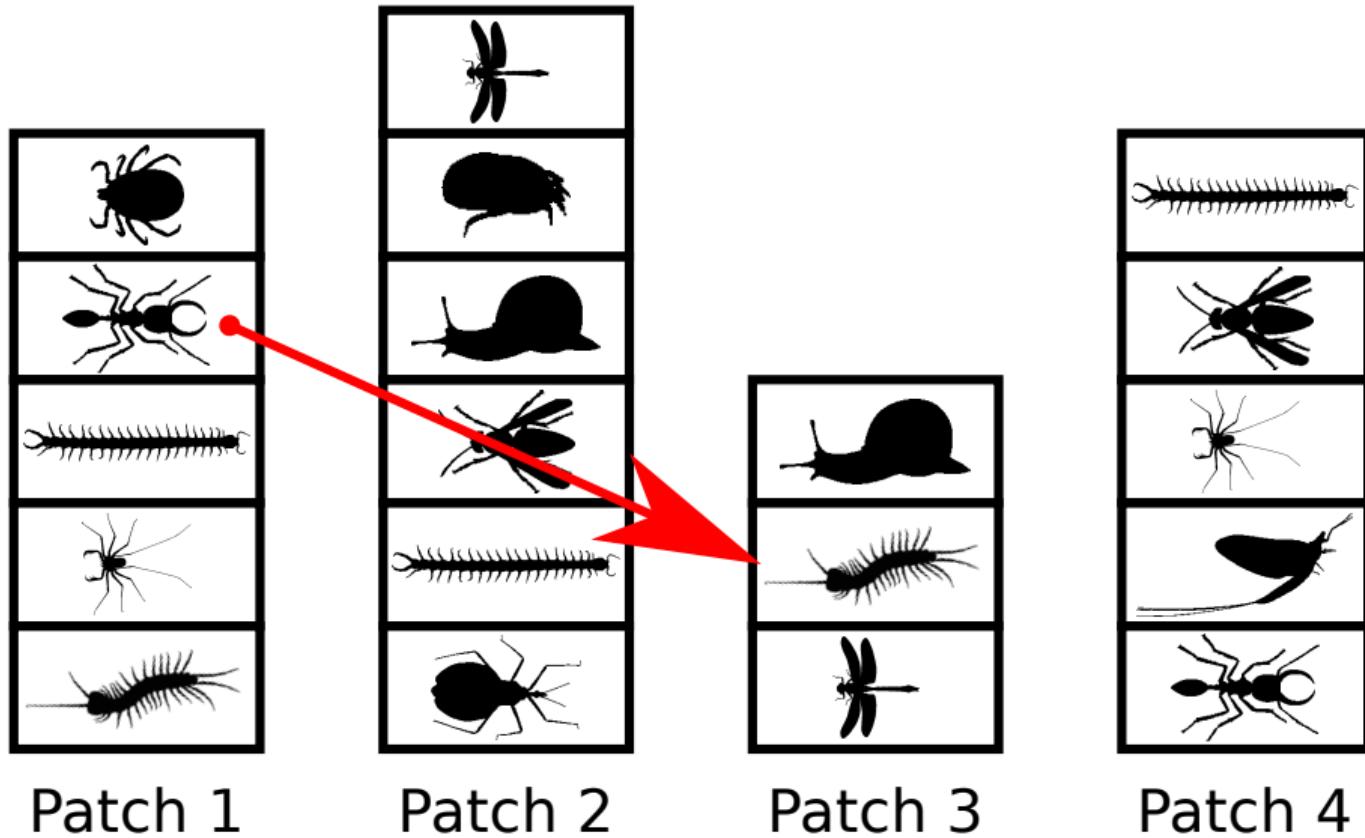
Assemblage Time Series Reveal Biodiversity Change but Not Systematic Loss

Maria Dornelas,^{1,*} Nicholas J. Gotelli,² Brian McGill,³ Hideyasu Shimadzu,^{1,4} Faye Moyes,¹
Caya Sievers,¹ Anne E. Magurran¹

