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Revision of *Pazala* Moore, 1888: The *Graphium* (*Pazala*) *mandarinus* (Oberthür, 1879) Group, with Treatments of Known Taxa and Descriptions of New Species and New Subspecies (Lepidoptera: Papilionidae)

SHAO-JI HU^{1, 2, 3}, ADAM M. COTTON⁴, FABIEN L. CONDAMINE⁵, KUANG DUAN³, RONG-JIANG WANG⁶, YU-FENG HSU⁷, XIN ZHANG⁸ & JUN CAO⁹

¹Yunnan Key Laboratory of International Rivers and Transboundary Eco-security, Yunnan University, Kunming, 650500, China ²Institute of International Rivers and Eco-security, Yunnan University, Kunming, 650500, China

³School of Agriculture, Yunnan University, Kunming, 650500, China

⁴86/2 Moo 5, Tambon Nong Kwai, Hang Dong, Chiang Mai, Thailand

⁵CNRS, UMR 5554 Institut des Sciences de l'Evolution, Université de Montpellier, Place Eugène Bataillon 34095 Montpellier, France ⁶College of Life Sciences, Peking University, Beijing, 100871, China

⁷Department of Life Sciences, National Taiwan Normal University, Taipei, 116, Taiwan, China

⁸Laboratory for Animal Genetic Diversity and Evolution of Higher Education in Yunnan Province, Yunnan University, Kunming, 650091, China

⁹Laboratory of Ecology and Evolutionary Biology, Yunnan University, Kunming, 650500, China ¹⁰Corresponding author. E-mail: juncao@vip.163.com

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Abstract

The previously recognised closely related species *Graphium (Pazala) mandarinus* (Oberthür, 1879) and *G. (P.) sichuanica* (Koiwaya, 1993) are shown to comprise seven species as a result of both molecular and morphological analysis. Molecular dating analysis is also performed on the *mandarinus* group in order to investigate the divergence time of the taxa. Two

taxa, G(P) garhwalica (Katayama, 1988) stat. nov. and G(P) paphus (de Nicéville, 1886) stat. nov., are raised from subspecific to specific status; G(P) hoeneanus Cotton & Hu nom. nov., stat. rev. is separated from sichuanica at species level; and two previously unrecognised new species, G(P) daiyuanae Hu, Zhang & Cotton **sp. nov.** and G(P) confucius Hu, Duan & Cotton **sp. nov.** are described from Vietnam and China respectively, the latter being sympatric with nominate G(P) mandarinus. The identity of the lectotype of G(P) mandarinus is confirmed and a lectotype is designated for the taxon Papilio Glycerion Gray, 1831. A new subspecies of G(P) mandarinus is described from western Yunnan and northern Myanmar, G(P) mandarinus stilwelli Cotton & Hu ssp. nov.

Key words: Leptocircini, Pazala, new species, India, China, Vietnam

Introduction

The genus *Graphium* Scopoli, 1777 (Papilionidae: Leptocircini) includes a number of species groups found in Asia, Australasia and Africa, and has relatives in Europe (genus *Iphiclides* Hübner, [1819]), Australia (genus *Protographium* Munroe, 1961) and the Americas (genus *Eurytides* Hübner, [1821]). Very few studies have been published on the genus as a whole in recent times, with major works dating from the 1980s. For instance, Saigusa *et al.* (1982) only studied the Asian species in subgenus *Graphium*, Hancock (1983) treated the genus as comprising 89 species in 4 subgenera, and Miller (1987) also recognised 4 subgenera without lower classification. This study concentrates on a single subgenus within the genus, and will be split into at least two parts.

The swordtail subgenus *Pazala* Moore, 1888 is a small group of Asian butterflies considered to comprise seven species enumerated by Racheli & Cotton (2009) as *eurous* (Leech, 1893), *mandarinus* (Oberthür, 1879), *sichuanica* (Koiwaya, 1993), *mullah* (Alphéraky, 1897), *alebion* (Gray, [1853]), *tamerlanus* (Oberthür, 1876) and *parus* (de Nicéville, 1900). Among these species, G(P) mandarinus (Oberthür, 1879) has the widest distribution range from N.W. India to C. China as well as N. Indochina (Racheli & Cotton 2009). Five subspecies have been recognised in G(P) mandarinus over the last 137 years, three of which were described in the 1980s (de Nicéville 1886; Murayama 1982; Okano 1986; Katayama 1988). Koiwaya (1993) described an allied species, G(P) sichuanica, from W. China based on differences of wing pattern and genitalic structures. Apart from its distinguishing characters, G(P) sichuanica shares a unique character with G(P) mandarinus — the two discal black bands on the hindwing underside twisted into an "8"-shaped pattern with a short line between the two rings, resembling a pair of glasses in appearance. Based on this shared character and the similarity of other morphological characters as well as genitalic structures, the authors use the term 'mandarinus group' to represent these taxa in the present study.

G. (P.) mandarinus is a variable species, with the identity of some populations previously uncertain (Cotton & Racheli 2007; Racheli & Cotton 2009). During DNA barcoding research on Chinese swallowtail butterflies in 2011, the authors of this study encountered a very strange anomaly, namely the samples of G (P.) mandarinus from a single locality in W. Sichuan split into two clades in the phylogenetic tree, implying that a cryptic species might be present in the samples. However, all co-authors were unable to draw any conclusions due to morphological similarity and limited samples available at that time. Subsequently in 2015, two specimens which appeared to be G (P.) mandarinus collected from Sapa, N. Vietnam were flagged as a distinct species by DNA barcoding analysis, different from both G (P.) mandarinus and the previously unidentified taxon. As a result, we decided to re-examine specimens in collections while obtaining more samples from different localities, in an attempt to form a better understanding of the mandarinus group.

After examining a long series of specimens across the distribution range of the *mandarinus* group with associated molecular data, seven species are now recognised within this group, including two new species and three previously recognised taxa elevated to specific status, one of which requires a new name. The name G(P) *mandarinus* (Oberthür, 1879) is assigned to the correct species after examining the lectotype specimen in the BMNH (Natural History Museum, London) collection and a lectotype is designated for a historically problematic taxon. The westernmost population of the true G(P) *mandarinus* is described as a new subspecies. The findings, a revision of the *mandarinus* group with descriptions of new taxa and a replacement name, are presented in this study.

Materials and methods

Taxon sampling. A large number of specimens of all taxa in the *mandarinus* group were examined and sampled from the authors' private collections, other private collections and specimen depositories of academic institutions, with permission. All the extensive material of *mandarinus* group taxa examined, including type material, together with the names and abbreviations of depositories, is documented in Appendix 1 at the end of this work. Subspecific designations for *Pazala* species followed Racheli & Cotton (2009). Sixty-two specimens of *mandarinus* group taxa were used in molecular analysis and the GenBank accession numbers are listed in Table 1. Since this research focuses on the *mandarinus* group, only the nominotypical subspecies of the other previously recognised *Pazala* species were included in the present study, and specimens of these taxa are not listed in Appendix 1.

Taxon (sample code)	Locality	Accession No.
G (P.) garhwalica (UT1)	Joshimath, Uttaranchal, India	MG197758
G. (P.) paphus (MT1–5)	Metok, Tibet, China	MG197757
G. (P.) sichuanica (LHG1–3)	Laohegou, Sichuan, China	MG197767
G. (P.) sichuanica (TJH1)	Tangjiahe, Sichuan, China	MG197768
G. (P.) sichuanica (TJH2)	Tangjiahe, Sichuan, China	MG197767
G. (P.) hoeneanus (ZY1)	Ziyuan, Guangxi, China	MG197762
G. (P.) hoeneanus (ZY2)	Ziyuan, Guangxi, China	MG197763
G. (P.) hoeneanus (LG1)	Lingui, Guangxi, China	MG197764
G. (P.) hoeneanus (LG2)	Lingui, Guangxi, China	MG197765
G. (P.) hoeneanus (LG3)	Lingui, Guangxi, China	MG197766
G. (P.) hoeneanus (TMS1)	W. Tianmu Shan, Zhejiang, China	MG197759
G. (P.) hoeneanus (TMS2)	W. Tianmu Shan, Zhejiang, China	MG197760
G. (P.) hoeneanus (TMS3)	W. Tianmu Shan, Zhejiang, China	MG197761
G. (P.) daiyuanae (SP1–4)	Sapa, Lao Cai, Vietnam	MG197773
G. (P.) daiyuanae (SP5)	Sapa, Lao Cai, Vietnam	MG197774
G. (P.) daiyuanae (SP6)	Sapa, Lao Cai, Vietnam	MG197775
G. (P.) confucius (HY1)	Hanyuan, Sichuan, China	MG197770
G. (P.) confucius (EB1)	Ebian, Sichuan, China	MG197770
G. (P.) confucius (BX1–2)	Baoxing, Sichuan, China	MG197769
G. (P.) confucius (DC1, DC3)	Dongchuan, Yunnan, China	MG197771
G. (P.) confucius (DC2)	Dongchuan, Yunnan, China	MG197772
G. (P.) confucius (KM1–3)	Kunming, Yunnan, China	MG197771
G. (P.) mandarinus kimurai (CM1)	Doi Inthanon, Chiang Mai, Thailand	MG197752
G. (P.) mandarinus kimurai (CM2–4)	Doi Inthanon, Chiang Mai, Thailand	MG197753
G. (P.) mandarinus kimurai (CM5)	Doi Inthanon, Chiang Mai, Thailand	MG197754
G. (P.) mandarinus fangana (CM1–3)	Doi Pu Meun, Chiang Mai, Thailand	MG197751
G. (P.) mandarinus stilwelli (WX1)	Weixi, Yunnan, China	MG197755
G. (P.) mandarinus stilwelli (YX1)	Yunxian, Yunnan, China	MG197755
G. (P.) mandarinus stilwelli (TC1–2)	Tengchong, Yunnan, China	MG197755
G. (P.) mandarinus stilwelli (YB1–2)	Yangbi, Yunnan, China	MG197755
G. (P.) mandarinus stilwelli (SG1)	Tarung Hka River, Sagaing, Myanmar	MG197755
G. (P.) mandarinus stilwelli (KC1)	Chudu Razi Hills, Kachin, Maynmar	MG197755
G. (P.) mandarinus mandarinus (BX1–4)	Baoxing, Sichuan, China	MG197749
G. (P.) mandarinus mandarinus (BX5)	Baoxing, Sichuan, China	MG197750

TABLE 1. Specimens used in molecular analysis with GenBank accession numbers, accession numbers shared between samples with the same *cox1* sequence.

Morphological comparison

Specimens were spread for examination, with the scent scales on their hindwings exposed.

Spread specimens were photographed using a Fujifilm S8600 digital camera (Fujifilm, Japan) with medium grey background. Photos were adjusted using Adobe Photoshop CS (Adobe, USA). Diagnostic characters of the new taxa were also denoted using Adobe Photoshop CS. For comparison between taxa, the lengths of forewing were measured to 0.5 mm precision. Average lengths of forewing were calculated, and the standard deviations were also calculated when $n \ge 3$.

To observe the male and female genitalia, the abdomen was taken from the specimen and placed into a 1.5 mL microcentrifuge tube, and 1 mL water was added to the abdomen to rehydrate the tissue at 50 °C for 30 min, then 1 mL 10% sodium hydroxide solution was used to digest soft tissue at 70 °C for 1 h. The treated abdomen was neutralised with 2% acetic acid and then dissected in a water-filled Petri dish under the stereoscope to remove residual tissues, scales, and hair. The genitalia were then transferred to 80% glycerol for 12 h to render them transparent. Photographs were taken with a Nikon DMX1200 digital camera (Nikon, Japan) mounted on a Nikon SMZ1500 stereoscope (Nikon, Japan) and automatically stacked using Helicon Focus 3.2 (Helicon Software, USA). The distance between socii of male genitalia is a useful morphometric character helpful to distinguish species (Koiwaya 1993); therefore this distance was measured to 0.2 mm precision for all dissected male genitalia. Similarly, average distances between socii were calculated, and standard deviation was also calculated when $n \ge 3$. After observation and photography, all parts of the genitalia were fixed on a glue card and pinned with the specimen.

Molecular phylogenetic analyses

For each specimen, one or two legs (except forelegs) on the same side were extracted, homogenised in protease buffer containing 100 μ L STE (10 mmol/L Tris-HCl, 1 mmol/L EDTA, 100 mmol/L NaCl, pH = 8.0) and 2 μ L Proteinase K (20 mg/mL) (O'Neill *et al.* 1992). Homogenised samples were treated at 37 °C incubation for 15 min to rehydrate the tissue and then at 95 °C incubation for 10 min to lyse the tissue. The supernatant was recovered through centrifugation at 6,000g and used directly as DNA template in polymerase chain reactions (PCR).

The PCR reaction was applied in a 25 μ L system by using TaKaRa Ex *Taq* Kit (TaKaRa Biotechnology Co., Ltd., Dalian, China) which contained 2.5 μ L of 10× PCR buffer, 2.0 μ L of MgCl₂ (25 mmol/L), 2.0 μ L of dNTP mixture (2.5 mmol/L each), 0.25 μ L of *Taq* DNA polymerase (5 U/ μ L), and 0.5 μ L of each of forward and reverse primers (20 μ mol/L). Primers LCO1490 (5'- GGT CAA CAA ATC ATA AAG ATA TTG G-3') and HCO2198 (5'- TAA ACT TCA GGG TGA CCA AAA AAT CA-3') (*cox1*; Folmer *et al.* 1994) were used. The thermal profile of PCR consisted of an initial denaturation at 95 °C for 3 min; 30 cycles of denaturation at 94 °C for 1 min, annealing at 50 °C for 1 min, and elongation at 72 °C for 5 min. Sequences were obtained by using an ABI Prism 3730 sequencer (Applied Biosystems, California, USA).

Raw sequences were proofread and aligned using Clustal W (Thompson *et al.* 1994) in BioEdit 7.0.9 (Hall 1999), and any sequence containing double peaks in the chromatograms was strictly excluded. The product sequences were checked by MEGABLAST against the genomic references and nucleotide collection in NCBI. Amino acid translation was realised with the invertebrate mitochondrial criterion in MEGA 6.0 (Tamura *et al.* 2013) to detect possible *Numts* (nuclear copies of mtDNA fragments). A search for nonsynonymous mutations, inframe stop codons, and indels was carried out to further minimise the existence of cryptic *Numts* (Song *et al.* 2008; Bertheau *et al.* 2011). The Kimura two-parameter (K2P) distances (Kimura 1980) between identified taxa were calculated by MEGA 6.0.

The phylogeny was reconstructed using Bayesian Inference (BI) method as implemented in MrBayes 3.2.6 (Ronquist *et al.* 2012), with the most appropriate nucleotide substitution model recovered by jModelTest 0.1 (Guindon & Gascuel 2003; Posada 2008). An individual of *Iphiclides podalirius* (Linnaeus, 1758) and of *Lamproptera meges* (Zinken, 1831) was used as outgroup, because we found that tree topology and node support were improved when using outgroup species which branched before *Pazala* split from the rest of *Graphium*. Initial trees using *Graphium chironides* (Honrath, 1884) as outgroup were poorly resolved. First, we reconstructed the phylogeny for a dataset containing the currently recognised species of *Pazala* (7 in total, subspecies not included)

in an attempt to obtain an overall phylogenetic framework for the subgenus. Second, we reconstructed phylogeny for a dataset containing only the currently recognised species and subspecies of the *Graphium mandarinus* group to produce phylogenetic relationships among the focal taxa in the present study. BI analyses consisted of two independent runs, each with four Markov chain Monte Carlo (MCMC) running for 10 million generations (sampled every 1000th generation) to calculate the clade posterior probabilities (PP).

Phyletic properties of each taxon in the *mandarinus* group were assessed using an online tool, Monophylizer (Mutanen *et al.* 2016; http://monophylizer.naturalis.nl/). Taxa identified as monophyletic were treated as good species or subspecies, while those identified as paraphyletic were further analysed using morphological characters and geographical ranges.

To estimate divergence times and infer their credibility intervals, we performed Bayesian relaxed-clock analyses using MrBayes on both datasets (for the *mandarinus* group and for the subgenus *Pazala*). For these analyses, we relied on the partitioning scheme and the substitution models as determined above. MrBayes analyses were performed with the same settings as above but running for 20 million generations with sampling tree every 2,000 generations. We used the autocorrelated clock model for two reasons. First, the autocorrelated model is more appropriate for our dataset because the rate along a given branch is more similar to its parent branch than a branch chosen at random, though autocorrelation models differ in the degree to which they restrict rate variation between parent and daughter branches (Thorne *et al.* 1998; Thorne & Kishino 2002). Second, Lepage *et al.* (2007) and Rehm *et al.* (2011) showed that the autocorrelated clock model generally offers the best fit, as compared to the uncorrelated model and a strict molecular clock model.

Calibration priors are based on the time-calibrated tree of Papilionidae (Condamine *et al.* 2012). We set four secondary calibrations using a (conservative) uniform prior with bounded by the minimum and maximum ages of the 95% credibility interval (CI) of the divergence times (a normal prior is not recommended, Schenk 2016). We could not use fossils in this study because the three fossils do not belong to the subfamily Papilioninae (Condamine *et al.* 2012). We calibrated the following nodes: (i) the root of the tree (crown of Leptocircini) set between 27.6 and 43.4 Ma; (ii) the crown between *Iphiclides* and *Lamproptera* set between 20.8 and 35.5 Ma; (iii) the crown of the genus *Graphium* set between 21.2 and 35.7 Ma; and (iv) the crown of the subgenus *Graphium* set between 14.4 and 29.7 Ma.

