#### Responses to Referee #1

### **Major Comment:**

[Comment] I read the manuscript "Detecting climatically driven phylogenetic and morphological divergence among spruce species (Picea) worldwide" with delight. The manuscript explored the relationship between climate and the phylogenetic and morphological divergence of Picea species in the Northern Hemisphere, based on 3388 georeferenced distribution sites. Temperature and precipitation parameters were the main driving factors for the primary divergence of spruce phylogeny and morphology, respectively. The climatic data extracted from current spruce locations captured the ecological divergence among spruces. These results suggested that the primary divergence of morphology and phylogeny among the spruces tended to be driven by different selective pressures. The data and methods are appropriate for this study; the manuscript is well organized and presented. I found that the manuscript has a merit for publication in the journal Biogeosciences, pending on the authors can address my following concerns. My major concern is that if the climatic data used in this manuscript can represent the local climate of the distribution sites. The Worlclim dataset has been widely used in biogeographic studies. It can be used to surrogate the local climate in plain areas. However, it cannot represent the local climate in the mountainous regions because of the coarse resolution (about 1km). In the mountainous regions, 1 km distance may cover an elevational interval of hundreds of meters (and therefore introduce several degrees of difference in temperature). The authors need to discuss the caveat of using this dataset.

[Response] This is a good question. The coarse resolution (about 1km) of climate data from The Worlclim dataset would likely weaken the potential to interpret spruce distribution and divergence. Discussion of the caveat of using this dataset is absolutely needed and will be done when we get the chance to revise this MS. Thank you!

Nevertheless, we have the confidence that the climate data from The WorlClim dataset used in this study is suitable for interpreting the overall pattern, i.e., the first several splits that represent "the primary trigger" that led to the divergence of among spruce, which are the major findings of this study. As we can see from Fig.1 (a, b), instead of elevation gradient, the geographical distribution of both the three phylogenetic clades and the morphological groups (quadrangular leaves versus flattened leaves) is largely determined by horizontal gradients (latitude and longitude). Specifically, clade-1 is a Eurasian clade and clade-2 is a North American clade, while clade-3 is an Asian clade with only one North American species. As for as the morphological groups, spruces flattened leaves tend to occur in eastern Asia and the beach area of the northern America, while spruce with quadrangular leaves distribute in the rest part of the whole distribution range. Given this base, the 1km-resolution of climate data we used in this study should be robust to interpret this large scale pattern.

We confess this dataset may give rise to some uncertainties in the context of the detection of some subtle variation such as within-species variation or among elevation-sensitive species. In this case, although the splits at the terminal nodes are between species, that is to say that we don't have any within-species variation, the caveat with respect to the dataset must be discussed. Further works that focus on the driving force underlying the variation of within-species or among elevation-sensitive species should use high resolution climate data.

# **Specific points:**

[Comment 1] Line 66: "Nearly 34 species" should be "Thirty-four species"

[Response] We will make this change as suggested.

[Comment 2] Line 83: "niche conservatism" is not a process, but a pattern (result of the processes)

[Response] Agree! We will change the "process" to "pattern" in the revised manuscript.

*[Comment 3]* Line 130: "Between 34 and 35 species" is conflict to the "34 species" (line 66) Line 132 (and other areas): "flora of China" should be "Flora of China"

[Response] Thank you. We will make this change as suggested.

[Comment 4] Line 145, 148, 152: delete "approximately"

[Response] Thank you. We will make this change as suggested.

[Comment 5] Line 159-166: set abbreviations for the climatic variables (and use the abbreviations afterwards)

[Response] Thank you. We will make this change as suggested.

[Comment 6] Line 216-224: move to the "Materials and Methods" section

# Reference

Harris I, Jones PD, Osborn TJ, Lister DH (2014) Updated high-resolution grids of monthly climatic observations - the CRU TS3.10 Dataset. International Journal of Climatology, **34**, 623-642.

Hijmans RJ, Cameron SE, Parra JL, Jones PG, Jarvis A (2005) Very high resolution interpolated climate surfaces for global land areas. International Journal of Climatology, **25**, 1965-1978.

Kriticos DJ, Webber BL, Leriche A, Ota N, Macadam I, Bathols J, Scott JK (2012) CliMond: global high-resolution historical and future scenario climate surfaces for bioclimatic modelling.

Methods in Ecology and Evolution, 3, 53-64.

#### Responses to Referee #2

### **Major Comment:**

[Comment 1] Wang et al., analyzed the relations between current climate and ecological (phylogenetic and morphological) divergence among spruce species at a global scale. The topic is suitable for Biogeosciences, but I do not think it is suitable for this special issue "Ecosystem processes and functioning across current and future dryness gradients in arid and semi-arid lands". The range of spruce (we could see in Figure 1) is not only limited in arid and semi-arid lands, but also covers a lot of other more wet regions. The main results of this paper are clear that phylogenetic and morphological divergence is driven by different climate variables, i.e., temperature for phylo and precipitation for morpho. But I have several questions/comments, which need carefully revised by the authors.

[Response] The major reason why we submitted this MS to the special issue "Ecosystem processes and functioning across current and future dryness gradients in arid and semi-arid lands" is that most spruce species are very important taxa in arid and semi-arid lands worldwide. Detecting climatically driven phylogenetic and morphological divergence among spruce species worldwide would deepen the understanding of ecosystem processes and functioning in arid and semi-arid lands.

To address this point, in this response, we extracted the Aridity Index (AI) for each point from the Global Aridity Index (Global-Aridity) and the Global Potential Evapo-Transpiration (Global-PET) Geospatial Database (<a href="http://www.cgiar-csi.org/2010/04/134/">http://www.cgiar-csi.org/2010/04/134/</a>), According the 1997 UNPE standard (Middleton& Thomas, 1997) climate zone classification, 8 spruce species are in arid and Semi-Arid areas, 11 spruce species in Dry sub-humid areas, and 14 spruce species in humid areas. According to the scenario of global climate change, there would have severe and widespread droughts in the next 30-90 years over land areas resulting from either decreased precipitation and/or increased evaporation, and the significant increases in aridity do occur in many subtropical and adjacent humid regions [1, 2]. When overlapping the spruce sampling point to the future Aridity Changes Map (Fig. 1, 2 in this response), nearly all the spruce species whose original distribution in sub-humid and humid areas would subject to drought stress.

The Special issue "Ecosystem processes and functioning across current and future dryness gradients in arid and semi-arid lands" aims to provided platform for researches in plant species associations, plant distribution along environmental gradients, which is not only applicable for species distributed in arid and semi-arid areas, but also for the species subjected to aridity stress in future. Our findings would be helpful for management strategies and inform policy to climate change in future.

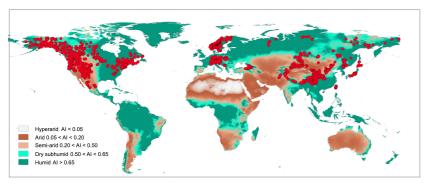


Fig. 1. The locations of sampling point in the study at different climate zone. The background image was the map of Global Aridity Index which obtained online (http://www.cgiar-csi.org)by the CGIAR-CSI with the support of the International Center for Tropical Agriculture (CIAT).

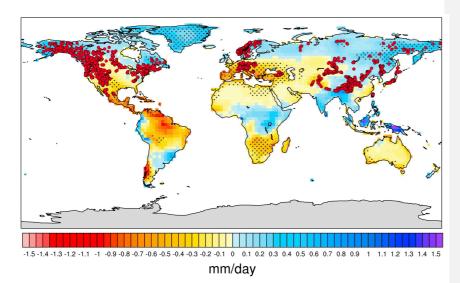


Fig. 2. The locations of sampling point in Aridity changes within the 21st century. The background image was the map of Changes in P–Rn/ $\lambda$  comparing present-day (1980–2000) and future climate (2080–2100) following the RCP8.5 pathway.(P, precipitation; Rn, net radiation;  $\lambda$ , latent heat of vaporization).[1].

*[Comment 2]* Firstly, the abstract is not well written. There are too much information on methods and results. Usually, we first need some background, importance of the study, come up with the question, and what we do, what we found, and finally the importance of our findings. Furthermore, some information in the abstract are repeated, e.g., line 30-34 and line 40-41. Other minor problems in Abstract include: bioclimatic or cli- matic (should be consistent here and other parts of this paper); global and northern hemisphere are different; there are ecological divergence, phylo divergence,

morpho divergence and divergence, should be consistent or clearly defined; younger nodes are called remaining/terminal/end nodes/splits, should be consistent.

[Response] Thank you for your comments. We will make these changes as suggested.

