

Article

Retrieving Foliar Traits of *Quercus garryana* var. *garryana* across a Modified Landscape Using Leaf Spectroscopy and LiDAR

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Abstract: Understanding the ecological effects of human activities on an ecosystem is integral to the implementation of conservation management plans. The plasticity of plant functional traits presents an opportunity to examine the capacity for intraspecific functional trait variations to be indicators of anthropogenic landscape modifications. The presence of intraspecific trait variation would indicate that plants of a single species could be used to evaluate and map functional diversity, a common metric used to measure biodiversity. This study uses leaf spectroscopy, light detection and ranging (LiDAR) and partial least squares regression (PLSR) to examine the intraspecific variation of functional traits in a population of 40 *Quercus garryana* experiencing varying levels of anthropogenic influence at the site level (<0.3 km²) in Duncan, B.C., Canada. These individuals vary in their spatial relationship to roads, agricultural land use change and an encroaching Coastal Douglas-fir forest. A total of 14 functional traits were estimated using pre-determined PLSR coefficients from a multi-species dataset. LiDAR data for each tree and were organized into functional categories based on their influence of plant lifeform, leaf growth or leaf structure. Principal components analysis was performed on each functional category to determine the relative influence of each trait. Results show that leaf growth and lifeform functional trait categories express significant variation in relation to three anthropogenic landscape modifications, while traits associated to leaf structure only varied between land use types ($p = 0.05$). Diameter at breast height (DBH), mass-based chlorophyll and leaf mass per area (LMA) showed the strongest variation across treatments. These findings support the hypothesis that trait variation exists in small populations of the same species and illustrate that spectroscopy can be used to indirectly sense land use via the leaf functional traits of a single tree species.

Keywords: spectroscopy; functional diversity; biodiversity; partial least-squares regression; functional traits; intraspecific; hyperspectral

1. Introduction

The current speed and scale of anthropogenic landscape modification is unprecedented [1,2]. With such high capacity for human actions to inflict biotic and abiotic change, it is increasingly necessary to accurately monitor the health of ecosystems in a timely manner [3]. The application of remote sensing techniques to evaluate relevant ecosystem characteristics, such as biodiversity and plant health, provide opportunities to achieve these goals [4] and improvements in sensor technology, deployment platforms and computing are enabling new monitoring methodologies at multiple spatial scales [5–7]. One technique that has benefited from these recent developments is the use of spectroscopy to identify

plant functional traits [8–11]. These traits, which relate to plant growth and development strategies, provide biologically relevant information relating to photosynthesis and environmental drivers such as nutrient, water and stress regimes [12–15]. When evaluated at a site- or landscape-level, plant functional traits can be used to map functional diversity, community assembly and other ecosystem characteristics linked with biodiversity [16].

A knowledge gap exists, however, with regards to intraspecific functional trait variation at the site-level ($<1 \text{ km}^2$) [17]. Intraspecific trait variation (ITV) has been largely underrepresented in spectroscopic studies of natural ecosystems as many ecological studies assume that interspecific variation is much larger than intraspecific variation [17]. This assumption inherently ignores the variation present between individuals of the same species by averaging their trait values into a species mean. Recent studies have shown that ITV exists and is an important factor when evaluating plant functional traits at the community-level [18–21].

Correct interpretation of plant health is especially important in fragmented ecosystems that contain high-levels of biodiversity, making spectroscopy-based ITV increasingly relevant for the management of these areas. The conventional approach for understanding the effects of these modifications in the ITV of a species across a single site would include complete soil and vegetation sampling over multiple years, as well as long-term plot monitoring [22–24]. Considering the expense and time associated with these tasks, it is unlikely that this approach is feasible for the majority of land managers. It is necessary, then, to identify methodologies that are both cost- and time-efficient, while still providing relevant information.

In landscapes with significant anthropogenic modification, the use of spectroscopy to estimate functional traits offers significant potential. However, there is an inherent risk of mixing the spectral properties of leaves when using remotely sensed imagery with a spatial resolution coarser than a single leaf [25,26]. Therefore, methodologies that establish relationships between leaf level spectra and individual traits are critical before attempting to scale these relationships to coarser pixels.

Leaf spectroscopy provides continuous spectral information across the visible, near-infrared (NIR) and shortwave infrared (SWIR) wavelengths relating to how light interacts with pigments and chemicals in leaves [27,28]. Partial least-squares regression (PLSR) [29], a key method in chemometrics, is widely used to estimate the functional traits of a sample from the values recorded in a continuous spectral signature [30–33]. An even more complete record of a tree's morphological and physiological traits can be accomplished through the combination of spectral data with measurements of a tree derived from light detection and ranging (LiDAR) analysis [34,35]. The addition of these structural metrics provides insight to the variation in lifeform associated with the leaves used to derive functional traits, such as the height of the leaves above the ground.

