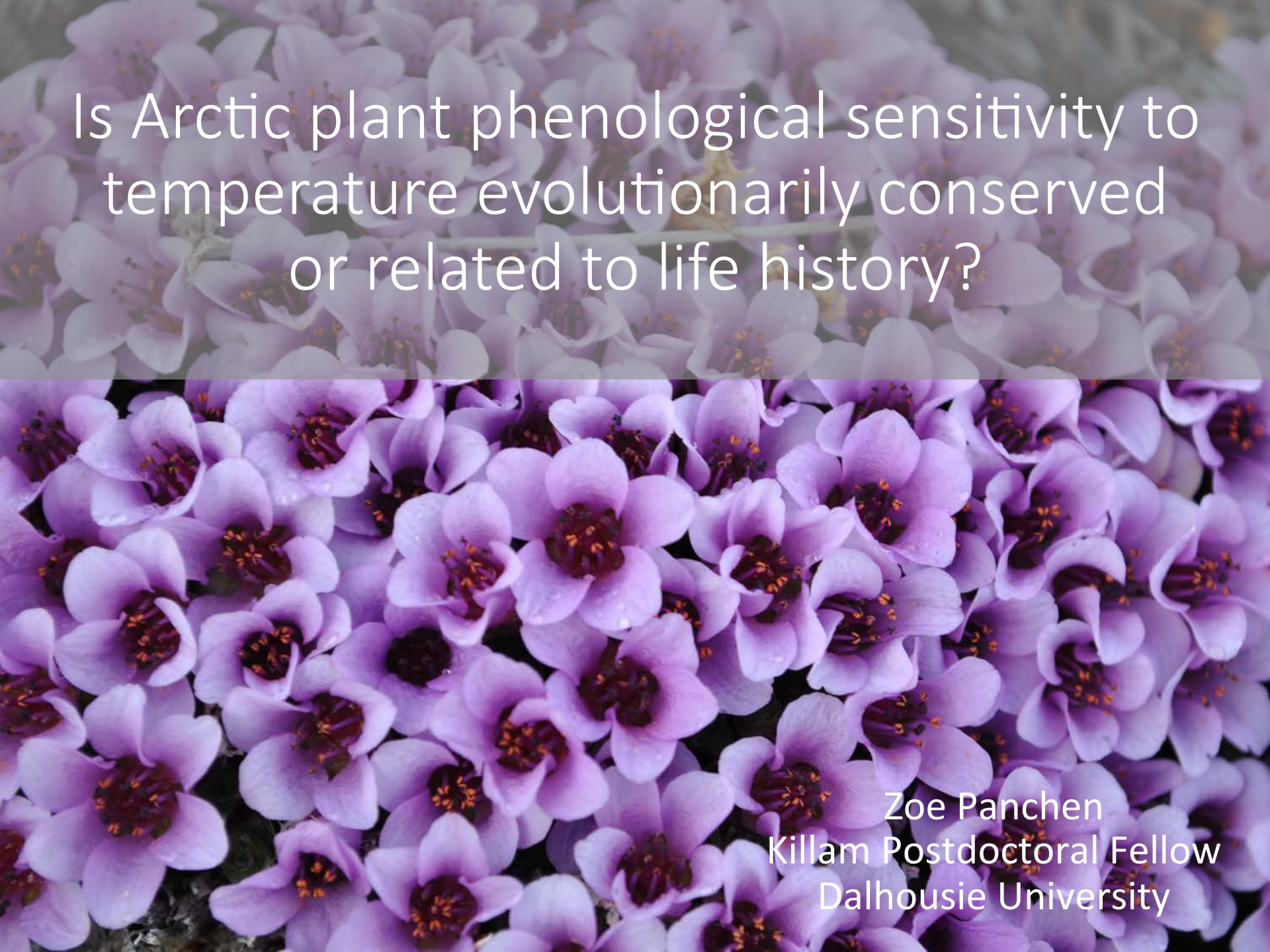


The background of the slide is a close-up photograph of numerous small, light purple flowers. The flowers have five petals each and dark, almost black, centers with visible orange-brown stamens. They are densely packed together, creating a textured, floral pattern. A semi-transparent dark grey horizontal band is positioned across the upper half of the image, serving as a backdrop for the title text.

Is Arctic plant phenological sensitivity to
temperature evolutionarily conserved
or related to life history?

Zoe Panchen
Killam Postdoctoral Fellow
Dalhousie University

Phenology timing versus sensitivity

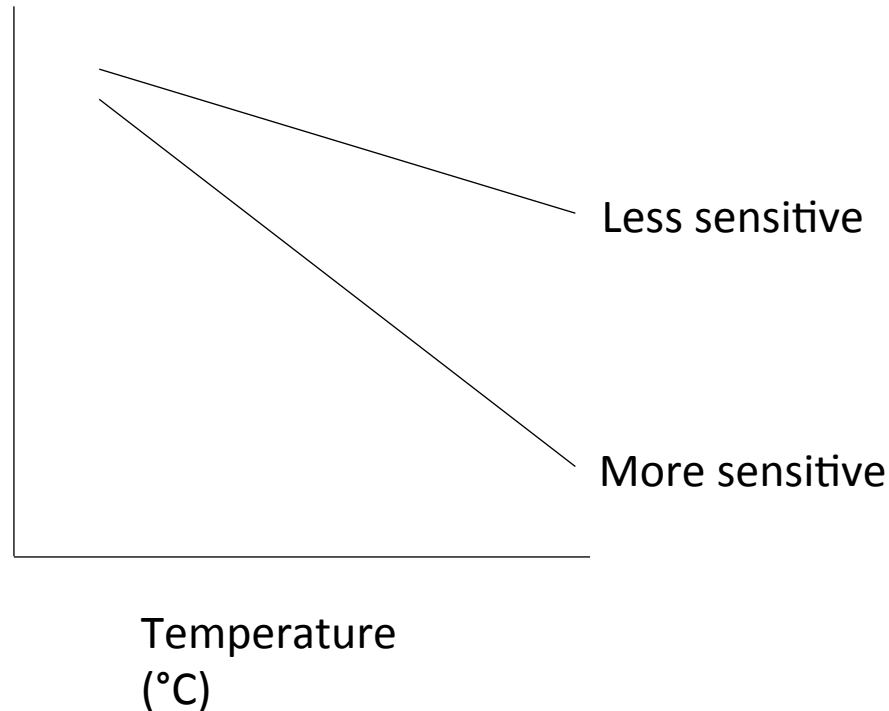


Time of
flowering

Or



Time of
green up
(DOY)



Timing: Day of Year of phenological event
(DOY in days)

Sensitivity: Phenological sensitivity to temperature
(slope — β in days/°C)

