

# Package ‘golden’

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**Type** Package

**Title** Framework for Patient-Level Microsimulation of Risk Factor Trajectories & Hazard-Based Events

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**Description** Fast, flexible, patient-level microsimulation. Time-stepped simulation with a 'C++' back-end from user-supplied initial population, trajectories, hazards, and corresponding event transitions. User-defined aggregate time series histories are returned together with the final population. Designed for simulation of chronic diseases with continuous and evolving risk factors, but could easily be applied more generally.

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**Imports** Rcpp (>= 1.1.0), data.table

**LinkingTo** Rcpp

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**Encoding** UTF-8

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---

bmi_fits	<i>Example BMI distribution</i>
----------	---------------------------------

---

### Description

details TODO

### Usage

bmi\_fits

### Format

TODO

**foo** bar

**Source**

TODO

---

check_column	<i>Validate an history column object If validation fails, an exception will be raised.</i>
--------------	--

---

**Description**

Validate an history column object If validation fails, an exception will be raised.

**Usage**

```
check_column(column, initPop = NULL)
```

**Arguments**

column	An S3 object of class "golden_history_column"
initPop	(Optional) data.table to check columns required by functions exist

**Value**

No return value, called for side effects.

**Examples**

```
library(data.table)
dt <- data.table(a = rep(0, 100))
# Create an S3 golden_history_column
col <- new_column("sum_a", sum, c("a"))
# check_column() will not throw an exception
# as col is a valid S3 golden_history_column
# and dt contains column "a"
check_column(col, dt)
```

---

check_hazard	<i>Validate an hazard object If validation fails, an exception will be raised.</i>
--------------	--

---

**Description**

Validate an hazard object If validation fails, an exception will be raised.

**Usage**

```
check_hazard(hazard, initPop = NULL)
```

**Arguments**

hazard            An S3 object of class "golden\_hazard"  
 initPop          (Optional) data.table to check columns required by functions exist

**Value**

No return value, called for side effects.

**Examples**

```
library(data.table)
N <- 100
dt <- data.table(a = runif(N, 0, 1), b = rep(0, N))
# Define a hazard function, which returns a vector of equal length uncertainties
test_hazard <- function(a) {
  ret <- (a < 0.5)
}
# Define a transition function, which sets all "b" columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Create an S3 golden_hazard
haz <- new_hazard(
  test_hazard,
  c("a"),
  new_transition(test_transition, c(), "b")
)
# check_hazard() will not throw an exception
# as haz is a valid S3 golden_hazard
# and dt contains column "a"
check_hazard(haz, dt)
```

---

check_history	<i>Validate an history object If validation fails, an exception will be raised.</i>
---------------	---

---

**Description**

Validate an history object If validation fails, an exception will be raised.

**Usage**

```
check_history(history, initPop = NULL)
```

**Arguments**

history            An S3 object of class "golden\_history"  
 initPop          (Optional) data.table to check columns required by functions exist

**Value**

No return value, called for side effects.

**Examples**

```
library(data.table)
dt <- data.table(a = rep(0, 100))
# Create an S3 golden_history, containing 1 golden_history_column
hist <- new_history(new_column("sum_a", sum, c("a")))
# check_history() will not throw an exception
# as hist is a valid S3 golden_history
# and dt contains column "a" used by the column
check_history(hist, dt)
```

---

check_parameters	<i>Validate the configuration passed to run_simulation() If validation fails, an exception will be raised.</i>
------------------	--

---

**Description**

Validate the configuration passed to run\_simulation() If validation fails, an exception will be raised.

**Usage**

```
check_parameters(parameters, initPop = NULL)
```

**Arguments**

parameters	An golden_parameters S3 object to be validated
initPop	data.frame which contains the columns required by parameters

**Value**

No return value, called for side effects.

**Examples**

```
library(data.table)
N <- 100
dt <- data.table(a = runif(N, 0, 1), b = rep(0, N))
# Define a hazard function, which returns a vector of equal length uncertainties
test_hazard <- function(a) {
  ret <- (a < 0.5)
}
# Define a transition function, which sets all "b" columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Create an S3 golden_hazard
```

```

haz <- new_hazard(
  test_hazard,
  c("a"),
  new_transition(test_transition, c(), "b")
)
# Define a trajectory function, which adds 2 to all members of the input vector
test_trajectory <- function(a) {
  return (a + 2)
}
# Create an S3 golden_trajectory
trj <- new_trajectory(test_trajectory, c("b"), "b")
# Create an S3 golden_history, containing 1 golden_history_column
hist <- new_history(new_column("sum_a", sum, c("a")))
# Create an S3 golden_parameters
params <- new_parameters(
  hazards = haz,
  trajectories = trj,
  steps = 10,
  debug = FALSE,
  history = hist
)
# check_parameters() will not throw an exception
# as params is a valid S3 golden_parameters
# and dt contains columns "a" and "b"
check_parameters(params, dt)

```

---

check_trajectory	<i>Validate an trajectory object If validation fails, an exception will be raised.</i>
------------------	--

---

## Description

Validate an trajectory object If validation fails, an exception will be raised.