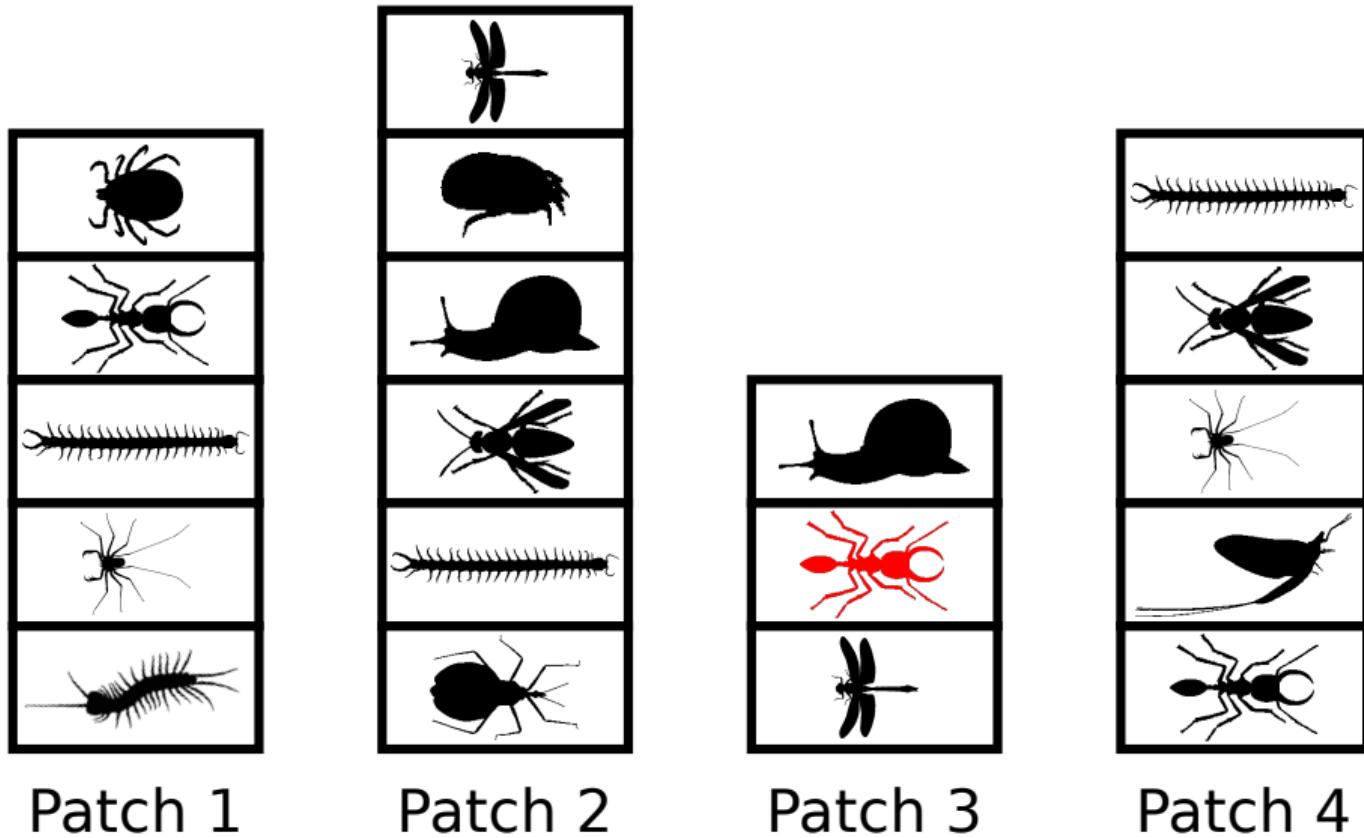
Dornelas et al. 2014, *Science*

The Locally Saturated Patch Occupancy Model

The LSPOM (Locally Saturated Patch Occupancy Model)

