

# *Syntormon* Loew (Diptera, Dolichopodidae) from Inner Mongolia, China, with the description of a new species

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## Abstract

Previously, no records of *Syntormon* Loew, 1857 species were known from Inner Mongolia (China). The genus is reported here from Inner Mongolia for the first time, with the description of a new species, *S. sinicum* **sp. nov.**, along with two previously described species, *S. dukha* Hollis, 1964 and *S. henanense* Yang & Saigusa, 2000. *Syntormon sinicum* **sp. nov.** and *S. dukha* Hollis, 1964 are barcoded for the first time to support the species delimitation. A key to *Syntormon* species in China is provided.

**Key words:** Barcoding, *Syntormon dukha*, *Syntormon henanense*, taxonomy

## Introduction

The genus *Syntormon* Loew contains more than 110 known species worldwide (Yang et al. 2006; Grichanov 2014; Grichanov 2021). Species of *Syntormon* can be recognized in both sexes by the antenna pedicel with one short finger-like projection projecting into the postpedicel. In recent years, *Syntormon* has been included in several molecular phylogenetic studies which have shed light on its phylogenetic position. *Syntormon* belongs to Sympycninae, within the Dolichopodidae *sensu lato*, and the intergeneric relationships within the Sympycninae have been demonstrated (Bernasconi et al. 2007; Lim et al. 2010; Germann et al. 2011). There have been 63 mitochondrial sequences from 10 species reported in these studies, of which 23 and 10 sequences were sequenced from *Syntormon flexibile* Becker, 1922 and *S. pallipes* Fabricius, 1794, respectively. These data have supported the molecular identification of *S. pallipes* and *S. pseudospicatum* (Chursina and Grichanov 2019; Tonguç et al. 2023). Still, most *Syntormon* species can only be identified by morphological characteristics (e.g. Drake 2020, 2021).

Thus far, 15 known species of *Syntormon* are known to occur China, with four species recorded in the Palearctic realm, nine species recorded in the Oriental realm, and two species recorded in both realms (Yang et al. 2011). The Palearctic species are recorded from areas with a temperate continental climate, which is typical of Inner Mongolia, a province that belongs to Palearctic China. The natural vegetation of Inner Mongolia encompasses a diverse range of ecosystems



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from forests, meadow steppe, steppe, desert steppe to Gobi Desert from northeast to southwest due to the greater precipitation in the northeast compared to the southwest and higher temperatures in the southwest compared to the northeast. However, there was no reports of *Syntormon* from Inner Mongolia.

The present study reports the distribution of *Syntormon* in Inner Mongolia for the first time, with records of a new species, *Syntormon sinicum* sp. nov., and two known species, *S. dukha* Hollis, 1964 and *S. henanense* Yang & Saigusa, 2000. This is also the first record of Sympycninae from Inner Mongolia. In addition, we provide the mitochondrial COI gene of *S. sinicum* sp. nov. and *S. dukha* Hollis, 1964 for the first time. A key to the species of the genus from China is provided.

## Materials and methods

### Morphological taxonomy

The specimens on which this study is based were collected in Inner Mongolia during 2013–2021 by sweep net. Hohhot, Bayan Nur, Baotou, Xilingol League, Ulanqab, Chifeng, Tongliao, Hinggan League in Inner Mongolia were investigated, except for Hulun Buir. We focused on wet biotopes such as river basins, lakes, and forests for collection. All specimens are deposited in the China Agricultural University (CAU), Beijing and Entomological Museum of Institute of Grassland Research, Chinese Academy of Agricultural Sciences (IGRCAAS). Morphological terminology follows Cumming and Wood (2017). Keys by Yang et al. (2011) for Chinese species were used to identify specimens collected. The following abbreviations are used: **acr** = acrostichal, **ad** = anterodorsal, **av** = anteroventral, **dc** = dorsocentral, **sc** = scutellars, **pd** = posterodorsal, **v** = ventral, **LI** = fore leg, **LII** = mid leg, **LIII** = hind leg, **CuAx ratio** = length of dm-cu / length of distal portion of CuA.

### DNA sequencing

Specimens used in this research were preserved in 95% ethanol at  $-20^{\circ}\text{C}$ . The mitochondrial genomic DNA was extracted from muscle tissue from thorax using the TIANamp Genomic DNA Kit (Tiangen) according to the manufacturer's protocol. All PCR reactions were performed in a 50  $\mu\text{L}$  volume: 2  $\mu\text{L}$  DNA extract, 25  $\mu\text{L}$  Taq PCR Master Mix, 2  $\mu\text{L}$  of each primer, and 19  $\mu\text{L}$  ddH<sub>2</sub>O. The reaction cycle was set as follows: 94  $^{\circ}\text{C}$  for 5 min of initial degeneration, 35 cycles for 94  $^{\circ}\text{C}$  for 30 s, 56  $^{\circ}\text{C}$  for 30 s, 72  $^{\circ}\text{C}$  for 1 min, and a final extension of 72  $^{\circ}\text{C}$  for 10 min.

