# Molecular and morphological evidence for a remarkable new species of *Leptopus* (Phyllanthaceae) from Southeast Yunnan, China (#60870)

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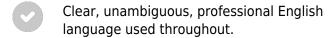
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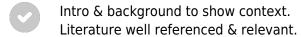
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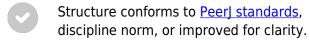
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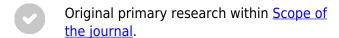


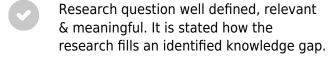


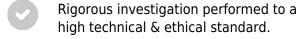




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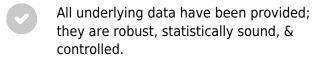


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I commend the authors for their extensive data set, compiled over many years of detailed fieldwork. In addition, the manuscript is clearly written in professional, unambiguous language. If there is a weakness, it is in the statistical analysis (as I have noted above) which should be improved upon before Acceptance.



# Molecular and morphological evidence for a remar ble new species of *Leptopus* (Phyllanthaceae) from Southeast Yunnan, China

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Leptopus malipoensis, a remarkable new species of Phyllanthaceae form Southeast Yunnan Province, China, is described. The phylogenetic position of the new species within the genus Leptopus was analyzed based on the nuclear ribosomal internal transcribed spacer (nrITS) and the plastid DNA region matK. The results show that L. malipoensis is highly supported to be the sister of L. fangdingianus (P. T. Li) Voronts. & Petra Hoff., a species endemic to Western Guangxi Province, China. Morphologically, the new species can be distinguished from all known congeneric taxa by its long and slim branches usually pendulous or procumbent, large leaves up to 15 cm long and 7 cm wide, and further differs from its sister species by its hirsute stems, leaves and pedicel of female flowers, longer pedicel of male flowers, 3-locular ovary and 3 styles. A key to all the Leptopus species is provided.

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### 19 Abstract

- 20 Leptopus malipoensis, a rema ple new species of Phyllanthaceae form Southeast Yunnan
- 21 Province, China, is described. The phylogenetic position of the new species within the genus
- 22 Leptopus was analyzed based on the nuclear ribosomal internal transcribed spacer (nrITS) and
- 23 the plastid DNA region *matK*. The results show that *L. malipoensis* is highly supported to be the
- 24 sister of L. fangdingianus (P. T. Li) Voronts. & Petra Hoff., a species endemic to Western
- Guangxi Province, China. Morphologically, the new species can be distinguished from all known
- 26 congeneric taxa by its long and slim branches usually pendulous or procumbent, large leaves up
- 27 to 15 cm long and 7 cm wide, and further differs from its sister species by its hirsute stems,
- leaves and pedicel of female flowers, longer pedicel of male flowers, 3-locular ovary and 3 styles.
- 29 A key to all the *Leptopus* species is provided.

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- 31 **Subjects** Biodiversity, Molecular Biology, Plant Science, Taxonomy
- 32 **Keywords:** Leptopus, Phyllanthaceae, Poranthereae, taxonomy, Yunnan, China

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### 34 Introduction

Leptopus Decne. is a small genus in the tribe Poranthereae, subfamily Phyllanthoideae of Phyllanthaceae and distributed mainly from the Caucasus to Malesia (Vorontsova & Hoffmann, 2009; Webster, 2014). In the latest taxonomic revision of *Leptopus*, nine species were accepted and six among them were recorded in China (Vorontsova & Hoffmann, 2009). Results from molecular phylogenetic analyses showed that *Leptopus* was sister to *Actephila* Blume with high support (Vorontsova et al., 2007; Vorontsova & Hoffmann, 2008). Morphologically, the genus Leptopus can be distinguished from its sister clade by its leaves membranous to paper (vs. leathery, rarely papery), disk lobed evidently (vs. annular), fruit less than 10 mm in diameter (vs. more than 10 mm in diameter), exocarp adnate to endocarp (vs. free from endocarp) and mature seed with copious endosperm (vs. seed without endosperm) (Li et al., 2008).

During the field investigation in Malipo Hsien, Southeast Yunnan Province of China, in March 2018, two of the authors (E.D. Liu and X.X. Zhu) collected a Phyllanthaceae specimen belongs to *Leptopus*, and the same species was collected again in the same locality by another author (G. Yao) in July 2020. The species superally differs from all the other members of *Leptopus* in morphology. After detailed morphological investigation and molecular phylogenetic analyses of *Leptopus*, it was concluded that the species is new to science, thus it is formally described here.

