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RESEARCH ARTICLE



Sinosenecio yangii (Asteraceae), a new species from Guizhou, China

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Abstract

A new species *Sinosenecio yangii* D.G. Zhang & Q. Zhou (Asteraceae, Senecioneae) from Guizhou Province, China, is described and illustrated based on its morphological characteristics and molecular evidence. It closely resembles *S. confervifer* and *S. guangxiensis*, the former in the scapigerous habit and smooth and glabrous achene surface, the latter in the calyculate involucre and purple abaxial leaf surface, and both in the shape and indumentum of leaf lamina, but differs markedly from the latter two in having fewer capitula and epappose achenes. Phylogenetic analysis based on nrITS and *ndhC-trnV* sequences shows that this new species belongs to the *S. latouchei* clade and is sister to *S. guangxiensis* with moderate support.

Keywords

molecular evidence, morphology, pappus

Introduction

Sinosenecio B. Nordenstam (1978) (Senecioneae, Asteraceae) contains 45 species mainly distributed in central and southwestern parts of China (Chen et al. 2011; Liu and Yang 2012; Liu et al. 2019; Zou et al. 2020; Chen et al. 2022). This genus is

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characterized by subscapiform or leafy stems, palmately or rarely pinnately veined leaf lamina, solitary to numerous capitula, and ecalyculate or sometimes calyculate involucres, etc. (Jeffrey and Chen 1984). *Sinosenecio* encompasses two species assemblages, i.e. the *Sinosenecio s.s.* group and the *S. oldhamianus* group, with different chromosome number (x = 30 vs. 24 or 13), patterns of endothecial cell wall thickenings (strictly polarized vs. polarized and radial), and phylogenetic affiliation (subtrib. Tussilagininae *s.s.* vs. subtrib. Thephroseridinae) (Liu 2010; Liu and Yang 2011a, b; Gong et al. 2016). These two groups also differ in geographical distribution. The former is restricted to mountainous regions around Sichuan Basin, southwestern China, and the latter is widely distributed in central and southern China, with two species extending to Indochina (Gong et al. 2016). However, a formal taxonomic adjustment is not yet proposed as phylogenetic relationships in subtrib. Thephroseridinae need to be further clarified (Nordenstam and Pelser 2011).

Libo County (Guizhou Province, China) belongs to the slope zone of transition from Guizhou Plateau to Guangxi Hilly Basin with typical karst topography and complex and diverse ecological environment (Tan 2010). In the past few years, some new species have been reported in this area, such as *Strobilanthes hongii* (Chen et al. 2019) and *Petrocodon luteoflorus* (Fan et al. 2020). During our field investigation at Lihua Town, Libo County in March 2021, we found several unusual *Sinosenecio* populations that morphologically resemble two members of the *S. oldhamianus* group, namely *S. confervifer* (H. Léveillé) Y. Liu & Q. E. Yang and *S. guangxiensis* C. Jeffrey & Y. L. Chen, but differs markedly from them in several morphological features, respectively. After examining herbarium specimens and relevant literature, we verified that it represents an undescribed species. Here, we described it as *S. yangii* D. G. Zhang & Q. Zhou with report on its chromosome number and phylogenetic position.

Materials and methods

Morphological observation

Morphological examination and comparison of the new species with *S. confervifer* and *S. guangxiensis* were based on fresh materials and herbarium specimens. Chromosome observation was conducted according to Meng et al. (2010).

Molecular analyses

To test the phylogenetic affiliation of *S. yangii*, we carried out phylogenetic analysis based on combined matrix of ITS and *ndhC-trnV* sequences. The matrix contained 23 accessions from 20 species, including the new species, 16 species of *S. oldhamianus* group, two of *Nemosenecio*, and an outgroup *Tephroseris flammea* (Turcz. ex DC.) Holub. The ITS and *ndhC-trnV* of *S. yangii* were sequenced in this study and the

Region	Name	Primer sequence (5'-3')
nrITS	ITS1	AGAAGTCGTAACAAGGTTTCCGTAGG
	ITS4	TCCTCCGCTTATTGATATGC
ndhC-trnV	ndhCretF	AAGTTTCTCCGGTCCTTTGC
	trnVretR	TCTACGGTTCGAGTCCGTATAG

Table 1. Primers used in this study.

