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Yoh, Natalie and Syme, Peter and Rocha, Ricardo and Meyer, Christoph F. J. and López-Baucells, Adrià (2020) Echolocation of Central Amazonian ‘whispering’ phyllostomid bats: call design and interspecific variation. *Mammal Research*, 65 (3). pp. 583-597. ISSN 2199-2401.

DOI

<https://doi.org/10.1007/s13364-020-00503-0>

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Echolocation of Central Amazonian ‘whispering’ phyllostomid bats: call design and interspecific variation

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Received: 13 September 2019 / Accepted: 30 April 2020 / Published online: 30 May 2020
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Abstract

Phyllostomids (New World leaf-nosed bats) are the ecologically most diverse bat family and have undergone the most extensive adaptive radiation of any mammalian family. However comprehensive, multi-species studies regarding phyllostomid echolocation are scarce in the literature despite abundant ecological research. In this study, we describe the call structure and interspecific variation in call design of 40 sympatric phyllostomid species from the Central Brazilian Amazon, focussing on general patterns within genera, subfamilies and between feeding guilds. All but one species utilized short, broadband FM calls consisting of multiple harmonics. As reported for other bat families, peak frequency was negatively correlated with body mass and forearm length. Twenty-five species alternated the harmonic of maximum energy, principally between the second and third harmonic. Based on PCA, we were unable to detect any significant differences in echolocation call parameters between genera, subfamilies or different feeding guilds, confirming that acoustic surveys cannot be used to reliably monitor these species. We present *Ametrida centurio* as an exception to this generalized phyllostomid structure, as it is unique in producing a mono-harmonic call. Finally, we discuss several hypotheses regarding the evolutionary pressures influencing phyllostomid call structure.

Keywords Bioacoustics · Chiroptera · Echolocation · Neotropical bats · Phyllostomidae

Introduction

The family Phyllostomidae is not only the second largest bat family, consisting of approximately 62 extant genera and 219 species, but it also remains unrivalled amongst mammalian families for its ecological and morphological radiation (Rex et al. 2011; Monteiro and Nogueira 2011; Simmons and Cirranello 2019). Intricate partitioning of resources has cultivated

sophisticated dietary specialization within the family which includes frugivory, folivory, insectivory, carnivory and nectarivory (Aguirre et al. 2003; Monteiro and Nogueira 2011; Rocha et al. 2017a), as well as the only three sanguivorous mammalian species (Zepeda Mendoza et al. 2018).

In the Central Brazilian Amazon, there are at least 84 known species of phyllostomids (López-Baucells et al. 2016). Many of these provide ecosystem services such as

Communicated by: Zuzanna Halat,

Electronic supplementary material The online version of this article (<https://doi.org/10.1007/s13364-020-00503-0>) contains supplementary material, which is available to authorized users.

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pollination, seed dispersal and arthropod suppression (Kunz et al. 2011), which are crucial to maintaining ecosystems and habitats that are under ever-increasing pressure from anthropogenic activities. Although mist netting is most often used to survey tropical bats, acoustic methods are being increasingly employed to sample and monitor aerial-hawking insectivorous species. This is due to the fact that acoustic sampling complements traditional capture techniques, which alone are insufficient for fully inventorying local species assemblages (Flaquer et al. 2007; MacSwiney et al. 2008; Pech-Canche et al. 2011; Rydell et al. 2002; Silva and Bernard 2017). The use and capabilities of bioacoustic surveys have advanced greatly over the past few decades (Clement et al. 2014; Deichmann et al. 2018), and sound is now utilized in the research of many taxonomic groups such as birds, amphibians, marine mammals and even plants (Deichmann et al. 2018; Jung et al. 2018). However, this technique has not been commonly applied to phyllostomid species which have been deemed unsuitable candidates for acoustic monitoring (Brinkløv et al. 2009; Fenton et al. 1992).

Whereas aerial-hawking insectivores rely predominantly on echolocation to forage, phyllostomids typically have more sophisticated sight and olfactory senses and have been shown to be less reliant on echolocation to locate food resources (Brinkløv et al. 2011; Korine and Kalko 2005; Fleming 1989). Most phyllostomids produce low-intensity calls (Brinkløv et al. 2009), and how species utilize them for foraging is not fully understood. However, there are several studies suggesting that sound may be an important means by which frugivorous species can quickly discriminate between desired floral species, e.g. fruit, within other vegetation (Brinkløv et al. 2009; Gonzalez-Terrazas et al. 2016; Korine and Kalko 2005). Echolocation call structure (including parameters such as duration, peak frequency, bandwidth and intensity) is closely associated with the affiliation to different types of food resources and the characteristics of the environment in which they are present (Denzinger and Schnitzler 2013). Bats foraging on aerial insects in open areas (e.g. above the canopy) utilize long, low-frequency, quasi-constant frequency (QCF) calls with narrow bandwidths for maximizing prey detection range but which provide minimal environmental information (Denzinger and Schnitzler 2013). In contrast, the majority of bats which glean insects within dense vegetation use either short, broadband, low-intensity frequency-modulated (FM) calls or almost entirely constant-frequency (CF) calls, which maximize environmental information, but attenuate quickly (Jennings et al. 2004; Russo et al. 2018).

Phyllostomids across different feeding guilds utilize calls of a similar structure, characterized as short, broadband, FM calls consisting of multiple harmonics (Brinkløv et al. 2009; Jennings et al. 2004). Divergence in nose leaf structure has been shown to be linked with dietary specialization within this group (Leiser-Miller and Santana 2020); however, this

morphologic variation and the associated variation in call structure is likely limited by phylogenetic constraints (Collen 2012). Therefore, despite their dietary diversity, phyllostomids rely on similar foraging strategies (Giannini and Kalko 2004)—typically gleaning their food from densely cluttered vegetation. The low intensity and high directionality of their echolocation calls make them difficult to record, and most studies to date have focussed on one or few species, leaving many phyllostomid calls undescribed (Arias-Aguilar et al. 2018; Jennings et al. 2004).

These difficulties, and therefore the limited capacity for bioacoustics to be used in their monitoring, have led to a research deficit concerning phyllostomid echolocation. Here, we address this information gap by describing and comparing the echolocation calls of 40 sympatric phyllostomid species (24 genera) from the Central Brazilian Amazon. We describe the interspecific variation within—and between—genera and subfamilies, as well as explore the influence of body size and feeding guild on call structure.

Material and methods

Study area

This study was carried out at the Biological Dynamics of Forest Fragments Project (BDFFP) ~ 80 km north of Manaus in the Central Amazon (2°20' S, 60°6' W; 80–160 m a.s.l.), Brazil. This reserve was established in the 1980s, with the main goal to investigate the effects of forest fragment size on Neotropical biota (Bierregaard and Lovejoy 1990). To do so, fragments of 1, 10 and 100 ha were isolated from continuous *terra firme* rainforest by distances of 80–650 m. Fragments are nowadays surrounded by a matrix of secondary forest at varying successional stages (Carreiras et al. 2014). Primary forest reaches 30–37 m in mean canopy height, with isolated trees up to 55 m tall (Laurance et al. 2011). Rainfall varies from 1900 to 3500 mm annually, with a dry season between July and November and a rainy season between November and June (Ferreira et al. 2017).

Mist netting

Capture sessions were carried out between August 2011 and October 2014, using both ground- and canopy- level mist nets placed in a variety of habitats: continuous primary forest, forest fragments and secondary forest in which standardized surveys were conducted and temporary lakes, rivers, streams and clearings where we sampled opportunistically (Farneda et al. 2015; Silva et al. 2020; Rocha et al. 2019; Torrent et al. 2018). Mist nets were exposed for six continuous hours after sunset. We identified all individuals using available field guides and morphological keys (López-Baucells et al. 2016). Taxonomy

follows López-Baucells et al. (2016) except for *Mimon crenulatum* and *Lonchophylla thomasi*, which are referred to as *Gardnerycteris crenulatum* and *Hsunycteris thomasi* respectively (Parlos et al. 2014; Hurtado et al. 2014). Standard morphological measurements (including forearm length and body mass) and demographic data were collected following Handley et al. (1991). Details relating to the habitat characteristics of where species were captured (e.g. vertical stratum) can be sourced from Farneda et al. (2015), Silva et al. (2020), Rocha et al. (2019) and Torrent et al. (2018).

