Disentangling heritability and plasticity effects on *Populus fremontii* leaf reflectance across

a temperature gradient

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Abstract

Globally, vegetation biodiversity is expected to decline as the rate of plant adaptation struggles

to keep pace with rising temperatures. To support conservation efforts through remote sensing,

we disentangled the nested effects of genetic and environmental influences on reflectance

spectra, leveraging spectroscopy to assess plant adaptations to temperature. Specifically, we

quantified the relative effect of plasticity and heritability on *Populus fremontii* (Fremont

cottonwood) leaf reflectance using clonal replicates propagated from 16 populations and grown

across three common gardens spanning a mean annual temperature gradient representing the

thermal range of *P. fremontii*. We used variance partitioning to decompose phenotypic variation

expressed in the leaf spectra into genotypic and environmental components to estimate broad-

sense heritability. Heritability was strongly expressed in the spectral red edge (~680-750nm) and

shortwave infrared (~1400-3000nm), though the heritability peak in the red edge was sensitive to

extreme temperatures. By comparing distances of group centroids in principal component space,

we determined that *P. fremontii* intraspecific spectral variation was shaped by the interaction

between common garden site conditions and source population. Support vector machine models

indicated pronounced environmental influence on spectral variation, as *P. fremontii* source

population and garden location were classified at 71.8% and 92.6% accuracy, respectively. These

findings emphasize the utility of reflectance data in separating genetic and environmental

influences on plant phenotypes, offering a pathway to scale these insights across broader

landscapes and aid in the conservation and management of vulnerable ecosystems in a warming

climate.

Introduction

Globally, forests are threatened by climate change through increasing temperatures and drought, leading to widespread tree mortalities (Allen et al., 2010). Many regions have already experienced declines in biodiversity, and these declines are predicted to accelerate (Pimm et al., 2014). Remote sensing, specifically imaging spectroscopy, has emerged as a powerful tool for addressing these challenges, offering insights into biodiversity, species distributions, and ecosystem health at the landscape scale (Rose et al., 2015; Seeley & Asner, 2021; Turner et al., 2003). For example, forest conservation applications of imaging spectroscopy include accurate, high-resolution species mapping (e.g. Balzotti et al., 2020; Seeley et al., 2023c), large-scale biodiversity estimations to inform and support protected area establishment (Asner et al., 2017; Féret & Asner, 2014), and forest physiological and chemical monitoring (Brodrick & Asner, 2017; Roberts et al., 2003; Weingarten et al., 2021). Imaging spectroscopy can also be used to explore plastic responses to changing environmental conditions and heritable, intraspecific trait variation that enables species to adapt to changing climates and novel conditions (Czyż et al., 2020; Seeley & Asner, 2023; Wang et al., 2022).

Previous research has successfully linked leaf and canopy spectra to both environmental and genetic factors. For example, visible through shortwave-infrared (VSWIR) reflectance data have been used to track environmental responses, such as changes in leaf traits due to stress or climatic variation (Czyż et al., 2020; Seeley & Asner, 2023). It has also been used to distinguish individuals within a species based on their genotype (Blonder et al., 2020; Cavender-Bares et al., 2016; Seeley et al., 2023b). However, many of these studies focus on either genetic or environmental influences, with limited attention to how environmental sensitivity of plant traits, and therefore spectra, can affect phylogenetic studies or how genetic differences in plasticity

influence spectral responses to environmental adaptation. Together, these considerations have the

potential to reveal insights into how plants adapt and respond to changing conditions.

We used a common garden experiment to assess plastic and genotypically constrained

responses of leaf reflectance spectra in *Populus fremontii* (Fremont cottonwood). *P. fremontii*

inhabits riparian zones in the southwest United States and Mexico and is recognized as one of the

most important foundation species in these regions (Ellison et al., 2005; Whitham et al., 2006)

due to its ability to structure communities across multiple trophic levels, drive ecosystem

processes, and influence biodiversity via genotypically based functional trait variation (Ferrier et

al., 2012; K. C. Grady et al., 2013). Despite active planting projects, the areal extent of *P.*

fremontii has declined dramatically over the last century due to water extraction, competition

with invasive species, and climate change (Braatne et al., 1996; Shafroth et al., 1995; Stromberg,

1998). To support management decisions for *P. fremontii* and protect riparian ecosystems

vulnerable to climate change, we attempted to link leaf-level spectral data to temperature

adaptations. By deconstructing VSWIR spectra into heritable and plastic components, we aimed

to map and monitor *P. fremontii* responses and tolerance to changing climate conditions and to

derive a method for quantifying the heritable resilience (intraspecific variation in plant traits that

could result in greater ecosystem resilience; Turner, 2023; Vahsen et al., 2023) of *P. fremontii*

populations to extreme temperatures at large spatial scales.

