

An evening bat (Chiroptera: Vespertilionidae) from the late Early Eocene of France, with comments on the antiquity of modern bats

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Abstract: Bats are among the most numerous and widespread mammals today, but their fossil record is comparatively meagre and their early evolution poorly understood. Here we describe a new fossil bat from dental remains recovered from late Early Eocene sediments at Prémontré, northern France. This 50 million-year-old bat exhibits a mosaic of plesiomorphic and apomorphic dental features, including the presence of three lower premolars, a single-rooted p3, short p4 with metaconid, myotodont lower molars and a tall coronoid process of the dentary. This combination of features suggests it is an early member of Vespertilionidae, today's most speciose and geographically widespread bat family. The Prémontré bat has bearing on hypotheses about the origins of vesper or evening bats (Family Vespertilionidae), as well as crown-group chiropterans.

Keywords: Western Europe, Ypresian, dental morphology, evolution, palaeobiogeography

Submitted 24 March 2016, Accepted 26 May 2016
 Published Online 1 August 2016, doi: [10.18563/pv.40.2.e2](https://doi.org/10.18563/pv.40.2.e2)
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INTRODUCTION

One fifth of all extant mammal species are bats (Simmons, 2005a) but their fossil record is poor, with an estimated 66–81% of their fossil history missing (based on ghost lineage analysis and molecular divergence dates as well as phylogeny-independent statistical methods; Teeling *et al.*, 2005; Eiting & Gunnell, 2009). Identifiable bats first appear in the fossil record in the earliest Eocene ~55 Ma in Europe (Tabuce *et al.*, 2009); older chiropterans have been predicted by molecular data and combined data analyses (e.g., Bininda-Emonds *et al.*, 2007; Meredith *et al.*, 2011; O'Leary *et al.*, 2013) but these remain unrecognized. The oldest fossil bats have unspecialised dentitions, similar to those of many other Paleocene-Eocene mammals (e.g., nyctitheriids, lipotyphlans, adapisoriculids), but skeletal adaptations indicate they could fly and most could echolocate (Haberstezer & Storch, 1987; Haberstezer *et al.*, 1994; Simmons *et al.*, 2008). These early bats are referred to several extinct families whose relationships to each other and modern lineages are unclear (Simmons & Geisler, 1998; Simmons *et al.*, 2008; O'Leary *et al.*, 2013). A probable Laurasian origin for Order Chiroptera is generally accepted (e.g., Gunnell & Simmons, 2005), but there remains debate about when, where, and how modern bat families originated (Eick *et al.*, 2005; Teeling *et al.*, 2005; Ravel *et al.*, 2011, 2014, 2015; Smith *et al.*, 2012; Yu *et al.*, 2014). Molecular data suggest that most of the 20 modern bat families had evolved by the end of the Eocene (Teeling *et al.*, 2005; Simmons, 2005b; Miller-Butterworth *et al.*, 2007) but empirical/fossil data are scarce.

The fossil locality of Prémontré in the department of Aisne, northern France (Fig. 1), is known for its rich assemblage of terrestrial and marine vertebrates of late Early Eocene age (late Ypresian, 50 Ma; Dégrémont *et al.*, 1985; Escarguel, 1999; Adnet & Cappetta, 2008). It has produced one of the

most diverse Eocene faunas in the Paris Basin (Dégrémont *et al.*, 1985; Louis, 1996) and is especially rich in small mammals, which are represented by numerous isolated teeth and dentary fragments. The most abundant mammals in the Prémontré deposit are primates (Godinot *et al.*, 1992), rodents (Escarguel, 1999), pantolestids (Smith, 2001) and bats, but rarer multituberculate, marsupial, nyctitheriid, adapisoricid, apatemyid, hyaenodontan, miacid, condylarth, perissodactyl and artiodactyl teeth also occur (Dégrémont *et al.*, 1985; Sudre & Erfurt, 1996).

The Prémontré vertebrate fossils were recovered by members of the Société Laonnoise de Paléontologie in 1980–90 from a deposit located within the grounds of the 18th Century Prémontré Abbey (now a hospital), in sediments regarded to be a lateral equivalent of the Argile de Laon and of latest Ypresian age (biozone NP12) (Dégrémont *et al.*, 1985; Lecomte, 1994; Steurbaut, 1998; Adnet & Cappetta, 2008). They occur in sandy estuarine sediments (Sables de Glennes) associated with numerous selachian teeth (Cappetta, 1992; Adnet & Cappetta, 2008) as well as bony fish, plants and invertebrates (Dégrémont *et al.*, 1985, fig. 1). The Prémontré mammal taxa are correlated with reference-level MP10 of the mammalian biochronological scale for the European Paleogene (~50 Ma; Godinot *et al.*, 1992; Lecomte, 1994; Sudre & Erfurt, 1996; Escarguel, 1999).

Dégrémont *et al.* (1985: p.16) noted that several bat taxa are represented in the Prémontré deposit as follows: *Icaronycteris* sp. indet. (common), cf. *Archaeonycteris* (rare), and a small bat whose lower molars possess a distinctive myotodont structure (rare). Here we describe the small Prémontré bat with myotodont molars as an early member of the family Vespertilionidae. Collectively known as vesper or evening bats, vespertilionids today have a global distribution (except polar regions) and number ~400 species or nearly a third of all living bats (Simmons, 2005a). The new fossil chiropteran extends the origin of this enormous radiation back to 50

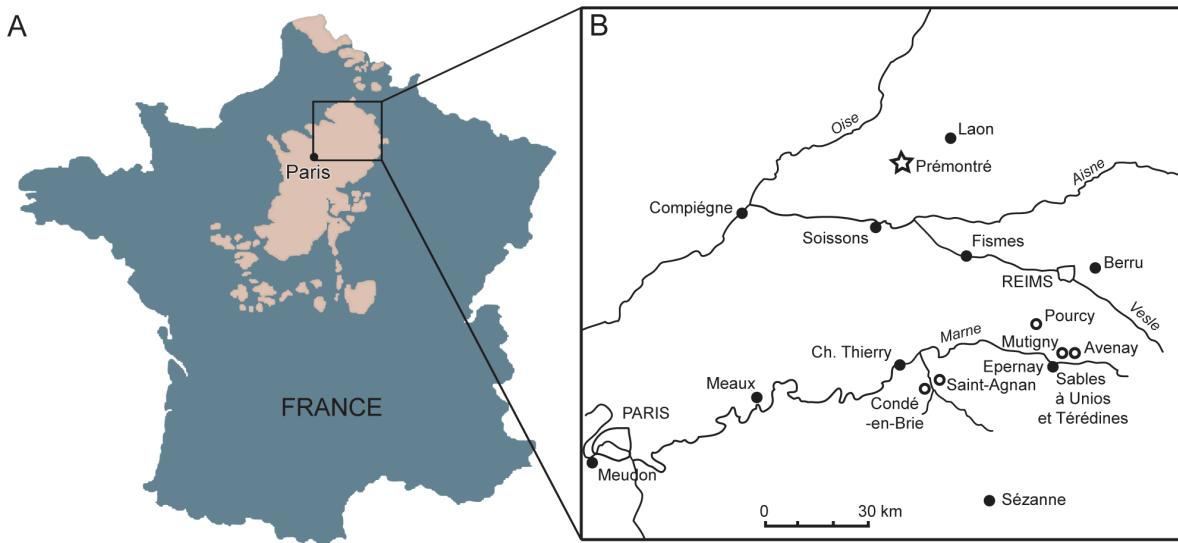


Figure 1. Map showing (A) Paleogene and Neogene sediments in the Paris Basin, and (B) location of Prémontré and other Ypresian mammal-bearing localities (open circles) noted in text (after Aubry *et al.*, 2005 and Louis, 1996).

Ma from 34 Ma (Gunnell *et al.*, 2008), with biogeographic inferences for modern bat origins, and provides an independent test of molecular clock divergence times for vespertilionids.

