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Research article

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Taxonomic revision of Australian *Amobia* Robineau-Desvoidy, 1830 (Sarcophagidae: Miltogramminae): integrating morphology and genetics finds a new species and tackles old problems

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Abstract. A taxonomic revision of the Australian species of *Amobia* Robineau-Desvoidy, 1830 (Diptera: Sarcophagidae: Miltogramminae) is completed using an integrated approach combining four molecular loci (three mitochondrial, COI, ND4 and CYTB; one nuclear, EF1 α) and morphological data. A new species, *Amobia* (s. str.) *serpenta* sp. nov., endemic to Australia, is described, and *Amobia auriceps* (Baranov, 1935) and *Amobia burnsi* (Malloch, 1930) are re-described. Molecular data are used to reconstruct inter-specific and generic relationships and support morphological species hypotheses. Phylogenetic analysis places all three Australian *Amobia* species together with *Amobia signata* (Meigen, 1824) (a Palaearctic species) in a single clade sister to *Senotainia* Macquart, 1846 (in part), which is in agreement with previous phylogenetic studies of the Miltogramminae. In addition to the description of species and molecular phylogenetics, general host associations for the Australian species of *Amobia* are discussed and evidence for the synonymisation of *A. pelopei* (Rondani, 1859) and *A. auriceps* is refuted.

Keywords. Phylogenetics, DNA barcoding, flesh flies, Miltogramminae.

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Introduction

The Miltogramminae (Diptera: Sarcophagidae), accounting for ~20% of global flesh fly diversity, are characterised by their mostly parasitic life histories. Unlike the many flesh flies that breed in carrion, most members of this subfamily are kleptoparasites in the nests of solitary wasps and bees (Barth 1908; McCorquodale 1986; Spofford *et al.* 1989; Spofford & Kurczewski 1990; Polidori *et al.* 2006). Seven genera are known to occur across the Australasian region (Pape 1996): *Aenigmetopia* Malloch, 1930 (five spp.; Johnston *et al.* 2020b), *Amobia* Robineau-Desvoidy, 1830 (two spp.), *Metopia* Meigen, 1803 (two spp.) *Miltogramma* Meigen, 1803 (four spp.), *Protomiltogramma* Townsend, 1916 (six spp.), *Senotainia* Macquart, 1846 (one sp.), and, most recently, *Macronychia* Rondani, 1859 (one sp.; Johnston *et al.* 2020a).

In an attempt to better understand the diversity of Australian sarcophagids, there has been a recent targeted effort to systematically revise the Australian Miltogramminae and place their constituent genera and species within a global phylogenetic context. This research has been targeted at specific genera, and revisions of *Aenigmetopia* and *Macronychia* have already been published (Johnston *et al.* 2020a, 2020b). This paper focuses on the genus *Amobia*, within the larger aim of a full treatment of the Australian Miltogramminae.

The genus *Amobia* is characterised morphologically by 1) the autapomorphic configuration of the anterior margin of the anepimeron, which shows a distinct swelling below the ampulla (Shewell 1987; Pape 1996), and 2) the reduction of the epiphallus, which most likely is apomorphic at the genus level (Pape 1996; Buenaventura *et al.* 2020). Additional diagnostic features, which are also found in other miltogrammine genera, are the relatively high number (> 5) of fine, hair-like proclinate fronto-orbital setae; the antennal insertion at or below the midpoint of the eye, and the claws longer than the fifth tarsomere in both sexes (Pape 1996). Similarly, species of *Amobia* can be characterised by the hosts of their inquilinous larvae, which are predominantly mud daubers and potter wasps (Hymenoptera: Sphecidae, Crabronidae; Vespidae: Eumeninae) (Itino 1988; Downing 1996; Coville 2000; Fateryga & Kovblyuk 2014).

The world fauna of *Amobia* consisted of 13 species prior to this study, with the highest species diversity in the Nearctic region (six spp.) and modest numbers known from the Afrotropical (two spp.), Australasian (two spp.), Neotropical (three spp.), Oriental (four spp.) and Palaearctic (four spp.) regions. *Amobia* comprises two subgenera: 1) *Sarcomacronychia* Townsend, 1892 restricted to the Nearctic and Neotropical regions and characterised by red-coloured terminalia, the male abdominal tergite 6 setose and free from syntergosternite 7+8, fewer than 10 proclinate fronto-orbital setae and posterior spiracles of the puparium not recessed*, and 2) the widespread *Amobia* (s. str.), characterised by brown terminalia, male abdominal tergite 6 bare* and fused to syntergosternite 7+8*, 10 or more proclinate fronto-orbital setae* and the posterior spiracles of the puparium recessed ('*' = possible synapomorphies; Zumpt 1961; Draber-Mońko 1966; Pape 1996; Zhang *et al.* 2011).

The evolutionary relationships between *Amobia* and the other miltogrammine genera have been hypothesised based on morphology (Rohdendorf 1967; Verves 1989) and more recently also molecular data (Piwczyński *et al.* 2017; Buenaventura *et al.* 2020). The molecular phylogenies of Piwczyński *et al.* (2017) and Buenaventura *et al.* (2020) are generally congruent and placed *Amobia* as sister to a part of *Senotainia* in a clade also containing *Macronychia** and *Oebalia* Robineau-Desvoidy, 1863, thereby

rendering *Senotainia* paraphyletic, with *Amobia signata* (Meigen, 1824) splitting *Senotainia tricuspis* (Meigen, 1838)* and *Senotainia albifrons* (Rondani, 1859) from *Senotainia conica* (Fallén, 1810). ('*' = included in the phylogeny of Piwczyński *et al.* 2017 only.) The early morphology-based evolutionary hypotheses favoured a tribal classification of the Miltogramminae, with *Amobia* first placed within the tribe Senotainiini along with *Aenigmetopia, Chorezmomyia* Rohdendorf, 1935, *Eumacronychia* Townsend, 1892, *Senotainia* and *Taxigramma* Perris, 1852, and later moved into its own monotypic tribe Amobiini (Verves 1989). A tribal classification has since been abandoned (Pape 1996; Piwczyński *et al.* 2017; Buenaventura *et al.* 2020).

The Australian fauna of *Amobia* has so far received little attention. Until now, the genus was represented in Australia by two species: *Amobia auriceps* (Baranov, 1935) (which is also distributed throughout the broader Australasian/Oceanian and Oriental regions) and the endemic *Amobia burnsi* (Malloch, 1930). This paper adds to this body of knowledge through the taxonomic revision of Australian *Amobia*, describing a new endemic species, providing new characters for the separation of species, and providing extensive photographic documentation of the adult morphology and terminalia. Molecular data (four loci: COI, CYTB, ND4 and EF1 α) are utilized to support morphological species hypotheses, allow further insight into the taxonomic distinctiveness of this genus and, through phylogenetics, place the Australian *Amobia* within the framework provided by Piwczyński *et al.* (2017).