Heritability is a statistic that estimates the degree of variation in a phenotypic trait that is

due to genetic variation between individuals of a population. Measuring heritability in plant traits

has long been a practice in selective breeding programs searching for heritable traits to maximize

yield and quality (Bradshaw, 2017). Broad-sense heritability is defined as $H^2 = v(G)/v(E)$, where

$v(G)$ is the proportion of genetic variance and $v(E)$ is the proportion of environmental variation,

typically quantified using clonal replicates (Hühn, 1975). Thus, broad-sense heritability can vary

according to the amount of environmental variation experienced by individuals. As spectroscopy

integrates a vast array of leaf traits in its analyses and is a better predictor of plant and ecosystem

function than a subset of foliar traits (Yan et al., 2021), quantifying heritability across the

reflectance spectrum may yield novel insights into genetic versus environmental controls on leaf

spectra, which are linked to functional traits, and reveal parts of the spectrum useful for

retrieving phylogenetic information.

By examining *P. fremontii* leaf reflectance across three common gardens spanning the

thermal range of *P. fremontii*, we sought to disentangle the genetic and environmental factors

shaping its adaptive capacity to varying temperatures. Specifically, our objectives were to: 1)

quantify the relative contributions of genetics (source population) and environmental influences

(common garden site) to intraspecific spectral variation, 2) explore adaptations to environmental

extremes by analyzing reflectance spectra, with a focus on phenotypic plasticity and genetic

differentiation under extreme temperatures, and 3) identify conserved and variable regions across

the VSWIR spectra of *P. fremontii* across the common gardens.

Materials and Methods

Study Sites

Three common garden sites were established across the thermal range of *P. fremontii* in

2014. Two were located at the thermal edges of the *P. fremontii* temperature range (Yuma,

Arizona: 22.8°C and Canyonlands, Utah: 10.7°C), and one garden was in the middle of this

range (Agua Fria, Arizona: 17.2°C; Figure 1; Table 1). As Yuma experiences extreme

temperatures where ambient temperatures could exceed the critical temperature at which

Photosystem II is disrupted (t_{crit}) (Moran et al., 2023), we were able to assess the responses of

leaf spectra when trees experienced air temperatures approaching and surpassing their critical

leaf temperature thresholds (Hultine et al., 2020; Moran et al., 2023).

Each garden hosted cuttings from sixteen geographically separate (minimum 13.5 km) *P.*

fremontii populations across Arizona spanning 12.2ºC Mean Annual Temperature. From each

cutting, clonal replicates were propagated in a greenhouse. 32 clonal replicates of each cutting

were planted in each common garden, across 4 stratified random blocks to minimize the effect of

within-garden variation on a population. Each common garden hosted 16 wild populations of *P.*

fremontii, eight genotypes (cuttings) from each population, and 32 clonal replicates from each

genotype, resulting in 4096 *P. fremontii* saplings per garden. Trees at the common gardens were

routinely watered to facilitate establishment in the arid southwest. A detailed description of the

common garden establishment can be found in Cooper et al. (2019, 2022).

P. fremontii in the common gardens represented two distinct ecotypes (Blasini et al.,

2021; (Bothwell et al., 2023; Ikeda et al., 2017). Three source populations were located on the

Mogollon Rim in the Colorado Plateau and are part of the Utah High Plateau/Mogollon Rim

ecotype (Blasini et al., 2021; Ikeda et al., 2017). The remaining thirteen populations represented

the environmental range of the Sonoran Desert. Two secondary ecotypes (northern and southern)

within the primary Sonoran Desert ecotype were represented (Bothwell et al., 2023). The larger,

northern clade includes nine source populations and is referred to as the main Sonoran Desert

cluster, while the southeast (SE) Sonoran Desert cluster includes three populations in the

southeastern portion of Arizona. Lastly, one population (KWF) did not fall into either Sonoran

Desert cluster.

Leaf Measurements

We measured leaf reflectance spectra from trees at the three common gardens in May 2018, four years after the saplings had been planted. Due to the differential survivorship of the saplings, the number of trees sampled from each population at each garden varied (0-52; Table S1). Three leaves were collected from each tree, and, where possible, leaves were harvested from different branches. Only full-expanded, sunlit adult leaves were chosen. Leaves were harvested by hand and transported in a cool box to the nearby field station for processing the same day.

Spectral measurements were collected using a leaf clip and Analytical Spectra Devices (ASD) field spectrometer from 350 to 2500 nm at 1-nm intervals (Analytical Spectra Devices Inc., Boulder, CO, USA). Three measurements were taken on the adaxial surface of each leaf in the center and averaged. The spectrometer was optimized using a white reference taken after every three leaves. To minimize noise, wavelengths with a low signal-to-noise ratio were removed (<400 nm and >2450 nm). Brightness normalization was applied to all spectral measurements to minimize variation not related to leaf chemistry and structure (Kruse et al., 1993; Myneni et al., 1989).