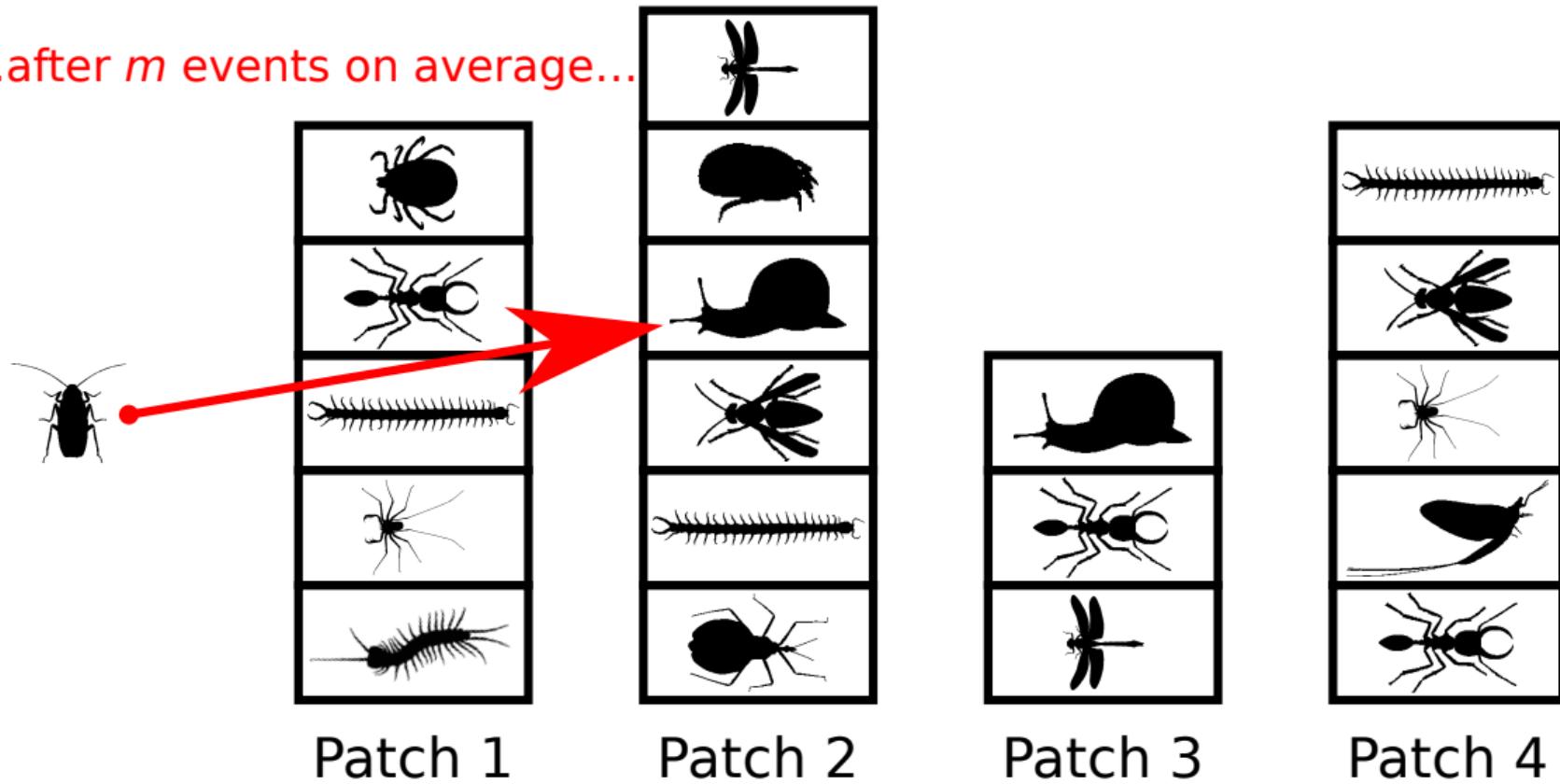


The LSPOM (Locally Saturated Patch Occupancy Model)

