

Potential distribution of *Monotropa uniflora* as a surrogate for range of Monotropeoideae (Ericaceae) in South Asia

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Abstract. Pradhan P. 2015. Potential distribution of *Monotropa uniflora* as a surrogate for range of Monotropeoideae (Ericaceae) in South Asia. *Biodiversitas* 16: 109-115. Monotropeoideae is a mycoheterotrophic subfamily of Ericaceae. Its members are highly specific to a particular fungal family, which has attributed to the rarity and limited distribution of Monotropeoideae. In the past two decades, there are considerable developments in understanding their biology and biogeography, among which, the distribution of *Monotropa uniflora* L. and *M. hypopitys* L. has been extensively studied. In this contribution, Ecological Niche Modeling of *M. uniflora* has been conducted to test its earlier proposed distribution in South Asia, to test the spatial scale of the said proposal, to test its potential distribution as a surrogate for range of Monotropeoideae in South Asia and to prioritize conservation areas for *M. uniflora* in the region. The model was built with five occurrence details of the rare plant *M. uniflora* in Western and Eastern Himalaya, in relation to 19 bioclimatic explanatory variables, performed in MaxEnt. The results show the good performance of the model with the training AUC of 0.994. 1,50,316 square Km. of suitable areas have been predicted for the growth of *M. uniflora* (IHS 0.5) in South Asia, many areas of which is in line with earlier distributional reports. The bioclimatic variables are able to predict and suitably justify the spatial distribution of *M. uniflora*. The predicted range of the species could be established for potential distribution of other Asian Monotropoids like *Monotropastrum* and *Cheilothecca*.

Key words: Ecological Niche Modeling, Himalaya, MaxEnt, Mycoheterotrophy, Russulaceae

Abbreviations: IHS = Index of Habitat Suitability; Bio = Bioclimatic variable

INTRODUCTION

Mycoheterotrophy includes an obligatory reliance of achlorophyllous and non-photosynthetic plants upon specialized mycorrhizal associates for carbon influx (Klooster and Culley 2009). Monotropeoideae is a mycoheterotrophic subfamily of Ericaceae, consisting of 10 genera and 15 species (Wallace 1975). Leake (1994) has reported 260 species of Mycoheterotrophic plants as endemic to Palaeotropics, extending from India in the West to Papua-New Guinea and Japan in the East. Endemic taxa of Monotropeoideae have been reported from Western North America (seven species) and Asia (two *Monotropastrum* species and two *Cheilothecca* species and a variety) (Tsukaya et al. 2008), preferring to grow mostly in shady old growth forests (Min et al. 2012). Their endemism and narrow geographic range have been linked to limited distribution of their taxa specific mycorrhizal fungal partners (Kruckeberg and Rabinowitz 1985; Bidartondo and Bruns 2001), and such trophic structure has left the subfamily with very isolated and rare taxa (Klooster and Culley 2009). *Monotropa uniflora* L. and *M. hypopitys* L. though distantly related but are the only species within Monotropeoideae which are distributed across Neotropics and Palaeotropics (Leake 1994). Although Leake (1994) has indicated broad range of genus *Monotropa* in Indian subcontinent, *M. uniflora* is considered regionally rare in

Indian states of West Bengal (WBBB 2012; FRLHT 2015) and Meghalaya (Mir et al. 2014).

Suitability of climate and presence of dense and shaded forest habitat has been attributed for the evolution of heterotrophy in paleotropics and neotropics (Leake 1994). Similarly, Ecological Niche/climate modeling has been proposed to act as a successful marker to infer the phylogeographic and demographic histories of mycoheterotrophic plants by Taylor et al. (2013). In this regard, paleo-distribution modeling of *M. hypopitys* in North America has been successfully conducted in MaxEnt using bioclimatic variables (Beatty and Provan 2011), however no study has been focused on Monotropeoideae as a whole or any species therein in South Asia regarding their distribution in bioclimatic envelope. For planning suitable conservation action for Monotropeoideae in this region, prime necessity is to prioritize areas for conservation having high suitability of the ecological niche of the taxa.