Sound recordings

Bats were recorded with a Pettersson D1000X bat detector (Pettersson Elektronik, Sweden) upon hand release. The detector was set to full spectrum, continuous recording (no trigger) and a sampling rate of 384 kHz. The detector was placed 5–10 m from the point of release (depending on the species), and once alight, we followed the bat's flight path for as long as possible.

Acoustic measurements

Sound files were analysed using Kaleidoscope V4.0.4 (Wildlife Acoustics, USA). Sound files consisting of low-quality recordings, or those with multiple species present, were discarded. Moreover, we discarded pulses recorded just after release due to the obvious stress signature (see Supplementary Fig. A.1). For each of the remaining files, ten pulses were identified for analysis. This was not possible for six species which had limited recordings (Table 1). In these cases, the maximum number of pulses obtained was analysed. Measurements were extracted from each of these pulses for six acoustic parameters on each harmonic present: the frequency of maximum energy (peak frequency), the frequency at the start of the harmonic (start freq.), the frequency where the harmonic ends (end freq.), the maximum frequency detected within a harmonic (maximum freq.), the minimum frequency detected within a harmonic (minimum frequency) and the duration in milliseconds between the start and end frequencies (pulse duration) (Fig. 1). Peak frequency was determined from the power spectrum, whereas other variables were measured manually from the spectrogram plots by comparing intensity. Maximum and minimum frequency and start and end frequency were considered 20 dB below peak. Furthermore, the peak frequency across harmonics was compared within each pulse to identify the harmonic of maximum energy. Kaleidoscope settings were as follows: FFT size 256, window size 128 and cache size 256 MB.

Due to pulse overlap, associated with the multi-harmonic nature of the echolocation calls produced by the species targeted by this study, it was neither possible to use an automated pulse-measuring software (e.g. Scan'R) nor to discern

call parameters such as start/end frequency and duration from the oscillogram. Although these approaches are preferable for measuring call characteristics as they reduce variability due to the analyst's bias present when parameters are measured via a sonogram, we believe that our methodology was well suited for our purposes and that any potential variability in the measurements did not substantially affect the conclusions drawn here. However, the call measurements we present here should be viewed in light of these measuring limitations.

Statistical analysis

Linear regression analysis, performed in R 3.4.4 (R Core Team 2018), was used to assess the relationship between body size and peak frequency. For this, the peak frequency of the third harmonic was regressed against the species' mean body mass and forearm length. The analysis was restricted to 23 species for which we had body mass and forearm length data based on our mist net captures. The third harmonic was used as it was the predominant harmonic of maximum energy for 22 of the 38 species included in the analysis. No echolocation calls of sufficient quality were obtained for *Lophostoma schulzi*. Therefore, it was excluded from the analysis, and the regression was performed for 22 species.

Principal component analysis (PCA) was conducted on the aforementioned call parameters measured on the third harmonic of each species' calls, using the R package 'stats', to summarize and visualize the similarity in echolocation call structure of all subfamilies. Log call variables were zero centred and were scaled to have unit variance.

Results

Call descriptions

Across the study period, 40 phyllostomid species from five subfamilies were captured. Acoustic analysis was conducted using a total of 2112 pulses from 135 individuals (Table 1). Sonograms were produced for all species except *Lophostoma schulzi*, as no representative release echolocation call of sufficient quality was obtained for this species. All species, apart from *Ametrida centurio*, produced multi-harmonic calls which were visible on the raw recordings. These were characterized as broadband, downward-sweeping, frequency-modulated calls with overlapping harmonic components. The frequency of maximum energy was almost exclusively located either in the second or third harmonic (88% of all recorded pulses). They also provided the best quality calls for analysis. Harmonics higher than the third were rare and were considered the harmonic of maximum energy for only 2% of recorded pulses. Moreover, these harmonics were extremely faint or corrupted (e.g. higher harmonics were

Table 1 Call parameter values (mean \pm standard deviation, range in brackets) for each harmonic of maximum energy (HME) for the 39 species of the subfamilies Phyllostominae, Glossophaginae, Caroliinae, Stenodermatinae and Desmodontinae, excluding *Lophostoma schulzi*

Subfamily	Species (acronym)	HME	Peak freq. (kHz)	Max freq. (kHz)	Min freq. (kHz)	Start freq. (kHz)	End freq. (kHz)	Duration (ms)	No. of bats recorded	No. of pulses	Primary feeding guild	
Phyllostominae	<i>Lamproncycteris brachyotis</i> (Lambra)	1	40 \pm 2 (37–43)	59	27	47 \pm 3 (43–53)	33 \pm 3 (30–38)	1.7 \pm 0.2 (1.3–2.0)	1	9	Insectivore	
		2	81 \pm 8 (72–86)	105	61	98 \pm 5 (92–101)	66 \pm 4 (62–69)	1.8 \pm 0.3 (1.5–2.0)	-	8		
	<i>Trinycteris nicefori</i> (Trinic)	2	57 \pm 6 (45–64)	77	37	67 \pm 5 (60–73)	41 \pm 2 (37–43)	2.6 \pm 0.6 (1.5–3.4)	3	26	Insectivore	
		3	73 \pm 2 (70–76)	100	57	92 \pm 3 (85–98)	64 \pm 3 (61–70)	1.7 \pm 0.4 (1.0–2.5)	-	30		
	<i>Lophostoma silvicolum</i> (Lopsil)	2	64 \pm 2 (61–67)	85	42	72 \pm 6 (64–85)	46 \pm 2 (44–50)	1.6 \pm 0.2 (1.1–2.0)	2	20	Insectivore	
	<i>Lophostoma brasiliense</i> (Lopbra)	3	91 \pm 5 (86–100)	116	72	105 \pm 4 (100–113)	79 \pm 2 (74–82)	1.9 \pm 0.2 (1.5–2.3)	2	20	Insectivore	
	<i>Lophostoma carrikeri</i> (Lopcar)	2	59 \pm 2 (55–61)	72	38	65 \pm 2 (60–68)	46 \pm 3 (43–53)	1.8 \pm 0.3 (1.3–2.3)	1	10	Insectivore	
	<i>Tonatia saurophila</i> (Tonsau)	2	43 \pm 1 (41–45)	54	27	47 \pm 2 (43–51)	33 \pm 2 (30–37)	1.8 \pm 0.5 (1.2–2.6)	4	36	Insectivore	
		3	64 \pm 4 (59–70)	84	47	74 \pm 3 (65–79)	54 \pm 3 (47–60)	1.6 \pm 0.3 (1.0–2.0)	-	37		
	<i>Phylloderma stenops</i> (Physte)	4	85 \pm 4 (80–90)	105	65	96 \pm 4 (91–101)	72 \pm 5 (67–78)	1.5 \pm 0.3 (1.2–1.9)	-	31		
		5	102 \pm NaN (-)	111	89	108 \pm NaN (-)	90.5 \pm NaN (-)	1.2 \pm NaN (-)	-	13	Insectivore	
	Glossophaginae	<i>Gardnerycteris crenulatum</i> (Garere)	2	52 \pm 1 (51–52)	61	36	58 \pm 1 (57–58)	37.8 \pm 2 (37–39)	1.7 \pm 0.1 (1.6–1.7)	3	24	Insectivore
			3	64 \pm 2 (61–68)	87	47	77 \pm 4 (68–87)	54 \pm 3 (47–61)	1.5 \pm 0.3 (1.0–2.0)	-	30	
		<i>Phyllostomus discolor</i> (Phydis)	3	69 \pm 3 (65–76)	91	52	82 \pm 3 (76–88)	60 \pm 2 (54–64)	2 \pm 0.4 (1.0–2.9)	4	40	Insectivore
			2	41 \pm 3 (36–44)	53	27	47 \pm 1 (45–50)	31 \pm 3 (27–36)	4.1 \pm 1.6 (1.4–5.9)	5	50	Insectivore
<i>Phyllostomus elongatus</i> (Phyelo)		3	62 \pm 2 (59–69)	88	38	75 \pm 3 (69–80.0)	48 \pm 3 (43–56)	2.7 \pm 0.5 (1.8–3.8)	-	49		
		4	74 \pm 2 (72–75)	94	60	88 \pm 2 (87–89)	65 \pm 2 (63–67)	2 \pm 0.5 (1.7–2.4)	-	37		
<i>Phyllostomus hastatus</i> (Phyhas)		3	61 \pm 2 (57–66)	76	47	68 \pm 3 (61–73)	52 \pm 2 (48–57)	1.9 \pm 0.4 (1.3–3.3)	5	50	Insectivore	
		4	75 \pm 2 (72–76)	94	57	88 \pm 4 (81–92)	61 \pm 2 (57–64)	1.9 \pm 0.3 (1.5–2.2)	-	49		
<i>Chrotopterus auritus</i> (Chraur)		3	47 \pm 2 (43–52)	60	34	54 \pm 2 (51–58)	40 \pm 3 (35–44)	2.3 \pm 0.5 (1.5–3.3)	3	30	Insectivore	
		2	77 \pm 1 (76–78)	88	67	83 \pm 3 (76–86)	71 \pm 3 (68–75)	1.1 \pm 0.2 (0.8–1.4)	4	25	Insectivore/carnivore	
<i>Glyptonycteris daviesi</i> (Glydav)		3	91 \pm 4 (84–102)	117	73	110 \pm 5 (98–117)	79 \pm 3 (74–85)	2.2 \pm 0.5 (1.4–2.9)	-	40		
		3	60 \pm 3 (57–67)	80	44	65 \pm 4 (60–73)	50 \pm 4 (44–57)	1.2 \pm 0.2 (0.8–1.4)	1	10	Insectivore/carnivore	
<i>Micronycteris hirsuta</i> (Michir)		2	52 \pm 1 (51–55)	74	42	63 \pm 3 (59–68)	48 \pm 2 (43–51)	1.3 \pm 0.2 (1.0–1.8)	3	25	Insectivore	
		3	73 \pm 5 (69–82)	101	46	88 \pm 3 (84–94)	51 \pm 6 (46–61)	3.0 \pm 0.8 (1.3–3.9)	2	16	Insectivore	
<i>Trachops cirrhosus</i> (Tracir)		2	98 \pm 3 (94–104)	131	80	115 \pm 6 (104–125)	85 \pm 3 (83–91)	1.4 \pm 0.1 (1.3–1.5)	-	16		
	2	61 \pm 4 (53–72)	90	37	84 \pm 5 (69–90)	46 \pm 3 (37–52)	1.9 \pm 0.5 (1.1–3.5)	5	49	Insectivore/carnivore		
<i>Glossophaga soricina</i> (Glosor)	2	81 \pm 3 (76–86)	107	34	83 \pm 4 (78–90)	45 \pm 6 (51–55)	3.0 \pm 0.3 (2.6–3.5)	1	10	Nectarivore		
	3	74 \pm 2 (72–76)	95	51	83 \pm 4 (79–87)	52 \pm 2 (51–55)	2.3 \pm 0.4 (1.9–2.6)	2	10	Nectarivore		
<i>Choeroniscus minor</i> (Chomin)	4	99 \pm 4 (88–104)	134	74	121 \pm 5 (113–130)	84 \pm 6 (73–97)	1.7 \pm 0.5 (1.1–2.6)	-	20			
	1	41 \pm 5 (36–51)	68	28	61 \pm 2 (59–64)	32 \pm 3 (28–35)	2 \pm 0.3 (1.6–2.4)	1	10	Nectarivore		
	3	105 \pm 7 (100–113)	159	87	149 \pm 11 (137–159)	94 \pm 7 (87–102)	2.1 \pm 0.3 (1.8–2.2)	-	9			

