Haplodontium altunense (Bryaceae, Bryopsida), a new moss species from Northwest China

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Abstract

Haplodontium altunense X.R. Wang & S. Mamtimin, a new moss species of the family Bryaceae from Xinjiang Uygur Autonomous Region, China is described and illustrated. Genetic analysis based on ITS sequences shows that this species is a member of the Bryaceae and in the same clade as Anomobryum. Particularly distinctive features of the new species include: double peristome; the exostome has raised and membranous chomata with united lamellae between two teeth proximally; the endostome is poorly developed and all the endostomial material tightly adherent to the exostome.

Keywords

Altun Mountain, Bryum, ITS, Plagiobryoides, Ptychostomum

Introduction

Haplodontium Hampe, classified in the family Bryaceae, has been confused with the genus Mielichhoferia Horisch. Shaw and Crum (1984) transferred all species in the genus to Mielichhoferia because of the similar peristome. Genetic research (Cox et al. 2000; Goffinet et al. 2001; Pedersen et al. 2003, 2007; Holyoak and Pedersen 2007) has clearly shown that some species originally in Haplodontium are nested within Bryaceae.

* The authors contributed equally to this study.
Spence (2005) resurrected the genus *Haplodontium* and transferred two North American species of *Mielichhoferia* to *Haplodontium*. In addition, the genus *Mielichhoferia* has been accommodated in the family Mielichhoferiaceae (Shaw 2009). Pedersen et al. (2007) showed that *Haplodontium*, *Acidodontium* Schwägr. and *Anomobryum* Schimp. are sister taxa and *Haplodontium* should be included in the family Bryaceae.

**Materials and methods**

**Morphological observations**

Microscopic examination was carried out using traditional methods. The collections of *Haplodontium* and relevant species of Bryaceae in the herbarium of Hebei Normal University (HBNU), Institute of Applied Ecology, Chinese Academy of Sciences (IFP), Kunming Institute of Botany, Chinese Academy of Sciences (KUN), Institute of Botany, the Chinese Academy of Sciences (PE) and Xinjiang University (XJU) were examined. Authors observed the plants under the dissecting microscope and examined the leaves, capsules and peristome under the compound light microscope and scanning electron microscope. Light micrographs were photographed using a Nikon E-800 microscope with a Nikon DXM1200F digital camera. The peristome and spores were mounted on double sided sticky tape on aluminium stubs, gold-coated and viewed using a Hitachi S-4800 field emission SEM. All line drawings were made using the drawing tube attachments of these optical microscopes.

**Phylogenetic analyses**

Twenty-one samples were used for the analyses (Table 1). To evaluate the systematic position of *Haplodontium altunense*, 20 representatives of allied genera in the family Bryaceae, including *Anomobryum*, *Bryum*, *Gemmabryum*, *Plagiobryum* and *Ptychostomum*, were also sampled as part of the ingroup (Cox et al. 2000; Goffinet et al. 2001; Pedersen et al. 2003, 2007; Kato et al. 2013). *Bryum argenteum* was selected as an outgroup. In addition to 10 sequences from GenBank, 11 sequences newly produced for the present study were included.

Genomic DNA was extracted from freshly collected and silica gel-dried plants using a Plant Genomic DNA Kit (TIANGEN Biotech (Beijing) Co., Ltd.) according to the manufacturer’s protocol. One nuclear marker ITS was chosen. The following primers were used to amplify the marker: ‘18SF’ and ‘26SR’ for the ITS region, or sometimes ‘18SF’ and ‘5.8SR’ for ITS1, and ‘5.8SF’ and ‘26SR’ for ITS2 (Hartmann et al. 2006). PCR cycles used an initial denaturation step of 3 minutes at 95 °C, followed by 35 cycles of 30 seconds at 95 °C, 30 seconds at 50 °C, 90 seconds at 72 °C, and a final elongation of 5 minutes at 72 °C. PCR products were purified with a Gel Extraction Kit (Cwbio, Shanghai, China) following the instruction manual. These purified PCR products were sequenced by Life Technologies Inc., China (http://www.lifetechnologies.com).
Sequence chromatograms were compiled using SeqMan II (DNASTAR Inc., Madison, WI, USA) and then aligned manually in PhyDE 0.9971 (Müller et al. 2010). Regions of partially incomplete data at the beginnings and ends of sequences were identified and excluded from subsequent analyses. Gaps were treated as missing data. The aligned ITS dataset was composed of 1213 bp.

