Original article

The patterns of vascular plant discoveries in China

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ABSTRACT

Botanical discovery has a long journey of revelation that contributes unparalleled knowledge

to shape our understanding about nature. Plant discovery in China is an immanent part of that

journey. To understand the patterns of plant discoveries in China and detect which taxa and

areas harbor most numbers of undiscovered species, we analyzed the discovery times of

31093 vascular plant species described in *Flora of China*. We found that species with larger

range size and distributed in northeastern part of China have a higher discovery probability.

Species distributed on the coast were discovered earlier than inland species. Trees and shrubs

of seed plants have the highest discovery probability and ferns have the lowest discovery

probability. Herbs hold the largest number of undiscovered species in China. Most

undiscovered species are found in southwest China, where three global biodiversity hotspots

locate. Spatial patterns of mean discovery year and inventory completeness are mainly driven

by the total number of species and human population density in an area and whether the area

is coastal or not. Our results showed that socio-economic factors dictate the discovery

patterns of vascular plants in China. Undiscovered species are mostly narrow-ranged,

inconspicuous endemic such as herbs, which are prone to extinctions and locate in

biodiversity hotspots in southwest China. We suggest that the future effort on plant discovery

in China be prioritized in southwest China.

INTRODUCTION

Despite 270 years' discovery of species since Linnaeus, our knowledge about the biological diversity in nature is still far from complete. According to a recent estimate, fewer than 20% of species on earth have been so far discovered (Mora *et al.*, 2011). With the ongoing biodiversity crisis, this lack of knowledge has become a major obstacle to biodiversity conservation as many species could go extinct without ever being known to science (Costello, May, & Stork, 2013; Humphreys *et al.*, 2019).

Early botanical and zoological discoveries are often biased towards large-sized, charismatic species with widespread geographic distributions (Gibbons *et al.*, 2005; Ferretti *et al.*, 2008; Stork *et al.*, 2008, 2015; Essl *et al.*, 2013; Randhawa, Poulin, & Krkošek, 2015). Positive correlations between species discovery probability and body size have been found in a variety of taxa including insects (Gaston & Hudson, 1994), birds (Blackburn & Gaston, 1995), mammals (Paxton, 1998; Medellín & Soberón, 1999; Collen, Purvis, & Gittleman, 2004), fishes (Zapata & Robertson, 2007), but not in taxa like reptiles (Reed & Roback, 2002) and marine holozooplanktons (Gibbons *et al.*, 2005). More recently discovered species are often of greater conservation interest because they are more likely to be narrow ranged and rare, and thus are prone to extinctions (Diniz-Filho *et al.*, 2005; Bebber *et al.*, 2007a, 2010; Tedesco *et al.*, 2014; Xu *et al.*, 2019).

In addition to biological factors, species discovery is also influenced by human factors such as taxonomic effort, technology innovations, and social-economic events. For example, it is commonly observed for many taxa that species discovery rates dropped during the two World Wars and peaked in the 1990s with the emergence of molecular techniques (Gaston,

1995; Bebber *et al.*, 2007b; Joppa, Roberts, & Pimm, 2011b; Lu & He, 2017). Geographically

speaking, Europe and North America have the most complete species inventories due to their

long histories of exploration (Gaston, 1995; Essl *et al.*, 2013), while species discovery in less

explored continents such as South America and Africa was affected by colonization histories

and indigenous knowledge (Gaston, 1995; Diniz-Filho *et al.*, 2005; Rosenberg *et al.*, 2013;

Ballesteros-Mejia *et al.*, 2013). As a result, biodiversity hotspots, most of which locate in less

developed countries (Myers *et al.*, 2000), often harbor the largest number of undiscovered

species (Joppa *et al.*, 2011a; Giam *et al.*, 2012). This imposes a more serious challenge for

biodiversity conservation in developing regions where economic growth is often achieved at

the expense of environmental degradation (He, 2009).

