Title: Species abundance distributions should underpin ordinal cover-abundance

transformations

Running Title: Ecological analysis using global plant data

Abstract

The cover and abundance of individual plant species have been recorded on ordinal

scales for millions of plots world-wide. Many ecological questions can be addressed

using these data. However ordinal cover data may need to be transformed to a

quantitative form (0 to 100%), especially when scrutinising summed cover of multiple

species. Traditional approaches to transforming ordinal data often assume that data

are symmetrically distributed. However, skewed abundance patterns are ubiquitous

in plant community ecology. A failure to account for this skew will bias plant cover

estimates, especially when cover of multiple species are summed. The questions

this paper addresses are (i) how can we estimate transformation values for ordinal

data that accounts for the underlying right-skewed distribution of plant cover; (ii) do

different plant groups require different transformations and (iii) how do our

transformations compare to other commonly used transformations within the context

of exploring the aggregate properties of vegetation? Using a continuous cover

dataset, each occurrence record was mapped to its commensurate ordinal value, in

this case, the ubiquitous Braun-Blanquet cover-abundance (BBCA) scale. We fitted a

Bayesian hierarchical beta regression to estimate the predicted mean (PM) cover of

each of six plant growth forms within different ordinal classes. We illustrate our

method using a case study of 2 809 plots containing 95 812 occurrence records with

visual estimates of cover for 3 967 species. We compare the model derived

estimates to other commonly used transformations. Our model found that PM

estimates differed by growth form and that previous methods overestimated cover,

especially of smaller growth forms such as forbs and grasses. Our approach reduced

the cumulative compounding of errors when transformed cover data were used to

explore the aggregate properties of vegetation and was robust when validated

against an independent dataset. By accounting for the right-skewed distribution of

cover data, our alternate approach for estimating transformation values can be

extended to other ordinal scales. A more robust approach to transforming floristic

data and aggregating cover estimates can strengthen ecological analyses to support

biodiversity conservation and management.

Keywords: aggregated**,** beta regression; Braun-Blanquet; growth form; midpoint;

ordinal transformation; species abundance distribution; sPlot; summed foliage cover;

VegBank; vegetation cover.

Abbreviations: Braun-Blanquet cover-abundance - BBCA

Introduction

Field-based assessment of the cover and abundance of individual plant species is complex. Observers making on-ground visual estimates of plant cover need to account for, and assess, foliage cover of different densities, dimensions, shapes and structures across multiple species, growth forms and strata. So too, counting cryptic, clonal, or copious numbers of plants can be complicated. Owing to this complexity, vast numbers of floristic plots across many continents have been surveyed using ordinal scales (Schaminée et al. 2009; Dengler et al. 2011; Chytrý et al. 2016). Whilst, in Braun-Blanquet (1932) originally described an abundance-dominance scale, the practical, on-ground application of this scale is to assess plant cover, and where cover is less than 5%, abundance is also assessed. The Braun-Blanquet cover-abundance (BBCA) scale is perhaps the most common ordinal scale used in plant ecology. For example, within the vegetation plot database sPlot v2.1 (www.idiv.de/splot), more than 745 000 plots (66%) have recorded plant occurrence using Braun-Blanquet cover-abundance (sPlot extract supplied by Borja Jiménez-Alfaro, 19th September 2017). This volume of data is testament that ground-based visual assessments of cover-abundance using ordinal scales provide a cost-effective, rapid and non-destructive approach to gathering the data needed to summarise the composition and structure of plant communities. These data represent a wealth of investment in field effort and have supported major advances in vegetation classification, mapping and distribution modelling.

The ever-growing access to global vegetation plot databases (Dengler et al. 2011; Schaminée et al. 2011) has opened pan-continental opportunities to explore many uses of floristic data. Some ecological questions may best be addressed using aggregate properties of vegetation, such as the summed total foliage cover within a

plot or across strata, the total summed cover or abundance of exotic or invasive

species, or the relative cover or abundance of plants within different functional,

taxonomic or growth form groups. Summing cover to derive aggregate properties of

floristic data have a multitude of uses in ecology including assessing presence and

diversity of faunal habitat, as covariates in species' distribution models (SDMs), for

assessing the spatial and temporal status of ecosystem baselines, predicting the

effects of shifts in climate, land use and land cover, or measuring site-scaled

responses to disturbance (e.g. Scholes & Biggs 2005; McElhinny et al. 2006; Pereira

et al. 2010). Aggregate properties of vegetation data are particularly relevant to

exploring ecological questions concerning the patterns, processes and prognoses at

a range of spatial scales in contemporary and predicted future landscapes.

There are many applications where ordinal data have been used successfully, such

as ordination, classification, modelling or mapping of vegetation communities (e.g.

