## **Resource Summary Report**

Generated by <u>NIF</u> on May 25, 2025

# **EpiEstim**

RRID:SCR\_018538 Type: Tool

**Proper Citation** 

EpiEstim (RRID:SCR\_018538)

#### **Resource Information**

URL: https://cran.r-project.org/web/packages/EpiEstim/index.html

Proper Citation: EpiEstim (RRID:SCR\_018538)

**Description:** Framework and software to estimate time varying reproduction numbers during epidemics. Tools to quantify transmissibility throughout epidemic from analysis of time series of incidence. Used to estimate time varying instantaneous reproduction numbers from incidence time series.

Synonyms: Estimate Time Varying Reproduction Numbers from Epidemic Curves

**Resource Type:** software application, software toolkit, data analysis software, data processing software, software resource

Defining Citation: PMID:24043437

**Keywords:** Estimate time varying, reproduction number, epidemics, quantify transmissibility, time series analysis, reproductionc number analysis, incidence time series, bio.tools

Funding: United Kingdom Medical Research Council

Availability: Free, Available for download, Freely available

Resource Name: EpiEstim

Resource ID: SCR\_018538

Alternate IDs: biotools:EpiEstim

Alternate URLs: https://bio.tools/EpiEstim

Record Creation Time: 20220129T080340+0000

Record Last Update: 20250525T031558+0000

### **Ratings and Alerts**

No rating or validation information has been found for EpiEstim.

No alerts have been found for EpiEstim.

### Data and Source Information

Source: <u>SciCrunch Registry</u>

### **Usage and Citation Metrics**

We found 28 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

Kehoe AD, et al. (2024) SARS-CoV-2 Transmission in Alberta, British Columbia, and Ontario, Canada, January 2020-January 2022. Emerging infectious diseases, 30(5), 956.

Kabami Z, et al. (2024) Ebola disease outbreak caused by the Sudan virus in Uganda, 2022: a descriptive epidemiological study. The Lancet. Global health, 12(10), e1684.

Wei F, et al. (2023) COVID-19 transmission driven by age-group mathematical model in Shijiazhuang City of China. Infectious Disease Modelling, 8(4), 1050.

Dai C, et al. (2023) A new method for the joint estimation of instantaneous reproductive number and serial interval during epidemics. PLoS computational biology, 19(3), e1011021.

Wei Y, et al. (2022) Global COVID-19 Pandemic Waves: Limited Lessons Learned Worldwide over the Past Year. Engineering (Beijing, China), 13, 91.

Ji H, et al. (2022) The effectiveness of travel restriction measures in alleviating the COVID-19 epidemic: evidence from Shenzhen, China. Environmental geochemistry and health, 44(9), 3115.

Rui J, et al. (2022) Optimal control strategies of SARS-CoV-2 Omicron supported by invasive and dynamic models. Infectious diseases of poverty, 11(1), 115.

Zhou Z, et al. (2022) Estimation of heterogeneous instantaneous reproduction numbers with application to characterize SARS-CoV-2 transmission in Massachusetts counties. PLoS

computational biology, 18(9), e1010434.

Chen Z, et al. (2022) Epidemiological characteristics and transmission dynamics of the outbreak caused by the SARS-CoV-2 Omicron variant in Shanghai, China: a descriptive study. medRxiv : the preprint server for health sciences.

Chen Z, et al. (2022) Epidemiological characteristics and transmission dynamics of the outbreak caused by the SARS-CoV-2 Omicron variant in Shanghai, China: A descriptive study. The Lancet regional health. Western Pacific, 29, 100592.

Petrie JG, et al. (2022) The variant-specific burden of SARS-CoV-2 in Michigan: March 2020 through November 2021. Journal of medical virology, 94(11), 5251.

Siddle KJ, et al. (2022) Transmission from vaccinated individuals in a large SARS-CoV-2 Delta variant outbreak. Cell, 185(3), 485.

Huisman JS, et al. (2022) Estimation and worldwide monitoring of the effective reproductive number of SARS-CoV-2. eLife, 11.

Hukic M, et al. (2021) SARS-CoV-2 virus outbreak and the emergency public health measures in Bosnia and Herzegovina: January - July, 2020. Bosnian journal of basic medical sciences, 21(1), 111.

Dainton C, et al. (2021) Quantifying the relationship between lockdowns, mobility, and effective reproduction number (Rt) during the COVID-19 pandemic in the Greater Toronto Area. BMC public health, 21(1), 1658.

Giovanetti M, et al. (2021) SARS-CoV-2 shifting transmission dynamics and hidden reservoirs potentially limit efficacy of public health interventions in Italy. Communications biology, 4(1), 489.

Zhao H, et al. (2021) Transmission dynamics and successful control measures of SARS-CoV-2 in the mega-size city of Guangzhou, China. Medicine, 100(48), e27846.

Komissarov AB, et al. (2021) Genomic epidemiology of the early stages of the SARS-CoV-2 outbreak in Russia. Nature communications, 12(1), 649.

Chu J, et al. (2021) A statistical analysis of the novel coronavirus (COVID-19) in Italy and Spain. PloS one, 16(3), e0249037.

Chadsuthi S, et al. (2021) Modelling the effectiveness of intervention strategies to control COVID-19 outbreaks and estimating healthcare demand in Germany. Public health in practice (Oxford, England), 2, 100121.