For all Bayesian runs (phylogenies and dating), convergence was ensured by checking average deviation of split frequencies (ADSF), potential scale reduction factor (PSRF) values, effective sample size (ESS) of all parameters, and by plotting log-likelihood of samples against number of generations in Tracer 1.6. To reach good convergence, the runs must have values of ADSF approaching zero, PSRF close to 1.00 and ESS above 200. Bayesian consensus trees were obtained using the 25% burn-in criterion (Ronquist *et al.* 2012), and the remaining samples were used to generate a 50% majority rule consensus tree.

All phylogenetic and dating analyses were performed on the computer cluster CIPRES Science Gateway (Miller *et al.* 2015), using BEAGLE (Ayres *et al.* 2012) with default parameters.

Results

Pazala Moore, 1888

Pazala Moore, 1888; Descr. Indian Lep. Ins. Atkinson, Pt. III: 283; TS: Papilio Glycerion Gray, 1831.

Non-mimetic without sexual dimorphism. Forewing short, triangular, glossy, and semitransparent in the basal, costal, and apical interspaces, with slightly curved costa, broad and obtuse apex, and slightly concave termen. Forewing creamy white or greyish white, with ten black bands. Hindwing triangular, with long and oblique costa, very oblique and sinuous termen, deeply scalloped tornal area, and long sword-like tails. Tornal area black, marked with three grey-blue crescents, and two connecting yellow spots above the black patch. Male anal scent fold narrow, with sparse long scales at the base, but no scent wool. The ventral lobes of ostium in female genitalia strongly forked, as stated by Miller (1987).

Phylogenetics of subgenus Pazala and the mandarinus group

The BI phylogenetic tree divided subgenus *Pazala* into four major clades, supported with maximal posterior probability (Figure 1). The first clade to branch off contains *Graphium (Pazala) alebion*, which is thus sister to all other *Pazala* species. A second clade comprises G(P) tamerlanus and G(P) parus (as tamerlanus group hereafter). The third clade contains G(P) eurous and G(P) mullah (as eurous group hereafter). The fourth clade includes G(P) mandarinus and G(P) sichuanica (hereafter referred to as the mandarinus group).



FIGURE 1. The Bayesian phylogenetic tree of subgenus *Pazala* Moore, 1888, with specimen photos (upperside on the left, underside on the right) near the end of each taxon clade. Values at nodes indicate the posterior probability. *Iphiclides podalirius* and *Lamproptera meges* as outgroup (specimen photos in the box at bottom right).

Phylogenetic analyses of the *mandarinus* group showed that species were well defined as monophyletic with high (if not maximal) posterior probabilities (Figure 2). It must be noted that within the *mandarinus* group, G(P) garhwalica stat. nov. is found to be sister to all taxa in the group, followed by G(P) paphus stat. nov., a further branch containing G(P) sichuanica and G(P) hoeneanus nom. nov., stat. rev., followed by the pairing of G(P) daiyuanae **sp. nov.** and G(P) confucius **sp. nov.**, and finally four subspecies of G(P) mandarinus. Recently other molecular analyses of Papilionidae have shown similar presence of cryptic species previously only recognised as subspecies, e.g. Shiraiwa *et al.* (2014) separated several Heraclides taxa at species level as a result of their studies.

The Monophylizer analysis identified most taxa in the *mandarinus* group as monophyletic except G(P) mandarinus stilwelli ssp. nov., which tangled (but not mixed) with G(P) mandarinus mandarinus forming a paraphyletic clade in the tree (Table 2). The phylogenetic structure and Monophylizer assessment of the mandarinus group indicates four main taxonomic results: (i) G(P) daiyuanae sp. nov. and G(P) confucius sp. nov. are both distinct species, differing from each other as well as from other known species; (ii) G(P) garhwalica and G(P) paphus are both distinctive species, instead of being subspecies of G(P) mandarinus as previously

recognised; (iii) G (P) hoeneanus nom. nov., stat. rev. is different from G (P) sichuanica and should be given full species status; and (iv) G (P) mandarinus stilwelli **ssp. nov.** is genetically rather close to the nominate subspecies of G (P.) mandarinus, but is morphologically a good subspecies.



FIGURE 2. The Bayesian phylogenetic tree of *Graphium (Pazala) mandarinus* group, with *Iphiclides podalirius* and *Lamproptera meges* as outgroup. Coloured rectangles delineate the species and subspecies of the *mandarinus* group. Values at nodes indicate the posterior probability.

The Kimura 2-parameter (K2P) distances (in percentage) between taxa ranged from 0.46% to 6.48%, with that between G (P.) mandarinus fangana and G (P.) mandarinus stilwelli ssp. nov. being the smallest, while that between G (P.) garhwalica stat. nov. and G (P.) mandarinus mandarinus being the greatest. Most K2P distances between identified species were greater than 2%, except that between G (P.) daiyuanae sp. nov. and G (P.) confucius sp. nov., being only 1.71% (Table 3).

Morphological examination, detailed in the revision section below, supported the species level phylogenetic analysis particularly in differences of both male and female genitalia.

TABLE 2. Monophylizer assessment	t of species and subspe	ecies in Graphium (Pazal	a) mandarinus group.

Taxon	Assessment	Tanglees	
G. (P.) garhwalica stat. nov.	monophyletic		
G (P.) paphus stat. nov.	monophyletic		
G (P.) sichuanica	monophyletic		
G (P.) hoeneanus nom. nov., stat. rev.	monophyletic		
G. (P.) daiyuanae sp. nov.	monophyletic		
G. (P.) confucius sp. nov.	monophyletic		
G. (P.) mandarinus	monophyletic		
ssp. kimurai	monophyletic		
ssp. fangana	monophyletic		
ssp. <i>stilwelli</i> ssp. nov.	paraphyletic	ssp. mandarinus	
ssp. mandarinus	monophyletic	_	

TABLE 3. The Kimura 2-parameter distances (shown in percentages) between species and subspecies in *Graphium* (*Pazala*) mandarinus group.

Taxon	1	2	3	4	5	6	7a	7b	7c	7d
1. garhwalica stat. nov.										
2. paphus stat. nov.	5.42									
3. sichuanica	4.75	5.10								
4. hoeneanus nom. nov., stat. rev.	5.10	4.82	2.03							
5. daiyuanae sp. nov.	4.54	5.32	3.94	3.46						
6. confucius sp. nov.	5.27	5.12	4.12	4.15	1.71					
7a. mandarinus kimurai	6.20	5.72	4.04	4.16	3.88	3.41				
7b. mandarinus fangana	5.76	5.96	3.29	4.09	3.46	2.67	1.64			
7c. mandarinus stilwelli ssp. nov.	5.93	6.13	3.46	4.26	3.62	3.12	1.48	0.46		
7d. mandarinus mandarinus	6.48	6.33	3.98	4.42	3.22	2.99	1.67	0.95	0.49	

Molecular dating analyses

For both the *mandarinus* group and the *Pazala* datasets, the Bayesian dating runs converged very well: almost all PSRF values were at 1.0 (average at 1.0 and maximum at 1.005) for both, and the ADSF was at 0.001 and 0.005 (maximum at 0.03), respectively. All the parameters have ESS well above 200, indicating good mixing of the MCMC.

The results of the molecular dating analyses are presented in Figures 3 and 4 for both the previously recognised seven *Pazala* species and the *mandarinus* group taxa designated in the present study, respectively. We estimated that the currently recognised subgenus *Pazala* originated in the early Miocene *ca.* 20.4 Ma (95% CI: 13.8–27.4 Ma). Most speciation events occurred before the Pliocene and we estimated no event in the Pleistocene (Figure 3), with the last speciation event taking place *ca.* 6.48 Ma (95% CI: 2.0–11.96 Ma). On the contrary in the *mandarinus* group, we estimated that most phylogenetic events happened in the Pliocene and Pleistocene (Figure 4). The crown age of the subspecies complex of *mandarinus* is estimated at 2.76 Ma (95% CI: 0.7–6.85 Ma). We note a good congruence between the age estimates obtained with the *Pazala* and the *mandarinus* datasets (e.g. crown age of Leptocircini is 35.8 Ma and 38.1 Ma, crown age of *Graphium* is 31.2 Ma and 31.5 Ma, crown age of *Pazala* is 20.4 Ma and 16.0 Ma, most recent common ancestor of *G* (*P.*) *mandarinus* and *G* (*P.*) *sichuanica* is 6.5 Ma and 5.8 Ma, respectively).



FIGURE 3. Bayesian molecular dating for the seven previously recognised species in the subgenus *Pazala* Moore, 1888. Values at nodes indicate the median divergence times, purple bars show 95% CI. Pleisto. = Pleistocene.



FIGURE 4. Bayesian molecular dating for species and subspecies in the *Graphium (Pazala) mandarinus* group found in this study. Values at nodes indicate the median divergence times, purple bars show 95% CI. Pleisto. = Pleistocene.

Revision of Graphium (Pazala) mandarinus group

Nomenclature of Pazala genitalia

Examination of Pazala male genitalia showed that extra names are needed (Figure 5) for the various structures of

the valve, since they are probably not homologous with those used for the Afrotropical *Graphium* species of Smith & Vane-Wright (2001), whereas nomenclature of female genitalia follows that publication (Figure 6). General descriptions of male and female genitalia are given below while only distinguishing characters are listed under each species.



FIGURE 5. Nomenclature of male genitalia of the *Graphium (Pazala) mandarinus* group used in this study, illustrated by *G*. *(P.) sichuanica* (Koiwaya, 1993). A: male genitalia with left valve removed, B: right valve, C: dorsal view of tegumen, socius, and uncus (subsequently as TSU in the text). Names proposed for the harpes and projections of the valve in italics.

General description of male genitalia. Moderately to highly sclerotised. Ring moderate in width, straight in the lower half and wavy in the upper half; saccus small to nearly reduced; socii small, ventrally curved with acutely pointed end, distance between the base of socii varies among taxa. Valve short, oval in general with five highly sclerotised harpes: the dorsal terminal harpe extends from apical 1/3 of the dorsal margin to the end of the valve, with curved, serrate edge and a tip of varying shape; the dorsal subterminal harpe, with serrate edge, of variable size located in the upper and outer 1/4 to 1/3 of the valve, opposite to the dorsal terminal harpe; the distal harpe large, the ventral margin with strongly serrate edge; the medial harpe long and curved, with a dorsal projection variable among taxa and a serrate sclerotised flange. Aedeagus long, curved ventrally, with a funnel-shaped base and pointed end. Juxta weakly sclerotised with hairy membrane on both sides.

Our study showed that the shape, size, and relative position of the dorsal terminal harpe, the dorsal subterminal harpe, and the shape of the dorsal projection are important diagnostic characters. Also, the uncus does not project noticeably from the tegumen, but contains a pair of socii with useful morphometric characters helpful in distinguishing species (Koiwaya, 1993).

General description of female genitalia. Papillae anales short, margin smooth and round; posterior apophysis slender, reaching approximately half of the 9th abdominal section; ostium partially covered by lamella antevaginalis and ostial lobe; lamella postvaginalis paired, approximately petal-shaped; lamella antevaginalis broad horizontally, lined with moderately sclerotised longitudinal striae; ostial lobe heavily sclerotised, with the posterior margin concave in the middle, separating it into a pair of spurs variable between taxa; ductus bursae of medium length; signum well defined by sclerotised granules.

Our study showed that the ostial lobe (i.e., bifurcation of posterior margin, shape of spurs) varies among species, and probably can be used as a diagnostic character. The characters of the ostial lobe of each species and the differences from other species are listed in detail below.



FIGURE 6. Nomenclature of female genitalia of the *Graphium (Pazala) mandarinus* group used in this study illustrated by *G. (P.) mandarinus mandarinus* (Oberthür, 1879). A: lateral view, B: ventral view.

Confirmation of the identity of *Papilio Glycerion* variété géographique *Mandarinus* Oberthür, 1879 and Lectotype Designation for *Papilio Glycerion* Gray, 1831 (section authors: Cotton & Hu)

Since this study has shown that the previously recognised nominate population of *Graphium (Pazala) mandarinus* (Oberthür, 1879) comprises two separate species, it is necessary to confirm which taxon the name *mandarinus* applies to.

On page 115 of Études d'Entomologie IV (1879) Oberthür described [*Papilio*] *Glycerion* variété géographique *Mandarinus* from an unspecified number of specimens 'de Moupin et de Kouy-Tchéou' without designating a holotype. On page 64 of the same publication he listed a single specimen of *Papilio glycerion* from Moupin (A. David [leg.]) and two males from 'Népaul'.

Hemming (1934: 195) designated the specimen from Moupin [= Baoxing, Sichuan, China] as lectotype of the taxon in accordance with article 74.5 of the ICZN Code (1999) by stating 'type from Moupin'. Since only a single Moupin specimen was listed by Oberthür in the syntypic series the specimen Hemming had designated was unambiguously identified despite the brevity of the statement. This action also fixed the type locality to Moupin under Code article 76.2 and rendered the status of the Kouy-Tchéou specimens as paralectotypes.

The lectotype (Figure 7) is clearly identifiable on morphological characters as conspecific with taxa previously treated as subspecies of *Graphium mandarinus* rather than the sympatric new species, *Graphium confucius* Hu, Duan & Cotton **sp. nov.**, recognised in this work; thus maintaining stability of names for the majority of relevant taxa. Two paralectotypes from Kouy-Tchéou are housed in BMNH, London. The Trustees kindly provided photos

of these specimens which clearly belong to *Graphium (Pazala) confucius* Hu, Duan & Cotton **sp. nov.**, and are designated as paratypes of the new taxon below.



FIGURE 7. Lectotype of *Papilio Glycerion* variété géographique *Mandarinus* Oberthür, 1879; upperside left, underside right, all labels below. All images © Reproduced with permission and with thanks to the Trustees of the Natural History Museum, London.



FIGURE 8. Lectotype of *Papilio Glycerion* Gray, 1831 designated in the present study; upperside left, underside right, all labels below. All images © Reproduced with permission and with thanks to the Trustees of the Natural History Museum, London.

From 2003 Campbell Smith worked on *Pazala* at the Natural History Museum, London (BMNH), intending to publish a revision of the subgenus. The first author of this section was in personal communication with Smith from 2006 until his retirement in early 2009 but the proposed revision of the group was never submitted for publication, as confirmed by R. I. Vane-Wright (pers. comm.) in 2016. Smith shared part of his manuscript with the first author, in which he was planning to designate the only known extant syntype of *Papilio Glycerion* Gray, 1831, housed in

BMNH, as lectotype of that taxon. Even though only one specimen is known it must be treated as a syntype since there was no indication of the number of specimens in the original description. Lectotype designation is necessary in order to fix the identity of the name due to long standing historical confusion detailed by Racheli & Cotton (2009: 26), even though the name is a junior homonym.

Since Smith never published his revision we herein designate as the Lectotype of *Papilio Glycerion* Gray, 1831 the male syntype chosen by Campbell Smith. This specimen (Figure 8) has the following labels attached: 1702 a [hand written on round white label] / Nepaul. Hardwicke Bequest. 1702 a. [hand written] / B. M. TYPE No. RH11072. Papilio glycerion, \bigcirc Gray / Type [round red edged label] / LECTO-TYPE [round dark blue edged label] / LECTOTYPE [purple print] *Papilio glycerion* G.R. Gray C.R. Smith det. 2003 / BMNH(E) # 149384.

Graphium (Pazala) garhwalica (Katayama, 1988) stat. nov.

(Figure 9, A–B)

Pazala glycerion garhwalica Katayama, 1988; Gekkan-Mushi, 203: 8; pl. 1, f. 1–2; TL: "Joshimath (2400 m), Uttar Pradesh [= Uttaranchal], NW. India".

Diagnostic characters: Small in size, forewing length: male 32.5–33.5 mm (n = 2), female 32.0 mm (only one female examined) [Katayama (1988) stated 36 mm for the female in the original description]. Both wings rather whitish. Forewing, glossy, apex blunt, termen straight (a); veins M₃ to Cu₂ discally completely free of black scales in both sexes (b). Hindwing, black discal band on the upperside completely reduced in both sexes (c), short submarginal black lines also reduced on the upperside in both sexes, only reaching space sc+r₁.



FIGURE 9. *G* (*P*) garhwalica (Katayama, 1988) stat. nov. (A–B) and *G* (*P*) paphus (de Nicéville, 1886) stat. nov. (C–F); upperside in the first row, underside in the second row; scale bar = 10 mm. A: \mathcal{J} , Joshimath, N.W. India, **PARATYPE**; B: \mathcal{Q} , ditto; C: \mathcal{J} , Tiger Hill, Darjeeling, N. India; D: \mathcal{Q} , ditto; E: \mathcal{J} , Metok, S. Tibet, China; F: \mathcal{J} , Tarung Hka River, N. Sagaing, Myanmar.



FIGURE 10. Male genitalia of *G*. (*P.*) garhwalica (Katayama, 1988) stat. nov. **PARATYPE** from Joshimath, N.W. India; scale bar = 1.0 mm. All: genitalia as a whole, R.: lateral view of ring, TSU: dorsal view of tegumen, socii and uncus, V. right valve, Ae.: lateral view of aedeagus, Ju.: ventral view of juxta.

Distribution: N.W. India (Uttaranchal Pradesh) to C. Nepal (Annapurna). **Phenology:** Univoltine in April-May.

Host plant: Unknown, presumably a plant species belonging to family Lauraceae.

Male genitalia (Figure 10): In total two male genitalia were dissected, the general characters are consistent. Highly sclerotised. Ring slightly wavy in the upper half; saccus very small, nearly reduced; distance between the base of socii 0.64-0.68 mm (n = 2). Valve short, oval in general, the end of dorsal terminal harpe extends into distinctly elongate acute tip; the dorsal subterminal harpe reaching the base of the elongate tip; the medial harpe long and curved, the dorsal projection flat and serrate; no tooth in the middle of the medial harpe. Juxta weakly sclerotised with hairy membrane on both sides.