The maximum likelihood (ML) method was performed using RAxML v.8.2.12 on the CIPRES Science Gateway (http://www.phylo.org/), and inferred under the default settings (Stamatakis 2014). The fast bootstrap option was used with 1000 replicates. TreeGraph 2 (Stöver and Müller 2010) was used to summarize the topologies and support values from the analyses.

Results

Taxonomic treatment

*Haplodontium altunense* X.R. Wang & S. Mamtimin, sp. nov.

urn:lsid:ipni.org:names:

Figs 1–3

**Type.** China. Xinjiang, Ruociiang County, Altun Mountain National Nature Reserve, 37°0.42’N, 88°36.35’E, 4290 m a.s.l., 22 July 2011, S Mamtimin 16752 (*holotype*: HBNU!; *isotype*: XJU!).

### Table 1. Voucher information and GenBank accession numbers of taxa used in the phylogenetic analyses.

| Taxon                      | Voucher (Herbarium) | Origin       | GenBank No. | Source                  |
|----------------------------|---------------------|--------------|-------------|-------------------------|
| *Anomobryum auratum 1*     | L.B. Li 20073626 (HBNU) | China        | MZ470251    | This study              |
| *Anomobryum auratum 2*     | L.B. Li 20073628 (HBNU) | China        | MZ470252    | This study              |
| *Anomobryum julaceum*      | L.B. Li 20072925 (HBNU) | China        | FJ796895    | Wang et al. 2011        |
| *Bryum argenteum*          | X.R. Wang 20150512031 (HBNU) | China        | MZ470253    | This study              |
| *Bryum paradoxum*          | J.C. Zhao 0610007 (HBNU) | China        | EU878207    | Wang et al. 2011        |
| *Bryum recurvatum*         | W.Q. Li 040900 (HBNU) | China        | EU878217    | Wang et al. 2011        |
| *Gemmabryum caespiticium*  | X.R. Wang 20156001 (HBNU) | China        | MZ470254    | This study              |
| *Haplodontium altunense*   | S. Mamtimin 16752 (XJU) | China        | MZ470255    | This study              |
| *Plagiothyrium zierii*     | W.Q. Li 000514 (HBNU) | China        | EU878219    | Wang et al. 2011        |
| *Ptychostomum arcticum*    | S. Mamtimin 15457 (HBNU) | China        | MZ470256    | This study              |
| *Ptychostomum binnum*      | L. Hedenäs B90015 (S) | Sweden       | DQ381780    | Holyoak and Hedenäs 2006 |
| *Ptychostomum cernuum*     | Y.L. Niu 110002 (HBNU) | China        | MZ470257    | This study              |
| *Ptychostomum inclinatum*  | N. Cao 20050085 (HBNU) | China        | EU878227    | Wang et al. 2011        |
| *Ptychostomum lunchoalun*  | N. Cao 20050187 (HBNU) | China        | FJ796878    | Wang et al. 2011        |
| *Ptychostomum neodamense*  | L. Hedenäs B65900 (S) | Sweden       | DQ381772    | Holyoak and Hedenäs 2006 |
| *Ptychostomum pallens*     | M.X. Xiao 20091246 (HBNU) | China        | MZ470258    | This study              |
| *Ptychostomum pallidescens*| S. Mamtimin 15265 (HBNU) | China        | MZ470259    | This study              |
| *Ptychostomum pendulum*    | J.C. Zhao 20060463 (HBNU) | China        | FJ796811    | Wang et al. 2011        |
| *Ptychostomum pseudotriquetrum* | D.T. Holyoak B90021 (S) | Ireland      | DQ381774    | Holyoak and Hedenäs 2006 |
| *Ptychostomum purpurascens*| Y.L. Niu 110045 (HBNU) | China        | MZ470260    | This study              |
| *Ptychostomum turbinatum*  | S. Mamtimin 15095 (HBNU) | China        | MZ470261    | This study              |
**Diagnosis.** Particularly distinctive features of the new species including: double peristome; the exostome has raised and membranous chomata with united lamellae between two teeth proximally; the endostome is poorly developed and all the endostomial material tightly adherent to the exostome.
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Description. Plants small, soft and dull, brown-green. Stems short, 2.5–6 mm high, weakly julaceous, branched, circular or pentagonal circular in transverse section with small and thick-walled peripheral cells surrounding 2–3 layers gradually larger and thin-walled cortical cells, central strand weakly developed. Leaves imbricate when

Figure 2. Light micrographs of Haplodontium altunense A–C plants (dry) D Capsule (dry) E capsule (wet) F annulus growing on the capsule mouth G annulus falling off the capsule mouth H dorsal views of median peristome showing the large papillae along the horizontal and median vertical lines I dorsal views of distal peristome showing adherent endostomial material and exostome teeth J transverse section of stem K transverse section of midleaf L transverse section of costa M leaf apex N median laminal cells O basal laminal cells P leaves. Photographed by Xiaorui Wang from the holotype (HBNU!).

