

中国𫌀翅目昆虫系统分类研究进展*

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摘要 𫌀翅目昆虫广泛分布在多种类型的水域中, 对水中的化学物质较为敏感, 是一类重要的水质监测指标生物。本文回顾了我国𫌀翅目分类的研究历史, 其中 Klapálek、Navás、Banks 及胡经甫对我国早期的𫌀翅目分类研究做出了重要贡献。同时总结了我国𫌀翅目物种多样性的研究现状, 即我国已记录𫌀翅目 10 科、66 属、657 种 (包括台湾分布的 31 种、香港分布的 4 种), 约占世界已记录种类的 17.8%; 我国学者定名的种类占全国记录种类的 74.7%, 其中 1949 年建国后的 70 年间定名的种类占 67.4%。此外, 还对我国𫌀翅目昆虫的地理分布及系统发育研究进行了总结分析, 其中我国学者共测序𫌀翅目昆虫全线粒体基因组 60 种, 占已测序种类的 90.9%, 为𫌀翅目的分子系统发育研究积累了重要分子数据; 同时利用这些数据对翅目的分子系统发育进行研究, 取得了较好的研究结果。最后, 文章分析了我国翅目系统分类研究存在的主要问题, 并对未来的研究进行了展望。

关键词 𫌀翅目; 系统分类; 系统发育; 物种多样性; 研究进展; 中国

Advances in Plecoptera systematics in China

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Abstract The Plecoptera are widely distributed aquatic insects that are sensitive to chemical substances in water, and therefore important indicators of water quality. This paper reviews the history of taxonomic research on the Plecoptera in China, a field in which Klapálek, Navás, Banks and Wu Chenfu have made important contributions to early taxonomic research. In addition, the species diversity of Chinese Plecoptera is summarized; there are a total of 657 species (including 31 in Taiwan and 4 in Hong Kong) representing 10 families and 66 genera, accounting for about 17.8% of all species worldwide. Species named by Chinese scholars account for 74.7% of all species recorded in China, and species named in the 70 years since the founding of the People's Republic of China in 1949 account for 67.4% of this total. In addition, the geographical distribution and phylogenetics of the Plecoptera in China are summarized and analyzed. Chinese scholars have sequenced the complete mitochondrial genomes of 60 species, 90.9% of all species that have been sequenced, and have accumulated important molecular data for molecular phylogenetic studies of this group. These data have allowed good progress to be made on the molecular phylogeny of the Plecoptera. The main obstacles impeding progress on Plecoptera systematics in China are analyzed, and prospects for future research discussed.

Key words Plecoptera; systematics; phylogeny; species diversity; research progress; China

翅目 (Plecoptera) 昆虫, 又称石蝇、

态系统中重要的一类水生昆虫,同时也是一些珍稀鱼类的食料(Hynes, 1976; Ward, 1992; Che et al., 2013)。此外,该类昆虫较广泛地分布在多种类型的水域中,并且对水中的化学物质反应较为敏感,是一类重要的水质监测指标生物,也是国际上常用的EPT(Ephemeroptera蜉蝣目、Plecoptera𫌀翅目和Trichoptera毛翅目)水质监测的三大水生昆虫之一(Morse et al., 1994; Rosenberg et al., 2009)。因此,开展该类昆虫的分类鉴定研究具有重要的实际意义。

目前已知𫌀翅目昆虫在除南极以外的世界各大陆上均有分布,但主要分布于东洋区、古北区及新北区(Illies, 1965; Zwick, 1973, 2000; 杜予州, 1999b)。由于我国世界动物地理区所处的重要位置以及横跨古北和东洋两大动物地理区,物种资源极为丰富,因此开展我国𫌀翅目昆虫区系分类及系统演化研究,可为世界动物地理学的研究提供可靠的区域性资料,也能更好地为水质环境监测提供服务。本文就我国翅目昆虫系统分类研究的历史、目前的研究现状以及存在的主要问题进行总结,并对未来的研究进行展望。

