



Contrasting Floristic Diversity of the Hengduan Mountains, the Himalayas and the Qinghai-Tibet Plateau *Sensu Stricto* in China

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The Qinghai-Tibet Plateau *sensu lato* (QTP *s.l.*) harbors an exceptionally high biodiversity, especially at its southeastern margin: this area encompasses the Hengduan Mountains and the eastern Himalayas, which have been listed as biodiversity hotspots. To the contrary, the plateau interior (namely the Qinghai-Tibet Plateau *sensu stricto*, QTP *s.s.*) is relatively species-poor because of its particularly harsh climate. With contrasting geological histories and environmental conditions of the Hengduan Mountains, the Himalayas, and the QTP *s.s.*, it would be expected that floristic compositions and diversity patterns of these three regions would differ between each other. To compare the floristic diversity of these three regions, we assembled data on seed plant's distribution in the three regions based on county-level mapping from published monographs and online databases, and we then analyzed their floristic features and species diversity patterns (horizontal and elevational). We found that the Hengduan Mountains hosted the most seed plant species (8,439), as expected. The highest percentage of shrub (22.88%) and tree species (9.80%) were in the Himalayas, whereas herbaceous species (81.50%) were relatively more prominent in the QTP *s.s.*. The Hengduan Mountains also had the most species-rich genera (10) with more than 50% of their total species diversity in China. Also, temperate genera dominated across these three regions, with a highest percentage (77.61%) within the QTP *s.s.*. Across the QTP *s.l.*, species diversity gradually decreased from the southeastern part to the northwest, and most of seed plants were distributed in the southern and eastern margin of the Hengduan Mountains and East Himalayas. Along elevational gradients, species richness all demonstrated a hump-shape curve, but the most species-rich elevation zone differed for each type of life-form across the three regions. Our study sets a base for exploring the origin and evolution of mountain taxa, as well as provides a snapshot of the current plant distribution, which will certainly be modified by climate change.

Keywords: biodiversity, flora, Hengduan Mountains, Himalayas, mountain plants, Qinghai-Tibet Plateau

INTRODUCTION

Mountainous regions harbor about one-quarter of terrestrial species, and inevitably, some of these areas have been listed as hotspots of biodiversity, including for example the two most diverse mountainous regions on Earth: the Andes and the Hengduan Mountains (Marchese, 2015; Hoorn et al., 2018). The establishment of high biodiversity levels in these remarkable mountain systems is likely to result from the combination of global and local abiotic conditions (e.g., geological and climatic processes) with biotic, often taxon-specific features (e.g., key innovations) (Lagomarsino et al., 2016). As stated in the “mountain geobiodiversity hypothesis” (Mosbrugger et al., 2018), the development of a full altitudinal zonation (i.e., thermal belts) and high ruggedness of the terrain is expected to foster the immigration of foreign pre-adapted lineages (biological interchange), whereas at the same time lowering the risk of climate-driven extinction by providing numerous alternative habitat within a short distance. Yet, high speciation rate would be achieved only during cyclical modifications of distribution ranges (i.e., species-pump effect). Indeed, radiations, which are major contributors to mountain biodiversity, appear to be concentrated in the last few million years (thus overlapping with climate oscillations), whereas mountain building started much earlier (Hughes and Atchison, 2015; Muellner-Riehl et al., 2019). Obviously, mountain building and past climate changes are crucial triggers of complex evolutionary processes accounting for species diversity in mountains (Antonelli et al., 2018; Rahbek et al., 2019). Thus, understanding the profile of species diversity in such a region could provide general and valuable insights into the recent evolution of local taxa, as well as help prioritizing conservation strategies of regional biota.

As the largest and highest plateau region on Earth, the Qinghai-Tibet Plateau *sensu lato* (QTP *s.l.*) harbors a great number of mountain plant species (Zhang et al., 2016). The QTP *s.l.* consists of three major regions: the Hengduan Mountains to the East, the Himalayas in the South and the Qinghai-Tibet Plateau *sensu stricto* (QTP *s.s.*). Among these, the Hengduan Mountains and the eastern Himalayas are two of the most important biodiversity hotspots in the world (Marchese, 2015). Over long period of time (from late Miocene onwards), the Hengduan Mountains have experienced a complex and rapid uplift that produced extreme ruggedness of the terrain as well as a remarkable environmental heterogeneity (Wang et al., 2012). In contrast, the QTP *s.s.* is most likely to be more ancient, with an uplift starting at least around the India-Asia collision [*c.* 55 Ma, (Favre et al., 2015)], with subsequent outward extensions (Mulch and Chamberlain, 2006). The QTP *s.s.* is particularly vast and high (*c.* 4,500 m in average) but displays only a low ruggedness. The Himalayas experienced a rather rapid orogeny starting during the Miocene (Deng and Ding, 2015; Ding et al., 2017), with the Qomolangma (Mt. Everest) attaining *c.* 5,000 m already during the early Miocene (Gébelin et al., 2013). The uplift of high mountain ranges resulted in modifications of the distribution of precipitations, with a probable intensifying

effect on the East Asian monsoon and the Indian monsoon, progressively generating modern climatic patterns in Asia (An et al., 2001). Currently, there is a strong climatic gradient from warm and humid conditions (in the southeast, in the Hengduan Mountains and East Himalayas) to cold and dry climatic regimes behind the rain shadow (in the center and northwest, in the QTP *s.s.*). The three regions differing from each other in their uplift history and topography, as well as in their climatic regime, distinctive evolutionary trajectories of species would be expected. For example, it is known that the Hengduan Mountains hosted higher *in situ* diversification rates than the other two regions over the past 10 Ma, during which time the overall dispersal rate from the Hengduan Mountains into the Himalayas and QTP *s.s.* increased gradually (Xing and Ree, 2017). A number of radiations of mountain genera occurred simultaneously with the uplift of the Hengduan Mountains and climate oscillations, including for example *Gentiana* (Favre et al., 2016) and *Saxifraga* (Ebersbach et al., 2017). Serving as a source area for multiple genera that colonized other areas of the world (e.g., Matuszak et al., 2016), and characterized by many radiations, the Hengduan Mountains are often depicted as a cradle of evolution for many temperate taxa (Chen et al., 2018; Lu et al., 2018).

Probably because of its roughly north-south orientation and rugged nature, the Hengduan Mountains appears to have experienced lower extinction rates during climatic oscillations. Hence, this region is usually viewed as important refugium preserving high levels of species and genetic diversity during glacials (Qiu et al., 2011; Yu et al., 2019b). In comparison, the east-west orientation of the Himalayas and the low ruggedness of the QTP *s.s.* rendered their fauna and flora more susceptible to be extinction during climate oscillations, leading to vacant niches and recolonization from the Hengduan Mountains during interglacials (e.g., Cun and Wang, 2010; Liu et al., 2013; Xing and Ree, 2017). However, there were several microrefugia in the last two regions that have probably preserved number of plant taxa (Liu et al., 2012). On the other hand, under current climate projections, plant species within the Hengduan Mountains and Himalayas are predicted to expand their range sizes as they shift upslope or westward into the QTP *s.s.*, and species richness in the Hengduan Mountains and QTP *s.s.* is expected to undergo great increases with expected climate warming [(Liang et al., 2018; He et al., 2019a,b)]. However, species richness in the Himalayas would remain almost stable (Liang et al., 2018).

Evolutionary studies targeting this region usually lump the entire area as one (the QTP *s.l.*). However, due to their distinct uplift histories and environmental conditions (i.e., climate and topography), the Hengduan Mountains, the Himalayas and the QTP *s.s.*, are likely to differ in terms of evolutionary processes as well as overall response of their vegetation to climate change. Hence, there could be profound differences in floristic diversity amongst these three regions. In this study, we aim to investigate the features and distributions of plant diversity in these regions, and among other aspects distinguishing between life-forms (i.e., herb, shrub, and tree).