Leaf trait measurements and analyses

Following spectral measurements, leaves were scanned using a digital scanner (Canon LiDE 110) to determine their area, oven dried at 70 °C for three days and weighed for their dry mass. Leaf mass per area (LMA) was calculated by dividing the leaf's dry mass (g) by its fresh one-sided area (m^2).

The degree of stress experienced by each tree was estimated by calculating the photochemical reflectance index (PRI) (Equation 1; Thénot et al., 2002).

174   Equation 1: $PRI = \frac{\rho_{531} - \rho_{570}}{\rho_{531} + \rho_{570}}$

PRI quantifies the normalized difference between ρ_{531}, the reflectance at 531 nm, and

ρ_{570}, that at 570 nm, to capture xanthophyll pigment content and epoxidation state in leaves. The

quantity and epoxidation state of xanthophyll pigments in leaves respond to plant stress and are

an indicator of thermal tolerance and photosynthetic efficiency (Gamon et al., 1992). PRI has

been used as an indicator of plant stress at both the leaf and airborne levels (Buddenbaum et al.,

2015; Hernández-Clemente et al., 2011; Rossini et al., 2013).

We next calculated solar-weighted reflectance, or albedo, for the visible (400-700nm)

and near-infrared (NIR)-shortwave infrared (SWIR; 700-2500nm). Reflectance spectra (not

brightness normalized) were weighted by the solar irradiance spectrum (ASTM G173-03

Reference Spectra derived from SMARTS version 2.9; Gueymard, 2004) to calculate broadband

albedo (Bartlett et al., 2011) according to Equation 2, where λ represents each wavelength:

Equation 2: $Albedo \approx \frac{\sum_{i=min}^{max} Reflectance\,(\lambda_i) \cdot Incident\ Irradiance\,(\lambda_i) \cdot \delta\lambda}{\sum_{i=min}^{max} Incident\ Irradiance\,(\lambda_i) \cdot \delta\lambda}$

To determine whether the common garden sites induced changes in leaf traits, a Tukey-Kramer

multiple comparison test from the scipy python package (Virtanen et al., 2020) was applied to

the albedo, PRI, and LMA data.

Phenotypic variation in leaf spectra

To explore phenotypic variation in leaf spectra, we first visualized spectral variation

across the three common gardens. Mean source population spectra at each common garden site

were plotted and colored according to site. Spectral differences between the same populations at

different common garden sites were further visualized using principal component analysis

(PCA). PCA reduces the dimensionality of the spectral data from over 2000 dimensions to a

handful of dimensions that explain the primary spectral variation across multiple channels

(Boardman & Green, 2000; Green & Boardman, 2000; Thompson et al., 2017). We randomly

selected 104 spectra from among the five populations that were represented in all three common

gardens (CLF, KWF, BCE, PSA, and SCT). Using scikit-learn (Pedregosa et al., 2011), we

applied PCA to spectra data, plotting the first two principal components (PC) along the x and y

axes. As higher PC are often important in spectroscopy applications for discriminating

differences in surface chemistry and other traits (Dai et al., 2022; Thompson et al., 2017), we

further visualized spectral differences across ecotypes and common gardens in the first 20 PCs.

As the effect of extreme heat at Yuma was obvious in the first two PCs (Figure 3), this second

analysis included spectral data from only Agua Fria and Canyonlands. Data were grouped by

ecotype (Colorado Plateau, Sonoran Desert Main, and Sonoran Desert Southeast (SE)) and

garden, randomly selecting spectra from 40 trees from each ecotype-garden combination. We

then calculated the centroids (group mean) of each ecotype-garden combination in the first 20

PCs and measured the Euclidean distances between these centroids. Distances were then

projected into 2D space using multidimensional scaling (MDS), following the approach of

Seeley et al. (2023c).

Decomposing genotypic and environmental variation across the spectra and estimating broad-

sense heritability

216   To quantify the relative effects of environmental versus genotypic variation across the

VSWIR, we used variance partitioning to decompose observed variance at each wavelength into

the relative contribution of site, ecotype, population, genotype, or clone. All sampled data was

used across all common garden sites. Following the method of Asner et al. (2014) and Fyllas et

al. (2009), variance partitioning was calculated using a linear mixed-effects model for each

wavelength across the VSWIR using the lme4 *R* package (Bates et al., 2015). The model was

fitted using Residual Maximum Likelihood (REML). All effects were treated as random.