1 我国

个种 (Chao, 1947)。在 20 世纪 30-40 年代, 由于胡经甫教授的杰出工作, 使我国的𫌀翅目分类研究处于世界先进水平。

但是, 自胡经甫教授去世后的近 17 年 (1973-1990), 国内对𫌀翅目的研究无人问津, 使我国的𫌀翅目分类研究处于停滞状态。20 世纪 90 年代初, 杨集昆教授及杨定博士发表了自胡经甫教授去世后由国内学者撰写的 2 篇𫌀翅虫的分类论文 (杨集昆和杨定, 1990a, 1990b), 重新开启了国内学者对𫌀翅目的分类研究; 此后, 他们两人共发表𫌀翅目分类论文 17 篇, 描述定名𫌀翅目 1 新属、35 新种 (表 1)。此外, 20 世纪 90 年代中后期, 在周尧教授指导下, 杜予州博士开始对中国𫌀翅目分类进行系统研究 (杜予州, 1995, 1999a, 1999b, 1999c; 2000a, 2000b; 杜予州和周尧, 1998, 1999; 杜予州等, 1999; Du et al., 1999)。进入 21 世纪, 杨定教授和杜予州教授培养了一批博士和硕士从事翅目的分类研究, 加之任东教授团队的翅目化石种类的研究, 形成了目前我国翅目分类研究的 3 个团队, 对推动我国

表 1 中国记载数目各科种类数量及定名人
Table 1 Describers and number of Chinese species described in each families of Plecoptera

(续表 1) (Table 1 continued)

定名人 Describer	科 Family										合计 Total
	黑𫌀科 Capniidae	带.DependencyInjection	叉.DependencyInjection	卷.DependencyInjection	大.DependencyInjection	扁.DependencyInjection	刺.DependencyInjection	.DependencyInjection	网.DependencyInjection	绿.DependencyInjection	
Sivec & Zhiltzova											1
Sivec & Zwick											(1)
Zhiltzova & Baumann	1										1
Zhiltzova & Levantova											1
Zhiltzova & Zwick											1
Tiemro de Figuerola & Fochetti											1
Wu (胡经甫)	44	6	4	3	64	7	7	3	131	1	1
Wu(胡经甫)&Classen			1	1	10	4	5		15		15
Chu (朱元鼎)	1	1	1								8
Chao (赵修复)	1	11		2	19	2					2
Yang (杨集昆) & Yang (杨定)				2	6						35
Du (杜予州) & Chou (周尧)											6
Du (杜予州), Wang (王志杰), Qian (钱昱含), Chen (陈志腾)等人	8	2	32	12	5	1	40	8	6	114	
Du (杜予州) & Sivec								5			5
Yang (杨定), Li (李卫海), Zhu (祝芳) 等人	2		84	15		1	43	1	5	151	
Li (李卫海), Yang (杨定) & Muñáni				11	2	2		8		24	
Li (李卫海) & DeWalt								1			1
合计 Total	19	4	191	67	1	18	12	277	45	25	659

表中的物种数主要依据 Wu, 1938; Claassen, 1940; Illies, 1966; Zwicky, 1973; Sivec *et al.*, 1997; 杜予州, 1999; Du *et al.*, 1999; 杨定和李卫海, 2018; 杨定等, 2015; DeWalt *et al.*, 2020 以及有关学者发表的论文所统计；统计时间截止 2020 年 2 月底。括号内的数字为台湾地区的种类数。

The number of the species in table is mainly based on the statistics of Wu, 1938; Claassen, 1940; Illies, 1966; Zwicky, 1973; Sivec *et al.*, 1997; Du, 1999c; Du *et al.*, 1999; Yang and Li, 2018; Yang *et al.*, 2020 and the papers published by the relevant scholars. The statistical time ended at the end of February 2020. The numbers in brackets are the number of the species in Taiwan of China.