In this contribution, Ecological Niche Modeling of *M. uniflora* has been conducted to test earlier proposed distribution of the species in South Asia, to test the spatial scale of the said proposal, to test potential distribution of the species as a surrogate for range of Monotropeoideae in South Asia and to prioritize conservation areas for *M. uniflora* in the region.

MATERIALS AND METHODS

Occurrence records and site characteristics

Occurrence records were obtained with the help of Garmin Etrex GPS machine for the locations of Jorepokhari (Figure 1); data from Rachela were obtained from Divisional Forest Office (Research Division), Darjeeling; occurrences in Phedkhal, Jakholi and Mandal of Uttarakhand were derived from Semwal et al. (2014) and in communication with Dr. Semwal (Figure 2). All the occurrence sites have montane topography. Vegetation wise, Rachela and Phedkhal have Oak dominated vegetation, Jakholi has mixed woodlands, Mandal has mixed forest with Oak, Birch and Cedrus, while Jorepokhari has mixed woodlands of *Cryptomeria japonica*, Oak and Bamboos. The geographical features of the occurrence sites are shown in Table 1.



Figure 2. *Monotropa uniflora* plants observed in Jorepokhari, Darjeeling District, West Bengal, India

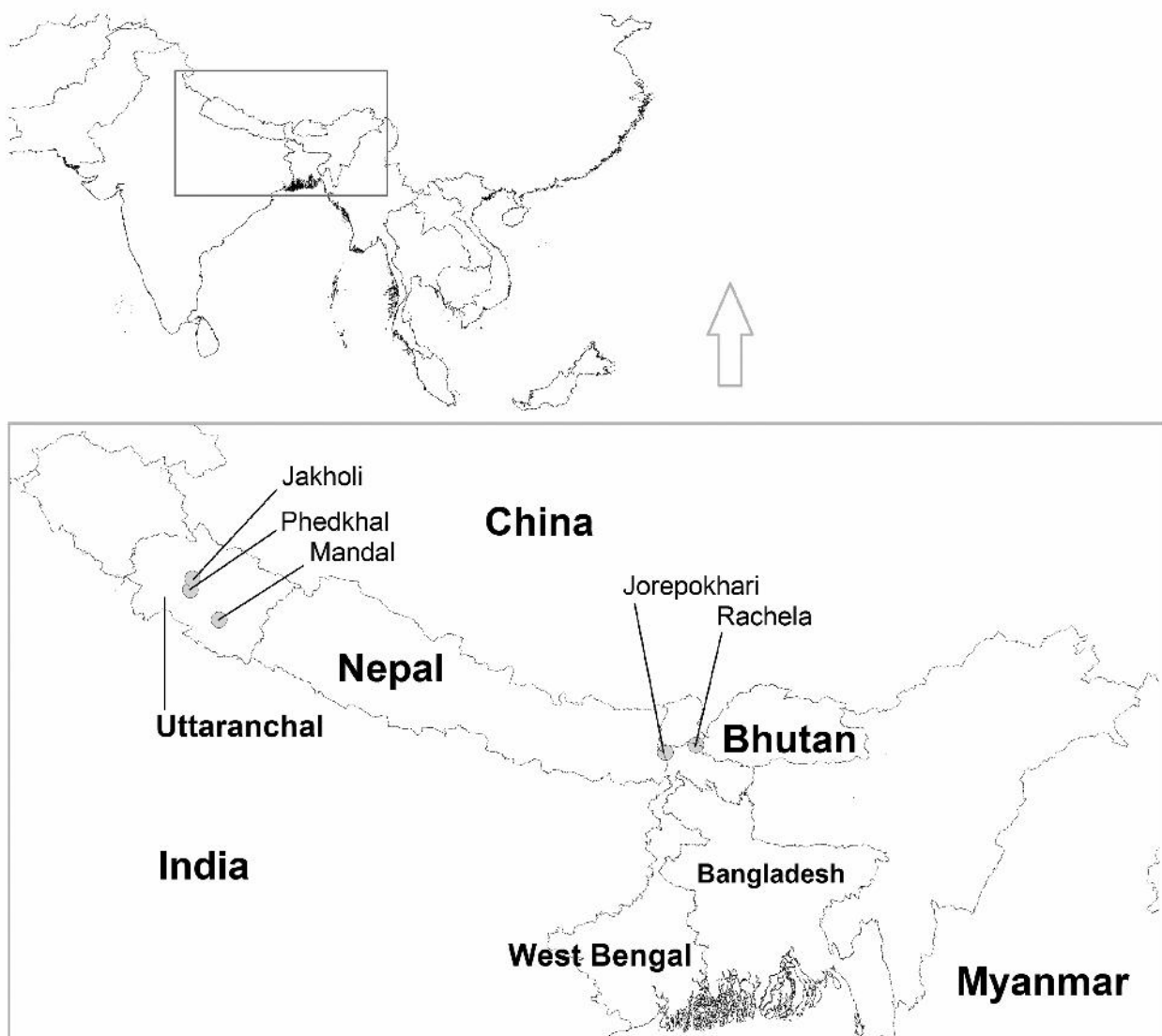


Figure 1. Occurrence records utilized for modelling potential distribution of *Monotropa uniflora*

Table 1. Geographical and climatic features of the occurrence sites; Tmin=mean monthly minimum temperature, Tmax=mean monthly maximum temperature, PPT=annual precipitation.

Site	Longitude	Latitude	Altitude (m)	Tmin avg (°C)	Tmax avg (°C)	PPT (mm)	Aspect
Jorepokhari	88.1490	26.9877	2239	9.3	16.1	2268	351.92 (N)
Rachela	88.7414	27.1378	3124	1.7	12.8	1068	312.45 (NW)
Phedkhal	78.8704	30.1682	1842	10.3	19.4	1634	308.95 (NW)
Jakholi	78.9080	30.3862	1706	10.9	20.4	1467	117.88 (SE)
Mandal	79.4331	29.5789	1621	9.6	19.5	1784	56.97 (NE)

Table 2. Range of Index of Habitat Suitability (IHS) along with predicted area and the respective percentage of the total suitable area.

IHS	Predicted area km ²	% of total suitable area
0.5 - 0.599	1,50,316	35.06
0.6 - 0.699	98,045	22.87
0.7 - 0.799	76,717	17.89
0.8 - 0.899	63,697	14.86
0.9 - 1	39,968	9.32

Modeling species distribution - Modeling method

