Predicting leaf traits across functional groups using reflectance spectroscopy

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Summary

- Plant ecologists use functional traits to describe how plants respond to and influence their environment. Reflectance spectroscopy can provide rapid, non-destructive estimates of leaf traits, but it remains unclear whether general trait-spectra models can yield accurate estimates across functional groups and ecosystems.

- We measured leaf spectra and 22 structural and chemical traits for nearly 2000 samples from 104 species. These samples span a large share of known trait variation and represent several functional groups and ecosystems. We used partial least-squares regression (PLSR) to build empirical models for estimating traits from spectra.

- Within the dataset, our PLSR models predicted traits like leaf mass per area (LMA) and leaf dry matter content (LDMC) with high accuracy (R^2>0.85; %RMSE<10). Models for most chemical traits, including pigments, carbon fractions, and major nutrients, showed intermediate accuracy (R^2=0.55-0.85; %RMSE=12.7-19.1). Micronutrients such as Cu and Fe showed the poorest accuracy. In validation on external datasets, models for traits like LMA and LDMC performed relatively well, while carbon fractions showed steep declines in accuracy.

31   • We provide models that produce fast, reliable estimates of several widely used functional traits

32   from leaf reflectance spectra. Our results reinforce the potential uses of spectroscopy in

33   monitoring plant function around the world.

35   Keywords: foliar chemistry; functional traits; leaf economics; partial least-squares regression; reflectance

36   spectroscopy

Introduction

Plant functional ecology relies on the measurement of traits that influence how plants interact with their abiotic and biotic environment. Such traits can be used to interpret or predict outcomes at the community- or ecosystem-scale (Funk et al. 2017), ranging from species sorting across environmental gradients (Kunstler et al. 2016) to rates of carbon and nutrient cycling (Cornwell et al. 2008; Ollinger et al. 2008). During the last 20 years, ecologists have developed standardized trait protocols and databases to enable broad comparisons of plant functional strategies across species (Perez-Harguindeguy et al 2013; Kattge et al. 2020). Nevertheless, because trait measurement campaigns can be costly and challenging, trait databases remain sparse and much of the world's plant diversity is poorly represented (Kattge et al. 2020). The resources needed to carry out intensive trait measurement campaigns using conventional methods has motivated the search for other approaches that could yield fast, reliable trait estimates.

One such approach is to predict plant traits from reflectance spectra, which report the fraction of light reflected from a surface across a range of wavelengths. For vegetation, the range most often studied is around 350-2500 nm, which includes >97% of solar radiation (American Society for Testing and Materials 2020). Because the structure and chemistry of plant tissues determine their optical properties, many traits can be estimated from reflectance spectra measured at leaf or canopy scales (Jacquemoud & Ustin 2019). At the leaf level, linking reflectance spectra and functional traits provides a fast way to estimate large numbers of leaf traits from individual plants. At the canopy level, it enables researchers to map traits using remote sensing, which may enable monitoring of plant community structure (Durán et al. 2019; Zheng et al. 2021) or biogeochemical cycles (Wessman et al. 1988; Chadwick & Asner 2018) over entire landscapes. Understanding leaf spectral variation is essential for scaling up to the canopy level (Asner et al. 2011), and leaf-level spectral estimates of traits are part of many workflows for canopy trait mapping (Wang et al. 2020). These uses of leaf spectra make it important to understand their functional drivers (Kothari & Schweiger 2022).

Methods to predict traits from leaf or canopy spectra can be summarized into three groups: (1) simple

indices calculated from reflectance at a few wavelength bands (e.g. Sims & Gamon 2002); (2) physics-

based radiative transfer models (Jacquemoud et al. 2009; Féret et al. 2017); and (3) multivariate empirical

methods that use large portions of the spectrum to build trait prediction models. Among this last group of

methods, partial least-squares regression (PLSR) is the most common, although others like stepwise

regression and (more recently) deep learning approaches like convolutional neural networks are also used.

These methods are less mechanistic but offer the flexibility often needed to predict traits associated with

complex optical properties (Curran 1989). For example, predicting leaf nitrogen concentrations using

radiative transfer models has been a persistent challenge because leaves contain nitrogen in many forms,

including proteins (e.g. RuBisCO) and pigments (Wang et al. 2018; Féret et al. 2021). Moreover, the

absorption features of nitrogen-containing bonds often overlap with others, especially after being

broadened by multiple scattering (Curran 1989). As a result, leaf nitrogen is not identifiable with unique,

distinct absorption features, and is often best predicted using empirical approaches like PLSR (Asner et

al. 2011; Yan et al. 2021). Researchers have likewise built PLSR models to estimate a wide variety of

foliar traits from fresh-leaf spectra, including pigments (Yang et al. 2016), non-structural carbohydrates

(Ely et al. 2019), and condensed tannins (Couture et al. 2016).

Estimating traits from leaf spectra has several advantages that may compensate for the sparsity of trait

datasets. Compared to measuring a large number of traits, measuring spectra is fast, has low marginal

cost, can be non-destructive, and requires relatively little training. But since PLSR is purely empirical

rather than mechanistic, a model calibrated using one dataset may yield inaccurate or biased predictions

on another dataset if the association between traits and optical properties varies among contexts (Yang et

al. 2016; Wang et al. 2020). The problem may be most severe when the model relies on absorption

features that are not causally associated with the target trait, but rather with other covarying traits

("constellation effects" *sensu* Chadwick & Asner 2016; Nunes et al. 2017). This situation may occur most

often when the target trait lacks major, well-defined absorption features, like concentrations of many

elements (Nunes et al. 2017). If trait covariance patterns vary among datasets, overfitting to the training

data could cause such models to produce inaccurate trait estimates from new spectral data (Kothari &

Schweiger 2022). Empirical models may be especially fallible beyond the range of data used to train them

(Schweiger 2020; Burnett et al. 2021).

To enable the widespread use of spectral models for trait prediction, it is essential to build general models

that encompass the broadest possible range of spectral and trait diversity (Serbin et al. 2019; Wang et al.

2020). Such models could allow researchers to estimate plant traits from spectra in their study systems

without going through the laborious process of building their own models. Asner et al. (2011) invoked

this goal in developing spectral models for many leaf traits in humid tropical forest canopies around the

world. In contrast, Serbin et al. (2019) focused on a single trait (leaf mass per area; LMA) but included

several functional groups and biomes from the tropics to the tundra. We sought to push this goal further

by including both a large number of leaf traits and a broad range of ecosystems and functional groups,

allowing us to compare which traits are the most practical targets for estimation by general spectral

models.

We compiled foliar traits and spectra measured using standardized protocols on nearly 2000 samples that

include trees, shrubs, graminoids, and other herbs from a range of ecosystems (Table 1). We included

structural and biochemical traits that relate to several aspects of leaf function, including nutrient

economics, water relations, photosynthesis and photoprotection, and structural defense. We focused on

building models to predict these traits from reflectance spectra, but we also compared the performance of

reflectance, transmittance, and absorptance spectra. Among these properties, reflectance is the simplest

and most common to measure because it bears the most relevance for interpreting remotely sensed canopy

reflectance and does not require an integrating sphere. But since the content of leaf chemical constituents

most directly alters the amount of light absorbed (Jacquemoud & Ustin 2019), we conjectured that

absorptance spectra might be better for estimating traits, and aimed to determine whether any potential

benefit transmittance or absorptance spectra provide in trait estimation outweighs their greater costs in

time and equipment.

The performance of statistical models is often evaluated by splitting a dataset at random into calibration

and validation subsets. This procedure represents a best-case scenario, since the data are all collected with

the same protocols and, based on the mathematics of random sampling, the two subsets must have similar

distributions of predictor and response variables. Therefore, we validated our models on both a random

subset of our dataset (internal validation) and on independent datasets (external validation)—the latter

perhaps representing a more realistic test of how they would perform in practical applications.

Concerns about the generality of spectral trait models also raise the question: What factors determine

whether a trait model yields accurate predictions on an external dataset? Many existing models to predict

leaf traits were built using samples belonging to a single functional group or ecosystem, leaving it an open

question how they would transfer to new settings. One might expect that models would show poorer

performance as the calibration samples (used to build the model) and the external dataset's samples

become less functionally similar (Yang et al. 2016). But if the models show consistently strong

performance, it would suggest that associations between traits and optical properties are robust, not

contingent on the particular samples included. To examine model transferability, we divided our dataset

by functional group and examined model accuracy using each functional group in turn as the validation

dataset for models calibrated using the remaining functional groups. We posed two hypotheses about

these analyses: (1) For a given trait, predictions are most accurate when the distributions for that trait are

most similar between the calibration and validation datasets; and (2) traits whose predictions are most

accurate across the full dataset will also yield the most accurate predictions (on average) when

transferring models onto left-out functional groups. We aimed to shed light on the factors that influence a

model's accuracy across a range of settings.

Methods

Sampling

We assembled a database (n = 1971; species = 104) of leaf spectra and functional traits measured as part

of the Canadian Airborne Biodiversity Observatory (CABO; www.caboscience.org). We aggregated data

from ten individual projects carried out during 2018 and 2019 at sites across temperate Canada and one

site in southwestern Australia (Table 1). The sites span a wide range of edaphic properties, from nutrient-

poor bogs and phosphorus-limited woodlands to highly fertile post-agricultural and wetland sites. Each

sample comprised a large, homogeneous group of sunlit leaves, which we divided up for spectral and trait

measurements. We classified samples into seven functional groups: broadleaf trees, coniferous trees,

ferns, forbs, graminoids, shrubs, and vines (Table 1). Further details on sampling protocols and functional

groups are found in the Supplementary Materials.