## Usage

```
check_trajectory(trajectory, initPop = NULL)
```

## Arguments

trajectory	An S3 object of class "golden_trajectory"
initPop	(Optional) data.table to check columns required by functions exist

## Value

No return value, called for side effects.

**Examples**

```

library(data.table)
dt <- data.table(b = rep(0, 100))
# Define a trajectory function, which adds 2 to all members of the input vector
test_trajectory <- function(a) {
  return (a + 2)
}
# Create an S3 golden_trajectory
trj <- new_trajectory(test_trajectory, c("b"), "b")
# check_trajectory() will not throw an exception
# as trj is a valid S3 golden_trajectory
# and dt contains column "b"
check_trajectory(trj, dt)

```

---

check_transition	<i>Validate an transition object If validation fails, an exception will be raised.</i>
------------------	--

---

**Description**

Validate an transition object If validation fails, an exception will be raised.

**Usage**

```
check_transition(transition, initPop = NULL)
```

**Arguments**

transition	An S3 object of class "golden_transition"
initPop	(Optional) data.table to check columns required by functions exist

**Value**

No return value, called for side effects.

**Examples**

```

library(data.table)
dt <- data.table(b = rep(0, 100))
# Define a transition function, which sets all columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Define an S3 golden_transition
trn <- new_transition(test_transition, c(), "b")
# check_transition() will not throw an exception
# as trn is a valid S3 golden_transition
# and dt contains column "b"
check_transition(trn, dt)

```

---

create_cohort	<i>Create a new cohort</i>
---------------	----------------------------

---

**Description**

Temporary testing method, probably replaced in future with R's simdata package or similar

**Usage**

```
create_cohort(demog, N)
```

**Arguments**

demog	Demographic information containing columns AgeGrp/PopMale/PopFemale/PopTotal
N	Size of the population to generate

**Value**

A sample population data.table, with columns male/age/bmi/death

**Examples**

```
library(data.table)
demog <- data.table(
  AgeGrp = c(0, 1, 2, 3),
  PopMale = c(1000, 1100, 1050, 980),
  PopFemale = c(950, 1020, 1005, 970)
)
demog[, PopTotal := PopMale + PopFemale]
cohort <- create_cohort(demog, 100)
```

---

globoRisk_coefs	<i>Example globoRisk coefficients</i>
-----------------	---------------------------------------

---

**Description**

details TODO

**Usage**

```
globoRisk_coefs
```

**Format**

TODO

**foo** bar

**Source**

TODO

---

globo-risk\_cvdr

*Example globo-risk baseline hazard*

---

**Description**

details TODO

**Usage**

globo-risk\_cvdr

**Format**

TODO

**foo** bar

**Source**

TODO

---

globo-risk\_rf

*Example globo-risk reference values*

---

**Description**

details TODO

**Usage**

globo-risk\_rf

**Format**

TODO

**foo** bar

**Source**

TODO

golden

*Golden: Framework for Patient-Level Microsimulation of Risk Factor Trajectories & Hazard-Based Events*

### Description

Fast, flexible, patient-level microsimulation. Time-stepped simulation with a 'C++' back-end from user-supplied initial population, trajectories, hazards, and corresponding event transitions. User-defined aggregate time series histories are returned together with the final population. Designed for simulation of chronic diseases with continuous and evolving risk factors, but could easily be applied more generally.

