

# Paraphlomis yingdeensis (Lamiaceae), a new species from Guangdong (China)

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Academic editor: E. Fischer | Received 15 November 2022 | Accepted 11 January 2023 | Published 8 February 2023

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**Citation:** Guo G-X, Zhao W-Y, Chen Y-P, Xiao J-H, Li Y-Q, Fan Q (2023) *Paraphlomis yingdeensis* (Lamiaceae), a new species from Guangdong (China). *PhytoKeys* 219: 107–120. <https://doi.org/10.3897/phytokeys.219.97547>

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## Abstract

*Paraphlomis yingdeensis* (Lamiaceae), a new species from the limestone area in northern Guangdong Province, China, is described and illustrated. Phylogenetic analyses, based on two nuclear DNA regions (ITS and ETS) and three plastid DNA regions (*rpl32-trnL*, *rps16* and *trnL-trnF*), suggest that *P. yingdeensis* represents a distinct species in *Paraphlomis*. Morphologically, *P. yingdeensis* is similar to *P. foliata* subsp. *montigena* and *P. nana*, but can be distinguished from the former by its densely villous lamina and calyx, not decurrent base of lamina and bristle-like-acuminate apex of calyx teeth, and distinguished from the latter by its significantly taller plant (15–20 cm vs. 1–5 cm) and larger lamina (6.2–16.5 × 4–11.5 vs. 2–7 × 1.5–4 cm), densely villous stem, lamina and calyx and yellow corolla.

## Keywords

endemics, limestone, new taxon, Paraphlomideae, phylogeny

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\* These authors contributed equally to this work.

## Introduction

As a member of tribe Paraphlomideae (Lamiaceae, Lamioideae) (Bendiksby et al. 2011; Zhao et al. 2021), the genus *Paraphlomis* Prain is characterised by its herbaceous habit, actinomorphic calyx with five lobes less than half as long as the tube, corolla 2-lipped (1/3) with hairy upper lip, but hardly bearded along the margin, included stamens and an apically truncate ovary (Wu and Li 1977; Bendiksby et al. 2011; Chen et al. 2021). A total of 36 species and seven varieties are recognised within *Paraphlomis*, most of which are distributed in southern China (Chen et al. 2022b; Yuan et al. 2022), with several species occurring in the Himalayas, Korea and Southeast Asia (Li and Hedge 1994; Ko et al. 2014; Chen et al. 2021). Many species of *Paraphlomis* are endemics of limestone soils, including the recently described *P. kuankuoshuiensis* R.B. Zhang, D. Tan & C.B. Ma (Zhang et al. 2020), *P. longicalyx* Y.P. Chen & C.L. Xiang (Chen et al. 2022a) and *P. hsiwenii* Y.P. Chen & X. Li (Chen et al. 2022b). This shows species richness of *Paraphlomis* has been quite underrated and more field investigations are needed to infer its diversity in limestone areas.

The botanical expedition to the Shimentai National Nature Reserve in Guangdong Province, China in October 2021, showed an unknown species of *Paraphlomis*. Based on other field observations (from May to August in 2022), morphological comparisons with congeneric species, as well as molecular phylogenetic studies, we confirmed that it represented a new species, here described and illustrated.

## Materials and methods

### Morphological study

Field observations and collections of the new species were carried out from May to August in 2022 in Boluo Town of Yingde City in northern Guangdong Province, China. Morphological comparisons of the putative new species with other *Paraphlomis* species were conducted firstly by consulting relevant taxonomic literature, included “Flora of China” (Li and Hedge 1994), “Flora of Guangdong” (Luo 1995) and other recently described species and infraspecies of *Paraphlomis* (Yan and Fang 2009; Ding et al. 2019; Zhang et al. 2020; Chen et al. 2021, 2022a, b, c; Zhao et al. 2022). We also carried out a check of herbarium specimens deposited in LBG, AU, IBK, FJFC, PE, ANUB, KUN, FJSI and SYS (herbarium acronyms following Thiers 2022). All morphological characters were measured using dissecting microscopes.

### Phylogenetic analyses

Previous molecular phylogenetic study revealed genus *Paraphlomis* is not monophyletic, because species of *Matsumurella* were recovered within it (Chen et al. 2021; Chen et al. 2022b). Thus, *Matsumurella* was also included in our phylogenetic analyses. A total of 37 accessions, representing 20 species and four varieties/subspecies of *Para-*

*phlomis* and two *Matsumurella* species were selected as ingroups. One species each of *Phlomis* L. and *Phlomoides* Moench were included as outgroups following Chen et al. (2022a, b). Except for the three accessions of the new species that were newly sampled here, sequences of the remaining accessions were all retrieved from our previous studies (Chen et al. 2021, 2022a, b, c). Genomic DNA of the potential new species was extracted from silica-gel-dried leaves using the modified 2× CTAB procedure of Doyle and Doyle (1987). We selected five DNA markers for the phylogenetic reconstruction, including two nuclear ribosomal regions (internal and external transcribed spacers, i.e. ITS and ETS) and three plastid DNA regions (*rpl32-trnL*, *rps16* and *trnL-trnF*). Primers used for the polymerase chain reaction (PCR) amplification and sequencing were the same as those of Chen et al. (2021), while PCR procedures followed those described in Chen et al. (2016). The specimen information of samples and GenBank accession numbers for all sequences are listed in Appendix 1.

Raw sequences were assembled and edited using Sequencher 4.1.4 (Gene Codes, Ann Arbor, MI, USA) and then aligned using MUSCLE (Edgar 2004) and manually adjusted in MEGA 6.0 (Tamura et al. 2013). Bayesian Inference (BI) (Ronquist et al. 2012) and Maximum Likelihood (ML) (Stamatakis 2014) analyses were used for phylogenetic reconstruction and detailed settings for the two analyses followed those described in Chen et al. (2021). The resulting trees with posterior probabilities (PP) and Bootstrap support (BS) values were visualised and annotated in TreeGraph 2 (Stöver and Müller 2010). The combined nuclear dataset and the combined plastid dataset were initially analysed separately. Topological incongruence between the two reconstructions was visually inspected, based on the thresholds of  $PP \geq 0.95$  and/or  $BS \geq 70\%$ . After excluding the taxa that exhibited strong conflicts between the nuclear tree and the plastid tree, the combined nuclear dataset and the combined plastid dataset were then concatenated for phylogenetic analyses.