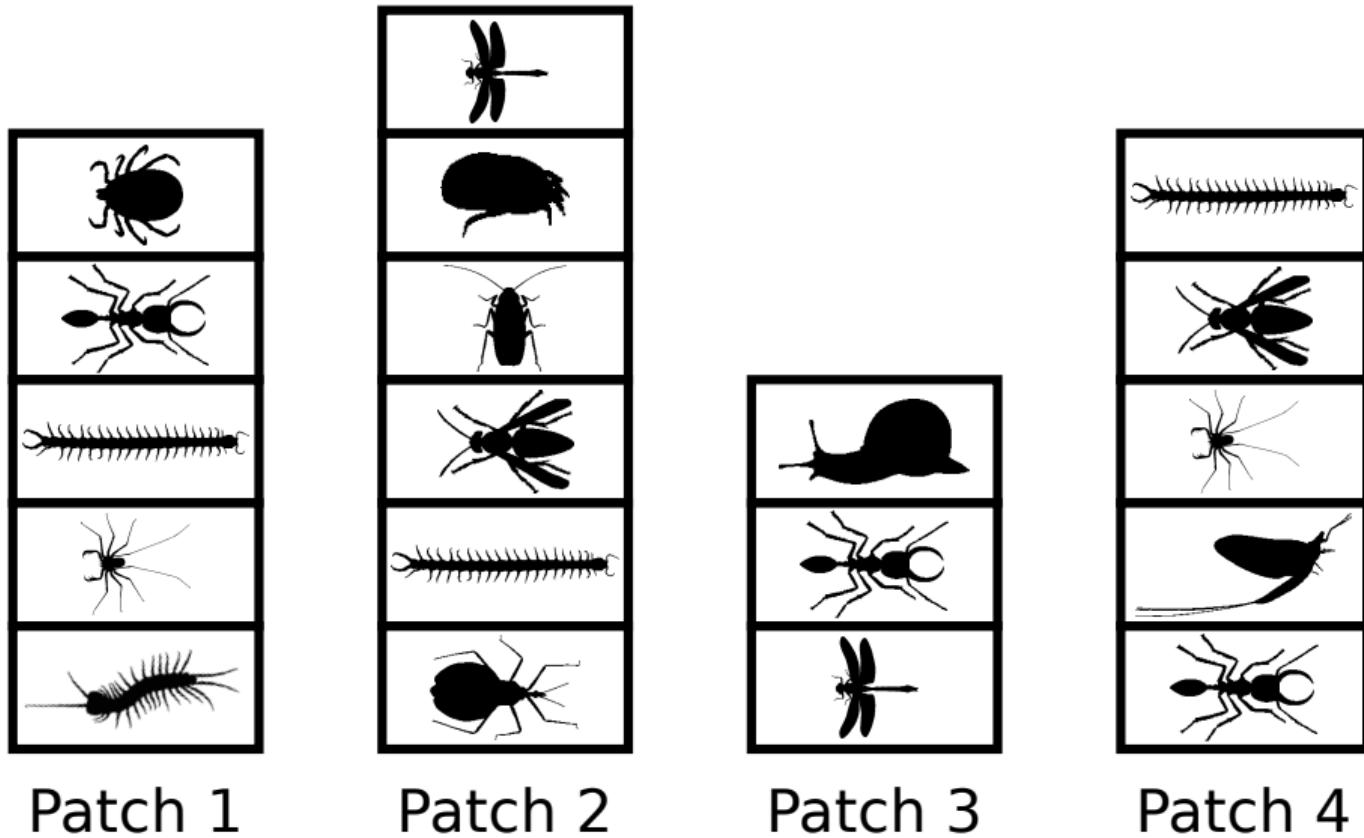


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