Here we address ITV using *Quercus garryana* var. *garryana* Douglas ex Hook (Garry Oak) in British Columbia (B.C.), Canada as a test species. In general, functional traits of the genus *Quercus* vary significantly based on environment [36–38], suggesting that ITV may be present at the site-level where significant variation in human manipulation of the environment is present. *Q. garryana* has considerable conservation value and utility to this study as a consequence of its role in biodiversity [39] and known variability throughout its range. *Q. garryana* is also the flagship species for one of the most biodiverse and highly threatened ecosystems in Canada, the Garry Oak meadow [39]. These meadows exist on the south-east coast of Vancouver Island and have experienced anthropogenic influence for the majority of their history [40]. Prescribed burning was used to reduce tree seedlings, such as *Pseudotsuga menziesii* var. *menziesii* (Mayr) Franco (Coastal Douglas-fir), as well as promote the growth of culturally important herbaceous and grass species [40]. At the turn of the 19th century, anthropogenic land use initiatives switched from preservation to urbanization. Since then, the highly arable meadows have transitioned into forest, agricultural or urban sites, disrupting the historic processes that enabled high biodiversity. The presence of these different landscapes offers the opportunity to examine the spatial variation of functional traits in relation to encroaching ecosystems. Due to their decreasing area

and ecological importance, Garry Oak meadows are considered one of Canada's most endangered ecosystems [39].

Infrastructure and fire suppression are two environmental influences controlled by anthropogenic landscape modifications that are related to the decrease of *Q. garryana* range area and quality. Roads, specifically, have multiple associated effects on adjacent ecosystems. Roadsides are intensively managed to preclude woody plant growth and are often dominated by exotic species, increasing the likelihood of their introduction to native systems [41]. Changes in nutrient composition are also common in landscapes adjacent to roads as automobile exhaust and de-icing agents, specifically sodium chloride (NaCl), having the potential to affect the composition of surrounding vegetation [42,43].

In conjunction with road infrastructure, fire suppression has enabled the encroachment of woody species, such as *P. menziesii*, into Garry Oak Ecosystems [40]. In an open meadow, these seedlings face low competition and are able to grow and recruit unimpeded [39,44]. Combined, road and fire suppression have the power to alter the traditional species, structure and nutrient composition of Garry Oak ecosystems, threatening the inherently high levels of biodiversity within. The analysis of ITV within *Q. garryana* will remain important as the influences of land use change on areas of high biodiversity are expected to continue [1,2].

In this paper, leaf-level spectroscopy, LiDAR and pre-determined PLSR coefficients are used to explore the intraspecific functional trait variation of *Q. garryana* in association with anthropogenic landscape modifications across a single site. Specifically, (1) does intraspecific functional trait variation exist between *Q. garryana* experiencing different land use regimes and (2) do *Q. garryana* experiencing the same land use regime exhibit functional trait variation in relation to the spatial distribution of anthropogenic landscape modifications. We conclude by discussing the operational implications of this work and how analysis of functional trait variation within a single species can be used to examine the functional diversity of a site as well as indirectly sense the effects of land use.

2. Materials and Methods

2.1. Study Site

The Cowichan Garry Oak Preserve (CGOP; Lat: 48°48'29.85"N, Long: 123°37'54.34"W) presents an ideal study site as it contains a large population of *Q. garryana* that has been managed for over 100 years and has experienced various levels of human influence. A portion of the site consists of a natural Garry Oak meadow which has been maintained through a mixture of burning and mowing (Figure 1). Other portions of the site have been used exclusively for the production of hay, providing the opportunity to examine the functional traits of leaves that developed in an agricultural environment (Figure 1). A large, heavily trafficked road spans the south-east length of the site providing a spatial reference from which to examine the association of functional trait values in relation to roads (Figure 1). The northern section of the CGOP consists of a Coastal Douglas-fir Forest (CDF) that was traditionally managed through fire, mowing and removal (Figure 1) [39].

2.2. Airborne Data Acquisition and Processing

Road vector information and LiDAR data were acquired from the B.C. Data Catalogue maintained by the B.C. Ministry of Forest, Lands and Natural Resource Operations and Rural Development and is freely available online (www.catalogue.data.gov.bc.ca). The road dataset was last updated April, 2017, while the LiDAR data of the CGOP was acquired by the Municipality of North Cowichan on 29 May 2017. The LiDAR data was collected using a Riegl Q1560 dual-channel LiDAR system at a nominal flight height of 1700m above ground level and has a density of 15 returns/m².

A canopy height model (CHM) was created from the LiDAR point cloud using the CanopyModel function in FUSION (Forest Service of the U.S. Department of Agriculture, Seattle, WA, USA, 2014). The CHM was used to select 30 individual *Q. garryana* trees for sampling with five additional individuals selected from both the agricultural and natural meadow sub-sites to enable comparison between

the agriculture and meadow sites ($n = 40$). Lifeform metrics of height, maximum crown width and height to crown base were calculated from the CHM using the ‘lascanopy’ tool provided by LAsTools (rapidlasso GmbH, Gilching, DE, 2018) in ArcGIS Pro (ESRI, Redlands, CA, USA, 2018). The ‘lascanopy’ tool uses the values of each cell with the CHM raster to derive the aforementioned lifeform traits and presents them in a single attribute table (rapidlasso GmbH, Gilching, DE, 2018). Diameter at breast height (DBH) was measured in the field.

