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Range extensions for the rare moss *Plagiothecium handelii*, and its transfer to the resurrected genus *Ortholimnobium*

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*Plagiothecium handelii* is newly recorded for Europe (Austria and Romania) and eastern North America (Tennessee, USA). This dainty species was previously known only from Yunnan and Sichuan Provinces, China. Morphologically, the disjunctive populations belong to a single species. A 27-taxon phylogeny of *Plagiothecium* based on nuclear ITS and plastid rpl16 intron DNA sequence data resolved Austrian and Chinese populations of *P. handelii* as sisters, in a clade with *P. paleaceum*, a julaceous Himalayan species with cochleariform leaves. In contrast, *P. handelii* is a filiform plant with distant, ovate-acuminate leaves. In sequence identity the three terminals have a similar level of variation, suggesting that the geographic disjunction between the two populations of *P. handelii* is quite old. Morphologically and genetically the clade is a well defined lineage (*Ortholimnobium*) that is transitional between *Plagiothecium s.str.* and *Struckia*. The new combinations *O. paleaceum* and *O. handelii* are made. In Europe, *O. handelii* should be classified as vulnerable.

Additional specimens of *P. handelii* from Romania and the eastern United States.

Collections of *Plagiothecium handelii* from Austria and China were included in Wynns et al.’s molecular study of the genus. Here, we performed phylogenetic analyses of combined nuclear ITS and plastid rpl16 intron DNA sequence data from 27 collections of *Plagiothecium* and related pleurocarpous mosses (Table 1), in order to establish the conspecificity of Chinese and Austrian populations of *P. handelii*, and to place the species in a phylogenetic framework.

**Methods**

For the DNA analyses we included 20 collections of *Plagiothecium* (sensu Zuo et al. 2011), six collections of other Plagiotheciacaeae (*Isopterygiopsis* Z. Iwats. and *Platydictya* Berk.), and one collection of *Fabronia pusilla* Raddi, which was used as the outgroup (Table 1). DNA extraction, PCR amplification and DNA sequencing were performed with the protocol of Wynns et al. (in press). Sequence alignment was performed manually in MEGA4 (Tamura et al. 2007). Next, indel data were generated for each alignment in SeqState ver. 1.4.1 (Müller 2005) using simple indel coding (Simmons and Ochoterena 2000). A single data file including both nucleotide and indel data for each DNA region was assembled in NEXUS format and analyzed by maximum parsimony (branch-and-bound search) in PAUP ver. 4.0.10b (Swofford 2002). A bootstrap (BS) analysis was also performed (2000 replicates) using branch-and-bound. A partition homogeneity test was performed (1000 branch-and-bound replicates),
to confirm that the two data sets were congruent and appropriate for a combined analysis.

In addition to the parsimony analysis, a Bayesian analysis was performed using the program MrBayes ver. 3.2 (Ronquist et al. 2012). The data were divided into four partitions, two of nucleotide sequence data (rpl16 and ITS) and two respective partitions of binary indel data. The partitions were unlinked and were allowed to evolve at different rates. Based on the Akaike information criterion and the results of the hierarchical likelihood ratio tests, the program Modeltest ver. 3.06 (Posada and Crandall 1998), selected the K81uf+I+Γ (Kimura 1981) model of DNA sequence evolution. However, this model is not implemented in MrBayes, so for the nucleotide data partitions we used the GTR+I+Γ model (cf. Lecocq et al. 2013), which in fact had the best overall log likelihood score. For the indel data partitions the default model was used, a Γ-shaped rate variation was assumed (Yang 1993), and the coding bias was set to variable. A Markov chain Monte Carlo (MCMC) analysis was then run for 11,000,000 generations under the default settings. The results of the Bayesian analysis including posterior probability (PP) support values were visualized as a 50%-compromise phylogram based on average branch lengths using the program TreeGraph 2.1.0-386 beta (Stöver and Müller 2010).

Results

The rpl16 data set included 986 characters, of which 54 were indel characters. The ITS data set included 682 characters, of which 22 were indel characters. Thus, the molecular analyses included 1744 characters in total, of which 1668 were nucleotide characters and 76 were indel characters (4.4%).