[Comment 3] The use of current climate: The author also discussed this problem. As far as I know, there are not only current climate data in worldclim, but also paleoclimate. Although the paleoclimate there only date back to LGM, it still could reflect the climate situation for a longer time to some extent. I am wondering if this paleoclimate could be a better choice than current climate.

[Response] Due to expansion and retreat occurred in the past, the present distribution of spruce is different from the distribution of the fossil locations. Thus, paleoclimate data does not necessarily match the present distribution. The 3388 data points of the 33 spruce species were sampled on present locations. Current climate data should be more appropriate to interpret current distribution pattern of spruce species.

[Comment 4] The authors did PCA analysis and found that the first three axes could explain 75.67% of the variance, but the following analysis used 8 separate climate variables. I want to know why they choose these 8 variables, and not using the first three axes. Generally, 75% variance is OK. I guess the 4 temperature variables the author used are highly correlated, as well as the four precipitation variables. So I doubt the necessary to use so many climate variables. By the way, the numbers in the main text is not consistent with the numbers in table 1. For instance, the first axis explains 43.52% of the variance in Table 1, but 29.8% in the main text; other numbers are also wrong. In table 1, the first column, how did the authors choose the bold variables. I mean temperature seasonality is -0.928, and mean temperature of the coldest quarter is 0.946, higher than the AMT. The use of elevation is also questionable. The author at list did not discuss the effect of elevation in discussion.

[Response] We actually ran the SEEVA by taking all the 16 climate factors into account. To illustrate the results briefly and clearly, we need to reduce the redundant variables. We focused on how mean value, extreme values of climate factors influence spruce divergence. In addition, The climatic variables must have higher divergence indices for the first split on the phylogeny and morphology of *Picea*, and relatively higher loading on the five component axes. As a result, we mapped eight climate factors in the histograms on the phylogeny and morphology tree. Take an example, Min Temperature of Coldest Month and Mean Temperature of Coldest Quarter both have high loading on axis-1of PCA: 0.931 vs. 0.946, but the former has higher divergence indices than the latter (0.0764 vs. 05524 in the phylogeny and 0.18 vs. 0.08 in the morphology). We therefore illustrated the results of the former variable.

Table-1 showed the eigenvalues, variance percentages, cumulative percentages and correlations of 19 bioclimatic factors but the rotated percentages were shown in the text. We will revise this inconsistency. Thanks.

Spruce is elevation-sensitive. We selected elevation as a variable because it can demonstrate a direct view with respect to spruce divergence, which would be helpful to understand how topography influences spruce divergence.

**Specific points:** 

[Comment 1]: The results do not need to be divided into 6 parts, I think the last 4 parts could be merged into 1.

[Response] Agree. We will reorganize the text. We think 3.3, 3.4 and 3.5 should be merged into one section

[Comment 2]: Some logic in the text is not reasonable. For instance, in line 87, information before "thus" and after "thus", I don't think they are well connected; line 178-189, the sequence of these parts is mess, line 188-189 should move to the front of the introduction of the SEEVA. The come up with several hypotheses in the introduction also feel not well connected with the text there. Anyway, the authors need to carefully check this throughout in the text.

[Response] Thanks. We will check these during the revision stage.

[Comment 3] Line 148, mainland China and Taiwan?

[Response] Agree. "mainland China and Taiwan" should be more formal. We will check these.

[Comment 4] Line 158-166, I am wondering if it's necessary to list all the climate variables here. [Response] Agree. A full list of climate variables has been shown in Table 1. We will check these during the revision stage.

[Comment 5] Line 349-350, how did the authors conclude like that? 6. Cannot or could not? [Response] This paragraph highlighted the exceptions observed for a few sister groups or species in the phylogenetic tree to the overall pattern. We explained these exceptions as a result of geographical isolation and the limitation of the selected climate parameters that do not adequately describe the climatic determinants of spruce distributions.

#### Reference

Budantsev LY (1994) The Fossil Flora of the Paleogene Climatic Optimum in North Eastern Asia,

Springer Berlin Heidelberg.

 $Dai\ A\ (2012)\ Increasing\ drought\ under\ global\ warming\ in\ observations\ and\ models.\ Nature\ Climate$ 

Change, 3, 52-58.

Farjón A (2001) World Checklist and Bibliography of Conifers (Second edn.) England, Cambridge

University Press.

Giesecke T (2004) The Holocene Spread of Spruce in Scandinavia.

Greve P, Seneviratne SI (2015) Assessment of future changes in water availability and aridity.

Geophys Res Lett, 42, 5493-5499.

- Hang S (2002) Evolution of Arctic-Tertiary flora in Himalayan-Hengduan mountains. Acta Botanica Yunnanica, 24, 671-688.
- Kullman L (1995) New and firm evidence for Mid-Holocene appearance of Picea abies in the Scandes Mountains, Sweden. Journal of Ecology, **83**, 439-447.
- Liu JQ, Gao TG, Chen ZD, Lu AM (2002) Molecular phylogeny and biogeography of the Qinghai-Tibet Plateau endemic Nannoglottis (Asteraceae). Molecular Phylogenetics & Evolution, 23, 307-325.
- Ran JH, Wei XX, Wang XQ (2006) Molecular phylogeny and biogeography of Picea (Pinaceae): implications for phylogeographical studies using cytoplasmic haplotypes. Mol Phylogenet Evol, **41**, 405-419.
- Spribille T, Chytry M (2002) Vegetation surveys in the circumboreal coniferous forests: A review.

  Folia Geobotanica, 37, 365-382.
- Struwe L, Smouse PE, Heiberg E, Haag S, Lathrop RG (2011) Spatial evolutionary and ecological vicariance analysis (SEEVA), a novel approach to biogeography and speciation research, with an example from Brazilian Gentianaceae. Journal of Biogeography, 38, 1841-1854.

  Wu S, Yang YP, Fei Y (1995) On the flora of the alpine region in the Qinghai-Xizang (Tibet) Plateau,

China. Acta Botanica Yunnanica.

# A list of all relevant changes made in the manuscript

- We have simplified and specified abstract.
- We add a describe of climate zone of spruce species distributed in the method section and discussion section.
- We Added a statement of significance of the findings in relation to climate change to abstract, and added more discussions about significance and implications of the findings of this research in relation to future climate change.
- We have improved the English writing and sent the manuscript to a language service, made the paper more formal.

# Detecting climatically driven phylogenetic and morphological divergence among spruce species (*Picea*) worldwide

Guo-Hong Wang<sup>1\*</sup>#, He Li<sup>1</sup>#, Hai-Wei Zhao<sup>1, 2</sup>, Wei-Kang Zhang<sup>1</sup>

<sup>1</sup> State Key Laboratory of Vegetation and Environmental Change, Institute of Botany,

Chinese Academy of Sciences, Beijing 100093, China

<sup>2</sup> University of the Chinese Academy of Sciences, Beijing 100049, China

\*Address for correspondence: State Key Laboratory of Vegetation and Environmental

Change, Institute of Botany, Chinese Academy of Sciences, No. 20 Nanxincun,

Xiangshan, Beijing 100093, China

**Tel:** +86-010-6283-6585

Fax: +86-010-6259-0833

E-mail: ghwangaq@ibcas.ac.cn

#These authors contributed equally to this paper and should be regarded as co-first authors.

- > Running title: Global ecological divergence of spruce
- Number of words in the abstract: 245

Number of words in the paper: <u>6334</u>

Number of references: 49

Number of tables and figures: 6

> Supporting information: 1

**删除的内容:** 332

删除的内容: 5996

删除的内容: 46

# 1 Abstract

- 2 This study aimed to elucidate the relationship between climate and the phylogenetic
- 3 and morphological divergence of spruces (Picea) worldwide. Climatic and
- 4 georeferenced data were collected from a total of 3388 sites distributed within the
- 5 global domain of spruce species. A phylogenetic tree and a morphological tree for the
- 6 global spruces were reconstructed based on DNA sequences and morphological
- 7 characteristics. The spatial evolutionary and ecological vicariance analysis (SEEVA)
- 8 method was used to detect the ecological divergence among spruces. A divergence
- 9 index (D) with (0, 1) scaling was calculated for each climatic factor at each node for
- both trees. The results indicated that the annual mean values, extreme values and annual
- 11 range of the climatic variables were among the major determinants for spruce
- divergence. The ecological divergence was significant (P<0.001) for 185 of the 279
- comparisons at 31 nodes in the phylogenetic tree and for 196 of the 288 comparisons at
- 14 32 nodes in the morphological tree. Temperature parameters and precipitation
- parameters tended to be the main driving factors for the primary divergence of spruce
- phylogeny and morphology, respectively. Generally, the  $D_{\text{max}}$  of the climatic variables
- was smaller in the basal nodes than in the remaining nodes. The primary divergence of
- morphology and phylogeny among the investigated spruces tended to be driven by
- different selective pressures. Given the <u>future</u> climate <u>of severe and widespread drought</u>
- during the next 30-90 years over land areas, our findings shed light on the prediction of
- 21 spruce distributions under future climate change.