Table 1 (continued)

Subfamily	Species (acronym)	HME	Peak freq. (kHz)	Max freq. (kHz)	Min freq. (kHz)	Start freq. (kHz)	End freq. (kHz)	Duration (ms)	No. of bats recorded	No. of pulses	Primary feeding guild
Carollinae	<i>Carollia brevicauda</i> (Carbre)	2	69 ± 1 (66–70)	95	39	79 ± 3 (74–84)	50 ± 6 (39–57)	1.9 ± 0.4 (1.2–2.4)	4	38	Frugivore
		3	86 ± 6 (78–101)	118	62	99 ± 18 (117–118)	72 ± 5 (62–81)	1.4 ± 0.2 (1.0–1.8)	-	40	
	<i>Carollia castanea</i> (Carcas)	3	81 ± 1 (80–82)	107	64	100 ± 4 (93–104)	69 ± 4 (65–76)	2.2 ± 0.3 (1.8–2.7)	1	10	Frugivore
Carollinae	<i>Carollia perspicillata</i> (Carper)	2	69 ± 3 (59–76)	91	47	85 ± 4 (75–91)	52 ± 3 (47–58)	1.7 ± 0.3 (1.2–2.2)	5	50	Frugivore
		3	92 ± 6 (82–100)	116	71	110 ± 4 (102–116)	75 ± 4 (71–86)	1.6 ± 0.3 (1.2–2.1)	-	46	
	<i>Rhinophylla pumilio</i> (Rhipum)	2	67 ± 6 (57–82)	99	41	87 ± 7 (76–99)	51 ± 5 (41–61)	1.6 ± 0.3 (0.9–2.1)	5	49	Frugivore
Stenodermatinae	<i>Artibeus planirostris</i> (Artpla)	3	91 ± 4 (86–102)	127	57	112 ± 7 (93–127)	73 ± 6 (57–84)	1.5 ± 0.4 (1.1–2.4)	-	48	
		2	52 ± 5 (43–63)	81	35	65 ± 7 (53–76)	43 ± 4 (35–49)	1.9 ± 0.5 (1.3–3.5)	6	60	Frugivore
		3	68 ± 5 (59–78)	101	46	87 ± 7 (73–101)	55 ± 5 (46–70)	1.8 ± 0.5 (1.1–3.2)	-	57	
Stenodermatinae	<i>Artibeus lituratus</i> (Artlit)	2	65 ± 6 (49–78)	94	32	77 ± 6 (57–90)	53 ± 6 (36–66)	1.7 ± 0.4 (0.8–3.0)	10	99	Frugivore
		2	53 ± 2 (51–57)	78	38	68 ± 4 (66–75)	46 ± 6 (40–58)	1.7 ± 0.2 (1.3–2.1)	8	58	Frugivore
	<i>Artibeus obscurus</i> (Artobs)	3	70 ± 5 (59–84)	103	42	87 ± 7 (70–98)	57 ± 6 (45–76)	1.5 ± 0.3 (1.0–2.5)	-	80	
Stenodermatinae	<i>Artibeus concolor</i> (Artcon)	4	88 ± 4 (78–94)	119	59	107 ± 4 (100–114)	66 ± 6 (59–75)	1.7 ± 0.3 (1.0–2.1)	-	50	
		2	95 ± 6 (78–104)	129	65	111 ± 6 (99–126)	79 ± 7 (67–125)	1.7 ± 0.4 (1.1–2.8)	9	90	Frugivore
	<i>Vampyriscus brocki</i> (Vambro)	2	69 ± 6 (64–88)	101	53	83 ± 6 (73–93)	60 ± 3 (56–65)	1.5 ± 0.4 (1.0–2.5)	2	19	Frugivore
Stenodermatinae	<i>Vampyriscus bidens</i> (Vambid)	3	107 ± 15 (89.9–115.4)	138	79	122 ± 15 (105–134)	89 ± 1 (88–90)	1.4 ± 0.2 (1.2–1.5)	-	10	
		2	71 ± 3 (69–80)	94	57	88 ± 2 (84–92)	60 ± 2 (57–65)	2.2 ± 0.2 (1.9–2.5)	2	13	Frugivore
		3	101 ± 2 (95.8–103.2)	127	81	118 ± 6 (106–126)	91 ± 5 (85–100)	1.2 ± 0.2 (0.5–1.5)	-	16	
Stenodermatinae	<i>Sturnira tilidae</i> (Stutil)	2	65.0 ± 4.3 (55.1–72.4)	100.4	34.5	81 ± 8 (69–98)	45 ± 4 (37–52)	2.2 ± 0.3 (1.5–3.1)	5	49	Frugivore
		3	84.1 ± 8.7 (75.9–97.7)	132.8	61.6	107 ± 16 (93–128)	70 ± 7 (62–77)	2.0 ± 0.3 (1.7–2.4)	-	47	
	<i>Sturnira lilium</i> (Stulil)	2	66.5 ± 5.0 (60.3–76.2)	90.8	42.2	78 ± 4 (72–84)	50 ± 3 (47–54)	1.7 ± 0.3 (1.2–2.2)	1	10	Frugivore
Stenodermatinae	<i>Mesophylla macconnelli</i> (Mesmac)	1	101 ± 4 (94–106)	123	81	112 ± 4 (106–119)	86 ± 4 (81–94)	1.6 ± 0.2 (1.3–2.0)	1	9	Frugivore
		2	118 ± 2 (117–119)	145	97	140 ± 0.0 (-)	103 ± 1 (103–104)	1.9 ± 0.0 (1.9–1.9)	-	7	
	<i>Dermanura cinereus/gnomus</i> (Dercingno)	1	51 ± 9 (35–61)	91	25	66 ± 10 (37–81)	36 ± 8 (25–68)	1.9 ± 0.7 (1.0–3.8)	8	63	Frugivore
Stenodermatinae	<i>Chiroderma trinitatum</i> (Chitri)	2	89 ± 14 (66–102)	135	53	112 ± 9 (95–128)	64 ± 6 (55–91)	1.6 ± 0.3 (1.0–2.8)	-	80	
		3	97 ± 2 (94–100)	138	71	126 ± 6 (113–137)	82 ± 7 (73–94)	2.2 ± 0.5 (1.7–3.3)	2	20	Frugivore
	<i>Anerida centurio</i> (Amece)	1	100 ± 6 (84–116)	199	52	149 ± 25 (114–196)	65 ± 4 (58–80)	2.8 ± 0.6 (1.8–4.6)	5	50	Frugivore
Stenodermatinae	<i>Uroderma bilobatum</i> (Urobil)	2	68 ± 2 (66–70)	91	52	83 ± 3 (81–86)	56 ± 4 (52–59)	2.7 ± 0.9 (2.1–3.7)	2	6	Frugivore
		3	90 ± 4 (86–96)	121	62	109 ± 5 (96–116)	78 ± 5 (65–85)	1.6 ± 0.4 (1.0–2.3)	-	20	
	<i>Desmodus rotundus</i> (Destot)	1	33 ± 0.0 (-)	45	23	42 ± 2 (40–43)	24 ± 2 (23–25)	1.8 ± 0.1 (1.8–1.9)	4	18	Sanguivore
Stenodermatinae		2	61 ± 5 (53–70)	84	43	77 ± 3 (72–84)	49 ± 3 (43–55)	1.7 ± 0.4 (1.1–2.9)	-	40	
		3	83 ± 7 (71–90)	111	66	105 ± 7 (94–111)	67 ± 2 (66–72)	1.8 ± 0.5 (1.4–2.8)	-	27	

