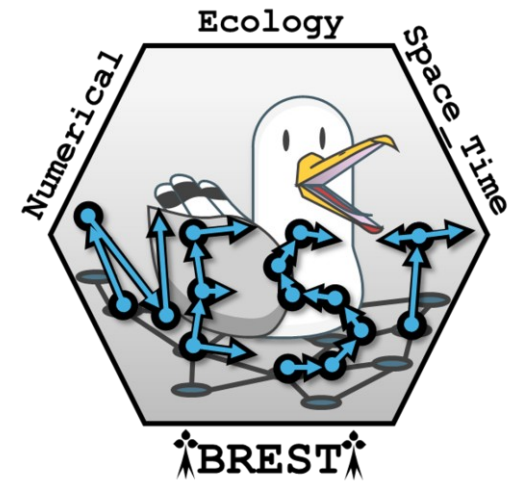


Network Tools for Ecology

David Cunillera-Montcusí
david.cunillera@dcm.cat

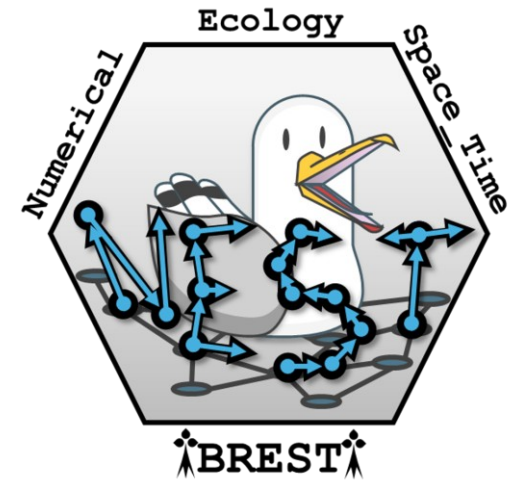


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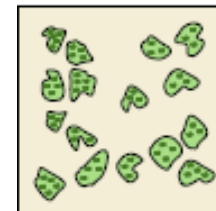
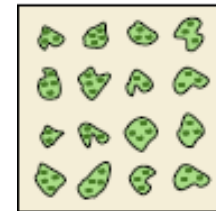
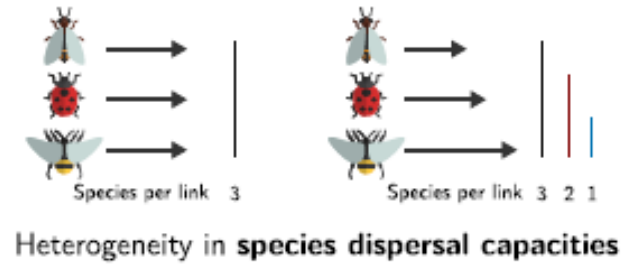
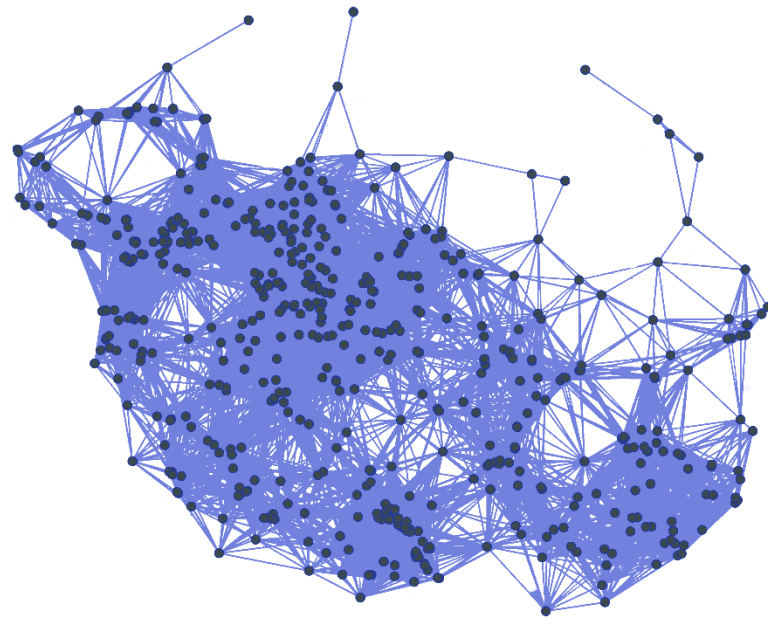
- First session (10:00-11:00)
 - Introduction to networks
 - Spatial networks and where to find them
 - R time through network examples
- Second session (11:30-12:30)
 - Connecting network structure with diversity
 - Coalescent runs and habitat loss



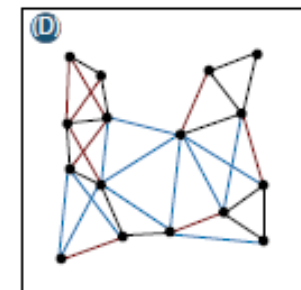
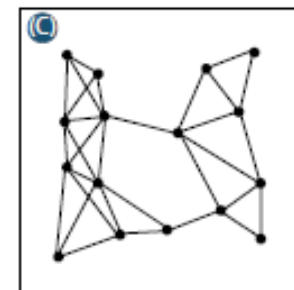
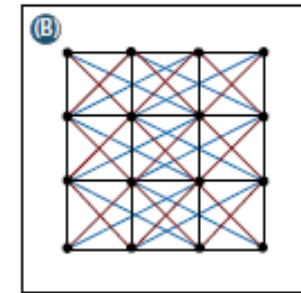
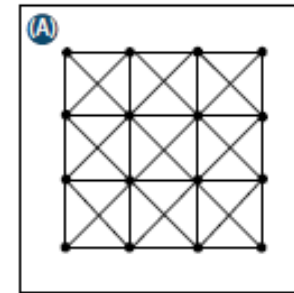
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Connecting network structure with diversity

Network structure

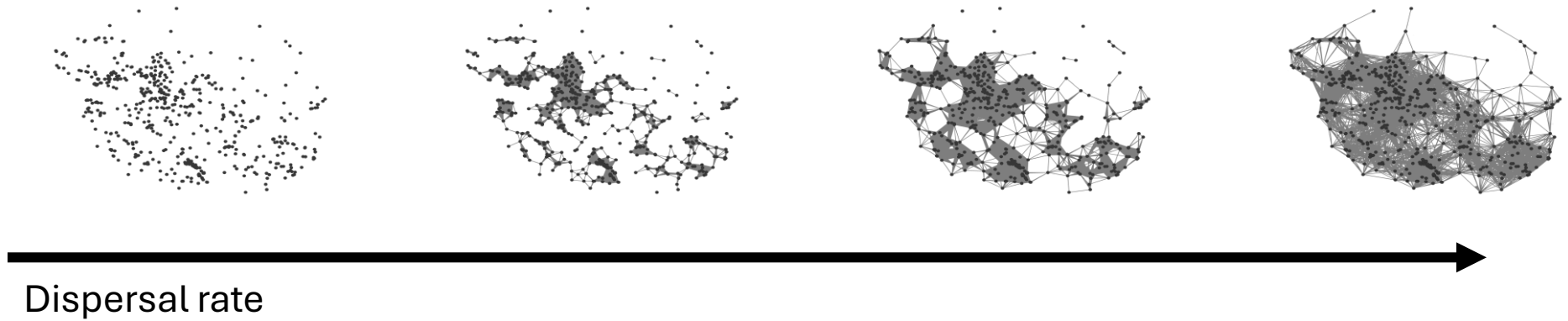


Heterogeneity in landscape structure



Trends in Ecology & Evolution

Network structure



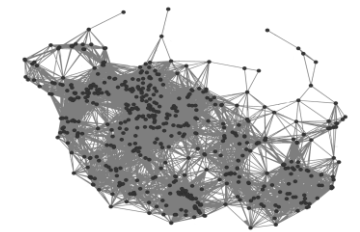
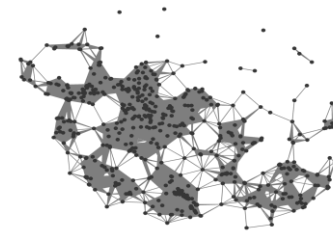
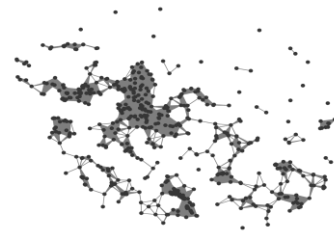
Connecting network structure with diversity



Biotic interactions
Abiotic filtering
Dispersal

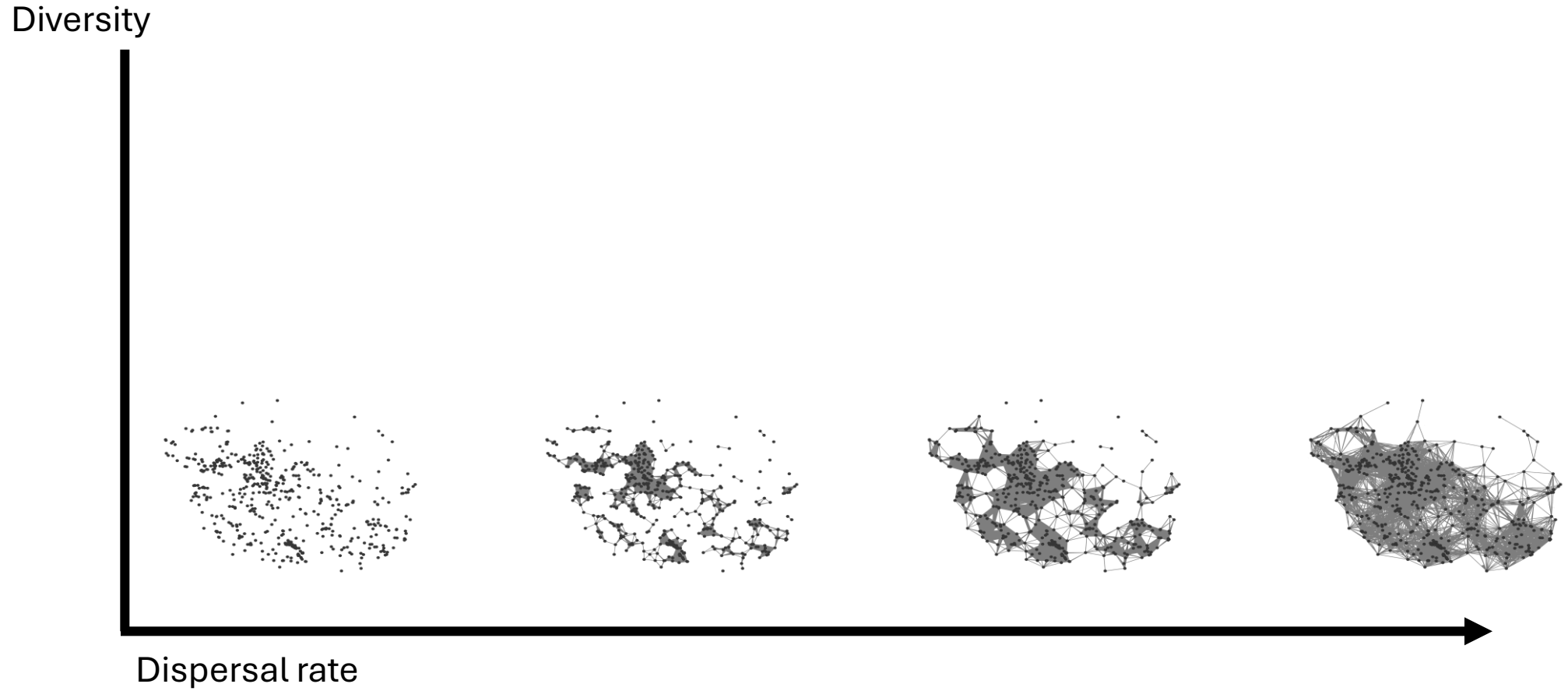
Biotic interactions
Abiotic filtering
Dispersal

