# **Package**

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data.agespecificfert Age-specific fertility rate

# Description

Dataset containing age-specific fertility rate by country, age range, year and gender

# Usage

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data.agespecificfert

# **Format**

A data frame with 20580 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value fertility rate
```

#### **Source**

data.births 3

data.births

Number of Births

# Description

Dataset containing number of births by country, age range, year and gender

# Usage

data.births

## **Format**

A data frame with 14700 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value number of births
```

## Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.birthsbymaternalage

Number of births by age of mother

# Description

Dataset with the number of births by maternal by country, year and gender

# Usage

data.birthsbymaternalage

#### **Format**

```
A data frame with 20580 observations on the following 8 variables.
```

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value number of births
```

#### Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

```
data.birthsquinquennial
```

Number of births - quinquennial

# Description

Dataset containing number of births occuring every 5 years by country, age range, year and gender

# Usage

```
data.birthsquinquennial
```

# Format

A data frame with 2940 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value number of births occuring every 5 years
```

#### **Source**

data.centraldeathrateASMR

data.centraldeathrateASMR

Central death rate

# Description

Dataset containing central death rate by country, age range, year and gender

# Usage

data.centraldeathrateASMR

#### **Format**

A data frame with 55860 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value central death rate
```

#### Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

# Description

Dataset containing crude birth rate by country, age range, year and gender

# Usage

data.crudebirthrate

6 data.crudedeathrate

#### **Format**

```
A data frame with 14700 observations on the following 8 variables.
```

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value crude birth rate
```

## Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.crudedeathrate

Crude death rate

## **Description**

Dataset containing crude death rate (CDR) by country, age range, year and gender

# Usage

data.crudedeathrate

## **Format**

A data frame with 14700 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value crude death rate
```

### Source

data.deathsbyage 7

data.deathsbyage

Number of deaths by age

# Description

Dataset containing number of deaths by country, age range, year and gender

# Usage

data.deathsbyage

#### **Format**

A data frame with 58800 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value number of deaths
```

#### Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.growthrate

Growth rate

## **Description**

Dataset containing growth rate by country, age range, year and gender

# Usage

data.growthrate

## **Format**

A data frame with 2940 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value growth rate
```

8 data.interpolatedpop

#### **Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.Hib\_DALY

Disability weights for Haemophilus influenzae type B

## **Description**

Dataset disability weights for Haemophilus influenzae type B by condition/sequelae

## Usage

```
data.Hib_DALY
```

#### **Format**

A data frame with 24 observations on the following 3 variables.

Disease disease

Condition condition/sequelae

GBD\_2015\_mean Mean disability weight

#### **Source**

General Guidance for DALYs calculation VIMC with input from DOVE 2017-11-24 11:03:46

Interpolated population (1-year time and age)

## **Description**

Dataset containing population - interpolated (1-year time and age) by country, age range, year and gender

## Usage

```
data.interpolatedpop
```

## **Format**

A data frame with 170480 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value population interpolated 1-year time and age
```

data.lifeexpectancy 9

#### Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.lifeexpectancy Life expectancy at birth

## **Description**

Dataset containing life expectancy at birth by country, age range, year and gender

# Usage

```
data.lifeexpectancy
```

#### **Format**

A data frame with 1470 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value expected remaining years of life
```

## **Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

```
data.neonatalmortality
```

Neonatal mortality rate

# **Description**

Dataset containing 28-day neonatal mortality rate by country, age range, year and gender

# Usage

```
data.neonatalmortality
```

10 data.netmigration

#### **Format**

```
A data frame with 14700 observations on the following 8 variables.
```

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value 28-day neonatal mortality rate
```

#### **Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.netmigration

Net migration rate

## **Description**

Dataset containing net migration rate by country, age range, year and gender

## Usage

```
data.netmigration
```

#### **Format**

A data frame with 2940 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value net migration rate
```

## Source

data.PCV\_DALY

cine)	data.PCV_DALY	Disability weights for Pneumococcus (Pneumococcal conjugate vaccine)
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### **Description**

Dataset disability weights for PCV by condition/sequelae

## Usage

```
data.PCV_DALY
```

#### **Format**

A data frame with 27 observations on the following 3 variables.

Disease disease

Condition condition/sequelae

GBD\_2015\_mean Mean disability weight

#### Source

General Guidance for DALYs calculation VIMC with input from DOVE 2017-11-24 11:03:46

data.pdeathbyage

Probability of dying by age

## **Description**

Dataset containing probability of dying by age by country, age range, year and gender

#### Usage