The predominant diet of each species was used to assign each species to a feeding guild. *Dermanura cinereus* and *D. gnomus*, as well as *Micronycteris microtis* and *M. megalotis*, were grouped together as these species are often indistinguishable from one another. Call parameter values for these species were averaged. Variation may exceed parameter ranges listed here due to differences in recording situation or environmental conditions

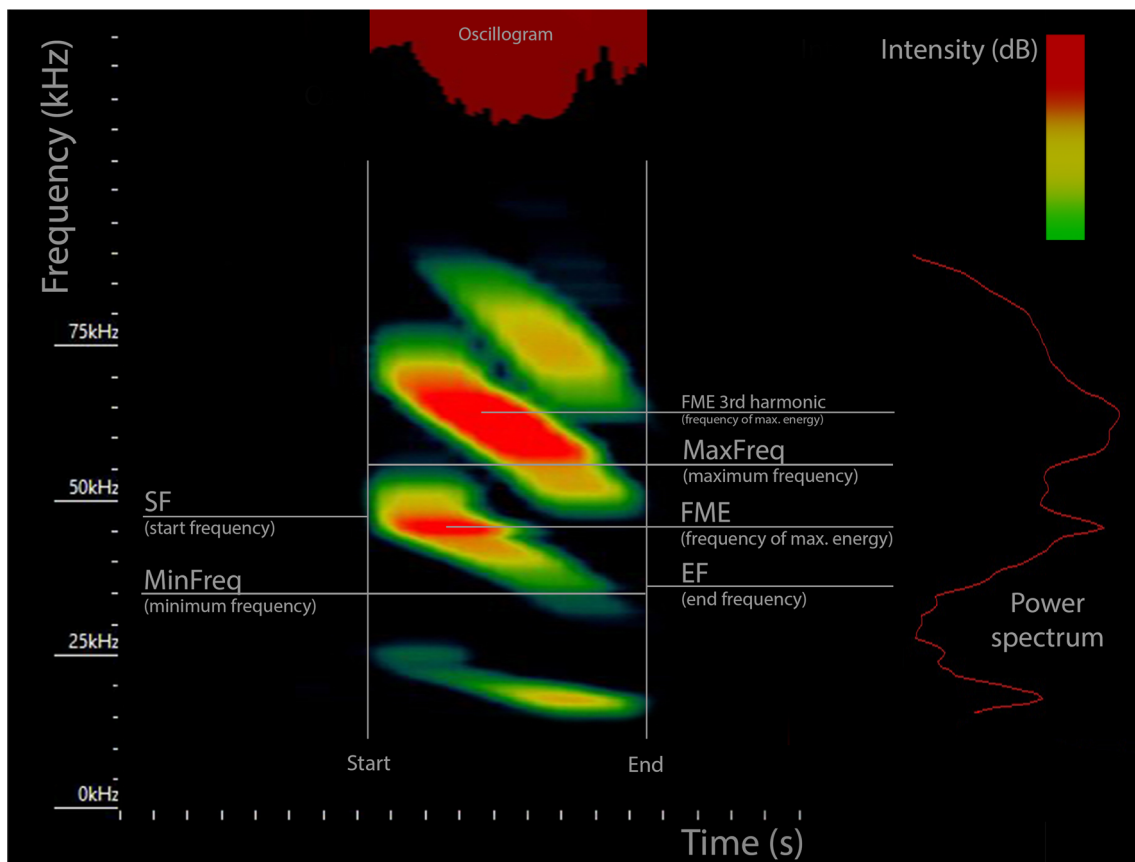


Fig. 1 Illustration of the points of measurement for call parameters used in this study in Kaleidoscope, including *Phyllostomus discolor* call. Peak frequency (FME) was measured from the power spectrum, whereas the

other parameters were measured from the sonogram. FFT size 5112 and win size 128

beyond the frequency range that could be accurately sampled) and therefore could not be measured accurately. Therefore, further call analysis focussed primarily on second and third harmonics. It should be noted that intraspecific variation may exceed parameter ranges due to differences in recording situation or environmental conditions beyond those within our study.

Peak frequency of the third harmonic was significantly inversely related to body size (Fig. 2); large bats, such as *Artibeus planirostris* (average body mass 51.0 g), generally had low-frequency calls, whereas smaller bats, such as *Vampyriscus bidens* (average body mass 12.0 g), exhibited high frequency calls.

Phyllostominae

A total of 17 species from 11 genera of the subfamily Phyllostominae were captured during the study period (Fig. 3). However, no release echolocation call of sufficient quality was obtained for *Lophostoma schulzi* (Fig. 4).

There is a high degree of similarity between the respective call harmonics of the genera *Lamproncycteris* and *Trinycteris* (Table 1). Within the genus *Lophostoma*, call frequencies are

reflective of relative body size as the largest of the three species examined; *L. silvicolum* utilized lower frequency calls than the smaller *L. brasiliense* and *L. carrikeri* (Table 1).

Body size was also negatively correlated with call frequency overall in the genus *Phyllostomus*. Despite being the largest species examined in this study, *Chrotopterus auritus* utilized calls that were higher in frequency than predicted for its body size (see Fig. 2 and Table 1).

Stenodermatinae

A total of 15 stenodermatids were captured during the study period. There was very little variation in the echolocation calls of *Artibeus planirostris*, *A. lituratus* and *A. obscurus*—all produced calls with similar peak frequencies of ~70 kHz on the harmonic of maximum energy (Fig. 3). *Artibeus concolor* (the smallest *Artibeus* examined during this study) emitted calls of comparatively higher frequencies than the larger congeners. However, an overlap occurred between the third harmonic of *A. planirostris* and *A. obscurus* and the second harmonic of the largest species, *A. lituratus*. Because of the overlap between the frequencies of