### Molecular identification

The newly sequenced and all available sequences of the mitochondrial COI gene of *Syntormon* were used to construct a distance-based neighbour-joining (NJ) phylogenetic tree, which was used for the species delimitation of the new species. Sequences that were significantly shorter than others, which were suspected to be incomplete, were pre-filtered. *Dolichopus ornatipennis* Van Duzee, 1921 and *D. nigrilineatus* Van Duzee, 1924 were used as outgroups. Sequences are all available in the GenBank of National Center for Biotechnology Information (NCBI, <https://www.ncbi.nlm.nih.gov>) and the Barcode of Life Data System (BOLD, <https://www.boldsystems.org/>) (Table 1). CLUSTAL W was used

**Table 1.** Sequences information used in molecular analysis.

Genus	Species	GenBank accession number
<i>Syntormon</i> Loew, 1857	<i>S. bicolorellum</i> Zetterstedt, 1843	<a href="#">MZ624427</a>
	<i>S. dukha</i> Hollis, 1964	<a href="#">OR762505</a>
	<i>S. flexibile</i> Becker, 1922	<a href="#">MG086951</a>
	<i>S. freymuthae</i> Loew, 1873	<a href="#">MZ626600</a>
	<i>S. pallipes</i> Fabricius, 1794	<a href="#">MZ611071</a>
	<i>S. pumilum</i> Meigen, 1824	<a href="#">MZ608991</a>
	<i>S. sinicum</i> sp. nov.	<a href="#">OR762504</a>
	<i>S. tarsatum</i> Fallén, 1823	<a href="#">MZ628445</a>
<i>Dolichopus</i> Latreille, 1796	<i>D. ornatipennis</i> Van Duzee, 1921	<a href="#">HM413216</a>
	<i>D. nigrilineatus</i> Van Duzee, 1924	<a href="#">KC502345</a>

to align the sequences (Kumar et al. 2018). The NJ tree was then constructed using the Kimura 2-parameter model and the 1000 rapid bootstrap replicates were performed using MEGA X (Kimura 1980; Trees 1987; Tamura et al. 2011).

## Taxonomy

### Key to the species of *Syntormon* from China

- 1 First tarsomere of LIII with ventral appendages.....**2**
- First tarsomere of LIII without ventral appendages .....**4**
- 2 Acr bristles uniseriate; 1<sup>st</sup> tarsomere of LIII with 2 curved spines .....  
..... ***S. beijingense*** (Beijing)
- Acr bristles biseriate; 1<sup>st</sup> tarsomere of LIII with 4 irregular processes.....**3**
- 3 Postpedicel 3.0 times longer than wide; hind tarsomere I with 2 same spines of same shape .... ***S. pallipes*** (Xinjiang, Qinghai, Beijing, Guizhou, Henan, Shaanxi)
- Postpedicel less than 3.0 times longer than wide; hind tarsomere I with 2 differently shaped spines, 1 strong 1 thin (Fig. 1A) .....  
..... ***S. dukha*** (Yunnan, Inner Mongolia)
- 4 Hind tibia and tarsus flattened and plumose ..... ***S. zhengi*** (Qinghai)
- Hind tibia and tarsus simple .....**5**
- 5 Hind tarsus slightly shortened, shorter than half length of tibia .....**6**
- Hind tarsus not shortened, longer than half length of tibia .....**7**
- 6 First tarsomere of LI distinctly longer than length of tarsomeres II–V.....  
..... ***S. guizhouense*** (Guizhou)
- First tarsomere of LI normal (not longer than tarsomeres II–V) ..... ***S. flexibile*** (Hebei, Fujian, Guangdong, Guizhou, Jiangsu, Zhejiang, Shanghai, Taiwan)
- 7 All coxae wholly black .....**8**
- Fore coxa yellow, mid and hind coxae black.....**9**
- 8 Postpedicel 1.8 times longer than wide; arista 2.0 times longer than postpedicel; squama with black hairs ..... ***S. luchunense*** (Yunnan)
- Postpedicel 3.0 times longer than wide; arista distinctly shorter than postpedicel, no longer than 1/2 of postpedicel; squama with white hairs .....  
..... ***S. xizangense*** (Xizang)
- 9 Postpedicel rather short, about as long as wide (Fig. 1C, D).....  
..... ***S. sinicum* sp. nov.** (Inner Mongolia)
- Postpedicel distinctly elongated, more than 3.0 times longer than wide ..... **10**

