

Forest Resource Sharing through Mycorrhizal Networks

The Fungal Friends

Chaitanya, Malou, Maurits, Steven

Theoretical Background



Mycorrhizal networks: The Wood Wide Web



The Biology

Obligate
symbiosis
fungi and trees

Networks link
roots different
trees

Scale-free
network

Theoretical Background



Mycorrhizal networks: The Wood Wide Web



The Biology

The Role of Mycorrhizal Networks

Resource Transfer

Sending defense signals

And more...

Theoretical Background



Mycorrhizal networks: The Wood Wide Web



The Biology

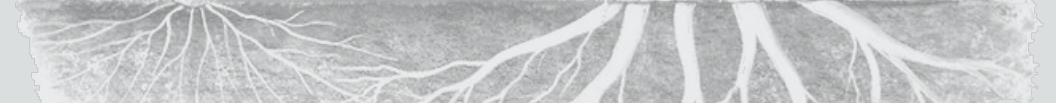
The Role of Mycorrhizal Networks

Motivation

Resilience and
Adaptation

Complex
Adaptive
System

Forestry in a
changing
climate

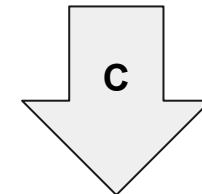


The Source-Sink Gradient Hypothesis

- Carbon Transfer along gradients
- Facilitation and Competition
- Survival young trees during stress



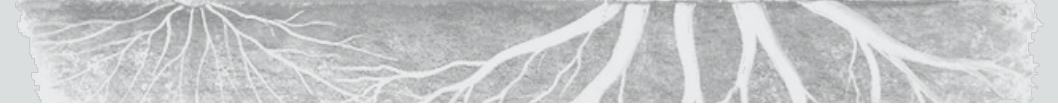
SOURCE



SINK



Hypothesis and Research Objective

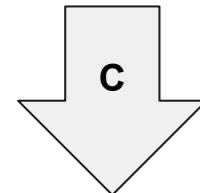


Research Objective

Investigate the dynamics of carbon source-sink gradients on a mycorrhizal network and how this can facilitate sapling growth and success.



SOURCE



SINK



Methods: Data Used



New Phytologist

Research

Architecture of the wood-wide web: *Rhizopogon* spp. genets link multiple Douglas-fir cohorts

Kevin J. Beiler^{1,2}, Daniel M. Durall¹, Suzanne W. Simard², Sheri A. Maxwell³ and Annette M. Kretzer⁴

¹Biology and Physical Geography Unit and SARAHs Centre, University of British Columbia Okanagan, Kelowna, BC V1V 1V7, Canada; ²Department of Forest Sciences, University of British Columbia, Vancouver, BC V6T 1Z4, Canada; ³Department of Biology, University of Western Ontario, London, ON N6A 5B8, Canada; ⁴SUNY College of Environmental Science and Forestry, Faculty of Environmental and Forest Biology, One Forestry Drive, Syracuse, NY 13210-2788, USA

Summary

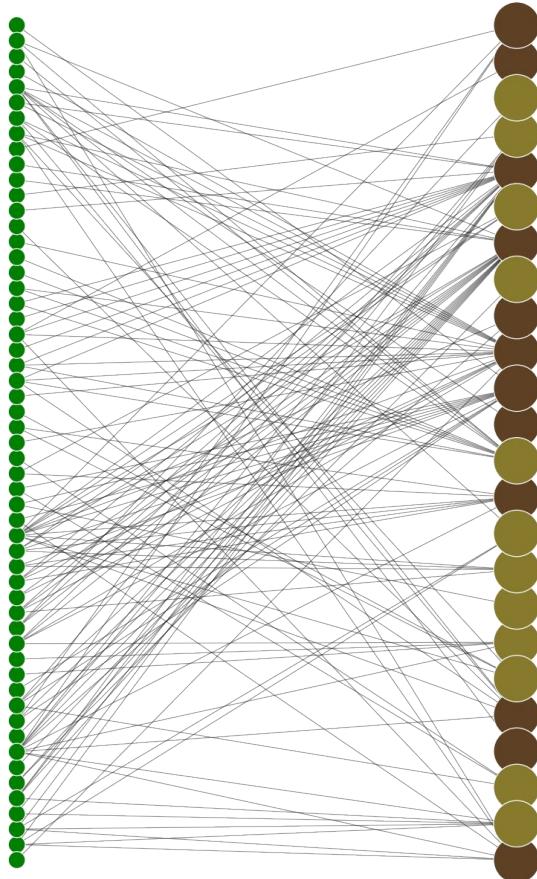
Author for correspondence:
Kevin J. Beiler
Tel: +1 250 826 1002
Email: Kevin.Beiler@gmail.com

Received: 17 June 2009
Accepted: 9 September 2009

- The role of mycorrhizal networks in forest dynamics is poorly understood because of the elusiveness of their spatial structure. We mapped the belowground distribution of the fungi *Rhizopogon vesiculosus* and *Rhizopogon vinicolor* and interior Douglas-fir trees (*Pseudotsuga menziesii* var. *glauca*) to determine the architecture of a mycorrhizal network in a multi-aged old-growth forest.
- Rhizopogon* spp. mycorrhizas were collected within a 30 × 30 m plot. Trees