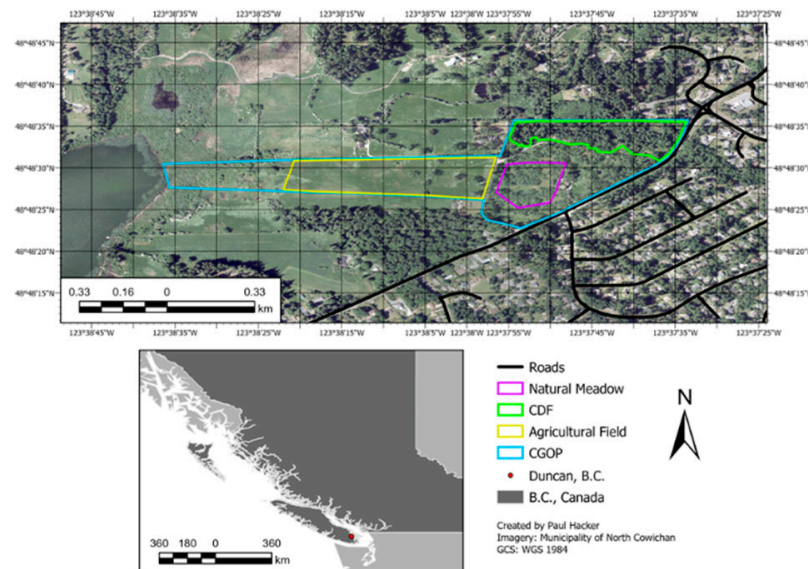


Figure 1. The Cowichan Garry Oak Preserve (CGOP) (light blue) and its associated sub-sites. The Coastal Douglas-fir Forest (CDF, green) places competitive pressure on the site as a whole, especially the Natural Meadow (purple). The Agricultural Field (yellow) was managed as a hay field for over 100 years and contains multiple *Quercus garryana* individuals.

2.3. Leaf Spectroscopy

Sun-lit leaf samples were collected from each of the 40 *Q. garryana* individual using 18m long pruners. All leaf samples were stored on ice for no longer than 1 h. Samples stored in the cooler underwent spectroscopy, scanning and weighing no more than 8 h after collection, after which they were flash frozen until drying. An ASD FieldSpec3 portable spectroradiometer and ASD Integrating Sphere, which is a closed illumination system (Analytical Spectral Devices Inc., Boulder, CO, USA) was used to measure the reflectance spectra of each sampled leaf. White and dark references were taken before each sample was scanned to normalize the recorded radiance, which enables the calculation of reflectance [45]. Each full scan comprised the average of 20-sub scans. Three leaves per tree underwent full scans an average of 20 times to provide a mean of approximately 400 scans per leaf sample. Reflectance was measured from 350–2500 nm and the values of each spectral band were the used as inputs for the partial least square regression (PLSR) that was used to estimate plant functional trait values.

2.4. Partial Least-Squares Regression

Recent studies have proven that PLSR is effective at estimating plant functional traits from a full spectral signature (350 nm–2500 nm) [33,46]. PLSR is a multivariate calibration method ideal for the analysis of data with high-dimensionality, such as that produced through spectroscopy [29,46,47]. The PLSR model is created using a training dataset which includes the spectral signatures and chemometric results of each functional trait for every leaf within [29,48]. The model produces output coefficients that are applied to a validation dataset, which also contains a spectral signature and

chemically determined values for each functional trait of every leaf within the dataset. To determine the accuracy of the PLSR coefficients, the estimated functional trait values produced by the validation dataset are compared to the chemical results [29]. For this study, the input variables are the spectral signature of 40 *Quercus garryana* leaves. This method is different to the traditional index-based approach of plant functional trait estimation [49].

This study utilizes PLSR leaf models and coefficients derived by Serbin 2012 using a dataset of various genera, including *Quercus*, from across the northern United States [50,51]. Chemical analysis of the sampled leaves was completed in order to validate the model [50,51]. The model was created using a similar fashion to those of other PLSR models designed for functional trait identification (Python, 2018) [46,50]. PLSR models for eleven functional traits were developed. The PLSR models were then applied to our spectra to provide trait estimates. This model was selected in favor of the index-based method as it employs the entire spectral signature of each sample and encompasses many genera, making it applicable for non-*Quercus* studies. This latter point is important as it enables the transference of methodologies employed in this study.

2.5. Principal Component Analysis

Functional traits of interest were separated into three categories based on their functional role: lifeform, leaf growth and leaf structure. Traits in the leaf growth and leaf structure categories were derived solely using coefficients from pre-existing, validated PLSR models (Table 1) [51]. Deriving functional traits in this manner highlights the ability of spectroscopy to provide information relating to a multitude of relevant plant traits. The functional traits in the lifeform category were measured principally using LiDAR (Table 1), again demonstrating the power of advanced remote sensing techniques. The PCA of each group enabled the trait with the highest amount of variation to be identified and its relationship with the other traits in the group to be extracted and analyzed. In total, 14 functional traits were evaluated in this study (Table 1).

Table 1. Categories of functional morphological traits used to analyze the variance between leaf samples and the agricultural, CDF and road anthropogenic landscape modifications.

Leaf Growth	Leaf Structure	Lifeform
Carbon (%)	Cellulose (%)	DBH (cm)
Carotenoid area (mmol/m ²)	Fiber (%)	Height (m)
Carotenoid mass (ng/mg)	Lignin (%)	Max Crown Width (m)
Chlorophyll ab area (mmol/m ²)	Leaf Mass Area (LMA) (g m ²)	Height to Crown base (m)
Chlorophyll ab mass (ng/mg)	Leaf Structure	
Nitrogen (%)	Cellulose (%)	

The values of each PC were extracted for all leaf samples in this study in order to enable their use as input values for statistical analysis. Normality and equal variance tests were completed in R using base functions of the software (R Core Team, 2018). Independence was achieved as the *Q. garryana* individuals in this study were not sampled more than once.