Biotic interactions
Abiotic filtering
Dispersal

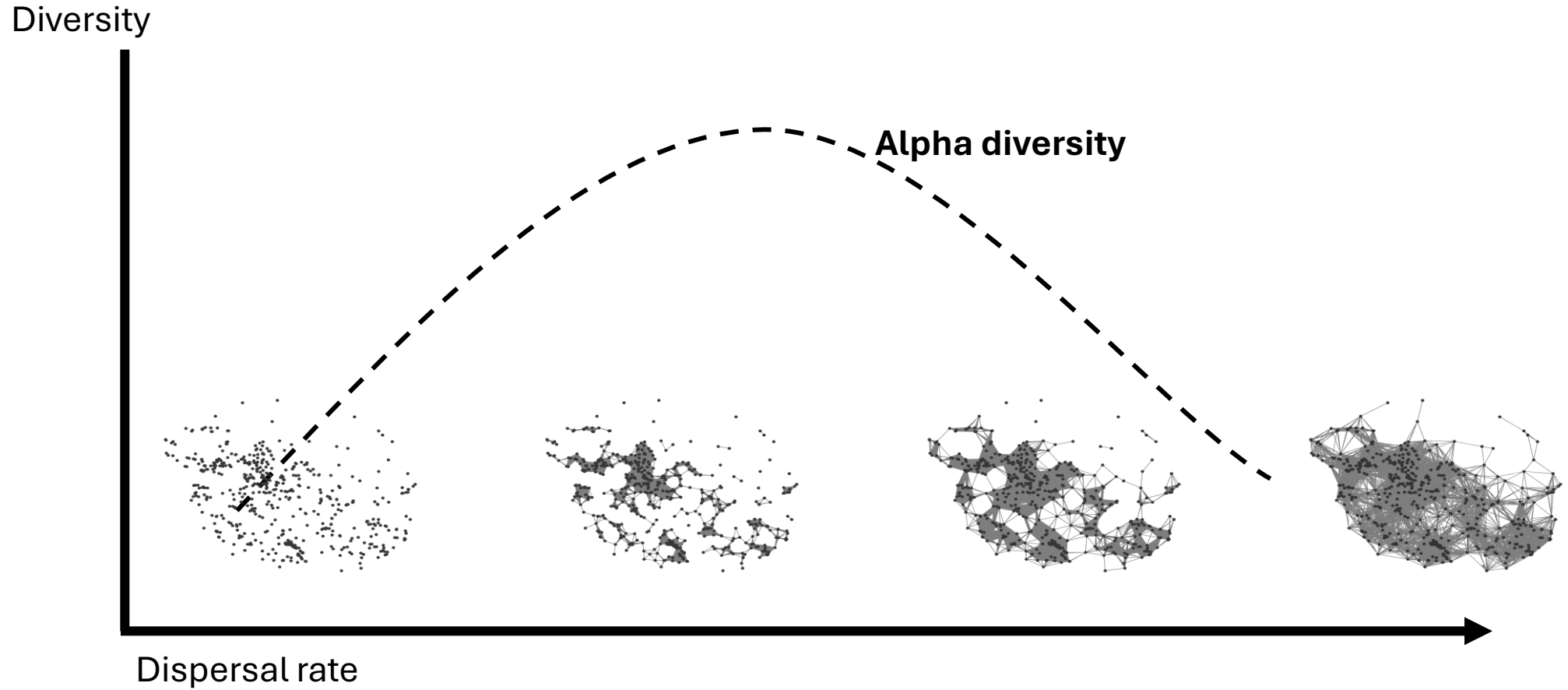


Dispersal rate

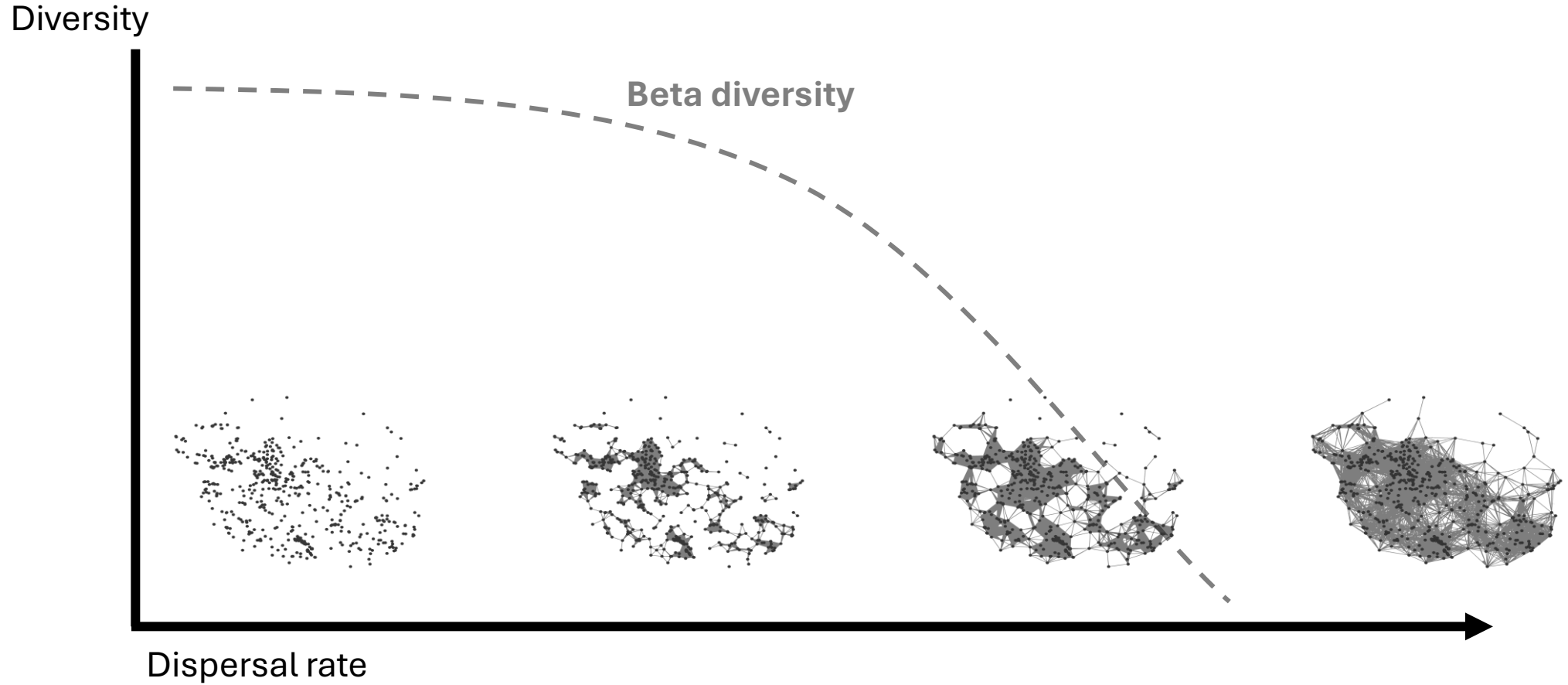
Connecting network structure with diversity



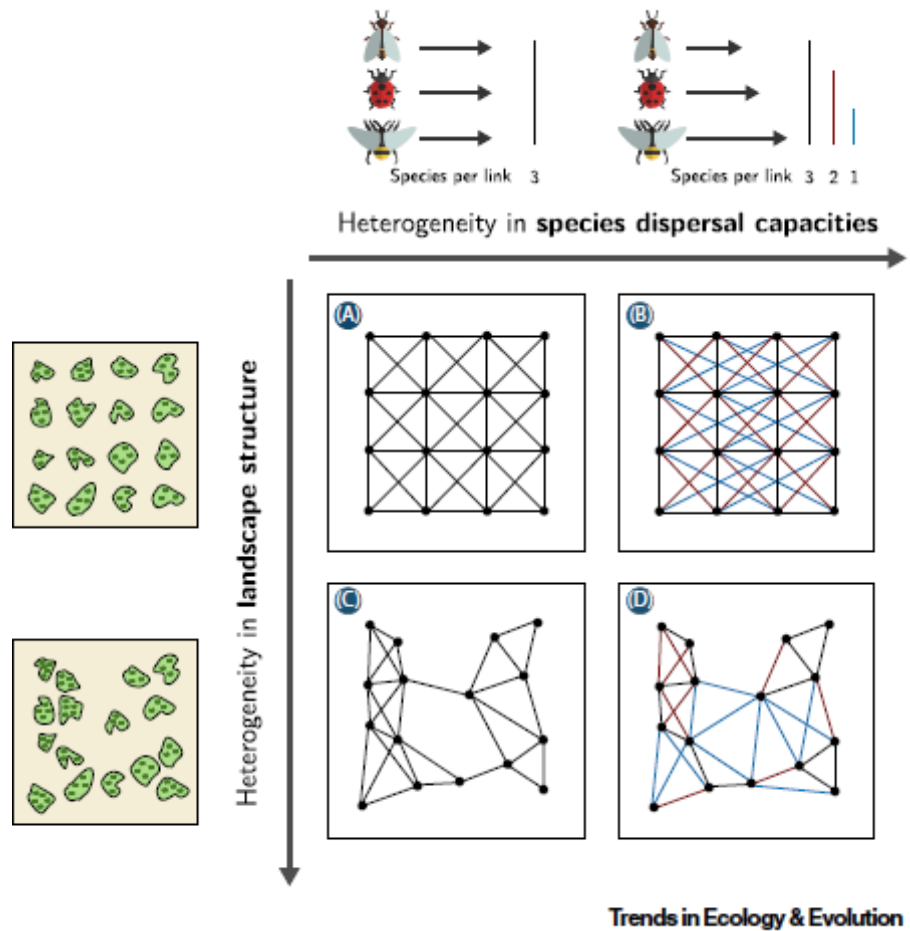
Connecting network structure with diversity



