Asianopis gen. nov., a new genus of the spider family Deinopidae from Asia

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Abstract
A new genus of the spider family Deinopidae C.L. Koch, 1850 is described from Asia: Asianopis Lin & Li gen. nov., with A. zhuanghaoyuni Lin & Li sp. nov. as the type species. The new genus is divided into two species groups, of which the liukuensis-group includes two species: A. dumogae (Merian, 1911) sp. reval. comb. nov. (♀) and A. liukuensis (Yin, Griswold & Yan, 2002) comb. nov. (♂♀); and the zhuanghaoyuni-group comprises five species: A. celebensis (Merian, 1911) comb. nov. (♂), A. konplong (Logunov, 2018) comb. nov. (♂), A. wangi Lin & Li sp. nov. (♂♀), A. wuchaoi Lin & Li sp. nov. (♂♀), and A. zhuanghaoyuni Lin & Li sp. nov. All previously described species are transferred from Deinopis MacLeay, 1839. Deinopis scrubjunglei Caleb & Mathai, 2014 is treated as a junior synonym of Asianopis liukuensis comb. nov.

Keywords
New combination, new species, species groups, systematics, taxonomy
Introduction

The spider family Deinopidae C.L. Koch, 1850 (Araneae, Deinopoidea), known as net-casting or ogre-faced spiders, is a small family that consisted of two genera and 64 species prior to the current study (World Spider Catalog 2019). The genus Deinopis was established by MacLeay (1839) based on Deinopis lamia MacLeay, 1839 (♀♂) from Cuba. The other genus, Menneus, was established by Simon (1876) based on Menneus tetragnathoides Simon, 1876 (♂) from Angola.

Ten species of Deinopidae were known from Asia: Deinopis aruensis Roewer, 1938 (♀) and D. celebensis Merian, 1911 from Indonesia; D. fasciculigera Simon, 1909 (♀), and D. konplong Logunov, 2018 (♀♂) from Vietnam; D. scrubjunglei Caleb & Mathai, 2014 (♀♂) from India; D. gubatmakiling Barrion-Dupo & Barrion, 2018 (juvenile), D. labangan Barrion-Dupo & Barrion, 2018 (♀), and D. luzonensis Barrion-Dupo & Barrion, 2018 (♀) from the Philippines; D. kollari Doleschall, 1859 (♂) from Myanmar and Malaysia; D. liukuensis Yin, Griswold & Yan, 2002 (♂♀) from China. Here, we describe a new genus and three new species, and present a molecular phylogenetic analysis of these spiders.

Material and methods

All specimens were preserved in 80% ethanol. Metatarsi and tarsi were removed for preservation in 100% ethanol for subsequent molecular work. Epigynes were cleared in proteinase K at 56 °C to dissolve non-chitinous tissues for three hours. Specimens were examined under a LEICA M205C stereomicroscope. Photomicroscope images were taken with an Olympus C7070 zoom digital camera (7.1 megapixels). Laboratory habitus photographs were taken with a Canon 5D Mark III digital camera equipped with a Canon MP-E 65 mm lens. Photos were stacked with Helicon Focus (version 6.7.1) or Zerene Stacker (version 1.04) and processed in Adobe Photoshop CC 2018. Photographs of Asianopis celebensis comb. nov. were taken by a KEYENCE. Photographs of Asianopis liukuensis comb. nov. from India (i.e., the type materials of D. scrubjunglei) were taken using a Leica DFC500 HD camera mounted on a Leica M205A stereomicroscope.

All measurements are in millimetres. Eye sizes are measured as the maximum diameter from either the dorsal or frontal view. Leg measurements are given as follows: total length (femur, patella+tibia, metatarsus, tarsus). Copulatory duct turns are defined by the number of apparent loops on the lateral margin of the copulatory/fertilization duct complex in dorsal view. The length of the embolic tip fold is measured as from the beginning of the fold to the embolic tip (Fig. 22D, E). The terminology used in the text and figures follows Coddington et al. (2012). Distribution maps were generated using ArcMap software (version 10.2).

