Reflectance spectroscopy allows rapid, accurate, and non-destructive estimates of functional traits from pressed leaves

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Running headline: Estimating traits from pressed-leaf spectra

Abstract

1. More than ever, ecologists seek to employ herbarium collections to estimate plant functional traits from the past and across biomes. However, many trait measurements are destructive, which may preclude their use on valuable specimens. Researchers increasingly use reflectance spectroscopy to estimate traits from fresh or ground leaves, and to delimit or identify taxa. Here, we extend this body of work to non-destructive measurements on pressed, intact leaves, like those in herbarium collections.

2. Using 618 samples from 68 species, we used partial least-squares regression to build models linking pressed-leaf reflectance spectra to a broad suite of traits, including leaf mass per area (LMA), leaf dry matter content (LDMC), equivalent water thickness, carbon fractions, pigments, and twelve elements. We compared these models to those trained on fresh- or ground-leaf spectra of the same samples.

3. Our pressed-leaf models were best at estimating LMA (R^2 = 0.932; %RMSE = 6.56), C (R^2 = 0.855; %RMSE = 9.03), and cellulose (R^2 = 0.803; %RMSE = 12.2), followed by water-related traits, certain nutrients (Ca, Mg, N, and P), other carbon fractions, and pigments (all R^2 = 0.514-0.790; %RMSE = 12.8-19.6). Remaining elements were predicted poorly (R^2 < 0.5, %RMSE > 20). For most chemical traits, pressed-leaf models performed better than fresh-leaf models, but worse than ground-leaf models. Pressed-leaf models were worse than fresh-leaf models for

estimating LMA and LDMC, but better than ground-leaf models for LMA. Finally, in a subset of

samples, we used partial least-squares discriminant analysis to classify specimens among 10

species with near-perfect accuracy (>97%) from pressed- and ground-leaf spectra, and slightly

lower accuracy (>93%) from fresh-leaf spectra.

4.   These results show that applying spectroscopy to pressed leaves is a promising way to estimate

leaf functional traits and identify species without destructive analysis. Pressed-leaf spectra might

combine advantages of fresh and ground leaves: like fresh leaves, they retain some of the spectral

expression of leaf structure; but like ground leaves, they circumvent the masking effect of water

absorption. Our study has far-reaching implications for capturing the wide range of functional and

taxonomic information in the world's preserved plant collections.

Keywords: functional traits; herbarium collections; leaf chemistry; partial least-squares regression

(PLSR); reflectance spectroscopy; species identification

Introduction

The world's herbaria together contain more than 390 million specimens (Thiers 2021) which are a rich source of information about global plant diversity. Herbarium specimens are collected for many reasons—often to document where a species is present, or to serve as vouchers for taxonomic studies. But these specimens are often repurposed for new ends, unforeseen by their collectors (Meineke et al. 2018). More than ever, ecologists and evolutionary biologists seek to use herbarium specimens to measure functional traits (Heberling 2021): for example, to evaluate the long-term imprint of human activity on plant communities (Lang et al. 2018; Meineke et al. 2018); to fill in gaps in sparse trait databases (Perez et al. 2020); or to conduct comparative studies of clades (Jardine et al. 2020). Measuring functional traits on herbarium specimens carries the promise of letting us reach the inaccessible, including the past or distant parts of the world. Using herbarium specimens also allows researchers to benefit from the expertise of taxonomists and refer back to the same specimens for further use—for example, as sources of genetic data, or as references for species identification from new collections (Heberling 2021). Using specimens, researchers can address ecological and evolutionary questions that require merging functional, genetic, and distributional data at global scales.

Many functional trait measurements require destructive sampling—for example, by grinding up tissue for chemical analyses. Such measurements include most protocols to determine the elemental or molecular composition of a sample (Pérez-Harguindeguy et al. 2013). Because herbarium specimens are irreplaceable—especially those from historical collections—curators may hesitate to let them be destroyed, even in part, for ecological research. Using specimens in functional ecology might be more feasible with new, non-destructive techniques to estimate their traits.

Reflectance spectroscopy is a technique often used to estimate foliar functional traits non-destructively (Curran 1989; Jacquemoud & Ustin 2020). Spectroscopy is the study of matter's interactions with electromagnetic radiation across wavelengths (Jacquemoud & Ustin 2020); spectroscopic studies of

leaves often target reflectance—the proportion of incident radiation that is reflected—as a particularly

revealing and easy-to-measure property. A typical leaf reflectance spectrum comprises reflectance at

many narrow wavelength bands between about 350 and 2500 nm, which includes over 98% of energy

from solar radiation reaching Earth's surface (American Society for Testing and Materials, 2006).

Because the leaf's chemical and structural makeup determines how it reflects, absorbs, and transmits

light, reflectance within this range carries information about many plant traits (Cavender-Bares et al.

2017).

Two main approaches exist to estimate traits using the full information in reflectance spectra. First,

physics-based radiative transfer models like PROSPECT can be inverted to estimate a handful of traits

with well-defined optical properties (Féret et al. 2017). Second, statistical models, often created using

machine learning techniques like partial least-squares regression (PLSR), can estimate an even wider

range of traits, albeit in a less mechanistic (and perhaps less general) way (Serbin & Townsend 2020).

This multivariate empirical approach gives researchers the flexibility to predict complex traits whose

absorption features might not be as strong or well-defined (Curran 1989). Likewise, multivariate

classification techniques like partial least-squares discriminant analysis (PLS-DA) use the full spectrum to

discriminate species, lineages, or other kinds of biological classes (Meireles et al. 2020b).

Empirical approaches like PLSR are widely used to estimate plant traits from spectroscopic data

measured on fresh or ground leaves. These traits include leaf N and leaf mass per area (LMA; Asner et al.

2011; Serbin et al. 2014; Serbin et al. 2019; Streher et al. 2020), pigments (Asner et al. 2011; Yang et al.

2016), defense compounds (Couture et al. 2016; Nakaji et al. 2019), non-structural carbohydrates (Ely et

al. 2019), and even photosynthetic capacity (Yan et al. 2021). Leaf-level PLSR models have been used to

address such varied ecological topics as defense responses to herbivory (Kula et al. 2020) and the role of

biodiversity in ecosystem function (Schweiger et al. 2018). Although this multivariate statistical approach

is flexible, it is sensitive to the kind of leaf tissue used to train the model. Existing PLSR models have

mostly been trained on reflectance spectra of fresh leaves (e.g. Serbin et al. 2019) or dried, ground leaves

(e.g. Serbin et al. 2014). Such models are not expected to transfer to dried, intact leaves like herbarium

specimens because both drying and grinding cause major changes in reflectance.

We built PLSR models to estimate traits from the reflectance spectra of pressed leaves, like herbarium

specimens, and compared their accuracy to models built from fresh or dried, ground leaves. Our pressed

leaf samples were prepared like herbarium specimens but not yet mounted on paper, and our analyses

might thus serve as a proof of concept for the technique by setting aside the methodological challenges

related to working with older mounted specimens. Previously, Costa et al. (2018) had shown that a related

spectroscopic technique, Fourier Transform-Near Infrared Spectroscopy (FT-NIR), could predict several

leaf structural traits from pressed leaves of tropical trees. Here, we explicitly compare the accuracy of trait

estimation from the spectra of fresh, pressed, and ground leaves for many leaf chemical and structural

traits from multiple biomes and functional groups.

For most chemical traits, such as elemental composition or carbon fractions, we conjectured that ground-

leaf spectral models would be the most accurate, as others have found (Serbin et al. 2014; Couture et al.

2016; Wang et al. 2020). Both drying and grinding might be important in achieving this accuracy. Drying

may reveal minor absorption features of compounds in dry matter within the short-wave infrared (SWIR)

range that, in fresh leaves, are obscured by the dominant effect of water absorption (Peterson et al. 1988).

Grinding homogenizes the variation in structure and composition throughout the leaf lamina, which may

allow us to capture a more even and representative sample of tissue (Richardson et al. 2021). Since

pressed leaves are dried but not ground, we predicted they would yield intermediate accuracy for

chemical traits.

For structural traits like LMA, we instead expected fresh and pressed leaves to outperform ground leaves

because grinding disrupts the leaf structure. For water-related traits like leaf dry matter content (LDMC;

dried mass divided by fresh mass), we expected fresh leaves to outperform both pressed and ground

leaves because they retain the water absorption features that allow direct prediction of water content

(Carter 1991). Likewise, because pigments tend to degrade after collection, we expected to estimate their

concentration best from fresh leaves. We also assessed sample discoloration and considered whether it

reduces the accuracy of trait estimates, which may indicate whether spectroscopy is useful on old or

degraded specimens.

Finally, we asked whether pressed-leaf spectra can be used to identify samples to species. Reflectance

spectra often show phylogenetic signal in certain wavelength ranges because of phylogenetic

conservatism in their underlying traits (McManus et al. 2016; Meireles et al. 2020a; Diniz et al. 2020).

