

Comparing C^r Values with Network Perturbation Results - Focal Species: Plants

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1 Plant Notes and Generalized C^r

We model a population of Buttercup plant building our model based on the formulation in *A general modeling framework for describing spatially structured population dynamics* by Sample et. al.

We track the population at eight breeding nodes. The population is modeled through three times steps each year:

- Summer, seeds and plants at all nodes.
- Autumn/Winter, seeds and plants at all nodes.
- Spring, seeds and plants at all nodes.

We use a generalized network model formulation where we define one time step as

$$\vec{\mathbf{N}}_{t+1} = \mathbf{A}_t \vec{\mathbf{N}}_t \quad (1)$$

Where $\vec{\mathbf{N}}_t$ contains the population at each of the n nodes and c classes at time t . The matrix \mathbf{A}_t contains all of the demographic node update and migration data for time t . (Christine and Joanna have a draft of a paper with this construction).

Within the plant code we calculate the C^r values for both the origin and the intermediate nodes during each season. We use a generalized one equation definition of C^r in matrix form.

$$\vec{\mathbf{C}}_t = \left(\prod_{\tau=t}^{t+s-1} \mathbf{A}_\tau^T \right) \vec{\mathbf{1}}_{nc} \quad (2)$$

where s is the total number of seasons in one annual cycle, for the plants $s = 3$ and $\vec{\mathbf{C}}_t$ is a column vector that contains the C^r values at each node for each class for focal season, or time, t .

We note that because the code is run to equilibrium, the overall growth rate of the network is equal to one. We can calculate the growth rate using:

$$\lambda_t = \frac{\vec{N}_t^T}{N_t^{tot}} \vec{C}_t \quad (3)$$

where N_t^{tot} is the sum of the equilibrium population values across all nodes and all classes in the network and is represented by the following summation

$$N_t^{tot} = \sum_{r=1}^n \sum_{x=1}^c N_{r,t}^x \quad (4)$$

2 Density Dependence in the Plant Population

The nodal update function f is density dependent. f is defined by:

$$f_{i,t}^p = \underbrace{s_{i,t}^p \cdot N_{i,t}^p}_{\text{plants that survive}} + \underbrace{\Psi_{i,t} \cdot T_{i,t} \cdot s_{i,t}^s \cdot N_{i,t}^s}_{\text{seeds that successfully transition to plants}} \quad (5)$$

$s_{i,t}^p$, $T_{i,t}$, and $s_{i,t}^s$ are all constants. $\Psi_{i,t}$ is the density-dependent postgermination survival rate. $\Psi_{i,t}$ is given by:

$$\Psi_{i,t} = \psi_{i,t} \cdot e^{-(N_{i,t}^p + T_{i,t} \cdot s_{i,t}^s \cdot N_{i,t}^s) / K_i} \quad (6)$$

K_i is the carrying capacity at node i and $\psi_{i,t}$ is the maximum germination rate, which is constant, but varies by season.

3 Baseline Simulation

The baseline simulation gives us identical results to *A general modeling framework for describing spatially structured population dynamics* by Sample et. al. We find that the long term carrying capacity for each node during the nonbreeding season, including all classes, is

```
season 1 : k= 43.50627 42.92571 43.32792 43.32185 43.3219 43.32762 42.92347 43.50624
season 2 : k= 470.9006 513.6243 493.1009 493.3224 493.3228 493.0962 513.5998 470.8987
season 3 : k= 145.3853 170.1022 158.4498 158.1963 158.1963 158.4482 170.0959 145.3845
```