Institutional abbreviation. PL PRE, Pierre Louis Prémontré collections, Muséum National d'Histoire Naturelle, Paris, France; SLP, Société Laonnaise de Paléontologie collections, Muséum National d'Histoire Naturelle, Paris, France. A list of comparative materials and other institutional abbreviations is given in Appendix 1.

Dental terminology follows Hand *et al.* (2015), in which the three lower premolars are numbered p₂, p₃ and p₄, and see Fig. 2.

SYSTEMATIC PALAEONTOLOGY

Class MAMMALIA Linnaeus, 1758
Order CHIROPTERA Blumenbach, 1779
Superfamily VESPERTILIONOIDEA Gray, 1821
Family VESPERTILIONIDAE Gray, 1821
Premonycteris gen. nov.

Type species. *Premonycteris vesper* sp. nov.

Generic diagnosis. As for the type species until others are recognized.

Generic etymology. After the fossil locality Prémontré and *nycteris*, bat, alluding to an early signal or premonition about modern bats

Premonycteris vesper sp. nov.

lsid:zoobank.org:act:7A1DEFCA-0B31-4B58-B260-6A37760521F0
(Figs 3-4)

Holotype. SLP 29 BS 193; partial left dentary with p₃, p₄ and m₁, alveoli for p₂, m₂₋₃, partial alveolus for c₁, and most of ascending ramus (Fig. 3).

Referred specimens. SLP 29 PE 875, right dentary fragment with m₂ and ascending ramus nearly complete (Fig. 4A-A', C, D); SLP 29 PE 110, right dentary fragment with m₁; SLP 29 PE 614, right dentary fragment with m₂ (Fig. 4B); SLP 27 PR 8, right m₂ (Fig. 4E, F-F', G, H); PL PRE 273, right m₁; PRE 347, right ?m₁; PL PRE 611, left m₂ (Fig. 4I, J-J', K, L); PL PRE 1136, left ?m₂; PL PRE 764, right m₁; SLP 43 PR 17, right m₂; PL PRE 1076, left m₁; SLP 29 PE 369, left m₃ (Fig. 4M, N-N', O, P); SLP 29 PE 290, left m₃.

Type locality and age. Prémontré, Aisne, north-eastern Paris Basin, France; Sables de Glennes; late Ypresian, Early Eocene (MP10 of the mammalian biochronological scale for the European Paleogene).

Etymology. *vesper*, from the common name for members of the family Vespertilionidae; Latin for evening.

Differential diagnosis. Differs from onychonycterids, icaronycterids, archaeonycterids, palaeochiropterygids, hassianycterids and mixopterygids (see included taxa and diagnoses in Smith *et al.*, 2012) and *Australonycteris clarkae*

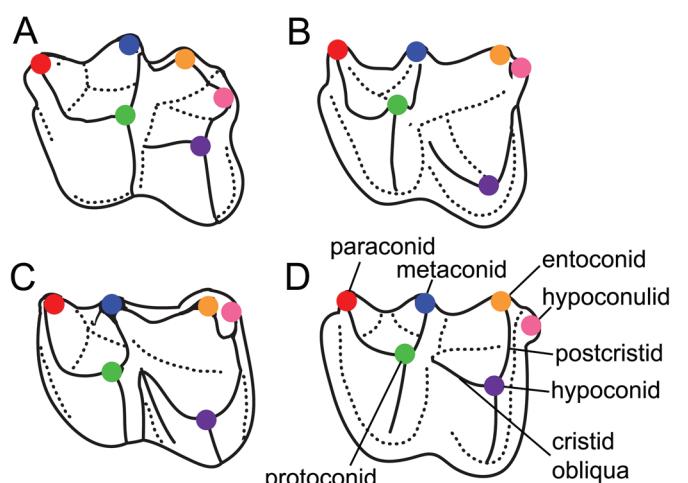


Figure 2. Lower molar structure: **A**, necromantodonty, **B**, nyctalodonty, **C**, submyotodonty, and **D**, myotodonty (after Maitre, 2014).

in its single-rooted p3, short p4 and myotodont lower molars (Hand *et al.*, 1994). Differs from *Stehlinia* spp. (regarded to be palaeochiropterygids but see Smith *et al.*, 2012 and below) in its single-rooted p3 and myotodont molars. Differs from *Necromantis* spp. (Hand *et al.*, 2012; Maitre, 2014) in its large (rather than tiny) p3, short p4 with metaconid and myotodont molars. Differs from *Tachypteron franzeni* Storch *et al.*, 2002 in its single-rooted p3 and myotodont molars. Differs in its myotodont lower molars from other early Eocene bat taxa known from m1 or m2: ?*Archaeonycteris precursor* Tabuce *et al.*, 2009, Silveirinha, Portugal; ?*Archaeonycteris* sp., Meudon, Paris Basin (Russell *et al.*, 1988); Eochiroptera indet. 2 (Marandat, 1991: pl. 5), Fournes, southern France; chiropteran indet. from Laguna Fria, Argentina (Tejedor *et al.*, 2005); eochiropteran from El Kohol, Algeria (Ravel *et al.*, 2011: fig. 4).

Differs from emballonuroids (emballonurids, nycterids), rhinopomatiids and most rhinolophoids (hipposiderids, megadermatids, craseonycterids) in having three lower premolars, short p4 with metaconid and myotodont lower molars. Differs from rhinolophids in its large p3, short p4 with metaconid and myotodont molars.

Among noctilionoids, differs from noctilionids and mystacinids in having three lower premolars. Differs from mormoopids in its large but single-rooted p3 and p4 with metaconid. Differs from phyllostomids in its p4 with tall metaconid and from most species additionally in its myotodont molars. Differs from myzopodids in its large p3 and p4 with metaconid. Differs from thyropterids in its single-rooted p3, and from furipterids in its myotodont lower molars. Differs from both in its condyloid process of dentary being lower than coronoid process. Differs from extinct speonycterids in its large p3 with one root and myotodont lower molars.

Among vespertilionoids, differs from miniopterids in its myotodont molars, $p4=p3>p2$ and tall coronoid process. Differs from natalids in its myotodont molars, single-rooted p3, $p4=p3>p2$, and condyloid process lower than coronoid process. Differs from molossids and extinct philisids in presence of p3.

Differs from other vespertilionids by the following combination of characters: three lower premolars; $p4=p3>p2$; p2 and p3 with one root each; p4 two roots; p4 molariform with paraconid and metaconid and short heel; p4 conspicuously shorter than m1; lower molars myotodont (where the posteristid directly joins the hypoconid to the entoconid isolating the hypoconulid; Fig. 2D); entoconid tall; posteristid relatively low; hypoconulid small, lingual and low; cristid obliqua meets trigonid at centre of crown; trigonid and talonid widths similar; m3 talonid only moderately reduced in width; m3 hypoconulid present; dentary with tall ascending ramus and coronoid process higher than condyloid process; well-developed, laterally inflected angular process.

Description

The horizontal (mandibular) ramus (SLP 29 BS 193, SLP 29 PE 875; Figs 3–4) is shallow, with a straight ventral margin, mental foramen beneath p2, a narrow symphysis extending posteriorly to p2, and has a maximum depth of 1.5 mm (approximately equivalent to m1 height). The dentary has a tall ascending ramus, with the condyloid process occurring at the level of the tooth row and conspicuously lower than the tall coronoid process. The anterior edge of the coronoid rises nearly vertically, then curves posterodorsally near its (missing) tip. The well-defined angular process is slightly laterally

inflected and occurs just dorsal to the ventral margin of the dentary. An elongate mandibular foramen opens at the level of the tooth row.