China harbors nearly one tenth of the plant species on earth (Joppa *et al.*, 2011b; Lu &

He, 2017). However, its rapid economic growth over the past three decades has resulted in

the colossal loss of millions of hectares of natural habitats (He, 2009). The sustainable

development of China depends on balancing economic growth and preservation of natural

habitats. Knowing where undiscovered species may locate is necessary for making decision

on habitat protection and conservation managements.

The interest for cataloging species in China has long predated the invention of Linnaeus'

binomial nomenclature, but in the light of modern taxonomy, much credit should be given to

western naturalists who diligently collected specimens and described species since the first

arrival of Jesuits in the 16th century, as reflected by the fact that nearly 70% of the type

specimens of Chinese plants are kept in herbaria in Europe and America (Chen, 1994). Due to

logistic constraints and political instability, most naturalists in the 18th and 19th centuries

made their botanical collections in the coastal areas (Bretschneider, 1898; Fan, 2004), which

likely had affected the patterns of discovery records.

There are more than 31000 vascular plant species documented in China, but this

inventory is not complete and many new species still await discovery (Lu & He, 2017).

Knowing what are those inconspicuous species and where they may locate is important to

discover them. Therefore, the objectives of this study are: (1) to model and map geographic

variation in botanical discovery of vascular plant species in China, (2) to find out which taxa

have the largest numbers of undiscovered species and where the undiscovered species most

likely are, and (3) to quantify what factors (e.g., species' range size and growth form) may

influence the spatial distribution of plant discoveries in China. This study will contribute to

understanding the pattern of species discoveries and their underlying factors and should be of

significance to botanical discoveries of other regions beyond China. The identification of

taxonomic and geographic gaps of undiscovered species will facilitate prioritizing our

taxonomic and conservational efforts in the future.

MATERIALS AND METHODS

Data

Data including species names, authorities, discovery years, province-level biogeographic

distributions, and genus-level growth forms were compiled from *Flora of China* (FOC) (Data

available in Appendix S2) which has a totol number of ~31000 species (http://efloras.org).

We treated discovery year as the time a species was first described in any name in a scientific

publication. When estimating the number of undiscovered species, data after 2000 were

excluded as a routine to avoid the effect of delayed entrance of newly discovered species

(Costello & Wilson, 2011; Costello, Wilson, & Houlding, 2012). Human population densities

on province-level were obtained from *2010 Population Census of The People's Republic of*

China (http://www.stats.gov.cn/), and *Monthly Bulletin of Interior Statistics*

(http://sowf.moi.gov.tw/stat/month/elist.htm). Province areas were obtained from *National*

Fundamental Geographic Information System of China (http://nfgis.nsdi.gov.cn/nfgis/). There

are in total 28 provinces after merging independent municipalities such as Beijing and

Shanghai to adjacent provinces. Range size, maximum/minimum latitude,

maximum/minimum longitude and whether or not a species is distributed in coastal areas

were obtained from province-level distributions. Genus-level growth forms were categorized

as fern, herb, shrubs/trees and vine/liana. Shrubs and trees were categorized as one group

because many species have both shrubs and trees as growthforms. Vines and liana include

herbaceous vines, woody lianas and all other plants with climbing forms. When a genus has

several different growth forms, we used the major growth form (which has the largest number

of species within the genus) as the genus-level growth form. Range size, maximum/minimum

latitude and maximum/minimum longitude were standardized to [0, 1] in order to calculate

the effect size on discovery probability. Range size was log-transformed before

standardization. Correlations between explanatory variables were checked prior to analysis.

No collinearity was found among explanatory variables (the highest correlation was 0.7).

