

A new species of broad-nosed bat *Platyrrhinus* Saussure, 1860 (Chiroptera: Phyllostomidae) from the Guianan Shield

PAÚL M. VELAZCO¹ & BURTON K. LIM²

¹Division of Paleontology, American Museum of Natural History, Central Park West at 79th Street, New York, NY 10024, USA.

E-mail: pvelazco@amnh.org

²Department of Natural History, Royal Ontario Museum, 100 Queen's Park, Toronto, Ontario, M5S 2C6, Canada.

E-mail: burtonl@rom.on.ca

Abstract

A new species of broad-nosed bat *Platyrrhinus* Saussure, 1860 (Chiroptera: Phyllostomidae: Stenodermatinae) from the Guianan Shield is described based on molecular and morphological data. Previously confused with *P. helleri* and *P. recifinus*, the new taxon is currently known from only Guyana and Suriname and is most closely related to *P. recifinus* from eastern Brazil and not to the two sympatric species (*P. fusciventris* and *P. incarum*) also recently recognized as distinct from *P. helleri*. Morphometrically the new taxon overlaps with the smaller species of the genus (*P. angustirostris*, *P. brachycephalus*, *P. fusciventris*, *P. helleri*, *P. incarum*, and *P. matapalensis*), but forms a different cluster from the larger *P. recifinus*. Morphologically the new taxon is distinguished from its congeners by a combination of external and craniodental characteristics. *Platyrrhinus* now includes 21 species making it the most speciose genus in the Neotropical family Phyllostomidae.

Key words: Guianan Shield, Phyllostomidae, Phylogeny, *Platyrrhinus*, Neotropics, systematics

Introduction

The genus *Platyrrhinus* Saussure, 1860, belongs to the Neotropical bat family Phyllostomidae that ranges from southwestern United States to Paraguay. It includes more than 160 species, the most diverse family of bats in the Neotropics, which together exhibits more variation in morphological features and feeding ecology than any other family-level group of mammals (Simmons 2005; Gardner 2008a; Baker *et al.* 2012). Phyllostomid bats exploit an unusually diverse array of feeding habits including sanguivory, insectivory, carnivory, omnivory, nectarivory, polinivory, and frugivory (Wetterer *et al.* 2000; Baker *et al.* 2012).

Platyrrhinus includes at least 20 species of frugivorous bats endemic to the Neotropics (Velazco *et al.* 2010). The combined geographic ranges of these species extends from southern Mexico into Paraguay and northern Argentina (Velazco & Patterson 2008; Velazco *et al.* 2010). Species of *Platyrrhinus* occur primarily in tropical lowland and montane forest from sea level to at least 2,550 m (Gardner 2008b; Velazco *et al.* 2010).

Platyrrhinus is diagnosed from other genera of the subfamily Stenodermatinae by a combination of three characters: two accessory cusps on the posterior face of P4 (Velazco 2005, figure 12), presence of three upper molars, and presence of a fringe of hair along the trailing margin of the uropatagium. Although other genera also have these characters, no other genera possess all three at the same time (Lim 1993).

During recent years, the use of morphometric, morphological, and molecular techniques to study *Platyrrhinus* has enormously improved our knowledge of its taxonomy and phylogenetic relationships (Velazco & Solari 2003; Velazco 2005; Velazco & Patterson 2008; Velazco & Gardner 2009; Velazco *et al.* 2010). But still some taxa remain undescribed.

The *Platyrrhinus helleri* species complex was reviewed by Velazco (2005), Velazco and Patterson (2008), and Velazco *et al.* (2010). They found that this once widespread species, occurring from Oaxaca and Veracruz (Mexico) to Peru, Bolivia, Amazonian Brazil, northern South America, and Trinidad, was a composite of at least five species:

and Donna L. Dittmann (LSUMZ, Collection of Genetic Resources); Víctor Pacheco (MUSM); James L. Patton and Carla Cicero (MVZ); Judith L. Eger and Liliane Lortie (ROM); Robert J. Baker and Heath J. Garner (TTU, TK); Alfred L. Gardner and Suzanne C. Peurach (USGS-PWRC). Sequencing was carried out in the Field Museum's Pritzker Laboratory for Molecular Systematics and Evolution, operated with support from the Pritzker Foundation. Valéria da Cunha Tavares and one anonymous reviewer read early drafts of this manuscript and made helpful suggestions for its improvement. We are grateful to Alex Borisenko for allowing us to use his wonderful photographs of live *Platyrrhinus guianensis*. PMV was supported by NSF grant DEB 0949859 to Nancy B. Simmons. Fieldwork by BKL in Guyana and Suriname was supported by several grants from the ROM Governors.