METHODS

Study Area

The QTP *s.l.* in China is the largest ($c. 2.5 \times 10^6 \text{ km}^2$) and highest plateau (average elevation $> c. 4,000 \text{ m}$) in the world (Zhang et al., 2002; **Figure 1A**). The relative abundance of its main biomes (forest, alpine meadow, alpine steppe, and alpine desert) varies along a southeast to the northwest gradient following climatic characteristics. The area can be divided into three regions (**Figure 1B**). First, the Hengduan Mountains (total area of $c. 3.9 \times 10^5 \text{ km}^2$, average elevation $> c. 4,000 \text{ m}$), are located at the southeastern edge of the plateau. This region is topographically complex with numerous north-south oriented mountain ranges and deep valleys. In terms of climate, this region is influenced heavily by both the East Asian monsoon and Indian monsoon (**Figure 1C**), and thus characterized by higher annual mean temperature and precipitation (**Figure 1D**). Then, the Himalayas (confined to the territory of China in this study) at the southern edge of the plateau, are a very long and narrow mountain range (east-west orientation) with a vast elevational gradient (**Figure 1E**), spanning over 20 degrees of longitude ($c. 79\text{--}99^\circ\text{E}$). It covers about $3.1 \times 10^5 \text{ km}^2$ with average elevation above 4,200 m. Because of the rain shadow effect, there is a contrasting climatic pattern in the Himalayas between their southern (warm and wet) and northern slopes (cold and dry). The third and largest area ($c. 1.8 \times 10^6 \text{ km}^2$) is the QTP *s.s.* itself, with a higher mean elevation ($> c. 4,500 \text{ m}$) and particularly harsh cold and dry climate in its northwestern part, whereas its eastern (e.g., the Qilian Mountains) and southeastern margins have relatively warm and wet climate influenced by the East Asian monsoon.

Collection of Species List and Distribution Data

First, we established the list of seed plant species (including intraspecific taxa) occurring in the QTP *s.l.* in China, and recorded their spatial distributions at the county level within each of the three regions (the Hengduan Mountains, the Himalayas and the QTP *s.s.*) investigated. Because species information within the southern parts of the Himalayas (e.g., Nepal, Bhutan, Sikkim, Kashmir) were difficult to obtain, we only considered the species distribution within the northern Himalayas (Chinese territory). We gathered information from the following monographs: *The Vascular Plants and Their Ecogeographical Distribution of the Qinghai-Tibetan Plateau* (Wu, 2008), *Flora Reipublicae Popularis Sinicae* (Editorial Committee of Flora Reipublicae Popularis Sinicae, 1959–2004), *Flora of China* (Wu et al., 1994–2013), floras of Tibet, Qinghai, Sichuan, Yunnan, Gansu and Xinjiang, and the *Vascular Plants of the Hengduan Mountains* (The Comprehensive Scientific Expedition to the Qinghai Xizang Plateau Chinese Academy of Sciences, 1993–1994), as well as field vegetation surveys in the Qiangtang Plateau. For species with only approximate distribution data, we further referred to several online databases, such as the Chinese Virtual Herbarium (<https://www.cvh.ac.cn/>) and the Global Biodiversity Information Facility (<https://www.gbif.org/>). For nomenclatural consistency across these sources, the preliminary

species list was based on the *Flora of China*, and unresolved or synonymous species names were excluded. The following information was recorded for each species: (i) life-form (tree, shrub, or herb), (ii) elevational range and (iii) whether it is endemic (only found in the QTP *s.l.*) or (iv) alpine (distributed above the tree line). We followed the APG IV classification system (The Angiosperm Phylogeny Group et al., 2016).

Basic Statistics and Floristic Composition Analysis

Basic analyses within the Hengduan Mountains, the Himalayas, the QTP *s.s.* and the QTP *s.l.* in China were conducted as follows: (1) we calculated the absolute and relative number of each life-form within each individual region; (2) we calculated similarity coefficients of all seed plant species and endemic species between the Hengduan Mountains, the Himalayas and the QTP *s.s.* at the family, genus and species levels; (3) we established a list of top 20 families/genera for each region and calculated their percentage of species in a family/genus in China; (4) Based on the areal types of the families and genera of seed plants proposed by Wu et al. (2006), we calculated the geographical floristic composition of the families and genera of seed plants within these regions.

Mapping Species Diversity Patterns

We counted the number of each life-form taxa in each county of the QTP *s.l.* in China and built their species richness patterns at the county level using the ArcGIS 10.1 (Environmental Systems Research Institute, Inc., Redlands, CA). We built the patterns of ratio of endemic and alpine species in a county across the QTP *s.l.* We also built the patterns of weighted endemism (WE; Linder, 2001) at the species and genus levels. Then, we built the elevational diversity patterns of each life-form taxa within the Hengduan Mountains, the Himalayas and the QTP *s.s.*, and the frequency line of species richness along the elevation was drawn using R 3.6.2 (R Core Team, 2016).

RESULTS

Floras of the Hengduan Mountains, the Himalayas, and the QTP *s.s.*

Among these three mountain regions, the Hengduan Mountains harbored the most seed plant species (8,439), belonging to 184 families and 1,273 genera (see **Appendix 1**), of which 5,817 (68.93%) species were herbaceous, 1,896 (22.47%) were shrubs, and 726 (8.60%) were trees (**Table 1**). In contrast, the Himalayas and the QTP *s.s.* had 5,468 (177 families and 1,146 genera) species and 3,908 (122 families and 746 genera) species (see **Appendixes 2, 3**), respectively. Within the Himalayas, the majority of species were herbaceous (3,681 species; 67.32%), also including the highest percentage of shrub (1,251, 22.88%) and tree species (536, 9.80%) across the three regions. The QTP *s.s.* had the largest proportion of herbaceous species (3,185; 81.50%), with additional 556 (14.23%) shrubs and 167 (4.27%) tree species. In total, the QTP *s.l.* recorded 12,091 seed plant species (200 families and 1,574 genera) with 8,513 (70.41%) herbs, 2,558 (21.16%) shrubs and 1,020 (8.43%) trees, of which 5,457 (45.13%) species were endemic, as well as 3,320 (27.46%) alpine species.

TABLE 1 | The basic floristic diversity of the Hengduan Mountains, the Himalayas, the Qinghai-Tibet Plateau s.s., and the Qinghai-Tibet Plateau s.l.

Region	Family	Genus	Tree	Shrub	Herb	Total
Hengduan	184	1273	726 (8.60%)	1896 (22.47%)	5817 (68.93%)	8439
Himalayas	177	1146	536 (9.80%)	1251 (22.88%)	3681 (67.32%)	5468
QTP s.s.	122	746	167 (4.27%)	556 (14.23%)	3185 (81.50%)	3908
QTP s.l.	200	1574	1020 (8.43%)	2558 (21.16%)	8513 (70.41%)	12091

The number in the bracket represents the proportion of life-form group of seed plants in each region. The dark number represents the highest proportion of life-form taxa among these four regions.

TABLE 2 | Similarity coefficients of floristic composition between the Hengduan Mountains, the Himalayas, and the Qinghai-Tibet Plateau s.s. at the family, genus and species levels.

(A)	Hengduan	Himalayas	QTP s.s.
Similarity coefficients at family-level			
Hengduan	1		
Himalayas	166/ 0.4548	1	
QTP s.s.	119/ 0.3851	120/ 0.3727	1
Similarity coefficients at genus-level			
Hengduan	1		
Himalayas	943/ 0.3889	1	
QTP s.s.	596/ 0.2943	572/ 0.3023	1
Similarity coefficients at species-level			
Hengduan	1		
Himalayas	3196/ 0.2984	1	
QTP s.s.	2059/ 0.2001	1611/ 0.2075	1
(B)	Hengduan	Himalayas	QTP s.s.
Similarity coefficients at family-level			
Hengduan	1		
Himalayas	94/ 0.4332	1	
QTP s.s.	62/ 0.3483	61/ 0.3609	1
Similarity coefficients at genus-level			
Hengduan	1		
Himalayas	364/ 0.3647	1	
QTP s.s.	221/ 0.2650	209/ 0.2935	1
Similarity coefficients at species-level			
Hengduan	1		
Himalayas	1059/ 0.2121	1	
QTP s.s.	693/ 0.1484	472/ 0.1651	1

A, for all seed plant species. **B**, for endemic plant species. The number in front of slashes represents the shared number of families, genera or species between two regions.