Leaf spectral and trait measurements

We measured directional-hemispherical reflectance and transmittance spectra (350-2500 nm) using an

HR-1024i spectroradiometer equipped with a DC-R/T integrating sphere from Spectra Vista Corporation

(Poughkeepsie, NY, USA). For each sample, we measured spectra from the adaxial surface of six leaves

or (for small or narrow leaves) leaf arrays. We resampled spectra to 1 nm resolution, averaged spectra

from the same sample, and trimmed them to 400-2400 nm by removing the ends, which are often noisy.

Finally, we calculated absorptance at each wavelength by subtracting the sum of reflectance and

transmittance from 1. Further details on spectral measurement and processing can be found in the

Supplementary Materials.

We measured the following traits on each leaf sample: Leaf mass per area (LMA), leaf dry matter content

(LDMC), equivalent water thickness (EWT), and concentrations of soluble cell components,

hemicellulose, cellulose, lignin, chlorophyll *a*, chlorophyll *b*, total carotenoids, and a number of elements

(Al, C, Ca, Cu, Fe, K, Mg, Mn, N, Na, P, and Zn). Elements other than C and N were only available for a

few projects that account for about one-third ($n = 678$) of the total samples. Further details on trait

measurement protocols can be found in the Supplementary Materials. We compared our trait distributions

to the TRY trait database to evaluate how well they span the global range of trait variation. Although it

has known geographic and taxonomic biases, TRY is the most comprehensive database of its kind (Kattge

et al. 2020). We also performed a principal components analysis (PCA) on the scaled and centered trait

data to visualize patterns of trait variation among our samples.

Within a dataset, chemical traits may vary more in proportion to the mass or the area of the leaf (Osnas et

al. 2013). When a trait is distributed mainly proportional to leaf area, mass-normalization induces a

negative correlation with LMA. Since interspecific variation in pigments tends to be area-proportional,

Kattenborn et al. (2019) used this principle, among others, to argue that pigment content (per unit area) is

a better target than concentration (per unit mass) for remote sensing-based estimation. We applied our

PLSR modeling approach to both mass- and area-based traits, but we focus on mass-based models since

conventional protocols measure most chemical traits on a mass basis. Chemical traits should be assumed

here to be mass-based when left unspecified. We also calculated the mass-proportionality of each

chemical trait in our dataset as the ordinary least squares slope between the log-transformed area-

normalized trait and log-transformed LMA (Osnas et al. 2013).

PLSR modeling

Trait estimation

Due to the high dimensionality and multicollinearity of spectral data, we used a PLSR modeling approach

to build trait estimation models. PLSR projects the high-dimensional input dataset (here, spectra) onto a

smaller number of latent components constructed to maximize their covariance with the variable(s) to be

predicted (traits). We built separate models using reflectance, transmittance, and absorptance as

predictors, in each case using the whole trimmed spectrum (400-2400 nm). We did not transform the

distributions of any trait to reduce skewness, as is sometimes recommended (Burnett et al. 2021); in

preliminary tests, such transformations did not yield any consistent improvement in model performance.

We performed all analyses in *R* version 3.6.3 (R Core Team 2020).

We calibrated and validated our main set of models ('primary models') largely following the example of

Serbin et al. (2014) and Burnett et al. (2021) using *R* package *pls* v. 2.7.1 (Mevik et al. 2019). First, we

divided the full dataset into 75% calibration and 25% validation sets at random, stratified by functional

group (see Table 1). We fit a preliminary PLSR model for each trait on the calibration subset, using 10-

fold cross-validation to select the number of components to use in our final models. We chose the

smallest number of components that brought the cross-validation root mean squared error of prediction

(RMSEP) within one standard deviation of the global minimum—a distinct number (4-27) for each trait

and type of spectrum. We calculated the variable influence on projection (VIP) metric (Wold et al. 2001)

to determine which spectral regions were most important for predicting each trait.

Next, we fit our final models through a jackknife resampling procedure in which we further divided the

75% calibration dataset 100 times into random 70% testing and 30% training subsets. For each iteration,

we built a model from the 70% testing dataset using the previously chosen number of components and

applied it to the 30% training dataset. This procedure allowed us to get a distribution of summary

statistics that shows how model performance varies based on random changes in the training and testing

datasets. The primary model comprises the 100 sets of coefficients for each trait—one from each

resampling iteration—which we applied to the 25% validation dataset to get a distribution of estimates for

each sample. We compared the measured values to the mean estimates for each trait to produce summary

statistics (R^2, RMSE, %RMSE) for internal validations. In all cases, we report RMSE relative to the 1:1

line. For robustness to outliers, we defined %RMSE as the RMSE divided by the range of the inner 95%

of trait values, in contrast to a more common definition (Burnett et al. 2021) that uses the full range.

External validation

We compiled data from three sources as external validation datasets for our primary models. The first two

are LOPEX (Hosgood et al. 1994) and ANGERS (Féret et al. 2008), which have been widely used to

validate methods for plant trait estimation from reflectance spectra. Between them, these datasets include

a very wide variety of temperate plants, many cultivated. The third source is an expanded version of the

Dessain dataset previously described in Kothari et al. (2022), which includes herbs and trees from

southern Québec. Unlike LOPEX and ANGERS, most of the Dessain dataset's trait data were measured

using the same protocols as the core CABO datasets. All three datasets were measured using integrating

spheres and include similar spectral ranges, but the instruments and their specifications differ from each

other and the core datasets we used to train the models, which can cause slight but important differences

in measured reflectance (Lukeš et al. 2017; Hovi et al. 2018).

Since we did not measure transmittance spectra for the Dessain project, we focused on reflectance spectra

from the validation datasets. We applied our sets of reflectance-based coefficients from resampling

analyses (100× for each trait) to the spectra in each dataset and took the mean of the resulting 100

estimates for each sample for comparison to the measured traits. Not all traits were available for each

dataset. Further details on our handling of external validation data are in the Supplementary Materials.

Model transferability among functional groups

Besides assessing the viability of our primary models, we also aimed to test whether and when models

could estimate traits accurately on a group of species functionally distinct from those used in calibration

(Hypotheses 1 and 2). To this end, we built another set of models: we selected a single functional group at

a time as a validation dataset and used the reflectance spectra of the remaining functional groups to

calibrate a PLSR model for each trait—using 10-fold cross-validation and component selection as above,

but omitting the jackknife procedure. Finally, we applied the model to the validation functional group. We

repeated this procedure using each functional group in turn as the validation dataset, besides ferns and

vines because of their low numbers of samples. Among elements, we only considered C and N because

other elements were only available for certain projects and we could not guarantee adequate coverage of

all functional groups. We quantified performance using R^2, RMSE, and %RMSE—calculating %RMSE

using the inner 95% trait range of the full dataset in the denominator rather than the inner 95% trait range

of only the validation functional group. This metric, which we call %RMSE$_{full}$ for clarity, avoids apparent

reductions in model performance due to a narrow trait range in the validation functional group.

We conjectured that for a given trait, model performance would decline as the calibration and validation

datasets became more dissimilar in their trait distributions (Hypothesis 1). We quantified dissimilarity

using the Hellinger distance between the calibration and validation datasets' distributions for each trait, as

estimated in *R* package *statip v. 0.2.3* (Poncet et al. 2019). Here we focused on %RMSE$_{full}$ because it is

normalized to the scale of the data (unlike RMSE) and incorporates bias (unlike R^2). We tested the

influence of trait dissimilarity on %RMSE$_{full}$ with analysis of covariance (ANCOVA) using Hellinger

distance (continuous), the trait identity (categorical), and their interaction as predictors.

Results

Trait and spectral variability

The ranges for most traits in our dataset spanned most of the trait ranges in TRY. Many traits, including

LMA, EWT, and all elements other than C and N, had distributions with strong positive skew in both

TRY and CABO data, resulting in a long upper tail of high values (Fig. 1). For certain elements (Al, Cu,

Ca, Fe, K, Mg), the CABO dataset had less representation of values along the upper tail than TRY. For

other elements (Na, P, Zn), as well as LDMC, N, and pigments, the CABO data had greater representation

of high values. For yet other traits (LMA and C), the CABO data had a somewhat narrower range towards

either extreme.

Within the dataset, functional groups often differed in their trait values, as shown by a principal

components analysis (PCA) of the centered and scaled trait data (excluding elements other than C and N;

Fig. S1). The first two PCs of the trait data explained 65.7% of total variance. Graminoids were

distinguished by high concentrations of hemicellulose and cellulose and low concentrations of soluble

carbon, lignin, and total C. Leaf mass per area and LDMC were negatively correlated with N and pigment

concentrations; conifers (besides deciduous *Larix laricina* (Du Roi) K. Koch) and the evergreen broadleaf

tree *Agonis flexuosa* (Willd.) Sweet had particularly high LMA and LDMC, while other functional groups

each contained significant variation along this axis. High values of EWT (>0.3 mm) were dominated by

evergreen conifers and graminoids adapted to wetlands, such as *Typha angustifolia* L. and *Phragmites*

australis (Cav.) Trin. ex Steud. *A. flexuosa* represented most of the samples with the lowest

concentrations of P, K, and Fe and the highest concentrations of Na, while bog shrubs and sedges had

particularly low Ca concentrations. The large fraction of variance explained by major trait axes indicates

that certain clusters of traits covary strongly, often in ways that correspond to functional groups.