Figure 1. Ortholimnobium handelii (Broth.) C. Schröck & J.T. Wynns. (A) (in oval) and (B) in situ, Austria. (C) Handel-Mazzetti 8314 (lectotype of Plagiothecium handelii Broth., S no. B160040). (D–I). Leaves: (D, E) Handel-Mazzetti 7817, Yunnan (syntype of P. handelii, S no. B160038); (F) Long 34930, Yunnan (E no. E00387864); (G) Vajda s.n., Transylvania, Romania (C no. C-M-9099); (H) Schröck 17477, Austria (C no. CP0010623); (I) Anderson 10524, Tennessee, USA (DUKE no. 78480); scale bar = 1 mm.
1384 characters were constant, 153 were variable but not parsimony-informative and 207 were parsimony-informative (20.6% variable characters). The strict consensus of two equally-parsimonious trees of 550 steps was well-resolved, with consistency index (CI, Kluge and Farris 1969) = 0.735, CI excluding uninformative characters = 0.627, retention index = 0.790, and rescaled consistency index = 0.581 (Farris 1989). The 50%-compromise Bayesian tree (Fig. 2) was identical to the parsimony tree in topology, with an additional unsupported node. The results of the partition homogeneity test (p = 0.190) indicated that the plastid and nuclear data sets reflect the same underlying phylogeny.

The two populations of Plagiothecium handelii formed a clade (BS = 79, PP = 0.88) that was sister to P. paleaceum (BS = 100; PP = 1). This clade was sister (BS = 100, PP = 1) to a Struckia Müll. Hal. sensu Ignatov et al. (2007) clade. The branches leading to these clades were relatively long (Fig. 2). All together, this early-diverging clade (Struckia sensu Huttunen et al. 2013) was intercalated between P. piliferum (Sw.) Schimp. and the rest of Plagiothecium.

The two populations of Plagiothecium handelii had some variation in both DNA regions (cf. Fig. 2). In ITS, the Chinese population had two autapomorphic transitions and four synapomorphic changes, while the Austrian population had a single synapomorphic transversion. In rpl16, the Chinese population had three autapomorphic changes (one transversion and two transitions) as well as some synapomorphic changes, while the Austrian population had a single autapomorphic transition. In the Chinese population, rpl16 included a variable 10–21 base pair insertion that was also present in P. laetum Schimp., P. draytonii (Sull.) E.B. Bartram, P. paleaceum, P. piliferum from Russia, Isopterygiopsis pulchella (Hedw.) Z. Iwats. and I. alpicola (Lindb.) Hedenäs (Table 1).
granite, gneiss or schist. Other plants found at these sites include ferns, Lycopodium annotinum L., Vaccinium myrtillus L., and bryophytes characteristic of montane boulderrich forests such as Hylocomium splendens (Hedw.) Schimp., Plagiothecium asplenioides (L.) Dumort., P. porelloides (Torr. ex Nees) Lindenb., Polytrichum formosum Hedw., P. longisetum Sw. ex Brid., Plagiothecium denticulatum (Hedw.) Schimp., P. succulentum (Wils.) Lindb., Rhytididiadelphus loreus (Hedw.) Warnst., R. subpinnatus (Lindb.) T.J. Kop., and R. triqueurus (Hedw.) Warnst. In contrast, P. neckeroides Schimp., which also has a main distribution in Asia and occurs disjunctively in the Alps, grows in colder, more dynamic sites with larger boulders on the same mountainsides. Unlike P. neckeroides, P. handelii also grows on (less acidic) shale, and thus the distribution of P. handelii in Europe is likely to be slightly wider.

The sister relationship between the Chinese and Austrian populations of Plagiothecium handelii that was found in the molecular analyses (Fig. 2) corroborates their conspecificity. Although the populations from Romania and USA were not included in the molecular sampling, the unusual growth form of this species (felt-like mats of etiolated stems, cf. Fig. 1, A, B) is quite easy to recognize, so we have confidence in their identity. Plagiothecium handelii thus has an unusual distribution pattern. While there are several bryophytes that occur disjunctively between the Alps and the Sino-Himalayan region, including P. neckeroides, Distichophyllum carinatum (Dixon & W.E. Nicholson) (Dixon and Nicholson 1909, Redfearn et al. 1994), Tayloria rudolphiana (Garov.) Bruch & Schimp, (Koponen 1992), and (possibly) Brotherella lorentziana (Molendo ex Lorentz) Loeske (Frahm 2013), none of these species occur disjunctively in the eastern USA. Similarly, some bryophytes occur disjunctively between the eastern USA and southwest China, for example Entodon macropodus (Hedw.) Müll. Hal. (Iwatsuki and Sharp 1967), Brotherella leana (Sull.) Müll. Hal., Grimmia pilifera P. Beauv. and Acroblus ciliatus (Mitt.) Schinnt. (Iwatsuki 1972), but these species are not found in Europe. This suggests either the European or American populations of P. handelii may have arisen through long-distance dispersal (cf. Frahm 2013).