Cox proportional hazard model

We first modeled the discovery time using survival analysis in which the discovery time was

treated as a random variable. Discovery time is the time it takes to discover a species, i.e.,

how long the species can "survive" not to be discovered. Because only discovered species

could be recorded, there are no censored data in our study. In this case, the empirical survival

curve is just the inverse of the accumulation curve (Steyskal, 1965; Essl *et al.*, 2013). Cox

proportional hazard (Cox PH) model was used to model the instantaneous discovery

probability which describes the probability that a species will be seen in the next step of time

($t+\Delta t$) if it remains unseen up to time t. This probability is expressed as a hazard function, $h(t)$,

given as

$$h(t) = \lim_{\Delta t \to 0} \frac{\Pr[(t \leq T < t + \Delta t) \mid T \geq t]}{\Delta t}.$$

The Cox PH regression model is:

$$h_i(t) = h_0(t) \exp(\beta X),$$

where matrix βX is the linear component of the model (i.e., coefficients and dependent

variables). $h_i(t)$ is the hazard function for species i (i.e., the "risk" to be discovered), and $h_0(t)$

is the baseline hazard function. Therefore, Cox PH model is a type of generalized linear

model. Compared to accelerated failure time model (AFT), Cox model is equivalent to a

semi-parametric model which makes no assumption about the underlying distribution of

survival time (i.e., discovery time), which is more appropriate for our data because discovery

process is highly influenced by historical events (Lu & He, 2017) and the probability

distribution of discovery time is unknown. Because it happened that multiple species were

discovered in the same year, Efron approximation was used to break ties in discovery time

(Hertz-Picciotto & Rockhill, 1997). The proportional hazard assumptions were examined by

plotting the scaled Schoenfeld residuals, denoted as $\beta(t)$, against time. A horizontal trend of

$\beta(t)$ implies that the time-independent coefficient assumption is met. Model selection was

done by choosing the model with the minimum AIC value. Concordance statistic (*C* statistic)

was used to show the discriminative ability of a model. It is equivalent to the area under the

Receiver Operating Characteristic Curve (AUC) in logistic regression, with the value of 0.5

indicating no discrimination power and the value of 1 indicating perfect discrimination. We

also presented the fitted survival curves for different treatments (i.e. inland or coastal

distributed species, and species with different growth forms) using strata models (in strata

model each treatment has a different baseline function $h_0(t)$). Effect sizes are regression

coefficients of the standardized predicting variables. All survival analyses were conducted

using R package 'survival' (Therneau, 2015).

Estimating species richness

Because models that take into account of taxonomic effort (Joppa *et al.*, 2011a,b; Lu & He,

2017) did not converge at the province level, we used a modified logistic discovery model

(Lu & He, 2017) to estimate species richness for different growth forms and for each

province:

$$\Delta S_t = (a + bS_t)(S_{tot} - S_t) + \varepsilon_t,$$

where ΔS_t is the number of species discovered per time interval (5 years in this study), S_{tot}

is the total number of species in a region, S_t is the accumulative number of species discovered

up to time t (=0, 5, 10, 15, 20,… years), *a* and *b* are fitting parameters and ε_t is the error

term. Our goal is to estimate S_{tot}. Note that the logistic discovery model is different from the

logistic regression model. The 'logistic' part of the discovery model derives from the logistic

shape of species discovery curve. The logistic discovery model provides only conservative

estimates in certain cases (Bebber *et al.*, 2007b; Essl *et al.*, 2013; Lu & He, 2017). Inventory

completeness was calculated as the ratio of the number of discovered species to the estimated

total number of species in a province.

Spatial patterns of discovery time and inventory completeness

We further examined the relationship between mean discovery time (i.e., the average number

of years taken to discover the recorded species in an area) of a province and human

population density, total number of species, whether a province is on the coast or inland, and

province area using ordinary linear regression. The spatial autocorrelation of mean discovery

year on the province scale was examined by Moran's *I*. The neighborhood structure of

provincial polygons is defined by contiguity (only polygons with shared borders are counted

as neighbours). We proceeded with ordinary linear regression after no spatial autocorrelation

was detected in the residuals of the model. Beta regression was used to model inventory

completeness with the same set of covariates as modeling discovery time (i.e., population

density, total number of species, whether coastal province or not, and area of each province)

using R package 'betareg' (Cribari-Neto & Zeileis, 2010). Beta regression is preferred

because it only assumes that the response variable ranges from 0 to 1 and is flexible to

accommodate the shape of the distribution (symmetric or skewed). Logistic regression is for

handling binary data, not for proportional data. If logistic regression is used for proportional

data, the base counts that lead to the calculation of proportion must be known. Without that

information, it is inappropriate to use logistic regression. In our data, the base population

which is the true total number of species is not known and hence logistic regression is not

applicable.

