

The systematic status of the Libyan bats of the genus *Plecotus* (Chiroptera: Vespertilionidae)

Systematická posice libyjských netopýrů rodu *Plecotus*
(Chiroptera: Vespertilionidae)

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Abstract. This paper briefly reviews the taxonomic history of the *Plecotus kolombatovici* complex, the bat populations which occur in isolated ranges surrounding the Mediterranean Basin, including Maghreb, Tripolitania, Cyrenaica, Anatolia, Balkans, and Italy, plus numerous Mediterranean islands. Since this complex exhibits genetic characters close to *Plecotus teneriffae* from the Canary Islands, the samples of the complex from Cyrenaica (NE Libya) were described as a subspecies *P. teneriffae gaisleri*. However, this name was later used for all populations of the complex inhabiting North Africa as *P. gaisleri*, a full species separated from *P. kolombatovici*. However, the available results of molecular genetic analyses of the complex support the opinion that these allopatric populations are conspecific and referable to *P. kolombatovici* and reopens the question regarding the systematic status of *P. teneriffae*.

Key words. *Plecotus teneriffae*, *Plecotus kolombatovici*, *Plecotus gaisleri*, Cyrenaica, taxonomy.

In the context of species protection issues, a question regarding the systematic status of long-eared bats occurring on Malta and Pantellaria arose. This article tries to clarify the complex taxonomy of *Plecotus kolombatovici*, a bat that occurs in often isolated Mediterranean landscapes around the Mediterranean Basin: Maghreb (Morocco, Algeria, Tunisia), Tripolitania and Cyrenaica (Libya) in the south, and Anatolia, Greece, Albania, Croatia, and Italy in the north (JUSTE et al. 2004, BENDA et al. 2004, SPITZENBERGER et al. 2006, ANCILLOTTO et al. 2019). Furthermore it has been recorded on numerous islands in the Adriatic and Aegean Seas, plus in Crete and Cyprus.

For about 150 years, systematists have referred all western Palaearctic long-eared bats to a single species, *Plecotus auritus* (Linnaeus, 1758), due to the lack of obvious phenetic differences within or among populations. *Plecotus christii* Gray, 1838 was the first taxon to be described as a species of its own; it was followed by *Plecotus teneriffae* Barret-Hamilton, 1907 and *P. balensis* Kruskop et Lavrenchenko, 2000. However, the former two names were considered just subspecies either of *P. auritus* or *P. austriacus* for a long time (HANÁK 1966, HORÁČEK et al. 2000). *Plecotus austriacus* (Fischer, 1829), originally described as a variety of *P. auritus*, was elevated to species rank by BAUER (1960), and twenty years later ĐULIĆ (1980) described a new subspecies of this bat from coastal Croatia, *P. austriacus kolombatovici* Đulić, 1980. Recognition of similarities of *P. teneriffae* and *P. austriacus kolombatovici* by DE PAZ (1994) was the last important contribution to the classification of the western Palaearctic long-eared bats based solely on morphological analysis.

Only the advent of genetic analyses shed light into the rich diversity of taxa within the genus *Plecotus* but also gave rise to considerable confusion and controversy regarding the taxonomy of these taxa. “It is

only with the recent use of molecular characters that the evolutionary picture of the Palearctic *Plecotus* has begun to be unveiled” (JUSTE et al. 2004: 1121).

Genetic analyses of populations of the *P. austriacus* clade established species rank of *P. kolombatovici* (MAYER & VON HELVERSEN 2001) and confirmed species status of *P. teneriffae* (JUSTE et al. 2004) as had been already suggested by IBÁÑEZ & FERNANDEZ (1985).

Based on analyses of mitochondrial genes, (1) JUSTE et al. (2004) classified the North African *Plecotus* samples as *P. cf. kolombatovici* (Cyrenaica) and *Plecotus* sp. (Morocco); (2) BENDA et al. (2004) assigned all *Plecotus* populations from Croatia, Turkey, Libya, and Morocco to *P. teneriffae*, described populations from Cyrenaica as a new subspecies, *P. teneriffae gaisleri* Benda, Kiefer, Hanák et Veith, 2004, and named the bats from Tripolitania and Morocco *P. teneriffae cf. gaisleri*; (3) SPITZENBERGER et al. (2006) treated *P. teneriffae* and *P. kolombatovici* as two separate species and found three distinct subspecific groups in the latter species: (a) Croatia and Turkey, (b) Cyrenaica, and (c) Morocco; (4) in a comprehensive study comprising material of two bat families and numerous genera, and *P. austriacus*, *P. christii*, *P. t. gaisleri*, *P. kolombatovici*, but not *P. teneriffae*, MAYER et al. (2007) suggested elevating *P. t. gaisleri* to the species rank; (5) based on analyses of a comprehensive multilocus molecular dataset (in total 96 samples of six *Plecotus* taxa, including *P. kolombatovici* complex) from a wide geographic scope, ANCILLOTTO et al. (2020) assigned samples from Malta and Pantelleria to *P. gaisleri* and provided ample information on intra- and interspecific divergences (p-distances) which clearly separated valid species (*P. austriacus*, *P. christii*, *P. kolombatovici*) from allopatric populations of *P. kolombatovici* plus *P. teneriffae*. While p-distances (16s gene) between the full species varied between 4.71% and 6.78%, the variation within *P. kolombatovici* plus *P. teneriffae* was between 1.31% and 2.75%.

This result supports the opinion that all allopatric populations (including those from Cyrenaica) of *P. kolombatovici* are conspecific and reopens the question regarding the systematic status of *P. teneriffae*. Not least because of the highly needed protection of endangered bat species living on islands and in isolated ranges, the systematics of *P. kolombatovici* should be subjected to a thorough study using nuclear markers. Such a study could elucidate the phylogeography of *Plecotus kolombatovici* and thus complement the series of studies of other bat species with circum-Mediterranean distribution (see BILGIN et al. 2016).

SOUHRN

Príspevek stručne shrnuje taxonomickou históriu komplexu ušana balkánskeho (*Plecotus kolombatovici*), tedy netopýřích populací, které v izolovaných areálech obývají území obklopující Středozemní moře, včetně Maghrebu, Tripolitánie, Kyrenaiky, Anatólie, Balkánskeho poloostrova a Itálie, a také mnoha středomořských ostrovů. Jelikož tento komplex vykazuje genetické znaky blízké ušanovi kanárskému (*Plecotus teneriffae*) z Kanárských ostrovů, byli jedinci tohoto komplexu obývající Kyrenaiku (severovýchodní část Libye) popsáni jako samostatný poddruh, *P. teneriffae gaisleri*. Později však toto jméno začalo být používáno jako jméno druhové pro všechny populace komplexu obývající severní Afriku, jako ušan berberský (*Plecotus gaisleri*), druh neodvislý od ušana balkánskeho. Dostupné výsledky molekulárně genetických analys mitochondriálního genomu celého komplexu však podporují názor, že tyto středomořské alopatrické populace představují jediný druh, *Plecotus kolombatovici*; tím také znovu otevírají otázku systematického postavení ušana kanárského (*Plecotus teneriffae*).

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