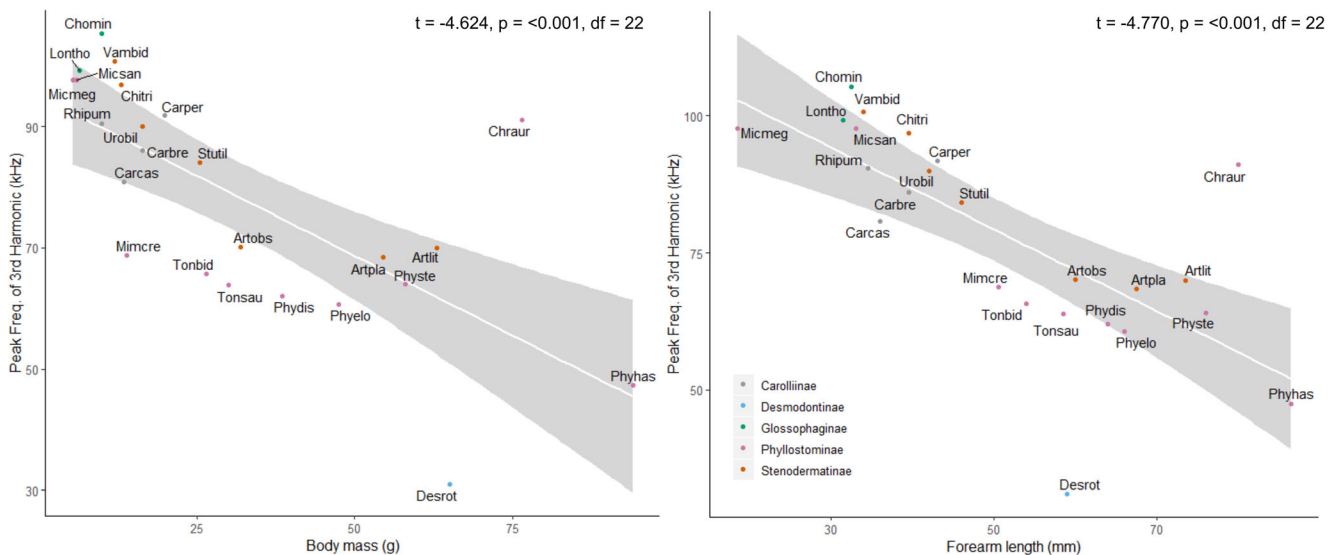


Fig. 2 Relationship between body size, represented by body mass and forearm length, and mean peak frequency of the third harmonic. The third harmonic was used as it was the predominant harmonic of maximum

energy across the species list. Line shows the results of a linear regression for each plot. See Table 1 for species abbreviations. Grey represents 95% confidence level interval

multiple harmonics, it is not possible to reliably distinguish between these species. The genera *Sturnira* and *Vampyriscus* produce calls that are indistinguishable and exhibit high similarity with the calls of *Dermanura*. There is a high degree of overlap between the different harmonics amongst this group (Table 1). The shortest bandwidths and highest frequencies are observed in the smaller species *Platyrrhinus helleri* and *Mesophylla macconnelli*. The species of these six genera produce calls similar to those present in other phyllostomid families. *Ametrida* is a monotypic genus consisting of *A. centurio*. Pulses are frequency-modulated resembling an inverted hockey stick shape, broad bandwidths and short durations. The species shows a high degree of variability within call frequencies (Table 1), particularly concerning the call start and end frequencies. It is also one of the few phyllostomid species to not emit a multi-harmonic call. This, combined with its high peak and maximum frequencies, enables *A. centurio* to be identified using acoustic analysis. *Uroderma bilobatum* also exhibits high frequency calls but overlaps with several species from other genera.

Desmodontinae

Desmodontinae is a monophyletic group comprising the species *Desmodus rotundus*, *Diaemus youngi* and *Diphylla ecaudata*. However, only *D. rotundus* was captured in this study. This species exhibited frequencies of maximum energy on each of the first three of the four visible harmonics (Table 1) and utilized low-frequency

calls for its body size. Calls produced by this species are typical of other phyllostomids in terms of their structure.

Carollinae

Four species from two genera of the subfamily *Carollinae* were captured. Although morphologically similar to the smallest species of the genus *Carollia* (*C. castanea*), *Rhinophylla pumilio* exhibited call frequencies similar to the largest species—*C. perspicillata* (Table 1). Call shape was similar across both genera (Fig. 3) and to that of other phyllostomids. Between *Carollia* species, frequencies reflected their relative body sizes, with larger species emitting echolocation calls of lower frequency.

Glossophaginae

Three species from three genera of the subfamily *Glossophaginae* were captured during the study. Unlike other subfamilies, glossophagines appear to exhibit variation in call shape amongst genera. *Glossophaga soricina* restricts the frequency of maximum energy to the second harmonic, utilized broader bandwidths and produced a distinctively longer call compared with all other phyllostomid species (Table 1). *Choeroniscus minor* exhibited frequencies of maximum energy on the first and third harmonic which were not expressed on the second harmonic.

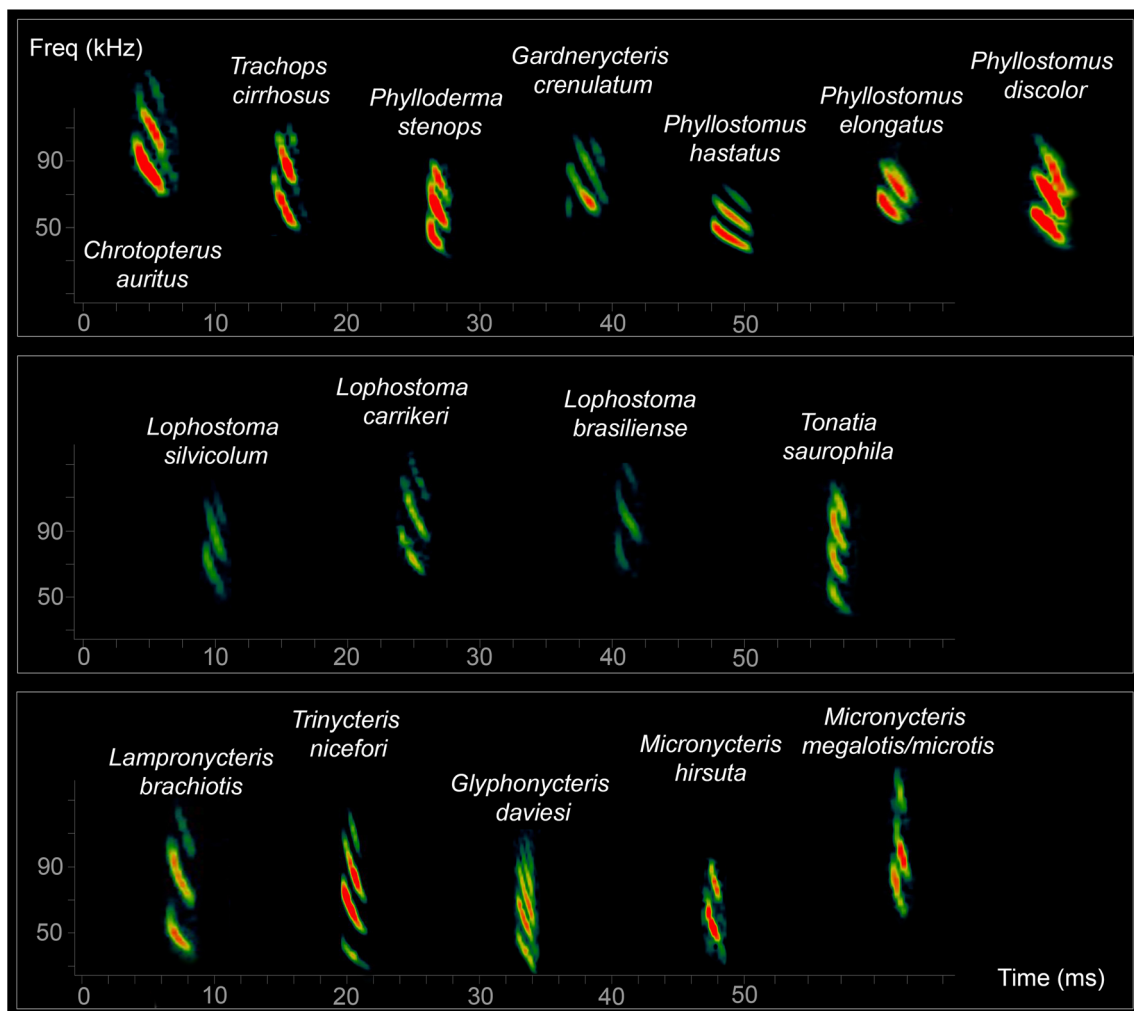


Fig. 3 Echolocation call design of 16 species (11 genera) from the subfamily Phyllostominae. *Lophostoma schulzi* is not included in this figure due to lack of a high-quality sonogram image. *Micronycteris megalotis* and *M. microtis* have been joined due to the difficulty in distinguishing between these two species. Call duration and frequency

have been scaled to allow comparison between species. Sonogram image is for illustrative purposes and should not be used in isolation for species identification. Call amplitude represented here is not standardized between species and therefore should not be used as a diagnostic feature

Call parameter overlap between phyllostomid subfamilies

When all species were considered together, no differentiation in call structure between subfamilies was perceivable (Fig. 5).