Podani 2005; Podani 2006; Lyons et al. 2016) and for modelling the cover of single

species (e.g. Damgaard 2014; Irvine et al. 2016). However, ordinal scaled cover

observations of individual species cannot be summed (Guisan & Harrell 2000;

Podani 2006; Chen et al. 2008b) and need to be transformed into a continuous scale

prior to aggregating.

Approaches to transforming Braun-Blanquet cover-abundance (BBCA) ordinal data

have been proposed by Tüxen and Ellenberg (1937) and Braun-Blanquet (1964)

(see Table 3 in van der Maarel 1979). In addition, van der Maarel (1979) proposed

the ordinal transform value (OTV) with different scale adjustments, as a solution for

converting ordinal data to percentage cover values. All these methods tend to

transform data to the approximate midpoint of the ordinal class range for

observations of cover greater than 5%. For classes with cover less than 5%, the

transformation values appear arbitrary and differ considerably (Table 1 columns 4–

6).

Transforming data to the approximate midpoint of the class ranges assumes that

data are symmetrically distributed within each class. Yet, patterns in plant

abundance including density, biomass (Chiarucci et al. 1999; Morlon et al. 2009),

frequency (Chiarucci et al. 1999), percentage cover (Damgaard 2009), size, energy

use and productivity (Whittaker 1965) have all been shown to have a right-skewed

distribution; skewed species abundance distributions occur in every known multi-

species community (McGill et al. 2007). Midpoint transformations are inflexible to the

underlying distribution of cover data and assume that the distribution does not vary

across species, groups of plant entities (such as growth forms, life forms, functional

or taxonomic groups), vegetation types or biomes. Due to the prevalence of right-

skewed distribution, we predict that midpoint transformations overestimate cover and

the compounding of these errors will result in gross overestimation of summed cover

for aggregated properties.

Here we develop a flexible approach to estimate cover transformations for ordinal

scaled data that can then be used to provide accurate estimates of summed

vegetation cover. The method we describe is applicable to data in any ordinal scale,

can be extended to allow for differences in vegetation type or among biomes and

can accommodate alternative aggregate properties of plant data such as growth

forms, life forms, functional or taxonomic groups. To demonstrate the potential

applicability of our approach we build and then validate the model using two

separate and independent datasets.

Given that diverse architectures and spatial arrangements of foliage lead to varied

patterns of plant cover (Damgaard 2013), we also predict that different plant growth

forms will require different transformation values. Growth forms are practical and

observable entities that can inform site-based assessment and monitoring, are

recognizable from remotely-sensed imagery and are used to report on broad-scale

biodiversity assessment or baselines (Pereira et al. 2013) with which we can

measure change in cover (Pettorelli et al. 2014; Abelleira Martínez et al. 2016).

Materials and Methods

We outline the key steps required to estimate transformation values within ordinal

classes for different plant groups. A pre-requisite for our method is cover data that

have been collected on a continuous cover scale, ideally sourced from the same

study region and vegetation types as the ordinal cover data. To prepare the input

data for the model, ordinal values need to be mapped, *a posteriori*, to this continuous

cover data as an intermediary variable (Figure 1, Step 1). Models, with a beta

distribution, are then used to predict the mean cover of each plant group within each

ordinal cover class. This predicted mean cover is the transformation value (Figure 1,

Step 2). Using a case study, we explore summed cover estimates for different plant

groups and evaluate the performance of the ordinal cover transformations. We

compare our transformation to existing approaches in the context of summed cover

for plant groups (Figure 1, Step 3). We evaluate the robustness of our predicted

mean transformations on an independent dataset (Figure 1, Step 4).

Estimate mean cover using parameters of a beta distribution

We used a generalised linear mixed model (GLMM) with a beta distribution to derive

estimates of the mean vegetation cover, within an ordinal class, given a plant's

growth form and random variation owing to plot identity. Individual species cover are

continuous proportional estimates, and once suitably transformed, fall within the

known range ($0<y<1$). Linear regression with a normal distribution is inappropriate

for the analysis of proportions, such as percent plant cover, because data often

violate assumptions such as normality and homogeneity of errors and furthermore

fitted values can fall outside of the range [0,1] (Ferrari & Cribari-Neto 2004). A

common approach to address these problems is to apply arcsine or logit

transformations to the response variable, prior to regression (Warton & Hui 2011),

although the results can be difficult to interpret (Ferrari & Cribari-Neto 2004).

Numerous authors have instead demonstrated that percent plant cover are more

appropriately analysed by assuming that cover approximates a two-parameter beta

distribution (Ferrari & Cribari-Neto 2004; Chen et al. 2008a; Cribari-Neto & Zeileis

2010; Herpigny & Gosselin 2015). Beta distributions are attractive because fitted

values are constrained between the interval $0<y<1$ and they can accommodate

asymmetrical distributions with left- or right-skew. This flexibility makes beta

distributions highly suitable for modelling diverse and often asymmetrical plant cover

data (Cribari-Neto & Zeileis 2010).