Common garden site was included as a crossed random effect, while clones were nested within

genotypes, which were nested within populations, which were nested within ecotype. The total

spectral variance about the mean for a given wavelength was, therefore, quantitatively parsed

into the variance explained by site ($\sigma2_s$), ecotype ($\sigma2_e$), populations within ecotype ($\sigma2_p$),

genotypes within populations ($\sigma2_g$), clones within genotypes ($\sigma2_c$) and residual variance ($\sigma2_r$)

according to Equation 3:

229   Equation 3: $\sigma2_{total} = \sigma2_s + \sigma2_e + \sigma2_p + \sigma2_g + \sigma2_c + \sigma2_r$

We next calculated broad-sense heritability across the VSWIR. Heritability at each

wavelength was calculated using the classic broad sense heritability equation ($H^2 = V_G/V_P$),

where V_G is genotypic variation, and V_P is total variation ($V_{G+}V_E$). Variance explained by

ecotypes, populations, and genotypes is categorized as genotypic variance (V_G), as the genetic

material within these groups differs. Variance explained by site and clones, as well as residual

variance, is categorized as environmental variation (V_E), or spectral plasticity, as the genetic

material within these groups is identical, so any observed differences in the spectra must be due

to environment or measurement error. Broad-sense heritability was plotted across the VSWIR

spectrum. This analysis was applied first to data from all three sites and then to data from

Canyonlands and Agua Fria alone. To obtain confidence intervals for H^2, we employed a

nonparametric bootstrap approach. Bootstrapping, a method commonly used for obtaining

uncertainty estimates for heritability (Schweiger et al., 2016), involves resampling the dataset

with replacement and recalculating the variance with each subsampled dataset. We performed

1000 iterations for each wavelength, sampling 31 spectra from each common garden x ecotype

combination, resulting in n=279 for each iteration. Mean H^2 of the 1000 iterations were

presented along with the 95% confidence interval calculated from the bootstrapped estimates.

Using leaf spectra to predict genotypic and environmental information

To explore whether leaf spectra could be used to extract both genotypic and

environmental information, we used a support vector machine (SVM) to predict *P. fremontii*

source population (genotypic variation) and garden location (environmental variation). SVMs

efficiently handle highly dimensional datasets by creating decision boundaries in n-dimensional

space that maximize the distance between class boundaries and sample points (Melgani &

Bruzzone, 2004). SVMs tend to outperform other machine learning algorithms such as random

forest for species and genotype classifications when applied to spectroscopy data (e.g. Dalponte

et al., 2012). For each radial basis function kernel SVM, we optimized hyperparameter selection

through a cross-validation grid search implemented in python with the scikit-learn package

(Pedregosa et al., 2011). As with the PCA analyses, 104 samples from each garden were

randomly selected from the same five source populations (CLF, KWF, BCE, PSA, and SCT).

For all models, data were divided into training/test datasets using a 70/30 stratified random split.

To assess the generalizability of models in unseen and extreme environments in

predicting genotypic information, we trained three additional SVM models. Each model was

trained to predict source population using *P. fremontii* spectra from two of the gardens and tested

on unseen spectra from the third garden. A different garden was withheld each time. This

allowed us to compare the applicability of the model in extreme environments and assess how

well the model can extrapolate outside of its training data.

Results

VSWIR leaf spectra responded to common garden treatment, particularly at Yuma, which

represents the upper end of the cottonwood thermal range. Brightness-normalized reflectance of

all populations at the Yuma site diverged from that at both Canyonlands and Agua Fria (Figures

2a & 3a). Albedo in both the visible and NIR-SWIR differed across the common gardens

($p<0.05$) and was highest at Yuma. In the visible, albedo was lowest at Canyonlands, and in the

NIR-SWIR, albedo was lowest at Agua Fria. PRI, the plant stress indicator where more negative

values indicate greater plant stress, was 0.018 for both Agua Fria and Canyonlands when

averaged across populations. At Yuma, mean PRI decreased to -0.011 (Figure 2b). While this

decrease was significant according to the Tukey-Kramer test ($p<0.05$), PRI did not vary

significantly between source populations.

LMA, an important driver of spectral variation, was highest in the Yuma common garden

and lowest in Canyonlands ($p<0.05$; Figure 2b). Mean LMA across all populations at

Canyonlands was 72.9 g/m^2 with a standard deviation of 10.2 g/m^2. At Agua Fria, the mean

LMA was 88.1 g/m^2, though it was the most variable with a standard deviation of 19.1 g/m^2.

Yuma exhibited the highest LMA at 108.4 g/m^2 and a standard deviation of 11.2 g/m^2.

In PC space, spectra differed across the common gardens and among the ecotypes (Figure

3). The first two PC of spectra from all common gardens explained 57.8 and 35.3% of the

variance, respectively. These components primarily separated Yuma spectra from those at

Canyonlands and Agua Fria (Figure 3a). When comparing spectra from different ecotypes at

Canyonlands and Agua Fria, the Colorado Plateau ecotype separated from both Sonoran Desert

ecotypes regardless of common garden in the first 20 PCs. Within the Sonoran Desert ecotypes

(Main and SE), common garden was the primary driver of spectral divergence in PC space.