## Results and discussion

The aligned length of the combined nuclear dataset was 1254 bp (810 bp for ITS, 444 bp for ETS) and that of the combined plastid dataset was 2479 bp (850 bp for *rpl32-trnL*, 812 bp for *rps16*, 817 bp for *trnL-trnF*). Since the placements of three taxa, *Paraphlomis albiflora* (Hemsl.) Hand.-Mazz., *P. nana* Y.P. Chen, C. Xiong & C.L. Xiang and *P. javanica* var. *pteropoda* D. Fang & K.J. Yan, showed hard incongruences in the nuclear tree (Appendix 2) and the plastid tree (Appendix 3), these taxa were excluded prior to the combination of the nuclear and plastid datasets. All the resulting trees (Fig. 1; Appendices 2–3) were topologically consistent with those in previous study (Chen et al. 2021). With the two species of *Matsumurella* deeply nested within *Paraphlomis*, both genera were shown to be non-monophyletic. The three individuals of the putative new species formed a strongly supported clade (Fig. 1:  $PP = 1.00 / BS = 100\%$ ), but its relationship with other species of *Matsumurella-Paraphlomis* was not resolved.

Our morphological study revealed that the new species *P. yingdeensis* W.Y.Zhao, Y.Q.Li & Q.Fan is most similar to *P. foliata* subsp. *montigena* X.H. Guo & S.B. Zhou

and *P. nana* for some morphological characters as they have short habits and triangular-lanceolate calyx teeth with apices acuminate or bristle-like-acuminate. *Paraphlomis foliata* subsp. *montigena* was classified by Guo (1993) as a subspecies of *P. foliata* (Dunn) C.Y. Wu & H.W. Li. However, previous molecular phylogenetic studies (Chen et al. 2022b, c) and our present analyses (Fig. 1; Appendices 2–3) indicated that *P. foliata* subsp. *montigena* might represent an independent species within the genus as it is distantly related to *P. foliata* subsp. *foliata*. The new species can be distinguished from *P. foliata* subsp. *montigena* in the morphology and indumentum of laminae and calyces. Both the laminae and calyces are densely villous in *P. yingdeensis*, but are sparsely strigose in *P. foliata* subsp. *montigena*; the base of lamina is broadly cuneate and not decurrent in the new species, but is cuneate and decurrent in *P. foliata* subsp. *montigena*; *P. yingdeensis* has bristle-like-acuminate apex of calyx teeth, in contrast, the apex of calyx teeth of *P. foliata* subsp. *montigena* is acuminate. The phylogenetic placement of *P. nana* was conflicting in the nuclear tree and plastid tree, but it was consistently sister to *P. albiflora* (Appendices 2–3). Both *P. nana* and *P. yingdeensis* have translucent and membranous calyces with bristle-like-acuminate apex of calyx teeth. The two species mainly differ in the height of plants, size and indumentum of laminae, as well as colour of corollae. Specifically, plants of *P. nana* are 1–5 cm tall, whereas those of *P. yingdeensis* are 10–20 cm tall. The stems and laminae are densely villous in *P. nana*, but are densely strigose in *P. yingdeensis*. Moreover, *P. yingdeensis* has significantly larger laminae than *P. nana* ( $6.2\text{--}16.5 \times 4\text{--}11.5$  cm vs.  $2\text{--}7 \times 1.5\text{--}4$  cm) and the corollae of *P. yingdeensis* are yellow, differing from the white corollae of *P. nana*. Detailed morphological comparisons amongst the three taxa were summarised in Table 1.

Geographically, *P. foliata* subsp. *montigena* is restricted to the Qingliangfeng Nature Reserve at the border area of Zhejiang and Anhui Provinces in eastern China (Guo 1993) and *P. nana* is now only known from Chongqing City in central China (Chen et al. 2022c). Both the two species are not karst-adapted. In contrast, the new species is distributed in the limestone area in Guangdong Province, southern China.