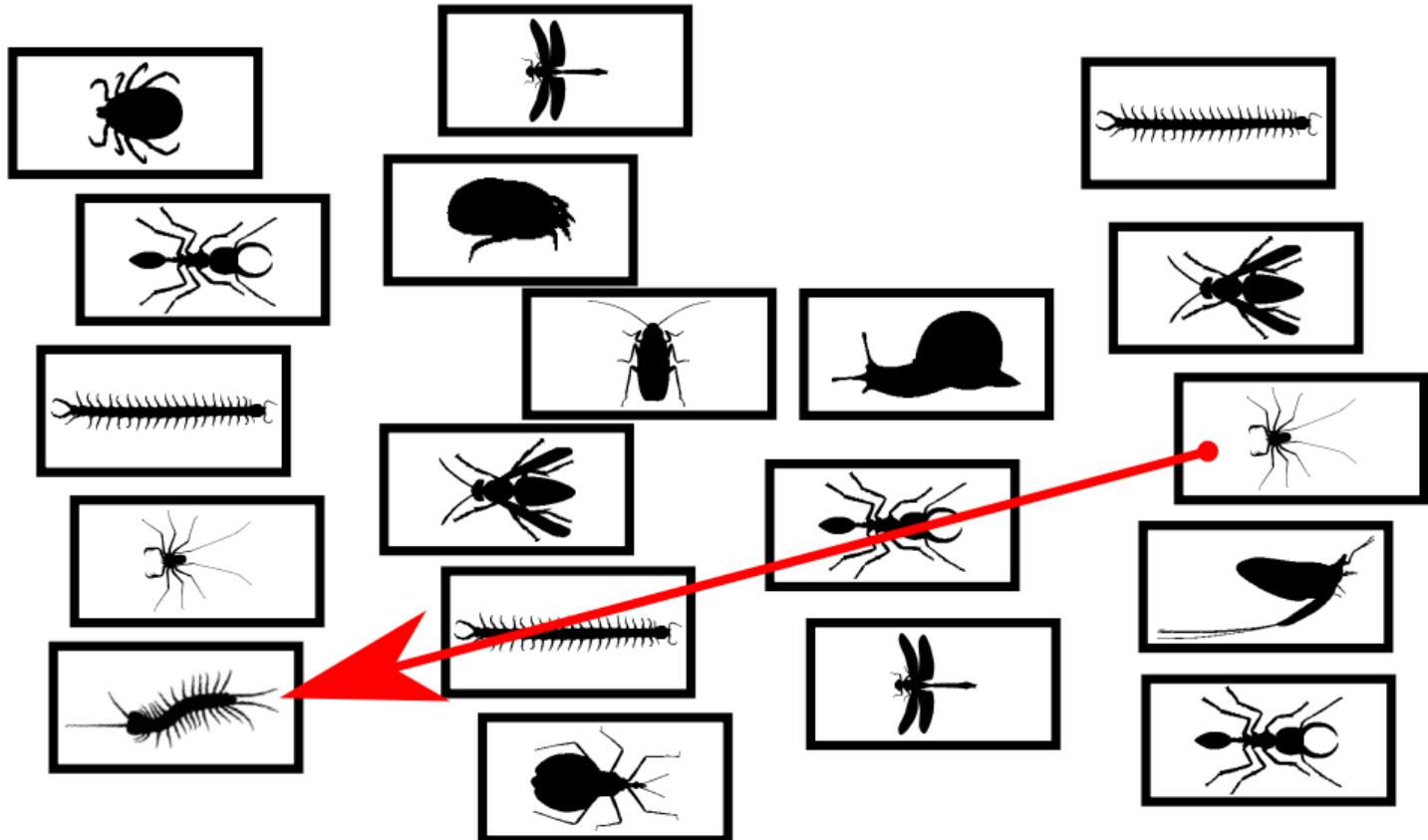
...after m events on average...