### Materials & methods

### **Ethics statements**

The collection location of the new species reported in this study is outside any natural conservation area and no specific permissions were required for the location. Since this species is currently undescribed, it is not currently included in the China Species Red List (Wang & Xie, 2004). Our field studies did not involve any endangered or protected species. No specific permits were required for the present study.

### Nomenclature

The electronic version of this article in Portable Document Format (PDF) will represent a published work according to the International Code of Nomenclature for algae, fungi, and plants (ICN), and hence the new names contained in the electronic version are effectively published under that Code from the electronic edition alone. In addition, new names contained in this work which have been issued with identifiers by IPNI will eventually be made available to the Global Names Index. The IPNI can be accessed and the associated information contained in this publication viewed through any standard web browser by using the web address "http://ipni.org/". The online version of this work is archived and available from the following digital repositories: PeerJ, PubMed Central, and CLOCKSS.

### **Material collection**

- 73 Flowering and fruiting specimens of the new species were collected in the mountain area, in
- Nandong to Bajiaoping, Laoshan, Malipo Hsien of Yunnan Province, China, for morphological



study. Leaf mather list for DNA extraction were collected and dried using silica gel in the field.

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### Morphological study

Specimens of *Leptopus* deposited in the herbaria GXMG, IBK, IBSC, KUN and PE were studies carefully in the present study. Field investigation of Chinese *Leptopus* species were also conducted in recent years. Morphological characters of stems, leaves, flowers and fruits of relevant species were photographed and measured. Additionally, morphological comparisons between the new species and all the nine *Leptopus* species accepted by Vorontsova & Hoffmann (2009) were also conducted.

### Phylogenetic study

Phylogenetic relationships among members of *Leptopus* were investigated by Vorontsova et al. (2007) based on analysis of two DNA fragments, viz. the nuclear ribosomal internal transcribed spacer (nrITS) and the plastid DNA region *matK*. To study the phylogenetic position of the new species within the genus *Leptopus*, the two above mentioned DNA regions of the new species were sequenced. Relevant DNA sequences of *Leptopus* species included in Vorontsova et al. (2007) were obtained from the website NCBI (*https://www.ncbi.nlm.nih.gov/*) and used in the present phylogenetic analyses. Additionally, outgroups were selected from the other seven genera of the tribe Poranthereae [*Actephila* Blume, *Andrachne* L., *Meineckia* Baill., *Notoleptopus* Voronts. & Petra Hoffm., *Phyllanthopsis* (Scheele) Voronts. & Petra Hoffm., *Pseudophyllanthus* (Müll. Arg.) Voronts. & Petra Hoffm. and *Poranthera* Rudge] and the genus *Heywoodia* Sim of the tribe Wielandieae, based on previously published phylogenetic frameworks (Kathriarachchi et al., 2005; Vorontsova et al., 2007). DNA sequences of outgroups were also downloaded from the website NCBI. Detailed information about the species sampled and DNA sequences are provided in Table 1.

Total DNA of the new species was extracted and then sequenced by genome skimming following the protocol of Zeng et al. (2018). Plastid and the nrITS sequence reads were assembled using the software GetOrganelle (Jin et al., 2020), with the reference plastid genome of *Glochidion chodoense* C.S. Lee & Im (GenBank accession number: NC\_042906) and nrITS sequence of *Leptopus chinensis* (Bunge) Pojark. (MH710764), respectively. Genes in the plastid genome obtained were annotated in the software PGA (Qu et al., 2019). Then the sequence *matK* was extracted from the assembled whole plastid genome.

Sequenced were aligned using MAFFT 7.221 (Katoh & Standley, 2013) and then three data sets were constructed: the *matK* dataset, the nrITS dataset and the combined dataset (including *matK* and nrITS). All the three datasets were analyzed using two approaches: Bayesian Inference (BI) and Maximum Likelihood (ML). Detailed information about the parameter setting in BI and ML analyses referred the phylogenetic analyses conducted in Yao et al. (2020). The models of nucleotide substitution of the two DNA fragments used were selected under the Akaike Information Criterion (AIC) using jModeTest v. 3.7 (Posada, 2008): TVM+G for *matK* and GTR+I+G for nrITS.



