Package 'prime'

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Title PRIME -- Papillomavirus Rapid Interface for Modelling and Economics

Version 2.0.1

Author Modelling and Economic Evaluation of Vaccines group, LSHTM

Maintainer http://meev.lshtm.ac.uk/

Description PRIME is a static model of HPV vaccination that uses proportional impact to estimate the health impact and cost-effectiveness of HPV vaccination 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

R topics documented:

`ageCoverage` *Get age-specific coverage-rates*

Description

Get age-specific coverage-rates

Usage

```
ageCoverage(ages, routine_coverage, vaccine_efficacy_nosexdebut,
  vaccine_efficacy_sexdebut, campaigns, lifetab, cohort, agevac,
  country_iso3 = NULL)
```

Arguments

`ages`	Numeric vector (required): ages in model
`routine_coverage`	Number (required): proportion of population that receives routine vaccination
`campaigns`	List or number (required): if a list, applies MAC vaccination (needs to change)
`lifetab`	Data.table (required): lifetable generated with lifeTable()
`cohort`	Number (required): cohort-size (only used in MAC campaigns)
`agevac`	Number (required): target age for vaccination
`vaccine_efficacy`	Number (required): proportion indicating vaccine-efficacy

Value

Data.table with coverage and effective coverage by age. Used in RunCohort()

Examples

```
ages <- c(0:100)
routine_coverage <- 0.75
vaccine_efficacy <- 0.8
lifetab <- lifeTable(unlist(data.mortall[iso3=="AFG", as.character(0:100), with=F], use.names=F), 9)
cohort <- unlist(data.popproj[iso3=="AFG", "2020"], use.names=F)
agevac <- 9
ageCoverage(ages, routine_coverage, vaccine_efficacy, -1, lifetab, cohort, agevac)
```

---

analyseCosts *Returns cost-effectiveness for a single birthcohort in a single country*

Description

Usually called using RunCountry(..., analyseCosts=TRUE)

Usage

```
analyseCosts(results, vaccine_cost, gdp_per_capita)
```

Arguments

results	Data.table (required): results from RunCohort()
vaccine_cost	Number (required): cost of a single vaccine
gdp_per_capita	Number (required): GDP per capita

Value

Data.table with cost-analysis

Examples

```
analyseCosts(RunCountry("AFG"), 100, 561)
```

BatchRun *Run multiple cohorts in a batch*

Description

Runs multiple cohorts in one batch, based on the data in .data.batch

Usage

```
BatchRun(countries = -1, coverage = -1, agevac = -1,
  agecohort = -1, canc.inc = "2018", daly.canc.diag = 0.288,
  daly.canc.control = 0.049, daly.canc.metastatic = 0.451,
  daly.canc.terminal = 0.54, sens = -1, unwpp_mortality = FALSE,
  year_born = -1, year_vac = -1, runs = 1,
  vaccine_efficacy_beforesexdebut = 1,
  vaccine_efficacy_aftersexdebut = 0, log = -1,
  by_calendaryear = FALSE, use_proportions = TRUE,
  analyseCosts = FALSE, psa = 0, psa_vals = ".data.batch.psa")
```

Arguments

countries	ignore, read from .data.batch
coverage	ignore, read from .data.batch
agevac	ignore, read from .data.batch
agecohort	ignore, read from .data.batch
canc.inc	year from where incidence data is read (2018; old data: 2008/2012) – with updated 2018 Globocan data, DALY weights from GBD, and DALY estimation based on prevalence instead of age of incidence, only 2018 is valid and 2008/2012 is not sensible
daly.canc.diag	disability weight of diagnosis and primary therapy phase of cervical cancer
daly.canc.control	Number: daly-weight for controlled phase of cervical cancer
daly.canc.metastatic	Number: daly-weight for metastatic phase of cervical cancer
daly.canc.terminal	disability weight of terminal phase of cervical cancer
sens	ignore, doesn't do anything anymore
unwpp_mortality	logical, whether to create lifetables based on UNWPP mx estimates or WHO data
year_born	ignore
year_vac	ignore
runs	ignore
vaccine_efficacy_beforesexdebut	vaccine efficacy before sexual debut
vaccine_efficacy_aftersexdebut	vaccine efficacy after sexual debut
log	name of log file
by_calendaryear	logical, output values by calendar year or by year of birth cohort
use_proportions	logical, output data as rates per capita or in totals
analyseCosts	logical, directly run cost-effectiveness analysis on output or not
psa	integer, number of runs for probabilistic sensitivity analysis (PSA)
psa_vals	data table with values to use in probabilistic sensitivity analysis, usually .data.batch.psa, generated by RegisterBatchData* functions (currently only RegisterBatchdata-Gavi)