This signal is what often makes it possible to classify species or higher-level taxa from fresh-leaf spectra

(Cavender-Bares et al. 2016; Meireles et al. 2020b). Studies with tropical forest species have also shown

that FT-NIR absorbance spectra of pressed leaves can be used to classify species or higher-level taxa

(Durgante et al. 2013; Lang et al. 2015; Prata et al. 2018). Based on these studies, Draper et al. (2020)

proposed using spectra of herbarium specimens as part of an integrative process of species delimitation

and identification. However, it remains uncertain whether fresh- or pressed-leaf spectra are better suited

to the task of classifying species. Here, we compared the accuracy of supervised classification from fresh-

, pressed-, and ground-leaf spectra among the common species in our dataset. Because they preserve some

of the leaf structure (unlike ground leaves) and reveal the distinctive SWIR absorption features of

macromolecules and other compounds (unlike fresh leaves), pressed leaves may represent the best of both

worlds for distinguishing species using spectroscopy.

Methods

Spectral and leaf trait measurements

We trained PLSR models on leaf reflectance spectra and traits measured as part of four projects

conducted by the Canadian Airborne Biodiversity Observatory (CABO). We validated the models both

internally and on an independent dataset of pressed tree and herb samples collected at Cedar Creek

Ecosystem Science Reserve (East Bethel, MN, USA). Table 1 describes the projects and lists how many

samples and species they include for each functional group. The leaf sampling procedure is described in

the Supplementary Materials.

For each CABO sample, we measured full-range reflectance spectra (350-2500 nm) of the leaves at three

stages: (1) freshly sampled, (2) pressed and (3) oven-dried and ground into a fine powder. We measured

fresh-leaf directional-hemispherical reflectance spectra on the adaxial surface of multiple leaves or leaf

arrays from each sample using spectroradiometers equipped with integrating spheres. We pressed a

portion of the sample, and measured reflectance spectra on the adaxial surface of pressed leaves between

six months and three years later using a spectroradiometer with a leaf clip (Fig. S1). Lastly, we measured

ground-leaf spectra using a spectroradiometer with a benchtop reflectance probe that pressed loose leaf

powder into an even pellet with very low transmittance. We trimmed all spectra to 400-2400 nm. Detailed

notes on measurement and post-measurement processing of reflectance spectra are found in the

Supplementary Materials.

While measuring pressed-leaf spectra, we inspected each pressed specimen by eye to note signs of

discoloration in preparation or storage. While all leaves have some changes in their appearance as they

dry, we were particularly interested in the loss of green color, such as blackening, browning, or the

development of a silvery or whitish finish on the leaf surface. We scored each leaf on a discrete scale

from 0 to 4 (see examples in Figs. S2-5). A score of 0 indicates no noticeable discoloration. Scores 1

through 4 indicate increasing discoloration, from 1 (either <10% blackening/browning or development of

a slight silvery finish to the leaf) to 4 (>75% blackening/browning).

We measured the following leaf structural and chemical traits on each CABO sample: Leaf mass per area

(LMA; kg m^{-2}), leaf dry matter content (LDMC; mg g^{-1}), equivalent water thickness (EWT; mm), carbon

fractions (soluble cell contents, hemicellulose, cellulose, and lignin; %), pigments (chlorophyll *a*,

chlorophyll *b*, and total carotenoids; mg g^{-1}), and concentrations of a variety of elements (Al, C, Ca, Cu,

Fe, K, Mg, Mn, N, Na, P, Zn; % or mg g^{-1}). Protocol summaries are in the Supplementary Materials.

PLSR modeling for trait estimation

We used a PLSR modeling framework to predict each trait from each of fresh-, pressed-, and ground-leaf

spectra across the full range (400-2400 nm). PLSR is suited to handle spectral datasets, which have many

collinear predictors, because it projects the spectral matrix onto a smaller number of orthogonal latent

components in a way that maximizes the ability to predict the response variable. We simply used

reflectance values as predictors, since calculating various common transformations of reflectance (see

Serbin et al. 2014) did not increase predictive accuracy in preliminary tests. We also did not transform

any trait values to reduce skewness, again finding in preliminary tests that it did not improve predictive

accuracy enough to warrant the added complexity. Some studies restrict the wavelengths used in

prediction, often to ranges known or assumed to contain features relevant to a given trait (e.g. Serbin et al.

2014). We considered that the 400-1300 nm range might be most liable to change in storage due to

degradation of photosynthetic pigments and accumulation of brown pigments that absorb in the near-

infrared range (NIR; Fourty et al. 1996). Thus, we also built pressed-leaf models restricted to 1300-2400

nm ('restricted-range models'), which might be expected to generalize better to datasets that include older

or more discolored leaves. We present these results mainly in the Supplementary Materials.

Our methods for model calibration and validation largely follow Burnett et al. (2021). First, we randomly

divided the data into calibration (75%) and validation (25%) datasets, stratified by functional group. We

began by fitting a model for each trait on the calibration dataset. We selected the smallest number of

components for which the root mean squared error of prediction (RMSEP) from 10-fold cross-validation

fell within one standard deviation of the global minimum. We used this number of components—a

different number for each trait—in further analyses to predict traits on the internal validation dataset. We

calculated the variable importance in projection (VIP) metric for calibration models to see which parts of

the spectrum were most important for predicting each trait (Wold et al. 1994).

To test how well we could predict traits on the internal validation subset, we first did a jackknife analysis

by iteratively (100×) dividing the 75% calibration data further into random 70% training and 30% testing

subsets. For each trait, we trained models on the 70% using the previously determined optimal number of

components and predicted the remaining 30%. This analysis gave us a distribution of model performance

statistics across the 100 iterates (R^2, %RMSE), which reveals the sensitivity of model performance to

randomly varying sets of training and testing data.

Next, we applied the 100 jackknife models for each trait to the 25% internal validation dataset, yielding a

distribution of 100 predictions for each validation sample. We quantified model performance using R^2 and

root mean squared error (RMSE) between measurements and mean predictions. We also report the RMSE

as a percentage of the 2.5% trimmed range of measured values (%RMSE), which we used rather than the

entire range (as in e.g. Burnett et al. 2021) for robustness to outliers. For each trait, we also tested whether

the magnitude of residuals (observed minus predicted) in the validation dataset varied among leaves with

different discoloration scores. We performed all statistical analyses in *R v. 3.6.3* (R Core Team 2020) and

used package *pls v. 2.7.1* (Mevik et al. 2019) for PLSR modeling.

In our main set of models, we kept chemical traits on a mass basis for consistency with the usual basis on

which such traits (except pigments) are measured and reported. However, some traits are most often

distributed proportionally to area, and some users may have reasons to prefer area-based estimates

(Kattenborn et al. 2019), so have we also made area-based models available (see Table S1 for

performance summary statistics). All models (mass- and area-based) are available to download (see *Data*

availability).

External validation

To test how well our PLSR models would transfer to a fully independent dataset, we applied the ensemble

of pressed-leaf models for five traits (LMA, LDMC, EWT, N, and C) to pressed-leaf spectra from Cedar

Creek, then compared the model-derived trait estimates to measured values. Like most of the CABO

dataset, the Cedar Creek dataset comprises trees and herbs from northeastern temperate North America,

but it includes an entire functional group (needleleaf conifers) absent among the CABO projects in this

study. We collected the spectra with the same instrument and foreoptic as the pressed-leaf spectra in the

CABO dataset, but used different sampling, preparation, and measurement protocols. We aimed to see

whether the inclusion of a new functional group and the various subtle differences in protocols would

affect the models' performance. Full details on this dataset are provided in Supplementary Materials.

PLS-DA modeling for species classification

We tested the potential to classify species with fresh-, pressed-, and ground-leaf spectra using partial

least-squares discriminant analysis (PLS-DA; Barker & Rayens 2003). We took spectra from the ten most

common species in our dataset—all of which are deciduous trees except *A. flexuosa*, which is evergreen.

Each was represented by at least 20 specimens (~480 total). For each tissue type, we divided the full

dataset into 60% calibration and 40% validation subsets, stratified by species. In the *R* library *caret v.*

6.0.84 (Kuhn 2020), we trained models on the calibration subset using 10-fold cross-validation repeated

10 times. We chose the number of PLS components during cross-validation by maximizing Cohen's

kappa (κ), which describes the agreement between the true and predicted species identities while

accounting for the probability of agreement by chance. Imbalanced training data can bias classification

algorithms (Sun et al. 2009), so we used a two-step procedure to balance classes while maintaining

enough training data and avoiding overfitting. First, we downsampled within better-represented species

classes at random so that all classes had equal size, then chose the number of components (n) that

maximized κ. Second, we upsampled from less-represented classes at random with replacement so that

classes had equal size—again maximizing κ, but restricting the range of components evaluated to no more

than the *n* chosen during the downsampling step. We applied the cross-validated PLS-DA model from the

upsampling step to the validation subset and summarized its performance using raw classification

accuracy and κ.

Results

Patterns in traits and reflectance spectra

We saw large variation among samples in each of our target traits within the CABO data, ranging from

1.4-fold variation in C to more than 20-fold variation in traits like lignin, P, K, and Zn (Table 2). The

ranges of most traits in our dataset covered a large portion of the global distributions in the TRY dataset,

but tended to be narrower at both extremes (Kattge et al. 2020). Many traits—including LMA, LDMC,

EWT, cellulose, and many elements—had distributions with a pronounced skew (most often positive).

