

# Package ‘univacr’

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**Title** UNIVAC -- Vaccine policy decision support model

**Version** 2.0.1

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**Description** UNIVAC is a universal framework for evaluating vaccine policy options in low- and middle-income countries.

**Depends** R (>= 3.4.3)

**Imports** data.table, foreach

**License** GPL-3

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.1.1

**Suggests** knitr,  
rmarkdown

**VignetteBuilder** knitr

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data.agespecificfert	<i>Age-specific fertility rate</i>
----------------------	------------------------------------

---

## Description

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

## Usage

```
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

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

data.births	<i>Number of Births</i>
-------------	-------------------------

---

**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	<i>Number of births by age of mother</i>
--------------------------	--

---

**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 occurring 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 occurring every 5 years

**Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

data.centraldeathrateASMR	<i>Central death rate</i>
---------------------------	---------------------------

---

**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)

---

data.crudebirthrate	<i>Crude birth rate (CBR)</i>
---------------------	-------------------------------

---

**Description**

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

**Usage**

```
data.crudebirthrate
```

**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.crudeddeathrate	<i>Crude death rate</i>
----------------------	-------------------------

---

**Description**

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

**Usage**

```
data.crudeddeathrate
```

**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**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

data.deathsbyage	<i>Number of deaths by age</i>
------------------	--------------------------------

---

**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	<i>Growth rate</i>
-----------------	--------------------

---

**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

**Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

data.Hib_DALY	<i>Disability weights for Haemophilus influenzae type B</i>
---------------	---

---

**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

---

data.interpolatedpop	<i>Interpolated population (1-year time and age)</i>
----------------------	--

---

**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



**Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

data.lifeexpectancy	<i>Life expectancy at birth</i>
---------------------	---------------------------------

---

**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	<i>Neonatal mortality rate</i>
------------------------	--------------------------------

---

**Description**

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

**Usage**

```
data.neonatalmortality
```

**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	<i>Net migration rate</i>
-------------------	---------------------------

---

**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**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

data.PCV_DALY	<i>Disability weights for Pneumococcus (Pneumococcal conjugate vaccine)</i>
---------------	---

---

**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	<i>Probability of dying by age</i>
------------------	------------------------------------

---

**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**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

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 region2** 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 region2** 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 region2** 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 region2** 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



**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**

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

---

data.sexratio	<i>Sex ratio at birth</i>
---------------	---------------------------

---

### 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	<i>Survivors from a birth-cohort of 100k</i>
---------------	--

---

### 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

**Source**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

data.totaldeaths	<i>Deaths in total</i>
------------------	------------------------

---

**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	<i>Total fertility rate</i>
----------------	-----------------------------

---

**Description**

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

**Usage**

```
data.totalfert
```

**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	<i>Total population</i>
---------------	-------------------------

---

**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**

Demographic data from the Vaccine Impact Modelling Consortium (VIMC)

---

data.u1mortality	<i>Under 1 mortality rate</i>
------------------	-------------------------------

---

**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	<i>Under 5 mortality rate</i>
------------------	-------------------------------

---

**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

**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](http://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/provac-toolkit/>.

**Vaccine impact model**

**Estimate the health impact and cost-effectiveness of vaccination.**

**Hib, pneumococcal and rotavirus vaccination can be separately analysed.**

---

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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