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	Depends:
	  R (≥ 4.0)
	Imports:
	  stats,
	  utils,
	  graphics,
	  methods,
	  curl,
	  deeptime (>= 1.0.0),
	  ape,
	  sf,
	  stringdist,
	  geosphere,
	  h3jsr (>= 1.3.0),
	  httr,
	  pbapply

Installation instructions	The palaeoverse package can be installed from CRAN using the install.packages function in R: install.packages("palaeoverse") If preferred, the development version of palaeoverse can be installed from GitHub via the remotes R package: remotes::install_github("palaeoverse-community/palaeoverse")
Test coverage	96%

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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	1.1.0 (12th January 2023)