RESULTS

Cox proportional hazard model for species discovery probability

The results in Table 1 show that ferns have the lowest discovery probability and trees and

shrubs have the highest discovery probability among all taxonomic groups. For example, in

year 1953 ferns have the highest proportion of undiscovered species and trees and shrubs

have the lowest proportion of undiscovered species (Fig. 1A). Coastal species were

discovered earlier than inland species (Fig. 1B). The discovery probability of a species

increases with its range size, maximum latitude and maximum longitude. The effect sizes of

maximum latitude, minimum latitude, maximum longitude and minimum longitude are larger

than the effect sizes of range size, coastal distribution and growth form, suggesting the

importance of geographic locations to discovery probability (Table 1).

Estimating species richness

The accumulation curve (i.e. the inverse of survival curve) of fern species has the steepest

slope at recent years (Fig. 2A), suggesting that a considerable number of fern species remain

to be discovered in the future. The logistic model estimates that the inventory completeness

of ferns is 0.62. For seed plants, the estimated inventory completeness is 0.73 for herbs, 0.75

for shrubs and trees, and 0.68 for vines/lianas (Table 2). Herbs harbor the largest number of

undiscovered species. The low ferns discovery rate is reflected by the fact that it took on

average the longest time to discovery a fern species, while herbs were earlier to be discovered

($p < 0.05$ for all pairs except between tree/shrub and vine; Fig. 2E).

Spatial patterns of mean discovery year and inventory completeness

The mean discovery year increases from northeast towards southwest (Fig. 3A). Besides

geographic variables (latitudes and coastal distribution), the best model shows that the total

number of species in a region also has a significant effect on mean discovery year (Table 3).

The total number of species is positively correlated with mean discovery year (Fig. 4B).

Mean discovery year decreases with latitude (Fig. 4C) and is earlier in coastal provinces than

in inland provinces (Fig. 4D).

We estimate that in 18 of the 28 provinces, plant species discoveries are more than 90%

complete. Provinces with the largest proportion of undiscovered species are in the southwest

China (75.1% in Guangxi province and 82.7% in Yunnan province; Fig. 3B). Beta regression

shows that inventory completeness is positively correlated with human population density,

and negatively correlated with area and latitude of a province (Table 4).

DISCUSSION

Today's knowledge about biodiversity is the result of arduous quest of generations of

naturalists for botanical and zoological discoveries. Although the nomenclatures of species

are universally binomial, the stories behind their discoveries are not and many of them are as

colorful as the species that were discovered (Kilpatrick, 2014). It is however surprising that

the literature in ecology is largely disconnected from this rich history of species discoveries

(but see Collen *et al.*, 2004; Diniz-Filho *et al.*, 2005; Randhawa *et al.*, 2015). Worse still is

that the data of species discoveries are not much used for answering questions important to

biogeography and biodiversity studies although there have been efforts in using such data for

estimating regional and global species richness (Gaston,1995; Joppa *et al*. 2011a, b; Bebber *et*

al. 2007a, b, 2014; Essle *et al*. 2013; Lu, & He 2017). The rich information provided by

discovery history is especially valuable for filling the knowledge gap in biodiversity research

(Meyer *et al.*, 2015; Meyer, Weigelt, & Kreft, 2016) because it provides guidance about when

and where future discoveries are going to be made and what traits are important to future

discovery (Collen *et al.*, 2004; Diniz-Filho *et al.*, 2005). Therefore, knowledge on species

discovery is of great value for species conversation if we strive to describe all species before

their extinction (Costello *et al.*, 2013; Tedesco *et al.*, 2014; Humphreys *et al*, 2019). In this

study, we compiled data on vascular plant species discovered over 250 years in China for

understanding the geographic variation of discovery rates and for assessing the completeness

of botanical inventory of the country.