Variance partitioning and broad-sense heritability

Across all gardens, spectral variance was highest in the visible (~500-670 nm) and red-

edge wavelengths (~680-750 nm). When Yuma was included in the analysis, variance in these

wavelengths was driven largely by the common garden site (Figures 2 and 3), and heritability in

these regions was correspondingly low with a maximum heritability of 0.08 and 0.43 for the

visible and red-edge, respectively (Figure 4). While heritability peaks near the red edge were

present regardless of the inclusion of Yuma in the analysis, the peak was shifted toward shorter

wavelengths when Yuma was excluded. Additionally, site-level variation in the visible (~500-

670 nm) and red-edge wavelengths (~680-750 nm) was minimal without Yuma. Instead,

variance at these wavelengths was driven more by genetic factors (population and genotype),

with corresponding high spectral heritability (~0.36 and ~0.66, respectively).

Across all gardens, heritability in *P. fremontii* reflectance was greatest in the SWIR

(1400 – 2450 nm), followed by the transition from red-edge to NIR (~750 nm), regardless of the

inclusion of Yuma (Figure 4). In the shorter wavelengths of the NIR (~750-1140 nm) and SWIR,

site-level variation explained less of the variance than genotypic factors in general, and the

spectral differences caused by Yuma were relatively minimal, resulting in high heritability (max:

0.51; min: 0.31). When Yuma was excluded, heritability peaked in the red-edge (740 nm;

heritability: 0.67). Heritability had a negative relationship with wavelength in the NIR until

~1320 nm where heritability increased and remained high in the SWIR, although it varied across

this region (max: ~0.56, min: ~0.37). Across the VSWIR, particularly heritable regions of the

spectra included the red-edge/NIR transition (~750 nm) and the SWIR. More environmentally

sensitive (plastic) regions included the visible and red-edge as well as much of the NIR.

SVM analysis and leaf predictions

SVM models were able to discriminate between spectra based on both common garden

site and source population at 92.6% and 71.8% accuracy, respectively (Figure 4). The SVM

trained to classify spectra based on the common garden site (environmental variation)

outperformed the SVM trained to discriminate spectra based on source population (genotypic

variation). The common garden site SVM accurately predicted spectra from the Yuma site 100%

of the time; confusion in this model occurred primarily between Canyonlands and Agua Fria,

with the true positive rate for Canyonlands and Agua Fria being 94% and 84%, respectively. For

the source population SVM, the true positive rate ranged from 61% for the KWF population to

84% for CLF. When predicting source population using one common garden site as a holdout for

testing, model performance declined. Source populations at Canyonlands were predicted with

36.0% accuracy when trained on data from Agua Fria and Yuma. At Agua Fria and Yuma,

accuracy was 32.8% and 37.5%, respectively, when SVM models were trained on data from the

other common garden sites.

Discussion

P. fremontii intraspecific variation in reflectance spectra was driven by the interaction

between common garden site conditions and source population. The heritability of spectra varied

across the VSWIR and depended on common garden site, with extreme temperatures, here

defined as the point where the ambient temperatures are at the upper threshold of the *P. fremontii*

thermal tolerance (Moran et al., 2023), at Yuma likely causing a threshold change in spectral

expression and heritability. Comparisons between the Agua Fria and Canyonlands sites revealed

greater spectral separation based on ecotype than between the common gardens. However, within

finer sub-ecotype categories, differences were primarily driven by the common garden,

reinforcing the interconnected roles of genetics and environment in shaping spectral traits. We

further demonstrated the strong environmental influence on spectral variation using SVM

models. While the SVMs classified *P. fremontii* source population with 78% accuracy among all

the common gardens, models built using spectra from two common gardens failed to generalize

to the third (<37% accuracy). Lastly, we determined that *P. fremontii* spectra had low heritability

in the visible and were most heritable in the SWIR. While heritability peaked in the red edge, the

location of the peak was sensitive to extreme temperatures.

Heritability and plasticity of reflectance spectra

P. fremontii reflectance demonstrated both plastic (responsive to common garden site)

and genotypic responses. The degree of plasticity exhibited by the spectra was influenced by the

inclusion of reflectance data from Yuma. In the first PCA analysis, the first two PCs primarily

separated the spectra according to whether the spectra were collected at Yuma or not, likely due

to the extreme heat experienced by the trees at Yuma. Similarly, Li et al. (2023) examined

genetic variation of *Nicotiana attenuata* (wilt tobacco) across common garden sites in Arizona

and Utah, and they found that PC1 of the spectra was primarily influenced by the location of the

common garden. *N. attenuata* genotype affected clustering of PC 2-4, indicating that

environment had the primary effect on leaf reflectance while genotype had a secondary effect (Li