### Author(s)

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- University of Sheffield [copyright holder]
- Horizon Europe [funder]

### Examples

```
## Not run:
# A full example can be found in the vignettes

## End(Not run)
```

lifetable\_data

*Example life table data*

### Description

details TODO

### Usage

```
lifetable_data
```

### Format

TODO

**foo** bar

**Source**

TODO

---

new_column	<i>Create a new golden_history_column</i>
------------	---

---

**Description**

Create a new golden\_history\_column

**Usage**

```
new_column(name, fn, args, filter_fn = NULL, filter_args = NULL)
```

**Arguments**

name	Name of the column in the output data-table
fn	Reduction function, which converts the input columns to a single value
args	Names of columns and special variables to be passed to fn
filter_fn	(Optional) Filter function, which returns a bool vector denoting which rows should be reduced
filter_args	(Optional) Names of columns and special variables to be passed to filter_fn. Required if filter_fn is

**Value**

An object of class "golden\_history\_column"

**Examples**

```
# Create an S3 golden_history_column named "sum_a", using sum(). with column "a"  
col <- new_column("sum_a", sum, c("a"))
```

---

new_hazard	<i>Create a new hazard object</i>
------------	-----------------------------------

---

**Description**

Create a new hazard object

**Usage**

```
new_hazard(  
  fn,  
  args,  
  transitions,  
  freq = 1,  
  first = 1,  
  last = 2147483647,  
  name = NULL  
)
```

**Arguments**

fn	Function which calculates the hazard likelihood
args	Character vector of parameter names expected by fn
transitions	Transition object(s) to be applied where the hazard is successful
freq	(Optional) The frequency of hazard execution, hazards always execute on first step
first	(Optional) First step the hazard should be enabled (initial step is index 1)
last	(Optional) Last step the hazard should be enabled (initial step is index 1)
name	(Optional) Name used in error messages and similar. Defaults to an automatic name

**Value**

An object of class "golden\_hazard"

**Examples**

```
# Define a hazard function, which returns a vector of equal length uncertainties  
test_hazard <- function(a) {  
  ret <- (a < 0.5)  
}  
# Define a transition function, which sets all "b" columns affected by the hazard to 100  
test_transition <- function() {  
  return (100)  
}  
# Create an S3 golden_hazard
```

```
haz <- new_hazard(  
  test_hazard,  
  c("a"),  
  new_transition(test_transition, c(), "b")  
)
```

---

new\_history                    *Create a new golden\_history*

---

### Description

Create a new golden\_history

### Usage

```
new_history(columns, frequency = 1)
```

### Arguments

columns                    golden\_history\_column S3 object(s)  
frequency                 The number of simulation steps per history collection.

### Value

An object of class "golden\_history"

### Examples

```
# Create an S3 golden_history, containing 1 golden_history_column  
hist <- new_history(new_column("sum_a", sum, c("a")))
```

---

new\_parameters                *Create a new golden\_parameters*

---

### Description

Create a new golden\_parameters

### Usage

```
new_parameters(  
  hazards = list(),  
  trajectories = list(),  
  steps,  
  random_seed = 0,  
  debug = TRUE,  
  print_timing = TRUE,  
  history = NULL  
)
```

**Arguments**

hazards	golden_hazard S3 object(s)
trajectories	golden_trajectory S3 object(s)
steps	Number of steps to run
random_seed	Seed to be used for random generation. If set 0, current R random state will be used.
debug	(TRUE/FALSE) flag indicating whether validation checks are enabled. These catch NaN, but reduce performance
print_timing	(TRUE/FALSE) flag indicating whether a per-function timing report should be printed after the simulation, this will always be suppressed for fast ( $\leq 1$ second) simulations.
history	golden_history S3 object representing the columns of data to be aggregated during simulation

**Value**

An object of class "golden\_parameters"

**Examples**

```
# Define a hazard function, which returns a vector of equal length uncertainties
test_hazard <- function(a) {
  ret <- (a < 0.5)
}
# Define a transition function, which sets all "b" columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Create an S3 golden_hazard
haz <- new_hazard(
  test_hazard,
  c("a"),
  new_transition(test_transition, c(), "b")
)
# Define a trajectory function, which adds 2 to all members of the input vector
test_trajectory <- function(a) {
  return (a + 2)
}
# Create an S3 golden_trajectory
trj <- new_trajectory(test_trajectory, c("b"), "b")
# Create an S3 golden_history, containing 1 golden_history_column
hist <- new_history(new_column("sum_a", sum, c("a")))
# Create an S3 golden_parameters
params <- new_parameters(
  hazards = haz,
  trajectories = trj,
  steps = 10,
  debug = FALSE,
  history = hist
)
```

---

new_trajectory	<i>Create a new trajectory object</i>
----------------	---------------------------------------

---

## Description

Create a new trajectory object

## Usage

```
new_trajectory(fn, args, property, name = NULL)
```

## Arguments

fn	Function defining the trajectory function
args	Character vector of parameter names expected by fn
property	Name(s) of the column(s) where the result(s) of the trajectory function is to be stored
name	(Optional) Name used in error messages and similar. Defaults to an automatic name