```
data.pdeathbyage
```

#### **Format**

A data frame with 52920 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value probability of dying
```

# Source

data.quinquennialpop Quinquennial population (5-year time and age)

# Description

Dataset containing population - quinquennial population (5-year time and age) by country, age range, year and gender

## Usage

data.quinquennialpop

#### **Format**

A data frame with 71442 observations on the following 8 variables.

```
country_code_numeric a numeric vector

country_code country code

country name of country

age_from starting age

age_to end age

year year

gender gender

value population quinquennial 5-year time and age
```

## **Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

```
data.remainingyearsoflife

Expected remaining years of life
```

# Description

Dataset containing expected remaining years of life by country, age range, year and gender

# Usage

```
data.remainingyearsoflife
```

#### **Format**

A data frame with 64680 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value expected remaining years of life
```

#### **Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

```
data.rotacases_nonsevereD1
```

Event rates (cases) for Rotavirus D1 non-severe RVGE

# Description

Dataset containing event rates (cases) for rotavirus D1 for non-severe RVGE

#### Usage

```
data.rotacases_nonsevereD1
```

#### **Format**

A data frame with 195 observations on the following 8 variables.

```
Country country
WHO region WHO region
WHO region2 WHO region country code
Income income level
Age age
Mid mid
Low low
High high
```

## **Source**

Bilcke J. et al. Estimating the Incidence of Symptomatic Rotavirus Infections: A Systematic Review and Meta-Analysis. PLOS One, June 2009, Volume 4, Issue 6, e6060. Note: random effects model resulted in a global incidence estimate of 0.24 [0.17; 0.34] symptomatic RV infections per person year of observation for children below 2 years of age. Crudely extrapolating to children aged <5yrs, and assuming minimal incidence aged 2+yrs, gives an under-five incidence rate of 0.10 [0.07 - 0.14] or 10,000 [7,000 - 14,000] per 100,000 per year <5yrs. Severe incidence rates derived from Fischer-Walker (see source for severe RVGE incidence), were then subtracted to give non-severe RVGE incidence.

data.rotacases\_severeD2

Event rates (cases) for Rotavirus D2 severe RVGE

## **Description**

Dataset containing event rates (cases) for rotavirus D2 for severe RVGE

#### Usage

data.rotacases\_severeD2

#### **Format**

A data frame with 195 observations on the following 8 variables.

Country country

WHO region WHO region

WHO region 2 WHO region country code

Income income level

Age age

Mid mid

Low low

High high

## Source

Fischer-Walker C. et al, Table 1: Global and regional burden of diarrhoea and pneumonia per year in children aged 0–4 years, by WHO region. Global burden of childhood pneumonia and diarrhoea. Lancet 2013; 381: 1405–16. Notes: Episodes per child per year <5yrs (2010) by WHO region were multiplied by the proportion of episodes that were severe by WHO region (approximately 2 by the rotavirus-positive proportion <5yrs by WHO region, reported among hospitalised diarrhoea cases in CHERG (Lanata C. et al, Global Causes of Diarrheal Disease Mortality in Children <5 Years of Age: A Systematic Review. PLOS One. September 2013, Volume 8, Issue 9, e72788). The uncertainty range only reflects uncertainty in the incidence of diarrhoea episodes.

data.rotadeaths\_severeD2

Event rates (deaths) for Rotavirus D2 severe RVGE

## **Description**

Dataset containing event rates (deaths) for rotavirus D2 for severe RVGE

### Usage

data.rotadeaths\_severeD2

#### **Format**

A data frame with 195 observations on the following 8 variables.

Country country

WHO region WHO region

WHO region 2 WHO region country code

**Income** income level

Age age

Mid mid

Low low

High high

## Source

See Clark et al, TRIVAC, Vaccine, Appendix

data.rotahosps\_severeD2

Event rates (hospital) for Rotavirus D2 severe RVGE

# Description

Dataset containing event rates (hospitals) for rotavirus D2 for severe RVGE

# Usage

data.rotahosps\_severeD2

## **Format**

A data frame with 195 observations on the following 8 variables.

Country country

WHO region WHO region

WHO region 2 WHO region country code

Income income level

Age age

Mid mid

Low low

High high

#### **Source**

See Clark et al, TRIVAC, Vaccine, Appendix

data.rotavisits\_nonsevereD1

Event rates (visits) for Rotavirus D1 non-severe RVGE

# Description

Dataset containing event rates (visits) for rotavirus D1 for non-severe RVGE

# Usage