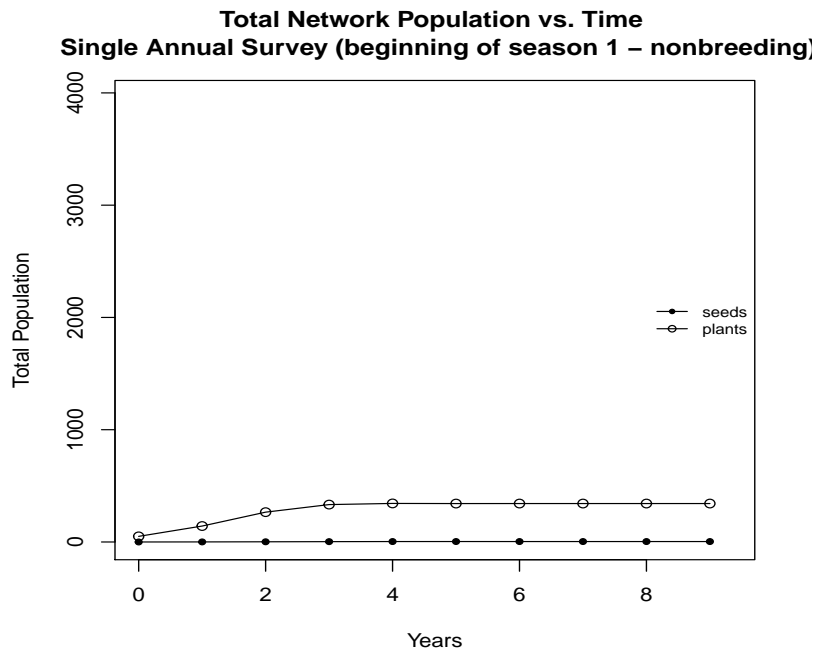


Figure 1: Baseline results: population over time at nonbreeding node.

4 Perturbation Experiment - Perturbing Node Survival Rates

To investigate the utility of the metrics as an indicator of the change in carrying capacity K we consider the following perturbations to the survival rate at each node:

$$PERT = .9, .8, .7, .6, .5$$

Some notes about the simulations:

- All simulations are run to equilibrium with an error tolerance of within 1 plant, meaning the network growth rate $\lambda = 1$.
- We only carry out negative perturbations because positive perturbations will cause survival rates to exceed 1.

5 Plant Plots - June 2018 - to Match Pintail Outputs

We will consider metrics for the x-axis of our graph:

- Baseline C^r vs K perturbations - this is the population weighted C^r value averaged across classes and seasons Figure 2
- Baseline W^r vs K perturbations - the population proportions averaged across seasons Figure 3
- Baseline D_s vs K perturbations - the proportional dependence as defined in Bagstad et al, generalizing to account for juveniles: Figure 4
- NEW - Baseline Criticality KR_i vs K perturbations - defined as the network growth rate in the absence of node i using a population weighted average across seasons: Figure 5. Note that because we are at equilibrium the baseline network growth rate is $\lambda = 1$.

5.1 Population weighted average C^r

In order to be consistent across case studies we decide to calculate C^r using a population weighted average. The general matrix form of C^r results in a C^r values for each node, class, and season. To combine these into a single full year network wide C^r value we first do a population weighted average by class within each node and then do a population weighted average across seasons. This results in a single C^r values for each node.

The plant model has three seasons (Summer, Autumn/Winter, Spring) and two classes (seeds and plants).