Our samples showed the typical shape of leaf spectra, including a sharp increase in reflectance and

transmittance ('red edge') between the visible and near-infrared (NIR; 700-1000 nm) ranges and

prominent water absorption bands in the short-wave infrared (SWIR; >1000 nm) range (Fig. 2; Fig. S2).

Compared to other functional groups, needleleaf conifers had low reflectance and transmittance and high

absorptance across the spectrum. Forbs also tended to have lower transmittance and higher absorptance in

the SWIR. In absolute terms, absorptance and transmittance spectra had greater variation than reflectance

spectra, especially in the SWIR.

PLSR modeling

Internal calibration and validation

The accuracy of trait estimates from reflectance spectra varied widely by trait (Table 2; Fig. 3-4; Fig. S2-

8). Our primary models estimated structural and water-related traits—LMA (R^2 = 0.892; %RMSE =

8.04), EWT (R^2 = 0.885; %RMSE = 8.46), and LDMC (R^2 = 0.867; %RMSE = 9.78)—with high

accuracy. Models for certain chemical traits such as carbon fractions (R^2 = 0.558-0.826; %RMSE = 12.7-

19.1), major elements like C, N, K, and Ca ($R^2 = 0.541$-0.747; %RMSE = 13.2-20.4), and pigments ($R^2 =$

0.644-0.689; %RMSE = 13.8-15.0) also showed intermediate-to-high accuracy. Finally, models for some

micronutrients like Cu ($R^2 = 0.295$; %RMSE = 25.9) or Mn ($R^2 = 0.254$; %RMSE = 22.8) performed very

poorly. For pigments, N, and some other elements, models underestimated trait values at the sparsely

represented upper tail of the measured range, while for C the models overestimated values at the sparse

lower tail (Fig. 3). Trait models with lower accuracy on the internal validation data also had greater

variability in accuracy across iterations in the resampling analysis—even when leaving out elements other

than C and N, which are only available for a few projects (Fig. S10-12).

The VIP metric revealed broad similarities in the regions of the spectrum that were important for

predicting different traits from reflectance spectra (Fig. S13-15). In particular, major features of the

visible region, including the green hump (centered at 550-560 nm) and especially the inflection point of

the red edge (710-720 nm) were important for predicting most traits besides EWT. These visible-range

importance peaks (especially at the red edge) were often higher for traits like Al, Zn, Mn, and

hemicellulose than for pigments, which are the dominant drivers of visible reflectance (Jacquemoud &

Ustin 2019). In general, VIP declined into the NIR but had several peaks in the SWIR, including

relatively distinct peaks for multiple traits at 1390 nm and 1880 nm, and weaker ones at 1720 nm, 2020

nm, 2150 nm, and 2290 nm. For EWT, LMA, Cu, Fe, and Na, VIP remained high throughout a larger

portion of the SWIR; uniquely, EWT had its global maximum VIP at 1390 nm.

In the dataset, most chemical traits were distributed mainly proportional to mass (Table S1). Exceptions

included all pigments and several macro- and micronutrients, which showed mass-proportionality less

than 0.5. We produced estimates of area-based chemical traits both by building models to estimate them

directly and by multiplying estimated mass-based traits by estimated LMA (Table S1; Fig. S16-19). For

most traits, estimating area-based traits directly yielded more accurate estimates; the difference was

greatest for pigments, Ca, Cu, N, P, which all have area-proportionality less than 0.6. However,

multiplying mass-based estimates by LMA estimates was slightly more accurate for many highly mass-

proportional traits, including Al, C, Na, and cellulose.

For most traits, transmittance and absorptance spectra yielded slight improvements in performance

relative to reflectance (Table S2; Fig. S3-12). Excluding traits that showed very poor performance in

general (average $R^2 < 0.35$), the greatest improvements between reflectance and either transmittance or

absorptance spectra (average Δ%RMSE > 1) occurred for LMA, LDMC, EWT, pigments, Ca, K, N, and

P. Using transmittance or absorptance spectra never worsened performance considerably (average

Δ%RMSE < -1).

External validation

We applied our reflectance-based primary models for each trait to three external validation datasets whose

spectral and trait measurement protocols differed from the CABO dataset. In general, models performed

less well on these independent datasets than the internal validation (Table 2; Fig. 6-7; Tables S4-5; Fig.

S20). The traits that had the highest R^2 between observed and predicted values in the internal validation

also had the highest R^2 in the external validation. As a result, structural and water-related traits (LMA,

LDMC, and EWT) had moderate-to-high R^2 in the external validation (0.589-0.842 averaged across

datasets). The RMSE for LDMC predictions was only slightly higher than in the internal validation (40.6-

46.4 vs. 32.9 mg g^{-1}). In contrast, both LMA and EWT were greatly overestimated for most samples, so

even though the R^2 was moderate-to-high, RMSE was also high (LMA: 0.0855-0.137 vs 0.0175 kg m^{-2};

EWT: 0.0395-0.0705 vs 0.0334 mm). Pigments, N, and particularly C and carbon fractions had

correspondingly lower prediction accuracy and varying amounts of bias. Many other macro- and

micronutrients (besides K) had very low prediction accuracy ($R^2 < 0.15$).

Model transferability among functional groups

For each trait, models generally performed worse when applied to a functional group left out from the

calibration dataset than when (as in our primary models) calibrated and applied to random selections of

samples (Fig. 5-6; Table S3). In particular, models calibrated using the other functional groups tended to

show poor performance when applied to graminoids or forbs. Carbon fractions tended to be estimated

with relatively low accuracy (high RMSE, low R^2); because these carbon fractions were predicted well

within the internal validation, this result weakened the relationship posited in Hypothesis 2 between

average model performance in internal validation and in functional group transferability analyses. Models

for carbon fractions applied to graminoids yielded particularly biased predictions, with %RMSE$_{full}$ > 50

for all but lignin. While graminoids were distinctively high in hemicellulose and cellulose and low in

solubles and lignin, the model predicted them to be much more similar to the other functional groups (Fig.

5). Models for LMA applied to conifers likewise showed %RMSE$_{full}$ > 50 because the distinctively high

values of evergreen (non-*Larix*) conifers were underestimated.

In our ANCOVA model, there were significant effects of trait ($F_{11,36}$ = 2.626; p = 0.014), Hellinger

distance ($F_{1,36}$ = 21.066; $p < 10^{-4}$), and their interaction ($F_{11,36}$ = 4.139; $p < 0.001$) on %RMSE$_{full}$. This

result led us to perform separate linear regressions between %RMSE$_{full}$ and Hellinger distance for each

trait (Fig. 6B). The relationship was significant and positive for LMA ($t(3)$ = 7.141; p = 0.006), but not

significant and highly variable in strength and direction among the other traits. Support for Hypothesis 1

was thus also weak and equivocal.

Discussion

We built PLSR models to predict widely used foliar traits from reflectance spectra across nearly 2000

samples of 104 species, including several functional groups and ecosystems. Our findings underscore that

leaf spectra integrate many aspects of leaf phenotypes in a single measurement (Cavender-Bares et al.

2017; Kothari & Schweiger 2022) and help explain why spectral variation can serve as a surrogate for

phenotypic variation (Schweiger et al. 2018; Frye et al. 2021).

Comparing PLSR model performance among traits

The performance of our primary models was highest for structural and water-related traits (LMA, LDMC,

and EWT), corroborating studies that have reported high accuracy using diverse modeling approaches

(Serbin et al. 2019; Guzmán Q. et al. 2021). Indeed, the contents of dry matter and water are strong

drivers of optical properties across much of the NIR and SWIR (Féret et al. 2008). Our models predicted

N, C, individual carbon fractions, and pigments with intermediate accuracy. These traits also have known

influences on leaf optical properties. Pigments control optical properties in the visible range (Jacquemoud

& Ustin 2019) and correlate with N. Cellulose, lignin, and N-containing compounds like proteins also

have bonds with known (if weak) absorption features, primarily in the NIR and SWIR (Curran 1989).

Models for elements at low concentrations often showed poor performance. Most of these elements have

no measurable absorption features in this range, so their concentrations can only be estimated indirectly,

often through symptoms of nutrient deficiency (Jacquemoud & Ustin 2019). In some cases, our models

could predict variation across, but not within, major biological groups. For example, Na could be

predicted with $R^2 = 0.519$, but mainly because Australian *A. flexuosa* has on average more than twice the

Na concentration of other species sampled. The model could have incorporated any optical features that

distinguish *A. flexuosa* from other specimens, whether or not those features truly result from Na

concentration. This kind of constellation effect may underlie the spectroscopic estimation of many

elements (Nunes et al. 2017; Kothari & Schweiger 2022).

Interpreting PLSR models

The VIP metric revealed strong similarities across traits in the major peaks of importance, including the

green hump, the red edge, and multiple features in the SWIR (e.g. 1390 and 1880 nm; Fig. S13-15). The

general pattern of high VIP along the green hump and at the red edge is shared by many traits across

studies (Yang et al. 2016; Ely et al. 2019; Yan et al. 2021; Kothari et al. 2022). In general, visible and red

edge reflectance are dominated by pigments and leaf structure (Richardson et al. 2002). Many traits show

isolated VIP peaks in the SWIR, which are harder to explain in terms of specific traits. Two peaks are in

off-center parts (1390 nm and 1880 nm) of major water absorption features, and the SWIR also contains

many shallow, overlapping absorption features of bonds in dry matter components, making it hard to

isolate the influence of any single one (Curran et al. 1989). In contrast to these distinct peaks, LMA and

EWT (as well as some elements) had high VIP throughout much of the SWIR, which might arise from the

strong and pervasive influence of LMA and EWT on SWIR reflectance.

