

Alpha-taxonomy in world mammals

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For A/Prof A.M. Muasya



Introduction

Herein I discuss the progress of mammalian alpha-taxonomy from 1975 to 2005. First, I discuss the new species and groups discovered in this period. Second, I focus on the methods employed to diagnose and delimit these mammal species and how these have changes over the the past 30 to 40 years. Third, I will attempt to draw conclusions, if possible, about the speciation mechanisms at work in mammal populations based on the evidence use to delimit those mammal species. And lastly, I will frame these discussions in the context of the species concepts (or lack thereof) typically used in mammalogy and mammalian taxonomy.

Mammal Species of the World, 3rd Edition (MSW3) (Wilson & Reeder 2005) is a reputable database of mammalian taxa (species, sub-specific, and super-specific) and information concerning their treatments (e.g. scientific name, authors' name and year described, original publication citation, distribution, etc. (Wilson & Reeder 2005)) from 1702 to 2005 AD. This data product is a boon to the discussions here. Using the .csv file from the MSW3 webpage (<https://www.departments.bucknell.edu/biology/resources/msw3/>), I explored this dataset in R (R Core Team 2017), primarily using the tidyverse suite of packages for exploration and visualisation (Wickham 2017).

There is a subset of mammal order that are most taxonomically “active” (Figure 1), such that there are only eight orders in which 10 new species were described in the period 1975—2005.

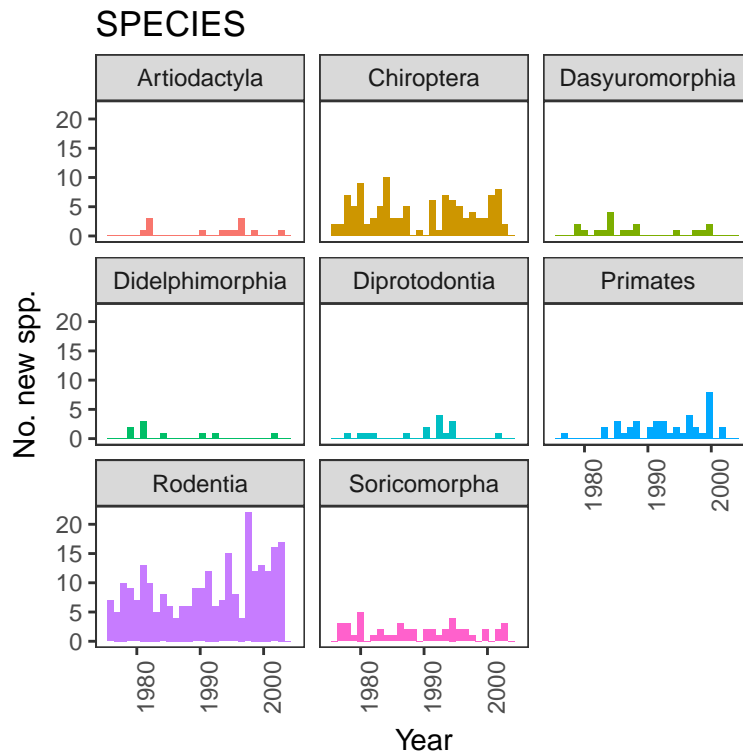


Figure 1: (ca. 54 hours range of sampling): **A)** Rain-fall input (m^3h^{-1}), **B)** stream-flow rate (m^3h^{-1}) of the Liesbeeck River, and **C)** $\delta^{18}O$ (‰) and **D)** δ^2H (‰) values for both the rain- and stream-water (keyed).

Mammal discoveries, 1975—2005

Lines of evidence

Modes of mammalian speciation

Mammalogical species concepts

Concluding remarks

e.g. Blah blah blah (Figures \ref{o_vs_h}B, \ref{o_vs_h}C).

e.g. {r timeline, warning = F, fig.height = 7, fig.width = 7, fig.cap = '\\label{timeline} \\footnotesize Timeline of isotopic and hydrological values throughout the winter storm-event during which sampling was carried out (ca. 54 hours range of sampling): \\normalsize \\textbf{A)} \\footnotesize Rain-fall input ($m^3 h^{-1}$), \\normalsize \\textbf{B)} \\footnotesize stream-flow rate ($m^3 h^{-1}$) of the Liesbeeck River, and \\normalsize \\textbf{C)} \\footnotesize $\delta^{18}O$ (‰) and \\normalsize \\textbf{D)} \\footnotesize δ^2H (‰) values for both the rain- and stream-water (keyed).'} }

References

- R Core Team. 2017. *R: A Language and Environment for Statistical Computing*. 2017. Vienna, Austria: R Foundation for Statistical Computing. Available: <https://www.r-project.org/>.
- Wickham, H. 2017. *tidyverse: Easily Install and Load 'Tidyverse' Packages*. 2017. Available: <https://cran.r-project.org/package=tidyverse>.
- Wilson, D. & Reeder, D.-A. Eds. 2005. *Mammal Species of the World. A Taxonomic and Geographic Reference*. 3rd edition. 2005. Baltimore: John Hopkins University Press. Available: <https://www.departments.bucknell.edu/biology/resources/msw3/>.