Female genitalia: The abdomen of the only available female specimen was unfortunately crushed prior to obtaining the specimen, the ostial lobe was still identifiable but split into three pieces. The first author photographed the three pieces and digitally rejoined them without alteration, obtaining a reconstructed image of the ostial lobe (Figure 11). Ostial lobe (Figure 11) heavily sclerotised, broad at the base with the posterior margin widely bifurcated into a pair of broad spurs in ventral view.



FIGURE 11. Ventral view of digitally reconstructed ostial lobe of female genitalia of *G.* (*P.*) garhwalica (Katayama, 1988) stat. nov. from Joshimath, Uttaranchal, N.W. India; scale bar = 0.5 mm.

Graphium (Pazala) paphus (de Nicéville, 1886) stat. nov.

(Figure 9, C–F)

Papilio paphus de Nicéville, 1886; J. Asiat. Soc. Bengal, 55 Pt. II (3): 254, pl. 11, f. 6; TL: Sikkim.
Papilio Glycerion Gray, 1831 (junior homonym of Papilio Glycerion Borkhausen, 1788 [Nymphalidae: Satyrinae]); Zool. Miscell., 1: 32; TL: "Nepaul" [= Nepal].

Diagnostic characters: Larger than the preceding species, forewing length: male 35.0–41.0 mm (mean = $37.1 \pm 1.3 \text{ mm}$, n = 29), female 39.5 mm (only one female examined). Both wings rather whitish. Forewing, apex more obvious than that of *garhwalica*, termen concave in the middle and slightly wavy at the end of each vein (a); veins Cu1 and Cu₂ mostly free of black scales except the base, while vein M₃ entirely black (b). Hindwing, black discal band on the upperside incomplete but usually present near and in the cell (c), short submarginal black lines reaching space sc+r₁.

Distribution: Nepal, Bhutan, N. India (Sikkim, Assam), N. Myanmar (Sagaing), and W. China (S. Tibet, i.e., Metok).

Phenology: Univoltine. Adults were collected from late April to mid May in Metok (Wa Da, pers. comm.). **Host plant:** Unknown, presumably a plant species belonging to family Lauraceae.

Male genitalia (Figure 12): In total eight male genitalia were dissected, the general characters are consistent, with slight variation in the harpes. Highly sclerotised. Ring wavy in the upper half; saccus small; distance between the base of socii 0.64-0.74 mm (mean = 0.70 ± 0.03 , n = 8). Valve short, oval in general, the dorsal terminal harpe with an acute but not elongate tip (elongate in *garhwalica*); the dorsal subterminal harpe small and isolated (reaching the dorsal terminal harpe in *garhwalica*), with serrate edge; the medial harpe long and straight, dorsal projection bayonet-shaped (in sharp and acute triangular pyramid shape) (flat in *garhwalica*); a variable number of teeth (0–2) in the middle of the medial harpe.

Female genitalia (Figure 13): Only one female was available for genitalia dissection. Lamella postvaginalis

small, round-shaped; ostial lobe heavily sclerotised, broad at the base and abruptly narrowing into a sharp spur in lateral view, while the posterior margin deeply bifurcated into a pair of acute spurs in ventral view (only shallowly bifid in *garhwalica*).



FIGURE 12. Male genitalia of G(P) paphus (de Nicéville, 1886) stat. nov. from Metok, S. Tibet, China; scale bar = 1.0 mm. All: genitalia as a whole, R.: lateral view of ring, TSU: dorsal view of tegumen, socii, and uncus, V. right valve, Ae.: lateral view of aedeagus, Ju.: ventral view of juxta.

Graphium (Pazala) sichuanica (Koiwaya, 1993)

(Figure 14, A–C)

Pazala sichuanica Koiwaya, 1993; Stud. Chinese Butt., 2: 77, f. 126–127, 132–133; TL: "Dujiang Yan, Sichuan" [= Dujiangyan, W. Sichuan, China]

Diagnostic characters: A small species, forewing length: male 32.0–40.0 mm (mean = 35.4 ± 2.3 mm, n = 24), female 42 mm [only one female examined; 37-38 mm stated by Koiwaya (1993)]. Both wings narrower, with a yellowish tinge and well developed black markings. Forewing, the 9th black band reaches vein 2A and joins with the 8th black band with equal width (a). Hindwing, short submarginal black lines well connected to each other, almost forming straight lines (b); the discocellular veins almost free of black scales on the underside (c).

Distribution: N. to W. Sichuan, China.

Phenology: Univoltine. The type specimens in Koiwaya (1993) were collected from April to May. Our examined specimens from N. Sichuan were all collected in mid to late April.

Host plant: Unknown, presumably a plant species belonging to family Lauraceae.

Male genitalia (Figure 15): In total six male genitalia were dissected, the general characters are consistent, with slight variation in the harpes. Moderately sclerotised. Ring wavy in the upper half; saccus well developed; distance between the base of socii 0.80-0.84 mm (mean = 0.81 ± 0.02 , n = 6). Valve elongate, oval in general, the dorsal terminal harpe extends obliquely outwards with produced but non-elongate tip; the dorsal subterminal harpe rather small and completely isolated; the medial harpe long and curved, the dorsal projection flat; no tooth in the middle of the medial harpe.

Female genitalia (Figure 16): Only one female was available for genitalia dissection. Lamella postvaginalis round petal-shaped; lamella antevaginalis covered with sclerotised wrinkles; ostial lobe heavily sclerotised, broad at the base and slightly curved into a sharp end in lateral view, while the posterior margin shallowly bifurcated into a pair of spurs in ventral view.



FIGURE 13. Female genitalia of G(P) paphus (de Nicéville, 1886) stat. nov. from Tiger Hill, Darjeeling, N. India; lateral view above, ventral view below; scale bar = 1.0 mm.



FIGURE 14. *G* (*P.*) *sichuanica* (Koiwaya, 1993) (A–C) and *G* (*P.*) *hoeneanus* Cotton & Hu nom. nov., stat. rev. (D–F); upperside in the first row, underside in the second row; scale bar = 10 mm. A and B: \bigcirc , Laohegou, Pingwu, Sichuan, W. China, \bigcirc Peking University & Laohegou Nature Reserve; C: \bigcirc , Tangjiahe, Qingchuan, Sichuan, W. China, \bigcirc Peking University & Tangjiahe Nature Reserve; D: \bigcirc , W. Tianmu Shan (350 m), Zhejiang, E. China; E: \bigcirc , Tiantong (400 m), Ningbo, Zhejiang, E. China, \bigcirc Jian-Qing Zhu (Shanghai Zoological Park, Shanghai, China); F: \bigcirc , Lingui (360 m), Guilin, Guangxi, S.W. China.

Graphium (Pazala) hoeneanus Cotton & Hu nom. nov., stat. rev.

(Figure 14, D-F)

Papilio tamerlanus hoenei Mell, 1935 (junior primary and secondary homonym of Papilio agetes hönei Mell, 1923 [Papilionidae: Graphium: Pathysa]); Mitt. D. Ent. Ges., 6 (3/4): 36; TL: "Chekiang: West- und Ost-Tienmoshan (Fukien, Südkiangsi), Nordkwangtung" [= West and East Tianmu Shan, Zhejiang (Fujian, S. Jiangxi), N. Guangdong].

This taxon has previously been treated as synonymous with *Graphium (Pazala) sichuanica* (Koiwaya, 1993) (Racheli & Cotton 2009), but is here shown both on molecular and morphological characters to be worthy of separate specific status (**stat. rev.**). Since the only name applicable to the taxon is a junior homonym and thus unavailable, we hereby propose the new replacement name *hoeneanus* Cotton & Hu **nom. nov.** as the valid taxon name.

Diagnostic characters: Larger than the preceding species, forewing length: male 34.0–41.0 mm (mean = 37.6 \pm 1.6 mm, n = 29), female 36.0–38.0 mm (mean = 37.0 \pm 0.91 mm, n = 4). Both wings broader, rather whitish, with reduced black markings. Forewing, the 9th black band only reaches space cu₂ and is only faintly indicated near the 8th black band, not clearly joining it (a). Hindwing, short submarginal black lines often disconnected from each other (b); the discocellular veins often tinged with black scales on the underside (c).

Distribution: Widely distributed in S. to E. China (i.e., Guangxi, Guangdong, Zhejiang; probably in Hunan, Hubei, Anhui, and Jiangxi).

Phenology: Univoltine. Adults were collected from early April to late May in Zhejiang, E. China (Jian-Qing Zhu, pers. comm.), as well as in Guangdong, S. China (Jia-Lin Chen, pers. comm.).



FIGURE 15. Male genitalia of G(P) sichuanica (Koiwaya, 1993) from Laohegou, Pingwu, Sichuan, W. China; scale bar = 1.0 mm. All: genitalia as a whole, R.: lateral view of ring, TSU: dorsal view of tegumen, socii, and uncus, V. right valve, Ae.: lateral view of aedeagus, Ju.: ventral view of juxta.

Host plant: According to Jian-Qing Zhu (pers. comm.), the host plant is *Lindera reflexa* of family Lauraceae in Zhejiang, E. China; while Wang & Gu (2017) recorded *Machilus phoenicis* of family Lauraceae in Nanling, S. China.

Derivatio nominis: The replacement name is dedicated to Hermann Höne, following the intention of Rudolf Mell in his original description of '*Papilio tamerlanus hoenei*', and is treated as a noun in apposition.

Male genitalia (Figure 17): In total eight male genitalia were dissected, the general characters are consistent, with the teeth on medial harpe variable. Moderately sclerotised. Ring slightly wavy in the upper half; saccus small; distance between the base of socii 0.84-0.90 mm (mean = 0.89 ± 0.02 , n = 8). Valve elongate, oval in general, the dorsal terminal harpe with produced but non-elongate tip (as in *sichuanica*); the dorsal subterminal harpe medium-sized (obviously larger than that in *sichuanica*) and completely isolated; the medial harpe straight, the dorsal projection flat (as in *sichuanica*); a variable number of small teeth (0–2) occur in the middle of the medial harpe.

Female genitalia (Figure 18): Only two females were available for genitalia dissection, the characters are consistent. Lamella postvaginalis round petal-shaped; lamella antevaginalis broad horizontally, covered with sclerotised wrinkles (as in *sichuanica*); ostial lobe heavily sclerotised, broad at the base and curved upwards into a sharp end in lateral view, while the posterior margin strongly bifurcated into a pair of large spurs deviating away from each other at the base in ventral view (smaller in *sichuanica*).



FIGURE 16. Female genitalia of G(P) sichuanica (Koiwaya, 1993) from Tangjiahe, Qingchuan, Sichuan, W. China; lateral view above, ventral view below; scale bar = 1.0 mm.

Graphium (Pazala) daiyuanae Hu, Zhang & Cotton sp. nov.

(Figure 19)

Description: Male: Forewing length: 32.0-35.5 mm (mean = $34.0 \pm 1.1 \text{ mm}$, n = 16). Forewing triangulate, broad, apex not produced, termen wavy or slightly dented at the end of veins, oblique outwards from apex to tornus. Forewing upperside: whitish, the upper and outer 1/3 (mostly the discocellular, the subterminal, and the terminal areas) hyaline; 10 black bands lined from the humeral to the termen, among which the 1st to the 6th almost run parallel; the 1st and the 2nd bands reach the tornal margin, the 3rd to the 5th bands extend beyond the discal cell along veins Cu₂, Cu₁, and M₃, the 6th band at the end of cell, often connected with the 5th band by a transverse black fine line, the 7th band joins the 8th band just at vein M₂, extends to the tornus and joins the 9th band at vein Cu₂ or just below it in space cu₂; the veins R₄₊₅ to Cu₁ are black after meeting the 6th and 7th bands and then divide the whitish-hyaline areas between the 6th to 10th bands into spots. Forewing underside: colour and markings

as upperside, but the hyaline areas are glossy. Hindwing triangulate in general, vein M_3 extends into a sword-like tail, termen slightly dented at the end of veins, the end of spaces m_1 to cu_1 expand like petals. Hindwing upperside: whitish with long white hair covering the inner 1/3; tornal margin blackish, with a small brown androconial patch near the humeral, a thick black band extends from the costal towards the tornus and joins the tornal blackish area just above two obliquely neighbouring yellow spots; the discal bands usually absent but can be seen through the underside; the submarginal black band coupled and interrupted by veins Rs and M_1 , reaching space rs or sc+r₁; the terminal band single, also interrupted by veins Rs and M_1 , reaching space rs or sc+r₁; all black bands mentioned above are rather discontinuous, breaking into several short bands in each space, and the ends of each short band seldom touch each other; all black with white tip. Hindwing underside: colour and markings similar to upperside but with a creamy yellow hue, all black bands and markings well defined, especially the discal band, two black lines twisting into an "8"-shaped pattern, the outer half of the upper of ring "8" yellow and the inner half white, the edge of the lower ring extends along veins M_1 to Cu_1 ; a square creamy white spot above each greyish-blue lunule in the tornal area, the yellow spots at the tornus somewhat reduced and crowned by fine white lines.



FIGURE 17. Male genitalia of $G_{(P.)}$ hoeneanus Cotton & Hu nom. nov., stat. rev. from W. Tianmu Shan, Zhejiang, E. China; scale bar = 1.0 mm. All: genitalia as a whole, R.: lateral view of ring, TSU: dorsal view of tegumen, socii, and uncus, V. right valve, Ae.: lateral view of aedeagus, Ju.: ventral view of juxta.



FIGURE 18. Female genitalia of $G_{(P)}$ hoeneanus Cotton & Hu nom. nov., stat. rev. from W. Tianmu Shan, Zhejiang, E. China; lateral view above, ventral view below; scale bar = 1.0 mm.

Female: Forewing length 33.5–38.5 mm (mean = 36.4 ± 1.7 mm, n = 14). General appearance similar to male but larger. the forewing termen oblique outwards from the apex to the tornus, the underside of forewing almost as glossy as that in male, the 4th black band on the forewing may be reduced or absent in some individuals, the hindwing androconial patch absent, discal band faintly defined or absent.

Male genitalia (Figure 20): In total 12 male genitalia were dissected, the general characters are consistent, with the teeth on the medial harpe variable. Moderately sclerotised. Ring wavy in the upper half; saccus small; distance between the base of socii 0.40–0.60 mm (mean = 0.49 ± 0.07 , n = 12). Valve oval in general, the dorsal terminal harpe with mostly straight (only in 1 case curved) edge and acute tip; the edge of dorsal subterminal harpe almost touching the base of the tip of the dorsal terminal harpe; the medial harpe mostly straight, the dorsal projection bayonet-shaped (some with blunt or dented end); a variable number of small teeth (0–3) occur in the middle of the medial harpe.



FIGURE 19. *Graphium (Pazala) daiyuanae* Hu, Zhang & Cotton, **sp. nov.**; red dot: holotype, yellow dots: paratypes; upperside on the first row, underside on the second row; A–C: \Diamond , D–F: \bigcirc , Sapa, Lao Cai, N. Vietnam; scale bar = 10 mm.



FIGURE 20. Male genitalia of G(P) daiyuanae Hu, Zhang & Cotton **sp. nov.**, scale bar = 1.0 mm. All: genitalia as a whole, R.: lateral view of ring, TSU: dorsal view of tegumen, socii, and uncus, V. right valve, Ae.: lateral view of aedeagus, Ju.: ventral view of juxta.

Female genitalia (Figure 21): In total 10 female genitalia were dissected, the general characters are consistent with slight variation as mentioned below in particular. Lamella postvaginalis round petal-shaped; lamella antevaginalis broad horizontally, lined with moderately sclerotised longitudinal striae; ostial lobe heavily sclerotised, forming a straight sharp spur in lateral view, while the posterior margin straight or slightly concave with a pair of acute spurs in ventral view, the length of these spurs can vary, but never absent in examined specimens.

Differential Diagnosis: The new species closely resembles G(P) mandarinus, but can be distinguished by careful examination of the following characters: 1) smaller than nominate mandarinus and the subsequent new taxon G(P) confucius Hu, Duan & Cotton **sp. nov.**; 2) forewing apex as G(P) mandarinus, termen oblique outwardly from apex to tornus; 3) upperside hindwing very pale, the discal band usually absent, especially in male; 4) the yellow tornal spots on the underside of hindwing obviously paler than G(P) mandarinus and the following species; 5) the submarginal and terminal bands on the underside of hindwing more interrupted and discontinuous than G(P) mandarinus and the following species. In male genitalia, the tip of dorsal terminal harpe acute but not elongated (as in mandarinus), dorsal subterminal harpe large, joining dorsal terminal harpe and forming a triangle, dorsal projection bayonet-shaped (Figure 20), distance between the base of socii 0.40–0.60 mm. The ostial lobe forming a straight sharp spur in lateral view, the posterior margin straight or slightly concave with a pair of acute spurs in ventral view (Figure 21).



FIGURE 21. Female genitalia of *G*. (*P*.) *daiyuanae* Hu, Zhang & Cotton **sp. nov.**; lateral view above, ventral view below; scale bar = 1.0 mm.

Type Material: Holotype: VIETNAM: ♂, Sapa (1,600 m), Lao Cai Province, 2015–VI, L. T. Le leg [KIZ, 0100000].

Paratypes: VIETNAM: 33, 29, same collecting data as holotype (1,600-1,700 m) [KIZ, 0100001–0100005]; 83, 89, same collecting data as holotype (1,600-1,700 m) [SJH]; 13, 49, the same locality, 2006–IV, local catcher leg. [AMC]; 23, the same locality, 2007–V, local catcher leg. [AMC]; 13, the same locality, 2007–VI, local catcher leg. [AMC]. 23, Mucangchai (1,700 m), Yen Bai Province, 2017–V, L. T. Le leg. [KIZ, 0100006– 0100007]; 33, 19, same collecting data, [SJH].

The holotype and seven paratypes were deposited in Kunming Institute of Zoology (KIZ), Chinese Academy of Sciences (Li *et al.* 2015).

