Managing for ecological surprises in metapopulations

Supplemental materials

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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

Resoure 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 a dult 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

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 $\sum \beta_i$, which is 300. This scale of monitoring generates the following local patch and metapopulation dynamics:

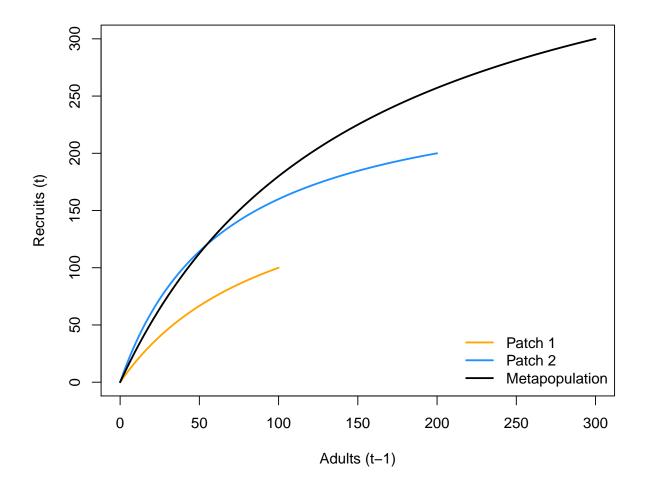


Figure 1: Metapopulation and local patch recruitment dynamics.

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:

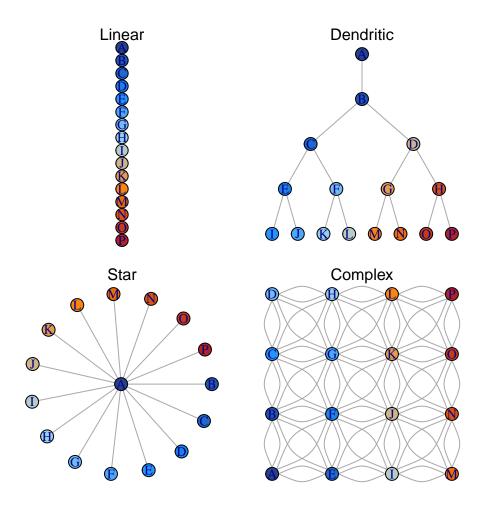


Figure 2: Four spatial network topologies.

Note that distances between neighbor patches in the above networks are equal.