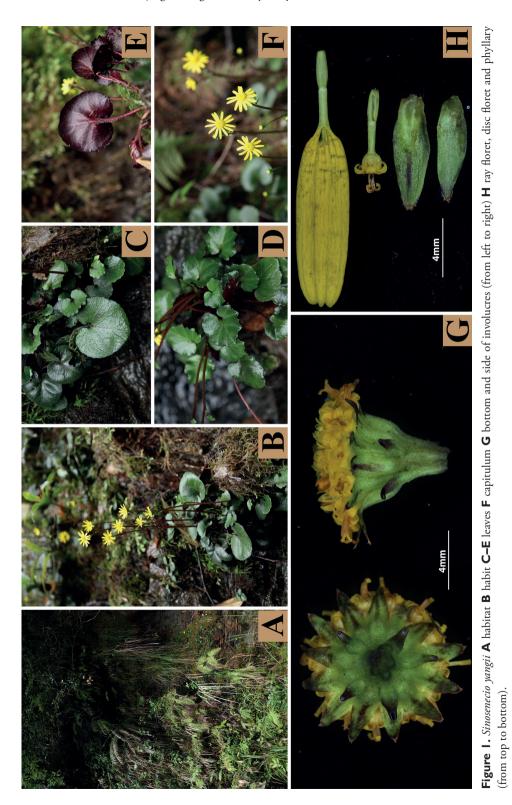
rest were downloaded from GenBank. The GenBank accession numbers are listed in Appendix 1. Total DNA was extracted from dried leaves using Plant Genomic DNA Kit DP305 (Beijing, China) and used as the template for polymerase chain reaction (PCR). The primers used in this study are listed in Table 1. Sequences obtained were edited using Sequencher-5.4.5 and then combined by Sequence Matrix-1.9 (Vaidya et al. 2011). Multi-sequence alignment and manual adjustment were conducted using programme CLUSTAL_W in Mega-X64 (Rédei 2008) and gaps were treated as missing data.

Phylogenetic trees were constructed using Bayesian Inference (BI) and Maximum Likelihood (ML) in CIPRES Portal (https://www.phylo.org/portal2). BI and ML analyses were performed using MrBayes version-3.2 (Ronquist et al. 2012) and RAxML-8.2.10 (Stamatakis 2014), respectively. For BI analysis, GTR+G was selected as best-fitting model using Akaike information criterion (AIC) in JmodelTest 2-2.1.6 (Posada 2008). The Markov chain Monte Carlo analyses were run with four simultaneous chains of 10,000,000 generations sampling one tree every 1,000 generations. After the first 25% of trees were discarded as burn-in, the remaining trees were used to construct a majority-rule consensus tree with Bayesian posterior probabilities. ML analysis was performed with GTRCAT model, support values was calculated with 1,000 bootstrap replicates using a fast bootstrapping algorithm (Stamatakis et al. 2008).

Results

Morphology and taxonomy

Morphological observation (Fig. 1) showed that *S. yangii*, *S. confervifer*, and *S. guangxiensis* share obvious resemblance in the leaf blade shallowly undulate and suborbicular, adaxially densely to sparsely villous and abaxially sparsely pubescent or nearly glabrous (Table 2). In addition, *S. yangii* is similar to *S. confervifer* in the stem leafless or with 1–2 bract-like leaf and smooth achene surface, and to *S. guangxiensis* in the calyculate involucre. Nevertheless, *S. yangii* differs from both species in having fewer capitula (usually 1–3) and epappose achenes. The metaphase chromosomes of this species were counted to be 2n = 48 (Fig. 2A). The achene surface was glabrous and smooth (Fig. 2B) and the anther endothecial cell wall thickenings were polarized and radial (Fig. 2C).