Broadleaf trees tended to have higher LDMC, C, and lignin than other growth forms. Among the herbs,

grasses had very high hemicellulose and cellulose and low lignin content, while forbs often had high N.

Some of the trait variation was driven by specific projects; for example, *A. flexuosa* in the Warren project

tended to have particularly high LMA, Na, and C and low N.

Both pressed and ground leaves had higher median reflectance across nearly the entire spectrum (Fig. 1),

as expected based on changes in water content and structure (Carter 1991). Indeed, water absorption

features (the largest of which are centered around 1450 and 1930 nm) largely disappeared in pressed and

ground leaves. The red edge between the visible and NIR regions was also blunted by both pressing and

grinding, causing the global maximum of median reflectance to shift from 872 nm (fresh) to 954 nm

(pressed) to 1313 nm (ground).

Within each tissue type, the coefficient of variation (CV) of reflectance was generally highest where

average reflectance was lowest. Across tissue types, pressed-leaf spectra tended to show greater absolute

variation in reflectance throughout much of the spectrum, particularly towards the tails of the distribution

(e.g. the middle 95 percent in Fig. 1B). The species that have the most exceptionally reflective pressed

leaves across the spectrum (mainly *Phragmites australis* (Cav.) Trin. ex Steud., *Phalaris arundinacea* L.,

and *Asclepias syriaca* L.) do not have particularly reflective fresh leaves, leaving it uncertain why their

pressed leaves are so reflective. In contrast, discolored leaves tended to have lower reflectance throughout

the visible and NIR ranges (Fig. S5). Unlike pressed leaves, ground leaves showed very low absolute

variation in reflectance throughout the SWIR, likely because grinding eliminates variation in leaf

structure. However, they showed high variation from 700 to 1100 nm, which may also result from varying

degrees of discoloration.

PLSR modeling for trait estimation

Tested on the internal validation dataset, pressed-leaf models performed best at predicting LMA ($R^2 =$

0.932; %RMSE = 6.56), C ($R^2 = 0.855$; %RMSE = 9.03), and cellulose ($R^2 = 0.803$; %RMSE = 12.2).

These traits were followed by a mixture of water-related traits, carbon fractions, and pigments, as well as

N and Ca (all $R^2 = 0.582$-0.790; %RMSE = 12.8-17.0). Although some other elements (Mg, P) could be

estimated with $R^2 > 0.5$ and %RMSE < 20, most models for other elements showed lower accuracy (Table

2; Figs. 2-3 and S6-12). In most cases, restricted-range (1300-2400 nm) models had similar predictive

accuracy to full-range models (Table S2).

We compared pressed-leaf models to both fresh-leaf and ground-leaf models. The optimal number of

components selected to predict each trait was between 3 and 26. For any given trait, ground-leaf models

usually had the most components, often followed by pressed-leaf models (Table 2). Fresh-leaf models

were best for predicting the structural and water-related traits—LMA, LDMC, and EWT (Figs. 2-3 and

S6-12). Ground-leaf models were best for predicting chemical traits, like carbon fractions and most

elements. For most traits, pressed-leaf models had intermediate performance, although for some (e.g.

pigments, LDMC) both fresh- and ground-leaf models performed better. Statistics from jackknife

analyses showed that model performance was more variable for traits that were predicted less accurately

(Figs. S13-15). There was no correlation between the magnitude of residuals from pressed-leaf models

and our discoloration index for any trait ($p > 0.05$).

For all traits except LMA and Fe, the VIP metric for fresh-leaf spectra showed a global maximum

between 710 and 720 nm (Fig. 4 and S16-18)—wavelengths slightly longer than the typical inflection

point of the red edge (Richardson et al. 2002). Many traits also show high VIP across the green hump at

~530-570 nm. Bands in the NIR range were less important for predicting most traits than much of the

visible range. The SWIR range was generally important for predicting LMA, EWT, Na, and pigments,

and many other traits showed several local peaks of importance, most prominently at about 1880 nm, but

also near 1480 and 1720 nm.

For predicting traits from pressed-leaf spectra, the general trend held that visible reflectance and certain

ranges in the SWIR were important for predicting most traits, while the NIR and much of the shorter

SWIR (800-1750 nm) were less important (Fig. 4 and S16-18). The red edge peak of importance for most

traits was near 705 nm. Other prominent local maxima for many traits lay close to 1440, 1720, 1920,

2130, and 2300 nm. We saw broadly similar patterns in ground-leaf spectra, except that VIP for most

traits was lower at longer SWIR wavelengths (2000-2400 nm).

External validation

For most traits, pressed-leaf model performance on the external validation dataset from Cedar Creek was

not quite as strong as the internal validation (Table 3; Fig. 5). For C, the models performed very poorly

($R^2 < 0.05$). Among the remaining traits, R^2 ranged from 0.345 (LDMC) to 0.876 (LMA). For N in

particular, %RMSE was high (37.6%) due to bias—N concentrations were underestimated for conifers

but slightly overestimated for remaining samples.

Since conifers were absent from the CABO training dataset, we considered whether the models we built

would extend to this new functional group. For LDMC, models performed better when excluding conifers

(R^2 = 0.406, %RMSE = 22.8) than when retaining them (R^2 = 0.345; %RMSE = 24.7). In contrast, for

LMA and EWT, models performed better when retaining conifers. For LMA in particular, estimates for

conifers were quite good, and their extension of the trait range raised R^2 (from 0.752 to 0.876) and

reduced %RMSE (from 16.5 to 10.3). Restricted-range models yielded somewhat better external

validation R^2 for N and LDMC both including and excluding conifers (Table S3; Fig. S19). For LDMC in

particular, this improvement resulted from improved estimates for extremely discolored samples of

Populus tremuloides Michx.

PLS-DA modeling for species classification

PLS-DA models using pressed- and ground-leaf spectra showed near-perfect performance at classifying

species (Fig. 6). Models using fresh-leaf spectra were slightly worse but still showed strong performance.

The optimal fresh-leaf model, which had 28 PLS components, correctly predicted the taxonomic identity

of 175 of the 188 samples in the training dataset (κ = 0.920; $p < 0.0001$). The best pressed-leaf model,

which had 37 PLS components, correctly predicted 184 of the 188 samples (κ = 0.975; $p < 0.0001$). The

best ground-leaf model, which had 48 PLS components, correctly predicted all 189 samples (κ = 1; $p <$

0.0001). The majority (>70%) of misclassifications were between congenerics.

Discussion

We show that we can estimate a wide range of leaf functional traits for 68 woody and herbaceous species

from reflectance spectra of pressed leaves (Table 2; Figs. 2-3 and S6-12). Model performance was highest

for LMA, C, and N, followed by a mixture of water-related traits, carbon fractions, pigments, and a few

important elements (Ca, Mg, and P). Other elements could only be estimated with fairly low accuracy.

These results show that pressed-leaf spectra provide an integrative measure of leaf phenotypes, much like

fresh-leaf spectra (Cavender-Bares et al. 2017), but with stronger potential to characterize variation in

chemical traits. Perhaps as a result, we could use pressed-leaf spectra to classify species as accurately as

ground-leaf spectra, and better than fresh-leaf spectra. Our results underscore the potential that using

reflectance spectroscopy on herbarium specimens could yield rapid and non-destructive estimates of

many functional traits, enabling more expansive studies of trait variation across space and time.

Comparing PLSR model performance

We compared pressed-leaf models to fresh-leaf and ground-leaf models from the same samples. Our

findings about which kind of tissue was best for predicting each trait mostly supported our hypotheses.

Ground-leaf spectra showed the strongest performance for most chemical traits, likely because grinding

homogenizes the lamina and removes the potentially confounding influence of leaf structure (Table 2).

Pressed-leaf spectra showed intermediate performance for most chemical traits, perhaps because, like

ground leaves, they lack the major water absorption features that mask the smaller features of other

compounds in the SWIR range (Peterson et al. 1988). Contrary to our predictions, ground-leaf spectra

performed about as well as fresh-leaf spectra, and better than pressed-leaf spectra, for estimating pigment

concentrations. The same factors that provided an advantage to ground-leaf spectra in estimating chemical

traits also perhaps explain why pressed- and (especially for LMA) ground-leaf spectra performed worse

for estimating water-related and structural traits (LMA, LDMC, and EWT). Pressed-leaf models may

represent a good compromise in allowing many traits to be estimated with mostly intermediate but

nonetheless quite high accuracy.

Our pressed-leaf models often performed as well as fresh- and ground-leaf models published here and

elsewhere. For example, our models for LMA had an RMSE (0.00970 kg m^{-2}) lower than many fresh-leaf

models from the literature, including Serbin et al. (2019; 0.015 kg m^{-2}), Nakaji et al. (2019; 0.015 kg m^{-2}),

and Streher et al. (2020; 0.051 kg m^{-2}). The ground-leaf models in Serbin et al. (2014) had a validation

RMSE of 1.4 and 2.4 for cellulose and lignin percentages, comparable to 1.38 and 1.81 for our pressed-

leaf models (Table 2). On the other hand, Serbin et al. (2014)'s ground-leaf models for N performed

better than our pressed-leaf models (2014; validation RMSE = 0.13 vs 0.297). Our models' error could be

within acceptable bounds for addressing many questions about large-scale ecological or evolutionary

patterns that encompass a wide range of trait variation. For some traits (e.g. EWT, N, K, Mn) many of the

samples with the greatest errors were at the poorly sampled tails of the measured trait distribution, which

suggests that more thorough sampling may be needed to ensure models can make reliable predictions at

these extremes (Figs. 2 and 3). Nevertheless, our external validation analyses indicate that our models for

some important traits—like LMA, LDMC, EWT, and N, but not C—can transfer reliably to other

datasets, and even sometimes to new functional groups (Fig. 5).