The dental formula is i?, c1, p2, p3, p4, m1-3, where p2 and p3 are single rooted and p4 has two roots, all aligned in the tooth row (SLP 29 BS 193, SLP 29 PE 110, SLP 29 PE 875, SLP 29 PE 614). The p4 is conspicuously shorter than m1 (55% of m1 length; Table 1) and of similar length to p3 but wider. The minimum length of p2 is 0.5 mm based on alveolus length. The p3 is single rooted (confirmed by microCT; Fig. 3C) but with a clear longitudinal groove in the lingual face of the root. The p3 protoconid is worn but similar in height to p4, and is surrounded by a more or less continuous cingulid interrupted by a small anterolingual cusp; p3 lacks a metaconid. The p4 has a metaconid and paraconid (both much lower than the protoconid), a short heel and continuous anterior, buccal and posterior cingulids.

The m1 (SLP 29 BS 193, SLP 29 PE 110, PL PRE 273, PL PRE 764, ?PRE 347; Fig. 3) is longer than wide, with the trigonid and talonid of similar widths. In the trigonid, the protoconid is the tallest cusp, and the paraconid well developed but not as tall as the metaconid. The talonid exhibits the myotodont condition (where the posteristid directly joins the hypoconid to the entoconid thereby isolating the hypoconulid); the hypoconulid is small and low. The posteristid is low and the cristid obliqua is inflected anteriorly before meeting the trigonid in the centre of the tooth. The anterior, buccal and posterior cingulids are narrow but continuous, and there is no lingual cingulid. The posterior root is thicker than the anterior root. The m2 (SLP 29 PE 875, SLP 29 PE 614, SLP 27 PR 8, PL PRE 611, PL PRE 1136, SLP 43 PR 17; Fig. 4) is very similar to m1, but differs in its better developed paraconid and more anteroposteriorly compressed trigonid. With respect to m1 and m2, m3 (SLP 29 PE 290, SLP 29 PE 369; Fig. 4) is similar in trigonid length and width, and only moderately reduced in talonid width. The entoconid is relatively lower, the entocristid at a slight angle to the long axis of the tooth and the postcristid at right angles to the entocristid.

Measurements of *Premonycteris vesper* are given in Table 1.

Gunnell *et al.* (2009) developed a set of algorithms to estimate body mass in extinct bats, based on dental, skeletal and weight measurements in 1,160 extant bats representing eight families (including Vespertilionidae). Using the proxy of lower first molar (m1) area and data in Gunnell *et al.* (2009: table 1), we estimated the weight of *Premonycteris vesper* to be 9.5 g, suggesting a relatively small bat, compared with the median value of 13.8 g for 905 extant bat species, and 3 to 50 g for living vespers (Smith *et al.*, 2004; Gunnell *et al.*, 2009).

DISCUSSION

A combination of dental features appears to exclude *Premonycteris vesper* from the extinct Eocene bat families Onychonycteridae, Icaronycteridae, Archaeonycteridae, Palaeochiropterygidae, Hassianycteridae, Mixopterygidae and Philisidae. This combination includes the presence of three lower premolars where $p4=p3>p2$, a large single-rooted p3, short p4 with metaconid, and myotodont lower molars (see Differential diagnosis above; and Table 2). A maximum of three upper and three lower premolars occurs in bats, but many lineages have subsequently lost one or two additional premolars (Table 2). In *Premonycteris vesper*, the retention of three lower premolars

is shared (probably plesiomorphically) with onychonycterids, icaronycterids, archaeonycterids, hessianycterids and palaeochiropterygids, as is presence of a metaconid on p4 shared with at least onychonycterids, icaronycterids, archaeonycterids and hessianycterids. The relative sizes of the premolars in *Premonycteris vespér* are more similar to those of icaronycterids, archaeonycterids, hessianycterids and palaeochiropterygids than to onychonycterids in which p3 is generally smaller than p4 and p2 (Smith *et al.*, 2012; Hand *et al.*, 2015; Table 2).

The higher level taxonomy of a number of Eocene bat taxa

remains in flux. *Stehlinia* species from the Middle Eocene to Late Oligocene of Europe were originally referred to the modern family Vespertilionidae (e.g., Revilliod, 1920) but subsequently transferred to the extinct Palaeochiropterygidae (e.g., Maitre, 2014). Palaeochiropterygidae has been included in Vespertilioidea (e.g., Russell & Sigé, 1970; Sigé, 1997; Maitre, 2014), but was found to be the immediate sister group of the modern crown-group Chiroptera in phylogenetic analyses (e.g., Gunnell & Simmons, 2005; Simmons *et al.*, 2008). *Lapichiropteryx xiei* from Middle Eocene of China is referred to Palaeochiropterygidae (Tong, 1997). However, the

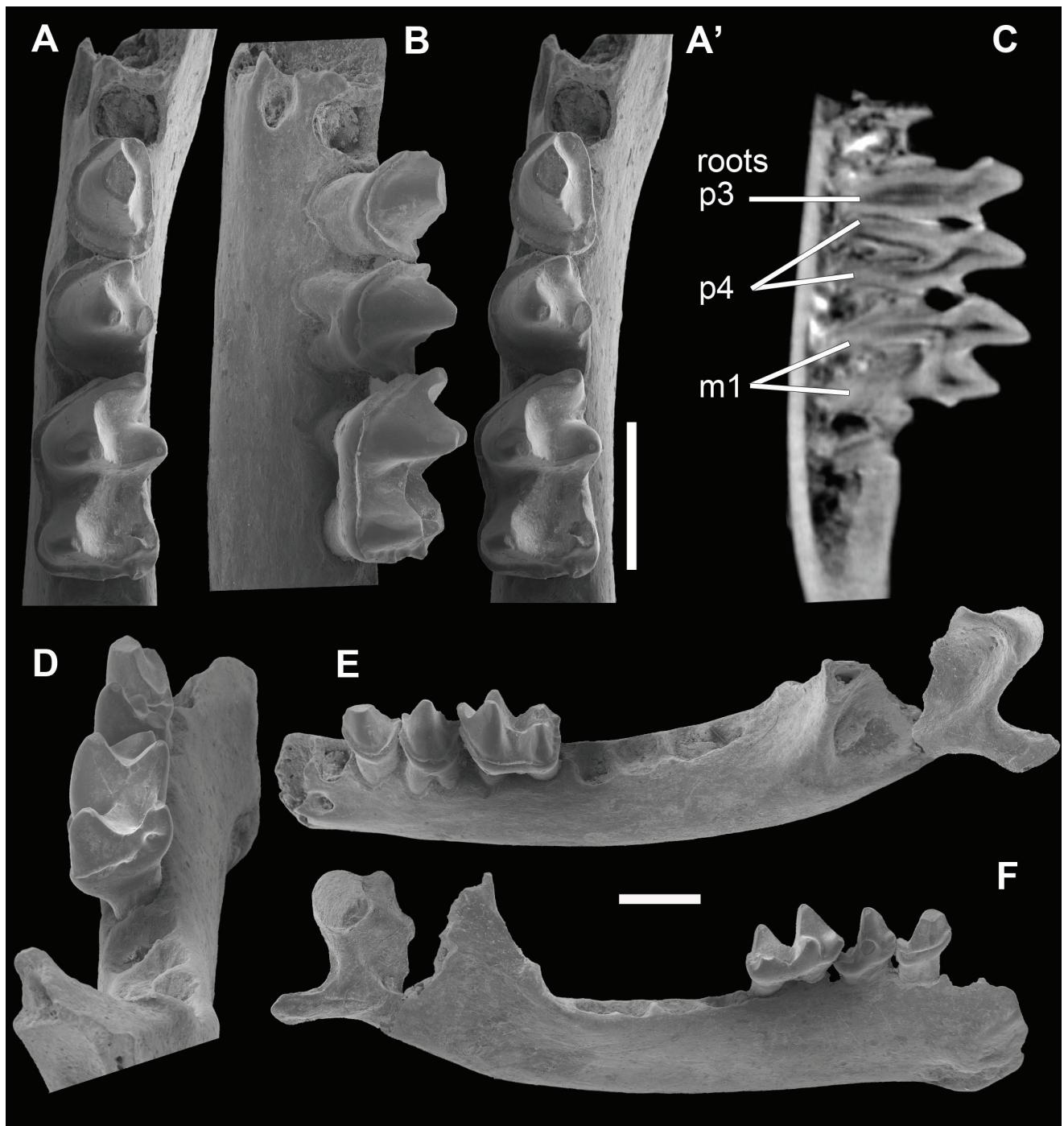


Figure 3. *Premonycteris vespér* gen. et sp. nov., Prémontre, Aisne, France; late Ypresian, Early Eocene. SLP 29 BS 193; partial left dentary with p3, p4 and m1, partial alveoli for c1, alveoli for p2, m2-3, and most of ascending ramus. **A-A'**, p3-m1, stereopair, occlusal view; **B**, buccal view; **C**, microCT axial section showing roots of p3-m1; **D**, posterior view; **E**, dentary, buccal view; **F**, dentary, lingual view. Scale bars = 1 mm.