Distribution: Currently known from the mountains in N. Vietnam (Sapa, Lao Cai; Mucangchai, Yen Bai), but very likely extends to the bordering area of S. Yunnan, China (Honghe Prefecture).

Phenology: Probably univoltine, but further survey on the biological and ecological aspects of the new taxon is required in future.

Host plant: Unknown, presumably a plant species belonging to family Lauraceae.

Derivatio nominis: The specific name of this new taxon was dedicated to the first author's beloved wife, Ms. Yuan Dai, with the order of surname and first name as per Chinese tradition. The species name is to be treated as a noun in apposition.

Graphium (Pazala) confucius Hu, Duan & Cotton sp. nov.

(Figure 22)

Description: Male: Forewing length: 35.0–46.0 mm (mean = 39.4 ± 2.2 mm, n = 79). Forewing triangulate, apex produced, termen almost smooth or slightly wavy at the end of veins, concave between apex and tornus. Forewing upperside: whitish, the upper and outer 1/3 (mostly the discocellular, the subterminal, and the terminal areas) hyaline; 10 black bands between the base and the termen, of which the 1^{st} to the 6^{th} almost run parallel; the 1^{st} and the 2nd bands reach the tornal margin, the 3rd to the 5th bands extend beyond the discal cell along veins Cu₂, Cu₁, and M₃, the 6th band at the end of cell is often connected with the 5th band by a transverse black fine line, the 7th band joins the 8th band just at vein M₂, extends to the tornus and joins the 9th band at vein Cu₂ or just below it in space cu₂, the 9th band curved inward in space r₄, interrupted by vein M₂ and the remaining section below displaced inwardly, the 10th band independent, extending along the margin from the apex to vein Cu₂ or just below it in space cu₂, the area between the 8th and 9th bands can be irrigated with blackish scales in some individuals; veins R₄₊₅ to Cu₁ are black after meeting the 6th and 7th bands and then divide the whitish-hyaline areas between the 6th to 10th bands into spots. Forewing underside: colour and markings as upperside, but the hyaline areas are glossy. Hindwing triangulate in general, vein M₃ extends into a sword-like tail, termen slightly dented at the end of veins, the ends of spaces m₁ to cu₁ expand like petals. Hindwing upperside: whitish with long white hair covering the inner 1/3; tornal margin blackish, with a small brown androconial patch near the base, a thick black band extends from the costa towards the tornus and joins the tornal blackish area just above two obliquely neighbouring yellow spots; the discal bands incomplete but well developed; the submarginal black band coupled and interrupted by veins Rs and M₁, reaching space rs or sc+r_i; the terminal band single, also interrupted by veins Rs and M_i , reaching space rs or sc+r_i; despite being interrupted by veins, the ends of each short band in each space touch or directly connect each other; all black bands mentioned above join in the black area at the end of spaces m_1 to cu_1 , with a greyish-blue lunule in each space; tail black with white tip. Hindwing underside: colour and markings similar to upperside but with a creamy yellow hue, all black bands and markings well defined, especially the discal band, two black lines twist into an almost "8"-shaped pattern, with a short connection between the rings, the outer half of the upper ring of the "8" yellow and the inner half white, the edge of the lower ring extends along veins M₁ to Cu₁; a square creamy white spot above each greyish-blue lunule in the tornal area, the yellow spots at the tornus somewhat reduced and crowned by fine white lines.

Female: Forewing length: 38.0-45.0 mm (mean = $40.5 \pm 1.8 \text{ mm}$, n = 30). General appearance similar to male but evidently larger and paler, the underside of forewing less glossy, the 4th black band on the forewing often reduced or even absent in some individuals.

Male genitalia (Figures 23): In total male genitalia of 12 specimens were dissected. Moderately sclerotised. Ring wavy in the upper half; saccus small; distance between the base of socii 0.50-0.64 mm (mean = 0.56 ± 0.05 , n = 12). Valve round shaped, the dorsal terminal harpe with curved edge and acute tip (less pointed than *daiyuanae* **sp. nov.**); the dorsal subterminal harpe mostly isolated; the medial harpe mostly straight, the dorsal projection bayonet-shaped (some with blunt or serrate tip) (as in *daiyuanae* **sp. nov.**); a variable number of small teeth (0–1) occur in the middle of the medial harpe.

Female genitalia (Figure 24): In total female genitalia of 13 specimens were dissected, the general characters are consistent, with slight variation as mentioned below. Lamella postvaginalis round petal-shaped; lamella antevaginalis broad horizontally, lined with moderately sclerotised longitudinal striae; ostial lobe heavily sclerotised, forming a triangular lobe with sharp end in lateral view, while the posterior margin concave in the



FIGURE 22. *Graphium* (*Pazala*) *confucius* Hu, Duan & Cotton, **sp. nov.**; red dot: holotype, yellow dots: paratypes; upperside on the first row, underside on the second row; A and B: \Im , Kunming, Yunnan, China; C: \Im , Kunming, Yunnan, China; D and E: \Im , Dongchuan, Yunnan, China; F: \Im , Guiyang, Guizhou, China; scale bar = 10 mm.



FIGURE 23. Male genitalia of $G_{(P)}$ confucius Hu, Duan & Cotton **sp. nov.**, scale bar = 1.0 mm. All: genitalia as a whole, R.: lateral view of ring, TSU: dorsal view of tegumen, socii, and uncus, V. right valve, Ae.: lateral view of aedeagus, Ju.: ventral view of juxta.

middle, separating it into a "W" shape in ventral view, a short or blunt spur may be present at the end of each triangular part in some specimens, making the end more acute.

Differential Diagnosis: The new species closely resembles G(P) mandarinus and G(P) daiyuanae sp. n., but can be generally distinguished by the following characters: 1) obviously larger size and distinctly produced forewing apex; 2) the black discal band on the upperside of hindwing well developed; 3) the tornal spots on the underside of hindwing are usually bright yellow. In male genitalia, dorsal subterminal harpe large, almost reaching dorsal terminal harpe and forming a heart-shape, dorsal projection bayonet-shaped (Figures 23), distance between the base of socii 0.50–0.62 mm. The ostial lobe forming a triangular lobe in lateral view, while the middle part of posterior margin concave in a "W" shape in ventral view with a very short or blunt spur at each tip (Figure 24).

Type Material: Holotype: CHINA: ♂, Xi Chong (2,000 m), Kunming, C. Yunnan, 2015–V–31, S. J. Hu leg [KIZ, 0100010].

Paratypes: CHINA: 1Å, Zhujiang Yuan (the headwater of Pearl River), Zhanyi, N.E. Yunnan, 2016–III–19, Z. Chang leg. [SJH]; 1Å, Qingjiang Er Qiao (near G318 highway) (1,500 m), Lianglu, Tianquan, W. Sichuan, 2015–VII–27, Z. B. Chen leg. [ZBC]; 3Å, Huanghu Ping (1,350–1,450 m), Ziyun, Baoxing, W. Sichuan, 2015–VII–15, Z. B. Chen leg. [ZBC]; 1Å, Kouy-Tchéou Abbe Largeteau [leg.], Ex Oberthür Coll., Brit. Mus. 1927–3, Paralecto-type [round pale blue edged label], **PARALECTOTYPE** [in blue] *Papilo* [sic] *glycerion* var. *Mandarinus*

Oberthür C.R. Smith det. 2003, BMNH(E) #146242 [BMNH]; 1∂, Kouy-Tchéou Abbe Largeteau [leg.], China Kouytcheou [hand written], Pap. glycerion, var. Mandarinus, Oberthür Etud. d' Entomol. IVelivraison - appendice variété géographique chinoise [hand written], Levick Bequest B.M.1941-83, Type [round red edged label], Paralecto-type [round pale blue edged label], PARALECTOTYPE [in blue] Papilo [sic] glycerion var. Mandarinus Oberthür C.R. Smith det. 2003, BMNH(E) #149385 [BMNH]; 13, Siao-Lou, 1893, Chasseurs indigènes, Rothschild Bequest, BMNH(E) # 220118 [BMNH]; 1♂, Siao-Lou, 1901, Chasseurs indigènes du P. Déjean, Ex Oberthür Coll., BMNH(E) # 146048 [BMNH]; 13, Siao-Lou, 1893, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146063 [BMNH]; 13, Siao-Lou, 1893, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146065 [BMNH]; 1♂, Siao-Lou, 1900, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146110 [BMNH]; 1♂, Siao-Lou, 1900, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146101 [BMNH]; 13, Siao-Lou, 1900, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146106 [BMNH]; 13, Chasseurs indigènes de Ta-tsien-lou. Récolte de 1910, Ex Oberthür Coll., BMNH(E) # 145814 [BMNH]; 13, Chasseurs indigènes de Ta-tsien-lou. Récolte de 1910, Ex Oberthür Coll., BMNH(E) # 145820 [BMNH]; 13, Chasseurs indigènes de Ta-tsien-lou. Récolte de 1910, Ex Oberthür Coll., BMNH(E) # 145821 [BMNH]; 1∂, Chasseurs indigènes de Ta-tsien-lou. Récolte de 1910, Ex Oberthür Coll., BMNH(E) # 145822 [BMNH]; 1♀, Ichang, Rothschild Bequest, BMNH(E) # 220120 [BMNH]; 1Å, Jinfushan, Exp. Stotzner leg. [ZFMK]; 1Å, Siao-Lou, 1893, Chasseurs indigènes [ZFMK]; 1Å, Rou Bi Gou, Baoxing, Sichuan, 2006–VII–10, Ming Yue leg. [AMC]; 2∂, 1♀, Ya'an, Sichuan, 2006–VII, local catcher leg. [AMC]; 22♂, Wu Shi He, Hanyuan County, Sichuan, 2005–VII–5-20, local catcher leg. [AMC]; 1♂, 1♀, Daba Shan, E Sichuan, 2006–VII, local catcher leg. [AMC]; 13, Gioncheng, Guangxi, 1996–VI–17, local catcher leg. [AMC]; 4∂, 1♀, Luzhou, Sichuan, 2007–V, local catcher leg. [AMC]; 2∂, Kangding, Sichuan, 2006–VII, local catcher leg. [AMC]; 1∂, Qingshuilang Shan, Yunlong, W. Yunnan, 1998–VII-VIII, Qin et al leg., ex Coll. Jan Moonen [NBC]; 1⁽²⁾, Jinxiu, Dayao Shan, Guangxi, 1998–VII-VIII, Liang et al leg., ex Coll. Jan Moonen [NBC]; 2♂, Xi Chong (2,000 m), Kunming, C. Yunnan, 2015–V–19, S. J. Hu leg. [KIZ]; 1♀, ditto, 2015–V–31, S. J. Hu leg. [KIZ, 0100011]; 13, Xiao Moyu, Kunming, C. Yunnan, 2015–IV–18, S. J. Hu leg. [SJH]; 23, Forest Park (1,570 m), Dongchuan, N.E. Yunnan, 2014–V–17, local catcher leg. [KIZ, 0100012–0100013]; 13, the same collecting data, [SJH]; 16♂, 1♀, Luzhou, Sichuan, 2013–VII, local catcher leg. [SJH]; 2♀, Xiaoche He Park, Guivang, Guizhou, 2013–VI–11, S. J. Hu leg. [SJH]; 1∂, 1♀, Zuogong, E. Tibet, 2013–VI, local catcher leg. [SJH]; 2♂, 3♀, Jigong Shan (1,940 m), Shimian, W. Sichuan, 2013–VI–13, Q. Zeng leg. [SFU]; 3♂, Ziyun Xiang, Baoxing, W. Sichuan, 2012–VIII, local catcher leg. [SJH]; 13, Panzhihua, W. Sichuan, 2001–VIII–5, E. T. Ye leg. [SJH]; 1∂, Huize, N.E. Yunnan, 1998–IV, local catcher leg. [SFU]; 1∂, Zhongdian, N.W. Yunnan, 1992–VI–20, D. Z. Dong leg. [KIZ]; 1Å, Oianjiang (1,780), Sichuan, 1989–VII–14, S. Lin leg. [KIZ]; 4Å, Lichuan (800 m), Hubei, 1988–VIII–3, X. C. Liang, D. Z. Dong, and S. Lin leg. [KIZ]; 1♀, Leigong Shan, Guizhou, 1988–VIII–15, S. Lin leg. [KIZ]; 3♂, Emei Shan (1,400 m), W. Sichuan, 1984–IV–13, local catcher leg. [DB]; 1♂, ditto, 1982–IV–13, local catcher leg. [DB]; 13, Xi Shan, Kunming, C. Yunnan, 1979-VIII-9, J. Xiong leg. [KIZ]; 13, Jinghong, Xishuangbanna, S. Yunnan, 1979–VII–28, D. Z. Dong leg. [KIZ]. VIETNAM: 8♂, 10♀, Sapa (1,500 m), Lao Cai, 2015–V, L. T. Le leg. [SJH], 2, Ha Giang, 2008–VII, local catcher leg. [SJH]; 6, 8, Ha Giang, 2006–V, local catcher leg. [AMC]; 2♂, Ha Giang, 2007–VII, local catcher leg. [AMC]; 1♂, Ha Giang, 2008–VII, local catcher leg. [AMC].

The holotype and three paratypes were deposited in Kunming Institute of Zoology (KIZ), Chinese Academy of Sciences (Li *et al.* 2015).

Distribution: This species is commonly found in China (E. Tibet, Sichuan, Yunnan, Guizhou, Guangxi, Hubei, Hunan, Jiangxi, and Zhejiang); it is also collected from Sapa and Ha Giang, N. Vietnam, which could be the southernmost point of its range.

The two Kouy-Tchéou paralectotypes of *Papilio glycerion* var. *mandarinus* Oberthür actually belong to *G* (*P*.) *confucius* Hu, Duan & Cotton **sp. nov.**, and are designated as paratypes of the new species above. The precise location of Kouy-Tchéou is unclear. Several alternatives have been included in publications, sometimes stating that the name is equivalent to the modern Guizhou Province, but this seems unusually vague for a locality cited by Oberthür. Racheli & Cotton (2009) stated that Kiunglai [= Qionglai], W. Sichuan is the modern equivalent of Kouy-Tchéou without citing a source, and Racheli (pers. comm.) is unable to recall the origin of this interpretation. Baker (1995) listed a male specimen of *Euaspis basalis* from Kouy-Tchéou, stating the location to be "Kwangsi, Ku-i chou [= Guyi, N. Guangxi], $25^{\circ}46'N$ $109^{\circ}26'E''$. Since Donald Baker is deceased it was not possible to ascertain the reasoning behind this conclusion; but James Hogan, currently Collections Manager at Oxford

University Museum of Natural History where Baker conducted his study, stated (pers. comm.) that probably this was based on Baker's investigations of the location where Abbé Largeteau (the Catholic missionary cited in Oberthür's original description of *mandarinus*) collected specimens. Hogan stated that Baker was particularly interested in the history of entomology and old collectors, so he may possibly have found evidence to pinpoint the identity of the location. The current authors refrain from choosing a location for Kouy-Tchéou due to lack of precise evidence in favour of any one possibility. All of the alternatives fall within the known range of the new taxon.

Phenology: This species is multivoltine (three generations) in its type locality, the 1st generation flies in late April to early May, the 2nd generation flies from early June to July, and the 3rd generation flies in August. The phenology in other localities requires further study.



FIGURE 24. Female genitalia of $G_{(P_i)}$ confucius Hu, Duan & Cotton **sp. nov.**; lateral view above, ventral view below; scale bar = 1.0 mm.

Host plant: A female was observed ovipositing on *Litsea rubescens* (Lauraceae) at Xi Shan, Kunming, and the subsequent rearing experiment confirmed the identity of the butterfly. The species may use other plants of genus *Litsea* in other localities across China.

Derivatio nominis: The specific name of this new taxon is dedicated to Confucius (Chinese name Qiu Kong, courtesy name Zhongni; 551–479 BC), a Chinese teacher, editor, politician, and philosopher of the Spring and Autumn Period of Chinese history. The species name is treated as a noun in apposition.

Graphium (Pazala) mandarinus (Oberthür, 1879)

(Figure 25)

[*Papilio*] *Glycerion* variété géographique *Mandarinus* Oberthür, 1879, Étud. d'Ent., 4: 115; TL: "Moupin" [= Baoxing, W. Sichuan, China]. Type locality restricted to Moupin by lectotype designation as stated above. The name *mandarinus* is treated as a noun meaning a Mandarin rather than an adjective, as recommended by Gerardo Lamas (pers. comm.) since there is no clear indication in the original description.



FIGURE 25. Three known subspecies of *G*. (*P.*) mandarinus (Oberthür, 1879) from Sichuan and Indo-China; upperside in the first row, underside in the second row; scale bar = 10 mm. A: ssp. kimurai Murayama, 1982, \Im , Doi Inthanon, Chiang Mai, N. Thailand; B: ssp. kimurai Murayama, 1982, \Im , Mt. Ashaemyinanaukmyin, W. Shan State, Myanmar; C: ditto, \Im ; D: ssp. fangana (K. Okano, 1986), \Im , Doi Pu Meun, Chiang Mai, N. Thailand; E: ditto, \Im ; F: ssp. mandarinus, \Im , Baoxing, W. Sichuan. China; G: ssp. mandarinus, \Im , Daba Shan, W. Sichuan. China.

Diagnostic characters: Size varies among subspecies, the nominate subspecies is the largest, ssp. *kimurai* very slightly larger than ssp. *fangana* which is the smallest, and ssp. *stilwelli* **ssp. nov.** is intermediate and variable in size, some specimens are quite small and some larger. Both wings whitish and less glossy. Forewing, termen straight or slightly concave in the middle; greyish scales scattered between the 8th and 9th forewing bands. Hindwing, the "8"-shaped twisted discal black bands complete on the underside.

