

Will different populations of Little Bluestem display different levels of phenotypic variation?

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Introduction

- One goal for seed sourcing is to foster trait diversity to maximize restoration outcomes
- Variation of functional traits, traits that indicate competitive ability, exist within and among populations
- When should we source seeds from one population, and when is it better to source seeds from multiple populations?
- How is trait variation structured both within and between populations?

Experimental Methods

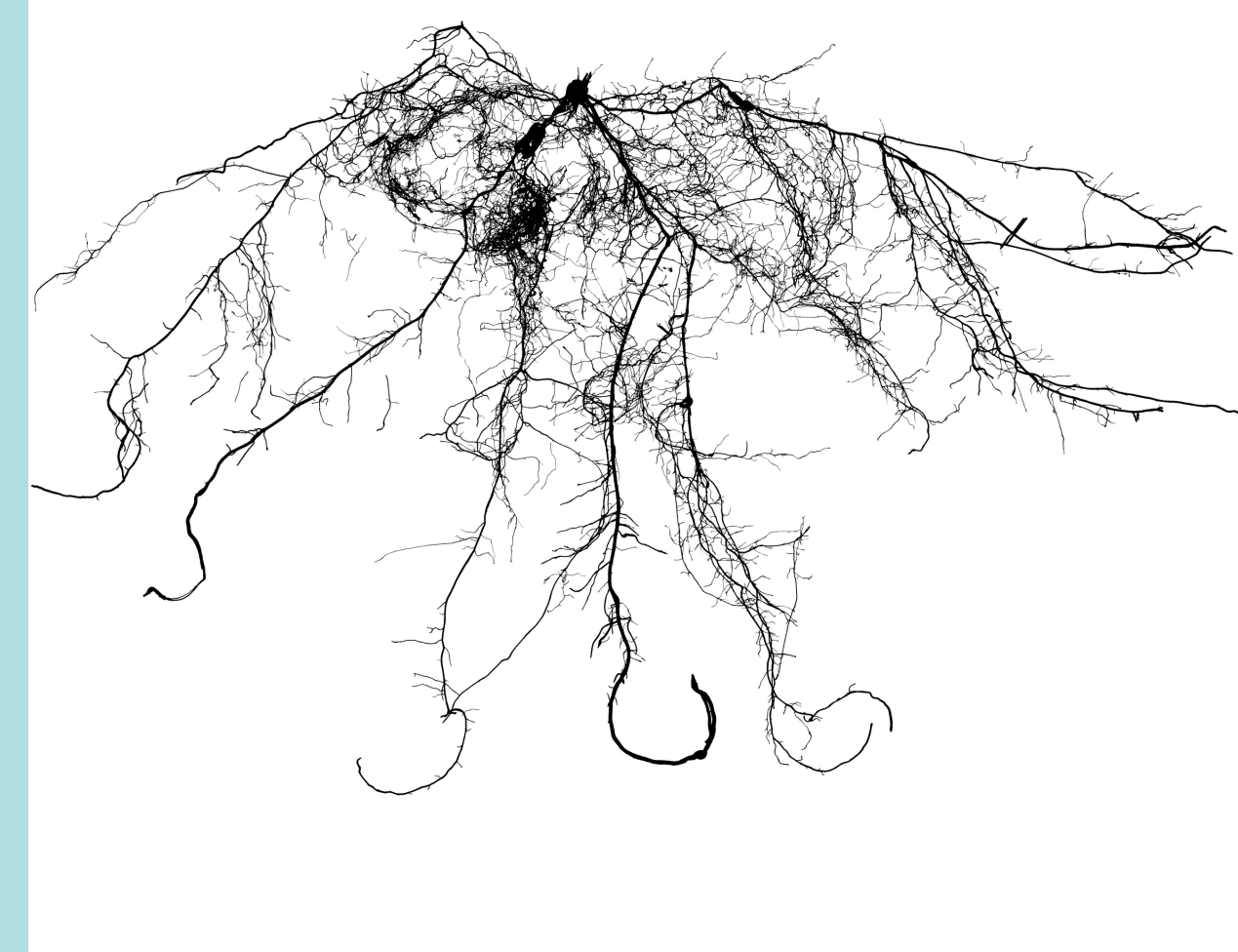
- Little Bluestem is a prairie grass native to IL
- Seeds were sourced from 4 populations in IL & WI
- Seeds were micropropagated, grown via tissue culture
- Sample size: 5-7 genotypes per population, 5-9 plants per genotype

Statistical Analysis

- **Coefficient of Variation (CV):** the extent of variability in relation to the mean of the population
- The higher the CV, the greater the variability
- We used linear models to understand differences in the CV of the genotypes based on populations

Traits

- **Root Dry Matter Content**
dry mass / fresh mass
 - Indicates lifespan and growth rate
- **Specific Leaf Area**
fresh mass / surface area
 - Regulates light capture and nutrient retention
- **Total Biomass**
 - Indicates plant productivity



Root scan

The different populations display significantly different levels of variation for root dry matter content (RDMC), but not for specific leaf area (SLA) or total biomass.