Figure 4. *Premonycteris vesper* gen. et sp. nov. SLP 29 PE 875, right dentary fragment with m2 and ascending ramus. **A-A'**, stereopair, occlusal view; **C**, buccal view; **D**, lingual view. SLP 29 PE 614, right dentary fragment with m2. **B**, oblique-occlusal view. SLP 27 PR 8, right m2; **E**, buccal view; **F-F'**, stereopair, occlusal view; **G**, lingual view; **H**, posterior view. PL PRE 611, left m2; **I**, buccal view; **J-J'**, stereopair, occlusal view; **K**, lingual view; **L**, posterior view. SLP 29 PE 369, left m3; **M**, buccal view; **N-N'**, stereopair, occlusal view; **O**, lingual view; **P**, posterior view. Scale bars = 1 mm.

familial assignment of *Lapichiropteryx* and *Stehlinia* species has been questioned by Smith *et al.* (2012), who noted that they share, among other features, a short p4 not found in other palaeochiropterygids (see below).

Irrespective, none of the archaic Eocene bats, including *Lapichiropteryx* and *Stehlinia* species, have a single-rooted p3, nor do they have myotodont lower molars (although the structure is not clear in *Onychonycteris finneyi*; Gunnell *et al.*, 2011). It is the combined presence of these features that persuades us *Premonycteris* does not belong in any of the extinct bat families.

In the myotodont lower molar structure of bats, the postcristid directly joins the hypoconid to the entoconid, and the posterolingual hypoconulid is typically greatly reduced with respect to the entoconid and isolated or linked to it by a small crest (as defined by Menu & Sigé, 1971; Sigé *et al.*, 2012; Fig. 2). The postcristid forms a blade that cuts against the anterior face of the paracone in the following upper molar. Myotodonty is less common than nyctalodonty among bats and may be more evolutionarily derived than the other dental patterns (nyctalodonty and necromantodonty; Menu & Sigé, 1971; Fig. 2). Myotodonty appears to have been independently acquired several times in various families, for example, among molossids (e.g., Hand, 1990; Maitre, 2014), phyllostomids (e.g., Sigé *et al.*, 2012), mystacinids (Hand *et al.*, 1998) and myzopodids (Gunnell *et al.*, 2014), but most frequently among vespertilionids with the majority of species and genera showing this pattern (Menu & Sigé, 1971; Menu, 1985, 1987; and see below). Several authors have noted that the pattern can be variable in some populations of some species (see examples in Sigé, 1995; Maitre, 2014); an intermediate structure, submyotodonty, is also recognized, in which the postcristid connects the hypoconid and entoconid as in myotodonty but a short crest may also connect the postcristid and hypoconulid (Legendre, 1984; Fig. 2). Generally, however, one pattern (nyctalodonty or myotodonty) dominates in a bat population (Menu & Sigé, 1971).

The combination of lower premolar and molar features noted above for *Premonycteris vesper* distinguishes it not only from extinct Eocene bat families but also from two of the four modern bat superfamilies, Rhinolophoidea and Emballonuroidea (Table 2). Rhinolophoids are characterized by p3 absent or tiny, relatively long p4 lacking metaconid, and presence of nyctalodont molars. Emballonuroids typically have two

lower premolars (except Eocene *Tachypteron* which retains a two-rooted p3; Table 2) and nyctalodont molars (although myotodonty is observed in some *Vespertiliavus* populations; Sigé, 1995; Maitre, 2014).

Noctilionoids exhibit other combinations of these features (see Differential diagnosis and Table 2). Myzopodids have three lower premolars and myotodont molars but reduced p3 and p4 without metaconid. Mormoopids have three lower premolars and myotodont molars but p3 either small and single-rooted or large and double-rooted. Noctilionids and mystacinids have myotodont molars but only two lower premolars. Phyllostomids have two or three lower premolars but generally a p4 lacking metaconid and nyctalodont molars (or modified for fruit, nectar or blood feeding). Thyropterids have myotodont molars and three premolars but p3 has two roots. Furipterids and extinct speonycterids have three premolars but nyctalodont lower molars. Thyropterids, furipterids and mormoopids (*Mormoops* spp.), like natalids and kerivoulines (below), differ additionally in having a low coronoid process of the dentary that is level with the condyloid process, reflecting significant dorsal cranial flexion (Morgan & Czaplewski, 2003) evidently not present in *Premonycteris vesper*.

The remaining, most speciose modern bat superfamily, Vespertilioidea, contains the families Vespertilionidae (~400 spp.), Miniopteridae (20 spp.), Cistugidae (2 spp.), Natalidae (11 spp.), Molossidae (~100 spp.) and extinct Philisidae (6 spp.) (Simmons, 2005a; Lack *et al.*, 2010; Ravel *et al.*, 2015). Molossids retain only two lower premolars, both with two roots, and generally have nyctalodont but sometimes myotodont lower molars (Table 2). Philisids have myotodont molars but only two lower premolars (premolars unknown in *Dizya*). Three lower premolars occur in natalids and miniopterids, but natalids have a two-rooted p3 and nyctalodont molars, while miniopterids have a single-rooted p3 but nyctalodont molars. Only (some) vespertilionids have three lower premolars, a single-rooted p3, short p4, and myotodont lower molars (e.g., species of *Myotis*, *Kerivoula*, *Plecotus*, *Idionycteris*; Table 2 and see below). The morphology of the dentary of *Premonycteris* (see Description above and Figs 3-4) is also consistent with that found in most vespertilionids (except kerivoulines as noted above).

Molecular and morphological data indicate that Vespertilionidae is monophyletic (e.g., Simmons *et al.*, 2008; Lack *et al.*, 2010) but no unambiguous dental synapomorphies for the family have been identified (Simmons & Geisler, 1998; Czaplewski & Morgan, 2001; Horáček, 2001; Hoofer & Van den Bussche, 2003; Eick *et al.*, 2005; Gunnell & Simmons, 2005; Simmons *et al.*, 2008; Ravel *et al.*, 2015). Vespertilionids are often characterized by their relatively unspecialized dentitions (e.g., Sigé, 1974; Menu, 1987; Horáček, 2001; Horáček *et al.*, 2006; Maitre, 2014). Slaughter (1970) noted that each dental feature considered primitive for bats can be found in some vespertilionid, although no modern species retains them all. Submyotodonty has been suggested as the plesiomorphic dental pattern among vespertilionoids (e.g., Ravel *et al.*, 2015), and myotodonty probably the primitive condition in vespertilionids (e.g., Sigé, 1985; Horáček, 2001; Gunnell *et al.*, 2012), but a thorough phylogenetic analysis of this speciose family is needed to formally establish this. In their molecular analysis of myotines, Lack *et al.* (2010) note that few informative taxonomic characters, extensive morphological convergence, and a likely rapid initial diversification have made it difficult to unravel the evolutionary relationships of vespertilionids on the basis of morphology.