2.6. Statistical Analysis

Statistical assumptions were completed to determine if the functional trait estimation data was parametric. Upon finding the data was parametric the Kruskal-Wallis rank sums test was used to evaluate ITV between the agricultural and meadow sites. ITV in spatial relation to the road and CDF anthropogenic landscape modifications was evaluated using Kendall's Tau rank correlation coefficient test. To ensure that ITV was not spatially significant at random within the meadow a randomization test was completed. Each sample was given a random distance before Kendall's Tau was completed. This was iterated 1000 times and the means of the resulting Tau estimates, z-statistics and p-values

were recorded (Table 2). It was determined that the samples used in this study did not have spatially significant ITV in comparison to the samples evaluated at random distances.

Table 2. Mean z - and p -values resulting from the randomization test. The insignificant p -values suggest that intraspecific trait variation (ITV) does not exist at random within the meadow.

Group	z	p -value
Lifeform	0.01	0.497
Leaf growth	−0.045	0.512
Leaf Structure	−0.013	0.508

Road, RGB and LiDAR data were all processed in ArcGIS Pro (ESRI, Redlands, CA, 2018). LiDAR data was manipulated using the LAStools ArcGIS toolbox (rapidlasso GmbH, Gilching, DE, 2018). Spectroscopic data was analyzed using a combination of ASD's ViewSpecPro software and R (Analytical Spectral Devices Inc., Boulder, CO, USA; R Core Team, 2018). Statistical analysis was completed using the R package 'prcomp' (R Core Team, 2018).

3. Results

3.1. Leaf-Level Spectroscopy

Mean spectral signatures were created for the agricultural, meadow and sample population leaf samples. Clear differences in the reflectance values are observed in association with various absorption features, and therefore functional traits (Figure 2). Spectral variation is especially notable from 500–650nm (Figure 2a), wavelengths dominated by pigments, and 1400nm to 1700nm (Figure 2b), wavelengths dominated by water absorption as well as leaf structural compounds and proteins (49).

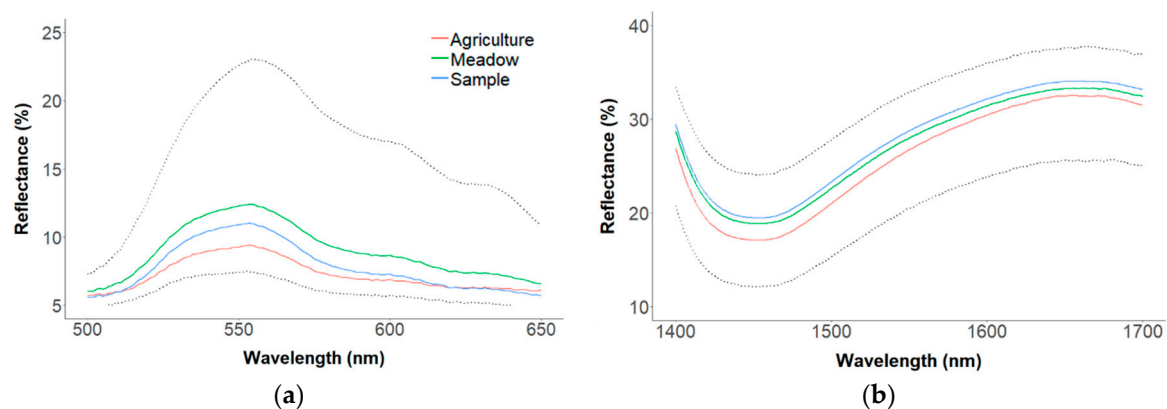


Figure 2. Mean spectral signatures of Agricultural ($n = 5$), Meadow ($n = 5$) and Sample ($n = 30$) population from (a) 500–650nm and (b) 1400–1700nm. Dotted lines signify the range of variation in the spectral signature evaluated.

3.2. Model Performance

PLSR model evaluation indicated that 10 of the 11 functional traits evaluated were well estimated by the PLSR model with predictive accuracies ranging ($R^2 = 0.30$ to 0.95 ; Table 3) [50,51]. Upon evaluation of the PLSR results, five meadow trees were removed from the analysis as the range of at least one of their functional trait estimations was outside the valid range of the model. The final sample population used for the analysis was 27 sample, 3 natural meadow and 5 agricultural *Q. garryana* individuals ($n = 35$).

Table 3. Mean and Standard deviation of the 10 functional traits calculated using partial least squares regression (PLSR).

Trait	Mean Value	Standard Deviation
Carbon (%)	50.082	0.219
Carotenoid Area (mmol/m ²)	181.786	3.383
Carotenoid Mass (ng/mg)	1266.166	38.087
Cellulose (%)	15.916	0.972
Chlorophyll ab Area (mmol/m ²)	589.166	24.965
Chlorophyll ab Mass (ng/mg)	7529.724	251.585
Fiber (%)	48.806	2.032
Lignin (%)	24.252	1.320
Nitrogen (%)	2.209	0.086
LMA (g/m ²)	139.895	13.114

3.3. Principal Component Analyses

3.3.1. PCA 1: ITV between Agricultural and Meadow Sub-Sites

The loadings of PC1 from each functional trait category were determined for the analysis of variation between the agricultural sub-site and the rest of the CGOP (Table 4). From the lifeform group, diameter at breast height (DBH) had the largest influence (−0.989) on PC1, which explained 99% of the variation. This suggests that DBH has the largest amount of variation between samples. Chlorophyll ab Mass (−0.989, 99% of variation) and LMA (0.995, 68% of variation) had the largest influence on the leaf growth and leaf structure functional categories respectively. All variables in the lifeform and structure PC1s are negatively correlated, while all variables in leaf growth's PC1 are positively correlated (Table 5).

