

Phylogeny and biogeography of the *Acanthodactylus scutellatus* species complex in North Africa

André Vicente Liz^{1,2,3*}, Dennis Rödder³, Duarte Vasconcelos Gonçalves^{1,4}, Guillermo Velo-Antón^{1,5}, Philippe Geniez⁶, Pierre-André Crochet⁷, José Carlos Brito^{1,2}

¹CIBIO/InBIO, Centro de Investigação em Biodiversidade e Recursos Genéticos da Universidade do Porto, Instituto de Ciências Agrárias de Vairão, Rua Padre Armando Quintas 7, 4485-661 Vairão, Portugal

²Departamento de Biologia, Faculdade de Ciências, Universidade do Porto, Rua Campo Alegre s/n, 4169-007 Porto, Portugal

³ZFMK, Zoologisches Forschungsmuseum Alexander Koenig, Adenauerallee 160, D-53113 Bonn, Germany

⁴CIIMAR, Centro Interdisciplinar de Investigação Marinha e Ambiental, Terminal de Cruzeiros de Leixões, Avenida G. Norton de Matos s/n, 4450-208 Matosinhos, Portugal

⁵Universidade de Vigo, Grupo GEA, Departamento de Ecología e Bioloxía Animal, Vigo, Spain

⁶CEFE, Univ Montpellier, CNRS, EPHE-PSL University, IRD, Biogéographie et Ecologie des Vertébrés, Montpellier, France

⁷CEFE, CNRS, Univ Montpellier, EPHE, IRD, Montpellier, France

*Corresponding author: A.V. Liz (andre.vicente.liz@cibio.up.pt)

Deserts and arid regions are perceived as homogeneous spaces poor in biodiversity. Yet, this contrasts with considerable endemism and the high degree of genetic structure observed in some species inhabiting the Sahara Desert, which have been frequently attributed to progressive aridification and climatic oscillations during the Plio-Pleistocene. Lizards of the *Acanthodactylus scutellatus* complex are among the most conspicuous representatives of the Saharan xeric fauna. The complex is composed of seven currently recognised species: *A. aegyptius*, *A. aureus*, *A. dumerilii*, *A. longipes*, *A. scutellatus*, *A. senegalensis*, and *A. taghitensis*. Despite being a promising model to assess historical biodiversity dynamics in arid regions, no comprehensive study on the evolutionary history of the complex has been performed to date. Our study addresses the phylogenetic relationships, potential cryptic diversity, and inter-specific contact zones within the *A. scutellatus* complex. Two mitochondrial markers and one nuclear gene were used to infer phylogenetic relationships and identify cryptic lineages, while species' ecological models based on climatic and remote-sensing variables were used to estimate potential contact zones. Phylogenetic relationships among species were generally recovered with good support. However, the lack of distinction between *A. dumerilii* and *A. senegalensis* calls for a taxonomic reassessment. The presence of



several highly-divergent lineages, particularly around mountain areas, reveals a high degree of cryptic diversity within the complex. Ecological models suggest a series of contact zones where potential inter-specific hybridization remains unstudied. This work contradicts the “empty-desert” conception and underlines the existence of still unexplored diversity across the Sahara.