Nevertheless, the combination of three lower premolars, a

Specimen no.	Position	Length	Width	Trig. width	Tal. width
SLP 29 BS 193*	p3-m1				
	p3	0.75	0.70		
	p4	0.75	0.80		
	m1	1.35		0.95	0.90
SLP 29 PE 110	m1	1.40		0.90	0.95
PRE 273	m1	1.40		0.90 (est.)	0.95
PRE 347	?m1	1.40		1.00	0.90
PRE 764	m1	1.35		1.00	1.00
PL PRE 1076	m1	1.35		0.95	0.90
SLP 29 PE 875*	m2	1.30		0.90	0.90
SLP 29 PE 614*	m2	1.35		1.00	0.95
PRE 1136	?m2	1.40		0.95	0.95
PRE 611*	m2	1.45		1.00	1.00
SLP 27 PR 8*	m2	1.40		0.90	0.85
SLP 43 PR 17	m2	1.35		0.90	0.90
SLP 29 PE 369*	m3	1.30		0.90	0.75
SLP 29 PE 290	m3	1.30		0.95	0.70

Table 1. Measurements (mm) of specimens of *Premonycteris vesper* gen. et sp. nov. Abbreviations: Trig., trigonid; Tal., talonid; *, figured; est., estimate.

single-rooted p3, short p4 with metaconid, myotodont lower molars, as well as tall coronoid process of the dentary, is most similar to features frequently seen among vespertilionionids (see below), and not other bat families, which is why, on the basis of available data, we refer *Premonycteris vesper* to Vespertilionidae. We recognise, however, that these features almost certainly represent a mosaic of plesiomorphic and apomorphic traits, and that it is possible, when more of the vespertilionoid fossil record is known, and/or dental synapomorphies for vespertilionids identified, that

Premonycteris vesper may be recognized as a derived member of a stem group rather than an early representative of the modern clade.

While *Premonycteris vesper* is interpreted here to be a vespertilionionid for the reasons noted above, it is not clear to which, if any, of the currently recognized four vespertilionionid subfamilies (sensu Roehrs *et al.*, 2010) it might belong. A member of the extant myotine subfamily, *Khonsunycteris aegyptiacus* from the late Eocene (~34 Ma) of the Fayum, Egypt, is the oldest, widely accepted fossil record for Vespertilionidae

Family	Species	Epoch	FAD (Ma)	Lower premolars	Lower premolar size	p3 roots	p4 metaconid	Molar form
Extinct Eocene families								
Onychonycteridae†	<i>Onychonycteris finneyi</i>	Early Eocene	53	3	p4>p3<p2	2	absent	myotodont?
	<i>Eppsinycteris anglica</i>	Early Eocene	55	3	p4>=p3	3	present	necromantodont
	<i>Honrovitis tsuwave</i>	Early Eocene	53	3	p4>p3<p2	2	absent	necromantodont
Icaronycteridae†	<i>Icaronycteris menui</i>	Early Eocene	54	3	p4>p3>p2	2	present	nyctalodont
Archaeonycteridae†	<i>Archaeonycteris</i> spp.	E-M Eocene	55	3	p4>p3>p2	2	present	necromantodont
	<i>Protonycteris gunnelli</i>	Early Eocene	54	3	p4>p3>p2	2	present	necromantodont
Hessianycteridae†	<i>Hessianycteris</i> spp.	E-M Eocene	54	3	p4>=p3>p2	2	absent	nyctalodont
Palaeochiropterygidae†	<i>Palaeochiropteryx</i> spp.	Middle Eocene	47	3	p4>p3>p2	2	present	nyctalodont
?Palaeochiropterygidae†	<i>Stehlinia</i> spp.	M Eoc-E Oligo	43	3	p4>=p3>p2	2	variable	nyctalodont
	<i>Lapichiropteryx xiei</i>	Middle Eocene	3				present	nyctalodont
Family indet.†	<i>Necromantis adichaster</i>	Middle Eocene	43	3	p4= p2>p3	1	absent	necromantodont
Mixopterygidae†		M Eoc - L Oligo	2		p4>p2	-	?absent	nyctalodont
	<i>Carcinopteryx maximensis</i>	M Eocene	43	2	p4>p2	-	?absent	nyctalodont
SF Emballonuroidea								
Emballonuridae	<i>Tachypteron franzeni</i> †	M Eocene	47	3	p4>p3>p2	2	?absent	nyctalodont
	<i>Vespertiliavus</i> spp. †	M Eoc- L Oligo	43	2	p4=p2	-	present	nyctalodont (var)
SF Rhinophoidea								
Rhinolophidae	<i>Protorhinolophus shanghuangensis</i> †	M Eocene	43	3	p4>p3<p2	1	absent	nyctalodont
Hipposideridae	<i>Hipposideros schlosseri</i> †	M Eocene	43	2	p4>p2	-	absent	nyctalodont
Megadermatidae	<i>Saharaderma pseudovampyrus</i> †	L Eocene	34	2	p4>p2	-	absent	nyctalodont
SF Noctilionoidea								
Myzopodidae	<i>Phasmatonycteris</i> spp. †	L Eoc-L Oligo	37.5	3	p4>p3<p2	1	absent	myotodont
Speonycteridae†	<i>Speonycteris aurantiadens</i>	E Oligocene	30	3	p4>p3>p2	2	?present	nyctalodont
Mystacinidae	<i>Icarops paradox</i> †	E Miocene	18	2	p4>p2	-	unknown	myotodont
Mormoopidae	Mormoopid indet.	E Oligocene	30	3	p4>p3=p2	2	absent	nyctalodont
	<i>Pteronotus</i> spp.	Pleist-Recent	2	3	p4>p3<p2	1	absent	myotodont
	<i>Mormoops</i> spp.	Pleist-Recent	2	3	p4>p3=p2	2	absent	nyctalodont
Noctilionidae	<i>Noctilio</i> spp.	M Miocene	13	2	p4>p2	-	absent	myotodont
Phyllostomidae	<i>Macrotis waterhousii</i>	Pleist-Recent	2	3	p4=p3=p2	2	absent	myotodont
	<i>Chrotopterus</i> spp.	Pleist-Recent	2	3	p4>p3>p2	1	?absent	nyctalodont
	<i>Notonycteris</i> spp. †	M Miocene	13	3	p4>p3<p2	1	unknown	nyctalodont
SF Vespertilioidea								
Natalidae		L Oligo-Recent						myot & nyctalodont
	<i>Natalus</i> spp.	Pliocene-Recent	5	3	p4=p3=p2	2	absent	nyctalodont
	<i>Primonatalus prattae</i> †	E Miocene	19	3	p4=p3>p2	2	?unknown	nyctalodont
Molossidae		L Eocene-Recent						myot & nyctalodont
	<i>Cuvierimops</i> spp. †	L Eoc-L Oligo	37	2	p4>p2	-	present	myot & nyctalodont
Miniopteridae	<i>Miniopterus</i> spp.	E Mioc-Recent	22	3	p4>p3>=p2	2	absent	nyctalodont
Philisidae†		E Eocene-E Oligo						myotodont
	<i>Dizya exultans</i>	Early Eocene	50	unknown	unknown	?	unknown	myotodont
	<i>Witwitia schlosseri</i>	Late Eocene	37.5	2	p4>p3	-	?present	myotodont
Cistugidae	<i>Cistugo</i> spp.	Recent	0.1	3	p4>p3<p2	1	?absent	myotodont
Vespertilionidae	<i>Premonycteris vesper</i> †	Early Eocene	50	3	p4=p3>p2	1	present	myotodont
	<i>Khonsunycteris aegyptiacus</i> †	Late Eocene	34	3	p4>p3<p2	2	absent	myotodont
	<i>Quinetia misonei</i> †	E Oligocene	32.4	3	p4>p3>=p2	1	unknown	nyctalodont
	<i>Hanakia sejafari</i> †	E Oligocene	32.4	3	p4>p3>p2	1	absent	myotodont
	<i>Myotis/Leucone</i> spp.	E Oligo-Recent	31	3	p4>p3<p2	1	absent	myotodont
	<i>Murina</i> spp.	Pliocene-Recent	5	2	p4>p3	-	present	nyctalodont
	<i>Kerivoula</i> spp.	Pliocene-Recent	5	3	p4=p3=p2	1	absent	myotodont
	<i>Idionycteris phyllotis</i>	Recent	0.1	3	p4>p3<p2	1	present	myotodont
	<i>Plecotus</i> spp.	M Mio-Recent	9	3	p4>p3>=p2	1	variable	myotodont
	<i>Lasionycteris</i> spp.	Pliocene-Recent	5	3	p4>p3>p2	1	?absent	nyctalodont
	<i>Nyctalus</i> spp.	L Oligo-Recent	32.4	2	p4=p2	-	absent	nyctalodont
	<i>Pipistrellus</i> spp.	L Mio-Recent	10	2	p4=p2	-	variable	myot & nyctalodont