Timing: evolutionary relatedness

Phylogenetic conservatism in plant phenology

T. Jonathan Davies^{1*}, Elizabeth M. Wolkovich², Nathan J. B. Kraft³, Nicolas Salamin^{4,5}, Jenica M. Allen⁶, Toby R. Ault⁷, Julio L. Betancourt⁸, Kjell Bolmgren^{9,10}, Elsa E. Cleland¹¹, Benjamin I. Cook^{12,13}, Theresa M. Crimmins¹⁴, Susan J. Mazer¹⁵, Gregory J. McCabe¹⁶, Stephanie Pau¹⁷, Jim Regetz¹⁸, Mark D. Schwartz¹⁹ and Steven E. Travers²⁰

“We show that more closely related species tend to flower and leaf at similar times”

A phylogenetic comparative study of flowering phenology along an elevational gradient in the Canadian subarctic

Malie Lessard-Therrien • T. Jonathan Davies • Kjell Bolmgren

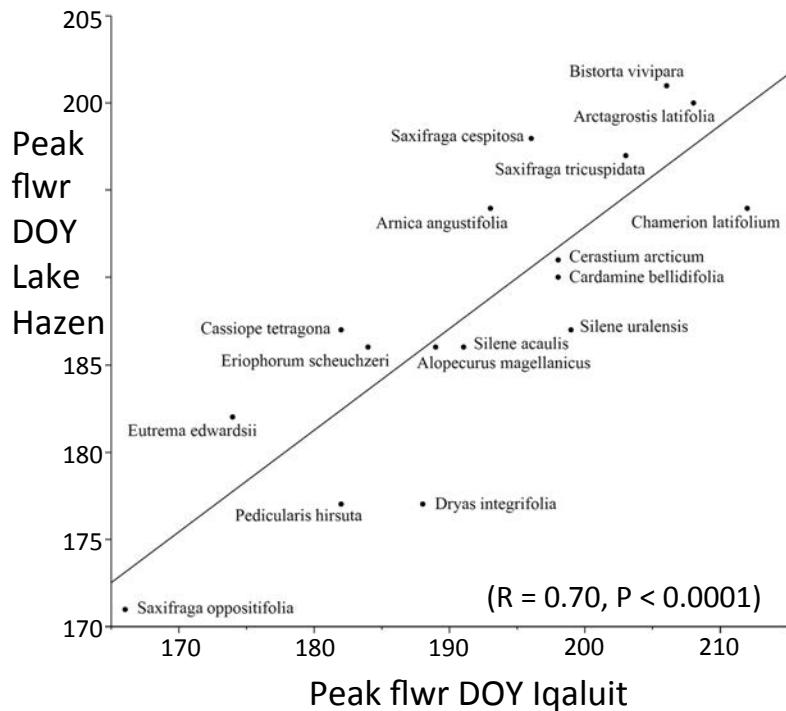
“We found that the phylogenetic signal for FFD was stronger at finer spatial scales”

Leaf out times of temperate woody plants are related to phylogeny, deciduousness, growth habit and wood anatomy

Zoe A. Panchen¹, Richard B. Primack², Birgit Nordt³, Elizabeth R. Ellwood⁴, Albert-Dieter Stevens³, Susanne S. Renner⁵, Charles G. Willis⁶, Robert Fahey⁷, Alan Whittemore⁸, Yanjun Du⁹ and Charles C. Davis¹⁰

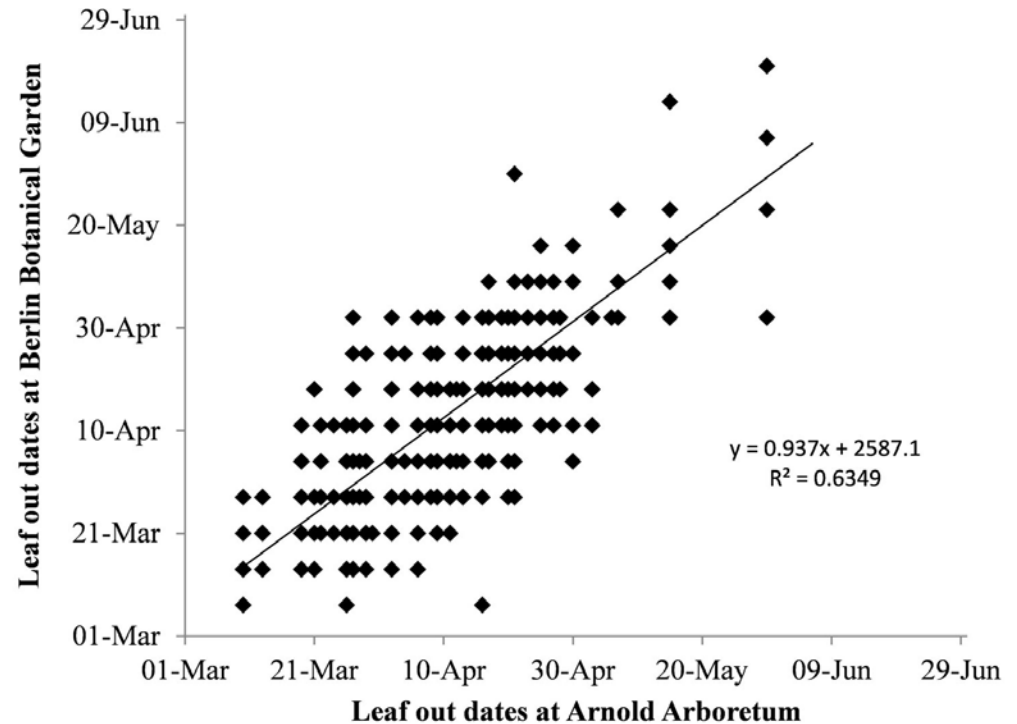
Species' flowering/leaf out sequence

Flowering times L. Hazen vs Iqaluit 2014



Panchen & Gorelick (2016)

Leaf out times Berlin vs Boston 2012



Panchen et al. (2014)

Species flower & leaf out in roughly the same order each year and at different locations

Sensitivity: evolutionary relatedness

Phenological responses to climate change do not exhibit phylogenetic signal in a subalpine plant community

PAUL J. CARADONNA^{1,2,4} AND DAVID W. INOUE^{1,3}

**FLOWERING DATE OF TAXONOMIC FAMILIES PREDICTS
PHENOLOGICAL SENSITIVITY TO TEMPERATURE:
IMPLICATIONS FOR FORECASTING THE EFFECTS OF
CLIMATE CHANGE ON UNSTUDIED TAXA¹**

SUSAN J. MAZER^{2,3,13}, STEVEN E. TRAVERS⁴, BENJAMIN I. COOK^{5,6}, T. JONATHAN DAVIES⁷,
KJELL BOLMGREN^{8,9}, NATHAN J. B. KRAFT¹⁰, NICOLAS SALAMIN¹¹, AND DAVID W. INOUE^{10,12}

