Package: epimdr2 (via r-universe)

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Title Functions and Data for ``Epidemics: Models and Data in R (2nd Edition)"

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Description Functions, data sets and shiny apps for ``Epidemics: Models and Data in R (2nd edition)" by Ottar N. Bjornstad (ISBN: 978-3-031-12055-8) <https://link.springer.com/book/9783031120558>. The package contains functions to study the S(E)IR model, spatial and age-structured SIR models; time-series SIR and chain-binomial stochastic models; catalytic disease models; coupled map lattice models of spatial transmission and network models for social spread of infection. The package is also an advanced quantitative companion to the Coursera Epidemics Massive Online Open Course <https://www.coursera.org/learn/epidemics>.

Depends R (>= 4.0.0), shiny, deSolve, plotly

Imports polspline, phaseR, ggplot2

Suggests ade4, bbmle, fields, forecast, imputeTS, lme4, ncf, nleqslv, nlme, nlts, plotrix, pomp, rootSolve, Rwave, rworldmap, statnet, scatterplot3d

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Encoding UTF-8

LazyData true

RoxygenNote 7.1.2

NeedsCompilation no

Config/pak/sysreqs make libicu-dev libssl-dev zlib1g-dev

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Repositoryhttps://epiverse-connect.r-universe.devRemoteUrlhttps://github.com/objornstad/epimdr2RemoteRefHEADRemoteShac359ca015f5c0aeadcbdf7da8cbe74cdf038e73d