Species richness, phylogenetic diversity and phylogenetic structure patterns of exotic and native plants along an elevational gradient in the Himalaya

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Abstract

Background: So far, macroecological studies in the Himalaya have mostly concentrated on spatial variation of overall species richness along the elevational gradient. Very few studies have attempted to document the difference in elevational richness patterns of native and exotic species. In this study, this knowledge gap is addressed by integrating data on phylogeny and elevational distribution of species to identify the variation in species richness, phylogenetic diversity and phylogenetic structure of exotic and native plant species along an elevational gradient in the Himalaya.

Results: Species distribution patterns for exotic and native species differed; exotics tended to show maximum species richness at low elevations while natives tended to predominate at mid-elevations. Native species assemblages showed higher phylogenetic diversity than the exotic species assemblages over the entire elevational gradient in the Himalaya. In terms of phylogenetic structure, exotic species assemblages showed majorly phylogenetic clustering while native species assemblages were characterized by phylogenetic overdispersion over the entire gradient.

Conclusions: The findings of this study indicate that areas with high native species richness and phylogenetic diversity are less receptive to exotic species and vice versa in the Himalaya. Species assemblages with high native phylogenetic overdispersion are less receptive to exotic species than the phylogenetically clustered assemblages. Different ecological processes (ecological filtering in case of exotics and resource and niche competition in case of natives) may govern the distribution of exotic and native species along the elevational gradient in the Himalaya.

Keywords: Exotic species, Native species, Himalaya, Elevational gradient, Richness pattern

Introduction

Exotic species (non-native species introduced by human activity to the areas outside their natural distribution) cause both ecological and economic harms by lowering biological diversity, resource availability, human health and altering ecosystem functions (Sala et al. 2000; Pimentel et al. 2005; Vilà and Hulme 2017). Consequently, their management and control becomes highly important. Generally, there are multiple and largely interactive factors that underlies the distribution of exotic species (Catford et al. 2009; Tarasi and Peet 2017) and it is equivocally debated as to what factors promote or reduce the spread of exotic species (Ibáñez et al. 2009; Gantchoff et al. 2018). In recent times, ecologists have increasingly used evolutionary relationships and phylogenetic metrics to analyze and investigate how exotic species are assembled and structured in new regions outside their areas of natural distribution (Cadotte et al. 2009, 2010; Lososová et al. 2015).

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Mountains occupy a total of 13.8 million km² and 12.5% of the Earth’s land surface (excluding Antarctica) and harbor one-quarter of total terrestrial species (Körner et al. 2016; Hoorn et al. 2018). Mountains not only act as harbingers of biodiversity and cradles of evolution, but they also provide numerous ecosystem services (Kier et al. 2016; Hoorn et al. 2018). In recent times, however, the spread of exotic species is threatening the quality and quantity of the ecosystem services accrued from mountain ecosystems (Pauchard et al. 2009; McDougall et al. 2011). In the lake cluster of Pokhara Valley of middle mountain Nepal, for example, yearly economic losses worth 0.19 million USD have been reported due to ecological invasion or spread of exotic species (Pathak et al. 2021). The mountain ecosystems that were earlier thought to be immune to the ecological invasion process (Averett et al. 2016) are now being increasingly invaded by exotic species due to climate and land-use changes, human population growth, road development, agricultural intensification and other anthropogenic activities (Dietz et al. 2006; Pauchard et al. 2009). There are increasing reports of colonization by non-native exotic species in the mountain areas around the world (Seipel et al. 2012; Morueta-Holme et al. 2015; Tecco et al. 2016).