Similarities in VIP may result from trait covariance (Fig. S1), which could allow models to leverage

constellation effects. Some cases are clear: the three pigment pools correlate tightly with each other ($R^2 =$

0.853-0.974) and with N ($R^2 = 0.457-0.517$), and their VIP patterns share an overall visual similarity.

Likewise, solubles, hemicellulose, and cellulose are closely linked ($R^2 = 0.643-0.834$) and have similar

VIP patterns. Other explanations are more elusive: for example, all four carbon fractions have high VIP

across the green hump and red edge, but correlate poorly ($R^2 = 0-0.107$) with pigments and LMA, which

account for most spectral variation in those regions. Such findings imply that strong covariance with a

few key traits cannot alone explain the striking similarities in VIP, nor the high accuracy of models for

traits like carbon fractions in the internal validation.

Alternate PLSR models

We focused on estimating chemical traits on a mass basis, but the remote sensing literature contains

powerful arguments for estimating certain constituents on an area basis. In particular, the optical

properties of leaves are driven more by the absolute quantity of a constituent than its quantity relative to

other constituents. This consideration should matter most for traits distributed in proportion to area

(Kattenborn et al. 2019). In our dataset, pigments and several nutrients were mostly area-proportional

(Table S1). As Kattenborn et al. (2019) suggested, estimating area-based traits directly was usually more

accurate than multiplying mass-based estimates by LMA estimates, especially for traits distributed in

proportion to area (Table S1). This consideration is important given that many research questions may

require traits expressed on an area basis.

Using transmittance or absorptance spectra instead of reflectance spectra increases accuracy for most

traits, but only slightly (Table S2). It may be relevant that the traits that show the greatest improvement

include most of the highly area-proportional traits. Given that transmittance or absorptance spectra

yielded little improvement for most traits, adding transmittance measurements may not aid much in trait

estimation, although they are still useful in endeavors like radiative transfer modeling (Spafford et al.

2021).

External validation and model transferability

Aside from our internal validation, we also considered model performance under less ideal scenarios: we

tested the primary models on external validation datasets, and built another set of models to be tested on a

functional group left out during their construction. When transferring models to left-out functional

groups, performance was usually worse (lower R^2, higher $RMSE_{full}$) than when models were evaluated on

a random internal validation dataset (Fig. 6; Table S3). This finding is consistent with previous studies

transferring models among species (Helsen et al. 2021) or sites (Yan et al. 2021). In general, trait models

performed worse when the trait distributions in the calibration and validation datasets were less similar,

but the strength and even the direction of the trend varied among traits (Fig. 6B). Thus, support for

Hypothesis 1 was equivocal. Predictions of LMA among conifers and carbon fractions among graminoids

were strongly biased and failed to capture the functional distinctiveness of these groups. These results

underscore that models must be built using samples whose diversity encompasses the models' expected

domain of application (Yang et al. 2016; Schweiger 2020; Beauchamp-Rioux 2022).

The external validation was a more severe test of model generality because the data were collected on

different species with different instruments using different trait measurement protocols. This kind of

scenario may better represent a realistic use case for such models. All traits showed declines in

performance from internal to external validation (Table 2). This result is expected and consistent with

previous findings: for example, Serbin et al. (2019)'s global model of LMA showed $R^2 = 0.89$ for internal

validation and 0.66 for external validation on LOPEX and ANGERS, as compared to $R^2 = 0.89$ (internal)

and 0.59 (external, averaged across datasets) here. These results further underscore the challenges that

arise when transferring models among settings and instruments.

The rank-order of R^2 across traits was broadly similar between the internal validation, external validation,

and model transferability analyses (Fig. 6A), which offers some general support for Hypothesis 2. Models

for LMA, LDMC, and EWT performed relatively well in all analyses, while models for most macro- and

micronutrients performed poorly in both internal and external validation (but were not included in model

transferability analyses). Pigments, N, C, and carbon fractions were generally intermediate. However, in

both the external validation and model transferability analyses, C and carbon fractions showed

particularly strong declines in R^2 compared to internal validation (Fig. 6A). Dry matter components like

cellulose and lignin are associated mainly with weak absorption features throughout the NIR and SWIR

(Curran et al. 1989). In contrast, traits without such strong declines—LMA, LDMC, EWT, N, and

pigments—have stronger influences on the spectrum, and many are part of mechanistic models like

PROSPECT (Féret et al. 2008). This contrast could imply that models are most transferable when their

target traits have strong absorption features. Many chemical traits may be estimated more reliably using

spectra of pressed or ground leaves, which by removing the water absorption features expose the subtler

features of dry matter components (Kothari et al. 2022).

Even though our estimates of LMA and EWT correlated well with measured values in the external

validation, they also had a strong bias towards overestimation, which could complicate their use in

practice. This bias may have resulted from varying measurement protocols, or from differences in

instrumentation that can have a subtle but pervasive influence on the measured spectra (Castro-Esau et al.

2006; Lukeš et al. 2017; Hovi et al. 2018). This bias could perhaps be corrected by measuring a subset of

trait values to determine how they correspond to model predictions, and then transforming the remaining

predictions accordingly. In the longer term, it might become easier to get accurate trait estimates using

existing models if techniques are developed to reconcile differences among instruments (Schweiger 2020;

Meireles et al. 2020). Deep learning approaches like convolutional neural networks may also address this

problem through their greater flexibility and capacity for transfer learning (Kattenborn et al. 2021).

Implications for global trait-spectra modeling

Our models draw from samples representing several functional groups and a wide range of temperate

ecosystems, including wetlands, grasslands, and closed-canopy forests (Table 1). However, these samples

represent just a portion of Earth's vast plant diversity—omitting, for example, tundra, drylands, and

tropical biomes. These biomes would surely contain combinations of optically important traits missing in

our dataset, which would need to be represented in a universal model that could be used in mapping

functional trait variation worldwide. We hope plant ecologists can use our data and models not only as a

tool to address ecological questions, but also as a foundation for building yet more general models.

Our results imply that some traits can consistently be estimated better than others. We can compare our

findings to Asner et al. (2011), who measured a similar set of traits on a vast number of samples from

canopy leaves in humid tropical forests. The two studies are broadly consistent despite representing

disparate subsets of the world's plant diversity. Both found that models for LMA and water content were

most accurate ($R^2 > 0.85$), followed by, in varying orders, C and N, pigments, and carbon fractions ($R^2 =$

0.55-0.85). In both studies, models for other elements were typically worse ($R^2 < 0.65$), but better for Ca

and K than many other elements. Both studies also found that transmittance spectra yielded slightly better

performance for most traits. The stability in rankings suggests a general hierarchy of traits that are the

most practical targets for reliable model-based estimation.

The emerging body of research showing that spectral models can yield fast, reliable estimates of many

leaf traits across functional groups and ecosystems could expand our ability to map and monitor plant

function around the world, bridging between conventional trait measurements and remote sensing. Our

findings clarify both the opportunities and challenges of using spectral trait estimates. Research has not

yet progressed to the point where plant ecologists can use existing empirical models to estimate traits in

their study systems without further validation. Aside from the issues of taxonomic and functional

representativeness, our external validation underscores the challenges of transferring models among

instruments, which might require new methodological developments to overcome. Even under ideal

conditions, some aspects of leaf function (e.g. elemental composition) may be hard to measure accurately

based on spectroscopy alone. We hope that the promise of rapid, large-scale monitoring of plant function

inspires the coordinated effort needed to understand and (when possible) overcome these challenges.

In sum, we built models to predict traits from reflectance spectra using nearly 2000 leaf samples from 104

species. Traits varied a great deal how well they could be estimated using spectral models. Our models for

structural and water-related traits were very accurate in internal validation and showed great promise in

external validation; in contrast, our models for many micronutrients performed very poorly. We suggest

that these patterns reflect a general hierarchy that emerges mainly from the varying degrees of influence

these traits have on the spectrum. Leaf reflectance spectra thus bear the imprint of many aspects of leaf

function—but not all to the same degree. Most importantly, we show that many traits can be estimated

accurately from spectra using general models that span multiple functional groups and ecosystems. This

finding represents a significant advance in leveraging spectra towards major goals in functional ecology.

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Author contributions

SK and EL designed the project with contributions from AB, NCC, MK, and MV. RBR, FB, ALC, AG, XGM, PWH, JP, AKS, and SDT collected and curated the spectral and trait data. SK analyzed the data, interpreted the results, and wrote the first draft with substantial contributions from EL. All authors contributed to further revisions of the paper.

Data availability

All fresh-leaf spectral data are available through the CABO data portal (https://data.caboscience.org/leaf). Upon publication, we will also upload all spectral data, as well as metadata and trait data, to the

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

552   Ecological Spectral Model Library (EcoSML, https://ecosml.org/). At that stage, we will update this

553   section accordingly. Analysis scripts are available as a repository on GitHub

554   (https://github.com/ShanKothari/CABO-trait-models).