On the other hand, Plagiothecium handelii is a dioicous species that is seldom collected in fruit, reducing the likeliness of dispersal by spores. Furthermore, it is restricted to pristine habitats which apparently have a similar ecology at all of the stations. Also, the plants occur in discontinuous localities, both in Europe and America. Last, the relatively large genetic divergence (Fig. 2) between Austrian and Chinese collections is suggestive of a very old disjunction. These facts support a hypothesis that extant populations of P. handelii are relicts of a pre-Pleistocene flora that once had a much broader distribution (Steere 1937, Iwatsuki 1972, Schuster and Damsholt 1974, Manos and Stanford 2001, Hedenäs 2008, Patiño et al. 2016). We believe this explanation is more likely. Many authors (Herzog 1926, Müller 1954, Schuster and Damsholt 1974, Schönswetter et al. 2005, Damsholt 2013) suggest that certain small Alpine localities must have remained uncovered during the Pleistocene glaciations and thus served as refugia for older floras.

**Phylogenetic position of Plagiothecium handelii**

The sister relationship between Plagiothecium handelii and P. paleaceum was first identified by Zuo et al. (2011), as was the sister relationship between this lineage and Struckia. Huttunen et al. (2013) subsequently transferred P. handelii and P. paleaceum to Struckia. Wynns et al. (in press) found P. handelii and P. paleaceum morphologically closer to...
Plagiothecium than to Struckia, and like Zuo et al. they included Struckia in Plagiothecium.

Struckia sensu Ignatov et al. (2007) includes two species. Struckia argenteata (Mitt.) Müll. Hal. is not uncommon on tree trunks in the Sino-Himalayan region, while S. enervis (Broth.) Ignatov, T.J. Kop. & D.G. Long occurs on rocks, and has a restricted distribution in boreal Asia with a disjunctive occurrence in southwest China (Tan et al. 1990, Hedenäs 1996). Struckia differs from Plagiothecium in several respects. In general, species of Plagiothecium are prostrate or semi-prostrate satiny mosses with spreading leaves, whereas both species of Struckia are small erect mosses with appressed leaves and a shaggy appearance (produced by the long-attenuate leaf tips). Microscopically, the leaves are hardly decurrent in Struckia, whereas this is a defining feature of Plagiothecium. Also unlike Plagiothecium, the leaves in Struckia have a large number of quadrate alar cells (reminiscent of Entodon Müll. Hal.). Struckia argenteata further differs from Plagiothecium in its epiphytic habit, denticulate upper leaf margins, and shiny, small-mouthed capsules with conic-mammillate opercula. On the other hand, S. argenteata resembles the sympatric species P. handelii and P. paleaceum in having short erect capsules and ovate-acuminate, concave, subplicate leaves that are often subtended by cells of the cortical hyalodermis when detached.

Struckia enervis is peculiar in that it forms erect defoliated shoots topped with clusters of shortly ligulate, costate gamemae (reduced leaves) (cf. Abramova and Abramov 1981). The specialized morphology of S. enervis promotes vegetative spread and probably arose with the transition away from the epiphytic habitat. It is presumably the more derived species of Struckia (Tan et al. 1990).

Paraphyletic taxa are the real result of evolution (Hörandl 2006) and should not automatically be rejected as the basis for names. In this case Plagiothecium would be rendered paraphyletic by retaining P. handelii and P. paleaceum but excluding Struckia as a distinct genus that evolved rapidly under adaptive pressures associated with a change to epiphytism. This is a plausible hypothesis that is consistent with the morphological and genetic (Fig. 2) differences between Plagiothecium and Struckia. However, this classification hides the close relationship between the P. handelii + P. paleaceum clade and Struckia. Morphologically P. handelii and P. paleaceum share features of both genera and are clearly a transitional taxon that can itself be treated as a distinct genus. The type of the genus Ortholimnobium Dixon belongs to P. paleaceum (Enroth et al. 1992), so this name is available.

**Taxonomy**


Plants glossy, pseudo-stipitate; branches more or less terete, with cortical hyalodermis; leaves imbricate to distant, symmetric, ovate-acuminate, concave, weakly plicate, decurrent, with short narrow cells; capsules short, erect.