References

- Baker, R.J., Bininda-Emonds, O.R.P., Mantilla-Meluk, H., Porter, C.A. & Van den Bussche, R.A. (2012) Molecular timescale of diversification of feeding strategy and morphology in New World leaf-nosed bats (Phyllostomidae): a phylogenetic perspective. In: Gunnell, G.F. & Simmons, N.B. (Eds.), *Evolutionary history of bats: fossils, molecules and morphology*. Cambridge University Press, Cambridge, pp. 385–409.
- Borisenko, A.V., Lim, B.K., Ivanova, N.V., Hanner, R.H. & Hebert, P.D.N. (2008) DNA barcoding in surveys of small mammal communities: a field study in Suriname. *Molecular Ecology Resources*, 8, 471–479.
<http://dx.doi.org/10.1111/j.1471-8286.2007.01998.x>
- Clare, E.L. (2011) Cryptic species? Patterns of maternal and paternal gene flow in eight Neotropical bats. *PLoS ONE*, 6, e21460.
<http://dx.doi.org/10.1371/journal.pone.0021460>
- Clare, E.L., Lim, B.K., Engstrom, M.D., Eger, J.L. & Hebert, P.D.N. (2007) DNA barcoding of Neotropical bats: species identification and discovery within Guyana. *Molecular Ecology Notes*, 7, 184–190.
<http://dx.doi.org/10.1111/j.1471-8286.2006.01657.x>
- Clare, E.L., Lim, B.K., Fenton, M.B. & Hebert, P.D.N. (2011) Neotropical bats: estimating species diversity with DNA barcodes. *PLoS ONE*, 6, e22648.
<http://dx.doi.org/10.1371/journal.pone.0022648>
- Darriba, D., Taboada, G.L., Doallo, R. & Posada, D. (2012) jModelTest 2: more models, new heuristics and parallel computing. *Nature Methods*, 9, 772.
<http://dx.doi.org/10.1038/nmeth.2109>
- Engstrom, M.D. & Lim, B.K. (2002) Mamíferos de Guyana. In: Ceballos, G. & Simonetti, J.A. (Eds.), *Diversidad y conservación de los mamíferos Neotropicales*. CONABIO-UNAM, México, D.F., pp. 329–375.
- Gardner, A.L. (2008a) Family Phyllostomidae Gray, 1825. In: Gardner, A.L. (Ed.), *Mammals of South America. Vol. 1. Marsupials, xenarthrans, shrews, and bats*. University of Chicago Press, Chicago, pp. 207–208.
- Gardner, A.L. (2008b) Genus *Platyrrhinus* Saussure, 1860. In: Gardner, A.L. (Ed.), *Mammals of South America. Vol. 1. Marsupials, xenarthrans, shrews, and bats*. University of Chicago Press, Chicago, pp. 329–342.
- Guindon, S. & Gascuel, O. (2003) A simple, fast, and accurate algorithm to estimate large phylogenies by maximum likelihood. *Systematic Biology*, 52, 696–704.
<http://dx.doi.org/10.1080/10635150390235520>
- Guttman, L. (1954) Some necessary conditions for common factor analysis. *Psychometrika*, 19, 149–161.
<http://dx.doi.org/10.1007/bf02289162>
- Huelsenbeck, J.P. & Ronquist, F. (2001) MRBAYES: Bayesian inference of phylogenetic trees. *Bioinformatics*, 17, 754–755.
<http://dx.doi.org/10.1093/bioinformatics/17.8.754>
- Jackson, D.A. (1993) Stopping rules in principal component analysis: a comparison of heuristical and statistical approaches. *Ecology*, 74, 2204–2214.
<http://dx.doi.org/10.2307/1939574>
- Larsen, P.A., Siles, L., Pedersen, S.C. & Kwiecinski, G.G. (2011) A new species of *Micronycteris* (Chiroptera: Phyllostomidae) from Saint Vincent, Lesser Antilles. *Mammalian Biology*, 76, 687–700.
<http://dx.doi.org/10.1016/j.mambio.2011.01.006>
- Lim, B.K. (1993) Cladistic reappraisal of stenodermatine bat phylogeny. *Cladistics*, 9, 147–165.
<http://dx.doi.org/10.1006/clad.1993.1009>
- Lim, B.K. (2009) Environmental assessment at the Bakhuis bauxite concession: small-sized mammal diversity and abundance in the lowland humid forests of Suriname. *The Open Biology Journal*, 2, 42–53.
<http://dx.doi.org/10.2174/1874196700902010042>
- Lim, B.K. (2012) Biogeography of mammals from the Guianas of South America. In: Patterson, B.D. & Costa, L.P. (Eds.) *Bones, clones, and biomes: an 80-million year history of modern Neotropical mammals*. The University of Chicago Press, Chicago, pp. 230–258.
- Lim, B.K. & Engstrom, M.D. (2000) Preliminary survey of bats from the upper Mazaruni of Guyana. *Chiroptera Neotropical*, 6 (1–2), 119–123.