	S. yangii	S. guangxiensis	S. confervifer
Height (cm)	15-25	10-30	10–65
Leaf shape	Suborbicular or reniform,	Suborbicular or reniform, mar-	Orbicular or suborbicular, margin repand
	margin irregularly deltoid or	gin coarsely repand or dentate	or lobed, with rounded or broadly deltoid
	rounded dentate, shallowly	with ovate-deltoid teeth	mucronulate or obscurely mucronulate shallow
	undulate or nearly entire		teeth or lobes
Leaf size (cm)	2.5-4.5 × 2.5-6.5	2-6 × 2.5-7	1.5–6 × 2–6
Adaxial surface of leaf	Green, densely or sparsely	Green or dark green, sparsely	Lustrous, green or deep green, densely or
lamina	pubescent	todensely villous or glabrous	sparsely villous or glabrous
Abaxial surface of leaf	Pale green or purplish red,	Deep purplish red, densely	Pale green or slightly purple with sparsely
lamina	sparsely arachnoid or nearly	white tomentose, sparsely	arachnoid, veins villous or pubescent
	glabrous	villous or glabrescent	
Cauline leaves	1-2, bract-like	1-5, similar to radical ones	1–2, bract-like
Petiole base of cauline	Expanded, not auriculate	Slightly expanded, not	Expanded, not auriculate
leaves		auriculate	
Number of capitula	Usually1, sometimes 2 or 3	2–7 or more, rarely 1	1-7 (-10) or more
Involucre	Calyculate	Calyculate	Not calyculate
Phyllaries	13	13	13
Chromosome number $2x$	48	48	48
Achene surface	Smooth, glabrous	Papillate, pubescent	Smooth, glabrous
Pappus	Absent	Present	Present
Geographical distribution	Guizhou	Guangxi, southwestern Hunan	Hunan, Sichuan, Chongqing, Guizhou, Yunnan

Table 2. Comparison of morphological characteristics among Sinosenecio yangii, S. guangxiensisand S. confervifer.

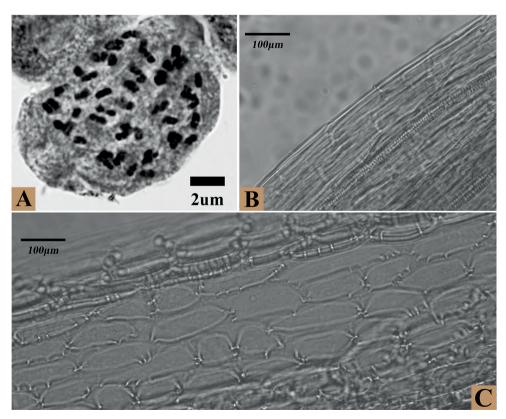


Figure 2. *Sinosenecio yangii* **A** metaphase chromosomes (2x = 48) **B** smooth and glabrous achene surface **C** polarized and radial endothecial cell wall thickenings.

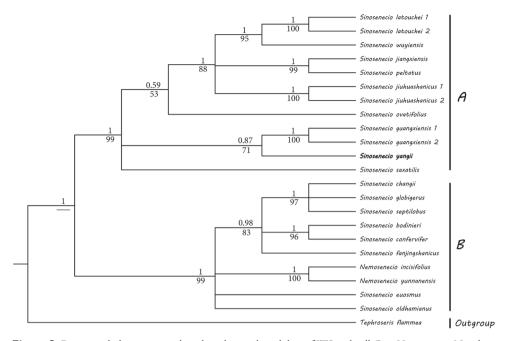


Figure 3. Bayesian phylogenetic tree based on the combined data of ITS and *ndhC-trnVs* equences. Numbers above and below branches are Bayesian posterior probabilities and ML bootstrap values, respectively. A and B represent the *S. latouchei* clade and *S. oldhamianus-Nemosenecio* clade. *Sinosenecio yangii* is noted in bold.

Phylogenetic analyses

The combined matrix of ITS and *ndhC-trnV* sequences contained 1,324 aligned bp. Bayesian (BI) and Maximum likelihood (ML) trees had similar topologies. The BI tree was presented in Fig. 3 with BI posterior probability (BP) and ML bootstrap support values (LP) labelled on the branches. Ingroups were resolved into two clades, viz. the *S. latouchei* clade (BP = 1, LP = 99) and the *S. oldhamianus-Nemosenecio* clade (BP = 1, LP = 99). *Sinosenecio yangii* was resolved as sister to *S. guangxiensis* (BP = 0.87, LP = 71) in the former clade, while *S. confervifer* was recovered as a member in the latter clade.

Discussion

Several lines of evidence demonstrated that *S. yangii* is a member of the *S. oldhamianus* group. *Sinosenecio yangii* has a base chromosome number of x = 24 (Fig. 2A) and polarized and radial anther endothecial cell wall thickenings (Fig. 2C), which are typical of the *S. oldhaminanus* group. Analyses of ITS and *ndhC-trnV* sequences also corroborated its phylogenetic affiliation, resolving it as sister to *S. guangxiensis* in the *S. latouchei* clade of *S. oldhaminanus* group.