- 10 Postpedicel 4.2 times longer than wide; arista 0.2 times as long as postpedicel.....**11**
  - Postpedicel 3.5 times longer than wide; arista as long as postpedicel.....**12**
- 11 Acr bristles 7–8 pairs; hind tibia with 1 ad bristle and 5 pd bristles (Fig. 1B)..... ***S. henanense*** (Henan, Shaanxi, Yunnan, Inner Mongolia)
  - Acr bristles 13 pairs; hind tibia without ad bristles and with 2 or 3 pd bristles ..... ***S. xinjiangense*** (Xinjiang)
- 12 Acr bristles 5 or 6 pairs; arista shorter than postpedicel.....**13**
  - Acr bristles 20 in line, uniseriate; arista longer than postpedicel.....  
.....***S. trisetum*** (Fujian)
- 13 Squama with yellowish hairs ..... **14**
  - Squama with black hairs.....***S. luchunense*** (Yunnan, Guizhou)
- 14 Five dc bristles; 10 or 11 pairs of acr bristles..... ***S. medogense*** (Xizang)
  - Six dc bristles; 5 or 6 pairs of acr bristles.....***S. emeiense*** (Sichuan, Guizhou)

#### ***Syntormon dukha* Hollis, 1964**

Fig. 1A

*Syntormon dukha* Hollis, 1964: 93. Type locality: Nepal: Sangu, Taplejung.  
*Syntormon dukha* Hollis, 1964. Yang et al. 2010: 1355.

**Diagnosis.** Antenna postpedicel distinctly elongated, 3.0 times longer than wide; arista subapical, as long as postpedicel. First tarsomere of LI with 1 unequal bifurcated ventral spine at base.

**Specimens examined.** CHINA: Inner Mongolia, 5 males 5 females, Mount Helan, Huangqokou, 1900 m, 31.VII.2010, Yan Li (CAU-SYMSYN001A01-SYMSYN001A10); 1 male 1 female, Mount Jiufeng, Erdaogou, 1400–1500 m, 3.VIII.2013, Xiao Zhang (CAU-SYMSYN001B01-SYMSYN001B02); 10 males 8 females ([OR762505](#)), Mount Helan, Halawu, 2000 m, 5.VIII.2021, Xingyang Qian (IGRCAAS-SYMSYN1A01-SYMSYN1A18); 1 male 2 females, Hohhot, Xiaojinggou, 1400 m, 22.VIII.2021, Xingyang Qian (IGRCAAS-SYMSYN1B01-SYMSYN1B03); 1 male, Chifeng, Heilihe, 1000 m, 19.VII.2022, Xingyang Qian (IGRCAAS-SYMSYN1A01).

**Habitat.** This species was collected in forests (Mount Jiufeng, Mount Helan, Xiaojinggou, Heilihe).

**Distribution.** China (Yunnan, Inner Mongolia)

#### ***Syntormon henanense* Yang & Saigusa, 2000**

Fig. 1B

*Syntormon henanense* Yang & Saigusa, 2000: 207. Type locality: China: Henan, Songxian, Baiyunshan Mountain.  
*Syntormon henanense* Yang & Saigusa, 2000. Yang et al. 2010: 1362.

**Diagnosis.** Antenna postpedicel distinctly elongated; arista very short. 7 or 8 acr in a line, short-haired. Hind tibia with 1 antero-dorsal bristle and 5 postero-ventral bristles.

