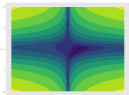
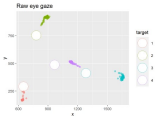


Computational Methods

[img2net](#) v0.1.0: Implements MAG3D, a Monomodal, Adaptive Dual-Averaged Gradient method for stochastic optimization. See [Dehrie & Jolani \(2021\)](#) for details and [REACME](#) to get started.



[TidNetRegression](#) v1.0.0: Provides functions to fit 2D and 3D transformations using [TidNet](#) which return posterior distributed for fitted parameters. There are vignettes on [Transformation Matrices](#), [Eye Gaze Mapping](#), and [Comparing Faces](#). See [REACME](#) to get started.



Data

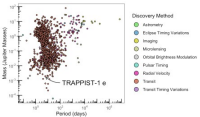
[AirClim](#) v0.1.0: Provides access to air quality and meteorological information from the China's National Air Quality System ([CNQA](#)). See [REACME](#) to get started.

[bepanque](#) v0.0.1: Provides a lightweight interface to access spatial benchmarks from open sources such as [OpenStreetMap](#), [Mapbox](#), and others. See [REACME](#) to get started.

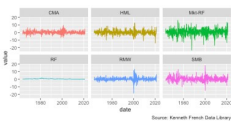


[causalrds](#) v0.1.1: Contains the data sets to run the example problems in the online causal inference textbooks: [The Effect](#) and [Causal Inference: What If](#) and more.

[exoplanets](#) v0.2.1: Provides access to NASA's [Exoplanet Archive](#). See the [vignette](#) to get started.



[franchise](#) v0.1.1: Provides access to Kenneth's French [franchise data library](#). See the [vignette](#) for basic usage.

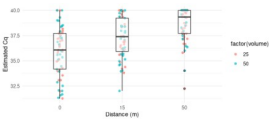


[Inequality](#) v0.5.0: Provides access to the data sets from [Tobin et al. \(2018\)](#) along with an [online book](#) containing commentary and the code to recreate the original analysis.



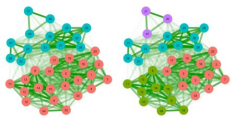
Genomics

[adonis](#) v0.0.1: Provides a modeling framework for the design and analysis of experiments collecting environmental DNA. There is an [Introduction](#) and also vignettes on [Unloading eDNA](#) and [gPCR Data and Simulating eDNA Data](#).



[MACE](#) v0.0.0: Provides functions to perform variant self-based main effect tests, gene-environment interaction tests, and joint tests for association, as proposed in [Weng et al. \(2020\)](#). See the [vignette](#) for details.

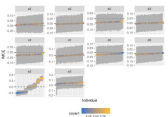
[MUTS](#) v0.5.1: Implements a bioinformatic approach to detect the multiple integration of viral vectors within the same clone. See the [vignette](#) for how to use the package.



[Region](#) v0.0.0: Provides functions to identify topological domains in genomes from Hi-C sequence data as described in [Shin et al. \(2016\)](#). See [REACME](#) to get started.

Machine Learning

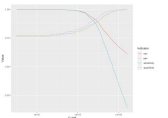
[rbrnet](#) v0.1.0: Implements a tool for analyzing conjoint experiments using Bayesian Additive Regression Trees (BART), a machine learning method developed by [Chipman & McCulloch \(2015\)](#). See the [vignette](#) for examples.



[TextTest](#) v0.0.1: Implements an interface to Facebook's [TextTest Library](#). See [Rogneswold et al. \(2017\)](#) for a description of the algorithm. There is a [Benchmark](#) vignette and an [Introduction](#).

Medicine

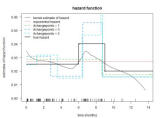
[vite](#) v0.1.1: Provides functions to estimate diagnostic performance (Sensitivity, Specificity, Positive predictive value, Negative predicted value) of a diagnostic test when there is no golden standard by estimating the attributable fraction using either a [logpenalized model](#) or a [latent class model](#).



[covidstat](#) v0.0.2: Provides an interface to [Johns Hopkins COVID-19 Dashboard](#) including tools for data access, maps and time series plotting, and a collection of numerous indicators relevant to the COVID-19 pandemic in the United States. There is a [Getting Started Guide](#), and vignettes on [Computing Signal Correlations](#), [Combining Data Sources](#), [Manipulating Multiple Signals](#), and [Plotting and Mapping Signals](#).



[eventTrack](#) v0.0.0: Implements the hybrid framework for event prediction in clinical trials as described in [Fang & Zhang \(2011\)](#). See the [vignette](#) for an example.



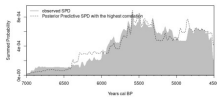
[gblibds](#) v0.0.0: Implements the Gblibds adaptive trial design for a time to event outcome using a piecewise exponential model and conjugate Gamma prior distributions as described in [Briggs et al. \(2014\)](#). See the [vignette](#) for an example.