Range of Network Carrying Capacities (K) vs Baseline CR

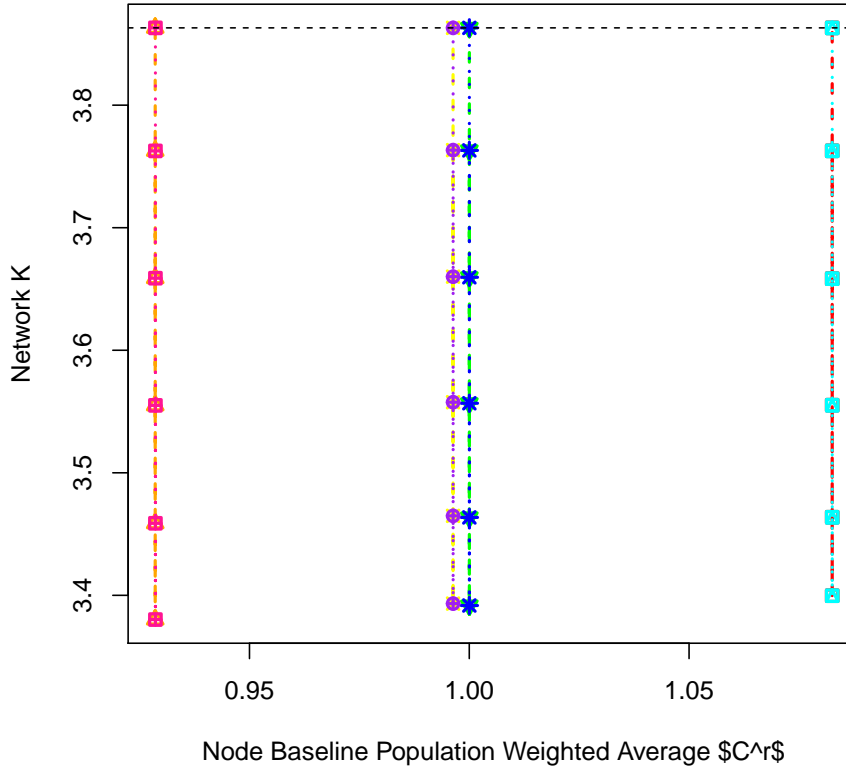


Figure 2: Perturbation results: Range of K values after perturbations to each node, X-axis represents **baseline population weighted C^r values** for each node

Node 1	□	year-round
Node 2	△	year-round
Node 3	×	year-round
Node 4	▽	year-round
Node 5	★	year-round
Node 6	⊕	year-round
Node 7	□	year-round
Node 8	▽	year-round

5.2 Population Distribution W^r

W^r is calculated as the percent of the total population residing at a node. This calculation results in W^r values for each node during each seasons. To get a consistent value for the network, we average (not weighted) across the seasons. The final numbers should sum to one.