We present a Bayesian GLMM with a logit link to estimate the parameters of the beta

distribution and allowed these parameters to vary among ordinal classes and plant

growth forms. Estimates of these parameters were used to derive the predicted

mean (PM) for each plant growth form in each ordinal class.

The proportional vegetation cover is given by the two-parameter beta distribution;

Proportion$_{ij}$ ~ Beta (a_{ij}, b_{ij})

Where a_{ij} and b_{ij} are shape parameters for species j in plot *i, and i = 1,…n plots.* The

shape parameters are further defined as

$a_{ij} = \theta \times \pi_{ij}$

$b_{ij} = \theta \times (1 - \pi_{ij})$

where θ allows for potential overdispersion to be incorporated in the model (Zuur et

al. 2013)*.*

π_{ij} is modelled with a logit link

logit (π_{ij}) = η_{ij}

The model consists of regression parameters (β) for each ordinal class, plant growth

form and their interactions, plot level random intercepts and variance (z_i):

$\eta_{ij} = X_{ij} \times \beta + z_i$

$z_i \sim N (0, \delta^2_{plot})$

Where z_i is a random intercept for plot, X_{ij} are the matrix of all covariates (ordinal

classes and their interaction with plant growth form) and *β* are the regression

parameters for each covariate. That is, for each ordinal class 1…6, separate *β* were

estimated for each plant growth form. For a simplified example with two growth forms

and two ordinal classes this can also be expressed as:

$\eta_{ij} = \beta_0 + \beta_1 \times fOrdinalClass_{ij} + \beta_2 \times fGrowthForm_{ij} + \beta_3 fOrdinalClass_{ij} \times fGrowthForm_{ij} +$

z_i

Where β_0 = predicted value of logit transformed cover if species *j* belongs to the

"reference" growth form and its' value in plot *i* has the "reference" level ordinal cover-

abundance class.

β_1 = departure of the predicted value for species j from β_0 if the observation is of

another ordinal cover-abundance class.

β_2 = departure of predicted value from β_0 if species j belongs to another growth form.

β_3 = departure of predicted value from $\beta_0 + \beta_1 + \beta_2$ when neither growth form nor

ordinal cover-abundance class are of the reference level.

In this example, $fOrdinalClass_{ij}$ and $fGrowthForm_{ij}$ are binary dummy variables

coding growth form and cover-abundance scale categories, thus Xij is a vector

containing values for these dummy variables (including their products) for species j

in plot i.

We included plot as a random intercept because although we assumed each plot

should follow the characteristic skewed species abundance curve, we expected

variation among plots and hence differences in the average cover of any given

ordinal class and plant growth form.

This basic model structure can be easily expanded to accommodate other possible

sources of variation, such as among vegetation types or owing to the richness of

plant species within a plot. In this case study, we decided not to include additional

covariates to minimise computational demands and simplify model interpretation and

operational complexity.

The model was fit via Markov chain Monte Carlo optimization in JAGS (http://mcmc-

jags.sourceforge.net) via the R2jags package (Su & Yajima 2015) within R 3.5.0 (R

Core Team 2018). Posterior parameter estimates and back-transformed predicted

means were derived from 3 chains, with a burn-in of 3000 iterations, 15 000

subsequent iterations per chain and with a thinning rate of 15. Autocorrelation and

mixing were visually inspected. The interaction models were compared to additive

models using Deviance Information Criteria. Appendix S1 contains R code for our

models.

Case study – New South Wales, Australia

We illustrate our model with a case study where we have used 1-6 BBCA as our

ordinal scale and grouped plants into six growth form categories. Following is a brief

description of how we prepared the case study dataset to build our model. We note

that randomly generated data from an appropriate beta distribution (for similar

example see Damgaard 2014) could also be used to demonstrate our approach.

However, we chose to use a large archival dataset from a range of bioclimatic

regions and vegetation types to demonstrate that, despite the underlying variation,

our approach still led to robust estimates of summed cover.

1. Preparation of observed percentage cover dataset

To demonstrate our modelled approach, we sourced case study data from archival

quantitative floristic data that met three considerations: (i) each species record

included a visual estimate of foliage cover on a continuous scale from 0.1% to 100%

and a count of abundance where cover was less than 5%; (ii) in each plot, full

species inventories were recorded from a fixed-area (400 m^2) and (iii) sites covered

a wide geographic distribution (Appendix S2—Figure 1) and included a wide range of

vegetation types with different structural complexity including rainforests, forests,

woodlands, shrublands, grasslands and wetlands (Keith 2004). A total of 2809 geo-

referenced plots containing 95 812 occurrence records with visual estimates of cover

for 3967 species met these criteria and were exported from the NSW BioNet Atlas

database (www.bionet.nsw.gov.au).