Across the five subfamilies, Stenodermatinae and Phyllostominae exhibited the largest variation in call structure amongst species.

Multiple species within and between phyllostomid subfamilies show substantial variation in peak frequency, such as *A. lituratus*, *Sturnira tildae* and *Vampyriscus brocki* (Fig. 6). However, despite the substantial difference between peak frequencies of certain species, almost every species has a large overlap in peak frequency range with multiple other species. The only exception to this is *P. hastatus*, which does not exhibit any overlap in peak frequency range of the third harmonic with any other study species.

Discussion

Our results demonstrate that phyllostomid subfamilies exhibit remarkably similar call structures even when belonging to different feeding guilds. Due to the high degree of overlap in call parameters as revealed by the PCA (Fig. 5), our results corroborate that it is not currently possible to reliably distinguish between most species, genera and subfamilies, based solely on the call parameters used in this study. The total frequency range observed (considering the peak frequencies for all harmonics of maximum energy) ranged from 33.4 kHz (*Desmodus rotundus*) to 118.2 kHz (*Mesophylla macconnelli*). Although some differences between certain subfamilies can be observed (Fig. 6), due to the continuous overlap along the gradient of all species, it is not possible to use acoustic data to reliably identify species or subfamilies.

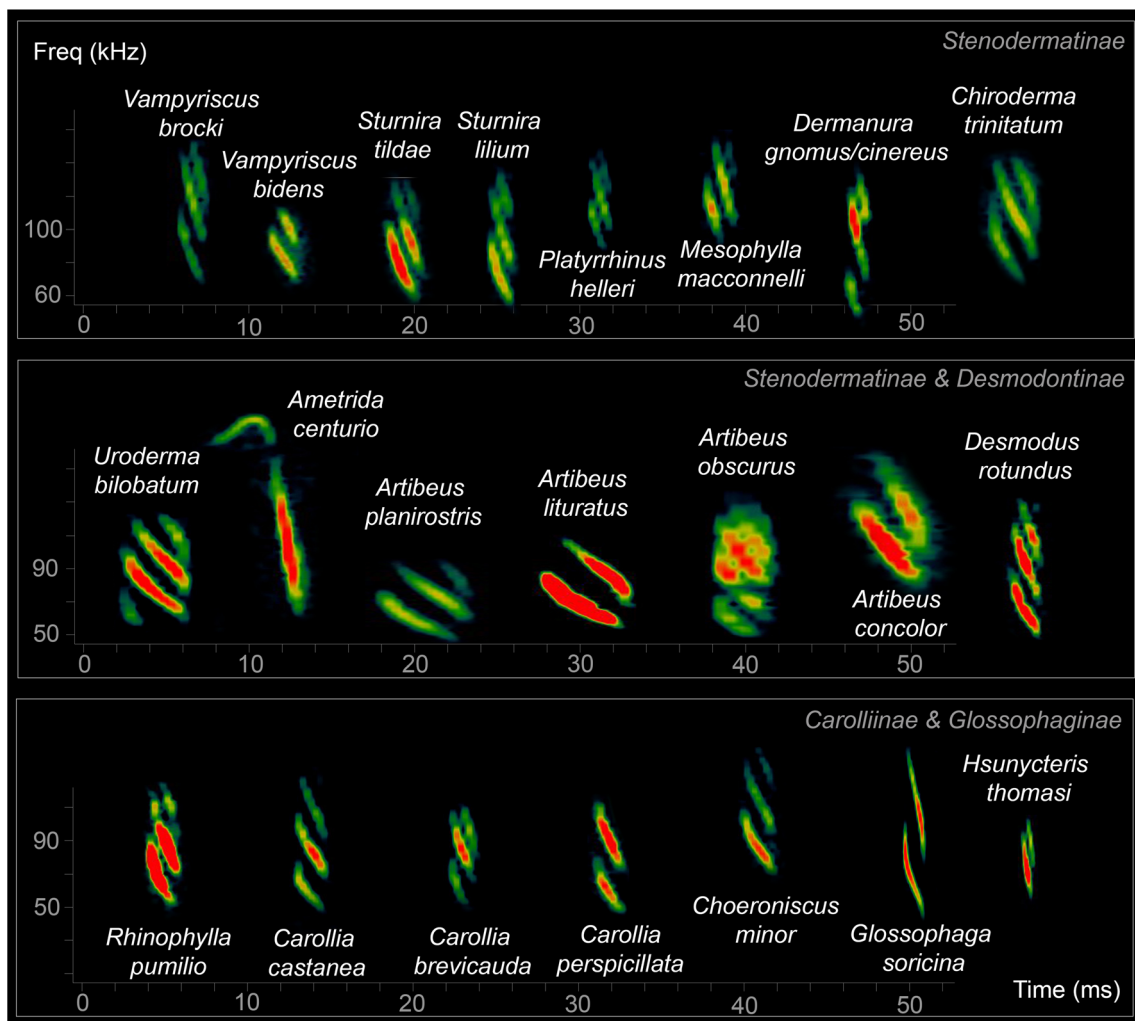


Fig. 4 Echolocation call design of 23 species (15 genera) from the subfamilies Stenodermatinae, Desmodontinae, Carollinae and Glossophaginae. *Dermanura gnomus* and *D. cinereus* are presented together due to the difficulty in distinguishing between these two species. Call duration and frequency have been scaled to allow

comparison between species. Sonogram image is for illustrative purposes and should not be used in isolation for species identification. Call amplitude represented here is not standardized between species and therefore should not be used as a diagnostic feature

Evolutionary advantage for multi-harmonic calls

All but one species (*Ametrida centurio*) produced pulses with multiple harmonic components, the majority of which showed a high degree of spectral overlap. By doing so, the production of frequency-modulated calls which extend their bandwidth possibly enables individuals to perceive a more complete representation of their environment than otherwise possible (Altringham 2011; Krumbholz and Schmidt 2001). This includes providing information on shape, size and texture of neighbouring vegetation. However, as they produce low-intensity calls, detection range is limited.

Phyllostomids are not unique in their use of low-intensity, multi-harmonic calls. The families Nycteridae and Megadermatidae not only exhibit convergent evolution in terms of call structure, but they also possess similarities

with phyllostomids concerning nose structure and foraging strategies (see Fig. 7) (Altringham 2011; Marimuthu et al. 2010; Schaer et al. 2015). Therefore, similar ecological pressures have resulted in similar evolutionary adaptations in both New and Old World species. By comparing wider groups of species assemblages, future research may be able to uncover the determining factors dictating call characteristics. In doing so, this would improve our understanding of the functionality of these characteristics and their purpose and—more broadly—expand our insights into sound ecology.

Species-specific exceptions to the general phyllostomid call structure

The phyllostomine *Ametrida centurio* was the only species which did not produce a multi-harmonic call, which is