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Fig. 1: Density plots comparing the distributions of various leaf structural and chemical traits in the

CABO dataset to the TRY database (Kattge et al. 2020). We filtered TRY data to include only values with

an error risk below 3. EWT is only shown for CABO because it is not a trait in TRY. Elements besides C

and N are only available for the Beauchamp-Rioux, Boucherville 2018, Girard, and Warren projects.

Fig. 2: The median leaf-level reflectance at each wavelength across the spectrum, separated by functional

group. We omit ferns and vines, which are only represented by one species each. The dashed line shows

the coefficient of variation across all samples.

Fig. 3: Plots of observations against reflectance-based PLSR predictions among internal validation data

for various leaf structural and chemical traits. The black dashed line in each panel is the 1:1 line. Colored

lines represent best-fit lines from OLS regression for each functional group. Error bars around each point

represent 95% confidence intervals based on the ensemble of models produced in the 100× jackknife

analysis.

Fig. 4: Plots of observations against reflectance-based PLSR predictions among internal validation data

for elements other than C and N. These data are only available for the Beauchamp-Rioux, Boucherville

2018, Girard, and Warren projects. The black dashed line in each panel is the 1:1 line. Colored lines

represent best-fit lines from OLS regression for each functional group. Error bars around each point

represent 95% confidence intervals based on the ensemble of models produced in the 100× jackknife

analysis.

Fig. 5: Plots of observations against reflectance-based PLSR predictions from model transferability

analyses for various leaf chemical and structural traits. Here, we estimated traits for samples of each

functional group from models calibrated on all the remaining functional groups—within panels,

predictions for each functional group come from distinct models. The black dashed line in each panel is

the 1:1 line. Colored lines represent best-fit lines from OLS regression for each functional group.

Fig. 6: (A) A summary of model R^2 for each trait. Colors indicate whether the R^2 describes performance

of models on: the random 25% internal validation dataset (black dots); specific functional groups used as

validation data in model transferability analyses; or external validation datasets. (B) The relationship

between %RMSE and Hellinger distance between calibration and validation trait distributions in model

transferability analyses. Functional group labels are shared between panels and refer to the validation

functional group.

B

833   Fig. 7: Plots of observations against reflectance-based PLSR predictions among external validation data

834   for various leaf structural and chemical traits. The black dashed line is the 1:1 line. Colored lines

835   represent best-fit lines from OLS regression for each dataset. Error bars around each point represent 95%

836   confidence intervals based on the ensemble of models produced in the 100× jackknife analysis.

Tables

Table 1: A summary of all projects with samples included in the study.

Project	Description	Latitude range	Longitude range	Species	Samples per functional group							Total
					Broadleaf	Conifer	Graminoid	Fern	Forb	Shrub	Vine	
Beauchamp-Rioux	Nine common broadleaf tree species measured from June to October 2018 at multiple forested sites in Québec (Beauchamp-Rioux 2022)	45.4-46.0 N	73.3-75.7 W	9	436							436
Blanchard	A sampling of nearly all tree species found in southern Québec, collected across multiple sites throughout summer 2019	45.1-46.0 N	71.1-74.0 W	43	270	68				3		341
Boucherville 2018	A variety of plants sampled in August 2018 from the Parc national des Îles-de-Boucherville in southern Québec	45.6-45.7 N	73.5-73.6 W	12			14		30	22	6	72
Boucherville 2019	A variety of plants sampled in July 2019 from the Parc national des Îles-de-Boucherville in southern Québec	45.6 N	73.5 W	19	40		38		59	26	10	173
CABO General	A small number of samples from southern Québec not collected as part of a specific project	45.5-46.0 N	71.9-74.1 W	13	33		6		3	39		81
Crofts	Trees sampled in July 2019 across an altitudinal gradient in southeastern Québec, along the transition from temperate deciduous to boreal forest	45.4-45.5 N	71.1-71.2 W	25	139	60				5		204
Girard	Four common bog plants sampled in summer 2018 in Ontario and southern Québec across a gradient of nitrogen deposition (Girard et al. 2020)	45.4-46.8 N	71.1-75.5 W	4			23			72		95
Hacker	Plants sampled in May 2019 to evaluate functional trait variation throughout an endangered oak	48.8-49.3 N	123.1-123.7 W	23	10		38	7	90	56		201

	Description	Lat	Long									Total
	savannah in British Columbia (Hacker et al. 2022)											
Phragmites Temporal	Six co-occurring wetland plant species sampled from June to September 2019 in Boucherville, Québec (Pardo 2021)	45.6 N	73.4-73.5 W	6			240		60			300
Warren	*Agonis flexuosa* sampled in November 2018 along a chronosequence of increasing soil age in southwestern Australia (Turner et al. 2018)	34.6 S	115.9 E	1	68							68
Total				104	996	128	359	7	242	223	16	1971

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844 Table 2: Summary statistics for the performance of reflectance-based models calibrated on CABO data and applied to internal and external validation datasets.
845 %RMSE is calculated as RMSE divided by the inner 95% trait range. The column '# obs' refers to the number of observations for the trait in the full CABO
846 dataset.
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	Internal					Dessain			LOPEX			ANGERS		
	# obs	# comps	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE
LMA (kg m^{-2})	1943	23	0.892	0.0175	8.04	0.593	0.0855	123	0.656	0.111	112	0.517	0.137	117
LDMC (mg g^{-1})	1930	20	0.867	32.9	9.78	0.824	40.6	13.6	0.860	46.4	12.0			
EWT (mm)	1929	13	0.885	0.0334	8.46	0.629	0.0395	24.2	0.888	0.0705	27.5	0.825	0.0665	34.8
N (%)	1963	19	0.709	0.403	13.7	0.417	0.664	21.6	0.535	0.854	20.2			
C (%)	1963	20	0.747	1.38	13.2	0.41	2.23	31.6	0.145	2.81	25.5			
Solubles (%)	1919	23	0.733	6.59	15.1	0.284	10.9	26.7						
Hemicellulose (%)	1915	20	0.695	4.35	16.1	0.324	5.28	19.7						
Cellulose (%)	1946	25	0.826	2.74	12.7	0.327	6.68	43.5	0.353	5.47	23.1			
Lignin (%)	1946	20	0.558	2.82	19.1	0.152	4.68	29.9	0.164	4.79	26.8			
Chl a (mg g^{-1})	1941	13	0.689	2.07	13.8	0.312	2.88	26.7	0.409	2.72	28.1	0.503	2.80	21.9
Chl b (mg g^{-1})	1941	14	0.679	0.716	15.0	0.318	1.05	31.6	0.441	0.927	23.4	0.409	1.32	28.7
Carotenoids (mg g^{-1})	1941	12	0.644	0.434	14.7	0.195	0.585	27.0	0.288	0.694	24.8	0.419	1.09	29.9
Al (mg g^{-1})	677	6	0.304	0.0293	24.6	0.006	0.0813	33.5						
Ca (mg g^{-1})	678	17	0.541	4.03	20.4	0.009	10.4	32.8						
Cu (mg g^{-1})	675	5	0.295	0.00311	25.9	0.004	0.0147	34.0						
Fe (mg g^{-1})	677	4	0.286	0.0295	23.0	0.045	0.053	26.6						
K (mg g^{-1})	678	10	0.55	2.32	15.7	0.443	9.13	28.9						
Mg (mg g^{-1})	678	18	0.394	0.714	22.8	0.149	2.3	53.6						
Mn (mg g^{-1})	678	10	0.254	0.164	22.8	0.005	0.504	54.6						
Na (mg g^{-1})	678	6	0.519	0.451	16.7	0.085	0.881	22.6						
P (mg g^{-1})	678	15	0.313	0.607	20.5	0.051	2.06	51.2						
Zn (mg g^{-1})	677	17	0.487	0.0486	24.2	0.045	0.118	34.9						

Supplementary Materials

Sampling

We collected spectra and traits for all specimens using consistent protocols. For each sample, we

collected sunlit leaves (>3 h per day of sun exposure) that were relatively homogeneous—for trees and

shrubs, at about the same canopy position in the same individual. When possible, we aimed to collect

enough leaves in each sample to carry out all destructive trait measurements while retaining several grams

of leaf tissue in long-term storage. We avoided leaves that showed clear signs of senescence, and

(particularly for spectral measurements) aimed to choose healthy-looking leaves without visible herbivore

or pathogen damage. On a subset of leaves, we measured reflectance and transmittance spectra and

punched leaf disks for pigment analyses. We also selected random subsets of visually similar leaves to

measure other structural or chemical traits. To prepare each sample for chemical trait measurements, we

dried several grams of leaves at 65 °C for three days and ground them in a 2 mm cyclone mill. All trait

measurements exclude petioles but include raches of compound leaves, since they are functionally

equivalent to the midrib of a simple leaf.

We identified species in the field and used the Database of Vascular Plants of Canada (VASCAN; Desmet

& Brouillet 2013) to classify each species by its family and growth form. VASCAN classified plants in

our dataset as either shrubs, trees, herbs, or vines. Some species were listed as adopting either shrub or

tree growth forms, in which case we manually classified them based on the habit of the individuals

sampled. We further divided species into functional groups as follows: we classified trees as either

needleleaf coniferous trees (here, families Pinaceae or Cupressaceae) or broadleaf trees; and we classified

herbs as either graminoids (families Poaceae, Cyperaceae, Typhaceae, or Juncaceae) or forbs. The

Australian species *Agonis flexuosa* (Willd.) Sweet was not in VASCAN, so we assigned it to the

broadleaf tree functional group. Broadleaf trees were by far the best-represented functional group ($n =$

996; species = 42), but the database also includes graminoids ($n = 359$; species = 15), forbs ($n = 242$;

species = 20), shrubs (n = 223; species = 20), conifers (n = 128; species = 12), vines (n = 16; species = 1),

and ferns (n = 7; species = 1). In total, the database includes samples from 31 families.