A total of 31 specimens of Deinopidae were collected for phylogenetic analysis (Suppl. material 1: Table S1). Sequences of seven specimens were from the National Center for Biotechnology Information (NCBI) public data, and the other 24 were from recent
field collections. Whole genomic DNA was extracted from 2–4 legs using a TIANamp Genomic DNA kit (TIANGEN Inc., Beijing, China) following the manufacturer’s protocol. Seven gene fragments were amplified in 20-μL reactions: COI (~640 bp), 12S (~330 bp), 16S (~470 bp), 18S (~1700 bp), 28S (~1200 bp), H3 (~310 bp) and wnt (~330 bp). Primers and PCR conditions for each locus are listed in Suppl. material 1: Table S2. Sequence chromatograms were proofed and edited using Sequencher version 4.2 Demo (Gene Codes Corporation, Ann Arbor, MI USA). The COI, H3 and wnt fragments were translated in MEGA version 7 (Kumar et al. 2016) to check for the presence of stop codons. A representative of the family Uloboridae was used as the outgroup, with the corresponding sequences downloaded from NCBI. The complete list of 32 taxa and GenBank accession numbers are provided in Suppl. material 1: Table S1.

Multiple sequence alignments were carried out with MAFFT version 7.243 (Katoh and Standley 2013). Alignments of the protein-coding COI, H3 and wnt genes were produced using the L-INS-i method. As for the highly variable ribosomal genes, the E-INS-i method was used to generate alignments of 12S, 16S, 18S, and 28S. To exclude the ambiguously aligned regions, alignments of the ribosomal genes were processed with the program trimAl version 1.3 (Capella-Gutiérrez et al. 2009). The alignments are shown in the supplementary data.

The concatenated gene matrix was partitioned by gene using PartitionFinder version 1.1.1 (Lanfear et al. 2012). The best partitioning scheme was selected based on the Akaike information criterion (AIC) (Suppl. material 1: Table S3). Maximum likelihood (ML) analysis was performed using RAxML version 8.2.9 with a GTR + Γ + I model applied to each partition (Stamatakis 2014). One thousand non-parametric bootstrap replicates were conducted to obtain the best-scoring ML tree.

Bayesian analysis was performed using MrBayes version 3.2.6 (Ronquist et al. 2012). Two independent runs, each with four independent chains, were carried out for 20,000,000 generations and were sampled every 1,000 generations with a burn-in of 25%. Partitions and models followed the result of PartitionFinder. Convergence of the runs was determined with the standard deviation of split frequencies (<0.01). Effective sampling sizes (>200) of all parameters were checked in Tracer version 1.6 (Rambaut et al. 2014). A 50% majority-rule consensus tree was then constructed from the post-burnin sampled trees to estimate posterior probabilities (PP).

### Abbreviations

| Abbreviation | Description                        |
|--------------|------------------------------------|
| ALE          | anterior lateral eye               |
| AME          | anterior median eye                |
| CD           | copulatory duct                    |
| CO           | copulatory opening                 |
| E            | embolus                            |
| EMA          | embolic middle apophysis           |
| EO           | embolic opening                    |
| ETA          | embolic terminal apophysis         |
| FD           | fertilization duct                 |
| MA           | median apophysis                   |
| MABL         | median apophysis–basal lobe        |
| MADL         | median apophysis–distal lobe       |
| MP           | median plate                       |
| PLE          | posterior lateral eye              |
Taxonomy

Family Deinopidae C.L. Koch, 1850

Genus *Asianopis* Lin & Li, gen. nov.
http://zoobank.org/C8CA3BB7-776C-4BB9-9E19-F819587E87AB

Type species. *Asianopis zhuanghaoyuni* Lin & Li, sp. nov.

**Etymology.** The generic name is a combination of the word “Asia”, referring to the distribution of the genus, and the generic name *Deinopis*. The gender is feminine.

**Diagnosis.** *Asianopis* gen. nov. can be easily distinguished from *Deinopis* by the following characters: a prominent setal fringe can be found above the posterior median eyes in both sexes of *Asianopis* species (Fig. 4A, B), which is absent in *Deinopis* (Coddington et al. 2012: fig. 3a); the embolic tip of male *Asianopis* has an embolic middle apophysis (*liukuensis*-group, Fig. 21A), an embolic terminal apophysis or is weakly folded apically (*zhuanghaoyuni*-group, Fig. 21B–E), whereas none of these characters is present in *Deinopis* (Coddington et al. 2012: fig. 11m); the MADL in *Asianopis* is small and has a basal lobe, while in *Deinopis*, the median apophysis is larger than the MABL and covers the entire base (Coddington et al. 2012: fig. 11m); female chelicerae with many denticles between the promarginal and retromarginal teeth (Fig. 2F) or female chelicerae without denticles (Fig. 2H), in contrast, denticles are only at the center of any two adjoining retromarginal teeth in *Deinopis* (Coddington et al. 2012: fig. 5c); femora I enlarged proximally in *Asianopis* gen. nov. (*liukuensis* group, Fig. 2I) or not enlarged (*zhuanghaoyuni*-group, Fig. 2J), but they are enlarged distally in *Deinopis* (Coddington et al. 2012: fig. 3b); epigynal median plate lateral margins anchor-shaped in *Asianopis* gen. nov. (Figs 3A, 6A), but ellipsoid in *Deinopis* (Coddington et al. 2012: fig. 9b); SpD is consistently narrow in *Asianopis* gen. nov. (Figs 3B, 6B) but tapering in *Deinopis* (Coddington et al. 2012: fig. 9d).

