

Title of submission	palaeoverse: a community-driven R package to support palaeobiological analysis
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Software location (including archive name and persistent identifier, e.g. doi)	GitHub: https://github.com/palaeoverse-community/palaeoverse CRAN: https://cran.r-project.org/web/packages/palaeoverse/
Operating system	Windows, macOS, Ubuntu (Linux)
Programming language and minimum version needed	R (≥ 4.0)
Additional system requirements	No additional requirements other than those required to run R (≥ 4.0).
Dependencies	R (≥ 4.0)
Installation instructions	The palaeoverse package can be installed from CRAN using the <code>install.packages</code> function in R: <code>install.packages("palaeoverse")</code> If preferred, the development version of palaeoverse can be installed from GitHub via the <code>remotes</code> R package: <code>remotes::install_github("palaeoverse-community/palaeoverse")</code>
Test coverage	97%
Continuous Integration	The palaeoverse repository uses GitHub Actions to perform R CMD check on Linux, Mac, and Windows operating systems every time a push (or pull request) to the repository is made.
License	GPL-3.0
Version and date published	1.0.0 (31 st October 2022)