Spatial mapping of key plant functional traits in terrestrial ecosystems across China

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Abstract

Trait-based approaches are of increasing concern in predicting vegetation changes and linking ecosystem structures to functions at large scales. However, a critical challenge for such approaches is acquiring spatially continuous plant functional trait ~~distribution~~maps. Here, ~~eight~~ six key plant functional traits were selected ~~to~~as they can reflect plant resource acquisition strategies and ecosystem functions~~represent two-dimensional spectrum of plant form and function~~, including specific leaf area (SLA)~~leaf area (LA)~~, leaf dry matter content (LDMC), leaf N concentration (LNC), leaf P concentration (LPC),~~, plant height, seed mass (SM),~~ leaf area (LA) ~~specific leaf area (SLA)~~ and wood density (WD). A total of ~~52477~~ 34589 in-situ trait measurements of ~~4291~~ 3447 seed plant species were collected from ~~1541~~ 1430 sampling sites in China and were used to generate ~~a~~ spatial plant functional trait ~~plant functional trait dataset~~ maps (~1 km), together with environmental variables and vegetation indices based on two machine learning models (random forest and boosted regression trees). The two models showed a good accuracy in estimating WD, LPC and SLA, with average R^2 values ranging from 0.45 to 0.66. In contrast, both the two models had ~~a~~ weak performance in estimating ~~SM and~~ LDMC, with average R^2 values below 0.25. Meanwhile, LA~~, SM and plant height~~ showed considerable differences between two models in some regions. To obtain the optimal estimates, a weighted average algorithm was further applied to merge the predictions of the two models to derive the final spatial plant functional trait ~~dataset~~maps. ~~The optimal estimates showed that c~~Climatic effects were more important than those of edaphic factors in predicting the spatial distribution of plant functional traits. Estimates of plant functional traits in northeast China and the Qinghai-Tibet Plateau had relatively high uncertainties due to sparse samplings, implying a need of more observations in these regions in future. Our spatial trait ~~dataset~~ maps could provide critical support for trait-based vegetation models and allow~~s~~ exploration into the relationships between vegetation characteristics and ecosystem functions at large scales. The ~~eight~~ six plant functional traits ~~datasets~~ maps for China with 1 km spatial resolution are now available at https://figshare.com/s/c527c12d310cb8156ed2 (An et al., 2023).

1 Introduction

Climate change has been affecting vegetation distributions and biogeochemical cycling globally and altering their feedbacks to the climate system (Kirilenko et al., 2000; Finzi et al., 2011; Jónsdóttir et al., 2022). Dynamic global vegetation models (DGVMs) are powerful tools for predicting changes in vegetation and ecosystem-atmosphere exchanges (e.g., water, carbon~~,~~ and nutrient cycling) in a changing climate (Foley et al., 1996; Peng, 2000). However, conventional DGVMs are still insufficient realistic, largely due to their dependence on the plant functional types (PFTs) assumption (Sitch et al., 2008; Yurova and Volodin, 2011; Scheiter et al., 2013). PFTs in conventional DGVMs commonly have fixed attributes (mostly trait values) (Van Bodegom et al., 2012; Wullschleger et al., 2014) that do not reflect plant adaptation to environments, limiting the quantification of carbon-water-nutrient feedback between terrestrial ecosystems and the atmosphere (Zaehle and Friend, 2010; Liu and Yin, 2013). Trait-based approaches can provide robust theoretical basis for developing the next generation of DGVMs (Van Bodegom et al., 2012; Sakschewski et al., 2015; Matheny et al., 2017). Plant functional traits, which are closely associated with ecosystem functions (Diaz et al., 2004; Yan et al., 2023), can effectively reflect response and adaptation of plants to environmental conditions (Myers-Smith et al., 2019; Qiao et al., 2023).

Attempts to predict spatially continuous trait ~~data~~ maps have been conducted at regional to global scales (Madani et al., 2018; Moreno-Martínez et al., 2018; Boonman et al., 2020; Loozen et al., 2020; Dong et al., 2023). Webb et al. (2010) proposed that the environment creates a filtered trait distribution along an environmental gradient, and such trait-environment relationships offer fundamental supports to predict the spatial distribution of plant functional traits through extrapolating local trait measurements. Boonman et al. (2020) mapped the global patterns of specific leaf area (SLA), leaf N concentration (LNC) and~~,~~ wood density (WD) ~~and plant height~~ based on a set of climate and soil variables. As the number of available global trait databases increases (~~Kattge et al., 2011;~~ Wang et al., 2018; Kattge et al., 2020), trait-environment relationships are becoming increasingly quantitative and accurate (Bruelheide et al., 2018; Myers-Smith et al., 2019). Alternatively, remote sensing approaches, such as empirical methods and physical radiative transfer models (e.g., partial least squares regression, PROSPECT model), have been developed to estimate plant physiological, morphological~~,~~ and chemical traits (e.g., leaf chlorophyll content, SLA, LNC and leaf dry matter content (LDMC)) (Darvishzadeh et al., 2008; Romero et al., 2012; Ali et al., 2016). Vegetation indices, such as the normalized difference vegetation index (NDVI) and the enhanced vegetation index (EVI), have been successful in estimating plant functional traits of crops, grasslands and forests (Clevers and Gitelson, 2013; Li et al., 2018; Loozen et al., 2018). Loozen et al. (2020) demonstrated that EVI was the most important predictor for mapping the spatial pattern of canopy nitrogen in European forests. Admittedly, recent studies have suggested that combining environmental variables and vegetation indices can improve the predictive accuracy of canopy nitrogen~~N~~ compared to those based on vegetation indices alone (Loozen et al., 2020).

Although there have been reports on plant functional trait distribution in China in some global or regional research (Yang et al., 2016; Butler et al., 2017; Madani et al., 2018; Moreno-Martínez et al., 2018; Boonman et al., 2020), they are still of large uncertainties in characterizing the spatial distribution of plant functional traits in China. First, global studies generally have relatively few, unevenly distributed sampling sites in China (Butler et al., 2017; Madani et al., 2018; Boonman et al., 2020), impeding our understanding of the true spatial characteristics of trait variability. Second, the spatial pattern of traits among these studies are usually inconsistent. For example, Moreno-Martínez et al. (2018) and Madani et al. (2018) demonstrated that SLA values were low in the southeast areas but high in the southwest areas of China, whereas Boonman et al. (2020) found the opposite. Third, most studies mainly focused on leaf traits (Yang et al., 2016; Loozen et al., 2018; Moreno-Martínez et al., 2018), whereas traits associated with the whole-plant ~~and reproductive~~ strategies, such as WD~~ and seed mass (SM)~~, were ignored. Therefore, mapping and verifying the spatial patterns of key functional traits that reflect the whole plant economics spectrum in China is a top priority.

In this study, our main objective was to generate ~~a~~ spatial ~~dataset~~maps~~–~~ for several key plant functional traits, through combining field measurements, environmental variables and vegetation indices. ~~To achieve this goal, w~~We ~~used a processing routine to predict the spatial distribution of plant functional traits.~~ selected six plant functional traits including SLA, LDMC, LNC, LPC, LA and WD. As key leaf economics traits, SLA, LDMC, LNC and LPC were selected because they are closely linked to plant growth rate, resource acquisition and ecosystem function (Wright et al., 2004; Diaz et al., 2016). LA is indicative of the trade-off between carbon assimilation and water-use efficiency (Wright et al., 2017), and WD reflects the trade-off between plant growth rate and support cost, with a higher WD linked to a lower growth rate, a higher survival rate and a higher biomass support cost (King et al., 2006). For each plant functional trait, we predicted spatial patterns at a 1 km resolution using an ensemble modelling algorithm based on two machine learning methods (i.e., random forest and boosted regression trees).

~~First, eight plant functional traits (i.e., SLA, LDMC, LNC, LPC, LA, plant height, WD and SM) were selected because they reflect plant adaptation to environment constraints and trade-offs between plant form and function (Reich and Cornelissen, 2014; Diaz et al., 2016). Second, we used random forest and boosted regression trees to predict the spatial patterns of plant functional traits by training the relationships between plant functional traits and environmental variables and vegetation indices. To obtain the optimal estimates, an ensemble model (i.e., weighted average algorithm) was further applied to merge the predictions of the two models. Finally, plant species were aggregated to PFTs, and the spatial abundance of PFTs at 1 km resolution was calculated using land cover map (100 m). We derived the spatial trait datasets via calculating community-weighted trait values within grid cells (1 km) based on these abundances of each PFT and predicted trait values.~~

2 Materials and Methods

2.1 Overview

The spatial maps of plant functional traits in China were generated based on machine learning algorithms trained by a large dataset of in-situ field measurements, environmental variables and vegetation indices in three steps (Fig. 1). First, in-situ field measurements of six plant functional traits were collected from TRY and China databases as well as published literature, and the PFTs of plant species were classified based on plant growth form, leaf type and leaf phenology. Multiple gridded predictors of climate, soil, topography and vegetation indices were used after avoiding the collinearity among them. Second, random forest and boosted regression trees were used to train the relationships between plant functional traits and predictors for each PFT individually. Third, the spatial abundance of each PFT within 1 km grid cell was calculated using land cover map (100 m). Community-weighted trait values within 1 km grid cell were calculated based on these abundances of each PFT and their predicted trait values in Step 2. To reduce the variability of different single-models, we derived the final spatial maps of plant functional traits using ensemble model to merge the predictions of random forest and boosted regression trees according to their cross-validated R^2 values.

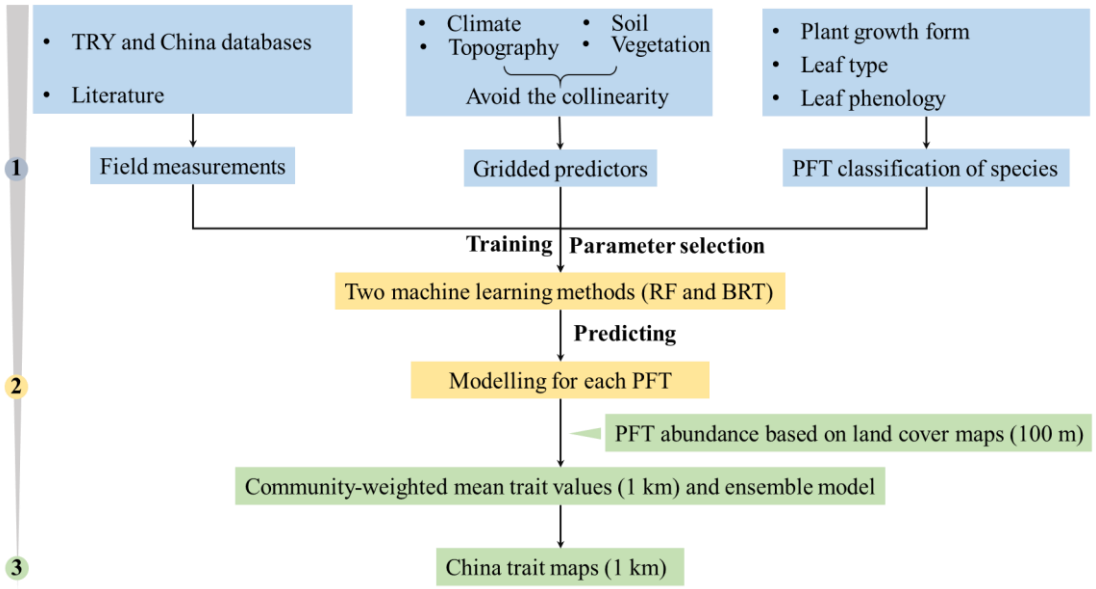


Figure 1. Methodological workflow for spatial mapping of plant functional traits. Trait mapping is performed in three steps. Step 1: in-situ field measurement of plant functional traits, PFT classification of plant species and gridded predictors were collected. Step 2: two machine learning methods were used to predict trait values by training the field measurements and predictors for each PFT. Step 3: spatialization of trait maps by calculating the abundance of each PFT using 100 m land cover map and predicted trait values within 1 km grid cells. PFT, plant functional type; RF, random forest; BRT, boosted regression trees.

2.~~1~~ 2 Plant functional trait collection and data processing

The information on the ~~eight~~ six plant functional traits and their ecological meanings are described in Table 1. Plant trait data was obtained and collected via two main sources. The first source was public trait databases, including the TRY database (Kattge et al., 2020) and the China Plant Trait Database (Wang et al., 2018). The second source was from literature (listed in Appendix A). To ensure data quality and comparability, we only included trait observations that met the following five criteria: 1) Measurements must be obtained from natural terrestrial fields in order to minimize the influences of management disturbance, and observations from cropland, aquatic habitat, control experiments or gardens were excluded; 2) According to the mass ratio hypothesis, the effect of plant species on ecosystem functioning is determined to an overwhelming extent by the traits and functional diversity of the dominant species and is relatively insensitive to the richness of subordinate species (Grime, 1998). Thus, we only included studies that measured plant trait observations from all species or dominant species within a community; 3) In order to consider the intraspecific trait variation, when the same species occurred in the same sampling site from different studies, we included all original observed data from different studies rather than averaging the values at the species level ~~In order to account for the trait variation within and between communities, we only considered real measurements of traits from individual plants, and not species-level averages~~ (Jung et al., 2010; Siefert et al., 2015); 4) Plant trait observations must be made on mature and healthy plant individuals, so some specific growth stages (e.g., seedling) and size classes (e.g., sapling) were excluded to reduce the confounding effect of ontogeny and seasonality (Thomas, 2010); 5) We only included studies with clear geographical coordinates to ensure alignment with predictor variables. The sampling location and sampling ~~sample~~ time~~time~~ information ~~from the original studies~~ were also included in the dataset. The sampling time mostly focused on the growing season of a year (i.e., May-October), which ensures the relative consistency of sampling time to minimize the effects of seasonality.

Plant functional traits must be sampled and measured according to standardized measurement procedures (Perez-Harguindeguy et al., 2013) to reduce the variation and uncertainty among different data sources. In this study, we included SLA measurements on ~~both~~ sun-leaves ~~and shade-leaves~~, and WD measurements on both heartwood and sapwood of tree species.~~, SM measurements on both seeds and fruits, and plant height measurements on both vegetative and generative organs.~~

Table 1 Description of plant functional traits selected in this study and their relevant ecosystem functions.

Trait	Abbreviation	Description	Relevant ecosystem functions
Specific leaf area	SLA	As a core leaf economics trait (Wright et al., 2004), it is related to trade-off between leaf lifespan and C acquisition as well as light competition (Reich et al., 1991)	Productivity, litter decomposition, competitive ability (Bakker et al., 2011; Smart et al., 2017)
Leaf dry matter content	LDMC	Strongly related to resource availability and potential growth rate (Hodgson et al., 2011)	Productivity, litter decomposition, herbivore resistance, and drought tolerance (Bakker et al., 2011; Smart et al., 2017; Blumenthal et al., 2020)
Leaf N concentration	LNC	As a core leaf economics trait, it is strongly related to photosynthetic capacity (Wright et al., 2004)	Productivity, nutrient cycling, litter decomposition (LeBauer and Treseder, 2008; Bakker et al., 2011)
Leaf P concentration	LPC	As a core leaf economics trait, it is strongly related to photosynthetic capacity (Wright et al., 2004)	Productivity, nutrient cycling, litter decomposition (LeBauer and Treseder, 2008; Bakker et al., 2011)
Leaf area	LA	Trade-off between C assimilation and water use efficiency, it is related to energy balance (Wright et al., 2017)	Productivity (Li et al., 2020)
~~Plant height~~	~~/~~	~~A major determinant of a plant's ability to compete for light (Moles et al., 2009)~~	~~C storage, animal diversity (Conti et al., 2013; Roll et al., 2015)~~
Wood density	WD	A measure of carbon investment, representing the trade-off between growth and mechanical support (Martínez-Vilalta et al., 2010)	Drought tolerance, productivity (Hoeber et al., 2014; Liang et al., 2021)
~~Seed mass~~	~~SM~~	~~Closely related to seed dispersal and seed predation, as well as seedling establishment, growth, and survival (Leishman et al., 2000)~~	~~Competitive ability (Zepeda and Martorell, 2019)~~

The plant trait data was checked for possible errors and corrected ~~as per~~in three steps as follows. First, species name and taxonomic nomenclature were corrected and standardized according to the Plant List (http://www.theplantlist.org/) using the "plantlist" package. Second, illogical values, repeated values and outliers were removed, which were defined by observations exceeding 1.5 standard deviations of the mean trait value for a given species (Kattge et al., 2011). Third, we appended information on plant growth form ~~(tree, shrub, and herb)~~, leaf type ~~(broadleaved and needle-leaved)~~ and leaf phenology ~~(evergreen and deciduous)~~ from the TRY categorical traits database (https://www.try-db.org/TryWeb/Data.php#3) and *Flora Reipublicae Popularis Sinicae* (http://www.iplant.cn/frps), which were used ~~. Furthermore, in order~~ to match species names to PFTs~~, .~~ ~~we~~ We associated each species ~~(i.e., plant growth form, leaf type and leaf phenology)~~ with a corresponding PFT based on plant growth form (tree, shrub and grass), leaf type (broadleaf and needleleaf) and leaf phenology (evergreen and deciduous). For example, the information on *Salix matsudana* is: tree, deciduous and broadleaf, thus, we were able to associate the PFT of deciduous

broadleaf forest (DBF) to this species. The species that did not correspond to any PFT were discarded. After these treatments, we collected a total of 3458952477 trait measurements from 1541 1430 sampling sites for our database, representing 4291 3447 species from 212 195 families and 1230 1066 genera (Fig. 12 and Fig. B1 in Appendix B). Information on the statistics for the eight six plant functional traits collected in this study is shown in Table B1 in Appendix B.

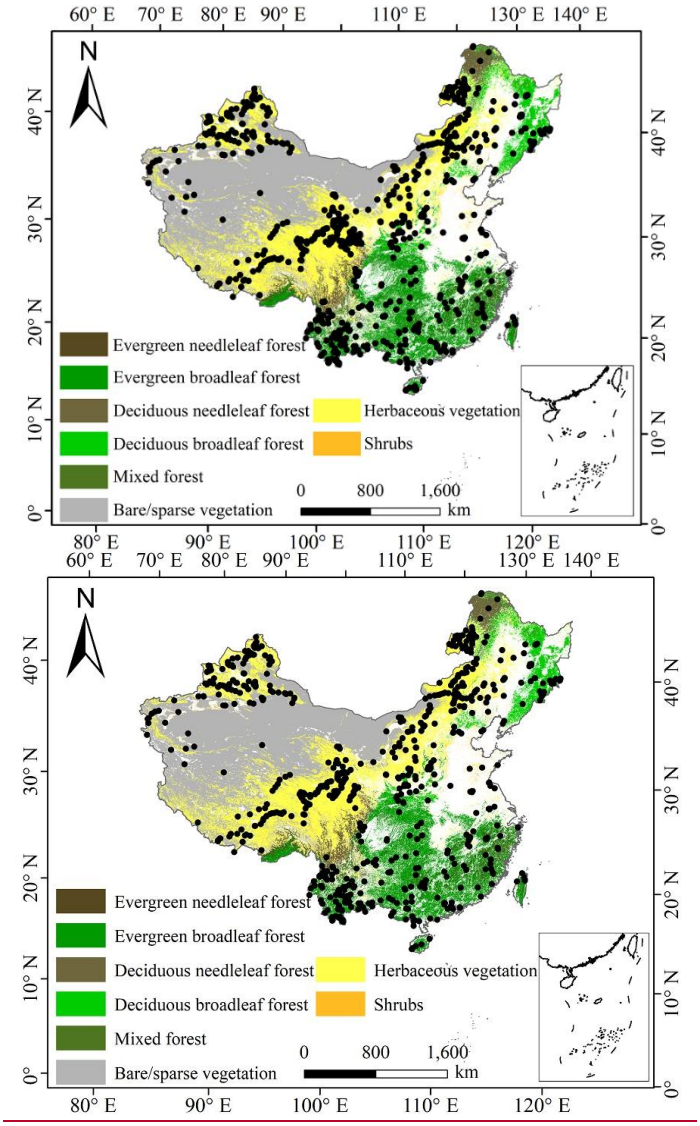


Figure 12. Location distribution and land cover map in China.

2.2 3 Preparing predictor variables Environmental data

2.23.1 Climate data

Twenty-one climate variables were used in this study, including 19 bioclimate variables, solar radiation (RAD), and aridity index (AI) (Table B2 in Appendix B). The 19 bioclimate variables and RAD were obtained from the WorldClim dataset (version 2.11) for the period from 1970 to 2000 (https://www.worldclim.org/data/worldclim21.html). The AI data was extracted from the CGIAR Consortium of Spatial Information (CGIAR-CSI) website for the period from 1950 1970 to 2000 (http://www.csi.cgiar.org) (Trabucco and Zomer, 2018). The spatial resolution of climate data was

is 1 km.

