

Phylogenetic relationships within the *cahirinus-dimidiatus* group of the genus *Acomys* (Rodentia: Muridae): new mitochondrial lineages from Sahara, Iran and the Arabian Peninsula

DANIEL FRYNTA^{1,7}, KLÁRA PALUPČÍKOVÁ¹, ERICA BELLINVIA², PETR BENDA^{1,3},
HANA SKARLANTOVÁ^{1,4}, LUCIE SCHWARZOVÁ^{1,5} & DAVID MODRÝ⁶

¹Department of Zoology, Faculty of Science, Charles University in Prague, Viničná 7, CZ-128 44 Praha 2, Czech Republic.

E-mail: klpr@post.cz

²Department of Plant Physiology, Faculty of Science, Charles University in Prague, Viničná 7, CZ-128 44 Praha 2, Czech Republic.
E-mail: popielica@yahoo.com

³Department of Zoology, National Museum (Natural History), CZ-115 79, Praha 1, Czech Republic. E-mail: petr_benda@nm.cz

⁴Present address: First Faculty of Medicine, Charles University in Prague, Kateřinská 32, CZ-121 08 Praha 2, Czech Republic.
E-mail: hanka.svraster@tiscali.cz

⁵Present address: Institute of Biology and Medical Genetics, 1st Faculty of Medicine and General Teaching Hospital, Albertov 4,
Prague 2, CZ-128 00, Czech Republic. E-mail: lucie.schwarzova@yahoo.com

⁶Institute of Parasitology, Veterinary University, Palackého 1/3, CZ- 612 42, Brno, Czech Republic. E-mail: ModryD@VFU.cz

⁷Corresponding author. E-mail: frynta@centrum.cz

Abstract

Spiny mice belonging to the *cahirinus-dimidiatus* group of the genus *Acomys* have become a widely used model in physiology and behaviour. To improve current knowledge concerning the phylogeny of this taxon, we analysed 24 samples from Libya, Chad, Egypt, Jordan, Cyprus, Crete, Turkey, Yemen and Iran. We sequenced the whole mitochondrial control region and part of the flanking tRNA genes for a total length of 986 to 996 bp and described 22 haplotypes. Our results confirmed that the Afro-Mediterranean and Asian clades are clearly distinct (p-distance = 6–8.1%). The former clade corresponds to *A. cahirinus sensu lato* (i.e. including also the Cretan *A. minous*, Cypriot *A. nesiotes* and Turkish *A. cilicicus*). Haplotypes of *A. cahirinus* from the E Sahara (S Egypt, SW Libya, N Chad) grouped with those of *A. cilicicus* and *A. minous* (p-distance ≤ 2.2%), while haplotypes of *A. nesiotes* grouped with one haplotype representing the commensal *A. cahirinus* from Cairo (p-distance = 1.2%). Close similarity among haplotypes from mainland Africa and NE Mediterranean (clade *A. cahirinus sensu stricto*) support the hypothesis that ancestors of *A. nesiotes*, *A. cilicicus* and *A. minous* dispersed most probably as commensal populations, thus questioning their status of valid species. The most surprising finding was the considerable genetic variation in Asia. In addition to a haplogroup from Sinai and Jordan (corresponding to *A. dimidiatus sensu stricto*), we detected two previously unknown haplogroups, from Yemen and Iran + United Arab Emirates. These clades are fairly distinct and separate species/subspecies status of these animals might be further considered.

Key words: spiny mice, mitochondrial DNA, mitochondrial control region, D-loop, phylogeography, commensalism, Yemen, Libya, Cyprus, Persian Gulf

Relazioni filogenetiche all'interno del gruppo *cahirinus-dimidiatus* nel genere *Acomys* (Rodentia: Muridae): nuove linee mitocondriali identificate nella regione del Sahara, in Iran e nella penisola araba

Sommario

I topi spinosi appartenenti al gruppo *cahirinus-dimidiatus* nel genere *Acomys* sono diventati animali modello ampiamente usati in studi fisiologici e comportamentali. Per migliorare le conoscenze attuali riguardanti la filogenesi di questo taxon, abbiamo analizzato 24 esemplari di topo spinoso provenienti da Libia, Chad, Egitto, Giordania, Cipro,