# 22 Keywords

**删除的内容:** R

删除的内容:,

删除的内容: A major finding is that t

删除的内容: scenario o

删除的内容: in

- Temperature, precipitation, natural selection, niche conservatism, parallel evolution,
- 29 speciation

# 1 Introduction

30

44

45

51

Environmental conditions play an important role in speciation (Mayr, 1947; Darnell and 31 Dillon, 1970; Wiens, 2004; Givnish, 2010; Schemske, 2010). However, quantitative 32 investigations of environmental influences on the origin and divergence of species are 33 less common than expected, especially in plants (Givnish, 2010). For example, although 34 taxonomic and phylogenetic studies have explicitly addressed phylogenetic and 35 morphological divergence among spruces (Farjón, 1990; Sigurgeirsson and Szmidt, 36 1993; Fu et al., 1999; Ran et al., 2006; Li et al., 2010; Lockwood et al., 2013), ecological 37 38 differentiation among sister groups or species remains unknown. Ecological vicariance differs from geographical vicariance (Wiley, 1988) and indicates the ecological 39 differentiation among sister groups or sister species within taxa, which provides 40 important information and ecological interpretations for the phylogenetic and 41 42 morphological divergence among taxa (Escudero et al., 2009; Struwe et al., 2011). 43 Spruce (Picea A. Dietrich) is an important component of boreal vegetation and

**删除的内容:** n

**删除的内容:** h

Spribille and Chytry, 2002). Nearly thirty-four species are recognized in the genus *Picea* worldwide (Farjón, 2001). Although taxonomic schemes of *Picea* based on

morphological characteristics differ slightly among authors, a consensus has been

reached for the criterion to determine the first several subdivisions (Liu, 1982; Farjón,

1990; Taylor, 1993; Fu et al., 1999). Accordingly, several sections within *Picea* have

subalpine coniferous forests and has a wide geographical range that covers the Northern

Hemisphere and extends from the Eurasian continent to North America (Farjón, 2001;

been classified based on morphological similarity. For example, section Picea and

section Casicta are characterized by quadrangular leaves and flattened leaves, respectively (Farjón, 1990). Alternatively, spruce species can be classified into phylogenetically distinct clades, namely clade-1, a Eurasian clade; clade-2, a North American clade; and clade-3, an Asian clade with one North American species (Ran et al., 2006; Lockwood et al., 2013). These chloroplast DNA sequence data-based classification schemes have the potential to reveal the phylogenetic affinity among spruces. We aimed to elucidate the ecological differentiations between sister groups or species identified based on their phylogenetic affinity and morphological similarity. A species' ecological niche depends on both the species' adaptation to its present habitat and the legacy of its ancestors (Wiens, 2004). Although species tend to retain similar ecological niches as their immediate ancestors, which is generally called phylogenetic niche conservatism, natural selection of ecologically important traits is the key process that determines the successful adaptation of incipient species (Peterson et al., 1999; Webb et al., 2002; Wiens and Graham, 2005). In addition, speciation tends to occur in geographic dimensions, whereas ecological differences evolve over time (Peterson et al., 1999). Thus, there should be tradeoff between niche conservatism and ecological differences among splits in the phylogeny of given taxa over evolutionary

54

55

56

57

58

59

60

61

62

63

64

65

66

67

68

69

70

71

72

73

74

75

time scales. Spruces likely originated in the early Tertiary or late Cretaceous era. The

fossil spruce species Picea burtonii Klymiuk et Stockey is regarded as the earliest fossil

spruces occurred over a long time scale, with a range of approximately 28 Ma to 1.8 Ma from the basal nodes to the terminal nodes (Lockwood et al., 2013). We hypothesize that there should be a relationship between the time since separation and the magnitude of ecological divergence or niche conservatism. Specifically, we expect to observe an increasing magnitude in terms of ecological divergence among sister groups from the basal nodes (older) to the terminal nodes (younger) on the evolutionary time scales because natural selection would favor species with high levels of ecological adaptation. Although phylogenetically close species are likely to be similar in appearance to one another, differences in the rate of evolution may substantially obscure these similarities (Baum et al., 2005). In the genus Picea, none of the morphology-based classification schemes are congruent with or supported by the schemes derived from cpDNA-based phylogenies. Therefore, spruce species within a taxonomic section are not always more similar in phylogenetic relatedness than those between sections or subsections, indicating that parallel evolution, i.e., the repeated appearance of similar characteristics that occur among distantly related species (Went, 1971; Hoekstra and Price, 2004; Schluter et al., 2004; Orr, 2005), occurs in *Picea*. Therefore, we hypothesize that the divergence of morphology and phylogeny among the investigated spruce species may be subject to different selective pressures under parallel evolution. Evolutionary trees indicate historical relationships among organisms (Baum et al., 2005). This "tree-thinking" approach has been used in almost all branches of biology to detect relatedness among organisms (Baum and Offner, 2008) and to examine

76

77

78

79

80

81

82

83

84

85

86

87

88

89

90

91

92

93

94

95

96

97

删除的内容: necessarily

删除的内容: which indicate indicates

ecological divergence between sister clades or species (Struwe et al., 2011). In this

100 study, tree-thinking methods were used to examine the ecological divergence among spruce species worldwide by reconstructing a phylogenetic tree and a morphological 101 tree. A dataset of spruce species was compiled to test our hypothesis by answering the 102 following three questions: are the climatic variables extracted from the current spruce 103 locations correlated with the divergence among spruces? If so, is there a relationship 104 between the time since separation and the magnitude of ecological divergence? Lastly, 105 is the morphological and phylogenetic divergence among spruce species subject to 106 different selective pressures? 107 108 2 Materials and Methods 2.1 Distribution data 109 The sampling sites were selected from within the entire natural range of spruce species 110

in the Northern Hemisphere, extending over a wide geographical range (latitude: 22.8-69.9°N; longitude: 53-165°W, 5-155°E; altitude: 103-4700 m a.s.l.) and exhibiting a steep climatic gradient, varying from a cold-temperate zone to a subtropical zone (Figure 1). Nearly 34 species are included in the genus Picea (Farjón, 2001). The global spruce checklist used in this study was primarily based on Farjón (2001) but refined according

111

112

113

114

115

116

117

120

121

to the flora of China (Fu et al., 1999). Specifically, because two species distributed in western China according to Farjón (1990), Picea retroflexa and P. aurantiaca, were 118 119 treated as a synonym and a variety of *P. asperata*, respectively, in the flora of China.

We followed the Chinese classification. Accordingly, the checklist used for this study contained 33 spruce species.

删除的内容: F

删除的内容: F

124 Georeferenced data for the 33 spruce species was partially downloaded from the Global Biodiversity Information Facility (GBIF), an international open data 125 infrastructure. Original data in the GBIF are derived from various sources, such as 126 natural history explorations (specimens or records) collected over the past 300 years, 127 current observations and automated monitoring programs (GBIF, 2015). We carefully 128 verified the original data downloaded from GBIF by excluding those data points with 129 geolocations outside of the natural distribution ranges (either horizontally or vertically 130 or both). As a result, 2397 point locations from the GBIF remained after the verification, 131 132 and they primarily represented spruce species in North America, Europe and East Asia (Japan and Korea Peninsula). Additional data for the spruce species from mainland 133 删除的内容: -referenced 134 China and Taiwan (991 locations for 16 species) were obtained from georeferenced herbarium collection records (490 sites) (Li et al., 2016) from the herbarium of the 135 136 Institute of Botany, Chinese Academy of Sciences; recent fieldwork (370 sites, 137 unpublished); and published sources (41 sites) (Tseng, 1991; Yang et al., 2002). As a result, 3388 point locations for the 33 spruce species were available for this analysis. 138 2.2 Climatic variables 139 140 A total of 19 climatic variables (Table 1) with a resolution of approximately 1 km<sup>2</sup> for 删除的内容: point 141 the 3388 Jocations were acquired and downloaded from WorldClim V. 1.4 (http://www.worldclim.org) (Hijmans et al., 2005). The values of each climate variable 142 删除的内容: software 143 at each site were extracted using the QGIS software package (http://qgis.osgeo.org),

144

145

and the final data were exported to an Excel worksheet for subsequent analysis.