Materials and methods

Morphology

Specimens were collected during several research expeditions across Australia during the summer months of 2016–2018. All specimens were collected individually using a hand net, killed in ethyl acetate vapour and either preserved in ethanol the same day or pinned for taxonomic identification. Photo-documentation of the adult morphology and male terminalia of *A. auriceps* and *A. burnsi* was completed using a BK lab system (Visionary Digital, VA, USA), with photographic stacking performed with Zerene Stacker (Zerene Systems, LLC, Richland, WA, USA). Photographs of dissected male terminalia were taken from preparations in glycerol. Photo-documentation of *A. serpenta* sp. nov. was completed using a Leica DFC295 digital camera (3M pixel) mounted on a Leica MZ16a stereo-microscope and analysed using Leica Application Suite ver. 3.8.0 (Leica Microsystems Limited, Germany).

For preparation of SEM images, male terminalia in glycerol were rinsed in demineralised water, dehydrated in 96% ethanol, air-dried, mounted on adhesive electrical tape attached to aluminium stubs, coated with ~100 nm of platinum/palladium and studied using a JEOL JSM-6335 F scanning electron microscope (JEOL, Tokyo, Japan). Due to the availability of only two male specimens of *A. serpenta* sp. nov., it was decided to avoid dissection of this species and to document and describe the relevant structures based on extended terminalia of the holotype.

For documentation of adult morphology, specimens were studied under a Leica MZ75 microscope (Leica Microsystems Limited) in conjunction with the photographs and SEM images. Terminology for morphological characters follows Cumming & Wood (2017), with additional characters relevant to Miltogramminae obtained from Pape (1996). For label data, Australian states and territories are denoted by standard abbreviations (ACT: Australian Capital Territory, NSW: New South Wales, NT: Northern Territory, QLD: Queensland, SA: South Australia, WA: Western Australia), additional information provided verbatim in square brackets after the label data, and repository by institutional abbreviation in parentheses.

Specimen repository abbreviations

AM	=	Australian Museum, Sydney, Australia	
ANIC	=	Australian National Insect Collection, Canberra, Australia	
BM	=	Bishop Museum, Hawaii, USA	
NMNH	=	National Museum of Natural History, Washington, D.C., USA	
QM	=	Queensland Museum, Brisbane, Australia	

DNA extraction, amplification and sequencing

Molecular analysis was employed to support the morphological species hypotheses and phylogenetically place the Australian *Amobia* among the global Miltogramminae. DNA extraction, amplification and sequencing methods followed Meiklejohn *et al.* (2011) and Piwczyński *et al.* (2017). In short: the total genomic DNA was extracted from a single leg of four specimens of *A. auriceps*, two of *A. burnsi* and a single specimen of *A. serpenta* sp. nov. using the salting-out method described by Aljanabi & Martinez (1997) (see Supplementary file 1 for a full list of taxa and specimens used in the molecular analysis, including accession numbers for genetic data originally sequenced by Piwczyński *et al.* (2017) for their global miltogrammine phylogeny, COI, CYTB, ND4 and EF1a, were also amplified for our *Amobia* specimens. Temperature cycling conditions for amplification followed the general structure of one cycle at 94°C for 2 min; 35 cycles at 95°C for 30 s; 45–65°C gradient for 15 s; 72°C for 30 s; and one cycle at 72°C for 5 min. Annealing temperature optimisation for gene fragments was as follows: 50.6°C for COI, 51°C for CYTB, 45°C for ND4 and 53°C for EF1a. The extension time for EF1a was increased from 30 s to 45 s to account for the >1 kb length of the target sequence. Primer pairs used for gene amplification and their relevant sequences are available in Table 1.

Amplification success was confirmed by resolving PCR products through agarose gel electrophoresis. Prior to sequencing, 5 µl aliquots of samples were treated with 2 µl of 2x diluted ExoSAPIT Express[®] (Applied Biosystems, Foster City, CA, USA), following the manufacturer's instructions, in order to digest unincorporated dNTPs and primers. ExoSAPIT Express[®] treated products were then sequenced using ABIPRISM[®] BigDyeTM Terminator ver. 3.1 Sequencing Kit (Applied Biosystems, Foster City, CA, USA). The sequencing products were separated using an ABI 3130*xl* Genetic Analyser (Applied Biosystems, Foster City, CA, USA).

Sequence electropherograms were edited using ChromasPro version 1.33 (Technelysium, South Brisbane, QLD, Australia; available online at www.technelysium.com.au/wp/chromaspro) and confirmed to be of miltogrammine origin by cross-checking species identity by submission to the Basic Local Alignment Search Tool (BLAST; National Center for Biotechnology Information, Bethesda, MD, USA; available online at blast.ncbi.nlm.nih.gov).

Automatic barcode gap discovery for species delimitation

Prior to phylogenetic analysis, COI sequences for all extracted *Amobia* (seven in total: four *A. auriceps*, two *A. burnsi* and a single *A. serpenta* sp. nov) were aligned using MUSCLE (Edgar 2004; MUSCLE parameters: gap opening penalty = -400, gap extension penalty = 0, Clustering Method = UPGMB, Maximum diagonal length = 24) in Geneious Prime 2019.0.3 (https://www.geneious.com) and submitted to the Automatic Barcode Gap Discovery online analysis tool (ABGD; Puillandre *et al.* 2012, available from: https://bioinfo.mnhn.fr/abi/public/abgd/). Analysis was completed using following Puillandre *et al.* (2012) using the Kimura two parameters model (Kimura 1980; default parameters). The number of input sequences used was lower than recommended for ABGD (three–five sequences per species); the developers (Puillandre *et al.* 2012) note that having less than the recommended number of sequences can result in the inability to accurately detect the "barcode gap". As such, this study did not use the prior

Primer name	Direction	Target gene	Sequence 5'–3'
LCO1490-L	Forward	COI	5'-GGTCWACWAATCATAAAGATA TTG-3'
HCO2198-L	Reverse	COI	5'-RAAACTTCWGGRTGWCCAAAR AATCA-3'
CB-J-10933	Forward	CYTB	5'-TATGTTTTACCTTGAGGACAAA TATC-3'
TSI-N-11683	Reverse	CYTB	5'-AAATTCTATCTTATGTTTTCAA AAC-3'
N4-N-9194	Forward	ND4	5'-ATTTTTTGAAAGAAGTTTAAT TCC-3'
N4-J-8502	Reverse	ND4	5'-GTTGGAGGAGCTGCTATATTA G-3'
EF1_21F	Reverse	EF1a	5'-TGAGCGYGARCGTGGTATCAC-3'
rcM4	Reverse	EF1a	5'-ACAGCVACKGTYTGYCTCATRTC-3'

Table 1. Primer sequences used for DNA amplification.

intraspecific divergence (barcode gap) as the basis for species delimitation, but rather the K2P distance matrix generated by ABGD was used to calculate inter- and intraspecific distance between species. The distance between COI barcodes was then used to support morphological species hypotheses.