MaxEnt program version 3.3.3 k (Phillips et al. 2006; Phillips and Dudik 2008) is a maximum entropy based general-purpose machine learning method and it was used for modeling species distribution in geographic space and creation of habitat suitability maps. Entropy in the context of probability theory and statistics measures the amount of information that is contained in a random variable or unknown quantity. MaxEnt uses the basic set of information to model the distribution of a species, i.e., a set of samples (species presence) available from a geographical region, which is linked to a set of explanatory variables (e.g. climatic).

Explanatory variables

Climatic variables of monthly precipitation and monthly mean, minimum and maximum temperature at a spatial resolution of 30 arc seconds (~1 × 1 km resolution) were derived from WORLDCLIM (Hijmans et al. 2005) (tile 18, 19, 28 and 29) which includes interpolation of climatic records from global network of 4000 climate stations, with time series of 1950–2000. Current investigation doesn't incorporate climatic information of tile 110 and 210 of WORLDCLIM, hence geographic space of East Asian countries like Taiwan, Japan etc. are excluded for potential species distribution. Using DIVA-GIS version 7.5 (Hijmans et al. 2001), tile data were converted into 19 bioclimatic variables i.e., annual mean temperature [Bio1], mean monthly temperature range [Bio2], isothermality [Bio3], temperature seasonality [Bio4], max temperature of warmest month [Bio5], min temperature of coldest month [Bio6], temperature annual range [Bio7], mean temperature of wettest quarter [Bio8], mean temperature of driest quarter [Bio9], mean temperature of warmest quarter [Bio10], mean temperature

of coldest quarter [Bio11], annual precipitation [Bio12], precipitation of wettest month [Bio13], precipitation of driest month [Bio14], precipitation seasonality [Bio15], precipitation of wettest quarter [Bio16], precipitation of driest quarter [Bio17], precipitation of warmest quarter [Bio18], precipitation of coldest quarter [Bio19] (Busby 1986; Nix 1986; Hijmans et al. 2005) in ESRI ASC format for use in MaxEnt program. These bioclimatic variables express spatial variation in annual means, seasonality and extreme or limiting climatic factors and represent biologically meaningful parameters for characterizing species distributions (Saatchi et al. 2008) hence they were used as explanatory variables.