DNA sequences were retrieved from the NCBI GenBank (www.ncbi.nlm.nih.gov) to reconstruct a phylogenetic tree of the 33 spruce species (Figure 2). This phylogenetic tree was constructed based on 3 plastid (trnL-trnF, trn-psbA, and trnS-trnG) and 2 mitochondrial (nad5 intron1 and nad1 intron2) DNA sequences, and it was equivalent to that of Lockwood et al. (2013), who proposed an improved phylogeny of Picea. In addition, we reconstructed a morphological tree of the 33 spruce species (Figure 3) based on Farjón (1990), Taylor (1993), and Fu et al. (1999). The first several splits in the tree primarily revealed divergence in the shape of the leaf cross section, the position of the stomatal line on the leaf surface, and the texture and arrangement of the seed scale, whereas traits such as the size of the leaf, seed cone and seed scale and the hairiness of the leaf or twig are important indicators for subsequent splits in the trees. The morphological tree and the phylogenetic tree contained 32 and 31 nodes, respectively. To detect ecological divergence among sister groups or species in the abovementioned trees, we used the spatial evolutionary and ecological vicariance analysis (SEEVA, Struwe et al., 2011), which can incorporate climatic data with phylogenetic data and morphological data using statistical methods to investigate ecological vicariance in speciation. The SEEVA compares the differences between each of the climatic variables for each node. A divergence index (D) with (0,1) scaling was calculated for each climatic factor at each node. D=0 indicates no difference between sister clades or groups, whereas D=1 indicates a maximum difference. Fisher's exact

149

150

151

152

153

154

155

156

157

158

159

160

161

162

163

164

165

166

167

168

169

170

test (Fisher, 1958), which generally provides a better P-value for tests with small

171 sample sizes, was performed to determine the significance of D. Because 31 and 32 172 independent tests were conducted for each of the climatic variables, a P-value less than 0.0016 indicated a significant difference in the ecological features for splits at a given 173 node after performing a Bonferroni correction, i.e., α=0.05/31 or 32≈0.0016. Details on 174 the calculations are available in Struwe et al. (2011). The SEEVA software can be 175 downloaded from http://seeva.heiberg.se. 176 A factor analysis was conducted to eliminate the redundant climatic variables, and a 177 principal component analysis (PCA) of the climatic variables was performed using the 178 删除的内容: we ran the SPSS statistical package (SPSS, Chicago, IL, USA). In addition, SEEVA was 179 删除的内容: all the performed by taking all 16 climate factors into account. To illustrate the results, we 180 删除的内容: briefly and clearly focused on how the mean, and extreme values of the individual climate factors 181 删除的内容: value. 删除的内容: must have both 182 influenced spruce divergence. The selected climatic variables were required to have 183 higher divergence indices for the first split on the phylogeny tree and the morphology 删除的内容: 184 tree and relatively higher loading on the five component axes. As a result, we mapped eight climate factors in the histograms, including four temperature variables (annual 185 mean temperature (AMT), minimum temperature of the coldest month (MTCM), 186 删除的内容: temperature 187 maximum temperature of the warmest month (MTWM) and annual temperature range 删除的内容: TAR 188 (ATR)) and four precipitation variables (annual precipitation (AP), precipitation of the 189 wettest month (PWM), precipitation of the driest month (PDM) and precipitation of the coldest quarter (PCQ)). In addition, elevation as a spatial variable was also used to 190 detect the ecological vicariance among sister groups because spruce is an elevation-191

sensitive taxon (Farjón, 1990; Taylor, 1993; Fu et al., 1999).

192

We compared the means of the nine abiotic variables among sister groups at several

删除的内容: the

key splits (i.e., the first two split levels) of both trees using a one-way analysis of

variance (ANOVA) to further interpret the observed ecological divergence.

3 Results

201

202

204

205

206

207

208

209

210

211

212

213

214

215

216

217

218

219

220

221

222

# 3.1 Variation in climatic variables

A factor analysis of the climatic variables across sampling sites revealed five dominant climatic gradients that accounted for 94.06% of the variance (Table 1). The first component, which had an eigenvalue of 8.27 and accounted for 43.52% of the variance, was a gradient characterized by variations in temperature variables. The second component, which had an eigenvalue of 3.60 and accounted for 18.93% of the variance, was a gradient characterized by variations in precipitation variables. The third, fourth and fifth components, which accounted for 13.21%, 11.89% and 6.51% of the variance, respectively, were characterized by variations in the precipitation of the driest month or quarter and precipitation seasonality; maximum temperature of the warmest month or quarter; and mean temperature of the wettest quarter and precipitation of the coldest quarter, respectively. 3.2 Ecological divergence among sister groups or species in the phylogeny of Picea Ecological divergence as indicated by the (0, 1) scaled index of D was significant (P<0.0016) for 185 of the 279 comparisons at 31 nodes in the phylogeny of Picea (see Table S1 in Appendix S1). The first split, which yielded node-2 (clade-1) and node-14 (clade-2 and clade-3), was significant for all 9 environmental variables. The annual temperature range (D=0.26\*) showed higher divergence, and it was followed by

elevation (D=0.25\*) and precipitation of the driest month (D=0.20\*). The spruce species in clade-1 tended to occur in climates with a lower annual temperature range and lower precipitation than the spruce species in node-14. The divergence within node-14 and between node-15 (clade-2) and node-22 (clade-3) was also significant for all 9 environmental variables. The parameters precipitation of the coldest quarter, precipitation of the driest month and precipitation of the wettest month had relatively high divergence (D=0.66\* to 0.42\*), elevation exhibited substantial divergence (D=0.46\*), whereas the temperature variables showed lower divergence (D=0.13\* to 0.31\*). Compared with clade-3, clade-2 occurred in climates with lower precipitation levels and a higher annual temperature range. Node-2 represented a split within clade-1 (the Eurasian clade) between a subclade at a higher elevational zone (in Caucasian area and Japan) with a warmer and wetter climate and a subclade at a lower elevational zone (esp. in boreal area) with a cold and dry climate. The elevation and temperature features showed relatively higher divergence (D=0.17\* to 0.38\*) than the precipitation variables (D=0.03\* to 0.23\*) (Figure 2, Table 2).

删除的内容: compared with

删除的内容: compared with

The ecological divergence for the subsequent 28 splits in the phylogeny of *Picea*, i.e., from node-3 to node-13 and from node-15 to node-31, was significant for approximately 63% of the comparisons. However, a universal pattern was not observed in terms of the ecological divergence for the remaining splits, which varied according to the sister groups or species. This finding suggests that a particular combination of environmental features is important for particular splits among sister groups or species (Figure 2, Table 2).

# 3.3 Ecological divergence among sister groups or species in the morphology of

# Picea

248

249

250

251

252

253

254

255

256

257

258

259

260

261

262

263

264

265

266

267

268

269

Ecological divergence was significant (P<0.0016) for 196 of the 288 comparisons at 32 nodes in the morphology tree of *Picea* (see Table S2 in Appendix S1). Of the 32 nodes, we focused on three splits that represent several key morphological divergence in Picea. Specifically, the split of node-1 represents divergence in the shape of the leaf cross section and the position of the stomatal line on the leaf surface, whereas the split of node-2 and node-25 represents divergence in the texture and seed scale arrangement. The remaining 29 splits, i.e., from node-3 to node-24 and from node-26 to node-32, reflect divergence in the leaf size, seed cone size, hairiness (pubescent vs. glabrous) and branchlet color, and these differences were significant for approximately 65% of the comparisons (Figure 3). The first split of the morphology-defined topology tree (Figure 3) yielded node-2 (leaf quadrangular) and node-25 (leaf flattened) and was significant for all 9 environmental variables. Precipitation features (D=0.16\*-0.54\*), predominantly precipitation of the wettest month, showed much stronger divergence than temperature features (D=0.05\*-0.18\*), with elevation showing a moderate divergence (D=0.30\*). Spruce species with quadrangular leaves tended to be favored by drier habitats with higher temperature annual ranges in lower elevational zones, which is inconsistent with the habitats for spruces with flattened leaves (Table 2). Such an overall pattern, however, does not always hold true for the sister groups or species that present different leaf cross