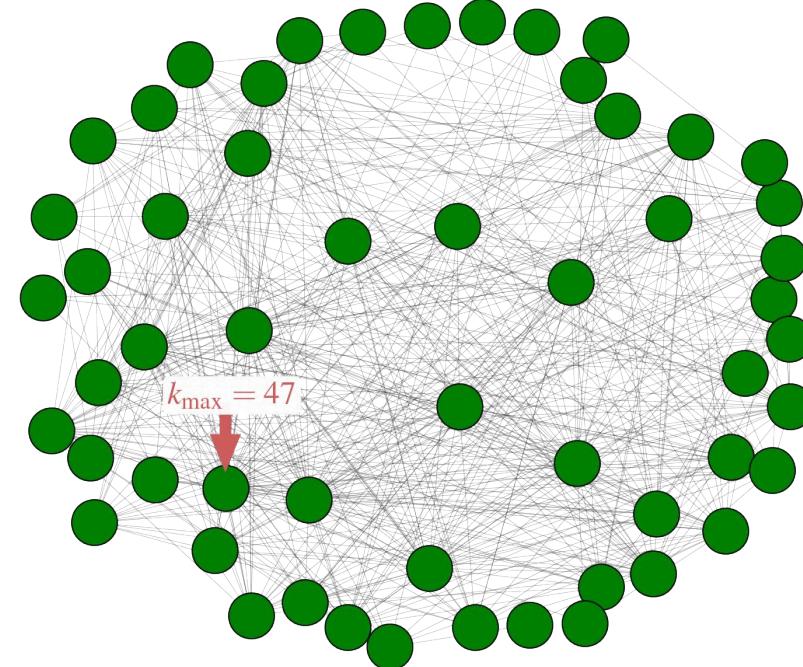
Methods: Network Set-up

Individual trees

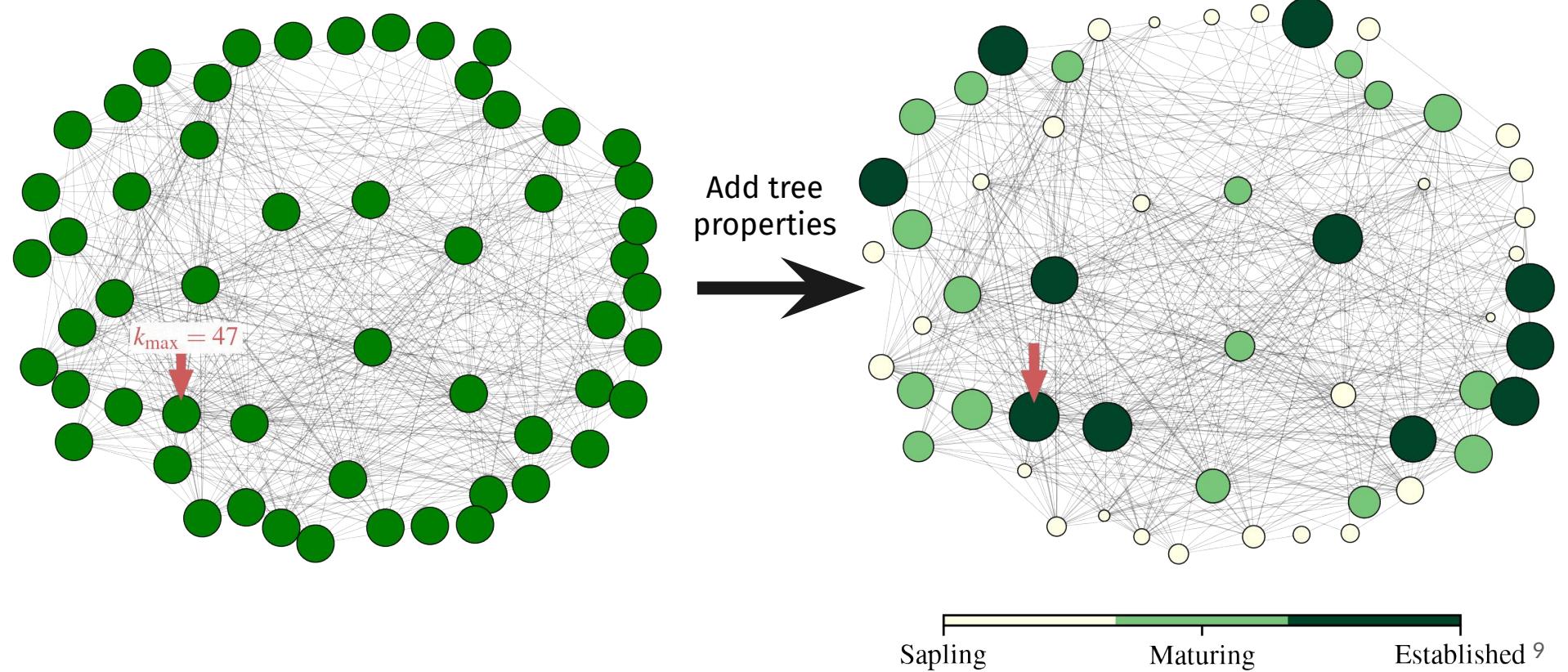


Individual fungi

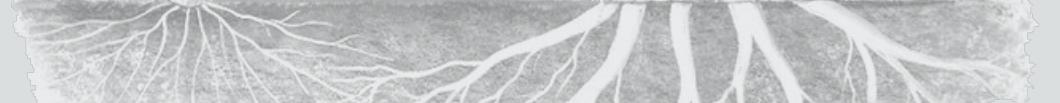
Tree
projection



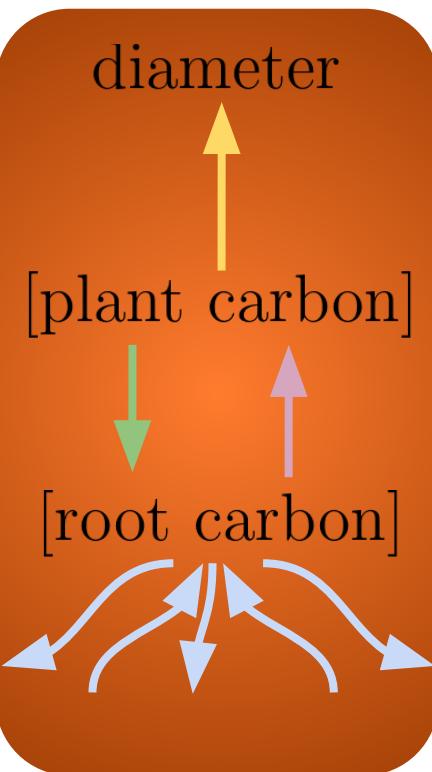
Methods: Network Set-up



Methods: The Model



Tree i



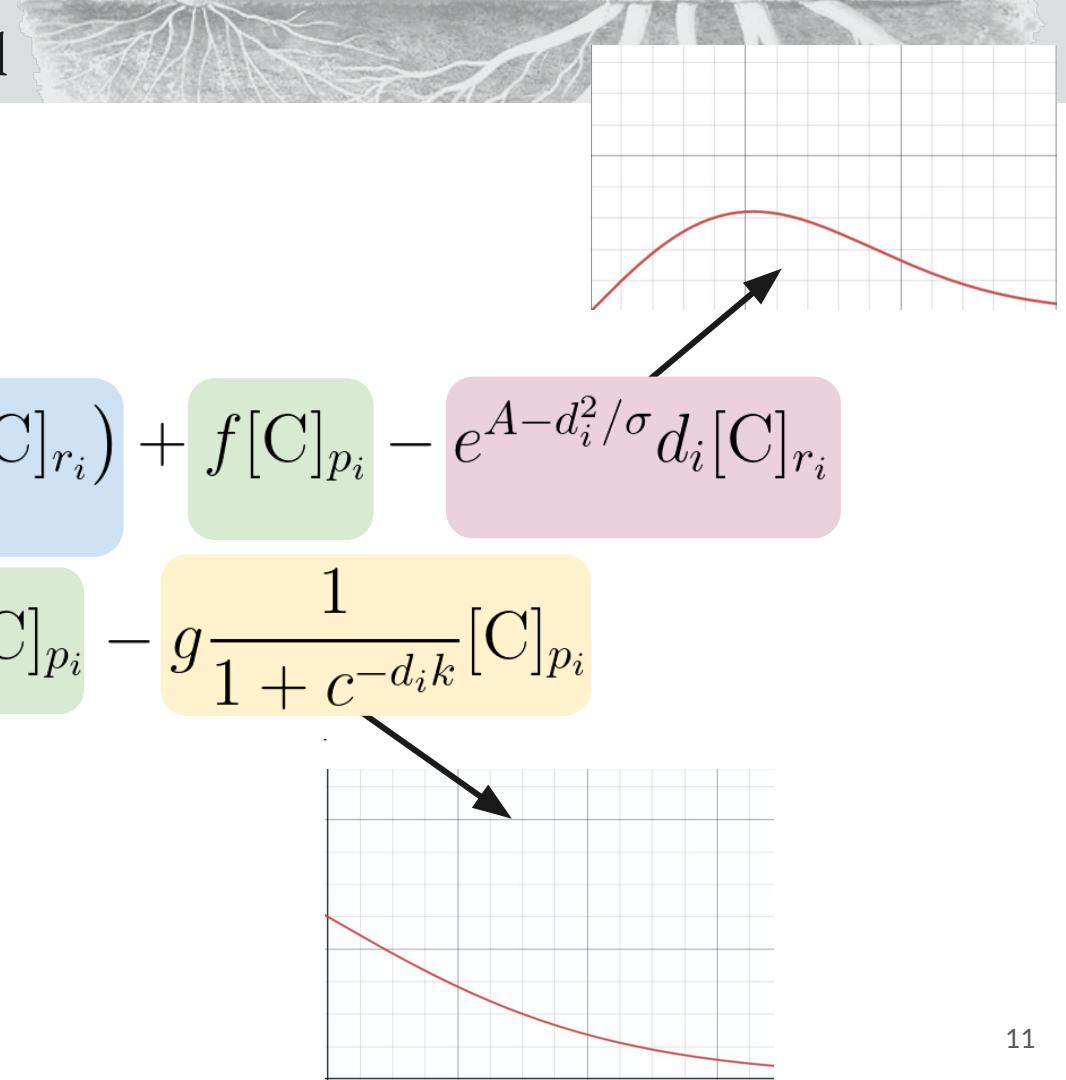
$$\frac{d[\text{root carbon}]_i}{dt} = \text{diffusion} + \text{deposition} - \text{uptake}$$

$$\frac{d[\text{plant carbon}]_i}{dt} = \text{uptake} - \text{deposition} - \text{growth}$$

$$\frac{ddiameter_i}{dt} = \text{growth}$$

Methods: Mathematical Model

$$\frac{d[\text{C}]_{r_i}}{dt} = D_C \sum_{j \in \mathcal{N}(i)} ([\text{C}]_{r_j} - [\text{C}]_{r_i}) + f[\text{C}]_{p_i} - e^{A-d_i^2/\sigma} d_i [\text{C}]_{r_i}$$
$$\frac{d[\text{C}]_{p_i}}{dt} = e^{A-d_i^2/\sigma} d_i [\text{C}]_{r_i} - f[\text{C}]_{p_i} - g \frac{1}{1 + c^{-d_i k}} [\text{C}]_{p_i}$$
$$\frac{dd_i}{dt} = \rho g \frac{1}{1 + c^{-d_i k}} [\text{C}]_{p_i}$$