The Himalayan ecosystems are increasingly experiencing more threats than any other mountain ecosystems of the world (Pandit 2017). The region has witnessed escalating anthropogenic pressures in recent decades, including extensive hydropower and infrastructure development coupled with urbanization, road construction, incessant tourism, deforestation and hunting (Pandit et al. 2014). Moreover, global warming and related climate changes have emerged as serious challenges to the Himalayan biodiversity (Telwala et al. 2013; Manish and Pandit 2019). Some estimates suggest that the region is warming at a rate three times faster than that of the global average (Shrestha et al. 2012; Pandit et al. 2014). Given the fact that the rich biodiversity of the Himalaya is critical for local livelihoods (Pandit et al. 2014; Pandit 2017) and spread of exotic species detrimentally affects the biodiversity (Pimentel et al. 2005), analyzing elevational distribution of exotic species in the Himalaya is very much needed (Khuroo et al. 2011; Ahmad et al. 2018). Most existing studies on the elevational species richness patterns in the Himalaya have focussed on documenting richness patterns for either native or endemic plant species (Vetaas and Grytnes 2002; Bhattarai and Vetaas 2003; Manish et al. 2017; Manish and Pandit 2018). Studies on exotic species richness patterns in the Himalaya have been generally been limited to preparing a checklist (Khuroo et al. 2007; Sekar 2012; Jaryan et al. 2013). Only a few studies have examined how patterns of native and exotic species differ in the Himalaya and even these have yielded contrasting results (see Kosaka et al. 2010; Khuroo et al. 2011; Bhattarai et al. 2014). This study attempts to address this knowledge gap by examining the elevational trends in species richness and phylogenetic diversity and phylogenetic structure of native and exotic species along a large elevational gradient (300–5500 m) in the Sikkim Himalaya by using a combination of traditional taxonomic metric (species richness), phylogenetic metric of diversity (phylogenetic diversity), and indices of phylogenetic structure (nearest taxon index and net relatedness index).

Materials and methods

Study area

The present study was conducted in Sikkim, a constituent state of India and a part of the Eastern Himalayan global biodiversity hotspot (Fig. 1). The study area is situated between 27° 00′46″ N and 28° 07′48″ N latitude and 88° 00′58″ E and 88° 55′25″ E longitude and possesses a completely mountainous terrain with a natural elevational gradient ranging from 246 to 7873 m (Fig. 1). The total geographical area of the state is 7096 km² and constitutes just 0.22% of India’s total geographical area. More than 59% of Sikkim’s geographic area lies above 3000 m, with slopes above 30% and 70% constituting more than 52% and 10% of the land area, respectively (CISMHE 2007). Sikkim receives a mean annual rainfall of about 2534 mm, with monthly maximum rainfall of 480 mm in July and minimum rainfall of 19 mm in December. The mean annual temperature of Sikkim is 18 °C, with minimum temperature of −13 °C in alpine zones and maximum temperature of 26 °C at low and mid-elevations (CISMHE 2007). In recent years, there have been reports of fast colonization and encroachment of exotic plant species in farmlands, pristine forests and grasslands, leading to productivity declines and economic losses (Sharma and Rai 2012; Pandit 2017).

Species data

Data on species occurrence, nomenclature, family and their respective elevational distribution and elevational range were collected from previously published records, databases, checklists and online resources (see Additional file 1: Appendix S1). The recorded plant species were categorized into exotic species and native species; exotic species included any non-native species, i.e., naturalized, casual, invasive and cultivated species that occur outside their center of origin and natural adapted ranges (see Jaryan et al. 2013; Pyšek et al. 2004; Raghubanshi et al. 2005), while native species included the native Himalayan species that are found in the Sikkim Himalayan region (see Additional file 1: Appendix S1). To determine the nativity of species, a host of published and online

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sources including efloras (http://www.efloras.org/), Flora Himalaya database (www.leca.univ-savoie.fr/db/florhy/infos.html), Plants of the World database (http://www.plantsoftheworldonline.org/), Flora of British India (Hooker 1875–1897), Flora of Sikkim (Hajra and Verma 1996; Srivastava 1998) and other published records, literature and checklists were used (see Additional file 1: Appendix S1).
Species richness patterns