The LSPOM (Locally Saturated Patch Occupancy Model)



The LSPOM (Locally Saturated Patch Occupancy Model)



Macroecology: Occupancy Frequency Distributions

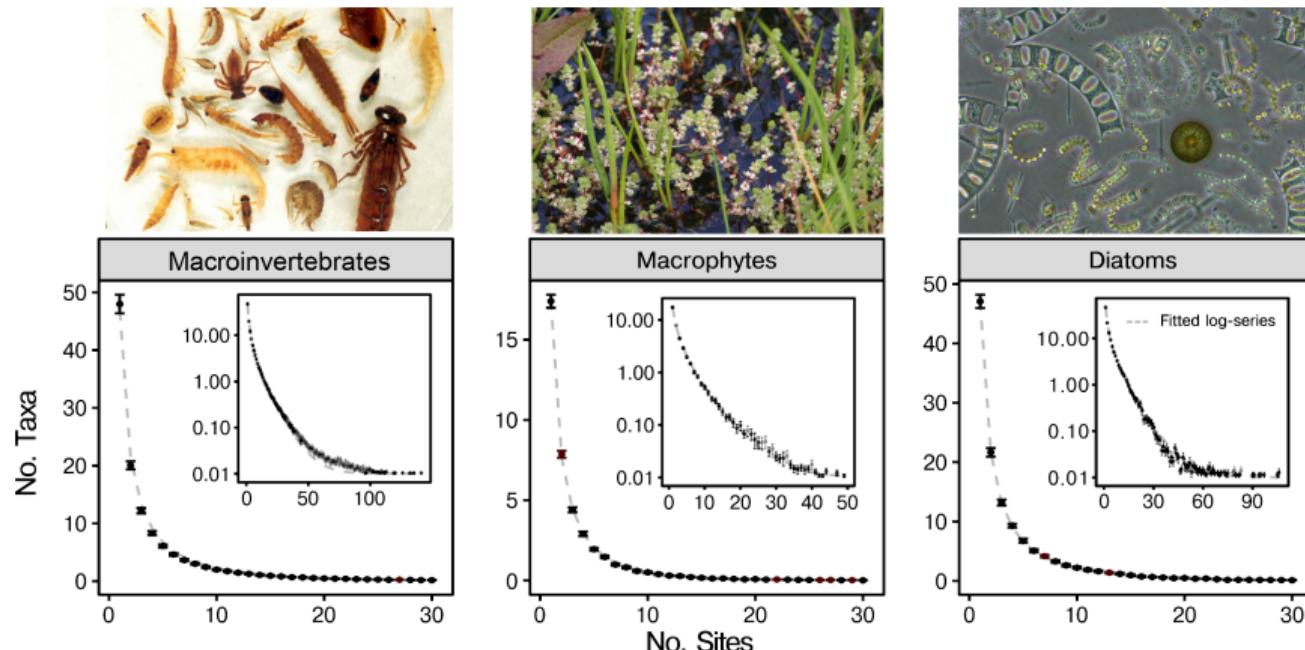
Log-series OFD:

$$(\text{number of taxa occupying } n \text{ sites}) \propto \frac{1}{n} \left(\frac{m}{m+1} \right)^n$$