“Family membership can be used to identify taxa of high and low sensitivity to temperature”

Phylogenetic patterns of species loss in Thoreau's woods are driven by climate change

Charles G. Willis^a, Brad Ruhfel^a, Richard B. Primack^b, Abraham J. Miller-Rushing^b, and Charles C. Davis^{a,1}

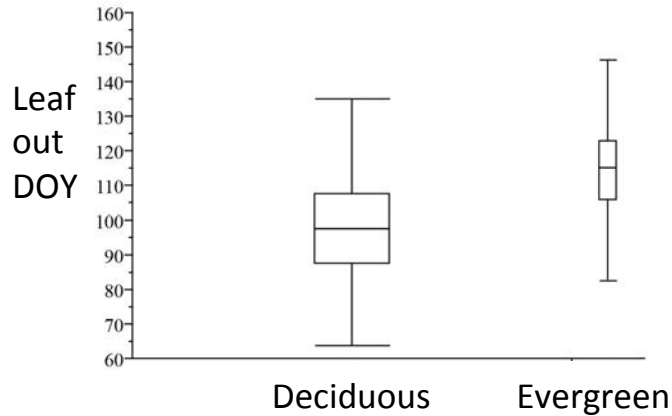
“.....flowering-time response traits are shared among closely related species.....”

Life history

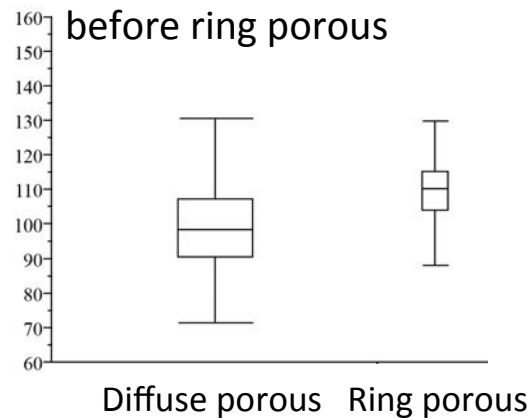
Timing



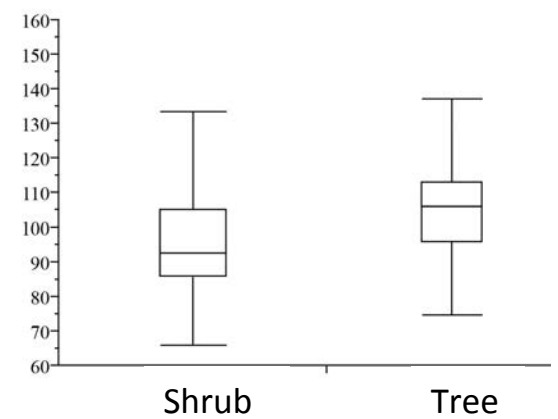
Deciduous leaf out before evergreen



Diffuse porous leaf out before ring porous



Shrubs leaf out before trees

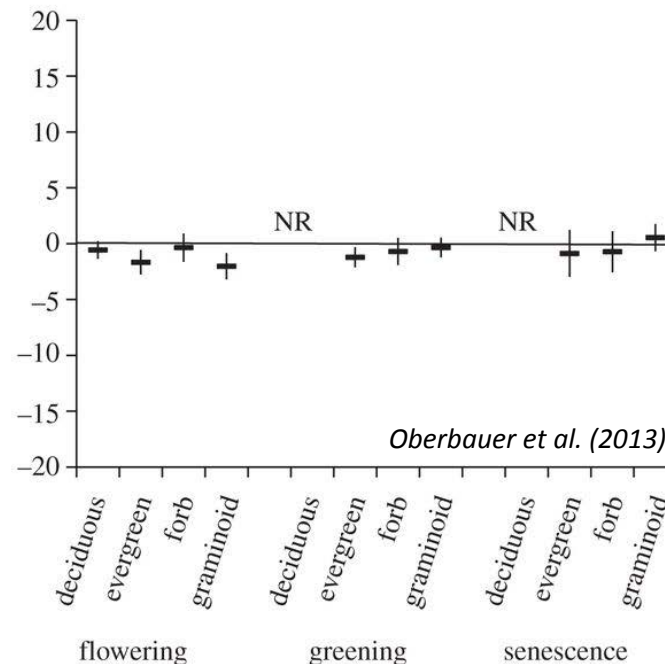


Panchen et al. (2014)

Sensitivity



Flower,
green up,
senescence
sensitivity to
temperature
(Days/°C)

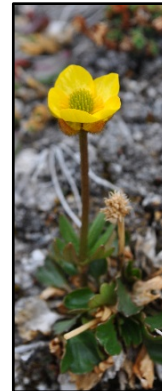
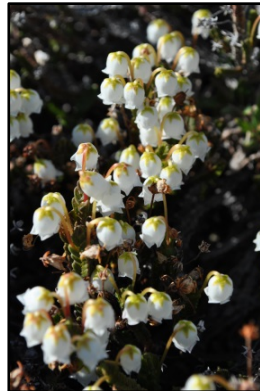


Oberbauer et al. (2013)

Questions

Arctic plant flower/green up timing and sensitivity

- evolutionarily conserved?
- related to life history?





Preliminary analysis

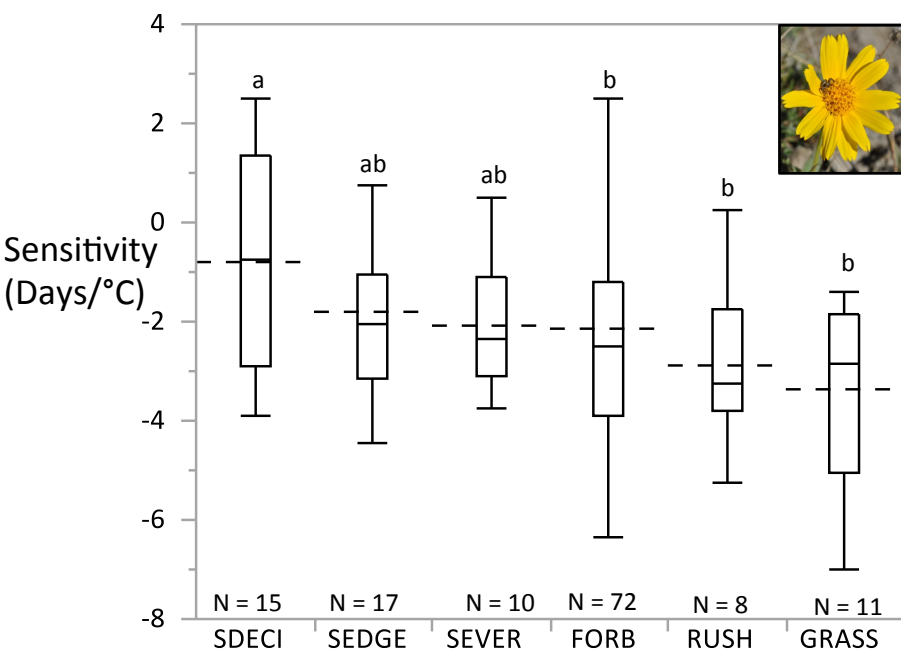


- Sensitivity to temperature (May, June, July) per species
 - $\text{DOY} \sim \alpha + (\beta^* \text{ monthly temp.}) + (\text{site}[\text{plot}])$
 - ITEX first flower & green up data set (Prevey et al. 2017)
 - β - most sensitive monthly temp. (June similar results)
- Life history patterns
- Bayesian hierarchical model