Besides, the similarity coefficients of all species and endemic species composition at the family level between each region were higher than that of the genus and species levels. Among these, the coefficients between the Hengduan Mountains and Himalayas at the three levels were the highest (**Tables 2A,B**).

Seventeen of the top 20 big families were common across all regions, but the number of species in these families varied in each region (**Table 3**). More than half of species diversity was located in the Hengduan Mountains for three families (Rosaceae, Orobanchaceae, and Gentianaceae). At the genus level, about 12

of the 20 largest genera were common throughout but the most represented genera differed between regions (**Table 4**).

The floras of the Hengduan Mountains and the Himalayas included relatively more typically tropical families than temperate ones, which was not the case in the QTP s.s. (**Table 5**). However, at the genus level, the percentage of typically temperate genera was higher than tropical ones in all three regions (**Table 6**). Large proportion of these temperate genera showed affinities with the flora of north temperate and East Asia.

Diversity Patterns

Across the QTP s.l., species diversity (also within each life-form taxa) gradually decreased from the southeastern part to the northwest (**Figures 2A–D**). Most parts of the Hengduan Mountains hosted a high percentage of endemics (**Figure 2E**), but the southern and eastern parts of the range were particularly species-rich. The most diverse part of the Himalayas was located in its eastern quarter. However, relatively high diversity was also found in the central Himalayas. Those mentioned areas also have a high level of weighted endemism (**Figures 2G,H**). In comparison, species diversity in the QTP s.s. was relatively low, except in the Qilian Mountains, the Anyemaqen Mountains and the Three-River Headwaters Region (East). The local flora is characterized by a high proportion of alpine plants (**Figure 2F**).

Along altitudinal gradients, species richness was always characterized by a hump-shape curve for different life-form types, but most species were located within distinct elevational zones within the three regions (**Figures 3A–D**). For example, most herbaceous species were located between 2,500 and 3,500 m (Hengduan Mountains), 2,700 and 3,700 m (Himalayas), and 3,100 and 4,100 m (QTP s.s.), respectively. In contrast, shrubs were mainly distributed between 2,100 and 3,100 m in the Hengduan Mountains, 2,200 and 3,200 m in the Himalayas, and 2,600 and 3,600 m in the QTP s.s.. Most tree species were distributed between 2,000 and 3,000 m across these three regions.

DISCUSSION

Although the Hengduan Mountains, the Himalayas and the QTP s.s. are adjacent to each other, their respective flora present distinct features and diversity patterns. The Hengduan Mountains host the most species and higher percentage of endemics, as expected, whereas the Himalayas and the QTP s.s. have relatively low species diversity. Across the QTP s.l., species diversity gradually decreased from the southeastern to

TABLE 3 | Top 20 big families for the Hengduan Mountains, the Himalayas, the Qinghai-Tibet Plateau s.s., and the Qinghai-Tibet Plateau s.l. and their percentage of species in their respective family in China (PF).

Hengduan		Himalayas		QTP s.s.		QTP s.l.	
Family	No./PF	Family	No./PF	Family	No./PF	Family	No./PF
Asteraceae	665/28.47	Asteraceae	459/19.65	Asteraceae	510/21.83	Asteraceae	1101/47.13
Rosaceae	498/ 52.42	Ericaceae	346/41.89	Poaceae	404/22.51	Poaceae	740/41.23
Poaceae	489/27.24	Poaceae	295/16.43	Fabaceae	257/15.36	Ranunculaceae	633/ 68.73
Ranunculaceae	434/47.12	Rosaceae	288/30.32	Ranunculaceae	200/21.72	Rosaceae	618/ 65.05
Ericaceae	378/45.76	Orchidaceae	257/18.52	Rosaceae	199/20.95	Fabaceae	557/33.29
Fabaceae	371/22.18	Ranunculaceae	227/24.65	Brassicaceae	127/30.83	Ericaceae	539/ 65.25
Lamiaceae	324/40.15	Fabaceae	212/12.67	Gentianaceae	127/30.31	Orchidaceae	429/30.91
Orchidaceae	323/23.27	Primulaceae	176/34.04	Caryophyllaceae	126/32.31	Lamiaceae	415/ 51.43
Orobanchaceae	311/ 72.49	Lamiaceae	151/18.71	Cyperaceae	112/12.95	Orobanchaceae	384/ 89.51
Apiaceae	293/47.72	Gentianaceae	146/34.84	Lamiaceae	110/13.63	Primulaceae	358/ 69.25
Gentianaceae	216/ 51.55	Apiaceae	138/22.48	Salicaceae	107/30.84	Apiaceae	347/ 56.51
Primulaceae	209/40.43	Caryophyllaceae	137/35.13	Orobanchaceae	99/23.08	Gentianaceae	331/ 78.80
Cyperaceae	205/23.70	Saxifragaceae	124/22.75	Apiaceae	91/14.82	Caryophyllaceae	279/ 71.54
Saxifragaceae	175/32.11	Cyperaceae	116/13.41	Boraginaceae	86/29.25	Cyperaceae	276/31.91
Caryophyllaceae	166/42.56	Orobanchaceae	108/25.17	Saxifragaceae	80/14.68	Saxifragaceae	251/46.06
Salicaceae	151/43.52	Papaveraceae	99/22.35	Papaveraceae	76/17.16	Papaveraceae	232/ 52.37
Papaveraceae	150/33.86	Urticaceae	95/27.86	Primulaceae	74/14.31	Salicaceae	228/ 65.71
Brassicaceae	119/28.88	Salicaceae	93/26.80	Crassulaceae	73/31.33	Brassicaceae	204/49.51
Crassulaceae	114/48.93	Brassicaceae	87/21.12	Polygonaceae	63/26.47	Boraginaceae	171/ 58.16
Liliaceae	114/15.70	Rubiaceae	82/11.70	Orchidaceae	61/4.39	Rubiaceae	159/22.68
Others	2734	Others	1932	Others	926	Others	3839

The common families among these regions are in bold. More than half of the percentage of species in a family was highlighted.

the northwest part, and most of seed plants were located in the southern and eastern margin of the Hengduan Mountains and East Himalayas. Along elevational gradients, species richness all demonstrated a hump-shape curve, but the maximum zones for each life-form differed across the three regions. These are probably due to local climatic and topographic complexity, as well as geological and climatic histories. Below we discussed these floristic findings more in detail.

The Flora of the Hengduan Mountains: As a Cradle of Evolution

Undoubtedly, the Hengduan Mountains host the most plant species in the region (see **Table 1**). This could partly be explained by current climatic and topographic conditions: the Hengduan Mountains are under a relatively generous climate, while their rugged terrain allows for a high diversity of habitats (Yu et al., 2019a). Such favorable conditions are currently also found in other species-rich mountainous areas, such as the Andes (Luebert and Weigend, 2014) and the Tropical Mountains of African (Marshall et al., 2016). However, as shown in Muellner-Riehl et al. (2019), historical factors have played an important role for species richness there. For example, extinction risks during climate oscillations were probably buffered by vast altitudinal gradients, allowing for a substantial overlap of altitudinal distribution of temperatures between glacials and interglacials. At the same time, climate oscillations are likely

to have generated a species-pump effect (Mosbrugger et al., 2018), fostering a number of radiations. The region is thus often depicted as a cradle of evolution (Lu et al., 2018), at least since the frequency of radiations has increased with the uplift of the Hengduan Mountains [c. 7–8 Ma, (Xing and Ree, 2017; Muellner-Riehl et al., 2019)]. These radiations predominantly occurred in herbaceous plant genera (see **Table 4**), such as *Pedicularis* (Yu et al., 2015), *Gentiana* (Favre et al., 2016), and *Saxifraga* (Ebersbach et al., 2017), and occasionally in shrub taxa such as *Rhododendron* (Yan et al., 2015). This is reflected in our results, with herbaceous plants accounting for a larger proportion of the flora of the Hengduan Mountains (see **Table 1**) than would be expected under these latitudes. Besides, amounts of genera in the Hengduan Mountains have experienced *in-situ* diversification, and most of them could colonize the East Asia and north temperate (e.g., Favre et al., 2016; Matuszak et al., 2016), generating large proportion of the East Asia and north temperate elements in this mountain region (see **Table 6**).

