

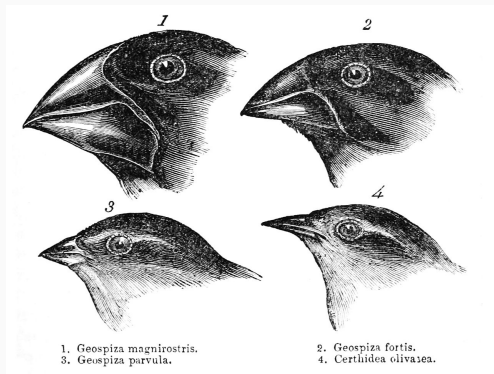
Trait dimensionality effects on model communities

Carlos A. Serván

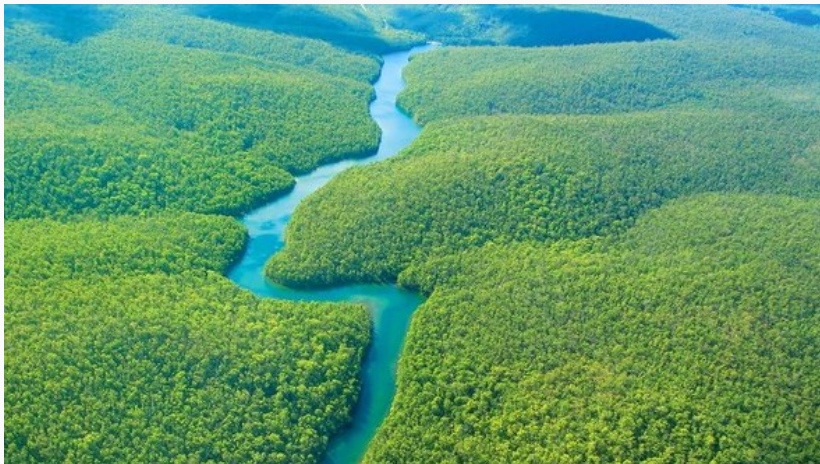
Joint work with Z. Miller, J.A. Capitán, T. Bodnar and S. Allesina

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Functional traits



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In our case $\mathcal{T} = \mathbb{R}^k$ and:

$$g : \mathcal{T} \rightarrow \mathbb{R}, t \rightarrow 1$$

$$f : \mathcal{T} \times \mathcal{T} \rightarrow \mathbb{R}, (t, t') \rightarrow \langle t, t' \rangle$$

For $x \in \mathbb{R}^n$:

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$$r = 1$$

$$A_{ij} = \frac{1}{k} f(t_i, t_j)$$

$$G = [t_{ji}] \in \mathbb{R}^{k \times n}$$

$$A = \frac{1}{k} G^T G$$

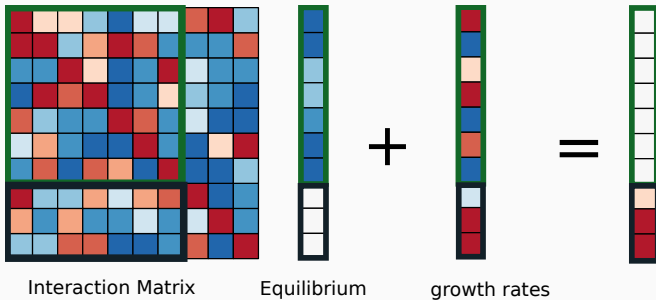
(Thm 15.3.1 Hofbauer and Sigmund 1998) Since A is symmetric and positive definite ($k \geq n$) we have a unique globally stable fixed point parameterized by $S \subset N = \{1, \dots, n\}$:

$$x_i > 0, \quad i \in S \quad (\text{Feasibility})$$

$$x_S(Ax_S + r) = 0 \quad (\text{Equilibrium})$$

$$(Ax_S + r)_i < 0, \quad i \notin S \quad (\text{Non-invasibility})$$

Non-invasive/saturated fixed point



Random zoo

- Jose A. Capitán (Universidad Politécnica de Madrid)
- Jacopo Grilli (ICTP)
- Kent E. Morrison (American Institute of Mathematics)
- Stefano Allesina (Chicago)



(Top Down approach)

- Take a pool of n species.
- Let dynamics elapse.
- $k \leq n$ species are coexisting.
- We want to determine $P(k|n)$.
- Study random ecosystems.