Flowering/green up β : Growth form

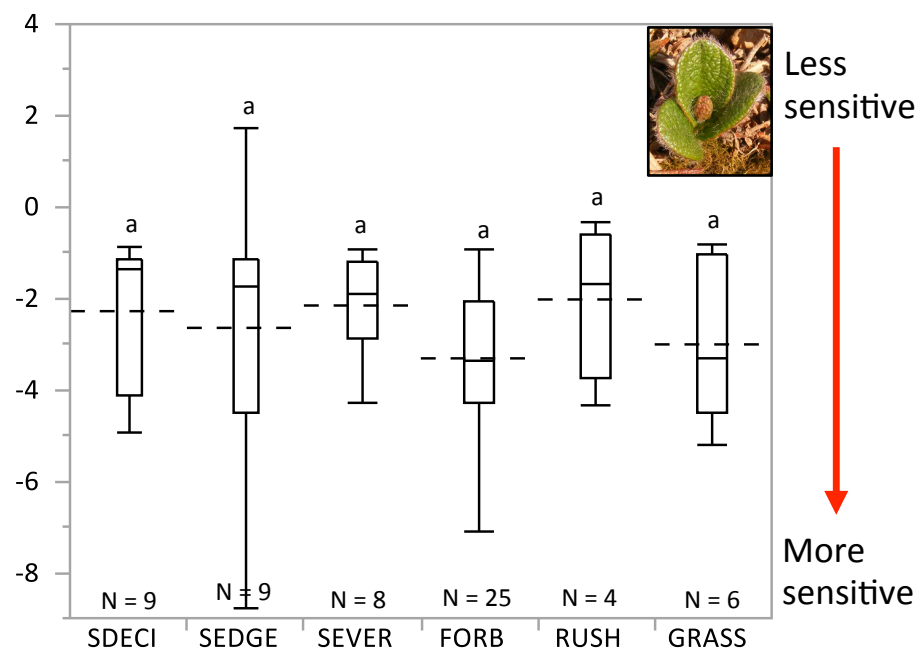
Flowering sensitivity

Deciduous shrubs least sensitive
Grass & rush most sensitive



Green up sensitivity

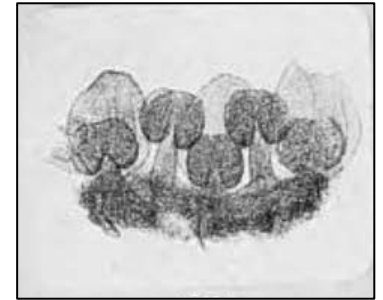
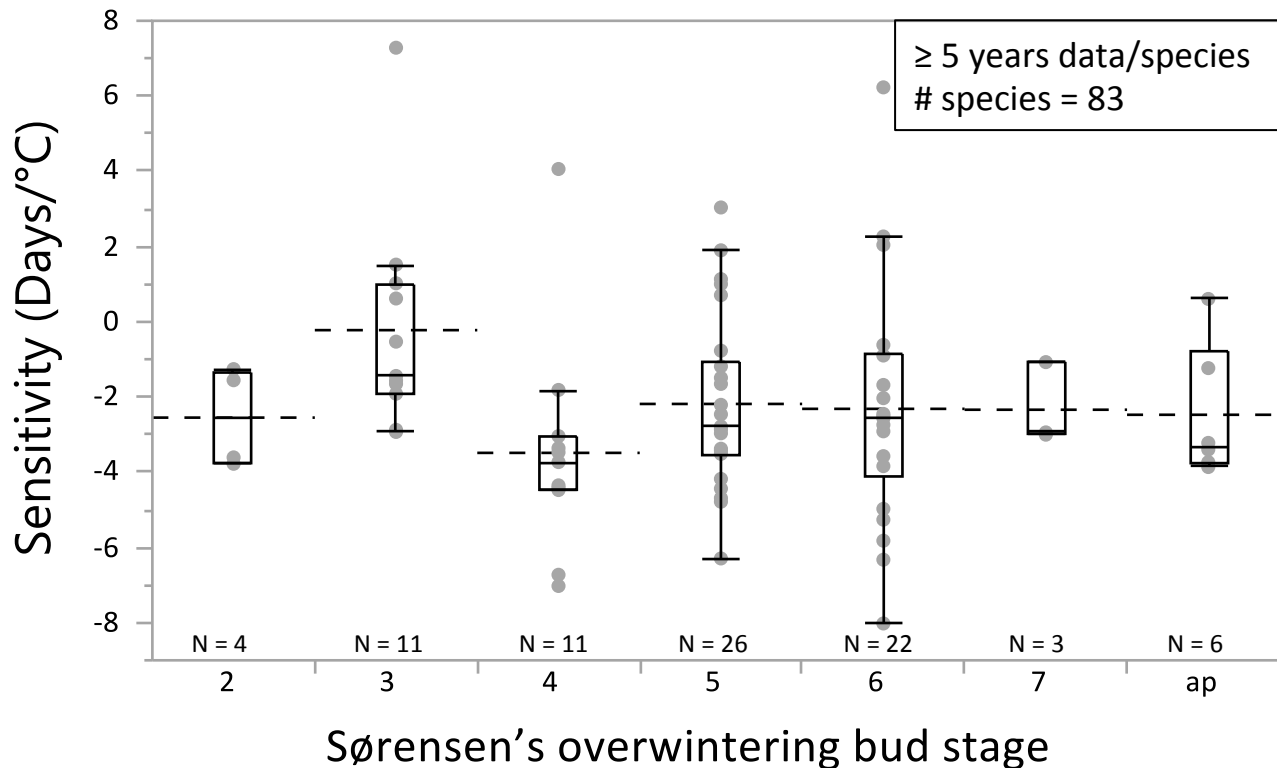
Little dif. between growth forms
Sensitivity differs from flowering