## Taxonomic treatment

### *Paraphlomis yingdeensis* W.Y.Zhao, Y.Q.Li & Q.Fan, sp. nov.

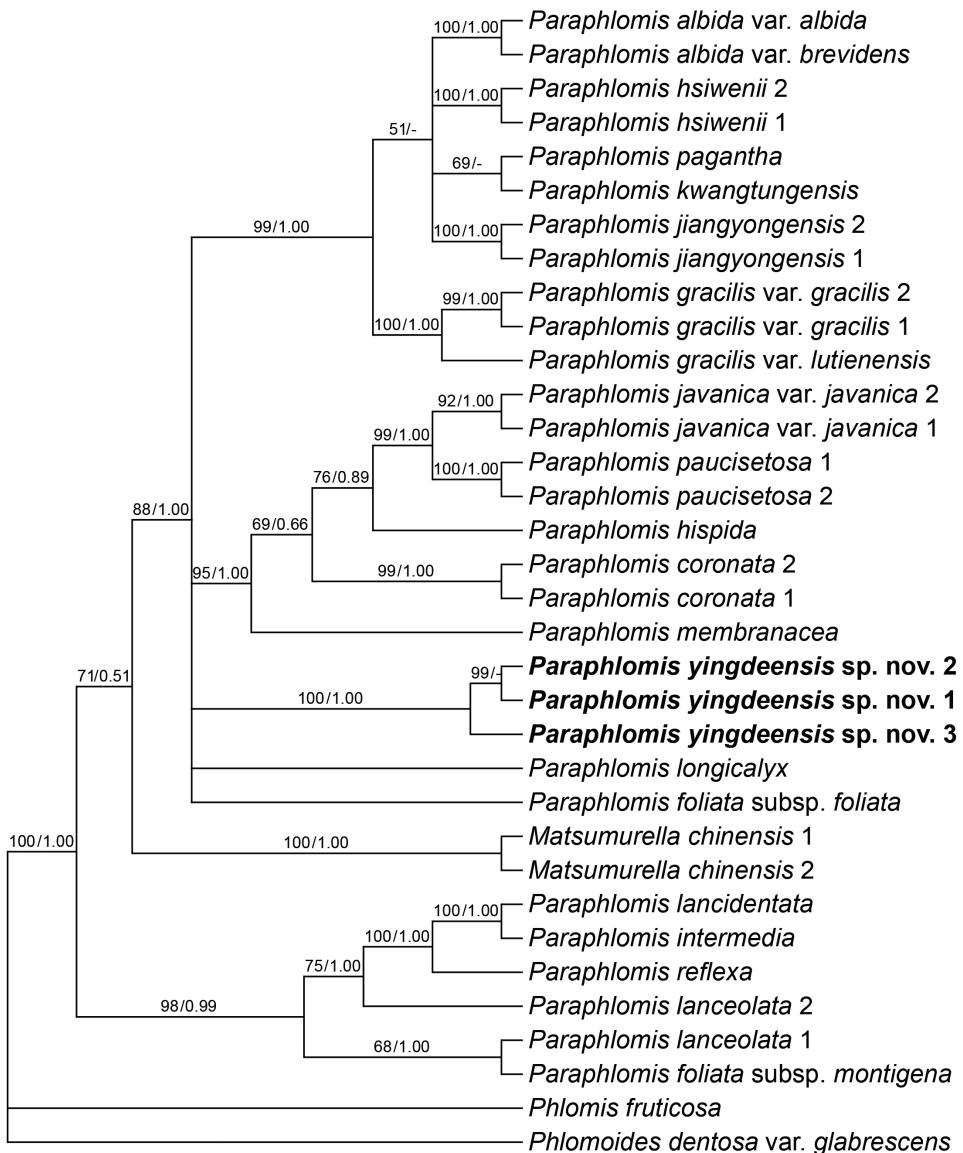
urn:lsid:ipni.org:names:77313387-1

Figs 2–4

Chinese name: 英德假糙苏

**Type.** CHINA. Guangdong Province: Yingde City, Boluo Town, on the way from Xinzhai Village to Changshan Village, on the limestone cliff at the roadside,  $24^{\circ}24'N$ ,  $113^{\circ}0'E$ , alt. 61 m, 29 May 2021, Zhao Wan-Yi, Li Yuan-Qiu, Pan Jia-Wen & Yang Ling-Han ZWY-2092 (holotype: SYS00236856!; isotypes: KUN!, SYS00236857!, SYS00236858!, SYS00236859!).

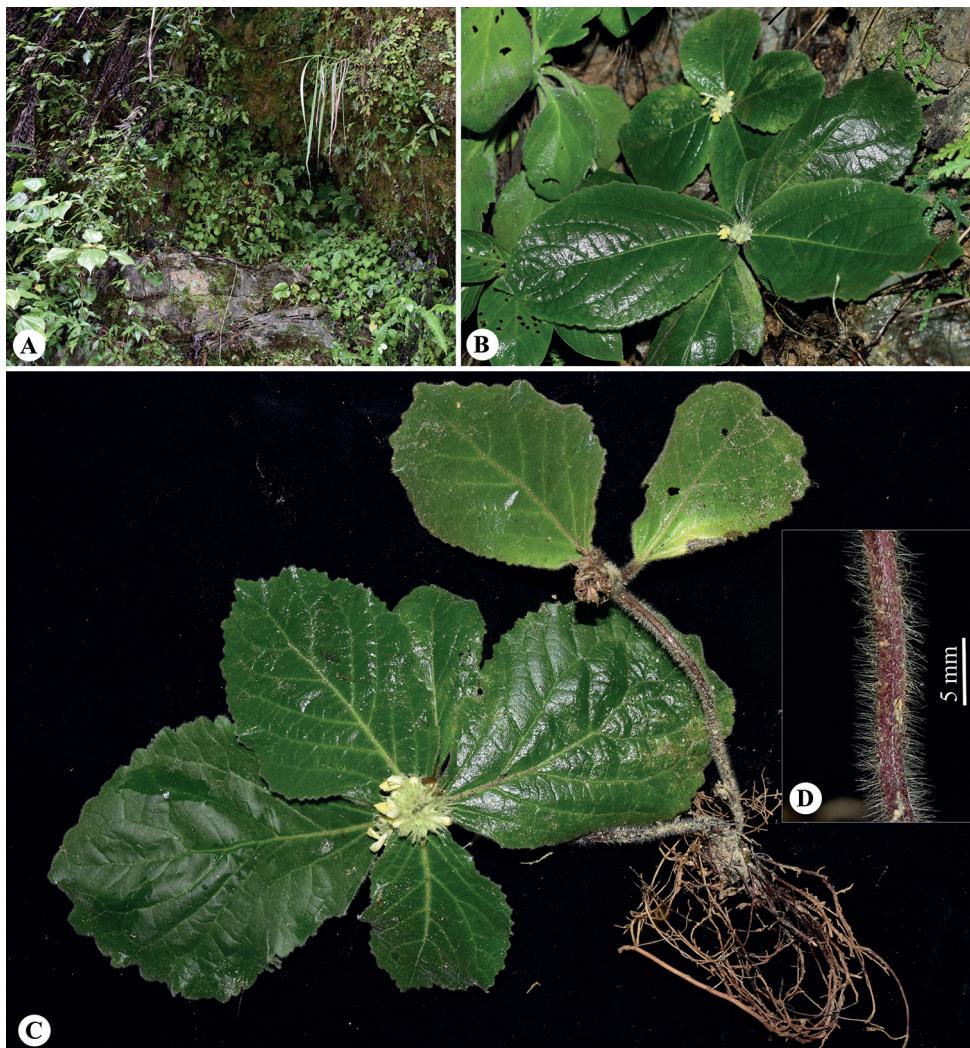
**Diagnosis.** *Paraphlomis yingdeensis* is morphologically similar to *P. foliata* subsp. *montigena* and *P. nana*, but differs from the former in its lamina and calyx densely villous (vs. sparsely strigose), base of lamina not decurrent (vs. decurrent) and apex



**Figure 1.** Optimal Maximum Likelihood tree of *Paraphlomis* inferred from combined nuclear (ETS and ITS) and plastid (*rpl32-trnL*, *rps16* and *trnL-trnF*) dataset. Support value  $\geq 50\%$  BS or 0.50 PP are displayed above the branches ("—" indicates a support value < 0.50 PP).