Science

[CopernicusDEM](#) v0.0.1: Provides an interface to the [Copernicus DEM](#) Digital Elevation Model of the European Space Agency with 90 and 30 meters resolution using the [WGS 84](#) coordinate line tool. See the [vignette](#) for an example.

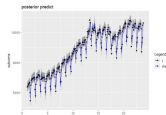


[HMMCalc](#) v0.1.2: Provides functions and a custom probability distribution for Bayesian analysis of redoxation data within the [HMMCalc](#) modeling framework, including a suite of functions for prior and posterior predictive checks for demographic inference as described in [Cerna & Shukla \(2021\)](#). See the [Introduction](#).

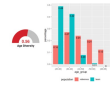


Statistics

bayesmodels v0.1.0: Implements a framework to bring a number of Bayesian models into the *tidymodels* ecosystem. See the [vignette](#) for an overview.



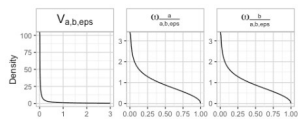
div v0.2.1: Provides functions to facilitate the analysis of teams in a corporate setting, assess the diversity per grade and job, search for bias and also provides methods to simulate the effects of bias. See [De Bruinier \(2021\)](#) and [De Bruinier \(2020\)](#) for background. Look [here](#) to get started.



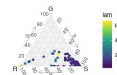
HoldingRings v0.1.1: Provides functions to compute the semi-axis lengths and coordinate points of holding ellipses. See [Bro & Snelis \(2016\)](#) and [Bretz \(2016\)](#) for background. Look [here](#) and at the [vignette](#) for examples.



mlcomparator v1.0.0: Provides tools to construct and visualize joint priors for variance parameters. *Vignettes* provide examples for Latin Square, i.i.d. models, neuronal mortality, and wheat breeding.



Prager v1.0.0: Provides functions for calculating the history metrics using matrix population models (MPMs) as described in [Jones et al. \(2021\)](#). There is a [Getting Started Guide](#) and *vignettes* on Vital Rates, Life History Traits, Denning Age, and Territory Plot.

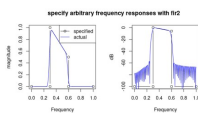


unsculptor v0.1.0: Provides functions to calculate *Mantel-like distances* for every row of a set of outcome variables. There is an [Introduction](#) and a *vignette* on the [calculations](#).



Time Series

signal v0.2-2: Implements the *Online signal* package which provides a variety of signal processing tasks, such as signal generation and measurement, correlation and convolution, filtering, filter design, filter analysis and conversion, power spectrum analysis, system identification, deconvolution and sample rate change, and windowing. See the [vignette](#) for an introduction.



signal v0.1.0: Provides functions for implementing multivariate data space models such as Vector Exponential Smoothing and Vector Error-Trend-Seasonal models, for time series analysis and forecasting as described in [de Souza et al. \(2010\)](#). There is a [Function Overview](#) and *vignettes* on *Vector ES* and *Vector ETS*.

Utilities

paranoid v0.1.2: Implements formal grammar and parser for R Markdown documents using the *Boost Spirit V3* library. It also includes a collection of high-level functions for working with the resulting abstract syntax tree. There is a [Getting Started Guide](#) and a *vignette* on [Rmd Templates](#).

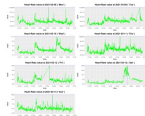
robustness v0.1.0: Provides facilities for assessing R packages against a number of metrics to help quantify their robustness. Look [here](#) for background on the package and [here](#) for background on the R Consortium, R Validation Hub project. There is a [Quick Start Guide](#) and a *vignette* on [Extending robustness](#).

robustness v0.1.0: Provides functions to improve the user experience of shiny apps by providing feedback when required inputs are missing, or input values are not valid. See [README](#) to get started.

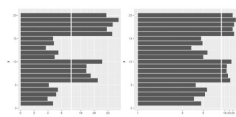
trv v1.0: Provides tools to create structured, formatted HTML tables. See the [vignette](#).

Visualization

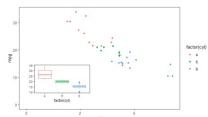
shiny v0.1.1: Implements a connection to the *PlotlyJS API* to provide *ggplot2*, *leaflet* and *mapbox* visualizations. See the [vignette](#) for examples.