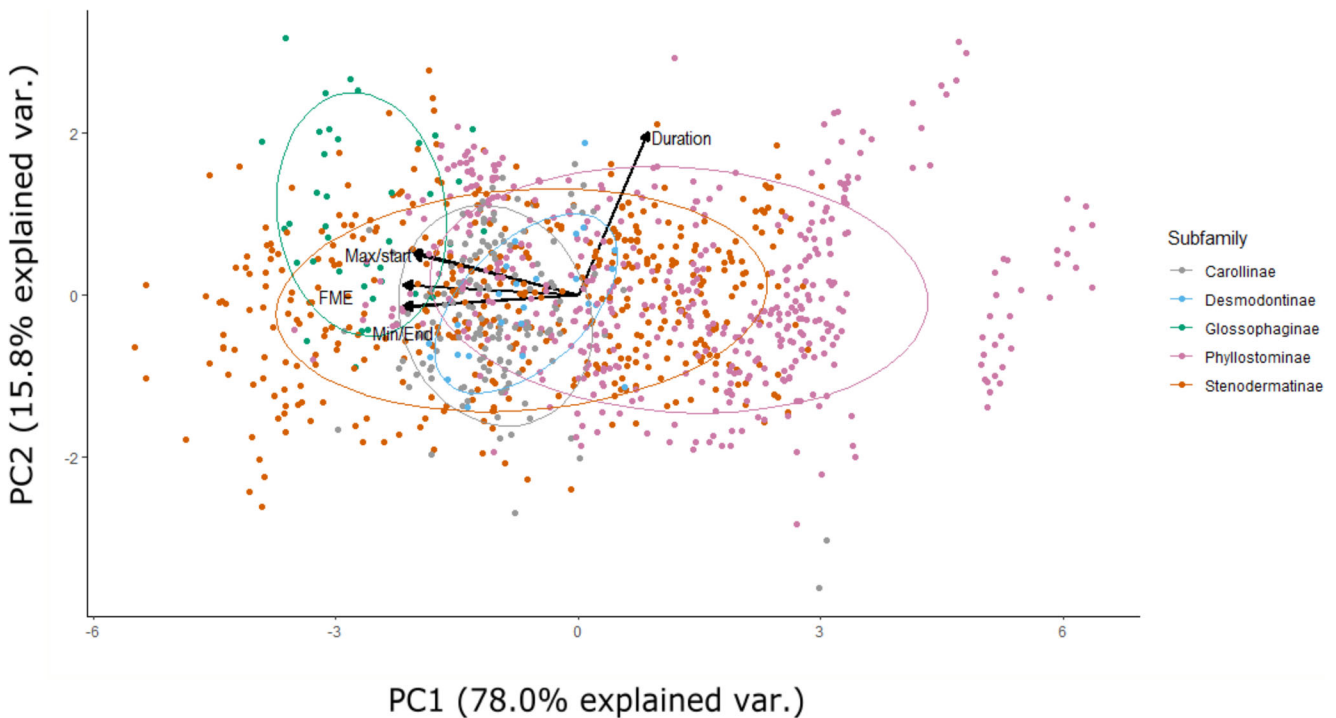


Fig. 5 Arrangement of the 38 phyllostomid species, excluding *L. schulzi*, according to call parameters of the third harmonic along the axes of a principal component analysis. Circles represent 0.68 confidence ellipse

probabilities. FME, frequency of maximum energy (peak frequency). Legend includes variable loadings

consistent with previous reports (Collen 2012). *Ametrida centurio* is considered to be a canopy species which is also associated with forest clearings and waterways (Lee and Dominguez 2000; Rocha et al. 2017b; Silva et al. 2020). It is primarily frugivorous and nectarivorous and favours less cluttered spaces (Lee and Dominguez 2000; Vilar et al. 2015). Being the smallest phyllostomid, it has fewer requirements in terms of manoeuvrability than larger bats (Altringham 2011). It is thus possible that the production of calls with singular harmonics reflects its habitat preferences. *Ametrida centurio* also expressed large variation in call frequency. This could also be a response to environmental requirements and can also be associated with sexual dimorphism in body size (Lee and Dominguez 2000). Further studies which differentiate between the calls of males and females could test this hypothesis. Research regarding the ecology and biology of this species is limited, in part due to its rarity across

its range as well as difficulties in surveying it (Vilar et al. 2015). Due to the unique nature of this species' echolocation, records of occurrence may emerge through future canopy surveys utilizing bioacoustics, subsequently improving our understanding of its ecological requirements. Although not present in our study area, *Lonchorhina aurita* represents another species from the subfamily Phyllostominae whose call Gessinger et al. (2019) recently demonstrated to differ fundamentally from the stereotypical echolocation call design of the family, likely reflecting an adaptation to aerial hawking.

Acoustic partitioning

Due to the degree of overlap between the echolocation call parameters of phyllostomid species, reliably distinguishing between subfamilies, genera and species is impossible using parameters adopted in this study, with few exceptions such as

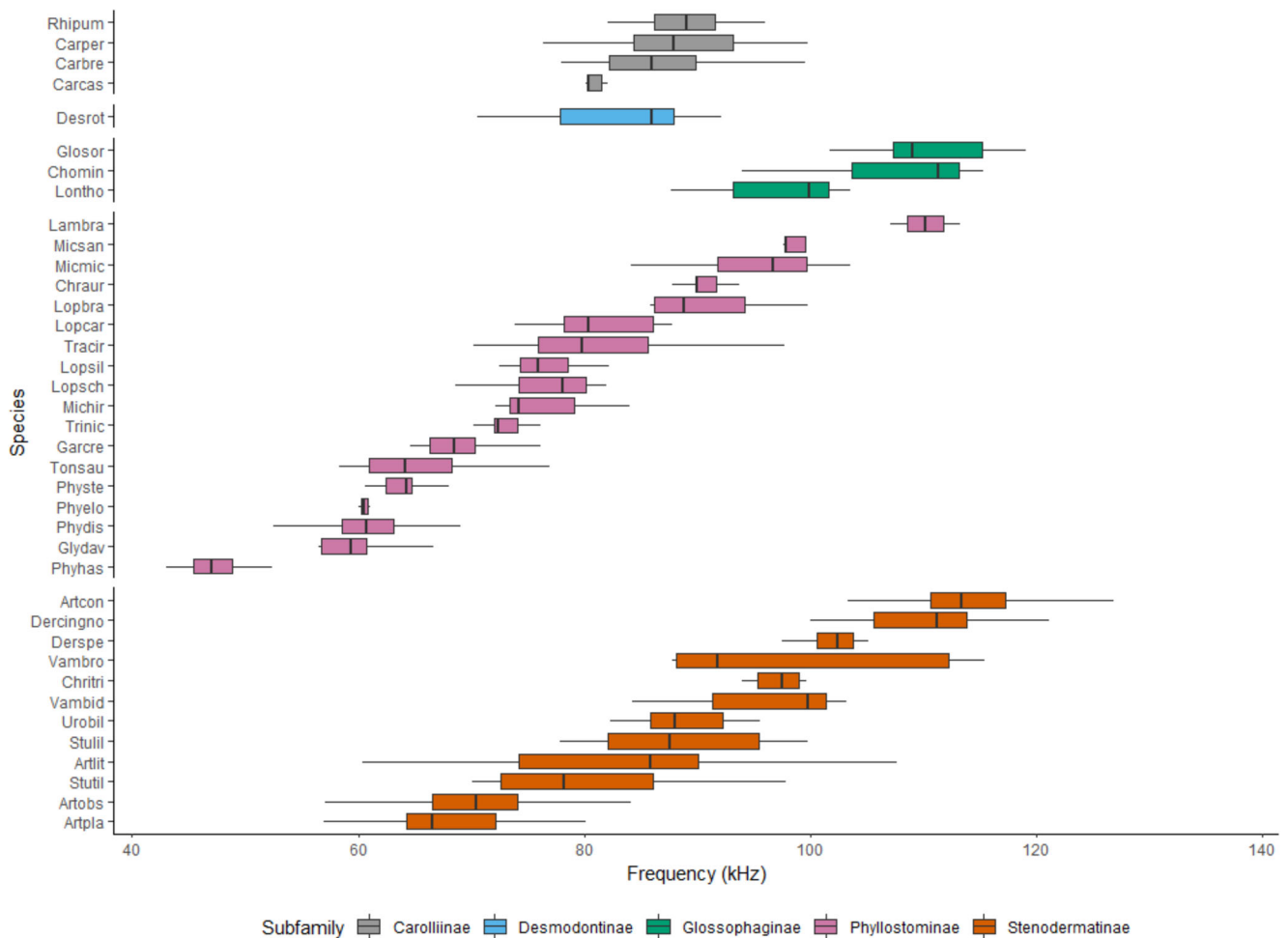


Fig. 6 Peak frequencies observed for the third harmonic of 38 species of the subfamilies Carollinae, Desmodontinae, Glossophaginae, Phyllostominae and Stenodermatinae. See Table 1 for species abbreviations

A. centurio, *P. hastatus* and *L. aurita*. In order to be able to exploit multiple resources, most phyllostomids exhibit a generalist call structure. This allows them to adapt their calls to suit the wide array of environmental conditions encountered and feeding resources exploited (Brinkløv et al. 2009; Fenton et al. 1992; Pio et al. 2010). Nevertheless, substantial differences in peak frequencies distributed across a continuous gradient of variation are evident. Acoustic partitioning between species is usually related to niche differences such as habitat preference and prey selection or to facilitate intraspecific communication (Brinkløv et al. 2009; Denzinger et al. 2018; Denzinger and Schnitzler 2013; Jones and Barlow 2004; Shi et al. 2009). Although multiple feeding guilds were examined in this study, all species considered are gleaners that forage in high-clutter situations and hence utilize broadly similar foraging strategies (Giannini and Kalko 2004). The majority of these species have flexible diets, demonstrating omnivorous selection in response to conditions such as seasonal fluctuations in resource availability (Giannini and Kalko 2004; Santana et al. 2011; Wilkinson and Wenrick Boughman 1998).