2.~~2~~3.2 Soil data

Twelve soil variables were included in this study, representing the different aspects of soil properties, i.e. soil texture, bulk density (BD), pH~~,~~ and soil nutrients (Table B2 in Appendix B). All soil variables were extracted from the Soil Database of China for Land Surface Modeling (http://globalchange.bnu.edu.cn/research/soil2) (Shangguan et al., 2013). Given the importance of topsoil properties on community composition (Bohner, 2005), we averaged the first four layers to ~~the soil data was averaged to~~ represent the topsoil ~~30 cm of soil~~ properties (~ 30 cm) in our study. The spatial resolution ~~was~~ is 1 km.

2.~~2~~3.3 Topography

The topographic variable was elevation. Elevation data was extracted from the STRM 90m dataset in China, based on the SRTM V4.1 database (https://www.resdc.cn/data.aspx?DATAID=123). The spatial resolution ~~was~~ is 1 km.

Given the collinearity among climate and soil variables, we reduced the number of environmental predictors based on Pearson's correlation coefficient (r) (Figs. ~~B1~~ B2 and ~~B2~~ B3 in Appendix B). Among a set of highly correlated variables ($r > 0.75$), only one variable was retained in subsequent analysis to ensure a combination of different environmental variables. The final selection of environment predictors included nineteen variables: mean annual temperature (MAT), mean diurnal range (MDR), min temperature of coldest quarter (Tmin), max temperature of ~~coldest~~ warmest quarter (Tmax), temperature seasonality (TS), mean annual precipitation (MAP), precipitation seasonality (PS), precipitation of wettest quarter (PEQ), precipitation of driest quarter (PDQ), AI, RAD, elevation, soil sand content (SAND), pH, BD, soil total N (STN), soil total P (STP), soil available P (SAP), soil alkali-hydrolysable N (SAN)~~,~~ and cation exchange capacity (CEC).

2.~~3~~ 3.4 Vegetation indices

Three categories of vegetation indices were included in this study (Table B2 in Appendix B). First, EVI ~~The first selected was EVI, which~~ was extracted from the MOD13A3 V006 product (https://lpdaac.usgs.gov/products/mod13a3v006/). This product is available as a monthly average with spatial resolution of 1 km, ranging from January 2000 to December 2018. Second, MODIS reflectance data was also extracted from the MOD13A3 V006 product~~. This, included~~ including MIR reflectance, NIR reflectance, red reflectance~~,~~ and blue reflectance. Third, the MERIS terrestrial chlorophyll index (MTCI) was extracted from the Natural Environment Research Council Earth Observation Data Centre (NERC-NEODC, 2005) (https://data.ceda.ac.uk/). MTCI data is available globally as a monthly average at 4.63 km spatial resolution, and ranges from June 2002 to December 2011. It is noted that valid MTCI values should be greater than 1, so our study deleted any values less than 1.

To avoid collinearity, we also reduced the number of vegetation indices based on Pearson's correlation coefficient (r) (Fig~~s~~. ~~B3~~ B4 in Appendix B). Most selected variables were related to

growing seasons due that plant functional traits were measured during the growing season. Furthermore, based on the results of Pearson's correlation coefficient (r), MTCI, MIR, NIR, red and blue in January showed low correlations with those in growing season, thus they were included in subsequent analysis. ~~Furthermore, given that most plant functional traits were measured during the growing season, the variables related to the growing season were determined to be important predictors.~~ The final selection included 36 variables: annual EVI, EVI (May, June, July, August and September), MTCI, MIR, NIR, red and blue (all for January, June, July, August and September).

Both environmental variables and vegetation indices variables were resampled to a consistent spatial resolution of 1 km using the nearest neighborhood method.

PFT is also an important factor in influencing the variation of plant functional traits (Verheijen et al., 2016; Loozen et al., 2020), thus the trait predictions were performed for each PFT individually.~~thus PFT was included as a predictor in this analysis.~~ We used the 2015 land cover map at a 100 m spatial resolution to calculate the relative abundance of each PFT within 1 km grid cells, which was extracted from the Copernicus Global Land Service (CGLS-LC100, Version 3) (https://land.copernicus.eu/global/products/lc) (Buchhorn et al., 2020). We focused on natural terrestrial vegetation, so all artificial or crop areas were thus eliminated in our dataset. Seven categories were included: evergreen needleleaf forest (ENF), evergreen broadleaf forest (EBF), deciduous needleleaf forest (DNF), deciduous broadleaf forest (DBF), shrubland (SHL), grassland (GRL) and bare/sparse vegetation. ~~Furthermore, in order to match species names to PFTs, we associated each species (i.e., plant growth form, leaf type and leaf phenology) with a corresponding PFT. For example, the information on *Salix matsudana* is: tree, deciduous and broadleaf, thus, we were able to associate the PFT of deciduous broadleaf forest (DBF) to this species. The species that did not correspond to any PFT were discarded.~~

2.~~4~~ 4 Model fitting and validation

To predict spatial patterns of plant functional traits, we used two machine learning models, i.e., random forest and boosted regression trees.

Random forest is an ensemble machine learning method based on classification and regression trees using collections of regression trees to classify observations according to a set of predictive variables (Breiman, 2001). This method repeatedly constructs a set of trees from random samples of training data, and the final prediction is produced by integrating the results of all individual trees, which makes it a robust method. The model~~s are~~ is controlled by two main parameters: the number of sampled variables (mtry) and the number of trees (ntree). The ~~parameter~~ mtry was set to range from 1 to 57 (at an interval of 1), and the ntree was set as 500, 1000, 2000, 5000 and 10000 in subsequent runs. This analysis was performed using the 'randomForest' function in the 'randomForest' package (Liaw and Wiener, 2002).

Boosted regression trees ~~is~~ are ~~a~~ machine learning method~~s~~ based on generalized boosted regression model~~s,~~ and using a boosting algorithm to combine many sample tree models to optimize

predictive performance (Elith et al., 2006). There is no need for prior data transformation or the elimination of outliers, and this method can fit complex non-linear relationships while automatically handling interaction effects between predictors (Elith et al., 2008). The four parameters to optimize in these models are the number of trees, interaction depth, learning rate and bag fractions. We varied the parameter settings to find the optimal parameter combination that achieves minimum predictive error. The number of trees was set to 3000, the interaction depth varied from 1 to 7 (at an interval of 1), the learning rate was set to 0.001, 0.01, 0.05 and 0.1, and the bag fraction was set to 0.5, 0.6, 0.7 and 0.75. PFT was used as a dummy variable in the boosted regression trees models. This analysis was conducted using the 'gbm' function in the 'gbm' package (Ridgeway, 2006).

We built separate predictive model for each plant functional trait. ~~We used a 10-fold cross validation~~ To select the optimal parameter combination and to evaluate the final model performance for each trait, we calibrated the models 10 times using randomly selected 80% of the data for training the models and validating against the remaining 20% based on cross-validation (Table B3 in Appendix B). ~~We spilt the data into two parts: 80% of the trait data was used to train the models, and the remaining 20% was used to assess model's performance.~~ The predictive performance was evaluated by regressing the predicted and observed trait values from all repetitions of the cross-validation. The fitting performances of the random forest and boosted regression trees methods were evaluated using determinate coefficient (R^2), normalized root-mean-square error (NRMSE)~~,~~ and mean absolution error (MAE). These scores are calculated following Eq. (1), Eq. (2) and Eq. (3):

$$R^2 = 1 - \frac{\sum_{i=1}^{n}(p_i - o_i)^2}{\sum_{i=1}^{n}(p_i - \hat{o_i})^2} \quad (1)$$

~~$RMSE = \sqrt{\quad}$~~ $$\text{NRMSE} = \frac{\sqrt{\frac{1}{n}\sum_{i=1}^{n}(p_i - o_i)^2}}{p_{max} - p_{min}} \quad (2)$$

$$MAE = \frac{1}{n}\sum_{i=1}^{n}|o_i - p_i| \quad (3)$$

where p_i and o_i are the predictive values and observed values, respectively~~,~~; $\hat{o_i}$ is the mean of the observed values.

To quantify the relative importance of each predictor ~~consistently~~ across the two models consistently, we used the method proposed by Thuiller et al. (2009). This method applies correlation between the standard predictions fitted with the original data and predictions where the variable under investigation has been randomly permutated. If the correlation is high, which indicates little difference between the two predictions, the variable permutated is considered not important for the model. ~~The correlations between fitted values and predictions were calculated using permuted values for the predictor of concern.~~ This step was repeated ~~a user-defined number of~~multiple times for each predictor, and the mean correlation coefficient over runs was recorded. Then the relative importance of each predictor was quantified as one minus the Spearman rank correlation coefficient (see Boonman et al., 2020). In addition, we used generalized additive models to fit the relationships between plant functional traits and the most important variables using the 'gam' function in the 'mgcv' package.

2.~~5~~ 5 Generation of plant functional trait maps and model performance

The generation of spatial maps of plant functional was performed in three steps. First, we predicted trait values for each PFT within 1 km grid cell separately. Second, the abundance of individual PFT within 1 km grid cell was estimated using a land cover map with a spatial resolution of 100 m. Third, refer to the Eq. (4) that has been widely applied in a community (Garnier et al., 2004), the ~~The~~ final ~~community weighted mean~~ trait values in a given 1 km grid cell were~~as~~ as calculated ~~according to~~as the sum of the predicted trait values ~~and~~multiplying by corresponding abundance of each PFT.

$$\mathrm{CWM} = \sum_{i=1}^{n} W_i X_i \quad (4)$$

where n is the total number of PFT in a given grid; W_i is the relative abundance of the ith PFT; and X_i is the predicted trait value of the ith PFT.~~To calculate community weighted mean trait values, the abundance of individual PFT within 1 km grid cell was estimated using a land cover map with a spatial resolution of 100 m. The final community weighted mean trait values were calculated according to the predicted trait values and corresponding abundance of each PFT.~~

To reduce the variability of different single-models and to construct a more stable and accurate model, ~~To obtain the optimal estimates,~~ the ensemble model was further applied to merge the predictions of random forest and boosted regression trees according to their cross-validated R^2 values. The predictive value of ensemble model was calculated in a given grid cell as described by Eq. (5) (Marmion et al., 2009). The model accuracy was calculated by regressing the predictive values of ensemble model against the observed trait values.~~The accuracy of the ensemble model was calculated by regressing the 20% of cross-validation data used for testing against the observed trait values.~~

$$Pred\_EM_t = \frac{\sum_{m=1}^{2}(pred_{m,t} \times r_{m,t}^2)}{\sum_{m=1}^{2} r_{m,t}^2} \quad (5)$$

where $Pred\_EM_t$ is the predictive values of t trait in the ensemble model; $pred_{m,t}$ is the predictive values of t trait in m model; $r_{m,t}^2$ is the cross-validated R^2 of t trait in m model.

To evaluate the model performance (i.e. the variability in the prediction across models), the coefficient of variation (CV) was calculated as the difference between the predictions of random forest and boosted regression trees methods and the ensemble prediction ~~weighted by the predictive performance of each of the models~~. CV is calculated as following Eq. (~~4~~6):

$$CV_t = \frac{\sqrt{\sum_{m=1}^{2}(pred_{m,t} - obs_t)^2 * r_{m,t}^2}}{\sum_{m=1}^{2} r_{m,t}^2} / obs_t \quad (~~4~~6)$$

where $pred_{m,t}$ is the predictive values of t trait in m model~~,~~; obs_t is the values of t trait in the ensemble model~~,~~; $r_{m,t}^2$ is the cross-validated R^2 of t trait in m model.

2.~~6~~ 6 Uncertainty assessments

Multivariate environmental similarity surface analysis (MESS) was used to identify the range of the extrapolated predictor values across the locations in the plant trait dataset (Elith et al., 2010). This

method is often used to evaluate the extent of extrapolation and the applicability domain. If the values are negative, this indicates that at a given grid cell, at least one predictor variable is outside the extent of referenced predictor layer. This analysis was conducted using the 'mess' function in the 'dismo' package.

All analyses were performed in R 4.0.2 (R Core Team, 2020).

3 Results

3.1 Performances of prediction models

Cross-validation showed that the performance of the predictive models differed greatly among the plant functional traits (Table 2, Tables C1 and C2 in Appendix C). WD had the best performance in all three models, with R^2 values of 0.64, 0.68 and 0.67 for random forest, boosted regression trees and ensemble model, respectively. SLA~~LPC, height~~ and LPC ~~SLA~~ had R^2 values greater than 0.45, while ~~SM and~~ LDMC performed the worst, with R^2 values below 0.25. ~~In addition, the ensemble model performed better than the random forest and boosted regression trees alone (Tables C1 and C2 in Appendix C).~~

Table 2 Results of plant functional traits for cross-validated R^2, NRMSE and MAE for random forest, boosted regression trees~~,~~ and ensemble model.

	Random forest			Boosted regression trees			Ensemble model		
Traits	R^2	NRMSE	MAE	R^2	NRMSE	MAE	R^2	NRMSE	MAE
SLA	0.48	~~7.02~~0.22	5.10	0.48	~~6.99~~0.20	5.08	0.49	~~6.98~~0.21	5.07
LDMC	0.23	~~0.10~~0.21	0.07	0.28	~~0.09~~0.18	0.07	0.24	~~0.09~~0.20	0.07
LNC	0.33	~~6.64~~0.19	4.92	0.34	~~6.52~~0.18	4.85	0.34	~~6.54~~0.19	4.85
LPC	0.51	~~0.80~~0.24	0.53	0.51	~~0.80~~0.22	0.53	0.51	~~0.80~~0.27	0.53
LA	0.37	~~68.94~~0.45	26.76	0.39	~~67.69~~0.51	27.47	0.40	~~67.28~~0.58	26.59
~~SM~~	~~0.24~~	~~4547.22~~	~~1228.07~~	~~0.26~~	~~4478.88~~	~~1183.61~~	~~0.25~~	~~4499.67~~	~~1201.83~~
~~Height~~	~~0.49~~	~~2.89~~	~~2.09~~	~~0.49~~	~~2.89~~	~~2.10~~	~~0.49~~	~~2.89~~	~~2.10~~
WD	0.64	~~0.13~~0.20	0.10	0.68	~~0.12~~0.13	0.10	0.67	~~0.13~~0.17	0.10

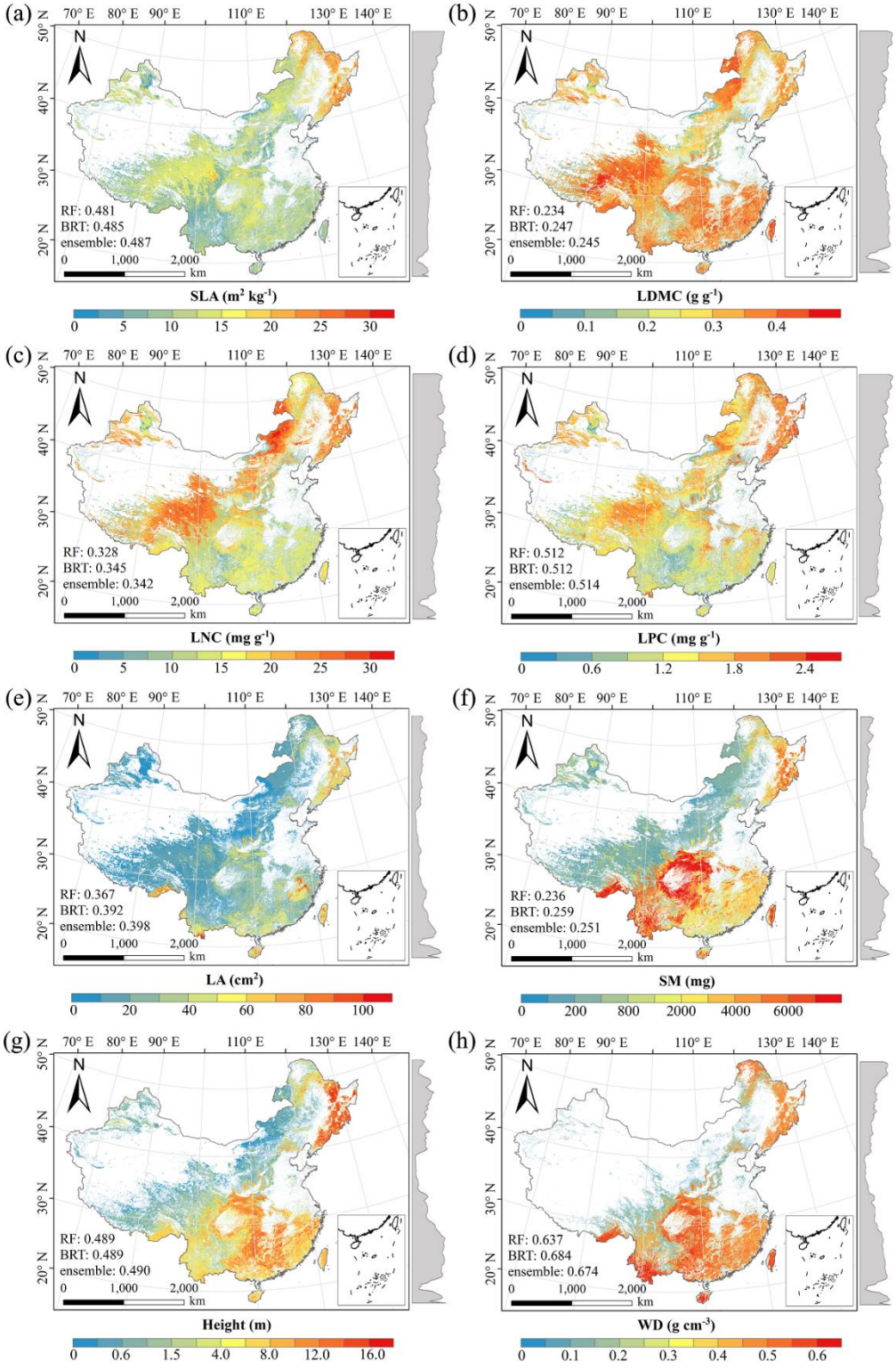
SLA, specific leaf area (m^2 kg^{-1}); LDMC, leaf dry matter content (g g^{-1}); LNC, leaf N concentration~~concertation~~ (mg g^{-1}); LPC, leaf P concentration~~concertation~~ (mg g^{-1}); LA, leaf area (cm^2); WD, wood density (g cm^{-3})~~;~~; R^2, determinate coefficient; NRMSE, normalized root-mean-square error; MAE, mean absolution error. ~~SM, seed mass.~~

3.2 Spatial patterns of predicted plant functional traits

There were relatively consistent spatial patterns for SLA, LNC and LPC, with high values in the northeastern and northwestern regions and the southeastern Qinghai-Tibet Plateau, and low values in southwestern China (Figs. ~~2a~~3a, ~~2c~~ 3c and ~~2d~~3d, Figs. D1 and~~,~~ D2 in Appendix D). SLA and LPC increased with latitude, while LNC did not vary significantly along the latitudinal gradient. For SLA, LNC and LPC, the variability was low among ~~the~~ random forest, boosted regression trees and ensemble model, with an overall CV less than 0.3 (Figs. ~~3a~~4a, ~~3c~~4c~~,~~ and ~~3d~~4d). LDMC values were

relatively high in most regions of China, and the low values were mainly located in eastern Yunnan and the Loess Plateau (Fig. ~~2b~~3b, Figs. D1 and~~,~~ D2 in Appendix D). LA ~~and SM showed consistent spatial patterns, with~~showed high values in the northeastern and southern regions (except for the Sichuan Basin), and the southeastern Qinghai-Tibet Plateau (Fig~~s~~. ~~2~~3e ~~and 2f~~, Figs. D1~~,~~ and D2 in Appendix D). The strong latitudinal gradient~~s~~ ~~were~~ was observed in LA ~~and SM~~, where values decreased with latitude.

The CV values of LPC ~~and SM~~ decreased with latitude, but other traits did not show latitudinal patterns (Fig~~s~~. ~~3~~4). The CV values of LA were relatively high, especially in the northwestern region~~,~~ and the Inner Mongolia ~~Plateau~~-Loess Plateau region ~~(only for LA) and Yunnan province (only for SM)~~ (Fig~~s~~. ~~3~~4e ~~and 3f~~). ~~Plant height and~~ WD had ~~consistent spatial patterns, with~~ high values in the northeastern and southern regions (Fig~~s~~. ~~2g~~ 2f~~and 2h~~, Figs. D1~~,~~ and D2 in Appendix D)~~. The CV values across models for plant height were higher in northwestern China and Inner Mongolia Plateau-Loess Plateau region, ~~,while CV values for WD in China were low throughout China (Fig~~s~~. ~~3g~~ 4f~~and 3h~~).