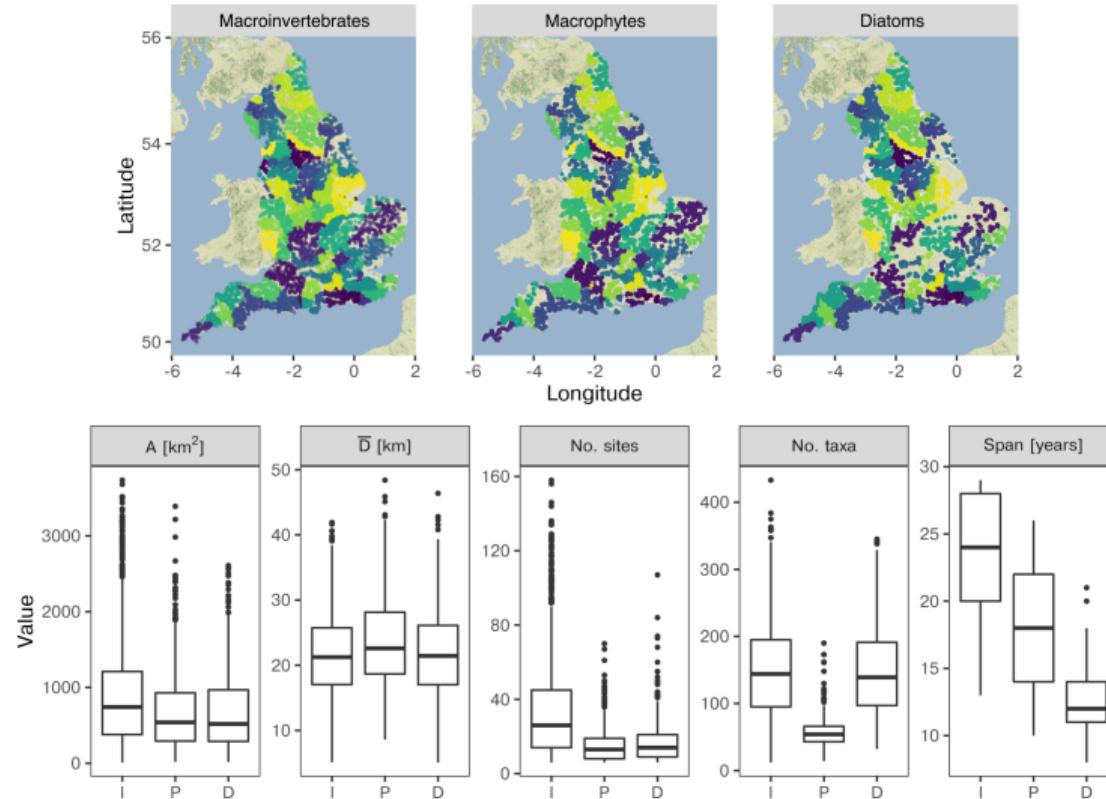
n : number of sites

m : mean number of colonisations per invasion

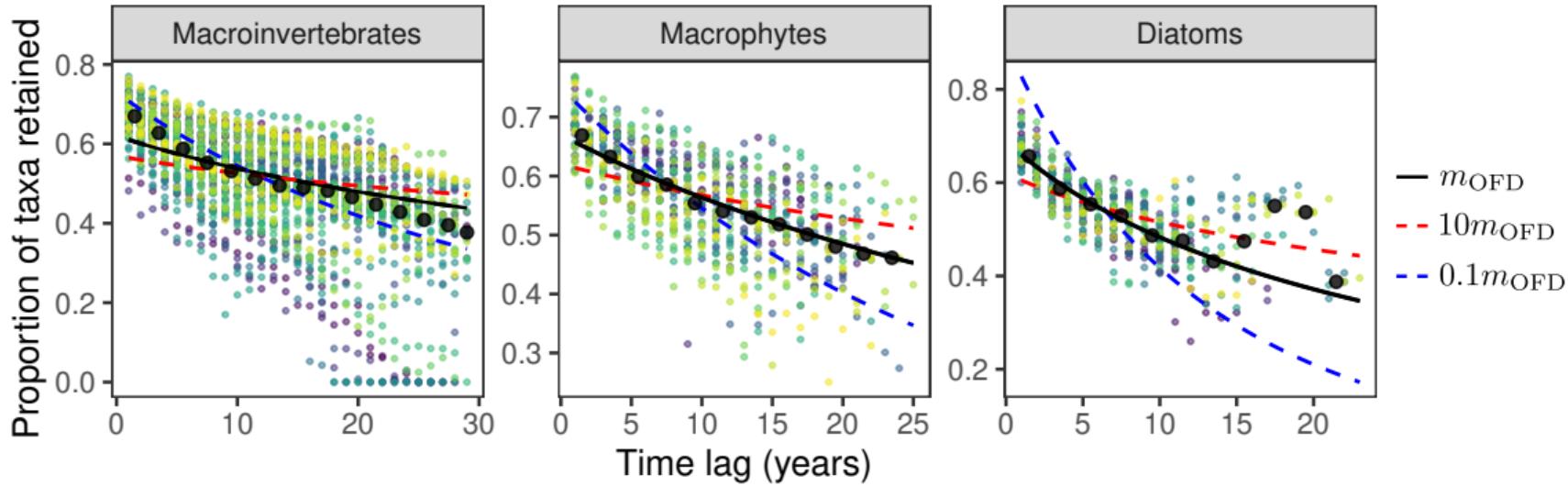
O'Sullivan, Terry, and Rossberg 2023, *Glob. Ecol. Biogeogr.*



Environmental Agency river surveys (Biosys)



Predicting Metacommunity-scale Turnover



Not bad!