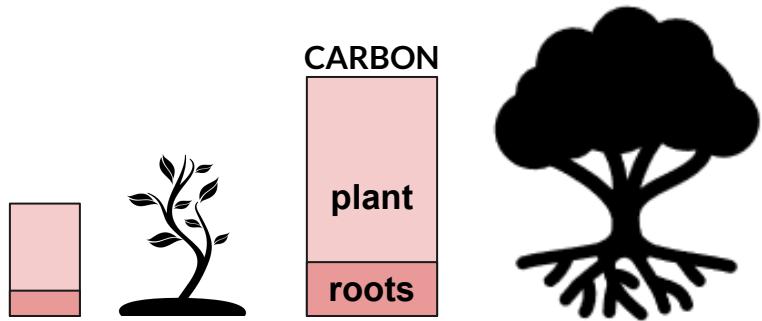


Methods: Experimental Set-up



Experimental Set-up

- Heterogeneous carbon distribution
- Numerically solve ODEs
 - Until convergence carbon reserves



Metrics

Biomass growth

In terms of plant diameter

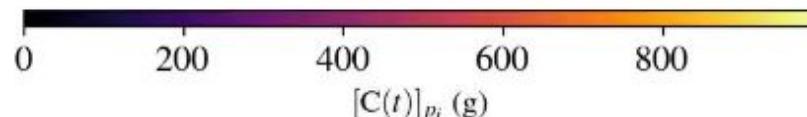
Sapling success

Minimum growth threshold

Results: Model Dynamics

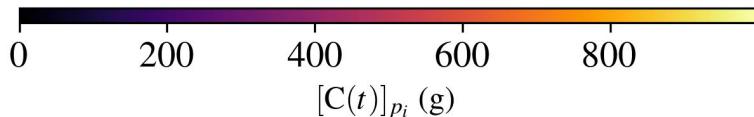
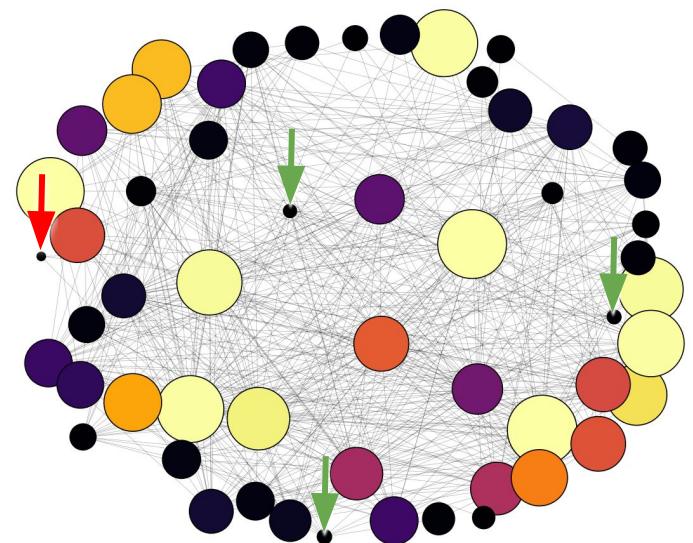
$t = 0.61$ days

**VIDEO IN ZIP-FILE.
network_evolution.mp4**

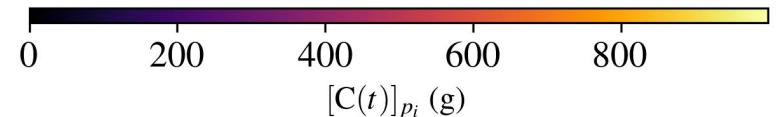
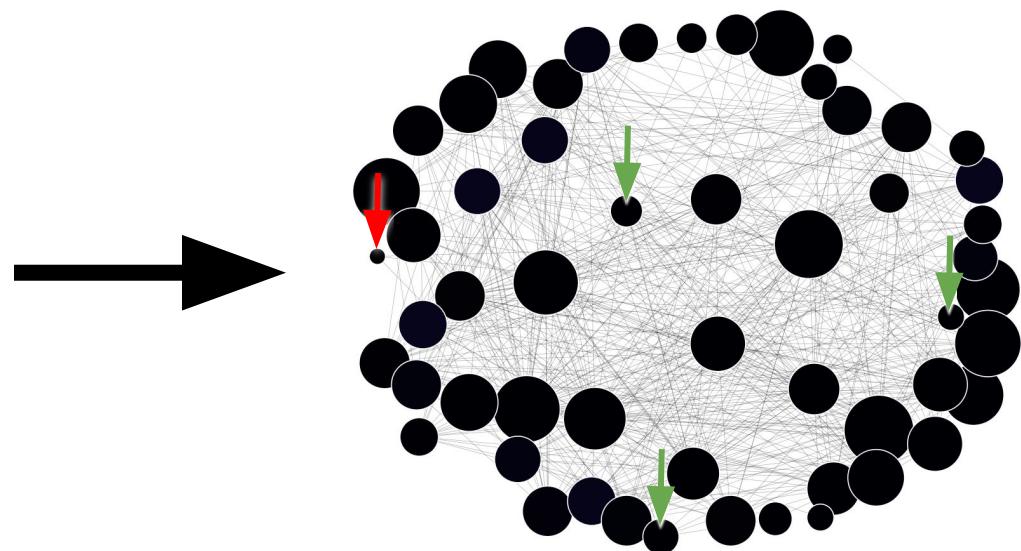


Results: Model Dynamics

$t = 0$ days

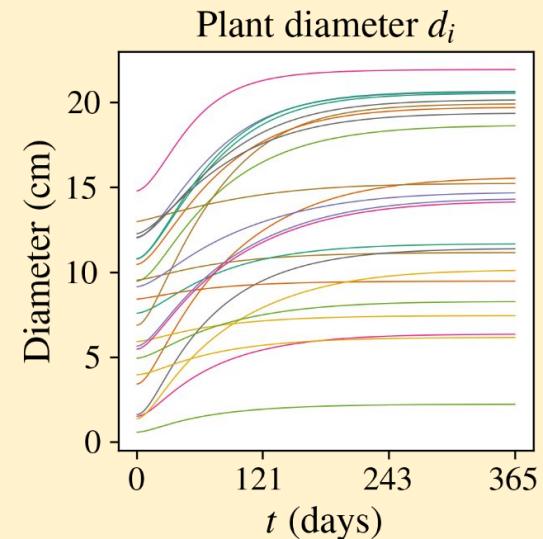
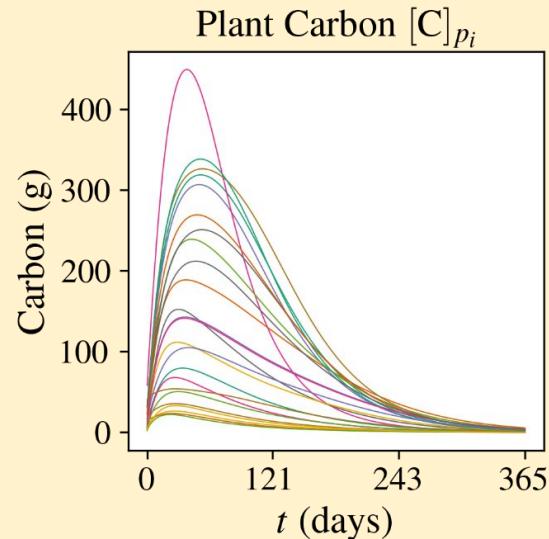
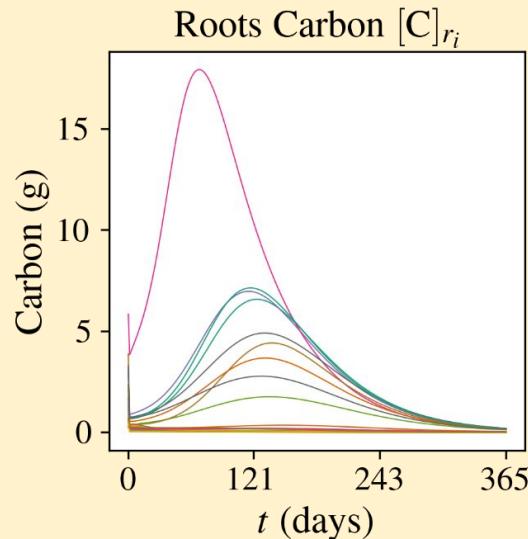