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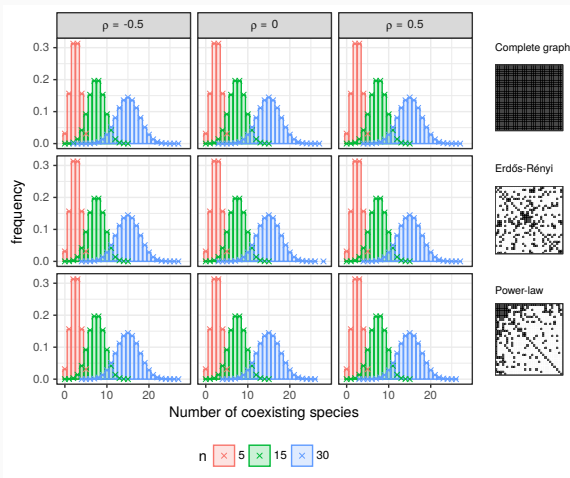
(For symmetric stable systems, equivalent to **Bottom up approach**)

Random Zoo

Computed the distribution of sizes of the survival community for the cases: (A_{ij}) and r_i *symmetric* about 0 random variables

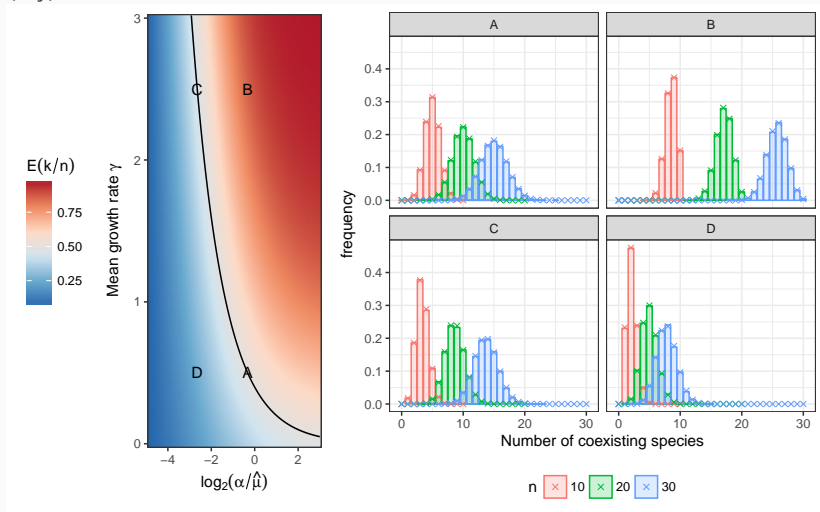
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Nonzero mean

(A_{ij}) deterministic and constant off diagonal with r gaussian.



Necessary condition for non-degenerate equilibrium $k \geq n$.

$$G_i \sim \mathcal{N}(0, \Sigma) \quad (\mathbf{Gaussian\ distribution})$$

Sample Covariance matrix

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$$A \sim \mathcal{W}_n\left(\frac{1}{k}\Sigma, k\right) \quad (\text{Wishart Distribution})$$

How does the community “look” as a function of the number of traits k ?

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- Distribution of the number of survivals : $\mathbb{P}(|S||k, n, \Sigma)$
- Mean number of survivals : $\varphi(k, n, \Sigma)$.
- Total biomass at the attractors : $W(k, n, \Sigma)$.