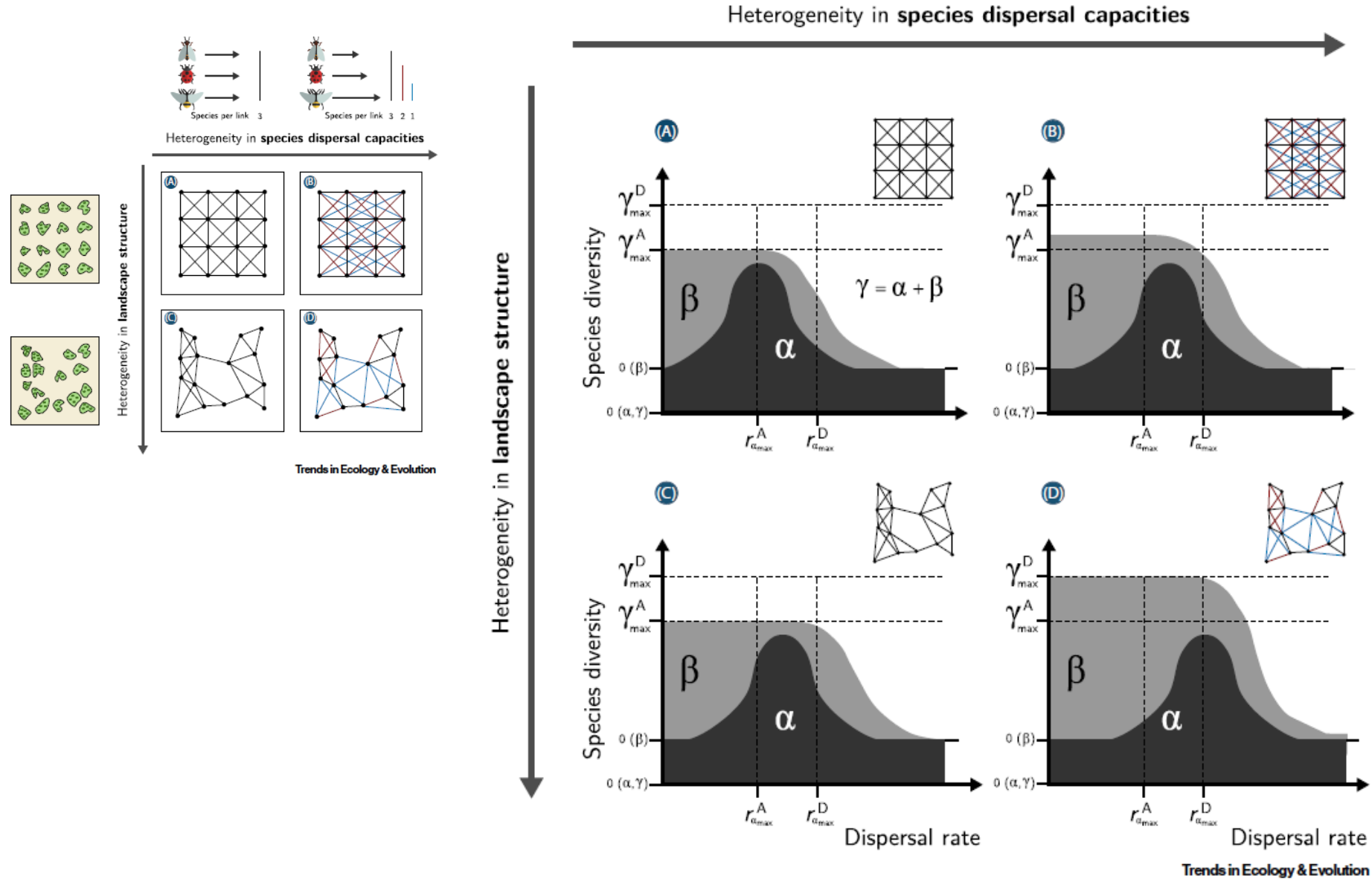
Connecting network structure with diversity



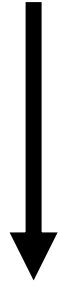
Connecting network structure with diversity



Connecting network structure with diversity

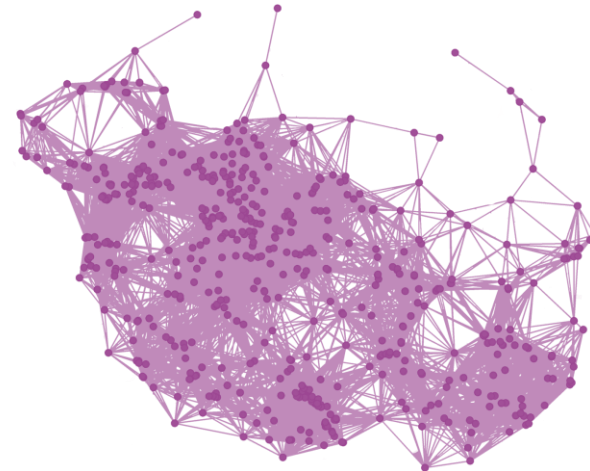
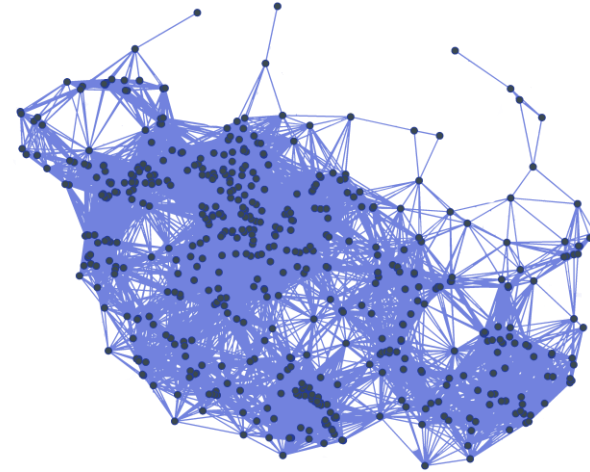


Spatial network



In silico

Metacommunity models




Connecting network structure with diversity





In silico

Metacommunity models

RESEARCH ARTICLE

Methods in Ecology and Evolution 


ecolottery: Simulating and assessing community assembly with environmental filtering and neutral dynamics in R

François Munoz¹  | Matthias Grenié² | Pierre Denelle² | Adrien Taudière² |
Fabien Laroche^{2,3} | Caroline Tucker^{2,4} | Cyrille Violle² 

Received: 19 April 2024 | Accepted: 4 March 2025

DOI: 10.1111/1365-2656.70033

RESEARCH ARTICLE

 Journal of Animal Ecology

Inferring riverscape dispersal processes from fish biodiversity patterns


Ana I. Borthagaray^{1,2}  | Franco Teixeira de Mello¹ | Matías Arim^{1,2} 

ECOLOGY LETTERS

Ecology Letters, (2020) 23: 1314–1329







doi:10.1111/ele.13568

IDEAS AND PERSPECTIVES

A process-based metacommunity framework linking local and regional scale community ecology 

Abstract

The metacommunity concept has the potential to integrate local and regional dynamics within a general community ecology framework. To this end, the concept must move beyond the discrete archetypes that have largely defined it (e.g. neutral vs. species sorting) and better incorporate local scale species interactions and coexistence mechanisms. Here, we present a fundamental reconception of the framework that explicitly links local coexistence theory to the spatial processes inherent to metacommunity theory, allowing for a continuous range of competitive community

Patrick L. Thompson,^{1*} 
Laura Melissa Guzman,^{1,2} 
Luc De Meester,^{1,4,5} 
Zsófia Horváth,^{1,4,7} 
Robert Ptáček,⁶ 
Bram Vanschoenwinkel,^{4,8} 
Duarte S. Viana,
Jonathan M. O.

ECOGRAPHY

Research

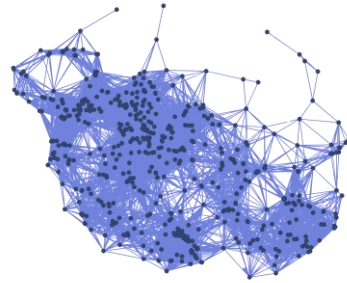
Metacommunity resilience against simulated gradients of wildfire: disturbance intensity and species dispersal ability determine landscape recover capacity

David Cunillera-Montcusí, Ana Inés Borthagaray, Dani Boix, Stéphanie Gascón, Jordi Sala, Irene Tornero, Xavier D. Quintana and Matías Arim

Connecting network structure with diversity

Observed landscape structure

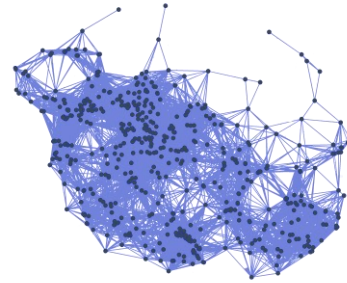
Spatial location of habitats



Connecting network structure with diversity

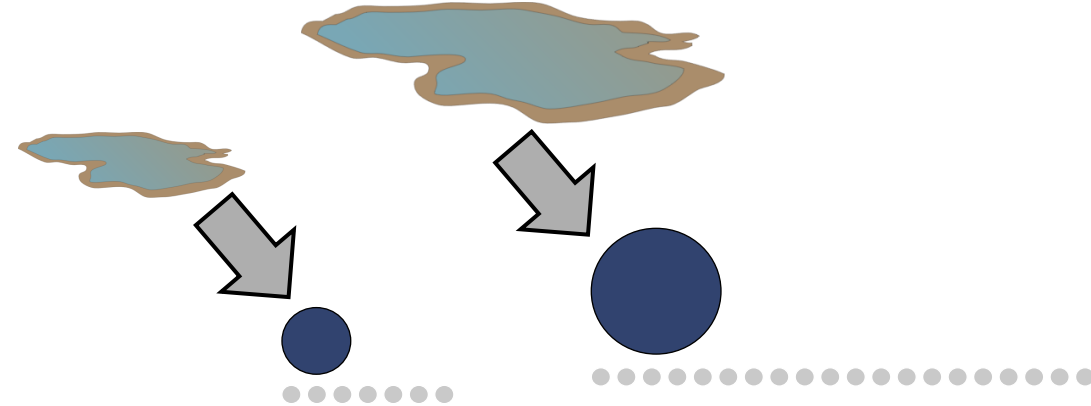
Observed landscape structure

Spatial location of habitats



Habitat size

Local community size (**J**) is a power function of Area ($J \sim \text{Area}^b$)



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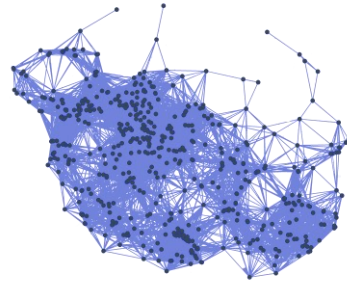
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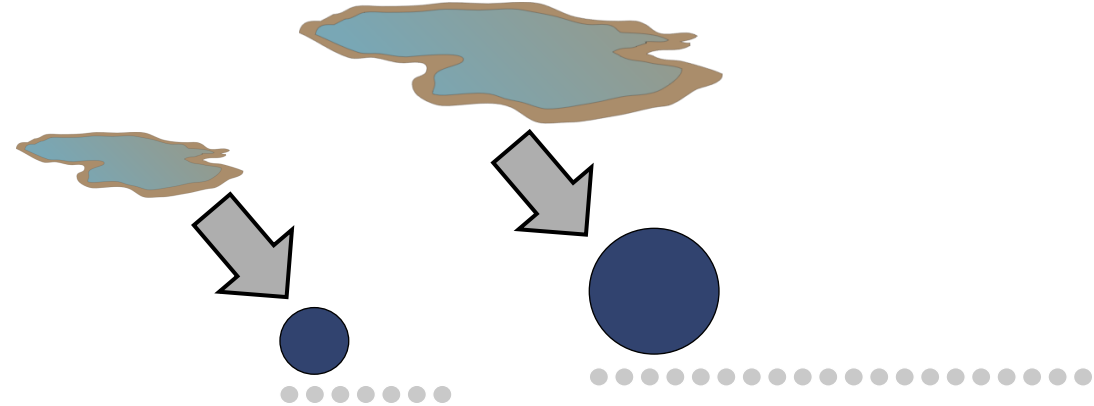
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Spatial location of habitats



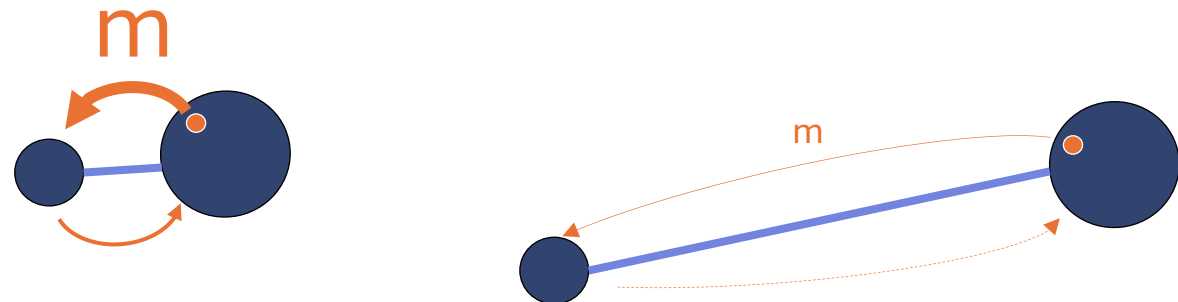
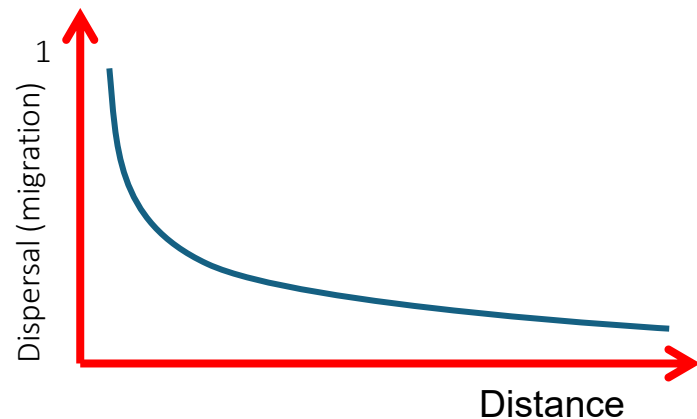
Habitat size

Local community size (**J**) is a power function of Area ($J \sim \text{Area}^b$)



Dispersal

Dispersal is function of the distance between local communities (dispersal Kernel).