The Flora of the Himalayas: Relatively Less Radiations but More Immigration

Although the number of species occurring in the Himalayas is only 65% of that in the Hengduan Mountains, the two regions present a high floristic similarity (see **Table 2**). This may indicate that relatively less species resulted from *in-situ*

TABLE 4 | Top 20 big genera for the Hengduan Mountains, the Himalayas, the Qinghai-Tibet Plateau s.s., and the Qinghai-Tibet Plateau s.l. and their percentage of species in their respective genus in China (PG).

Hengduan		Himalayas		QTP s.s.		QTP s.l.	
Genus	No./PG	Genus	No./PG	Genus	No./PG	Genus	No./PG
<i>Pedicularis</i>	287/ 81.53	<i>Rhododendron</i>	231/40.46	<i>Saussurea</i>	94/32.53	<i>Rhododendron</i>	385/ 67.43
<i>Rhododendron</i>	280/49.04	<i>Primula</i>	124/41.33	<i>Salix</i>	91/33.09	<i>Pedicularis</i>	348/ 98.86
<i>Saxifraga</i>	145/ 67.13	<i>Saxifraga</i>	108/ 50.00	<i>Pedicularis</i>	83/23.58	<i>Primula</i>	229/ 76.33
<i>Gentiana</i>	137/ 55.24	<i>Pedicularis</i>	97/27.52	<i>Astragalus</i>	82/20.45	<i>Saxifraga</i>	213/ 98.61
<i>Aconitum</i>	133/ 63.03	<i>Gentiana</i>	83/33.47	<i>Artemisia</i>	76/40.86	<i>Gentiana</i>	199/ 80.24
<i>Primula</i>	124/41.33	<i>Salix</i>	80/29.09	<i>Gentiana</i>	75/30.24	<i>Salix</i>	184/ 66.91
<i>Salix</i>	121/44.00	<i>Corydalis</i>	69/19.33	<i>Saxifraga</i>	71/32.87	<i>Aconitum</i>	175/ 82.94
<i>Carex</i>	114/21.63	<i>Saussurea</i>	67/23.18	<i>Carex</i>	65/12.33	<i>Saussurea</i>	174/ 60.21
<i>Corydalis</i>	112/31.37	<i>Carex</i>	59/13.09	<i>Corydalis</i>	54/15.13	<i>Corydalis</i>	173/48.46
<i>Saussurea</i>	104/35.99	<i>Potentilla</i>	58/ 67.44	<i>Oxytropis</i>	52/39.10	<i>Carex</i>	157/29.79
<i>Delphinium</i>	99/ 57.23	<i>Astragalus</i>	55/13.72	<i>Delphinium</i>	49/28.32	<i>Delphinium</i>	153/ 88.44
<i>Rubus</i>	89/42.79	<i>Polygonum</i>	55/48.67	<i>Potentilla</i>	49/ 56.98	<i>Astragalus</i>	145/36.16
<i>Astragalus</i>	81/20.20	<i>Aconitum</i>	51/24.17	<i>Poa</i>	47/ 58.02	<i>Berberis</i>	124/ 57.67
<i>Berberis</i>	76/35.35	<i>Berberis</i>	51/23.73	<i>Rhododendron</i>	45/7.88	<i>Artemisia</i>	110/ 59.14
<i>Sedum</i>	75/ 61.98	<i>Artemisia</i>	47/25.27	<i>Primula</i>	43/14.33	<i>Rubus</i>	101/48.56
<i>Arenaria</i>	65/ 63.73	<i>Aster</i>	43/34.96	<i>Rhodiola</i>	38/ 69.10	<i>Arenaria</i>	96/ 97.12
<i>Ligularia</i>	65/ 52.85	<i>Rubus</i>	40/19.23	<i>Roegneria</i>	38/ 54.29	<i>Potentilla</i>	84/ 97.67
<i>Poa</i>	64/ 79.01	<i>Agapetes</i>	38/ 71.70	<i>Ranunculus</i>	37/29.60	<i>Sedum</i>	83/ 68.60
<i>Potentilla</i>	61/ 70.93	<i>Arenaria</i>	38/37.25	<i>Arenaria</i>	36/35.29	<i>Aster</i>	82/ 66.67
<i>Aster</i>	60/48.78	<i>Impatiens</i>	37/16.30	<i>Allium</i>	35/25.36	<i>Impatiens</i>	80/35.24
Others	6147	Others	4038	Others	2748	Others	8786

The common genera among these regions are in bold. More than half of the percentage of species in a genus was highlighted.

radiations, and proportionally more from immigration in the Himalayas. In fact, Xing and Ree (2017) have found that biological interchange between the Hengduan Mountains and the Himalayas has increased in the last few million years (simultaneously with climate oscillations), and in an asymmetrical manner: immigration from the Hengduan Mountains to the Himalayas is more frequent than the other way around. This may be explained by the East-West orientation of the Himalayas and the presence of a sharp rain shadow, both of which might have rendered species of this region more prone to extinction during climate oscillations than in the Hengduan Mountains. Vacant niches were then filled from the neighboring species stock surviving in the Hengduan Mountains (Ren et al., 2015; Yan et al., 2015; Ebersbach et al., 2017). However, because species in the southern Himalayas (more favorable habitats and plenty altitudinal gradients) were not considered in this study, it should be aware of that the flora of the Himalayas is likely to be underestimated, and there could be several micro-refugia preserving amounts of species in the southern Himalayas. Also, most parts of South Himalayas belong to subtropical and tropical regions, an increasing number of tropical elements could be found there. Thus, exploring the flora of southern Himalayas (beyond the Chinese territory) would provide an important supplement to understand the floristic diversity of the QTP s.l..

The Flora of the QTP s.s.: Fewer but More Specialized Species

Although the QTP s.s., occupies 72% of the QTP s.l., its species diversity is less than half that of the Hengduan Mountains (c. 46%). Most of these species are herbaceous (81.50%, see Table 1). This clearly is the result of an extremely cold and dry climate, preventing most tree and shrub species to grow. Such harsh and alpine habitats resulted in large proportion of temperate elements of genus (see Table 6), these dominant temperate elements were also found in the alpine subnival belt of the Hengduan Mountains and Shangri-La in northwestern Yunnan (Xu et al., 2014; Zhu, 2015). Little evidence exists for its biological interchange with the Himalayas and the Hengduan Mountains, and the species similarity between the QTP s.s. and the Hengduan Mountains, the Himalayas are only c. 20% (see Table 2). This might be further explained by the absence of some thermal belts of lower elevation, and the low ruggedness of the terrain. In fact, during climate oscillations, vegetation types (e.g., grasslands and forests) are more likely to have replaced each other over time rather than accumulated (Qiu et al., 2011), thus causing the local extinction of entire communities. Furthermore, only few radiations are centered on the QTP s.s., with a significant number of local endemics (e.g., in *Poa*, *Rhodiola*, *Saussurea*, *Salix*, and *Astragalus*) (see Table 4), all of these genera being strongly specialized to extreme alpine habitats. Some other diverse genera, such as *Salix*, would benefit from azonal vegetation near bodies of water.

TABLE 5 | Geographical elements of seed plants of the Hengduan Mountains, the Himalayas, the Qinghai-Tibet Plateau s.s., and the Qinghai-Tibet Plateau s./ at the family level.