Interpreting PLSR model performance

It may seem perplexing that we could succeed at all in predicting LMA from ground-leaf spectra, or

LDMC and EWT from pressed- and ground-leaf spectra. The ability to estimate these traits must not

result from the optical expression of the traits themselves. We suggest that we instead sense these traits

via their correlations with other traits that have a stronger optical expression. This kind of effect—a

"constellation effect" (*sensu* Chadwick & Asner 2016; Nunes et al. 2017)—has been invoked to explain

the ability to estimate traits like rock-derived nutrients (Nunes et al. 2017) and $\delta^{15}N$ (Serbin et al. 2014)

that are not known to have strong absorption features in the measured range of wavelengths. However,

models that rely on such constellation effects could break down when patterns of trait covariance vary

(Kothari & Schweiger 2022), which could make models for certain traits fail beyond the domain of the

training data.

The potential role of constellation effects makes it important to interpret how our models work. The VIP

metric helps us understand what features are most important, but interpreting it can be challenging

because reflectance at a given band is never driven by a single trait. Fresh-leaf VIP for most traits had

peaks in the visible and SWIR ranges, with a global maximum at 705 nm (Fig. 4 and S16-18). The pattern

of high VIP along the green hump and red edge is common in PLSR models from fresh leaves (Yang et

al. 2016; Ely et al. 2019; Streher et al. 2020; Yan et al. 2021). The red edge (particularly 700-725 nm)

may be so important because of its sensitivity to both chlorophyll content and leaf structure (Richardson

et al. 2002). Much of the visible range was proportionally even more important for pressed- and ground-

leaf models, and most of the NIR was less important (except for LMA; Fig. 4 and S16-18). There were

multiple small VIP peaks in the SWIR. Although some (e.g. at 1440 and 1920 nm) lie within major water

absorption features, any causal link to the leaf's fresh water content is unlikely for pressed and ground

leaves. Many of these peaks also lie near broad absorption features for many components of dry matter,

including protein, cellulose, lignin, and starch, which complicates their interpretation (Curran et al. 1989;

Fourty et al. 1996).

With some exceptions, the VIP metric showed that the same bands are often important for predicting

different traits. This pattern might be taken as an artifact of trait covariance: For example, the three

pigment pools covaried strongly ($R^2 = 0.827$-0.969) and had nearly identical VIP across the spectrum

(Figs. S16-18). One might take similarities in VIP further to imply that there are a small number of traits

whose tight coordination with others underlies the performance of all models through constellation

effects. Nevertheless, across the whole dataset, many traits covaried only weakly but still shared VIP

patterns. For example, EWT, cellulose, N, and K were not tightly coordinated ($R^2 = 0.003$-0.152) but

shared similar patterns of pressed-leaf VIP across the spectrum (Fig. 4), including peaks at 705 and 1920

nm. While VIP is a useful heuristic, it does not show the direction in which a band's reflectance alters

trait estimates; the same bands may matter for different traits in different ways. Here, similarities in VIP

do not appear to result solely from strong networks of trait covariance. Nevertheless, the fact that we can

estimate traits like LDMC and EWT from pressed-leaf spectra appears to imply some role for trait

covariance, perhaps in a more diffuse way.

PLS-DA modeling for species classification

PLS-DA models showed that fresh-, pressed-, and ground-leaf spectra alike could be used to classify

species with perfect accuracy for ground leaves, near-perfect accuracy (>97%) for pressed leaves, and

excellent accuracy (>93%) for fresh leaves (Fig. 6). In contrast to prior work that deliberately selected

many congenerics (Lang et al. 2015), our most common species were often distantly related. Among the

misidentified samples, most were mistaken for congenerics, which implies that related species are more

spectrally similar (Schweiger et al. 2018; Meireles et al 2020a). However, past studies using dry leaves

have shown great success with closely related species (Lang et al. 2015; Prata et al. 2018) or even

populations (Stasinski et al. 2021).

Our analysis reinforces that that pressed- or ground-leaf models might be particularly suited to the task of

classifying or delimiting species (Fig. 6). This finding is notable because measuring spectra of pressed

leaves in an herbarium is also much simpler than measuring spectra of fresh leaves through an intensive

field campaign across the range of a clade. We conjecture that these models have an advantage because

drying reveals the absorption features of multiple compounds in the SWIR range that might together

allow finer discrimination of species than water content does. Indeed, ground-leaf spectra have greater

intrinsic dimensionality than fresh-leaf spectra (Kothari & Schweiger 2022), which suggests they have

more independent axes of variation along which species may separate. Our results support the growing

practice of using spectra of pressed herbarium specimens in species delimitation and identification (Prata

et al. 2018; Draper et al. 2020). However, classification models from pressed or ground leaves have less

relevance for research using remotely sensed imagery, which is dominated by fresh leaves.

The future of spectroscopic trait estimation

Although trait predictions from spectral models are not perfect, they have a few advantages over

conventional trait measurements: they (1) can be non-destructive; (2) are fast and require relatively little

training; and (3) have very low marginal cost, despite the high capital cost of buying a spectrometer

(Costa et al. 2018). These advantages could make it easier to address questions that require large datasets

of functional traits. But researchers may be deterred if they must each build their own models tailored to

particular uses—and for herbarium specimens, it might not be possible to do the destructive trait

measurements often needed to train the models. Ideally, spectral models would be general enough that

researchers could confidently use them without further validation, but this aim is not easy to achieve: for

several reasons, a model trained on any particular spectral dataset may make poor trait predictions on new

data. As with any other technique, the goal for spectroscopic trait estimation is to improve model

accuracy and generality as much as they can be jointly improved. Below, we discuss some challenges one

by one, particularly as they concern pressed leaves.

One concern is that the new data could be outside the range of traits or optical properties in the training

dataset (Schweiger 2020). A general model, if such a thing is possible, would need to represent the vast

range of leaf functional traits and optical properties. Another kind of concern about model generality

concerns sample preparation before spectral measurements. For example, particle size influences ground-

leaf spectra (Foley et al. 1998). For pressed leaves, it may be particularly important to prepare samples in

consistent ways that preserve the leaves' anatomical integrity. In our external validation analyses, we

found that pressed-leaf models yielded reasonably accurate predictions of most traits, even though the

validation dataset differed in sample preparation protocols and included conifers, which were absent from

the training dataset. Nevertheless, even setting aside conifers, external validation for one trait (C) was

very poor, and for another (N) showed noticeable bias—enough that researchers might need to develop

their own correction factors to use the model in practice.

Another class of challenges concerns spectrometers and their foreoptics. Spectra of fresh leaves can be

measured with different foreoptics, including integrating spheres, contact probes, or leaf clips. We used a

leaf clip with pressed specimens because mounting delicate pressed leaves in an integrating sphere could

damage them. While leaf clips and contact probes often have a higher signal-to-noise ratio, they are less

likely to produce consistent measurements among instruments or replicate samples due to variation in

viewing geometry and anisotropic surface reflectance (Petibon et al. 2021). The logistical constraint of

having to use them on pressed leaves could thus make it harder to compare data among instruments. In

theory, the greater inconsistency of leaf clip measurements could have reduced the performance of our

pressed-leaf models compared to our integrating sphere-based fresh-leaf models, but we still found that

the former performed better for most chemical traits.

Another challenge is that while many herbarium specimens are glued to a paper backing, measuring

reflectance with a leaf clip or probe usually requires placing a black absorbing background under the

sample to keep transmitted light from being reflected back into the sensor. When unattached leaves are

not available, using spectra from these specimens may require new methods to correct for reflectance

from the mounting paper.

Spectrometers and their software also vary subtly in their sensors and techniques for processing spectra

(Castro-Esau et al. 2006), and in some cases researchers must take steps to reconcile spectra measured

from different instruments (Meireles et al. 2020b). Assuming that these kinds of technical challenges can

be overcome, our results and others encourage confidence about building general models to estimate traits

from a wide variety of plants (Serbin et al. 2019). The creation of open libraries for spectral data (like

EcoSIS, https://ecosis.org/; or the CABO data portal, https://data.caboscience.org/leaf) and spectral

models (like EcoSML; https://ecosml.org/) will contribute to this goal. Lastly, we note that many of the

same concerns about discrepancies among sampling and measurement protocols could arise when using

existing spectral libraries to aid in species identification (Draper et al. 2020).