Model building and evaluation

Presence-only data were used for model building. However, in order to evaluate models on the basis of error rates, absence details are needed. To overcome this, 25,000 random points throughout the study area were assumed as absence 'pseudo-absence' (Zaniewski et al. 2002). Linear regularization feature was used for model building; default value of 1 for regularization which gives consistent AUC peaks (Phillips et al., 2004) was used; prevalence or the probability of presence was taken to be 0.5.

Threshold-independent analysis

In the threshold-independent analysis, model performance/strength (the power to discriminate between sites where a species is present, versus those where it is absent) was evaluated using the Area under the Curve (AUC) of the Receiver Operating Characteristic (ROC) curve (Fielding and Bell 1997; Elith and Burgman 2002). The AUC statistic ranges from 0 to 1, where a score of 0.5 implies discrimination that is no better than random, and a score of 1 indicates perfect discrimination (Fielding and Bell 1997; Pearce and Ferrier 2000).

Explanatory variable importance

The importance of each bioclimatic factors explaining the distribution of species was determined by Jackknife analysis of the average gain with training and test data. 500 iterations were conducted for training algorithm, the increase or decrease in regularized gain was added or subtracted, respectively, to the input of the corresponding variable, giving a heuristic estimate of bioclimatic variable contribution for the model (Phillips et al. 2006).

Model output and Mapping

Logistic format was selected which categorizes estimates of probability of occurrence or Index of Habitat Suitability (IHS) within 0-1 (Anderson et al. 2003; Baldwin 2009). In DIVA-GIS, the logistic output grid was reclassified and exported to raster. IHS regions < 0.5 were discarded as random prediction, > 0.5 IHS were treated as suitable areas of species distribution, 0.7 IHS were treated as the core areas of species distribution (Kuemmerle et al. 2010), and prioritized areas for conservation were taken for IHS 0.9 .

RESULTS AND DISCUSSION

The potential distribution of *Monotropa uniflora* in South Asia have been found to be in the longitudinal range of 68.94-117.8 decimal degrees and latitudinal range of 6.76-36.81 decimal degrees, covering 1,50,316 square Km

(Table 2). Suitable areas for the growth of the species (IHS 0.5) have been found in the countries of Tajikistan, Afghanistan, Pakistan, India, Nepal, China, Bangladesh, Myanmar, Srilanka, Laos and Vietnam. Whereas, conservation areas that could be prioritized (IHS 0.9) have been found in India, Nepal, China, Afghanistan, Sri Lanka, Myanmar and Vietnam covering are of 39,968 square Km (Figure 3).

The model performed well with the training AUC value above 0.9 (0.994) with the training omission rate of zero for evaluation localities. The jackknife test of variable importance showed the environmental variable with the highest gain when used in isolation (which has the most useful information by itself) to be temperature annual range [Bio7] (maximum temperature of the warmest month - minimum temperature of the coldest month). The environmental variable that decreases the gain the most when it is omitted is precipitation of driest quarter [Bio17], which therefore appears to have the most information that isn't present in the other variables.