## Value

An object of class "golden\_trajectory"

## Note

If a list is passed to property, fn must return a list of equal length

## Examples

```
# Define a trajectory function, which adds 2 to all members of the input vector
test_trajectory <- function(a) {
  return (a + 2)
}
# Create an S3 golden_trajectory
trj <- new_trajectory(test_trajectory, c("b"), "b")
```

---

`new_transition`      *Create a new transition object*

---

### Description

Create a new transition object

### Usage

```
new_transition(fn, args, state, name = NULL)
```

### Arguments

<code>fn</code>	Function defining the transition functions
<code>args</code>	Character vector of parameter names expected by <code>fn</code>
<code>state</code>	Name(s) of the column(s) where the result of the transition function is to be stored
<code>name</code>	(Optional) Name used in error messages and similar. Defaults to an automatic name

### Value

An object of class "golden\_transition"

### Examples

```
# Define a transition function, which sets all columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Define an S3 golden_transition
trn <- new_transition(test_transition, c(), "b")
```

---

`pop_snapshot`      *Example population structure*

---

### Description

details TODO

### Usage

```
pop_snapshot
```

**Format**

```
TODO

foo bar
```

**Source**

```
TODO
```

---

```
print.golden_hazard Print the contents of a golden_hazard type S3 object
```

---

**Description**

Print the contents of a golden\_hazard type S3 object

**Usage**

```
## S3 method for class 'golden_hazard'
print(x, ..., indent = 0L)
```

**Arguments**

x	The object to be printed
...	Not used. Included for S3 method compatibility.
indent	(Optional) The level the printing is indented, useful if nested within another S3 object

**Value**

No return value, called for side effects.

**Examples**

```
# Define a hazard function, which returns a vector of equal length uncertainties
test_hazard <- function(a) {
  ret <- (a < 0.5)
}
# Define a transition function, which sets all "b" columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Create an S3 golden_hazard
haz <- new_hazard(
  test_hazard,
  c("a"),
  new_transition(test_transition, c(), "b")
)
print(haz)
```

---

```
print.golden_history
```

*Print the contents of a golden\_history type S3 object*

---

**Description**

Print the contents of a golden\_history type S3 object

**Usage**

```
## S3 method for class 'golden_history'  
print(x, ..., indent = 0L)
```

**Arguments**

x	The object to be printed
...	Not used. Included for S3 method compatibility.
indent	(Optional) The level the printing is indented, useful if nested within another S3 object

**Value**

No return value, called for side effects.

**Examples**

```
# Create an S3 golden_history  
hist <- new_history(new_column("sum_a", sum, c("a")))  
print(hist)
```

---

```
print.golden_history_column
```

*Print the contents of a golden\_history\_column type S3 object*

---

**Description**

Print the contents of a golden\_history\_column type S3 object

**Usage**

```
## S3 method for class 'golden_history_column'  
print(x, ..., indent = 0L)
```

**Arguments**

x	The object to be printed
...	Not used. Included for S3 method compatibility.
indent	(Optional) The level the printing is indented, useful if nested within another S3 object

**Value**

No return value, called for side effects.

**Examples**

```
# Create an S3 golden_history_column
col <- new_column("sum_a", sum, c("a"))
print(col)
```

---

```
print.golden_parameters
```

*Print the contents of a golden\_parameters type S3 object*

---

**Description**

Print the contents of a golden\_parameters type S3 object

**Usage**

```
## S3 method for class 'golden_parameters'
print(x, ..., indent = 0)
```

**Arguments**

x	The object to be printed
...	Not used. Included for S3 method compatibility.
indent	(Optional) The level the printing is indented, useful if nested within another S3 object

**Value**

No return value, called for side effects.

**Examples**

```

# Define a hazard function, which returns a vector of equal length uncertainties
test_hazard <- function(a) {
  ret <- (a < 0.5)
}
# Define a transition function, which sets all "b" columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Create an S3 golden_hazard
haz <- new_hazard(
  test_hazard,
  c("a"),
  new_transition(test_transition, c(), "b")
)
# Define a trajectory function, which adds 2 to all members of the input vector
test_trajectory <- function(a) {
  return (a + 2)
}
# Create an S3 golden_trajectory
trj <- new_trajectory(test_trajectory, c("b"), "b")
# Create an S3 golden_history, containing 1 golden_history_column
hist <- new_history(new_column("sum_a", sum, c("a")))
# Create an S3 golden_parameters
params <- new_parameters(
  hazards = haz,
  trajectories = trj,
  steps = 10,
  debug = FALSE,
  history = hist
)
print(params)

```

---

```
print.golden_timing
```

*Print the contents of a golden\_timing type S3 object*

---

**Description**

Print the contents of a golden\_timing type S3 object

**Usage**

```
## S3 method for class 'golden_timing'
print(x, ...)
```

**Arguments**

x	The object to be printed
...	Not used. Included for S3 method compatibility.