Spectral measurements and processing

We measured directional-hemispherical reflectance and transmittance spectra (350-2500 nm) using an

HR-1024i spectroradiometer equipped with a DC-R/T integrating sphere from Spectra Vista Corporation

(Poughkeepsie, NY, USA). The instrument's nominal bandwidth varies among sensors: ≤ 1.5 nm through

the range of the silicon detector array (~350-1000 nm), ≤ 3.8 nm through the first InGaAs array (~1000-

1890 nm), and ≤ 2.5 nm through the second InGaAs array (~1890-2500 nm). The spectral resolution

likewise ranges from 3.3 to 9.5 nm, depending on the spectral region. For each sample, we measured

spectra from the adaxial surface of six leaves or leaf arrays (Laliberté & Soffer 2018a). For leaves that

were too small or narrow to cover the port of the integrating sphere, we followed a protocol adjusted from

Noda et al. (2013) to measure single-layer composites of leaves (Laliberté & Soffer 2018b). We either

sealed leaves in plastic bags or left them attached to the branch to reduce water loss during the short

interval between collection and measurement. Reflectance measurements were calibrated against a white

Spectralon 99% reflectance standard and corrected for stray light.

We processed reflectance and transmittance spectra following Schweiger & Laliberté (2020). We

resampled the spectra linearly to 1 nm resolution and interpolated linearly over the sensor overlap regions.

We then averaged spectra from leaves of the same sample and reduced noise by applying a Savitzky-

Golay filter 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. We trimmed spectra to 400-2400 nm to remove the particularly noisy ends of the spectrum.

Finally, we calculated absorptance at each wavelength by subtracting the sum of reflectance and

transmittance from 1. We performed all spectral processing using *spectrolab* v. 0.0.10 (Meireles et al.

2017) in *R* version 3.6.3 (R Core Team 2020).

Spectral measurement protocols are given in other sources for external validation data, including LOPEX

(Hosgood et al. 1993; Hosgood et al. 1994), ANGERS (Jacquemoud et al. 2003; Féret et al. 2008;

http://opticleaf.ipgp.fr/index.php?page=database), and Dessain (Kothari et al. 2022). The Dessain dataset

is expanded from our previous description in Kothari et al. (2022); it has 200 samples, including trees (n

= 107; species = 36), shrubs (n = 53; species = 29), forbs (n = 28; species = 19), graminoids (n = 6;

species = 2), ferns (n = 4; species = 3), and vines (n = 2; species = 2). Each dataset contained spectra that

included the 400-2400 nm range already resampled to 1 nm resolution. For consistency, we processed the

external validation spectra much like our core dataset. First, we either interpolated over the sensor overlap

region (Dessain) or matched sensors using *match_sensors()* in *spectrolab*. Based on a visual assessment,

we matched sensors at 862 nm for LOPEX, but did not match sensors for ANGERS because there was no

conspicuous offset at the sensor overlap region. Next, we applied a Savitzky-Golay filter to the three

datasets, using the same parameters we used on the CABO dataset. Finally, we trimmed all spectra to

400-2400 nm.

Trait measurements

Leaf mass and area

We measured the fresh mass of a subset of leaves from each sample in the field shortly after collection.

We rehydrated the leaves in sealed plastic bags with damp paper towels and stored them in a refrigerator

for at least 12 h. We then measured their rehydrated mass, scanned them, and measured their area using

WinFOLIA (Regent Instruments, Québec, QC, CA). Finally, we dried them at 65 °C for at least 72 h

before measuring their dry mass. We used total masses and areas, summed across all leaves from a

sample, in trait calculations. We calculated leaf mass per area (LMA) as the dry mass divided by leaf

area; leaf dry matter content (LDMC) as dry mass divided by fresh mass; and equivalent water thickness

(EWT) as fresh mass minus dry mass, all divided by leaf area (Laliberté 2018).

Chemical traits

We estimated leaf carbon and nitrogen concentration on ground samples using an Elementar Vario

MICRO Cube (Langenselbold, Hesse, Germany; Ayotte et al. 2019). We measured concentrations of ten

other nutrients (Al, Ca, Cu, Fe, K, Mg, Mn, Na, P, and Zn) using nitric acid digestion followed by

inductively coupled plasma optical-emission spectrometry (ICP-OES) on an Optima 7300DV

(PerkinElmer, Waltham, MA, USA). These ICP-OES data are available only for the Beauchamp-Rioux,

Boucherville, Girard, and Warren projects. We removed six data points from the ICP-OES output as

unrepresentative outliers because their reported values were >30% greater than all others for their

respective elements. We also set concentrations below the detection limit, which sometimes returned as

negative, to zero.

We measured carbon fractions on ground samples using an ANKOM2000 Fiber Analyzer (ANKOM

Technology, Macedon, NY, USA; Ayotte & Laliberté 2019). The procedure is a sequential digestion in a

series of solutions that wash away fractions of the leaf tissue, leaving behind fractions that are

increasingly recalcitrant. The first digestion at 100 °C in a neutral detergent solution washes away soluble

cell contents. The second digestion at 100 °C in a mildly acidic detergent solution washes away bound

proteins and hemicellulose, which we simply call hemicellulose. The third digestion at room temperature

in 72% sulfuric acid removes cellulose, leaving behind lignin and other recalcitrant materials. Finally, we

ashed all samples in a furnace to separate inorganic recalcitrant materials (ash) from organic ones, which

we simply call lignin. By weighing the remaining leaf tissue after each stage, we could determine the

mass of each fraction. We excluded solubles and hemicellulose from three species (*Acer platanoides*

Linnaeus, *Ulmus americana* Linnaeus, and *Ulmus rubra* Muhlenberg) and all carbon fractions from *Alnus*

incana subsp. rugosa (Du Roi) R.T. Clausen because of leaf properties that caused unreliable results. For

example, *A. platanoides* has leaf exudates that wash off inconsistently in neutral detergent; *A. incana*

samples often returned negative values of hemicellulose.

We stored leaf disks in a cooler with ice in the field and transferred them first to a -20 °C freezer in the

lab, then to a -80 °C freezer for long-term storage. When we needed to ship them (e.g. the Warren

project), we freeze-dried them first. We extracted pigments from frozen leaf disks (or cut pieces, for

narrow-leaved plants) under darkness in an $MgCO_3$-MeOH solution and estimated pigment

concentrations on a SPECTROstar Nano microplate reader (BMG LABTECH, Ortenburg, Germany)

using a spectrophotometric protocol (Girard et al. 2020). While concentrations were measured per unit

fresh mass, we used LDMC as a conversion factor to express them per unit dry mass.

We occasionally removed specific trait values when an error was noted during the measurement protocol.

External validation

Trait measurement protocols are given in the aforementioned sources for external validation data,

including LOPEX (Hosgood et al. 1994), ANGERS (Féret et al. 2008;

http://opticleaf.ipgp.fr/index.php?page=database), and Dessain (Kothari et al. 2022). In the Dessain

dataset, we did not measure initial fresh mass of leaves and instead used the rehydrated mass. This

difference could cause EWT to be overestimated and LDMC to be underestimated. However, relative

water content in the remaining projects was usually close to 100% (median: 87.7%, 2.5–97.5[th] percentile:

68.3–97.6%). We also removed solubles and hemicellulose values from the Dessain dataset for the same

species whose values we removed from the main CABO dataset. Finally, we used a conservative outlier

detection procedure—the same as we used for the main CABO dataset—to remove eight data points from

Dessain ICP-OES data. In LOPEX, there were two estimates of both cellulose and lignin for each sample,

so we took the average for each one.

Besides applying the main set of models described in the main text, we also tested whether certain

transformations of the spectral data could improve performance on the external validation data—as, for

example, if certain instruments tended to measure higher or lower reflectance across the spectrum. We

used: (1) brightness normalization, which normalizes each spectrum to a unit vector while preserving its

shape (Feilhauer et al. 2010), and (2) continuum removal, which performs an albedo normalization by

reporting the difference between the actual spectrum and a linear interpolation of its convex hull (Clark &

Roush 1984). We first transformed the spectra in the CABO dataset using either brightness normalization

or continuum removal and built models using the calibration and validation approaches described in the

main text (model performance for internal validation not shown). We then applied the models to the

external validation data transformed likewise. Neither transformation caused any systematic improvement

in trait estimates (Tables S4-5).

Supplementary References

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Supplementary figures

 Fig. S1: (top) The first two dimensions in a principal components analysis (PCA) of CABO trait data,

 excluding all elements other than C and N. All chemical trait data are expressed on a mass basis, as in the

 core PLSR analyses of this paper. (bottom) A PCA of CABO trait data with chemical trait data expressed

 on a normalization-independent basis (Osnas et al. 2013), which is meant to eliminate covariance induced

 by jointly normalizing variables by mass. Here, we also remove solubles, since carbon fractions are nearly

 constrained to sum to 1, as well as chlorophyll *b* and carotenoids, since the three pigment pools covary

 strongly.