Value

Returns combined results

Examples

```
#
```

checkSize *Checks whether the size of a variable is larger than 0*

Description

Used to determine that all required variables are passed to a function Checks whether a vector has length > 0 or a data.table/data.frame has nrow > 0

Usage

```
checkSize(v)
```

Arguments

v	Variable (required)

Value

Logical: TRUE if size is not 0, false if size is 0

Examples

```
x <- c()
checkSize(x)

x <- c(2,5)
checkSize(x)

A <- c()
B <- c(1,2,3)
sapply(c("A","B"),function(x){checkSize(get(x))})
```

data.cecx_1y_prevalence *1-year prevalence of cervical cancer*

Description

A dataset containing the 1-year prevalence (proportion) of cervical cancer in 185 countries, as reported by IARC's Globocan 2018 database.

Usage

```
data.cecx_1y_prevalence
```

Format

A data table containing 185 observations of 103 variables.

Country Country name

0..100 Age 0-100

iso3 ISO3 country code

Details

As per IARC definition – The (1-year) prevalence of a given cancer is the number of individuals within a defined population who have been diagnosed with that cancer (within 1 year) and who are still alive at a given point in time (i.e. the survivors).

Source

https://gco.iarc.fr/today/online-analysis-table

---

data.cecx_3y_prevalence

3-year prevalence of cervical cancer

Description

A dataset containing the 3-year prevalence (proportion) of cervical cancer in 185 countries, as reported by IARC's Globocan 2018 database.

Usage

```
data.cecx_3y_prevalence
```

Format

A data table containing 185 observations of 103 variables.

Country Country name

0..100 Age 0-100

iso3 ISO3 country code

Details

As per IARC definition – The (3-year) prevalence of a given cancer is the number of individuals within a defined population who have been diagnosed with that cancer (within 3 years) and who are still alive at a given point in time (i.e. the survivors).

Source

https://gco.iarc.fr/today/online-analysis-table

data.cecx_5y_prevalence *5-year prevalence of cervical cancer*

Description

A dataset containing the 5-year prevalence (proportion) of cervical cancer in 185 countries, as reported by IARC's Globocan 2018 database.

Usage

```
data.cecx_5y_prevalence
```

Format

A data table with 185 observations of 103 variables.

Country Country name

0..100 Age 0-100

iso3 ISO3 country code

Details

As per IARC definition – The (5-year) prevalence of a given cancer is the number of individuals within a defined population who have been diagnosed with that cancer (within 5 years) and who are still alive at a given point in time (i.e. the survivors).

Source

https://gco.iarc.fr/today/online-analysis-table

data.costcecx *Cost of cervical cancer treatment*

Description

A dataset containing the cost of cervical cancer treatment.

Usage

```
data.costcecx
```

Format

A data table with 194 observations of 4 variables.

country Country name

cancer_cost cost per cancer episode, in $US

cancer_cost_adj cost per cancer episode – adjusted, in international/PPP $

iso3 ISO3 country code

data.countryname *Country names and codes*

Description

A dataset containing the country names and codes (ISO/WHO/WB/UN).

Usage

```
data.countryname
```

Format

A data table with 251 observations of 13 variables.

name1, name2, name3, name4 Country names

iso2, iso3 ISO2 and ISO3 country codes

isonum ISO number

WHOcode WHO country code

who_region WHO region

who_mort WHO mortality stratum - A/B/C/D/E

WBincome WB income classification of countries

UNgroup UN group classification of countries

GDPpc2011id GDP per capita (2011)

data.global *Global data table*

Description

A dataset containing a global range of variables.

Usage

```
data.global
```

Format

A data table with 194 observations of 35 variables.