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### Results & discussion

### 117 Phylogenetic analysis

The matK dataset, nrITS dataset and combined dataset contained 2000 bp, 812 bp and 2812 bp, 118 respectively. Conflicted topologies were found between the *matK* and nrITS frameworks (Fig. 1), 119 120 but relevant conflicted phylogenetic nodes were all lowly supported in analyses of the nrITS dataset (Fig. 1B). Phylogenetic relationships derived from the combined dataset were much 121 better resolved compared with those obtained from analyses based on the other two datasets, and 122 phylogenetic relationships among Leptopus species sampled here were all resolved with high 123 supports (Fig. 2). Thus we focus on describing phylogenetic relationships based on the result 124 derived from the combined dataset. 125

Phylogenetic results showed that the genus *Leptopus* was sister to *Actephila* with high support values (MLBS = 100%, PP = 1.00), and the monophyly of *Leptopus* was strongly supported (MLBS = 100%, PP = 1.00). Within *Leptopus*, three major clades were recovered. *Leptopus australis* (Zoll. & Moritzi) Pojark. represents the earliest divergent clade within this genus. The new species is strongly supported to be the sister of *L. fangdingianus* (P.T. Li) Voronts. & Petra Hoffm. (MLBS = 100%, PP = 1.00) and then sister to *L. clarkei* (Hook. f.) Pojark. (MLBS = 100%, PP = 1.00). *Leptopus chinensis* (Bunge) Pojark. and *L. cordifolius* Decne. formed the third clade and sister to the (new species *L. fangdingianus*)-*L. clarkei* clade with strong support (MLBS = 100%, PP = 1.00). Furthermore, the sister relationship between the new species and *L. fangdingianus* was highly supported in both of the *matK* (Fig. 1A) and nrITS (Fig. 1B) frameworks.

### **Morphological comparisons**

Morphologically, the new species has pendulous or prophologically, the new species has pendulous or prophologically, the new species has pendulous or prophologically, the new species has pendulous or prophologically. leaves up to 15 cm long and 7 cm wide. These characters could distinguish it easily from all the other Leptopus members, which usually have ascendant or erect stem and smaller leaves less than 10 cm long and 5 cm wide. The procumbent habit is also recorded in L. clarkei, a species widely distributed from southern China, extending west to Assam and Ruma, and south to northern Vietnam (Vorontsova & Hoffmann, 2009), and also is closely related to the pair of the new species and L. fangdingianus phylogenetically (Figs. 1 and 2). However, the new species differs from L. clarkei by having straight and not ribbed stems (Fig. 3C-F) (vs. flexuous and longitudinally strongly ribbed stems), larger leaves up to 15 cm long and 7 cm wide (vs. less than 10 cm long and 3 cm wide), leaves hirsute on both surfaces (Figs. 3C-E & 4D) (vs. glabrous adaxially and glabrous to sparsely hirsute abaxially), margin of leaves densely hirsute (Figs. 3C-E & 4D) (vs. glabrous to hirsute). While the new species differs from its sister L. fangdingianus by its leaves hirsute on both surfaces (Figs. 3C-E & 4D) (vs. glabrous to sparsely hirsute on both surfaces), pedicel of male flowers 10–25 mm long (vs. usually less than 10 mm long), pedicel of female flowers hirsute (Fig. 4D) (vs. glabrous), ovary 3-locular (vs. 4–5) and styles 3 (vs. 4-5).

Four species of *Leptopus* [viz. *L. micans* (Dunn) Pojark., *L. hainanensis* (Merr. & Chun) Pojark., *L. pachyphyllus* X.X. Chen and *L. robinsonii* Airy Shaw] were not sampled in the



present phylogenetic analyses. Morphologically, the new species also can be easily distinguished 157 from these four congeneric species based on its above-mentioned pendulous or procumbent 158 stems and large leaves. Additionally, the new species has hirsute indumentum, not ribbed 159 branches, chartaceous leaves with 4 (or rarely 5) pairs of secondary veins. In contrast, L. micans 160 has glabrous branches longitudinally ribbed and 8-10 pairs of secondary veins in leaves. 161 Leptopus hainanensis also has glabrous branches and leaves, as well as 2–3 pairs of secondary 162 veins in leaves. While the other two species L. pachyphyllus and L. rasponii both have 163 glabrous and bilaterally flattened branches, and the leaves of L. pachyphyllus are glabrous on 164 both sides and almost succulent. 165