1069   Fig. S2: The median leaf-level (A) reflectance, (B) transmittance, and (C) absorptance at each wavelength

1070   across the spectrum, separated by functional group. We omit ferns and vines, which are only represented

1071   by one species each. The dashed line shows the coefficient of variation across all samples.

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1073 Fig. S3: Plots of observations against PLSR predictions among internal validation data for structural and

1074 water-related traits, comparing predictions from reflectance, transmittance, and absorptance spectra. The

1075 black dashed line in each panel is the 1:1 line. Colored lines represent best-fit lines from OLS regression

1076 for each functional group. Error bars around each point represent 95% confidence intervals based on the

1077 ensemble of models produced in the 100× jackknife analysis.

1078

Fig. S4: Plots of observations against PLSR predictions among internal validation data for N, C, and P,

comparing predictions from reflectance, transmittance, and absorptance spectra. P is only available for the

Beauchamp-Rioux, Boucherville 2018, Girard, and Warren projects. The black dashed line in each panel

is the 1:1 line. Colored lines represent best-fit lines from OLS regression for each functional group. Error

bars around each point represent 95% confidence intervals based on the ensemble of models produced in

the 100× jackknife analysis.

1085

Fig. S5: Plots of observations against PLSR predictions among internal validation data for carbon

fractions, comparing predictions from reflectance, transmittance, and absorptance spectra. The black

dashed line in each panel is the 1:1 line. Colored lines represent best-fit lines from OLS regression for

each functional group. Error bars around each point represent 95% confidence intervals based on the

ensemble of models produced in the 100× jackknife analysis.

Fig. S6: Plots of observations against PLSR predictions among internal validation data for pigments,

comparing predictions from reflectance, transmittance, and absorptance spectra. The black dashed line in

each panel is the 1:1 line. Colored lines represent best-fit lines from OLS regression for each functional

group. Error bars around each point represent 95% confidence intervals based on the ensemble of models

produced in the 100× jackknife analysis.

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Fig. S7: Plots of observations against PLSR predictions among internal validation data for Al, Ca, and

Cu, comparing predictions from reflectance, transmittance, and absorptance spectra. These data are only

available for the Beauchamp-Rioux, Boucherville 2018, Girard, and Warren projects. The black dashed

line in each panel is the 1:1 line. Colored lines represent best-fit lines from OLS regression for each

functional group. Error bars around each point represent 95% confidence intervals based on the ensemble

of models produced in the 100× jackknife analysis.

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Fig. S8: Plots of observations against PLSR predictions among internal validation data for Fe, K, and Mg,

comparing predictions from reflectance, transmittance, and absorptance spectra. These data are only

available for the Beauchamp-Rioux, Boucherville 2018, Girard, and Warren projects. The black dashed

line in each panel is the 1:1 line. Colored lines represent best-fit lines from OLS regression for each

functional group. Error bars around each point represent 95% confidence intervals based on the ensemble

of models produced in the 100× jackknife analysis.

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Fig. S9: Plots of observations against PLSR predictions among internal validation data for Mn, Na, and

Zn, comparing predictions from reflectance, transmittance, and absorptance spectra. These data are only

available for the Beauchamp-Rioux, Boucherville 2018, Girard, and Warren projects. The black dashed

line in each panel is the 1:1 line. Colored lines represent best-fit lines from OLS regression for each

functional group. Error bars around each point represent 95% confidence intervals based on the ensemble

of models produced in the 100× jackknife analysis.

1118

    Fig. S10: The distributions of reflectance-based model performance statistics for each trait based on 100

1120   jackknife resampling iterations from the calibration data set. The red dots show the R^2 and %RMSE from

1121   applying the ensemble of models to the internal validation subset. Abbreviations: sol = solubles, hemi =

1122   hemicellulose, cell = cellulose, lign = lignin, car = total carotenoids.

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1125   Fig. S11: The distributions of transmittance-based model performance statistics for each trait based on

1126   100 jackknife resampling iterations from the calibration data set. The red dots show the R^2 and %RMSE

1127   from applying the ensemble of models to the internal validation subset. Abbreviations: sol = solubles,

1128   hemi = hemicellulose, cell = cellulose, lign = lignin, car = total carotenoids.

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1131   Fig. S12: The distributions of absorptance-based model performance statistics for each trait based on 100

1132   jackknife resampling iterations from the calibration data set. The red dots show the R^2 and %RMSE from

1133   applying the ensemble of models to the internal validation subset. Abbreviations: sol = solubles, hemi =

1134   hemicellulose, cell = cellulose, lign = lignin, car = total carotenoids.

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1137   Fig. S13: The variable influence on projection (VIP) metric for reflectance-based models (Wold et al.

1138   2001). The dashed line at 0.8 represents a heuristic threshold for band importance recommended by

1139   Burnett et al. (2021).

1140

Fig. S14: The variable influence on projection (VIP) metric for transmittance-based models (Wold et al.

2001). The dashed line at 0.8 represents a heuristic threshold for band importance recommended by

Burnett et al. (2021).

Fig. S15: The variable influence on projection (VIP) metric for absorptance-based models (Wold et al.

2001). The dashed line at 0.8 represents a heuristic threshold for band importance recommended by

Burnett et al. (2021).

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1149   Fig. S16: Plots of observations against reflectance-based PLSR predictions among internal validation data

1150   for various area-based leaf chemical traits. The black dashed line in each panel is the 1:1 line. Colored

1151   lines represent best-fit lines from OLS regression for each functional group. Error bars around each point

1152   represent 95% confidence intervals based on the ensemble of models produced in the 100× jackknife

1153   analysis.

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Fig. S17: Plots of observations against reflectance-based PLSR predictions among internal validation data

for area-based contents of elements other than C and N. The black dashed line in each panel is the 1:1

line. Colored lines represent best-fit lines from OLS regression for each functional group. Error bars

around each point represent 95% confidence intervals based on the ensemble of models produced in the

$100\times$ jackknife analysis.

Fig. S18: The distributions of reflectance-based model performance statistics for each area-based

chemical trait based on 100 jackknife resampling iterations 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.

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Fig. S19: The variable influence on projection (VIP) metric for reflectance-based models (Wold et al.

2001) for area-based chemical traits. The dashed line at 0.8 represents a heuristic threshold for band

importance recommended by Burnett et al. (2021).

Fig. S20: Plots of observations against reflectance-based PLSR predictions among external validation

data for elements other than C and N. The black dashed line is the 1:1 line. Error bars around each point

represent 95% confidence intervals based on the ensemble of models produced in the 100× jackknife

analysis. The colored line represents the best-fit line from OLS regression.

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Supplementary tables

Table S1: Summary statistics for internal validations of models predicting area-based leaf chemical traits from reflectance spectra. %RMSE is calculated as RMSE divided by the inner 95% trait range. Mass-proportionality is calculated as in Osnas et al. (2013) as the slope fitting the area-normalized trait vs. LMA, both log-transformed.

Trait	Mass-proportionality	# comps	Area-based estimates			Mass-based estimates × LMA estimates		
			RMSE	%RMSE	R^2	RMSE	%RMSE	R^2
N (mg cm^{-2})	0.591	20	0.0279	11.9	0.792	0.0459	19.5	0.542
C (mg cm^{-2})	1.057	24	0.856	7.70	0.902	0.830	7.47	0.908
Solubles (mg cm^{-2})	1.000	22	1.10	7.76	0.903	1.18	8.30	0.891
Hemicellulose (mg cm^{-2})	0.834	22	0.404	13.7	0.733	0.454	15.4	0.685
Cellulose (mg cm^{-2})	1.010	25	0.442	10.4	0.804	0.398	9.39	0.840
Lignin (mg cm^{-2})	1.361	22	0.272	10.4	0.836	0.281	10.8	0.827
Chl a (µg cm^{-2})	0.316	14	11.9	14.7	0.741	16.7	20.5	0.609
Chl b (µg cm^{-2})	0.308	14	3.96	15.6	0.722	5.63	22.2	0.574
Carotenoids (µg cm^{-2})	0.311	15	2.68	16.6	0.671	3.69	22.8	0.518
Al (µg cm^{-2})	0.865	4	0.265	22.8	0.253	0.253	21.8	0.295
Ca (µg cm^{-2})	0.435	17	35.7	16.2	0.515	41.3	18.7	0.366
Cu (µg cm^{-2})	-0.092	6	0.0223	23.1	0.321	0.0262	27.1	0.089
Fe (µg cm^{-2})	0.203	1	0.230	29.6	0.006	0.234	30.2	0.021
K (µg cm^{-2})	0.335	7	19.9	19.9	0.468	21.5	21.5	0.372
Mg (µg cm^{-2})	0.544	15	6.97	20.7	0.446	7.69	22.8	0.389
Mn (µg cm^{-2})	0.905	11	1.60	21.5	0.321	1.70	22.8	0.263
Na (µg cm^{-2})	1.906	6	5.79	13.9	0.696	5.54	13.3	0.717
P (µg cm^{-2})	0.421	6	5.51	22.9	0.307	6.13	25.5	0.238
Zn (µg cm^{-2})	-0.087	12	0.428	24.4	0.338	0.436	24.9	0.347

Table S2: Summary statistics for internal validations of models predicting leaf traits from reflectance (R), transmittance (T), and absorptance (A) spectra. %RMSE is calculated as RMSE divided by the inner 95% trait range.