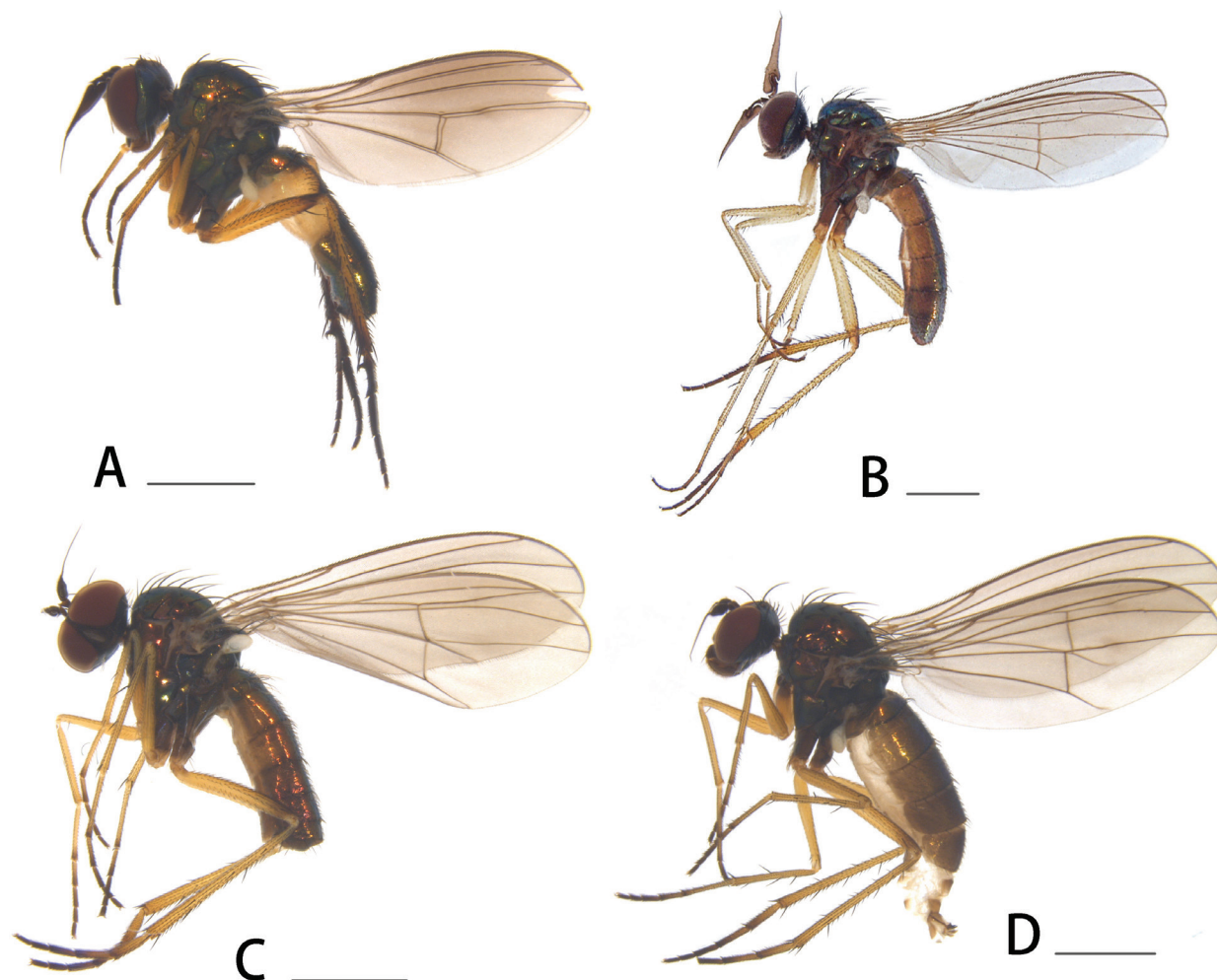


Figure 1. Habitus, lateral view **A** *S. dukha* Hollis, 1964, male **B** *S. henanense* Yang & Saigusa, 2000, male **C** *S. sinicum* sp. nov., male **D** *S. sinicum* sp. nov., female. Scale bars: 1 mm.

**Specimens examined.** CHINA: Inner Mongolia, 3 males, Mount Helan, Halawu, 13.VIII.2010, Yan Li (CAU-SYMSYN002A01-SYMSYN002A03); 2 males, Mount Jiufeng, Erdaogou, 1400–1500 m, 3.VIII.2013, Xiumei Lu (CAU-SYMSYN002B01-SYMSYN002B02).

**Habitat.** This species was collected in forests (Mount Helan and Mount Jiufeng)

**Distribution.** China (Henan, Shaanxi, Yunnan, Inner Mongolia).

***Syntormon sinicum* sp. nov.**

<https://zoobank.org/5C27BA90-3670-4A7A-9C2D-170B4C28D9E1>

Figs 1C, D, 2A–D

**Diagnosis.** Antennal scape with dorsal hairs; postpedicel 1.2 times longer than wide; arista long, much longer than postpedicel, subapical. Fore coxa yellow; mid and hind coxae black; hind femur brown at tip.

**Description. Male.** Body length 3.1–3.3 mm, wing length 3.6–4.0 mm, based on three specimens.

**Head.** Frons wide, metallic green; face narrowed downwards, with gray pollinosity. Antenna (Fig. 2A) black; scape with 2 or 3 dorsal hairs; pedicel with 1