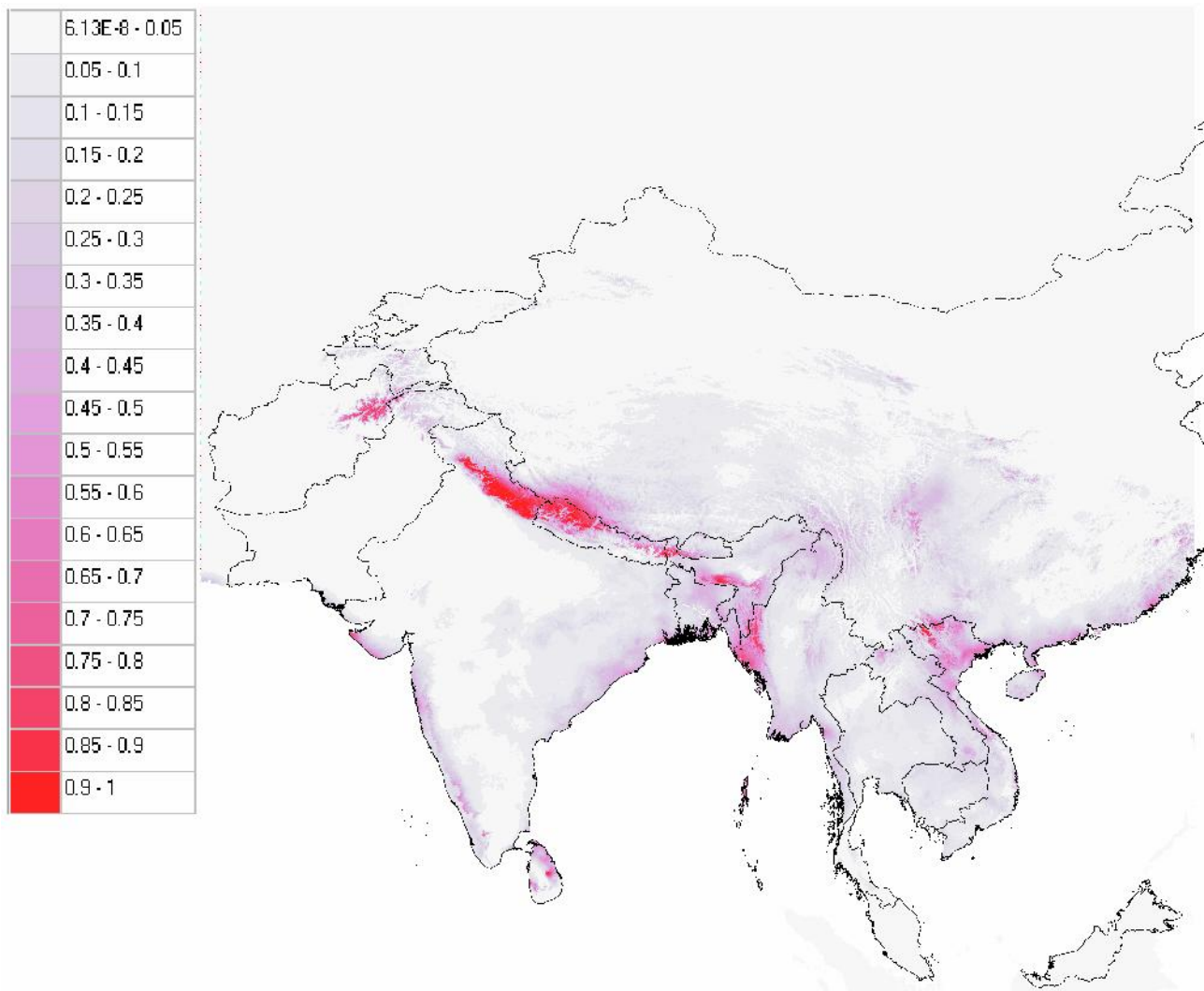


Figure 3. Potential distribution of *Monotropa uniflora* in South Asia with graded Index of Habitat Suitability; warmer colour are indicative of more suitable areas.