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### **Taxonomic treatment**

- Leptopus malipoensis W.H. Zhang & Gang Yao, sp. nov. (Figs. 3-4) 168
- 169
- Type. CHINA. Yunnan Province, Wenshan State, Malipo Hsien, Laoshan, Nandong to 170
- Bajjaoping, on stone slopes near the roadsides of the semi-shady forests, at the elevation of ca. 171
- 1200 m, 15 July 2020, G. Yao YGYN2020071501 (holotype: IBSC; isotypes: KUN, CANT). 172
- Diagnosis. The species is similar to <u>L. fangdingianus</u> (P.T. Li) Voronts & Petra Hoffm. in 173
- morphology, but differs from the latting y its procumbent habit with long and slim branches 174
- usually pendulous, larger leaves up to 15 cm long and 7 cm wide, hirsute stems, leaves and 175
- pedicel of female flowers, longer pedicel of male flowers, 3-locular ovary and 3 styles. 176
- **Description.** Shrub, monoecious. Stems straight, terete, hirsute; branchlets long and slim, 177
- sometimes up to 1.5 m long, usually pendulous or procumbent, hirsute. Leaves alternate, 178
- chartaceous, elliptic to ovate, 2-15 cm long and 1.2-7 cm wide, both surfaces hirsute, densely 179
- 180 hirsute when young, margin densely hirsute, base cuneate to round, apex acuminate; midvein
- adaxially impressed, abaxially raised; secondary veins usually 4 pairs, rarely 3 or 5 pairs, 181
- adaxially slightly impressed, abaxially raised, obliquely ascending, sometimes arcuately 182
- anastomosing near margins. Petiole 4–12 mm long, hirsute. Stipules narrowly triangular, apically 183 acuminate. Infloresce unisexual or bisexual. The flowers 1- er fascicle, ca. 3 mm in 184
- diameter, light yellow to slightly green; pedicel 10–25 mm long, glabrous; pals 5, oblong, 185
- apically round, adaxially glabrous, abaxially sparsely hirsute, 0-3-veined; petals 5, clavate to 186
- slightly linear, alternating with sepals; disk with 5 contiguous regular segments deeply biobed,
- 187 apical of lobes truncate to round; stamens 5, opposite sepals; filaments free; anthers
- 188 longitudinally denscent. He pale flowers usually 1 per fascicle, 3.5–4 mm in diameter; pedicels 189
- usually 15–25 mm long, sparsely hirsute, apically dilated evidently; sep 5, oblong to ovate-190
- triangular, apically acute to round, adaxially glabrous, abaxially glabrous or sparsely hirsute, 191
- 192 usually 0-5-veined; petals 5, slightly linear, alternating with sepals; disk with 5 contiguous
- regular segments deeply biobed, apical of lobes truncate to round; ovary 3-locular, globose, 193 glabrous; styl free, deeply bifid, lobes usually recurved. Fruiting pedicel 2–3 cm long, hirsute; 194
- capsules depressed globose, smooth, glabrous or sparsely hirsute when young, 4-6 mm in 195
- diameter, 2.5–3 mm high, persistent sepals oblong; seeds 6, brown to dark-brown, hemispheric 196
- or laterally compressed, ca. 2.5 mm long and 2 mm wide. 197



Etyr Degy. Leptopus malipoensis is named after its type locality, Malipo Hsien. Malipo Hsien is 198 a hotspot for biodiversity research in Yunnan Province, China, and many new species were 199 described recently from there, such as Bredia malipoensis D. H. Peng, S. Jin Zeng & Z.Y. Wen 200 (Wen et al., 2019), Habenaria malipoensis Q. Liu & W.L. Zha Zhang et al., 2017), Primulina 201 malipoensis L.H. Yang & M. Kang (Yang et al., 2018), Salacia malipoensis X.D. Ma & J.Y. 202 Shen (Ma et al., 2020). 203 **Phenology:** Flowering in April to August, fruiting in May to October. 204 Paratype: CHINA. Yunnan Province, Wenshan State, Malipo Hsien, Laoshan, Nandong to 205 Bajjaoping, under the semi-shady forests, at the elevation of 1171 m, 5 March 2018, Z.D. Wei, 206 F.Z. Shangguan, X.X. Zhu, et al. LiuED8755 (KUN). 207 Distribution and habitat: The species is known only from its type locality, Malipo Hsien in 208 Southeast Yunnan Province, China (Fig. 5). 209 Habitat. The species grows on stone slopes near the roadsides of the semi-shady forests or under 210 the semi-shady forests, in limestone environments, at the elevation of 1171–1200 m. 211 Chinese name. Ma Li Po Que She Mu (麻栗坡雀舌木) 212 213 Key to species of *Leptopus*, modified from Vorontsova & Hoffmann (2009) 214 1a. Leaf blades coriaceous, almost succulent; endemic to Guangxi Province, China 215 216 Leaf blades membranaceous to thick chartaceous. 1b. succulent 217 never 2 218 2a. Leaf blades with 8-10 visible pairs of secondary veins; fruit strongly reticulate; seed with 219 220 2b. Leaf blades with 0–6 (7) visible pairs of secondary veins; fruit smooth to faintly reticulate; 221 222 3a. Ascendant herb to subshrub up to 0.5 m high; feel pedicels 2–5 mm in flower, 5–9 mm 223 224 3b. Erect to procumbent herb or shrub 0.5-4 m; female pedicels 5-30 mm in flower, 7-36 mm 225 226 227 4b. Male pedicels no less than 3 in length, and usually up to 10 mm or longer ..... 6 228 5a. Branches white to light brown; petals, filaments and styles mostly glabrous; seeds 229 endemic smooth: to Hainan Province. China 230 ......L. hainanensis 231 5b. Branches reddish; male petals, filaments and styles hirsute; seeds transversely to 232 irregularly ridged, sometimes pitted; endemic to Khanh Hoa Province, Vietnam 233 .....L. robinsonii 234 235 236 7a. Leaves usually less than 3 cm in length, and never longer than 5 cm 237 L. chinensis 238