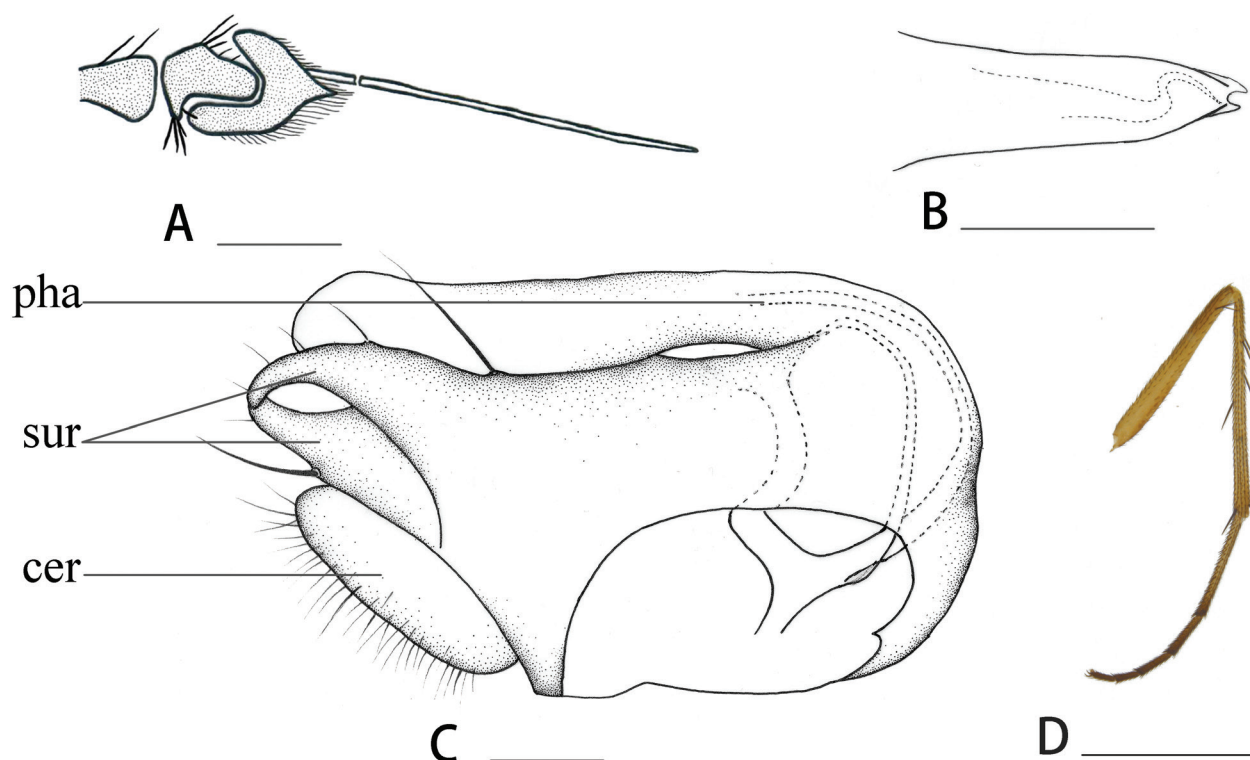


Figure 2. *Syntormon sinicum* sp. nov., male **A** antenna, lateral view **B** hypandrium, lateral view **C** genitalia, lateral view **D** mid leg, lateral view. Abbreviations: pha = phallus, sur = surstylus, cer = male cercus. Scale bars: 0.1 mm.

short, finger-like projection into postpedicel; postpedicel small, short, 1.2 times longer than wide; arista black, subapical, basal segment 0.15 times as long as apical segment. Proboscis and palpus dark brown, with black hairs, palpus with 1 blackish apical bristle.

**Thorax.** Metallic green with pale gray pollinosity. Hairs and bristles on thorax black. 5 acr bristles uniserate; 6 dc bristles, long and strong; scutellum with 2 strong bristles and 2 much smaller bristles. Propleuron with yellowish hairs.

**Legs.** Mainly yellow; coxa of LI yellow, coxa of LII and LIII black; trochanters of LII and LIII brownish yellow or dark brownish yellow; femur of LIII brown apically; tarsi of LI and LII dark brown from tip of tarsomere I onwards; leg of LIII brown to dark brown onwards entirely. Hairs and bristles on legs mostly black; coxa of LI with yellowish hairs and brown apical bristles, coxa of LII with brown hairs and bristles (1 strong long apical bristle dark brown), coxa of LIII with 1 blackish outer bristle. femora of LII and LIII each with 1 ad preapical bristle. tibia of LI with 1 pd bristle at middle and 2 short apical bristles; tibia of LII with 3 ad bristles, 1 pd bristle, 1 av bristle and 3 apical bristles; tibia of LIII with 3 ad bristles, 5 pd bristles and 4 apical bristles. Relative lengths of tibia and 5 tarsomeres of legs LI: 1.95: 1.0: 0.5: 0.4: 0.3: 0.25; LII: 2.5: 1.2: 0.6: 0.45: 0.25: 0.2; LIII: 3.4: 0.8: 0.7: 0.5: 0.35: 0.3. Wing nearly hyaline; veins blackish,  $R_{4+5}$  and M slightly convergent apically, CuAx ratio 0.55. Squama yellow with brownish yellow hairs. Halter yellow.

**Abdomen.** Metallic greenish with pale gray pollinosity. Hairs and bristles on abdomen mostly black, tergite 1 with yellow lateral hairs and bristles, hairs and bristles on tergites I–V yellow. Male genitalia (Fig. 2C): epandrium distinctly

longer than wide; surstylus with dorsal lobe wide, lateral side with long bristles, ventral lobe very wide with apical bristles; male cercus blunt at tip.