**Description.** **Male.** Total length 12.14–16.10 (n = 8), carapace pear-shaped, yellow-brown (*liukuensis*-group) or brown (*zhuanghaoyuni*-group) with white edge, white line extending from cephalic area to posterior margin and small spines sparsely dis-
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...tributed; fovea longitudinal, indistinct. Chelicerae with a promarginal tooth and one or two retromarginal teeth (liukuensis-group) or with four promarginal teeth and 2–6 retromarginal teeth (zhuanghaoyuni-group), no denticles. Endites and labium brown, distally white; sternum diamond-shaped, brown with median light band and few small spines. Legs brown, ventrally with black pattern and short spines, leg formula 1243. Opisthosoma cylindrical, brown or dark-brown with small black spots and irregular pattern. Cribellum entire, spinnerets brown (Figs 4, 10, 13, 16).

Female. Total length 14–24 (n = 13). Chelicerae with four promarginal teeth and seven retromarginal teeth, many denticles in between the promarginal and retromarginal teeth (liukuensis-group) or four promarginal teeth and 8–13 retromarginal teeth, without denticles (zhuanghaoyuni-group). Appearance of carapace, opisthosoma and legs as in male but femora of legs I enlarged basally (liukuensis-group) (Fig. 2I).

Male palpal tibia longer than cymbium; cymbium almost round; tegulum distinctly wider than the diameter of embolic coil (liukuensis-group) or tegulum obscured by embolic coil (zhuanghaoyuni-group) (Figs 17, 18); embolus long and strongly coiled around MA, embolic base beginning at 7–8 o’clock position, coiled 1200° (liukuensis-group) or more than 1500° (zhuanghaoyuni-group), embolic tip straight (liukuensis-group) or widened subapically, folded and without apophysis (zhuanghaoyuni-group); MA small, directed at 7–8 o’clock position, with two lobes, a small lobe at the base, and a narrow distal lobe with two apophyses (liukuensis-group) or large, with two lobes, a large lobe at the base and a kidney-shaped distal lobe (zhuanghaoyuni-group).

Epigyne with anchor-shaped median plate, CO distinct, CD with three turns, S oval, SpD consistently wide (liukuensis-group) or with a well-developed MP, obscuring CO, CD with 7–8 turns, S oval, SpD consistently thin (zhuanghaoyuni-group).

Molecular phylogeny. The molecular phylogenetic analysis indicates with strong support that all the species in this study do not belong to Deinopis. Based on the 4893 bp-aligned sequences of seven gene fragments, the ML and Bayesian analyses produced the same topology, showing a split of a Southwest China clade from other clades and is strongly supported (Bootstrap value: 88; PP: 0.98) (Fig. 1). Our results are consistent with the results of Chamberland et al. (2018) who conducted a global phylogenetic analysis of Deinopis. Therefore, the Southwest China clade can be classified as a new genus with strong support (Bootstrap value: 100; PP: 1). Although intraspecific support values are low in both ML and Bayesian analyses results, basal nodes are strongly supported, including the sister relationship of A. wangi Lin & Li, sp. nov. & A. zhuanghaoyuni Lin & Li, sp. nov. (Bootstrap value: 95; PP: 1).

Natural habitat. All the species of Asianopis gen. nov. were collected from bushes in low-elevation forests.

Composition. This new genus comprises two species groups: the liukuensis-group with two species: A. dumogae (Merian, 1911) sp. reval. comb. nov. and A. liukuensis (Yin, Griswold & Yan, 2002) comb. nov. and the zhuanghaoyuni-group with five species: A. celebensis (Merian, 1911) comb. nov., A. konplong (Logunov, 2018) comb. nov., A. wangi sp. nov., A. wuchaoi sp. nov., and A. zhuanghaoyuni sp. nov.

Distribution. China (Fujian, Yunnan, Hong Kong, Guangxi, Hainan), India, Indonesia, and Vietnam.
Figure 1. Phylogenetic tree of Deinopidae spiders based on 31 specimens. Numbers on nodes indicate Maximum Likelihood bootstrap values and Bayesian posterior probabilities.