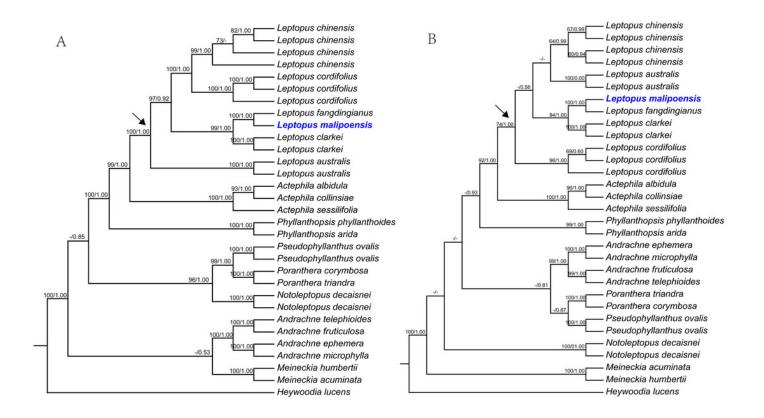


239	7b. Leaves usually longer than 5 cm in length, sometimes even longer than 10 cm in length
240	So I ar loss than 10 am in langth and loss than 2.5 am in width, loss blades with 4.7 nairs
<ul><li>241</li><li>242</li></ul>	8a. Leave less than 10 cm in length and less than 3.5 cm in width; leaf blades with 4–7 pairs of secondary veins; leaf apex rounded
243	8b. Leaves sometimes longer than 10 cm and wider than 3.5 cm; leaf blades 4 or 5 pairs of
244	secoandary veins; leaf apex acute or acuminate
245 246	9a. Stems and branchlets usually pendulous or procumbent; ovary 3-locular; styles 3
247	9b. Stems and branchlets ascendant or erect; ovary 4–5-locular; styles 4–5
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249	2. Junguing
250	Adhikari et al. (2010) described the new species Leptopus nepalensis B. Adhikari, R.P.
251	Chaudhary & S.K. Ghimire from Nepal, the holotype ( <i>B. Adhikari 224</i> , TUCH) of which had
252	been identified as <i>Phyllanthus griffithii</i> Müll.Arg On the basis of the morphological description
253	provided by Adhikari et al. (2010), the new species is characterized by its 6 petals in two whorls,
254	6 stamens (or 3 as observed from the linear picture drawn based on the holotype), 3
255	connated into a column up to about halfway, however all of these characters are much different
256	from those of the genus <i>Leptopus</i> as currently circumscribed (Vorontsova & Hoffmann, 2009;
257	Webster, 2014). Thus the species <i>L. nepalensis</i> is not included in the above key, and the
258	taxonomic status of the species needs to be clarified in further study, especially inolecular
259	phylogenetic analysis.
260	F-17-1-80-1-11-11-11-11-11-11-11-11-11-11-11-11-
261	Acknowledgements
262	The authors are grateful to the curators and staff of the herbaria GXMG, IBK, IBSC, KUN and
263	PE from hosting our visits or providing images of specimens, to Mr. Long in Malipo Hsien,
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265	
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Maximum likelihood (ML) trees of *Leptopus* and its relatives inferred from the *matK* dataset (A) and nrITS dataset (B).