Our analyses showed that vascular plant species with a larger range size is more likely to

be discovered in China, consistent with the previous findings that widespread species are

discovered earlier in the history (Bebber *et al.*, 2007a; Essl *et al.*, 2013). Tree and shrub

species of seed plants have the highest discovery probability while fern species have the

lowest discovery probability. Herb and vine/liana species of seed plants all have similar

discovery probabilities. The unique history of botanical discovery in China is revealed by the

fact that species distributed on the coast have a higher discovery probability than inland

species even when geographic information such as the maximum and minimum latitude range

of a species is included in the model (Table 1), and is also shown by the result that

province-level mean discovery time is negatively correlated with coastal distribution (Table

3). This is likely because coastal areas in China were most economically developed and much

more accessible to western naturalists since the Opium War (Bretschneider, 1898; Fan, 2004).

Our results indicate that fern is likely to be the most under-discovered taxon in China

because the species discovery curve for fern shows little sign of level-off (Fig. 2A, Table 2).

Herbs have the largest number of undiscovered species (Table 2) and the second lowest

discovery probability estimated from the Cox proportional hazard model (Table 1). Higher

discovery probability usually leads to higher inventory completeness, which is shown by the

concordance between ranks of discovery probability and ranks of inventory completeness

among taxa in our results (Table 1, Table 2). Although the inventory completeness estimated

from this study varies among the growth forms, the difference in inventory completeness is

relatively small in magnitude (~ 2% between herb and tree/shrub). Herbs have the largest

number of undiscovered species likely because the total number of herb species is larger than

that of any other growth form of seed plants. We suspect that the effect of growth form on

discovery probability and inventory completeness at least partly reflects the difference in the

availability of taxonomic expertise, especially for ferns. The description of fern species

started relatively late in China (~1920s, Chen, 1994) compared to other groups likely because

of the difficulty in distinguishing subtle morphological characters, different species concept

compared to seed plants and the lack of taxonomic expertise at that time. Given that herbs

also contain the largest number of undiscovered species and that most specimens of

undescribed species have already been preserved in herbaria (Bebber *et al.*, 2010), our study

suggests that the limiting factor of discovering new species in China may be the lack of

taxonomic expertise, which resonates with the call to address the challenge of "taxonomic

impediments" (Ebach, Valdecasas, & Wheeler, 2011; Bebber *et al.*, 2014).

The spatial patterns of mean species discovery time and inventory completeness are

largely driven by human population density and species diversity in an area. Although human

history did affect the spatial patterns of species discovery (Diniz-Filho *et al.*, 2005;

Rosenberg *et al.*, 2013; Ballesteros-Mejia *et al.*, 2013), geographic sampling bias does not

change the prioritization of the current conservation efforts because the total number of

species and number of discovered species are highly correlated (Joppa *et al.*, 2011a; Giam *et*

al., 2012; Fig. S3). We expect that new discoveries in the future will be most likely made in

interior southwestern provinces with high species richness such as Xizang, Guangxi and

Yunnan.