The liukuensis-group

Asianopis dumogae (Merian, 1911), sp. reval. comb. nov.

Fig. 3

Dinopis dumogae Merian, 1911: 171 (♀ only, ♂ mismatched).

Type material examined. 1 ♀ (NMB-ARAN-00514a), “Wald bei Duluduo”, Sulawesi Utara, forest near Duluduo, 00°31'33"N, 123°57'10"E, Sulawesi, Indonesia.

Diagnosis. This species can be distinguished from A. liukuensis comb. nov. by the MP nearly covering the CO, S round, and the overall equal thickness of the CD (Figs 4, 6).

Description. See Merian (1911). Photos of the epigyne of the syntype are given in Figure 6.

Distribution. Indonesia (North Sulawesi).

Comments. Merian (1911) reported D. celebensis based on three specimens from different localities in Sulawesi, Indonesia. One male (NMB-ARAN-00514b, “Zen-
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Figure 2. Prosoma (frontal view, upper ♂, lower ♀) (A–D), chelicerae (E–H) and leg I (I–J). Figures A and C modified from Coddington et al. (2012). A Deinopis spinosa B Asianopis liukuensis comb. nov. C Menneus dromedarius D Asianopis zhuanghaoyuni sp. nov. E Chelicerae of male A. liukuensis comb. nov. F Chelicerae of female A. liukuensis comb. nov. (Arrows indicate the denticles) G Chelicerae of male A. zhuanghaoyuni sp. nov. H Chelicerae of female A. zhuanghaoyuni sp. nov. I Left leg I of female A. liukuensis comb. nov. Arrow shows enlarged femur J Left leg I of female A. zhuanghaoyuni sp. nov.

Central-Celebes, nördlich vom Golf von Bone”, South Sulawesi, north of the Gulf of Boni (precise locality not known), one female from North Sulawesi (NMB-ARAN-00514a, “Wald bei Duluduo”, Sulawesi Utara, forest near Duluduo, 00°31’33”N, 123°57’10”E and one female from Central Sulawesi (NMB-ARAN-00514c, Larga, südlich vom Posso-See, unterhalb Patiro Rano, bei 900 m, Central Sulawesi, south of Lake Poso at an elevation of 900 m (the localities “Larga” and “Patiro Rano” could not be located on maps; the epigyne of this specimen is missing, but the specimen is clearly larger than the others).
Merian (1911) stated that the male and the females may not represent the same species and suggested the name *D. celebensis* for the male, and *D. dumogae* for the female. According to the International Code of Zoological Nomenclature (International Commission on Zoological Nomenclature 1999: Article 11.5.1), such conditionally proposed species names are potentially available as valid names if published before 1961. The species has not been listed in any of the catalogues. We examined the types and concluded the male and the two females are indeed three different species. The palp of the male *D. celebensis* exhibits features of the *zhuanghaoyuni* group: the tegulum is obscured by the embolic coil, and the embolus is long and strongly coiled around the MA. The female from North Sulawesi (Doloduo) has features of the *liukuensis* group: an anchor-shaped median plate, CO distinct, CD with three turns. Thus, we revalidated the female *D. dumogae* as *Asianopis dumogae* (Merian, 1911), sp. reval. comb. nov.

*Asianopis liukuensis* (Yin, Griswold & Yan, 2002), comb. nov.
Figs 2B, E, F, I, 4–8, 19, 21A, 22A, G, 23

*Deinopis liukuensis* Yin et al., 2002: 610, figs 1–7 (♂♀)
*Deinopis liukuensis* Zhang & Wang, 2017: 238 (♂♀)
*Deinopis scrubjunglei* Caleb & Mathai, 2014: 2, figs 1–20 (♂♀) syn. nov.

**Type. Holotype.** ♂ (HNU, no. 00-LK-1, lost), China, Yunnan Province, Liuku, Mt Gao-ligong, 25°30’48”N, 98°30’36”E, elevation ca 800 m, 26.VI.2000, Heng-Mei Yan leg.