(a)
RF: 0.481
BRT: 0.485
ensemble: 0.487
SLA (m² kg⁻¹)
0 5 10 15 20 25 30
(b)
RF: 0.234
BRT: 0.247
ensemble: 0.245
LDMC (g g⁻¹)
0 0.1 0.2 0.3 0.4
(c)
RF: 0.328
BRT: 0.345
ensemble: 0.342
LNC (mg g⁻¹)
0 5 10 15 20 25 30
(d)
RF: 0.512
BRT: 0.512
ensemble: 0.514
LPC (mg g⁻¹)
0 0.6 1.2 1.8 2.4
(e)
RF: 0.367
BRT: 0.392
ensemble: 0.398
LA (cm²)
0 20 40 60 80 100
(f)
RF: 0.236
BRT: 0.259
ensemble: 0.251
SM (mg)
0 200 800 2000 4000 6000
(g)
RF: 0.489
BRT: 0.489
ensemble: 0.490
Height (m)
0 0.6 1.5 4.0 8.0 12.0 16.0
(h)
RF: 0.637
BRT: 0.684
ensemble: 0.674
WD (g cm⁻³)
0 0.1 0.2 0.3 0.4 0.5 0.6
70° E 80° E 90° E 110° E 130° E 140° E
50° N 40° N 30° N 20° N
N
0 1,000 2,000 km

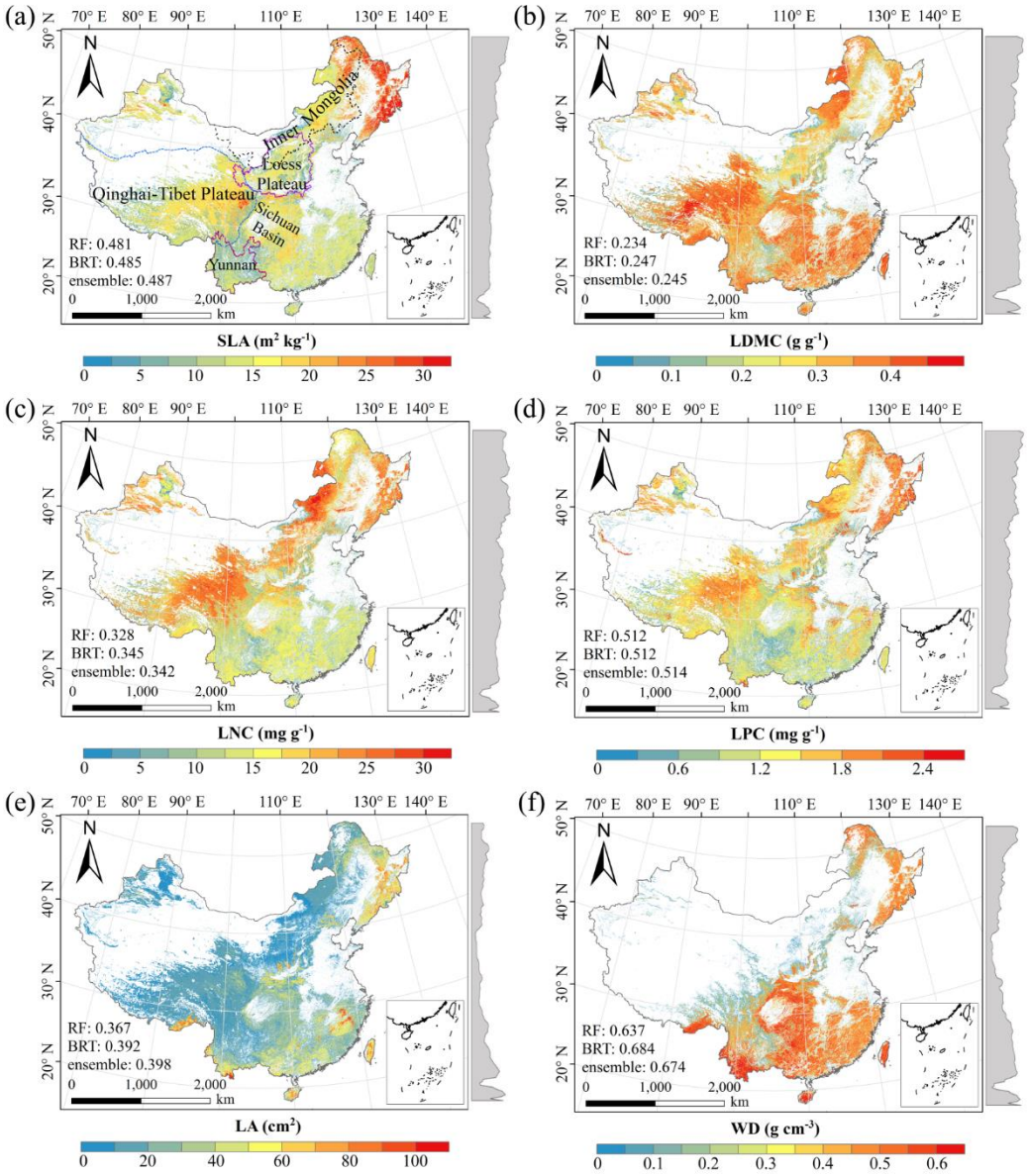
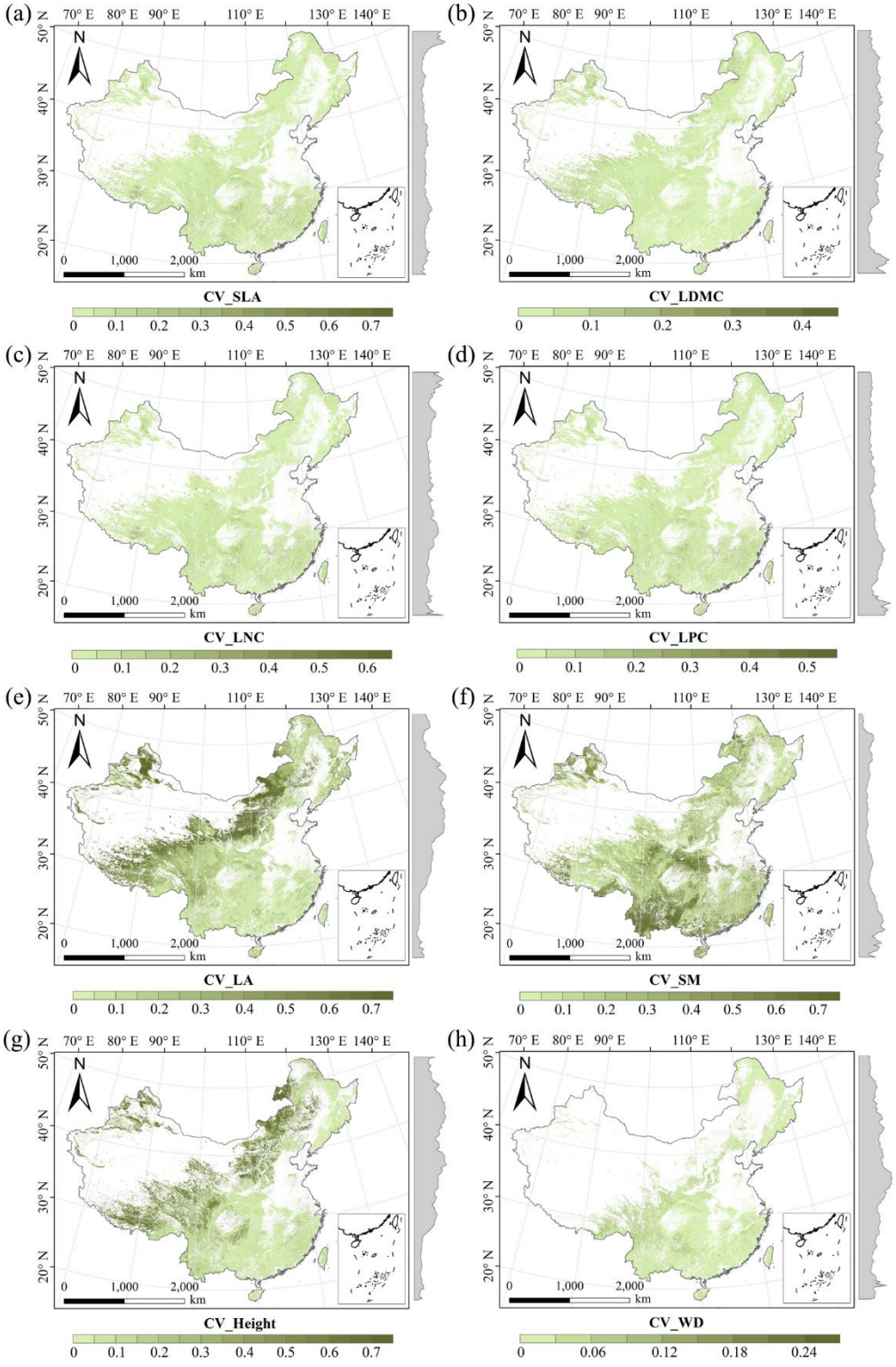


Figure ~~2~~3. Spatial patterns of predicted plant functional traits in China based on the ensemble model. The grey curves to the right of the maps display trait distribution along with latitude. RF, random forest; BRT, boosted regression trees; ensemble, ensemble model; SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density~~; SM, seed mass~~.


(a)
CV_SLA
0 0.1 0.2 0.3 0.4 0.5 0.6 0.7
(b)
CV_LDMC
0 0.1 0.2 0.3 0.4
(c)
CV_LNC
0 0.1 0.2 0.3 0.4 0.5 0.6
(d)
CV_LPC
0 0.1 0.2 0.3 0.4 0.5
(e)
CV_LA
0 0.1 0.2 0.3 0.4 0.5 0.6 0.7
(f)
CV_SM
0 0.1 0.2 0.3 0.4 0.5 0.6 0.7
(g)
CV_Height
0 0.1 0.2 0.3 0.4 0.5 0.6 0.7
(h)
CV_WD
0 0.06 0.12 0.18 0.24
70° E 80° E 90° E 110° E 130° E 140° E
50° N 40° N 30° N 20° N
N
0 1,000 2,000 km

397

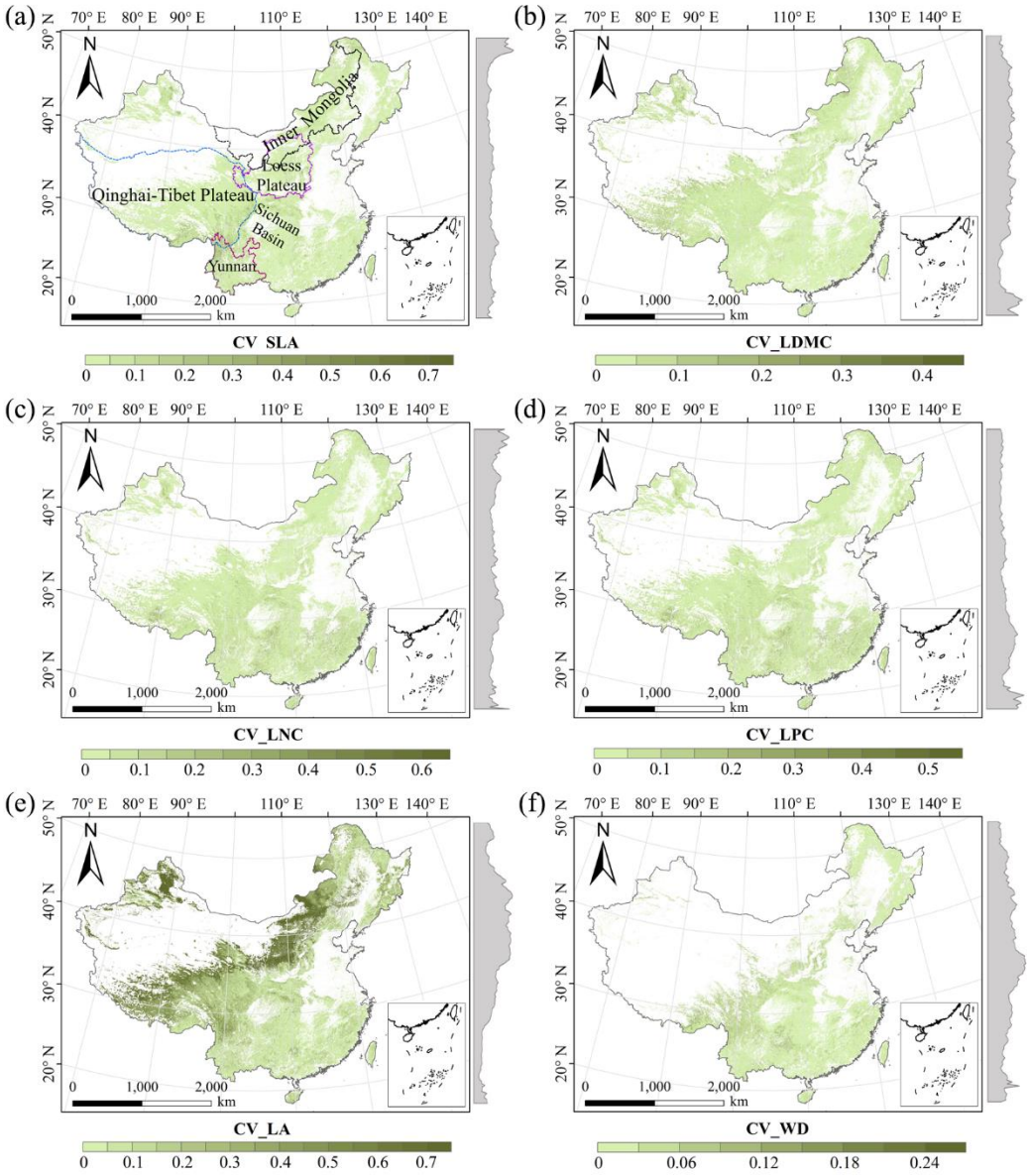


Figure ~~3~~4. The variability in plant functional trait predictions among random forest, boosted regression trees and ensemble model. The grey curves to the right of the maps display coefficient of variation along with latitude. SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density~~; SM, seed mass~~.

3.3 Relative importance of predictive variables

The dominant factors explaining spatial variation differed greatly among ~~the eight~~ plant functional traits (Table 3). Overall, climate variables were more important for predicting plant functional traits than were soil variables. Temperature variables (i.e., MAT, MDR and TS) showed close relationships with SLA, LDMC, LPC and WD, while precipitation variables (i.e., PS, PEQ, MAP and PDQ) were more important for predicting the spatial patterns of LNC, LPC~~,~~ and LA ~~and plant~~

~~height~~. RAD was the fourth most dominant factor in predicting the spatial patterns of SLA ~~, SM,~~ and WD ~~and plant height~~. Elevation also played an important role in the LDMC and LPC predictions. Within soil variables, soil nutrients ~~[~~(i.e., pH and ~~ and soil available P (~~SAP~~)]~~) showed close associations with SLA~~,~~ and LNC ~~and SM~~. In addition to the environmental variables, MTCI emerged as an important predictor for explaining SLA, LDMC~~,~~ and LA ~~and plant height~~. Finally, EVI was the most important predictor for LA ~~and SM~~, and MIR in January and May were the primary predictors of WD. The relationships between plant functional traits and the most important variables were shown in Figs. E1 and E2 in Appendix E.

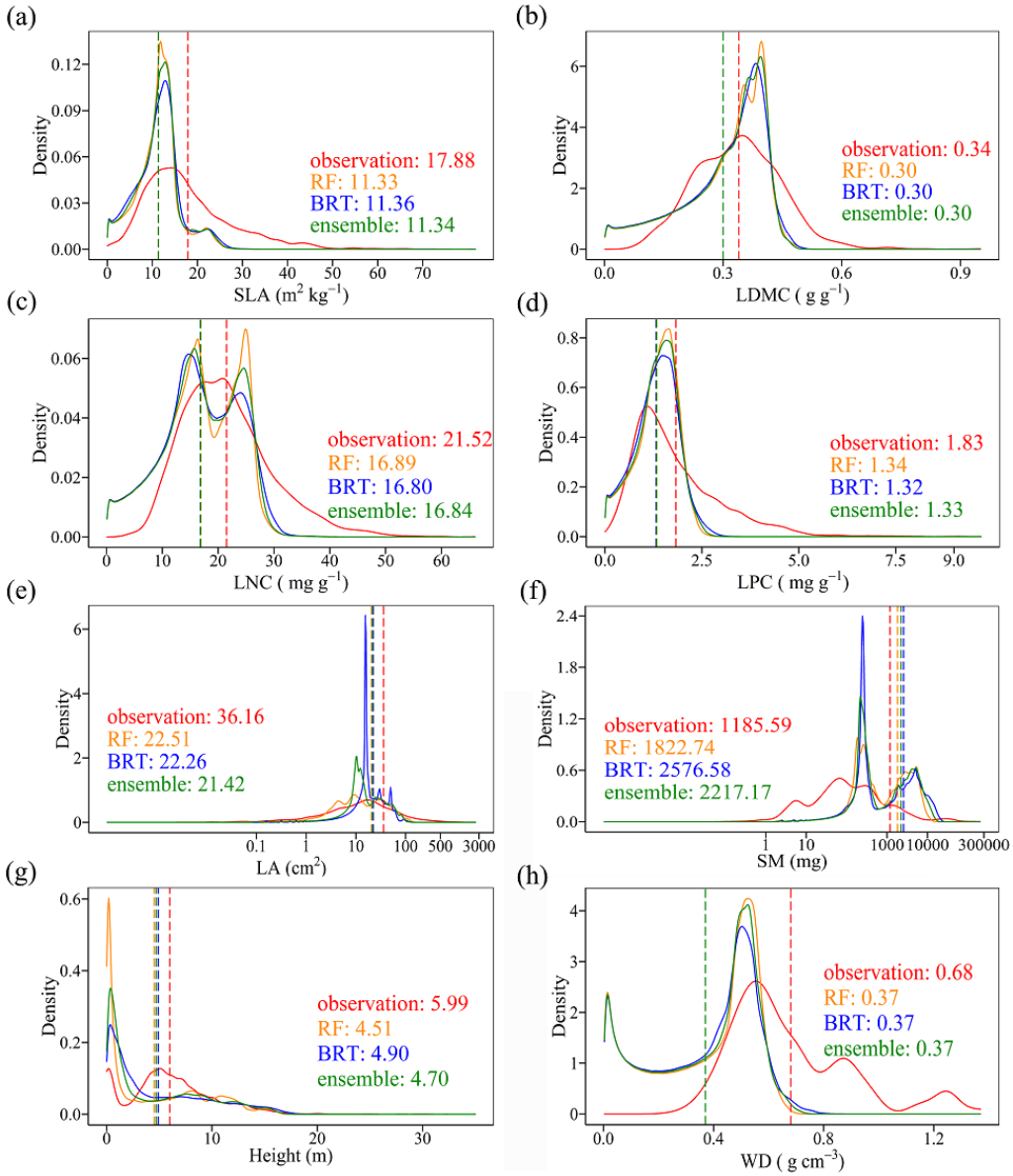
Table 3 List of the eight most important variables for plant functional trait predictions.

Rank	SLA	LDMC	LNC	LPC	LA	~~SM~~	~~Height~~	WD
1	SAP	MAT	PS	MDR	EVI5	~~EVI8~~	~~PEQ~~	MIR1
2	TS	Elevation	SAP	PDQ	PEQ	~~SAP~~	~~EVI6~~	TS
3	blue9	MTCI5	pH	Elevation	MTCI9	~~MAT~~	~~MTCI6~~	MIR5
4	RAD	blue8	MDR	MIR8	NIR9	~~RAD~~	~~RAD~~	RAD
5	MTCI4	MTCI4	MAP	Tmax	AI	~~NIR1~~	~~pH~~	MIR6
6	MTCI6	MTCI6	PEQ	MTCI6	MTCI6	~~NIR8~~	~~MAP~~	pH
7	Elevation	NIR1	MIR1	MIR7	MAP	~~SAND~~	~~red9~~	red5
8	MTCI7	CEC	Tmax	MIR9	red5	~~BD~~	~~red5~~	PS

SLA, specific leaf area (m^2 kg^{-1}); LDMC, leaf dry matter content (g g^{-1}); LNC, leaf N concentration (mg g^{-1})~~concertation~~; LPC, leaf P concentration (mg g^{-1})~~concertation~~; LA, leaf area (cm^2); WD, wood density (g cm^{-3})~~; SM, seed mass~~; SAP, soil available P; TS, temperature seasonality; blue, blue reflectance; RAD, solar radiation; MTCI, MERIS terrestrial chlorophyll index; MAT, mean annual temperature; NIR, near-infrared reflectance; CEC, cation exchange capacity; PS, precipitation seasonality; MDR, mean diurnal range; MAP, mean annual precipitation; PEQ, precipitation of wettest quarter of a year; MIR, middle infrared reflectance; Tmax, max temperature of warmest month of a year; PDQ, precipitation of driest quarter of a year; EVI, enhanced vegetation index; AI, aridity index; red, red reflectance~~; SAND, soil sand content; BD, bulk density~~.