Body size is a strong predictor of call frequency for echolocating bat species (see Fig. 2) (Altringham 2011; Jones 1999; Shi et al. 2009). However, this relationship is not consistent amongst phyllostomids (Jones 1999; Russo et al. 2018), and several deviations from this pattern were evident in our data. Although both *C. auritus* and *D. rotundus* produce calls which follow the basic phyllostomid structure (low-intensity, multi-harmonic), it is possible that more specialized diets result in increased selection pressures on acoustic parameters, as observed in *L. aurita* (Gessinger et al. 2019). *D. rotundus* utilized frequencies lower than expected relative to its body size (see Table 1). This is likely due to its unique feeding habits, but whether this is related to foraging strategy or prey type remains unclear. Variation in foraging strategy amongst different species has been observed to affect call structure (Altringham 2011; Denzinger and Schnitzler 2013). However, the vast majority of studies relate to aerial insectivores (but see Halfwerk et al. 2014). *D. rotundus* has an extended forearm length relative to species of the same weight (Altenbach 1979) as it uses its forearm for approaching prey on the ground. It is therefore

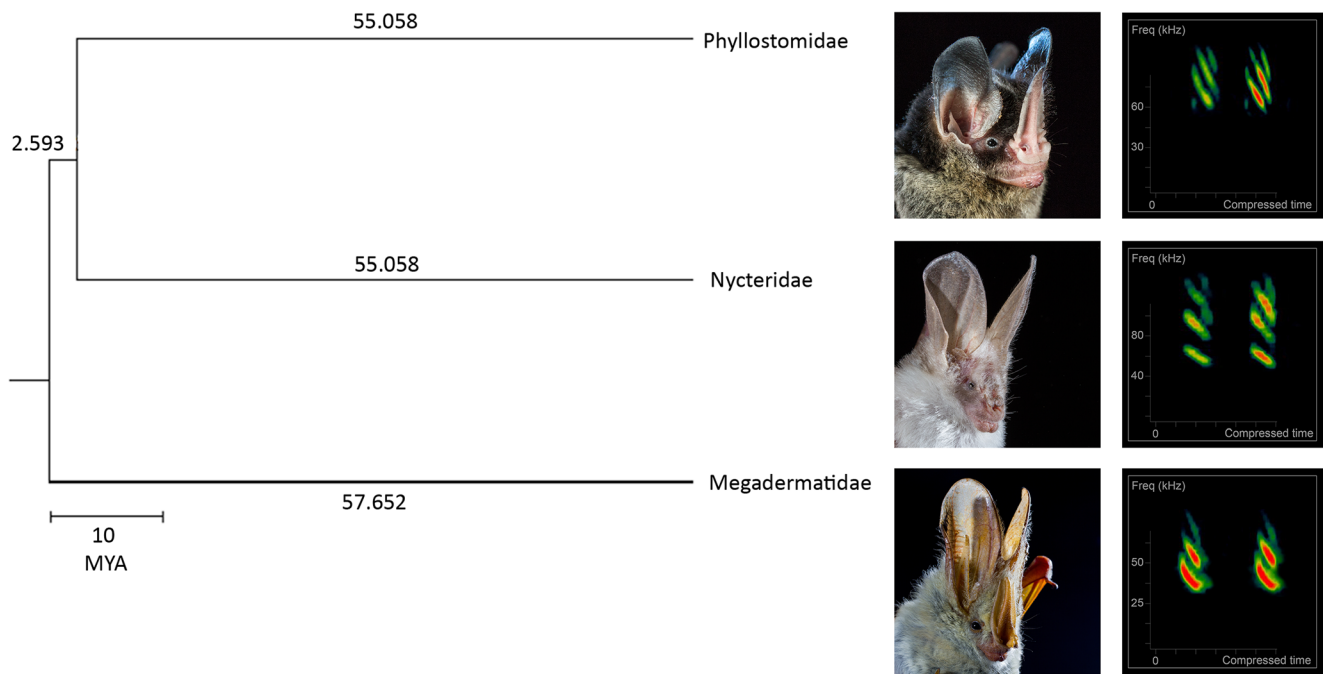


Fig. 7 Despite Phyllostomidae, Nycteridae and Megadermatidae only sharing a common ancestor over 55 million years ago, each family has independently evolved complex nose structures and multi-harmonic calls, a clear example of convergent evolution. Examples of a species call from each family: **a** *Gardnermyotis crenulatum* (Phyllostomidae), **b** *Nycteris*

thebaica (Nycteridae) and **c** *Lavia frons* (Megadermatidae) (b and c were recorded by Adrià López-Baucells in Sibiloi National Park, Kenya). Numbers on branches indicate the amount of time passed since the previous speciation event in millions of years. Phylogenetic reconstruction pruned from Shi and Rabosky (2015)

possible that the frequencies are not low but rather the forearm is relatively oversized. However, this hypothesis appears unlikely as *D. rotundus* still exhibits low-frequency calls when this variable is excluded from analysis. Another possible explanation for this lies in the fact that *D. rotundus* does not echolocate when approaching prey, its main use thus being navigation during commuting (Schmidt and Schmidt 1977). A high wing loading for long-distance commuting to find mammalian prey (Wilkinson 1985) supports the theory for a competitive advantage in producing decreased call frequencies (Altringham 1996). In contrast, *C. auritus* exhibited a higher than predicted call frequency relative to its body size (see Table 1) despite gleaning its prey. This species, unlike other large carnivorous bats such as *Trachops cirrhosus*, feeds almost exclusively on vertebrates (Smith 2008; Wilkinson and Wenrick Boughman 1998). Echolocating at lower frequencies may increase the likelihood of being detected by mammalian prey, hence the higher frequency echolocation observed (Norberg and Fenton 1988).

Study limitations

O'Farrell et al. (1999) showed that bat echolocation calls recorded shortly after hand release are similar to those recorded from bats emerging from roosts. However, phyllostomid bats usually emit highly modulated pulses of short ranges, and therefore the effect of handling should not be as severe as in

other aerial insectivorous bats with narrow-band calls. For this study, we minimized this potential bias by only considering pulses recorded at a distance greater than 5 m away from the point of release and, whenever possible, only when the released bat started circling around it. Although the initial screening process ensured maximum quality of acoustic recordings, this reduced the number of recordings available for analysis. Thus, the number of recordings for several less common species (e.g. *A. caudifer*) was relatively small compared with more commonly captured species (e.g. *A. lituratus*) (Table 1).

Conclusion

In this study we have aimed to both expand and update current knowledge regarding phyllostomid echolocation call design. It is clear that due to the considerable overlap between phyllostomid call parameters, live-trapping still provides the best method for surveying these species. However, future technological developments may allow minute differences in the echolocation call designs of phyllostomid bats to be distinguished from one another and thus allow these species to be monitored using remote sensing techniques.

Acknowledgements We would like to thank the BDFFP for logistical support and the many colleagues, volunteers and field assistants that

helped collecting data: Jorge M. Palmeirim, Fábio Z. Farneda, Oriol Massana, Diogo F. Ferreira, Marta Acácio, Gilberto Fernandez, Milou Groenenberg, Rodrigo Marciente, Madalena Boto, Inês , Julia Treitler, Solange Farias, Ubirajara Capaverde Jr., Alaércio Reis, Luiz Queiroz, Josimar Menezes, Osmaildo Silva and José Tenaçol. Paulo E.D. Bobrowiec also helped with fieldwork and provided logistic support.

Funding information This work was supported by the Portuguese Foundation for Science and Technology under research grant PTDC/BIABIC/111184/2009 (C.F.J.M.) and scholarships SFRH/BD/80488/2011 (R.R.), PD/BD/52597/2014 (A.L.B.) and NE/L002582/1 (N.Y.). Additional funding was provided by the Bat Conservation International to A.L.B. and R.R. and by ARDITI—Madeira's Regional Agency for the Development of Research, Technology and Innovation (grant M1420-09-5369-FSE-000002)—to R.R. This is publication 787 in the Technical Series of the BDFFP.

Compliance with ethical standards

Human and animal rights and informed consent All applicable international, national and/or institutional guidelines for the care and use of animals were followed. All procedures performed in studies involving animals were in accordance with the ethical standards of the institution or practice at which the studies were conducted (Instituto Chico Mendes de Conservação da Biodiversidade (ICMBio) under permit 26877-2).

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