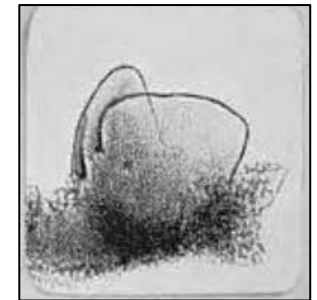
SDECI – Deciduous shrub FORB – Forbs
SEDGE – Cyperaceae GRASS – Poaceae
SEVER – Evergreen shrub RUSH – Juncaceae

Flwr ≥ 10 ; Green up ≥ 5 years of data
Flowering # species = 133
Green up # species = 61

Flowering sensitivity: Sørensen's overwintering bud stage



Saxifraga oppositifolia (stage 6)



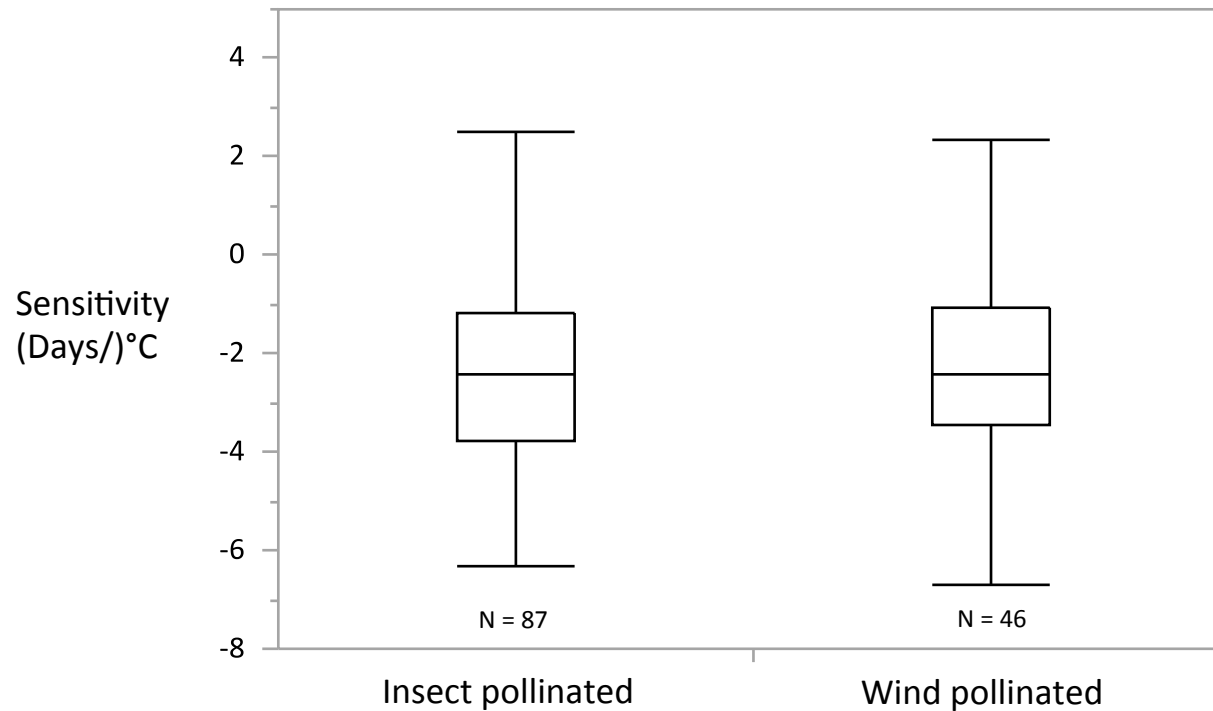
Pinguicula vulgaris (stage 2)

Sørensen, 1941

Species with less advanced bud stage 3 less sensitive than
more advanced bud stages 4, 5, 6
BUT more data required

Flowering sensitivity: Pollination

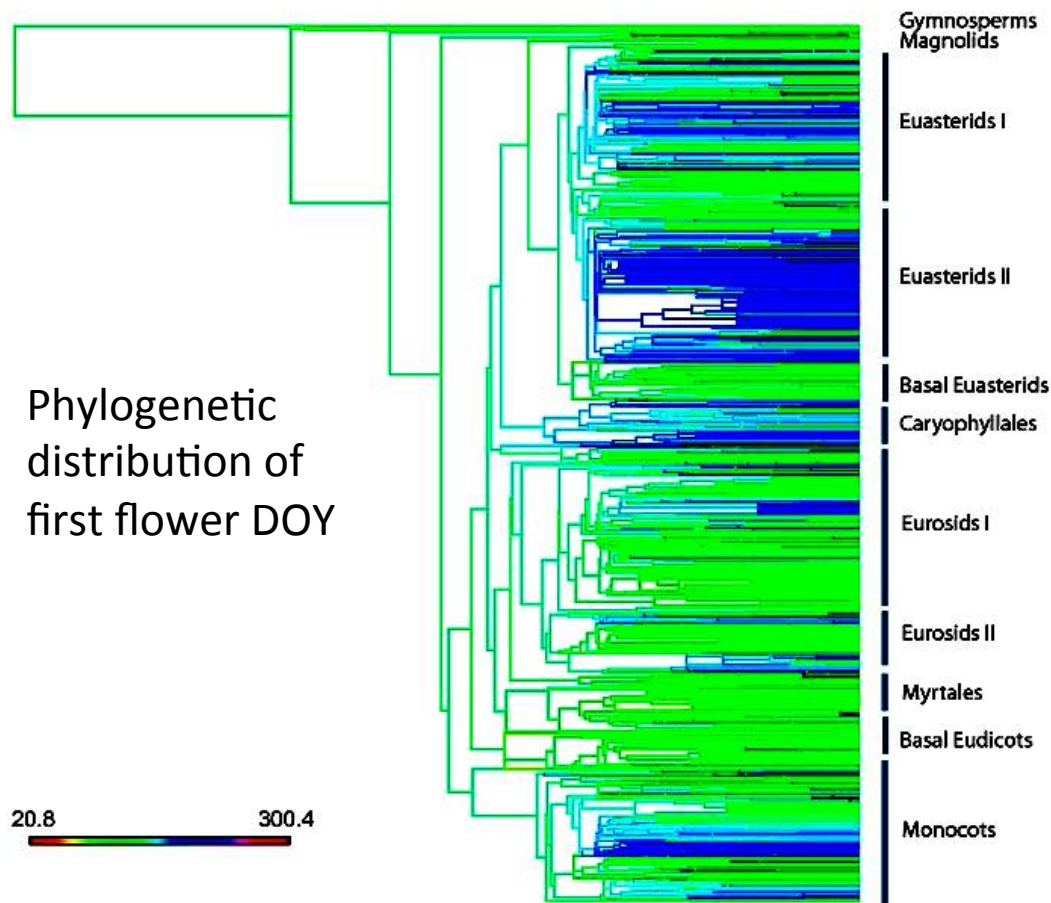
No difference in flowering time sensitivity of wind vs insect pollinated species