Implications for herbarium-based research

A particular challenge for herbarium specimens is that their optical and chemical properties (especially

light-sensitive pigments) may degrade during preparation or storage. Such degradation could make it hard

to distinguish changes in the traits of living plants over time from changes in storage. Even in this study,

where no specimens were collected before 2017, many underwent visible changes in color, including

browning or blackening; ~12% were scored at 2 or higher, with large variation among species (e.g. 42%

of *Populus grandidentata* specimens, but 0% of *Betula papyrifera* specimens). We found little evidence

that such discoloration hinders trait estimation: Both the results of our discoloration analyses and the

similar performance of full- and restricted-range models suggest that PLSR is flexible enough to predict

traits despite the variable influences of discoloration in our specimens. This capacity likely depends on

using samples for model calibration that show a similarly wide range of discoloration.

Our specimens were collected no more than three years before measurement, but ecologists may want to

use specimens collected decades ago. While our findings give reason to be optimistic that properly

calibrated models could return accurate estimates of many traits from old or discolored specimens, it

remains untested whether there are any limits to this potential. In general, not much is known about long-

term changes in specimen chemical (aside from DNA) or optical properties (Lang et al. 2018). Color

changes are known to accelerate under certain preparation and storage conditions—including exposure to

some chemical preservatives and high light, drying temperature, or humidity (Bridson & Forman 1999;

Metsger & Byers 1999)—which it may be worth avoiding when possible. Long-term studies of

specimens—perhaps subjected to varying preparation and storage techniques—could establish how

chemical and optical properties change over time and help refine these guidelines further.

Some of the challenges we describe pertain to projects that would measure spectra on samples already

collected, but spectroscopy—like other novel uses for herbarium specimens—could also prompt changes

in collection practice. For example, it underscores the potential value of gathering and storing extra leaf

material (e.g. in fragment packets), which would circumvent the challenge of measuring mounted leaves

and aid destructive analyses of herbarium specimens (Heberling 2021). We propose that herbaria could

also incorporate spectroscopy into their operations by measuring incoming specimens shortly after

pressing, which could mitigate the challenges caused by mounting and degradation.

Linking spectral data measured on herbarium specimens to the digital record of the voucher could be a

powerful tool to enable data synthesis, but it may require new informatic tools (Draper et al. 2020). The

hyperdimensionality of the reflectance spectrum could make it hard to accommodate within existing

standards like the Darwin Core (Wieczorek et al. 2012), at least without extensions. One could link

records to external spectral databases like EcoSIS or SPECCHIO (Hueni et al. 2020), which are also

designed to store metadata about instrumentation and processing. We would advocate for coordination

between herbarium managers and researchers who use reflectance spectroscopy, which could build

agreement about best practices for spectral measurement and curation and allow data to be synthesized

and compared across research groups.

We show that non-destructively measured pressed-leaf spectra retain much of the information about many

leaf functional traits found in fresh-leaf spectra. While validating this technique on older specimens will

require extensive further research, our findings suggest that reflectance spectroscopy could allow herbaria

to take on a greater role in plant functional ecology and evolution. Our study has far-reaching

implications for capturing the wide range of functional and phenotypic information in the world's

preserved plant collections.

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Conflict of interest statement

The authors have no conflicts of interest to declare.

Author contributions

SK, JCB, and EL conceived the ideas and designed the methodology. RBR and SK collected the data. SK

analyzed the data and wrote the first draft of the manuscript. All authors made substantial contributions to

further drafts and gave final approval for publication.

Data availability

All fresh-leaf spectral data are available through the CABO data portal (https://data.caboscience.org/leaf).

Upon publication, we will upload other spectral data, as well as metadata and trait data, to the Ecological

Spectral Information System (EcoSIS, https://ecosis.org/), and upload models to the Ecological Spectral

Model Library (EcoSML, https://ecosml.org/). At that stage, we will update this section accordingly.

Tables

Table 1: A summary of CABO projects used for model building and internal validation (above the thick black line) and the Cedar Creek dataset used for external

validation (below the thick black line). The column heading 'broadleaf' refers to broadleaf trees only.

Project	Description	Samples (species) per functional group					
		Broadleaf	Conifer	Herb	Shrub	Vine	Total
Beauchamp-Rioux	Deciduous forest trees sampled throughout the growing season in 2018 at several sites in southern Québec and Ontario	405 (9)					405 (9)
Dessain	Forbs, grasses, shrubs, and broadleaf trees from forests and open areas sampled throughout the growing season in 2017 at four sites in southern Québec	31 (24)		15 (13)	26 (22)		72 (59)
Boucherville	Forbs, shrubs, and grasses sampled throughout the growing season in 2018 from the Parc national des Îles-de-Boucherville in southern Québec			45 (8)	22 (3)	6 (1)	73 (12)
Warren	A single broadleaf evergreen tree species (*Agonis flexuosa* (Willd.) Sweet) collected across a soil chronosequence (Turner et al. 2018) in southwestern Australia	68 (1)					68 (1)
Total		504 (25)		60 (19)	48 (23)	6 (1)	618 (68)
Cedar Creek	Herbs and broadleaf and coniferous trees sampled at Cedar Creek LTER in Minnesota in late summer and fall 2018	177 (10)	101 (4)	55 (14)			333 (28)

599 Table 2: Summary statistics of PLSR model internal validations from full-range fresh-, pressed-, and ground-leaf models. %RMSE is calculated as RMSE divided

600 by the 2.5% trimmed range of measured values within the validation subset. Global ranges are derived from the TRY database (Kattge et al. 2020; accessed 13

601 April 2021), omitting data with error risk greater than 3. Carbon fractions (solubles, hemicellulose, cellulose, and lignin) and carotenoids are each represented by

602 <1250 points in TRY, which may limit the estimates of the global range. EWT is not available on TRY.

Trait	Dataset range	Global range	Fresh				Pressed				Ground			
			No. comps	R^2	RMSE	%RMSE	No. comps	R^2	RMSE	%RMSE	No. comps	R^2	RMSE	%RMSE
LMA (kg m^{-2})	0.0233-0.217	0.00866-0.354	15	0.961	0.00740	5.02	9	0.932	0.00970	6.56	18	0.858	0.0139	9.44
LDMC (mg g^{-1})	155-573	68.2-857	16	0.830	27.5	9.25	13	0.671	38.5	12.9	19	0.800	29.9	10.1
EWT (mm)	0.0488-0.282	NA	7	0.868	0.0153	10.4	10	0.790	0.0193	13.2	18	0.733	0.0213	14.5
Solubles (%)	38.5-88.9	20.0-74.6	10	0.591	5.03	16.3	13	0.743	3.97	12.8	22	0.810	3.54	11.5
Hemicellulose (%)	2.77-34.1	8.71-45.0	10	0.577	3.15	17.6	12	0.730	2.49	13.8	19	0.707	2.73	15.3
Cellulose (%)	5.19-25.8	7.65-42.6	22	0.645	1.85	16.5	16	0.803	1.38	12.2	24	0.934	0.791	7.05
Lignin (%)	0.918-21.5	2.27-32.4	8	0.551	2.10	18.4	13	0.675	1.81	15.8	22	0.818	1.35	11.8
C (%)	39.5-53.6	25.5-60.3	11	0.751	1.09	11.7	17	0.855	0.839	9.03	22	0.932	0.562	6.06
N (%)	0.883-5.62	0.14-7.27	12	0.692	0.329	15.1	15	0.754	0.297	13.6	21	0.961	0.117	5.38
Chl a (mg g^{-1})	1.24-15.5	1.38-22.1	5	0.724	1.09	14.2	8	0.675	1.18	15.4	16	0.744	1.04	13.6
Chl b (mg g^{-1})	0.486-5.29	(total Chl)	5	0.693	0.402	14.5	4	0.582	0.469	16.8	7	0.671	0.413	14.9
Carotenoids (mg g^{-1})	0.193-3.05	0.00102-5.1	5	0.672	0.235	15.9	8	0.632	0.251	17.0	20	0.790	0.189	12.8
Al (mg g^{-1})	0-0.379	0.004-19.7	6	0.326	0.0423	23.0	8	0.350	0.0418	22.7	10	0.504	0.0362	19.7
Ca (mg g^{-1})	1.64-36.9	0.68-69	14	0.436	4.83	19.7	17	0.752	3.23	13.1	26	0.877	2.22	9.02
Cu (mg g^{-1})	0-0.052	0.00154-0.064	6	0.428	0.00471	21.6	20	0.314	0.00508	23.1	9	0.500	0.00443	20.3
Fe (mg g^{-1})	0.0150-0.266	0.0128-1.1	3	0.165	0.0399	21.7	8	0.209	0.0390	21.2	8	0.394	0.0341	18.6
K (mg g^{-1})	1.06-38.2	0.3-78.1	7	0.553	3.23	19.3	16	0.488	3.33	21.4	14	0.710	2.61	15.6
Mg (mg g^{-1})	0.482-7.86	0.277-15.6	14	0.243	0.890	24.5	10	0.514	0.714	19.6	18	0.603	0.659	18.2
Mn (mg g^{-1})	0-1.02	0.00560-5.37	4	0.242	0.182	22.1	13	0.386	0.165	20.0	17	0.487	0.148	18.0
Na (mg g^{-1})	0-4.87	0.00499-48.7	9	0.480	0.508	21.8	10	0.494	0.503	21.5	8	0.353	0.565	24.3
P (mg g^{-1})	0.222-7.18	0.1-9.9	10	0.449	0.859	17.3	20	0.621	0.714	14.3	12	0.550	0.776	15.6

| Zn (mg g⁻¹) | 0.00142-0.647 | 0.00221-0.242 | 8 | 0.361 | 0.0658 | 25.1 | 11 | 0.444 | 0.0615 | 23.4 | 11 | 0.517 | 0.0571 | 21.8 |

Table 3: Summary statistics for external validation of pressed-leaf PLSR models. The models were trained on CABO data (see Table 2) and applied to a dataset

605   collected at Cedar Creek.