Alignment and phylogenetic analysis (Fig. 1)

Following ABGD analysis, a phylogeny was prepared to determine the position of Australian Amobia species among the global miltogrammines. Sequence data from the phylogeny of Piwczyński et al. (2017) for COI, CYTB, ND4 and EF1a were obtained from GenBank (taxa list: Supplementary file 2), concatenated, and aligned along with the new sequences of Amobia using the MUSCLE algorithm in Geneious Prime 2019.0.3 (https://www.geneious.com). Aligned sequences were then submitted for analysis in PartitionFinder2 (Lanfear et al. 2016), which simultaneously determines the best partitioning scheme and model. Partitioning by gene and codon position (a total of 12 partitions) produced the highest corrected Akaike Information Criterion (AICc) value and was used in Maximum Likelihood (ML) and Bayesian inference (BI) analyses. The model with the highest calculated AICc was GTR+I+G; this model was utilised for all partitions in the BI analysis. The creator of RAxML recommends not estimating invariant sites in RAxML; as such, the GTR+G model was used for all partitions in the ML analysis. ML analysis was performed in RAxML ver. 8.0 (Stamatakis 2014) (GTR+G model, 1000 bootstrap iterations; Miller et al. 2010) and BI analysis was performed in MrBayes ver. 3.2.6 utilising the BEAGLE library. Four simultaneous Bayesian analyses were completed, each utilising four Markov Chain Monte Carlo (MCMC) chains (three heated, one cold; temperature default = 0.1). Each chain was run for 20 million generations, sampling every 2000 generations (for a total of 40,000 samples). All priors for the analysis remained at their default settings. To confirm run success and effective sampling of the priors, MCMC convergence was confirmed by examining effective sample size (ESS) and potential scale reduction factors (PRSF) (which were confirmed to be greater than 100 and ~1, respectively). A consensus tree was then prepared from the resultant trees with the first 25% of all trees discarded as burn-in. Paritionfinder2, RAxML and MrBayes analyses were performed within the online CIPRES platform (Miller et al. 2010).

A final tree graphic summarising both analyses (ML and BI) was produced using FigTree (ver. 1.4.3; Rambaut 2010) and edited in Adobe Illustrator[®] (available from https://www.adobe.com/au/products/illustrator).

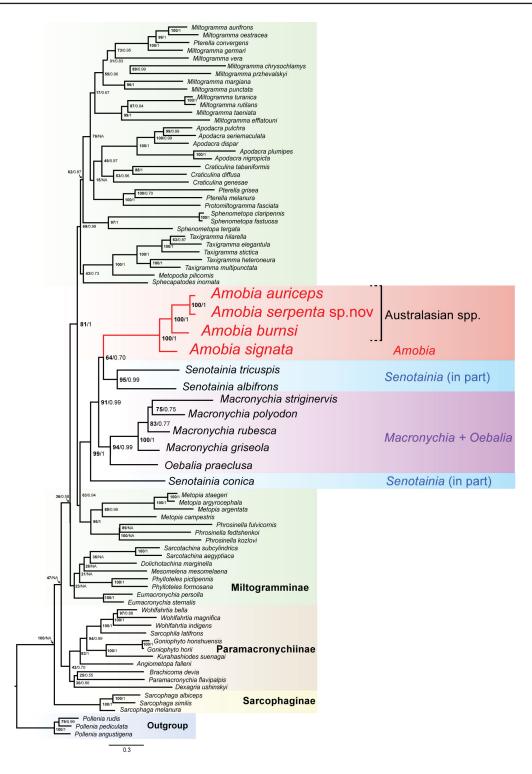


Fig. 1. Maximum likelihood phylogenetic tree indicating the placement of *Amobia* among the global Miltogramminae, inferred from COI, CYTB, ND4 and EF1 α sequence data. With the exception of the three newly sequenced Australian *Amobia* and *Macronychia rubesca*, data was acquired from Piwczyński *et al.* (2017). Node support values are shown for both maximum likelihood analysis (bootstrap support, 1000 iterations; in bold font) and Bayesian analysis (posterior probability, 40 million generations), 'NA' indicates nodes resolved that were not resolved by the Bayesian analysis. Branch length scale = 0.3 nucleotide substitutions per site (calculated by the GTR+G nucleotide substitution model).

Results

Taxonomy

Class Insecta Linnaeus, 1758 Order Diptera Linnaeus, 1758 Family Sarcophagidae Macquart, 1834 Subfamily Miltogramminae Lioy, 1864 Genus *Amobia* Robineau-Desvoidy, 1830

Amobia (s. str.) auriceps (Baranov, 1935) Figs 2–3

Pachyophthalmus auriceps Baranov, 1935: 558.

Diagnosis

The males of this species can be distinguished from the other two Australian species of *Amobia* by the combination of dull yellow microtomentum present on the fronto-orbital and parafacial plates, brown basicosta and bare proepisternum (the latter feature, however, is variable in the two other Australian species and therefore not by itself fully diagnostic).

Material examined

Lectotype (designated by Sabrosky & Crosskey 1970: 430) SRI LANKA • ♂; Colombo; 6 Feb. 1931; N. Baranov leg.; NMNM.