3.4 Model performance

The distributions of the predictive trait values based on random forest, boosted regression trees, and ensemble model were consistent with the original trait observations, especially the peak values (Fig. ~~4~~5). ~~Except for SM, t~~The mean values of trait observations were relatively higher than those of the predictive values.


(a)
Density
SLA (m² kg⁻¹)
observation: 17.88
RF: 11.33
BRT: 11.36
ensemble: 11.34
(b)
Density
LDMC (g g⁻¹)
observation: 0.34
RF: 0.30
BRT: 0.30
ensemble: 0.30
(c)
Density
LNC (mg g⁻¹)
observation: 21.52
RF: 16.89
BRT: 16.80
ensemble: 16.84
(d)
Density
LPC (mg g⁻¹)
observation: 1.83
RF: 1.34
BRT: 1.32
ensemble: 1.33
(e)
Density
LA (cm²)
observation: 36.16
RF: 22.51
BRT: 22.26
ensemble: 21.42
(f)
Density
SM (mg)
observation: 1185.59
RF: 1822.74
BRT: 2576.58
ensemble: 2217.17
(g)
Density
Height (m)
observation: 5.99
RF: 4.51
BRT: 4.90
ensemble: 4.70
(h)
Density
WD (g cm⁻³)
observation: 0.68
RF: 0.37
BRT: 0.37
ensemble: 0.37

434

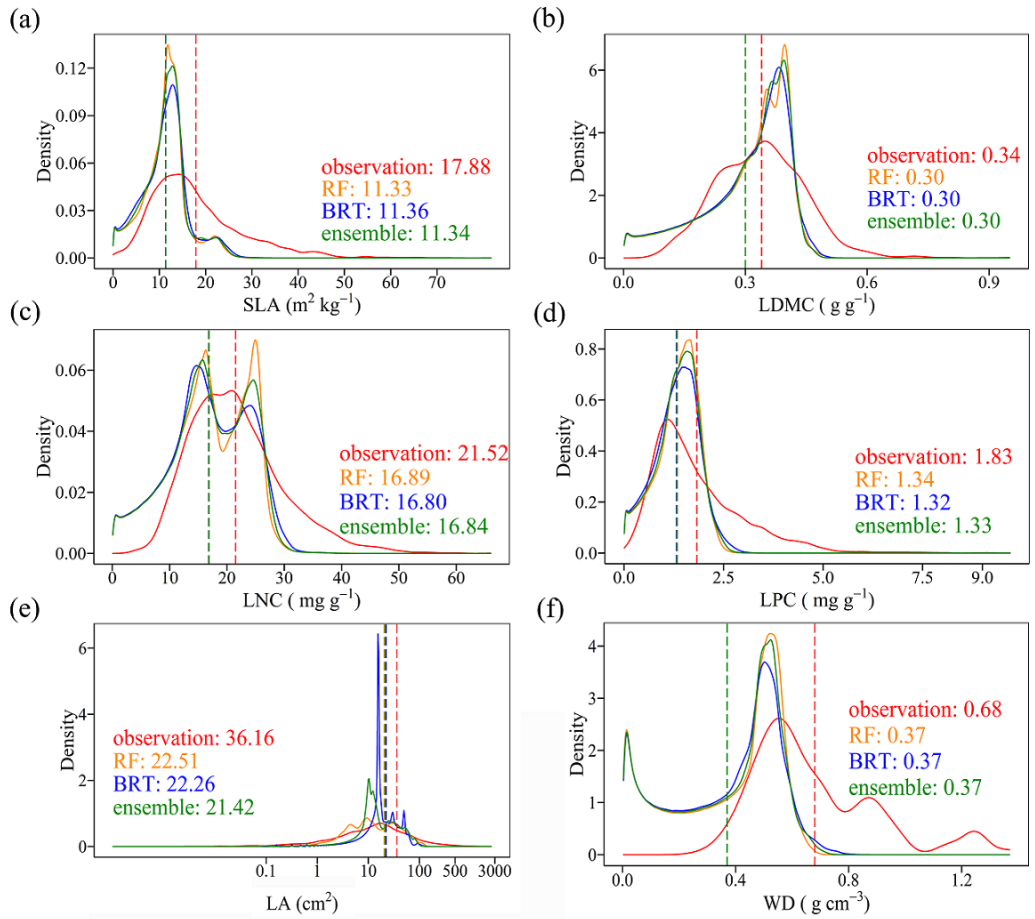
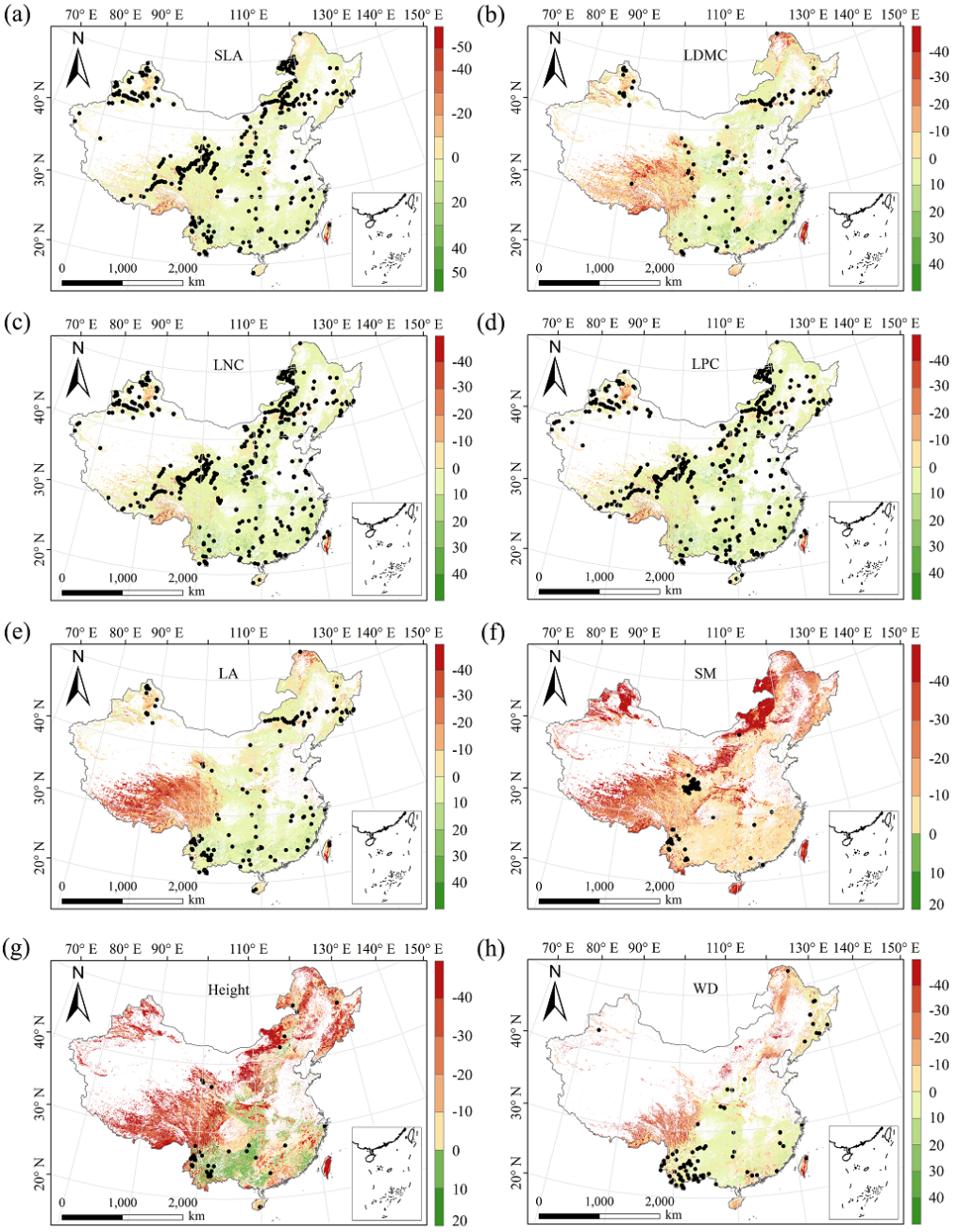


Figure 45. Comparison of trait distribution between observations and predictive values in each of the different models. Each panel depicts the distribution of observations in solid red, of the random forest (RF) model in yellow, of the boosted regression trees (BRT) model in blue, and of the ensemble model in green. The dashed vertical lines indicate mean values. SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density~~; SM, seed mass~~.

3.5 Uncertainty assessments

The MESS values of ~~five leaf traits and WD~~all plant functional traits were positive in most regions, indicating a wide applicability domain of our models (Fig. ~~5~~6). Nevertheless, trait predictions should be interpreted carefully for northeastern China and the Qinghai-Tibet Plateau due to the sparse samplings in these regions.~~ In addition, spatial predictions for SM and plant height were extrapolated to a larger extent than were the other plant functional traits.~~


(a)
SLA
(b)
LDMC
(c)
LNC
(d)
LPC
(e)
LA
(f)
SM
(g)
Height
(h)
WD
70° E 80° E 90° E 110° E 130° E 140° E 150° E
40° N
30° N
20° N
N
0 1,000 2,000 km

449

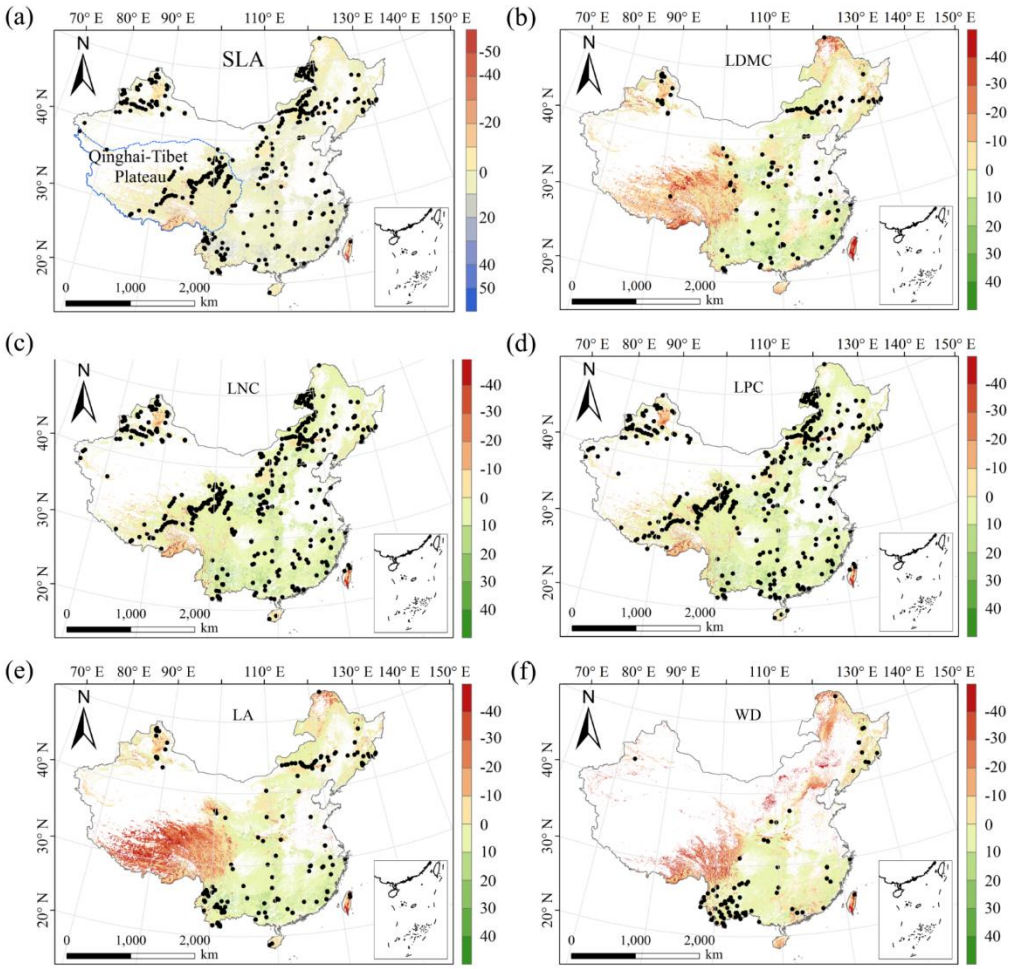


Figure ~~5~~6. Multivariate environmental similarity surface (MESS) assessments for the ~~eight~~ six plant functional traits. The black dots represented the locations of trait observations. More intense shades indicate greater similarity (blue) or difference (red) in environmental conditions of the location compared to the predictive factors covered by the training dataset. SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density~~; SM, seed mass~~.

4 Discussion

4.1 Comparison with previous work

Our study predicted the spatial patterns of ~~eight~~ six key plant functional traits across China using machine learning methods and identified the applicability domain of the models. WD had the highest precision with a~~n~~ R^2 of 0.66, which was higher than the global WD prediction (Boonman et al., 2020). This improvement in precision may be attributed to the large number and dense occurrence of sample sites as well as the inclusion of vegetation indices in our study. In addition, SLA and LPC also showed good accuracy with R^2 values of 0.50, which was higher than that of Boonman et al. (2020) and consistent with that of Moreno-Martínez~~Martínez~~ et al. (2018). However,

LNC and LA showed relatively poor performance, which may be related to the reason that these two traits were more influenced by phylogeny than environmental variables (Yang et al., 2017; An et al., 2021).

The frequency distribution of plant functional traits in China differed between our study and previous studies (Fig. 7, Fig. F1, Table F1 in Appendix F). Given that the spatial resolution of trait maps in most previous studies is 0.5° (except for Moreno-Martínez et al. (2018) and Vallicrosa et al. (2022)), we resampled the data products of previous studies and our study to 0.5° spatial resolution. The distribution in our study contained more predictions at lower values of SLA, LNC and LPC and was broader than those for SLA and LNC in previous global studies. However, the distribution of LNC in our study was consistent with that in Vallicrosa et al. (2022) at the 1 km spatial resolution (Fig. F1 in Appendix F). LA in our study contained more predictions at higher values and was also broader than those in previous global studies. WD did not show the lower and higher predictive values, however, the WD values in the studies of Boonman et al. (2020) and Schiller et al. (2021) had more predictions at higher values and no lower values (< 0.3 g cm^{-3}). Our predicted values of SLA showed the highest spatial correlation with those of Dong et al. (2023), and LNC showed the strongest spatial correlation with those of Butler et al. (2017) (Table 5). LA and WD showed the best spatial correlation with those of Schiller et al. (2021), but LPC showed relatively weak spatial correlation with those of published studies.

~~Our study also predicted the spatial pattern of SM with an R^2 of 0.24, which was lower than in the global study of Madani et al. (2018) in which environmental variables explained 45.5% of SM variation. The low precision of SM may be explained in two ways. First, with few sampling sites included in our study, the environmental gradients were narrow. Second, previous studies and our unpublished study have suggested that SM variation is primarily controlled by a phylogenetic effect, whereas the environmental effect is weak (Moles et al., 2006). Therefore, phylogenetic relatedness among species should be considered to improve the predictive precision of the spatial pattern of SM in further work.~~

~~There was no consensus in the spatial patterns of plant functional traits among the global studies. Thus~~In addition, we compared our results to the other studies focused on China. Yang et al. (2016) predicted the spatial distribution of leaf mass per area (1/SLA) and LNC based on trait-environment relationships in China and had an R^2 of 0.13-0.16. The lower predictive precision may be because Yang et al. (2016) only used MAT, MAP~~,~~ and RAD as predictors in estimating the spatial patterns of leaf mass per area and LNC, which likely led to poor performance and low heterogeneity. These results also demonstrated the advantage of our methods in mapping the spatial patterns of plant functional traits at a regional scale.

Table 5 Spatial correlations for SLA, LNC, LPC, LA and WD between this study and other previous trait maps, labelled by the first author of the corresponding publication (see Table F1 in Appendix F for citations)

Spatial correlation	Dong	Vallicrosa	Schiller	Boonman	Moreno	Madani	Butler	Bodegom
SLA	0.398		-0.082	0.327	0.242	0.136	-0.042	0.319
LNC	0.156	0.359	0.229	0.252			0.394	
LPC		0.136					0.057	
LA			0.514					
WD			0.647	0.107				

The spatial correlation of LDMC between our study and previous study was not included, as the LDMC maps were not available. SLA, specific leaf area (m^2 kg^{-1}); LNC, leaf N concentration (mg g^{-1}); LPC, leaf P concentration (mg g^{-1}); LA, leaf area (cm^2); WD, wood density (g cm^{-3}).

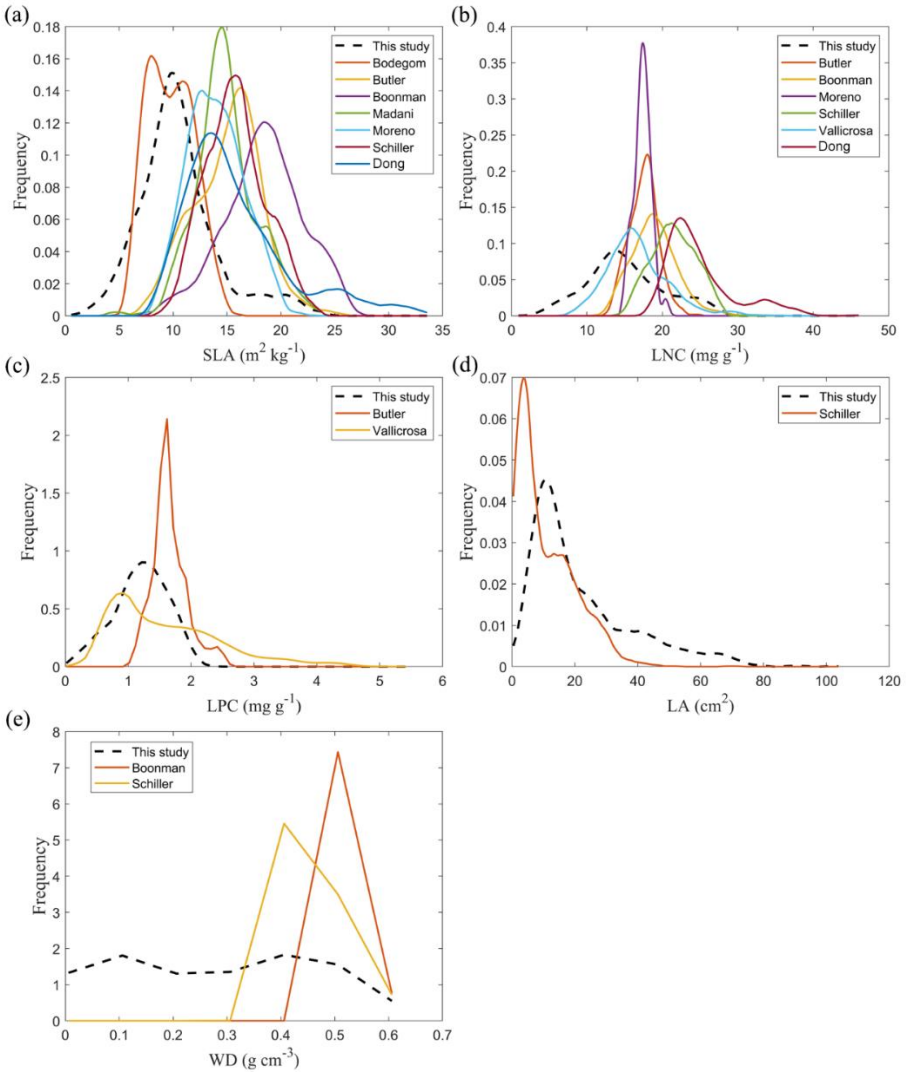


Figure 7. Frequency distributions of plant functional traits in our study ("This study", dashed black lines) and other trait maps, identified by the first author of the corresponding publication (see Table F1 for citations). SLA, specific leaf area (m^2 kg^{-1}); LNC, leaf N concentration (mg g^{-1}); LPC, leaf P concentration (mg g^{-1}); LA, leaf area (cm^2); WD, wood density (g cm^{-3}).