**Value**

No return value, called for side effects.

**Examples**

```

library(data.table)
N <- 100
dt <- data.table(a = runif(N, 0, 1), b = rep(0, N))
# Define a hazard function, which returns a vector of equal length uncertainties
test_hazard <- function(a) {
  ret <- (a < 0.5)
}
# Define a transition function, which sets all "b" columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Create an S3 golden_hazard
haz <- new_hazard(
  test_hazard,
  c("a"),
  new_transition(test_transition, c(), "b")
)
# Define a trajectory function, which adds 2 to all members of the input vector
test_trajectory <- function(a) {
  return (a + 2)
}
# Define an S3 golden_trajectory
trj <- new_trajectory(test_trajectory, c("b"), "b")
# Create an S3 golden_history, containing 1 golden_history_column
hist <- new_history(new_column("sum_a", sum, c("a")))
# Define an S3 golden_parameters
params <- new_parameters(
  hazards = haz,
  trajectories = trj,
  steps = 10,
  debug = FALSE,
  history = hist
)
# Run the simulation to collect results
results <- run_simulation(dt, params)
print(results$timing)

```

---

```
print.golden_trajectory
```

*Print the contents of a golden\_trajectory type S3 object*

---

**Description**

Print the contents of a golden\_trajectory type S3 object

**Usage**

```
## S3 method for class 'golden_trajectory'  
print(x, ..., indent = 0L)
```

**Arguments**

x	The object to be printed
...	Not used. Included for S3 method compatibility.
indent	(Optional) The level the printing is indented, useful if nested within another S3 object

**Value**

No return value, called for side effects.

**Examples**

```
# Define a trajectory function, which adds 2 to all members of the input vector  
test_trajectory <- function(a) {  
  return (a + 2)  
}  
# Create an S3 golden_trajectory  
trj <- new_trajectory(test_trajectory, c("b"), "b")  
print(trj)
```

---

```
print.golden_transition
```

*Print the contents of a golden\_transition type S3 object*

---

**Description**

Print the contents of a golden\_transition type S3 object

**Usage**

```
## S3 method for class 'golden_transition'  
print(x, ..., indent = 0L)
```

**Arguments**

x	The object to be printed
...	Not used. Included for S3 method compatibility.
indent	(Optional) The level the printing is indented, useful if nested within another S3 object

**Value**

No return value, called for side effects.

**Examples**

```
# Define a transition function, which sets all columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Define an S3 golden_transition
trn <- new_transition(test_transition, c(), "b")
print(trn)
```

---

run_simulation	<i>Execute a patient trajectory simulation</i>
----------------	--

---

**Description**

Execute a patient trajectory simulation

**Usage**

```
run_simulation(initPop, parameters)
```

**Arguments**

initPop	data.table containing initial population for simulation
parameters	Simulation configuration

**Value**

An list containing final population, history and timing data.tables

**Examples**

```
library(data.table)
N <- 100
dt <- data.table(a = runif(N, 0, 1), b = rep(0, N))
# Define a hazard function, which returns a vector of equal length uncertainties
test_hazard <- function(a) {
  ret <- (a < 0.5)
}
# Define a transition function, which sets all "b" columns affected by the hazard to 100
test_transition <- function() {
  return (100)
}
# Create an S3 golden_hazard
haz <- new_hazard(
  test_hazard,
```

```
        c("a"),
        new_transition(test_transition, c(), "b")
    )
# Define a trajectory function, which adds 2 to all members of the input vector
test_trajectory <- function(a) {
  return (a + 2)
}
# Define an S3 golden_trajectory
trj <- new_trajectory(test_trajectory, c("b"), "b")
# Create an S3 golden_history, containing 1 golden_history_column
hist <- new_history(new_column("sum_a", sum, c("a")))
# Define an S3 golden_parameters
params <- new_parameters(
  hazards = haz,
  trajectories = trj,
  steps = 10,
  debug = FALSE,
  history = hist
)
# Run the simulation to collect results
results <- run_simulation(dt, params)
```

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