Table 2. Summary of dental features for representative bat taxa noted in text. Abbreviations: FAD, first appearance date ; E, early; Eoc, Eocene; L, late; M, middle; Ma, millions of years ago; Mio, Miocene; myot, myotodont; Oligo, Oligocene; †, extinct taxon; SF, superfamily. Data from Czaplewski & Morgan, 2001, 2012; Eiting & Gunnell, 2009; Gunnell *et al.*, 2008, 2014; Gunnell & Simmons, 2005; Harrison & Hooker, 2010; Hooker, 1996; Horáček, 2001; Maitre, 2014; Morgan & Czaplewski, 2003, 2012; Ravel *et al.*, 2011, 2012, 2014, 2015; Sigé, 1991; Smith *et al.*, 2012; Storch *et al.*, 2002; and pers. obs. (S. J. Hand, 2016).

(Gunnell *et al.*, 2008; Figs 5-6). *Premonycteris vesper* shares with extant myotines and *Khonsunycteris aegypticus* presence of three premolars and myotodont lower molars, but differs in its taller and longer, but narrower p3 and shorter p4 with metaconid. It differs additionally from *K. aegypticus* in p3 having one root, and an evidently less reduced m3 (known only by alveoli in *Khonsunycteris*). Kerivoulines, like *Premonycteris vesper*, have three lower premolars and myotodont molars, but p2 is larger, p3 double-rooted, p4 lacks a metaconid and the coronoid process of the dentary is relatively low with respect to the condyle. Murinines have two lower premolars and generally nyctalodont lower molars, and are least similar to *Premonycteris vesper*. In vespertilionines with three premolars and myotodont molars (e.g., *Plecotus*, *Idionycteris* and allies; Table 2), p2 and p3 are usually conspicuously smaller than p4, with p3 often smallest. Overall, the mosaic of traits observed in the lower dentition of *Premonycteris vesper* does not precisely match any modern vespertilionid subfamily, but could be ancestral to vespertilionines. Until the upper dentition of *Premonycteris* is known, we refrain from assigning it to an existing modern vespertilionid subfamily.

Premonycteris vesper extends the fossil record of the family Vespertilionidae back to 50 Ma (Fig. 5). This late Early Eocene age is similar to the oldest record for the modern bat families Hipposideridae and Emballonuridae, undescribed taxa from the late Early Eocene of Chambi, Tunisia (Sigé, 1991; Ravel *et al.*, 2011, 2014, 2015; Figs 5-6), and only 5 million years younger than the oldest currently known record for Chiroptera: *?Archaeonycteris precursor* from Silveirinha, Portugal (Tabuce *et al.*, 2009; Figs 5-6). Globally, Early Eocene bat faunas are characterized by species referred to several extinct families (see Smith *et al.*, 2012 for a review of taxa) that apparently represent stem taxa forming progressively closer sister-groups to the modern bat clade (e.g., Simmons & Geisler,

1998; Gunnell & Simmons, 2005; Simmons *et al.*, 2008). Alternatively, some could represent a separate archaic clade (O’Leary *et al.*, 2013). During the Middle Eocene, there was a decline of archaic bat families and increasing representation of modern families in Eurasia, Africa and North America (i.e., where the record is sufficiently known; Fig. 5). At Prémontre, the vespertilionid *Premonycteris vesper* lived alongside at least two archaic bat species (*Icaronycteris* sp. indet. and cf. *Archaeonycteris*; Dégrémont *et al.*, 1985, and pers. obs. [S. J. Hand, 2016]).

Discovery of *Premonycteris vesper* in the late Early Eocene of northern France adds complexity to the evolutionary history of vespertilionoids. The presence of modern bat groups in the late Early Eocene of Afro-Arabia, and their abrupt, taxonomically-diverse appearance in Middle Eocene Europe, has suggested an African source for the group with subsequent immigration into Europe (Sigé, 1991; Gunnell *et al.*, 2008; Ravel *et al.*, 2011, 2015). The extinct philisids of Paleogene Afro-Arabia have been identified as an early vespertilionoid offshoot, and the Late Eocene *Khonsunycteris aegypticus* as an ancestor for crown vespertilionids (Ravel *et al.*, 2015: fig. 6). *Premonycteris vesper* appears to be equally or more derived in some features (myotodont lower molars, p3 single rooted) than the younger myotine vespertilionid *Khonsunycteris* but less derived in other features (larger p3, molariform p4), suggesting these two Eocene vespertilionids shared an earlier common ancestor. Further, Afro-Arabian philisids appear to be more derived than the similarly-aged European *Premonycteris vesper* in at least some dental features (two lower premolars, m3 lacking hypoconulid). Taken together, these observations suggest that an African origin for superfamily Vespertilioidea is less certain.

Molecular estimates for the time of divergence of vespertilionids from other vespertilionoids (i.e., natalids,

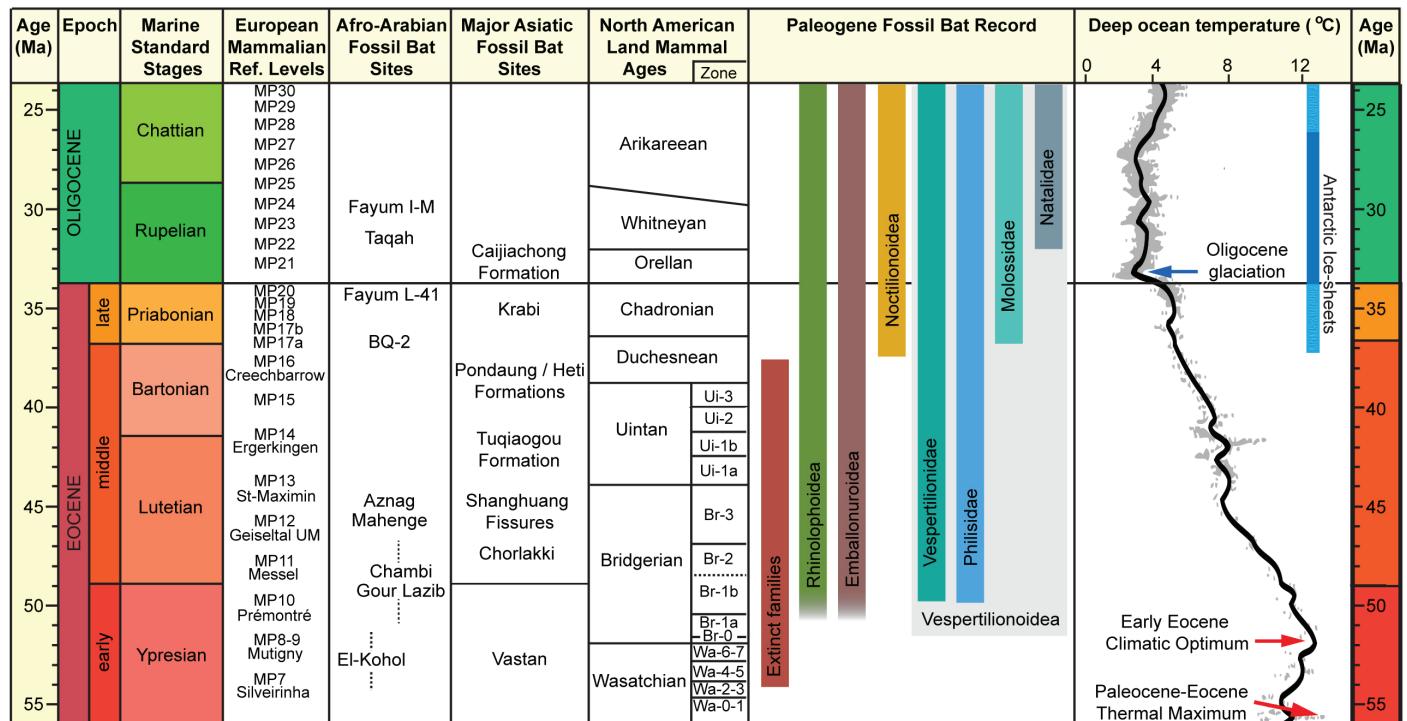


Figure 5. Correlation chart of the Paleogene fossil bat record with marine standard stages, European Mammalian Paleogene (MP) Reference Levels, North American Land Mammal Ages, European, Afro-Arabian and Asiatic fossil bat sites and Paleogene climate change. Modified from Godinot (2014). Deep ocean temperatures modified from Zachos *et al.* (2001).