Trait	Including conifers				Excluding conifers			
	Range	R^2	RMSE	%RMSE	Range	R^2	RMSE	%RMSE
LMA (kg m⁻²)	0.0179-0.229	0.876	0.0186	10.3	0.0179-0.136	0.752	0.0133	16.5
LDMC (mg g⁻¹)	133-525	0.345	69.9	24.7	133-504	0.406	65.7	22.8
EWT (mm)	0.0602-0.372	0.666	0.0423	17.1	0.0602-0.363	0.587	0.0338	18.7
C (%)	35.7-59.8	0.035	4.48	25.6	35.7-54.1	0.043	4.23	27.2
N (%)	0.75-4.68	0.546	0.721	37.6	0.75-4.68	0.548	0.792	31.0

  Figures

Fig. 1: Distributions of spectral reflectance and its coefficient of variation (CV) among fresh (A), pressed (B), and ground

(C) leaf samples. A solid black line connects the median reflectance at each wavelength. Dark blue and light blue ribbons

denote the middle 50% and 95%. The solid red line shows the coefficient of variation at each wavelength. In (D), the

medians of all three are shown for comparison.

Fig. 2: Internal validation results for LMA, EWT, cellulose, and chlorophyll *a* predicted from fresh- (left), pressed-

(middle), and ground-leaf (right) spectra. In each panel, each functional group has a separate ordinary least-squares

regression line overlaid on top of the thick dashed 1:1 line. The error bars for each data point are 95% confidence intervals

calculated from the distribution of predictions based on the ensemble of 100 models produced from jackknife analyses.

Figs. 2 and 3 display a subset of traits that represent various functions and span the range of model performance. Plots for

remaining traits are in the Supplementary Materials.

Fig. 3: Internal validation results for three elements (N, K, and Mn) predicted from fresh- (left), pressed- (middle), and

ground-leaf (right) spectra, displayed as in Fig. 2.

Fig. 4: The variable importance of projection (VIP) metric calculated based on fresh- (left), pressed- (middle), and ground-leaf (right) models. As in Figs. 2 and 3, we selected seven traits that represent a range of patterns in VIP across the tissue types (cell = cellulose). The dashed horizontal line at 0.8 represents a heuristic threshold for importance suggested by Burnett et al. (2021). VIP plots for remaining traits are in the Supplementary Materials.

646   Fig. 5: External validation results for various traits in Cedar Creek data predicted using models trained on CABO data.

647   Panels are displayed as in Figs. 2 and 3.

Fig. 6: Confusion matrices for partial least-squares discriminant analysis from (A) fresh-, (B) pressed-, and (C) ground-

leaf spectra. Rows specify the true species identity, while columns specify the models' predictions. Each row sums to 100:

numbers on the diagonal represent the percentage of specimens of each species that were correctly classified, while off-

diagonals represent the percentage misclassified as other species. Full binomials: *Agonis flexuosa*, *Acer rubrum* L., *Acer*

saccharinum L., *Acer saccharum* Marshall, *Betula papyrifera* Marshall, *Betula populifolia* Marshall, *Fagus grandifolia*

Ehrh., *Populus grandidentata* Michx., *Populus tremuloides* Michx., and *Quercus rubra* L.

B

C

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 Supplementary Materials

Sampling

For each sample, we collected a large group of sunlit leaves (>3 h estimated sun exposure per day) at

about the same vertical position—for trees, the same branch or neighboring branches from the same

individual. We avoided leaves with noticeable damage from herbivores or pathogens. We measured fresh-

leaf spectra and fresh-leaf mass in the field, froze leaf disks for pigment analyses, and transported samples

in a cooler to the lab. We then measured other whole-leaf traits and divided the remainder of the sample

into two groups: (1) a portion to be pressed, and (2) a portion whose leaves were dried and ground. We

measured pressed-leaf spectra on the former portion prior to mounting, and ground-leaf spectra and

chemical traits on the latter. Although these are distinct subsets of leaves, we treat them as identical in

their trait values since they are divided at random from the larger, relatively homogeneous bulk sample. A

minimum of one representative pressed specimen of each species at each site and date is prepared as a

specimen at the Marie-Victorin Herbarium in Montréal, Québec, Canada.

We classified species' growth forms following the Database of Vascular Plants of Canada (VASCAN;

Desmet & Brouillet 2013). We classified *Agonis flexuosa* as a broadleaf tree and manually disambiguated

some species listed as adopting either shrub or three growth forms.

Spectral measurements

For the Beauchamp-Rioux, Boucherville, and Warren projects, we measured fresh-leaf directional-

hemispherical reflectance spectra on each sample in 2018 using an HR-1024i spectroradiometer outfitted

with a DC-R/T integrating sphere from Spectra Vista Corporation (SVC; Poughkeepsie, NY, USA). We

measured spectra on the adaxial surface of six leaves or, for small or narrow leaves, single-layer leaf

arrays following the protocol in Noda et al. (2013) as adapted for our instrument (Laliberté & Soffer

2018a; Laliberté and Soffer 2018b). For the Dessain project, we measured fresh-leaf reflectance spectra in

2017 using a FieldSpec 4 spectroradiometer equipped with an RTS-3ZC integrating sphere from

Analytical Spectral Devices, Inc. (ASD; Boulder, CO, USA) following the leaf spectroscopy protocol

from the Global Airborne Observatory (https://gao.asu.edu/spectranomics-protocols). We collected

spectra from a variable number of leaves or arrays per sample—six in most cases, but as few as three or

as many as eighteen from certain samples. For all projects, each reflectance measurement was calibrated

against a white Spectralon 99% reflectance panel and corrected for stray light.

Each pressed specimen was dried in a plant press between sheets of newspaper in a setting with constant

air flow and at most gentle heating ($\leq 30\ °C$) for a minimum of 48 h. For all projects, we measured

reflectance spectra on the adaxial surface of pressed, intact leaf samples using a PSR+ 3500

spectroradiometer with a leaf clip foreoptic (Spectral Evolution, Haverhill, MA) between six months and

three years after collection. We calibrated reflectance against a white Spectralon reflectance panel and

measured 3-7 spectra per sample. Although we pressed samples until dry, some nevertheless dried in a

way that created an uneven leaf surface. At the time of measurement, we noted any visible discoloration

that samples underwent during drying or storage, including browning or blackening (Figs. S2-5). In

general, we aimed to avoid uneven or discolored areas when measuring spectra, although it was

sometimes not entirely possible when discoloration was pervasive.

Each ground-leaf bulk sample was oven-dried at 65 °C for three days, ground using a cyclone mill with 2

mm mesh, redried at 65 °C, and stored for chemical analyses. For all projects, we also measured ground-

leaf spectra using the PSR+ 3500 spectroradiometer with a benchtop reflectance probe foreoptic that

pressed loose powder into a smooth, even pellet. For each sample, we added at least 0.6 g of leaf powder

into a sample tray; in preliminary tests, adding additional material did not change the spectra, suggesting

that transmittance was close to zero. We calibrated reflectance against a white Spectralon reflectance

panel placed in an identical sample tray. We measured three spectra per sample on the same pellet,

turning the sample tray 120° between the first two spectra, and loosening and mixing the powder before

reshaping the pellet between the second and third spectra.

A small number of specimens did not have usable spectra at all three stages, usually either due to lack of

enough pressed or ground material or to poor spectral quality at any stage. To ensure fair comparisons

among the stages, we removed any sample from the dataset that was not available for at least two stages.

Among the remaining samples, any sample that was not available for all three stages (<4%) was placed in

the training subset for PLSR analyses.

Our spectral processing pipeline varied by the type of data. Although the actual spectral resolution was

more variable and coarser, the ASD Field Spec 4 and Spectral Evolution PSR+ 3500 spectrometers'

software automatically resampled the spectra to 1 nm resolution and interpolated over the overlap region

between sensors (Schweiger & Laliberté 2020). For data from the SVC HR-1024i spectrometer, we

performed these steps using linear interpolation. For pressed-leaf spectra only, we detected jumps in

reflectance at the sensor overlap region and removed them using the function *match_sensors()* in *R*

package *spectrolab v. 0.0.10* (Meireles et al. 2017). Next, we averaged all spectra for a given sample and

tissue type (fresh, pressed, or ground). Spectra measured with integrating spheres tend to have greater

noise, especially at the ends. To reduce noise in fresh-leaf spectra, we applied a Savitzky-Golay filter

using *R* package *signal 0.7.6* (Signal Developers, 2013), with varying order and length: order 3 and length

21 from 350-715 nm, order 3 and length 35 from 715-1390 nm, order 3 and length 75 from 1390-1880

nm, and order 5 and length 175 from 1880-2500 nm. Finally, we trimmed all spectra to 400-2400 nm,

removing the extremes where sensors show greater noise. We did all spectral processing in *R v. 3.6.3* (R

Core Team 2020) using *spectrolab v. 0.0.10* (Meireles et al. 2017).

