

Package Ggthemes R

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Package Ggthemes R
ggBRT is designed for use with object of S3 class gbm, obtained from the gbm.step routine in the dismo package. Examples of each function using data from Jouffray et al. (2019) can be found in this ...

Explore and visualise the results of boosted regression trees
Build complex and customized plots from data in a data frame. We start by loading the required packages. ggplot2 is included in the tidyverse package. ggplot2 is a plotting package that provides ...

Data visualization with ggplot2
Build complex and customized plots from data in a data frame. We start by loading the required package. ggplot2 is also included in the tidyverse package. — Column specification ...

Data Visualisation with ggplot2
install.packages("drt") drat::add("epiforecasts") install.packages("covidregionaldata") install.packages("data.table") install.packages("ggplot2") install.packages ...

CoMix summary analysis for the United Kingdom
These files were imported into R and merged with a metadata file into a single Phyloseq object. Alpha diversity indices were computed using phyloseq package in R and the implementation of linear mixed ...

Gastrointestinal microbiota composition predicts peripheral inflammatory state during treatment of human tuberculosis
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Estrogen receptor—positive breast cancer survival prediction and analysis of resistance-related genes introduction
The variables in the dataset (after some processing shown below) are: estimates how much of the variation in herd size can be explained by all of these predictors combined (via the coefficient of ...

Hypothesis Testing
Analysis of HSP mRNA Levels in Pan-Cancer Tissues and Cells Deseq2 R package was used to analyze expression data in TCGA to identify differentially expressed HSP family members in pan-cancer tissues.

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