Areal types	Hengduan		Himalayas		QTP s.s		QTP s./	
	No.	%	No.	%	No.	%	No.	%
1 Cosmopolitan	49	26.63	46	25.99	43	35.25	50	25.00
2 Pantropic	52	28.26	58	32.77	27	22.13	61	30.50
3 Tropical Asia and Tropical America disjuncted	13	7.07	11	6.21	4	3.28	13	6.50
4 Old World Tropics	2	1.09	3	1.69	0	0.00	4	2.00
5 Tropical Asia to Tropical Australasia	4	2.17	4	2.26	2	1.64	4	2.00
6 Tropical Asia to Tropical Africa	1	0.54	2	1.13	1	0.82	2	1.00
7 Tropical Asia (Indo-Malaysia)	4	2.17	3	1.69	0	0.00	4	2.00
2–7 Total tropical elements	76	41.30	81	45.76	34	27.87	88	44.00
8 North Temperate	32	17.39	29	16.38	27	22.13	33	16.50
9 East Asia and North America disjuncted	11	5.98	7	3.95	6	4.92	11	5.50
10 Old World Temperate	1	0.54	1	0.56	1	0.82	1	0.50
11 Temperate Asia	0	0.00	0	0.00	0	0.00	0	0.00
12 Mediterranean, West Asia to Central Asia	2	1.09	2	1.13	4	3.28	4	2.00
13 Central Asia	2	1.09	2	1.13	2	1.64	2	1.00
14 East Asia	10	5.43	9	5.08	4	3.28	10	5.00
15 Endemic to China	1	0.54	0	0.00	1	0.82	1	0.50
8–15 Total temperate elements	59	32.07	50	28.25	45	36.88	62	31.00
Total	184	100	177	100.00	122	100.00	200	100.00

TABLE 6 | Geographical elements of seed plants of the Hengduan Mountains, the Himalayas, the Qinghai-Tibet Plateau s.s., and the Qinghai-Tibet Plateau s./ at the genus level.

Areal types	Hengduan		Himalayas		QTP s.s		QTP s./	
	No.	%	No.	%	No.	%	No.	%
1 Cosmopolitan	82	6.44	75	6.54	79	10.59	94	5.97
2 Pantropic	171	13.43	152	13.26	48	6.43	196	12.45
3 Tropical Asia and Tropical America disjuncted	17	1.34	18	1.57	3	0.40	21	1.33
4 Old World Tropics	64	5.03	60	5.24	7	0.94	81	5.15
5 Tropical Asia to Tropical Australasia	38	2.99	36	3.14	9	1.21	50	3.18
6 Tropical Asia to Tropical Africa	63	4.95	55	4.80	10	1.34	71	4.51
7 Tropical Asia (Indo-Malaysia)	127	9.98	150	13.09	11	1.47	188	11.94
2–7 Total tropical elements	480	37.71	471	41.10	88	11.80	607	38.56
8 North Temperate	233	18.30	211	18.41	213	28.55	251	15.95
9 East Asia and North America disjuncted	66	5.18	50	4.36	31	4.16	69	4.38
10 Old World Temperate	81	6.36	70	6.11	93	12.47	106	6.73
11 Temperate Asia	21	1.65	17	1.48	24	3.22	28	1.78
12 Mediterranean, West Asia to Central Asia	18	1.41	28	2.44	54	7.24	62	3.94
13 Central Asia	32	2.51	35	3.05	53	7.10	60	3.81
14 East Asia	181	14.22	149	13.00	75	10.05	204	12.96
15 Endemic to China	79	6.21	40	3.49	36	4.83	93	5.91
8–15 Total temperate elements	711	55.85	600	52.36	579	77.61	873	55.46
Total	1273	100.00	1146	100.00	746	100.00	1574	100.00

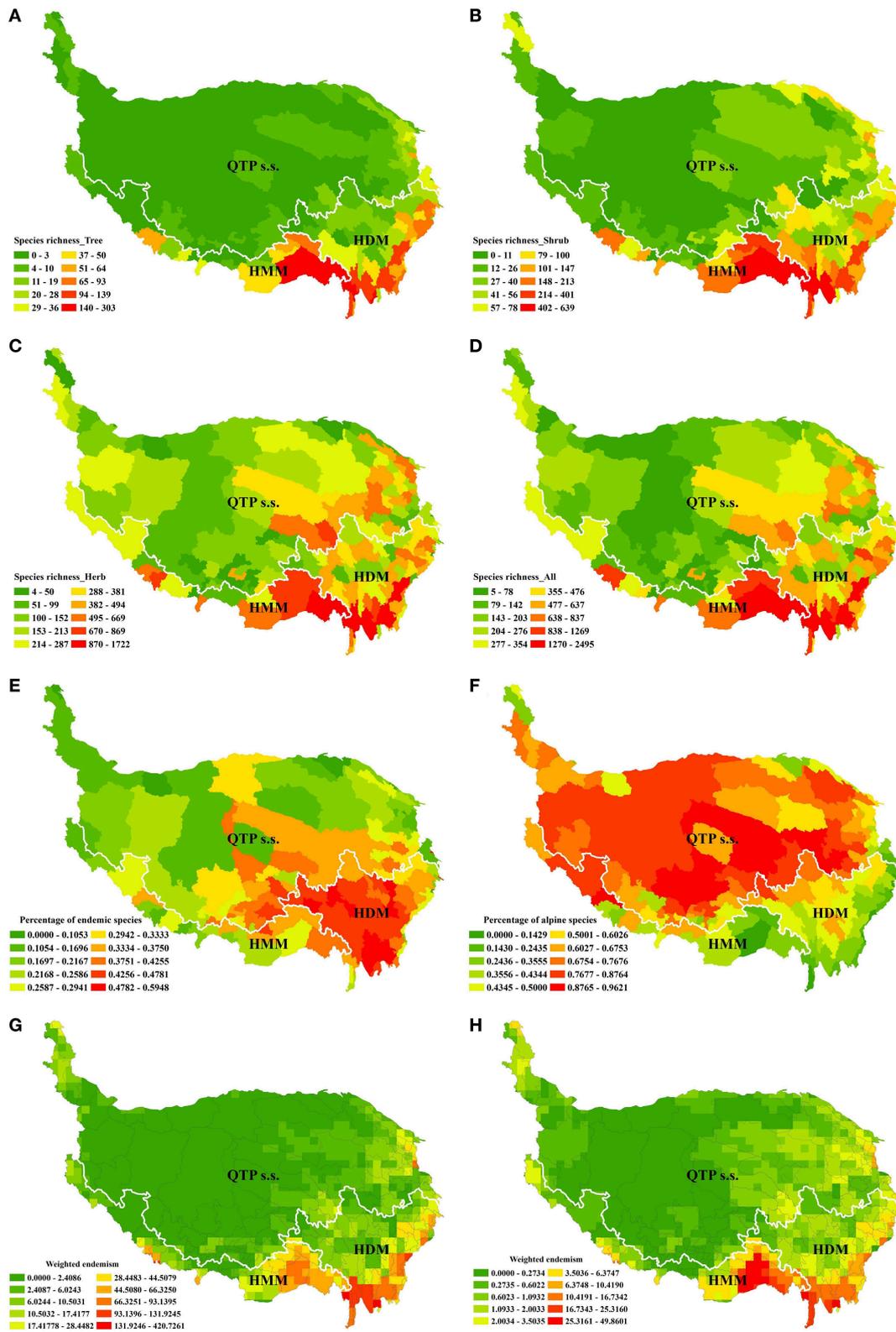
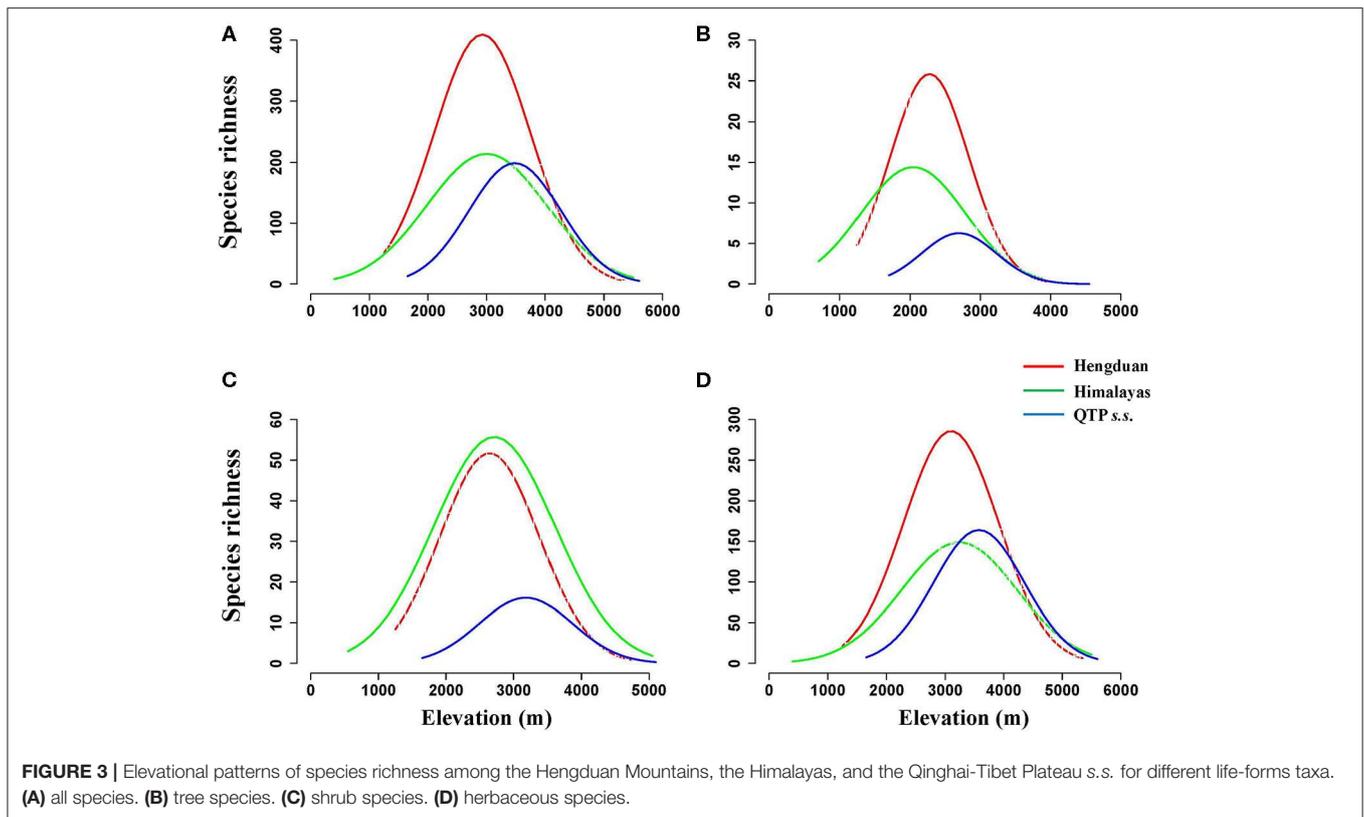