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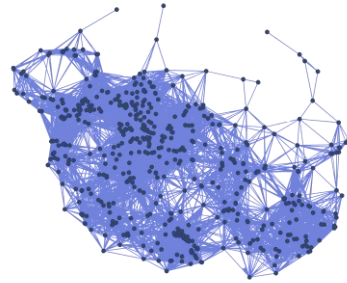
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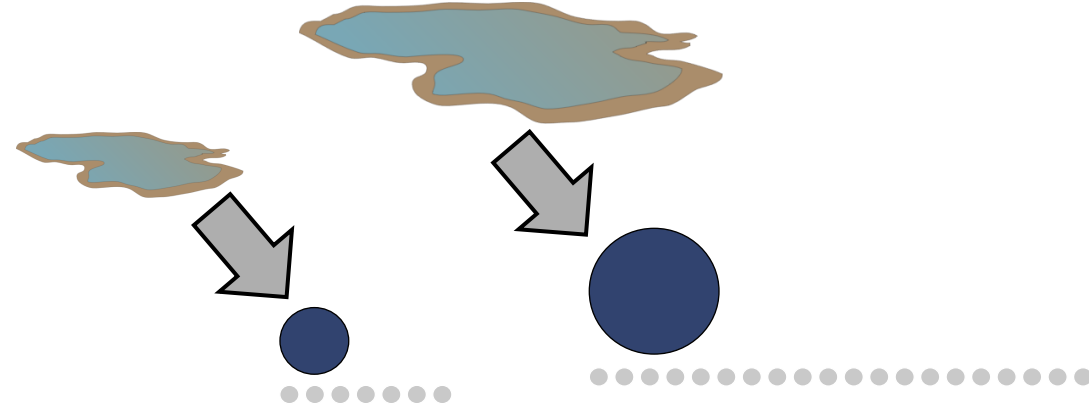
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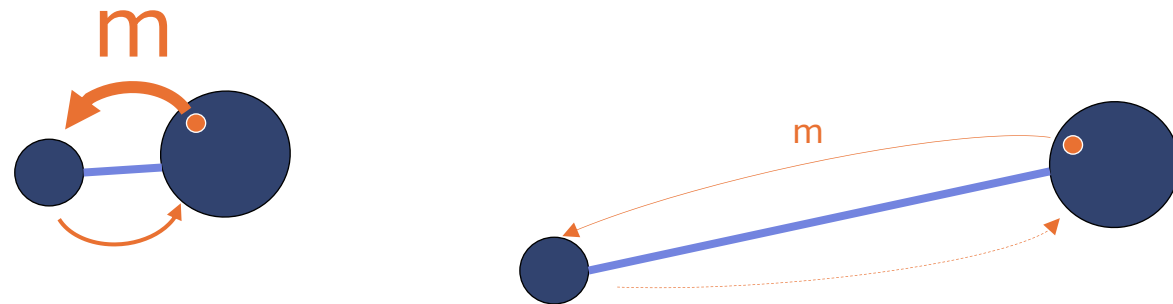
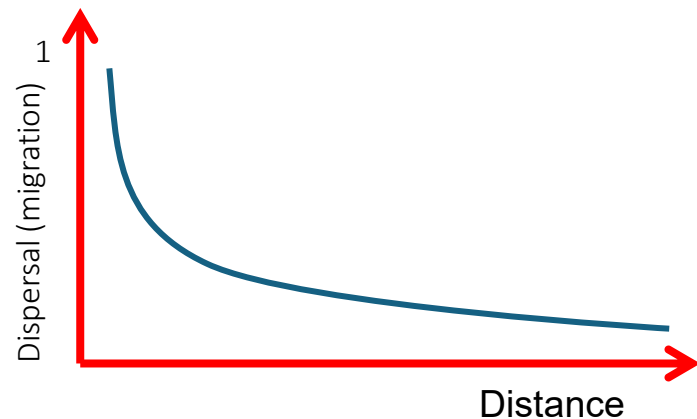
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Dispersal

Dispersal is function of the distance between local communities (dispersal Kernel).



The “**weight**” of links is determined by the expected dispersal, times the size of the source community (Hanski 1999)

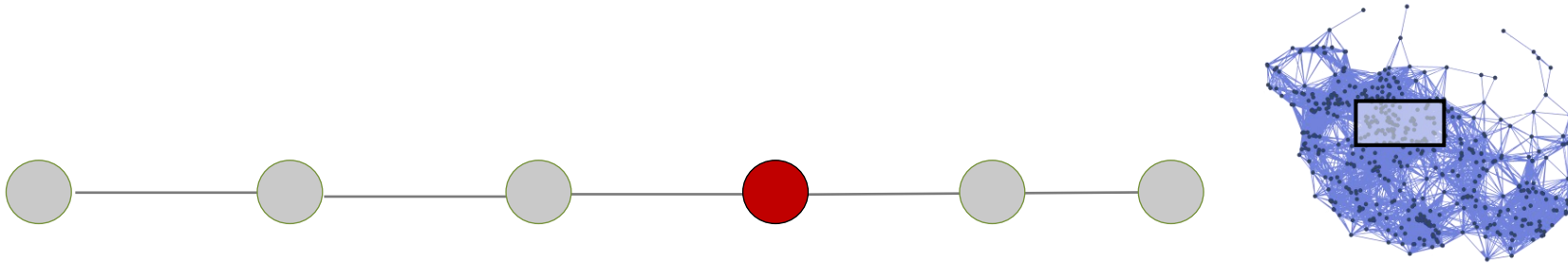
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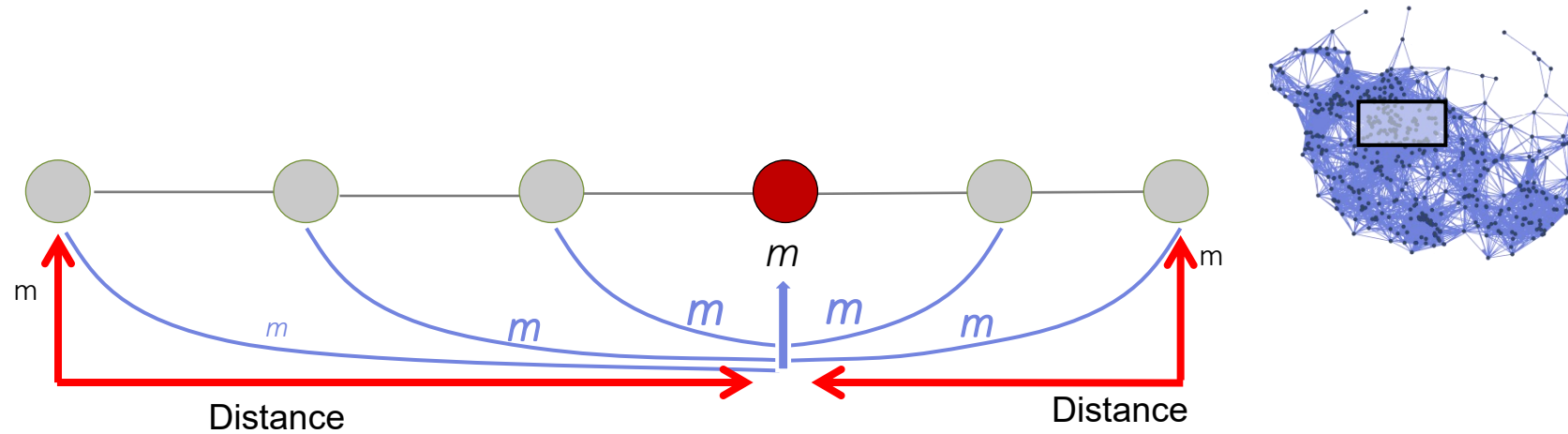
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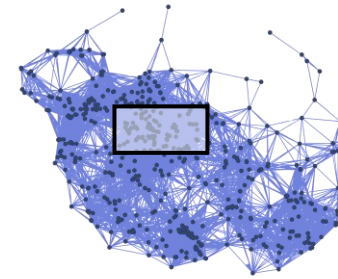
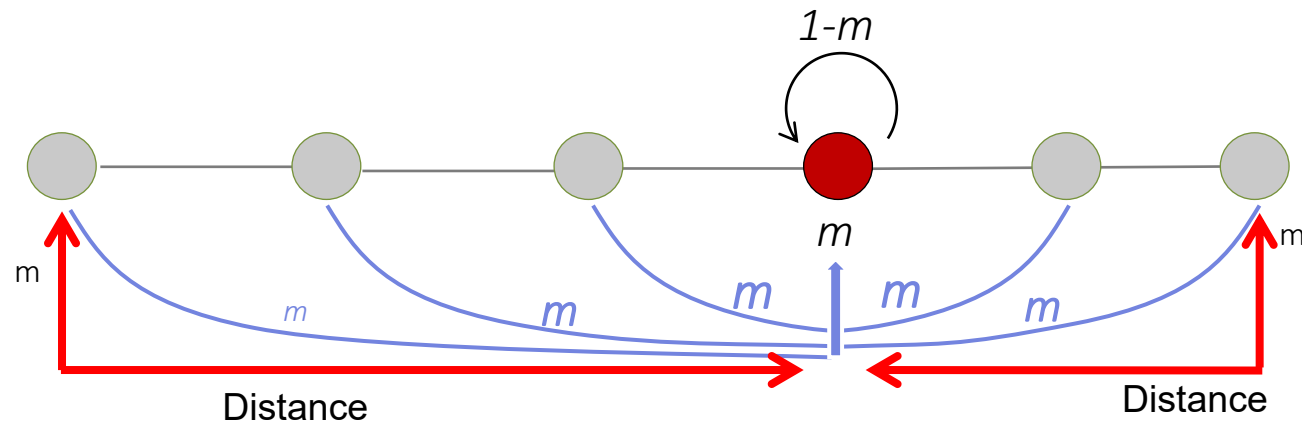
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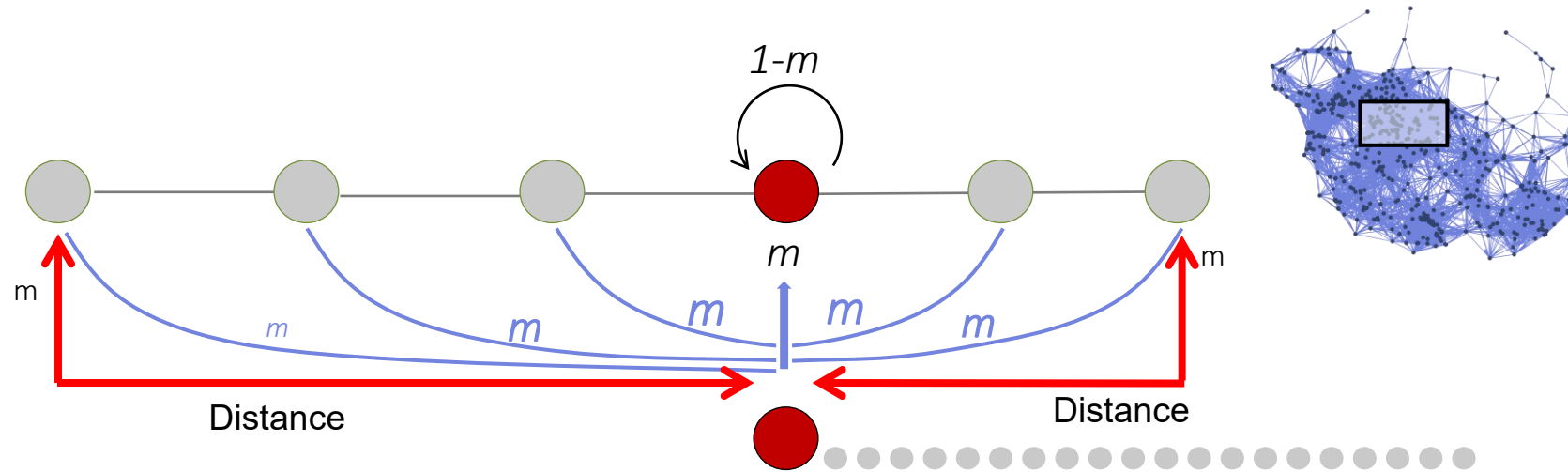