**Table I.** Morphological comparisons amongst *Paraphlomis yingdeensis*, *P. foliata* subsp. *montigena* and *P. nana*.

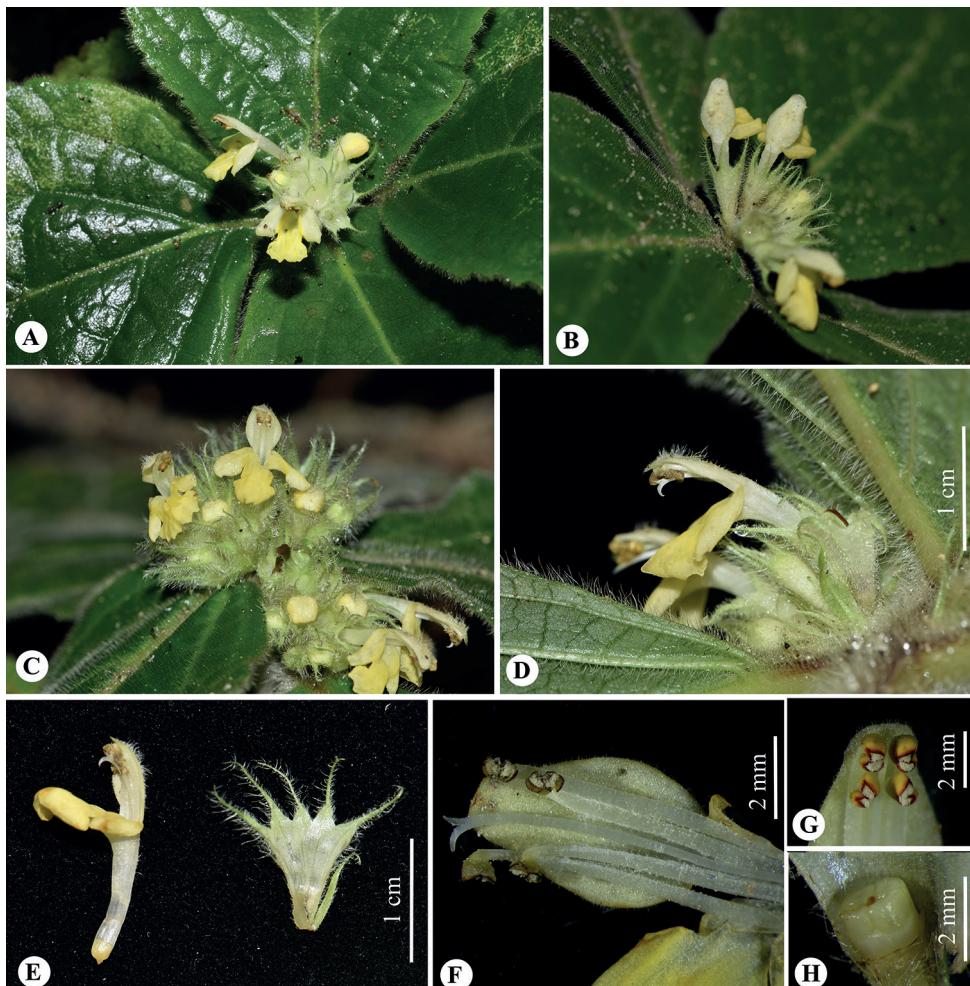
Characters	<i>P. yingdeensis</i>	<i>P. foliata</i> subsp. <i>montigena</i>	<i>P. nana</i>
Stem	10–20 cm tall, densely villous	15–20 cm tall, densely villous	1–5 cm tall, densely retrorse strigose
Lamina	6.2–16.5 $\times$ 4–11.5 cm, base broadly cuneate, not decurrent, densely villous	5–16 $\times$ 2.5–6.5 cm, base cuneate, decurrent, sparsely strigose	2–7 $\times$ 1.5–4 cm, base cuneate to broadly cuneate, decurrent, densely to sparsely strigose
Calyx	Densely villous outside, teeth 3–4 mm long, apex bristle-like-acuminate	Sparingly strigose outside, teeth ca. 2.5 mm long, apex acuminate	Appressed strigose outside, teeth ca. 3 mm long, apex bristle-like-acuminate
Corolla	yellow	yellow	white



**Figure 2.** *Paraphlomis yingdeensis* from the type locality **A** habitat **B**, **C** plants **D** stem. (Photographs: **A**, **C, D** by W.-Y. Zhao; **B** by Y.-Q. Li).