Analysis of the empirical cover distribution

To confirm our assumption of right-skewed distribution of cover data we plotted our

data and used the 'skewness' function in the e1071 package (Meyer et al. 2017)

within R 3.3.3 (R Core Team 2018) to calculate the adjusted Fisher-Pearson

skewness coefficient (G_1) (Joanes & Gill 1998) for the whole distribution, and for

distributions within each BBCA class. Skewness is a diagnostic tool usually used to

test the symmetry of the data distribution. Here, we interpret skewness coefficients

as being strongly and positively skewed when the G_1 coefficient is greater than 0.5

(Bulmer 1979; Doane & Seward 2011).

2. Preparation of plant group entities

All taxa were allocated to one of six growth form categories: tree, shrub, grass and

grass-like (hereafter referred to as grass), forb, fern and other (remaining growth

forms) (Oliver et al. submitted). For each growth form in each plot, total cover was

estimated by summing the observed quantitative estimates of cover and the

estimates of cover derived from the transformations of the ordinal data.

3. Allocating an intermediary variable

We created an intermediary variable by matching each quantitative estimate of cover

for every floristic record (n = 95 812) to its commensurate ordinal value. Any ordinal

scale can be used to partition data, but here we demonstrate our approach by

allocating data to 1–6 BBCA (Table 1). BBCA1 and BBCA2 were assigned based on

their observed foliage cover (<5%) and abundance; where BBCA1 ≤ 10 and BBCA2

> 10 individuals. The pragmatic choice of ten individuals provides an explicit

quantitative abundance threshold between classes BBCA1 and BBCA2. BBCA3–

BBCA6 were assigned based on observed foliage cover (≥5%) (Mueller-Dombois &

Ellenberg 1974). The ordinal dataset created by this process approximates the form

of many data held within vegetation databases.

In our case study dataset, observations of 5% cover were more prevalent than

expected from a typical theoretical beta distribution (Figure 2). This bias was

detected in preliminary model convergence diagnostics and model fit suggested that,

for our case study, it would be preferable to split the data and separately model (i)

BBCA1 and BBCA2 bounded between 0 and less than 5% cover and (ii) BBCA3 to

BBCA6 bounded between 5% and 100% cover inclusive. To ensure the response

variable was bounded by 0 and 1, percent cover was transformed using (y-a)/(b-a)

where in (i) a = 0 and b = 5 and in (ii) a = 5 and b = 100 (Cribari-Neto & Zeileis

2010). In the second model, the response variable was further transformed using (y *

(n-1))/n where n = sample size (Cribari-Neto & Zeileis 2010). This split-model

approach may not be necessary for all datasets, especially where data are derived

from less subjective cover methods (e.g. point intercept or pin frame) but is included

here to support the handling of datasets with similar patterns in distribution (see

Appendix S3—Figures S1–S4 for other datasets that appear to show similar pattern).

Evaluation of past and proposed approaches to transforming ordinal data

We transformed each of the 1–6 BBCA records using three different approaches

outlined in Table 1. We then evaluated these past approaches proposed by Tüxen

and Ellenberg (1937), Braun-Blanquet (1964) and van der Maarel (2007) to the PM

estimated from a beta distribution.

For each plot, growth form cover and total cover were calculated by summing the

observed continuous cover estimates (%) and the estimates of cover derived from

the various transformations. Linear regression models with zero-intercept were fitted

to the sum of observed continuous cover data (y) and sum of transformed cover data

(x) cover data in R 3.5.0 (R Core Team 2018). We can justify using a regression

through the origin because we are most interested in comparing the slope of the

regression line to the 1:1 line of best fit to determine if our PM models were over or

underpredicting summed cover. We compared the root mean squared deviation

(RMSD) (see Eq. 1) as an estimate of the deviation of the transformed cover values

from the 1:1 line.

RMSD =

$$\sqrt{\frac{1}{n-1}\sum_{i-1}^{n}(\hat{y}_i - y_i)^2}$$

288   (Eq. 1)

Where $\hat{y}_i$ are the predicted cover values; y_i are the observed cover values and n is

the number of observations.

The RMSD estimate represents the mean deviation of transformed cover values with

respect to the true observed cover values. We also compared estimates of the slope

with lower and upper 95% confidence intervals expecting that robust transformations

would result in a slope = 1 and transformations that overestimate summed cover will

have a slope <1. We include the adjusted coefficient of variation (R^2) to evaluate how

much of the linear variation of observed cover values is explained by the variation of

transformed cover values.