Country Country name

iso2 ISO2 country code

WHO Region WHO regions

WHO Mortality Stratum WHO mortality stratum

World Bank Income Group (2011) World Bank income group levels

GAVI Eligibility GAVI eligibility status

PAHO Revolving Fund PAHO revolving fund status

Cohort size (2010) [1] Cohort size

Coverage (3 doses at year 10) [2] Vaccination coverage at age 10

Vaccine efficacy vs vaccine type infection [2] Vaccine efficacy

Duration of protection [2] Duration of vaccine protection

Age group [3] Age group

Vaccine price [4] Vaccine price

Vaccine delivery/ operational/ admin costs [5] Vaccine delivery, operational and administration costs

Cancer treatment costs - primary level hospital i$ (per episode, over lifetime) [6] Cancer treatment costs - primary level hospital i$ (per episode, over lifetime)

Cancer treatment costs - primary level hospital US$ (per episode, over lifetime) [6] Cancer treatment costs - primary level hospital US$ (per episode, over lifetime)

Cancer treatment costs - secondary level hospital i$ (per episode, over lifetime) [6] Cancer treatment costs - secondary level hospital i$ (per episode, over lifetime)

Cancer treatment costs - secondary level hospital US$ (per episode, over lifetime) [6] Cancer treatment costs - secondary level hospital US$ (per episode, over lifetime)

Cancer treatment costs - teaching hospital i$ (per episode, over lifetime) [6] Cancer treatment costs - teaching hospital i$ (per episode, over lifetime)

Cancer treatment costs - teaching hospital US$ (per episode, over lifetime) [6] Cancer treatment costs - teaching hospital US$ (per episode, over lifetime)

Discount rate [2] Disount rate

Perspective [2] Perspective

Costs [2]

Time horizon [2] Time horizon

"Percent" CeCx due to 16/18 Percentage of cervical cancer due to HPV strains 16 and 18

Vaccine programme Vaccine programme

Vaccine programme Oct2013 Vaccine programme Oct2013

Econ evaluation Economic evaluation

GDP per capita (2011 US$) [7] GDP per capita (2011 US$)

GDP per capita (2011 i$) [7] GDP per capita (2011 i$)

GNI per capita (2011 i$) [7] GNI per capita (2011 i$)

GNI per capita (2011 US$) [7] GNI per capita (2011 US$)

V33 International$

V34 US$

iso3 ISO3 country code

data.incidence *Incidence of cervical cancer*

Description

A dataset containing the incidence of cervical cancer in 185 countries, as reported by IARC's Globocan 2018 database.

Usage

```
data.incidence
```

Format

A data table with 185 observations of 103 variables.

Country Country name

0..100 Age 0-100

iso3 ISO3 country code

Source

https://gco.iarc.fr/today/online-analysis-table

data.mortall *WHO life table*

Description

A dataset containing the WHO life table.

Usage

```
data.mortall
```

Format

A data table with 196 observations of 107 variables.

Country ¦ Age [12] Country name

0..100 Age 0-100

V103..V106 na

iso3 ISO3 country code

Source

https://www.who.int/gho/mortality_burden_disease/life_tables/life_tables/en/

data.mortall.unwpp.mx *UNWPP life table*

Description

A dataset containing the UNWPP life table (World Population Prospects 2017).

Usage

```
data.mortall.unwpp.mx
```

Format

A data table with 115710 observations of 8 variables.

country_code_numeric Country code numeric
country_code ISO3 country code
country Country name
age_from age from (start-age)
age_to age to (end-age)
year Year
gender Gender
value mx mortality rate in year = proportion of individuals of age x dying by age x+1

Source

https://population.un.org/wpp/

data.mortcecx *Mortality from cervical cancer*

Description

A dataset containing the mortality from cervical cancer in 185 countries, as reported by IARC's Globocan 2018 database.

Usage

```
data.mortcecx
```

Format

A data table with 185 observations of 103 variables.

Country Country name
0..100 Age 0-100
iso3 ISO3 country code

Source

https://gco.iarc.fr/today/online-analysis-table

data.pop *UNWPP population estimates*

Description

A dataset containing the UNWPP population estimates – World Population Prospects 2017.

Usage

```
data.pop
```

Format

A data table with 3306163 observations of 8 variables.

country_code_numeric Country code numeric

country_code ISO3 country code

country Country name

age_from age from (start-age)

age_to age to (end-age)

year Year

gender Gender

value Population size

Source

https://population.un.org/wpp/Download/Standard/Population/

VIMC

data.popproj *Population projections of 5-year old girls*

Description

A dataset containing population projections of 5-year old girls.