The elevational gradient of Sikkim Himalaya between 300 and 5500 m was divided into 100 m equal elevational bands. The lowermost elevational limit (300 m) represents the southernmost point of the study region (Melli Bazar, 246 m rounded to the nearest upper 100 m elevation), while the highest elevational limit (5500 m) marks the limit beyond which no species were recorded. Thus, there were a total of 53 elevational bands (100 m each) for both exotic and native species. Interpolation method was used to estimate the number of species or species richness in each elevational band. A species was assumed to be present in each elevational band between its recorded lowermost and uppermost elevational limits and no species was assumed to have an elevational distribution range of less than 100 m (see Grytnes and Vetaas 2002; Manish et al. 2017). Interpolation method has been reported to be the most accurate estimator of species richness among all parametric and non-parametric estimators in the Himalayan region (see Manish et al. 2017). A species recorded at a single site was given an elevational range of 100 m and included in the analyses in accordance with the standard literature (see Bhattarai and Vetaas 2006; Manish et al. 2017). Friedman’s super smoother function in TIBCO Spotfire 5 + 8.1 software was used to decipher the species richness patterns along the elevational gradient and analyze the relationship between species richness and elevation for both exotic and native species. Friedman’s super smoother function creates a smooth curve around the data points by automatically determining the level of smoothness using cross-validation technique (Friedman and Silverman 1989). In this way, the Friedman’s super smoother function has an advantage over the normal regression techniques like generalized linear models (GLM) and generalized additive models (GAM) that are commonly used to document species richness patterns (see Vetaas and Grytnes 2002; Bhattarai and Vetaas 2006; Bhatt et al. 2012; Manish et al. 2017) as the curve generated by the super smoother function is dictated by the data contained in the dataset itself while curves generated by GLM and GAM are generally limited by the statistical parameters and distribution of the models (Pedersen et al. 2019).

Phylogenetic analyses

A megatree approach was used to construct family-level phylogenetic trees for exotic and native plant species using the Phylomatic tool of Phylocom ver. 4.2 (Webb et al. 2008) and the modified protocol of Webb et al. (2008) and Manish and Pandit (2018). Such a megatree approach has been successfully used in a number of previous studies for estimating the phylogenetic diversity and structure in different regions where molecular data for species do not exist (Li et al. 2014; Qian et al. 2016; Manish et al. 2021). For preparing the family-level phylogenetic trees, the data on the exotic and native species occurrences in different elevational bands of the study area and a backbone family-level Angiosperm Phylogeny Group IV seed plant supertree (R20160415.new) of Gastauer et al. (2017) in the Phylocom software were used. The Phylomatic tool in Phylocom software generated uncalibrated family-level phylogenetic trees for exotic and native species with equal branch lengths. The branch lengths in these uncalibrated trees were calibrated using the Branch Length Adjustment algorithm (BLADJ) tool (Webb 2000) in Phylocom based on the angiosperm nodes contained in the dated tree of Bell et al. (2010). The BLADJ calibrated family-level phylogenetic trees of exotic and native species were then used as a base for determining phylogenetic diversity and phylogenetic structure patterns of exotic and native species along the elevational gradient in the study area.

Phylogenetic diversity patterns

Several ways to quantify phylogenetic diversity (PD) exist in literature (see Kellar et al. 2015). Of these, some are based on abundance measures (Hill numbers, Chao et al. 2010), incidence data (Faith’s PD, Faith 1992), or entropy calculations (Rao’s quadratic entropy, Rao 1982). In this study, I used Faith’s PD measure to quantify PD of each elevational band as it can be used with species occurrence data (Faith 1992; Morlon et al. 2011). Faith’s PD can be defined as the minimum total length of all the existing phylogenetic branches required to span across all taxa in a given phylogenetic tree (Faith 1992; Faith and Baker 2006).

PD values of exotic and native species were calculated for each 100 m elevational band using the respective BLADJ calibrated family-level phylogenetic trees in R 2.14.0 software (R Development Core Team 2011) utilizing the “picante” package (Kembel et al. 2010). Friedman’s super smoother function in TIBCO Spotfire 5 + 8.1 software was used to explore the relationship between PD and elevation in the study area. Pearson’s correlation coefficient was used to analyze the significance and strength of association between PD and species richness of exotic and native species along the elevational gradient of the study area.

