

Managing for ecological surprises in metapopulations

Supplemental materials

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01 May 2019

Metapopulation model

Local & metapopulation dynamics

Our metapopulation is defined by a set of local populations N_p with time-dynamics that follows birth (i.e., recruitment R), immigration, death, and emigration (BIDE) processes:

$$N_{it} = R_{it}\epsilon_{it} + I_{it} - D_{it} - E_{it}$$

where N_{it+1} is the number of adults in patch i at time t , R_{it} is number of recruits, I_{it} is number of recruits immigrating into patch i from any other patch, D_{it} is number of recruits that die due to disturbance regime, E_{it} is the number of recruits emigrating from patch i into any other patch, and ϵ_{it} is stochasticity in recruitment.

Resource monitoring often occurs at the scale of the metapopulation, hence we define metapopulation adults as:

$$MN_t = \sum_{i=1}^{N_p} N_{it}$$

with metapopulation recruits:

$$MR_t = \sum_{i=1}^{N_p} R_{it}$$

Local patch recruitment at time t depended on adult densities at $t-1$ and followed a reparameterized Beverton-Holt function:

$$R_{it} = \frac{\alpha_i N_{it-1}}{1 + \frac{\alpha_i - 1}{\beta_i} N_{it-1}}$$

where α_i is the recruitment compensation ratio and β_i is local patch carrying capacity.

For example, in a two patch model that varies α_i and β_i parameters such that

```
alpha <- c(2, 4)
beta <- c(100, 200)
```

Management often monitors metapopulation resources as the aggregate of all local populations. In this way, recruitment compensation from local patches α_i gets averaged across the metapopulation leading mean compensation $\bar{\alpha}$ of 3. Likewise, the total carrying capacity of the metapopulation $\bar{\beta}$ becomes the summation of local patch carrying capacities β_i , which is 300. This scale of monitoring generates the following local patch and metapopulation dynamics:

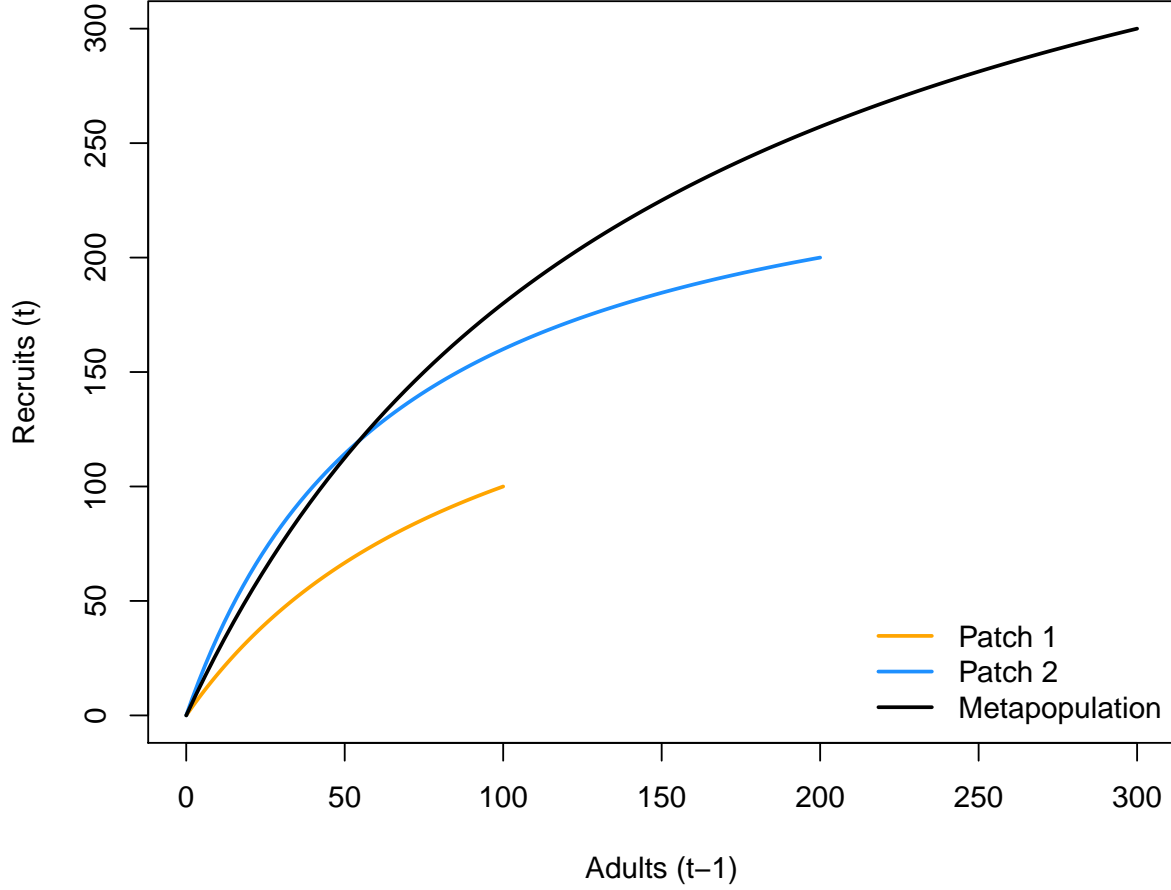


Figure 1: Metapopulation and local patch recruitment dynamics.