4.2 Spatial patterns of plant functional traits in China

Our study revealed the spatial patterns of different plant functional traits across China, and the variability among the two machine learning methods was relatively low. We compared the spatial differences of trait maps between our study and previous studies at the global scale ~~In some regions, there were consistent patterns in plant functional traits between our study and the previous studies~~(Figs. F2-F6 in Appendix F). For example, our study showed high SLA values in the southeastern Qinghai-Tibet Plateau, which concurred with the global study of Boonman et al. (2020). The spatial difference of SLA between our study and Bodegom et al. (2014) was relatively low, and the predictive values in most regions were slightly lower in our study than those in Bodegom et al. (2014). The spatial pattern of difference in SLA between our study and Moreno et al. (2018), Bulter et al. (2017) and Bodegom et al. (2020) was consistent, and the values were higher in northeastern China and southwestern Qinghai-Tibet Plateau in our study than those studies. Our study showed higher LNC values in the northern Inner Mongolia-the Loess Plateau-the eastern Qinghai-Tibet Plateau and northwestern China than those studies at the global studies (Butler et al., 2017; Moreno-Martínez et al., 2018; Boonman et al., 2020; Vallicrosa et al., 2022; Dong et al., 2023), reflecting the consistent spatial pattern among these studies. However, Yang et al. (2016) predicted high LNC values in northeastern and northwestern China, northern Inner Mongolia and the entire Qinghai-Tibet Plateau, and SLA and LNC had low heterogeneity overall. The discrepancy with Yang et al. (2016) may be attributed to spatial extrapolation based on trait-climate relationships with a low predictive precision. There was no consistent spatial pattern in LPC between our study and previous studies. Consistent with the global pattern (Wright et al., 2017), LA was larger in southern regions than in northern regions and showed a decreasing trend with latitude. In addition, LA and WD values in our study were lower in most regions than those ones at the global scale. These discrepancies between our study and previous studies at the global scale may be related to three reasons. First, there is bias in the available in-situ field measurement data from China in these global studies, with large gaps in western China for SLA and no data in China for WD (Boonman et al., 2020). Second, some trait-environment relationships may be scale dependent (Bruelheide et al., 2018), and these studies we compared are from the global scale due to the trait maps in China are not available. Third, the methods used for trait mapping were different among studies, including eco-evolutionary optimality models (Dong et al., 2023), Convolutional Neural Networks based on RGB photographs (Schiller et al., 2021), machine learning algorithms (Vallicrosa et al., 2022; Boonman et al., 2020) and multiple regression analysis (Bodegom et al., 2014).

~~Consistent with the global pattern (Wright et al., 2017), LA was larger in southern regions than in northern regions and showed a decreasing trend along a latitudinal gradient. Potapov et al. (2021) mapped global forest canopy height at a spatial resolution of 30 m by integrating GEDI and Landsat data, and their resulting spatial pattern of plant height in China was consistent with our study. This confirms the accuracy and reliability of predicting plant height through spatial extrapolation as in our study. However, in some regions there were contradictory patterns in the plant functional traits~~

~~between our study and previous ones. Our study showed high LNC values in the northern Inner Mongolia Plateau-Loess Plateau-eastern Qinghai-Tibet Plateau and high heterogeneity in southern China, whereas Yang et al. (2016) predicted high LNC values in northeastern and northwestern China, northern Inner Mongolia Plateau, and the entire Qinghai-Tibet Plateau, but low heterogeneity overall. In addition, Yang et al. (2016) predicted low SLA variability in China, especially in the Qinghai-Tibet Plateau. These discrepancies may be attributed to spatial extrapolation based on trait-climate relationships with a low predictive precision.~~

Moreover, our study also identified the applicability domain of our models for predicting the spatial patterns of plant functional traits across China. Five leaf traits and WD appeared to have poor applicability in northeastern China and the Qinghai-Tibet Plateau, primarily due to sparse samplings. ~~Although the predictions of plant height and SM were obtained in most regions through spatial extrapolation, the results were consistent with previous studies. This demonstrates the reliability of our methods.~~ Future studies predicting plant functional traits across a large scale through remote sensing observations or other supplementary data will be needed to re-evaluate our results.

4.3 The role of predictive variables

Our study indicates that environmental variables are important for predicting the spatial patterns of plant functional traits, especially climate variables. Temperature variables were primary predictors for SLA, LDMC, LPC and WD. The relationships between leaf traits and temperature have been widely discussed in global and regional studies (Reich and Oleksyn, 2004; Bruelheide et al., 2018). The positive linkage between WD and temperature may be driven by changes in the viscosity of water. Plants can adapt to the low water viscosity at high temperatures by reducing the diameter and density of their vessels and by thickening cell walls (Roderick and Berry, 2002; Thomas et al., 2004). Precipitation variables were important predictors for leaf nutrient traits~~,~~ and LA~~, and plant height~~. For example, precipitation of wettest quarter of a year was the factor that most influenced LA variation, which has been confirmed by a previous study (An et al., 2021). A smaller LA could be an adaptive strategy to decrease water loss via reducing the surface area for transpiration under dry environmental conditions (Du et al., 2019). ~~Water availability emerged as the primary predictor of plant height, which may be explained by the hydraulic limitation hypothesis. Relative to shorter plants, taller plants tend to be at a higher risk of embolisms due to their increased xylem length and conduit width. Thus, tall plants may be forced to close their stomata and reduce the amount of photosynthesis (Renninger et al., 2009; Wang et al., 2019).~~ Although the effects of soil on trait predictions were relatively weak, we found that SAP and pH played key roles in SLA~~,~~ and LNC ~~and SM~~ predictions. These results were similar with the previous studies that reported that soil pH was an important driver of trait variation at the global scale and in tundra regions (~~Kemppinen et al., 2021;~~ Maire et al., 2015; Kemppinen et al., 2021). Additionally, from the perspective of cost-efficient theory, the strong effects of SAP reflected that high SLA ~~and small seeds~~ may be an adaptation for facilitating soil exploration more efficiently in fertile soils (Freschet et al., 2010).

Vegetation indices have recently been proposed as important predictors of spatial patterns of plant functional traits (Loozen et al., 2018). Our results corroborate these findings and further suggest that EVI, MTCI and MIR reflectance are important predictors in models. Here, the underlying mechanisms between vegetation indices and plant functional traits are not further discussed due to their complexity and uncertainty. However, our results indicated that vegetation indices and NIR reflectance are not key predictors of LNC estimation, which contrasts the findings from global and regional studies (Wang et al., 2016; Loozen et al., 2018; Moreno-Martínez et al., 2018~~; Wang et al., 2016~~). This may be related to the multitude of factors that influence the relationship between LNC and vegetation indices and NIR reflectance, such as forest type and canopy structure (Dahlin et al., 2013).

4.4 Uncertainties

Although our study mapped the spatial patterns of key functional traits of seed plants in China through large-scale field investigations and compared the predictions with previous studies performed at global and regional scales, there persists some uncertainties in the interpretation of these results. First, the predictive ability of ~~machine learning method~~models was relatively worse for certain traits, especially ~~for SM and~~ LDMC. Beyond the environmental effects, the variation in plant functional traits is also regulated by phylogenetic structure among plant species (e.g., family, order and phylogenetic clade) (Li et al., 2017). Consequently, incorporating the phylogenetic information will be a promising avenue ~~in future studies~~ for further improving the accuracy of spatial predictions of plant functional traits (Butler et al., 2017). A second potential issue is sampling bias; there were major spatial gaps in field investigation in both the northeastern China and the Qinghai-Tibet Plateau.~~, especially for SM and plant height. There are an increasing number of available datasets and studies on SM and plant height, but many did not provide the corresponding geographic coordinates, thus rendering the data unusable. In addition, d~~Due to the ~~challenges~~ lack of measurements for small shrubs and low vegetation, WD data is mainly confined to eastern forests ~~(Perez-Harguindeguy et al., 2013)~~, and the overall quantity of WD data was much lower than that of leaf ~~and reproductive~~ traits, even in the TRY database~~, which is the largest trait database in the world (Kattge et al., 2020)~~. The environmental information of sampling sites was not always obtained from original literature, thus using the public environmental products is a common resolution in large-scale plant trait studies (Boonman et al., 2020; Vallicrosa et al., 2022). Such mismatch between in-situ trait measurements and predictors should be resolved in further work. Finally, additional key challenges in data availability must be resolved to scale up from the species to the community levels, in particular with data surrounding species co-occurrence and their relative cover or abundance in ecological communities (He et al., 2023). Global biodiversity data (e.g., sPlot and Global Biodiversity Information Agency databases) that contains information on species occurrence or the proportion of species in a community has the potential for enabling the calculation of community-weighted trait values and the re-evaluation of our results in future work (Telenius,

2011; Bruelheide et al., 2019). The lack of consistent time period and spatial resolution of predictors due to limitation of data availability is another key challenges in the spatial mapping of plant functional traits. In addition, although Worldclim version 2.1 product has high spatial resolution and includes various aspects of climatic parameters, there exists certain limitation and uncertainty in predicting trait maps. ~~In addition,~~ Therefore, integrating satellite remote sensing monitoring methods with ~~in~~ in-situ trait data collection can also provide an effective way to estimate and assess the species diversity at a large scale (Cavender-Bares et al., 2022).

4.5 Potential applications

Maps of these key functional traits of seed plants highlighted large-scale variability in space, which will significantly advance ecological analyses and future interdisciplinary research. First, using the spatially continuous trait ~~datasets~~maps, one can optimize and develop trait-flexible vegetation models, which allows for the exploration of the community assembly rules based on how plants with different trait combinations perform under a given set of environmental conditions (Berzaghi et al., 2020). When trait-flexible vegetation models are available, incorporating trait maps into models will bridge the gap for vegetation classifications and predictions of vegetation distribution under global change, which is not feasible in conventional vegetation models (Van Bodegom et al., 2012; Yang et al., 2019). Second, the assessments of China's terrestrial ecosystem carbon sink have had large uncertainties so far (Piao et al., 2022), but the spatial continuous trait ~~datasets~~ maps will provide an effective way to link ecosystem characteristics to ecosystem carbon sink estimates in China (Madani et al., 2018; Šímová et al., 2019). These analyses will help shed light on the mechanisms underlying plant functional traits and terrestrial ecosystem carbon storage at a large scale.

5 Data availability

The original ~~eight~~ plant functional trait~~s~~ ~~dataset~~ data collected in this study that were used for machine learning models (named by Data file used for machine learning models.csv) and final maps of ~~eight~~ plant functional traits in terrestrial ecosystems in a GeoTIFF format across China (named by plant functional trait category) are now available for the private link https://figshare.com/s/c527c12d310cb8156ed2 (An et al., 2023). Once the article is accepted, we will publicly publish these ~~datasets~~ maps at the figshare website.

6 Conclusions

We ~~created~~ generated a set of spatial continuous trait ~~datasets~~ maps at a 1-km spatial resolution using machine learning methods in combination with field measurements, environmental variables and vegetation indices. Models for leaf traits (except for LDMC~~),~~) and WD ~~and plant height~~ showed good accuracy and robustness, whereas models of ~~SM and~~ LDMC had relatively poor precision and robustness. Temperature variables were the most important predictors for leaf traits (except for LA)

and WD, and precipitation variables were the most important predictors for leaf nutrient traits~~,~~ and LA ~~and plant height~~. We caution that plant functional trait predictions should be interpreted carefully for northeastern China and the Qinghai-Tibet Plateau. The spatial continuous trait ~~datasets~~ maps generated in our study are complementary to current terrestrial in-situ observations and offer new avenues for predicting large-scale changes in vegetation and ecosystem function under climate scenarios in China.

Appendix A Data collection from literature

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Appendix B

Table B1 Summary of statistics in plant functional traits, environmental variables and geographical distribution in China.

Trait	Unit	Range	Mean	CV (%)	No. of species	Entries	Sites
SLA	$m^2\ kg^{-1}$	0.06–81.68	17.88	54.96	2463	9195	1032
LDMC	$g\ g^{-1}$	0.06–0.95	0.34	100.00	1582	3957	193
LNC	$mg\ g^{-1}$	3.41–66.02	21.52	37.44	2335	7407	567
LPC	$mg\ g^{-1}$	0.09–9.70	1.83	62.19	2074	6266	515
LA	cm^2	0.0033–2553.33	36.16	259.64	1838	5976	691
~~Height~~	~~m~~	~~0.01–35.00~~	~~5.99~~	~~67.58~~	~~1171~~	~~16324~~	~~636~~
WD	$g\ cm^{-3}$	0.25–1.37	0.68	33.16	768	1788	639
~~SM~~	~~mg~~	~~0.10–201300.00~~	~~1185.59~~	~~562.32~~	~~1163~~	~~3080~~	~~134~~
Altitude	m	-144–5454					~~1541~~1430
MAT	°C	-12.07–24.32					~~1541~~1430
MAP	mm	15–2982					~~1541~~1430
Soil total N	$g\ kg^{-1}$	0.11–10.25					~~1541~~1430
Bulk density	$g\ cm^{-3}$	0.83–1.45					~~1541~~1430

SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density; ~~SM, seed mass;~~ MAT, mean annual temperature; MAP, mean annual precipitation.

987 **Table B2** List of all the predictors including environment and remote sensing variables used in this study.

Type of variables	Variable name	Abbreviations	Units	Time periods	Spatial resolution	Source
Climate	Mean annual temperature	MAT	°C	1970-2000	1 km	WorldClim version 2.1
	Mean diurnal range	MDR	°C	1970-2000	1 km	WorldClim version 2.1
	Temperature seasonality	TS	°C	1970-2000	1 km	WorldClim version 2.1
	Max temperature of warmest month	Tmin	°C	1970-2000	1 km	WorldClim version 2.1
	Min temperature of coldest month	Tmax	°C	1970-2000	1 km	WorldClim version 2.1
	Temperature annual range	TAR	°C	1970-2000	1 km	WorldClim version 2.1
	Isothermality	IS	%	1970-2000	1 km	WorldClim version 2.1
	Mean temperature of wettest quarter	MTEQ	°C	1970-2000	1 km	WorldClim version 2.1
	Mean temperature of driest quarter	MTDQ	°C	1970-2000	1 km	WorldClim version 2.1
	Mean temperature of warmest quarter	MTWQ	°C	1970-2000	1 km	WorldClim version 2.1
	Mean temperature of coldest quarter	MTCQ	°C	1970-2000	1 km	WorldClim version 2.1
	Mean annual precipitation	MAP	mm	1970-2000	1 km	WorldClim version 2.1
	Precipitation of wettest month	PEM	mm	1970-2000	1 km	WorldClim version 2.1
	Precipitation of driest month	PDM	mm	1970-2000	1 km	WorldClim version 2.1
	Precipitation seasonality	PS	%	1970-2000	1 km	WorldClim version 2.1
	Precipitation of wettest quarter	PEQ	mm	1970-2000	1 km	WorldClim version 2.1
	Precipitation of driest quarter	PDQ	mm	1970-2000	1 km	WorldClim version 2.1
	Precipitation of warmest quarter	PWQ	mm	1970-2000	1 km	WorldClim version 2.1
	Precipitation of coldest quarter	PCQ	mm	1970-2000	1 km	WorldClim version 2.1
	Aridity index	AI	/	1970-2000	1 km	Global CGIAR-CSI
	Solar radiation	RAD	kJ m^{-2} day^{-1}	1970-2000	1 km	WorldClim version 2.1
Topography	Elevation	/	m		1 km	SRTM 90m V4.1
Soil	Soil sand content	SAND	%	/	1 km	Shangguan et al. (2013)
	Soil silt content	SILT	%	/	1 km	Shangguan et al. (2013)
	Soil clay content	CLAY	%	/	1 km	Shangguan et al. (2013)
	Bulk density	BD	g cm^{-3}	/	1 km	Shangguan et al. (2013)
	Soil pH	pH	/	/	1 km	Shangguan et al. (2013)

Continued

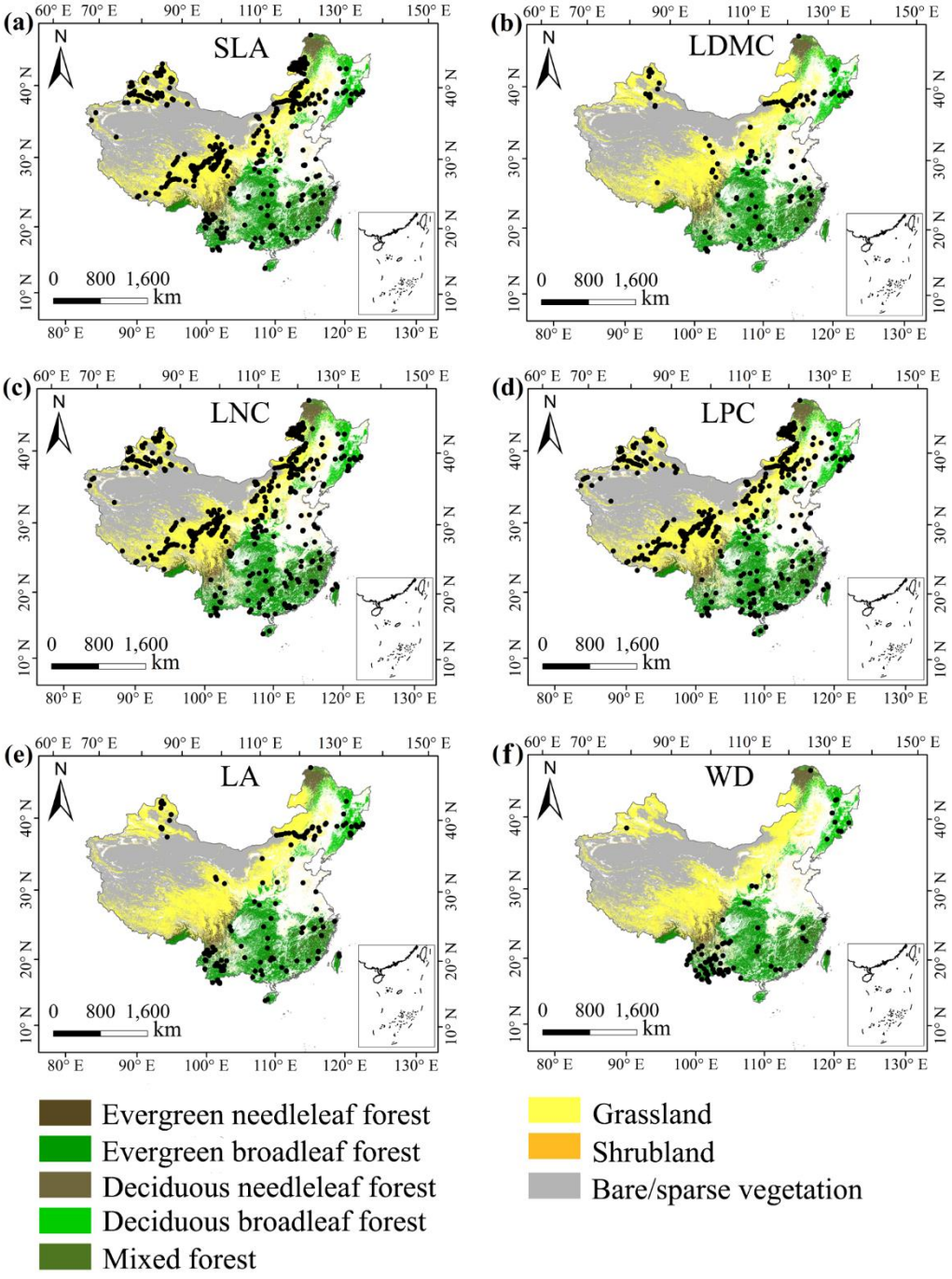
Type of variables	Variable name	Abbreviations	Units	Time periods	Spatial resolution	Source
	Soil organic matter	SOC	g kg^{-1}	/	1 km	Shangguan et al. (2013)
	Soil total N	STN	g kg^{-1}	/	1 km	Shangguan et al. (2013)
	Soil total P	STP	g kg^{-1}	/	1 km	Shangguan et al. (2013)
	Soil alkali-hydrolysable N	SAN	mg kg^{-1}	/	1 km	Shangguan et al. (2013)
	Soil available P	SAP	mg kg^{-1}	/	1 km	Shangguan et al. (2013)
	Soil available K	SAK	mg kg^{-1}	/	1 km	Shangguan et al. (2013)
	Cation exchange capacity	CEC	me kg^{-1}	/	1 km	Shangguan et al. (2013)
EVI	MODIS EVI long-term monthly averages		/	2001-2018	1 km	MOD13A3 V006
NIR	MODIS NIR long-term monthly averages		/	2001-2018	1 km	MOD13A3 V006
MIR	MODIS MIR long-term monthly averages		/	2001-2018	1 km	MOD13A3 V006
Red	MODIS red long-term monthly averages		/	2001-2018	1 km	MOD13A3 V006
Blue	MODIS blue long-term monthly averages		/	2001-2018	1 km	MOD13A3 V006
MTCI	MTCI long-term monthly averages		/	2003-2011	4.63 km	MTCI level 3 product
Land cover	Land cover map		/	2015	100 m	Copernicus Global Land Service Collection 3

The remote sensing variables are calculated as long-term monthly averages from 2001 to 2018. Thus 12 variables of each remote sensing category are obtained.