**Female.** Body length 3.0–3.1 mm, wing length 3.6–4.4 mm based on 3 specimens. Postpedicel slightly short, as long as wide, face wide.

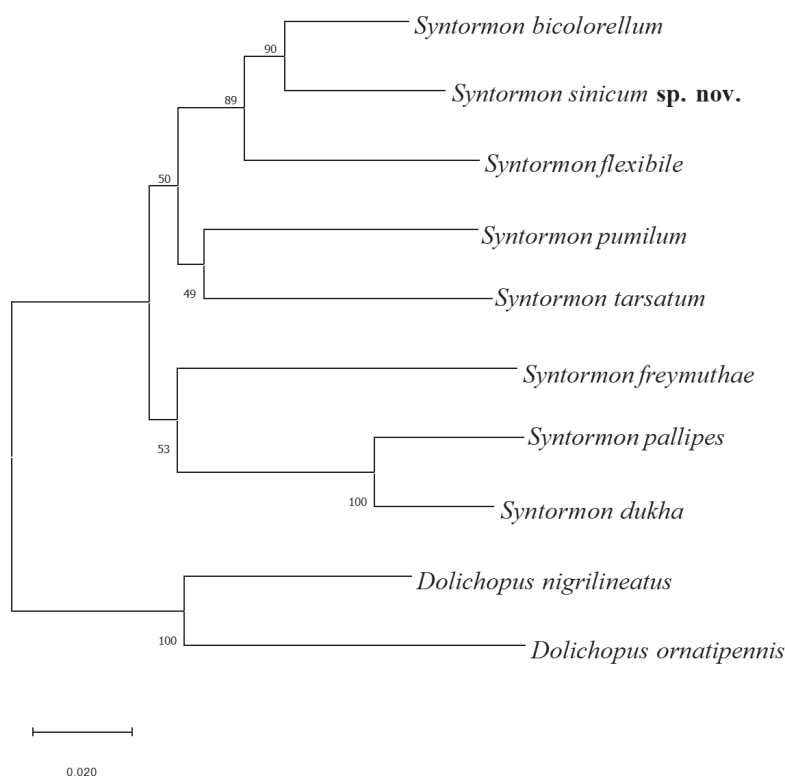
**Molecular delimitation.** In the NJ phylogenetic tree, *S. sinicum* sp. nov. is sister to *S. bicolorellum* Zetterstedt, 1843, forming a clade with *S. flexible* Becker, 1922 (Fig. 3). The interspecific genetic distance between *S. sinicum* sp. nov. and other *Syntormon* species ranged from 5.72% to 20.47% (Table 2). These results support the classification of *S. sinicum* as a separate species (Hebert et al. 2003).

**Type material.** **Holotype**, male, CHINA: Inner Mongolia, Mount Helan, Xiangchizi, 1900 m, 30.VII.2013, Xiao Zhang (CAU-SYMSYN003A01). **Paratypes**, 2 females, same data as holotype (CAU-SYMSYN003B01); 5 males 5 females, Mount Helan, Xiangchizi, 1900 m, 7.VIII.2021, Liang Wang (CAU-SYMSYN003C01-SYMSYN003C10); 3 males 3 females (OR762504), Hohhot, Xiaojinggou, 1400 m, 21.VIII.2021, Xingyang Qian (IGRCAAS-SYMSYN2A01-SYMSYN2A06).

**Habitat.** This species was collected in forests (Mount Helan, Xiaojinggou).

**Distribution.** China (Inner Mongolia).

**Remarks.** The new species is somewhat similar to *S. luchunense* Yang & Saigusa, 2001, but it can be distinguished by postpedicel, which is 1.2 times longer than wide. In *S. luchunense* Yang & Saigusa, 2001, the postpedicel is 2.5 times longer than wide (Yang et al. 2011). The new species is also somewhat similar to *S. brevicornis* Frey, 1936, but it has one long bristle on the ventral side of the mid tibia (Fig. 2D), and its fore tibia is yellow. In *S. brevicornis*, the mid tibia lacks ventral bristles, and the fore tibia is brown at the base (Frey 1936; Negrobov 1975).