- Lim, B.K. & Engstrom, M.D. (2001a) Species diversity of bats (Mammalia: Chiroptera) in Iwokrama Forest, Guyana, and the Guianan subregion: implications for conservation. *Biodiversity and Conservation*, 10, 613–657.
<http://dx.doi.org/10.1023/A:1016660123189>
- Lim, B.K. & Engstrom, M.D. (2001b) Bat community structure at Iwokrama Forest, Guyana. *Journal of Tropical Ecology*, 17, 647–665.
<http://dx.doi.org/10.1017/s0266467401001481>
- Lim, B.K. & Engstrom, M.D. (2005) Mammals of Iwokrama Forest. *Proceedings of the Academy of Natural Sciences of Philadelphia*, 154, 71–108.
[http://dx.doi.org/10.1635/0097-3157\(2004\)154\[0071:moif\]2.0.co;2](http://dx.doi.org/10.1635/0097-3157(2004)154[0071:moif]2.0.co;2)
- Lim, B.K., Engstrom, M.D., Genoways, H.H., Catzeffis, F.M., Fitzgerald, K.A., Peters, S.L., Djosetro, M., Brandon, S. & Mitro, S. (2005a) Results of the Alcoa Foundation-Suriname expeditions. XIV. Mammals of Brownsberg Nature Park, Suriname. *Annals of Carnegie Museum*, 74, 225–274.
[http://dx.doi.org/10.2992/0097-4463\(2005\)74\[225:rotafe\]2.0.co;2](http://dx.doi.org/10.2992/0097-4463(2005)74[225:rotafe]2.0.co;2)
- Lim, B.K., Engstrom, M.D. & Ochoa G., J. (2005b) Mammals. In: Hollowell, T. & Reynolds, R. P. (Eds.), Checklist of the terrestrial vertebrates of the Guiana Shield. *Bulletin of the Biological Society of Washington*, 13, 77–92.
- Lim, B.K. & Norman, Z. (2002) Rapid assessment of small mammals in the eastern Kanuku Mountains, Lower Kwitaro River area, Guyana. In: Montambault, J.R. & Missa, O. (Eds.), *A biodiversity assessment of the eastern Kanuku Mountains, Lower Kwitaro River, Guyana*. Conservation International, Arlington, Virginia, pp. 51–58.
- Lim, B.K., Peterson, A.T. & Engstrom, M.D. (2002) Robustness of ecological niche modeling algorithms for mammals in Guyana. *Biodiversity and Conservation*, 11, 1237–1246.
<http://dx.doi.org/10.1023/a:1016038501986>
- Lim, B.K. & Tavares, V.C. (2012) Review of species richness and biogeography of bats (Chiroptera) from the Guiana subregion of South America with comments on conservation. *Ecotropica*, 18, 195–118.
- Rambaut, A. & Drummond, A.J. (2007) Tracer. Available from: <http://beast.bio.ed.ac.uk/Tracer> (accessed 30 April 2014)
- Simmons, N.B. (2005) Order Chiroptera. In: Wilson, D.E. & Reeder, D.M. (Eds.), *Mammal species of the World: a taxonomic and geographic reference*. Johns Hopkins University Press, Baltimore, pp. 312–529.
- Smith, P.G. & Kerry, S.M. (1996) The Iwokrama Rain Forest Programme for sustainable development: How much of Guyana's bat (Chiroptera) diversity does it encompass? *Biodiversity and Conservation*, 5, 921–942.
<http://dx.doi.org/10.1007/bf0054742>
- Swofford, D.L. (2002) *PAUP*. Phylogenetic Analysis Using Parsimony (*and other methods)*. Sinauer Associates, Sunderland, Massachusetts.