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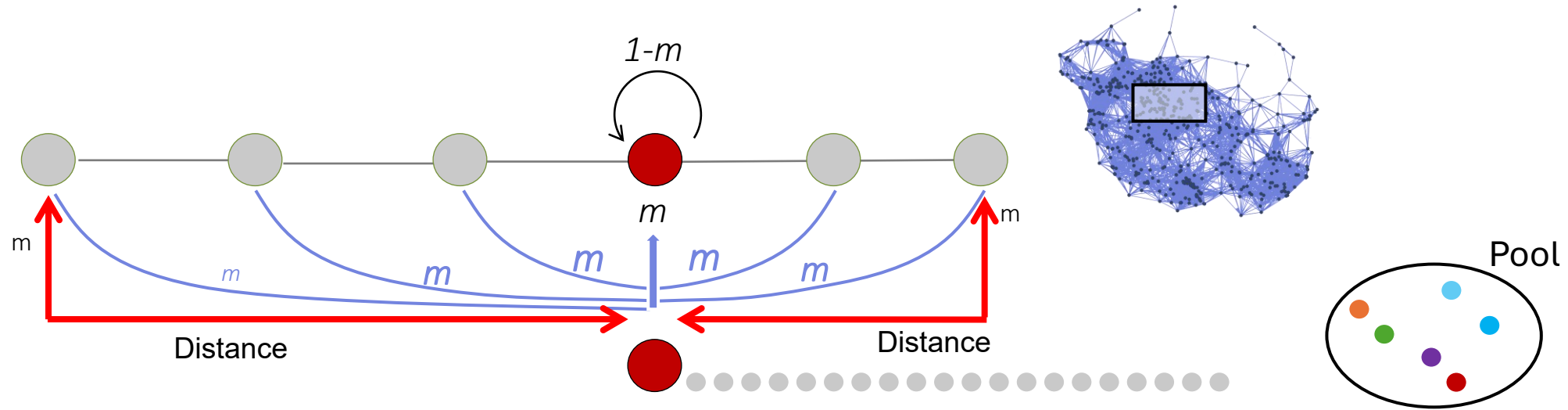
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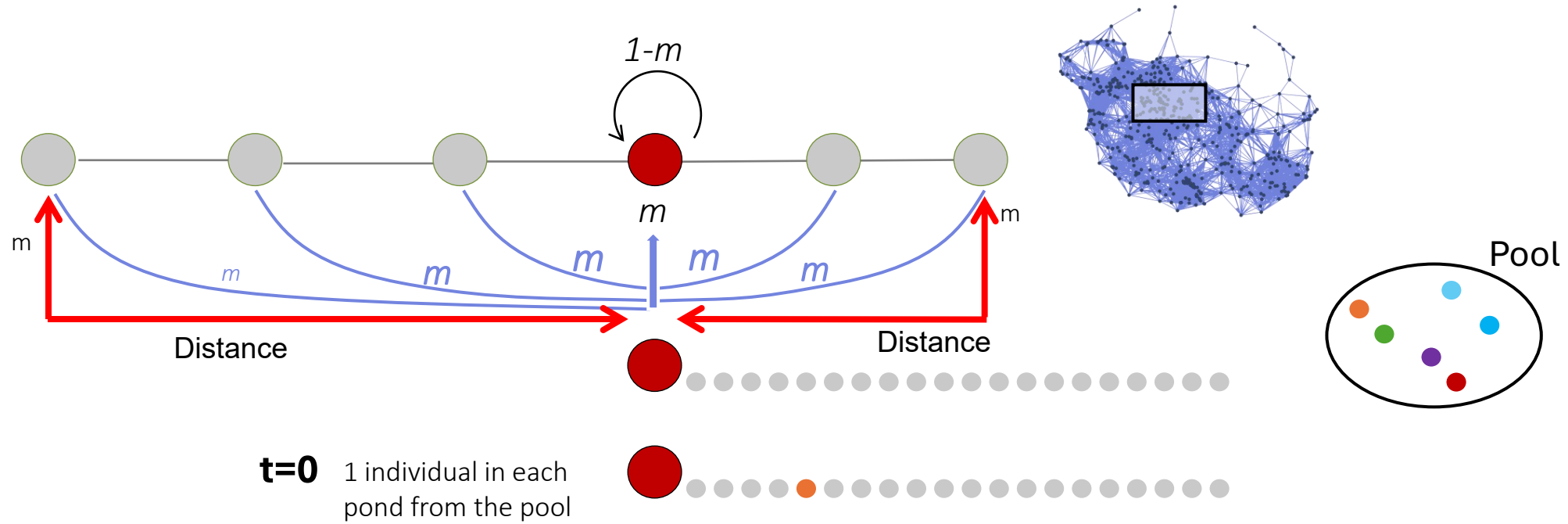


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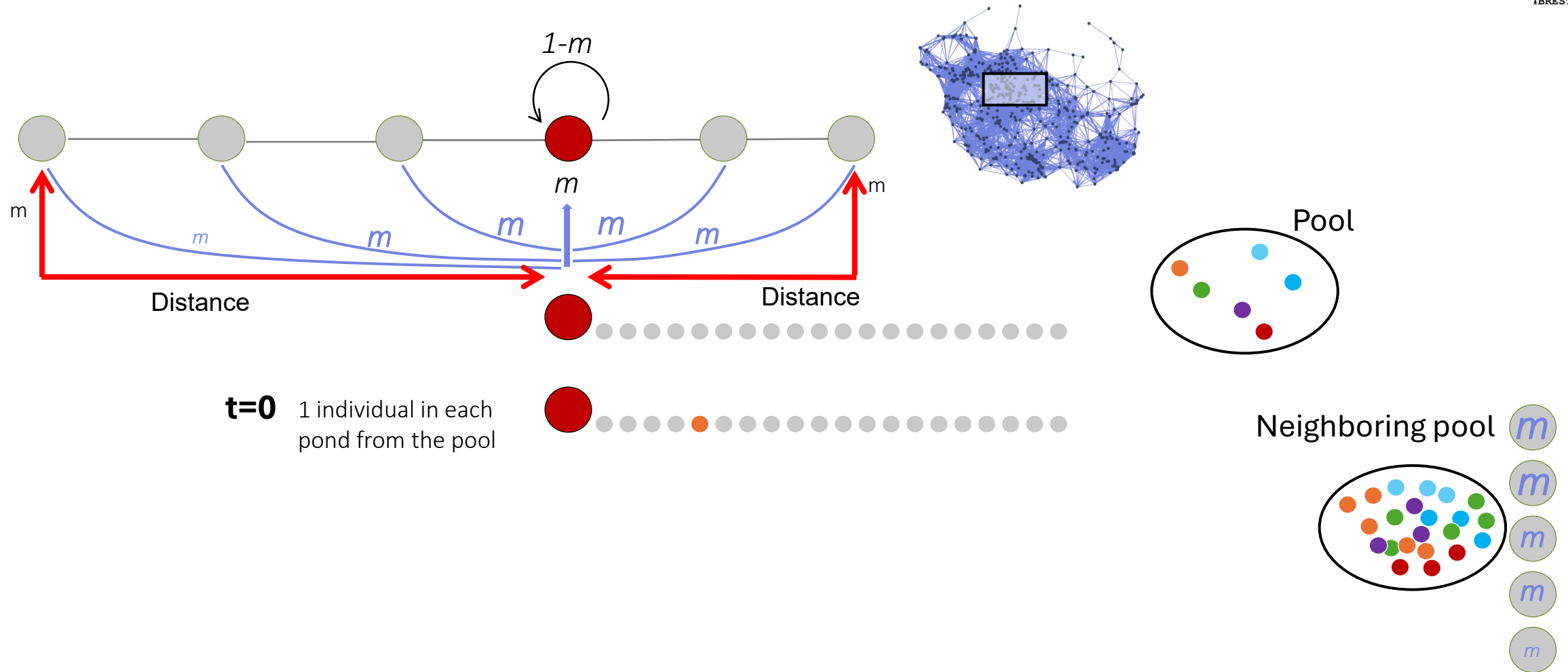