Other material

AUSTRALIA • 1 \Diamond ; QLD, Gordonvale; 1919; J.F. Illingworth leg.; [label indicates 'collected ex mudwasp']; ANIC • 2 \Diamond \Diamond ; QLD, Cairns; 1919; J.F. Illingworth leg.; [label indicates 'collected ex mudwasp']; BM); • 6 \Diamond \Diamond ; QLD, Cairns; 1919; J.F. Illingworth leg.; [label indicates 'collected ex mudwasp']; BM • 1 \Diamond ; QLD, Gordonvale; 1919; J.F. Illingworth leg.; [label indicates 'collected ex mudwasp']; BM • 1 \Diamond ; QLD, Gap Creek Reserve; 6 Dec. 2016; Johnston, Butterworth and Wallman leg.; ANIC • 1 \Diamond ; QLD, Bowling Green National Park; 9 Jan. 2017; Johnston, Pape and Wallman leg.; ANIC • 1 \Diamond ; NT, Track to Lost City, Litchfield National Park; 10 Nov. 2017; Johnston, Wallman and Szpila leg.; ANIC • 1 \Diamond ; QLD, 26 km W 'Fairview'; 24 Apr. 1989; G. Daniels and A. Daniels leg.; QM • 1 \Diamond ; QLD, Mt Abbott; 12 Apr. 1997; C.J. Burwell leg.; QM • 1 \Diamond ; QLD, Redlands, Hillard Creek; 20 Jan. 2009; QM Party leg.; [label indicates collected in 'scribbly gum open forest']; QM • 1 \Diamond ; QLD, Teddington Weir, Maryborough; 13 Nov. 1994; G. Daniels and A. Daniels leg.; [label indicates collected in 'vine forest margin']; QM.

Redescription

SIZE. 6.0-9.0 mm (n = 19).

Male

HEAD (Fig. 2B, D, G). With dull yellow microtomentum on fronto-orbital and parafacial plates; frontal stripe dark brown and concave; fronto-orbital plate setose with a single distinct row (at least 15 setae) and additional unordered fine fronto-orbital setulae; third aristomere tapered, fine and hair-like distally, black and twice as long as first flagellomere; pedicel and scape setose; first flagellomere black and 1.5 \times the length of pedicel; gena and postgena grey with black setae; genal groove present, dark brown; vibrissa located midway between tip of first flagellomere and mouth margin; numerous supra-vibrissal

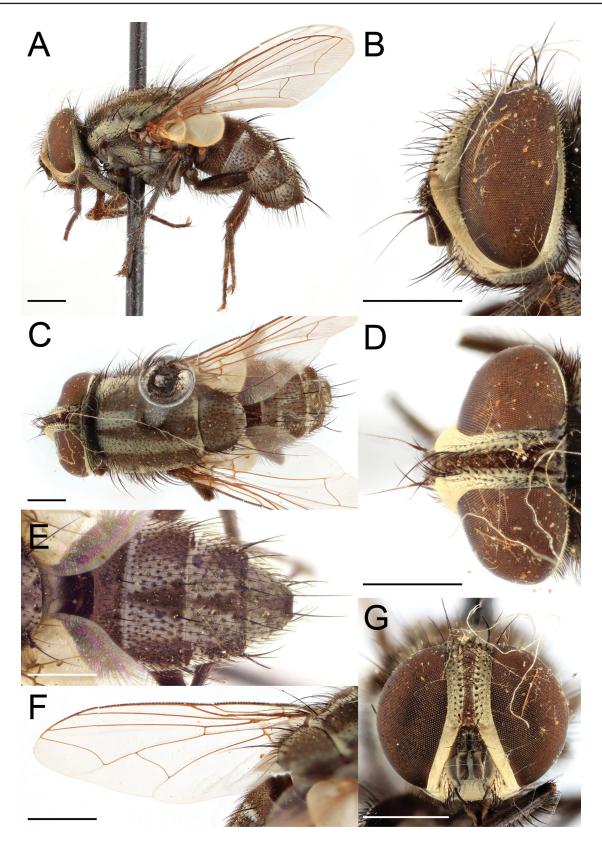


Fig. 2. *Amobia auriceps* (Baranov, 1935), \mathcal{E} , QLD, Cairns, 1919, J.F. Illingworth leg. (BM). **A**. Habitus, lateral view. **B**. Head, lateral view. **C**. Habitus, dorsal view. **D**. Head, dorsal view. **E**. Abdomen, dorsal view. **F**. Left wing dorsal view. **G**. Head anterior view. Scale bars: 1 mm.

setae; vibrissal setae and supra-vibrissal setae present in two rows, inner row strong and ordered, outer row weak and unordered.

THORAX (Fig. 2A, C). Grey with dull yellow-brown microtomentum (visible only at certain angles); one major median and two major lateral stripes; major stripes ending at anterior margin of scutellum; scutellum with some darkening on lateral margin; two notopleural setae with additional finer setae on entire surface; proepisternum bare; katepisternal setae 1+1.

WING (Fig. 2F). Hyaline; vein R_1 bare; vein R_{4+5} with several setae at base; cell r_{4+5} open; tegula black; basicosta, brown.

LEGS (Fig. 2A).Black; claws of fore tarsus slightly longer than tarsomere 5; mid tibia with one anterodorsal and one ventral seta.

ABDOMEN (Fig. 2A, C, E). T1+2 dark brown; setose and with two distinct median marginal setae; T3 setose, with two distinct median marginal setae and additional finer setae along entire posterior margin; T4–T5: setose with marginal setae on entire posterior margin; black ground colour most clearly visible towards the posterior margin; dull grey microtomentum present (occupying the majority of each tergite); three distinct black spots (two lateral and one median) on T3 and T4; T5 with only median black spot.

MALE TERMINALIA (Fig. 3A–B). Cercus sickle-shaped in lateral view and narrowest in middle portion (middle portion ~20% of cercus length); setose dorsally; cerci straight in posterior view, separated by a small gap between tips of cercal prongs; surstylus straight in lateral view, shorter than cercus; curved medially towards cercus in posterior view; phallus blunt with small spines on distal lobe.

Female

As male except microtomentum generally paler, especially on fronto-orbital plate where it appears silvery grey. The authors have chosen not to redescribe the female genitalia.

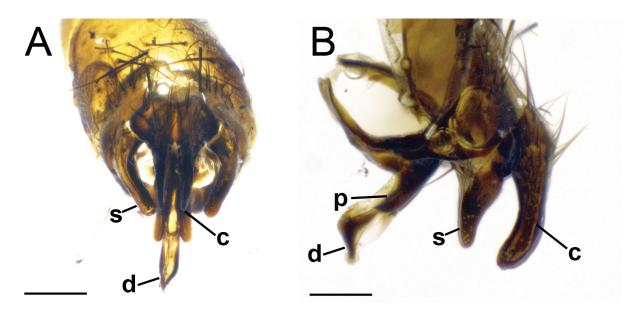


Fig. 3. *Amobia auriceps* (Baranov, 1935), male terminalia, QLD, Cairns, 1919, J.F. Illingworth leg. (BM). A. Epandrium, cerci, surstyli and phallus, posterior view. **B**. Epandrium, cerci, surstyli, phallus and pre-gonite, lateral view. Abbreviations: c = cercus; d = distal lobe of phallus; <math>s = surstylus; p = pre-gonite. Scale bars: 100 µm.

Biology

Label data confirm an association with 'mudwasps', probably family Sphecidae.