Phylogenetic structure

Phylogenetic structure indicates the phylogenetic distribution of taxa in a community or species assemblages (Manish and Pandit 2018). Net relatedness index (NRI) and nearest taxon index (NTI) are used as indices of phylogenetic structure to analyze the variation of phylogenetic structure of exotic and native plant species
assemblages along the Sikkim Himalayan elevational gradient. Both the indices have been quite frequently used in literature to document the phylogenetic structure of plant communities (Li et al. 2014; Manish and Pandit 2018). NRI is used to analyze phylogenetic relatedness among different species in a species assemblage by measuring the mean phylogenetic distance between taxa in a phylogenetic tree, while NTI is used to analyze phylogenetic relatedness by calculating mean nearest taxon distance between terminal taxa in a phylogenetic tree (Kamilar et al. 2015; Manish and Pandit 2018). The following modified formulae of Webb (2000) and Webb et al. (2008) were used to calculate the values of NRI and NTI in each 100 m elevational band for exotic and native species in R 2.14.0 using the “picante” package (Kembel et al. 2010):

\[
\text{NRI} = -1 \times \left( \frac{\text{MPD}_o - \text{MPD}_r}{\text{sdMPD}_r} \right),
\]

\[
\text{NTI} = -1 \times \left( \frac{\text{MNTD}_o - \text{MNTD}_r}{\text{sdMNTD}_r} \right),
\]

where NRI = net relatedness index, MPD<sub>o</sub> = observed mean phylogenetic distance in the phylogenetic tree, MPD<sub>r</sub> = expected mean phylogenetic distance of species assemblages that were randomized, sdMPD<sub>r</sub> = standard deviation of mean phylogenetic distance of species assemblages that were randomized, NTI = nearest taxon index, MNTD<sub>o</sub> = observed mean nearest taxon distance in the phylogenetic tree, MNTD<sub>r</sub> = expected mean nearest taxon distance of species assemblages that were randomized, sdMNTD<sub>r</sub> = standard deviation of mean nearest taxon distance of species assemblages that were randomized. Randomized species assemblages were created using a null model that shuffled the phylogeny tip labels across all taxa included in a phylogenetic tree using the R package “picante” (Kembel et al. 2010). The statistical significance of the NRI and NTI values in each elevational band was assessed against this null model with a total of 999 iterations (see Manish and Pandit 2018).

The positive values of NRI and NTI (>0) in the current study indicated presence of phylogenetic clustering (i.e., more species belonged to closely related evolutionary families than that expected by chance), while negative values of NRI and NTI (<0) indicated phylogenetic overdispersion (i.e., more species belonged to distantly related evolutionary families than that expected by chance). The variation of NRI and NTI along the elevational gradient of the study area was analyzed with the help of Friedman’s super smoother function in TIBCO Spotfire S + 8.1 software.

The results of phylogenetic structure analyses revealed that exotic species showed phylogenetic clustering (NRI and NTI values >0) between 300–1400 m, 1900–2500 m, 4100–4500 m and phylogenetic overdispersion (NRI and NTI values <0) between 1500–1800 m, 2600–4000 m and 4600–5500 m elevations (Fig. 5). On the other hand, native species showed phylogenetic overdispersion throughout the elevational gradient except between 1200–1500 m and 2500–3200 m elevations where phylogenetic clustering was observed (Fig. 5). This indicated that the exotic and native species exhibited different phylogenetic structures within the same study area and also revealed that the indices of phylogenetic structure (NRI
Fig. 2 Variation of (A) Species richness (SR) and (B) Phylogenetic diversity (PD) of exotic and native plant species along the elevational gradient in Sikkim Himalaya. Both exotic and native plant species showed unimodal hump-shaped curves for PD and SR along the elevational gradient. The primary Y-axis in both the plots represents SR and PD of native species, while secondary Y-axis represents SR and PD of exotic species. The fitted lines in both the plots were generated using Friedman’s super smoother function and do not imply any cause–effect relationship.
and NTI) did not differ much for both exotic and native taxa (Fig. 5). Thus, the choice of the metric of phylogenetic structure did not influence the overall results in the current study.