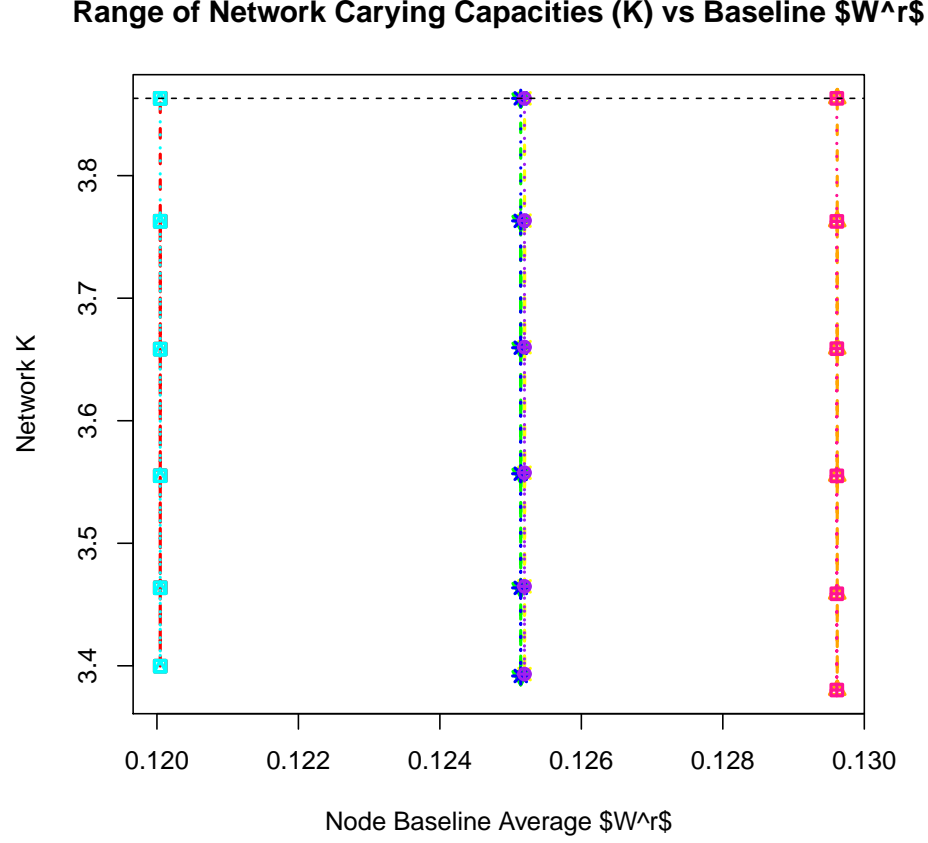


Figure 3: Perturbation results: Range of K values after perturbations to each node, X-axis represents baseline average W^r value for each node

Node 1	□	year-round
Node 2	△	year-round
Node 3	×	year-round
Node 4	▽	year-round
Node 5	★	year-round
Node 6	⊕	year-round
Node 7	□	year-round
Node 8	▽	year-round

5.3 Proportional Dependence D_s

D_s is calculated following Bagstad et al. We first find the population weighted C^r values for each of the seasons and then average across the annual cycle.

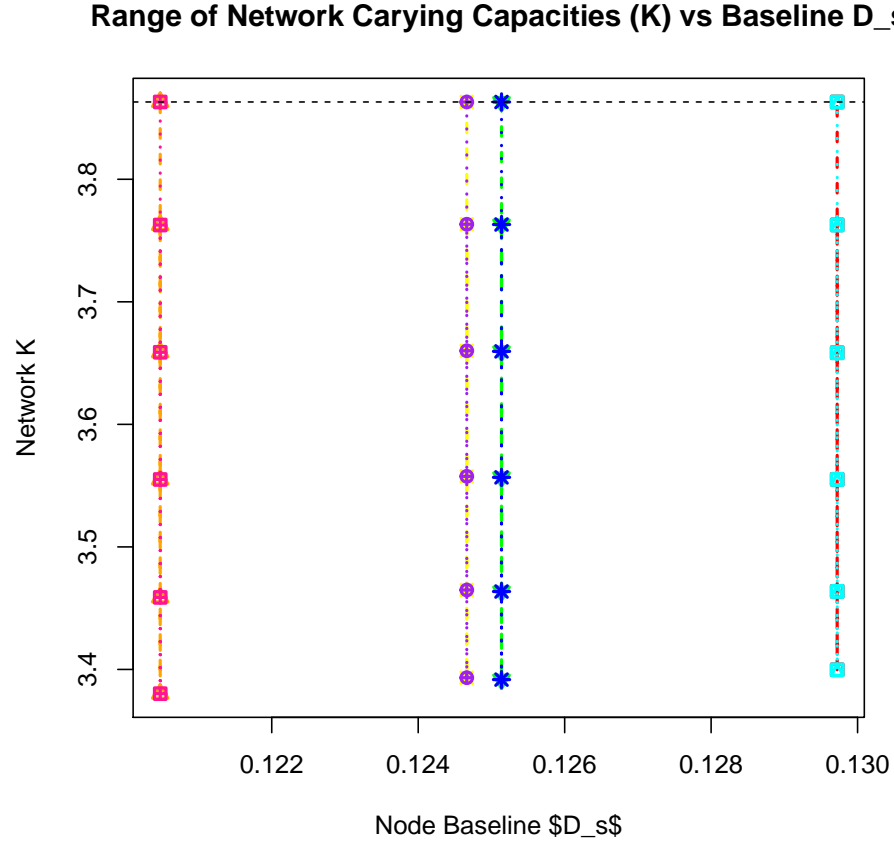


Figure 4: Perturbation results: Range of K values after perturbations to each node, X-axis represents baseline D_s value for each node

Node 1	□	year-round
Node 2	△	year-round
Node 3	×	year-round
Node 4	▽	year-round
Node 5	★	year-round
Node 6	⊕	year-round
Node 7	□	year-round
Node 8	▽	year-round