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Sinosenecio yangii was morphologically and phylogenetically close to S. guangxiensis in the S. latouchei clade. However, there are differences in morphology, distribution and ecology between the two species. S. yangii is easily distinguished from S. guangxiensis in the stem leafless or with 1-2 bract-like leaf (vs. 1-5 cauline leaves), fewer capitula (vs. 2-7 or more), smooth and glabrous achene surface (vs. papillate and pubescent), and epappose achenes (vs. present pappus). From the perspective of distribution area, the former is restricted to Libo County in Guizhou, appearing on the wet rock cliff, and the geographical location is adjacent to the border with Guangxi province. The latter is distributed in the Guangxi and southwestern Hunan, growing on the damp, shady places or rocky places at mountain summits. To some extent, the close relationship between these two species may also be related to their distributional ranges adjacent to each other. Additionally, it is worth noting that the epappose achenes of S. yangii is a character previously never recorded in the S. latouchei clade.

Taxonomic treatment

Sinosenecio yangii D. G. Zhang & Q. Zhou, sp. nov. urn:lsid:ipni.org:names:77305747-1

Figs 4, 5

Type. CHINA. Guizhou: Libo County, Lihua Town, 25°36'53"N, 108°12'63"E, on rock cliff by the side of a rural road, elev. 347 m, 16 March 2021, D. G. Zhang & T. Deng 14231. (holotype: JIU!; isotype: JIU!).

Description. Scapigerous herbs. Rhizomes short and stout with many fibrous roots. Stems slender, scapiform, erect or declining, solitary or several, 13-22 cm long, basally reddish-brown and sparsely white villous, almost smooth in upper part. Radical leaves several; petiole ca. 3-6.5 cm long, densely villous or glabrescent, basally expanded, not auriculate; lamina suborbicular or reniform, ca. $2.5-4.5 \times 2.5-6.5$ cm, base cordate, margin irregularly triangular dentate, shallowly undulate or entire, apex slightly acute; adaxially green, densely or sparsely pubescent, abaxially pale green or purplish red, sparsely arachnoid or nearly glabrous. Upper leaves 1 or 2, bract-like, shortly petiolate, lanceolate. Capitula usually 1-3, peduncles slender, ca. 2-3.5 cm long, with a basal linear bracteole, or with 1-2 small linear bracteoles in the upper part. Involucres campanulate, calyculate with 2-3 bracteoles or more; phyllaries ca. 13, lanceolate, ca. 6 mm long, with ciliate margin, apically acute or obtuse and sometimes purplish. Ray florets ca. 13, corolla tube 3 mm long, glabrous; ray yellow, oblong, ca. 12 mm long, 4-veined, apically 3-denticulate. Disc florets numerous; corolla yellow, 4 mm, with ca. 1.5 mm glabrous tube and 0.85 mm limb. Anthers oblong, 5, ca. 1.2 mm long, basally obtuse. Style branches ca. 0.5 mm long, puberulent. Achenes ca. 1 mm long, smooth and glabrous. Pappus absent.

Phenology. Flowering from March to May, fruiting from April to June.



Figure 4. Holotype sheet of Sinosenecio yangii D. G. Zhang & Q. Zhou.

Etymology. The species was named after Professor Qin-er Yang, an expert in the field of *Asteraceae* at the Chinese Academy of Sciences. The Chinese name is given as " 亲二蒲儿根" (qīn èr pú ér gēn).

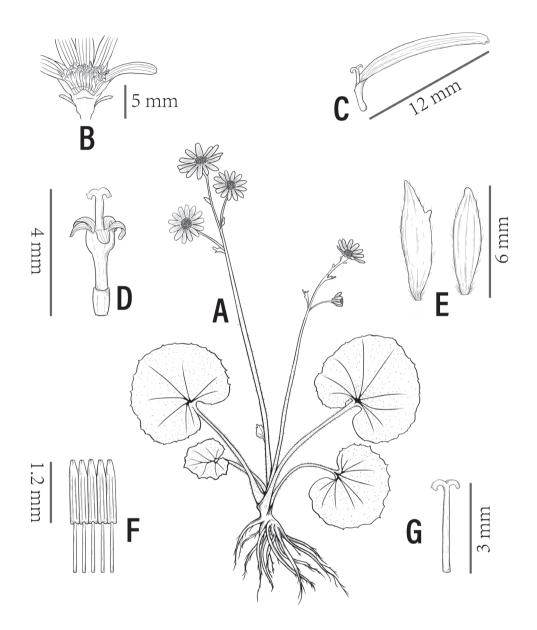


Figure 5. *Sinosenecio yangii* **A** habit **B** capitulum **C** ray floret **D** disk floret **E** phyllary **F** stamens **G** style (drawing by Chu-miao Xie).