FIGURE 2 | Patterns of floristic diversity in the Qinghai-Tibet Plateau *s.l.*. **(A–D)** patterns of species richness for tree, shrub, herbaceous plants, and all species. **(E)** pattern of ratio of endemic species in a county. **(F)** pattern of ratio of alpine species in a county. **(G,H)** patterns of weighted endemism at the species and genus levels. HDM, the Hengduan Mountains; HMM, the Himalayas; QTP *s.s.*, the Qinghai-Tibet Plateau *sensu stricto*. White lines represent the boundary between the Hengduan Mountains, the Himalayas and the QTP *s.s.*.



Molecular studies for *Saussurea* and *Rhodiola* have suggested that these genera radiated because of geological processes, such as the uplifts of the QTP s.s. (Zhang et al., 2014; Xu et al., 2019). This has been strongly criticized (see Renner, 2016), and an increasing number of publications tend to associate radiations with faster and more profound changes in the environment such as climate oscillations (among others, Mosbrugger et al., 2018; Muellner-Riehl et al., 2019).

Gradients of Species Diversity Within the Hengduan Mountains, the Himalayas and the QTP s.s.

Over the entire region, plant species diversity decreases gradually, roughly from the southeastern edge of the QTP s.l.. This pattern follows general trends in terms of climate (warm and wet to cold and dry), mean elevation (low to high) and topography (high to low ruggedness). This overall gradient superimposes with more local ones in each of the three regions (see Figures 2A–D). For example, the southern part of the Hengduan Mountains (higher precipitation and ruggedness) is richer in term of species than the northern part (relatively less wet and rugged). The southern Hengduan Mountains is also a transition zone with incursion of numerous species of the Yunnan Plateau (subtropical taxa) (Zhu, 2012), for example in the Gaoligong Mountains and the Nujiang Valley. Furthermore, this area is a vital glacial refugium (Yu et al., 2019b). To the contrary, vast expands of the northern part of the Hengduan Mountains are

more high plateau-like, such as the Litang and the Songpan Plateaus, which were likely more prone to extinction during glaciations and less permeable to the immigration of non-local floristic elements.

In the Himalayas, there is a clear decrease of diversity pattern from the east to the west. The eastern Himalayas have similar climatic and topographic conditions as the southern Hengduan Mountains. However, in the central and western Himalayas, because most areas are located within the rain shadow zone and influenced slightly by the monsoon system, species become less (drought-tolerant plant dominate), only several valleys in the southern slope (plentiful rainfall) such as the Zhangmu Valley, Jilong Valley, Yadong Valley have remarkable amounts of species.

The interior of the QTP s.l. (i.e., QTP s.s.) is extremely high with average elevation above 4,500 m. Alpine species dominate this region (see Figure 2F). Duo to climatic and elevational gradients from the southeast to the northwest, species diversity reduces gradually along this direction. However, the southeastern and eastern margin areas (e.g., the Three-River Headwaters Region, the Qilian Mountains) have a higher topographic heterogeneity and are partially affected by the monsoons, leading to the occurrence of relatively more species there. Moreover, several glacial micro-refugia were located within these areas that could preserve multiple species, as suggested in Yu et al. (2019b). However, that study included a large spectrum of species without considering their ecological preferences, and was not specific to any thermal belts. Although Yu et al. (2019b) probably revealed some hotspots of genetic diversity on the QTP platform, their

importance may vary whether or not only species with similar ecology were included in the analysis.

Altitudinal Gradients

Along elevational gradients, species richness for each life-form taxa presents a hump-shape curve across all three regions (see **Figure 3**). Such pattern was also discussed in the Gaoligong Mountain (Wang et al., 2007), the Gongga Mountain (Zu et al., 2019), the eastern Himalayas (Sharma et al., 2019), and other montane regions in the world (e.g., Quintero and Jetz, 2018). Usually, ecological and evolutionary factors account for this elevational pattern of species diversity (Laiolo et al., 2018). For example, within the Hengduan Mountains, the other side of large valleys such as the Jinsha River, Mekong River (Lancang) and Salween River (Nujiang) are rain shadow zones, indicating that the foot of a valley is much drier, and thus only the mid-elevation zone is wetter and can sustain more species. From an evolutionary standpoint, habitats of intermediate elevation could have had more time for speciation/diversification and the accumulation of species, thus generating high species richness at mid-elevations [mountain museum hypothesis; (Stephens and Wiens, 2002)]. This hypothesis was for example supported by a study targeting the high endemic diversity of the QTP *s.l.* (Yu et al., 2019a). Muellner-Riehl et al. (2019) also argued that mountain systems hosting relatively more species were those characterized by the largest overlap in altitudinal gradients of temperatures between glacials and interglacials. The mid-elevation thermal belts might thus have acted as refugium in general. In addition, occurrence of high species richness along the elevation varies among these three regions. We assumed that this could be related to climatic variation of these regions. For example, the Hengduan Mountains and Himalayas have more suitable climatic conditions (abundant rainfall and varied temperature profiles), leading to a high species richness at lower elevation than that of the QTP *s.s.*

CONCLUSIONS

The vast QTP *s.l.* is highly heterogeneous (e.g., topography, climate) and its biodiversity is the result of complex and contrasting evolutionary histories. The Hengduan Mountains, the Himalayas and the QTP *s.s.* demonstrate distinct floristic features and diversity patterns, which in parts reflects their individual process of species assemblage deriving from their contrasting geological and climatic histories. Exploring the

current distribution of biodiversity and its evolutionary origin is essential to forecasting threats to biodiversity under current and future climate changes and human activities. Our study thus represent a first necessary step to suggest valid conservation strategies to protect this unique and precious plateau region.