Potential conservation areas for *M. uniflora* (IHS 0.9) in India are predicted in Eastern Kathua in the state of Jammu and Kashmir; Charua, Chamba, Dalhousie, Kangra, Brahmaur, Dharmshala, Palampur, Baijnath, Kullu, Mandi, Thunag, Chachyot, Karsog, Banjar, Nermand, Ani, Rampur, Rohru, Kotkhai, Theog, Rajgarh, Chaupal, Renuka, Shilla, Jubbal, Paonta in the state of Himachal Pradesh; Chakrata, Purola, Rajgarhi, Dunda, Tehri Garhwal, Dehradun, Northern Lansdown, Pauri, Pauri Garhwal, Devprayag, Narendranagar, Chamoli, Karnaprayag, Ranikhet, Naini Tal, Almora, Champawat, Bageshwar, Southern Joshimath in the state of Uttaranchal; Darjeeling, Kurseong and Kalimpong subdivisions of Darjeeling District in the state of West Bengal; South Eastern Nongstoin area of West Khasi Hills, South Eastern area of Shillong in the state of Meghalaya.

IHS 0.9 have been predicted in the Kuran Wa Munjan and Wakhan areas of Afghanistan; Chitral of N.W.F.P. in Pakistan; Central and Northern Matupi, Southern Thlangtang areas of Myanmar; Baitadi, Dadeldhura, Doti, Western Darchula, Bajhang, Bajura, Kalikot, Northern Jajarkot, South Eastern Jumla, Mugu, South Eastern Humla, Western Dolpa, Northern Myagdi, Northern Rapti, Southern Manang, Northern Kaski, Eastern Ilam of Nepal; Elahera, Thamankadua, Dimbulagala of Sri Lanka; Eastern Lai Chau and Western Lao Cai areas of Vietnam.

Suitable areas (IHS 0.5) in China are predicted mostly in South-Central and South-Western China with IHS ranging upto 0.80 in Yunnan, 0.81 in Sichuan and 0.94 in Xizang provinces. Wenshan, Pingbiang Miao, Hekou Yao, Maguan, Malipo, Jinping Yao, Miao and Dai areas of Yunnan province are predicted in continuum of the potential areas from Northern Vietnam; Zanda, Burang and Zhongba area of Xizang province are in continuum of Western and Central Himalaya; Kangding, Luding, Hongya, Hanyuan, Meishan, Ebian Yi, Meigu, Mabian Yi, Ganluo, Jinkouhe areas of Sichuan have been predicted as potential habitats in central China.

Not only is the present prediction takes into account for high altitude habitats, but also coastal vegetation with IHS ranging upto 0.82 in Sagar Island, West Bengal, India. Possibility of suitable habitats of the taxa in coastal areas (IHS 0.5) which await prioritized inventory are presently predicted along the coast of Western India, Western Ghats, Bay of Bengal, Gujrat in India; Maungtaw, Buthidaung, Sitwe, Chaungzon, Mudon, Paung, Moulmein areas of Myanmar, Patuakhali, Noakhali, Cox's Bazar, Bandarban areas of Bangladesh; Thua Thien - Hue, Nghe An, Thai Binh, Hai Phong, Quang Ninh area of Vietnam; Hainan, Haikou, Zhanjiang, Maoming, Yangjiang, Jiangmen, Zhuhai, Zhangzhou areas of China.

Genus *Monotropa* was represented by Leake (1994) to cover Indian, Nepalese and Chinese part of Western, Central, Eastern Himalaya and coastal Indian Subcontinent including other parts of India and Bangladesh and Japan in the east as well. Some of the Monotropeidae reported from Asia include *Cheilothea malayana* from Japan (Matsuda and Yamada 2003); *M. uniflora* L. from damp deciduous or mixed forests of Anhui, Gansu, Guizhou,