An example dispersal matrix for the complex network:

```
ABEFCGDHIJKLMNOP
## A O 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
  0 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:

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

where E_{ij} is the total dispersing animals from patch i into patch j and probability of dispersal between patches p_{ij} :

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

where d_{ij} is the pairwise distance between patches. 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:

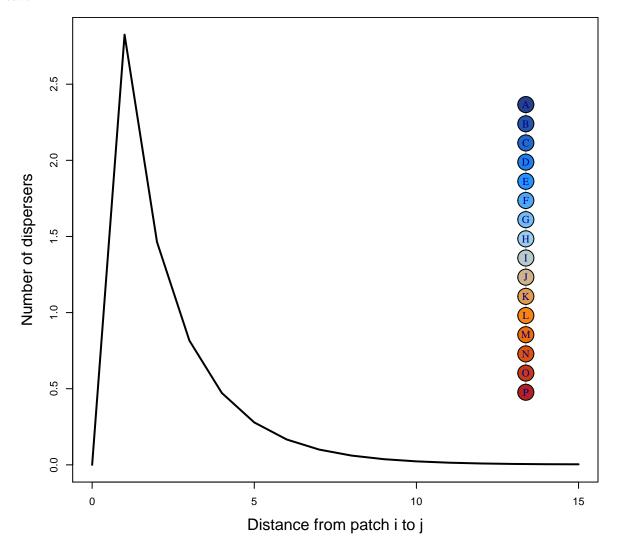


Figure 3: Example dispersal patterns across linear network.

Recruitment stochasticity

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Spatio-temporal correlations

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Disturbance regimes

In all scenarios, disturbance was applied after 50 years of equilibrating the metapopulaton at pristine conditions. At year 50+1, we applied the disturbance regime (the regime varied from uniform, localized, random, localized, extirpation, and localized, targeted - see Scenarios below). Disturbance immediately removes a set proportion of the metapopulation adults at that time (i.e., 0.9 of $MN_{t=51}$). Once applied, the metapopulation is no longer disturbed and spatio-temporal dynamics emerge naturally from these new conditions.

Emergent outcomes

We measured the following ecological outcomes that have direction application for conservation metrics.

- 1. Recovered & recovery rate after disturbance the number of simulations where metapopulation abundance averaged 1.0 carrying capacity for 5 consecutive years post-disturbance and the number of years it took to get there.
- 2. Extinction & extinction rate the number of simulations where metapopulation abundance was <0.05 carrying capacity and the mean number of years it took to get there.
- 3. Patch occupancy the mean number of patches with >0.05 local carrying capacity.
- 4. Spatial or temporal variance
- 5. Temporal variation in source, sink, pseudo-sink defined as:
 - a. Sources provide surplus recruits and net emigrants such that: $(R_{it} > N_{it})$ & $(E_{it} > I_{it})$
 - b. Sinks consume recruits and net immigrants such that: $(R_{it} < N_{it}) & (E_{it} < I_{it})$
 - c. Pseudo-sinks would provide surplus recruits in the absence of dispersal such that $(R_{it} > N_{it})$ but $(R_{it} + E_{it}) < (R_{it-1} + I_{it-1} E_{it-1})$
- 6. Fit stock-recruitment model to aggregate of metapopulation to estimate:
 - a. Relative bias in recruitment compensaton ratio compared to true metapopulation average
 - b. Relative bias in metapopulation carrying capacities compared to true sum of carrying capacities across metapopulation
 - c. Relative bias in expected recruitment production to true recruitment production across all patches

Monitoring & management at aggregate-scale

While true metapopulation dynamics are controlled by local patch dynamics and dispersal such that:

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

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

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

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

natural resoure managers often monitors and manages at the scale of the metapopulation. Hence, management at this scale inherently defines the stock-recruitment dynamics of the aggregate complex of patches (i.e., metapopulation) as:

$$MR_t = \frac{\hat{\alpha_t} M N_{t-1}}{1 + \frac{\hat{\alpha_t} - 1}{\hat{\beta_t}} M N_{it-1}}$$

where $\hat{\alpha}_t$ is the estimated compensation ratio averaged across the metapopulation at time t and $\hat{\beta}_t$ is the estimated carrying capacity of the entire metapopulation. Necessarily, these estimates emerge from monitoring data collected across all patches and are sensitive to the quality of the data and how local patches perform through time. For example, temporal shifts in productivity regimes may be masked if most of the data were sampled before the regime shift. To help surmount these issues, modern resource assessments use data weighting and penalties (i.e., priors) when fitting models to data.

In our assessment, we weighted recent years of sampling over more distant years such that:

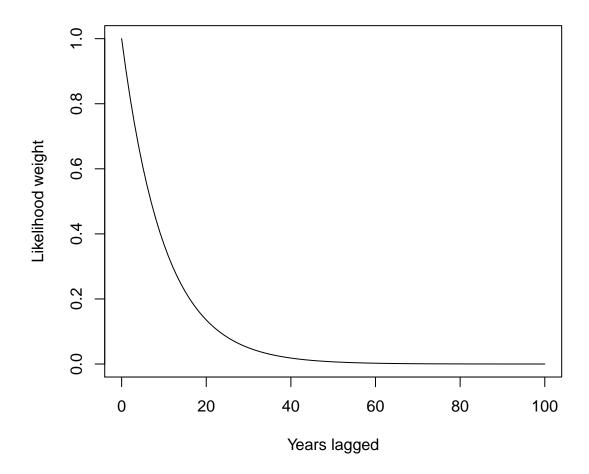


Figure 4: Likelihood weighting for samples collected over time from current year of sampling.

Furthermore, we used penalized normal likelihoods on both $\hat{\alpha}_t$ and $\hat{\beta}_t$ such that:

```
\begin{split} \hat{\alpha_t} \sim N(\mu = \hat{\alpha_{t-1}}, \sigma = 3\hat{\alpha_{t-1}}) \\ \text{and} \\ \hat{\beta_t} \sim N(\mu = \hat{\beta_{t-1}}, \sigma = 3\hat{\beta_{t-1}}) \end{split}
```

where $\mu = \hat{\alpha_{t-1}}$ and $\mu = \hat{\beta_{t-1}}$ represents the best estimates from the previous assessment and the 3 in the σ term represents a 300% coefficient of variation. We used these penalized likelihoods to fit the above aggregate stock-recruitment model with lognormal error to the metapopulation stock-recruit data collected at time t. We used the following function and fitted to the below θ parameters (termed theta in the function optim() using the L-BFGS-B optimizer with a lower bound on $\hat{\alpha}$ of 1.01 (i.e., constrained to be at least above replacement).