删除的内容: compared with

删除的内容: that of

删除的内容: necessarily

sections (flattened vs. quadrangular) but close phylogenetic relationships. Sister groups

are relevant examples (Figure 3). For example, although elevation was important for 274 275 the divergence between P. jezoensis and P. glehnii (node-10), temperature parameters were important for the divergence between P. wilsonii and P. purpurea (node-31). 276 The second-level splits in the morphological tree (Figure 3) yielded two pairs of sister 277 groups, namely node-3 vs. node-24 (within node-2) and node-26 vs. node-29 (within 278 node-25). These two pairs of spruce sister groups collectively indicated divergence in 279 the seed scale characteristics, i.e., closely arranged seed scales with a rigid woody 280 281 texture vs. loosely arranged seed scales with a thin, flexible, leathery or papery texture. For the split within node-2, elevation showed the highest divergence (D=0.51\*) and 282 was followed by annual temperature range (D=0.48\*) and precipitation of the driest 283 month (D=0.35\*), whereas the remaining climatic variables had significant but relative 284 285 low divergence (D=0.06\*-0.25\*). Compared with the results for node-24 (loosely 286 arranged seed scales), the species in node-3 (closely arranged seed scales) tended to occur in lower elevational zones with higher precipitation of the driest month and a 287 wider variation of annual temperature range (Table 2). For the split within node-25, 288 both the minimum temperature of the coldest month (D=0.46\*) and precipitation of the 289 driest month (D=0.43\*) showed substantial divergence, with a moderate divergence for 290 elevation (D=0.35\*). Compared with the results for node-26 (loosely arranged seed 291 scales), the species in node-29 (closely arranged seed scales) tended to occur in lower 292 elevational zones with higher temperature and greater precipitation in the coldest 293 quarter (Table 2). 294

or species at node-10, node-13, node-18, node-26 and node-31 in the phylogeny tree

273

# 3.4 Magnitude of ecological divergence and time since separation

Nine levels of splits occurred in the phylogenetic tree. From level 1 to 3, the (0,1) scaled index of divergence (D) tended to increase in terms of the median value, maximum value and interquartile range. From level 3 to 9, the maximum value of D for most cases (except level 8) was approximately 1, whereas the median and the interquartile range were less structured (Figure 4a). There were 10 levels of splits in the morphological tree. The maximum value of D, which was even slightly higher for level 1 (D=0.54) than level 2 (D=0.48), was approximately 1 for the remaining levels. The median tended to increase from level 1 to 7 and then decrease from level 7 to 10. The interquartile range tended to increase from level 1 to 9 (Figure 4b).

4 Discussion.

# 4.1 Climatic data extracted from current spruce locations captures the ecological

# divergence among spruces

In this study, we used climatic data extracted from the current locations of spruce populations to examine the ecological divergence among spruce species at various time scales from approximately 28 Ma to 1.8 Ma. Our results showed significant divergence for the ecological niches among sister groups throughout the phylogenetic tree and the morphological tree, which indicated the overall relevance of the climatic data on spruce ecological divergence at various time scales. However, the magnitude of ecological divergence (as indicated by the divergence index (D)) decreased with the time since the separation of species and became much more specific, i.e., variation of D among the nine environmental variables was larger in the more recent splits than in the basal splits.

删除的内容: s

This finding is likely associated with the incompatibility of the time scale between environmental data and ecological divergence because the environmental data extracted from the current locations tended to be more relevant to the divergence of younger nodes than older nodes. The low ecological divergence observed at the first split in both trees should be an indicator of high ecological niche conservatism (Struwe et al., 2011); however, it is likely related to the strong species interactions that obscure the splits. The higher divergence observed for the younger sister groups or sister species might suggest a strong selective effect of climate on extant spruce species derived from more recent splits, and might also be associated the fact that the fewer species of the sister groups within each node in the more recent splits have relatively less complicated trait compositions and hence weak interactions. Our first hypothesis is largely verified by the findings of our study and those of a previous case study (Struwe et al., 2011).

Exceptions to the above-mentioned trend were observed for a few sister groups or species in the phylogenetic tree. Specifically, within clade-3, significant ecological divergence was not detected for the split (node-29 in Figure 2) between *P. spinulosa* and *P. brachytyla*. These two sister species are distributed in the Circum-Tibetan Plateau and their geographical ranges are adjacent. *P. spinulosa* is distributed in the Mt. Himalaya region and has a narrow range (S Xizang, Bhutan, Nepal and Sikkim), whereas *P. brachytyla* is distributed in the SE to E Tibetan Plateau and has a wide range. These differences suggest that instead of ecological divergence, geographical isolation caused by the deep valleys and high mountain peaks in this area, which act as barriers to gene flow between species, might have played a major role in the speciation of these

删除的内容:, on the one hand,

删除的内容: on the other hand,

删除的内容: However(Struwe et al., 2011). Thus, t

删除的内容:

344 two sister species (Li et al., 2010). In addition, we cannot rule out the possibility that 345 the selected climate parameters and their relatively coarse resolution (approximately 1 346 km) do not adequately describe the climatic determinants of spruce distributions. 347 In mountainous regions, a distance of 1 km may cover an elevation interval of hundreds of meters, introducing several degrees of difference in temperature, which 348 349 may give rise to uncertainties in the detection of subtle variations, such as withinspecies variation, or among elevation-sensitive species. In this case, based on Fig. 1 (a, 350 351 b), instead of the elevation gradient, the geographical distributions of the three 352 phylogenetic clades and the morphological groups (quadrangular leaves versus flattened leaves) are largely determined by horizontal gradients (latitude and longitude). 353 Therefore, although the 1-km resolution climate data should be robust to interpret this 354 355 large-scale pattern, further research that focus on the driving force underlying the 356 variation of within-species or among elevation-sensitive species should use high-357 resolution climate data. 4.2 Temperature features tend to be the main driving factors of the primary 358 divergence of spruce phylogeny 359

	删除的内容: about
	删除的内容: -
	删除的内容: the
	删除的内容: distance
	删除的内容: and therefore
******	删除的内容: e
	删除的内容: some
	删除的内容: context of the
	删除的内容: some
	删除的内容: as we can see from
	删除的内容: Fig.1
Ì	删除的内容: both
	删除的内容: is
	删除的内容: Given this base
	删除的内容: -
	删除的内容: of
	删除的内容: we used in this study
	删除的内容:
	删除的内容: work
	删除的内容: s
	删除的内容:

删除的内容: s well as

Of the 31 splits in the phylogeny tree of *Picea*, the first split is much more important

360

361

362

363

364

365

then warmed during the late Oligocene and early Miocene (Lockwood et al., 2013). This oscillation may provide an explanation for the higher divergence of temperature features. The divergence among the nine environmental variables for the subsequent splits, however, varied according to the sister groups or species.

It is well established that the variations in the historical climate associated with the

advancement and retreat of ice sheets during the late Tertiary and Quaternary periods played an important role in determining plant distributions (Walker, 1986; Hewitt, 2000). In this process, old taxa became extinct or survived in refugia, whereas derived taxa dispersed to new locations and underwent severe selection by climate (Hewitt, 2000; Hampe and Petit, 2010). Therefore, the formation of biogeographical plant patterns is a product of interactions among these processes (Wolf et al., 2001).

In fact, considerable variations in geology and climate have occurred since spruce originated in the late Oligocene. For example, the earliest spruce pollen fossil is from the late Oligocene to the early Miocene in Asia and was found on the Tibetan Plateau (Wu et al., 2007), and spruce pollen has frequently been found in sediments originating from the late Pliocene and the Pleistocene in northern, eastern and southwestern China (Xu et al., 1973; Xu et al., 1980; Shi, 1996) and Taiwan (Tsukada, 1966). A higher proportion of spruce pollen in specific sediments is generally assumed to indicate a cold period, whereas a lower proportion of spruce pollen indicates a warmer period (Xu et al., 1980). The proportion of spruce pollen in the sediments varied substantially with the geological age of the sediments, suggesting that spruce underwent frequent expansion and retreat during glacial cycles. In North America, fossils of Brewer spruce

(*P. breveriana*) have been observed in northeastern Oregon in Miocene deposits that date to more than 15 Ma years ago; however, the present distribution of Brewer spruce is different from the distribution of the fossil locations, indicating that expansion and retreat occurred in the past (Waring et al., 1975). It is difficult to match all the details of paleo-geological or paleo-climatic events to the ecological divergence observed for specific nodes, although our findings offer a quantitative interpretation with respect to the influence of climate on spruce speciation.

# 4.3 Precipitation features tend to be the main driving factors of the primary

# divergence of spruce morphology

The morphological tree in this study was based on spruce taxonomic schemes and highlights the divergence between leaf cross sections in spruce. Although this morphological tree is an artificial scheme, our results indicate that precipitation features were "the primary trigger" of the divergence between quadrangular leaves and flattened leaves among spruce species. A universal pattern was not observed for the climatic variables with respect to the ecological divergence of spruce morphology, which varied according to the specific nodes or splits.

