Community Phylogenetics of Lichens in the Flat Tops Wilderness Area of Colorado

Piper Boudart | Dr. Rachel Jabaily | Colorado College

Organismal Biology & Ecology | April, 2020

**01** Field Work & Specimen Collection **02** Specimen Identification & Deposition 03 Phylogeny Creation 04 Community Phylogenetic & Statistical Analyses

## Lichens

- Symbiotic relationship between fungus and algal photobiont
- Classification based on fungal symbiont
- Demonstrate substrate-specificity

   Present study includes saxicolous species only

   Assemblages vary based on microenvironment

   In part dictated by aspect and rock type





in the as Diver Blatianal Cornet M

Lichens of Colorado Flat Tops Wilderness Area

Rhizocarpon disporum (Nann F

Lichens of Colorado Flat Tops Wilderness Area

Physcia dubia (Hoffm.) Lettau

Lichens of Colorado Flat Tops Wilderness Area

Lecidella carpathica Körb

Lichens of Colorado Flat Tops Wilderness Area

Rusavskia elegans (Link) S. Y. Kondr. & Kärnefelt

Det. Piper Boudart, 2019

LOCALITY: Garfield Co., Colorado: Flat Tops Wilderness Area in the White River National Forest, M. E. Hale site no. 4 (1982). Large talus field on NE side of

> Lichens of Colorado Flat Tops Wilderness Area

Xanthoparmelia cumberlandia (Gyelnik) Hale

Det Diger Roudart 2010





U.S. National Herbarium





**U.S. National Herbarium** 

# Community Phylogenetics

- Intersection of ecology and phylogenetics
  Seeks to determine if local communities are random subsets of a broader regional species pool
  Analyzes evolutionary relationships within species assemblages
  Infere mechanisme of ecompletes
- Infers mechanisms of assemblage

#### **Regional Pool**

**Competitive Exclusion:** Local extinctions as the result of competition with evolutionarily similar neighbors







Fig. 1 Maximum likelihood consensus tree of the regional pool used to conduct community phylogenetic analyses.

### Phylogenetic Diversity Metrics

**Faith's Phylogenetic Diversity (PD):** the sum of the branch lengths of all species in the community phylogeny

**Mean Pairwise Distance (MPD)**: the mean branch length between each pair of species in community phylogeny

**Mean Nearest Taxon Distance (MNTD):** the mean branch length between each species and its nearest relative in the community phylogeny





Quadrat+Year	SES PD	SES MPD	SES MNTD
Q1_1982	-0.626	-0.529	-1.465
Q1_1992	-0.282	-0.179	-0.929
Q1_2009	0.420	0.356	-0.316
Q1_2019	-0.185	0.032	-0.710
Q2_1982	1.133	0.833	1.202
Q2_1992	-0.063	0.325	-0.402
Q2_2009	0.235	0.550	-0.111
Q3_1982	0.158	0.271	0.385
Q3_1992	-0.270	0.018	-0.350
Q3_2009	0.292	0.125	0.381
Q3_2019	0.743	0.530	0.000
Q4_1982	-0.343	-0.084	-0.320
Q4_1992	-0.315	-0.158	-0.284
Q5_1982	-0.111	-0.242	0.141
Q5_1992	-0.357	-0.491	0.081
Q6_1982	-0.071	-0.324	0.118
Q6 1992	-1.067	-0.473	-1.267
Q7 1982	+2.162	+2.218	1.144
Q7 1992	2.124	\$1,904	1.146
Q8 1982	0.657	0.387	1.078
Q8 1992	0.665	1.154	1.073
Q8 2009	0.763	0.922	1.212
Q9_1982	0.450	0.219	0.680
Q9 1992	0.778	0.161	1.105
Q9 2009	0.778	0.161	1.105
Q10_1982	1.568	1.369	+2.118
Q10 1992	0.046	0.315	-0.497
Q10 2009	-0.206	0.047	-0.586
Q11_1982	♦1.897	1.465	♦1.849
Q11_1992	1.264	0.755	0.755
Q11_2009	♦1.897	1.465	♦1.849
Q12 1982	0.036	0.058	-0.309
Q12 1992	0.161	0.328	-0.893
Q12 2009	-0.370	-0.008	-1.635
013 2019	0.086	-0.655	0.022

PD, MPD, and MNTD SES values for each quadrat, each inventory period

### **Phylogenetic Diversity Metrics**



Fig. 2 Average SES value for each quadrat over all available inventory periods with standard error. Triangles represent MNTD, squares MPD, and circles PD.

### Multivariate Analysis of Variance

Table 1

#### Repeated-Measures MANOVA for Time, Aspect, and Rock Type on Phylogenetic and Taxonomic Diversity Measures

PD	MPD	MNTD	Species Richness	% Crustose
0.2208	0.0633*	0.2433	0.6466	0.9143
0.2612	0.7148	0.2335	0.1019	0.2877
0.6637	0.8048	0.7858	0.6394	0.7626
0.1945	0.256	0.2647	0.0695*	0.7446
0.1719	0.6579	0.2102	0.0441**	0.2591
0.2828	0.5851	0.6027	0.5194	0.187
	PD 0.2208 0.2612 0.6637 0.1945 0.1719 0.2828	PD     MPD       0.2208     0.0633*       0.2612     0.7148       0.6637     0.8048       0.1945     0.256       0.1719     0.6579       0.2828     0.5851	PD         MPD         MNTD           0.2208         0.0633*         0.2433           0.2612         0.7148         0.2335           0.6637         0.8048         0.7858           0.1945         0.256         0.2647           0.1719         0.6579         0.2102           0.2828         0.5851         0.6027	PD         MPD         MNTD         Species Richness           0.2208         0.0633*         0.2433         0.6466           0.2612         0.7148         0.2335         0.1019           0.66637         0.8048         0.7858         0.6394           0.1945         0.256         0.2647         0.0695*           0.1719         0.6579         0.2102         0.0441**           0.2828         0.5851         0.6027         0.5194

\* P < 0.10

\*\* P < 0.05

#### Table 2

#### ANOVA for Aspect and Rock Type on Phylogenetic Diversity Measures

Effect	PD	MPD	MNTD
Aspect			
Prob > F	0.0072**	0.0065**	0.0203*
RSquare	0.626951	0.634664	0.541409
Rock type			
Prob > F	0.2895	0.3290	0.1820
RSquare	0.10119	0.086622	0.155772

\* P < 0.05 \*\* P < 0.01

### Discussion

• Quadrats 7 and 10 demonstrate overdispersion Indicative of competitive exclusion • Strong correlation between quadrat aspect and phylogenetic diversity metrics Most overdispersed quadrats located on horizontal rock faces flush with ground Significant effect of rock type on species richness

## Discussion

• Limited impact of habitat filtering • Continued increase in species richness • Coupled with decrease in available space • Phylogenetic structure likely to vary with successional stage Stronger habitat filtering in earliest stages 0 • Greater overdispersion in intermediate stages

## **Future Directions**

• Photographs provide opportunity to study lichen growth, neighbor dynamics, and successional patterns over time • Herbarium specimens provide physical inventory and opportunities for elemental analyses Added time point continues legacy of Hale and provides continuity for a long-term study

# Acknowledgments

This research was made possible by the Hevey Family Fund for Student Research, as well as the Keller Venture Grant program and OBE Departmental Funds. I am very grateful for the academic support of Lauren Trotta and Dr. Shane Heschel and the mentorship and advice of Dr. Rachel Jabaily and the rest of the Jabaily Lab.