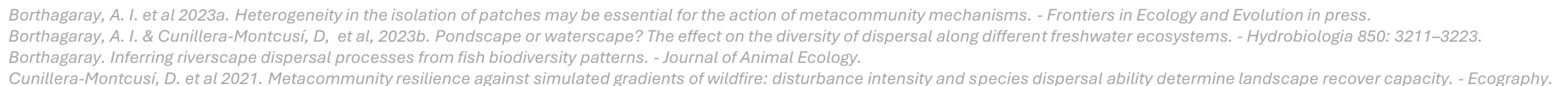
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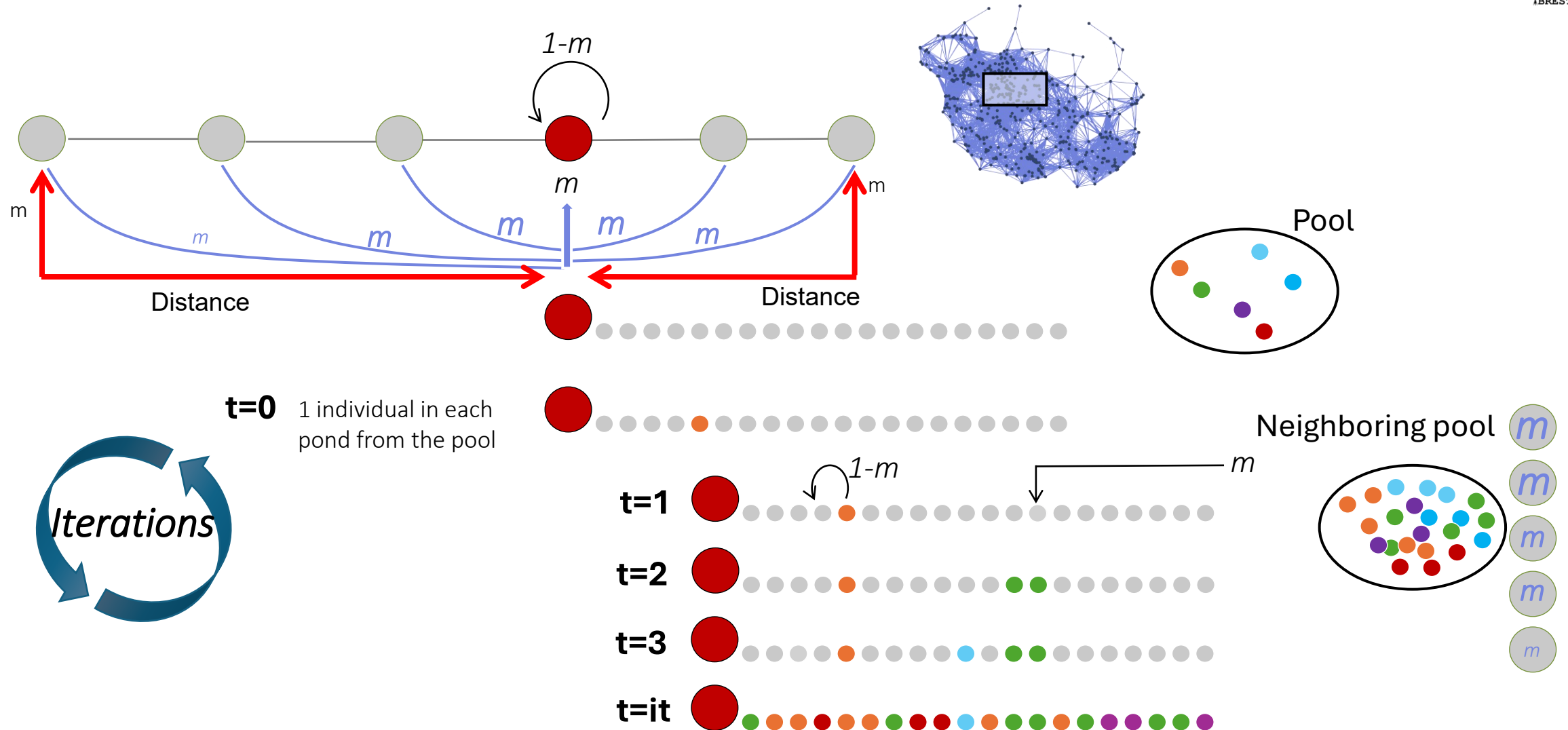


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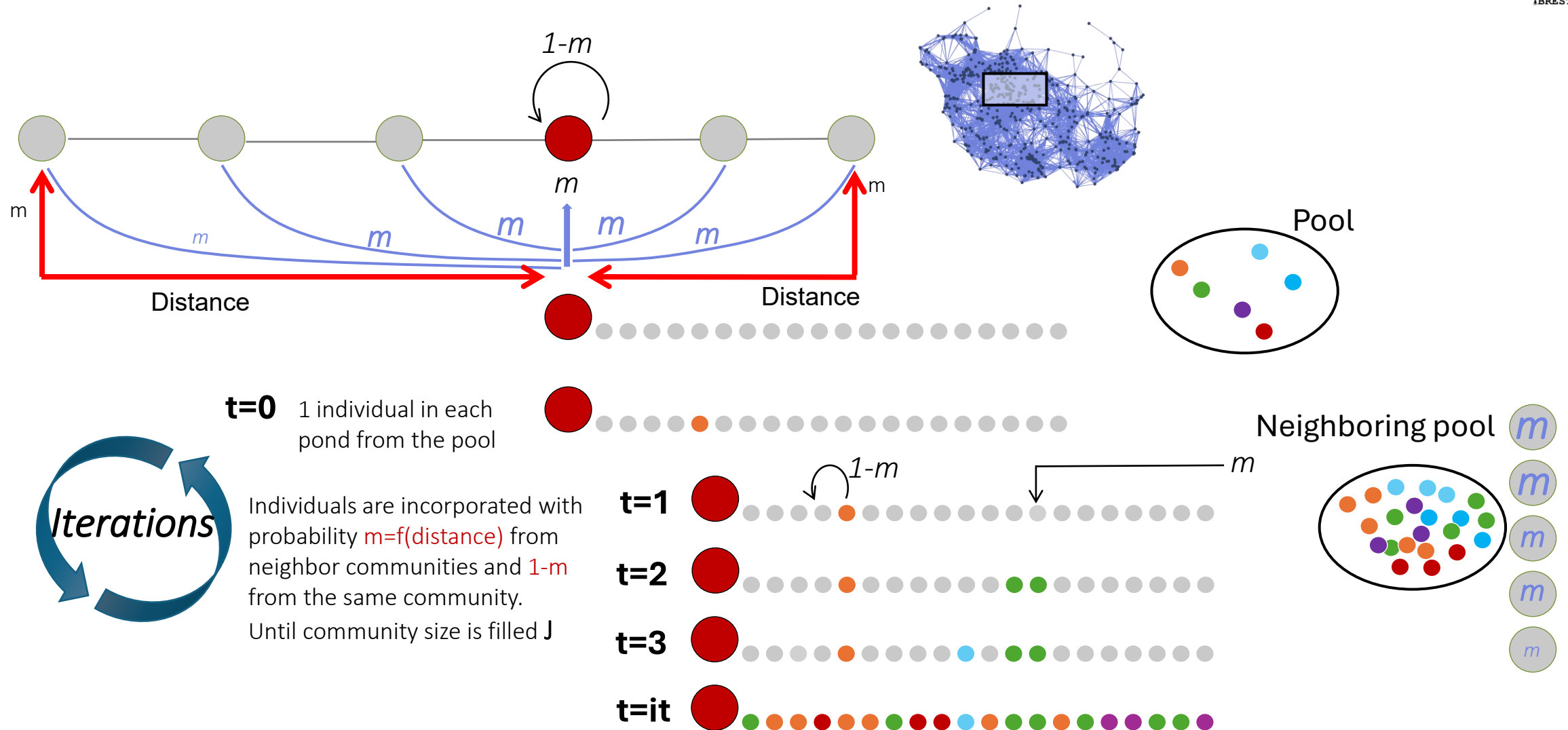




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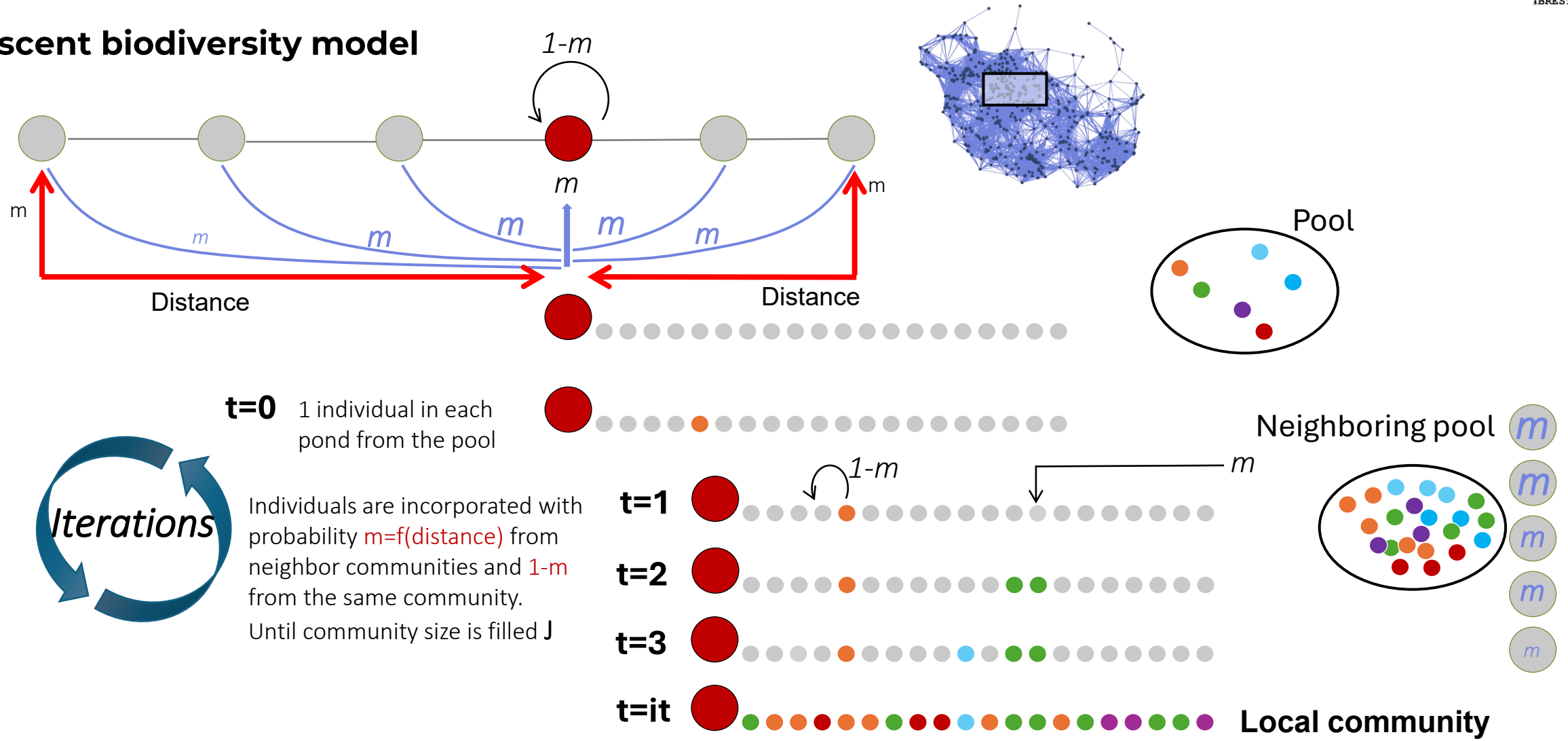


Connecting network structure with diversity



Connecting network structure with diversity

Coalescent biodiversity model



Connecting network structure with diversity

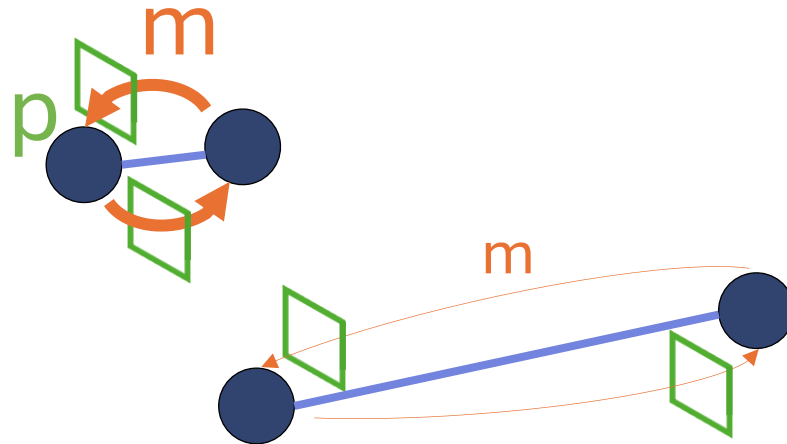
Coalescent biodiversity model

Local conditions

Penalty imposed to species that **diminishes their probability of establishment.**

Habitats with determined conditions (e.g. polluted) may impose a greater environmental filter for sensitive species.

