Package ‘shinySIR’

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Type Package

Title Interactive Plotting for Mathematical Models of Infectious Disease Spread

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Description Provides interactive plotting for mathematical models of infectious disease spread. Users can choose from a variety of common built-in ordinary differential equation (ODE) models (such as the SIR, SIRS, and SIS models), or create their own. This latter flexibility allows ‘shinySIR’ to be applied to simple ODEs from any discipline. The package is a useful teaching tool as students can visualize how changing different parameters can impact model dynamics, with minimal knowledge of coding in R. The built-in models are inspired by those featured in Keeling and Rohani (2008) <doi:10.2307/j.ctvcm4gk0> and Bjornstad (2018) <doi:10.1007/978-3-319-97487-3>.

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Encoding UTF-8

LazyData true

RoxygenNote 6.1.1

Depends dplyr (>= 0.8.0.1), tidyr (>= 0.8.3), ggplot2 (>= 3.1.1), shiny (>= 1.3.2), deSolve (>= 1.2.1)

Suggests knitr (>= 1.22), rmarkdown (>= 1.12), testthat (>= 2.2.0)

VignetteBuilder knitr

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Description

This function prints a list of all built in models, along with their parameter arguments and a short description.

Usage

```r
default_models()
```

Value

- data frame of model descriptions.

Examples

```r
default_models()
```
get_ics

Get default initial conditions

Description
This function returns the default parameter vectors for a particular built-in model.

Usage
get_ics(model)

Arguments
model name of the model to be solved. Examples include: SIR and SIR vaccination.

Value
named vector of default initial conditions.

get_name

Get model display names

Description
This function gets the display name for any built-in model.

Usage
get_name(model)

Arguments
model character specifying the name of the built-in model.

Value
character of the corresponding display name.

Examples
get_name(model = "SIR")
get_params

Get default parameters

Description
This function returns the default parameter vectors for a particular built-in model.

Usage
get_params(model)

Arguments
model name of the model to be solved. Examples include: SIR and SIR vaccination.

Value
list of default parameter vectors.

Examples
get_params(model = "SIR")

plot_model

Plot model output. This function plots the output of a fitted model data frame.

Description
Plot model output. This function plots the output of a fitted model data frame.

Usage
plot_model(output, linesize, textsize, xlabel, ylabel, legend_title,
levels, values, ...)

Arguments
output data frame output from solve_eqns().
linesize numeric value for line width in ggplot.
textsize numeric value for textsize in ggplot.
xlabel character string for x axis label.
ylabel character string for y axis label.
legend_title character string for legend title.
run_shiny

levels character vector of the variable names in the order they should be plotted. Default is to obtain the order from the initial conditions vector 'ics'.

values vector specifying manual color scale. Length must equal the number of model variables.

... extra argument to be passed through to ggplot scale_colour_manual: use 'labels' to change the legend names

Value

ggplot object

Description

This function solves an ODE model using 'deSolve' and returns the output as a data frame.

Usage

run_shiny(model = "SIR", neweqns = NULL, ics = NULL, tstart = 0,
timestep = 1, tmax = 365, parm0 = NULL, parm_names = NULL,
parm_min = NULL, parm_max = NULL, sigfigs = 4, showtable = TRUE,
linesize = 1.2, textsize = 14, xlabel = "Time",
ylabel = "Number of individuals", legend_title = "Compartment",
slider_steps = NULL, values = NULL, ...)

Arguments

model name of the model to be solved. Examples of built-in models are: "SIR", "SIR vaccination". Default is "SIR".

neweqns function specifying the equations of the user-defined model. Only to be used if a model is required that is not built-in. Default is NULL.

ics named numeric vector specifying the initial conditions i.e. the initial values of all model variables. Default is c(S = 9999, I = 1, R = 0) for the SIR model.

tstart numerical value of form c(tmin, tmax) indicating the time to start simulations. Default value is 0.

timestep numerical value indicating time step be used when solving equations. Default value is 1/365.

tmax numerical value indicating maximum time point to be considered.

parm0 named numeric vector of starting parameter values. Names must correspond to those used in the model equations.

parm_names character vector of parameter names to be displayed in shiny menu. Must be in the same order as 'parm0'.
parm_min  named numeric vector of minimum parameter values.
parm_max  named numeric vector of maximum parameter values.
sigfigs   number of significant figures to round parameter input vectors. Default is 4.
showtable logical TRUE/FALSE. Should the table of transformed parameters be shown? Only applies to built-in models. Default is TRUE.
linesize numeric value for line width in ggplot output. Default is 1.2.
textsize numeric value for textsize in ggplot output. Default is 14.
xlabel    character string for x axis plotting label. Default is "Time".
ylabel    character string for y axis plotting label. Default is "Number of individuals".
legend_title character string for legend title. Default is "Compartment".
slider_steps numeric vector of step size to include between slider input values. Should be NULL or a vector with an entry for each parameter input. Default is NULL.
values    vector specifying manual color scale (if desired). Length must equal the number of model variables.
... extra argument to be passed through to ggplot scale_colour_manual: use 'labels' to change the legend names.