```
SRfn <- function(theta) {
    a.hat <- theta[1]
    b.hat <- exp(theta[2])
    sd.hat <- exp(theta[3])
    rec.mean <- (a.hat * spawnRec$spawners)/(1 + ((a.hat - 1)/b.hat) * spawnRec$spawners)
    # negative log likelihood on recruitment parameters
    nll <- -1 * sum(dlnorm(spawnRec$recruits, meanlog = log(rec.mean), sdlog = sd.hat,
        log = TRUE) * spawnRec$weights, na.rm = TRUE)
    # penalized likelihood on estimated alpha
    penalty1 <- -dnorm(a.hat, alphaLstYr, 3 * alphaLstYr, log = TRUE)
    # penalized likelihood on estimated carrying capacity
    penalty2 <- -dnorm(b.hat, metaKLstYr, 3 * metaKLstYr, log = TRUE)
    jnll <- sum(c(nll, penalty1, penalty2), na.rm = TRUE)
    return(jnll)
}</pre>
```

This above function allows us to assess the bias in $\hat{\alpha}_t$, $\hat{\beta}_t$, and \hat{MR}_t compared to true $\bar{\alpha}$, $\bar{\beta}$, and MR_t across the metapopulation. This then allows us to see how much information management & monitoring programs are missing when they assess metapopulations at the aggregate (rather than local) scales.

Scenarios

We tested all combinations of the following eight processes (below) and ran a 100 bootstraps per scenario to estimate the mean for each of the above outcomes.

- 1. Homogenous and spatially variable recruitment compensation ratio across patches, i.e. intrinsic rate of population growth (α_i) .
- 2. Homogenous and spatially variable local carrying capacity across patches, i.e. asymptote of expected recruits at high adult densities (β_i)
- 3. Disturbances where a proportion of individuals removed from metapopulation (e.g., 0.90) occurs.
 - a. uniform random individuals removed at equal vulnerability across all patches.
 - b. *localized, random* random individuals removed from randomly selected subset of patches (as long as target loss can be achieved in subset)
 - c. *localized, extirpation* total extirpation of randomly selected subset of patches (as long as target loss can be achieved in subset)
 - d. *localized*, *targeted* random individuals removed from subset of patches selected based on productivity (i.e., most productive patches are targeted so long as target loss can be achieved in subset).
- 4. Density-independent dispersal rates ω from 0 to 20% of individuals within a patch will disperse.

- 5. Topology of the spatial networks with linear, dendritic, star, and complex networks. Each network with N_p of 16 and distance between patches \bar{d} of 1.
- 6. Stochastic recruitment deviates from low, medium, high coefficient of variation on lognormal error. Generate stochasticity in time-dynamics via random recruitement deviates away from expected.
- 7. Temporal correlation in recruitment deviates from low, medium, high correlation (i.e., good year at time t begets good year at time t+1).
- 8. Spatial correlation in recruitment deviates among patches from low, medium, to high correlation (i.e., neighboring patches go up or down together).

Example results

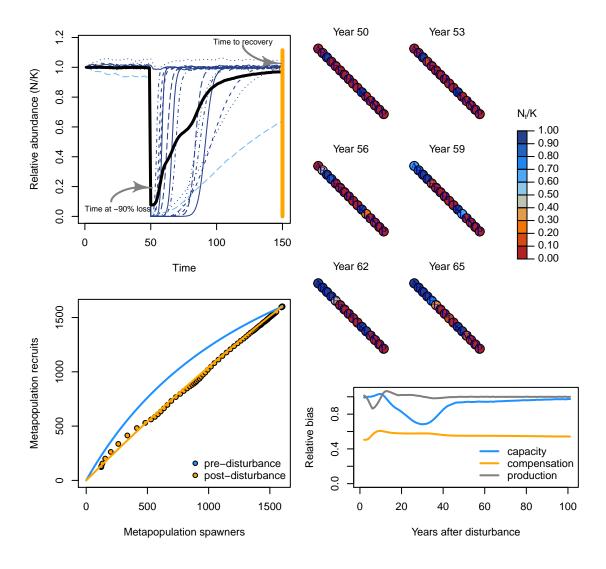


Figure 5: Spatial recovery regime of metapopulation with linear topology through time (top left) and space (top right). Recruitment dynamics before and 10 years after disturbance (bottom left). Relative bias in aggregate-scale estimates of carrying capacity, compensation ratio, and recruitment production in recovery phase (bottom right).

Simulation test & bootstrap

General patterns