Male genitalia (Figure 26): In total male genitalia of 19 specimens, comprising two individuals of ssp. *fangana*, five individuals of ssp. *kimurai*, six individuals of ssp. *stilwelli* ssp. nov., and six individuals of ssp.

mandarinus, were dissected. Moderately to heavily sclerotised. Ring wavy in the upper half; saccus well developed; distance between the base of socii 0.50–0.64 mm (mean = 0.56 ± 0.05 , n = 19), without significant variation between subspecies. Valve round shaped, the dorsal terminal harpe with almost straight edge and acute but not elongate tip (blunt in *sichuanica* and *hoeneanus* nom. nov., stat. rev. and elongate in *garhwalica* stat. nov.); the dorsal subterminal harpe moderate in size (similar to that of *daiyuanae* **sp. nov.**), isolated or reaching the base of the tip of the dorsal terminal harpe (joining the tip of dorsal terminal harpe in *garhwalica* stat. nov.; smaller and completely isolated in *paphus* stat. nov., *sichuanica*, and *hoeneanus* nom. nov., stat. rev.; larger in *confucius* **sp. nov.**), the medial harpe straight, the dorsal projection finger-shaped (some with blunt or serrate end) (similar to that of *paphus* stat. nov. and *daiyuanae* **sp. nov.**; flat in *garhwalica* stat. nov., *sichuanica*, and *hoeneanus* nom. nov., stat. rev.); with or without a small tooth in the middle of the medial harpe.

Female genitalia (Figure 27): In total female genitalia of 4 specimens (including two ssp. *kimurai*, one ssp. *fangana*, and one ssp. *mandarinus*) were dissected, the general characters among subspecies are consistent. Lamella postvaginalis small; lamella antevaginalis broad horizontally, lined with weakly sclerotised longitudinal striae; ostial lobe heavily sclerotised, forming a triangular lobe with sharp end in lateral view, while the posterior margin shallowly concave in the middle, with a small short spur at each tip.



FIGURE 26. Male genitalia of $G_{(P)}$ mandarinus mandarinus (Oberthür, 1879) from Baoxing, Sichuan, W. China; scale bar = 1.0 mm. All: genitalia as a whole, R.: lateral view of ring, TSU: dorsal view of tegumen, socii, and uncus, V. right valve, Ae.: lateral view of aedeagus, Ju.: ventral view of juxta.

Graphium (Pazala) mandarinus kimurai Murayama, 1982

(Figure 25, A–C)

Graphium glycerion kimurai Murayama, 1982; New Entomologist, 31 (4): 1, f. 1; TL: "Doi Inthanon, N. Thailand" [Doi Inthanon, Chiang Mai Province, N. Thailand].

Diagnostic characters: Very small in size, forewing length: male 27.0–32.0 mm (mean = 29.5 ± 1.4 mm, n = 33), female 32.0–33.0 mm (only two females examined). Both wings rather whitish. Forewing, apex acute but not produced, termen straight; veins M₃ to Cu₂ usually free of black scales in male, but with only trace of black in female. Hindwing, black discal band on the upperside usually completely reduced in both sexes, submarginal black short lines also reduced on the upperside in both sexes, only reaching space m₁.

Distribution: C. Myanmar, N. Thailand, and C. Laos.

Phenology: Subspecies *kimurai* flies from late October through the cool season, peaking at the type locality in December and again in February. Whether this is a staggered single generation with two peaks of emergence or represents two separate generations is unknown. Murayama (1982), in the original description of the taxon, also lists two paratypes from Doi Inthanon caught on 7 April 1978 and a third paratype from Thung Salengluang, N. Thailand [= Thung Salaeng Luang National Park, Phitsanulok Province, N. Thailand] with date 10 July 1976. This is the only known record of a specimen of ssp. *kimurai* captured anywhere during the rainy season.

Host plant: Unknown, presumably a plant species belonging to family Lauraceae.



FIGURE 27. Female genitalia of $G_{(P)}$ mandarinus mandarinus (Oberthür, 1879) from Daba Shan, Sichuan, W. China; lateral view above, ventral view below; scale bar = 1.0 mm.

Graphium (Pazala) mandarinus fangana (K. Okano, 1986)

(Figure 25, D and E)

Pazala glycerion fangana K. Okano, 1986; Tokurana, 11 (3): 1, f. 3; TL: "near Fang, alt. 2,200 m, (about 20°20' N, 99°50' E), Chieng Mai Prov., North Thailand" [The co-ordinates stated in the original description are inaccurate, the correct coordinates are approximately 20°05' N, 99°08' E].

Diagnostic characters: Very similar to the preceding subspecies in appearance, forewing length: male 26.0–31.0 mm (mean = 29.2 ± 1.2 mm, n = 29), female 32.0 mm (only one female). Forewing, apex slightly blunt, also not produced, termen straight; vein M₃ and the base of vein Cu₁ tinged with black scales in both sexes. Hindwing, submarginal black short lines better developed than the preceding subspecies, which may reach space sc+r₁ in some specimens.

Distribution: Only known from a few mountains in northern Thailand near Mae Ai and Fang districts. The restricted range of this taxon effectively inside the potential range of the previous subspecies needs further study. It is notable that ssp. *kimurai* is absent from the mountains where *fangana* occurs, and the taxa are genetically distinct as well as separable on wing pattern.

Phenology: This subspecies is univoltine, flying only in February and March, unlike ssp. *kimurai*. Recently deceased Prasobsuk Sukkit (pers. comm.) confirmed that he and local collectors have looked for ssp. *fangana* throughout the year over many years, and never found it other than in February to March.

Host plant: Unknown, presumably a plant species belonging to family Lauraceae.

Graphium (Pazala) mandarinus stilwelli Cotton & Hu ssp. nov.

(Figure 28)

Description: Male: Forewing length: 29.0–37.0 mm (mean = 33.8 ± 1.5 mm, n = 81). Forewing triangulate, apex blunt and not produced, termen straight or slightly concave in the middle. Forewing upperside: whitish, the upper and outer 1/3 (mostly the discocellular, the subterminal, and the terminal areas) hyaline and glossy; 10 black bands between the base and the termen, of which the 1st to the 6th almost run parallel; the 1st and the 2nd bands reach the tornal margin; the 3rd band does not cross the cell vein and the 4th band crosses just beyond the discal cell but usually does not extend along vein CuA₁; the 5th band reaches the margin of the discal cell and usually extends along vein M_3 ; the 6th band at the end of cell is often connected with the 5th band by a transverse black fine line; the 7th band joins the 8th band just at vein M₂, extends to the tornus and joins the 9th band at vein Cu₂ or just below it in space cu_2 ; the 9th band curved inward in space r_4 ; the 10th band independent, extending along the margin from the apex to space cu_2 ; the area between the 8th and 9th bands without black scales; veins R_{4+5} to Cu_1 are black after meeting the 6th and 7th bands and then divide the whitish-hyaline areas between the 6th to 10th bands into spots. Forewing underside: colour and markings as upperside, but the hyaline areas are glossier. Hindwing triangulate in general, vein M₃ extends into a thin sword-like tail, termen slightly indented at the end of veins, the ends of spaces m, to cu, expand into lobes. Hindwing upperside: whitish with long white hair covering the inner 1/3; tornal margin blackish, with a small brown and roconial patch near the base, a thick black band extends from the costa towards the tornus and joins the tornal blackish area just above two obliquely neighbouring yellow spots; the discal bands absent or indicated by a faint black streak near the cell; the submarginal black band coupled, reaching space rs or m_i ; the terminal band single, reaching space $sc+r_i$; despite being interrupted by veins, the ends of each short band in each space touch or directly connect each other; all black bands mentioned above join in the black area at the end of spaces m, to cu, with a greyish-blue lunule in each space; tail black with white tip. Hindwing underside: colour and markings similar to upperside but with a creamy yellow hue, all black bands and markings well defined, especially the discal band, two black lines twist into an almost "8"-shaped pattern, with a short connection between the rings, the outer half of the upper ring of the "8" pale yellow and the inner half white, the edge of the lower ring extends along veins M_1 to Cu; a square creamy white spot above each greyish-blue lunule in the tornal area, the yellow spots at the tornus somewhat reduced and crowned by fine white lines.



FIGURE 28. *Graphium (Pazala) mandarinus stilwelli* Cotton & Hu ssp. nov.; red dot: holotype, yellow dots: paratypes; upperside on the first row, underside on the second row; A: \Diamond , Weixi, N.W. Yunnan, China; B: \Diamond , Yunxian, S.W. Yunnan, China; C: \Diamond , Tengchong, W. Yunnan, China, \mathbb{O} Philip Yik-Fui Lo (Kadoorie Farm and Botanic Garden, Hong Kong, China); D: \Diamond , Yingjiang, W. Yunnan, China; E: \Diamond , Chudu Razi Hills, Kachin State, Myanmar; F: \bigcirc , Loutsekiang, \mathbb{O} Zoologisches Forschungsinstitute und Museum Alexander Koenig (ZFMK), Bonn, Germany; scale bar = 10 mm.



FIGURE 29. Male genitalia of G(P) mandarinus stilwelli Cotton & Hu s**sp. nov.** from Tacheng, Weixi, N.W. Yunnan, China; scale bar = 1.0 mm. All: genitalia as a whole, R.: lateral view of ring, TSU: dorsal view of tegumen, socii, and uncus, V. right valve, Ae.: lateral view of aedeagus, Ju.: ventral view of juxta.

Extensive examination and comparison between the new subspecies of *Graphium mandarinus* and *Graphium paphus* stat. nov., which may be sympatric, showed the following difficulties of distinguishing them on morphological characters: (1) variable size: although on average *stilwelli* **ssp. nov.** is smaller than *paphus* stat. nov., the smallest individuals of the latter taxon are almost the same size as the largest individuals of the former, making it rather hard to apply size solely to identify individual specimens; (2) variable shape of forewing termen: generally the middle part of the forewing termen in *paphus* stat. nov. is concave while in all subspecies of *mandarinus* this is straighter, however some individuals of *stilwelli* **ssp. nov.** also showed this character, and some *paphus* stat. nov. have a straight termen. Fortunately, genitalic dissection showed no overlapping characters between the two taxa. Hence, currently, the most reliable method in distinguishing *Graphium paphus* stat. nov. and *Graphium mandarinus stilwelli* **ssp. nov.** is genitalic dissection.

Female: Only a single specimen from Loutsekiang [= upper Salween or upper Mekong valleys (Fisher, 1948), N.W. Yunnan] in ZFMK, Bonn, known to us. Similar to the male, wings broader and more rounded. Forewing length ~40 mm, estimated from photograph (Figure 34F).

Male genitalia (Figure 29): Similar to those of the other three subspecies, namely ssp. *mandarinus*, ssp. *kimurai*, and ssp. *fangana*.

Differential Diagnosis: Intermediate between the nominotypical subspecies, ssp. *kimurai* and ssp. *fangana*, from which it can be distinguished by the following characters: 1) forewing apex blunt, not produced, termen straight or slightly concave in the middle; 2) the postdiscal area of forewing very clear, mostly with only a trace of black scales near the base of vein M_3 , although occasionally well marked; 3) the black discal band on the upperside of hindwing absent or indicated before the end of the cell; 4) hindwing submarginal black lines reaching spaces m_1 or sc+ r_1 .

While molecular analysis does not definitively separate this taxon from the nominate subspecies it is morphologically distinct and thus worthy of subspecific status.

Type Material: Holotype: CHINA: ♂, Tacheng (1,900 m), Weixi, W. Yunnan, 2015–IV–29, S. J. Hu leg., [KIZ, 0100008].

Paratypes: CHINA: 1Å, Gazhi River (1,050 m), Yunxian, W. Yunnan, 2016–V–9, Z. C. Xiong leg., [KIZ, 0100009]; 2Å, Gaoligong Shan, Tengchong, W. Yunnan, 2015–V–20, P. Y. F. Lo leg. [SJH]; 1Å, Liuku (100 m), W. Yunnan, 2002–IV–20, unknown leg. [SFU]; 1Å, Zhongdian, N.W. Yunnan, 1992–VI–20, D. Z. Dong leg. [KIZ]; 1Å, Ailao Shan, Jingdong, W. Yunnan, 1992–V–29, D. Z. Dong leg. [KIZ]; 1Å, Xujia Ba, Jingdong, W. Yunnan, 1992–V–29, D. Z. Dong leg. [KIZ]; 1Å, Xujia Ba, Jingdong, W. Yunnan, 1984–III–16, unknown leg. [KIZ]; 6Å, west of Fugong, NW. Yunnan, 2009–IV–2-29, local catcher leg. [AMC]; 3Å, Pingpo, Dali, Yunnan, 2016–IV–25-V–20, local catcher leg. [AMC]; 1Å, Qingshuilang Shan, Yunlong, W. Yunnan, 1998–VII-VIII, Qin et al. leg., ex Coll. Jan Moonen, [NBC]; 1Å, Loutsekiang [ZFMK]. MYANMAR: 3Å, Chudu Razi Hills, Kachin State, 2008–IV–21, local catcher leg. [SJH]; 1Å, ditto, 2008–IV–23, local catcher leg. [SJH]; 1Å, ditto, 2008–IV–24, local catcher leg. [SJH]; 2Å, ditto, 2007–V–5-9, local catcher leg. [AMC]; 5Å, ditto, 2006–V–9, local catcher leg. [AMC]; 8Å, ditto, 2007–V–5-9, local catcher leg. [AMC]; 5Å, ditto, 2006–V–9, local catcher leg. [AMC]; 5Å, ditto, 2007–V–5-9, local catcher leg. [AMC]; 1m#, ditto, 2012–V–15, local catcher leg. [AMC]; 1m#, ditto, 2012–V–16, local catcher leg. [AMC]; 8Å, ditto, 2007–V–5-9, local catcher leg. [AMC]; 1m#, ditto, 2012–V–15, local catcher leg. [AMC]; 1m#, ditto, 2012–V–15, local catcher leg. [AMC]; 5Å, ditto, 2012–V–26, local catcher leg. [AMC]; 8Å, Kushin, E. of Renam, N. Kachin State, 1998–IV–25-V–4, local catcher leg. [AMC]; 5Å, ditto, Kushin, E. of Renam, N. Kachin State, 1998–IV–25-V–4, local catcher leg. [AMC]; 5Å, ditto, Kushin, E. of Renam, N. Kachin State, 1998–IV–27, local catcher leg. [PS].

The holotype and a paratype were deposited in Kunming Institute of Zoology (KIZ), Chinese Academy of Sciences (Li *et al.* 2015).

Distribution: Currently recorded in W. Yunnan and N.E. Myanmar.

Phenology: It appears to be bivoltine based on the collecting data of specimens examined, the first generation from April to June, and the second from July to August. More surveys are required to confirm the phenology.

Host plant: Unknown, presumably a plant species belonging to family Lauraceae.

Derivatio Nominis: The subspecific name of this new taxon is dedicated to General Joseph W. Stilwell (1883–1946) of the U.S. Army, who served in the China Burma India (CBI) Theatre and was instrumental in the liberation of W. Yunnan and Myanmar during World War II. The name is to be treated as a noun in apposition.

Graphium (Pazala) mandarinus mandarinus (Oberthür, 1879)

(Figure 25, F and G)

[Papilio] Glycerion variété géographique Mandarinus Oberthür, 1879, Étud. d'Ent., 4: 115; TL: "Moupin" [= Baoxing, W. Sichuan, China].

Pazala mandarina; Chou, 1994; Monographia Rhopalocerorum Sinensium: 55, 176 (unjustified emendation).

Diagnostic characters: The largest subspecies of *mandarinus*, forewing length: male 31.0–39.0 mm (mean = 36.4 \pm 1.5 mm, n = 48), female 39.5 mm (only one female examined). Forewing, apex blunt, not produced, termen straight or slightly concave in the middle; veins M₁ to Cu₂ usually tinged with black scales, especially near the base. Hindwing, black discal band on the upperside well developed in male but often reduced in female, submarginal black short lines well developed on the upperside in both sexes, reaching costa or at least space sc+r₁.

Distribution: Sichuan and W. Hubei, China.

Phenology: Probably bivoltine like the preceding subspecies; further surveys are required in future.

Host plant: This subspecies feeds on *Litsea populifolia* (Lauraceae) in W. Sichuan, China (Zhi-Bing Chen, pers. comm.).

Discussion

This study divided the two previously recognised species in the *mandarinus* group into seven species, based on morphological evidence and molecular phylogeny. It is also the first publication to provide photos of female genitalia of subgenus *Pazala*, and a case example for the classification of such morphologically similar species.

After such a revision, it is important to re-establish the distribution range of all *mandarinus* group species. On commencement of our study we recognised two species, *Graphium (Pazala) mandarinus* as a Sino-Himalayan element with five subspecies, plus G(P) sichuanica restricted to China proper (Racheli & Cotton 2009). Our study remapped the distribution range of the seven herein recognised species in the *mandarinus* group (Figure 30), and the pattern can be concluded as follows. 1) The entire species group remains a Sino-Himalayan element, extending from W. Himalaya to E. and S. China and N. Indochina. 2) Strong geographical endemism of taxa was observed: G(P) garhwalica stat. nov. in W. Himalaya of N.W. India and Nepal; G(P) paphus stat. nov. in E. Himalaya of Nepal, N. India, Bhutan, N.W. Myanmar, and S. Tibet of China; G(P) sichuanica in N. Sichuan and N. Hubei of China; G(P) hoeneanus nom. nov., stat. rev. in S. to E. China proper; G(P) daiyuanae **sp. nov.** currently only known from N. Vietnam; G(P) confucius **sp. nov.** in most of C. and W. China and N. Vietnam; and G(P) mandarinus in Myanmar, N. Indochina, W. Yunnan, W. Sichuan to N. Hubei of China with four subspecies.



FIGURE 30. Distribution map of the seven species in the *Graphium (Pazala) mandarinus* group, only the ten relevant countries, comprising India, Nepal, Bhutan, Bangladesh, China, Myanmar, Thailand, Laos, Vietnam, and Cambodia, are shown on this map.