Creating the spatial networks

The next aspect to our metapopulation model is connecting the set of patches to one another. We need to specify the number of patches, their arrangements (i.e., connections), and how far apart they are from one another. We followed some classic metapopulation and source-sink arrangements to create four networks that generalize across a few real-world topologies: a linear habitat network (e.g., coastline), a dendritic or branching network (e.g., coastal rivers), a star network (e.g., mountain & valley), and a complex network (e.g., terrestrial plants).

To make networks comparable, each spatial network type needs the same leading parameters (e.g., N_p and \bar{d}). In this case for number of patches, we set N_p to 16 and \bar{d} to 1 unit (distance units are arbitrary). We used the `igraph` package and some custom code to arrange our spatial networks as the following:

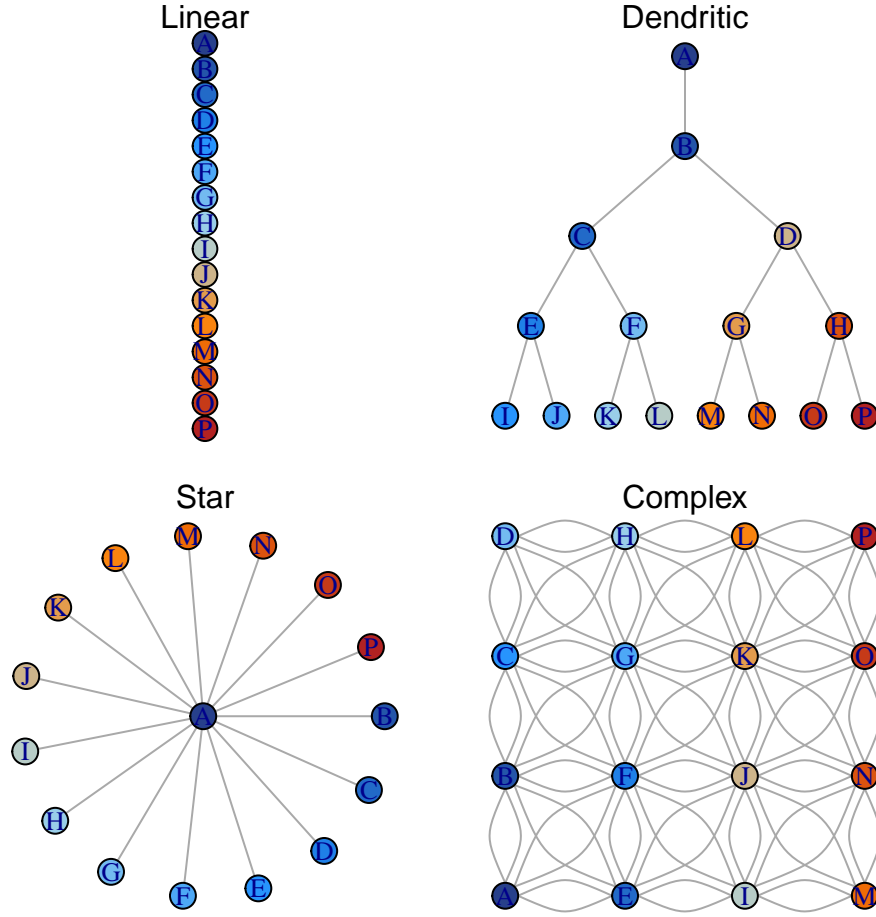


Figure 2: Four spatial network topologies.

Note that distances between each connecting patch (the links between nodes) in the above networks are equal.

An example dispersal matrix for the complex network:

##	A	B	E	F	C	G	D	H	I	J	K	L	M	N	O	P
## A	0	1	1	1	2	2	3	3	2	2	2	3	3	3	3	3
## B	1	0	1	1	1	1	2	2	2	2	2	2	3	3	3	3
## E	1	1	0	1	2	2	3	3	1	1	2	3	2	2	2	3
## F	1	1	1	0	1	1	2	2	1	1	1	2	2	2	2	2
## C	2	1	2	1	0	1	1	1	2	2	2	2	3	3	3	3
## G	2	1	2	1	1	0	1	1	2	1	1	1	2	2	2	2
## D	3	2	3	2	1	1	0	1	3	2	2	2	3	3	3	3
## H	3	2	3	2	1	1	1	0	3	2	1	1	3	2	2	2
## I	2	2	1	1	2	2	3	3	0	1	2	3	1	1	2	3
## J	2	2	1	1	2	1	2	2	1	0	1	2	1	1	1	2
## K	2	2	2	1	2	1	2	1	2	1	0	1	2	1	1	1
## L	3	2	3	2	2	1	2	1	3	2	1	0	3	2	1	1
## M	3	3	2	2	3	2	3	3	1	1	2	3	0	1	2	3
## N	3	3	2	2	3	2	3	2	1	1	1	2	1	0	1	2
## O	3	3	2	2	3	2	3	2	2	1	1	1	2	1	0	1
## P	3	3	3	2	3	2	3	2	3	2	1	1	3	2	1	0

