# Stability in ecological meta-community models

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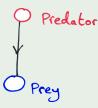
# Ecological models: overview

We may be familiar with two-species models in population dynamics:

### Predator-prey model

Prey: 
$$\frac{\mathrm{d}x}{\mathrm{d}t} = f(x)x - g(x,y)y$$

Predator: 
$$\frac{\mathrm{d}y}{\mathrm{d}t} = \lambda g(x, y)y - dy$$



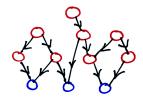
e.g. with logistic prey growth and Lotka-Volterra predation. . .

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

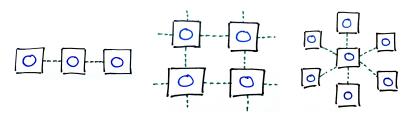
Predator: 
$$\frac{\mathrm{d}y}{\mathrm{d}t} = \lambda cxy - dy$$

# Ecological models: overview

• Community (foodweb) model: multiple interacting species.

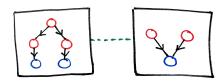


• Meta-population model: single species, multiple patches.



## Ecological models: overview

• Meta-community model: multiple species, multiple patches.



 Eco-evolutionary model: mutation/speciation, adaptive dynamics.



 Community assembly model: multiple species, new species invade from pre-existing pool.

# Eco-evolutionary meta-community models

#### Combine all three features:

- Mutation and evolutionary effects
- Space
- Multiple species

### **Conceptual challenges:**

- How to define species (one vs. multiple traits)?
- Dynamically determine feeding relationships (based on traits)?
- Network structure of the spatial landscape?
- Dispersal mechanism (diffusion, probabilistic, adaptive)?
- Population dynamics (reproduction, functional response)?
- Does the ruleset result in plausible foodweb structures?

## 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 on a square  $N \times N$  lattice. Simulation occurs in nested loops:

- **Evolutionary loop:** choose a 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 adaptively decide feeding strategies (avoid competition, prefer populous prey).

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

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

At time t, population dynamics of species i in patch (x, y) obeys:

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

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 ratio-dependent response on j.

Subsequently, migration from and to neighbouring patches:

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

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

## Example ensembles

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

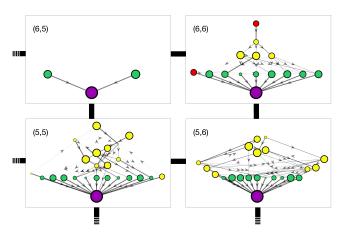


Figure 1: Foodwebs in  $6 \times 6$  spatial network (colours denote trophic role)

# Foodweb structure and stability

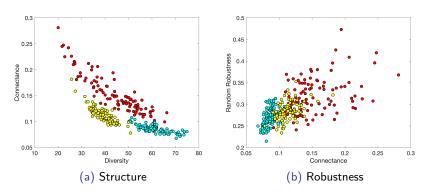


Figure 2: Local foodweb properties (colours relate to model configuration)

- How does the number of realised feeding links L depend on diversity S? Connectance (L/(S(S-1))) decreases with diversity.
- **Stability:** Community robustness increases with connectance.

# Species extinction and invasion

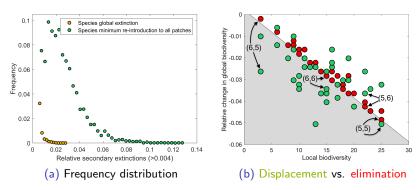


Figure 3: Effects of extinction, invasion and displacement

- Each species is deleted, (max loss 2.1%) then re-introduced to *all* patches (max loss 12%).
- Displacement to neighbouring patches can be more damaging than extinction.

### 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?

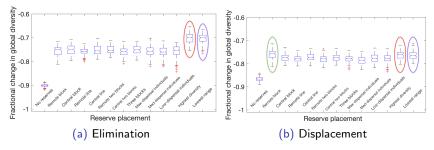


Figure 4: Biodiversity loss due to perturbation of random patch sequences

- 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<sup>1</sup>

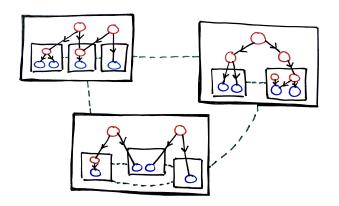
### **Summary:**

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

## **Challenges:**

- Validation from empirical data?
- Species-area curve and mixed range of species (neither spatially-homogeneous, nor total species-sorting).
- Limited conclusions from abstract models for specific ecosystems.

## Future work: strand I



- Simplified trait-space with explicit specialist-generalist trade-off;
- Multi-scale species perceptions of habitat;
- Role of scaling patterns in habitat structure and its removal.

## Future work: strand II

Application to management of water vole, mink and otter meta-communities in South Yorkshire:

- Collaboration with Sheffield Hallam University and the Sheffield and Rotherham Wildlife Trust;
- Simplified community model three-species population dynamics;
- Highly-resolved spatial network from GIS data;
- Application to predicted impact of habitat perturbation.

## References

- Original model: 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.
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