**Discussion**

The results of the present study indicate that the exotic and native plant species show different species distribution and phylogenetic diversity patterns along the Sikkim Himalayan elevational gradient, albeit both of them showed hump-shaped pattern. Hump-shaped species distribution patterns have been reported to be the most generalized and common pattern of species distribution along the elevational gradient in the Himalaya (see Vetaas and Grytnes 2002; Grytnes and Vetaas 2002; Carpenter 2005; Bhatt et al. 2012; Manish et al. 2017; Manish 2019). The hump-shaped patterns in species distributions may be correlated to a combination of different ecological factors such as climatic constraints, dispersal limitations or environmental filters that governs the distribution of various taxa in the study area. For example, ambient energy and water availability have been reported to govern the overall distribution of plant species richness patterns and both these factors also show hump-shaped patterns when plotted along the elevational gradient in the Sikkim Himalaya region (Manish et al. 2017). Similarly, productivity and environmental heterogeneity have a unimodal relationship with species richness that may

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**Fig. 3** Plots showing the zones of maximum and minimum species richness and phylogenetic diversity of (A, B) exotic and (C, D) native plant species in the Sikkim Himalayan region. Exotics were found to predominantly occupy the low elevations, while natives were predominantly limited to the mid-elevations in the study area both in terms of species richness and phylogenetic diversity.
result in hump-shaped patterns (Stein et al. 2014; Fraser et al. 2015). Exotic plant species showed maximum species richness and phylogenetic diversity at low elevations (700–1000 m and 500–1000 m, respectively), while maximum native plant species and native phylogenetic diversity were recorded at high elevations (3300–3700 m and 2800–3500 m, respectively). Similar results have been reported earlier in the Himalaya by Kosaka et al. (2010), Ahmad et al. (2018) and Dar et al. (2018). Kosaka et al. (2010) found maximum exotic species richness in the low-elevation tropical forests (100–1600 m) in the neighboring state of Arunachal Pradesh (a part of the same Eastern Himalayan biogeographic region as that of Sikkim). Ahmad et al. (2018) found maximum exotic species richness in 1000–1100 m elevational bands in Himachal Pradesh (Western Himalaya), while Dar et al. (2018) reported a general decline in exotic plant species richness with increasing elevation in the Kashmir Himalayan region (Western Himalaya). More exotic species richness is expected at low elevations in the mountains because of favorable climates, low geographic and climate isolation, high human disturbance, high road and human population densities, less environmental heterogeneity and less extreme habitats (Becker et al. 2005; Pauchard et al. 2009). Low temperature, low ecosystem productivity, short growing season and low precipitation at high elevations limit spread of exotic species in the Himalaya (Manish et al. 2017). In a way, the present results conform to the “directional ecological filtering” hypothesis proposed by Alexander et al. (2011) that posits its uniform decline of exotic species along the elevational gradient in the mountain areas as exotic species cannot successfully establish under the extreme environmental conditions that keep on increasing with elevation. Due to greater stress at high elevations, only exotic species with narrow climatic ranges and specialists with adaptation to the extreme ecophysiological conditions such as low temperature, slow growth rate, etc., are able to survive (Alexander et al. 2011; Marini et al. 2013). Another reason that can be attributed to the decline of exotic species richness at high elevations is the lack of pre-adaptation of exotic species to survive in extreme conditions at high elevations and lack of colonization of high elevation areas (Alexander et al. 2016). Native species on the other hand showed maximum preponderance and phylogenetic diversity at the mid-elevations. The mid-elevations of the study area have been previously reported to be the regions of highest endemic plant species richness and endemic phylogenetic diversity (Manish and Pandit 2018). This is likely due to multiple evolutionary diversifications and adaptive radiations linked to habitat isolation and vicariance driven by isolated mountain peaks and highly dissected river valleys at the mid-elevations in the Sikkim Himalaya (see Pandit 2017; Manish and Pandit 2018). The mid-elevations of Sikkim Himalaya are also the harbingers of most ancestral/primitive plant species in the region and are the areas where speciation, diversification and accumulation events have been occurring for the longest periods of time in the region (Manish and Pandit 2018). Thus, the occurrence of native plant species richness and phylogenetic diversity maxima at higher elevations than the exotic species in the present study seems to be justified. By contrary, Bhattarai et al. (2014) reported that both exotic and native species richness peaked in the same elevational range (600–1100 m) in the Nepal Himalaya. A probable reason for this can be that Bhattarai et al. (2014) included just one life-form (trees) for comparison of distribution patterns of exotic and native species in their study. Trees irrespective of whether they are endemic, native or exotic generally tend to dominate at low elevations in the Himalaya due to the ecophysiological and climatic limitations (Manish et al. 2017). The present study is more comprehensive than Bhattarai et al. (2014) as it includes all life-forms (trees, shrubs, climbers and herbs) and presents
Fig. 5 Plots showing the variation of phylogenetic structure of exotic and native plant species along the Sikkim Himalayan elevational gradient. Phylogenetic structure was measured using net relatedness index (NRI) and nearest taxon index (NTI). Overall, exotics showed phylogenetic clustering (NRI and NTI > 0) and natives showed phylogenetic overdispersion (NRI and NTI < 0) along the elevational gradient. In both the plots, the primary Y-axis represents NRI and NTI values of native species, while secondary Y-axis represents NRI and NTI values of exotic species. The fitted lines in both the plots were generated using Friedman's super smoother function and do not imply any cause–effect relationship.
a more generalized pattern. Elsewhere, species richness and diversity patterns can vary with life-forms and geographic status (native and endemic) in the Himalaya (see Manish et al. 2017; Manish 2019). Therefore, it is more meaningful to analyze the diversity and distribution patterns only after including the entire life-form spectrum of the taxa under investigation. An important finding of the present study is that the areas with low native species richness and phylogenetic diversity (low elevations in the current study) offer greater ecological opportunities for the establishment of exotic species. This study therefore tends to agree with the hypothesis that ‘low native species diversity promotes exotic species diversity’ (Elton 1958; Naeem et al. 2000; Dark 2004) as opposed to the ‘high native species diversity promotes exotic species diversity’ hypothesis (Stohlgren et al. 1999; Levine 2000; Gilbert and Lechowiez 2005). High native species richness leads to decreased exotic receptivity mainly because of increased resource competition, high functional diversity and lack of empty niches (Elton 1958; Kennedy et al. 2002; Sol et al. 2012).