It is worth noting that the spatial pattern of species inventory completeness at the

province level is opposite to the spatial pattern at the county level which shows that eastern

part of China has lower inventory completeness (Yang, Ma, & Kreft, 2013, 2014). Yang *et al.*

measured county-level inventory completeness using the slope of sample-based accumulation

curves. Their assessment of inventory completeness might be biased by a sampling strategy

that aims at collecting as many novel species as possible for a given amount of samples

(Chen, 1994). We also argue that the slope of species accumulation curve is not a genuine

measure of inventory completeness. Rather, it measures the variation in species composition

in the samples used to construct the species-accumulation curve. Another possible reason for

this discrepancy is that inventory completeness is scale-dependent. In a hypothetical scenario,

even if the inventory completeness at the county level is on average 90%, the inventory

completeness at the province level could be lower than 90% if most of the recorded species at

the county level are common species. While Yang *et al.* (2014) argued that more efforts

should be devoted to increasing botanical collections in eastern densely populated areas

where county-level inventories are less complete, our study here does not support this

advocacy. Instead, we suggest that future botanical collection efforts should be more allotted

to the provinces of southwest China where there is high species diversity and the botanical

inventory is least complete. Recent findings support our conlusion by showing that the

majority of newly discovered species in China after the completion of *Flora of China* in 2013

came from the global biodiversity hotspots of Indo-Burma and Mountains of Southwest

China (Cai *et al.*, 2019), and southwest China (Yunnan, Guangxi, Sichuan, Xizang) and

Taiwan are places where most new species were discovered during 2000 - 2019 (Du *et al.*,

2020).

There are two major limitations in this study. The first one is that species discovery year

is not the time when the species was first collected in the field but is the time when the

species was first described in a scientific publication. This could lead to inflating discovery

probability for widespread species if their type specimens were collected outside of China. To

address this problem, we reran our Cox proportional hazard model using species only

endemic to China (12917 species). The results show that the key drivers of discovery

probability are mostly consistent but with some noticeable differences (Table S1): the effect

size of coastal distribution became negative (Table S1) because the majority of these endemic

species were discovered after 1900 (Fig. S1A), which removed the effect of early discoveries

made by western naturalists. For the province-level analysis, excluding non-endemic species

in China does not significantly change the spatial pattern of mean discovery year (Fig. S2A).

Human population density became negatively correlated with the mean discovery year of

endemic species; area became positively correlated with mean discovery year; the effect of

coastal distribution and number of species are no longer significant (Table S2). For

province-level inventory completeness, the effects of predicting variables are mostly

consistent with the model that includes all species except that coastial distribution and

longitude also became significant variables (Table S3).

The second limitation is that our data do not distinguish the discovery of a new species

from a species renamed from a known synonym. As data including a full list of all synonyms

at each time step are not available, we are not able to model the transition rate from

synonyms to valid names (Alroy, 2002).

In summary, our study shows that most underdiscovered vascular plant species in China

are ferns and herbs of seed plants, which are mostly narrowly distributed endemic species in

the southwest biodiversity hotspots of China. Given the "taxonomic impediments" we are

facing, more resources should be channeled to the training and recruitments of taxonomic

expertise in these two particular groups. There is an urgency of cataloging undiscovered

species in southwest mountainous areas for future conservation designs and botanical study.

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478   Table 1. Cox proportional hazard model with all variables (growth form as categorical data

479   and tree/shrub used as the baseline variable).

	Effect size	Standard error	*P*-value
Growth form.fern	-0.82	0.03	<0.001
Growth form.herb	-0.13	0.01	<0.001
Growth form.vine.liana	-0.13	0.03	<0.001
Range size	0.51	0.06	<0.001
Coast	0.13	0.02	<0.001
Maximum longitude	1.43	0.05	<0.001
Minimum longitude	-1.05	0.05	<0.001
Maximum latitude	0.83	0.07	<0.001
Minimum latitude	-0.77	0.04	<0.001
N = 30944			
Concordance = 0.653 (se = 0.002)			

481   Table 2. Estimated species richness for different growth forms.

	Number of discovered species	Estimated total number of species	Lower 95% bound	Upper 95% bound	Completeness (lower 95% -)
Fern	1921	2712	749	4676	0.622 (0.41-)
Vine/Liana	1202	1710	856	2563	0.685 (0.47-)
Herb	18370	23867	16982	30752	0.737 (0.59-)
Tree/Shrub	9451	12998	8855	15741	0.754 (0.60-)

