

Inheritance and inequality under communal land tenure: convergence on cultural–evolutionary regimes

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R packages

An alphabetical list of all R packages used in this work is provided below, indicating version numbers used at time of writing.

ade4 (1.7.23), adegenet (2.1.11), adiv (2.2.1), ape (5.8.1), bayesplot (1.15.0), Biodem (0.5), brms (2.23.0), cluster (2.1.8.2), conflicted (1.2.0), cowplot (1.2.0), dendextend (1.19.1), dplyr (1.2.0), e1071 (1.7.17), factoextra (2.0.0), forcats (1.0.1), fpc (2.2.14), geiger (2.0.11), geosphere (1.6.5), ggplot2 (4.0.2), ggpubr (0.6.3), ggtree (4.0.4), graph4lg (1.8.0), gridExtra (2.3), here (1.0.2), hillR (0.5.2), Hmisc (5.2.5), installr (0.23.4), lattice (0.22.9), lubridate (1.9.5), magick (2.9.1), magrittr (2.0.4), maps (3.4.3), MASS (7.3.65), NbClust (3.0.1), nlme (3.1.168), ouch (2.20), permute (0.9.10), phangorn (2.12.1), phylogram (2.1.0), phytools (2.5.2), picante (1.8.2), pkgbuild (1.4.8), purrr (1.2.1), RColorBrewer (1.1.3), Rcpp (1.1.1), readr (2.2.0), readxl (1.4.5), REAT (3.0.3), rr2 (1.1.1), sf (1.1.0), spData (2.3.4), spdep (1.4.2), stargazer (5.2.3), stringr (1.6.0), surface (0.6), tibble (3.3.1), tidyr (1.3.2), tidyverse (2.0.0), treebalance (1.2.0), vegan (2.7.2), viridisLite (0.4.3), writexl (1.5.4).