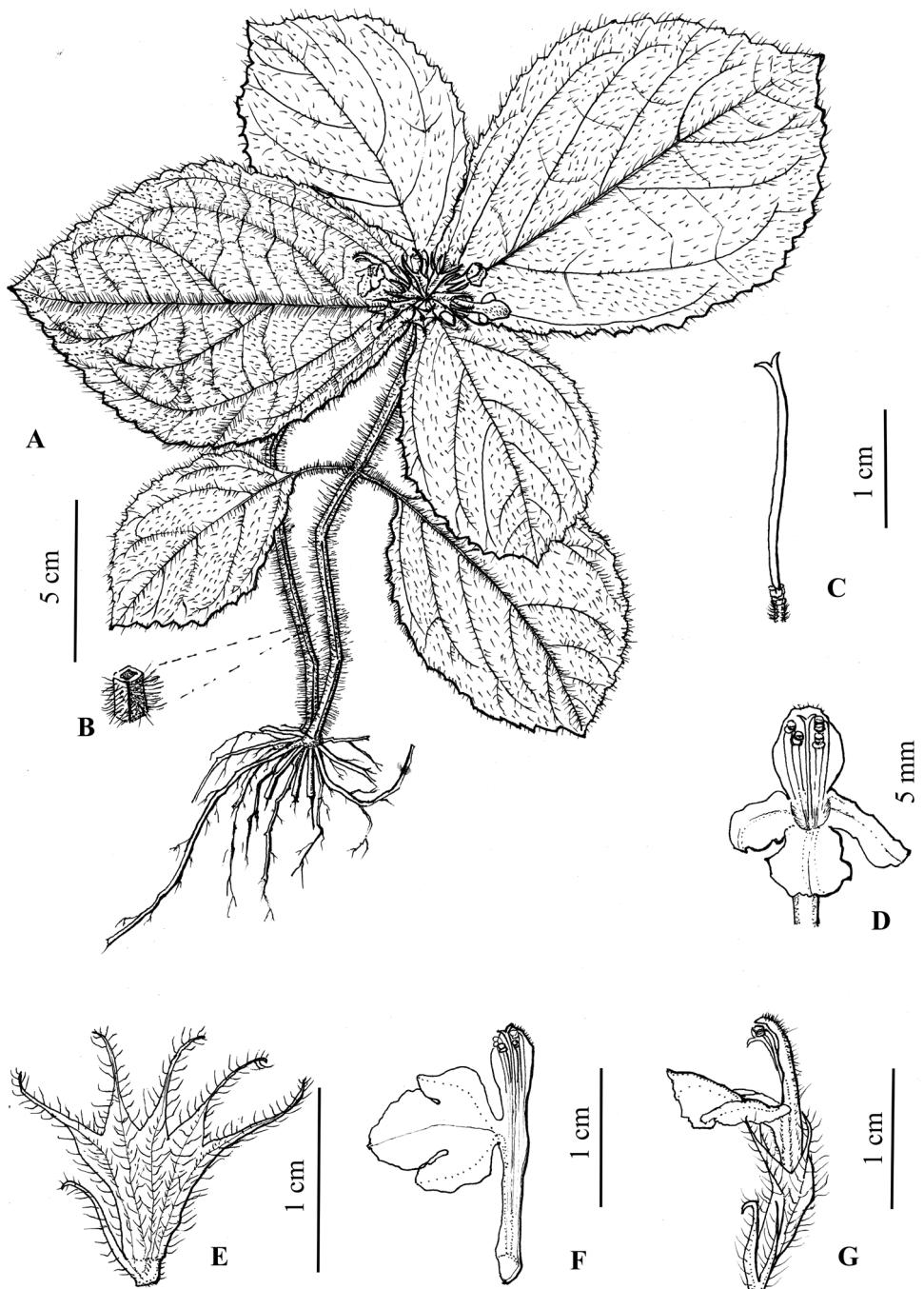
of calyx teeth bristle-like-acuminate (vs. acuminate) and from the latter in its plants 10–20 cm tall (vs. 1–5 cm tall), lamina 6.2–16.5 × 4–11.5 cm and densely villous (vs. 2–7 × 1.5–4 cm and densely strigose) and corolla yellow (vs. white).

**Description.** **Herbs** perennial, 10–20 cm tall. **Rhizomes** short; roots fibrous, yellowish-brown, glabrous. **Stems** erect or prostrate, 4-angled, green (young branch) to purplish-red, densely villous. **Leaves** opposite, leafless towards base, upper two pairs crowded and rosulate; petiole 0.3–2.5 cm long, densely villous; lamina obovate, papery, 6.2–16.5 cm long, 4–11.5 cm wide, apex obtuse, base broadly cuneate, margin crenate-serrate; adaxially green, abaxially light green, densely villous on both sides; lateral veins 5–7-paired, obviously raised abaxially. **Verticillasters** in compact, sometimes capitate-like



**Figure 3.** Floral traits of *Paraphlomis yingdeensis* **A, B** inflorescences **C** frontal view of flower **D** lateral view of flower **E** corolla and dissected calyx (inner view) **F** pistil and stamens **G** anthers **H** ovary. (Photographs: **A, B** by Y.-Q. Li; **C–H** by W.-Y. Zhao).

inflorescences, 8–16-flowered, 2.2–3.0 cm in diam.; bracteoles lanceolate to linear, 7–8 mm long, densely villous. **Calyx** light green, translucent, membranous, campanulate, 6–7 mm long, densely villous outside, glabrous inside, conspicuously 10-veined; teeth 5, subequal, triangular lanceolate, 3–4 mm long, apex bristle-like-acuminate. **Corolla** yellow, 1.5–1.8 cm long; tube 1.0–1.1 cm long, ca. 1.5 mm in diam., straight, pubescent annulate in throat inside; 2-lipped, villous outside, upper lip oblong, margin entire, erect, ca. 6 mm long, ca. 3.5 mm wide; lower lip spreading or reflexed, 4–5 mm long, 3-lobed, medium lobe suborbicular, apex emarginate, lateral lobes oblong, apex obtuse. **Stamens** 4, inserted above middle and upper of corolla tube, straight, included, filaments flat, sparsely puberulent-villous; anther cells 2, ovoid, glabrous. **Style** filiform, included, glabrous, apex subequally 2-lobed. **Ovary** 4-loculed, glabrous. **Nutlets** not seen.



**Figure 4.** Line drawing of *Paraphlomis yingdeensis* **A** plant **B** transverse section of stem **C** pistil **D** frontal view of flower **E** dissected calyx (outside view) **F** dissected corolla **G** lateral view of flower. (Drawn by Rong-En Wu).

**Distribution and habitat.** Currently, only one population of *P. yingdeensis* was found in Boluo Town, Yingde City, in northern Guangdong Province. This town was located in the subtropical monsoon climate region, with development of a large area of karst landform. *Paraphlomis yingdeensis* usually grows on moist limestone cliffs in evergreen broad-leaved forests in association with *Tectaria devexa* Copel., *Primulina yingdeensis* Z.L. Ning, M. Kang & X.Y. Zhuang, *Begonia leprosa* Hance and *Ficus* spp.

**Phenology.** Flowering from May to June and fruiting from June to August.

**Etymology.** The specific epithet “*yingdeensis*” is derived from the type locality of the new species, i.e. Yingde City in Guangdong, China.

**Additional specimens examined (paratypes).** CHINA. Guangdong Province: Yingde City, Boluo Town, on the way from Xinzhai Village to Changshan Village, 24°24'N, 113°0'E, alt. 61 m, 9 June 2021, Q. Fan 19013 (SYS); *ibid.*, 5 June 2022, Li Yuan-Qiu ZWY-2020 (SYS); *ibid.*, 14 August 2022, Ye Fan ZWY-2032 (SYS).

**Specimens of *P. foliata* subsp. *montigena* examined.** CHINA. Anhui Province: Xi County, Qingliangfeng, alt. 1300 m, 29 October 1980, Guo Xin-Hu 800023 (ANUB 13030926); *ibid.*, 16 July 1989, Guo Xin-Hu & Zhou Shou-Biao 89011 (KUN 778733).