Leaf trait measurements

We performed all leaf trait measurements excluding petioles, but including the rachis for compound

leaves, since the rachis is functionally analogous to the midrib of a simple leaf. We measured the

following leaf structural and chemical traits: Leaf mass per area (LMA; $kg\ m^{-2}$), leaf dry matter content

(LDMC; $mg\ g^{-1}$), equivalent water thickness (EWT; mm), carbon fractions (soluble cell contents,

hemicellulose, cellulose, and lignin) on a mass basis, pigments (chlorophyll a, chlorophyll b, and total

carotenoids) on a mass basis, and concentrations of a variety of elements (Al, C, Ca, Cu, Fe, K, Mg, Mn,

N, Na, P, Zn).

From a subset of the leaves for each sample whose fresh-leaf spectra we measured, we used a 7 mm cork

borer to punch leaf disks (or cut fragments of narrow leaves) in the field and stored them on ice in the

darkness before transferring them to a -80 °C freezer in the lab. We extracted pigments under darkness in

an $MgCO_3$-MeOH solution and estimated pigment concentrations using a spectrophotometric protocol on

a SPECTROstar Nano microplate reader (BMG LABTECH, Ortenburg, Germany; Girard et al. 2020).

From a separate subset of leaves from each sample, we weighed fresh mass in the field shortly after

collection. We then rehydrated the leaves in sealed plastic bags with damp paper towels for at least 12 h

and weighed them for rehydrated mass. We scanned leaves and estimated their area using WinFOLIA

(Regent Instruments, Québec, QC, CA). Lastly, we dried them for at least 72 h in a drying oven at 65°C

before reweighing for dry mass. We calculated LDMC as total dry mass divided by total fresh mass and

LMA as total dry mass divided by total rehydrated area (Laliberté 2018). We further calculated EWT

from LDMC and LMA. For the Dessain project alone, fresh mass in the field was not available, so we

used rehydrated mass to calculate LDMC and EWT, which may lead our reported values to underestimate

LDMC and overestimate EWT. These biases are likely to be relatively small since, in the remaining

projects, relative water content was generally close to 100% (median: 89.5%, 2.5–97.5th percentile: 79.2–

97.8%).

We analyzed ground-leaf bulk samples for N and C using a Vario MICRO Cube combustion analyzer

(Elementar, Langenselbold, Germany; Ayotte et al. 2019). We analyzed the same samples for other

elements (Al, Ca, Cu, Fe, K, Mg, Mn, Na, P, Zn) using nitric acid digestion followed by inductively

coupled plasma-optical emission spectrometry (ICP-OES) using an Optima 7300CV (PerkinElmer,

Waltham, MA, USA). We removed five data points from the ICP-OES output as extreme outliers that

were not representative of the range of concentrations of their respective elements across our samples. We

also set values below the detection limit, which sometimes returned as negative concentrations, to zero.

We also analyzed the ground-leaf samples for carbon fractions using an ANKOM 2000 Fiber Analyzer

(Ankom Technology, Macedon, New York, USA) to perform a sequence of digestions (Ayotte &

Laliberté 2019). The first digestion in a heated neutral detergent washes off soluble cell contents. The

second digestion in a heated acidic detergent removes hemicellulose and bound proteins—which, for

simplicity, we just refer to as hemicellulose. The third digestion in 70% sulfuric acid removes cellulose,

leaving behind lignin and recalcitrant compounds. We used the change in each bag's mass between stages

to determine the mass of each fraction. We ashed all samples and used ash content as a correction factor

to calculate the ash-free lignin fraction.

1007

External validation dataset

From August to November 2018, we collected 333 samples for an external validation dataset at Cedar

Creek Ecosystem Science Reserve (East Bethel, MN, USA). Most were from the Forests and Biodiversity

(FAB) experiment (Grossman et al. 2017), the Biodiversity in Willows and Poplars (BiWaP) experiment

(Grossman & Cavender-Bares 2019), and the Big Biodiversity (BioDIV or e120) experiment (Tilman et

al. 1997), although some were from other parts of the site.

1014

Each sample included multiple leaves from the same individual, taken when possible from a nearby

position on the plant. Unlike in the CABO dataset, we collected samples from any part of an individual,

not just sunlit leaves. From each sample, we designated one leaf for fresh mass to be measured

immediately following collection; this leaf ('fresh-mass subsample') was stored separately and its spectra

were labeled to keep track of its identity. (For conifers or other plants with small or narrow leaves, we

often included multiple leaves in this fresh mass sample.) We took and flash-froze 3-5 hole punches from

another leaf (or group of leaves) for pigment analyses ('pigment subsample'; pigment data not shown);

this leaf and its spectra were also labeled separately. We collected more leaves to have enough tissue for a

bulk chemical sample, but did not keep track of them individually.

1024

In the field, we measured multiple fresh-leaf spectra per sample (not shown here) using a PSR+ 3500

spectroradiometer with a leaf clip foreoptic (Spectral Evolution, Haverhill, MA) and placed the leaves in

sealed plastic bags with a damp paper towel. These bags were kept in coolers until the end of the

sampling day. We weighed the fresh-mass subsample immediately upon return to the lab and kept all

leaves in a refrigerator until we scanned the fresh-mass and pigment subsamples within the next few days.

After scanning, we placed all leaves in flat paper bags and pressed them for several hours in a plant press

before removing them and placing them in a drying shed at 40 °C for at least one week. This deviates

from most standard herbarium protocols, including the one used in the CABO project, where leaves are

kept in the press while drying. After we dried the leaves, we weighed the fresh-mass and pigment

subsamples. Certain species, especially *Populus tremuloides*, seemed especially prone to discoloration

during sample preparation.

1036

In May 2020, we measured pressed-leaf spectra using a PSR+ 3500 spectroradiometer with a leaf clip

foreoptic. We did not use a specific protocol for small or narrow leaves, but aimed to arrange leaves in a

way that minimized overlaps or gaps. We removed spectra with especially low (<40%) or high (>80%)

peak reflectance, and any with major visible gaps in the sensor overlap regions. We measured three

spectra per subsample except in cases where getting high-quality spectra proved especially difficult. We

then removed the petioles (when applicable) and put leaves in scintillation vials with ball bearings to

grind the tissue in a paint shaker, combining all subsamples from a given sample. We then dried samples

again and measured C and N with an elemental analyzer (Costech ECS 4010 Analyzer, Valencia,

California, USA).

1046

1047   For estimating LMA, LDMC, and EWT, we only used the spectra from the fresh-mass subsample. For C

1048   and N, we used all spectra from a given sample, ignoring the subsample labels. As among the CABO

1049   pressed-leaf spectra, we first matched sensors, then averaged spectra within the relevant sample or

1050   subsample, and lastly trimmed them to 400-2400 nm.

1051

1052   We applied the ensemble of models resulting from our 100× jackknife analysis on the pressed-leaf CABO

1053   data and applied them to the Cedar Creek data. We also applied the model ensemble from the pressed-leaf

1054   restricted-range (1300-2500 nm) CABO data to the restricted-range Cedar Creek data. We calculated

1055   summary statistics and plotted results as in the internal validation.

1056   **Supplementary references**

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 Supplementary tables

1108   Table S1: Summary statistics of PLSR model internal validation for area-based chemical traits from full-

1109   range fresh- and pressed-leaf models. %RMSE is calculated as RMSE divided by the 2.5% trimmed range

1110   of measured values within the validation subset.

Trait	Fresh				Pressed			
	No. comps	R^2	RMSE ($\times 10^6$)	%RMSE	No. comps	R^2	RMSE ($\times 10^6$)	%RMSE
Solubles (g cm^{-2})	9	0.948	614	5.88	8	0.915	791	7.56
Hemicellulose (g cm^{-2})	17	0.688	258	15.8	11	0.754	228	14.0
Cellulose (g cm^{-2})	21	0.858	158	10.3	15	0.902	133	8.58
Lignin (g cm^{-2})	13	0.898	167	8.17	16	0.916	151	7.41
C (g cm^{-2})	16	0.969	349	4.48	10	0.942	475	6.07
N (g cm^{-2})	19	0.753	20.7	13.9	14	0.761	19.9	13.3
Chl a (g cm^{-2})	6	0.588	6.93	17.2	7	0.570	7.00	17.3
Chl b (g cm^{-2})	6	0.535	2.56	18.6	6	0.502	2.63	19.0
Carotenoids (g cm^{-2})	9	0.562	1.36	17.6	9	0.477	1.48	19.0
Al (g cm^{-2})	16	0.281	0.362	31.3	7	0.255	0.372	31.9
Ca (g cm^{-2})	15	0.529	39.6	18.3	17	0.75	29.1	13.4
Cu (g cm^{-2})	5	0.273	0.0302	21.6	8	0.132	0.0317	31.6
Fe (g cm^{-2})	2	0.040	0.265	25.3	7	0.080	0.261	24.9
K (g cm^{-2})	9	0.524	18.1	17.2	18	0.553	16.0	16.6
Mg (g cm^{-2})	7	0.564	6.72	16.8	11	0.707	5.50	13.7
Mn (g cm^{-2})	4	0.162	1.44	23.5	15	0.402	1.23	20.1
Na (g cm^{-2})	5	0.701	6.06	15.2	11	0.767	5.46	13.7
P (g cm^{-2})	3	0.338	8.20	24.1	13	0.553	6.59	19.2
Zn (g cm^{-2})	10	0.336	0.495	22.6	14	0.480	0.44	20.0

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1112   Table S2: Summary statistics of PLSR model internal validation from pressed-leaf spectra restricted to

1113   1300-2500 nm.