$t = 365$ days



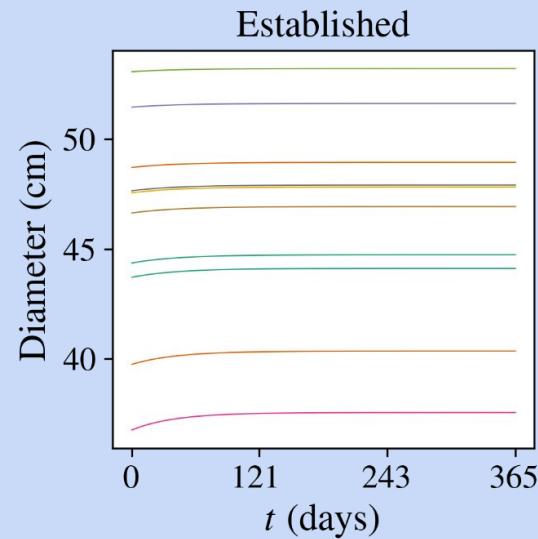
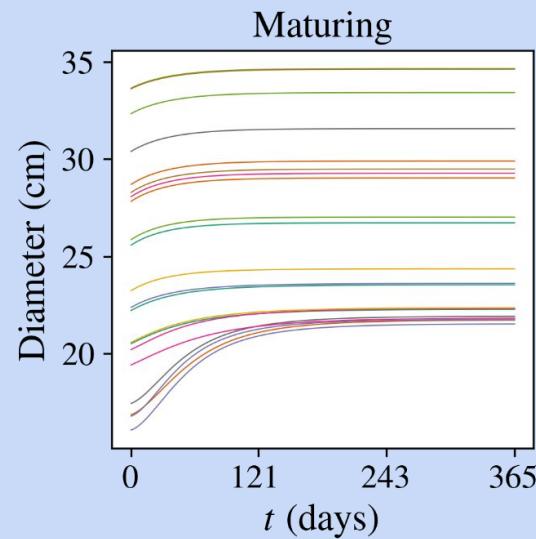
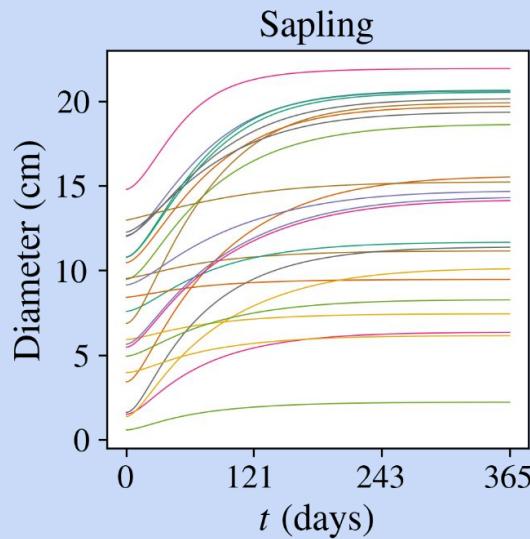
Results: Model Dynamics

Saplings

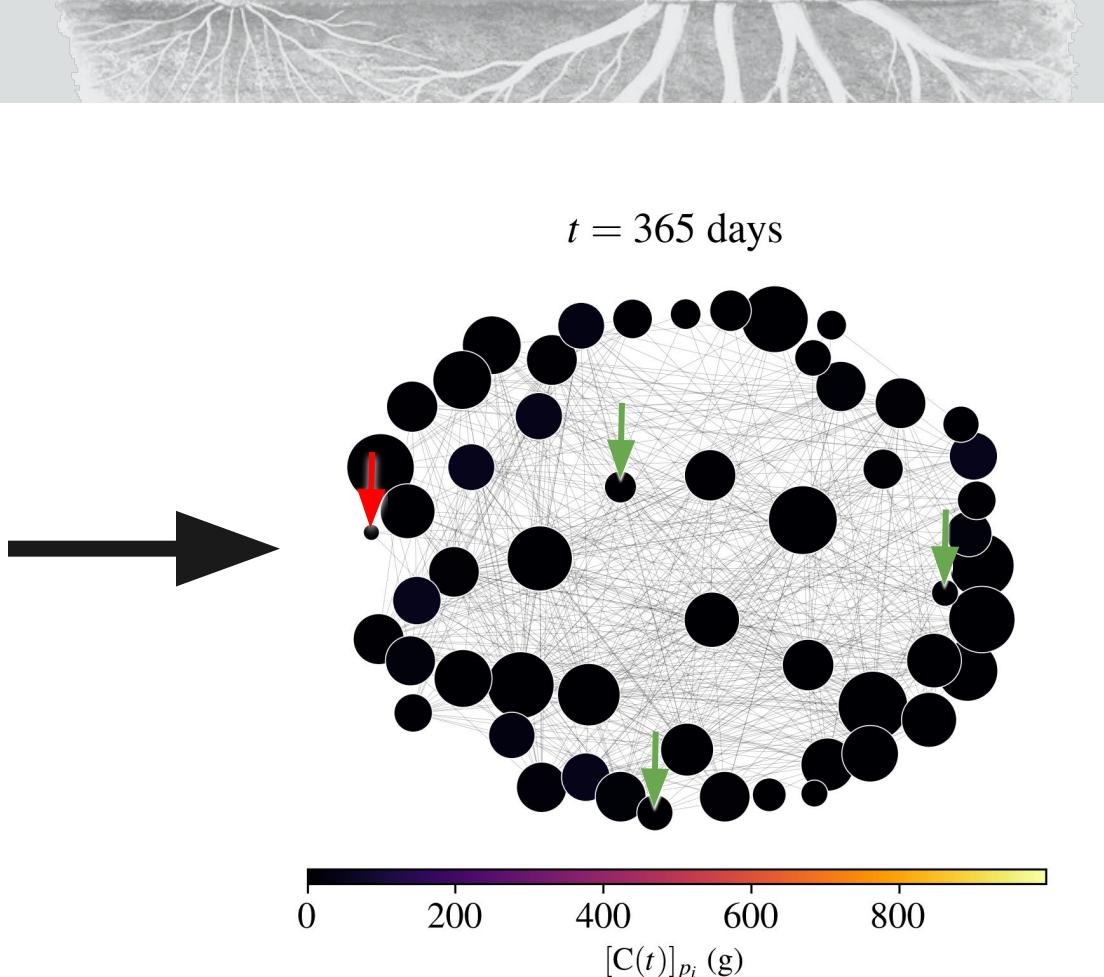
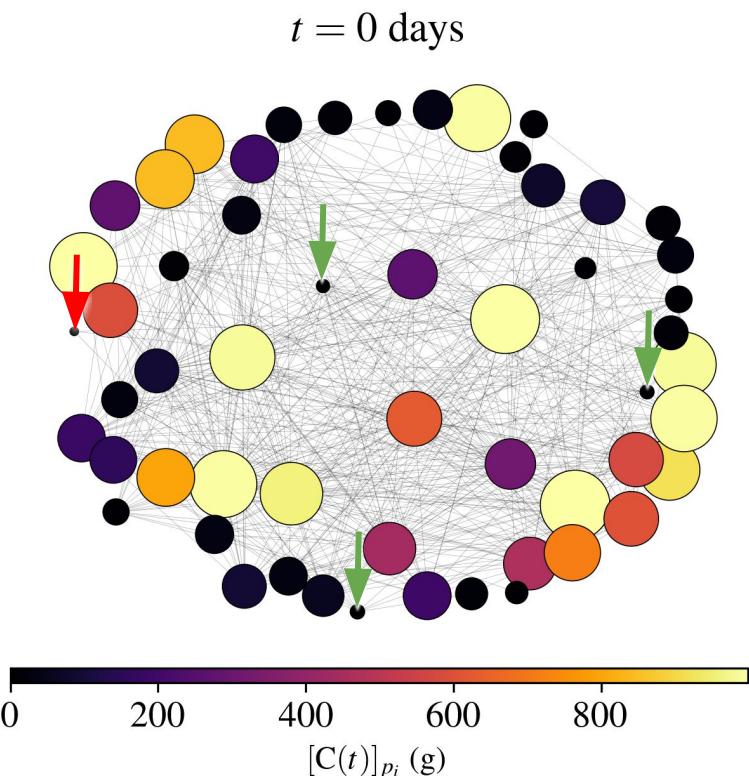