The results of phylogenetic structure analyses in the present study revealed that exotic species generally showed phylogenetic clustering and native species showed phylogenetic overdispersion in the areas with respective maximum species richness and phylogenetic diversity (Fig. 5). These findings are similar to that of Cadotte et al. (2010) and Carvallo et al. (2014) who reported low phylogenetic diversity and phylogenetic clustering in the exotic clades and phylogenetic overdispersion in the native communities. Various authors have pointed out that the presence of phylogenetic clustering indicates ‘ecological filtering’ while phylogenetic overdispersion indicates ‘competition’ (Webb et al. 2002; Li et al. 2014; Manish and Pandit 2018). It is likely that ‘directional ecological filtering’ (Alexander et al. 2011) shapes the phylogenetic structure and overall species composition of exotic species in Sikkim Himalaya by only selecting lineages with ability to tolerate anthropogenic disturbances, high stress and fluctuating resources at low elevations and minimum temperature at high elevations. The overrepresentation of a few families such as Poaceae, Asteraceae, Fabaceae, Cyperaceae and Amaranthaceae that together make up 56% of the exotic flora is also likely to dictate the phylogenetic clustering of exotic species with majority of the species belonging to these families being annuals with rapid growth and having the ability of producing copious seeds with good dispersal ability (Additional file 1: Table S1). The results also reveal that less number of exotic species is found in those areas where there is high phylogenetic overdispersion of native species assemblages (Fig. 5). Phylogenetic overdispersion is the result of strong competition between closely related species due to the dependence on the same resources as a result of their shared ancestry (Webb et al. 2002; Lososová et al. 2015). Thus, the intense resource competition and lack of empty niches drive out the exotic species in areas with high native phylogenetic diversity and overdispersion.