Description. Plants small, soft and dull, brown-green. Stems short, 2.5–6 mm high, weakly julaceous, branched, circular or pentagonal circular in transverse section with small and thick-walled peripheral cells surrounding 2–3 layers gradually larger and thin-walled cortical cells, central strand weakly developed. Leaves imbricate when
dry, erect when moist, enlarged towards stem apex, ovate to broadly ovate, concave, 0.5–1.1×0.3–0.7 mm; base not decurrent; margins plane or recurved medially, 1-stratose, limbidium absent, smooth or finely serrulate distally; apex broadly acute; costae not reaching apex, guide cells weakly developed, 2–4 in one layer in costal transverse section, ventral and dorsal stereid bands present; alar cells not differentiated from juxtacostal cells; laminal cells lax; distal laminal cells rhomboidal, 30–44×11–21 μm, with slightly thickened walls; medial laminal cells long rhomboidal to rectangular,

**Figure 3.** Scanning electron micrographs of *Haplodontium altunense* A dorsal views of peristome B dorsal views of peristome with annulus C dorsal surface of exostome teeth proximally showing the pores and the raised and membranous chomata horizontal lines with united lamellae between two teeth D dorsal surface of exostome teeth distally showing the large papillae along the median vertical lines E dorsal surface of exostome teeth distally showing the endostomial material adherent to the teeth F ventral views of peristome G ventral surface of peristome proximally showing the smooth endostome basal membrane H ventral views of peristome distally I, J ventral surface of peristome medially and distally showing the endostomial material adherent to the teeth K spore L exine ornamentation of spore. Photographed by Xiaorui Wang from the holotype (HBNU!).
Haplodontium altunense new species China

37–69×12–20 μm, somewhat narrower in 2 or 3 rows toward the margins but not forming a distinct border; proximal laminal cells long rectangular, 37–56×20–28 μm. Dioicus(?). Perigonía not seen. Perichaetia at the end of short, inconspicuous stems, appearing laterally because of well-developed innovations; perichaetial leaves larger than vegetative leaves. Setae single, light brown, 15–19 mm long. Capsules nutant and symmetric, reddish brown, obovoid, 1.5–2 mm, neck short and indistinct, mouth small, stomata abundant in the neck, superficial; opercula long-conic with short rostrate; annulus present, consisting of two rows of cells, revoluble and cells with slit-like lumen; peristome double, exostome inserted below the mouth, teeth lanceolate, red-brown and pored, raised and membranous chomata with united lamellae between two teeth proximally, pale yellow to hyaline and largely papillose along horizontal and median vertical lines distally; endostome poorly developed, basal membrane smooth, segments and cilia rudimentary, all the endostomial material strongly adhere to the exostome. Spores spherical, 20–22 μm in diameter, minutely papillose.

**Etymology.** The specific epithet altunense refers to the type locality in Altun Mountain National Nature Reserve in the Northwestern China.

**Distribution and habitat.** China (Xinjiang). Only known from the type locality, on soil substrates at 4290 m in the Altun Mountain National Nature Reserve. The population grows in a dry, cold, and windy habitat with intense evaporation. The companion species include some xerophytic mosses of the family Pottiaceae.

**Chinese name.** 阿尔金拟缺齿藓 (ā ěr jīn nǐ quē chǐ xiān)

Phylogenetic analyses

The present phylogenetic analysis, based on the nuclear ribosomal internal transcribed spacer region ITS1–5.8S–ITS2 (hereafter, ITS) region, included 20 species from six genera, as well as Bryum argenteum Hedw. as outgroup (Fig. 4). The only sample of Haplodontium altunense is sister to the Anomobryum clade (81 MLBS), which is monophyletic with three members. Twelve samples of Ptychostomum Hornsch. formed a monophyletic clade (91 MLBS), in which Gemmabryum caespiticium (Hedw.) J.R. Spence was nested. Plagiobryum zierii (Hedw.) Lindb. is closely related to the Ptychostomum clade. The Bryum Hedw. clade (100 MLBS) with three species were basal to the main clades.