Results: Model Dynamics

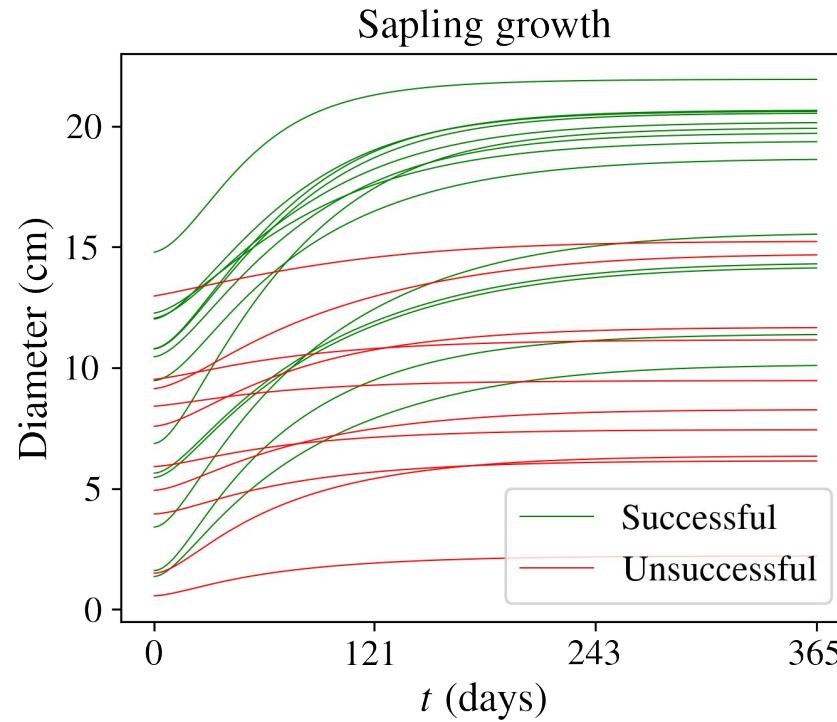
Diameter evolution per cohort



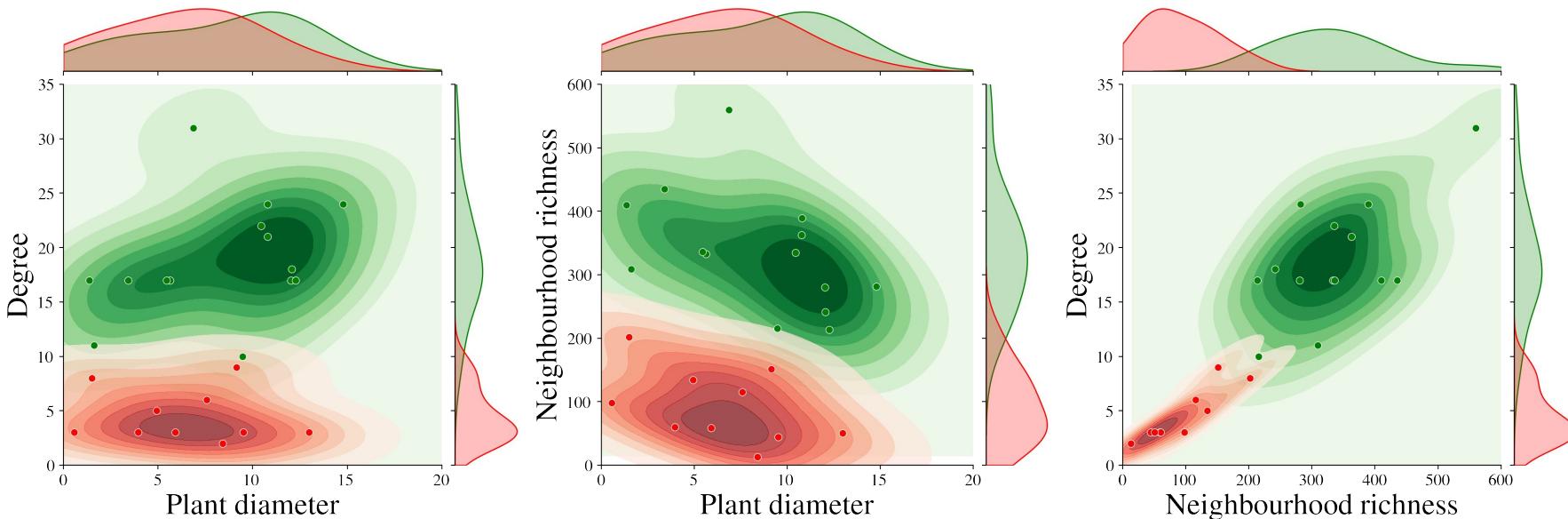
Results: Successful Saplings



Results: Successful Saplings

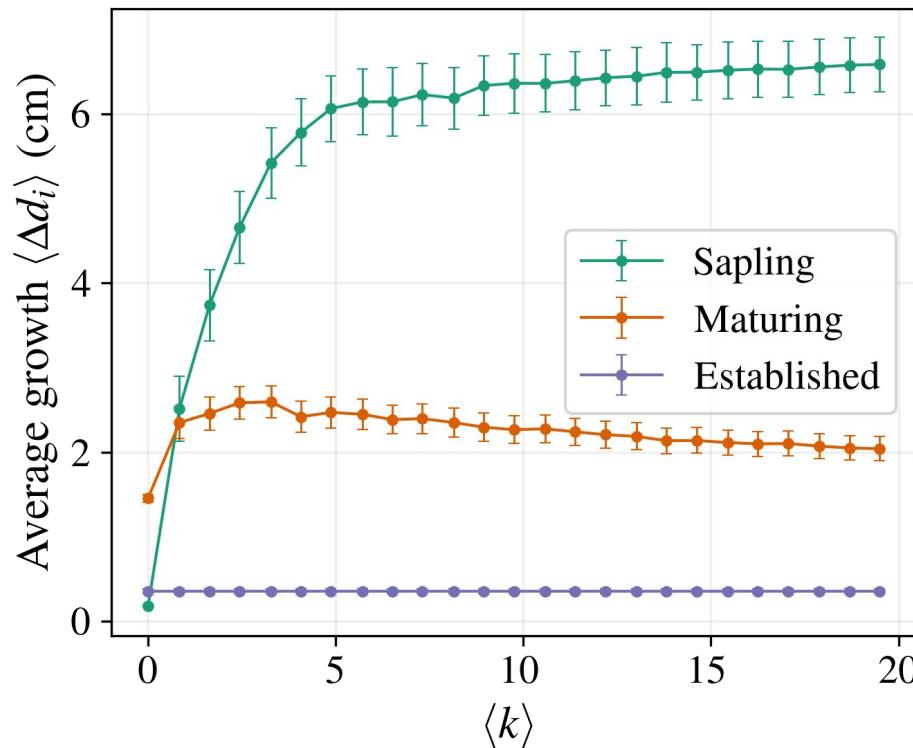


Results: Successful Saplings

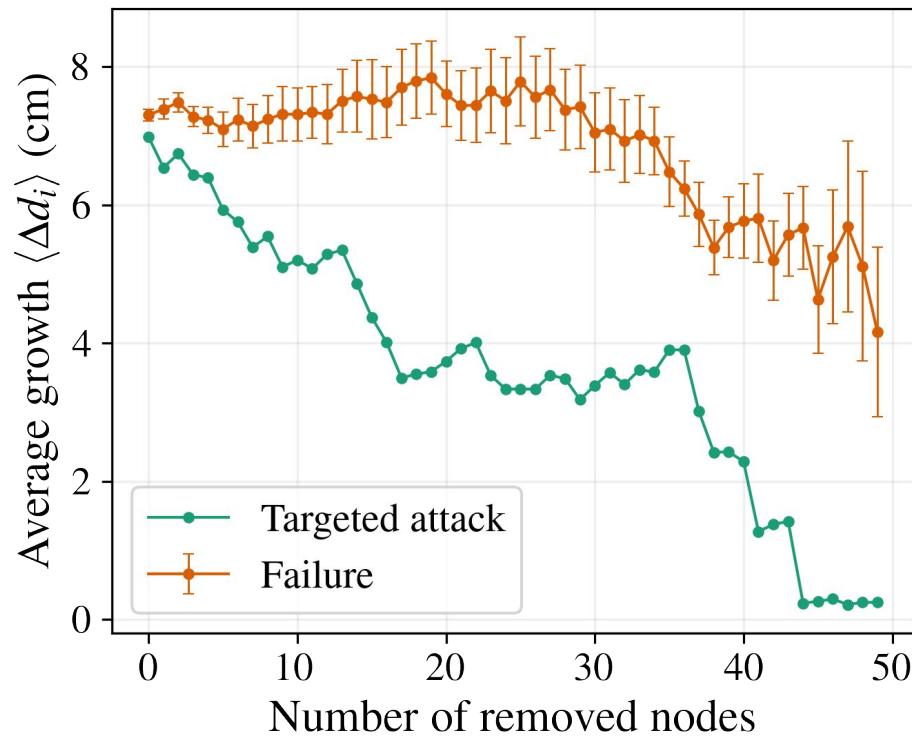


Diameter	Degree	Neighbourhood richness
$p\text{-value}$	< 0.001	< 0.001

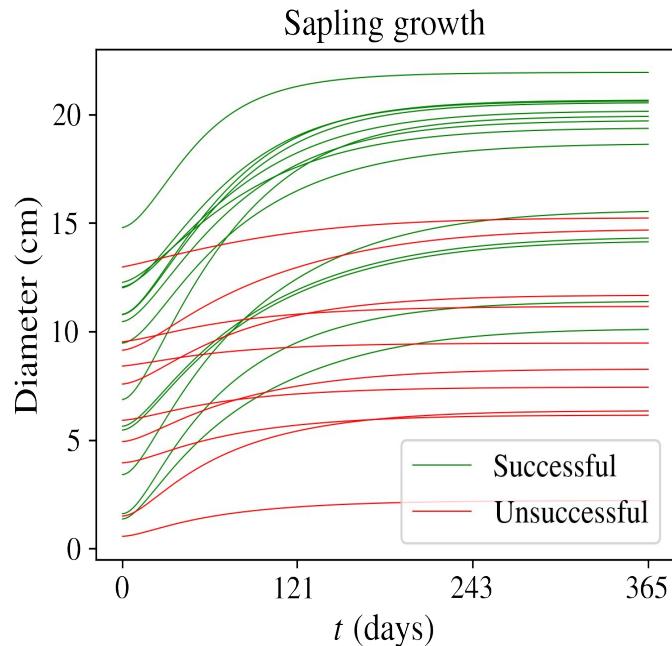
Results: Changing network properties



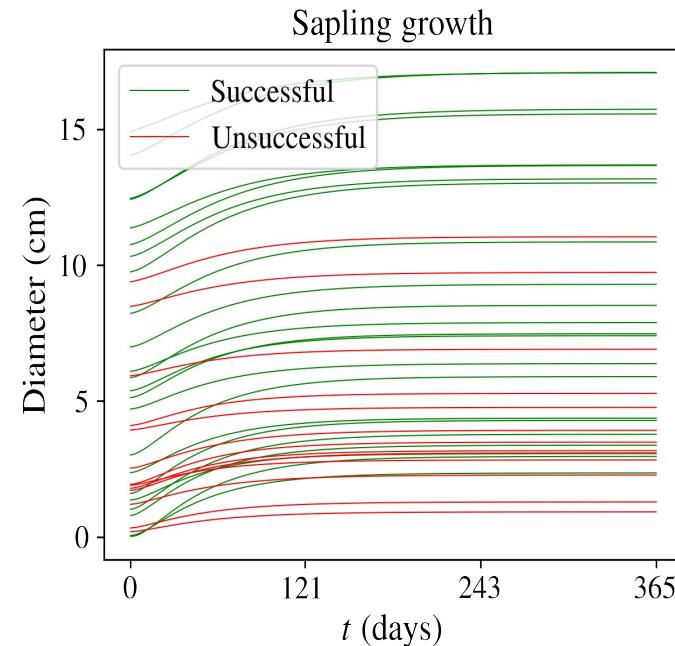
Results: Changing network properties



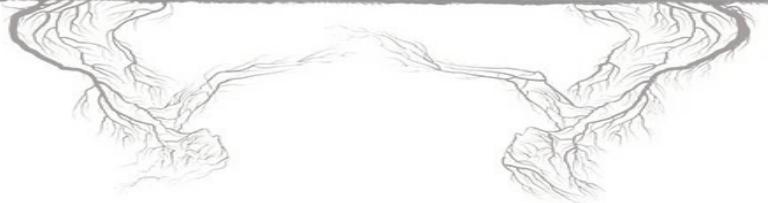
Results: Scale Free Network



Forest



Barabasi-Albert



Discussion and Future Work

Discussion

- Correspondence with source-sink gradients hypothesis
- Degree and neighbourhood richness are good predictors of sapling success.
- Benefit for saplings and maturing trees
- Sensitive to targeted attack

Future work

- Better understanding network topology of forests
- Other hypotheses: *Fungi as decision-maker*





References

- Babikova, Z., Gilbert, L., Bruce, T. J., Birkett, M., Caulfield, J. C., Woodcock, C., ... & Johnson, D. (2013). Underground signals carried through common mycelial networks warn neighbouring plants of aphid attack. *Ecology letters*, 16(7), 835-843.