襀翅目 16 科、66 种的全线粒体基因组, 其中我国学者(主要是杨定和李卫海团队及杜予州团队)测序 60 种(9 种与国外学者合作)、德国和美国学者各测序 2 种, 加拿大和新西兰学者各测定了 1 种(表 2)。中国学者在襀翅目昆虫的全线粒体基因组数据积累中做出了重要贡献, 也为更好地开展襀翅目的分子系统学研究奠定了基础。

3.2 襀翅目在昆虫纲的系统地位

襀翅目隶属六足总纲、昆虫纲、复新翅类(Polyneoptera), 是新翅类昆虫中较早分化出来的一个相对原始的目, 而且该目的单系性也得到几个共同衍征的支持(Zwick, 2000, 2009)。早期基于形态学的襀翅目系统地位的研究较多, 多数学者认为襀翅目与直翅类或纺足目的亲缘关系最近(Crampton, 1938; Martynov, 1938; Henning, 1953, 1981; Bitsch and Ramond, 1970; Kristensen, 1981), 也有少数学者认为与蛩蠊目最近(Rasnitsyn, 1976; Rodendorf and Rasnitsyn, 1980)。随着分子数据的应用, 基于分子标记和形态数据的襀翅目系统地位的研究报道也日益增多, 但得出的结果仍有较大分歧(Liu and Beckenbach, 1992; Burmester et al., 1998; Flook and Rowell, 1998; Wheeler et al., 2001; Rasmus et al., 2002; Terry and Whiting, 2005; Yoshizawa and Johnson, 2005; Trautwein et al., 2012; Cameron, 2014; Misof et al., 2014)。

近年来, 我国学者应用分子标记及全线粒体基因组数据进行襀翅目系统地位的探讨(Zhang and Zhou, 2008; Xie et al., 2009; Qian et al., 2014; Wu et al., 2014; Song et al., 2016), 如 Wu 等(2014)和 Song 等(2016)研究的结果与其他多数学者研究的结果基本一致, 即襀翅目与革翅目的亲缘关系最近。总之, 襀翅目的系统地位仍然存在较大的争议, 有必要继续发掘可靠的形态和分子数据进行更深层次的研究。

3.3 襀翅目的系统发育研究

基于形态特征的襀翅目高级阶元系统发育与进化国外已有较多的研究报道(Tillyard,

1921; Ricker, 1950; Illies, 1965; Kawai, 1967; Zwick, 1973, 1980, 2000, 2009; Nelson, 1984, 1988; Uchida and Isobe, 1989)。在国内, 杜予州(1995, 1999a)应用支序分析法对襀亚科 18 个属的系统发育关系进行了初步研究, 分析了各属之间的系统发育关系。这也是国内仅有的 1 篇基于形态学的襀翅目系统发育研究报道。这方面的研究仍是国内襀翅目研究的薄弱环节, 有待进一步加强。

近年来, 应用分子标记, 特别是应用全线粒体基因组进行昆虫的系统发育研究越来越多(Cameron, 2014)。基于形态和分子数据的襀翅目高级阶元系统发育已有一些研究报道(Thomas et al., 2000; Terry, 2004; Weiss et al., 2012; Mynott et al., 2011; Davis, 2013; McCulloch et al., 2016), 这些研究对了解襀翅目高级阶元的亲缘关系有了进一步认识。我国学者在测序襀翅目昆虫全线粒体基因组的基础上, 也对襀翅目的分子系统发育及进化做了大量研究工作, 在探讨襀翅目的科级系统发育关系方面取得了较好的研究成果(Chen and Du, 2017a, 2017b, 2018; Chen et al., 2018; Wang et al., 2017, 2018, 2019; Cao et al., 2019a, 2019b; Ding et al., 2019; Shen and Du, 2020)。

4 展望

通过多年的调查发现, 我国的襀翅目昆虫多样性极为丰富。但由于襀翅目不同类群的区系分布特点不同, 加之调查采集的地区不平衡, 如我国东北地区、青藏高原、横断山脉以及台湾等地区的调查较为薄弱, 因此各类群记录的种类数量也不平衡, 如襀科已记录 23 属、275 种, 叉襀科 7 属、191 种, 而大襀科和带襀科仅记录 1 种和 2 种; 我国台湾也仅记录由外国人定名的 31 种(表 1)。此外, 由于襀翅目昆虫不同类群的生物学特性, 特别是成虫的羽化时间的不同(Harper, 1976; Hynes, 1976), 需要调查采集的时间也不同, 如黑襀科和带襀科的种类成虫主要在冬季羽化(夏季在高海拔雪山和冷凉地区也可羽化), 故又称冬襀(Winter stonefly),