Penalty imposed to species & habitats. If all species & habitats have the same penalty, there is no filter.



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Connecting network structure with diversity

Coalescent built

Community fully determined by the landscape structure (habitat position + size).
It summarizes regional structure **from a source-sink** point of view.

*Lottery biodiversity model

Will “exaggerate” the coalescent pattern and bring “extinction + colonization” dynamics.
Adds a temporal component. The more iterations the model is run the more submitted to drift communities are.



Network&Diversity_Part3.R

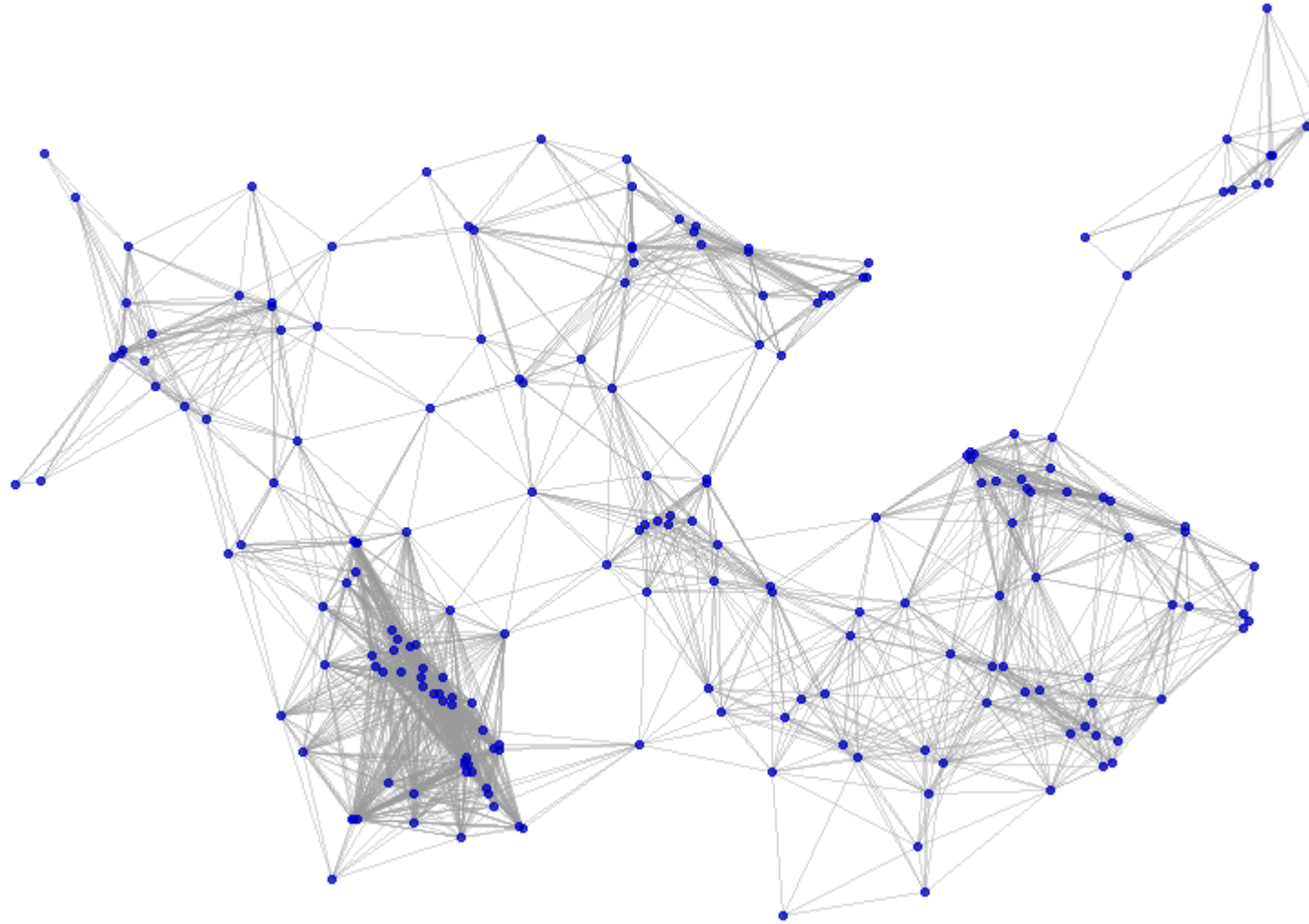
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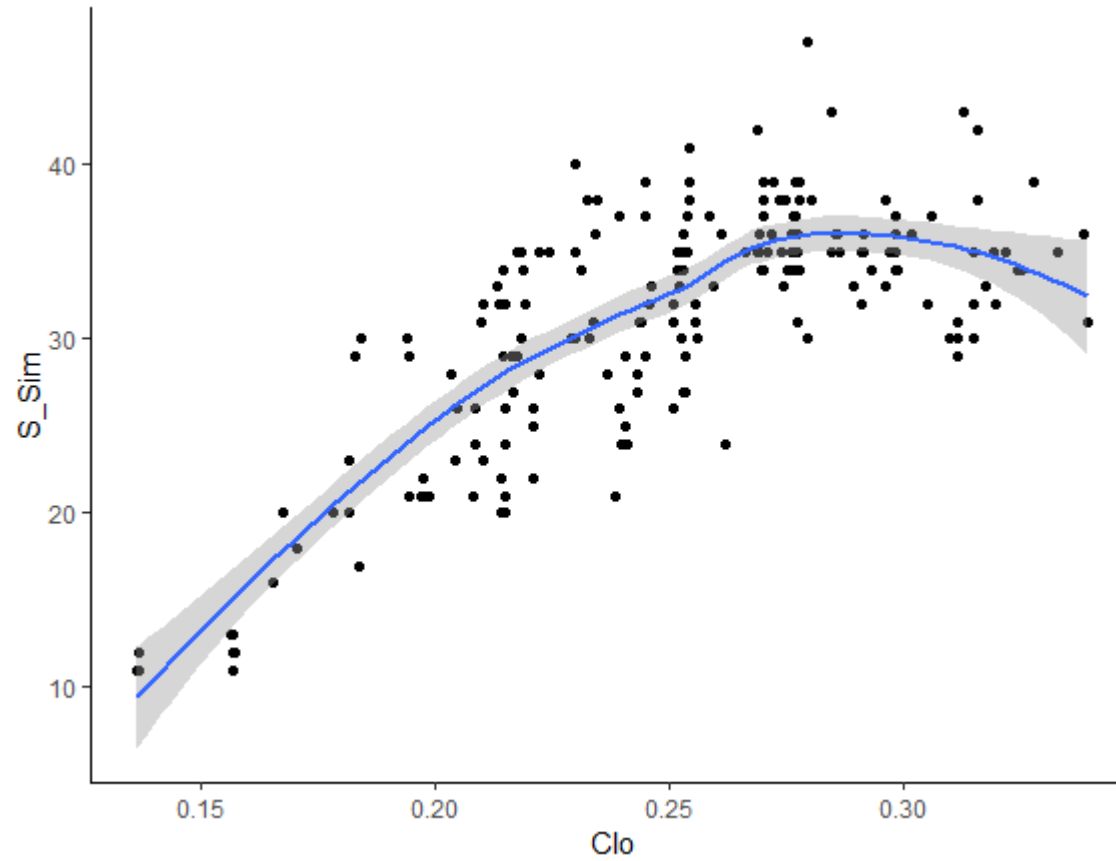
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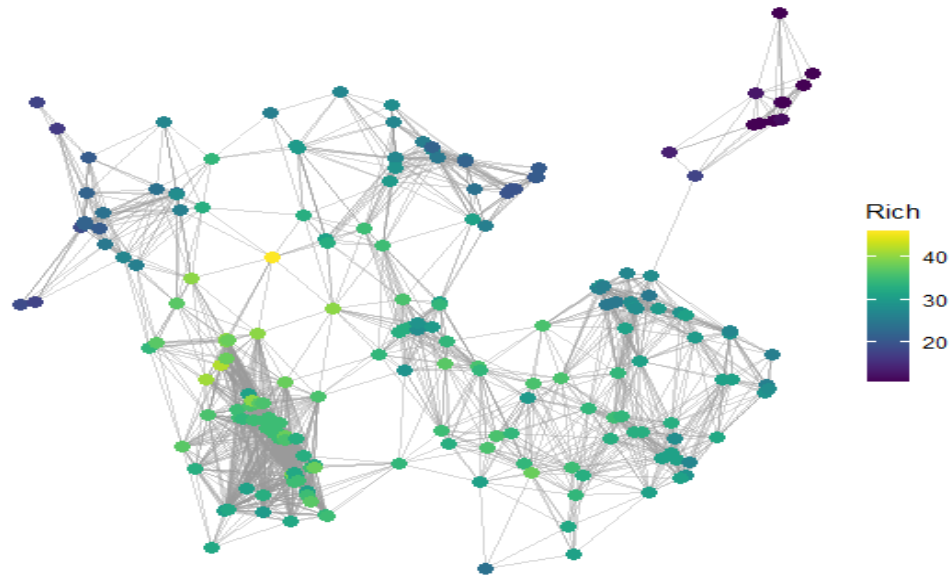
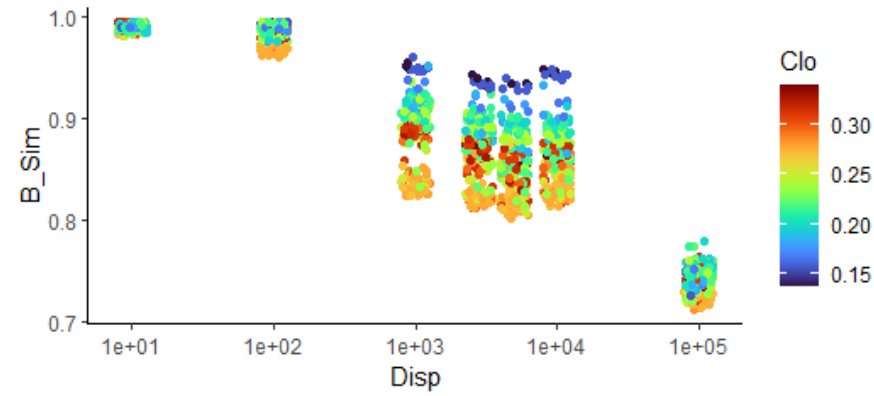
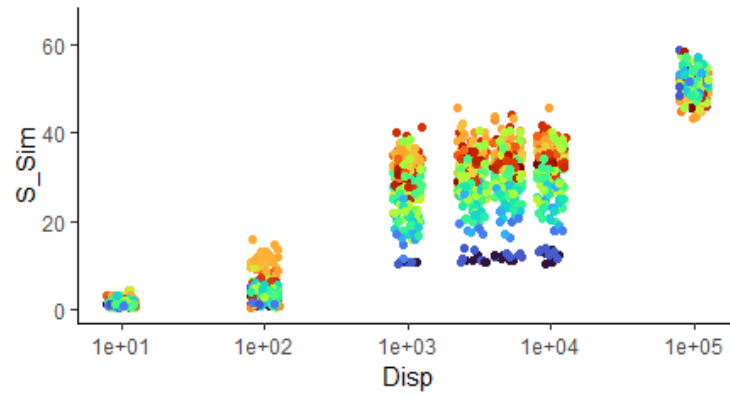
Connecting network structure with diversity