Trait	No. comps	R^2	RMSE	%RMSE
LMA (kg m^{-2})	8	0.922	0.0105	7.08
LDMC (mg g^{-1})	11	0.703	36.9	12.4
EWT (mm)	11	0.781	0.0194	13.3
Solubles (%)	11	0.759	3.88	12.5
Hemicellulose (%)	10	0.725	2.51	13.9
Cellulose (%)	10	0.783	1.44	12.6
Lignin (%)	11	0.670	1.83	15.9
C (%)	11	0.864	0.825	8.89
N (%)	11	0.741	0.303	13.9

Chl *a* (mg g^{-1})	7	0.631	1.26	16.4
Chl *b* (mg g^{-1})	7	0.570	0.475	17.0
Carotenoids (mg g^{-1})	7	0.652	0.244	16.5
Al (mg g^{-1})	6	0.231	0.0452	24.5
Ca (mg g^{-1})	11	0.743	3.30	13.4
Cu (mg g^{-1})	10	0.354	0.00490	22.3
Fe (mg g^{-1})	3	0.135	0.0407	22.2
K (mg g^{-1})	12	0.548	3.13	20.2
Mg (mg g^{-1})	8	0.538	0.712	19.6
Mn (mg g^{-1})	9	0.327	0.171	20.8
Na (mg g^{-1})	11	0.461	0.519	22.1
P (mg g^{-1})	11	0.511	0.809	16.2
Zn (mg g^{-1})	10	0.428	0.0624	23.7

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1115   Table S3: Summary statistics for external validation of pressed-leaf restricted-range PLSR models. The

1116   models were trained on CABO data (see Table S2) and applied to a dataset collected at Cedar Creek.

	Including conifers			Excluding conifers		
Trait	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE
LMA (kg m^{-2})	0.857	0.0195	10.8	0.686	0.0127	15.8
LDMC (mg g^{-1})	0.427	54.7	19.3	0.494	55.7	19.3
EWT (mm)	0.679	0.0436	17.6	0.629	0.0315	17.4
C (%)	0.017	4.46	25.5	0.026	4.34	27.9
N (%)	0.665	0.615	32.1	0.661	0.712	27.9

1117

118 **Supplementary figures**

119 Fig. S1: Two researchers (A. Scott and I. Clark) demonstrate a typical setup for measuring pressed-leaf spectra with a

120 PSR+ 3500 spectroradiometer and leaf clip on unmounted pressed specimens. The specimen shown here was not part of

121 this study.

Fig. S2: Scans of specimens that exemplify scores 0 and 1 on our discoloration scale. Specimens evaluated at 0 (A and B)

showed no visible discoloration, as would be indicated by loss of green color (e.g. browning or blackening). Specimens

evaluated at 1 (C and D) showed <10% discoloration throughout the leaf, or in some cases a more subtle, pervasive

silvery or matte white appearance to the leaf surface in areas that remained otherwise green. On close inspection, these

two samples both show some browning in a small portion of the total leaf area. Note that we do not consider spots or

marks due to pathogen damage that were present at sampling time to be a form of discoloration; we also exclude some

immature anthocyanic leaves as in (A). When in doubt, we examined scans of fresh leaves taken shortly after they were

collected in the field. (Species: A, *Betula populifolia* Marshall; B, *Acer saccharinum* L.; C, *Acer rubrum* L.; D, *Diervilla*

lonicera Miller.)

All photos in Figs. S2-4 depict specimens included in the study, but were taken in March 2022, nearly three years

after we took pressed-leaf spectral measurements. Due to continuing discoloration, their assigned scores as part of this set

of example images may be larger than they were at the time we measured their spectra; the latter are what we used for

data analysis.

Fig. S3: Scans of specimens that exemplify scores 2 and 3 on our discoloration scale. Specimens evaluated at 2 (A and B)

showed 10-50% discoloration throughout the leaf. Here, A shows a number of dark streaks as well as more general

browning on some leaves, while B shows more general darkening on some leaves. Specimens evaluated at 3 (C and D)

showed 50-75% discoloration throughout the leaf. Here, C shows considerable browning across much of its leaf area, as

well as a silvery appearance across much of the leaf surface. D shows considerable blackening across much of its leaf

area. (Species: A, *Prunus virginiana* L.; B, *Viburnum lentago* L.; C, *Fagus grandifolia* Ehrhart; D, *Viburnum lantana* L.)

Fig. S4: Scans of specimens that exemplify score 4 on our discoloration scale. These specimens showed nearly complete

(>75%) discoloration, often making it difficult to take spectral measurements on a spot that retained green color. In some

cases (A), the leaves retained patches of green color, while in others (B) the discoloration was nearly complete. (Species:

A, *Populus grandidentata* Michaux; B, *Populus deltoides* W. Bartram ex Marshall.)

160   Fig. S5: (A) The wavelength-wise mean reflectance spectrum of leaves at each discoloration score. In general, more

161   discolored specimens have lower visible and NIR reflectance and a blunted red edge. This representation does not

162   separate changes in spectra due to discoloration from aspects of spectral variation that might correlate with predisposition

163   to discoloration. (B) A histogram of the number of pressed specimens evaluated at each discoloration stage at the time of

164   spectral measurements.

170   Fig. S6: Internal validation results for predictions of LMA, LDMC, and EWT from fresh- (left), pressed- (middle), and

171   ground-leaf (right) spectra. In each panel, each functional group has a separate ordinary least-squares regression line

172   overlaid on top of the thick dashed 1:1 line. The error bars for each data point are 95% confidence intervals calculated

173   from the distribution of predictions based on the ensemble of 100 models produced from jackknife analyses.

187   Fig. S8: Internal validation results for predictions of soluble cell contents, hemicellulose, cellulose, and lignin from fresh-

188   (left), pressed- (middle), and ground-leaf (right) spectra, displayed as in Fig. S6.

190   Fig. S9: Internal validation results for predictions of chlorophyll *a*, chlorophyll *b*, and total carotenoids from fresh- (left),

191   pressed- (middle), and ground-leaf (right) spectra, displayed as in Fig. S6.

206 Fig. S11: Internal validation results for predictions of Fe, K, and Mg from fresh- (left), pressed- (middle), and ground-leaf

207 (right) spectra, displayed as in Fig. S6.

214   Fig. S12: Internal validation results for predictions of Mn, Na, and Zn from fresh- (left), pressed- (middle), and ground-

215   leaf (right) spectra, displayed as in Fig. S6.

222 Fig. S13: The distributions of fresh-leaf spectral model performance statistics for each trait based on 100 jackknife

223 resamples from the calibration data set. The red dots show the R^2 and %RMSE from applying the ensemble of models to

224 the internal validation subset. Abbreviations: sol = solubles, hemi = hemicellulose, cell = cellulose, lign = lignin, car =

225 total carotenoids.

  Fig. S14: The distributions of pressed-leaf spectral model performance statistics for each trait based on 100 jackknife

  resamples from the calibration data set. The red dots show the R^2 and %RMSE from applying the ensemble of models to

  the internal validation subset. Abbreviations: sol = solubles, hemi = hemicellulose, cell = cellulose, lign = lignin, car =

  total carotenoids.

246 Fig. S15: The distributions of ground-leaf spectral model performance statistics for each trait based on 100 jackknife

247 resamples from the calibration data set. The red dots show the R^2 and %RMSE from applying the ensemble of models to

248 the internal validation subset. Abbreviations: sol = solubles, hemi = hemicellulose, cell = cellulose, lign = lignin, car =

249 total carotenoids.

Fig. S16: The variable importance of projection (VIP) metric calculated for all traits from fresh-leaf spectra.

Abbreviations: sol = solubles, hemi = hemicellulose, cell = cellulose, lign = lignin, car = total carotenoids. The dashed

horizontal line at 0.8 represents a heuristic threshold for importance suggested by Burnett et al. (2021).

Fig. S17: The variable importance of projection (VIP) metric calculated for all traits from pressed-leaf spectra.

Abbreviations: sol = solubles, hemi = hemicellulose, cell = cellulose, lign = lignin, car = total carotenoids. The dashed

horizontal line at 0.8 represents a heuristic threshold for importance suggested by Burnett et al. (2021).

Fig. S18: The variable importance of projection (VIP) metric calculated for all traits from ground-leaf spectra.

Abbreviations: sol = solubles, hemi = hemicellulose, cell = cellulose, lign = lignin, car = total carotenoids. The dashed

horizontal line at 0.8 represents a heuristic threshold for importance suggested by Burnett et al. (2021).

Fig. S19: External validation results for predictions of various traits in Cedar Creek data based on models trained on

CABO data, restricted to 1300-2500 nm. Each panel is displayed as in Figs. S6-12.

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