Distribution

Australasia/Oceania – Australia (NT, QLD), Hawaiian Is., Papua New Guinea. Oriental – India, Laos, Malaysia, Philippines, Sri Lanka, Taiwan.

Remarks

There has been some contention as to the validity of *A. auriceps*. Kurahashi (1974) considered *A. auriceps* to be conspecific with *Amobia pelopei* (Rondani, 1859), a species known only from the western Palaearctic region. He did, however, note differences in the shape of the cerci and surstyli, as well as in the colouration of the basicosta. Verves (1979) and Pape (1996), by comparison, considered *A. pelopei* and *A. auriceps* as distinct species, separated through the morphology of the terminalia and by the colouration of the fronto-orbital plates. This study is in agreement with Verves (1979) and Pape (1996). It should be noted that perhaps the most reliable character for the separation of these species is the shape of the cercus in lateral view, which is gradually tapering in *A. pelopei* (Kurahashi, 1974: fig. 1) and distinctly narrowed in the middle in *A. auriceps* (Fig. 3B).

Austrometopia burnsi Malloch, 1930: 438.

Material examined

Holotype

AUSTRALIA • ♂; QLD, Meringa; 28 Jun. 1926; A.N. Burns leg.; [label indicates 'Parasitic on Eumenid']; ANIC.

Other material

AUSTRALIA • 2 33, 2 99; NT, 15 km N Katherine; 16 May 2005, R.W. Matthews leg.; [label indicates 'Reared: ex. Nest of *Delta latreillei* (Saussure)']; ANIC • 1 9; QLD, Lockhart R; Rev. Hint leg.; ANIC • 1 3; NSW, North Head, Sydney Harbour; 14 Feb. 1978; G. Daniels leg.; AM • 19; [neither locality, date nor collector; label indicates 'parasitic on *Sceliphron laetum*']; QM.

Diagnosis

Males of this species can be distinguished from the other two Australian species of *Amobia* by the combination of distinct bright golden microtomentum on the fronto-orbital as well as parafacial plates and yellow-brown basicosta.

Redescription

SIZE. 8.0–10.0 mm (n = 7).

Male

HEAD (Fig. 4B, D, G). With golden microtomentum on fronto-orbital and parafacial plates; frontal stripe dark brown and concave; fronto-orbital plate setose with two distinct rows of setae, one row of frontal setae (> 10 setae) and one row of proclinate fronto-orbital setae (> 10 setae) and additional fine fronto-orbital setulae, setal rows ending before lunule; third aristomere broad at base and tapering to fine hair-like tip distally, black and twice as long as first flagellomere; pedicel and scape setose; first flagellomere

Amobia (s. str.) burnsi (Malloch, 1930) Figs 4–5

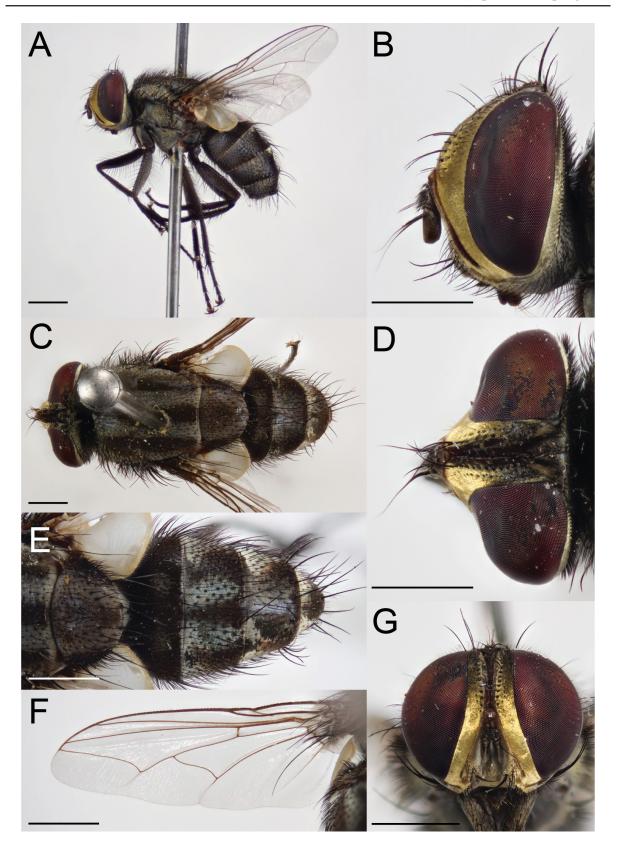


Fig. 4. *Amobia burnsi* (Malloch, 1930), ♂, NSW, North Head, Sydney Harbour, 14 Feb. 1978, G. Daniels leg. (AM) **A**. Habitus, lateral view. **B**. Head, lateral view. **C**. Habitus, dorsal view. **D**. Head, dorsal view. **E**. Abdomen, dorsal view. **F**. Left wing, dorsal view. **G**. Head, anterior view. Scale bars: 1 mm.

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black and slightly longer than pedicel; parafacial plate with fine setae near facial ridge in multiple disordered rows; gena and postgena grey with black setae; genal groove present, dark brown; vibrissa located midway between tip of first flagellomere and mouth margin; numerous supra-vibrissal setae present in two rows, inner row strong and ordered, outer row weak and unordered.

THORAX (Fig. 4A, C). Grey with some dull yellow microtomentum anteriorly; one major median, two major lateral and two faint minor lateral stripes (located mediad to lateral major stripes and approximately one quarter of their width); major stripes ending at anterior margin of scutellum; minor stripes ending just anterior to suture; scutellum with some darkening on lateral margins; two notopleural setae, with numerous additional fine setae; proepisternum setose; katepisternal setae 1+1.