The first split of the basal node of the morphological tree was based on the leaf cross section (i.e., quadrangular vs. flattened); however, each sister group is actually a combination of multiple traits, including the size, shape, color and pubescent/glabrous state of the seed cones, seed scales, bud scales, leaves, leaf apex, and first- and second-year branchlets (Farjón, 1990; Sigurgeirsson and Szmidt, 1993; Fu et al., 1999). The morphological and morphometric traits of spruce species have been demonstrated to

produce strong climatic signals; however, specific traits for different species do not

432

433

434

435

436

437

438

439

440

441

442

443

444

445

446

447

448

449

451

452

453

删除的内容: necessarily

删除的内容: necessarily

<u>always</u> exhibit the same response to specific environmental gradients (Wang et al., 2015; Li et al., 2016). This inconsistency in response is likely due to parallel evolution because morphological similarity among species does not <u>always</u> coincide with the phylogenetic relatedness of species (Went, 1971; Orr, 2005). Accordingly, spruce species with similar morphological characteristics but distant phylogenetic relatedness may differ because of the tradeoff between niche conservatism and ecological divergence. In addition, the composition of traits within a species is also species specific. For example, the shape of the leaf cross section co-varies along with the stomatal line position on the leaf surface, seed scale arrangement and seed scale texture. However, evidence in support of the co-evolution between the leaf cross section (quadrangular (Q) vs. flattened (F)) and seed scale arrangement (closely (C) vs. loosely (L)) has not been observed. Trait combinations such as Q+C, Q+L, F+C and F+L are found in 22, 2, 4 and 5 of the 33 species in *Picea*, respectively (Farjón, 2001). Therefore, without providing additional details, a universal pattern of ecological divergence cannot be predicted for the entire morphological tree of Picea.

4.4 Divergence of morphology and phylogeny among spruce species is affected by

different selective pressures under parallel evolution

Closely related species in a phylogenetic tree tend to be similar in appearance, although 450

this may not be so under parallel evolution (Hoekstra and Price, 2004; Baum et al., 2005;

Orr, 2005), and both cases can be observed in spruce. First, of the three clades in the

phylogenetic tree, most of the spruce species (19 of 22) in clade-1 and clade-2 tended

to have quadrangular leaves, whereas nearly half of the spruce species (6 of 11) in clade3 tended to have flattened leaves. In addition, two North American species, *P. rubens*and *P. mariana*, are sister species in both constructed trees. Accordingly, the
morphological divergence and phylogenetic divergence of these species are subject to
the same selective pressures. Second, cases of parallel evolution are quite obvious. For
example, two Asian species, *P. purpurea* and *P. wilsonii*, are sister species in the
phylogenetic tree but are located in different sections of the morphological tree; this
scenario is also observed for another two North American species, *P. glauca* and *P. engelmannii*. As a result, the morphological and phylogenetic divergences for these
species pairs are subject to different selective pressures, which suggests that the
divergence of morphology and phylogeny among the species in question may or may
not be subject to different selective pressures depending on the process of speciation.

# 4.5 Significance and implications of the findings of this research in relation to

# future climate change

According to the 1997 UNPE standard climate zone classification (Middleton and Thomas, 1997), 8 spruce species are in arid and semi-arid areas, 11 in dry sub-humid areas, and 14 in humid areas. Global climate change predictions have shown that severe and widespread droughts are expected during the next 30-90 years due to either decreased precipitation or increased evaporation, and the significant increases in aridity are expected occur in many subtropical and adjacent humid regions (Dai, 2012; Greve and Seneviratne, 2015). When overlapping the spruce sampling point to the future aridity change, map (data not shown), nearly all spruce species whose original

删除的内容: &

删除的内容: The scenario of global

删除的内容: there would have

删除的内容: in

删除的内容: over land areas resulting from

删除的内容: and/or

删除的内容: do

删除的内容: &

删除的内容: &

删除的内容: C

删除的内容: S

删除的内容: of the

distribution encompasses sub-humid and humid areas is subject to drought stress.

Therefore, our findings suggest that spruces with quadrangular leaves and in clade-1 are expected to expand, whereas those with flattened leaves and in clade-2 and clade-3 are expected to retreat. This should be taken into account in the context of forming strategies in response to future climate change.

# 5 Summary and conclusions

6 Data availability

In summary, the influence of climate on the divergence of the morphology and phylogeny of spruces is mediated by a number of biotic and abiotic factors, such as geographical isolation, niche conservatism and ecological adaptation. A major finding from this study is that temperature and precipitation parameters tend, to be the main driving factors for the primary divergence of spruce phylogeny and morphology, respectively. Our hypotheses are largely verified by the findings of the present study. However, exceptions to the overall pattern cannot be ignored. For example, although most spruce species with quadrangular leaves tend to occur in drier habitats, Taiwan spruce (*P. morrisonicola*) presents quadrangular leaves and is naturally distributed in subtropical areas with abundant rainfall; thus, its present distribution is likely within a refugium from the postglacial period (Tsukada, 1966; Xu et al., 1980). Further work that considers all determinants is required to understand the forces driving ecological divergence among spruce sister groups or species. In addition, our findings shed light on the management issues with respect to spruce distributions under future climate change.

删除的内容: in sss

删除的内容: would

删除的内容: Given this

删除的内容: predicted

删除的内容: while

删除的内容: predicted

删除的内容: ed

删除的内容: y-making

**删除的内容:** of the

523	The relevant data are within the paper and its Supporting Information section.		删除的内容: files
524	7 Author contributions		
525	GHW conceived and designed the experiments. All authors performed the experiment.		
526	GHW and HL analyzed and interpreted the data, and wrote the paper. The authors		
527	declare they have no conflict of interest.		
528	Supporting Information	, and a second	删除的内容: 8
529	Additional supporting information may be found in the online version of this article:		
530	Appendix S1 Index of divergence (D) from the phylogeny-based and morphology-		
531	based SEEVA evaluation of spruce species.		
532	Table S1 Index of divergence (D) from the phylogeny-based SEEVA evaluation of		
533	spruce species worldwide.		
534	Table S2 Index of divergence (D) from the morphology-based SEEVA evaluation of spruce		
535	species worldwide.		
536	9 Acknowledgements		
537	We thank Xing Bai, Lijiang Zhou, Miao Ma, Qinggui Wang, Hongchun Wang, Zhi Ma,		
538	Ziying Chen and Tiancai Chen for providing field assistance. This work was supported		
539	by National Natural Science Foundation of China (41571045), the Chinese National		
540	Basic Research Program (2014CB954201), and the National Natural Science		
541	Foundation of China (30870398).		
542	References		
543	Baum, D. A., Smith, S. D., and Donovan, S. S.: Evolution. The tree-thinking challenge, Science,		
544	310, 979-980, 10.1126/science.1117727, 2005.		

548	Dai, A.: Increasing drought under global warming in observations and models, Nat. Clim. Change
549	3, 52-58, 2012.
550	Darnell, R. M., and Dillon, L. S.: Ecology and the origin of species. Introductory statement, Am
551	Zool., 10, 7-8, 1970.
552	Escudero, M., Valcarcel, V., Vargas, P., and Luceno, M.: Significance of ecological vicariance and
553	long-distance dispersal in the diversification of Carex sect. Spirostachyae (Cyperaceae)
554	Am. J. Bot., 96, 2100-2114, 10.3732/ajb.0900134, 2009.
555	Farjón, A.: Pinaceae: Drawings and Descriptions of the Genera Abies, Cedrus, Pseudolarix
556	Keteleeria, Nothotsuga, Tsuga, Cathaya, Pseudotsuga, Larix and Picea, Cambridg
557	University Press, Konigstein, Germany, 1990.
558	Farjón, A.: World Checklist and Bibliography of Conifers, 2nd ed., Cambridge University Press
559	Cambridge, UK, 2001.
560	Fisher, R. A.: Statistical Methods for Research Workers, 13th ed., Hafner Press, Hafner, NY, 1958
561	Fu, L., Li, N., and Mill, R. R.: Picea, in: Flora of China, edited by: Wu, ZY., and Raven, P. H.
562	Science Press, Beijing, China, 25-32, 1999.
563	GBIF: Global Biodiversity Information Facility: http://doi.org/10.15468/dl.mdqygv10, last access
564	Oct. 13, 2015.
565	Givnish, T. J.: Ecology of plant speciation, Taxon, 59, 1326-1366, 2010.
566	Greve, P. and Seneviratne, S. I.: Assessment of future changes in water availability and aridity
567	Geophys. Res. Lett., 42, 5493-5499, 2015.

Baum, D. A., and Offner, S.: Phylogenies & tree-thinking, Am. Biol. Teach., 70, 222-229, 2008.