We note that RMSD is useful for evaluating models as it represents an absolute

measure of fit to the 1:1 line and reports the prediction error in the same units as the

data (i.e. summed cover). Whereas adjusted R^2 gives a relative measure of

proportion of total variance that is explained by the model on a scale between 0 and

1.

We validated the PM transformation values on an independent dataset (2 227 sites

with 51 497 observations) from West Virginia Natural Heritage Program (Vanderhorst

et al. 2012) accessed from VegBank (Peet et al. 2013 accessed 28th Aug 2018).

Whilst VegBank has a primary role for enabling the vegetation classification, large

volumes of individual floristic observations are available for ecoinformatic synthesis

and analysis. Owing to the ease of access and completeness of datasets stored in

VegBank we were able to validate our model estimates on a geographically distinct

dataset containing cover estimates of plants from entirely different vegetation

communities. Details outlining the data preparation are included in Appendix S6.

Results

The empirical cover distribution

The source continuous cover data were right-skewed and dominated by low cover—

85% of observations were between 0.1 and 4%, and 60% of these observations

were of cover less than 1% (Figure 2). Data were heavily right-skewed for the whole

distribution (G_1 = 5.62) and right-skewed within five of the six BBCA classes (BBCA1

G_1 = 2.64, BBCA2 G_1 = 1.57, BBCA3 G_1 = 1.04, BBCA4 G_1 = 0.61 and BBCA6 G_1 =

0.95). Only BBCA5 had a skewness coefficient less than 0.5 (G_1 = 0.36). We also

note potential observer bias for 5% cover. These patterns are similar to other visually

estimated floristic cover data from other archived datasets (see Appendix S3—

Figures 1-4).

Estimate mean cover using parameters of a beta distribution

Table 1 (columns 7 and 8) shows the predicted mean transformations and their lower

2.5% and upper 97.5% credible interval for each ordinal class, independent of

growth forms. The most marked differences are noted in BBCA2 and BBCA3, where

the predicted means are well below the previous approaches. The predicted mean

for class BBCA6 is lower than the midpoint but was derived from relatively few

observations (n = 138).

Table 1: Class divisions for the 1–6 Braun-Blanquet ordinal cover-abundance (BBCA) scale (columns 1–3), previous proposals for transforming them to percentage cover (columns 4–6), and proposed transforms (independent of growth form) based on estimating the predicted mean (PM) from a beta distribution of observed quantitative cover data and the lower 2.5% and upper 97.5% credible intervals. Number of observations (n) for BBCA1 (n = 54 811); BBCA2 (n = 26 968); BBCA3 (n = 11 946); BBCA4 (n = 1583); BBCA5 (n = 366) and BBCA6 (n = 138).

Column 1	2	3	4	5	6	7	8
BBCA Class	Range of cover (%)	Qualitative abundance terms	Tüxen & Ellenberg (1937) [1]	Braun-Blanquet (1964) [1]	van der Maarel (2007) [2]	PM	Credible interval
1	<5	e.g. present, few, rare, erratic, occasional, uncommon	0.1	0.1	1	0.49	0.48–0.51
2	<5	e.g. common, abundant, many, several	2.5	5	2	0.74	0.72–0.76
3	5–25		15	17.5	17.5	8.95	8.84–9.07
4	26–50		37.5	37.5	35	38.77	37.97–39.57
5	51–75		62.5	62.5	70	62.43	60.69–64.13
6	76–100		87.5	87.5	140	81.24	79.10–83.26

[1] adapted from van der Maarel (1979).
[2] ordinal transform values (OTV) using 1.415 weighting factor (van der Maarel 2007).
Column 3 shows some of the qualitative descriptors used by field surveyors to divide observations between BBCA1 and BBCA2.

Table 2: Proposed transformation values, tailored to different growth forms, based on estimates of the predicted mean (PM) from a beta distribution of observed data. Lower 2.5% and upper 97.5% credible intervals (CI) are shown in square brackets; n = number of individual observations for each Braun-Blanquet cover-abundance (BBCA) class.