et al., 2023). Extreme heat additionally caused an increase in visible and SWIR albedo across all

populations. Yuma is near the southern edge of the distribution of *P. fremontii* where it may not

survive future climatic conditions (e.g., Blasini et al., 2021; Moran et al., 2023), so identifying

environmental indicators of extreme environmental conditions may allow us to monitor stress at

the shifting range edge of *P. fremontii.*

Below this physiological temperature threshold, spectral differences were predominantly

governed by genetic factors, with environmental conditions playing a significant role in shaping

phenotypic expression within genetically similar groups. Excluding Yuma in the PCA resulted in

the source population (genotype) being the primary driver of separability, observed by the

separation of the Sonoran Desert and Colorado Plateau ecotypes in PC space. For the finer-scale

comparison of sub-ecotypes within the Sonoran Desert ecotypes, the environmental influence

became dominant; the two Sonoran Desert ecotypes were separated first by common garden site

(environment) and second by ecotype (genotype). This nested effect of genotype and

environment on phenotypic variation can help us understand how ecotypes adapt to future

climate scenarios, where genetic predispositions and environmental variability shape ecological

outcomes.

Across the VSWIR, we observed greater sensitivity to temperature in the visible and red-

edge regions as broad sense heritability across those wavelengths shifted in response to the

inclusion of Yuma in the analysis. As the visible region of the spectra evolved largely to

optimize light capture, this region has been strongly influenced by evolutionary forces (Meireles

et al., 2020). As such, heritability in the visible region has been well-documented in other species

(Čepl et al., 2018; Feng et al., 2017; McManus et al., 2016). For example, *Metrosideros*

polymorpha varieties that organize along environmental gradients diverged in chlorophyll

content, which was associated with differences in the visible wavelengths (Seeley et al. 2023b).

Here and in other studies, the visible region has also been highly sensitive to changing

environmental conditions, likely due to mechanisms that help protect plants from heat stress

(Mathur et al., 2014), suggesting that the visible range may not only be indicative of heat stress

but also potential adaptability to future increases in temperature.

Similarly, the red edge can be both heritable (Čepl et al., 2018; Czyż et al., 2020; Feng et

al., 2017) and indicative of stress (Abdullah et al., 2019; Dobrowski et al., 2005; Smith et al.,

2004). For example, Feng et al. (2017) found that the red edge was predictive of rice genotypes,

and both Čepal et al. (2018) and Czyż et al. (2020) observed a red edge signal in the heritability

of *Pinus sylvestris* (Scotts pine) and *Fagus sylvatica* (European beech) leaf reflectance,

respectively. We observed high heritability in the red edge relative to the rest of the VSWIR

spectra, though this peak was narrower and shifted toward longer wavelengths when Yuma was

included in the analysis, likely due to thermal stress. The shift in the red-edge heritability peak

requires further investigation as it may challenge large-scale health monitoring where vegetation

is genetically diverse and genotype classifications where stress varies.

P. fremontii heritability in the NIR also peaked at 1350 nm. These heritability peaks are

similar to those of both *Pinus sylvestris* and *F. sylvatica* (Čepl et al., 2018; Czyż et al., 2020),

though *Pinus sylvestris* had an additional region of high heritability around 1130-1330 in the

NIR that corresponded to low heritability in *P. fremontii*. As much of the NIR corresponds to

leaf structure (Knipling, 1970; Slaton et al., 2001), differences in heritability between *P.*

fremontii and *Pinus sylvestris* may relate to differences between broad and needle leaves.

Overall, heritability in the SWIR was high relative to other wavelengths, especially when

Yuma was included in the analysis. Spectral variation due to source population and genotype

(genetic factors) was greater than variation stemming from common garden sites (environmental

factors) across the NIR and SWIR, regardless of the inclusion of Yuma. When predicting the

genetic structure of *F. sylvatica* using the airborne prism experiment imaging spectrometer,

longer wavelengths (> ~1900nm) in the SWIR were more predictive (Czyż et al., 2020).

McManus et al. (2016) likewise found that the phylogenetic signal was greatest in the SWIR

across many Neotropical tree species, and Meireles et al. (2020) observed a phylogenetic signal

in Rosids, a clade of flowering plants that includes *Populus*, in the shorter wavelengths of the

SWIR. The high heritability in the SWIR, consistent regardless of temperature stress, suggests

that plant traits not directly related to photosynthesis (e.g. lignin, cellulose, and tannins; Asner

and Martin, 2016, 2009) may be driving, in part, phylogenetic signals.

