

# Temporal patterns of and relationships between floral color and phylogenetics in North American wildflower communities

Chris Talbot & Marjorie Weber

## Background

Floral color assemblage in wildflower communities may be driven by myriad factors. Co-flowering assemblages are hypothesized to be **phylogenetically clustered**. Many factors driving floral color may **interact** with the phylogenetic composition of co-flowering assemblages.

As abiotic & biotic factors **vary over the season**, so too might the **relationship** between floral color and phylogenetics in assemblages. We ask:

**Q1:** In common Eastern North American wildflowers, what are the **temporal** patterns of  
**a) phylogenetic** diversity?  
**b) floral color** diversity?

**Q2:** Is there a **correlation** between floral color dispersion and phylogenetic dispersion?

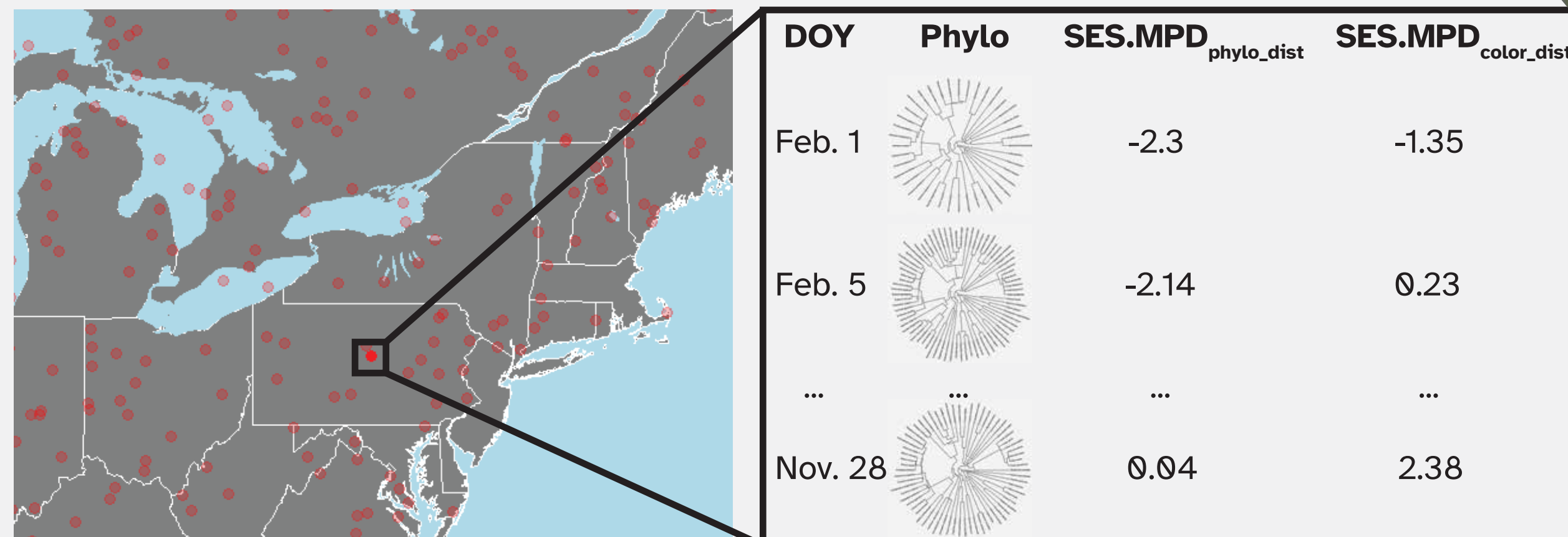
## Methods

For 985 species of common **Eastern North American wildflowers**, we collected the following data:



To compare floral colors, we calculated **Floral color dissimilarity**<sub>sp1,sp2</sub> = Euclidean distance in HSL colorspace

We calculated dispersion metrics using mean pairwise distance (MPD) using co-flowering assemblages from 1,000 sites across 4-day intervals::



We designed a generalized linear mixed model:  
**floral color dispersion** ~ phylogenetic dispersion + (1 | site) + (1 | day of year) + spatial autocorrelation + temporal autocorrelation

## Discussion

**Q1)** Floral color and phylogenetics show **strong temporal patterns** potentially linked to pollinator diversity or seasonal weather. Further research is needed.