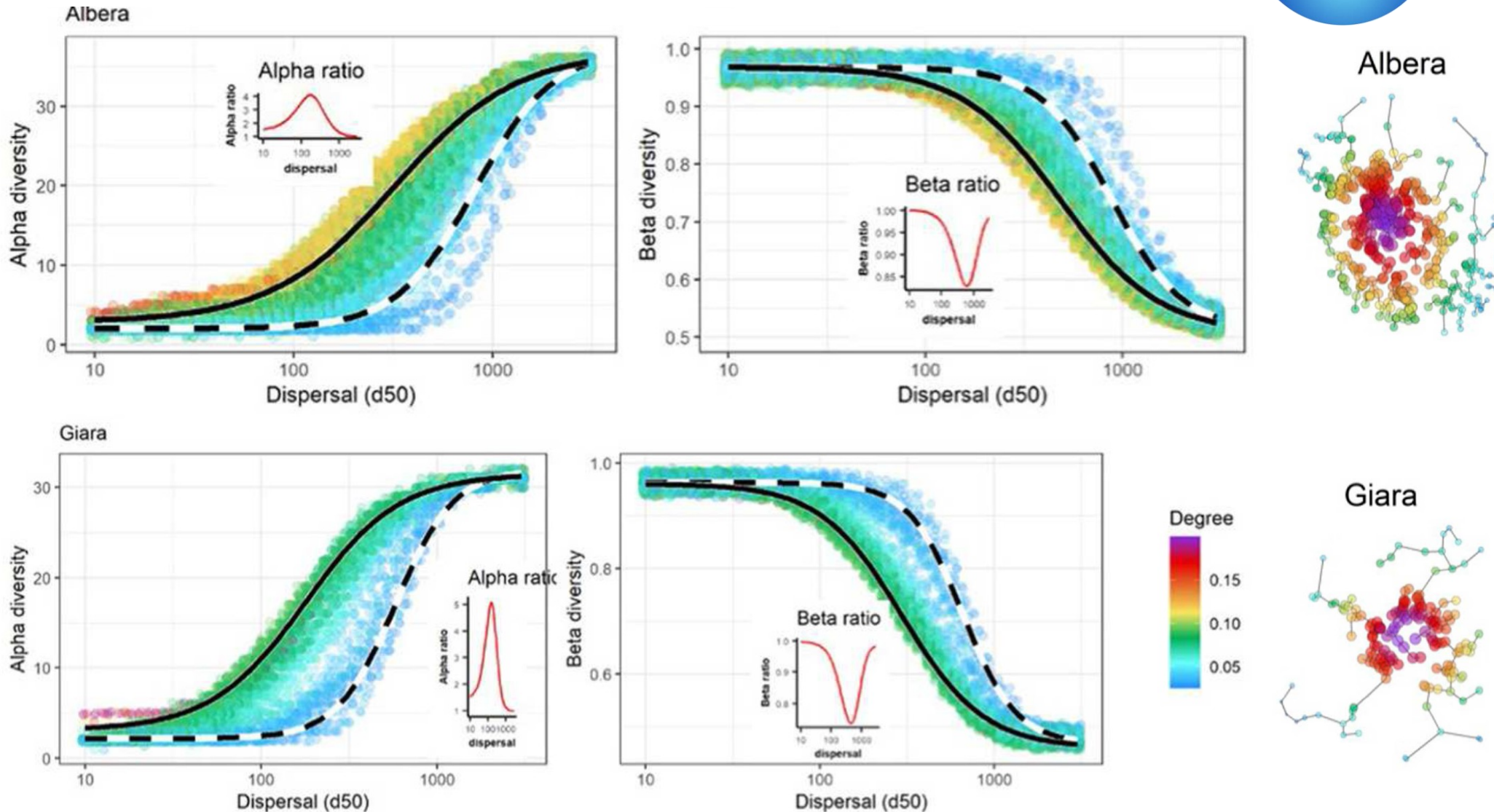
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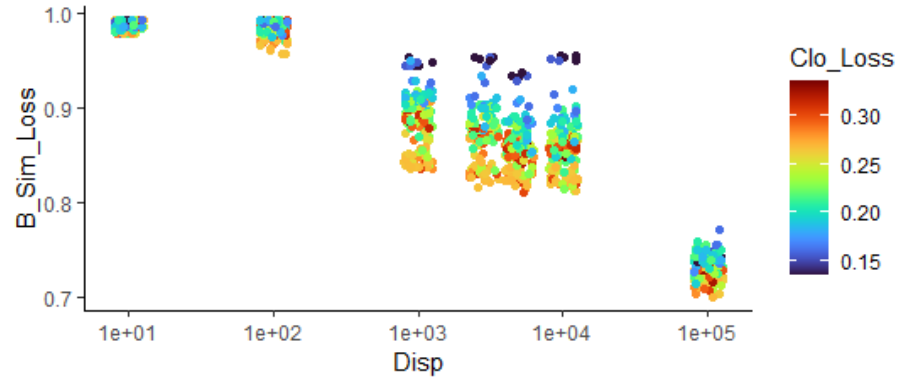
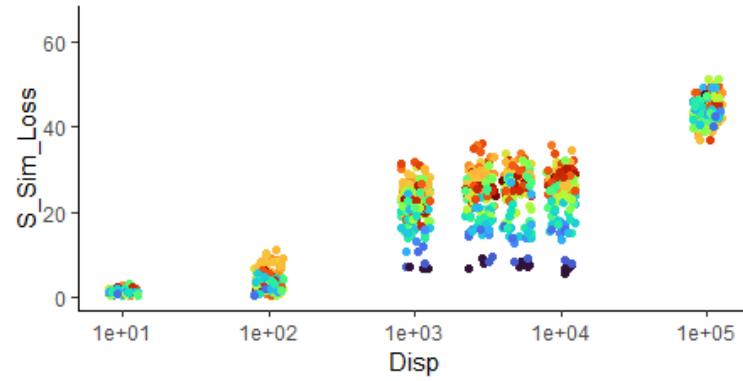
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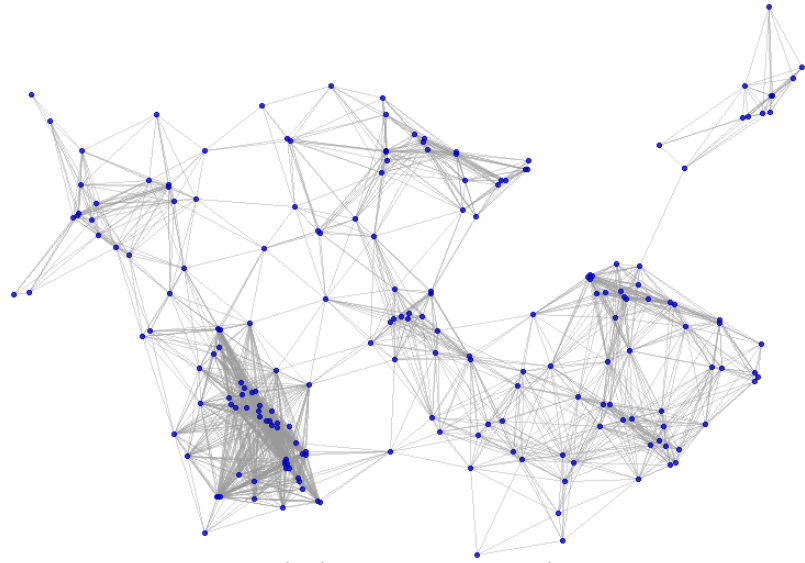
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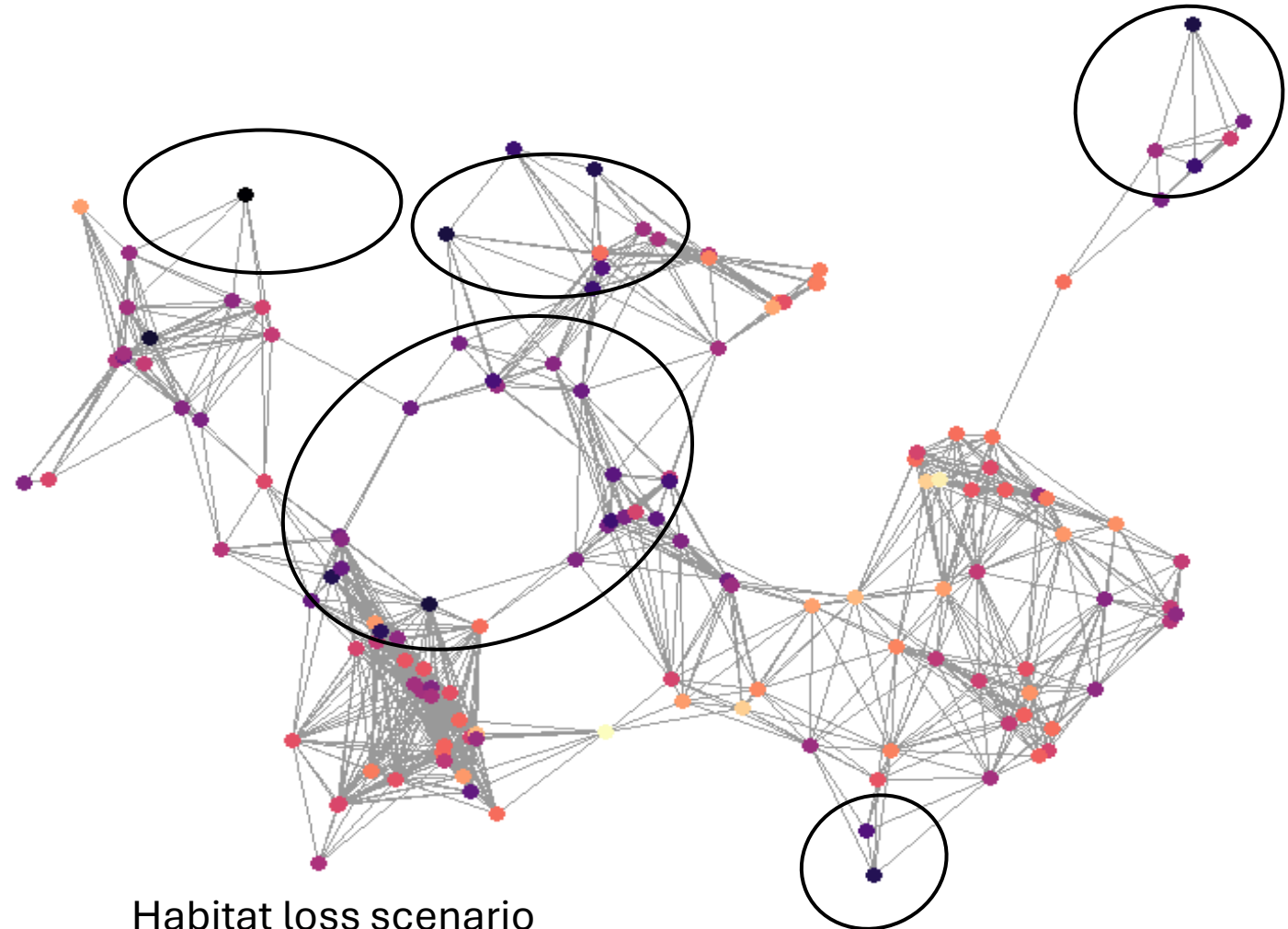
Connecting network structure with diversity



Connecting network structure with diversity



Original scenario



Habitat loss scenario