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Let $A \sim \mathcal{W}_n(\Sigma, k)$, $1_n \in \mathbb{R}^n$ a vector of ones and $L_{n-1} = (I_{n-1} 0)$, then
(Proof of Thm 1. Bodnar and Okhrin 2011)

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$$\tilde{x} = \frac{L_{n-1} A^{-1} 1_n}{1_n^T A^{-1} 1_n}$$

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$$\tilde{x} = \frac{L_{n-1} A^{-1} \mathbf{1}_n}{\mathbf{1}_n^T A^{-1} \mathbf{1}_n}$$

$$\tilde{x} \sim t_{n-1}(k - n + 2; \frac{L_{n-1} \Sigma^{-1} \mathbf{1}_n}{\mathbf{1}_n^T \Sigma^{-1} \mathbf{1}_n}, \frac{1}{(k - n + 2) \mathbf{1}_n^T \Sigma^{-1} \mathbf{1}_n} L_{n-1} R_1 L_{n-1}^T)$$

Probability of feasibility

$$Ax = 1$$

$$x_i > 0$$

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$$R_1 = \Sigma^{-1} - \Sigma^{-1} 1_n 1_n^T \Sigma^{-1} / 1_n^T \Sigma^{-1} 1_n$$

$$P_f(n) = \int_{\mathbb{R}^{n-1}} d\tilde{x}^{n-1} \rho(\tilde{x}) \Theta(1 - \mathbf{1}_{n-1}^T \tilde{x}) \prod_i \Theta(\tilde{x}_i)$$

$$P_f(n) = \int_{\mathbb{R}^{n-1}} d\tilde{x}^{n-1} p(\tilde{x}) \Theta(1 - \mathbf{1}_{n-1}^T \tilde{x}) \prod_i \Theta(\tilde{x}_i)$$
$$P_f(n) = \int_{\mathbb{R}} du g(u) \mathbb{P}(y_u > 0, \mathbf{1}_{n-1}^T y_u < 1) \quad (1)$$

Where:

$$u \sim \chi_{k-n+2}^2$$
$$y_u \sim \mathcal{N}\left(\frac{L_{n-1} \Sigma^{-1} \mathbf{1}_n}{\mathbf{1}_n^T \Sigma^{-1} \mathbf{1}_n}, \frac{1}{u \mathbf{1}_n^T \Sigma^{-1} \mathbf{1}_n} L_{n-1} R_1 L_{n-1}^T\right) \quad (2)$$

$$A = \begin{pmatrix} A_{11} & A_{12} \\ A_{21} & A_{22} \end{pmatrix}, \quad \Sigma = \begin{pmatrix} \Sigma_{11} & \Sigma_{12} \\ \Sigma_{21} & \Sigma_{22} \end{pmatrix}$$

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(Thm 3.2.10 Muirhead 1982):

$$A_{21}|A_{11} \sim \mathcal{N}(\Sigma_{21}\Sigma_{11}^{-1}A_{11}, \Sigma_{22.1} \otimes A_{11})$$
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In particular:

$$A_{21}A_{11}^{-1}\mathbf{1}|A_{11} \sim \mathcal{N}(\Sigma_{21}\Sigma_{11}^{-1}\mathbf{1}, \mathbf{1}^T A_{11}^{-1} \mathbf{1} \Sigma_{22.1})$$

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Probability of Non-Invasibility

Let $z = 1 - A_{21}A_{11}^{-1}\mathbf{1}$, and $W = \mathbf{1}^T A^{-1}\mathbf{1}$.

$P_{ni}(m) = \mathbb{P}(z < 0)$:

$$P_{ni}(m) = \int_{\mathbb{R}_+} dw g(w) O^-(n-m, 1 - \Sigma_{21}\Sigma_{11}^{-1}\mathbf{1}, w\Sigma_{22.1})$$

Theorem 3.2.11 Muirhead 1982 implies that :

$$\frac{\mathbf{1}^T \Sigma_{11}^{-1} \mathbf{1}}{W} \sim \chi_{k-m+1}^2$$

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$$P_{ni}(m) = \int_{\mathbb{R}_+} dw f(w) O^-(n-m, 1 - \Sigma_{21} \Sigma_{11}^{-1} \mathbf{1}, \frac{\mathbf{1}^T \Sigma_{11}^{-1} \mathbf{1}}{w} \Sigma_{22.1}) \quad (3)$$

For f the density function of χ_{k-m+1}^2 .