Usage

```
data.popproj
```

Format

A data table with 98 observations of 91 variables.

iso3 ISO3 country code

2011..2100 Year – 2011..2100

data.quality *Data quality of incidence and mortality*

Description

A dataset indicating data quality of cervical cancer incidence and mortality.

Usage

```
data.quality
```

Format

A data table with 186 observations of 4 variables.

Country Country name

Incidence Quality of cervical cancer incidence data

Mortality Quality of cervical cancer mortality data

iso3 ISO3 country code

data.sexual_debut *Sexual debut data*

Description

A dataset containing sexual debut data and (2) parameters for the sexual debut curve (logistic model).

Usage

```
data.sexual_debut
```

Format

A data table with 94 observations of 14 variables.

V1 Row number

iso2 ISO2 country code

country Country name

iso3 ISO3 country code

who WHO region

X15 Proportion of people who have sexually debuted at age 15

X18 Proportion of people who have sexually debuted at age 18

X20 Proportion of people who have sexually debuted at age 20

X22 Proportion of people who have sexually debuted at age 22

X25 Proportion of people who have sexually debuted at age 25

Never Proportion of people who had not sexually debuted

cluster.id Clustering countries with similar characteristics

a Parameter for sexual debut curve (logistic model)

b Parameter for sexual debut curve (logistic model)

`data.valid` *Model validation*

Description

A dataset containing data for validation.

Usage

```
data.valid
```

Format

A data table with 26 observations of 49 variables.

Country Country name

iso2 ISO2 country code

WHO Region WHO region

World Bank Income Group (2011) World Bank income group classification (2011)

Author Author

Year Year of publication

Title Title of publication

Currency Currency

Currency year Currency year

Conversion to I$2011 International dollar (I$2011)

ICER vs no prevention Incremental cost-effectiveness ratio of vaccination versus no prevention (CHECK)

ICER vs screen Incremental cost-effectiveness ratio of vaccination versus no prevention (CHECK)

Denominator Denominator for health impact

Vaccine total costs Vaccine total costs

Vaccine coverage Vaccine coverage

Vaccine efficacy vs vaccine type infection Vaccine efficacy versus vaccine type infection

Duration of protection Duration of protective immunity from vaccination

Cohort size Cohort size

Age at vaccination Age at vaccination, years

Cancer treatment cost per episode Cancer treatment cost per episode, dollars

Discount rate: costs Discount rate for costs

Discount rate: benefits Discount rate for benefits

Perspective Perspective of economic evaluation

Comparator is no screening Comparator refers to no scereening, logical (Y/N)

Time horizon Time horizon of analysis

GDP per capita GDP per capita

Cervical cancer due to 16/18 Proportion of cervical cancer due to HPV types 16 and 18

CeCx cost low original Cervical cancer cost / low / original

CeCx cost high original Cervical cancer cost / high / original

CeCx cost low Cervical cancer cost / low

CeCx cost high Cervical cancer cost / high

CeCx data available Cervical cancer data available, logical (Y/N)

0-4 0-4 years

5-9 5-9 years

9-14 9-14 years

15-19 15-19 years

20-24 20-24 years

25-29 25-29 years

30-34 30-34 years

35-39 35-39 years

40-44 40-44 years

45-49 45-49 years

50-54 50-54 years

55-59 55-59 years

60-64 60-64 years

65-69 65-69 years

70-74 70-74 years

75-79 75-79 years

80+ 80+ years

`dtAggregate` *Collapse data-tables*

Description

Collapse data-tables

Usage

```
dtAggregate(DT, aggr_on, measure.vars = c(), id.vars = c(),
  func = "sum", na.rm = TRUE)
```

Arguments

DT	Data-table (required)
aggr_on	Character string (required): column-name that will be used to collapse on (i.e. combine all age-strata)
measure.vars	Character string (optional): column-names that will be collapsed (function will be applied to all these columns)
id.vars	Character string (optional): column-names that will remain stratified N.b. if measure.vars is not provided, all columns that are not in id.vars and aggr_on will be assumed to be assumed
func	Character string (optional): function that will be applied to data (if optional, values will be summed)
na.rm	Logical (optional): if TRUE, removes NA from measure.vars columns before applying function (or passes na.rm=TRUE to function)

Value

Returns collapsed data.table

Examples

```
dtAggregate(data.popproj,"iso3",id.vars="")
```

---

dtColMatch *Match two data-tables on multiple columns*

Description

Returns vector with column-of-interest where columns match

Usage

```
dtColMatch(input, input_match_on, reference, reference_match_on,
  reference_return)
```