DATA AVAILABILITY STATEMENT

All datasets for this study are included in the article/**Supplementary Material**.

AUTHOR CONTRIBUTIONS

HY, AF, ZC, SM, and GX conceived the study. HY and XG collected the data. HY led data analysis and writing. AF contributed to the writing. All authors contributed critically to the drafts and gave final approval for publication.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fevo.2020.00136/full#supplementary-material>

Appendix 1 | The seed plant list for the Hengduan Mountains.

Appendix 2 | The seed plant list for the Himalayas in China.

Appendix 3 | The seed plant list for the QTP *s.s.*

REFERENCES

- An, Z., Kutzbach, J. E., Prell, W. L., and Porter, S. C. (2001). Evolution of Asian monsoons and phased uplift of the Himalaya-Tibetan plateau since late miocene times. *Nature* 411, 62–66. doi: 10.1038/35075035
- Antonelli, A., Kissling, W. D., Flantua, S. G., Bermúdez, M. A., Mulch, A., Muellner-Riehl, A. N., et al. (2018). Geological and climatic influences on mountain biodiversity. *Nat. Geosci.* 11, 718–725. doi: 10.1038/s41561-018-0236-z
- Chen, Y., Deng, T., Zhou, Z., and Sun, H. (2018). Is the East Asian flora ancient or not? *Natl. Sci. Rev.* 5, 920–932. doi: 10.1093/nsr/nwx156
- Cun, Y., and Wang, X. (2010). Plant recolonization in the Himalaya from the southeastern Qinghai-Tibetan Plateau: Geographical isolation contributed to high population differentiation. *Mol. Phylogenet. Evol.* 56, 972–982. doi: 10.1016/j.ympev.2010.05.007
- Deng, T., and Ding, L. (2015). Palealtimetry reconstructions of the Tibetan Plateau: progress and contradictions. *Natl. Sci. Rev.* 2, 417–437. doi: 10.1093/nsr/nwv062
- Ding, L., Spicer, R. A., Yang, J., Xu, Q., Cai, F., Li, S., et al. (2017). Quantifying the rise of the Himalaya orogen and implications for the South Asian monsoon. *Geology* 45, 215–218. doi: 10.1130/G38583.1

- Ebersbach, J., Muellner-Riehl, A. N., Michalak, I., Tkach, N., Hoffmann, M. H., Röser, M., et al. (2017). In and out of the Qinghai-Tibet Plateau: divergence time estimation and historical biogeography of the large arctic-alpine genus *Saxifraga* L. *J. Biogeogr.* 44, 900–910. doi: 10.1111/jbi.12899
- Editorial Committee of Flora Reipublicae Popularis Sinicae (1959–2004). *Flora Reipublicae Popularis Sinicae*. Beijing: Science Press.
- Favre, A., Michalak, I., Chen, C., Wang, J., Pringle, J. S., Matuszak, S., et al. (2016). Out-of-Tibet: the spatio-temporal evolution of *Gentiana* (Gentianaceae). *J. Biogeogr.* 43, 1967–1978. doi: 10.1111/jbi.12840
- Favre, A., Päckert, M., Pauls, S. U., Jähnig, S. C., Uhl, D., Michalak, I., et al. (2015). The role of the uplift of the Qinghai-Tibetan Plateau for the evolution of Tibetan biotas. *Biol. Rev.* 90, 236–253. doi: 10.1111/brv.12107
- Gébelin, A., Mulch, A., Teysseier, C., Jessup, M. J., Law, R. D., and Brunel, M. (2013). The miocene elevation of mount everest. *Geology* 41, 799–802. doi: 10.1130/G34331.1
- He, X., Burgess, K. S., Gao, L., and Li, D. (2019a). Distributional responses to climate change for alpine species of *Cyananthus* and *Primula* endemic to the Himalaya-Hengduan mountains. *Plant Divers.* 41, 26–32. doi: 10.1016/j.pld.2019.01.004
- He, X., Burgess, K. S., Yang, X., Ahrends, A., Gao, L., and Li, D. (2019b). Upward elevation and northwest range shifts for alpine *Meconopsis* species in the Himalaya-Hengduan mountains region. *Ecol. Evol.* 9, 4055–4064. doi: 10.1002/ece3.5034
- Hoorn, C., Perrigo, A., and Antonelli, A. (2018). “Mountains, climate and biodiversity: an introduction,” in *Mountains, Climate, and Biodiversity*, eds C. Hoorn, A. Perrigo, and A. Antonelli (New York, NY: Wiley) 1–14.
- Hughes, C. E., and Atchison, G. W. (2015). The ubiquity of alpine plant radiations: from the Andes to the Hengduan Mountains. *New Phytol.* 207, 275–282. doi: 10.1111/nph.13230
- Körner, C., Jetz, W., Paulsen, J., Payne, D., Rudmann-Maurer, K., and Spehn, E. M. (2017). A global inventory of mountains for bio-geographical applications. *Alpine Bot.* 127, 1–15. doi: 10.1007/s00035-016-0182-6
- Lagomarsino, L. P., Condamine, F. L., Antonelli, A., Mulch, A., and Davis, C. C. (2016). The abiotic and biotic drivers of rapid diversification in Andean bellflowers (*Campanulaceae*). *New Phytol.* 210, 1430–1442. doi: 10.1111/nph.13920
- Laiolo, P., Pato, J., and Obeso, J. R. (2018). Ecological and evolutionary drivers of the elevational gradient of diversity. *Ecol. Lett.* 21, 1022–1032. doi: 10.1111/ele.12967
- Liang, Q., Xu, X., Mao, K., Wang, M., Wang, K., Xi, Z., et al. (2018). Shifts in plant distributions in response to climate warming in a biodiversity hotspot, the Hengduan Mountains. *J. Biogeogr.* 45, 1334–1344. doi: 10.1111/jbi.13229
- Linder, H. P. (2001). Plant diversity and endemism in sub-Saharan tropical Africa. *J. Biogeogr.* 28, 169–182. doi: 10.1046/j.1365-2699.2001.00527.x
- Liu, J., Möller, M., Provan, J., Gao, L., Poudel, R. C., and Li, D. (2013). Geological and ecological factors drive cryptic speciation of yews in a biodiversity hotspot. *New Phytol.* 199, 1093–1108. doi: 10.1111/nph.12336
- Liu, J., Sun, Y., Ge, X., Gao, L., and Qiu, Y. (2012). Phylogeographic studies of plants in China: advances in the past and directions in the future. *J. Syst. Evol.* 50, 267–275. doi: 10.1111/j.1759-6831.2012.00214.x
- Lu, L., Mao, L., Yang, T., Ye, J., Liu, B., Li, H., et al. (2018). Evolutionary history of the angiosperm flora of China. *Nature* 554, 234–238. doi: 10.1038/nature25485
- Luebert, F., and Weigend, M. (2014). Phylogenetic insights into Andean plant diversification. *Front. Ecol. Evol.* 2:27. doi: 10.3389/fevo.2014.00027
- Marchese, C. (2015). Biodiversity hotspots: a shortcut for a more complicated concept. *Glob. Ecol. Conserv.* 3, 297–309. doi: 10.1016/j.gecco.2014.12.008
- Marshall, C. A., Wieringa, J. J., and Hawthorne, W. D. (2016). Bioquality hotspots in the tropical African flora. *Curr. Biol.* 26, 3214–3219. doi: 10.1016/j.cub.2016.09.045
- Matuszak, S., Muellner-Riehl, A. N., Sun, H., and Favre, A. (2016). Dispersal routes between biodiversity hotspots in Asia: the case of the mountain genus *Tripterosperrum* (Gentianinae, Gentianaceae) and its close relatives. *J. Biogeogr.* 43, 580–590. doi: 10.1111/jbi.12617
- Mosbrugger, V., Favre, A., Muellner-Riehl, A. N., Päckert, M., and Mulch, A. (2018). “Cenozoic evolution of geo-biodiversity in the Tibeto-Himalayan region,” in *Mountains, Climate, and Biodiversity*, eds C. Hoorn and A. Antonelli (New York, NY: Wiley-Blackwell), 111–122.
- Muellner-Riehl, A. N., Schnitzler, J., Kissling, W. D., Mosbrugger, V., Rijdsdijk, K. F., Seijmonsbergen, A. C., et al. (2019). Origins of global mountain plant biodiversity: testing the ‘mountain-geobiodiversity hypothesis’. *J. Biogeogr.* 46, 2826–2838. doi: 10.1111/jbi.13715
- Mulch, A., and Chamberlain, C. P. (2006). The rise and growth of Tibet. *Nature* 439, 670–671. doi: 10.1038/439670a
- Qiu, Y., Fu, C., and Comes, H. P. (2011). Plant molecular phylogeography in China and adjacent regions: tracing the genetic imprints of Quaternary climate and environmental change in the world’s most diverse temperate flora. *Mol. Phylogenet. Evol.* 59, 225–244. doi: 10.1016/j.ympev.2011.01.012
- Quintero, I., and Jetz, W. (2018). Global elevational diversity and diversification of birds. *Nature* 555, 246–250. doi: 10.1038/nature25794
- R Core Team. (2016). *R: A Language and Environment for Statistical Computing*. Vienna: R Foundation for Statistical Computing. Available online at: <http://www.R-project.org/>
- Rahbek, C., Borregaard, M. K., Antonelli, A., Colwell, R. K., Holt, B. G., Nogues-Bravo, D., et al. (2019). Building mountain biodiversity: geological and evolutionary processes. *Science* 365, 1114–1119. doi: 10.1126/science.aax0151
- Ren, G., Conti, E., and Salamin, N. (2015). Phylogeny and biogeography of *Primula* sect. *Armerina*: implications for plant evolution under climate change and the uplift of the Qinghai-Tibet Plateau. *BMC Evol. Biol.* 15:161. doi: 10.1186/s12862-015-0445-7
- Renner, S. S. (2016). Available data point to a 4-km-high Tibetan Plateau by 40 Ma, but 100 molecular-clock papers have linked supposed recent uplift to young node ages. *J. Biogeogr.* 43, 1479–1487. doi: 10.1111/jbi.12755
- Sharma, N., Behera, M. D., Das, A. P., and Panda, R. M. (2019). Plant richness pattern in an elevation gradient in the Eastern Himalaya. *Biodivers. Conserv.* 28, 2085–2104. doi: 10.1007/s10531-019-01699-7
- Stephens, P. R., and Wiens, J. J. (2002). Explaining species richness from continents to communities: the time-for-speciation effect in emydid turtles. *Am. Nat.* 161, 112–128. doi: 10.1086/345091
- The Angiosperm Phylogeny Group, Chase, M. W., Christenhusz, M., Fay, M. F., Byng, J. W., Judd, W. S., Soltis, D. E., et al. (2016). An update of the Angiosperm Phylogeny Group classification for the orders and families of flowering plants: APG IV. *Bot. J. Linn. Soc.* 181, 1–20. doi: 10.1111/boj.12385
- The Comprehensive Scientific Expedition to the Qinghai Xizang Plateau Chinese Academy of Sciences (1993-1994). *Vascular Plants of the Hengduan Mountains*. Beijing: Science Press.
- Wang, E., Kirby, E., Furlong, K. P., Van Soest, M., Xu, G., Shi, X., et al. (2012). Two-phase growth of high topography in eastern Tibet during the Cenozoic. *Nat. Geosci.* 5, 640–645. doi: 10.1038/ngeo1538
- Wang, Z., Tang, Z., and Fang, J. (2007). Altitudinal patterns of seed plant richness in the Gaoligong mountains, south-east Tibet, China. *Divers. Distrib.* 13, 845–854. doi: 10.1111/j.1472-4642.2007.00335.x
- Wu, Y. (2008). *The Vascular Plants and Their Eco-geographical Distribution of the Qinghai-Tibetan Plateau*. Beijing: Science Press.
- Wu, Z., Raven, P. H., and Hong, D. (1994-2013). *Flora of China*. St. Louis, MO: Missouri Botanical Garden Press.
- Wu, Z., Zhou, Z., Sun, H., Li, D., and Peng, H. (2006). *The Areal Types of Seed Plants and Their Origin and Differentiation*. Kunming: Yunnan Science and Technology Press.
- Xing, Y., and Ree, R. H. (2017). Uplift-driven diversification in the Hengduan Mountains, a temperate biodiversity hotspot. *Proc. Natl. Acad. Sci. U.S.A.* 114, 3444–3451. doi: 10.1073/pnas.1616063114
- Xu, B., Li, Z., and Sun, H. (2014). Plant diversity and floristic characters of the alpine subnival belt flora in the Hengduan Mountains, SW China. *J. Syst. Evol.* 52, 271–279. doi: 10.1111/jse.12037
- Xu, L., Herrando-Moraira, S., Susanna, A., Galbany-Casals, M., and Chen, Y. (2019). Phylogeny, origin and dispersal of *Saussurea* (Asteraceae) based on chloroplast genome data. *Mol. Phylogenet. Evol.* 141:106613. doi: 10.1016/j.ympev.2019.106613
- Yan, L., Liu, J., Möller, M., Zhang, L., Zhang, X., Li, D., et al. (2015). DNA barcoding of *Rhododendron* (Ericaceae), the largest Chinese plant genus in biodiversity hotspots of the Himalaya-Hengduan Mountains. *Mol. Ecol. Resour.* 15, 932–944. doi: 10.1111/1755-0998.12353
- Yu, H., Deane, D. C., Sui, X., Fang, S., Chu, C., Liu, Y., et al. (2019a). Testing multiple hypotheses for the high endemic plant diversity of the Tibetan Plateau. *Glob. Ecol. Biogeogr.* 28, 131–144. doi: 10.1111/geb.12827