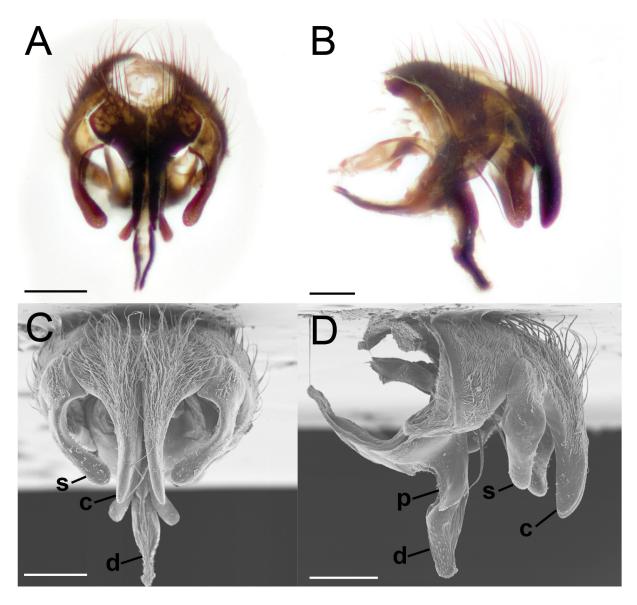


Fig. 5. *Amobia burnsi* (Malloch, 1930), male terminalia, NT, 15 km N Katherine, 16 May 2005, R.W. Matthews leg (ANIC). **A**. Epandrium, cerci, surstyli and phallus, posterior view. **B**. Epandrium, cerci, surstyli, phallus and pre-gonite, lateral view. **C**. SEM image of epandrium, gonites, cerci and surstyli, posterior view. **D**. SEM image of epandrium, gonites, phallus, cerci and surstyli, lateral view. Abbreviations: c = cercus; d = distal lobe of phallus; <math>s = surstylus; p = pre-gonite. Scale bars: 100 µm.

WING (Fig. 4F). Hyaline; vein R_1 bare; vein R_{4+5} with several setae at base; cell r_{4+5} open; tegula black; basicosta light yellow brown.

LEGS (Fig. 4A). Black; claws of fore tarsus slightly longer than tarsomere 5; pulvilli slightly shorter than fore tarsal claws; mid tibia with one anteroventral and one anterodorsal seta.

ABDOMEN (Fig. 4A, C, E). T1+2 black; setose, with two distinct median marginal setae; T3–T5: setose, with two distinct median marginal setae and additional finer setae along entire posterior margin of each tergite; black ground colour most clearly visible towards posterior margin; dull grey microtomentum present (occupying almost the entirety of each tergite); three distinct black spots (two lateral and one median) on T3 and T4; T5 with only one median black spot.

MALE TERMINALIA (Fig. 5A–D). Cercus sickle-shaped in lateral view, broad at base and pointed apically; densely setose dorsally; cerci diverging in posterior view with a weak curve away from one another; surstylus straight in lateral view, shorter than the cerci; in posterior view, surstyli slightly curved medially towards cerci; phallus blunt with small spines on distal lobe, swollen apically.

Female

As male except with generally paler microtomentum, especially on head where it appears dull grey. The authors have chosen not to redescribe the female genitalia.

Biology

Label data indicate [klepto]parasitism of wasps of the families Vespidae (*Delta latreillei* (Saussure, 1852)) and Sphecidae (*Sceliphron laetum* (Smith, 1856)).

Distribution

Australasia – Australia (NSW, NT, QLD, WA).

Remarks

Amobia burnsi was described in the monotypic genus *Austrometopia* Malloch, 1930. This classification has since been rejected as it rendered the genus *Amobia* paraphyletic (Kurahashi 1970; Pape 1996). Pape (1996) specified that, in the genus *Amobia*, the fore-tarsal claws are longer than the fifth tarsomere in both sexes; however, this was not observed in females of *A. burnsi* examined in this study, and as such may not be a character of all *Amobia*.

It should be noted that proepisternal setosity has been observed to be variable in this species and as such cannot be used to diagnose it.

Amobia (s. str.) serpenta sp. nov. urn:lsid:zoobank.org:act:884490D3-8113-4F75-B4BD-E0E88209BC88 Figs 6–7

Diagnosis

Males of this species can be distinguished from the other two Australian members of *Amobia* by the unique dual head colouration (silver microtomentum on the fronto-orbital and parafacial plates, and golden microtomentum on upper mouth edge, gena and postgena) in combination with the area of golden microtomentum on the notopleuron.

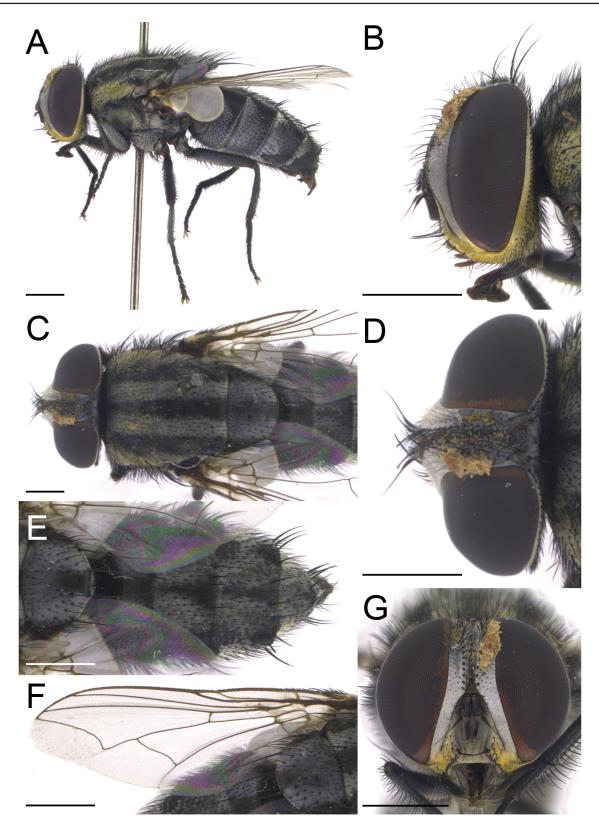


Fig. 6. *Amobia serpenta* sp. nov., holotype, ♂, NT, Serpentine Gorge, West MacDonnell National Park, 13 Nov. 2017, Johnston, Wallman and Szpila leg. (ANIC). **A**. Habitus, lateral view. **B**. Head, lateral view. **C**. Habitus, dorsal view. **D**. Head, dorsal view. **E**. Abdomen, dorsal view. **F**. Left wing, dorsal view. **G**. Head, anterior view. Scale bars: 1 mm.

Etymology

The species epithet '*serpenta*', which should be treated as a noun in apposition, refers to Serpentine Gorge, the type locality of this species.

Type material

Holotype

AUSTRALIA • ♂; NT, Serpentine Gorge, West MacDonnell National Park; 13 Nov. 2017; Johnston, Wallman & Szpila leg.; ANIC.

Paratypes

AUSTRALIA • 1 \Diamond ; WA, 2 miles S of Dempster; 19 Oct. 1970; D.H. Colless leg.; ANIC • 1 \Diamond ; NT, Alice Springs; 21–27 Sep. 1972; R.W. Matthews leg.; ANIC • 1 \Diamond ; QLD, Birdsville Track; Sep. 1936; A. Muncham leg.; [label indicates: 'reared from clay nest of mason wasp (Eumenidae)']; ANIC • 1 \Diamond ; WA, White Spring Ruins NE of Wittenoom; 16 Aug. 1970; P. Ferrar leg.; ANIC • 1 \Diamond ; SA, Nullarbor; 25 Oct. 1958; E.F. Riek leg.; ANIC • 1 \Diamond ; WA, Carson Escarpment; 15 Aug. 1975; I.F.B. Common and M.S. Upton leg.; ANIC.