Included species:


Syntypes: INDIA. Himalaya, Sikkim, Tonglo, 7–8000 ft., Hooker 1006, ex herb. Mitten (NY nos. 00913411, 00913412), Singalehah, 11000 ft., Hooker 1007 (NY nos. 00913413, 00913414, P no. PC0132613).

Enroth et al. (1992) reported that the lectotype specimen was probably collected by T. Thomsen, not J. D. Hooker. Here we have simply followed the information given by Mitten (1859), which corresponds with the label data.


NEPAL. Sankhuwasabha Distr., E bank of Saldim Khola, near bridge, on bank under stump in mossy *Tsuga* forest, 27°46’N, 87°16’E, ca. 2920 m, 12 Oct 1991, D. G. Long 21152 (E no. E00387890).


Lectotype (designated here): AUSTRIA. Steiermark, bei Schladming, Wald am Rissachfall [Riesachfall], 1080 m, 30 July 1876, sub *P. denticulatum*, *P. roeseanum* var. gracile (GJO 0071175).

Syntypes (all leg. and det. J. Breidler as *P. roeseanum* var. gracile): AUSTRIA. Salzburg, Pinzgau, 12–1500 m, Wald am Krimmler Fall, 1879, sub *P. denticulatum* (GJO 00711660), Felsstrümmerwerk im Walde in der Ammerthal Öd n. Mittersill, 1879, sub *P. denticulatum* (GJO 00711649), Steiermark, bei Schladming, Wald unterhalb der Ursprungalm im Preuneggthald bei Schladming, Wald unterhalber der Ursprungalm im Preuneggthald, 1880 (GJO 00711577), ca. 1300 m, 7 Aug 1876, sub *P. denticulatum* (GJO 00711749), Unterthal, 1870, sub *P. sylvaticum* (GJO 00711711), Felsstrümmer beim Kreuztrog [Kreuztrog südlich Bräutal] b. St. Nikolai in der Sölk, 1200 m, 1883, sub *P. roeseanum* var. tenellum (GJO 00711766), Seewieggraben bei Aich, 8–900 m, 1888, sub *P. roeseanum* var. tenellum (GJO 00717127).


Syntypes: GERMANY. Algiu, in Sandsteinklüften beim Zwintegst 3100’ nw. und am Freiberge 2800’ nö, leg. Molendo [?].

Plants bright green, delicate, pseudo-stipitate, in loose trailing felt-like mats over rock or humus. Rhizoids fasciculate, smooth, reddish-brown. Stems with cortical hyalodermis, often etiolated and threadlike with leaves poorly developed, or appearing thickened by small clasping leaves; branches frequent, somewhat flattened. Leaves distant, or somewhat overlapping, 1–1.4 × 0.35–0.65 mm, concave, erect-spreading, twisted when dry, shortly ovate- to oblong-lanceolate, weakly striate, acuminate with slender tip, clasping at base, decurrent; leaf margins entire, erect toward base; median leaf cells narrow, (75)85–125 × 7.5–10 μm; basal cells sometimes brownish; decurrent cells inflated, ovoid, rectangular, irregularly thick-walled, flanking thin-walled rectangular stem cortical cells that detach with leaves. Dioicus [fide Brothertus 1929]. Seta orangish- or reddish-brown. Capsule erect or slightly inclined, cylindric, slightly constricted below mouth, not or weakly furrowed when dry, ca 1 mm long without lid. Operculum shortly rostrate. Annulus present.


**Variation**

In general *Ortholimnobium handelii* is quite stenotypic. However, a form with wider cells occurs at high elevations. We noticed this both in Austrian (Breidler s.n., 2100 m, GJO 0071156) and American (Anderson 10524, ca 2000 m, DUKE no. 78480) populations. Occasional collections of most species of *Plagiothecium* s.l. have a laxer leaf areolation, perhaps in response to environmental factors.

**Conservation**

In Austria, *Ortholimnobium handelii* is apparently rare to scattered in mountainous areas with siliceous bedrock. It has a wider distribution than the sympatric *Plagiothecium neckeroides*, which is rare or absent in the western Alps. Most populations of *O. handelii* are not currently threatened, but the complex local climate of scree slopes can certainly be disturbed by road construction and logging. Although numerical population data are missing, *O. handelii* can be...
categorized as VU D1 (Vulnerable) under current IUCN criteria, on the basis of small population sizes. Now that *O. handelii* is known to occur in Europe, future floristic studies should improve our knowledge of the frequency and distribution of this species.

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