In overall external appearance, the seven species in the *mandarinus* group recognised in this revision are similar, except for the shape of the discal band on the hindwing, which has long been used for distinguishing the old *'mandarinus'* and *'sichuanica'* (Racheli & Cotton 2009). Our analysis of a long series of specimens has discovered that some 'micro' morphological characters, including ground colour (dark or light), the extent of black tinge on forewing veins M_1 to Cu_2 , the extent of hindwing subterminal and terminal black bands, the colour of hindwing tornal yellow spots, size (measured as forewing length), and wing shape, can be adopted in association with genitalic structure to distinguish species and subspecies in the *mandarinus* group.

Similar examples were also commonly reported in genera like Heliophorus Geyer, 1832 (Riley 1929; Eliot

1963; Yago 2002; Yago *et al.* 2000, 2002), *Ahlbergia* Bryk, 1946, *Novosatsuma* Johnson, 1992, and *Cissatsuma* Johnson, 1992 of Lycaenidae, as well as genera like *Ypthima* Hübner, 1818 (Elwes & Edwards 1893; Shirôzu 1979; Takahashi 2000), *Euthalia* Hübner, [1819] (Yokochi 2010, 2011, 2012), and *Neptis* Fabricius, 1807 (Bozano 2008) of Nymphalidae. Many new taxa were recognised in these butterfly groups in recent decades based on thorough analysis of certain known taxa with wide distribution range and morphological variation (e.g., Huang 1999; Huang 2002; Yago 2002; Yago *et al.* 2002; Yago 2003; Yokochi 2011, 2012; Huang *et al.* 2016; Huang & Sun 2016; Huang & Zhu 2016). Within subgenus *Pazala*, *Graphium* (*Pazala*) *eurous* (Leech, 1893) and *G* (*P.*) *mullah* (Alphéraky, 1897) also exhibit morphological variation in their distribution range, which in total almost mirror that of the *mandarinus* group (Racheli & Cotton 2009).

The present study benefited largely from molecular biology to highlight and initially distinguish morphologically similar species of the *mandarinus* group. The partial mitochondrial gene (*cox1*) used in our study, namely the 'DNA barcode' sequence, is a powerful tool in identifying biological species (Hebert et al. 2003) and unveiling cryptic species (Hebert et al. 2004). The 'DNA barcoding' proposed a threshold for species delineation at 2-3% of nucleotide differentiation (K2P distance), and that for subspecies between 1-2% (Hebert *et al.* 2003). In our genetic distance analysis, the distances between most taxon pairs were above the threshold, except for a pair of species, G. (P.) daiyuanae sp. nov. and G. (P.) confucius sp. nov., with the K2P distance at 1.71% (Table 2). Nonetheless, the K2P distances between subspecies of G. (P.) mandarinus, except for those related to ssp. kimurai, were also under the threshold, ranging from 0.46–0.95% (Table 2). Judging from this criterion solely, it seemed that daiyuanae sp. nov. and confucius sp. nov. should be conspecific, and the three subspecies of mandarinus were inseparable. However, apart from their constant morphological and genitalic differences between each other, geographical distribution also supplied additional evidence to separate them. We examined specimens of both daiyuanae sp. nov. and confucius sp. nov. from Sapa, N. Vietnam, indicating that they cannot be conspecific with such constant morphological and genitalic differences. The geographic range of each of the three subspecies of mandarinus is clearly separated from the next, without any cline of morphological transition; therefore, these populations are isolated from each other by geographical barriers such as low altitude areas. Based on the abovementioned evidence, we were eventually able to separate them as distinct species or subspecies, regardless of limited genetic distances. Similar cases in Asian Papilionidae were also reported for Papilio maackii Ménétriès, 1859 vs. P. syfanius Oberthür, 1886 and P. elwesi Leech, 1889 vs. P. maraho Shiraki & Sonan, 1934, subspecies of Bhutanitis thaidina (Blanchard, 1871) and P. bianor Cramer, 1777 (Tsao & Yeh 2008; Lu et al. 2009; Zhu & Wu 2011; Condamine et al. 2013b; Zhao et al. 2017). Hence, although 'DNA barcoding' is a powerful analytical tool for taxonomy, one must keep in mind that the resolution of that sole short molecular marker can be limited in some congeneric species groups, especially infraspecific taxa. As widely discussed before, morphological, genitalic, and geographical evidence should also be taken into consideration equally with molecular evidence in modern taxonomy (Moritz & Cicero 2004; Will & Rubinoff 2004; Hajibabaei et al. 2007).

Limited genetic distances between certain taxa might be explained by our phylogenetic and molecular dating analyses. Using molecular dating, we estimated that the *mandarinus* group originated in the late Miocene (ca. 11.8 Ma), and that most phylogenetic events occurred in the late Pliocene to early Pleistocene: the divergence times between G. (P.) sichuanica and G. (P.) hoeneanus nom. nov., stat. nov. occurred ca. 2.9 Ma, the divergence times between G (P.) daiyuanae sp. nov. and G (P.) confucius sp. nov. occurred ca. 2.6 Ma, and the divergence times of the subspecies complex of G. (P.) mandarinus occurred ca. 2.8 Ma (Figure 4). This period was characterised by synchronous speciation in many other terrestrial organisms, as demonstrated by previous works in butterflies (Condamine et al. 2013a; Condamine et al. 2013b). Such speciation pattern in the late Pliocene and throughout the Pleistocene can be largely attributed to past geological and climatic events such as the change in the Eastern Himalayas or glaciation cycles initiated at the end of the Pliocene (see Sanmartín et al. 2001 for a review). Frequent tectonic shifts coupled with climatic oscillations contributed to allopatric (vicariance) divergences either due to physical barriers like mountains and valleys or isolations caused by dispersal within the distribution range of ancestral species. Such pattern of allopatric divergences has likely played the role of an initial driving force of speciation during this period. From the distribution map, we can see that most phylogenetic events that occurred in the late Pliocene throughout the Pleistocene are concentrated in the E. Himalaya-Hengduan Mountains and the adjacent N. Indochinese highlands (Figure 36). This area has been impacted by tectonic shifts and climate oscillations, which could have led to high speciation (Xing & Ree 2017). Taking the subspecies divergence of G (P.) mandarinus for instance, all infraspecific divergence times estimated by the molecular dating happened

between 0.8 and 2.8 Ma, which coincides with tectonic and climatic events that occurred on the region (Lü *et al.* 2014).

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References

- Alphéraky, S. (1897) Lépidoptères des provinces chinoises Sé-Tchouen et Kham recueillis, en 1893, par M-r G. N. Potanine. *In*: Romanoff, N.M. (Ed.), *Mémoires sur les Lépidoptères. Vol. 9.* P.P. Soïkine, St.-Pétersbourg, pp. 83–149.
- Ayres, D.L., Darling, A., Zwickl, D.J., Beerli, P., Holder, M.T., Lewis, P.O., Huelsenbeck, J.P., Ronquist, F., Swofford, D.L., Cummings, M.P., Rambaut, A. & Suchard, M.A. (2012) BEAGLE: an application programming interface and highperformance computing library for statistical phylogenetics. *Systematic Biology*, 61, 170–173. https://doi.org/10.1093/sysbio/syr100
- Baker, D.B. (1995) A review of the Asian species of the genus *Euaspis* Gerstäcker (Hymenoptera: Apoidea: Megachilidae). *Zoölogische Mededelingen, Leiden*, 69, 281–302.
- Bertheau, C., Schuler, H., Krumböck, S., Arthofer, W. & Stauffer, C. (2011) Hit or miss in phylogenetic analyses: the case of the cryptic NUMTs. *Molecular Ecology Resources*, 11, 1056–1059. https://doi.org/10.1111/j.1755-0998.2011.03050.x
- Borkhausen, M.B. (1788) *Naturgeschichte der Europäischen Schmetterlinge nach Systematischer Ordnung. 1.* Varrentrapp und Wenner, Frankfurt, 36 + 288 pp.
- Bozano, G.C. (2008) *Guide to the Butterflies of the Palearctic Region, Nymphalidae part III, Subfamily Limenitidinae, Tribe Neptini.* Omnes Artes, Milano, 77 pp.
- Bryk, F. (1946) Type Lycaena ferra for Satsuma Murray nec. Adams (trans.). Arkiv för Zoologi, 38, 50.
- Chou, I. (1994) *Monographia Rhopalocerorum Sinensium*. Henan Science and Technology Publishing House, Zhengzhou, 9 + 852 pp.
- Condamine, F.L., Sperling, F.A.H., Wahlberg, N., Rasplus, J.-Y. & Kergoat, G.J. (2012) What causes latitudinal gradients in species diversity? Evolutionary processes and ecological constraints on swallowtail biodiversity. *Ecology Letters*, 15, 267– 277.

https://doi.org/10.1111/j.1461-0248.2011.01737.x

Condamine, F.L., Sperling, F.A.H. & Kergoat, G.J. (2013a) Global biogeographical pattern of swallowtail diversification demonstrates alternative colonization routes in the Northern and Southern hemispheres. *Journal of Biogeography*, 40, 9–23.

https://doi.org/10.1111/j.1365-2699.2012.02787.x

- Condamine, F.L., Toussaint, E.F.A., Cotton, A.M., Genson, G.S., Sperling, F.A.H. & Kergoat, G.J. (2013b) Fine-scale biogeographical and temporal diversification processes of peacock swallowtails (*Papilio* subgenus *Achillides*) in the Indo-Australian Archipelago. *Cladistics*, 29, 88–111.
 - https://doi.org/10.1111/j.1096-0031.2012.00412.x
- Cotton, A.M. & Racheli, T. (2007) A preliminary annotated checklist of the Papilionidae of Laos with notes on taxonomy, phenology, distribution and variation (Lepidoptera, Papilionoidea). *Fragmenta Entomologica*, 38, 279–378. https://doi.org/10.4081/fe.2006.10
- de Nicéville, L. (1886) On some new Indian butterflies. *Journal of the Asiatic Society of Bengal*, 55, Part II (3), 249–256, pl. XI.
- de Nicéville, L. (1900) On new and little-known Lepidoptera from the Oriental Region. *Journal of Bombay Natural History Society*, 13 (1), 157–176, pls. CC–EE.
- Eliot, J.N. (1963) The Heliophorus epicles (Godart, 1823) (Lepidoptera: Lycaenidae) complex. Entomologist, 96, 169–180.
- Elwes, H.J. & Edwards, J. (1893) A revision of the genus *Ypthima*, with especial reference to the characters afforded by the male genitalia. *Transactions of the Entomological Society of London*, 41, 1–54. https://doi.org/10.1111/j.1365-2311.1893.tb02051.x
- Fabricius, J.C. (1807) Die neueste gattungs-eintheilung der schmetterlinge aus den Linnéichen Gattungen *Papilio* und *Sphinx*. *Magazin für Insektenkunde*, 6, 277–285.
- Fisher, K.J. (1948) Some geographical notes on West China localities. Entomologist, 81, 213-219.
- Folmer, O., Black, M.B., Hoch, W., Lutz, R.A. & Vrijehock, R.C. (1994) DNA primers for amplification of mitochondrial cytochrome *c* oxidase subunit I from diverse metazoan invertebrates. *Molecular Marine Biology and Biotechnology*, 3, 294–299.
- Geyer, C. (1832) Zuträge zur Sammlung exotischer Schmettlinge. Vol. 4. im Verlag der Hübner'schen Werke, Augsburg, 48 pp. [in German]
- Gray, G.R. (1831) Description of eight new species of Indian butterflies, (*Papilio*, Lin.) from the collection of General Hardwicke. *Zoological Miscellany*, 1, 32–33.
- Gray, G.R. [1853] Catalogue of Lepidopterous Insects in the Collection of the British Museum. Part I. Papilionidae. Taylor and Francis, London, iii + 84 pp.
- Guindon, S. & Gascuel, O. (2003) A simple, fast and accurate method to estimate large phylogenies by maximum-likelihood. *Systematic Biology*, 53, 696–704.

https://doi.org/10.1080/10635150390235520

- Hajibabaei, M., Singer, G.A.C., Hebert, P.D.N. & Hickey, D.A. (2007) DNA barcoding: how it complements taxonomy, molecular phylogenetics and population genetics. *Trends in Genetics*, 23, 167–172. https://doi.org/10.1016/j.tig.2007.02.001
- Hall, T.A. (1999) BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. *Nuclear Acids Symposium Series*, 41, 95–98.
- Hancock, D.L. (1983) Classification of the Papilionidae (Lepidoptera): a phylogenetic approach. Smithersia, 2, 1–48.
- Hebert, P.D.N., Cywinska, A., Ball, S.L. & de Waard, J.R. (2003) Biological identifications through DNA barcodes. *Proceedings of the Royal Society of London*, 270, 313–321. https://doi.org/10.1098/rspb.2002.2218
- Hebert, P.D.N., Penton, E.H., Burns, J.M., Janzen, D.H. & Hallwachs, W. (2004) Ten species in one: DNA barcoding reveals cryptic species in the neotropical skipper butterfly *Astraptes fulgerator*. *Proceedings of the National Academy of Sciences* of the United States of America, 101, 14812–14817. https://doi.org/10.1073/pnas.0406166101
- Hemming, F. (1934) Revisional notes on certain species of Rhopalocera (Lepidoptera). *Stylops*, 3, 193–200. https://doi.org/10.1111/j.1365-3113.1934.tb01578.x

Huang, H. (1999) Some new butterflies from China 1 (Rhopalocera). Lambillionea, 94, 642-676.

- Huang, H. (2002) Some new nymphalids from the valley of Nujiang and Dulongjiang, China (Lepidoptera, Nymphalidae). *Atalanta*, 33, 339–360.
- Huang, H. & Sun, W.H. (2016) Ahlbergia bijieensis spec. nov. from Guizhou, China. Atalanta, 47, 151-160.
- Huang, H., Wang, J.Y. & Chen, Z. (2016) A new species of the *Euthalia* subgenus *Limbusa* from the Yarlung Tsangpo Grand Canyon, South-Eastern Tibet (Lepidoptera, Nymphalidae). *Atalanta*, 47, 205–210.
- Huang, H. & Zhu, J.Q. (2016) *Ahlbergia maoweiweii* sp. n. from Shaanxi, China with revisional notes on similar species (Lepidoptera: Lycaenidae). *Zootaxa*, 4114 (4), 409–433.

https://doi.org/10.11646/zootaxa.4114.4.3

- Hübner, J. (1818) Zuträge zur Sammlung exotischer Schmettlinge. Vol. 1. bey dem Verfasser zu finden, Augsburg, 40 pp., 35 pls. [in German]
- Hübner, J. [1819] Verzeichniss Bekannter Schmettlinge. Vol. 3. bey dem Verfasser zu finden, Augsburg, 16 pp. [in German]
- Hübner, J. [1819] Verzeichniss Bekannter Schmettlinge. Vol. 6. bey dem Verfasser zu finden, Augsburg, 16 pp. [in German]
- Hübner, J. [1821] Sammlung Exotischer Schmetterlinge. Vol. 2. im Verlag der Hübner'schen Werke, Augsburg, 24 pp. [in German]

Johnson, K. (1992) The Palaearctic "elfin" butterflies (Lycaenidae, Theclinae). Neue Entomologische Nachrichten, 29, 5-141.