Constant Correlation $\rho \geq 0$ - Feasibility

$$\Sigma = (1 - \rho)I + \rho \mathbf{1}\mathbf{1}^T$$

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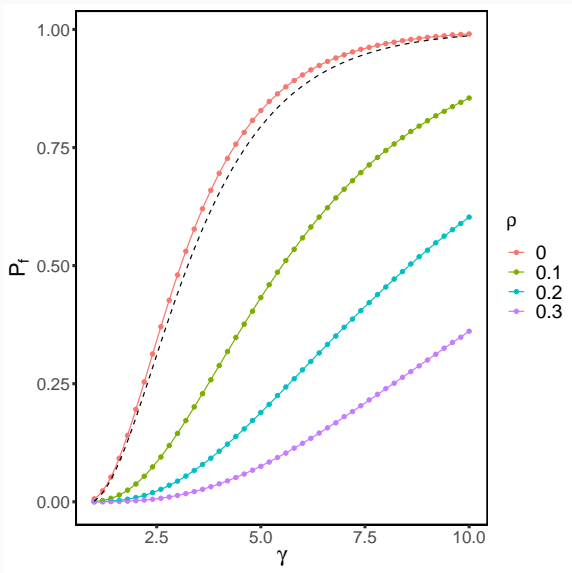
$$\Sigma = (1 - \rho)I + \rho \mathbf{1}\mathbf{1}^T$$

$$P_f(n) = \int_{\mathbb{R}_+} du g(u) \frac{-i\sqrt{n\alpha_u}}{\sqrt{2\pi}} \int_{\Gamma} d\zeta e^{\frac{n\zeta^2\alpha_u}{2}} \Phi\left(\frac{1/n + \zeta\alpha_u}{\sqrt{\alpha_u}}\right)^n \quad (4)$$

$$\alpha_u := \frac{1 + (n-1)\rho}{un(1-\rho)}$$

$$\beta_u := \frac{\alpha_u}{n}$$

Constant Correlation $\rho \geq 0$ - Feasibility



Constant Correlation $\rho \geq 0$ - Invasibility

$$P_{ni}(m, n) = \int_{\mathbb{R}^+} dw f(w) \int_{\mathbb{R}} dy \phi(y) \Phi\left(\frac{-1/m + y\sqrt{\beta_w}}{\sqrt{\alpha_w}}\right)^{n-m} \quad (5)$$

$$\alpha_w = \frac{1 + (m-1)\rho}{mw(1-\rho)}$$

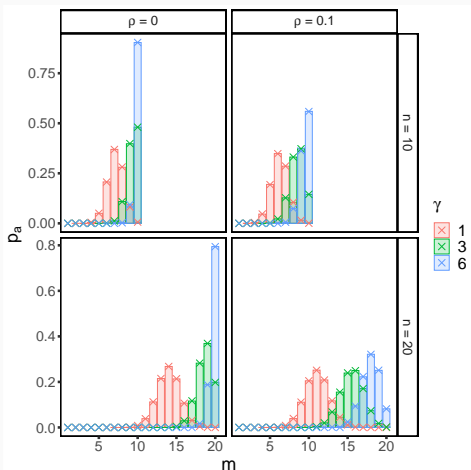
$$\beta_w = \frac{\rho\alpha_w}{1 + (m-1)\rho}$$

Distribution of survivors

$$\mathbb{P}(|S| = m | n, k, \rho) = \binom{n}{m} P_{f, n_i}(m) = \binom{n}{m} P_f(m) P_{n_i}(m) = P_a(m) \quad (6)$$

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Approximations for ϑ

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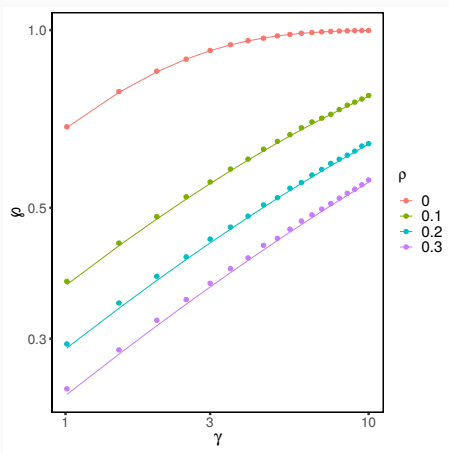
$$\lambda_q = 1 + \frac{nq\rho}{1-\rho}$$

$$\frac{\gamma - q^*}{q^*} - \left(\frac{\phi(\hat{q})}{q^*} + \hat{q} \right) \left((\lambda_q - 1) \frac{\phi(\hat{q})}{q^*} + \lambda_q q^* \right) = 0 \quad (7)$$