minipterids, molossids, cistugids) range from 55 to 30 Ma (e.g., Teeling *et al.*, 2005 [53–42 Ma]; Eick *et al.*, 2005 [49–41 Ma]; Miller-Butterworth *et al.*, 2007 [49–38 Ma]; Meredith *et al.*, 2011 [52–42 Ma]; Lack *et al.*, 2010 [38–30 Ma]; Yu *et al.*, 2014 [55–51 Ma]). Fossils used to calibrate the molecular clock for vespertilionoid evolution, and bats more broadly, include the early Middle Eocene appearance of *Stehlinia* spp. and the late Middle Eocene *Wallia scalopidens* (e.g., Teeling *et al.*, 2005; Miller-Butterworth *et al.*, 2007; Meredith *et al.*, 2011; O’Leary *et al.*, 2013). Use of these calibration points leads to a minimum date of 37 Ma (end of Middle Eocene) for the split between vespertilionids and molossids. As noted above, the familial identity of *Stehlinia* species among bats remains uncertain (e.g., Smith *et al.*, 2012) and *Wallia*, originally described as a non-bat (lipotyphlan) by Storer (1984) and interpreted as a possible molossid by Legendre (1985), is currently regarded as a non-chiropteran mammal (Smith *et al.*, 2012). Nevertheless, widely accepted molossids (*Cuvierimops* spp.) do appear in the early Late Eocene record (Maitre, 2014), and 37 Ma is a reasonable *minimum* date for this divergence. If we are correct in our assignment of *Premonycteris vesper* to Vespertilionidae, this fossil provides a new calibration point for the family, and crown Vespertilioidea more broadly, in dated molecular phylogenies of bats. At 50 Ma, *Premonycteris* is similar in age or older than the time of vespertilionid origins predicted by analyses of genomic data.

The fossil record for vespertilionoids, despite evidently being only ~25% complete, is better than that for most bat superfamilies (Eiting & Gunnell, 2009). Vespertilionid generic diversity is lowest in the Paleogene (5 genera), increasing to 10 – 20 genera after the early Miocene, and highest at 51 genera today (Eiting & Gunnell, 2009; Simmons, 2005b; pers. comm. N. Simmons, 2016). Molecular analyses detect an increase in diversity at the end of the Eocene, as well as a Miocene explosion of vespertilionid lineages (e.g., Jones *et al.*, 2005; Stadelman *et al.*, 2007; Lack *et al.*, 2010; Yu *et al.*, 2014). These authors suggest that these diversifications probably relate to global cooling at the end of the Eocene (Fig. 5) and the later Miocene, with lower sea levels and new land connections emerging, as well as the decline of non-hibernating rhinolophoid and emballonuroid bats in temperate

zones. These events would have provided new opportunities for vespertilionids able to exploit energy-conserving metabolic and behavioural capabilities, such as torpor, migration and delayed fertilization, to cope with increasingly unpredictable conditions and resources (e.g., Horáček, 2001; Stadelman *et al.*, 2007; Lack *et al.*, 2010; Yu *et al.*, 2014).

With an estimated weight of 9.5 g (see Description above), *Premonycteris vesper* was a relatively small bat, compared with an average 13.8 g for 905 extant bat species, and 3 to 50 g for living vespers (Smith *et al.*, 2004; Gunnell *et al.*, 2009). It was significantly smaller than many Early Eocene bats (e.g., 24 – 41 g for *Icaronycteris*, *Onychonycteris* and *Archaeonycteris* spp.; Giannini *et al.*, 2012) but similar to estimates for the Middle Eocene Messel palaeochiropterygid *Palaeochiropteryx tupaiodon* (Giannini *et al.*, 2012), and larger than the early-middle Early Eocene palaeochiropterygid *Microchiropteryx folieae* from Vastan, India (Smith *et al.*, 2007: table 1 dental measurements; Rose *et al.*, 2014; Smith *et al.*, 2015).

Premonycteris lived during the Early Eocene Climatic Optimum (EEOC) when the climate was globally warmer than today (53–50 Ma; Fig. 5), with an estimated mean annual sea surface temperature 50 Ma in the Paris Basin of ~22°C (Huyghe *et al.*, 2015). Warm, humid megathermal rainforest extended to high latitudes in the EEOC and this was associated with high diversification rates among mammals (Collinson, 2000). In the Prémontré assemblage, the abundant remains of primates, arboreal rodents and bats, and a rarity of large mammals, suggest a locally densely forested habitat (Degrémont *et al.*, 1985). There is no evidence of regional karst or caves and the evidence collectively suggests *Premonycteris* was probably a tree-dweller like many extant vespertilionids (e.g., *Lasiurus* and *Pipistrellus* species; Menzel *et al.*, 1998; Veilleux *et al.*, 2003). Its simple dilambodont dentition indicates that *Premonycteris* ate insects, perhaps caught by aerial hawking or gleaning like many vespertilionids past and present (Norberg & Rayner, 1987; Norberg, 1989; Horáček, 2001), and differs significantly from the specialized dentition and proposed carnivorous diet of contemporary philisids (Ravel *et al.*, 2015). Simmons & Geisler (1998) have argued that continuous aerial hawking using echolocation was the primitive foraging strategy for crown group bats, and a key innovation in their evolution.

In conclusion, we remain in the dark about whether *Premonycteris*, Europe’s oldest vespertilionid, was an immigrant or evolved *in situ* during the EEOC from one of the extinct groups such as palaeochiropterygids. *Premonycteris* is still poorly known but does not appear to be a member of any modern vespertilionid subfamily. As noted above, it is also possible that *Premonycteris vesper* may be a derived member of a stem group rather than an early representative of the modern clade Vespertilionidae. Much of the fossil record, phylogenetic relationships and evolutionary history of today’s most successful family of bats remains to be revealed.

ACKNOWLEDGMENTS

For access to specimens and casts, we thank: D.E. Russell and M. Godinot, Muséum National d’Histoire Naturelle, Paris, France; B. Marandat and R. Tabuce, Université de Montpellier, Montpellier, France; T. Smith, Royal Belgian Institute of Natural Sciences, Brussels, Belgium; G. Storch, J. Habersetzer, E. Schlosser-Sturm, and R. Rabenstein, Forschungsinstitut Senckenberg, Frankfurt am Main, Germany; N.



Figure 6. Eocene fossil bat localities noted in text and Fig. 5 (modified from Smith *et al.*, 2012: figs 16 & 17); palaeogeographic reconstruction of 50 Ma after Scotese (2006). Filled circle, Early Eocene; open circle, Middle Eocene; open triangle, Middle-Late Eocene; filled triangle, Middle Eocene-Oligocene site.