**Type materials of Deinopis scrubjunglei examined.** ♂ (SRC-ZSI I/SP 19), Madras Christian College, Chennai, Tamil Nadu, 12°55’12.7”N, 80°07’24.6”E, elevation ca 32 m, 5.XII.2013, John Caleb T.D. leg.; ♂ (SRC-ZSI I/SP 20), 22.IV.2014, same location, John Caleb T.D and Karthy leg.
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Figure 4. *Asianopis liukuensis* comb. nov., male from Xishuangbanna and female from Jianfengling. 
A Male prosoma, frontal view 
B Female prosoma, frontal view 
C Male habitus, dorsal view 
D Male habitus, ventral view 
E Female habitus, dorsal view 
F Female habitus, ventral view.
Figure 5. *Asianopis liukuensis* comb. nov., left palp, male from Xishuangbanna. **A** Prolateral view **B** Retrolateral view.
Figure 6. *Asianopis liukuensis* comb. nov., female from Jianfengling. **A** Epigyne **B** Vulva, dorsal view.
Other material examined. 2♂, China, Yunnan Province, Xishuangbanna Dai Autonomous Region, rubber tree plantation near Jinhong City, 28.IV.2016, Chao-tai Wei leg.; 1♀, China, Hainan Island, Ledong County, Jianfengling National Park, 13.VII.2019, Zixuan Lin leg.

Diagnosis. This species can be distinguished from other congeners by the distinct female copulatory opening, oval S, and CD tapering from the copulatory opening to spermatheca (Figs 6, 8).

Description. See Yin et al. (2002) and Caleb and Mathai (2014).
Distribution. China (Yunnan, Guangxi, Hainan), India.

Comments. Type materials of *D. scrubjunglei* syn. nov. were examined and no differences between *A. liukuensis* and *D. scrubjunglei* were observed. Thus, we consider *D. scrubjunglei* to be a synonym of *A. liukuensis*, and the figures of *D. scrubjunglei* are given for comparison (Figs 7, 8, 19C).
The **zhuanghaoyuni-group**

*Asianopis celebensis* (Merian, 1911), comb. nov.

Fig. 9A–F

*Dinopis celebensis* Merian, 1911: 167, figs A, B (♂ only, ♀ mismatched).

**Type material examined.** ♂ (NMB), NMB-ARAN-00514b, “Zentral-Celebes, nördlich vom Golf von Bone”, South Sulawesi, north of the Gulf of Boni (precise locality not known).

**Diagnosis.** The male can be distinguished from other congeners by having the distal lobe of the MA distinctly smaller than the basal lobe; in other *Asianopis* spp., the distal lobe is slightly smaller than the basal lobe (Fig. 9A, C).

**Description.** See Merian (1911). Photos of holotype male habitus and palps are shown in Fig. 9A–F.

**Distribution.** Indonesia (Sulawesi).

**Comments.** One male and two females were types for *Asianopis celebensis* (Merian, 1911) comb. nov. after Merian (1911). Based on the current study, one type female from North Sulawesi is *Asianopis dumogae* (Merian, 1911) sp. reval. comb. nov., and the other type female from South Sulawesi is a member of the **zhuanghaoyuni-group**, but its status at the species level is uncertain because of the missing epigyne.

*Asianopis konplong* (Logunov, 2018), comb. nov.

*Deinopis konplong* Logunov, 2018: 141, figs 1–7 (♂).

**Type.** Holotype ♂ (MMUE, G7579.37) from Vietnam, Kon Tum Province, Kon Plong District, 14 km north of Kon Plong, 14°43’20”N, 108°18’59”E, elevation ca 1030 m, 3–12.VI.2016, A.A. Abramov leg. Not examined.

**Diagnosis.** This species can be distinguished from other *Asianopis* species by the short palp (ratio of the length of the palpal tarsus to the length of the cymbium: 1:1) and upturned embolic tip (Logunov 2018: fig. 4).

**Description.** See Logunov (2018).

**Distribution.** Vietnam (Kon Tum).

*Asianopis wangi* Lin & Li, sp. nov.

http://zoobank.org/64A4C3D1-03A5-4D7A-B2E6-E30EA28DC41C

Figs 10–12, 20B, 21C, 22C, D, H, 23

**Type.** Holotype ♂ (IZCAS-Ar39681), China, Hainan Province, Wuzhishan City, Wuzhishan Nature Reserve, Diewupo, 17.V.2019, Dongdong Wang leg.
Figure 9. *Asianopis celebensis* comb. nov., male type. A Male right palp, prolateral view B Embolic tip C male left palp (embolic tip detached), retrolateral view D Male prosoma, lateral view E Male opisthosoma, dorsal view F Male prosoma, frontal view.
**Paratypes.** 1♂1♀ (IZCAS-Ar39682-Ar39683), same data as holotype; 1♂2♀ (IZCAS-Ar39684-Ar39686) China, Hainan Province, Wuzhishan City, Nansheng Town, Maoxiang Village, 18.V.2019, Dongdong Wang leg.