Arguments

input	Data.table (required): input-table to match
input_match_on	Character vector (required): column-names in input-table to match
reference	Data.table (required): reference-table to match
reference_match_on	Character vector (required): column-names in reference-table to match
reference_return	Character string (required): column-name in reference-table that is returned (where values match)

Details

If at least one value in any of the input_match_on columns matches with a value in any of the reference_match_on columns, the two rows will match

Value

Character vector with values from reference_return column in reference_match_on data.table where values match

Examples

```
dtColMatch(data.global,c("Country"),data.countryname,c("name1","name2","name3","name4"),"iso3")
```

---

getISO3 *Retrieve ISO3-code of country*

Description

Retrieve ISO3-code of country

Usage

```
getISO3(countryname, name = FALSE)
```

Arguments

countryname	Character string (required): Full name of the country
name	Logical (optional): If TRUE, returns full name and alternative names of returned country (may be useful to double-check that it is the correct country)

Value

Character string with ISO3 code. Will also return full name if name=TRUE.

Examples

```
getISO3("Afghanistan")
getISO3("Congo",name=TRUE)
```

---

lifeTable *Construct lifetable based on qx-column*

Description

qx = age-specific probability of dying

Usage

```
lifeTable(qx = NULL, mx = NULL, agecohort = 0)
```

Arguments

qx	Numeric vector (required): Age-specific probabilities of dying
agecohort	Number (optional): Age at which cohort is started

Value

Data.table with lifetable

Examples

```
qx <- unlist(data.mortall[iso3=="AFG", as.character(0:100), with=F], use.names=F)
lifeTable(qx, 9)
```

`monetary_to_number` *Convert monetary character-strings to numeric values*

Description

Convert monetary character-strings to numeric values

Usage

```
monetary_to_number(x)
```

Arguments

`x`	Character string to convert

Value

Returns number with value, stripped from any currency symbols and thousand-seperators (i.e. "B#2,010.50" becomes 2010.5)

Examples

```
monetary_to_number("$2,200.20")

#Note that values using German or Dutch notation (i.e. using a comma to separate decimals and a dot to seperate t
monetary_to_number("$2.200,20")
```

`OutputGavi` *Formatting output for VIMC Montagu*

Description

`OutputGavi` takes result of BatchRun and outputs it in format to be uploaded to VIMC Montagu.

Usage

```
OutputGavi(DT, age_stratified = TRUE, calendar_year = FALSE,
  gavi_template = -1)
```

Arguments

DT	data table with results
age_stratified	logical, whether output should be stratified by age
calendar_year	logical, whether output should be given by calendar year of event OR by year of birth of cohort
gavi_template	data table with template file downloaded from montagu

Value

#

Examples

#

---

prime *prime: Papillomavirus Rapid Interface for Modelling and Economics (PRIME).*

Description

PRIME stands for "Papillomavirus Rapid Interface for Modelling and Economics". The R package is based of the spreadsheet-based tool (see http://primetool.org).

PRIME provides estimates of

The magnitude of the burden of cervical cancer.

The impact of introducing HPV vaccination for girls prior to sexual debut.

Healthcare costs incurred as a result of cervical cancer treatment.

Costs associated with vaccination.

Long-term savings which may result from a vaccination program.

---

propSexDebut *Proportion of girls sexually debuted*

Description

propSexDebut returns proportion of girls sexually debuted in country country_iso3 at age age.

Usage

```
propSexDebut(age, country_iso3)
```

Arguments

age	age of girls
country_iso3	ISO3 country code

Value

Returns proportion of girls in a given country that has sexually debuted at a given age.

Examples

```
propSexDebut (20, "IND")
propSexDebut (30, "ETH")
```

`RegisterBatchData` *Creates .data.batch for running multiple birth cohorts*

Description

Creates .data.batch which is used when running/looping over multiple birth cohorts (runCohort()) at once.

Usage

```
RegisterBatchData(coverage_data, reporting_years = -1, force = FALSE)
```

Arguments

`coverage_data`	data table with columns country_code, year (of vaccination), age_first, age_last, coverage
`reporting_years`	numeric_vector, years that should be reported (parameter: not required)
`force`	logical, whether .data.batch should be overwritten if it already exists (parameter: not required)

Details

.data.batch is based on the data.table (DT) coverage_data, which is a DT with columns country_code (ISO3), year (of vaccination), age_first (age at vaccination), age_last (age at vaccination), coverage (in proportion, for all the agegroups specified).