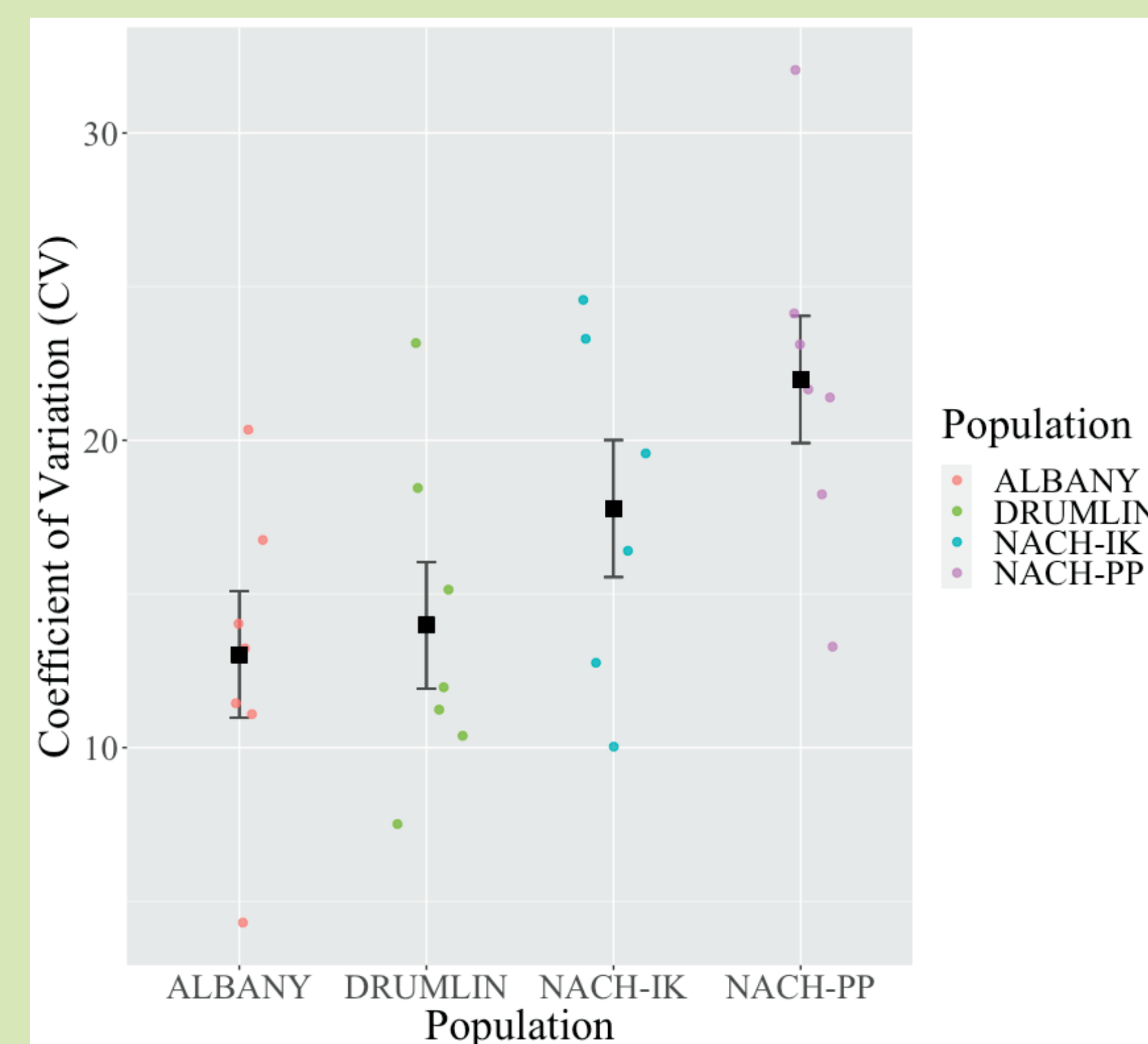


Figure 1: Coefficient of variation (CV) of RDMC per populations. Black, square points represent estimated mean population-level CV, while circular points represent estimated genotype-level CV.

Population	Coefficient of Variation
Albany	13
Drumlin	14
Nach-IK	17.8
Nach-PP	22

Figure 2: Estimated mean population level CV for RDMC

Trait	P-value	F-statistic	Df
SLA	0.397	1.03	3, 22
RDMC	0.022	3.89	3, 23
Total Biomass	0.617	0.61	3, 22

Figure 3: Statistical analysis of CV between populations for SLA, RDMC, and total biomass



Little Bluestem grown in the lab

References

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Discussion

- The NACH-PP population had the most RDMC variation
 - This was the only mesic soil population, suggesting more trait syndromes may be viable in a resource rich or less stressful environment
- It may be viable to source seeds from only one population if it has enough trait variation, but, as this study shows, the amount of trait variation differs between populations
- Future research could include more populations from mesic sites to see if the pattern of increased variation holds true
 - Additionally, it would be beneficial to repeat this study with other restoration relevant prairie species

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