**Etymology.** The species is named after Mr Dongdong Wang, the collector of the holotype; noun (name) in genitive case.

**Diagnosis.** The males resemble *A. zhuanghaoyuni* sp. nov. but can be distinguished from other species by the ratio of the length of the embolic opening to the length of the embolic tip fold, which is 1:6 in *A. wangi* sp. nov. and 1:8 in *A. zhuanghaoyuni* sp. nov. The fold is more developed in *A. wangi* sp. nov. (Fig. 21C, D). The median plate is triangular in *A. wangi* sp. nov. and subtriangular in *A. zhuanghaoyuni* sp. nov. (Figs 12, 19).

**Description.** Male holotype (Figs 10A, C, D, 11, 20B, 21D, 22C). Total length 15.31, carapace 6.22 long, 4.60 wide, opisthosoma 9.32 long, 2.10 wide. Eye sizes and interdistances: AME 0.30, ALE 0.38, PME 0.65, PLE 0.34, AME–AME 0.30, AME–ALE 0.97, PME–PME 0.23, PME–PLE 0.69, AME–PME 0.24, ALE–PLE 1.82. Clypeus height 0.10. Chelicerae with four promarginal and 10–13 retromarginal teeth. Leg measurements: leg I: 84.08 (21.13 + 26.50 + 29.53 + 6.92), leg II: 59.70 (18.39 + 19.55 + 15.80 + 5.96), leg III: 36.14 (12.05 + 11.79 + 10.26 + 2.04), leg IV: 37.23 (11.92 + 12.37 + 11.28 + 1.66). Leg formula: 1243.

Male palp (Figs 11, 20B, 21D). Cymbium hemispherical; tegulum flat, obscured by embolic coils; embolus long and strongly coiled, originating at 10 o’clock and coiling 1500° around MA; embolic tip widened subapically, strongly folded and without apophysis. MA large, with two lobes.

Female paratype (Figs 10B, E, F, 12, 22D). Total length 24.04, carapace 7.56 long, 5.32 wide, opisthosoma 16.28 long, 6.86 wide. Eye sizes and interdistances: AME 0.28, ALE 0.38, PME 1.34, PLE 0.42, AME–AME 0.13, AME–ALE 1.03, PME–PME 0.39, PME–PLE 1.30, AME–PME 0.22, ALE–PLE 1.92. Clypeus height 0.34 (n = 1). Chelicerae with four promarginal and 8–13 retromarginal teeth (8(n = 1), 13(n = 1)). Leg measurements: Leg I: 54.24 (16.22 + 16.83 + 17.63 + 3.56), leg II: 50.59 (15.90 + 16.41 + 15.00 + 3.28), leg III: 30.84 (10.96 + 10.38 + 7.88 + 1.62), leg IV: 30.28 (10.13 + 10.58 + 8.27 + 1.30). Leg formula: 1234.

Epigyne (Fig. 12) with a median plate, CD with 7 or 8 turns, S oval, SpD consistently narrow.

**Distribution.** China (Hainan).

*Asianopis wuchaoi* Lin & Li, sp. nov.
http://zoobank.org/F05E46B7-98E7-4DA1-B7DF-AD440C2E05B6
Figs 13–15, 21B, 22B, 23

**Type.** Holotype. ♂ (IZCAS-Ar39687), China, Yunnan Province, Jinghong City, Mount Jino, 10.V.2019, Chao Wu leg.

**Paratypes.** 2♀ (IZCAS-Ar39688-Ar39689), China, Yunnan Province, Jinghong City, Mengla County, Mengxing Village, 16.VI.2019, Yi Li leg.; 1♀ (IZCAS-
Figure 10. *Asianopis wangi* sp. nov., male holotype and female paratype. **A** Male prosoma, frontal view  
**B** Female prosoma, frontal view  
**C** Male habitus, dorsal view  
**D** Male habitus, ventral view  
**E** Female habitus, dorsal view  
**F** Female habitus, ventral view.
Figure 11. *Asianopus wangi* sp. nov., left palp, male holotype. A Prolateral view B Retrolateral view.

Ar39690), China, Yunnan Province, Jinghong City, Situlaozhai Village, 20.V.2019, Chaotai Wei leg.