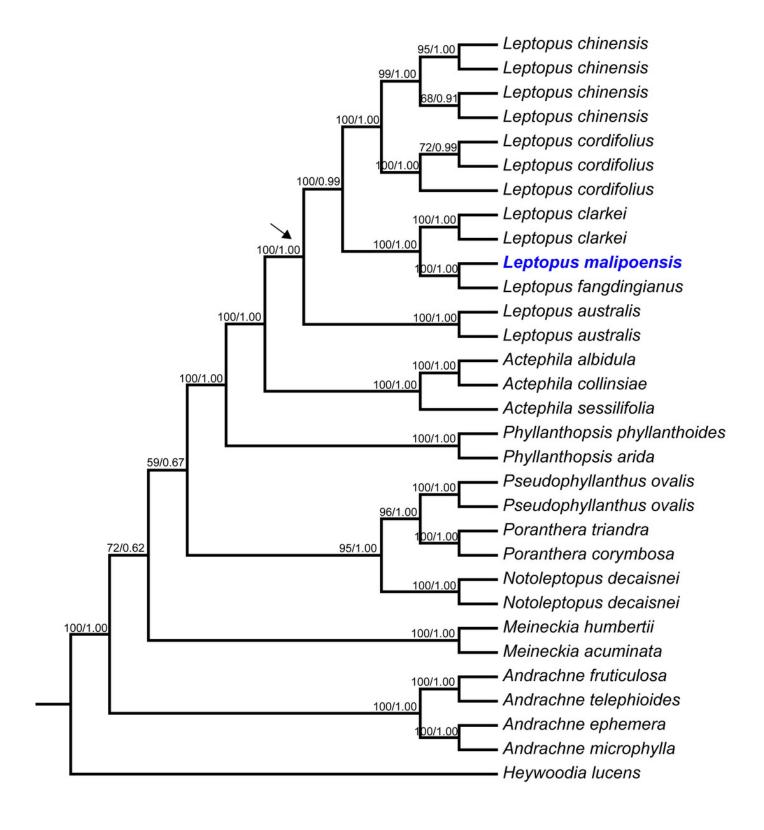
**Fig. 1.** Maximum likelihood (ML) trees of *Leptopus* and its relatives inferred from the *matK* dataset (A) and nrITS dataset (B). Bootstrap (BS) value  $\geq 50\%$  in ML analysis and posterior probability (PP)  $\geq 0.50$  in Bayesian inference (BI) is indicated on the left and right of slanting bar associated with phylogenetic node, respectively. Dashes denote that the phylogenetic node associated was not supported or the BS value is < 50% in ML analysis or PP > 0.50 in BI. The crown node of *Leptopus* is shown by the arrowhead.





Maximum likelihood (ML) trees of *Leptopus* and its relatives inferred from the combined data set (including nrITS and *matK*).

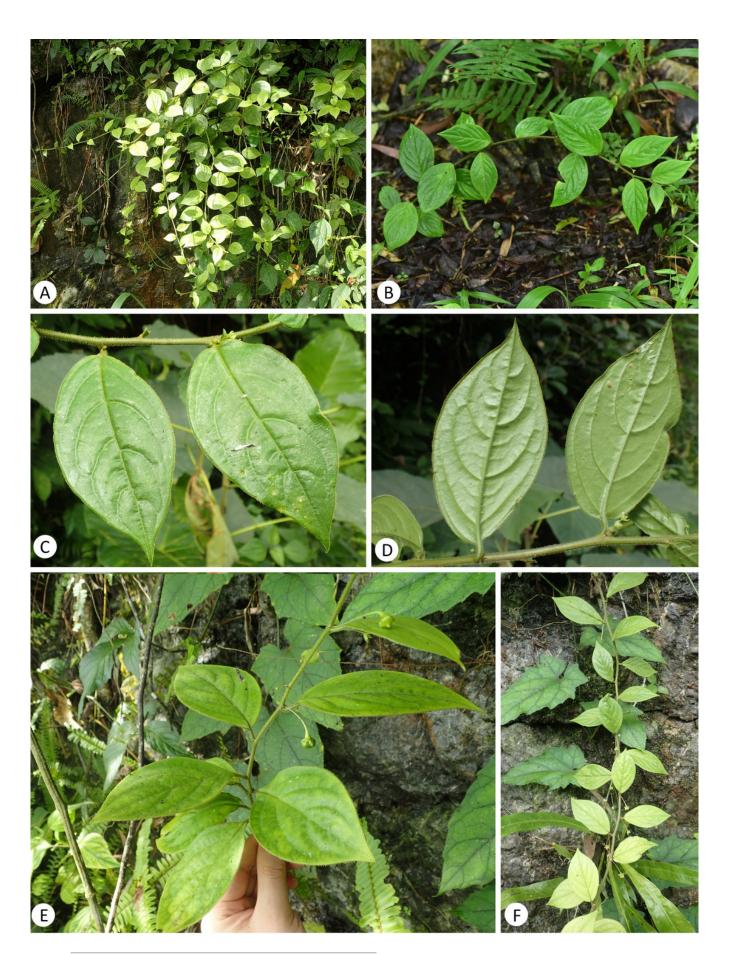
**Fig. 2.** Maximum likelihood (ML) trees of *Leptopus* and its relatives inferred from the combined data set (including nrITS and *matK*). Bootstrap (BS) value in ML analysis and posterior probability (PP) in Bayesian inference (BI) is indicated on the left and right of slanting bar associated with phylogenetic node, respectively. The crown node of *Leptopus* is shown by the arrowhead.