If you only want to run 1 age in this country/coverage combination, age_first==age_last

Value

None

Examples

```
#
```

`RegisterBatchDataGavi` *Creates .data.batch for running multiple birth cohorts (Gavi runs)*

Description

Creates .data.batch which is used when running/looping over multiple birth cohorts (runCohort()) at once. Similar to RegisterBatchData, but for when we make runs for Gavi.

Usage

```
RegisterBatchDataGavi(gavi_coverage, gavi_template, use_campaigns,
  use_routine, restrict_to_coverage_data = FALSE, force = FALSE,
  psa = 0)
```

Arguments

`gavi_coverage`	data table with coverage estimates as downloaded from VIMC montagu
`gavi_template`	data table with reporting template as downloaded from VIMC montagu
`use_campaigns`	logical, whether campaigns as stated in coverage files should be modelled
`use_routine`	logical, whether routine vaccination as stated in coverage file should be modelled
`restrict_to_coverage_data`	logical, whether the first birth-cohort should be the first cohort that is mentioned in the coverage data. If TRUE, restrict to coverage data. If FALSE, restrict to cohorts provided in gavi_template.
`force`	logical, whether .data.batch should be overwritten if it already exists
`psa`	integer, indicating how many runs for probabilistic sensitivity analysis (PSA). 0 to run no PSA.

Details

.data.batch is based on the data.table (DT) coverage_data, which is a DT with columns country_code (ISO3), year (of vaccination), age_first (age at vaccination), age_last (age at vaccination), coverage (in proportion, for all the age groups specified).

Value

None

Examples

```
#
```

RunCohort *Run PRIME for a single birth-cohort*

Description

Runs PRIME for one birth-cohort. Usually called by another function such as RunCountry().

Usage

```
RunCohort(lifetab, cohort, incidence, mortality_cecx, prevalence, agevac,
  coverage, campaigns, vaccine_efficacy_nosexdebut,
  vaccine_efficacy_sexdebut, daly.canc.diag, daly.canc.seq,
  daly.canc.control, daly.canc.metastatic, daly.canc.terminal, cost_cancer,
  disc.cost = 0.03, disc.ben = 0.03, discounting = TRUE,
  country_iso3 = NULL, run_country = FALSE)
```

Arguments

lifetab	Data.table: The life-table for this cohort. Can be created using the lifeTable() function.
cohort	Number: The cohort-size of this birth-cohort at the time where the lifetable starts.
incidence	Numeric vector: Age-specific CeCx(16/18) incidence-rates.
mortality_cecx	Numeric vector: Age-specific CeCx(16/18) mortality-rates.
prevalence	Numeric vector: Age-specific CeCx(16/18) prevalence rates (5-year prevalence) – referring to people who are alive within 5 years of diagnosis.
agevac	Number: Age at which the cohort is vaccinated.
coverage	Number: Proportion of the cohort that will receive a vaccination.
campaigns	List or number: MAC cohort-vaccinations (needs to be changed).
vaccine_efficacy_nosexdebut	Number: proportion indicating vaccine-efficacy before sexual debut.
vaccine_efficacy_sexdebut	Number: proportion indicating vaccine-efficacy after sexual debut.
daly.canc.diag	Number: daly-weight for cancer diagnosis.
daly.canc.seq	Number: daly-weight for cancer treatment.
daly.canc.control	Number: daly-weight for controlled phase of cervical cancer
daly.canc.metastatic	Number: daly-weight for metastatic phase of cervical cancer
daly.canc.terminal	Number: daly-weight for death from cancer.
cost_cancer	Number: total per capita cost of cancer.
disc.cost	Number (optional): discounting for cancer cost.
disc.ben	Number (optional): discounting for...
discounting	Logical: should discounting be applied?

Value

Returns a data.table with size of the birth-cohort and age-specific incidence-rates, mortality-rates, years-of-life-lost, years-of-healthy-life-lost, and cancer-costs before and after vaccination. Also displays whether discounting has been used ("type" column).