		Growth Form					
		Tree	Shrub	Grass	Forb	Fern	Other
BBCA1	PM	0.78	0.52	0.45	0.42	0.43	0.45
	CI	(0.76 – 0.80)	(0.51 – 0.53)	[0.44 – 0.46]	[0.41 – 0.44]	[0.41 – 0.45]	[0.44 – 0.46]
	n	6465	13366	8731	16981	1863	7405
BBCA2	PM	1.78	0.96	0.82	0.58	0.75	0.74
	CI	[1.7 – 1.86]	[0.93 – 0.99]	[0.8 – 0.84]	[0.57 – 0.6]	[0.72 – 0.78]	[0.71 – 0.77]
	n	441	2821	8488	12065	1324	1829
BBCA3	PM	9.53	8.60	8.86	8.02	8.80	8.48
	CI	[9.38 – 9.7]	[8.42 – 8.76]	[8.71 – 9.01]	[7.81 – 8.24]	[8.46 – 9.19]	[8.23 – 8.74]
	n	4347	2070	3500	893	412	724
BBCA4	PM	38.06	39.18	39.32	37.86	39.30	38.72
	CI	[36.74 – 39.38]	[36.98 – 41.26]	[38.05 – 40.59]	[34.01 – 41.81]	[34.92 – 43.72]	[34.76 – 42.64]
	n	582	217	599	68	52	65
BBCA5	PM	61.71	63.15	62.67	62.80	62.01	62.11
	CI	[58.15 – 65.16]	[58.23 – 67.78]	[60.24 – 65.08]	[49.59 – 75.8]	[54.16 – 70.05]	[54.66 – 69.15]
	n	89	48	184	6	18	21
BBCA6	PM	80.80	83.9	80.81	78.87	77.59	80.55
	CI	[76.04 – 85.31]	[78.99 – 88.41]	[77.82 – 83.65]	[64.83 – 90.78]	[55.35 – 93.93]	[70.09 – 89.57]
	n	29	23	73	4	2	7

Evaluation of past and proposed transform values for summed growth form cover

Estimates of the PM suggest that accounting for growth form within each ordinal class

results in more robust summed cover estimates. Credible intervals suggest that in classes

BBCA1 to BBCA3, trees typically have higher mean cover and warrant higher

transformation values (Table 2). Credible intervals also suggest the need for separate

transformation values for shrubs in BBCA1 and BBCA2 and a lower value for forbs in

BBCA2 and BBCA3 (Table 2).

When these growth form specific transformations were evaluated using the summed cover

estimates RMSD did not exceed 9.50 (trees) (Figure 3 and Appendix S4—Table 1). In

contrast, estimates based on past transformations frequently resulted in RMSD exceeding

10. Slope ranged from 0.91 (forbs) to 1.05 (others), whereas past transformations slopes

were <0.85, suggesting considerable overestimation of summed cover (see Appendix S4—

Table 1 and Appendix S5—Figures 1-4).

Evaluation of past and proposed transform values for total summed cover

Evaluation of summed total cover revealed that when transformations are tailored to growth

forms, the PM performed better than existing approaches (Figure 3). The PM reduced the

overestimation of total summed cover by up to 4 times. The evaluation of model fit for

summed total cover using past approaches generally revealed a poorer model fit: RMSD

ranged from 41.47–79.37 (PM = 18.21) (see Appendix S4—Table 1) and slope ranged from

0.57 to 0.74 (PM = 1.01) and adjusted R^2 ranged from 0.61-0.96 (PM – 0.97).

Evaluation of the growth form specific PM transformation on an entirely independent

validation dataset from West Virginia Natural Heritage Program (Vanderhorst et al. 2012)

show that transformations were robust, although tended to underestimate summed cover of

most growth forms (Appendix 6—Table 1). RMSD ranged between 1.59 (others) and14.97

(trees); slope ranged between 1 (others) and 1.12 (forbs) and adjusted R^2 were high and

ranged between 0.97 (trees and shrubs) and 0.93 (others). When compared to the

transformation proposed by Tüxen and Ellenberg (1937), the PM transformation values

were marginally better. Tuxen and Ellenberg (1937) transformation values tended to result

in an overestimate summed cover of all growth forms (Appendix S7—Figures 2a-f); RMSD

was consistently higher than PM transformations for all growth forms; slopes were further

from 1 between 0.76 (forbs) and 0.95 (shrubs) and adjusted R^2 ranged between 0.86

(others) and 0.98 (trees) (Appendix 6—Table 1).

Evaluation of total cover, using the PM transformation values, showed RMSD was less than

that estimated if the transformation was undertaken using estimates of Tüxen and Ellenberg

(20.54 *cf.* 27.01) (Appendix 6—Table 1) and PM transformation values show a slight

underestimation (slope = 1.1; adjusted R^2 = 0.98) when tested on the independent dataset.

Scatter plots showing the relationships between visual estimates of summed cover for all

six growth form groups using the PM model and for Tüxen and Ellenberg (1937)

transformations are provided in Appendix S7—Figures 1a-f and Figures 2a-f.