**Specimens of *P. nana* examined.** CHINA. Chongqing: Chongkou County, Mingzhong Town, Jinchi Village, Longmenxi, Dabashan National Natural Reserve, on the moist cliff, alt. 996 m, 7 July 2021, Chi Xiong XC21097 (holotype: KUN; isotypes: CQNM, IBK); Wushan County, Zhuxian Town, Shizhuzi Village, Daguling, Wulipo National Natural Reserve, in the moist valley, alt. 1310 m, 18 July 2021, Chi Xiong & Hou-Lin Zhou XC21126 (KUN); *ibid.*, 11 September 2021, Hou-Lin Zhou s.n. (KUN).

## Acknowledgements

We thank Fan Ye and Ling-Han Yang for their help in the fieldwork, and Rong-En Wu for the line drawing. This work was supported by the Guangdong Provincial Special Research Grant for the Creation of National Parks (2021GJGY034).

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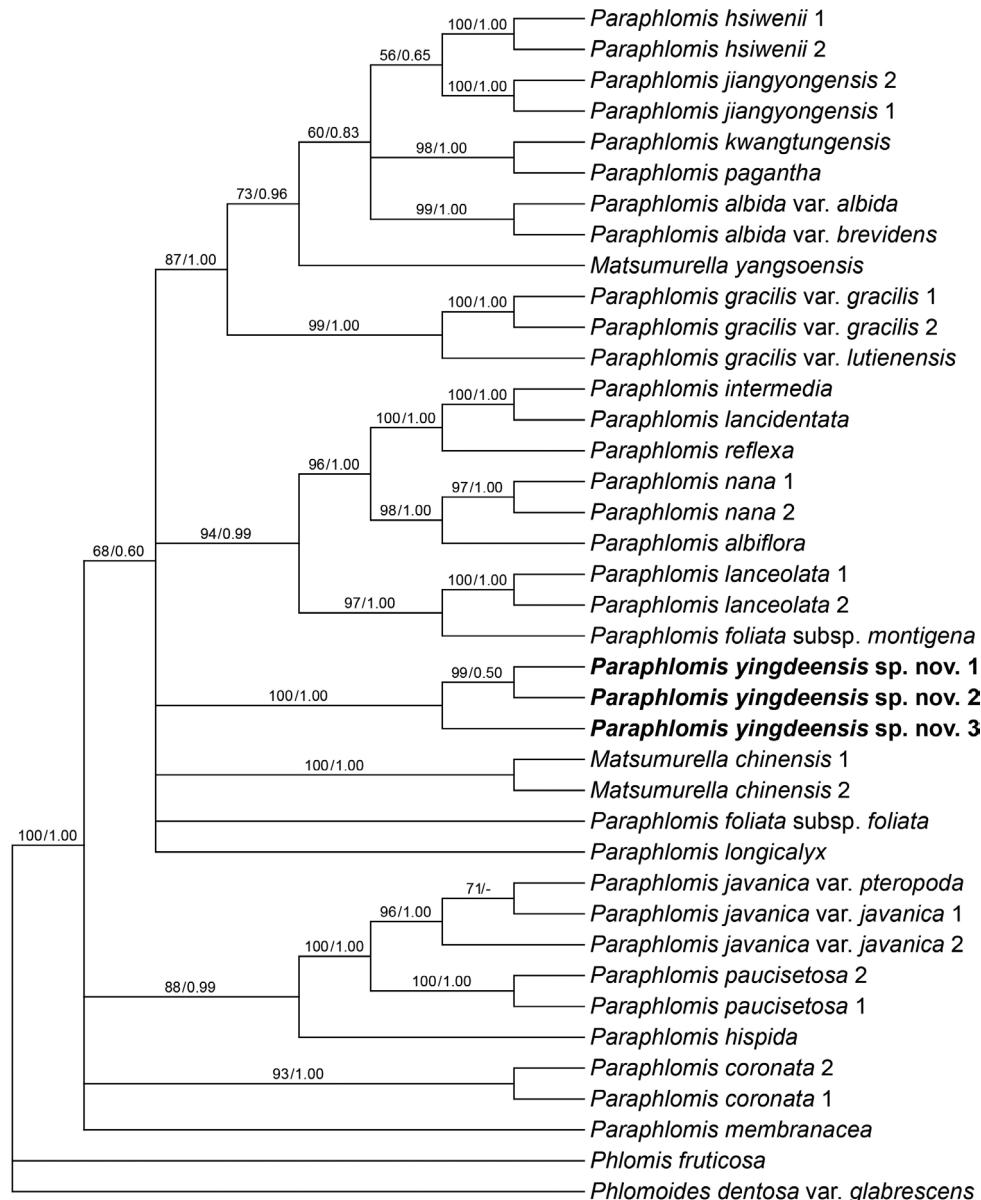
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## Appendix I

**Table A1.** Sequence information for all samples used in present study. A “/” indicates a missing sequence. Herbarium abbreviations are listed after the vouchers.  
The accession numbers marked in bold represent sequences newly generated.