- Yu, H., Favre, A., Sui, X., Chen, Z., Qi, W., and Xie, G. (2019b). Mapping the genetic patterns of plants in the region of the Qinghai-Tibet Plateau: implications for conservation strategies. *Divers. Distrib.* 25, 310–324. doi: 10.1111/ddi.12847
- Yu, W., Liu, M., Wang, H., Mill, R. R., Ree, R. H., Yang, J., et al. (2015). Towards a comprehensive phylogeny of the large temperate genus *Pedicularis* (*Orobanchaceae*), with an emphasis on species from the Himalaya-Hengduan mountains. *BMC Plant Biol.* 15:176. doi: 10.1186/s12870-015-0547-9
- Zhang, D., Ye, J., and Sun, H. (2016). Quantitative approaches to identify floristic units and centres of species endemism in the Qinghai-Tibetan Plateau, southwestern China. *J. Biogeogr.* 43, 2465–2476. doi: 10.1111/jbi.12819
- Zhang, J., Meng, S., Allen, G. A., Wen, J., and Rao, G. (2014). Rapid radiation and dispersal out of the Qinghai-Tibetan Plateau of an alpine plant lineage *Rhodiola* (*Crassulaceae*). *Mol. Phylogenet. Evol.* 77, 147–158. doi: 10.1016/j.ympev.2014.04.013
- Zhang, Y., Li, B., and Zheng, D. (2002). A discussion on the boundary and area of the Tibetan Plateau in China. *Geogr. Res.* 21, 1–8. doi: 10.11821/yj2002010001
- Zhu, H. (2012). Biogeographical divergence of the flora of Yunnan, southwestern China initiated by the uplift of Himalaya and extrusion of Indochina block. *PLoS ONE* 7:e45601. doi: 10.1371/journal.pone.0045601
- Zhu, H. (2015). Biogeography of Shangri-la flora in Southwestern China. *Phytotaxa* 203, 231–244. doi: 10.11646/phytotaxa.203.3.2
- Zu, K., Luo, A., Shrestha, N., Liu, B., Wang, Z., and Zhu, X. (2019). Altitudinal biodiversity patterns of seed plants along Gongga Mountain in the southeastern Qinghai-Tibetan Plateau. *Ecol. Evol.* 9, 9586–9596. doi: 10.1002/ece.3.5483

Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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