Hubei, Jiangxi, Qinghai, Shaanxi, Shanxi, Sichuan, Xizang, Yunnan, Zhejiang regions of China (100-1500 m), Bangladesh, Bhutan, India, Korea, Myanmar, Nepal, Sikkim (FOC 2014; Min et al. 2012); Japan (Bidartondo and Bruns 2001); Murree hills in Pakistan (33.9043° N, 73.3942° E) (FOP 2014); *Monotropa hypopitys* L. from India (Barik et al. 2009), Japan (Bidartondo and Bruns 2001); *Monotropastrum humile* (D. Don) Hara from Himalaya (East Asia) to Japan (Kitamura et al. 1975; Bidartondo and Bruns 2001), Yunnan, China. (Min et al. 2012), *M. humile* var. *humile* from Fengshan, Chi-Tou Region of Taiwan, Mt. Mirokusan of Korea, Shiouपुरi, Kathmandu, Nepal (Tsukaya et al. 2008); *Monotropastrum sciaphilum* from Yunnan, China (Min et al. 2012); *Monotropastrum macrocarpum* from Eastern Himalaya (Maheshwari 1969) etc. The potential distribution of *M. uniflora* is in line with the reported distribution of other Monotropoids. Large tracts of Central and Western Himalayas are predicted to be bioclimatically suitable for *M. uniflora*, with disjunct distribution in various countries reaffirming previous reports. However, many areas viz. Zhejiang, Fujian, Guangdong and Hainan of China are predicted in addition to earlier reports (E florae 2014a). Previously, members of Monotropeidae were reported to grow in coastal dunes in the Pacific North West USA with prostrate *Arctostaphylos* mats (Leake 1994). Current prediction of *M. uniflora* in many coastal areas of South Asia therefore necessitates further insight into such habitat of the species. Though temperature annual range have singly important role to play in *M. uniflora* distribution, precipitation of driest quarter may be important in density dependent feedback of fungal biomass in the substratum of mycorrhizal partner of *M. uniflora* (Krivtsov et al. 2006).

Monotropastrum humile and *Cheilothea malayana* which are distributed in South East Asia from the Himalaya to Japan (Wallace 1975; Kitamura et al. 1975) are similar to *M. uniflora* regarding specialization to fungal family Russulaceae (Young et al. 2002; Matsuda and Yamada 2003). In fact, *M. humile* has been shown to be phylogenetically close to *M. uniflora*, therefore current potential distribution of *M. uniflora* could be functionally predictive for the said species of Monotropeidae. However, currently described potential distribution should be carefully applied to derive cross-continent inference for the species, as collections of *M. uniflora* from Asia, North America, and Central America are indicated to be molecularly diverged and phylogenetically distinct (Bidartondo and Bruns 2001; Neyland and Hennigan 2004).

Monotropa uniflora has been reported in association with *Lithocarpus fenestrata* (Roxb.) Rehd., from temperate forests of Meghalaya and adjoining forests of Shillong plateau, Nagaland, Mizo and Mikir Hills (1800-3500 m.) during winter season from November to March (Singh et al. 2002). Report of *M. hypopitys* L. from North Eastern India (Barik et al. 2009) has indicated that this belt might indeed be abode for the Genera. The projected range of *Monotropa* in Northeast Hills (Meghalaya and Mizoram-Manipur-Kachin forest zones) has also been proposed for

prioritized conservation of amphibians and reptiles, highlighting importance of said areas for conservation, which are yet currently out of protected area network (Pawar et al. 2007). One of the collection sites (Jorepokhari) of *M. uniflora* in the present study is located within plantation of *Cryptomeria japonica* (introduced from Japan), while collection of *M. humile* by Matsuda et al. (2011) from ecotone of natural forests of *Cryptomeria japonica* in Tateyama, Japan indicates prevalence and competence of Monotropeoideae in allelopathic environment.

The climatic relationship of Monotropoid distribution predicted by earlier workers is in line with the present work and the present study is able to justify the same by refining spatial extent of their distribution in climatically suitable areas. The congruence of the potential range of the *M. uniflora* and other Asian Monotropoids like *Monotropastrum* and *Cheilothea*, indicates that the potential distribution of *M. uniflora* could act as a surrogate for understanding the range of the Monotropeoideae (Ericaceae) in South Asia. The identified novel distributional areas may add to the effective conservation efforts of the subfamily in the region.

One of the prerequisite of realizing niche of a mycoheterotrophic species is the range maps of many ectomycorrhizal host trees, which unfortunately are of limited availability in public domain of a developing nation. Furthermore, modeling species distribution with more occurrence records would be helpful in refining spatial scale of suitable areas. Therefore the currently predicted distribution of Monotropeoideae in South Asia has to be further tested with more occurrence records and calibrated with the range maps of ectomycorrhizal fungi and their host trees for consolidation.

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