Discussion

Transforming ordinal data to a quantitative form is common practice in plant ecology and

extends across disciplines including restoration (Fill et al. 2017), classification (Cawsey et

al. 2002; Faber-Langendoen et al. 2007; Wiser & De Cáceres 2013); and for assessing

disturbance (Scott & Kirkpatrick 2008; Knapp & Ritchie 2016). Similarly, universal skewed

patterns in the species abundance distribution are a long standing and well recognised

pattern in ecology (e.g. MacArthur 1960). The data we present here are no exception. Yet

the integration of these two concepts, underpinned by a robust modelling approach has

received little attention, especially in the context of synthesizing information on aggregate

properties of vegetation data. We demonstrate, using two large quantitative independent

datasets that when the underlying right-skewed cover distribution is accounted for, a more

robust set of transformations are generated. Where the aggregate properties of floristic data

are of interest, our method, unlike previous approaches to transformation of ordinal data,

does not overinflate cover.

Where possible, we advocate that others replicate this approach and source continuous

cover data, so that the means within each ordinal class can be estimated accounting for the

underlying distribution. Ideally, the continuous cover datasets will encompass the same

temporal and spatial variation as that of the ordinal data. Notwithstanding these

recommendations for best-practice, we have demonstrated our modelling approach can

produce robust estimates of summed cover using floristic data from geographically distinct

dataset containing observations of entirely unrelated vegetation communities. We expect

the estimates of summed cover would further improve had we used representative data

from that region and vegetation to model specific estimates of the parameters for the beta

distribution. Undoubtedly there will be circumstances where appropriate continuous data

will not be available and the parameters of the beta distribution cannot be estimated for a

specific study or region. In these situations, adopting the PM transformations provided in

Tables 1 and 2 would be preferable to application of ordinal class midpoints. When plant

cover are right-skewed, midpoint transformations will bias and overestimate total cover.

Hierarchical models are useful for handling complex interactions in observational data.

Despite the size of the initial dataset, some plant groups were poorly represented in the

higher cover classes. By appropriately specifying the hierarchical model, estimates for

these combinations could still be obtained, because they draw from the full model structure.

We have identified that different growth forms have different cover distributions. Our

empirical evidence strongly suggests that in plots where there are many small entities from

the same growth form, such as for forbs and grasses, the cumulative cover of that growth

form (when derived from transformations of ordinal data) may amplify and inaccurately

describe the structural complexity of vegetation communities. Identifying and accounting for

these distributions in other grouped entities has the potential to further improve summed

cover estimates.

We also note potential observer bias for cover estimates of 5%. We acknowledge that

visual estimates of cover and counts are subject to inter- and intra-operator error and bias

(see Morrison 2016 for review) and this may account for the data digressions from a

smooth shaped abundance curve. This is no doubt an artefact of observer preference for

regular intervals when estimating cover, rather than a true representation of plant cover. In

our case study analyses, the high frequency of estimates of 5% cover required a split-

model approach where cover data were treated in two separate models. Given this right-

skewed distribution and potential bias among disparate datasets (Appendix S3—Figures 1-

4), we propose the split-model approach may serve wider applications. Simulated beta

distribution data may not be entirely appropriate when using visually-estimated cover data,

but may be useful where other less subjective methods for estimating cover are used (such

as point-intercept methods). Given that visual estimates of cover-abundance are the

assessment protocol for many floristic surveys, our approach offers a way these data can

still be transformed and used with greater confidence, despite the underlying variability and

bias. The approach we outline here can rapidly generate robust and defensible

transformation estimates that are less prone to inflating summed cover estimates.

We envisage that our method may be useful when combined with emerging technologies

such as 3-dimensional LiDAR or radar sensors that can penetrate vegetation canopies and

assess complex structural elements. Furthermore, where large-scale biodiversity

assessments, that rely on terrestrial vegetation as indicators of change, seek to integrate

site observations to validate or train imagery, vegetation cover data collected in an ordinal

scale will be of little benefit. Summation of midpoint transformations have been widely used

in remote-sensing applications, no doubt often resulting in overinflated cover estimates.

While transformations derived from a beta distribution will reduce the problem of over

inflation, summed cover estimates can still exceed 100%. Jennings *et al.* (2009) offer an

approach to rescaling summed cover so that cover estimates do not exceed 100% and their

approach may be useful where site-based data are integrated to inform remote sensing

applications.

Our approach to transforming ordinal estimates of cover using a beta distribution can

extend the application of these data beyond the realm of vegetation classification and can

salvage information from many millions of floristic records. We expect most large

repositories of floristic data will contain cover estimates with multifarious and nuanced

ordinal scales. Here we provide a method that can be applied to floristic data in different

ordinal scales for transforming and integrating datasets with much greater confidence. We

have demonstrated a pan-continental approach to transforming ordinal cover estimates

needed to build robust and accurate aggregated cover estimates. We foresee this approach

supporting the synthesis of multiple datasets containing legacy data collected in different

ordinal scales, especially where the aggregate properties of vegetation cover for different

plant groups are of interest. These transformations and the resultant aggregated properties

of cover data can support a multitude of uses in ecology from site-scaled, to landscape-

scaled and for global applications.

