

Package ‘attrib’

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Title Attributable Burden of Disease

Version 2021.1.2

Description Provides functions for estimating the attributable burden of disease due to risk factors. The posterior simulation is performed using `arm::sim` as described in Gelman, Hill (2012) <[doi:10.1017/CBO9780511790942](https://doi.org/10.1017/CBO9780511790942)> and the attributable burden method is based on Nielsen, Krause, Molbak <[doi:10.1111/irv.12564](https://doi.org/10.1111/irv.12564)>.

Depends R (>= 3.5.0)

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Encoding UTF-8

LazyData true

Imports data.table, magrittr, glue, pbs, dlnm, lubridate, mvmeta, tsModel, stats, lme4, arm, tibble, stringr, ggplot2, utils, progress

Suggests testthat, knitr, rmarkdown

RoxygenNote 7.1.1

VignetteBuilder knitr

NeedsCompilation no

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data_fake_county	<i>Fake data for mortality in Norway</i>
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Description

Fake data for mortality in Norway

Usage

data_fake_county

Format

location_code Location code of the Norwegian municipalities

week Week

season Season used for influenza like illnesses

yrwk Year and week

x Number of weeks from the start of the season

pop Population size

pr100_ili Per hundred ILI, percentage of consultations diagnosed as influenza like illnesses

pr100_ili_lag_1 pr100_ili_lag_1

temperature temperature

temperature_high temperature_high

deaths deaths

data_fake_nation	<i>Fake data for mortality in Norway nationally</i>
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Description

Fake data for mortality in Norway nationally

Usage

data_fake_nation

Format

location_code Location code
week Week
season Season used for influenza like illnesses
yrwk Year and week
x Number of weeks from the start of the season
pop Population size
pr100_ili Per hundred ILI, percentage of consultations diagnosed as influenza like illnesses
pr100_ili_lag_1 pr100_ili_lag_1
temperature temperature
temperature_high temperature_high
deaths deaths

 est_attrib

Estimates simulations of expected responses

Description

For each exposure the dataset is copied and the original value replaced by the reference value. Then the sim function is used to generate 500 simulations of expected responses for each row. Finally the dataset is transformed to obtain expected response for original and reference values of the given exposures for each original row of the dataset.

Usage

```
est_attrib(fit, data, exposures, n_sim = 500)
```

Arguments

fit	A model fit constructed by fit_attrib
data	The observed data
exposures	The exposures that will get reference expected mortalities
n_sim	Number of simulations

For more details see the help vignette: `vignette("intro", package="attrib")`

Details

The burden method is based on Nielsen, Krause, Molbak <doi:10.1111/irv.12564>.

For more details see the help vignette: `vignette("intro", package="attrib")`

Value

Dataset with expected responses for all simulations including expected responses given the exposure reference values

Examples

```

response <- "deaths"
fixef <- "pr100_ili_lag_1 + sin(2 * pi * (week - 1) / 52) + cos(2 * pi * (week - 1) / 52)"
ranef <- " (pr100_ili_lag_1| season)"
offset <- "log(pop)"

data <- attrib::data_fake_nation

fit <- fit_attrib(data = data, response = response, fixef = fixef, ranef = ranef, offset = offset)
exposures <- c(pr100_ili_lag_1 = 0)
n_sim <- 5
new_data <- est_attrib(fit, data, exposures, n_sim)
new_data[]

```

fit_attrib

Data fit

Description

Data fit using glmer from lme4 with family poisson to fit the dataset with the given formula.

Usage

```
fit_attrib(data, response, fixef, ranef, offset = NULL)
```

Arguments

data	The observed data to be fitted.
response	The response
fixef	The fixed effects
ranef	The random effects
offset	The offsets.

Value

The model fit of the data with additional attributes offset, response and fit_fix. Offset and response are the same as in the input and fit_fix is the linear model of the fix effects.

For more details see the help vignette: vignette("intro", package="attrib")

Examples

```
response <- "deaths"

fixef <- "pr100_ili_lag_1 + sin(2 * pi * (week - 1) / 52) + cos(2 * pi * (week - 1) / 52)"
ranef <- " (pr100_ili_lag_1| season)"
offset <- "log(pop)"

data <- attrib::data_fake_nation

fit_attrib(data = data, response = response, fixef = fixef, ranef = ranef, offset = offset)
```

sim	<i>Generates simulations of expected mortality by simulating the model coefficients.</i>
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Description

With the given fit from `fit_attrib` the function `sim`, from package `arm` as described in Gelman, Hill (2012) <doi:10.1017/CBO9780511790942>, is used to generate 500 simulations of all the coefficients, from their respective posterior distributions. This is then used to compute the expected response for all simulations and rows in the input dataset.

Usage

```
sim(fit, data, n_sim)
```

Arguments

<code>fit</code>	A model fit created by <code>fit_attrib</code>
<code>data</code>	The data with either observed values or reference values.
<code>n_sim</code>	Number of simulations

Details

```
vignette("intro", package="attrib")
```

Value

A dataset with 500 simulations of the expected response for each row in the original dataset.

Examples

```
response <- "deaths"
fixef <- "pr100_ili_lag_1 + sin(2 * pi * (week - 1) / 52) + cos(2 * pi * (week - 1) / 52)"
ranef <- " (pr100_ili_lag_1| season)"
offset <- "log(pop)"

data <- attrib::data_fake_nation

fit <- fit_attrib(data = data, response = response, fixef = fixef, ranef = ranef, offset = offset)

n_sim <- 5
sim(fit, data, n_sim)
```

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