Micklich, Hessisches Landesmuseum, Darmstadt, Germany; G. F. Gunnell, Duke University, Durham, USA; N. B. Simmons and E. Westwig, American Museum of Natural History, New York, USA. For microCT scan data, we thank T.-t. Hung, Lowy Cancer Centre, University of New South Wales. We thank N. B. Simmons, G. F. Gunnell, T. Smith and R. Tabuce for their constructive comments on drafts of the manuscript. This research was supported by Australian Research Council DP130100197.

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APPENDIX 1

List of Comparative Material Used in this Study

Institutional abbreviations: AMNH American Museum of Natural History, New York, USA; BMNH The Natural History Museum, London, UK; CB Chambi collection, Office National des Mines, Tunis, Tunisia; CM Carnegie Museum of Natural History, Pittsburgh, USA; FNR Université de Montpellier, Fournes Locality collection, France; GMH Geiseltal Museum, Halle, Germany; GPIMUH Geologisch-Paläontologisches Institut und Museum, Universität Hamburg, Germany; GU/RSR/VAS Garhwal University, Srinagar, India; HLMD Hessisches Landesmuseum, Darmstadt, Germany; IIITR/SB/VLM Indian Institute of Technology, Roorkee, India; IRSNB M Institut Royal des Sciences Naturelles de Belgique, Brussels, Belgium; IVPP V Institute of Vertebrate Paleontology and Paleoanthropology, Beijing, China; LIEB-PV Laboratorio de Investigaciones en Evolución y Biodiversidad, Universidad Nacional de la Patagonia —San Juan Bosco, Argentina; MNHN Muséum National d'Histoire Naturelle, Paris, France; QM Queensland Museum, Brisbane, Australia; ROM Royal Ontario Museum, Toronto, Canada; SMF Me Senckenberg Museum, Messel Department, Frankfurt, Germany; SMNK Staatliches Museum für Naturkunde, Karlsruhe, Germany; UM Université de Montpellier, Montpellier, France; UNLSNC Universidade Nova de Lisboa, Portugal, Silveirinha new collection; UT University of Tlemcen, Tlemcen, Algeria; UW University of Wyoming Museum, Laramie, Wyoming, USA; YPM-PU Princeton University (collections now housed in the Yale Peabody Museum, New Haven, CT, USA).

In the list below, *denotes original material examined in addition to (otherwise) primary literature, supplementary photographs and/or scan data for all specimens. Families listed alphabetically.

Archaeonycteridae: *Archaeonycteris trigonodon* Revilliod, 1917 Grube Messel near Darmstadt, Germany HLMD 1398-Me 33a, b*, HLMD 16069; *A. brailloni* Russell *et al.*, 1973, Mutigny and Avenay, Marne, France MNHN Louis-410 Mu, MNHN Louis-432 Mu*, MNHN Bn-751 Av*, MNHN Bn-757 Av* (cast SMF 77/262); *A. pollex* Storch & Habersetzer, 1988 Grube Messel near Darmstadt, Germany SMNK Me 982*; *A. relicta* Harrison & Hooker, 2010 Creechbarrow, Dorset, England BMNH M35710/12; ?*A. precursor* Tabuce *et al.*, 2009 Silveirinha Portugal UNLSNC-447*; ?*A. storchi* Smith *et al.*, 2007 Vastan lignite mine, Gujarat, India GU/RSR/VAS 140; cf. *Archaeonycteris* sp. (*sensu* Smith *et al.*, 2012; cf. *Icaronycteris* sp. in Russell *et al.*, 1988) Meudon, France MNHN 16065 Me*; *Protonycteris gunnelli* Smith *et al.*, 2007 Vastan lignite mine, Gujarat, India GU/RSR/VAS 436. **Hassianycteridae:** *Hassianycteris revillioidi* (Russell & Sigé, 1970) Grube Messel near Darmstadt, Germany HLMD 4294 Me 16* (cast SMF 77/269). *H. messelensis* Smith & Storch, 1981, Grube Messel near Darmstadt, Germany HLMD Me 7480*; *H. magna*

Smith & Storch, 1981, Grube Messel near Darmstadt, Germany HLMD Me 7539*; *H. kumari* Smith *et al.*, 2007 Vastan lignite mine, Gujarat, India GU/RSR/VAS 56, GU/RSR/VAS 561; *Cambaya complexus* Bajpai *et al.*, 2005 Vastan lignite mine, Gujarat, India IITR/SB/VLM 435. **Icaronycteridae:** *Icaronycteris index* Jepsen, 1966; Green River Formation, Wyoming, USA, YPM-PU 18150* and UW 21481*; *I. menui* Russell *et al.*, 1973 Mutigny and Avenay, Marne, France MNHN Louis-333*, MNHN Louis-Mu*, MNHN Louis-331*, MNHN Louis-380*, MNHN Louis-384*, MNHN Louis-353*, MNHN Louis-357*, MNHN Louis-117*, MNHN Av7201*, MNHN Av6797*, MNHN Av7200*, MNHN Louis-360 Mu*, MNHN Bn-547 Av*, MNHN Bn-340 Av*, MNHN Bn-319 Av* (SMF 77/259-261); *I. sigei* Smith *et al.*, 2007 Vastan lignite mine, Gujarat, India GU/RSR/VAS 137. **Mixopterygidae:** *Carcinopteryx maximinensis* Maitre *et al.*, 2008 from St-Maximin, Gard Phosphorites, France, UM-SMXCm A.1.3*, UM-SMXCm A.1.5*. **Onychonycteridae:** *Onychonycteris finneyi* Simmons *et al.*, 2008 Green River Formation, Wyoming, USA, ROM 55351A,B*, AMNH FM 142467*; *Ageina tobieni* Russell *et al.*, 1973 Mutigny, Marne, France MNHN Mu 5112*, MNHN Louis-481 Mu* (cast SMF 77/263); *Eppsinycteris anglica* (Cooper, 1932) Abbey Wood, London Basin, England BMNH M13776; *Honrovits tsuwave* Beard *et al.*, 1992 Wind River Formation, Wyoming, USA

CM62640*, CM 62641*; *?Honrovits joeli* (Smith & Russell, 1992) Evere, Belgium IRSNB M 1567 (cast); *Marnenycteris michauxi* Hand *et al.*, 2015 Pourcy, France UM POY-11. **Palaeochiropterygidae:** *Palaeochiropteryx tupaiodon* Grube Messel near Darmstadt, Germany HLMD Me25* (cast SMF 77/271 and 272), Me 266* (cast SMF 77/273 and 274); *Palaeochiropteryx spiegeli* Grube Messel near Darmstadt, Germany HLMD Me32 * (cast SMF 77/270); *Cecilionycteris prisca* Heller, 1935 Geiseltal Obere Mittelkohle, Germany, GMH 3965*; *Matthesia germanica* Sigé & Russell, 1980, Geiseltal Obere Mittelkohle, Germany, GMH 3940*; *Lapichiropteryx xiei* Tong, 1997 Tuqiaogou, Shanxi province, China IVPP V10204; *Microchiropteryx foliae* Smith *et al.*, 2007 Vastan lignite mine, Gujarat, India GU/RSR/VAS 459. **Philisidae:** *Dizzya exsultans* Sigé, 1991 Chambi, Tunisia CB 1-15*; *Witwavia schlosseri* Gunnell *et al.*, 2008 Fayum, Egypt; *Philisia sphingis* Sigé, 1985 Fayum, Egypt. **Family incertae sedis:** *Australonycteris clarkae* Hand *et al.*, 1994 Murgoon, Queensland, Australia QM F19147, QM F19149*; *Jaegeria cambayensis* Bajpai *et al.*, 2005 Vastan lignite mine, Gujarat, India IITR/SB/VLM/585; Chiropteran (Tejedor *et al.*, 2005) Laguna Fría, Chubut Province, Argentina, LIEB-PV 999; Chiroptera indet. 2 (Marandat, 1991) Fournes, Minervois, France FNR-02*; Eochiroptera (Ravel *et al.* 2011) El Kohol, Algeria UT-KD-02, UT-KD-03.