Species with ≥ 10 years of data
Number of species = 133

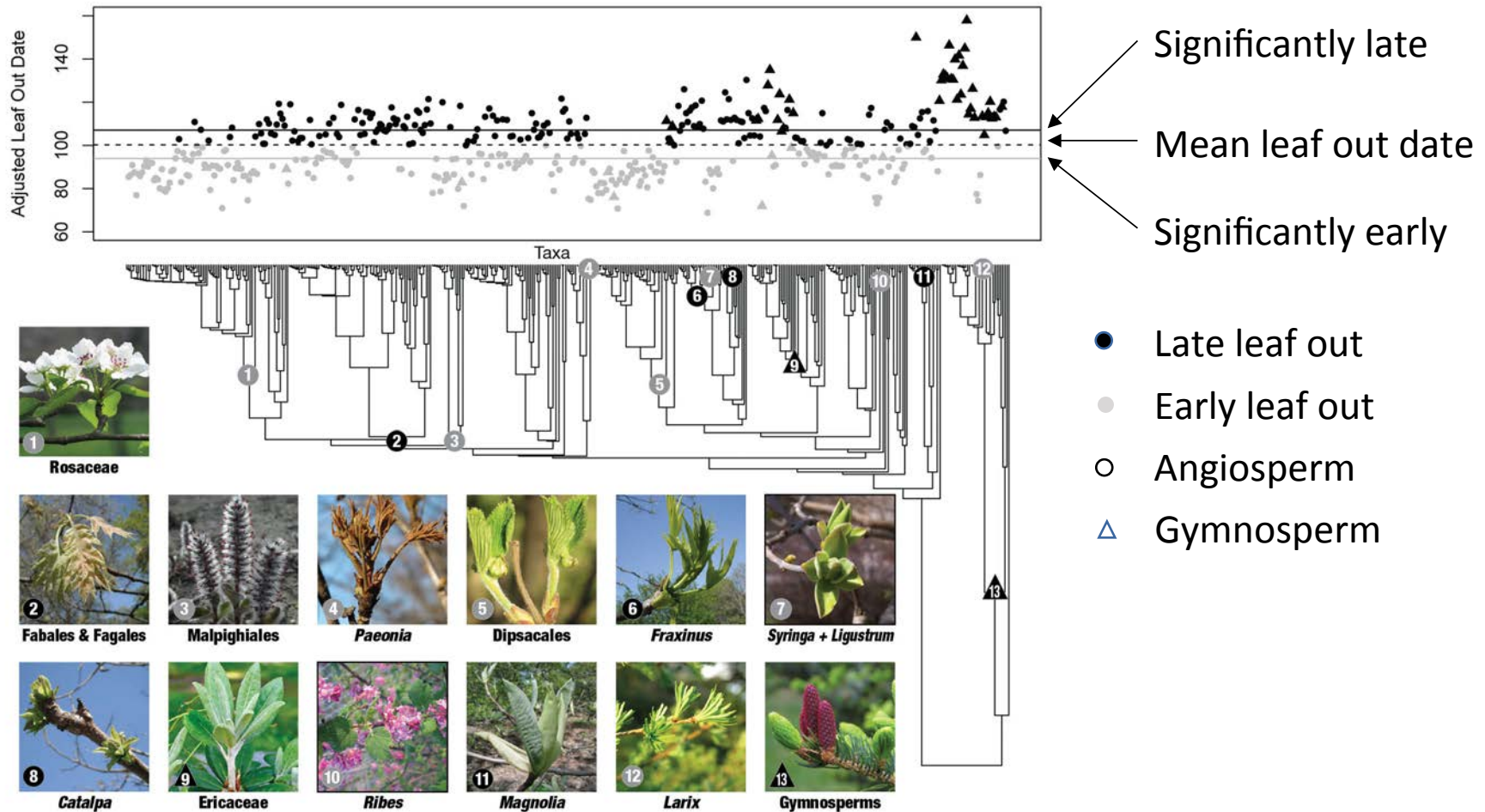
Phylogenetic analysis plan

- Determine phylogenetic signal in flowering and green up timing and sensitivity



Phylogenetic analysis plan

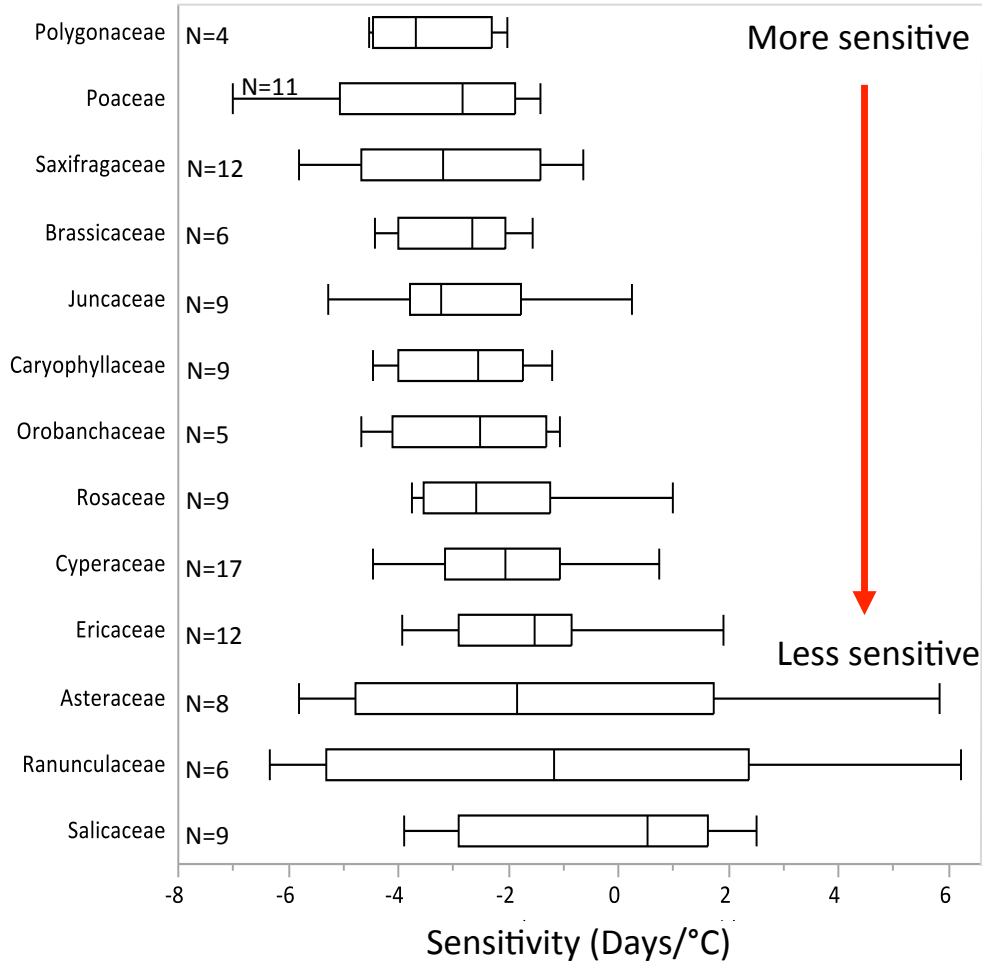
- Clades flowering/greening up significantly early/late
- Clades significantly more/less sensitive