5.4 Criticality - NEW K^r

Criticality K^r is a new metric conceived of by Christine and her colleagues. The basic idea is that from C^r we can calculate the network growth rate λ . In the case of a population in equilibrium, like our simulations, $\lambda = 1$. Then one could also calculate a theoretical C^r value as if one of the nodes was removed. From this theoretical C^r value we could calculate a new network growth rate γ . We define criticality as

$$K^r = \lambda - \gamma$$

In other words K^r represents the amount of the network growth rate that flows through the focal node r . In our equilibrium case is $K^r = 1$ then all of the network growth rate must flow through r and removal of r reduces the new growth rate γ to zero. Alternatively, if $K^r = 0$ then none of the original network growth rate flows through r and removal of r does not change the growth rate, $\lambda = \gamma = 1$.

The K^r calculation results in criticality values for each node during each season. To get a single yearly value we use a population weighted average.

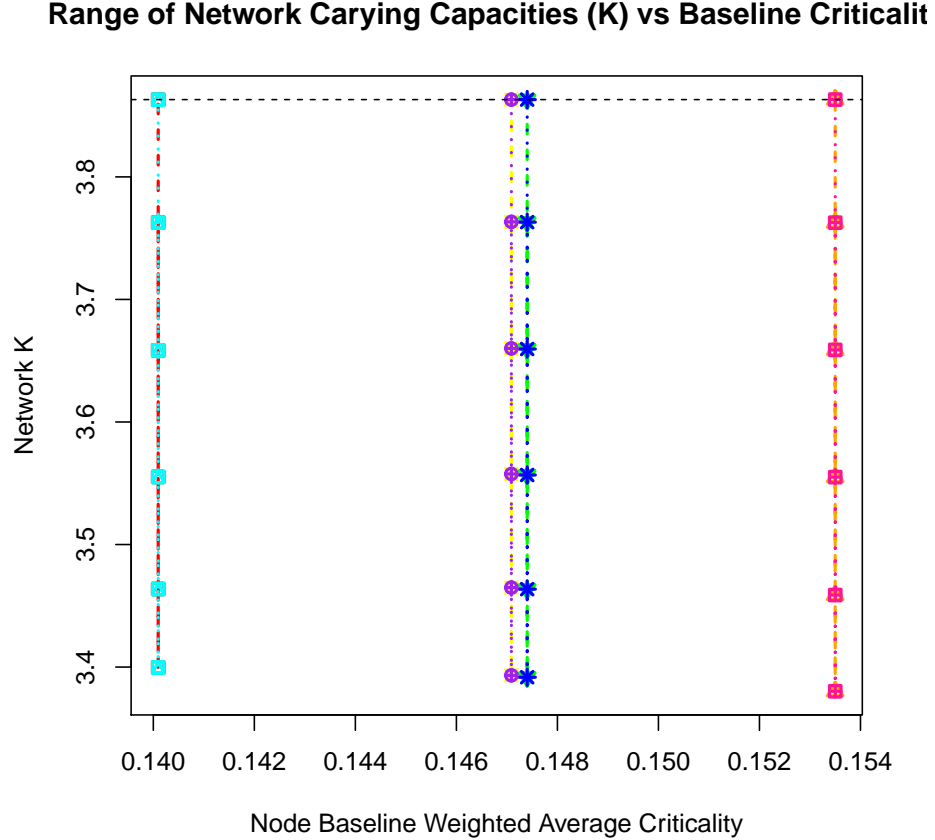


Figure 5: Perturbation results: Range of K values after perturbations to each node, X-axis represents baseline Criticality value for each node

Node 1	\square	year-round
Node 2	\triangle	year-round
Node 3	\times	year-round
Node 4	∇	year-round
Node 5	\star	year-round
Node 6	\oplus	year-round
Node 7	\square	year-round
Node 8	∇	year-round