Table 4. PCA 1 completed using all leaf samples (agricultural and natural meadow sub-sites) collected at the CGOP in July 2018. The loadings from the first principal component for each functional trait category are shown as it accounted for ≥68% of the variations in the analysis. Variations are presented in brackets.

PCA 1					
Lifeform	PC1 (>0.99)	Leaf Growth	PC1 (>0.99)	Leaf Structure	PC1 (0.68)
DBH	−0.995	Chlorophyll ab mass	0.989	LMA	−0.995
Height	−0.086	Carotenoid mass	0.137	Lignin	−0.068
Height to crown base	−0.043	Chlorophyll ab area	0.053	Fiber	−0.068
Max crown width	−0.031	Carotenoid area	0.005	Cellulose	−0.027
		Nitrogen	<0.0001		
		Carbon	<−0.0001		

3.3.2. PCA 2: ITV in Spatial Relation to Anthropogenic Landscape Modifications

PCA 2 (Table 5), which did not include the *Q. garryana* leaf samples from the agricultural site, differed from the PC1 values of PCA 1 (Table 5). DBH (−0.987, 98% of variation), chlorophyll ab mass (−0.989, 99% of variation) and LMA (−0.999, 90% of variation) still accounted for the largest amount of variation in the lifeform, leaf growth and leaf structure functional categories respectively. The fact that the same functional traits were dominant in the PC1s of both PCA 1 and PCA 2 indicates that they are more variable than other traits across the species, regardless of context.

Table 5. PCA 2 completed using only non-agricultural leaf samples collected at the CGOP in July 2018. The loadings from the first principal component for each functional trait category are shown as it accounted for $\geq 90\%$ of the variations in the analysis. Variations are presented in brackets.

PCA 2					
Lifeform	PC1 (0.98)	Leaf Growth	PC1 (>0.99)	Leaf Structure	PC1 (0.90)
DBH	−0.987	Chlorophyll ab mass	−0.989	LMA	−0.999
Height	−0.138	Carotenoid mass	−0.139	Fiber	0.013
Height to crown base	−0.069	Chlorophyll ab area	−0.045	Lignin	0.018
Max crown width	−0.033	Carotenoid area	−0.007	Cellulose	<−0.001
		Nitrogen	<−0.001		
		Carbon	<−0.001		

3.4. Statistical Analysis

3.4.1. PCA 1: ITV between Agricultural and Meadow Sub-Sites

The Kruskal-Wallis test yielded significant results ($\alpha = 0.05$) for the lifeform (<0.001), leaf growth ($p = 0.05$) and structure ($p = 0.01$) functional categories (Table 6). This indicates significant variation between functional traits of individuals at the two land use sites. The large Chi-square (χ^2) value associated with the lifeform group indicates that there is large variation between the agricultural and non-agricultural individuals. Leaf growth has a χ^2 of 3.80, which is ten times less than that of the lifeform group, indicating that although significant, the variation in leaf growth traits between sub-sites is considerably lower (Table 6). The χ^2 of structure is 6.25, suggesting that leaf structure is more variable between sub-sites than leaf growth, but less variable than lifeform (Table 6).

Table 6. Results of the Kruskal-Wallis test evaluating the differences between agricultural and non-agricultural leaf samples.

	Chi-Square (χ^2)	p-Value
Lifeform	38.24	<0.001
Leaf Growth	3.80	0.05
Leaf Structure	6.25	0.01

Due to the nature of the Kruskal-Wallis test, it is not possible to identify how each functional trait category differs between land-use from this analysis alone. To determine this, the statistical mean of the functional trait with the highest loading in each of the functional trait categories must be examined. Table 7 shows that DBH, chlorophyll ab mass and LMA are lower in the meadow sub-site compared to the agricultural sub-site.

Table 7. Statistical means of the trait with the largest loading value in the PC1 of each functional trait category for the agricultural and meadow sub-sites of the CGOP.

Trait	Functional Trait Category	Agricultural (n = 5)	Meadow (n = 30)
DBH (cm)	Lifeform	125.14	32.12
Chlorophyll ab mass (mmol/m ²)	Leaf growth	9463.48	7207.43
LMA (g/m ²)	Leaf Structure	170.85	134.74

3.4.2. PCA 2: ITV in Spatial Relation to Anthropogenic Landscape Modifications

A significant correlation was identified between the PC1 loadings of lifeform and leaf growth functional categories and the distance to both the CDF and road anthropogenic landscape modifications

using Kendall's Tau (Table 8). The leaf structure functional category showed a suggestive relationship association with the CDF anthropogenic landscape modification, but not the road (Table 8a,b). Leaf growth's relationship with CDF is the strongest in comparison with lifeform, while leaf structure has a suggestive rather than significant relationship (Table 8a). When associated with the road, lifeform has a less significant relationship compared to leaf growth, with leaf structure's relationship being insignificant (Table 8b). For both the road and CDF anthropogenic landscape modifications the leaf growth functional category has the largest significant association (Table 8a,b).