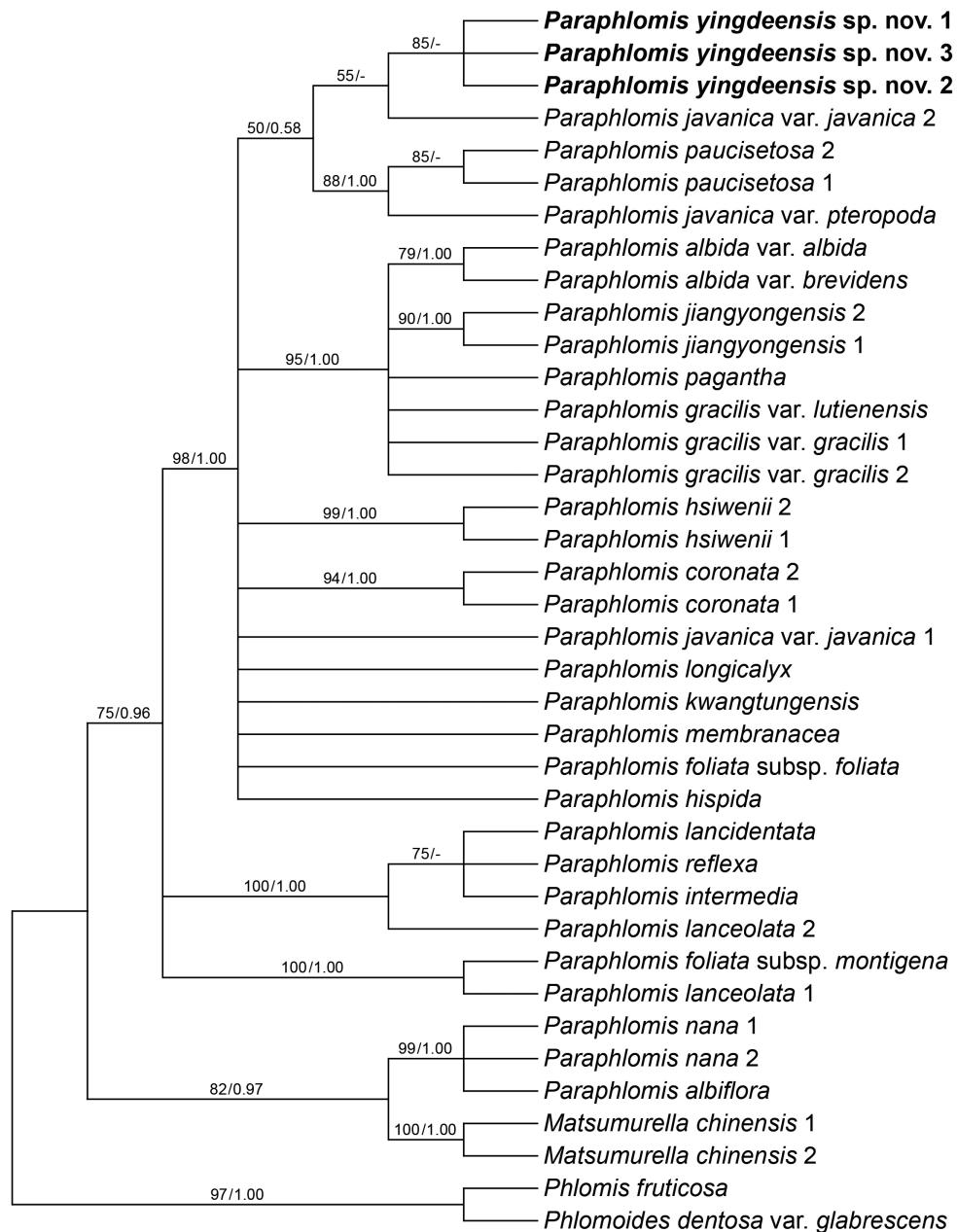
Taxon	Voucher	Country	ITS	ETS	<i>rib32-trnL</i>	<i>rpS16</i>	<i>trnL-trnF</i>
<i>Matumurella chinensis</i> (Benth.) Bendiksby 1	Y. Yang OYY00316 (KUN)	Pingxiang, Jiangxi, China	MW602147	MW602117	MW602021	MW602053	MW602084
<i>Matumurella chinensis</i> (Benth.) Bendiksby 2	Y. Yang OYY0031 (KUN)	Gulin, Guangxi, China	MW602148	MW602118	MW602022	MW602054	MW602085
<i>Matumurella yanggeensis</i> (Y.Z. Sun) Bendiksby	L. Wu & W.B. Xu 10965 (IBK)	Yangshuo, Guangxi, China	MW602142	MW602112	/	/	/
<i>Paraphlomis albida</i> Hand.-Mazz. var. <i>albida</i>	A. Liu et al. LK0841 (CSFI)	Ningyuan, Hunan, China	MW602124	MW602101	MW601996	MW602028	MW602060
<i>Paraphlomis albida</i> var. <i>brevirena</i> Hand.-Mazz.	Y.P. Chen EM312 (KUN)	Hezhou, Guangxi, China	MW602130	MW602098	MW602003	MW602035	MW602067
<i>Paraphlomis albiflora</i> (Hemsl.) Hand.-Mazz.	C.M. Tran et al. 1806393 (JPF)	Jiujiang, Jiangxi, China	/	MW602101	MW602006	MW602038	MW602069
<i>Paraphlomis ornata</i> (Vaniot) Y.P. Chen & C.L. Xiang 1	E.D. Liu et al. 3043 (KUN)	Enmeishan, Sichuan, China	MW602137	MW602107	MW602012	MW602044	MW602075
<i>Paraphlomis cornuta</i> (Vaniot) Y.P. Chen & C.L. Xiang 2	C.L. Xiang XCL135 (KUN)	Jiangkou, Guizhou, China	MW602123	MW602090	MW601995	MW602027	MW602059
<i>Paraphlomis foliata</i> (Dunn) C.Y. Wu & H.W. Li subsp. <i>foliata</i>	S.P. Chen s.n. (KUN)	Jiangle, Fujian, China	/	MW602097	MW602002	MW602034	MW602066
<i>Paraphlomis foliata</i> subsp. <i>montigena</i> X.H. Guo & S.B. Zhou	Y.C. Dai s.n. (KUN)	Hangzhou, Zhejiang, China	OMR84664	OMR84453	OMR84456	OMR84459	OMR84462
<i>Paraphlomis gracilis</i> (Hemsl.) Kudo var. <i>gracilis</i> 1	A.J.in LK0931 (CSFI)	Changsha, Hunan, China	MW602134	MW602104	MW602009	MW602041	MW602072
<i>Paraphlomis gracilis</i> (Hemsl.) Kudo var. <i>gracilis</i> 2	C.L. Xiang XCL1315 (KUN)	Chongqing, China	MW602141	MW602111	MW602016	MW602048	MW602079
<i>Paraphlomis gracilis</i> var. <i>indutinensis</i> (Y.Z. Sun) C.Y. Wu	C.L. Xiang XCL1881 (KUN)	Shibing, Guizhou, China	MW602131	MW602099	MW602004	MW602036	MW602068
<i>Paraphlomis hispida</i> C.Y. Wu	X. Li LX200702 (GXF)	Napo, Guangxi, China	MW602132	MW602102	MW602007	MW602039	MW602070
<i>Paraphlomis hispina</i> Y.P. Chen & Xiong Li 1	W.H. Wu et al. D0426 (KUN)	Jingxi, Guangxi, China	OP605346	OP609841	OP609848	OP609855	OP609862
<i>Paraphlomis hispina</i> Y.P. Chen & Xiong Li 2	W.H. Wu et al. D0426 (KUN)	Jingxi, Guangxi, China	OP605347	OP609842	OP609849	OP609863	OP609863
<i>Paraphlomis intermedia</i> C.Y. Wu & H.W. Li	X. Zhong et al. ZX16823 (CSH)	Suichang, Zhejiang, China	MW602135	MW602105	MW602010	MW602042	MW602073
<i>Paraphlomis javanica</i> (Blume) Prain var. <i>javanica</i> 1	Y.P. Chen s.n. (KUN)	Kunming, Yunnan, China	MW602121	MW602088	MW601993	MW602025	MW602057
<i>Paraphlomis javanica</i> (Blume) Prain var. <i>javanica</i> 2	L.B. Liu et al. JLB0029 (KUN)	Magnan, Yunnan, China	MW602133	MW602113	MW602017	MW602049	MW602080
<i>Paraphlomis javanica</i> var. <i>perforata</i> D. Fang & K. Yan	X. Li 202009501 (GXF)	Jingxi, Guangxi, China	MW602140	MW602100	MW602047	MW602078	MW602081
<i>Paraphlomis javanica</i> var. <i>perforata</i> D. Fang & K. Yan	A. Liu et al. LK0104 (CSFI)	Jiangyong, Hunan, China	MW602128	MW602095	MW602000	MW602032	MW602064