- Beiler, K. J., Durall, D. M., Simard, S. W., Maxwell, S. A., & Kretzer, A. M. (2010). Architecture of the wood-wide web: Rhizopogon spp. genets link multiple Douglas-fir cohorts. *New Phytologist*, 185(2), 543-553.
- Figueiredo, A. F., Boy, J., & Guggenberger, G. (2021). Common Mycorrhizae Network: A Review of the Theories and Mechanisms Behind Underground Interactions. *Frontiers in Fungal Biology*, 48.
- Hinckley, T. M., Lachenbruch, B., Meinzer, F. C., & Dawson, T. E. (2011). A lifespan perspective on integrating structure and function in trees. In *Size-and age-related changes in tree structure and function* (pp. 3-30). Springer, Dordrecht.
- Issartel, J., & Coiffard, C. (2011). Extreme longevity in trees: live slow, die old?. *Oecologia*, 165(1), 1-5.
- Simard, S. W. (2018). Mycorrhizal networks facilitate tree communication, learning, and memory. In *Memory and learning in plants* (pp. 191-213). Springer, Cham.
- Simard, S. W., Beiler, K. J., Bingham, M. A., Deslippe, J. R., Philip, L. J., & Teste, F. P. (2012). Mycorrhizal networks: mechanisms, ecology and modelling. *Fungal Biology Reviews*, 26(1), 39-60.