Table 8. Associations between the functional traits with the largest loading in PC1 of each functional trait category and the distance to CDF and road for all non-agricultural *Q. garryana* individuals ($n = 30$). The dataset of each trait is arranged based on the individual's distance to the (a) CDF and (b) road anthropogenic landscape modifications, starting at 0m. Quartiles are the median value of the first quartile (25%), the median (50%) and the third quartile (75%) of each dataset. Significance was calculated using Kendall's Tau.

a) Distance from CDF (m)				
Trait	First Quartile (25%)	Median Value (50%)	Third Quartile (75%)	p-value
DBH (cm)	30.50	27.70	22.0	<0.001
Chlorophyll ab mass (ng/mg)	6674.25	6777.98	7035.07	<0.001
LMA (g/m ²)	135.14	131.35	130.38	0.069
b) Distance from Road (m)				
Trait	First Quartile (25%)	Median Value (50%)	Third Quartile (75%)	p-value
DBH (cm)	22.00	27.70	29.00	0.004
Chlorophyll ab mass (ng/mg)	7608.99	7208.69	7028.97	0.002
LMA (g/m ²)	131.22	131.35	131.48	0.243

4. Discussion

4.1. Remote Sensing of Intraspecific Trait Variation

The accurate use of remote sensing technologies to estimate plant functional traits has been well documented [4,10,22]. Physiological traits, such as leaf pigment content, can be derived from spectroscopy, while morphological traits, such as plant height, can be measured from LiDAR [22,52]. This paper uses a combination of these two methodologies to derive 14 functional traits from 40 individual trees of the same species. Spectroscopy at the leaf level provided the spectral information required to apply PLSR coefficients acquired from open-source, pre-validated models (Table 1) [51]. The functional trait estimations resulting from this analysis were within the acceptable range of the validated model, demonstrating that validated PLSR models are transferable (Table 3) [51]. This is significant as the cost and time required to collect and validate PLSR models is considerable.

The LiDAR data used to estimate morphological functional traits in this study was obtained freely from the Municipality of North Cowichan. With a density of 16 points/m², this dataset provides excellent representation of tree crowns and allows for the accurate derivation of traits like tree height [52]. Due to the open-source nature of this LiDAR data considerable time and cost were avoided when compared to measuring these attributes in the field.

The efficiency of acquiring remotely sensed and open-source data makes it possible to derive accurate functional traits within weeks of collecting leaf level spectral information. This reality means that land managers could have access to quantifiable information about functional traits across their site within a timeframe appropriate to undertake meaningful action. To make use of this potential, the focus of our research was on a relatively underrepresented area of functional trait analysis: intraspecific

variation. This research demonstrates the ability of remotely sensed technologies to enable the analysis of intraspecific functional trait variation with respect to anthropogenic landscape modifications and identify the main drivers of functional diversity across the site.

4.2. Spectral Separation

Visual examination of the mean spectral signatures of the meadow, agriculture and sample populations indicates that the agricultural and meadow trees vary in multiple areas of the electromagnetic spectrum between 350 nm and 2500 nm. Separation between the signatures in Figure 2a from 500 nm to 600 nm suggests that there are differences in the pigments associated with leaf growth, such as chlorophyll and anthocyanin, between individuals living in the two land use types [10,49,53]. Variation in structural components, such as cellulose or lignin, and possibly proteins, is suggested by differences in the reflectance values between 1400nm and 1700nm in Figure 2b (10,49,53). This simple examination of the mean spectral signatures from two sub-population in relation to the mean signature of the sample population supports further examination of the intraspecific variation of *Q. garryana* through the two hypotheses of this study. The use of leaf based spectroscopy and derivation of traits is replicable and can be applied across species, enabling an efficient exploration of variance for managers interested in evaluating the ITV in a species of concern.

4.3. Model Performance

In order to derive traits, we utilized a large North American database of plant spectral, pigment and chemical traits, including samples from a range of Oaks, the key genus in this study. These coefficients are derived from PLSR models with validation R^2 ranged from 0.30–0.95, while validation nRMSE ranged from 4.6–19.15% [50,51]. Only $\delta^{15}\text{N}$ was removed from the analysis due to low validation R^2 (<0.30). The ten traits determined through PLSR demonstrated reasonable mean values and standard deviations (Table 3). By using this model, this study demonstrates that a multi-species, pre-validated dataset can be used to explore intraspecific variation within a single population. This is important as it suggests a single, multi-genera dataset can be used to estimate plant functional traits within a population and across a variety of landscapes.

4.4. Functional Trait Categories

The results of both PCA 1 and PCA 2 determined that the greatest variation in lifeform, leaf growth and leaf structural functional trait categories was found in each of their PC1s, with DBH, chlorophyll ab mass and LMA having the highest influence respectively. The PC1 of each functional category explained over 68% of the variation between samples, so it was used exclusively in the statistical analysis. However, it is also possible to evaluate the presence of ITV for a desired trait by using the values of the principal component in which said trait explains the highest proportion of the variance. This could prove useful for managing the drivers of a specific trait across a landscape.