<i>Paraphlomis jiangyongensis</i> X.L. Yu & A. Liu 1	A. Liu et al. LK1104 (CSFI)	Jiangyong, Hunan, China	MW602129	MW602096	MW602001	MW602033	MW602065
<i>Paraphlomis jiangyongensis</i> C.Y. Wu & H.W. Li	Y.P. Chen & Y. Zhao EM1391 (KUN)	Huihai, Guangdong, China	MW602126	MW602093	MW601998	MW602030	MW602062
<i>Paraphlomis lancolata</i> Hand.-Mazz. 1	C.Z. Huang s.n. (KUN)	Guidong, Guangdong, China	MW602145	MW602115	MW602019	MW602051	MW602082
<i>Paraphlomis lancolata</i> Hand.-Mazz. 2	A. Liu et al. LK0845 (CSFI)	Ningyuan, Hunan, China	MW602116	MW602020	MW602052	MW602083	MW602083
<i>Paraphlomis lancolata</i> Hand.-Mazz. 2	X. Zhong et al. ZX16824 (CASH)	Suichang, Zhejiang, China	MW602136	MW602106	MW602011	MW602043	MW602074
<i>Paraphlomis lancolata</i> Hand.-Mazz. 2	Y.P. Chen et al. EM583 (KUN)	Huanjiang, Guangxi, China	OK104771	OK104774	OK104778	OK104780	OK104783
<i>Paraphlomis membranacea</i> C.Y. Wu & H.W. Li	M.S. Nuralev 1057 (MW)	Thanh Son, Phu Tho, Vietnam	/	MW602100	MW602005	MW602037	/
<i>Paraphlomis membranacea</i> C.Y. Wu & H.W. Li	C. Xiong XC21097 (KUN)	Chengkou, Chongqing, China	OMR84451	OMR84454	OMR84457	OMR84460	OMR84460
<i>Paraphlomis membranacea</i> C.Y. Wu & H.W. Li	C. Xiong & H.L. Zhou XC21126 (KUN)	Wushan, Chongqing, China	OMR84663	OMR84452	OMR84455	OMR84458	OMR84461
<i>Paraphlomis membranacea</i> C.Y. Wu & H.W. Li	L.X. Yuan et al. s.n. (KUN)	Qionghai, Hainan, China	OP605345	OP609840	OP609847	OP609854	OP609861
<i>Paraphlomis membranacea</i> C.Y. Wu & H.W. Li	X.X. Zhu s.n. (KUN)	Malipo, Yunnan, China	MW602125	MW602092	MW601997	MW602029	MW602061
<i>Paraphlomis membranacea</i> C.Y. Wu & H.W. Li	X. Li LX200704 (GXF)	Napo, Guangxi, China	MW602133	MW602103	MW602008	MW602040	MW602071
<i>Paraphlomis reflexa</i> C.Y. Wu & H.W. Li	Z.Z. Yang et al. s.n. (HB)	Tongshan, Hubei, China	MW602122	MW602089	MW601994	MW602026	MW602058
<i>Paraphlomis yingdeensis</i> WYZhao, Y.Q.Li & Q.Fan 1	Q. Fan et al. 19013 (SY)	Yingde, Guangdong, China	OP605348	OP609843	OP609850	OP609857	OP609864
<i>Paraphlomis yingdeensis</i> WYZhao, Y.Q.Li & Q.Fan 2	Q. Fan et al. 19013 (SY)	Yingde, Guangdong, China	OP605349	OP609844	OP609851	OP609858	OP609865
<i>Paraphlomis yingdeensis</i> WYZhao, Y.Q.Li & Q.Fan 3	Q. Fan et al. 19013 (SY)	Yingde, Guangdong, China	/	OP609845	OP609852	OP609859	OP609866
<i>Phomoides tratinus</i> L.	Y. Tong s.n. (KUN)	Shanghai, China (cultivated)	MW602119	MW602086	MW601991	MW602023	MW602055
<i>Phomoides denotata</i> var. <i>glabrescens</i> (Danguy) C.L. Xiang & H.	Y.P. Chen EM360 (KUN)	Beijing, China (cultivated)	MW602120	MW602087	MW601992	MW602024	MW602056

## Appendix 2



**Figure A1.** Optimal Maximum Likelihood tree of *Paraphlomis* inferred from combined nuclear (ITS and ETS) dataset. Support value  $\geq 50\%$  BS or 0.50 PP are displayed above the branches ("—" indicates a support value  $< 0.50$  PP).

### Appendix 3



**Figure A2.** Optimal Maximum Likelihood tree of *Paraphlomis* inferred from combined plastid (*rpl32-trnL*, *rps16* and *trnL-trnF*) dataset. Support value  $\geq 50\%$  BS or 0.50 PP are displayed above the branches (“-” indicates a support value  $< 0.50$  PP).