Dispersal

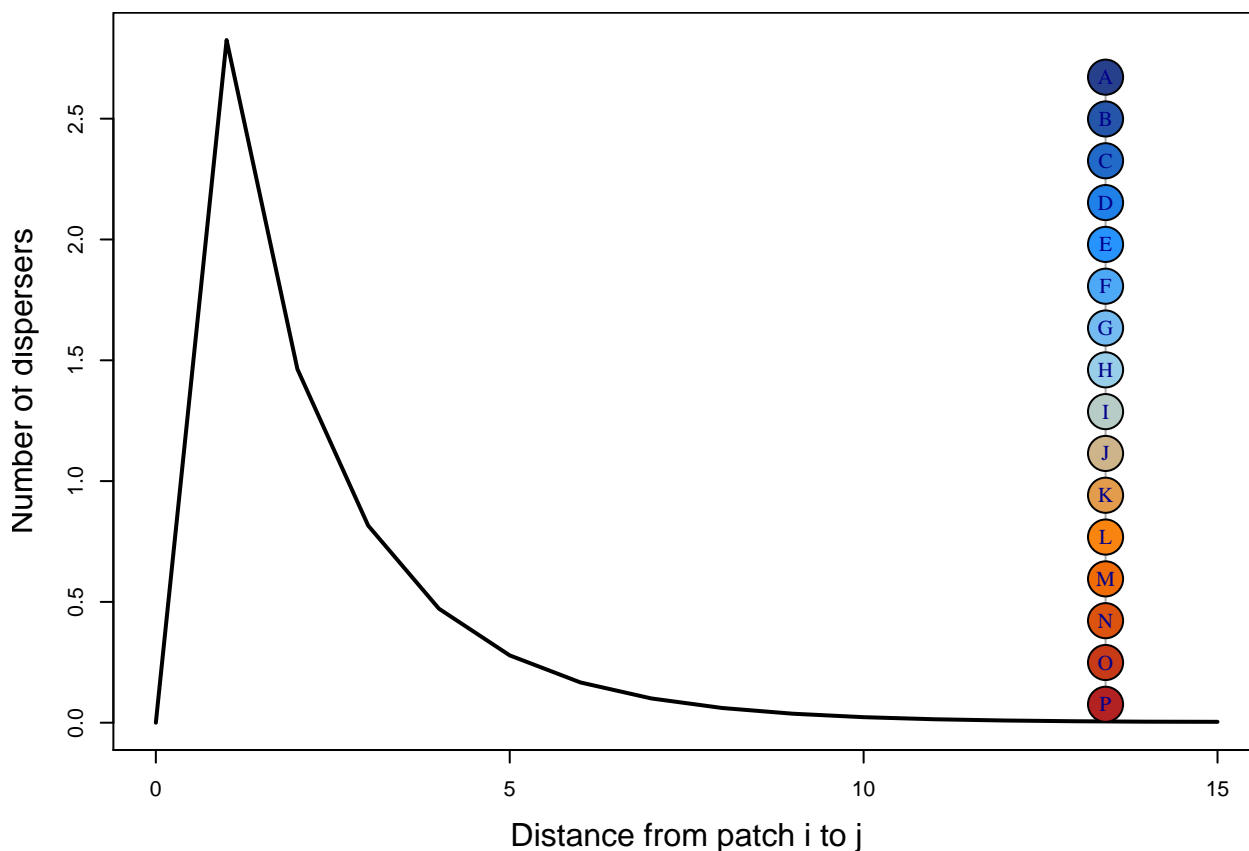
Dispersal from patch i into patch j depends on constant dispersal rate ω (defined as the proportion of total local recruits that will disperse) and an exponential distance-decay function between i and j with distance cost to dispersal m following:

$$E_{ij(t)} = \omega R_{it} p_{ij}$$

with probability of dispersal from patch i into patch j :

$$p_{ij} = \frac{e^{-md_{ij}}}{\sum_{\substack{j=1 \\ j \neq i}}^{N_p} e^{-md_{ij}}}$$

where d_{ij} is the pairwise distance between patches and E_{ij} is the total dispersing animals from patch i into patch j . The summation term in the denominator normalizes the probability of moving to any patch to between 0 and 1. With $\bar{d} = 1$, $m = 0.5$, $\omega = 0.1$, $R_{it} = 100$ in a linear network:



Recruitment stochasticity

Disturbance

Emergent outcomes

Scale of management & monitoring

Example scenarios