Table B3 The number of samples of eight plant functional trait used for model training (80%) and validation (20%).

Traits	No. of samples	No. of samples used for model training	No. of samples used for model validation
SLA	9195	7356	1839
LDMC	3957	3166	791
LNC	7407	5926	1481
LPC	6266	5013	1253
LA	5976	4781	1195
WD	1787	1430	357

SLA, specific leaf area (m^2 kg^{-1}); LDMC, leaf dry matter content (g g^{-1}); LNC, leaf N concentration (mg g^{-1}); LPC, leaf P concentration (mg g^{-1}); LA, leaf area (cm^2); WD, wood density (g cm^{-3}).



994

995 **Figure B1.** The distribution of sampling site of each plant functional traits across China. The black

996 dots represented the locations of trait observations.

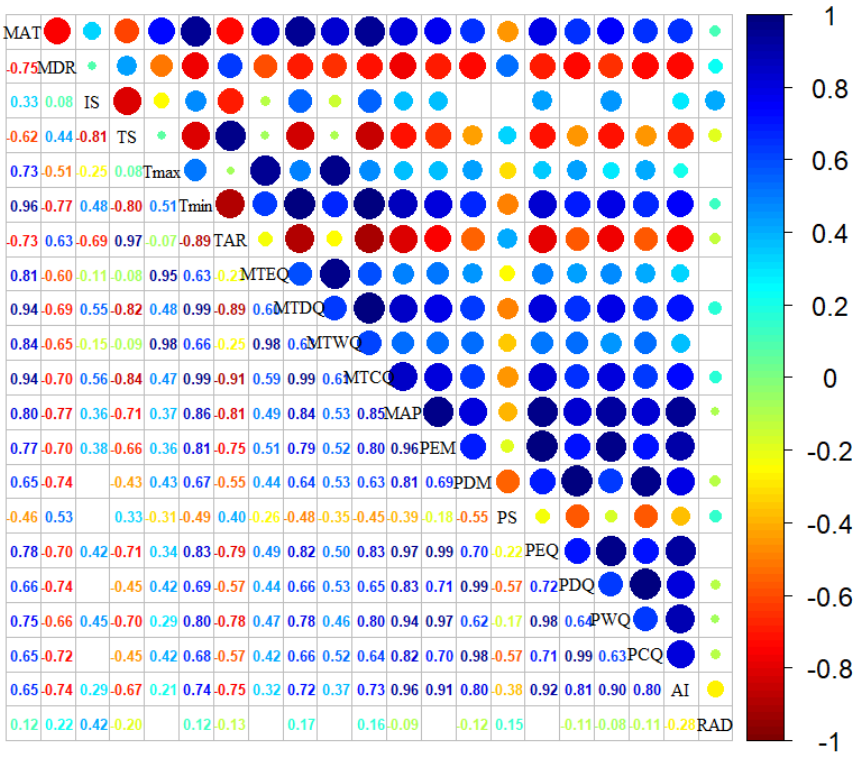


Figure ~~B1~~B2. Correlations among climate variables. The blank indicates that the correlations are not significant ($P > 0.05$). The size of the circles is proportional to the correlation coefficient. The abbreviation of climate variables is seen in Table B2.

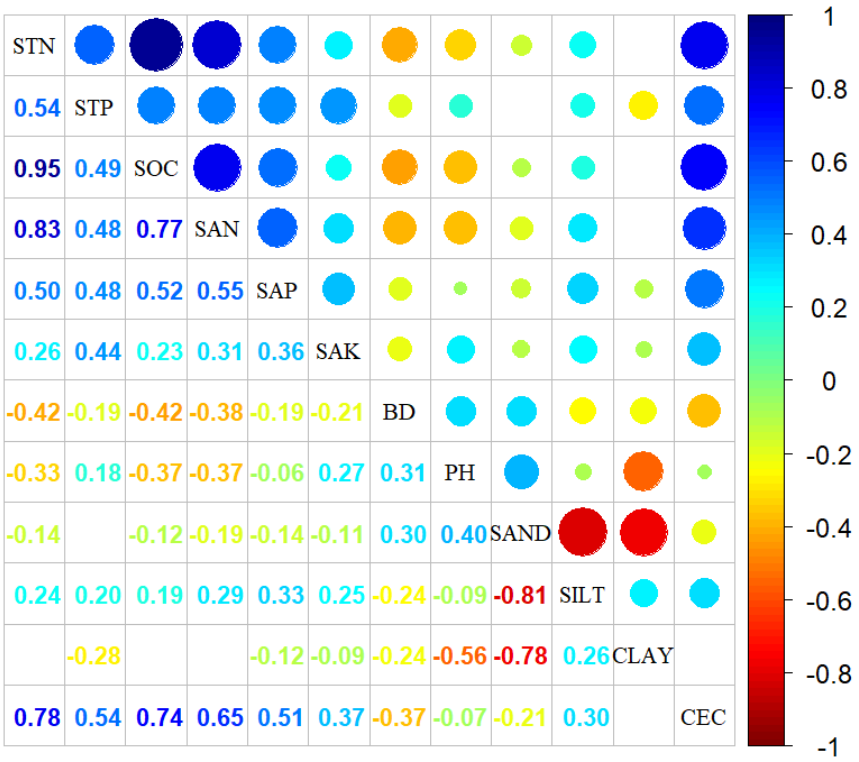


Figure ~~B2~~B3. Correlations among soil variables. The blank indicates that the correlations are not significant ($P > 0.05$). The size of the circles is proportional to the correlation coefficient. The abbreviation of soil variables is seen in Table B2.

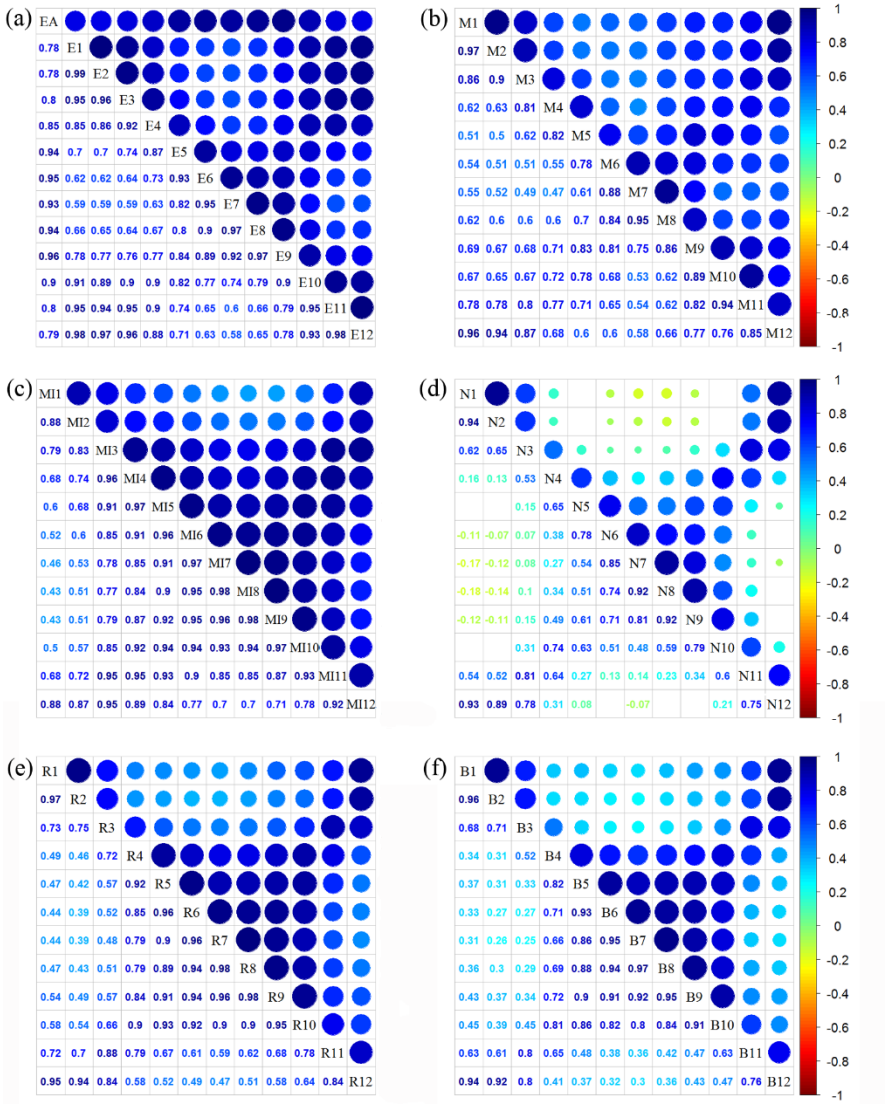


Figure ~~B3~~B4. Correlations among monthly remote sensing variables. The blank indicates that the correlations are not significant ($P > 0.05$). The size of the circles is proportional to the correlation coefficient. (a) enhanced vegetation index (EVI); (b) MERIS terrestrial chlorophyll index (MTCI); (c) MIR reflectance; (d) NIR reflectance; (e) red reflectance; (f) blue reflectance.

Appendix C

Table C1 Optimal parameter combination and model performance of random forest ~~(RF)~~ for plant functional traits

Traits	ntree	mtry	R^2	NRMSE	MAE
SLA	1000	24	0.476	~~7.049~~0.22	5.134
LDMC	1000	11	0.234	~~0.095~~0.20	0.072
LNC	1000	57	0.392	~~0.129~~0.00	0.098
LPC	1000	20	0.587	~~0.176~~0.05	0.129
LA	1000	18	0.278	~~72.996~~0.48	26.622
~~Height~~	~~1000~~	~~38~~	~~**0.871**~~	~~0.234~~	~~0.178~~
WD	1000	9	0.531	0.0~~9~~2	0.072
~~SM~~	~~1000~~	~~22~~	~~**0.197**~~	~~6043.95~~	~~1290.866~~

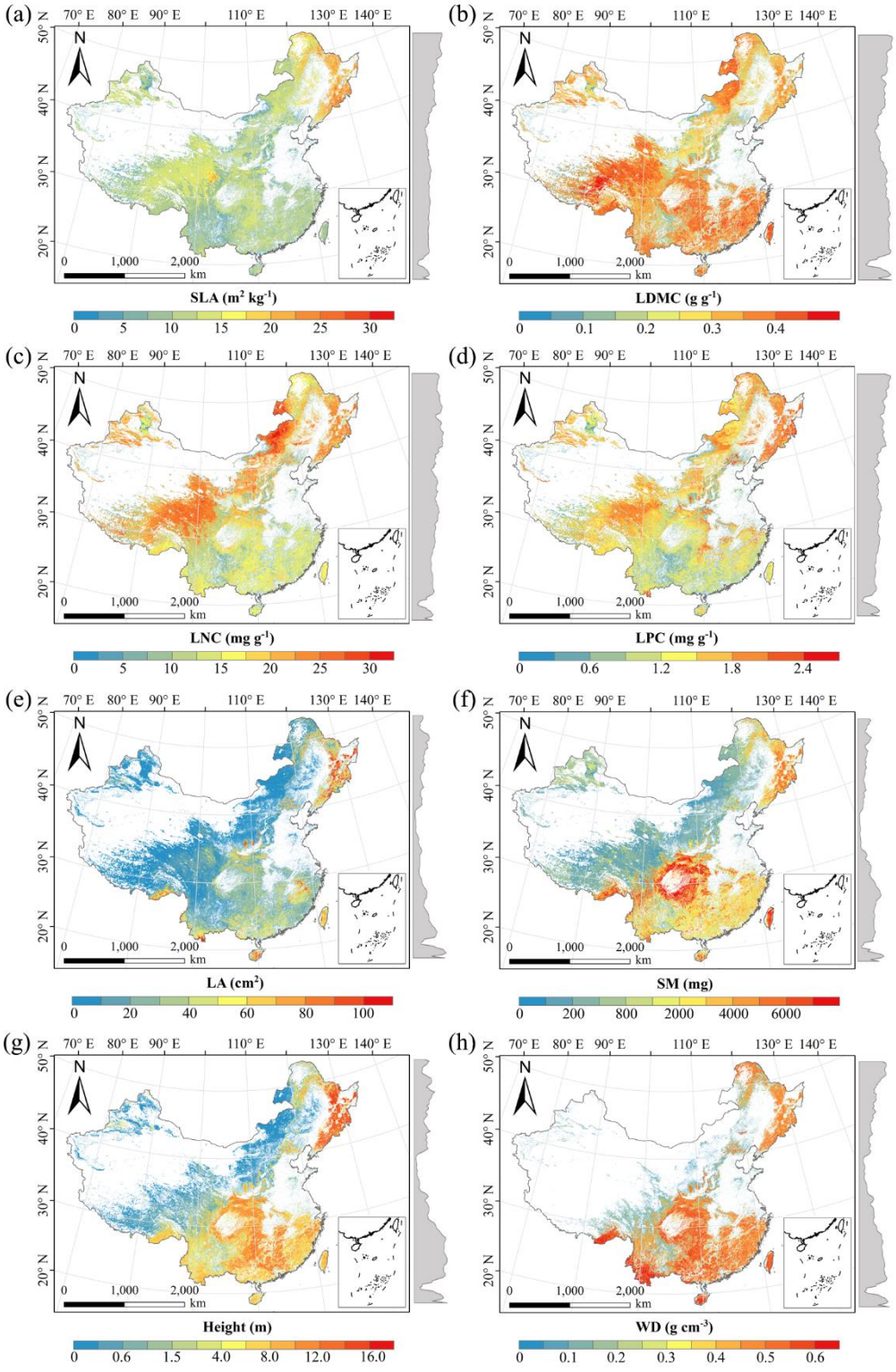
SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density~~; SM, seed mass~~.

Table C2 Optimal parameter combination and model performance of boosted regression trees ~~(BRT)~~ for plant functional traits

Traits	n.tree	interaction. depth	shrinkage	learning rate	bag fractions	R^2	NRMSE	MAE
SLA	3000	6	0.01	10	0.75	**0.486**	~~6.986~~0.20	5.082
LDMC	3000	2	0.01	10	0.75	**0.247**	~~0.094~~0.19	0.071
LNC	3000	6	0.01	10	0.70	**0.414**	~~0.126~~0.00	0.096
LPC	3000	7	0.01	10	0.75	**0.591**	~~0.175~~0.05	0.129
LA	3000	3	0.001	10	0.75	**0.282**	~~72.308~~0.55	27.556
~~Height~~	~~3000~~	~~3~~	~~0.05~~	~~10~~	~~0.6~~	~~0.871~~	~~0.234~~	~~0.178~~
WD	3000	4	0.01	10	0.70	**0.627**	~~0.082~~0.01	0.066
~~SM~~	~~3000~~	~~7~~	~~0.001~~	~~10~~	~~0.50~~	~~0.192~~	~~6070.703~~	~~1268.386~~

SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density~~; SM, seed mass~~.

Appendix D


(a)
70° E 80° E 90° E 110° E 130° E 140° E
50° N
40° N
30° N
20° N
N
0 1,000 2,000 km
SLA (m² kg⁻¹)
0 5 10 15 20 25 30
(b)
LDMC (g g⁻¹)
0 0.1 0.2 0.3 0.4
(c)
LNC (mg g⁻¹)
0 5 10 15 20 25 30
(d)
LPC (mg g⁻¹)
0 0.6 1.2 1.8 2.4
(e)
LA (cm²)
0 20 40 60 80 100
(f)
SM (mg)
0 200 800 2000 4000 6000
(g)
Height (m)
0 0.6 1.5 4.0 8.0 12.0 16.0
(h)
WD (g cm⁻³)
0 0.1 0.2 0.3 0.4 0.5 0.6

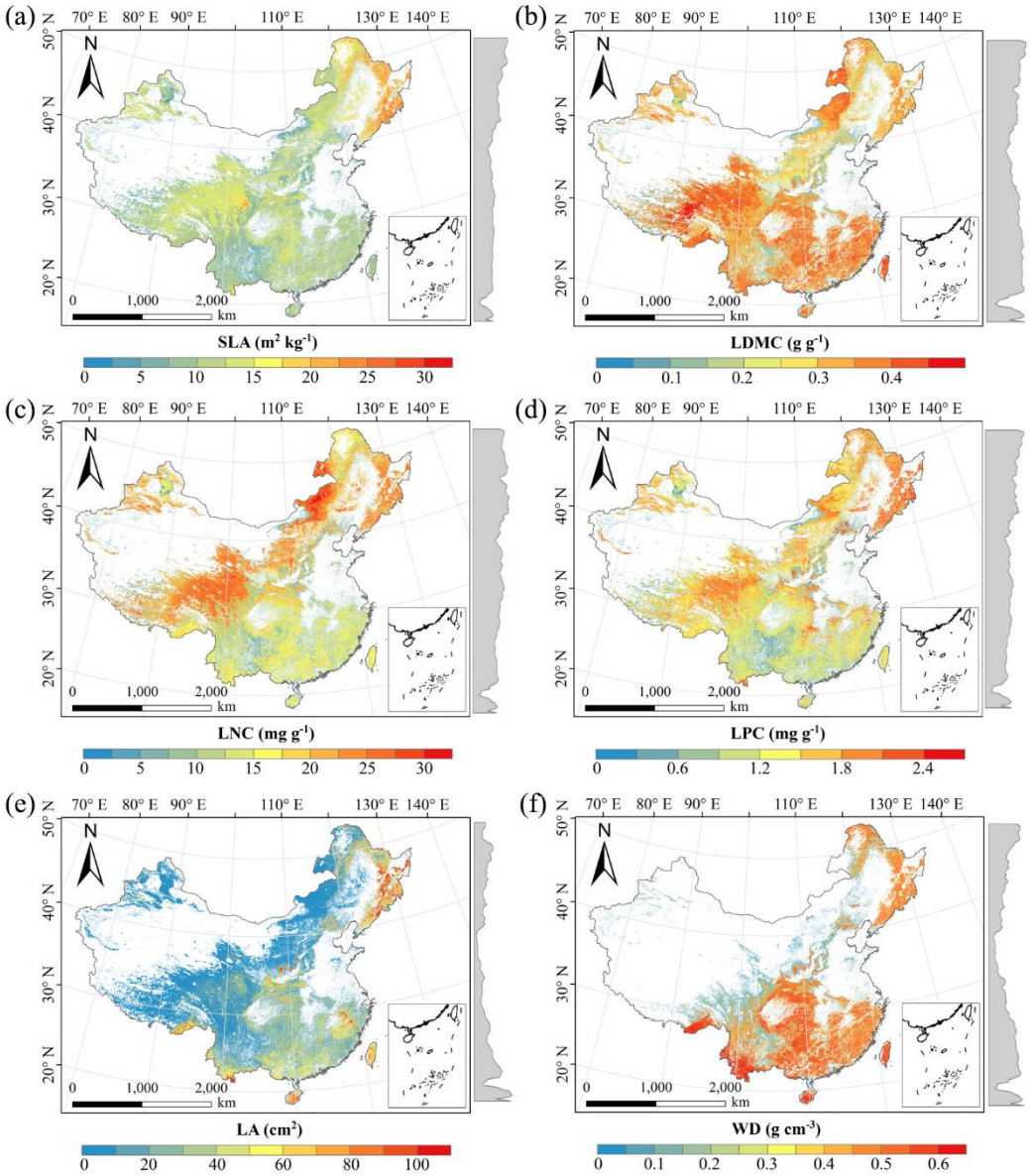
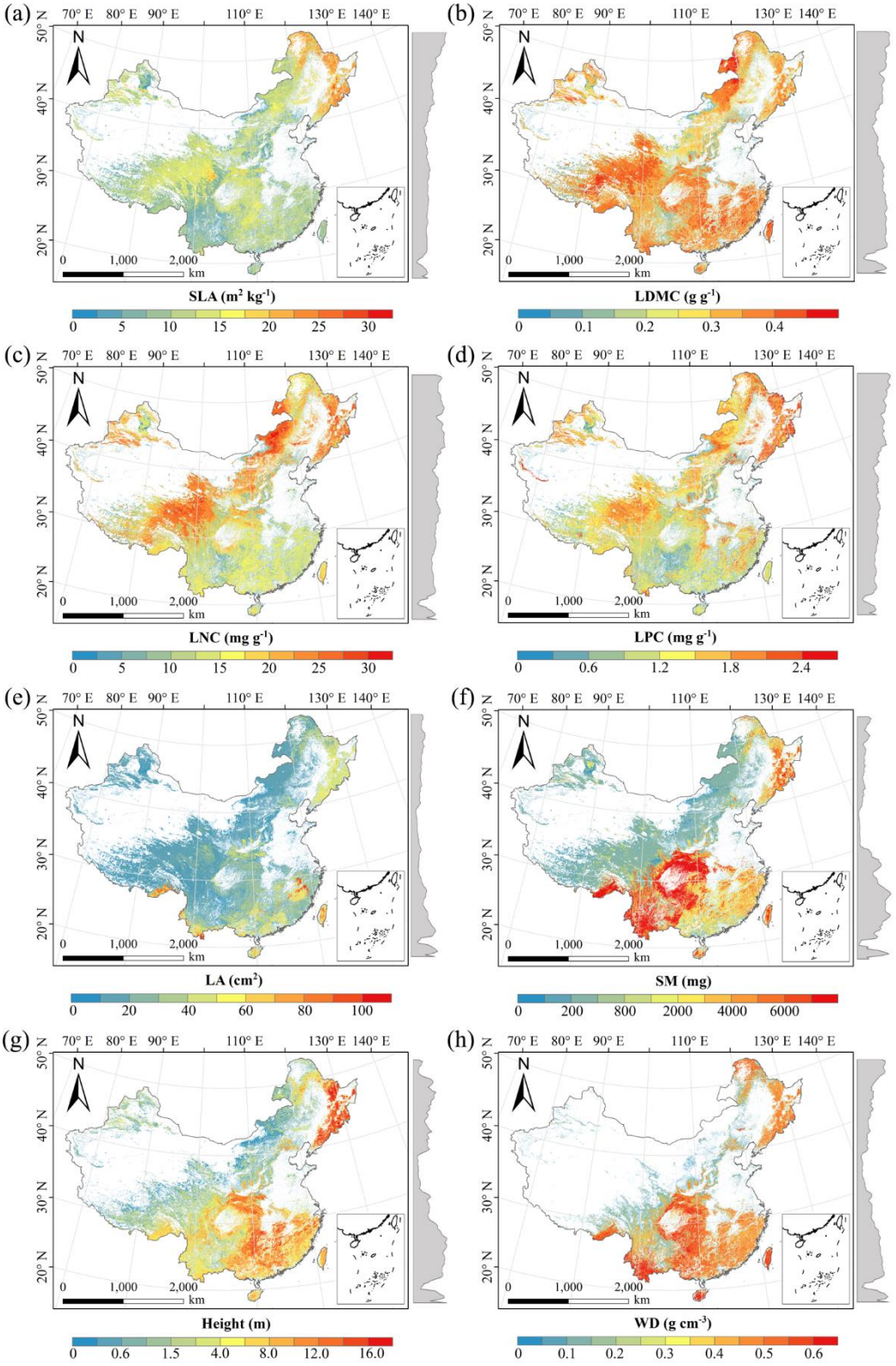


Figure D1. Spatial distribution of plant functional traits based on random forest ~~(RF)~~. The grey curves on the right of maps were trait distribution along with latitude. SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density~~; SM, seed mass~~.