**Q2)** Co-flowering assemblages with more close relatives display more disparate floral colors. **Divergent floral color** between co-flowering close relatives may be key in generating floral color diversity locally and globally. Floral color may also be a key floral trait generating prezygotic isolation.

Future studies should evaluate these patterns at this scale using **pollinator vision**.

## Acknowledgements

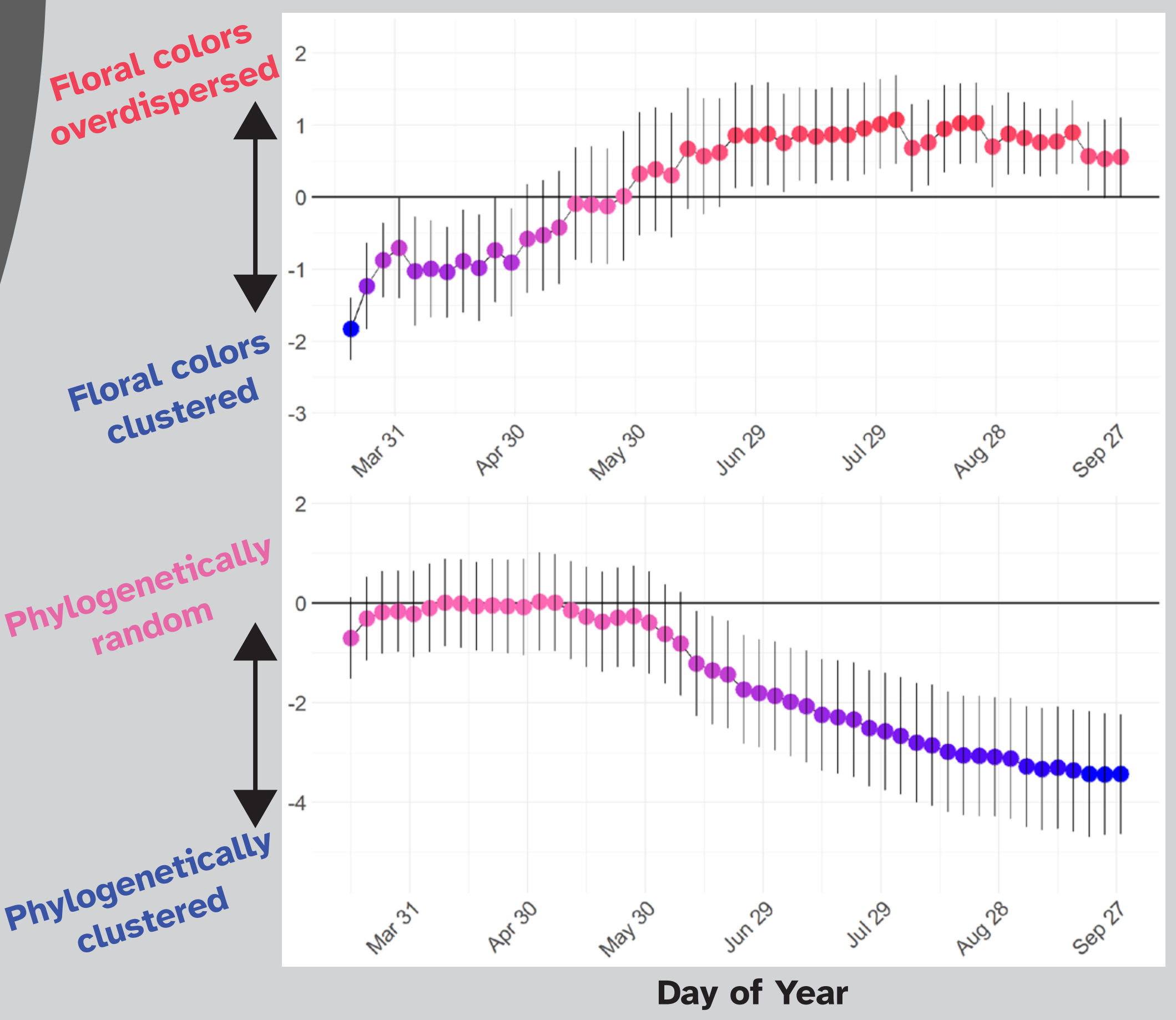
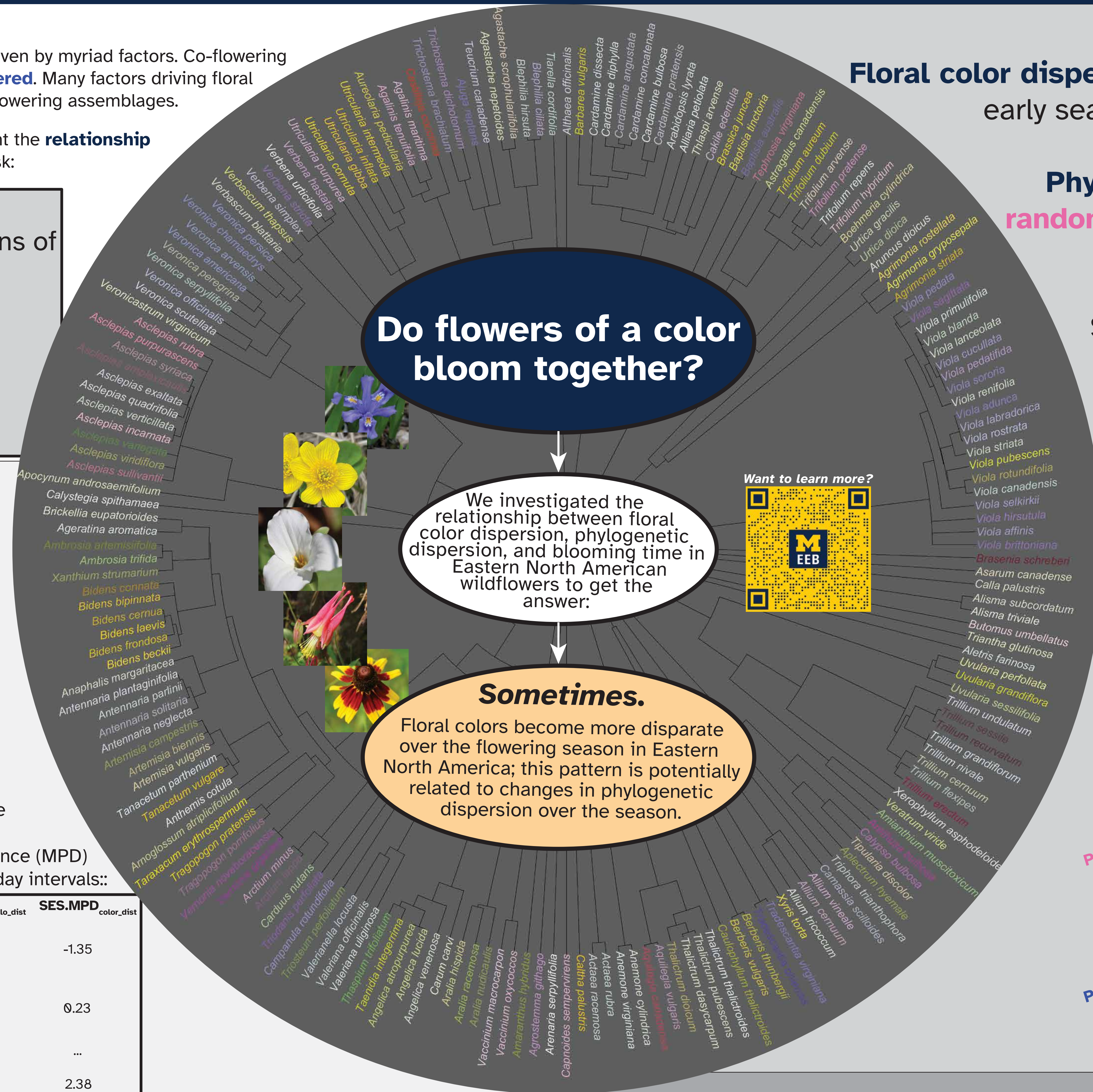
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## Results: Q1

**Floral color dispersion** transitions from **clustered** in the early season to **overdispersed** later in the year.

**Phylogenetic dispersion** transitions from **random** in the early season to **very clustered** later in the year.

Side by side, it seems that when groups are more **closely related**, they display more **disparate floral colors**. This may indicate diverging floral color between close relatives generates prezygotic isolation and floral color diversity.



## Results: Q2

Our mixed model reveals a small, **significant negative effect** of phylogenetic dispersion on floral color dispersion with strong temporal autocorrelation.

**Effect size: -0.11**  
**95% CI: [-0.13, -0.10]**