It is amply clear from the present study that the exotic species are currently limited to the low-elevation tropical regions in the Himalaya. This is however likely to change in near future with climate change and increased human disturbance including urbanization, road development, agricultural intensification, tourism, etc. With global warming, the low temperature constraint on exotic species might be reduced and lead to the spread of exotic species towards high elevations. Unplanned road development can cause recurring disturbances including changing species composition, local hydrology, soil ecology, microclimate, and fragmenting habitats (Forman et al. 2003; Müllerová et al., 2011). These changes can both serve as important sources of propagules and facilitate the establishment of exotic species (Johnston and Pickering 2001; Godfree et al. 2004). In many studies, it has been shown that exotic species first colonize the low elevations and then spread upwards with increasing disturbances and anthropogenic pressures with time (Guo et al. 2018; Yang et al. 2018). This poses a serious threat to the alpine ecosystems of the Himalaya that are important harbingers of endemic biodiversity and medicinal plants and are important sources of local livelihoods (see Pandit 2017). The conservation managers and policy-makers would have to carefully and pragmatically manage development including road and infrastructure development, urbanization and tourism in areas with high native biodiversity and perhaps it would be better to leave these areas as undisturbed and excluded from any kind of anthropogenic disturbance.

Conclusions
This study presents five broad ideas: (i) species distribution patterns for exotic and native plant species differ in the Himalaya; exotics tend to show maximum species richness at low elevations while natives tend to predominate at mid-elevations; (ii) native plant species assemblages show higher phylogenetic diversity than the exotic species assemblages over the entire elevational gradient in the Himalaya; (iii) exotic plant species assemblages show majorly phylogenetic clustering while native species assemblages are characterized by phylogenetic overdispersion over the entire Himalayan elevational gradient; (iv) areas with high native species richness and phylogenetic diversity are less receptive to exotic species and vice versa in the Himalaya; (v) the assembly of exotic plant species in the Himalaya is shaped by ecological
filtering while native species assemblages are dictated by resource and niche competitions. Future studies in the region should be directed towards examining the critical functional traits that influence community assembly and species assemblage patterns of the natives and exotics, and how niche partitioning occurs in the invaded and un-invaded habitats. It is also necessary to do long-term monitoring of the changes in the distribution of exotic species along the elevational gradient and determine the temporal shift in their distribution patterns in order to assess the effect of climate change in the region. Climate change in particular is expected to favor the expansion of low-elevation exotic plant species into the temperate and alpine habitats. This can greatly threaten the existence of endemic species that predominate in these habitats due to their low tolerance to changing environmental conditions and encroaching exotic species. Studies along these lines would help the conservation managers and policymakers to effectively limit the spread of exotic species in the Himalaya.

Supplementary Information
The online version contains supplementary material available at https://doi.org/10.1186/s13717-021-00335-z.

Additional file 1: Appendix S1. List of published records used for collating data on exotic and native species occurrences, nomenclatures, families and their respective elevational distribution and elevational ranges in Sikkim Himalaya. Table S1. List of exotic angiosperm species of the Sikkim Himalayan region. The species have been listed alphabetically according to their names. Table S2. List of native angiosperm species of the Sikkim Himalayan region. The species have been listed alphabetically according to their names.

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Authors’ contributions
KM conceptualized the study, collected and analyzed the data and wrote the manuscript. The author read and approved the final manuscript.

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Availability of data and materials
The raw data for this manuscript (species list) are included in the Additional Material.

Declarations

Ethics approval and consent to participate
No formal approval is required for this study since this article does not contain any studies with human participants, animals or on-site data collection.

Consent for publication
Not applicable.

Competing interests
The author declares that he has no competing interests.
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