(a)
SLA (m² kg⁻¹)
0 5 10 15 20 25 30
(b)
LDMC (g g⁻¹)
0 0.1 0.2 0.3 0.4
(c)
LNC (mg g⁻¹)
0 5 10 15 20 25 30
(d)
LPC (mg g⁻¹)
0 0.6 1.2 1.8 2.4
(e)
LA (cm²)
0 20 40 60 80 100
(f)
SM (mg)
0 200 800 2000 4000 6000
(g)
Height (m)
0 0.6 1.5 4.0 8.0 12.0 16.0
(h)
WD (g cm⁻³)
0 0.1 0.2 0.3 0.4 0.5 0.6
70° E 80° E 90° E 110° E 130° E 140° E
50° N 40° N 30° N 20° N
N
0 1,000 2,000 km

1027

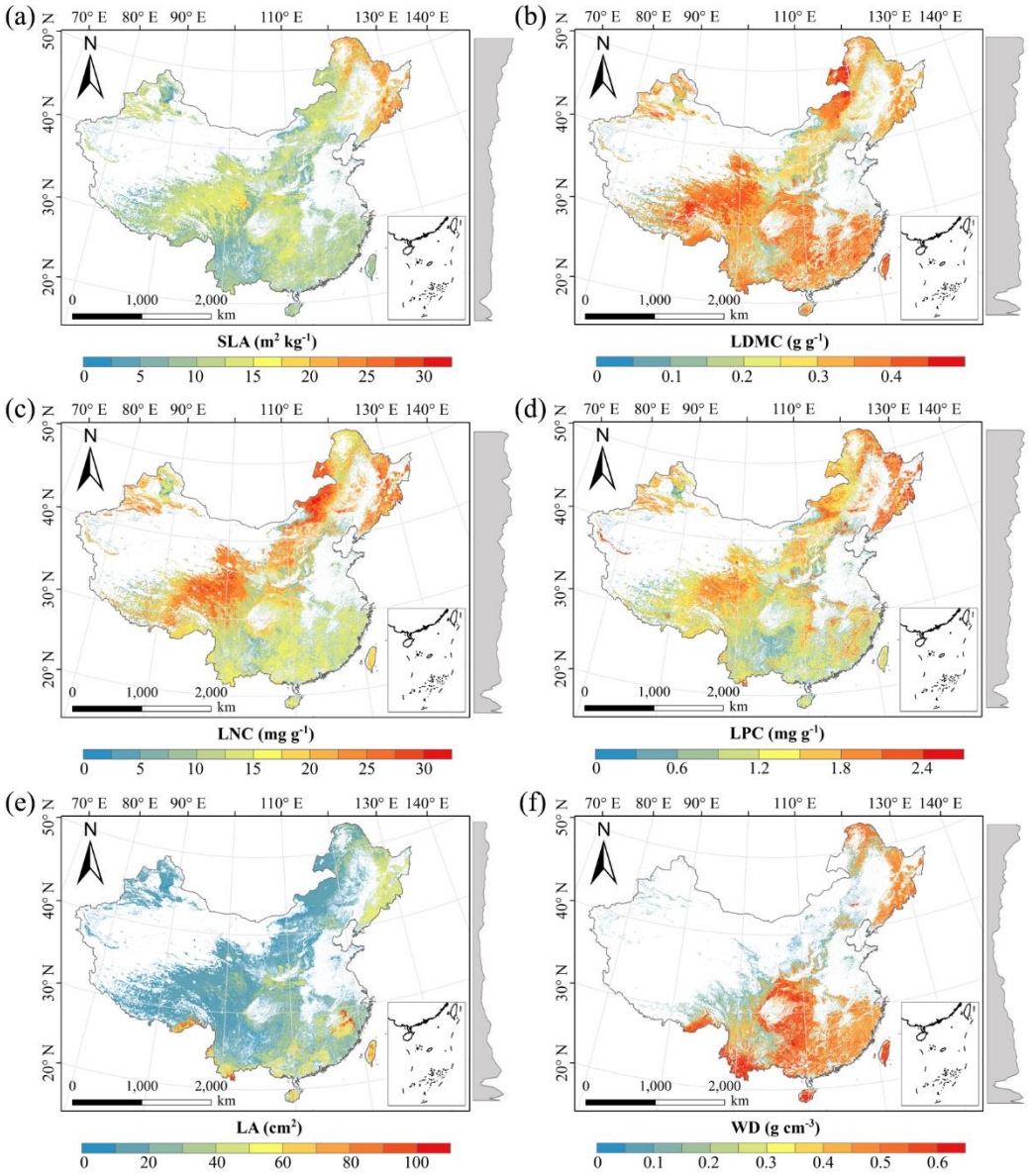


Figure D2. Spatial distribution of plant functional traits based on boosted regression trees ~~(BRT)~~. The grey curves on the right of maps were trait distribution along with latitude. SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration~~concertation~~; LPC, leaf P concentration~~concertation~~; LA, leaf area; WD, wood density~~; SM, seed mass~~.

Appendix E

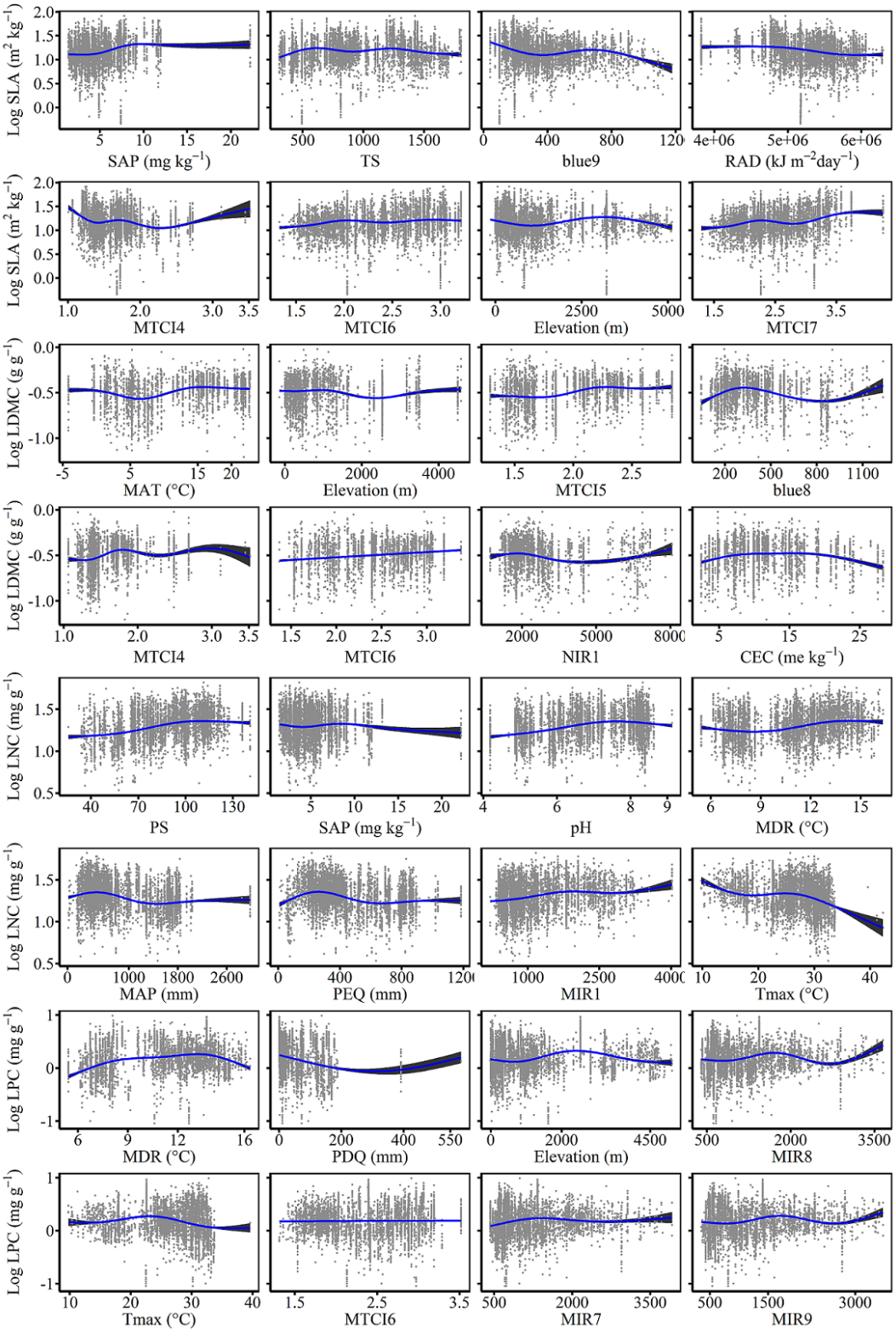
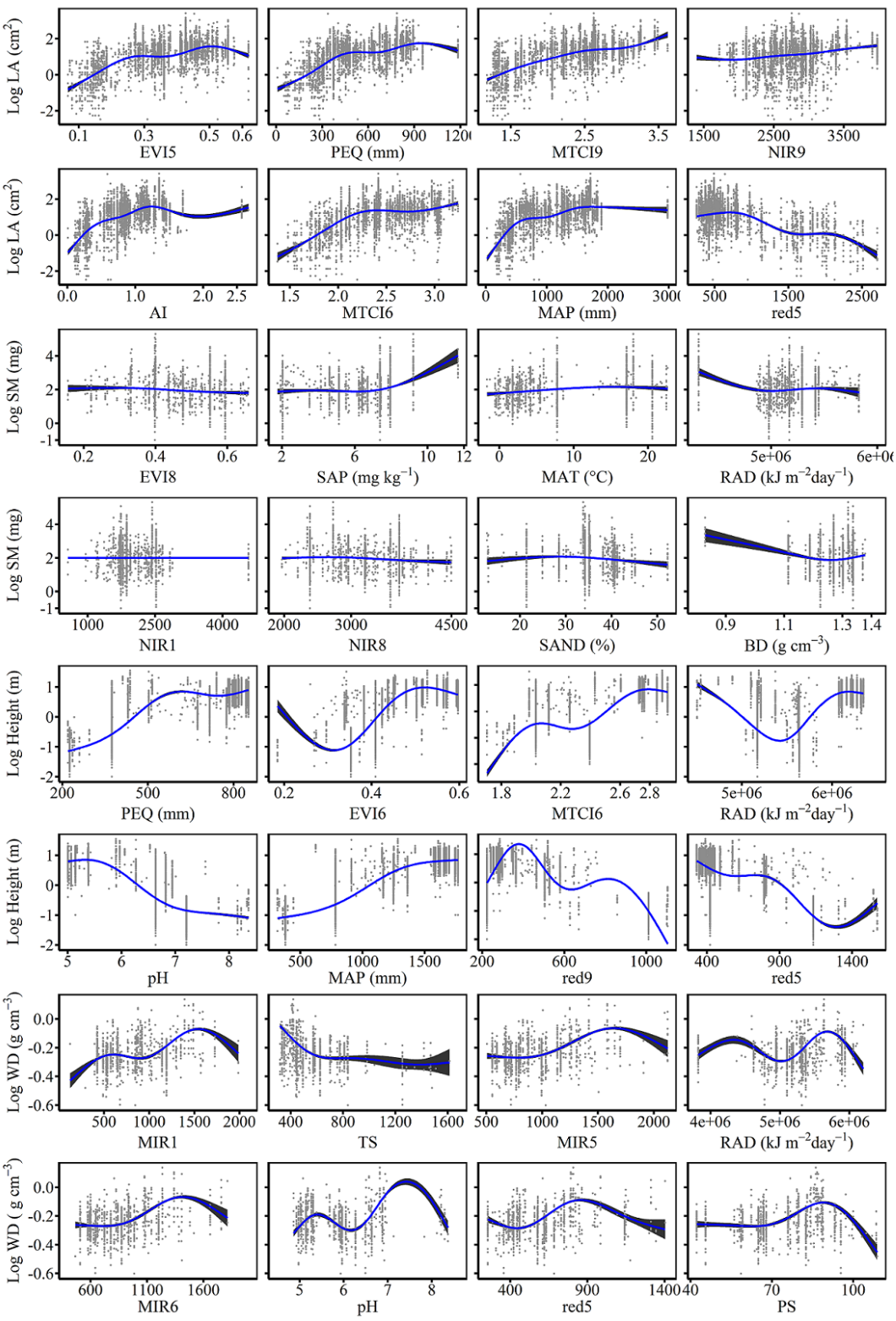


Figure E1. The relationships between SLA (specific leaf area), LDMC (leaf dry matter content), LNC (leaf N concentration), LPC (leaf P concentration) and their eight most important predictors.


Log LA (cm^2)
EVI5
PEQ (mm)
MTCI9
NIR9
AI
MTCI6
MAP (mm)
red5
Log SM (mg)
EVI8
SAP (mg kg^{-1})
MAT (°C)
RAD (kJ m^{-2}day^{-1})
NIR1
NIR8
SAND (%)
BD (g cm^{-3})
Log Height (m)
PEQ (mm)
EVI6
MTCI6
RAD (kJ m^{-2}day^{-1})
pH
MAP (mm)
red9
red5
Log WD (g cm^{-3})
MIR1
TS
MIR5
RAD (kJ m^{-2}day^{-1})
Log WD (g cm^{-3})
MIR6
pH
red5
PS

1037

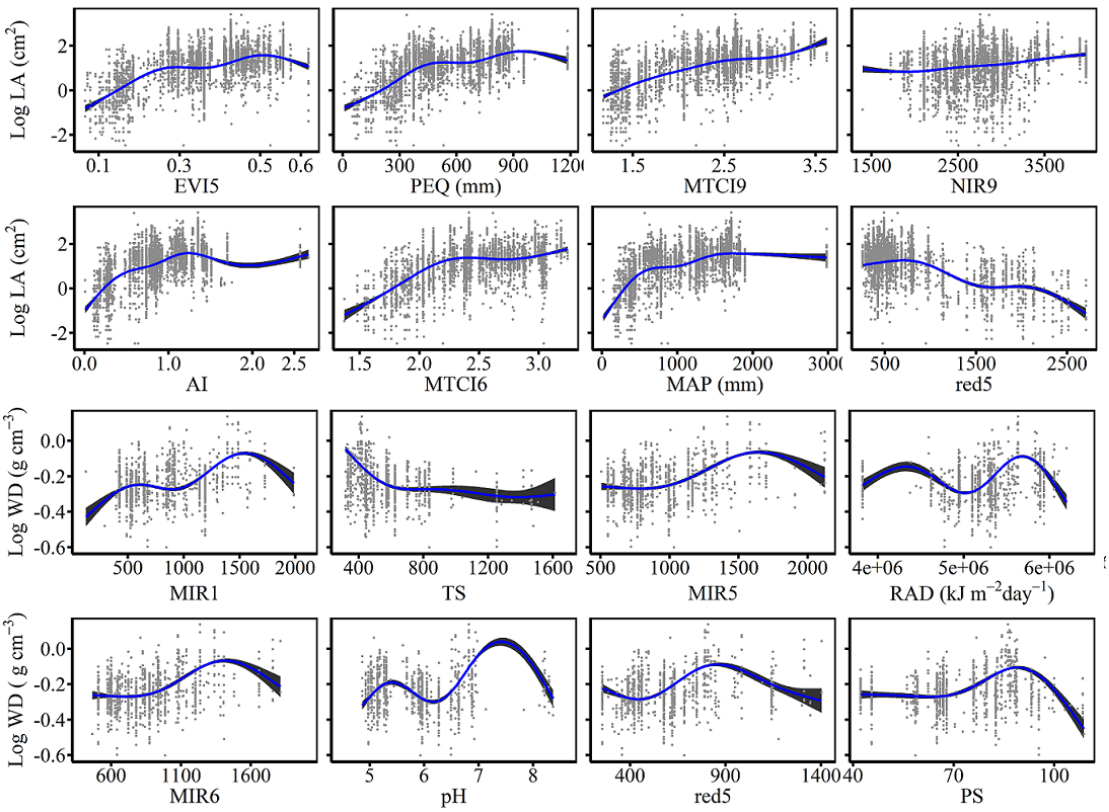


Figure E2. The relationships between LA (leaf area)~~, SM, Height,~~ WD (wood density) and their eight most important predictors.

Appendix F Comparisons between our study with trait maps from previous studies

Given that the trait maps predicted for China were not available from the literature and authors, we compared our study with those studies performed at the global scale (see Table F1). Thus, we extracted the data in China from global trait maps. Before the quantitative comparisons with previous studies, we performed two steps to make the data products as comparable as possible and improve the consistency between different studies. First, due to different spatial resolution of global trait maps (mainly 0.5°) and our study, we resampled the data products of previous studies and our maps to 0.5° spatial resolution. In addition, Vallicrosa et al. (2022) generated the global maps of LNC and LPC with a 1 km spatial resolution, we also compared the frequency distribution of Vallicrosa et al. (2022) with that of our study at a 1 km spatial resolution. Second, our study focused on natural vegetation, so the global trait maps were used to filter out non-natural vegetation (e.g., croplands). For example, Madani et al. (2018) predicted the spatial distributions of SLA that included croplands. We quantitatively compared our maps with previous studies from two perspectives. The comparisons among trait maps were made using frequency plots and spatial correlations (Figure 7 and Table 5). And the maps of spatial differences between our study and previous studies were displayed as Figs F1-F5 in Appendix F.

Table F1. Summary table of related trait maps of previous studies used in this study.