表 2 已测序全线粒体基因组的𫌀翅目昆虫种类
Table 2 Species of the complete mitochondrial genome sequenced in Plecoptera

科 Family	种 Species	序列登录号 Sequence accession number	序列公 布时间 Sequence release time	登录 国家 Login sequence country	论文作者 Author of paper	论文发表杂志 Journal of publication
𫌀科 Perlidae	<i>Dinocras cephalotes</i> <i>Kamimuria wangi</i>	KF484757 KC894944	2013.12.6 2014.4.25	德国 中国	Elbrecht V, Poettker L, John U, Leese F	Mitochondrial DNA Part A, 2015, 26: 469–470. PLoS ONE, 2014, 9(1): e86328.
	<i>Togoperla</i> sp.	KM409708	2014.10.26	中国	Qian YH, Wu HY, Ji XY, Du YZ	Mitochondrial DNA Part A, 2016, 27: 1703–1704.
	<i>Kamimuria chungnanshana</i>	KT186102	2015.9.29	中国	Wang K, Wang Y, Yang D	Mitochondrial DNA Part A, 2016, 27: 3810–3811.
	<i>Acroneuria hainana</i>	KM199685	2015.2.12	中国	Huang M, Wang Y, Liu X, Li W, Kang Z, Wang K, Li X, Yang D	Gene, 2015, 557(1): 52–60.
	<i>Claassenia</i> sp.	MN419914	2020.1.15	中国	Chen M, Cao J, Li W, Wang Y	Mitochondrial DNA Part B, 2019, 4(2): 3790–3791.
	<i>Flavoperla</i> sp.	MN419916	2020.1.15	中国	Wang Y, Cao J, Chen M, Li W	Mitochondrial DNA Part B, 2019, 4(2): 3902–3903.
	<i>Neoperllops gressitti</i>	MN400756	2020.1.15	中国	Zhang G, Wang Y, Cao J	Mitochondrial DNA Part B, 2019, 4(2): 3324–3325.
	<i>Kamimuria klapaleki</i>	MN400755	2020.1.15	中国	Chen J, Cao J, Li W, Wang Y	Mitochondrial DNA Part B, 2019, 4(2): 3416–3417.
	<i>Eurocorema hochii</i>	MK905888	2020.1.15	中国	Liu Z, Wang Y, Li W, Cao J	Mitochondrial DNA Part B, 2019, 4(2): 2690–2691.
	<i>Niponiella limbatella</i>	MK686067	2020.1.15	中国	Li J, Cao J, Wang Y, Kong F	Mitochondrial DNA Part B, 2019, 4(1): 1666–1667.
	<i>Sinacneuria dabieshana</i>	MK492253	2020.1.15	中国	Cao J, Li W, Wang Y	Mitochondrial DNA Part B, 2019, 4(1): 1327–1328.
	<i>Caleneuria stigmatica</i>	MG677941	2020.1.15	中国	Cao J, Wang Y, Li N, Li W, Chen X	Mitochondrial DNA Part B, 2019, 4(1): 553–554.
	<i>Caroperla siveci</i>	MG677942	2019.8.31	中国	Cao J, Wang Y, Li N, Li W, Chen X	Mitochondrial DNA Part B, 2019, 4(1): 553–554.
	<i>Neoperla</i> sp.	KX091859	2020.1.1	中国	Song F, Liu Y, Jiang P, Li H, Cai W	Unpublished
	<i>Neoperla ignacivici</i>	KX091858	2020.1.1	中国	Song F, Liu Y, Jiang P, Li H, Cai W	Unpublished
	<i>Flavoperla biocellata</i>	MK905206	2020.3.9	中国	Shen Y, Du YZ	Peer J, 2020, 8: e8762.
	<i>Isoperla bilineata</i>	MF716959	2018.7.16	中国	Chen ZT, Zhao MY, Xu C, Du YZ	Int. J. Biol. Macromol., 2018, 111: 542–547.
	<i>Isoperla eximia</i>	MG910457	2018.7.31	中国	Wang Y, Cao J, Li WH	Gene, 2018, 675: 254–264.
	<i>Perloides</i> sp.	MF197377	2018.7.25	中国	Chen ZT, Zhao MY, Xu C, Du YZ	Int. J. Biol. Macromol., 2018, 111: 542–547.
	<i>Pseudomegarctys japonica</i>	MG910458	2018.7.31	中国	Wang Y, Cao J, Li WH	Unpublished
	<i>Suwalla telekensis</i>	MF198253	2018.4.23	中国	Wang Y, Cao J, Li WH	Int. J. Mol. Sci., 2018, 19(3): 680.
网𫌀科 Perlodidae	<i>Suwalla bimaculata</i>	KX091865	2020.1.1	中国	Cao J, Wang Y, Chen M, Yuan M, Li W	Unpublished
	<i>Suwalla bimaculata</i>	MN121757	2020.1.15	中国	Cao J, Wang Y, Chen M, Yuan M, Li W	Mitochondrial DNA Part B, 2019, 4(2): 2828–2829.
绿𫌀科 Chloroperlidae	<i>Sweltsa longistyla</i>	KM216826	2015.2.12	中国	Chen ZT, Du YZ	Gene, 2015, 558(1): 82–87.