- Katayama, T. (1988) Notes on *Pazala glycerion* (Gray), with description of a new subspecies from North-west India. *Gekkan-Mushi*, 203, 8–9, pl. 1. [in Japanese with English description]
- Kimura, M. (1980) A simple method for estimating evolutionary rates of base substitutions through comparative studies of nucleotide sequences. *Journal of Molecular Evolution*, 16, 111–120. https://doi.org/10.1007/BF01731581
- Koiwaya, S. (1993) Description of three new genera, eleven new species and seven new subspecies of butterflies from China. *Studies of Chinese Butterflies*, 2, 43–111, pl. 9–27. [in Japanese with English descriptions]
- Leech, J.H. (1893–94) Butterflies from China, Japan, and Corea. Part II. R. H. Porter, London, 385 pp., 15 pls. [pp. 297–681, pls. 29–43]
- Lepage, T., Bryant D., Philippe H. & Lartillot N. (2007) A general comparison of relaxed molecular clock models. *Molecular Biology and Evolution*, 24, 2669–2680. https://doi.org/10.1093/molbev/msm193
- Li, K.Q., Wang, Y.Z., Dong, D.Z. & Zhang, L.K. (2015) Catalog of insect type specimens preserved at the Kunming Institute of Zoology, Chinese Academy of Science with corrections of some specimens. *Zoological Research*, 36, 263–284. https://doi.org/10.13918/j.issn.2095-8137.2015.5.263.
- Lu, C.C., Wu, L.W., Jiang, G.F., Deng, H.L., Wang, L.H., Yang, P.S. & Hsu, Y.F. (2009) Systematic status of *Agehana elwesi f. cavaleriei* based on morphological and molecular evidence. *Zoological Studies*, 48, 270–279.
- Lü, J., Hu, S.J., Ma, X.Y., Chen, J.M., Li, Q.Q. & Ye, H. (2014) Origin and expansion of the Yunnan shoot borer, *Tomicus yunnanensis* (Coleoptera: Scolytinae): a mixture of historical natural expansion and contemporary human-mediated relocation. *PLoS ONE*, 9, e111940.

https://doi.org/10.1371/journal.pone.0111940

- Mell, R. (1923) Noch unbeschriebene Lepidopteren aus Südchina. II. Deutsche Entomologische Zeitschrift, 1923, 153–160.
- Mell, R. (1935) Noch unbeschriebene chinesische Lepidopteren. IV. Mitteilungen der Deutschen Entomologischen Gesellschaft, 6, 36-38.
- Miller, J. S. (1987) Phylogenetic studies in the Papilioninae (Lepidoptera: Papilionidae). Bulletin of the American Museum of Natural History, 186, 365–512.
- Miller, M.A., Schwartz, T., Pickett, B.E., He, S., Klem, E.B., Scheuermann, R.H., Passarotti, M., Kaufman, S. & O'Leary, M.A. (2015) A RESTful API for access to phylogenetic tools via the CIPRES science gateway. *Evolutionary Bioinformatics Online*, 11, 43–48.

https://doi.org/10.4137/EBO.S21501

Moritz, C. & Cicero, C. (2004) DNA barcoding: promise and pitfalls. *PLoS Biology*, 2, e354. https://doi.org/10.1371/journal.pbio.0020354

- Moore, F. (1888) Additional Species Family Papilionidae. In: Hewitson & Moore (1879–1888), Description of New Indian Lepidopterous Insects from the Collection of the Late Mr. W.S. Atkinson, M.A., F.L.S., & c., Director of the Public Instruction, Bengal. Rhopalocera. Part III. Asiatic Society of Bengal, Calcutta, pp. 283–285.
- Mutanen, M., Kivelä, S.M., Vos, R.A., Doorenweerd, C., Ratnasingham, S., Hausmann, A., Huemer, P., Dincă, V., van Nieukerken, E.J., Lopez-Vaamonde, C., Vila, R., Aarvik, L., Decaëns, T., Efetov, K.A., Hebert, P.D.N., Johnsen, A., Karsholt O., Pentinsaari, M., Rougerie, R., Segerer, A., Tarmann, G., Zahiri, R. & Godfray, H.C.J. (2016) Species-level para- and polyphyly in DNA barcode gene trees: strong operational bias in European Lepidoptera. *Systematic Biology*, 65, 1024–1040.

https://doi.org/10.1093/sysbio/syw044

- Munroe, E. (1961) The classification of the Papilionidae (Lepidoptera). Canadian Entomologist Supplement, 17, 1–51.
- O'Neill, S.L., Giordano, R., Colbert, A.M.E., Karr, T.L. & Robertson, H.M. (1992) 16S rRNA phylogenetic analysis of the bacterial endosymbionts associated with cytoplasmic incompatibility in insects. *Proceedings of the National Academy of Sciences of the United States of America*, 89, 2699–2702. https://doi.org/10.1073/pnas.89.7.2699
- Oberthür, C. (1876) Lépidoptères nouveaux de la Chine. Études d'Entomologie, 2, 13–34, pl. 1–4. [in French]
- Oberthür, C. (1879) Catalogue raissoné des Papilionidae de la collection de Ch. Oberthür. Études d'Entomologie, 4, 19–102, 107–117, pls. I–VI. [in French]
- Okano, K. (1986) Two new subspecies of butterflies from Thailand. Tokurana, 11, 1-8.
- Posada, D. (2008) jModelTest: phylogenetic model averaging. *Molecular Biology and Evolution*, 25, 1253–1256. https://doi.org/10.1093/molbev/msn083
- Racheli, T. & Cotton, A.M. (2009) *Guide to the Butterflies of the Palearctic Region. Papilionidae. Part I.* Omnes Artes, Milano, 70 pp.
- Rehm, P., Borner, J., Meusemann, K., von Reumont, B.M., Simon, S., Hadrys, H., Misof, B. & Burmester, T. (2011) Dating the arthropod tree based on large-scale transcriptome data. *Molecular Phylogenetics and Evolution*, 61, 880–887. https://doi.org/10.1016/j.ympev.2011.09.003
- Riley, N.D. (1929) Revisional notes on the genus *Heliophorus* with descriptions of new forms. *Journal of the Bombay Natural History Society*, 33, 384–402.

Murayama, S. (1982) Notes on some butterflies of North Thailand and China. New Entomologist, 31, 1-6.

- Ronquist, F., Teslenko, M., Van Der Mark, P., Ayres, D.L., Darling, A., Höhna, S., Larget, B., Liu, L., Suchard, M.A. & Huelsenbeck, J.P. (2012) MrBayes 3.2: efficient Bayesian phylogenetic inference and model choice across a large model space. *Systematic Biology*, 61, 539–542. https://doi.org/10.1093/sysbio/sys029
- Saigusa, T., Nakanishi, A., Shima, H. & Yata, O. (1982) Phylogeny and Geographical Distribution of the Swallow-Tail Subgenus Graphium (Lepidoptera: Papilionidae). *Entomologia Generalis*, 8, 59–69.
- Sanmartín, I., Enghoff, H. & Ronquist, F. (2001) Patterns of animal dispersal, vicariance and diversification in the Holarctic. *Biological Journal of the Linnean Society*, 73, 345–390. https://doi.org/10.1111/i.1095-8312.2001.tb01368.x
- Schenk, J.J. (2016) Consequences of secondary calibrations on divergence time estimates. *PLoS ONE*, 11, e0148228. https://doi.org/10.1371/journal.pone.0148228
- Scopoli, J.A. (1777) Introductio ad historiam naturalem sistens genera lapidum, plantarum, et animalium. Gerle, Prague, x + 506 pp.
- Shiraiwa, K., Cong, Q. & Grishin, N.V. (2014) A new *Heraclides* swallowtail (Lepidoptera, Papilionidae) from North America is recognised by the pattern on its neck. *Zookeys*, 468, 85–135. https://doi.org/10.3897/zookeys.468.8565
- Shirôzu, T. (1979) On the natural groups and their phylogenetic relationships of the genus *Ypthima* Hübner mainly from Asia (Lepidoptera: Satyridae). *Sieboldia*, 4, 231–295.
- Smith, C.R. & Vane-Wright, R.I. (2001) A review of the Afrotropical species of the genus *Graphium* (Lepidoptera: Rhopalocera: Papilionidae). *Bulletin of the British Museum (Natural History) Entomology*, 70, 503–719.
- Song, H., Buhay, J.E., Whiting, M.F. & Crandall, K.A. (2008) Many species in one: DNA barcoding overestimates the number of species when nuclear mitochondrial pseudogenes are coamplified. *Proceedings of the National Academy of Sciences of the United States of America*, 105, 13486–13491. https://doi.org/10.1073/pnas.0803076105
- Takahashi, M. (2000) A revision of the *Ypthima sakra* group (Lepidoptera, Satyridae) in Taiwan, China, with description of a new species. *Transactions of the Lepidopterological Society of Japan*, 51, 1–18. [in Japanese with English summary]
- Tamura, K., Stecher, G., Peterson, D., Filipski, A. & Kumar, S. (2013) MEGA6: Molecular evolutionary genetics analysis version 6.0. *Molecular Biology and Evolution*, 30, 2725–2729. https://doi.org/10.1093/molbev/mst197
- Thompson, J.D., Higgins, D.G. & Gibson, T.J. (1994) CLUSTAL W: improving the sensitivity of progressive multiple sequence alignment through sequence weighting, position-specific gap penalties and weight matrix choice. *Nuclear Acids Research*, 22, 4673–4680.

https://doi.org/10.1093/nar/22.22.4673

Thorne, J.L. & Kishino, H. (2002) Divergence time and evolutionary rate estimation with multilocus data. *Systematic Biology*, 51, 689–702.

https://doi.org/10.1080/10635150290102456

Thorne, J.L., Kishino, H. & Painter, I.S. (1998) Estimating the rate of evolution of the rate of molecular evolution. *Molecular Biology and Evolution*, 15, 1647–1657.

https://doi.org/10.1093/oxfordjournals.molbev.a025892

- Tsao, W.C. & Yeh, W.B. (2008) DNA-Based discrimination of subspecies of swallowtail butterflies (Lepidoptera: Papilioninae) from Taiwan. *Zoological Studies*, 47, 633–643.
- Wang, X. & Gu, M.B. (2017) Species of butterflies and its host in Nanling National Nature Reserve. *Journal of Zhejiang* Forestry Science and Technology, 36, 37–45.
- Will, K.W. & Rubinoff, D. (2004) Myth of the molecule: DNA barcodes for species cannot replace morphology for identification and classification. *Cladistics*, 20, 47–55. https://doi.org/10.1111/j.1096-0031.2003.00008.x
- Xing, Y. & Ree, R.H. (2017) Uplift-driven diversification in the Hengduan Mountains, a temperate biodiversity hotspot. Proceedings of the National Academy of Sciences of the United States of America, 114, E3444–E3451. https://doi.org/10.1073/pnas.1616063114
- Yago, M. (2002) Comparative morphology and identification of the subgenus Kulua, with description of a new species from Vietnam (Lepidoptera, Lycaenidae, Heliophorus). Tijdschrift voor Entomologie, 145, 145–171. https://doi.org/10.1163/22119434-900000107
- Yago, M. (2003) A new species of *Heliophorus* Geyer from Nepal, with a key to the Nepalese species (Lepidoptera, Lycaenidae). *Bulletin de la Société Entomologique de France*, 108, 27–34.
- Yago, M., Saigusa, T. & Nakanishi, A. (2000) Rediscovery of *Heliophorus yunnani* D'Abrera and its systematic position with intrageneric relationship in the genus *Heliophorus* (Lepidoptera: Lycaenidae). *Entomological Science*, 3, 81–100.
- Yago, M., Saigusa, T. & Nakanishi, A. (2002) A Revision of the *Heliophorus kohimensis* group (Lepidoptera: Lycaenidae). *Entomological Science*, 5, 375–388.
- Yokochi, T. (2010) Revision of the subgenus *Limbusa* Moore, [1897] (Lepidoptera, Nymphalidae, Adoliadini) Part 1. Systematic arrangement and taxonomic list. *Bulletin of the Kitakyushu Museum of Natural History and Human History*, Series A, 8, 19–67.

- Yokochi, T. (2011) Revision of the subgenus *Limbusa* Moore, [1897] (Lepidoptera, Nymphalidae, Adoliadini) Part 2. Group division and description of species (1). *Bulletin of the Kitakyushu Museum of Natural History and Human History*, Series A, 9, 9–106.
- Yokochi, T. (2012) Revision of the subgenus *Limbusa* Moore, [1897] (Lepidoptera, Nymphalidae, Adoliadini) Part 3. Description of species (2). *Bulletin of the Kitakyushu Museum of Natural History and Human History*, Series A, 10, 9–100.
- Zhao, J., Hu, S.J., He, Q.J., Yi, C.H., He, J., Feng, Z.W., Yang, J.H. & Chen, P. (2017) Genetic diversity of *Bhutanitis thaidina* Blanchard, based on mitochondrial COI and ND1. *Journal of Fujian Agricultural and Forestry University, Natural Science Edition*, 46, 387–391. [in Chinese with English abstract]
- Zhu, L.X. & Wu, X.B. (2011) Phylogenetic evaluation of the taxonomic status of *Papilio maackii* and *P. syfanius* (Lepidoptera: Papilionidae). *Zoological Research*, 32, 248–254. https://doi.org/10.3724/SPJ.1141.2011.03248

APPENDIX 1. List of Graphium (Pazala) mandarinus group specimens examined.

Names of depositories are given in alphabetical order, with institutions listed after private collections, and are abbreviated as follows: AMC: collection of Adam M. Cotton (Chiang Mai, Thailand); DB: collection of Danny Burk (South Bend, Indiana, USA); SJH: collection of Shao-Ji Hu (Kunming, China); JQZ: collection of Jian-Qing Zhu (Shanghai, China); PKU: Peking University, Beijing, China; PS: collection of Prasobsuk Sukkit (†) (Chiang Mai, Thailand), now housed in AMC collection; PYFL: collection of Philip Yik-Fui Lo (Kadoorie Farm and Botanic Garden, Hong Kong, China); TR: collection of Tommaso Racheli (Rome, Italy); XZ: collection of Xin Zhang (Kunming, China); ZBC: collection of Zhi-Bing Chen (Shanghai, China); BMNH: collections of the Natural History Museum, (London, United Kingdom); KIZ: Kunming Institute of Zoology, Chinese Academy of Sciences (Kunming, Yunnan); NBC: Naturalis Biodiversity Center, Leiden, The Netherlands; SFU: collection of Southwest Forest University (Kunming, China); ZFMK: collection of Zoologisches Forschungsinstitute und Museum Alexander Koenig (Bonn, Germany).

Graphium (Pazala) garhwalica (Katayama, 1988) stat. nov.

- INDIA: 1[↑], **PARATYPE**, Joshimath, Uttaranchal, 1987–IV–26, Toshihiko Katayama leg. [AMC]; 1[♀], Joshimath, Uttaranchal, 2007–IV–28, Peter Smetacek leg. [AMC].
- NEPAL: 1⁽²⁾, Banthanti, 2300 m., Annapurna District, 1988–V–5-7, Toshihiko Katayama leg. [AMC].

Graphium (Pazala) paphus (de Nicéville, 1886) stat. nov.

NEPAL: 1 SYNTYPE [of *Papilio glycerion* Gray, 1831], 'Nepaul, Hardwicke Bequest', BMNH(E) # 149384, [BMNH] [designated as **LECTOTYPE** herein]; 1 Shivapuri Hill, 2700 m., Kathmandu, 1988–IV–29-V–2, Toshihiko Katayama leg. [AMC].

CHINA: 2⁽⁷⁾, Jiefang Qiao (700–800 m), Bipung, Metok, S.E. Tibet, 2016–IV–15-17, X. W. Zhao leg. [SJH];

- INDIA: ♂ HOLOTYPE [of Papilio paphus de Nicéville, 1886], Sikkim, BMNH(E) # 145642, [BMNH]; 1♂, Air to Gna-tong, Sikkim, 7-12500ft, 1894–IV, Bingham leg., Rothschild Bequest, BMNH(E) # 220230 [BMNH]; 1♂, Sikkim, Dudgeon leg., Moore Bequest [BMNH]; 1♀, Sikkim, 1-4000 ft., Möller leg., Rothschild Bequest [BMNH]; 1♂, Tiger Hill, Darjeeling, 1979–V–19, Yutaka Inayoshi leg. [AMC], 10♂, 1♀, ditto, 1996–IV–21-V–5, Prasobsuk Sukkit leg. [AMC].
- MYANMAR: 2⁽³⁾, Tarung Hka River, N. Sagaing State, 2008–V–6-9, local catcher leg. [SJH]; 1⁽³⁾, ditto, 2008–IV–28, local catcher leg. [SJH], 8⁽³⁾, ditto, 2008–IV–22-V–17, local catcher leg. [AMC].

Graphium (Pazala) sichuanica (Koiwaya, 1993)

CHINA: 1∂, Bashan (Chengkou Xian), Daba Shan Mts. 1600-1900 m, 32° 09' N/ 108° 04' E, 1994–V–9, Native collector leg. [BMNH]; 1♀, Chengkou, Daba Shan, 1994–V–17, local catcher leg. [BMNH]; 1♂, Baokang, Hubei, 1996–V-VI, local catcher leg. [BMNH]; 2♂, Shabazi, Laohegou, Pingwu, N.E. Sichuan, 2013–IV–15, Y. Lei leg., [PKU]; 1♂, Shuijing Ping, Laohegou, Pingwu, N.E. Sichuan, 2013–IV–16, Y. Lei leg., [PKU]; 1♂, Roadside of Shuangshitou, Laohegou, Pingwu, N.E. Sichuan, 2013–IV–16, Q. Wen leg., [PKU]; 1♂, Xiao Hezigou, Laohegou, Pingwu, N.E. Sichuan, 2013–IV–20, J.L. Li leg., [PKU]; 1♂, Xiao Hezigou, Laohegou, Pingwu, N.E. Sichuan, 2013–V–20, J.L. Li leg., [PKU]; 1♂, Xiao Hezigou, Laohegou, Pingwu, N.E. Sichuan, 2013–V–9, Y. Lei leg., [PKU]; 3♂, Sigou, Tangjiahe, Qingchuan, N. Sichuan, 2014–IV–16, D. Zhang & C. Fan leg., [PKU]; 4♂, Guozishugou, Tangjiahe, Qingchuan, N. Sichuan, 2014–IV–18, D. Zhang & C. Fan leg., [PKU]; 2♂, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–IV–19, D. Zhang & C. Fan leg., [PKU]; 1♂, Sichuan, 2014–V–4, C. Fan leg., [PKU]; 1♂, Baixiong Ping, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–5, C. Fan leg., [PKU]; 3♂, ditto, 2014–V–17, C. Fan leg., [PKU]; 1♂, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–17, C. Fan leg., [PKU]; 1♂, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg., [PKU]; 1⊘, Shibangou, Tangjiahe, Qingchuan, N. Sichuan, 2014–V–18, C. Fan leg

Graphium (Pazala) hoeneanus Cotton & Hu nom. nov., stat. rev.