References	Related traits	Methods	Predictors	Consideration of PFT	Resolution
Dong et al. (2023)	SLA LNC	Optimality models	Climate	Yes	0.5°
Vallicrosa et al. (2022)	LNC LPC	Neural networks	Climate Soil N and P deposition	Yes	0.0083°
Schiller et al. (2021)	SLA LNC LA WD	Convolutional Neural Networks	Climate In-situ RGB images	No	0.5°
Boonman et al. (2020)	SLA LNC WD	Generalized linear model, Generalized additive model, Random forest, Boosted regression trees, Ensemble model	Climate Soil	No	0.5°
Moreno et al. (2018)	SLA LNC LPC LDMC	Regularized linear regression, Random forest, Neural networks, Kernel networks	Climate Elevation Reflectance	Yes	0.0045°
Madani et al.	SLA	Generalized additive	Climate	No	0.5°

(2018)		model			
Butler et al. (2017)	SLA LNC LPC	Bayesian model	Climate Soil	Yes	0.5°
Bodegom et al. (2014)	SLA WD	Multiple regression analysis	Climate Soil	No	0.5°

The resolutions 0.5°, 0.0083° and 0.0045° correspond to square grid cell sizes of about 50 km, 1 km and 500 m at the equator. PFT, plant functional type; SLA, specific leaf area; LDMC, leaf dry matter content; LNC, leaf N concentration; LPC, leaf P concentration; LA, leaf area; WD, wood density.

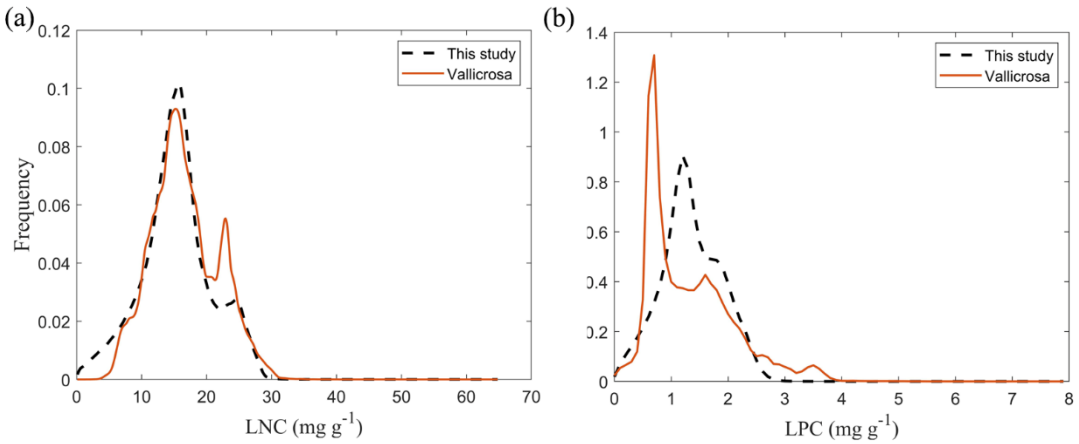


Figure F1. Frequency distributions of plant functional traits in our study ("This study", dashed black lines) and Vallicrosa et al. (2022) at 1 km spatial resolution. LNC, leaf N concentration (mg g^{-1}); LPC, leaf P concentration (mg g^{-1}).

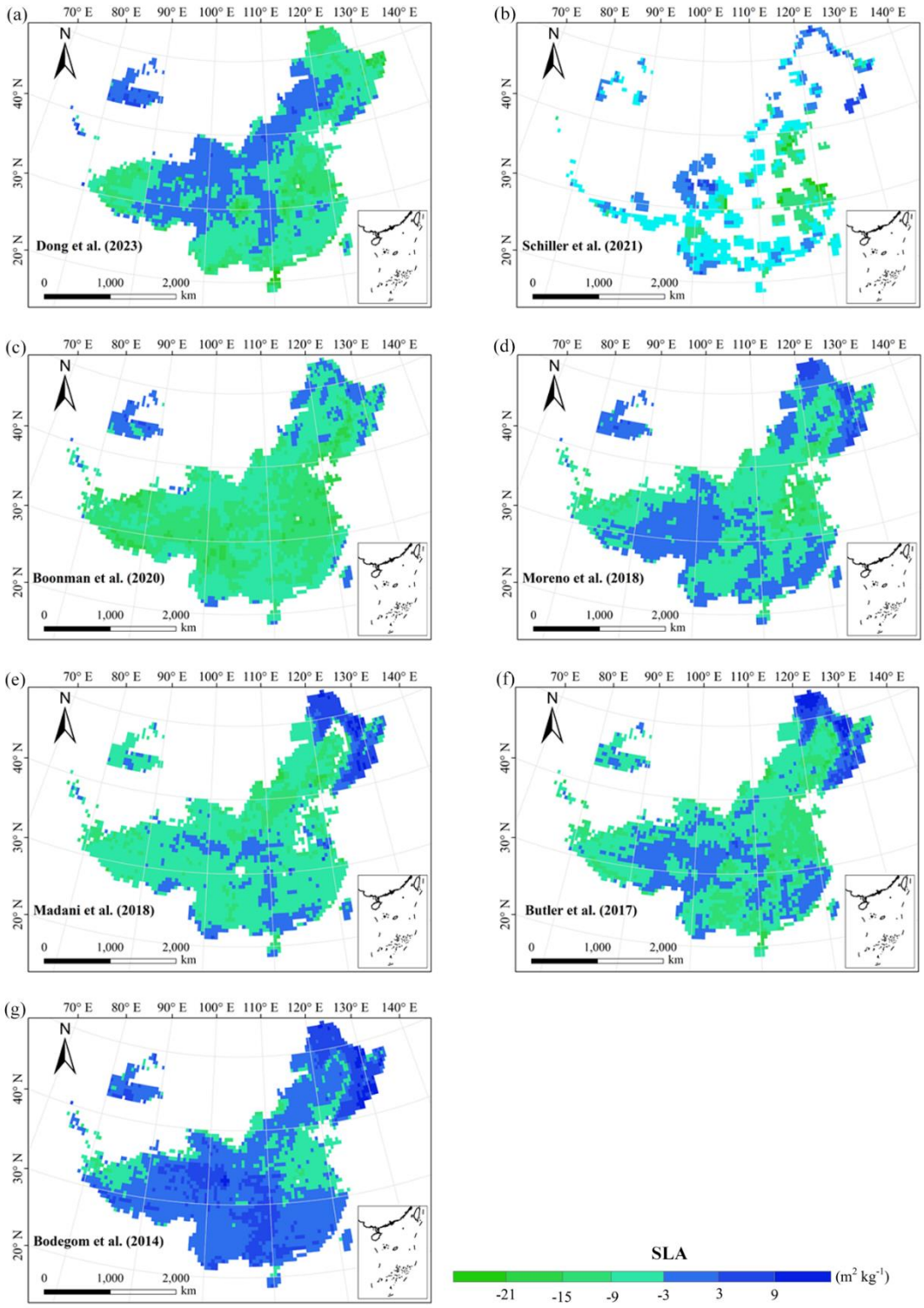


Figure F2. Spatial differences in SLA (specific leaf area, $m^2\ kg^{-1}$) between our study and trait maps from previous studies (see Table F1 for citations).

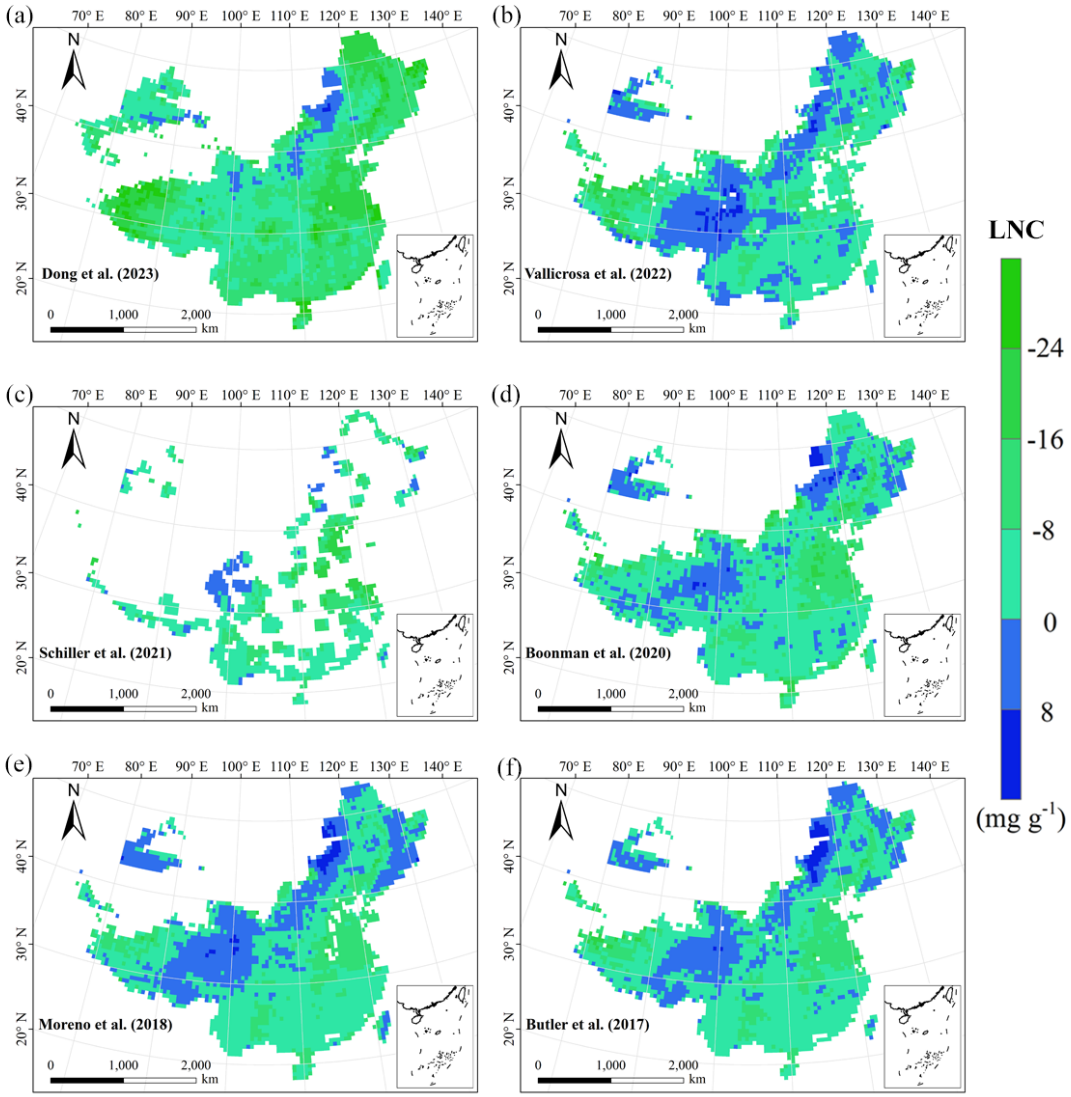


Figure F3. Spatial differences in LNC (leaf N concentration, mg g^{-1}) between our study and trait maps from previous studies (see Table F1 for citations).

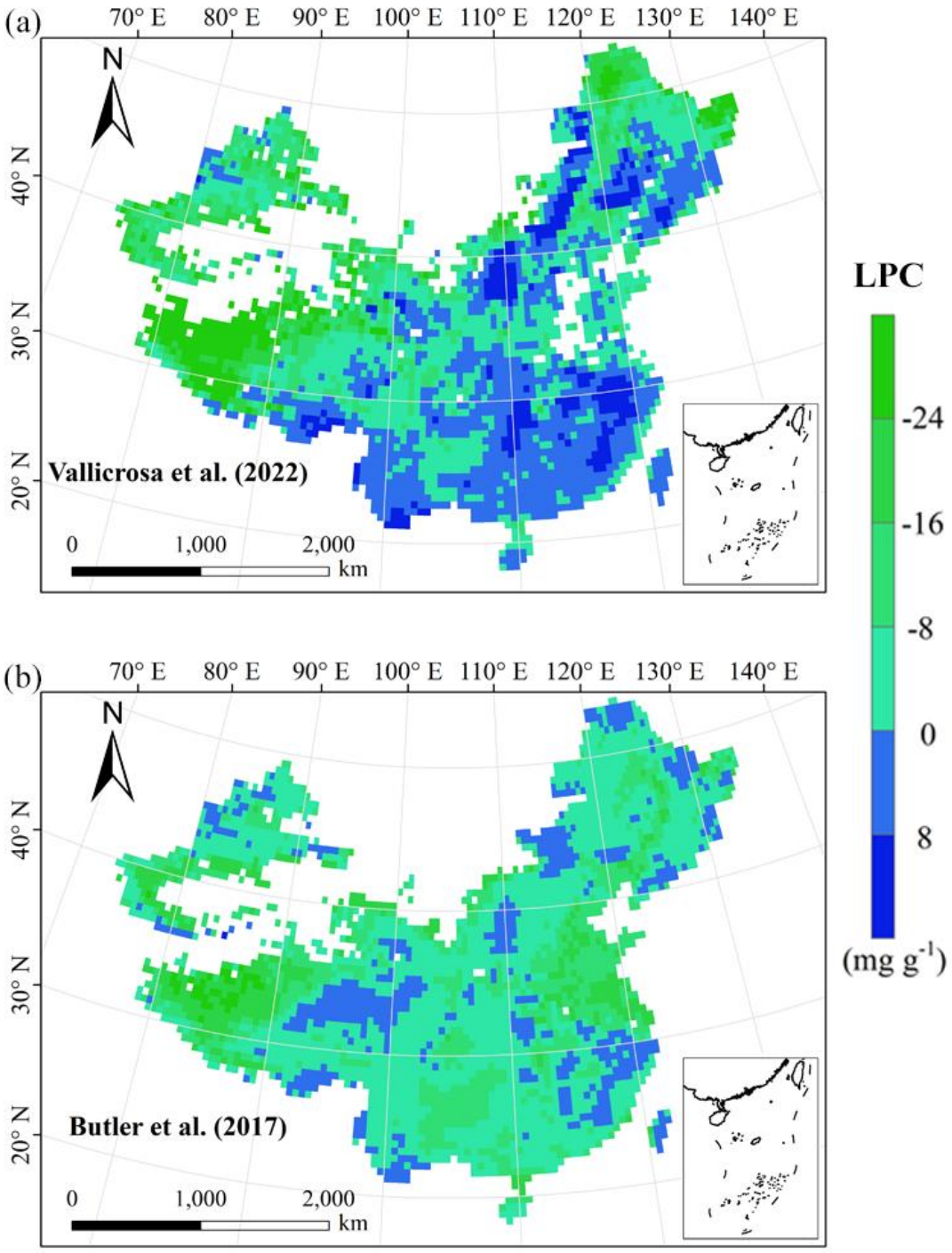


Figure F4. Spatial differences in LPC (leaf P concentration, mg g^{-1}) between our study and trait maps from previous studies (see Table F1 for citations).

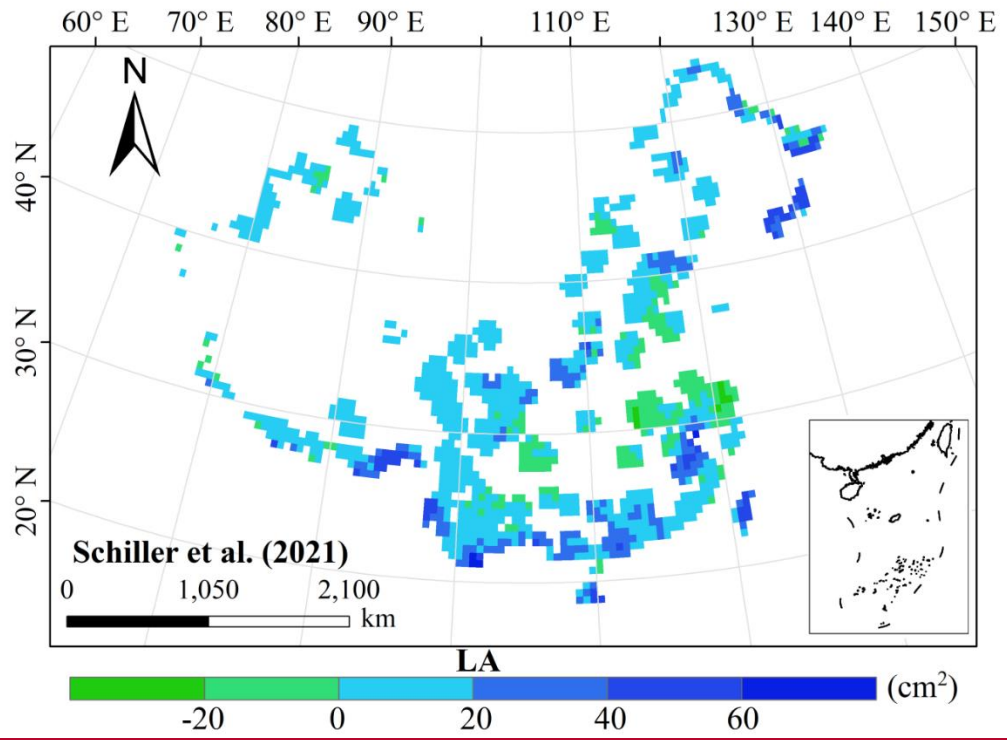


Figure F5. Spatial differences in LA (leaf area, cm^2) between our study and trait maps from previous studies (see Table F1 for citations).

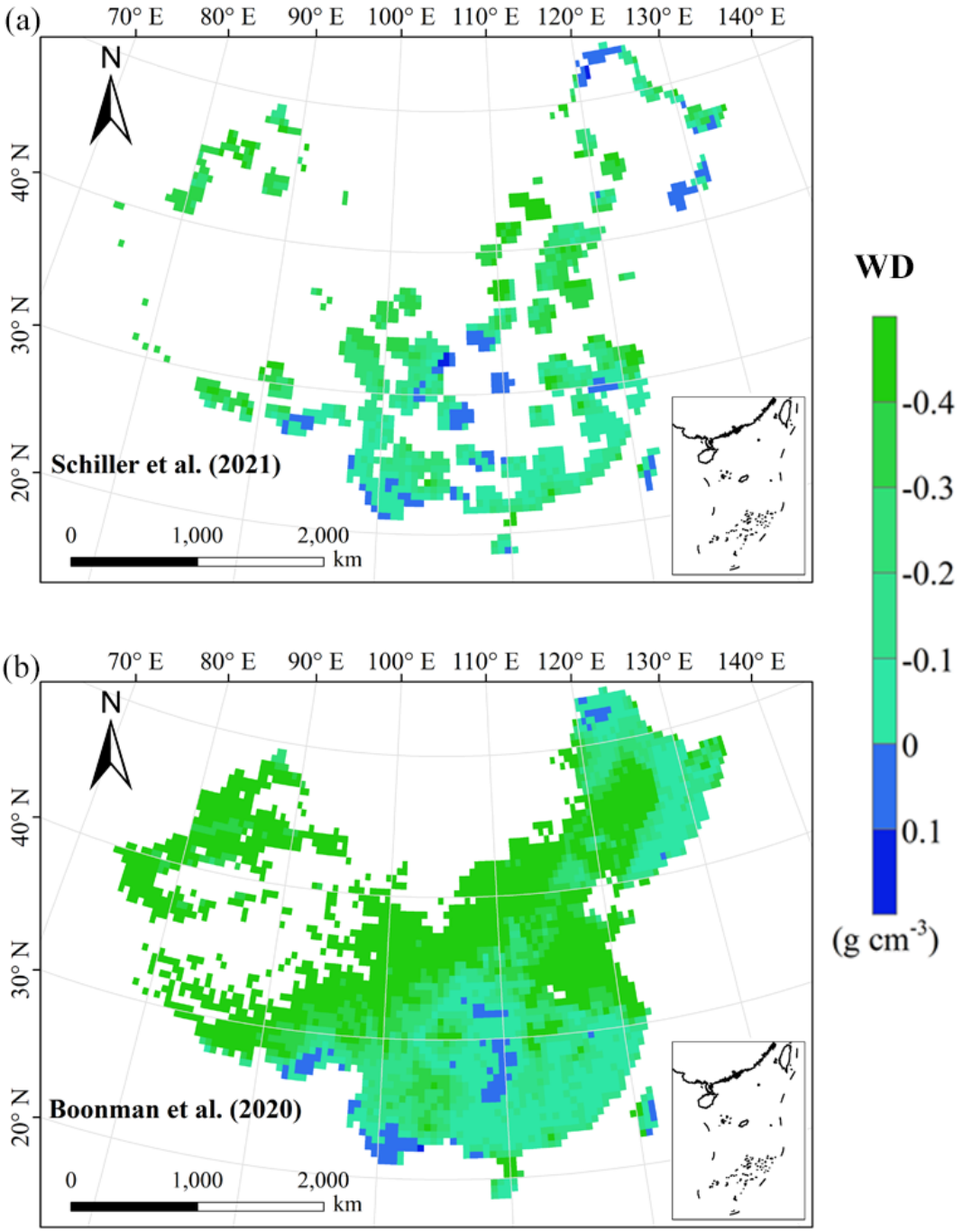


Figure F6. Spatial differences in WD (wood density, g cm^{-3}) between our study and trait maps from previous studies (see Table F1 for citations).

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