**Etymology.** The species is named after Mr Chao Wu, the collector of the holotype male; noun (name) in genitive case.

**Diagnosis.** The males can be easily distinguished by the length of the palpal tibia which is almost equal to the length of the cymbium; simple embolic tip with ETA (Fig. 21B); embolus coiling almost 3300° around MA. Epigyne with a well-developed, subtriangular median plate, obscureing CO, and CD with 9 turns (Fig. 14).
Figure 12. *Asianopis wangi* sp. nov., female paratype. **A** Epigyne **B** Vulva, dorsal view.
Figure 13. *Asianopis wuchaoi* sp. nov., male holotype and female paratype. A Male prosoma, frontal view B Female prosoma, frontal view C Male habitus, dorsal view D Male habitus, ventral view E Female habitus, dorsal view F Female habitus, ventral view.
Description. Male holotype (Figs 13A, C, D, 14, 21A). Total length 12.14, carapace 4.00 long, 3.40 wide, opisthosoma 8.14 long, 2.4 wide. Eye sizes and interdistances: AME 0.15, ALE 0.26, PME 0.52, PLE 0.29, AME–AME 0.17, AME–ALE
Figure 15. *Asianopis wuchaoi* sp. nov., female paratype. A Epigyne B Vulva, dorsal view.
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0.70, PME–PME 0.16, PME–PLE 0.61, AME–PME 0.11, ALE–PLE 0.95. Clypeus height 0.05. Chelicerae with four promarginal and six retromarginal teeth. Leg measurements: leg I: damaged, leg II: damaged, leg III: (6.92 + 6.86 + ? + 1.44), leg IV: 21.82 (6.91 + 7.18 + 6.35 + 1.38).

Male palp (Figs 14, 21A). Cymbium hemispherical; tegulum flat, obscured by embolic coils; embolus long and strongly coiled, originating at five o’clock and coiling 3300° around MA. MA large, with two lobes.

Female paratype (Figs 13B, E, F, 15). Total length 14.60, carapace 6.28 long, 4.10 wide, opisthosoma 9.29 long, 3.72 wide. Eye sizes and interdistances: AME 0.11, ALE 0.34, PME 0.94, PLE 0.29, AME–AME 0.30, AME–ALE 1.03, PME–PME 0.64, AME–PME 0.14, ALE–PLE 1.33. Clypeus height 0.13 (n = 1). Chelicerae with four promarginal and 8–13 retromarginal teeth (8(n = 1), 10(n = 1), 13(n = 1)). Leg measurements: Leg I: 39.82 (12.11 + 11.67 + 13.01 + 3.03), leg II: 36.81 (11.47 + 11.79 + 10.83 + 2.72), leg III: 23.53 (9.47 + 6.79 + 5.83 + 1.44), leg IV: 21.71 (7.18 + 7.76 + 5.70 + 1.07). Leg formula: 1234.

Epigyne (Fig. 15) with a median plate, obscuring CO, CD with 9 turns, S oval, SpD is consistently thin.

Distribution. China (Yunnan).

Note. The male died during ecdysis so some legs are damaged or curled, and the palps are expanded.

Asianopis zhuanghaoyuni Lin & Li, sp. nov.
http://zoobank.org/21A5E514-F8EE-4479-9338-51D419AA6E4A
Figs 2D, G, H, J, 16–18, 20, 21D, 22E, F, H, 23

Type. Holotype. ♂ (IZCAS-Ar39691), China, Fujian Province, Fuzhou City, Minhou County, Xiyuan Reservoir, 26°03′15.5″N, 119°06′05.4″E, elevation ca 102 m, 25.VI.2018, Haoyun Zhuang and Zhuoheng Jiang leg.

Paratypes. 1 ♀ (IZCAS-Ar39692), same data as holotype, Haoyun Zhuang leg.; 1♂1♀ (IZCAS-Ar39693-Ar39694), same locality data as holotype, but 15.V.2018, Haoyun Zhuang leg.; 1♂4♀ (IZCAS-Ar39695-Ar39699), same locality data as holotype, but 19.VI.2019, Haoyun Zhuang leg.; 1♂1♀ (IZCAS-Ar39700-Ar39701), same locality data as holotype, but 26.V.2019, Haoyun Zhuang leg.

Etymology. The species is named after Mr Haoyun Zhuang, the collector of the type specimens; noun (name) in genitive case.

Diagnosis. The males resemble A. konplong (Logunov, 2018) comb. nov. but can be distinguished by the embolus originating at five o’clock in A. zhuanghaoyuni sp. nov. (9 o’clock in A. konplong (Logunov, 2018) comb. nov.); the ratio of the length of the palpal tarsus to the length of the cymbium is 11:9 in A. zhuanghaoyuni sp. nov., while in A. konplong (Logunov, 2018), comb. nov. it is 1:1 (Figs 18, 22A; Logunov 2018, figs 4–6).
Figure 16. *Asianopis zhuanghaoyuni* sp. nov., male holotype and female paratype. 