O'Sullivan, Terry, and Rossberg 2023, *Glob. Ecol. Biogeogr.*
Pigolotti et al. 2005, *Proc. Natl. Acad. Sci.*

Take home messages

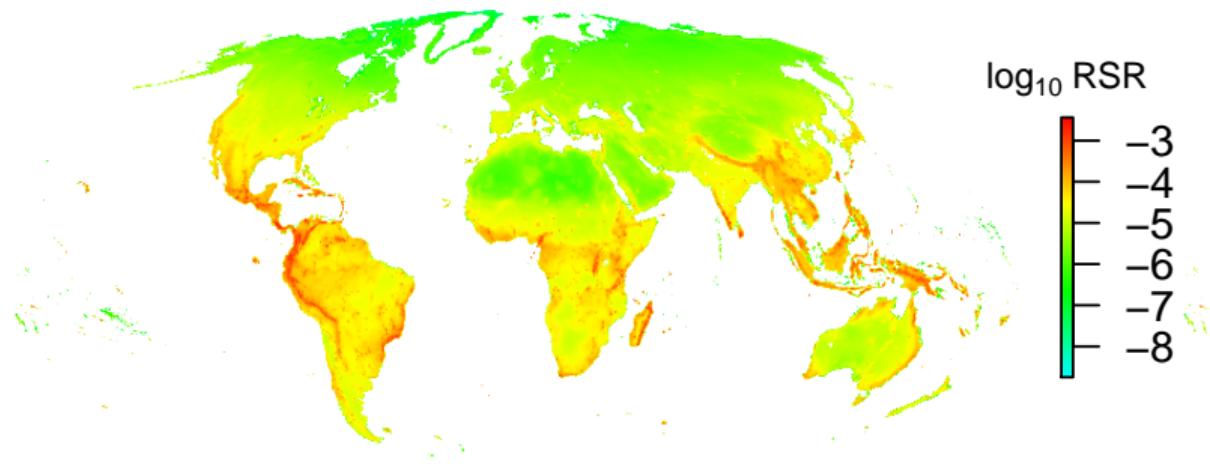
- The asymmetric random Lotka-Volterra competition assembly model and
- the Locally Saturated Patch Occupancy Model (LSPOM)

have both been solved analytically (in some limit). Their behaviour is well understood.

- Both models predicate empirically observed macroecological phenomena to high numerical accuracy.
- Both models invoke self-organised ESI.
- This adds to already strong empirical evidence for self-organised ESI from previous studies (Paris 2019).
- Self-organised ESI is *the* mechanisms controlling complexity and stability of real-world ecological networks.
- The fundamental principles governing ‘complexity’ and ‘stability’ of real-world ecological networks are understood. **Let's take it from there!**

-
- ① Which structure in interaction matrices matters, and for what? What are appropriate “guilds” of species?
(Barbier et al. 2018, PNAS)
 - ② How close is Nature to the UFP/MA transition?
 - ③ Community assembly with environmental filtering and variability [$dB_j/dt = (\textcolor{red}{s}_j - \sum_k^S G_{jk}B_k)B_j$].
 - ④ Spatio-temporally explicit variants of LSPOM.
 - ⑤ Species re-supply by allopatric speciation in long-separate lineages.
 - ⑥ Spatially continuous models (maths of neutral theory?). Predict, e.g., global range-size rarity ($RSR_j = \sum_i \text{at patch } j | \text{patches occupied by } i |^{-1}$).

Explain This!



Data: <https://www.iucnredlist.org/resources/other-spatial-downloads>

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