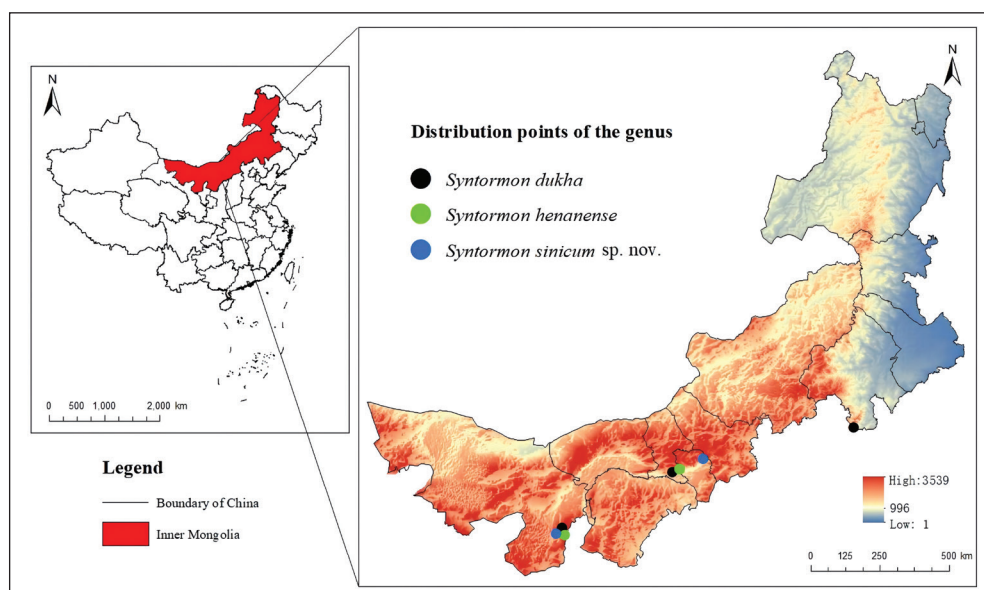
**Figure 3.** The neighbour-joining (NJ) tree of *Syntormon* species based on 10 mitochondrial COI sequences.

**Table 2.** Interspecific genetic distance between *Syntormon* COI sequences.

	<i>S. bicolorellus</i>	<i>S. flexibile</i>	<i>S. freymuthae</i>	<i>S. pallipes</i>	<i>S. pumilum</i>	<i>S. tarsatus</i>	<i>S. sinicum</i> sp. nov.	<i>S. dukha</i>	<i>D. nigrilineatus</i>	<i>D. ornatipennis</i>
<i>S. bicolorellus</i>										
<i>S. flexibile</i>	0.0805									
<i>S. freymuthae</i>	0.1258	0.1530								
<i>S. pallipes</i>	0.1330	0.1451	0.1388							
<i>S. pumilum</i>	0.1131	0.1205	0.1293	0.1420						
<i>S. tarsatus</i>	0.1062	0.1168	0.1459	0.1480	0.1132					
<i>S. sinicum</i> sp. nov.	0.0572	0.0877	0.1277	0.1276	0.1203	0.1222				
<i>S. dukha</i>	0.1131	0.1356	0.1312	0.0542	0.1365	0.1460	0.1150			
<i>D. nigrilineatus</i>	0.1612	0.1668	0.1858	0.1726	0.1760	0.1760	0.1669	0.1923		
<i>D. ornatipennis</i>	0.1825	0.1936	0.2086	0.1904	0.1943	0.2058	0.2047	0.1985	0.1145	

## Discussion

The study reports *Syntormon* from Inner Mongolia for the first time. Records of the genus in Inner Mongolia are shown in Fig. 4. The province has a temperate continental climate over more than one million square kilometers, with a variety of natural landscapes, including forests, meadows, and grasslands, which is suitable for Dolichopodidae. There are several reports of the genus from the neighbouring regions, such as *S. flexibile* Becker, 1922 from Hebei province and *S. henanense* Yang & Saigusa, 2000 and *S. pallipes* Fabricius, 1794 from Shaanxi province. In addition, *S. beijungense* Yang, 1998 and *S. pallipes* Fabricius, 1794 are recorded from Beijing and *S. pallipes* Fabricius, 1794 and *S. xinjiangense* Yang, 1999 are recorded from Xinjiang province (Yang et al. 2011). Furthermore, the previous insect investigations in Inner Mongolia mainly focused on Mount Helan (Alax League), Mount Jiufeng (Baotou), and Saihanwula (XilinGol League), and were limited in scope. Therefore, the species diversity of Sympycninae in Inner Mongolia is undoubtedly underestimated.

**Figure 4.** Distribution of *Syntormon* in Inner Mongolia.



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## Additional information

### Conflict of interest

The authors have declared that no competing interests exist.

### Ethical statement

No ethical statement was reported.

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### Author contributions

Conceptualization: NW. Data curation: XQ. Funding acquisition: NW. Project administration: NW. Resources: DY. Writing - original draft: XQ. Writing - review and editing: DY and CT.