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Author Contribution Statement

• conceived the ideas and designed methodology – ALL

• extracted the case study and validation data – MM

• designed predictive model – JD

• analysed and interpreted the data – ALL

• led the writing of the manuscript – MM

• contributed critically to drafting and revising the manuscript and gave final approval

for publication – ALL

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STEP 1. Identify primary dataset, map continuous cover estimates to an ordinal scale

1. Extract cover (%) and abundance (count) floristic data, from fixed area quadrats

2. Allocate each taxon to plant group e.g. growth form

→ 3. Generate an intermediary variable by allocating each continuous cover observation *a posteriori* to an ordinal scale →

PRIMARY DATASET	continuous cover data			Intermediary variable
Species list	GF	cover(%)	abund.	BBCA
Species 1	tree	25	na	3
Species 2	grass	55	na	5
Species 3	grass	15	na	3
Species 4	shrub	3	16	2
Species 5	forb	35	na	4
Species 6	forb	15	na	3
Species 7	fern	0.1	4	1
Species 8	other	80	na	6
et cetera				

STEP 2. Model patterns of plant cover within ordinal classes

1. Split dataset into model building and validation subsets or (see Step 4.1) source an independent validation dataset

2. Fit GLMM, estimate parameters and derive predicted mean transformation values for each plant group within each ordinal band

→ 3. Allocate transformation (mean cover) values to each intermediary ordinal value →

PRIMARY DATASET	continuous cover data			Intermediary variable	
Species list	GF	cover(%)	abund.	BBCA	PM
Species 1	tree	25	na	3	9.53
Species 2	grass	55	na	5	62.67
Species 3	grass	15	na	3	8.86
Species 4	shrub	3	16	2	0.96
Species 5	forb	35	na	4	37.86
Species 6	forb	15	na	3	8.02
Species 7	fern	0.1	4	1	0.43
Species 8	other	80	na	6	80.55
et cetera					

STEP 3. Test the fit of predicted transformation values for summed growth form cover and total cover

1. Sum cover of each growth form group, at each site, using
a) the original continuous cover data and
b) the transformed ordinal data

2a. Plot summed continuous (y) and summed transformed (x)

2b. Linear regression to estimate fit and RMSD to estimate error and evaluate transformations

→

STEP 4. Validate predicted mean transformation values for summed growth form cover and total cover

1a. Validate model with hold-out subset (see Step 3.1)

OR

1b. Source independent dataset, prepare data as per Step1 & Steps 2.2-2.3

→ 2. Repeat Step 3 →

3 Figure 1: Workflow showing the major elements required to estimate transformation

4 values for ordinal data using continuous cover estimates. Here we use Braun-

5 Blanquet cover-abundance (BBCA) 1-6 scale, although this approach could be

6 extended to any ordinal scale. Note, this flow diagram represents data from one plot,

7 but many plots are needed to obtain robust estimates of mean cover.

Visual estimate of cover with corresponding cover-abundance class

Figure 2: Distribution of visual estimates of cover for 95 812 observations, and their

corresponding Braun-Blanquet cover-abundance (BBCA) class for our case study.

Dashed vertical lines show cut points between each BBCA class. Number of

observations (n) for BBCA1 (n = 54 811); BBCA2 (n = 26 968); BBCA3 (n = 11 946);

BBCA4 (n = 1 583); BBCA5 (n = 366) and BBCA6 (n = 138). Numbers between the

dashed lines show the percentage of each class in the dataset. BBCA1 and BBCA2

(both represent <5% cover) are shown as stacked histograms; BBCA1 (≤ 10

individuals) sits above BBCA2 (> 10 individuals). See Appendix S3—Figures 1-4 for

comparison with other archival datasets.

Figure 3: Results of linear regression with zero-intercept to compare root mean squared deviation (RMSD) and slope for each growth form and for total summed cover under previous transformations compared to the predicted mean transformation. Lower and upper 95% confidence intervals are shown for slope. Vertical dashed lines represent the perfect regression fit where RMDS = 0 and slope = 1. (data table supplied in Appendix S4—Table 1). Number of observations (n) for trees (n = 11 953); shrubs (n = 18 545); grasses (n = 21 575); forbs (n = 30 017); ferns (n = 3671); other (n = 10 051) and total (n = 2809). See Appendix S5 — Figures 1a-f to 4a-f for plots of all growth forms and three previous approaches to transformation proposed by Tüxen and Ellenberg (1937); Braun-Blanquet (1964) and van der Maarel (2007).