CHINA: 1♂ **SYNTYPE** [of *Papilio tamerlanus hoenei* Mell, 1935], Ost Tien Mu Shan, 1500 m., Prov. Chekiang, 1931–IV–18, H. Höne leg., BMNH(E) # 149381, [BMNH]; 1♂ **SYNTYPE** [of *Papilio tamerlanus hoenei* Mell, 1935], Ost Tien Mu Shan, Prov. Chekiang, 1931–IV–16, H. Höne leg. [TR]; 24♂ **SYNTYPES** [of *Papilio tamerlanus hoenei* Mell, 1935], Ost

Tien Mu Shan, Prov. Chekiang, 1931–IV–6-V–16, H. Höne leg. [ZFMK]; 6 $\stackrel{\circ}{\circ}$ **SYNTYPES** [of *Papilio tamerlanus hoenei* Mell, 1935], West Tien Mu Shan, Prov. Chekiang, 1931–IV–20-29, H. Höne leg. [ZFMK]; 4 $\stackrel{\circ}{\circ}$, Lingui (360 m), Guilin, Guangxi, 2013–IV–22, local catcher leg. [SJH]; 5 $\stackrel{\circ}{\circ}$, Lingui County, Guilin, Guangxi, 2013–III, local catcher leg. [AMC]; 15 $\stackrel{\circ}{\circ}$ Yue Cheng Ling, 1000 m., Ziyuan (Dahe), Guangxi, 2007–IV-V, local catcher leg. [AMC]; 1 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (400 m), Zhejiang, 2009–IV–10, J. Q. Zhu leg. [JQZ]; 2 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (350 m), Zhejiang, 2009–IV–11, J. Q. Zhu leg. [JQZ]; 2 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (350 m), Zhejiang, 2009–IV–11, J. Q. Zhu leg. [SJH]; 1 $\stackrel{\circ}{\circ}$, W. Tianmu Shan, Zhejiang, 2009–IV–21 (reared), J. Q. Zhu leg. [JQZ]; 2 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (400 m), Zhejiang, 2009–V–7, J. Q. Zhu leg. [JQZ]; 3 $\stackrel{\circ}{\circ}$, 1 $\stackrel{\circ}{\circ}$, Ningbo, Zhejiang, 2009–IV–18, J. Q. Zhu leg. [JQZ]; 3 $\stackrel{\circ}{\circ}$, 1 $\stackrel{\circ}{\circ}$, Tiantong (400 m), Ningbo, Zhejiang, 2008–IV–8, J. Q. Zhu leg. [JZQ]; 2 $\stackrel{\circ}{\circ}$, 1 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (400 m), Zhejiang, 2008–IV–8, J. Q. Zhu leg. [JZQ]; 2 $\stackrel{\circ}{\circ}$, 1 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (400 m), Zhejiang, 2008–IV–8, J. Q. Zhu leg. [JZQ]; 2 $\stackrel{\circ}{\circ}$, 1 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (400 m), Zhejiang, 2008–IV–8, J. Q. Zhu leg. [JZQ]; 2 $\stackrel{\circ}{\circ}$, 1 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (400 m), Zhejiang, 2008–IV–8, J. Q. Zhu leg. [JZQ]; 2 $\stackrel{\circ}{\circ}$, 1 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (400 m), Zhejiang, 2008–IV–8, J. Q. Zhu leg. [JZQ]; 2 $\stackrel{\circ}{\circ}$, 1 $\stackrel{\circ}{\circ}$, W. Tianmu Shan (400 m), Zhejiang, 2008–IV–24-25, J. Q. Zhu leg. [JQZ].

Graphium (Pazala) daiyuanae Hu, Zhang & Cotton sp. nov.

VIETNAM: ♂ HOLOTYPE, Sapa (1,600 m), Lao Cai Province, 2015–VI, L. T. Le leg. [KIZ, 0100000]; 3♂, 2♀ PARATYPES, same collecting data as holotype (1,600–1,700 m) [KIZ]; 8♂, 8♀ PARATYPES, same collecting data as holotype (1,600–1,700 m) [SJH]; 1♂, 4♀ PARATYPES, Sapa, Lao Cai Province, 2006–IV, local catcher leg. [AMC], 2♂ PARATYPES, ditto, 2007–V, local catcher leg. [AMC], 1♂ PARATYPE, ditto, 2007–VI, local catcher leg. [AMC]. 2♂ PARATYPES, Mucangchai (1,700 m), Yen Bai Province, 2017–V, L. T. Le leg. [KIZ, 0100006–0100007]; 3♂, 1♀ PARATYPES, same collecting data, [SJH].

Graphium (Pazala) confucius Hu, Duan & Cotton sp. nov.

CHINA: 13 HOLOTYPE, Xi Chong (2,000 m), Kunming, C. Yunnan, 2015–V–31, S. J. Hu leg. [KIZ, 0100010]; 13 PARATYPE, Zhujiang Yuan (the headwater of Pearl River), Zhanyi, N.E. Yunnan, 2016–III–19, Z. Chang leg. [SJH]; 13 PARATYPE, Qingjiang Er Qiao (near G318 highway) (1,500 m), Lianglu, Tianquan, W. Sichuan, 2015-VII-27, Z. B. Chen leg. [ZBC]; 3 PARATYPES, Huanghu Ping (1,350–1,450 m), Ziyun, Baoxing, W. Sichuan, 2015–VII–15, Z. B. Chen leg. [ZBC]; 2♂ **PARATYPE**, Xi Chong (2,000 m), Kunming, C. Yunnan, 2015–V–19, S. J. Hu leg. [SJH]; 1♀ PARATYPES, ditto, 2015-V-31, S. J. Hu leg. [KIZ, 0100011]; 1 PARATYPE, Xiao Moyu, Kunming, C. Yunnan, 2015–IV–18, S. J. Hu leg. [SJH]; 23 PARATYPES, Forest Park (1,570 m), Dongchuan, N.E. Yunnan, 2014–V–17, local catcher leg. [KIZ, 0100012–0100013]; 1 β **PARATYPES**, the same collecting data, [SJH]; 6 β , 1 ς **PARATYPES**, Luzhou, Sichuan, 2013–VII, local catcher leg. [SJH]; 2^O PARATYPES, Xiaoche He Park, Guiyang, Guizhou, 2013–VI– 11, S. J. Hu leg. [SJH]; 1, 1, 1, PARATYPES, Zuogong, E. Tibet, 2013–VI, local catcher leg. [SJH]; 2, 3, 3PARATYPES, Jigong Shan (1,940 m), Shimian, W. Sichuan, 2013–VI–13, Q. Zeng leg. [SFU]; 3 PARATYPES, Ziyun Xiang, Baoxing, W. Sichuan, 2012–VIII, local catcher leg. [SJH]; 13 PARATYPE, Panzhihua, W. Sichuan, 2001–VIII–5, E. T. Ye leg. [SJH]; 13 PARATYPE, Huize, N.E. Yunnan, 1998–IV, local catcher leg. [SFU]; 13 PARATYPE, Zhongdian, N.W. Yunnan, 1992–VI–20, D. Z. Dong leg. [KIZ]; 1 d PARATYPE, Qianjiang (1,780), Sichuan, 1989–VII– 14, S. Lin leg. [KIZ]; 4 PARATYPES, Lichuan (800 m), Hubei, 1988–VIII–3, X. C. Liang, D. Z. Dong, and S. Lin leg. [KIZ]; 1♀ **PARATYPE**, Leigong Shan, Guizhou, 1988–VIII–15, S. Lin leg. [KIZ]; 3♂ **PARATYPES**, Emei Shan (1,400 m), W. Sichuan, 1984–IV–13, local catcher leg. [DB]; 1 d PARATYPE, ditto, 1982–IV–13, local catcher leg. [DB]; 1 d PARATYPE, Xi Shan, Kunming, C. Yunnan, 1979–VIII–9, J. Xiong leg. [KIZ]; 1∂ PARATYPE, Jinghong, Xishuangbanna, S. Yunnan, 1979–VII–28, D. Z. Dong leg. [KIZ]; 1 PARATYPE, Kouy-Tchéou Abbe Largeteau [leg.], Ex Oberthür Coll., Brit. Mus. 1927-3, Para-lecto-type [round pale blue edged label], PARALECTOTYPE [in blue] Papilo [sic] glycerion var. Mandarinus Oberthür C.R. Smith det. 2003, BMNH(E) #146242 [BMNH]; 13 PARATYPE, Kouy-Tchéou Abbe Largeteau [leg.], China Kouytcheou [hand written], Pap. glycerion, var. Mandarinus, Oberthür Etud. d' Entomol. IV^e livraison - appendice variété géographique chinoise [hand written], Levick Bequest B.M.1941-83, Type [round red edged label], Para-lecto-type [round pale blue edged label], PARALECTOTYPE [in blue] Papilo [sic] glycerion var. Mandarinus Oberthür C.R. Smith det. 2003, BMNH(E) #149385 [BMNH]; 13 PARATYPE, Siao-Lou, 1893, Chasseurs indigènes, Rothschild Bequest, BMNH(E) # 220118, [BMNH]; 18 PARATYPE, Siao-Lou, 1901, Chasseurs indigènes du P. Déjean, Ex Oberthür Coll., BMNH(E) # 146048, [BMNH]; 1 PARATYPE, Siao-Lou, 1893, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146063, [BMNH]; 1∂ PARATYPE, Siao-Lou, 1893, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146065, [BMNH]; 13 PARATYPE, Siao-Lou, 1900, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146110, [BMNH]; 13 PARATYPE, Siao-Lou, 1900, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146101, [BMNH]; 1 PARATYPE, Siao-Lou, 1900, Chasseurs indigènes, Ex Oberthür Coll., BMNH(E) # 146106, [BMNH]; 1³ **PARATYPE**, Chasseurs indigènes de Ta-tsien-lou. Récolte de 1910, Ex Oberthür Coll., BMNH(E) # 145814, [BMNH]; 13 PARATYPE, Chasseurs indigènes de Ta-tsien-lou. Récolte de 1910, Ex Oberthür Coll., BMNH(E) # 145820, [BMNH]; 13 PARATYPE, Chasseurs indigènes de Ta-tsien-lou. Récolte de 1910, Ex Oberthür Coll., BMNH(E) # 145821, [BMNH]; 1 PARATYPE, Chasseurs indigènes de Ta-tsien-lou. Récolte de 1910, Ex Oberthür Coll., BMNH(E) # 145822, [BMNH]; 1♀ **PARATYPE**, Ichang, Rothschild Bequest, BMNH(E) # 220120, [BMNH]; 1 PARATYPE, Jinfushan, Exp. Stotzner leg. [ZFMK]; 1 PARATYPE, Siao-Lou, 1893, Chasseurs indigènes [ZFMK]; 1♂ PARATYPE, Rou Bi Gou, Baoxing, Sichuan, 2006–VII–10, Ming Yue leg. [AMC]; 2♂, 1♀ PARATYPES, Ya'an, Sichuan, 2006–VII, local catcher leg. [AMC]; 22 PARATYPES, Wu Shi He, Hanyuan County, Sichuan, 2005–VII–5-20, local catcher leg. [AMC]; 1♂, 1♀ PARATYPES, Daba Shan, E Sichuan, 2006–VII, local catcher leg. [AMC]; 1♂ PARATYPE, Gioncheng, Guangxi, 1996–VI–17, local catcher leg. [AMC]; 4♂, 1♀ PARATYPES, Luzhou, Sichuan, 2007–V, local catcher leg. [AMC]; 2d PARATYPES, Kangding, Sichuan, 2006–VII,

local catcher leg. [AMC]; 1⁽³⁾ **PARATYPE**, Qingshuilang Shan, Yunlong, W. Yunnan, 1998–VII-VIII, Qin et al leg., ex Coll. Jan Moonen [NBC]; 1⁽³⁾ **PARATYPE**, Jinxiu, Dayao Shan, Guangxi, 1998–VII-VIII, Liang et al leg., ex Coll. Jan Moonen [NBC].

VIETNAM: 8³, 10⁹ PARATYPES, Sapa (1,500 m), Lao Cai, 2015–V, L. T. Le leg. [SJH]; 2³ PARATYPES, Ha Giang, 2008–VII, local catcher leg. [SJH]; 6³, 8⁹ PARATYPES, Ha Giang, 2006–V, local catcher leg. [AMC]; 2³ PARATYPES, Ha Giang, 2007–VII, local catcher leg. [AMC]; 1³ PARATYPE, Ha Giang, 2008–VII, local catcher leg. [AMC].

Graphium (Pazala) mandarinus kimurai Murayama, 1982

LAOS: 3⁽⁷⁾, Nam Cha (1250 m), ~5 km N of Xaysomboun, 2006–II–28, Hiroyuki Wakahara leg. [AMC].

- MYANMAR: 1Å, Mt. Kennedy, N. Chin State, 2000–III–21, local catcher leg. [AMC]; 1Å, ditto, 2000–V–5, local catcher leg. [AMC]; 5Å, Mt. Ashaemyinanaukmyin (2,363 m), W. Shan State, 2008–III–18-22, local catcher leg. [SJH]; 11Å, 2¢, ditto, 2008–III–15-29, local catcher leg. [AMC]; 1Å, Mong Pat, N. Shan State, 2000–III–20, local catcher leg. [AMC].
- THAILAND: 1⁽³⁾, Doi Internon [sic], North Thailand, 1985–II–10, K. Okano leg., BMNH(E) # 220231 [BMNH]; 2⁽³⁾, Doi Inthanon (2200 m), Chiang Mai Province, 1976–XII–4-5, Bro. Amnuay Pinratana leg. [AMC]; 5⁽³⁾, Observatory entry road (2,446 m), Doi Inthanon, Chiang Mai Province, 2012–XII–14, local catcher leg. [SJH]; 5⁽³⁾, ditto, 2012–XII–14-15, local catcher leg. [AMC].

Graphium (Pazala) mandarinus fangana (K. Okano), 1986

THAILAND: 2³, Doi Pu Meun, Chiang Mai, 2007–II, local catcher leg. [SJH]; 1³1², ditto, 2016–III, local catcher leg. [SJH]; 1³, ditto, 1989–II, local catcher leg. [AMC]; 3³, ditto, 1990–II, local catcher leg. [AMC]; 4³, ditto, 1992–III, Prasobsuk Sukkit leg. [AMC]; 3³, ditto, 1988–II–10-14, local catcher leg. [AMC]; 5³, ditto, 2002–II, local catcher leg. [AMC]; 3³, ditto, 2007–II, local catcher leg. [AMC]; 4³, ditto, 2013–II–26, Prasobsuk Sukkit leg. [AMC]; 1³, Chiang Mai, 1989–II, local catcher leg., ex Coll. Jan Moonen [NBC].

Graphium (Pazala) mandarinus stilwelli Cotton & Hu ssp. nov.

- CHINA: 1♂ HOLOTYPE, Tacheng (1,900 m), Weixi, W. Yunnan, 2015–IV–29, S. J. Hu leg. [KIZ, 0100008]; 1♂
 PARATYPE, Gazhi River (1,050 m), Yunxian, Lincang, S.W. Yunnan, 2016–V–9, Z. C. Xiong leg. [KIZ, 0100009]; 2♂
 PARATYPES, Gaoligong Shan, Tengchong, W. Yunnan, 2015–V–20, P. Y. F. Lo leg. [SJH]; 1♂
 PARATYPE, Liuku (100 m), W. Yunnan, 2002–IV–20, unknown leg. [SFU]; 1♂
 PARATYPE, Zhongdian, N.W. Yunnan, 1992–VI–20, D. Z. Dong leg. [KIZ]; 1♂
 PARATYPE, Ailao Shan, Jingdong, W. Yunnan, 1992–V–29, D. Z. Dong leg. [KIZ]; 1♂
 PARATYPE, Xujia Ba, Jingdong, W. Yunnan, 1984–III–16, unknown leg. [KIZ]; 6♂
 PARATYPES, west of Fugong, NW. Yunnan, 2009–IV–2-29, local catcher leg. [AMC]; 3♂
 PARATYPES, Pingpo, Dali, Yunnan, 2016–IV–25-V–20, local catcher leg. [AMC]; 1
 PARATYPE, Qingshuilang Shan, Yunlong, W. Yunnan, 1998–VII-VIII, Qin et al leg., ex Coll. Jan Moonen, [NBC]; 1♀
- MYANMAR: 3 PARATYPES, Chudu Razi Hills, Kachin State, 2008–IV–21, local catcher leg. [SJH]; 1 PARATYPE, ditto, 2008–IV–23, local catcher leg. [SJH]; 1 PARATYPE, ditto, 2008–IV–28, local catcher leg. [SJH]; 1 PARATYPE, ditto, 2008–IV–28, local catcher leg. [SJH]; 1 PARATYPE, ditto, 2008–IV–28, local catcher leg. [SJH]; 2 PARATYPES, ditto, 2008–IV–10-V–5, local catcher leg. [AMC]; 4 PARATYPES, ditto, 2009–IV–30, local catcher leg. [AMC]; 8 PARATYPES, ditto, 2007–V–5-9, local catcher leg. [AMC]; 5 PARATYPES, ditto, 2006–V–9, local catcher leg. [AMC]; 2 PARATYPES, ditto, 2006–VI–7, local catcher leg. [AMC]; 8 PARATYPES, Kushin, E. of Renam, N. Kachin State, 1998–IV–25-V–4, local catcher leg. [AMC]; 1 m# PARATYPE, ditto, 2012–V–15, local catcher leg. [AMC]; 1 m# PARATYPE, ditto, 2012–V–26, local catcher leg. [AMC]; 5 PARATYPES, ditto, Kushin, E. of Renam, N. Kachin State, 1998–IV–19-27, local catcher leg. [PS].

Graphium (Pazala) mandarinus mandarinus (Oberthür, 1879)

CHINA: 1♂, LECTOTYPE, Moupin [=Baoxing, Sichuan], A. David leg., ex Oberthür Coll., BMNH(E) # 146241 [BMNH];
7♂, Ziyun, Baoxing, W. Sichuan, 2012–VIII, local catcher leg. [SJH]; 2♂, Lokshan [= Leshan], W. Sichuan, 1991–VI–15, local catcher leg. [DB]; 1♂, Rou Bi Gou, Baoxing, Sichuan, 2006–V–19, Ming Yue leg. [AMC]; 1♂, ditto, 2006–VII–16, Ming Yue leg. [AMC]; 6♂, ditto, 2006–VIII–10, Ming Yue leg. [AMC]; 23♂, Lu Shan, 800-1500 m, Sichuan, 2006–IV–15-20, local catcher leg. [AMC]; 2♂, Emei Shan, Sichuan, 1996–VI, local catcher leg. [AMC]; 1♂, Ya'an, Sichuan, 2007–IV, local catcher leg. [AMC]; 2♂, Songbai (S. of Fang Xian), 1800 m, Shennongjia, W. Hubei, 1998–VI–1-18, Tang leg., ex Coll. Jan Moonen [NBC]; 1♂, Fongtongzhai, Baoxing, Sichuan, 1998–VII–1-15, Wan et al. leg., ex Coll. Jan Moonen [NBC].