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## References

- Bernasconi MV, Pollet M, Ward PI (2007) Molecular systematics of Dolichopodidae (Diptera) inferred from COI and 12S rDNA gene sequences based on European exemplars. *Invertebrate Systematics* 21(5): 453–470. <https://doi.org/10.1071/IS06043>
- Chursina M, Grichanov IY (2019) Analysis of the differences between *Syntormon pallipes* and *S. pseudospicatus* (Diptera: Dolichopodidae): morphological and molecular data. *Zoosystematica Rossica* 28(2): 305–316. <https://doi.org/10.31610/zsr/2019.28.2.305>
- Cumming J, Wood DM (2017) Adult morphology and terminology. In: Kirk-Spriggs AH, Sinclair BJ (Eds) *Manual of Afrotropical Diptera: Vol. 1, Suricata 4*, SANBI Publications, Pretoria, 89–133.
- Drake CM (2020) The identity of *Syntormon pseudospicatum* Strobl (Diptera, Dolichopodidae). *Dipterists Digest* 27: 61–82.
- Drake CM (2021) Comments on the taxonomic status of some British species of *Syntormon* Loew, 1857 (Diptera, Dolichopodidae). *Dipterists Digest* 28: 17–44.

- Frey R (1936) Die Dipterenfauna der Kanarischen Inseln und ihre Probleme. *Societas Scientiarum Fennica Commentationes Biologicae* 6: 1–237.
- Germann C, Pollet M, Wimmer C, Bernasconi MV (2011) Molecular data sheds light on the classification of long-legged flies (Diptera: Dolichopodidae). *Invertebrate Systematics* 25(4): 303–321. <https://doi.org/10.1071/IS11029>
- Grichanov IYa (2014) Alphabetic list of generic and specific names of predatory flies of the epifamily Dolichopodidae (Diptera). *Plant Protection News (Supplements 14)*: 1–544.
- Grichanov IYa (2021) Alphabetic list of generic and specific names of predatory flies of the epifamily Dolichopodidae (Diptera). Saint Petersburg. All-Russian Research Institute of Plant Protection 23: 188–191.
- Hebert PD, Cywinska A, Ball SL, DeWaard JR (2003) Biological identifications through DNA barcodes. *Proceedings of the Royal Society of London, Series B, Biological Sciences* 270(1512): 313–321. <https://doi.org/10.1098/rspb.2002.2218>
- Hollis D (1964) On the Diptera of Nepal (Stratiomyidae, Therevidae and Dolichopodidae). *Bulletin of the British Museum (Natural History). Entomology* 15: 83–116. <https://doi.org/10.5962/bhl.part.20536>
- Kimura M (1980) A simple method for estimating evolutionary rates of base substitutions through comparative studies of nucleotide sequences. *Journal of Molecular Evolution* 16(2): 111–120. <https://doi.org/10.1007/BF01731581>
- Kumar S, Stecher G, Li M, Knyaz C, Tamura K (2018) MEGA X: Molecular evolutionary genetics analysis across computing platforms. *Molecular Biology and Evolution* 35(6): 1547–1549. <https://doi.org/10.1093/molbev/msy096>
- Lim GS, Hwang WS, Narayanan Kutty S, Meier R, Grootaert P (2010) Mitochondrial and nuclear markers support the monophyly of Dolichopodidae and suggest a rapid origin of the subfamilies (Diptera: Empidoidea). *Systematic Entomology* 35(1): 59–70. <https://doi.org/10.1111/j.1365-3113.2009.00481.x>
- Negrobov O (1975) A review of the genus *Syntormon* Meigen (Diptera, Dolichopodidae) from Palaearctic. *Entomologicheskoe Obozrenie* 54: 652–664.
- Tamura K, Peterson D, Peterson N, Stecher G, Nei M, Kumar S (2011) MEGA5: Molecular evolutionary genetics analysis using maximum likelihood, evolutionary distance, and maximum parsimony methods. *Molecular Biology and Evolution* 28(10): 2731–2739. <https://doi.org/10.1093/molbev/msr121>
- Tonguç A, Yüzer Ö, Doğaç E (2023) Genetic diversity of the cosmopolitan long-legged fly *Syntormon pallipes* in Turkey. *Zoology in the Middle East* 69(2): 193–200. <https://doi.org/10.1080/09397140.2023.2203983>
- Trees RP (1987) The Neighbor-joining Method: A New Method for. *Molecular Biology and Evolution* 4: 406–425.
- Yang D (1999) New species of Dolichopodidae from Henan (Diptera: Empidoidea). The fauna and taxonomy of insects in Henan. In: Shen XC, Pei HC (Eds) *Insects of the Mountains Funiu and Dabie Regions*. China Agriculture, Science and Technology Press, Beijing, 189–210.
- Yang D, Zhu YJ, Wang MQ, Zhang LL (2006) World catalog of Dolichopodidae (Insecta: Diptera). China Agricultural University Press, Beijing, 502–510.
- Yang D, Zhang L, Wang M, Zhu Y (2011) *Fauna Sinica Insecta, Vol. 53. Diptera Dolichopodidae*. Science Press, Beijing, 1352–1374.