483 Table 3. Linear regression of mean discovery year.

	Coefficient	Standard Error	*P*-value
Intercept	1941	1.5	<0.001
Number of species	0.0029	0.0008	<0.01
Coast	-13.8	4.49	<0.01
Mean latitude	-2.03	0.36	<0.001

Multiple R^2 = 0.82, Adjusted R^2 = 0.80

486   Table 4. Beta regression of province level inventory completeness.

	Coefficient	Standard Error	*P*-value
Intercept	-0.39	0.23	<0.05
Human population density	8.70e-04	3.17e-04	<0.001
Area	-5.23e-07	1.33e-07	<0.001
Latitude	0.07	6.8e-03	<0.001
Pseudo $R^2 = 0.85$			

Figure Legends

Figure 1. Fitted survival curves of the Cox proportional hazard models (A) stratified on variable 'growth form' and (B) stratified on variable 'coast'. Dashed lines show 95% confidence intervals.

Figure 2. (A) – (D) Species accumulation curves for four different taxonomic groups (based on 5 years-interval data). (E) Boxplot for discovery time of the four growth forms. "a" labels the groups with no significant difference in the TurkeyHSD test.

Figure 3. (A) Distribution of mean province-level species discovery year in China. (B) Distribution of species inventory completeness. Mollweide projection is used for the distributions.

Figure 4. (A) – (C) Mean species discovery year against human population density, number of discovered species, and latitude. (D) Boxplot of mean discovery year for inland and coastal provinces. (E) – (G) Species inventory completeness against human population density, number of discovered species, and latitude. (H) Boxplot of inventory completeness for inland and coastal provinces. Solid curves show the fitted smooth spline curves.

507   Figure 1

509　　Figure 2

511   Figure 3

514   Figure 4

Appendices: **The patterns of vascular plant discoveries in China**
Muyang Lu, Lianming Gao, Hongtao Li, Fangliang He

Table S1. Cox proportional hazard model with all variables for endemic species in China (growth forms were coded as a categorical variable in which tree/shrub was used as the baseline variable).

	Effect size	Standard error	P-value
Growth form.fern	-0.83	0.04	<0.001
Growth form.herb	-0.082	0.02	<0.001
Growth form.vine	-0.23	0.10	<0.001
Range size	1.21	0.03	<0.001
Coast	-0.12	0.03	<0.001
Maximum longitude	1.30	0.10	<0.001
Minimum longitude	0.09	0.09	0.32
Maximum latitude	0.30	0.11	<0.01
Minimum latitude	-0.70	0.08	<0.001

N = 12917

Concordance = 0634 (se = 0.003)

Table S2. Linear regression of mean discovery time for endemic species.

	Coefficient	Standard Error	P-value
Intercept	1964	6.8	<0.001
Human population density	-0.013	0.0071	0.074
Area	2.30e-05	3.91e-06	<0.001
Mean latitude	-1.46	0.18	<0.001
Multiple R^2: 0.80, Adjusted R^2: 0.77			

Table S3. Beta regression of province-level inventory completeness for endemic species.

	Coefficient	Standard Error	P-value
Intercept	-1.49	5.18	<0.01
Human population density	2.06e-03	5.53e-04	<0.001
Coast	-0.49	0.17	<0.01
Area	-8.58e-07	3.16e-07	<0.01
Mean longitude	-0.13	0.047	<0.01
Mean latitude	0.68	0.14	<0.001
Longitude:latitude	-5.45e-03	1.23e-03	<0.001
Pseudo R^2: 0.76			

Figure S1. Histograms of discovery date (A) and the number of distributed provinces of endemic species to China (B).

Figure S2. (A) Mean discovery time for province-level endemic species in China. (B) Species inventory completeness for province-level endemic species. Mollweide projection is used.

Figure S3. (A) Estimated total number of species against the number of discovered species at the province-level. (B) Estimated inventory completeness against the number of discovered species at the province-level.