Distribution and habitat. *Sinosenecio yangii* is known from Lihua Town, Libo County, Guizhou Province, China (Fig. 6). It was collected from a rock cliff by the side of a rural road in this town, at an altitude of 347 m.

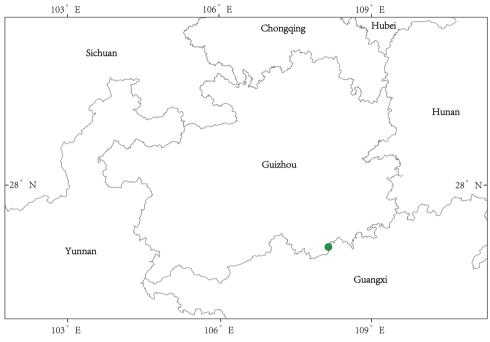


Figure 6. Distribution of *Sinosenecio yangii*.

Key to species of the S. latouchei clade

1	Pappus absent
_	Pappus present
2	Leaf lamina peltate
_	Leaf lamina not peltate
3	Involucres calyculate
_	Involucres ecalyculate
4	Cauline leaf absent or 1 and bract-like; base of petiole of cauline leaf slightly
	auriculate; capitula solitary, rarely 2 or 3
_	Cauline leaves 1-5, similar to radical ones; base of petiole of cauline leaves
	never auriculate; capitula 1–5 or moreS. guangxiensis
5	Ovaries and achenes glabrous
_	Ovaries and achenes pubescent7
6	Leaf lamina broadly flabellate or suborbicular, dentate or palmately lobed to
	1/2, lobes apically 2 or 3-denticulate, both surfaces glabrous S. wuyiensis
_	Leaf lamina reniform or suborbicular, regularly 5-7-palmatilobed, lobes
	ovate-triangular, both surfaces glabrous or sometimes white tomentose abaxi-
	ally and later glabrescent
_	Leaf lamina ovate, broadly ovate, rarely ovate-orbicular, inconspicuously
	undulate-dentate, adaxial surface villous, sometimes sparsely arachnoid, and
	abaxial surface villous and densely white arachnoid

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Appendix I

Species	GenBank accession number of ITS / ndhC-trnV		
Tephroseris flammea	KU696137 / KU750769		
Nemosenecio yunnanensis	KU696047 / KU750695		
Nemosenecio incisifolius	KU696045 / KU750694		
Sinosenecio latouchei	JQ797428 / KU750748		
Sinosenecio latouchei	JQ797429 / KU750749		
Sinosenecio wuyiensis	JQ797431 / KU750764		
Sinosenecio jiangxiensis	KT149879 / KU750743		
Sinosenecio peltatus	MK818500 /		
Sinosenecio jiuhuashanicus	JQ797426 / KU750746		
Sinosenecio jiuhuashanicus	JQ797425 / KU750745		
Sinosenecio ovatifolius	MT522620 / -		
Sinosenecio guangxiensis	JQ797432 / KU750738		
Sinosenecio guangxiensis	JF978599 / KU750739		
Sinosenecio yangii	OM413747 / OM371331		
Sinosenecio saxatilis	JQ797430 / KU750757		
Sinosenecio changii	AY176164 / KU750721		
Sinosenecio globigerus	AY176159 / KU750736		
Sinosenecio septilobus	AY176161 / KU750758		
Sinosenecio bodinieri	KT149888 / KU750720		
Sinosenecio confervifer	KT149891 / KU750723		
Sinosenecio fanjingshanicus	KT149886 / KU750732		
Sinosenecio euosmus	JF978589 / KU750730		
Sinosenecio oldhamianus	JF978616 / KU750753		

Table A1. GenBank accessions of species used in this study.