4.5. Functional Trait Variation due to Land Use

Results support the hypothesis that functional trait variation exists in *Q. garryana* of the same population experiencing different land use pressures. These results are consistent with other studies exploring intraspecific variation through spectroscopy [18,53]. The Kruskal-Wallis rank sums test found a significant level of variation when comparing each of the functional trait groups of individuals persisting in an agricultural environment with those living throughout the rest of the CGOP. Specifically, traits related to lifeform ($p = <0.001$) and leaf structure ($p = 0.01$), such as DBH and LMA respectively, had higher values in the agricultural sub-site, while individuals from the meadow had higher values for leaf growth ($p = 0.05$) (Figure 3). Increased LMA values for individuals in the agricultural sub-site suggests that these trees are growing thicker leaves to increase their resistance to stress, such as agro-centric management practices including the removal of competitors, the addition of fertilizers and consistent watering.

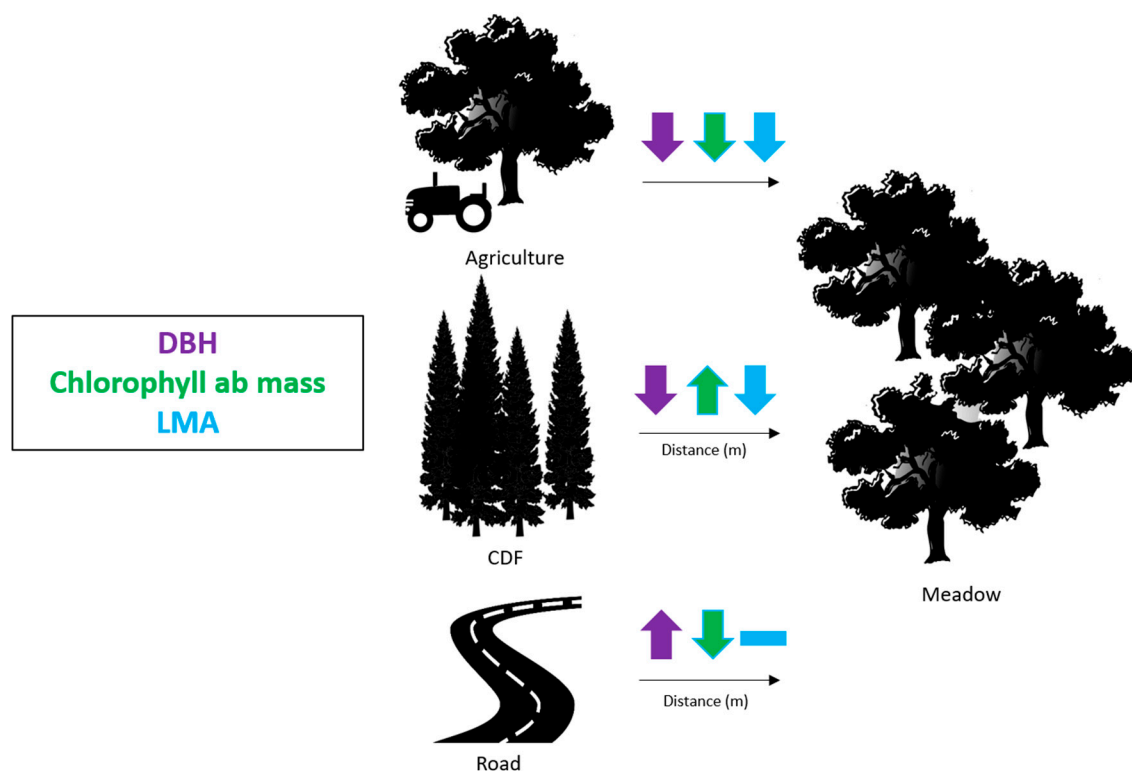


Figure 3. Schematic depicting the relative increase or decrease in values for the traits with the highest PCA loadings from lifeform (DBH, purple), leaf growth (Chlorophyll ab mass, green) and leaf structure (LMA, blue) functional categories in relation to each anthropogenic landscape modification. Arrows are arranged in accordance with Table 4 for the agricultural vs. meadow analysis and on the Tau values in Table 8 for the CDF and road analyses. Coastal Douglas-fir forest (CDF) and Road values are measured on a continuous scale of distance (m) starting at the anthropogenic landscape modification border (0 m) and ending at 248 m and 273 m respectively. Values associated with the Agriculture anthropogenic landscape modification are Boolean.

Lifeform functional traits proved to be the most variant between sites ($p < 0.001$, χ^2 of 38.24), with mean DBH being significantly lower in the meadow (32.12 cm) compared to the agricultural (125.14 cm) sub-site (Table 7). Management practices such as the removal of competitors could be a possible reason for this. These findings suggest that it is important to acknowledge the impacts of land use change on the functional strategies related to overall tree size and shape. The leaf growth functional trait category also showed significant differences between the agricultural and meadow sub-sites ($p = 0.05$, $\chi^2 = 3.80$), with *Q. garryana* individuals in the meadow (7207.43 ng/mg) exhibiting lower chlorophyll ab mass values than those in the agricultural (9463.48 ng/mg) sub-site (Table 7). These variations could be due to the consistent addition of fertilizers to the soil over the last century. ITV related to leaf structure was also proven significant ($p < 0.001$, $\chi^2 = 6.25$), with LMA in the agricultural sub-site (170.85 g/m²) being larger than LMA in the meadow (134.74 g/m²) sub-site. This suggests that functional traits such as LMA can vary based on land use techniques, recognizing that LMA is indicative of overall plant health and a leaf's investment in photosynthesis versus longevity [31].