Discussion

Pending a careful examination of Bryaceae in the Altun Mountains, we discovered an unusual collection with distinctive morphological features of the peristome: endostome poorly developed and strongly adherent to the exostome. We thought this collection could belong to the genus Synthetodontium Cardot at first sight based on its fused peristome. Phylogenetic analysis showed that this collection was nested into Bryaceae clade. It is genetically distinct from the closely related Anomobryum group (Fig. 4). Further examination revealed that it has morphological characters of Haplodontium
such as stems nearly julaceous, leaf margins entire, distal and medial laminal cells laxly rhomboidal to rectangular. However, the collection is distinguished from the other previously recognized species in the region by its double peristome, and raised and membranous chomata with united lamellae between two exostome teeth proximally. We thus describe it as a new species of the genus *Haplodontium*.

The gametophytes of *Haplodontium* species are similar to those of *Plagiobryum* and *Plagiobryoides* J.R.Spence (Spence 2015). Genetic research has shown that the type species of *Haplodontium* is closely related to *Acidodontium* and *Anomobryum* (Cox and Hedderson 2003; Pedersen et al. 2007). Peristome reduction is common and complex in *Haplodontium*, from double to single to absent (Spence 2005).

The gametophyte characters of *Haplodontium altunense* are similar to *Plagiobryoides brachyneura* (Kindberg) J.R.Spence (Spence 2015). At the same time, both species have double, reduced and fused peristome. In *P. brachyneura*, setae are red-brown and 5–15 mm long, capsules are inclined to nutant, elongate-pyriform and 2–4 mm long with elongate-neck, opercula are weakly convex, exostome teeth are short and irregular in shape. However, those characters of *H. altunense* are as follows: setae are light brown and 15–19 mm long, capsules are nutant, pyriform to obovoid and 1.5–2 mm long with indistinct short-neck, opercula are long-conic with short rostrate, exostome teeth are regular long lanceolate.

Figure 4. Phylogenetic relationships of *Haplodontium altunense* sp. nov. from ITS rDNA with related genera in Bryaceae based on Maximum Likelihood analysis. Support values are given above branches. *Bryum argenteum* was served as an outgroup.
Haplodontium altunense is also similar to Ptychostomum pendulum Hornschuch (Spence 2015) (≡Bryum algovicum Sendtnr ex Müller Hal. (Li 2006; Zhang et al. 2007) in that the endostome adheres to the exostome teeth. While the former species differ from the latter one in length of stems (2.5–6 mm vs. 5–20 mm), leaf apex (broadly acute or obtuse vs. acuminate), leaf margin (indistinct bordered vs. strong limbidium), costae (ending at or near the apex vs. long-excurrent in denticulate awn), exostome teeth (united at the base, large papillose above vs. separate, smooth above), endostome (segments reduced vs. segments with ovate perforations).

Wang et al. (2017) reported one new species, Haplodontium zangii X.R.Wang & J.C.Zhao, from Tibet, China and transferred two Chinese species in Mielichhoferia to Haplodontium as new combinations: H. himalayanum (Mitt.) X.R.Wang & J.C.Zhao and H. sinensis (Dix.) X.R.Wang & J.C.Zhao. The most significant difference among the four species of Haplodontium in China is that the first three species all have single peristome, while H. altunense has double and fused peristome.

Key to the Haplodontium species in China

1 Peristome double, exostome teeth lanceolate, raised and membranous chomatata with united lamellae between two teeth proximally, endostome reduced, all the endostomial material strongly adhere to the exostome........... H. altunense
– Peristome single, exostomial........................................................................ 2
2 Leaves lanceolate; costae excurrent, ending in long denticulate awns, awns 140–310 μm long.................................................................H. himalayanum
– Leaves ovate to oblong-ovate; costae subpercurrent to ending in short awns, awns 0–130 μm long ............................................................H. zangii
3 Leaf apex cucullate; capsules pyriform; exostome teeth regularly lanceolate, not perforate, vertically striped on dorsal surface, smooth on ventral surface. .................................................................H. sinensis
– Leaf apex plane; capsules subglobose to short pyriform; exostome teeth irregularly linear-lanceolate, sometimes perforate, smooth or finely papillose .. ............................................................................H. sinensis

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