Description

SIZE. 6-9 mm (n = 7).

Male

HEAD (Fig. 6B, D, G). Grey with silver microtomentum on fronto-orbital and parafacial plates, and golden microtomentum on upper mouth edge, gena and postgena; frontal stripe black and distinctly concave; fronto-orbital plate setose with one distinct row of fronto-orbital setae (>10 setae) and additional unordered fine fronto-orbital setulae; setal rows ending at height of lunule; third aristomere broad in basal third then tapering to a fine hair-like tip distally, black and $2 \times$ length of the first flagellomere; pedicel and scape setose; first flagellomere black and slightly longer than pedicel; parafacial plate with fine setae near facial ridge, in multiple disordered rows; gena yellow and postgena light grey with black

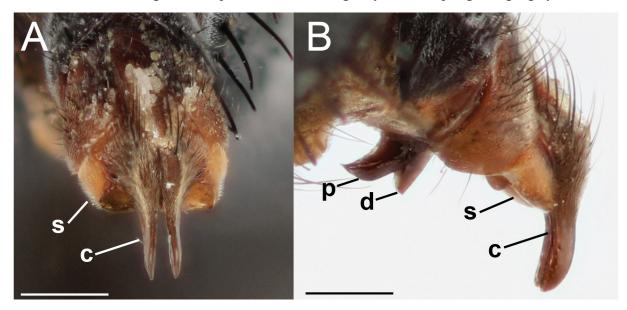


Fig. 7. *Amobia serpenta* sp. nov., holotype, male terminalia, NT, Serpentine Gorge, West MacDonnell National Park, 13 Nov. 2017, Johnston, Wallman and Szpila leg. (ANIC). **A**. Epandrium, cerci, surstyli, and phallus, posterior view. **B**. Epandrium, cerci, surstyli, phallus and pre-gonites, lateral view. Abbreviations: c = cerci; d = distal lobe (broken); s = surstylus; p = pre-gonite. Scale bars: 100 µm.

setae; genal groove present, dark brown; vibrissa located midway between tip of first flagellomere and mouth margin; numerous supra-vibrissal setae.

THORAX (Fig. 6A, C). Grey with golden microtomentum, particularly antero-dorsally and forming distinct patch on notopleuron; one major median, two major lateral and two faint minor lateral stripes (minor stripes placed medially to major stripes and approximately one quarter of their width); major stripes continuing full length of scutellum, minor stripes ending just anterior to suture; two notopleural setae, with numerous additional fine setae on entire notopleural surface; proepisternum bare or setose; katepisternal setae 1+1.

WING (Fig. 6F). Hyaline; vein R_1 bare; vein R_{4+5} with several setae at base; cell r_{4+5} open; tegula black; basicosta yellow white.

LEGS (Fig. 6A). Black; claws of fore tarsus slightly longer than tarsomere 5; pulvilli as long as fore tarsal claws; mid tibia with one anteroventral and one anterodorsal seta.

ABDOMEN (Fig. 6A, C, E). T1+2: black; setose and with two distinct median marginal setae; T3–T5: setose, each with one pair of distinct median marginal setae; T5 with additional marginal setae along entire posterior surface; black ground colour most clearly visible towards posterior margin; dull grey microtomentum present, occupying entirety of tergites; three distinct black stripes (two lateral and one median) on T3; T4 with median stripe and two lateral spots; T5 with only one median black spot.

MALE TERMINALIA (Fig. 7A–B). Cercus (lateral view) sickle-shaped broad at base and tapered; setose on dorsal surface; posteriorly, cerci slightly diverging apically; surstylus shorter than cercus, single lobed on posterior margin; posteriorly, surstyli curved medially.

Female

As male except for slightly darker microtomentum on the fronto-orbital plate, appearing dull yellow. Parafacial plate, gena and post gena remain golden and distinctly different to fronto-orbital plate. The authors have chosen not to describe the female genitalia.

Biology

The holotype of this species was collected on a hilltop, perching on an exposed rock. Paratype label data indicate an association with vespid wasps (Vespidae: Eumeninae).

Distribution

Australasia - Australia (NT, QLD, SA, WA).

Remarks

Amobia serpenta sp. nov. is placed within the subgenus *Amobia* s. str. due to the brown coloured terminalia and more than 10 proclinate fronto-orbital setae (Zumpt 1961; Pape 1996). Due to the limited number of male specimens (two), it was decided to avoid dissection of the male terminalia and rather to describe the extended male terminalia of the pinned holotype. For this reason, and as the phallus was broken off completely in the holotype, only the cerci and surstyli could be accurately described.

Key to males of Australian Amobia

- Fronto-orbital plate with bright golden microtomentum (Fig. 4A–B, G); basicosta yellow-brown; male terminalia: cerci in lateral view gradually tapering from base to tip (Fig. 5B, D); in posterior view cerci weakly curved away from one another (Fig. 5A, C)....A. (s. str.) burnsi (Malloch, 1930)

Phylogenetic analysis

Molecular analysis

BLAST searches of COI sequences from *A. auriceps*, *A. burnsi* and *A. serpenta* sp. nov. all showed the closest matches with a previously submitted sequence erroneously identified as a *Protomiltogramma* sp. (> 96% similarity; JN964698). Fortunately, the original voucher specimen could be retrieved from the University of Wollongong Diptera collection and re-identified as *A. auriceps*. The sequences of *Amobia* were also a good match with a previously identified *Amobia signata* originally sequenced for the phylogeny of Piwczyński *et al.* (2017) (> 92% similarity; KY749709).

Representative COI sequences from each species were analysed using the ABGD online tool. Preliminary analysis of COI data supported the morphological species hypothesis, despite the low number of input sequences. All three species were molecularly distinct, with mean inter- and intraspecific distance calculated as 4.00% and 0.24%, respectively (calculated using Kimura two parameters model; Kimura 1980). The greatest between-species distance was observed between *A. burnsi* and *A. auriceps* (mean interspecific distance = 4.85%, n = 8). The next greatest distance was between *A. auriceps* and *A. serpenta* sp. nov. (mean interspecific distance = 4.64%, n = 2). The smallest distance is ~10 × higher than the mean intraspecific distance for all *Amobia* species (0.24%) and 13 × higher than the mean intraspecific distance calculated for *A. burnsi* (0.15%).