4.6. Functional Trait Variation in Relation to Landscape Modifications

Significant correlations of ITV were found with respect to the spatial relationship of *Q. garryana* individuals and both CDF and road anthropogenic landscape modifications, supporting the second hypothesis of this study (Table 8a,b). Kendall's tau rank sum test also found that functional trait categories are affected differently by each of the anthropogenic landscape modifications and suggests

that each modification invokes unique environmental changes. This supports the findings of other studies that anthropogenic landscape features can alter the functional strategies of plants [2].

Lifeform traits changed significantly depending on an individual's distance to the CDF ($p = <0.001$) (Table 8a). DBH, for example, decreased as distance to the CDF increased, suggesting that individuals are shifting their lifeform strategies based on the association with this competing ecosystem (Figure 3). The growth functional trait category ($p < 0.001$) also shows a significant relationship to the distance of an individual *Q. garryana* from the CDF and suggests that leaf chlorophyll increases as distance from the CDF increases. Correlation between the leaf structure ($p = 0.069$) functional trait category and the CDF is suggestive, implying that *Q. garryana* individuals may also change their leaf structure based on proximity to competitors, with leaf structural values associated with stress tolerance decreasing with distance (Figure 3) [39,40]. This is interesting as leaf structure is related to a leaf's defensive capabilities and longevity [13,54,55]. These findings suggest that individuals closer to the CDF have a higher stress tolerance, which could be caused by a more favorable microclimate compared to individuals living in the open meadow.

Similar correlations were found between the functional trait category of each sample and their distance to the main road (Table 8b), supporting the hypotheses that intraspecific functional trait variation exists in relation to the spatial distribution of various anthropogenic landscape modifications. Lifeform ($p = 0.004$) values decreased as distance from the road increased, while leaf growth ($p = 0.002$) values increased (Figure 3). This could be due to changes in the nutrient composition of the soil closer to roads, with runoff and de-icing agents like NaCl affecting leaf growth [42,43]. The relationship between the leaf structure functional trait category ($p = 0.243$) and the road anthropogenic landscape modification was found to be insignificant. The associations between the three functional trait categories and the road anthropogenic landscape modification was inverse to the relationship of those groups with the CDF. This suggests that the influence of the road and CDF affect the lifeform, leaf growth and leaf structure strategies of *Q. garryana* in different manners. If this is the case, managers should deploy mitigation efforts for each anthropogenic influence separately, such as salt loading in the winter or excess ground water due to runoff from the road, rather than prescribing a site-wide plan [42]. The presence of significant intraspecific variation also suggests that *Quercus* trees should be considered individually when using functional traits to determine biodiversity metrics as the values of many traits vary in association with different anthropogenic landscape modifications.

In a biodiversity management context, where this project has key implications, our results imply that functional traits of *Q. garryana* should be considered on an individual-plant basis. When mapping functional traits across a site, for example, intraspecific variations among trees may provide insight relating to the relative functional diversity of one sub-site in relation to another. This information could be used to determine which treatment is necessary at specific locations as well as provide general, site-level context to other biodiversity metrics such as species diversity and species composition across sub-sites.

4.7. Future Directions

This research advances the use of image spectroscopy for conservation science. First, as universal ITV approaches are becoming more prevalent, through the use of standardized equations, it is always important to ensure the predictions of traits are valid and accurate. Ongoing research to develop models between measured traits and spectra thus remains an ongoing concern.

Importantly, by utilizing advanced remote sensing techniques we are able to extend these observations in both time and space to make the results more relevant in a conservation framework. Acquiring data from a sensor on an unmanned aerial vehicle (UAV), as opposed to ground-based measurements, would allow more species to be sampled over a broader landscape. The collection of imaging spectroscopy from UAVs would also enable the hypotheses presented in this study to be tested at a larger spatial scale, enhancing the context in which results can be interpreted. This would be

the most time and cost effective image acquisition protocol, allowing agencies to assess species traits over time and across scales relevant to management goals.

This research lays the ground work for space based applications. With the imminent launch of EnMAP (Environmental Mapping and Analysis Program) by the German Aerospace Center and the potential for future missions from other nations, imaging spectroscopy from space offers tantalizing possibilities. While spatial resolutions will not allow individual canopy traits to be assessed at the individual crown scale, forest stand or community traits could be, allowing these hypothesis to be tested over larger spatial extents and routinely over time.

Examination of these differences at a larger spatial resolution would also be beneficial to this study. This would enhance the timeliness of similar studies by improving the temporal resolution of data collection and enable rapid assessment of plant functional traits. The ability of managers to quickly and accurately identify changes in plant health related to anthropogenic landscape modifications, such as functional traits is key to improving on the research presented in this study.

5. Conclusions

Overall, the results presented in this study support the hypotheses that intraspecific functional trait variation exists within a population experiencing different land uses and in relation to various anthropogenic landscape modifications. This study also successfully begins to bridge the knowledge gap between intraspecific functional trait variations at the landscape- and site-levels. The methodology presented in this research provides a simple, replicable and relatively inexpensive baseline on which future studies examining functional trait variation can expand. For example, the coefficients used to estimate each of the leaf growth and leaf structure functional traits were determined by conducting PLSR on a spectral database containing various different genera, rather than *Quercus* alone. This variety suggests that examination of intraspecific trait variation is possible for the other genera, or species, represented in the database. For many land managers, this could improve efficiency as they would not be required to create and validate their own spectral datasets.

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