Leptopus malipoensis (from the type locality).

**Fig. 3.** Leptopus malipoensis (from the type locality). (A–B & E–F) habit; (C) adaxial side of leaves; (D) abaxial side of leaves. (Photo credit: Gang Yao).

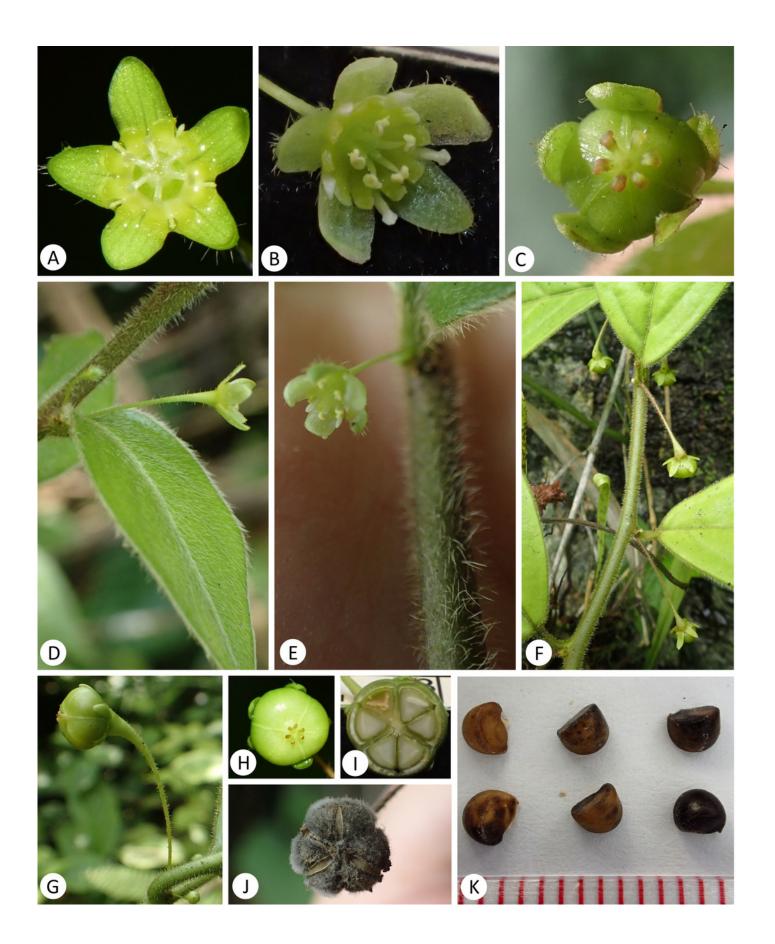


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Leptopus malipoensis (from the type locality).