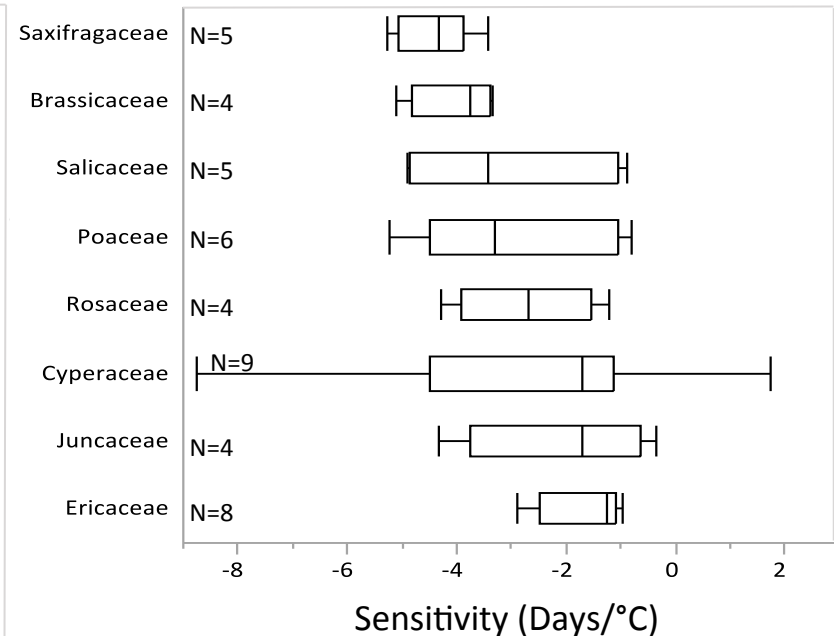
Flowering/green up β : Family



Flowering sensitivity



Green up sensitivity



Flwr ≥ 10 ; Green ≥ 5 years of data
Families with >3 species



Bayesian hierarchical model

- $\text{FlwrDOY}_{\text{sp}[ij]} \sim \text{Normal}(\mu_{\text{sp}[j]}, \sigma_{\text{sp}[j]})$
- $\mu_{\text{sp}[j]} = \alpha_{\text{sp}} + \beta_{\text{sp}} * \text{MonthlyTemp}_{\text{site}[j]}$
- $\beta_{\text{sp}} \sim \text{Normal}(\mu_{\text{sp}}, \sigma_{\text{sp}})$
- $\mu_{\text{sp}} = \beta_{\text{gf}[n]} + \beta_{\text{bs}[m]} + \beta_{\text{ps}[o]} + \beta_{\text{site}[j]}$
- $\beta_{\text{site}[j]} \sim \text{Normal}(\mu_{\text{site}}, \sigma_{\text{site}})$
- $\mu_{\text{site}} = \beta_{\text{plot}[p]} + (\alpha_{\text{lat}} + \beta_{\text{lat}} * \text{Latitude}) + (\alpha_{\text{site}} + \beta_{\text{site}} * \text{MonthlyTemp}_{\text{site}})$
- $\beta_{\text{plot}[p]} \sim \text{Normal}(\mu_{\text{plot}}, \sigma_{\text{plot}})$
- $\mu_{\text{plot}} = \alpha_{\text{habitat}[h]} + \beta_{\text{habitat}[h]}$

β = flowering sensitivity to temperature

α = intercept

observation = i

species (**sp**): s = (*Agrostis mertensii*, ...)

growth form (**gf**): n = (forb, grass, rush, ...)

bud stage (**bs**): m = (1, 2, 3, ..., 7, ap)

pollinator syndrome (**ps**): o = (insct, wnd)

Site (**site**): j = (Alex, ..., Zackenberg)

plot (**plot**): p = (1, 2, ...)

habitat (**habitat**): h = (dry, moist, wet)



Next Steps

- Feedback and input
- Gather/compile/add 2015-2017 ITEX phenology data
- Build phylogeny
- Run analysis using Bayesian hierarchical modelling
- Add non-ITEX datasets to the analysis