Examples

```
lifetab <- lifeTable(unlist(data.mortall[iso3=="AFG", as.character(0:100), with=F], use.names=F), 9)
cohort <- unlist(data.popproj[iso3=="AFG", "2020"], use.names=F)
incidence <- unlist(data.incidence[iso3=="AFG",as.character(0:100),with=F],use.names=F)
mortality_cecx <- unlist(data.mortall[iso3=="AFG",as.character(0:100),with=F],use.names=F)
prevalence <- unlist(data.cecx_5y_prevalence[iso3=="AFG",as.character(0:100),with=F],use.names=F)
agevac <- 9
coverage <- 0.8
campaigns <- -1
vaccine_efficacy_nosexdebut <- 0.95
vaccine_efficacy_sexdebut <- 0
daly.canc.diag <- 0.002
daly.canc.seq <- 0.002
daly.canc.control <- 0.05
daly.canc.metastatic <- 0.05
daly.canc.terminal <- 0.1
cost_cancer <- 100

RunCohort(lifetab, cohort, incidence, mortality_cecx, prevalence, agevac, coverage, campaigns,
 vaccine_efficacy_nosexdebut, vaccine_efficacy_sexdebut, daly.canc.diag, daly.canc.seq, daly.canc.control,
 daly.canc.metastatic, daly.canc.terminal, cost_cancer, disc.cost=0.03, disc.ben=0.03, discounting=FALSE,
 country_iso3="AFG", run_country=FALSE)
```

RunCountry *Run PRIME for a specific country*

Description

Runs RunCohort() using country-specific estimates. If year_born and year_vac are not provided, assumes vaccination occurs in the current year.

Usage

```
RunCountry(country_iso3, vaceff_beforesexdebut = 1,
  vaceff_aftersexdebut = 0, disc.cost = 0.03, disc.ben = 0.03,
  cov = 1, agevac = 10, agecohort = 10, cohort = -1,
  canc.cost = "unadj", canc.inc = "2018", daly.canc.diag = 0.288,
  daly.canc.control = 0.049, daly.canc.metastatic = 0.451,
  daly.canc.terminal = 0.54, sens = -1, unwpp_mortality = FALSE,
  year_born = -1, year_vac = -1, campaigns = -1,
  analyseCosts = FALSE, discounting = TRUE, run_batch = FALSE,
  psadat = -1)
```

Arguments

country_iso3	Character string (required): ISO3 code of the country
disc.cost	Number (optional): Discounting for costs (only if discounting=TRUE)
disc.ben	Number (optional): Discounting for ... (only if discounting=TRUE)
cov	Number (optional): Proportion with routine coverage
agevac	Integer (optional): Target age for HPV vaccination
agecohort	Integer (optional): Reference age for cohort-size (only used when 'cohort' is not provided)
cohort	Integer (optional): Cohort-size. -1 if unknown
canc.cost	Character (optional): Is cost of cancer adjusted ("adj") or not ("unadj")
canc.inc	Integer (optional): Reference year for cancer incidence rates (2018 or 2012 or 2008)
daly.canc.diag	Number (optional): Daly weight for cancer diagnosis
daly.canc.control	Number: daly-weight for controlled phase of cervical cancer
daly.canc.metastatic	Number: daly-weight for metastatic phase of cervical cancer
daly.canc.terminal	Number (optional): Daly weight for cancer death
sens	Numeric-vector (optional): Specific values to be used in a PSA. -1 if PSA's are not used
unwpp_mortality	Logical (optional): If TRUE, uses year-specific UNWPP mortality estimates to construct life-tables. If FALSE, use WHO based mortality estimates
year_born	Integer (optional): Year in which cohort is born
year_vac	Integer (optional): Year in which cohort is vaccinated
campaigns	List (optional): Multi-Age-Cohort campaigns (needs to be changed)
analyseCosts	Logical (optional): If FALSE, returns result from RunCohort() function. If TRUE, runs analyseCosts() with country-specific results
discounting	Logical (optional): If TRUE, runs analysis undiscounted and discounted. If FALSE, only uses undiscounted
vaceff	Number (optional): Proportion indicating vaccine-efficacy

Value

data.table with country-specific results of HPV vaccination. Returns cost-analysis if analyseCosts=TRUE

Examples

```
RunCountry("AFG")
RunCountry("AFG", year_vac=2020, agevac=10, cov=0.75, vaceff=0.88)
RunCountry("AFG", year_vac=2020, agevac=10, cov=0.75, vaceff=0.88, analyseCosts=TRUE)
```

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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