(续表2) (Table 2 continued)

科 Family	种 Species	序列登录号 Sequence accession number	序列公布时间 Sequence release time	登录国家 Login country	论文作者 Author of paper	论文发表杂志 Journal of publication
大𫌀科 Pteronarcidae	<i>Pteronarcys princeps</i> <i>Pteronarcella badia</i>	AY687866 KU182360	2016.7.26 2015.12.27	加拿大 美国	Stewart JB, Beckenbach AT Sproul JS, Houston DD, Nelson CR, Evans RP, Crandall KA, Shiozawa DK	Genome, 2006, 49(7): 815–824. BMC Evolutionary Biology, 2015, 15(1): 279.
刺𫌀科 Stylopertidae	<i>Styloperla spinicercia</i> <i>Cerconychia sapa</i>	KX845569 MF100783	2017.5.22 2019.12.12	中国	Wang Y, Cao J, Li W	Zootaxa, 2017, 4243 (1): 125–138.
扁𫌀科 Peltopertidae	<i>Styloperla</i> sp. <i>Soliperla</i> sp. <i>Peltoperlopsis cebuano</i>	KR088971 MF716958	2015.9.26 2018.7.16	中国	Chen ZT, Wu HY, Du YZ	Conservation Genetics Resources, 2018, 10(2): 145–148. Mitochondrial DNA, 2015, 27(4): 1–2.
黑𫌀科 Capniidae	<i>Microperla geei</i> <i>Cryptoperla</i> sp. <i>Apteroperlta tikumana</i> <i>Capnia zijinshana</i>	MK387068 MN096323 KC952026 KR604721	2020.1.15 2019.12.21 2014.3.24 2015.8.1	中国	Cao J, Li W, Yan F, Wang Y Cao J, Wang Y, Wei X, Chen S, Li W Wu HY, Ji XY, Yu WW, Du YZ Zhou C, Tan M, Du S, Zhang R, Machida R, Zhou X	Int. J. Boil. Macromol., 2017, 111: 542–547. Mitochondrial DNA Part B, 2019, 4 (1): 1103–1104. Mitochondrial DNA Part B, 2019, 4 (2): 2679–2680. Gene, 2014, 537(2): 177–183. Mitochondrial DNA Part A, 2016, 27(4): 1940–1944. Journal of Asia-Pacific Entomology, 2017, 20(2): 305–312.
带𫌀科 Taeniopterygidae	<i>Mesocapnia daxingana</i> <i>Mesocapnia arizonensis</i>	KY568983 KP642637	2019.12.12 2015.3.24	中国 德国	Wang Y, Cao J, Li W, Chen X Elbrecht V, Leese F	Conservation Genetics Resources, 2017, 9(4): 639–642. Mitochondrial DNA Part A, 2016, 27(5): 3365–3366.
叉𫌀科 Nemouridae	<i>Taeniopteryx ugola</i> <i>Doddgia occidentalis</i> <i>Nemoura rankinensis</i> <i>Amphinemura longispina</i>	MG589786 MG589787 KY940360 MH085446	2018.7.12 2018.5.12 2018.3.27 2019.9.1	中国	Chen ZT, Du YZ Chen ZT, Du YZ Chen ZT, Du YZ Cao JJ, Wang Y, Li WH	Int. J. Boil. Macromol., 2018, 111: 70–76. Int. J. Mol. Sci., 2018, 18(5): 996. Int. J. Boil. Macromol., 2019, 138: 292–301. Unpublished
	<i>Amphinemura yao</i> <i>Indonemoura jacobsoni</i> <i>Indonemoura nohirei</i>	MH085447 MH085448 MH085449	2019.9.1 2019.9.1 2019.9.1	中国	Wang Y, Cao JJ, Li WH Wang Y, Cao JJ, Li WH Wang Y, Cao JJ, Li WH	Unpublished Unpublished Unpublished
	<i>Mesonemoura metafiligera</i> <i>Mesonemoura tritaenia</i> <i>Protoneamura kohnae</i> <i>Protoneamura orbiculata</i>	MH085450 MH085451 MH085452 MH085453	2019.8.31 2019.8.31 2019.9.1 2019.9.1	中国	Cao JJ, Wang Y, Huang YR, Li WH Wang Y, Cao JJ, Li WH Wang Y, Cao JJ, Li WH Wang Y, Cao JJ, Li WH	Zookeys, 2019, 835: 43–63. Unpublished Unpublished Unpublished