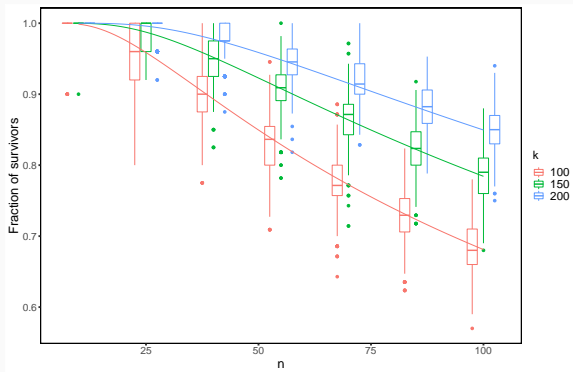
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Signature of Trait space dimension



Let T be a rooted phylogenetic tree, with total time 1.

$$\Sigma_T(i, j) = 1 - d(i, j) \quad (8)$$

Phylogenetic correlations

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In the limit of $\gamma \rightarrow \infty$, then $\frac{1}{k}A \rightarrow \Sigma_T$.

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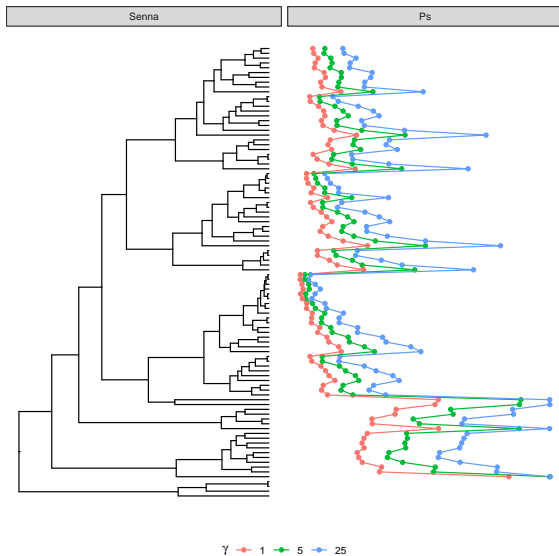
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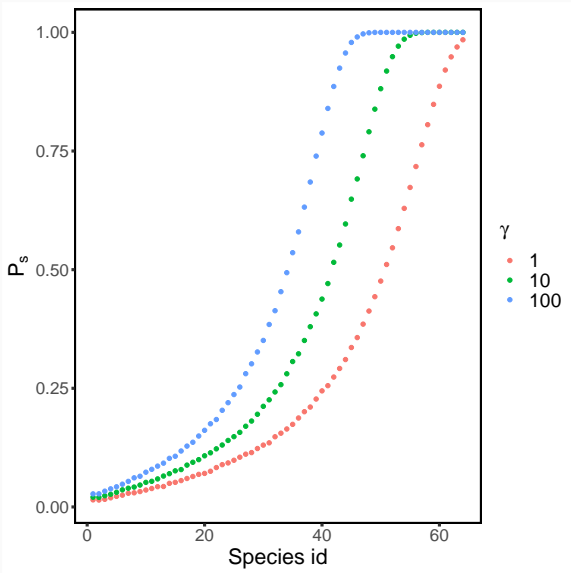
What happens in the case of γ *finite*?

Species sorting

(Weber and Agrawal 2014)



Perfectly unbalanced tree



- $k = n$ is necessary for a non-degenerate equilibrium, but almost never gives full coexistence.

Summary

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- We can detect a signature of the dimension of the trait space (under independent trait values).
- Our framework reproduces species sorting under phylogenetic correlation.

Acknowledgements

- Stefano Allesina
- Collaborators: José Capitán, Jacopo Grilli, Kent Morrison, Taras Bodnar, Zachary Miller.
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Thank you!