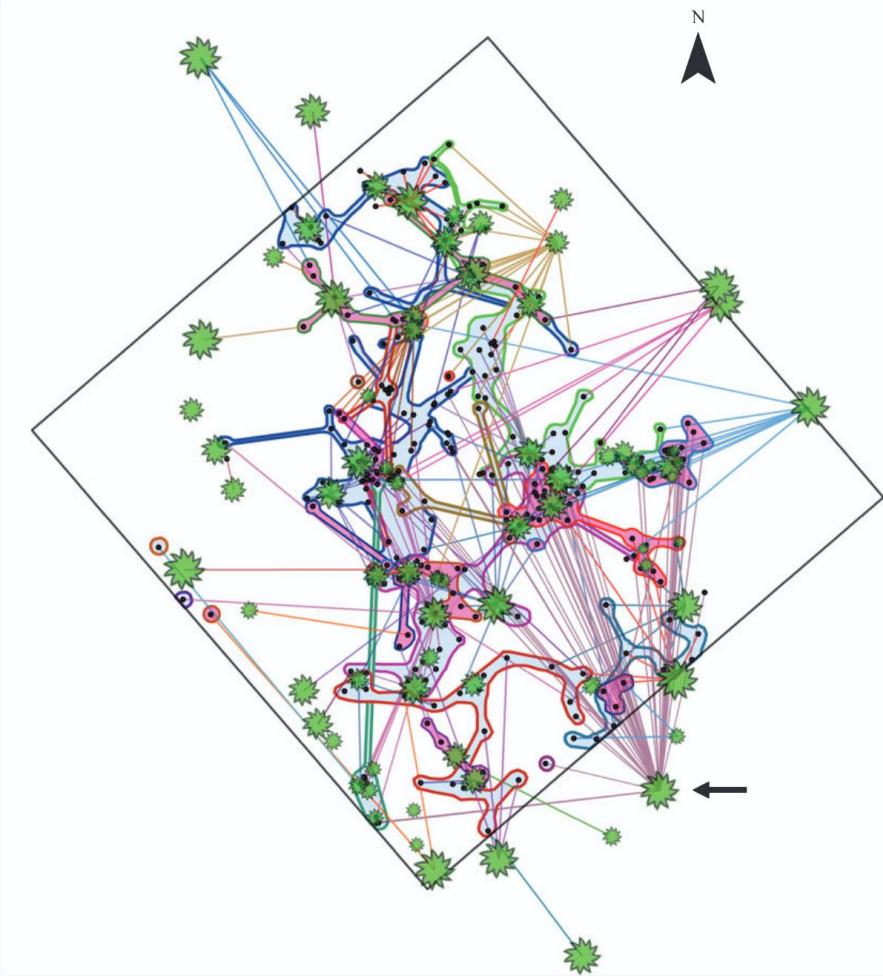


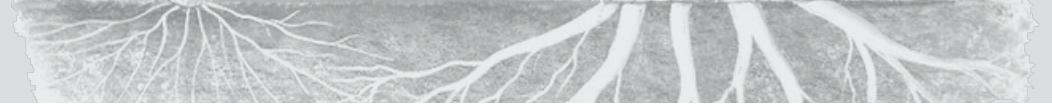


Appendices



Appendix I: Forest Sampling Map

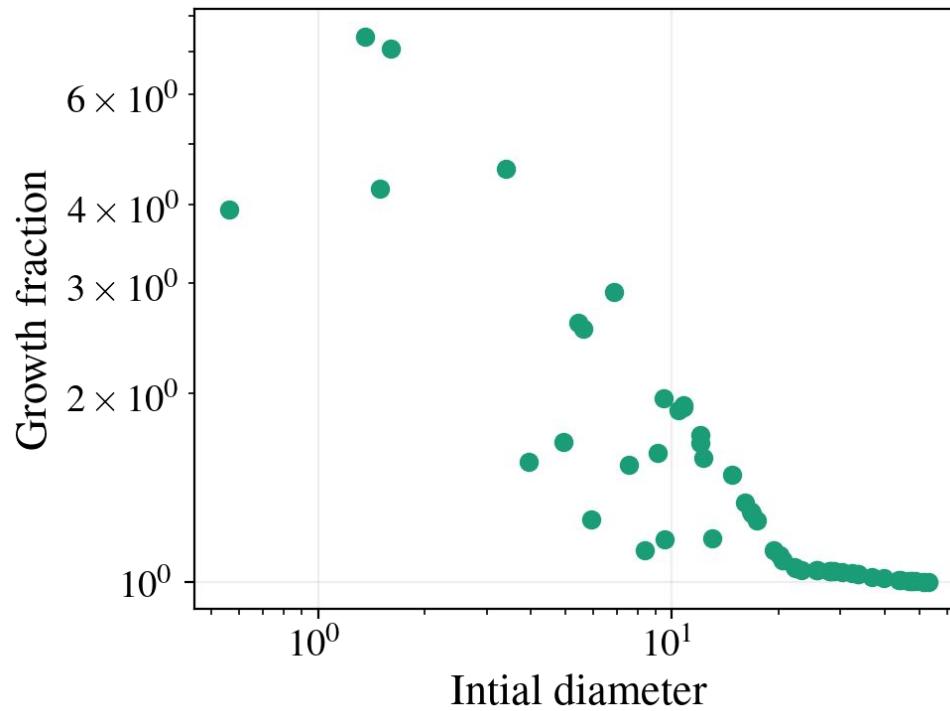




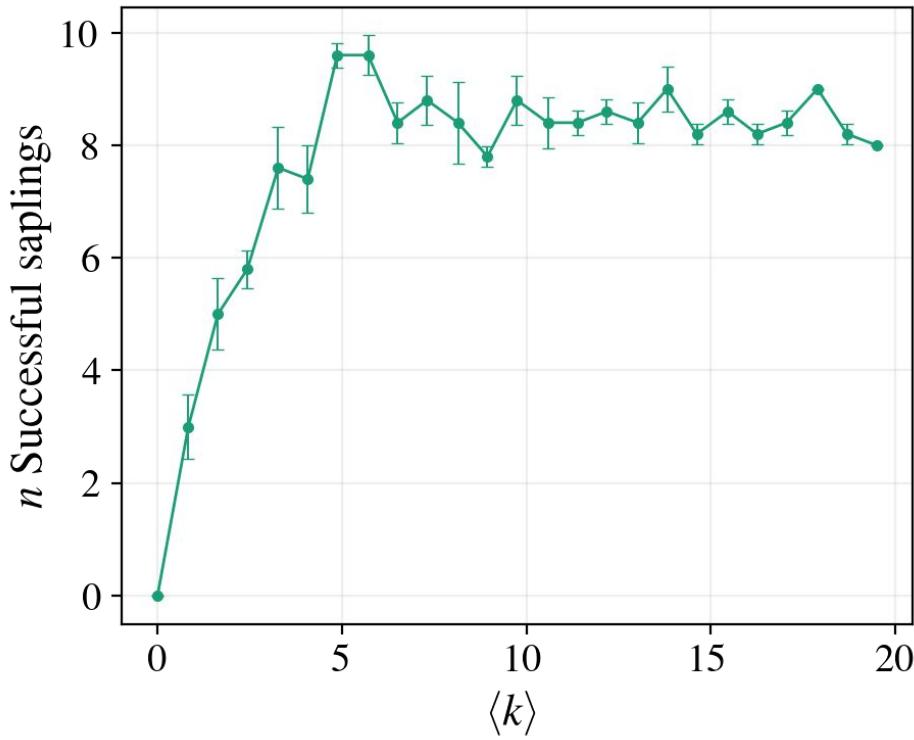
Appendix II: Model Parameters

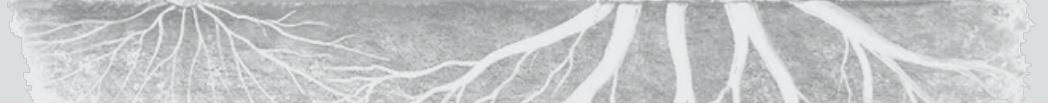
Symbol	Description	Value
D_C	Carbon diffusion coefficient	0.0005
f	Fraction of plant carbon transferred to root	0.0005
A	Root uptake coefficient	0.55
σ	Root uptake conversion factor	55
g	Growth speed	0.004
c	Logistic growth base	0.82
k	Growth conversion factor	0.6
ρ	Carbon to diameter conversion factor	0.01