删除的内容: Nature Climate

547

568

Hampe, A., and Petit, R. J.: Cryptic forest refugia on the 'Roof of the World', New Phytol., 185, 5-

570	7, 10.2307/25609586, 2010.
571	Hewitt, G.: The genetic legacy of the Quaternary ice ages, Nature, 405, 907-913, 10.1038/35016000,
572	2000.
573	Hijmans, R. J., Cameron, S. E., Parra, J. L., Jones, P. G., and Jarvis, A.: Very high resolution
574	interpolated climate surfaces for global land areas, Int. J. Climatol., 25, 1965-1978,
575	10.1002/joc.1276, 2005.
576	Hoekstra, H. E., and Price, T.: Parallel evolution is in the genes, Science, 303, 1779-1781, DOI
577	10.1126/science.1096413, 2004.
578	Klymiuk, A. A., and Stockey, R. A.: A lower Cretaceous (Valanginian) seed cone provides the
579	earliest fossil record for Picea (Pinaceae), Am. J. Bot., 99, 1069-1082, 10.3732/ajb.1100568,
580	2012.
581	LePage, B. A.: New species of Picea A. Dietrich (Pinaceae) from the middle Eocene of Axel Heiberg
582	Island, Arctic Canada, Bot. J. Linn. Soc., 135, 137-167, DOI 10.1111/j.1095-
583	8339.2001.tb01088.x, 2001.
584	Li, H., Wang, G., Zhang, Y., and Zhang, W.: Morphometric traits capture the climatically driven
585	species turnover of 10 spruce taxa across China, Ecol. Evol., 6, 1203-1213,
586	10.1002/ece3.1971, 2016.
587	Li, Y., Stocks, M., Hemmila, S., Kallman, T., Zhu, H., Zhou, Y., Chen, J., Liu, J., and Lascoux, M.:
588	Demographic histories of four spruce (Picea) species of the Qinghai-Tibetan Plateau and
589	neighboring areas inferred from multiple nuclear loci, Mol. Biol. Evol., 27, 1001-1014,
590	10.1093/molbev/msp301, 2010.

591 Liu, T.: A new proposal for the classification of the genus Picea, Acta Phytotaxonomica et

592	Geobotanica 33, 227-245, 1982.
593	Lockwood, J. D., Aleksic, J. M., Zou, J., Wang, J., Liu, J., and Renner, S. S.: A new phylogeny for
594	the genus Picea from plastid, mitochondrial, and nuclear sequences, Mol. Phylogenet. Evol.,
595	69, 717-727, 10.1016/j.ympev.2013.07.004, 2013.
596	Mayr, E.: Ecological factors in speciation, Evolution, 1, 263-288, Doi 10.2307/2405327, 1947.
597	Middleton, N. and Thomas, D. S. G.; World atlas of desertification, Geogr. J., 160, 210, 1997.
598	Orr, H. A.: The probability of parallel evolution, Evolution, 59, 216-220, 10.2307/3449009, 2005.
599	Peterson, A. T., Soberón, J., and Sanchez-Cordero, V. V.: Conservatism of ecological niches in
600	evolutionary time, Science, 285, 1265-1267, 1999.
601	Ran, J. H., Wei, X. X., and Wang, X. Q.: Molecular phylogeny and biogeography of Picea (Pinaceae):
602	Implications for phylogeographical studies using cytoplasmic haplotypes, Mol. Phylogenet.
603	Evol., 41, 405-419, 10.1016/j.ympev.2006.05.039, 2006.
604	Schemske, D. W.: Adaptation and the origin of species, Am. Nat., 176 Suppl 1, 4-25,
605	10.1086/657060, 2010.
606	Schluter, D., Clifford, E. A., Nemethy, M., and McKinnon, J. S.: Parallel evolution and inheritance
607	of quantitative traits, Am. Nat., 163, 9-22, 10.1086/383621, 2004.
608	Shi, N.: Development of spruce and fir in north China during the Pliocnen and the early Pleistonene:
609	palaeoclimatic implications, Quaternary Sci., 16, 319-328, 1996.
610	Sigurgeirsson, A., and Szmidt, A. E.: Phylogenetic and biogeographic implications of chloroplast
611	DNA variation in Picea, Nord. J. Bot., 13, 233-246, DOI 10.1111/j.1756-
612	1051.1993.tb00043.x, 1993.

删除的内容:, and Programme, U. N. E.

删除的内容: The

删除的内容: aphical

删除的内容: Journal

613 Spribille, T., and Chytry, M.: Vegetation surveys in the circumboreal coniferous forests: a review,

619	Folia Geobot., 37, 365-382, Doi 10.1007/Bf02803253, 2002.			
620	Struwe, L., Smouse, P. E., Heiberg, E., Haag, S., and Lathrop, R. G.: Spatial evolutionary and			
621	ecological vicariance analysis (SEEVA), a novel approach to biogeography and speciation			
622	research, with an example from Brazilian Gentianaceae, J. Biogeogr., 38, 1841-1854,			
623	10.1111/j.1365-2699.2011.02532.x, 2011.			
624	Taylor, R. J.: Picea, in: Flora of North America North of Mexico; volume 2: pteridophytes and	**	<b>削除的内容:</b>	
625	gymnosperms, edited by: Flora of North America Editorial Committee, Oxford University	#	<b>1除的内容:</b> 369 – 373,	
626	Press, New York, 369-373, 1993.		<b> 除的内容:</b> , NY	
627	Tseng, Y. S.: Studies on the vegetation ecology of Salihsianhsi watershed in central Taiwan II:			
628	studies on the forest dynamics and population structure of Taiwan spruce forest, Master's			
629	Thesis, Institute of Forestry, National Taiwan University, Taipei, 1991.			
630	Tsukada, M.: Late pleistocene vegetation and climate in taiwan (formosa), Proc. Natl. Acad. Sci.			
631	U.S.A., 55, 543-548, 10.2307/57266, 1966.			
632	Walker, D.: Late Pleistocene early Holocene vegetational and climatic changes in Yunnan Province,			
633	southwest China, J. Biogeogr., 13, 477-486, Doi 10.2307/2844968, 1986.			
634	Wang, G. H., Liu, J. L., and Meng, T. T.: Leaf trait variation captures climate differences but differs			
635	with species irrespective of functional group, J. Plant Ecol., 8, 61-69, 10.1093/jpe/rtu009,			
636	2015.			
637	Waring, R. H., Emmingham, W. H., and Running, S. W.: Environmental limits of an endemic spruce,			
638	Picea-breweriana, Can. J. Bot., 53, 1599-1613, 1975.			
639	Webb, C. O., Ackerly, D. D., McPeek, M. A., and Donoghue, M. J.: Phylogenies and community			
640	ecology, Annu. Rev. Ecol. Syst., 33, 475-505, 10.1146/annurev.ecolysis.33.010802.150448,			

644	2002
044	2002

- 645 Went, F. W.: Parallel evolution, Taxon, 20, 197, 10.2307/1218877, 1971.
- 646 Wiens, J. J.: Speciation and ecology revisited: phylogenetic niche conservatism and the origin of
- species, Evolution, 58, 193-197, 10.2307/3449309, 2004.
- 648 Wiens, J. J., and Graham, C. H.: Niche conservatism: integrating evolution, ecology, and
- conservation biology, Annu. Rev. Ecol. Evol. S., 36, 519-539,
- 650 10.1146/annurev.ecolsys.36.102803.095431, 2005.
- 651 Wiley, E. O.: Vicariance biogeography, Annu. Rev. Ecol. Syst., 19, 513-542, 10.2307/2097164, 1988.
- Wolf, P. G., Schneider, H., and Ranker, T. A.: Geographic distributions of homosporous ferns: does
- dispersal obscure evidence of vicariance?, J. Biogeogr., 28, 263-270, 10.2307/2656102,
- 654 2001.
- 655 Wu, Z., Wu, Z., Hu, D., Ye, P., and Zhou, C.: Geological evidences for the Tibetan Plateau uplifted
- in late Oligocene, Acta Geol. Sin.-Engl., 81, 577-587, 2007.
- 657 Xu, R., Tao, J. R., and Sun, X. J.: On the discovery of a Quercus semicarpifolia Bedin Mount Shisha
- Pangma and its significance in botany and geology, J. Integr. Plant Biol., 15, 103-119, 1973.
- 659 Xu, R., Kong, Z. C., and Du, N. Q.: Plant assemblages of Picea and Abies in the Pleistocene and
- implications for Quarternary study Quaternary Sci., 5, 48-56, 1980.
- 661 Yang, G. Z., Chen, Y. F., Zhao, W. C., Chen, X. Y., Wu, L. T., Zhao, G. R., and Lu, Z. F.: Plant
- Resource Investigation in Nantzuhsien Creek Watershed in Yushan of National Park,
- Yushan of National Park, Nantou, Taiwan, 2002.

**Table 1.** Factor analysis showing the eigenvalues, variance percentages, cumulative percentages and correlations of 19 climatic variables with the five components.

Climatic variables in bold were selected for further analysis.