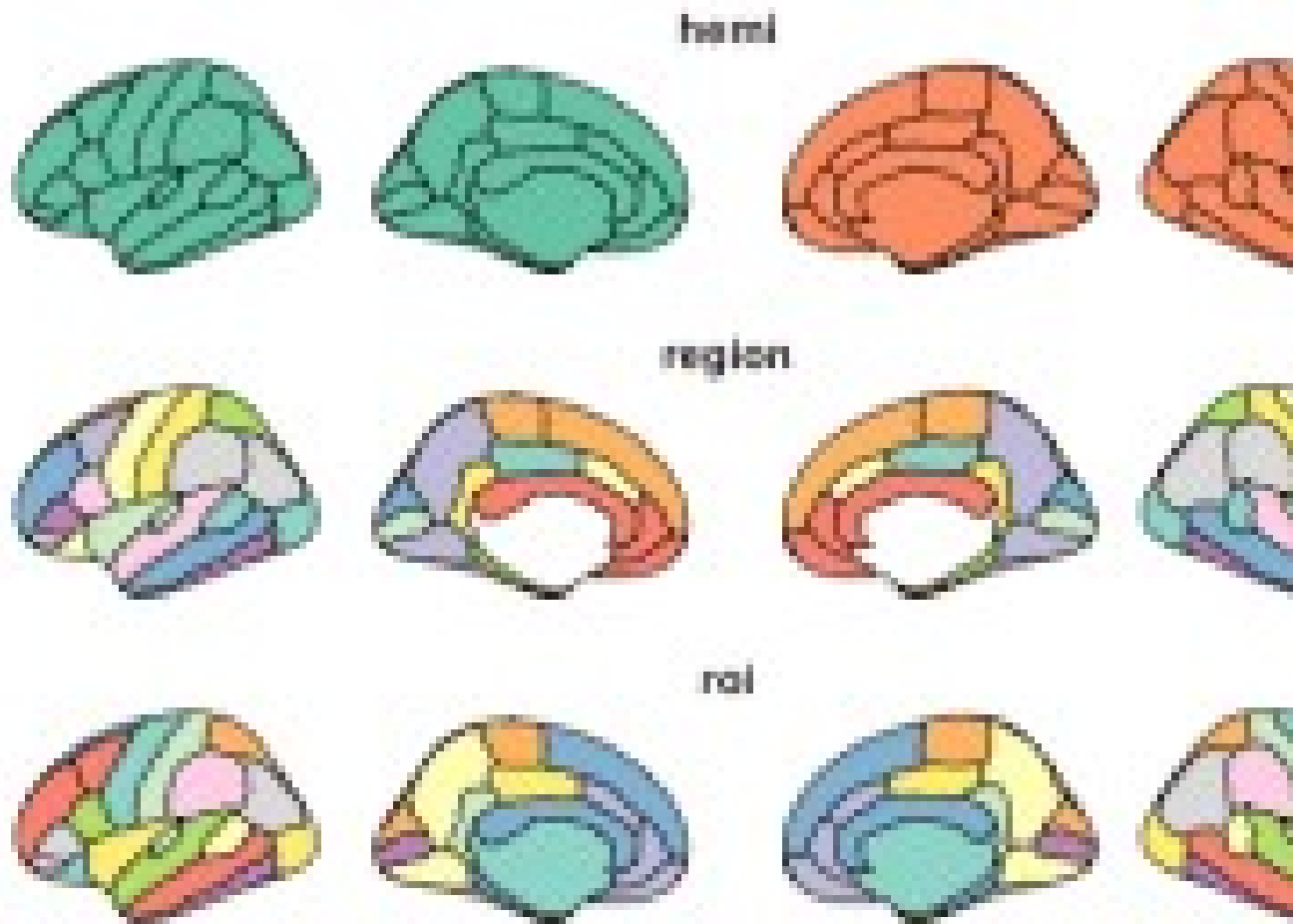
ggplot v0.0.3: Implements scale functions for setting area breaks for *ggplot2*. See the [vignette](#).



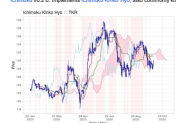
ggpr v0.4.0: Provides extensions to *ggplot2* to add trends to plots using both native and *ggpr* data coordinates. See the [vignette](#) for examples.



ggpr v1.0.0: Implements a *ggplot* geom for plotting brain atlases using simple features. The target component of the package is the data for two built-in atlases. See [Mouskoulas & Veld-Peters \(2020\)](#) for background. There is an [Introduction](#) along with *vignettes* on external data, *FreeSurfer files*, using *atlases*.



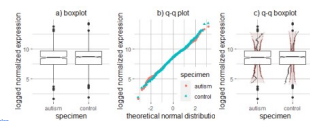
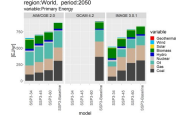
[shinobu v0.2.0](#) implements [Shinobu Kikuchi](#) [figs](#), also commonly known as [cloud charts](#), including static and interactive visualizations with tools for creating, backfitting and developing quantitative shinobu strategies. There is a [Reference](#) and a [signature](#) on [Stratagem](#).



[shinobu v0.1.0](#) provides functions for comparing interactive visualizations and creating linked interactive graphics for exploratory high-dimensional data analysis. See [Lee et al. \(2020\)](#) for background. There is a [signature](#) on [Lightning Non-linear Scatterplots](#) and another on the geometry of [Parameter Space](#).



[shinobu v0.1.1](#) provides generic functions to produce area, bar, box, and line plots following Integrated Assessment Modeling Consortium ([IAMC](#)) submission format in order to visualize climate migration scenarios. See the [signatures](#) for that stage.



[shinobu v0.1.0](#) implements Q-Q boxplots as an extension to [ggplot2](#). There is a [signature](#) on [Basic Usage](#) and another that provides [Examples](#).