Value

data frame of model solutions in long format.

Examples

run_shiny(model = "SIR")

---

seir.app  Launch a shiny-app simulating the seasonal SEIR model

Description

# ' This launches an app running the SEIR model i.e. a model incorporating latency and seasonal forcing in transmission.

Usage

seir.app

Format

An object of class shiny.appobj of length 5.
seirs.app

**Details**
Launch app for details

**Examples**
```r
## Not run: seir.app
```

---

**seirs.app**

*Launch a shiny-app simulating the SEIRS model*

---

**Description**
This launches an app running the SEIRS model i.e. a model incorporating latency and loss of immunity.

**Usage**

```r
seirs.app
```

**Format**
An object of class `shiny.appobj` of length 5.

**Details**
Launch app for details

**Examples**
```r
## Not run: seirs.app
```

---

**SIR**

*SIR model*

---

**Description**
These equations describe the classic SIR model with no births or deaths.

**Usage**

```r
SIR(t, y, parms)
```

**Arguments**

- `t` numeric vector of time points.
- `y` numeric vector of variables.
- `parms` named vector of model parameters.
**SIRbirths**  
*SIR model with demography*

**Description**

These equations describe the classic SIR model with equal births and deaths.

**Usage**

SIRbirths(t, y, parms)

**Arguments**

- **t**: numeric vector of time points.
- **y**: numeric vector of variables.
- **parms**: named vector of model parameters.

**Value**

equation list

---

**SIRS**  
*SIRS model*

**Description**

These equations describe the classic SIRS model without births or deaths.

**Usage**

SIRS(t, y, parms)

**Arguments**

- **t**: numeric vector of time points.
- **y**: numeric vector of variables.
- **parms**: named vector of model parameters.

**Value**

equation list
**SIRSbirths**  

**Description**  
These equations describe the classic SIRS model with equal birth and death rates.

**Usage**  

\[ \text{SIRSbirths}(t, y, \text{parms}) \]

**Arguments**  
- \( t \) numeric vector of time points.  
- \( y \) numeric vector of variables.  
- \( \text{parms} \) named vector of model parameters.

**Value**  
equation list

---

**SIRSvaccination**  

**Description**  
These equations describe the classic SIRS model with equal birth and death rates and vaccination at birth.

**Usage**  

\[ \text{SIRSvaccination}(t, y, \text{parms}) \]

**Arguments**  
- \( t \) numeric vector of time points.  
- \( y \) numeric vector of variables.  
- \( \text{parms} \) named vector of model parameters.

**Value**  
equation list
**SIRvaccination**  
*SIR model with vaccination at birth*

**Description**
These equations describe the classic SIR model with births and deaths, constant population size, and (optional) vaccination at birth.

**Usage**

```r
SIRvaccination(t, y, parms)
```

**Arguments**
- **t**  
  numeric vector of time points
- **y**  
  numeric vector of variables
- **parms**  
  named vector of model parameters.

**Value**
equation list

---

**SIS**  
*SIS model*

**Description**
These equations describe the classic SIS model with no births or deaths.

**Usage**

```r
SIS(t, y, parms)
```

**Arguments**
- **t**  
  numeric vector of time points
- **y**  
  numeric vector of variables
- **parms**  
  named vector of model parameters.

**Value**
equation list
SISbirths  

Description  
These equations describe the classic SIR model with equal births and deaths.

Usage  
SISbirths(t, y, parms)

Arguments  
t numeric vector of time points.  
y numeric vector of variables.  
parms named vector of model parameters.

Value  
equation list

solve_eqns  
Solve equations

Description  
This function solves an ODE model using ‘deSolve’ and returns the output as a data frame.

Usage  
solve_eqns(eqns, ics, times, parms)

Arguments  
eqns name of the model to be solved. Examples include: SIR and SIR vaccination.  
ics named numeric vector specifying the initial conditions i.e. the initial values of all model variables.  
times numerical vector indicating the time points at which the equation should be solved.  
parms named numeric vector of parameter values.

Value  
data frame of model solutions in long format.
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