Overall, environmental conditions exerted a stronger control over *P. fremontii*

intraspecific spectral variation. Much of the evidence presented above demonstrates the strong

influence of the environment on spectral variation. Additionally, we found that SVM models

based on common garden sites were ~14% more accurate than those separating source

population spectra. Cavender-Bares et al. (2016) similarly found that while the source population

of *Quercus oleoides* could be distinguished using PLS-DA models (kappa score = 0.342),

watering treatment was predicted with greater accuracy (kappa score = 0.61). When the SVM

was trained on data from two common garden sites and tested on the third, accuracy in predicting

P. fremontii source population declined by over 34%, supporting other studies that have

demonstrated how the environmental sensitivity of spectra limits large-scale species

classifications (e.g. Seeley et al., 2024). While plastic responses to heat stress may decrease

spectral heritability, environmentally sensitive regions of the spectra offer opportunities to

understand physiological responses to extreme temperatures at large scales and may serve as

environmental indicators of stress events.

Environmental thresholds

When exposed to extreme heat, plants employ a variety of mechanisms to increase their

tolerance. Plastic responses at the individual level include increased LMA (observed at Yuma),

increased heat shock proteins (HSP) to help repair heat damage, concentrations of which have

been found to correlate with LMA (Knight & Ackerly, 2003), trichome development to promote

self-shading (Banowetz et al., 2008), an accumulation of phenolics and isoprenoids for their

ability to protect photosynthetic apparatuses and antioxidant properties (Bartwal et al., 2013;

Smirnoff, 2005), and increased production of soluble sugars (Wahid & Close, 2007) and

saturated fatty acids (Ahmad & Prasad, 2011; Tang et al., 2016). As many of the physical and

chemical changes such as increased LMA, plant phenols, soluble sugars, and fatty acids can be

estimated using spectroscopy (Akkaya, 2018; Asner & Martin, 2009, 2016), it is unsurprising

that *P. fremontii* at Yuma differed spectrally from those at the other sites. For *Panax ginseng*,

differences between VSWIR reflectance of heat-stressed and control plants corresponded to

spectral regions linked to fatty acids, sugars, protein, chlorophyll, and nitrogen (Park et al.,

2021). Similarly, other studies have found differences between control and heat-stressed plants

using fluorescence (Faqeerzada et al., 2023) and visible-NIR (Kim et al., 2022) spectroscopy.

P. fremontii LMA increased along the common garden temperature gradient, contrasting

with many studies reporting a negative relationship between LMA and temperature (e.g. Poorter

et al., 2009). This increase in LMA may reflect a response to water stress, as higher LMA is a

strategy to reduce water loss under drought conditions (Poorter et al., 2009). While all gardens

were irrigated, high temperatures and drought often co-occur. Knight and Ackerly (2003)

observed increased LMA, which correlated with increases in photosynthetic recovery rates, in

desert plants when thermally stressed, identifying high LMA as a convergent trait in species like

P. fremontii that are adapted to elevated temperatures and low precipitation.

451   The shift in reflectance between Yuma trees and those at the cooler sites primarily

occurred in the visible and red-edge wavelengths, indicating that photosystems were likely

disrupted. Together, the visible and red edges are predictive of chlorophyll and other

photosynthetic pigments (Horler et al., 1983; Vogelmann et al., 1993). Both as a mechanism of

photosynthetic apparatus protection and as a side effect of the stress response, trees at Yuma had

higher reflectance in the visible range, similar to that of heat-stressed rice (Xie et al., 2019) and

maize (Tirado et al., 2021). As photosystem II is vulnerable to heat, photosynthesis is often the

first mechanism influenced by temperature (Mathur et al., 2014). Shifts in the red edge have been

linked to plant stressors such as heat and water stress (Dobrowski et al., 2005), bark beetles

(Abdullah et al., 2019), and natural gas leakage (Smith et al., 2004), all of which induced

changes in chlorophyll (Carter, 1993; Carter & Miller, 1994). PRI, a plant stress indicator

(Thénot et al., 2002), as well as the stunted growth of trees at Yuma compared to that at the other

sites, supports the conclusion that trees at Yuma were more stressed than those at the other sites.

464   In the SWIR, site-level responses in spectral variation remained regardless of the

inclusion of Yuma. Park et al. (2021) noted the importance of the SWIR for separating heat-

stressed and control plants using spectroscopy. Further, temperature-dependent leaf traits of

other *Populus* species, *P. tremuloides* and *P. deltoides*, were predictable using wavelengths in

visible and SWIR regions (Serbin et al., 2012). Both of these studies linked temperature-related

responses in the SWIR with plant traits known to respond to heat stress (Park et al., 2021; Serbin

et al., 2012). Here, site-level variation between Agua Fria and Canyonlands was primarily

observed in the SWIR and, to a lesser degree, the NIR, emphasizing the importance of using the

full spectrum to distinguish between temperature treatments.

Across the entire VSWIR spectra, albedo increased in response to extreme temperature,

likely lowering canopy temperatures and helping mitigate further heat stress. In the NIR – SWIR,

albedo decreased from Canyonlands to Agua Fria, supporting the results from Doughty et al.