Questions - Suggestions



Questions

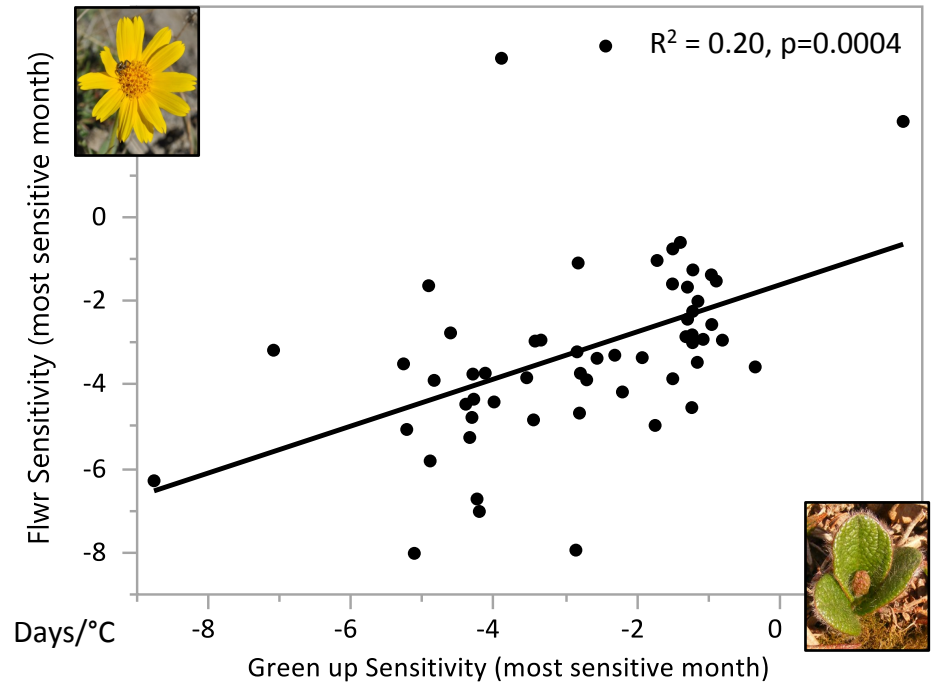
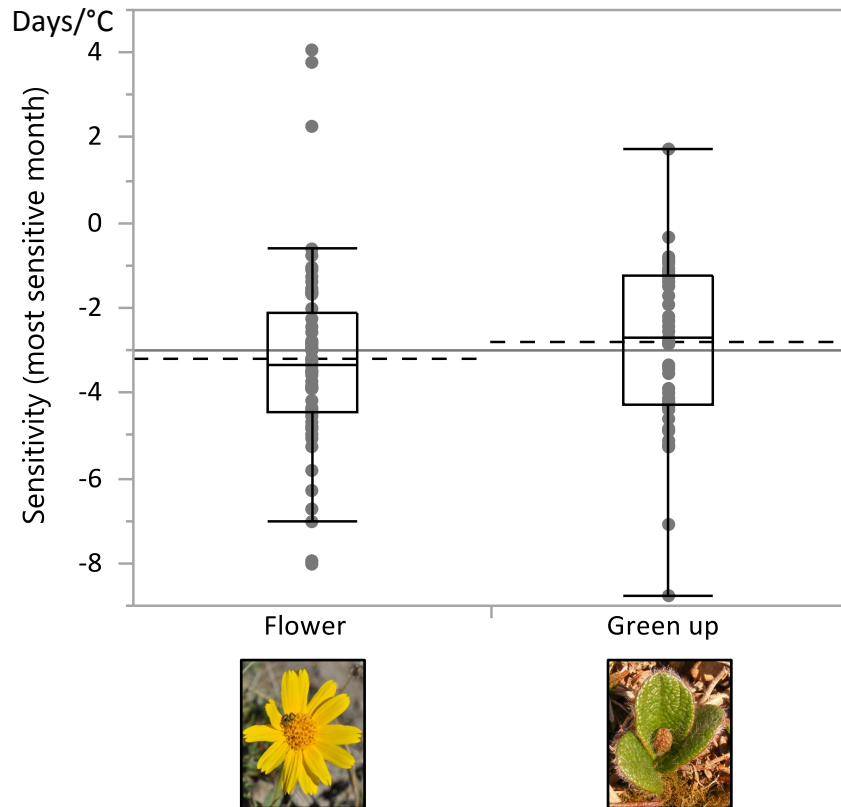
- Any comments/suggestions for Bayesian hierarchical model
- What temperatures (months) to use



Flowering versus green up time β

No significant difference in β s
Greater range flower time β

Green up time β 1.6 days/ $^{\circ}\text{C}$ more
sensitive than flower time β

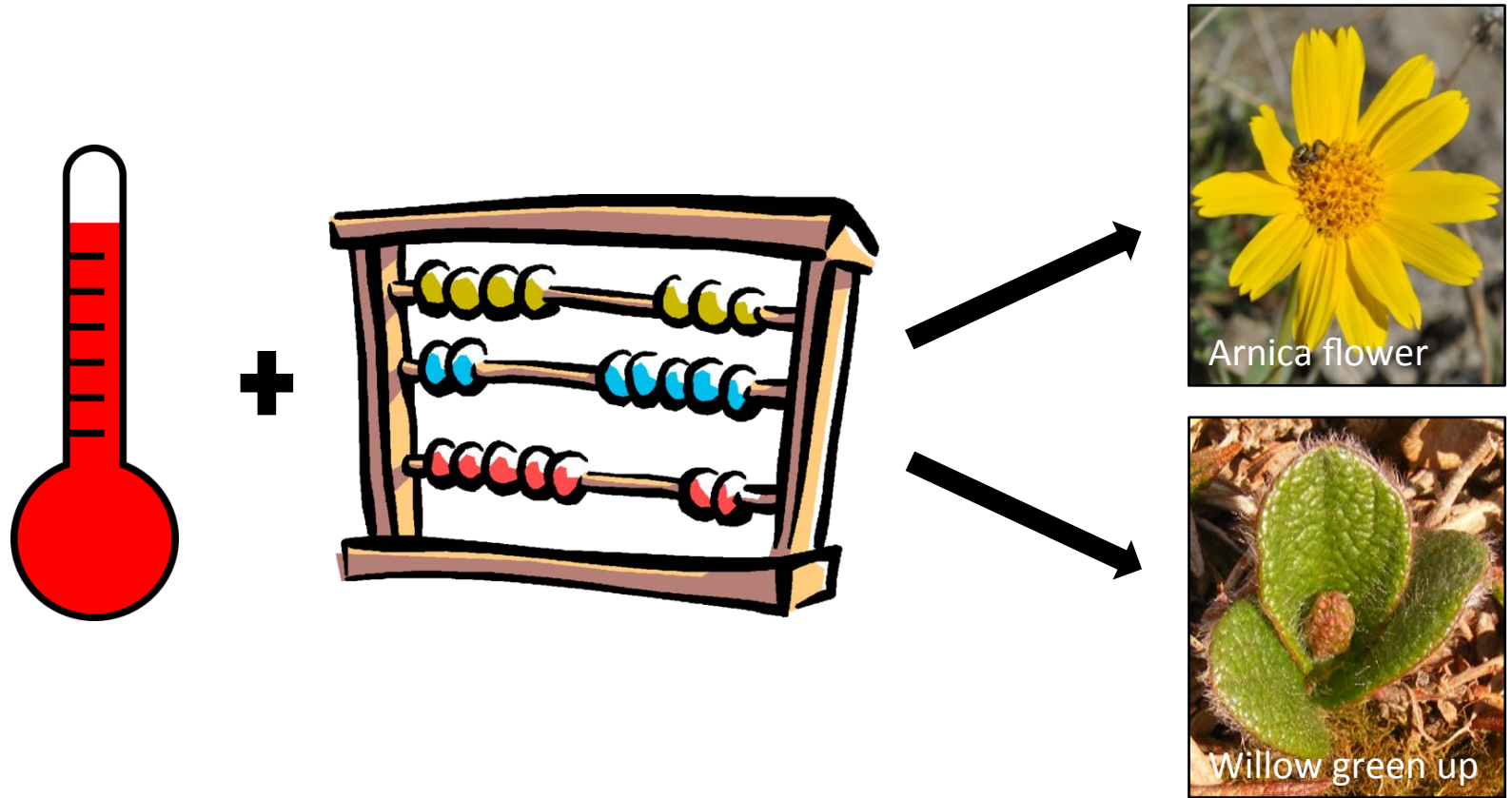


> 5 years of data
species = 57

Phenology



Timing of flowering and green up



Flowering time and green up time related to temperature
(monthly temperature or growing degree days)