```
data.rotavisits_nonsevereD1
```

#### **Format**

A data frame with 195 observations on the following 8 variables.

Country country

WHO region WHO region

WHO region 2 WHO region country code

**Income** income level

Age age

Mid mid

Low low

High high

## **Source**

Assume 1 visit for every 2 non-severe cases

data.rotavisits\_severeD2

Event rates (visits) for Rotavirus D2 severe RVGE

# Description

Dataset containing event rates (visits) for rotavirus D2 for severe RVGE

# Usage

```
data.rotavisits_severeD2
```

data.Rota\_DALY

#### **Format**

A data frame with 195 observations on the following 8 variables.

Country country

WHO region WHO region

WHO region 2 WHO region country code

Income income level

Age age

Mid mid

Low low

High high

#### **Source**

Clark A et al, unpublished update of the IHME/MCEE/WHOCDC estimates. Median/Min/Max 2015 estimates were used or the estimate for the most recent pre-vax year (using WUENIC 15th July 2017).

data.Rota\_DALY

Disability weights for Rotavirus

## Description

Dataset disability weights for Rotavirus by condition/sequelae

## Usage

data.Rota\_DALY

## **Format**

A data frame with 3 observations on the following 3 variables.

Disease disease

Condition condition/sequelae

GBD\_2015\_mean Mean disability weight

## Source

General Guidance for DALYs calculation VIMC with input from DOVE 2017-11-24 11:03:46

18 data.survival

data.sexratio

Sex ratio at birth

## **Description**

Dataset containing sex-ratio at birth by country, age range, year and gender

#### Usage

```
data.sexratio
```

#### **Format**

A data frame with 2940 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value sex-ratio
```

#### **Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.survival

Survivors from a birth-cohort of 100k

# Description

A data frame with 64680 observations on the following 8 variables.

## Usage

```
data.survival
```

## **Format**

A data frame with 20580 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value number of survivors
```

data.totaldeaths 19

#### Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.totaldeaths

Deaths in total

# Description

Dataset containing total number of deaths by country, age range, year and gender

# Usage

data.totaldeaths

## **Format**

A data frame with 14700 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value number of deaths
```

## Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.totalfert

Total fertility rate

# Description

Dataset containing total fertility rate by country, age range, year and gender

# Usage

data.totalfert

20 data.totalpop

#### **Format**

```
A data frame with 14700 observations on the following 8 variables.
```

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value total fertility rate
```

# Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.totalpop

Total population

## **Description**

Dataset containing total population country, age range, year and gender

## Usage

```
data.totalpop
```

### **Format**

A data frame with 14798 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value total population
```

## Source

data.u1mortality 21

data.u1mortality

Under 1 mortality rate

# Description

Dataset containing under 1 mortality rate by country, age range, year and gender

# Usage

```
data.u1mortality
```

#### **Format**

A data frame with 14700 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value under 1 mortality rate
```

#### Source

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.u5mortality

Under 5 mortality rate

## **Description**

Dataset containing under 5 mortality rate by country, age range, year and gender

# Usage

```
data.u5mortality
```

## **Format**

A data frame with 20580 observations on the following 8 variables.

```
country_code_numeric a numeric vector
country_code country code
country name of country
age_from starting age
age_to end age
year year
gender gender
value fertility rate
```

22 univacr

#### **Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

data.vaccine\_schedules

Disability weights for Haemophilus influenzae type B

## **Description**

Dataset containing disability weights for Haemophilus influenzae type B by condition/sequelae (UNIVAC model)

### Usage

data.vaccine\_schedules

#### **Format**

A data frame with 195 observations on the following 6 variables.

Country country

BCG BCG target age in weeks

DTP1 DTP1 target age in weeks

DTP2 DTP2 target age in weeks

DTP3 DTP3 target age in weeks

Meas1 Measles target age in weeks

# Source

www.who.int/immunization/monitoring\_surveillance/data/schedule\_data.xls

univacr

univacr: UNIVAC decision support model

## Description

A universal framework for evaluating vaccine policy options in low- and middle-income countries. This R package is based of the spreadsheet-based tool that is accessible at https://www.paho.org/provactoolkit/.

## Vaccine impact model

Estimate the health impact and cost-effectiveness of vaccination.

Hib, pneumococcal and rotavirus vacccination can be separately analysed.

writelog 23

writelog

Simulation log reporting

# Description

Appends message of simulation run (x) to log file (logname).

# Usage

```
writelog(logname, x)
```

# Arguments

logname log filename

x message of simulation run

# Value

None

# **Examples**

#

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