(2018), where albedo declined across a temperature gradient in Peru. Regardless of which region

of the VSWIR was analyzed, albedo at Yuma was higher than at the other common gardens.

While additional evidence is required to confirm whether the increase in leaf albedo is a

universal plant response to extremely high temperatures, this phenomenon aligns with plant

ecophysiological principles, as many stress factors are known to decrease chlorophyll

concentration (Carter & Knapp, 2001).

Conclusions

Reflectance spectra of *P. fremontii* contain valuable information about both the

environmental conditions in which they grow and their genetic identities. Intraspecific spectral

variation of *P. fremontii* was shaped by both common garden site and source population, with

environmental conditions playing a dominant role under extreme stress, such as the high

temperatures at the Yuma site. At the upper thermal threshold of *P. fremontii*, the spectral

response was most pronounced in the visible and red-edge regions, corresponding to documented

heat stress responses. In these regions, spectral heritability decreased when Yuma was

considered in the analysis, suggesting that extreme environmental conditions may drive a

threshold shift in spectral expression. However, genetic information was retained, and

heritability was more consistent in parts of the NIR and SWIR.

These results underscore the potential for using VSWIR reflectance data to disentangle

genetic and environmental contributions to plant phenotypes, providing a path forward for

scaling these insights to larger landscapes. As airborne and satellite-based imaging spectroscopy

data continue to become more widely available, they offer a powerful opportunity to assess

phenotypic plasticity and heritable resilience of *P. fremontii* to extreme conditions across broad

spatial scales. Spectroscopy may be a powerful tool for monitoring the adaptive capacity of this

foundation species in the face of rapid environmental change, supporting efforts to preserve and

manage at-risk ecosystems in a warming climate.

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https://github.com/MegsSeeley/temperature_cottonwood.

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    Tables

Source Population	Elevation	Mean Annual Temperature (°C)	Mean Annual Precipitation (mm)	Ecotype
MRN	1774	10.4	593	Sonoran Desert Main
KKH	1920	10.7	258	Colorado Plateau
JLA	1507	12.3	212	Colorado Plateau
CLF	1299	14.1	176	Colorado Plateau
KWF	1126	15	243	Sonoran Desert KWF
BCE	1109	15.6	432	Sonoran Desert Main
PSA	1234	15.7	471	Sonoran Desert SE
TSZ	1219	16.9	322	Sonoran Desert SE
CAF	988	17.2	440	Sonoran Desert Main
TSE	986	17.5	402	Sonoran Desert SE
WHY	575	19.6	284	Sonoran Desert Main
CCU	696	19.9	349	Sonoran Desert Main
NRV	666	19.9	337	Sonoran Desert Main
SCT	26	22.1	88	Sonoran Desert Main
LBW	143	22.3	137	Sonoran Desert Main
CCR	70	22.6	97	Sonoran Desert Main
Common Garden Site				
Canyonlands	1581	10.7	225	
Agua Fria	998	17.2	453	
Yuma	49	22.8	93	

Table 1: Source population and study site environmental conditions. For source populations, the

ecotype is listed.

Figure Captions

Figure 1: Source population and common garden site locations in Arizona and southern Utah.

Background colors represent Köppen climate types.

Figure 2: a) Mean brightness normalized reflectance of each source population, colored by

common garden site. See Figure S1 for reflectance prior to brightness normalization. Boxplots of

albedo calculated across the b) visible and c) near-infrared-shortwave infrared, d) photosynthetic

reflectance index (PRI), and e) leaf mass per area (LMA). Within each garden, boxplots are

ordered according to source population mean annual temperature, with cooler populations on the

left and warmer populations on the right. Gaps correspond to populations that were not sampled

at a given garden. Boxplots represent a different number of sampled trees per population per site.

Figure 3: a) First and second principal components (PC) of leaf spectra colored according to the

common garden site. b) Distance between the centroids of source population grouped by ecotype

x common garden in PC space. Colors represent common garden (blue: Canyonlands, grey:

Agua Fria). Shapes correspond to the ecotype (x: Colorado Plateau, circle: Sonoran Desert,

triangle: Sonoran Desert SE).

Figure 4: Heritability of reflectance spectra (solid line) and its 95% bootstrapped confidence

interval (grey fill), mean brightness normalized reflectance spectra (dotted line), and the source

of variability (filled colors) from the visible to shortwave infrared for *P. fremontii* at a) all sites

and b) Canyonlands and Agua Fria.

Figure 5: Support vector machine confusion matrix results for predicting a) common garden site

and b) source population. Source populations are ordered left to right and top to bottom from

lowest to highest mean annual temperature.

Figures

Figure 1

Figure 2

Figure 3

Figure 4

Figure 5