(续表 2) (Table 2 continued)

科 Family	种 Species	序列登录号 Sequence accession number	序列公 布时间 Sequence release time	登录 国家 Login sequence country	论文作者 Author of paper	论文发表杂志 Journal of publication
叉襀科 Nemouridae	<i>Sphaeronomoura grandicuda</i>	MH085454	2019.9.1	中国	Wang Y, Cao JJ, Li WH	Unpublished
	<i>Sphaeronomoura hamistyla</i>	MH085455	2019.9.1	中国	Wang Y, Cao JJ, Li WH	Unpublished
	<i>Lednia tumana</i>	MH1374046	2019.4.30	美国	Hotaling S, Kelley JL, Weistrock DW	Aquatic Insects, 2019, 40(4): 362–369.
	<i>Indonemoura auriformis</i>	MN419915	2020.1.15	中国	Chen M, Wang Y, Chen J, Cao J	Mitochondrial DNA Part B, 2019, 4(2): 3392–3393.
	<i>Nemoura papilla</i>	MK290826	2020.1.15	中国	Cao J, Wang Y, Ma G, Li W	Mitochondrial DNA Part B, 2019, 4(1): 806–807.
	<i>Sphaeronomoura hainana</i>	MK111420	2019.12.21	中国	Ding SM, Li WH, Wang Y, Cameron SL, Muranyi D, Yang D	Mol. Phylogenet. Evol., 2019, 135: 123–135.
背襀科 Notonemouridae	<i>Amphinemura</i> sp.	KX091847	2020.1.1	中国	Song F, Liu YQ, Jiang P, Li H, Cai WZ	Unpublished
	<i>Neonemura barrosi</i>	MK111418	2019.4.7	中国	Ding S, Li W, Wang Y, Cameron SL, Muranyi D, Yang D	Mol. Phylogenet. Evol., 2019, 135: 123–135.
卷襀科 Leuctridae	<i>Rhopalopsole bulbifera</i>	MK111419	2019.4.7	中国	Ding S, Li W, Wang Y, Cameron SL, Muranyi D, Yang D	Mol. Phylogenet. Evol., 2019, 135: 123–135.
裸襀科 Scopuridae	<i>Leuctra</i> sp.	MK568475	2019.11.25	中国	Shen Y, Du YZ	Zootaxa, 2019, 4671(4): 571–580.
	<i>Scopura longa</i>	MH510071	2019.1.28	中国	Wang Y, Cao JJ, Li N, Ma GY, Li WH	Int. J. Biol. Macromol., 2019, 122: 893–902.
纬襀科 Gripopterygidae	<i>Zelandoperla fenestrata</i>	KY522907	2017.5.27	新西兰	Veale AJ, Deaden PK, Waters JM	Unpublished
	<i>Antarctoperla michaelensi</i>	MK111413	2019.3.27	中国	Ding S, Li W, Wang Y, Cameron SL, Muranyi D, Yang,D	Mol. Phylogenet. Evol., 2019, 135: 123–135.
始襀科 Diamphipnoidae	<i>Diamphipnoa annulata</i>	MK111416	2019.4.7	中国	Ding S, Li W, Wang Y, Cameron SL, Muranyi D, Yang,D	Mol. Phylogenet. Evol., 2019, 135: 123–135.
	<i>Diamphipnopsis</i> sp.	MK111417	2019.4.7	中国	Ding S, Li W, Wang Y, Cameron SL, Muranyi D, Yang,D	Mol. Phylogenet. Evol., 2019, 135: 123–135.
原襀科 Eustheniidae	<i>Neuroperla schedingii</i>	MK111415	2019.3.27	中国	Ding S, Li W, Wang Y, Cameron SL, Muranyi D, Yang,D	Mol. Phylogenet. Evol., 2019, 135: 123–135.
澳襀科 Austroperlidae	<i>Klapopteryx armillata</i>	MK111414	2019.3.27	中国	Ding S, Li W, Wang Y, Cameron SL, Muranyi D, Yang D	Mol. Phylogenet. Evol., 2019, 135: 123–135.