Trait	# comps			RMSE			%RMSE			R^2		
	R	T	A	R	T	A	R	T	A	R	T	A
LMA (kg m^{-2})	23	27	26	0.0175	0.0146	0.0141	8.04	6.69	6.49	0.892	0.928	0.930
LDMC (mg g^{-1})	20	21	19	32.9	28.3	29.1	9.78	8.4	8.66	0.867	0.901	0.895
EWT (mm)	13	13	12	0.0334	0.0292	0.0298	8.46	7.38	7.54	0.885	0.913	0.910
N (%)	19	22	19	0.403	0.342	0.351	13.7	11.6	12.0	0.709	0.791	0.780
C (%)	20	25	23	1.38	1.38	1.35	13.2	13.2	13.0	0.747	0.748	0.757
Solubles (%)	23	24	25	6.59	6.41	6.35	15.1	14.7	14.6	0.733	0.747	0.751
Hemicellulose (%)	20	23	22	4.35	4.12	4.17	16.1	15.3	15.5	0.695	0.725	0.719
Cellulose (%)	25	26	26	2.74	2.76	2.53	12.7	12.8	11.7	0.826	0.822	0.851
Lignin (%)	20	23	22	2.82	2.93	2.77	19.1	19.8	18.7	0.558	0.522	0.576
Chl a (mg g^{-1})	13	20	16	2.07	1.77	1.87	13.8	11.8	12.5	0.689	0.772	0.744
Chl b (mg g^{-1})	14	23	18	0.716	0.607	0.63	15.0	12.7	13.2	0.679	0.769	0.751
Carotenoids (mg g^{-1})	12	16	13	0.434	0.396	0.404	14.7	13.4	13.7	0.644	0.703	0.691
Al (mg g^{-1})	6	9	10	0.0293	0.0283	0.0261	24.6	23.7	21.9	0.304	0.328	0.441
Ca (mg g^{-1})	17	22	17	4.03	3.32	3.88	20.4	16.8	19.7	0.541	0.686	0.566
Cu (mg g^{-1})	5	4	5	0.00311	0.00315	0.00307	25.9	26.2	25.6	0.295	0.278	0.311
Fe (mg g^{-1})	4	6	5	0.0295	0.0301	0.0295	23.0	23.5	23.0	0.286	0.257	0.283
K (mg g^{-1})	10	18	18	2.32	1.87	1.94	15.7	12.7	13.1	0.55	0.712	0.688
Mg (mg g^{-1})	18	17	16	0.714	0.708	0.74	22.8	22.6	23.6	0.394	0.410	0.354
Mn (mg g^{-1})	10	8	11	0.164	0.177	0.158	22.8	24.6	22.0	0.254	0.157	0.304
Na (mg g^{-1})	6	4	5	0.451	0.472	0.452	16.7	17.5	16.8	0.519	0.461	0.520
P (mg g^{-1})	15	20	20	0.607	0.550	0.579	20.5	18.6	19.6	0.313	0.460	0.410
Zn (mg g^{-1})	17	12	13	0.0486	0.0491	0.0493	24.2	24.4	24.5	0.487	0.473	0.462

Table S3: Summary statistics from model transferability analyses among functional groups. Each functional group in turn, other than ferns and vines, was left out of the dataset used to train a PLSR model for each trait using 10-fold cross-validation. The functional group left out (column names) was then used for validation.

	Broadleaf			Conifer			Forb			Graminoid			Shrub		
	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE
LMA (kg m^{-2})	0.764	0.0249	12.0	0.413	0.104	50.3	0.300	0.0278	13.4	0.552	0.0252	12.2	0.879	0.0230	11.1
LDMC (mg g^{-1})	0.668	37.6	10.7	0.517	40.2	11.5	0.814	39.0	11.1	0.675	57.2	16.3	0.832	33.7	9.61
EWT (mm)	0.453	0.0362	9.2	0.575	0.104	26.4	0.572	0.0555	14.1	0.906	0.0652	16.6	0.600	0.0329	8.34
N (%)	0.599	0.592	21.3	0.577	0.418	15.1	0.393	0.673	24.3	0.409	0.566	20.4	0.734	0.584	21.1
C (%)	0.468	1.54	14.5	0.239	1.88	17.7	0.035	2.13	20.0	0.089	2.16	20.3	0.746	2.28	21.4
Solubles (%)	0.357	8.15	18.3	0.266	9.67	21.7	0.108	15.4	34.6	0.086	22.4	50.2	0.292	8.99	20.2
Hemicellulose (%)	0.262	4.43	16.0	0.084	5.51	19.9	0.041	9.37	33.8	0.067	16.0	57.8	0.255	5.68	20.5
Cellulose (%)	0.285	4.08	18.4	0.358	5.64	25.5	0.090	6.74	30.5	0.061	11.6	52.4	0.002	4.63	20.9
Lignin (%)	0.182	3.44	22.4	0.358	2.76	18.0	0.116	3.60	23.4	0.034	5.46	35.6	0.466	4.06	26.4
Chl a (mg g^{-1})	0.667	2.31	16.5	0.535	1.60	11.5	0.210	4.63	33.1	0.316	2.88	20.6	0.750	2.41	17.2
Chl b (mg g^{-1})	0.668	0.690	14.5	0.481	0.620	13.1	0.157	1.78	37.5	0.332	0.958	20.2	0.737	0.736	15.5
Carotenoids (mg g^{-1})	0.629	0.478	16.5	0.406	0.358	12.3	0.196	0.841	28.9	0.187	0.620	21.3	0.566	0.634	21.8

Table S4: Summary statistics for the performance of models calibrated on continuum-removed CABO data and applied to external validation datasets.

	Dessain			LOPEX			ANGERS		
	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE
LMA (kg m^{-2})	0.364	0.0907	131	0.435	0.0976	98.3	0.444	0.130	111
LDMC (mg g^{-1})	0.840	33.1	11.1	0.893	55.2	14.3			
EWT (mm)	0.617	0.0477	29.2	0.861	0.0661	25.8	0.774	0.0751	39.3
N (%)	0.409	0.741	24.1	0.505	0.837	19.8			
C (%)	0.447	3.21	45.5	0.163	3.81	34.5			
Solubles (%)	0.222	10.9	26.8						
Hemicellulose (%)	0.263	5.98	22.3						
Cellulose (%)	0.324	3.86	25.2	0.243	9.05	38.3			
Lignin (%)	0.140	4.44	28.4	0.158	4.98	27.9			
Chl a (mg g^{-1})	0.354	2.38	22.1	0.411	3.53	36.4	0.473	4.06	31.7
Chl b (mg g^{-1})	0.369	0.842	25.2	0.431	1.23	31.1	0.347	1.99	43.1
Carotenoids (mg g^{-1})	0.213	0.517	23.9	0.307	0.64	22.8	0.376	0.934	25.6
Al (mg g^{-1})	0.001	0.0763	31.5						
Ca (mg g^{-1})	0.011	18.1	57.2						
Cu (mg g^{-1})	0.001	0.0144	33.4						
Fe (mg g^{-1})	0.000	0.0594	29.8						
K (mg g^{-1})	0.322	8.04	25.5						
Mg (mg g^{-1})	0.107	2.89	67.3						
Mn (mg g^{-1})	0.024	0.568	61.5						
Na (mg g^{-1})	0.062	0.903	23.2						
P (mg g^{-1})	0.041	3.30	82						
Zn (mg g^{-1})	0.054	0.107	31.8						

Table S5: Summary statistics for the performance of models calibrated on brightness-normalized CABO data and applied to external validation datasets.

	Dessain			LOPEX			ANGERS		
	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE	R^2	RMSE	%RMSE
LMA (kg m^{-2})	0.658	0.0700	101	0.735	0.0879	88.6	0.540	0.127	108
LDMC (mg g^{-1})	0.812	41.7	14	0.881	45.7	11.8			
EWT (mm)	0.614	0.0422	25.8	0.905	0.0619	24.1	0.797	0.0751	39.2
N (%)	0.475	0.659	21.4	0.541	0.918	21.8			
C (%)	0.413	1.91	27.1	0.114	2.85	25.8			
Solubles (%)	0.306	12.2	30.1						
Hemicellulose (%)	0.353	4.91	18.3						
Cellulose (%)	0.307	7.41	48.3	0.358	5.37	22.7			
Lignin (%)	0.187	4.59	29.3	0.153	4.89	27.4			
Chl a (mg g^{-1})	0.333	2.75	25.5	0.392	3.44	35.5	0.548	2.8	21.9
Chl b (mg g^{-1})	0.342	0.964	28.9	0.451	1.17	29.6	0.426	1.45	31.4
Carotenoids (mg g^{-1})	0.193	0.568	26.3	0.270	0.656	23.4	0.491	1.01	27.7
Al (mg g^{-1})	0.003	0.0742	30.6						
Ca (mg g^{-1})	0.005	11.7	37						
Cu (mg g^{-1})	0.001	0.0145	33.4						
Fe (mg g^{-1})	0.046	0.0526	26.4						
K (mg g^{-1})	0.468	9.02	28.6						
Mg (mg g^{-1})	0.123	2.37	55.1						
Mn (mg g^{-1})	0.004	0.503	54.5						
Na (mg g^{-1})	0.084	0.907	23.3						
P (mg g^{-1})	0.063	2.33	58						
Zn (mg g^{-1})	0.037	0.107	31.7						