# Invasion and robustness in ecological meta-community models

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## Ecological models: overview

We may be familiar with two-species predator-prey models:

#### Predator-prey model with logistic growth and Lotka-Volterra FR

$$\frac{\mathrm{d}x}{\mathrm{d}t} = rx(1-x) - cxy,$$
  $\frac{\mathrm{d}y}{\mathrm{d}t} = \lambda cxy - dy$ 

Generalising these relationships:

- **Community (foodweb) model:** multiple interacting species. Food chains; specialists vs. generalists.
- Meta-population model: single species, multiple patches.
  Environmental heterogeneity; dispersal.
- Meta-community model: multiple species, multiple patches.

## Meta-community eco-evolutionary models

Considerations for models of ecosystem development:

- How to define species (one vs. multiple traits)?
- How to specify the feeding relationships?
- Is network structure of the spatial landscape (regular, fractal, small worlds or large worlds) important?
- Can foodweb structure be generated by simple evolutionary rules with ecological feedback (eco-evolutionary dynamics)?

## An eco-evolutionary meta-community model

Species are defined by a bodysize and discrete set of traits that score against other traits. Begin with one species, and resources in each patch. Simulation occurs in nested loops:

- Evolutionary loop: select parent species and introduce mutant (with 9/10 of parent's traits and similar bodysize).
  - Ecological loop: feeding, reproduction, death, dispersal.
    - Foraging loop: local populations select feeding strategies.

Developed from the Webworld foodweb model (Drossel et al 2001).

# Ecological loop (feeding, reproduction, death, dispersal)

During ecological timestep t, population dynamics of species i in patch (x, y) updated:

$$N_{i,x,y}^{t} \mapsto N_{i,x,y}^{t} + \Delta \left( -2s_{i}^{-0.25}N_{i,x,y}^{t} + \lambda s_{i}^{-1}N_{i,x,y}^{t} \sum_{j=0}^{n} g_{i,j}s_{j} - \sum_{k=1}^{n} N_{k,x,y}^{t}g_{k,i} \right)$$

Loss due to mortality. Gains due to feeding. Loss due to predation.  $s_i$  is bodysize;  $\lambda = 0.3$  is ecological efficiency;  $g_{i,j}$  is functional response on j

Subsequently, migration to and from neighbouring patches:

$$\textit{N}_{i,\mathsf{x},\mathsf{y}}^t \mapsto \textit{N}_{i,\mathsf{x},\mathsf{y}}^t + \sum_{j=1}^{\mathsf{x}_{\max}} \sum_{k=1}^{\mathit{y}_{\max}} \mu_{i,j,k,\mathsf{x},\mathsf{y}} \textit{N}_{i,j,k}^t - \sum_{j=1}^{\mathsf{x}_{\max}} \sum_{k=1}^{\mathit{y}_{\max}} \mu_{i,\mathsf{x},\mathsf{y},j,k} \textit{N}_{i,\mathsf{x},\mathsf{y}}^t$$

Dispersal rate  $\mu$  increases in patches where population is declining.

## Foraging loop

Between updating population sizes, the local population of species i in patch (x, y) iterates feeding strategies in two stages:

• Update feeding efforts distributed across prey *j*:

$$f_{i,j,x,y} = \frac{g_{i,j,x,y}}{\sum_{k \in K_i} g_{i,k,x,y}}$$

• Update ratio-dependent functional responses:

$$g_{i,j,x,y} = \frac{S_{i,j}f_{i,j,x,y}N_{j,x,y}}{0.005N_{j,x,y} + \sum_{k \in P_i} \alpha_{i,k}S_{k,j}f_{k,j,x,y}N_{k,x,y}}$$

 $K_i$  is the set of possible prey of i;  $P_j$  is the set of possible predators of j;  $S_{i,j}$  is trait-dependent feeding score of i on j;  $\alpha_{i,k}$  controls competition based on level of similarity.

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### Example ensembles

Simulation generates a meta-community of co-evolved species.

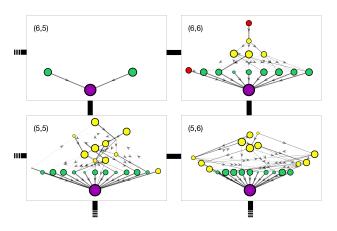


Figure 1: Foodwebs in  $6 \times 6$  spatial network (colours denote trophic role)

# Foodweb structure and stability

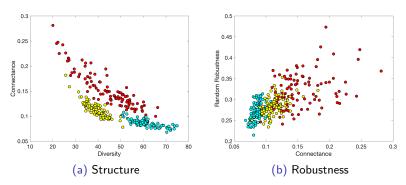


Figure 2: Local foodweb properties (colours relate to model configuration)

- Scaling of number of feeding links L with number of species S.
- Connectance (L/(S(S-1))) decreases with diversity.
- Community robustness increases with connectance.

## Invasion of patches in the meta-community

Practical stability: **robustness** of existing ecosystems to species invasion.

Each species is deleted, (max loss 2.1%) then re-introduced to *all* patches (max loss 12%).

Number of patches invaded is increased by:

- Co-evolved traits.
- Larger bodysizes.

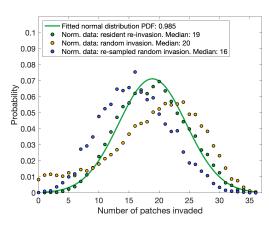


Figure 3: Probability of patch invasion

## Invasion - role of bodysize

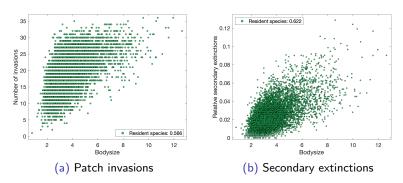


Figure 4: Bodysize correlations with invasion and secondary extinctions

- Larger-bodysize species prefer to predate at higher trophic levels, avoiding strong competition to feed on resources.
- Benefit more from uniform-population re-introduction.

#### Habitat loss and nature reserves

Patches are subjected to repeated random disturbances, except for 6 (of 36) designated as nature reserves. What is the best choice?

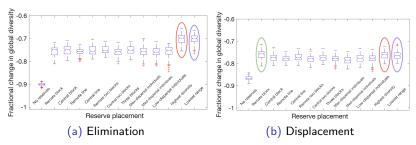


Figure 5: Biodiversity loss due to perturbation of random patch sequences

- Reliable to select patches with the greatest biodiversity or to protect the rarest species.
- If disturbed populations are displaced (rather than eliminated), isolating large, remote areas from invasion also effective.

#### Outlook

#### **Summary:**

- Models with simple evolutionary rules re-create complex network structure with greater stability.
- Can be used to determine principles for which sites and species are most vulnerable to perturbation, and hence inform environmental efforts.

#### **Challenges:**

- Empirical data limited for validation.
- Abstract models hard to apply to specific ecosystems focus on simpler, bespoke models?
- Exact mechanisms not easily identifiable in complex systems.

#### References

- Drossel, B., Higgs, P. G., & McKane, A. J. (2001). The influence of predator—prey population dynamics on the long-term evolution of food web structure. *Journal of Theoretical Biology*, 208(1), 91-107.
- Abernethy, G. M. (2020). Allometry in an eco-evolutionary network model. *Ecological Modelling*, 427, 109090.
- Abernethy, G. M. (2021). Sequences of patch disturbance in a spatial eco-evolutionary model. Communications in Nonlinear Science and Numerical Simulation, 97, 105746.