Appendix III: Initial Diameter vs. Growth

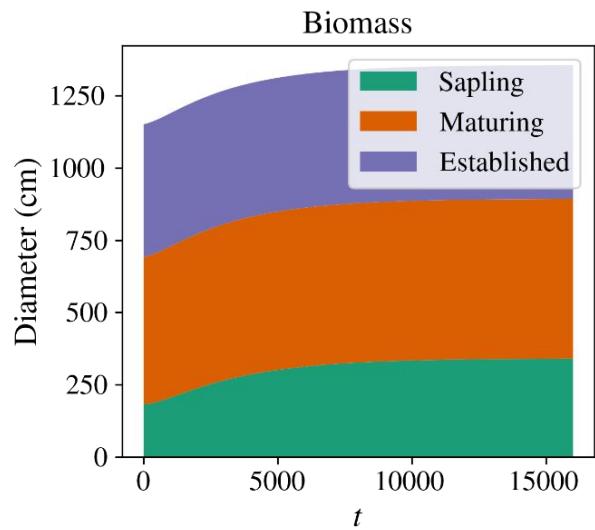
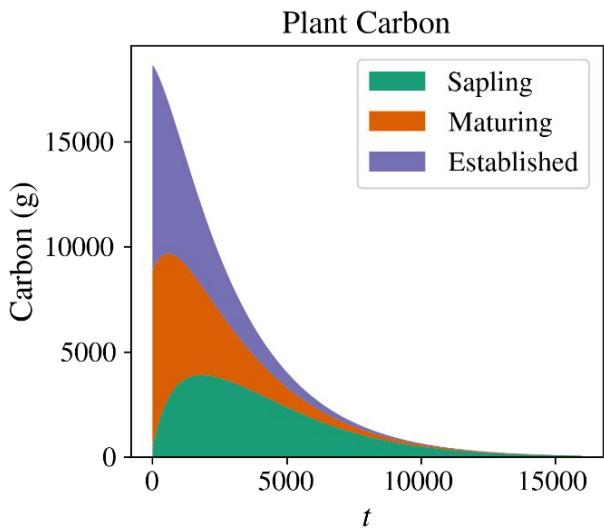
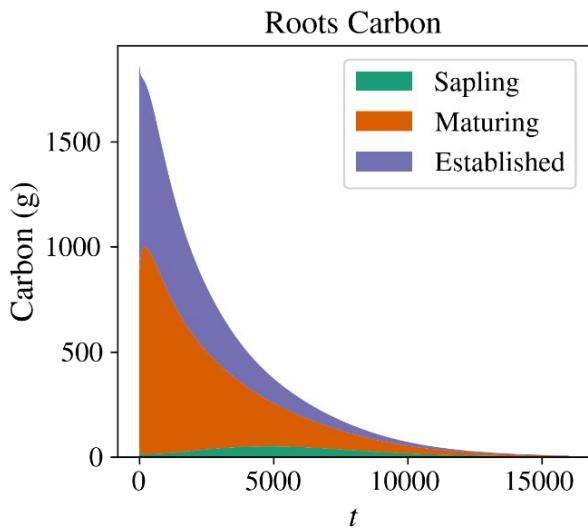


Appendix IV: Average Degree vs. n Successful Saplings

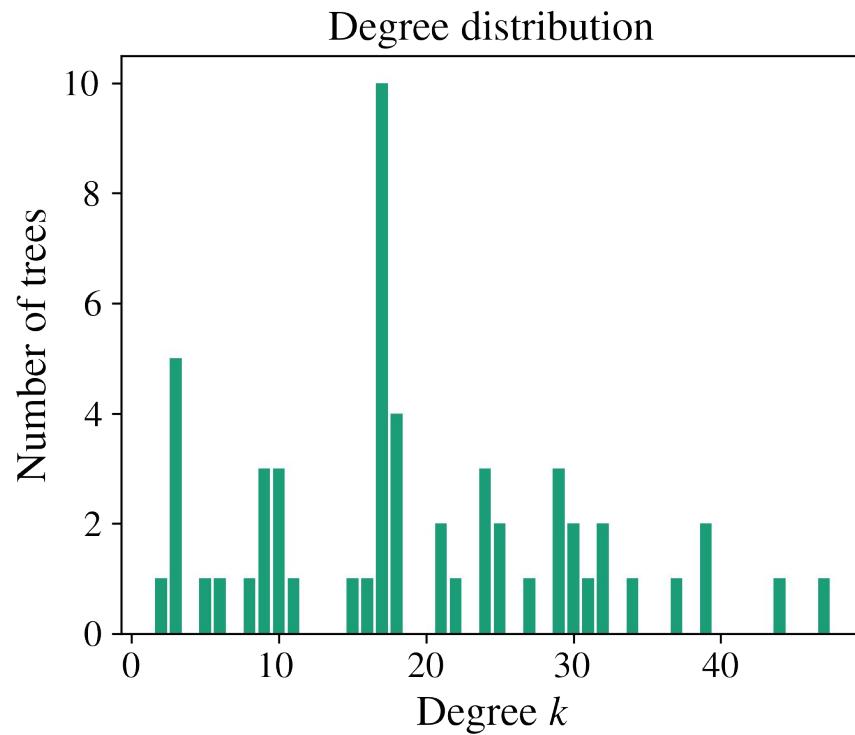




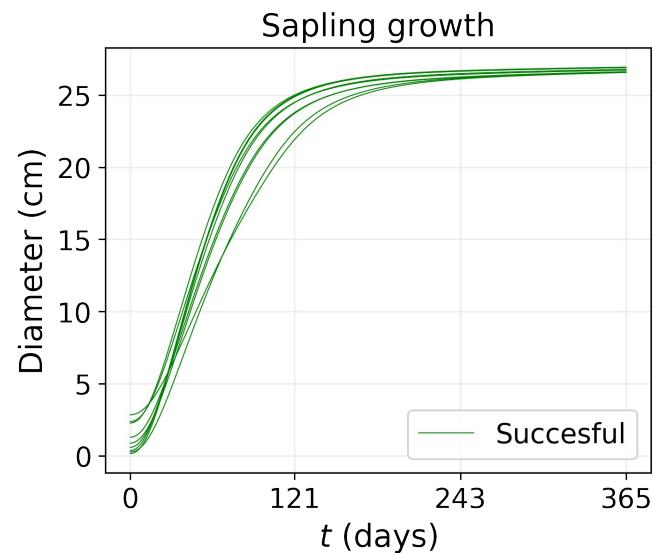
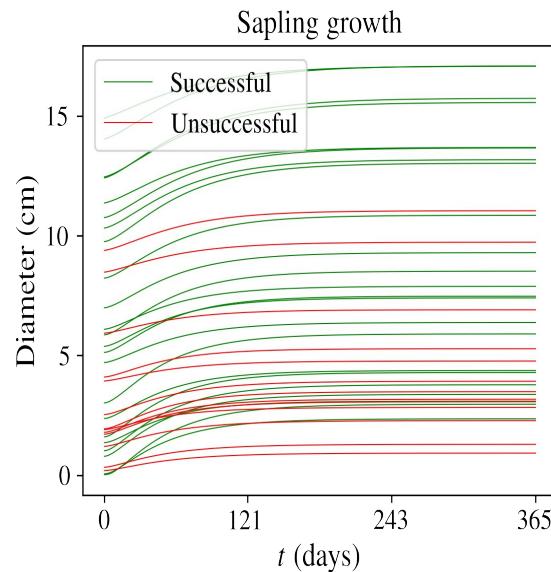
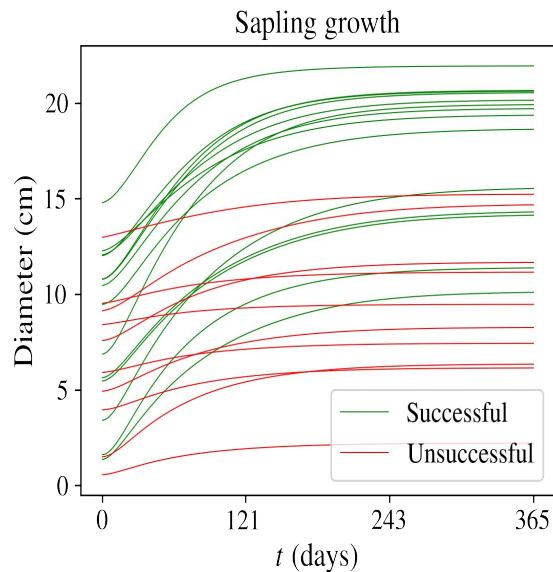
Appendix V: Stacked Plots Compartments



Appendix VI: Network Degree Distribution



Appendix VII: Network Topology Comparison



Forest

Barabasi-Albert

Random

Appendix VIII: Characteristic Time