**A** Male prosoma, frontal view  
**B** Female prosoma, frontal view  
**C** Male habitus, dorsal view  
**D** Male habitus, ventral view  
**E** Female habitus, dorsal view  
**F** Female habitus, ventral view.
**Description.** Male holotype (Figs 2G, 16A, C, D, 17, 20A, 21E, 22E). Total length 16.54, carapace 5.58 long, 3.84 wide, opisthosoma 11.40 long, 1.90 wide. Eye sizes and interdistances: AME 0.25, ALE 0.30, PME 0.59, PLE 0.30, AME–AME 0.25, AME–ALE 0.85, PME–PME 0.23, PME–PLE 0.59, AME–PME 0.19, ALE–PLE 1.28. Clypeus height 0.20. Chelicerae with four promarginal teeth and a retro-marginal tooth. Leg measurements: leg I: 66.35 (18.50 + 22.55 + 18.95 + 6.35), leg II: 52.87 (16.54 + 17.65 + 13.10 + 5.58), leg III: 30.39 (10.78 + 10.83 + 7.18 + 1.60), leg IV: 30.06 (10.42 + 11.12 + 7.18 + 1.34). Leg formula: 1234.
Figure 18. *Asianopis zhuanghaoyuni* sp. nov., female paratype. A Epigyne B Vulva, dorsal view.
Figure 19. *Asianopis liukuensis* comb. nov., left palp, ventral view. **A, B** Male from Xishuangbanna. **C** Male from India, type of *Deinopis scrubjunglei* syn. nov.
Male palp (Figs 18, 22A). Cymbium hemispherical; tegulum flat, obscured by embolus coils; originating at five o’clock, coiling 1500° around MA, embolic tip widened subapically, folded and without apophysis. MA large, with two lobes.

**Female** paratype (Figs 2H, J, 16B, E, F, 18, 22F, H). Total length 22.60, carapace 5.90 long, 4.55 wide, opisthosoma 15.40 long, 5.90 wide. Eye sizes and interd-
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Interstices: AME 0.22, ALE 0.35, PME 1.08, PLE 0.33, AME–AME 0.37, AME–ALE 1.22, PME–PME 0.16, PME–PLE 0.98, AME–PME 0.081, ALE–PLE 1.61. Clypeus height 0.59. (n = 1). Chelicerae with four promarginal and 10 or 11 (10 (n = 2), 11 (n = 1)) retromarginal teeth. Leg measurements: Leg I: 49.68 (14.80 + 15.83 + 16.02 + 3.03), leg II: 46.08 (14.71 + 15.20 + 13.33 + 2.84), leg III: 27.79 (9.73 + 9.41 + 7.18 + 1.47), leg IV: 26.78 (9.02 + 9.61 + 6.86 + 1.29). Leg formula: 1234.

Epigyne (Fig. 18) with a median plate, CD with 7–8 turns, S oval, SpD consistently thin.

**Distribution.** China (Fujian).

*Figure 21.* Embolic tips of four species of *Asianopis* gen. nov. **A** *A. liukuensis* (Yin, Griswold & Yan, 2002) comb. nov. **B** *A. wuchaoi* sp. nov. **C** *A. zhuanghaoyuni* sp. nov. **D** *A. wangyi* sp. nov.
Figure 22. Photos of four live spiders of *Asianopis* gen. nov., including webs of two species of *Asianopis* gen. nov. **A**. *A. liukuensis* comb. nov., female **B**. *A. wuchaoi* sp. nov., female **C**. *A. wangi* sp. nov., male **D**. *A. wangi* sp. nov., female **E**. *A. zhuanghaoyuni* sp. nov., male **F**. *A. zhuanghaoyuni* sp. nov., female **G**. Web of *A. liukuensis* comb. nov. **H**. Web of *A. wangi* sp. nov.
Figure 23. Distribution records of seven species of *Asianopsis* gen. nov. in Asia. 1 *A. liukuensis* comb. nov. 2 *A. dumogae* sp. reval. comb. nov. 3 *A. celebensis* comb. nov. 4 *A. konplong* comb. nov. 5 *A. wangi* sp. nov. 6 *A. wuchaoi* sp. nov. 7 *A. zhuanghaoyuni* sp. nov.

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Supplementary material I

Asianopis gen. nov., a new genus of the spider family Deinopidae from Asia
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Data type: specimen/primer/DNA sequence
Explanation note: Table S1. List of voucher information and GenBank accession numbers. Table S2. Primers and PCR conditions for the genetic markers used in this study (modified after Zhao and Li unpublished). Table S3. Sequence characteristics and models of DNA evolution selected for the seven sequence regions analyzed. References cited in supplementary tables.
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