由于该类襀翅虫羽化的特殊性,对这类昆虫的调查采集极为困难,这也是我国黑襀科和带襀科昆虫种类记录较少的原因。因此,今后一段时间需要加强对过去调查较少的地区进行系统调查采集,同时还要根据不同襀翅虫类群的羽化特性在不同的时间和季节进行采集调查,这样才能完整地了解我国襀翅目昆虫物种多样性及地理分布特征。

虽然我国学者积累了大量襀翅昆虫的全线粒体基因组数据,并根据这些数据对襀翅目科级分子系统发育进行了分析,取得了较好的研究成果,但是基于形态特征和分子数据结合的襀翅目高级阶元和科级以下分类阶元的系统发育研究至今未见报道;此外,在襀翅目昆虫的生物地理学和系统演化方面我国也缺乏研究,因此加强这些方面的深入研究是我国襀翅目分类学者今后努力的方向。

早期我国的襀翅目昆虫分类研究主要是由外国人做的,所有模式标本分散存放在国外有关单位、私人收藏或下落不明。此外,我国学者胡经甫教授早期定名的绝大多数襀翅目昆虫模式标本已因战争丢失(胡经甫,1962),即使在1962年和1973年发表定名的新种模式标本也因“文革”时期管理不善而损坏或丢失(杜予州于1994和1998年两次对存放在中国科学院动物研究所的襀翅目昆虫模式标本作过检查);朱元鼎(1928a, 1928b, 1928c, 1929)和赵修复描述定名的模式标本为私人收藏,现在也下落不明(Chao, 1947)。此外,早期的襀翅目分类学者所描述定名的种类,主要描述及绘图依据的是干标本,而襀翅虫身体较软,标本干后容易变形,并且在鉴定时未进行阳茎解剖,描述和绘制的特征图也很简单,加之模式标本丢失或保存在国外,因此给这些早期定名的种类鉴定造成了很大困难。因此,需要对现在仍然保存的襀翅目昆虫的模式标本,特别是还保存在科学院动物所和国外的模式标本重新进行整理和补充描述,同时尽可能在已丢失模式标本的模式产地采集到相同种类的标本,重新指定新模,以提高其科学价值。

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