Phylogenetic reconstruction

A Maximum Likelihood phylogeny was reconstructed based on the new sequences obtained from Australian *Amobia* (three concatenated sequences, one for each of the three species) and the sequence data of Piwczyński *et al.* (2017) (Fig. 1). The Bayesian inference phylogeny was generally concordant with the Maximum Likelihood phylogeny (Fig. 1; only support values shown).

The phylogeny recovered the genus *Amobia* as monophyletic. In terms of interspecific relationships, *A. auriceps* emerged as sister to *A. serpenta* sp. nov. (ML: bootstrap support = 100%, BI: posterior probability = 1) and *A. auriceps* + *A. serpenta* sp. nov. were sister to *A. burnsi* (ML: bootstrap support = 100%, BI: posterior probability = 1). *Amobia signata* was sister to all other *Amobia* species (ML: bootstrap support = 100%, BI: posterior probability = 1).

The intergeneric relationships presented in both the ML and BI phylogenetic hypotheses agreed with Piwczyński *et al.* (2017), with all *Amobia* resolved as a single clade sister to *Senotainia* (in part: *Senotainia albifrons* + *Senotainia tricuspis*) (ML: 64% bootstrap support, BI: posterior probability = 0.70) and with *Amobia* + *Senotainia* (in part) sister to *Macronychia* + *Oebalia* (ML: 91% bootstrap support, BI: posterior probability = 0.99). The *Amobia, Senotainia, Macronychia* and *Oebalia* cluster was sister to *Metopia, Phrosinella* Robineau-Desvoidy, 1863 and the rest of the higher miltogrammines (Piwczyński *et al.* 2017; ML: 63% bootstrap support, BI: posterior probability = 0.94).

Discussion

Species diagnosis

This study is the first to attempt to integrate morphological and molecular data to establish a strongly supported species hypothesis for the Australian *Amobia*. Morphology supported a three-species hypothesis, with the description of a new species, *A. serpenta* sp. nov., and the re-description of both *A. auriceps* and *A. burnsi*. The most diagnostic characters for the Australasian species of *Amobia* were colouration of the fronto-orbital plate and colouration of the basicosta, characters also known to be diagnostic for species of this genus in the Oriental, Nearctic and Palaearctic regions (Kurahashi 1974; Pape 1996). Unlike other regions, the presence and colouration of the microtomentum on the notopleuron was also found to be diagnostic in Australian *Amobia*.

DNA barcoding and molecular phylogenetics

Despite the low number of input sequences, preliminary ABGD analysis provided evidence supporting all three Australian species of *Amobia* identified by morphology. Between-species genetic distance was $\sim 17 \times$ higher than within-species genetic distance (4.00% compared to 0.23%). The most closely related COI barcodes were observed between *A. burnsi* and *A. serpenta* sp. nov. (2.08% interspecific distance) however this was still $\sim 10 \times$ higher than the mean intraspecific distance calculated for all species.

Both the Maximum Likelihood and Bayesian phylogenetic analyses clustered all three Australian species of *Amobia* together with *A. signata*. Within this *Amobia* clade *A. serpenta* sp. nov. + *A. auriceps* is sister to *A. burnsi*. Other than providing insight into the taxonomy and phylogeny of the Australian fauna of *Amobia*, our results agree with the intergeneric relationships presented by Piwczyński *et al.* (2017); *Amobia* is sister to *Senotainia* (in part; *S. tricuspis* and *S. albifrons*) then *Amobia* + *Senotainia* (in part) is sister to *Macronychia*. Thus, the addition of additional *Amobia* species in this analysis provides additional support to the previous phylogenetic hypotheses for the subfamily Miltogramminae.

Rejection of synonymisation of A. auriceps (Baranov, 1935) and A. pelopei (Rondani, 1859)

This revision supports the separation of *A. auriceps* and *A. pelopei* as distinct species. Verves (1979) and Pape (1996) separated *A. pelopei* and *A. auriceps* through the colouration of the fronto-orbital plates (*A. pelopei* with silver microtomentum, *A. auriceps* with yellow gold microtomentum). Verves (1979) additionally suggested that the terminalia of *A. auriceps* could be distinguished from those of *A. pelopei* by the terminally angular cerci and an apically dilated phallus. We follow Verves (1979) and Pape (1996) in considering *A. auriceps* and *A. pelopei* as separate species and confirm the dull yellow colouration of the fronto-orbital plate in *A. auriceps* as an important character for the separation of these species.

In addition to the morphological evidence for separation, the barcode library of Australian *Amobia* prepared in this study will help to either support or refute the separation of *A. auriceps* and *A. pelopei* once molecular data have been obtained for *A. pelopei*.

All Australian representatives identified by Meiklejohn *et al.* (2012) as *A. pelopei* were examined as part of this study and are re-identified as *A. burnsi* through the diagnostic combination of golden fronto-orbitals and male terminalia. The COI barcode fragment of these specimens also matched to *A. burnsi*.

Wasp associations and the distribution of species of Amobia

Label data from Australasian species provides evidence for the association of *A. burnsi* and *A. serpenta* sp. nov. with vespid wasps (recorded from the nests, or closely following 'potter' wasps; subfamily Eumeninae), and *A. auriceps* and *A. burnsi* with sphecid wasps (recorded from 'mud' wasps; family Sphecidae). This is generally consistent with previously recorded host associations for this genus in other parts of the world. Species of *Amobia* have been recorded as kleptoparasites of sphecid and vespid wasps in the Nearctic, Oriental and Palaearctic regions (Fye 1965; Itino 1988; Wcislo 1996; Fateryga & Kovblyuk 2014) and of vespids and crabronid wasps in the Neotropical region (Freeman & Jayasingh 1975; Buys 2007; Nascimento & Garófalo 2014). No records are available that associate any Australasian species with wasps of the family Crabronidae.

While known from locations hundreds of kilometres from the nearest city or town (e.g., Litchfield NP, Northern Territory), Australia's species of *Amobia* also appear to occur with their hosts in urban environments. *Amobia burnsi* and *A. auriceps* species have been observed usurping wasp nests located in tunnels, overpasses and attached to buildings (pers. obs.). Australian *Amobia* should be expected to occur wherever suitable wasp hosts are found.

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Supplementary file 1. Vouchers, citations and accession numbers for sequences included in the phylogenetic analysis.

https://doi.org/10.5852/ejt.2020.722.1135.3003

Supplementary file 2. Bayesian inference phylogenetic tree displaying the placement of *Amobia* among the global Miltogramminae, inferred from COI, CYTB, ND4 and EF1α sequence data https://doi.org/10.5852/ejt.2020.722.1135.3005