Climatic variables			Comp	onents		
		1	2	3	4	5
	Eigenvalues	8.27	3.60	2.51	2.26	1.24
	Variance %	43.52	18.93	13.21	11.89	6.51
	Cumulative %	43.52	62.46	75.67	87.55	94.06
Annual Mean Temperature (AMT)		0.803	0.222	0.082	0.513	-0.152
Mean Diurnal Range (MDR)		-0.118	-0.155	-0.686	0.476	0.31
Isothermality (I)		0.687	0.283	-0.45	0.158	0.307
Temperature Seasonality_(TS)		-0.928	-0.237	-0.099	0.204	-0.12
Max Temperature of Warmest Month	(MTWM)	0.037	-0.155	-0.129	0.968	0.01
Min Temperature of Coldest Month (M	ИТСМ)	0.931	0.216	0.257	0.086	0.006
Annual Temperature Range (ATR)		-0.854	-0.267	-0.294	0.329	-0.001
Mean Temperature of Wettest Quarter_(N	MTWeQ)	-0.123	0.091	-0.066	0.48	-0.788
Mean Temperature of Driest Quarter_(M	ΓDQ)	0.841	0.093	0.138	0.116	0.408
Mean Temperature of Warmest Quarter_(	MTWaQ)	0.14	0.02	0.04	0.918	-0.294
Mean Temperature of Coldest Quarter_(N	MTCQ)	0.946	0.24	0.108	0.179	0.007
Annual Precipitation (AP)		0.306	0.856	0.365	-0.041	0.178
Precipitation of Wettest Month (PWM	)	0.288	0.942	-0.006	-0.033	0.109
Precipitation of Driest Month (PDM)		0.147	0.255	0.911	0.008	0.087
Precipitation Seasonality (PS)		-0.109	0.255	-0.887	-0.006	-0.131
Precipitation of Wettest Quarter (PWeQ)		0.297	0.937	0.026	-0.038	0.134
Precipitation of Driest Quarter (PDQ)		0.175	0.302	0.894	-0.003	0.152
Precipitation of Warmest Quarter (PWaQ	))	0.144	0.888	0.086	-0.057	-0.313
Precipitation of Coldest Quarter (PCQ	2)	0.323	0.402	0.418	-0.016	0.652

	删除的内容: Temperature
	删除的内容: Annual
Ì	删除的内容: TAR
Ì	带格式表格

Table 2. Mean comparisons of the elevation and 8 climatic variables (mean±SD, 删除的内容:
abbreviations are the same as in Table 1) between sister groups at the first two split
levels of both the phylogeny tree and the morphology tree. Mean±SD values marked
删除的内容:
删除的内容:

indicates a non-significant difference (*P*>0.05).

with different letters indicate a significant difference at P<0.05, and the same letter

	N	Elevation (m)	AMT (℃)	MTWM (°C)	MTCM (℃)	<u>ATR</u> (℃)	AP (mm)	PWM (mm)	PDM (mm)	PCQ (mm)
Phyl	ogeny N	odes								
Siste	r Group	s: node-2 (clade-1	) vs. node-14 (	(clade-2 + clade-	3)					
2	1568	964±750°	3.2±4.2 a	19.6±3.7 a	-12.6±8.6 a	32.1±9.5 a	845.8±416.9°	117.1±52.3 a	38.0±25.7°	158.9±124.2°
14	1820	1721±1150 b	$3.8{\pm}5.0^{\rm \ b}$	21.8±3.9 b	-13.9±8.8 <sup>b</sup>	35.7±8.8 <sup>b</sup>	910.7±727.6 b	143.6±119.0 b	$26.9{\pm}27.8^{b}$	186.5±209.3 b
Siste	r Group	s: node-15 (clade-	2) vs. node-22	2 (clade-3)						
15	1100	1176±906 a	2.5±5.0°	22.5±3.6 a	-16.6±8.2 a	39.1±7.3 a	784.1±442.6°	106.3±61.6 a	35.7±27.7°	190.7±180.0°
22	720	2554±971 b	5.9±4.3 <sup>b</sup>	20.6±4.0 <sup>b</sup>	-9.9±8.1 <sup>b</sup>	30.6±8.4 <sup>b</sup>	1104.0±989.0 <sup>b</sup>	200.8±157.0 b	13.5±21.8 b	180.0±247.4°
Siste	r Group	s: node-3 vs. node	-11 (two sister	r groups within c	lade-2)					
3	1502	951±755°	3.0±4.2 a	19.4±3.6 a	-12.8±8.6 a	32.2±9.7 °	834.5±411.2 °	116.2±51.3 a	37.4±25.8 a	157.2±126.0°
11	66	1275±542 b	7.1±2.8 b	22.9±2.6 <sup>b</sup>	-7.5±3.7 b	30.4±2.8 b	1101.8±464.7 b	137.8±70.0°	52.3±16.7 b	196.3±63.3 b
Mor	phology	Nodes								
Siste	r Group	s: node-2 vs. node	-25 (i.e., quad	rangular leaf gro	up vs. flattened le	eaf group)				
2	2857	1191±915°	3.1±4.7°	20.8±4.0 a	-14.0±8.8 a	34.8±9.7 °	849.4±624.2 a	120.0±95.2 a	35.3±27.2°	163.8±146.4°
25	531	2337±1222 b	5.8±3.7 <sup>b</sup>	20.7±3.7 a	-9.3±6.6 b	29.9±5.5 b	1048.5±452.1 b	192.2±67.9 b	14.5±21.0 b	226.8±279.7 b
Siste	r Group	s: node-3 vs. node	-24 (i.e., withi	in quadrangular l	eaf group: seed so	cale closely arrange	ed group vs. loosely	arranged group)		
3	2530	1059±850°	3.0±4.8°	20.5±3.9 a	-14.3±9.2°	34.8±10.2 a	864.7±646.3 a	121.6±97.8 a	36.6±28.4 a	155.8±135.1 °
24	327	2219±729 b	3.7±3.7 b	22.8±4.0 b	-12.1±4.8 b	34.8±4.2 a	730.9±396.0 b	107.7±70.6 b	25.7±10.8 b	225.9±204.9 b
Siste	r Group	s: node-26 vs. nod	le-29 (i.e., witl	hin flattened leaf	group: seed scale	closely arranged g	group vs. loosely arr	anged group)		
26	283	2806±1301°	4.6±4.1°	19.0±3.3 a	-12.4±7.3 a	31.4±6.7 °	996.1±564.2 a	190.1±77.4°	15.1±23.7°	125.5±252.6°
29	248	1802±854 b	7.2±2.5 b	22.5±3.2 <sup>b</sup>	-5.7±3.0 b	28.2±2.9 b	1108.4±261.7 b	194.6±55.3 a	13.8±17.4°	342.4±264.2 b
4	2118	1124±890°	3.0±4.9 a	20.0±3.9 a	-13.8±9.5 °	33.8±10.5 a	853.8±682.2 a	124.6±105.6 a	33.3±26.2 a	149.0±139.0°
21	412	724±487 b	3.2±4.3 a	23.2±2.9 b	-17.0±6.9 b	40.1±6.0 b	921.0±412.3 a	106.2±33.1 b	53.2±33.0 b	190.8±105.7 b

# Figure legends:

682

684

686

688

690

692

694

Figure 1. Sites were sampled across the entire range of spruces worldwide. Sites

marked with different symbols represent three phylogenetically distinct clades (a), and

two morphological groups (b), Elevation gradients are indicated by colored fields.

Figure 2. Divergence indices (scales range from 0-1) shown as histograms for elevation

and for the 8 climatic variables for each node of the phylogeny of *Picea* worldwide.

\*Indicates a significant difference in ecological features after Bonferroni correction

689 (*P*<0.0016).

**Figure 3.** Divergence indices (scales range from 0-1) shown as histograms for elevation

and for the 8 climatic variables for each node of the morphology of *Picea* worldwide.

\*Indicates a significant difference of ecological features after Bonferroni correction

693 (*P*<0.0016).

Figure 4. Box plots showing the index of divergence at each of the splitting levels in

the phylogenetic tree (a) and the morphological tree (b) of spruce species worldwide.

The central box in each box plot indicates the interquartile range and median, whereas

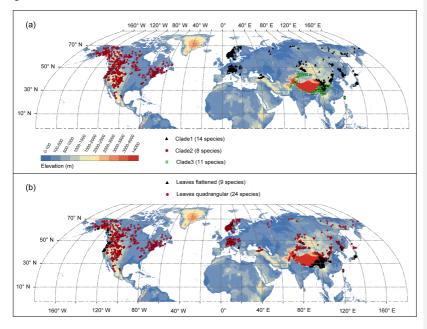
the whiskers show the 10th and 90th percentiles. Mean values marked with different

letters indicate a significant difference at P < 0.01.

删除的内容:

删除的内容:, respectively

# 701 Figure 1



批注 [E1]: Add a hyphen between 'Clade'.

批注 [E2]: Changed "temperature annual range" to "annual temperature range".