- Tavares, V.C. & Velazco, P.M. (2010) *Platyrrhinus recifinus* (Chiroptera: Phyllostomidae). *Mammalian Species*, 42, 119–123.
<http://dx.doi.org/10.1644/859.1>
- Vaidya, G., Lohman, D.J. & Meier, R. (2011) SequenceMatrix: concatenation software for the fast assembly of multi-gene datasets with character set and codon information. *Cladistics*, 27, 171–180.
<http://dx.doi.org/10.1111/j.1096-0031.2010.00329.x>
- Velazco, P.M. (2005) Morphological phylogeny of the bat genus *Platyrrhinus* Saussure, 1860 (Chiroptera: Phyllostomidae) with the description of four new species. *Fieldiana Zoology, New Series*, 105, 1–53.
<http://dx.doi.org/10.5962/bhl.title.2689>
- Velazco, P.M. (2009) *Historical diversification in the Neotropics: evolution and variation of the bat genus Platyrhinus*. Ph.D. dissertation, University of Illinois at Chicago, Chicago, 329 pp.
- Velazco, P.M. & Gardner, A.L. (2009) A new species of *Platyrrhinus* (Chiroptera: Phyllostomidae) from western Colombia and Ecuador, with emended diagnoses of *P. aquilus*, *P. dorsalis*, and *P. umbratus*. *Proceedings of the Biological Society of Washington*, 122, 249–281.
<http://dx.doi.org/10.2988/08-40.1>
- Velazco, P.M., Gardner, A.L. & Patterson, B.D. (2010) Systematics of the *Platyrrhinus helleri* species complex (Chiroptera: Phyllostomidae), with descriptions of two new species. *Zoological Journal of the Linnean Society*, 159, 785–812.
<http://dx.doi.org/10.1111/j.1096-3642.2009.00610.x>
- Velazco, P.M. & Patterson, B.D. (2008) Phylogenetics and biogeography of the broad-nosed bats, genus *Platyrrhinus* (Chiroptera: Phyllostomidae). *Molecular Phylogenetics and Evolution*, 49, 749–759.
<http://dx.doi.org/10.1016/j.ympev.2008.09.015>
- Velazco, P.M. & Patterson, B.D. (2013) Diversification of the yellow-shouldered bats, genus *Sturnira* (Chiroptera, Phyllostomidae), in the New World tropics. *Molecular Phylogenetics and Evolution*, 68, 683–698.
<http://dx.doi.org/10.1016/j.ympev.2013.04.016>
- Velazco, P.M. & Solari, S. (2003) Taxonomía de *Platyrrhinus dorsalis* y *Platyrrhinus lineatus* (Chiroptera: Phyllostomidae) en Perú. *Mastozoología Neotropical*, 10, 303–319.
- Wetterer, A.L., Rockman, M.V. & Simmons, N.B. (2000) Phylogeny of phyllostomid bats: data from diverse morphological systems, sex chromosomes, and restriction sites. *Bulletin of the American Museum of Natural History*, 248, 1–200.
[http://dx.doi.org/10.1206/0003-0090\(2000\)248<0001:popbmc>2.0.co;2](http://dx.doi.org/10.1206/0003-0090(2000)248<0001:popbmc>2.0.co;2)
- Zwickl, D.J. (2006) *Genetic algorithm approaches for the phylogenetic analysis of large biological sequence datasets under the maximum likelihood criterion*. Ph.D. dissertation, The University of Texas at Austin, Texas, 125 pp.