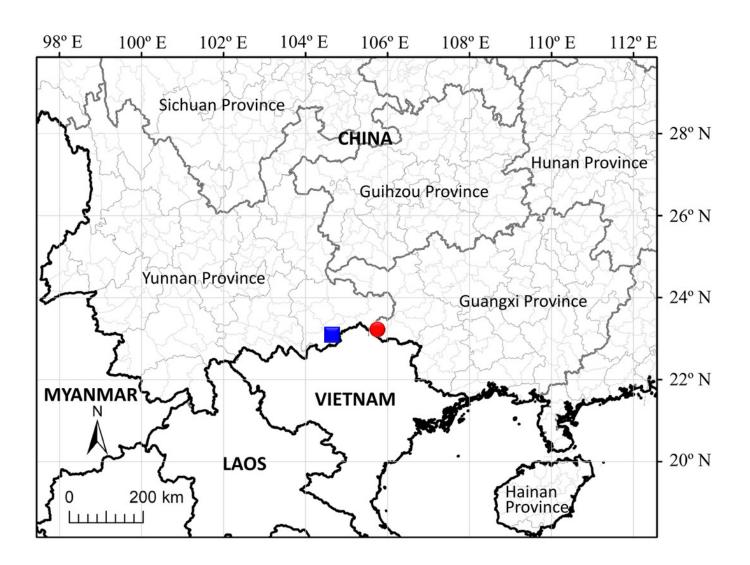
**Fig. 4.** Leptopus malipoensis (from the type locality). (A) female flower; (B) male flower; (C) young fruit; (D) young leaf, stem and female flower with pedicel; (E) stem and male flower with pedicel; (F) stem and young fruits with fruiting pedicels; (G) fruit and fruiting pedicel; (H) fruit; (I) fruit transection; (J) capsule with fungus on its surface; (K) seeds. (Photo credit: Gang Yao and Xin-Xin Zhu).





Distribution of Leptopus malipoensis (blue square) and L. fangdingianus (red circular).

Fig. 5. Distribution of Leptopus malipoensis (blue square) and L. fangdingianus (red circular).





## Table 1(on next page)

Sequences information for all samples used in the present study.

**Table 1.** Sequences information for all samples used in the present study. Sequences newly generated in this study are marked in bold.



1 Table 1. Sequences information for all samples used in the present study. Sequences newly

2 generated in this study are marked in bold.

Taxon	nrITS	matK
Leptopus australis (Zoll. & Moritzi) Pojark.	AM745811	AM745812
Leptopus australis (Zoll. & Moritzi) Pojark.	AM745813	AM745814
Leptopus chinensis (Bunge) Pojark.	MN722097	MN722149
Leptopus chinensis (Bunge) Pojark.	MH710764	MH659095
Leptopus chinensis (Bunge) Pojark.	AM745819	AM745820
Leptopus chinensis (Bunge) Pojark.	AM745821	AM745822
Leptopus clarkei (Hook. f.) Pojark.	AM745938	AM745939
Leptopus clarkei (Hook. f.) Pojark.	AM745940	AM745941
Leptopus cordifolius Decne.	AM745826	AY552433
Leptopus cordifolius Decne.	AM745827	AM745828
Leptopus cordifolius Decne.	AM745829	AY552433
Leptopus fangdingianus (P.T. Li) Voronts. & Petra		
Hoffm.	AM745809	AM745810
Leptopus malipoensis W.H. Zhang & Gang Yao	MW962203	MZ062211
Actephila albidula Gagnep.	AM745910	AM745911
Actephila collinsiae W. Hunter ex Craib	AM745912	AM745913
Actephila sessilifolia Benth.	AM745931	AM745932
Andrachne ephemera M.G. Gilbert	AM745767	AM745768
Andrachne fruticulosa Boiss.	AM745773	AM745774
Andrachne microphylla (Lam.) Baill.	AM745787	AM745788
Andrachne telephioides L.	AM745802	AM745803
Heywoodia lucens Sim	AM745935	AM745937
Meineckia acuminate (Verdc.) J.F. Brunel	AM745894	AM745895
Meineckia humbertii G.L. Webster	AM745846	AM745847
Notoleptopus decaisnei (Benth.) Voronts. & Petra		
Hoffm.	AM745830	AM745831
Notoleptopus decaisnei (Benth.) Voronts. & Petra		
Hoffm.	AM745832	AM745833
Phyllanthopsis arida (Warnock & M.C. Johnst.)		
Voronts. & Petra Hoffm.	AM745762	AM745763
Phyllanthopsis phyllanthoides (Nutt.) Voronts. &		
Petra Hoffm.	AM745836	AM745837
Poranthera corymbosa Brongn.	AM745872	AM745873
Pseudophyllanthus ovalis (E. Mey. ex Sond.) Voronts.		
& Petra Hoffm.	AM745789	AY830260
Pseudophyllanthus ovalis (E. Mey. ex Sond.) Voronts.		
& Petra Hoffm.	AM745790	AM745791
Poranthera triandra J.M. Black	AM745892	AM745893



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