

# Package: microsimulation (via r-universe)

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**Type** Package

**Title** Discrete Event Simulation in R and C++, with Tools for  
Cost-Effectiveness Analysis

**Version** 1.4.4

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**Description** Discrete event simulation using both R and C++ (Karlsson et al 2016; <[doi:10.1109/eScience.2016.7870915](https://doi.org/10.1109/eScience.2016.7870915)>). The C++ code is adapted from the SSIM library <<https://www.inf.usi.ch/carzaniga/ssim/>>, allowing for event-oriented simulation. The code includes a SummaryReport class for reporting events and costs by age and other covariates. The C++ code is available as a static library for linking to other packages. A priority queue implementation is given in C++ together with an S3 closure and a reference class implementation. Finally, some tools are provided for cost-effectiveness analysis.

**License** GPL (>= 3)

**Depends** Rcpp (>= 0.10.2), methods

**Imports** parallel, grDevices, ascii, survival

**Suggests** testthat

**LinkingTo** Rcpp, RcppArmadillo

**LazyData** true

**URL** <https://github.com/mclements/microsimulation>

**BugReports** <https://github.com/mclements/microsimulation/issues>

**Encoding** UTF-8

**RoxygenNote** 7.3.2

**Repository** <https://mclements.r-universe.dev>

**RemoteUrl** <https://github.com/mclements/microsimulation>

**RemoteRef** HEAD

**RemoteSha** fd767bb3aa2f335f5deb44965085a7468d1769e0

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microsimulation-package  
*microsimulation*

---

### Description

Discrete event simulations in both R and C++ with Tools for Cost-Effectiveness Analysis.

### Introduction

Discrete event simulations in both R and C++ with Tools for Cost-Effectiveness Analysis.

### Author(s)

Mark Clements <mark.clements@ki.se>

### References

<https://github.com/mclements/microsimulation>

### See Also

[sourceCpp](#)

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.microsimulationLdFlags

*Internal function*

---

### **Description**

Is this function needed? We could define the current stream in open code.

Again, is this needed?

### **Usage**

```
.microsimulationLdFlags()

inlineCxxPlugin(...)

LdFlags()

microsimulation.init(PACKAGE = "microsimulation")

microsimulation.exit(PACKAGE = "microsimulation")

unsigned(seed)

signed(seed)

rnormPos(n, mean = 0, sd = 1, lbound = 0)

set.user.Random.seed(seed, PACKAGE = "microsimulation")

advance.substream(seed, n, PACKAGE = "microsimulation")

next.user.Random.substream(PACKAGE = "microsimulation")

user.Random.seed(PACKAGE = "microsimulation")

enum(obj, labels, start = 0)

enum(obj) <- value

RNGstate()

frontier(x, y, concave = TRUE, convex = NULL)

lines_frontier(x, y, pch = 19, type = "b", ...)

discountedPoint(y, time, dr)
```

ICER(object1, object2, ...)

.onLoad(lib, pkg)

.onUnload(libpath)

### Arguments

...	other arguments
PACKAGE	package for the seed
seed	random number seed
n	number of sub-streams to advance
mean	numeric for the mean of the (untruncated) normal distribution (default=0)
sd	numeric for the sd of the (untruncated) normal distribution (default=1)
lbound	numeric for the lower bound (default=0)
obj	integer or logical for factor levels
labels	labels for the factor levels
start	first value of the levels
value	labels for the factor levels
x	vector of x coordinates
y	the undiscounted value
concave	logical for whether to calculate a concave frontier (default=TRUE)
convex	logical for whether to calculate a convex frontier (default=NULL)
pch	type of pch for the plotted symbols (default=19)
type	join type (default="b")
time	the time of the event
dr	discount rate, expressed as a percentage
object1	first object
object2	second object
lib	library string
pkg	package string
libpath	library path string

### Value

No return value, called for side effects

No return value, called for side effects

No return value, called for side effects

unsigned seed

signed seed

numeric vector  
invisibly returns the new seed  
the advanced seed  
invisibly returns TRUE – called for side effect  
random seed  
the new factor  
update the factor  
a list with oldseed (the old value of `.Random.seed`), and `reset()`, which resets `.Random.seed`  
a list with components `x` and `y` for the frontier  
No return value, called for side effects  
numeric vector

---

`callCalibrationPerson` *call CalibrationPerson example*

---

## **Description**

Example that uses the `RngStream` random number generator  
Example that uses the Mersenne-Twister random number generator  
Example that uses the Mersenne-Twister random number generator  
Example that uses the Mersenne-Twister random number generator

## **Usage**

```
callCalibrationPerson(  
  seed = 12345,  
  n = 500,  
  runpar = c(4, 0.5, 0.05, 10, 3, 0.5),  
  mc.cores = 1  
)  
  
callPersonSimulation(n = 20, seed = rep(12345, 6))  
  
callSimplePerson(n = 10)  
  
callSimplePerson2(n = 10)  
  
callIllnessDeath(n = 10L, cure = 0.1, zsd = 0)
```

**Arguments**

seed	random number seed
n	number of simulations (default=10)
runpar	parameters
mc.cores	number of cores
cure	probability of cure
zsd	frailty standard deviation

**Value**

data-frame  
data-frame  
data-frame  
data-frame  
data-frame

---

discountedInterval     *Integrate a discounted value*

---

**Description**

Integrate a discounted value

**Usage**

```
discountedInterval(y, start, finish, dr)
```

**Arguments**

y	the undiscounted value
start	the start time
finish	the finish time
dr	discount rate, expressed as a percentage

**Value**

numeric discounted value

---

fhcrcData	<i>Old data used in the prostata model</i>
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---

**Description**

Old data used in the prostata model

**Usage**

fhcrcData

**Format**

An object of class `list` of length 10.

---

pqueue	<i>S3 priority queue implementation using C++</i>
--------	---

---

**Description**

This provides a priority queue that is sorted by the priority and entry order. The priority is assumed to be numeric. The events can be of any type. As an extension, events can be cancelled if they satisfy a certain predicate. Note that the inactive events are not removed, rather they are marked as cancelled and will not be available to be popped.

Based on C++ code. See also the S3 implementation `pqueue`.

This event queue is simple and useful for pedagogic purposes.

Inherit from this class to represent a discrete event simulation. The API is similar to that for `Omnet++`, where an `init` method sets up the initial events using the `scheduleAt(time, event)` method, the messages are handled using the `handleMessage(event)` method, the simulation is run using the `run` method, and the `final` method is called at the end of the simulation.

**Usage**

```
pqueue(lower = TRUE)
```

**Arguments**

lower	boolean to determine whether to give priority to lower values (default=TRUE) or higher values
-------	---

**Details**

The algorithm for pushing values into the queue is computationally very simple: simply rank the times using `order()` and re-order times and events. This approach is probably of acceptable performance for smaller queue. A more computationally efficient approach for pushing into larger queues would be to use a binary search (e.g. using `findInterval()`).

For faster alternatives, see `pqueue` and `PQueueRef`.

**Value**

a list with

**push** function with arguments priority (numeric) and event (SEXP). Pushes an event with a given priority

**pop** function to return a list with a priority (numeric) and an event (SEXP). This pops the first active event.

**cancel** function that takes a predicate (or R function) for a given event and returns a logical that indicates whether to cancel that event or not. This may cancel some events that will no longer be popped.

**empty** function that returns whether the priority queue is empty (or has no active events).

**clear** function to clear the priority queue.

**ptr** XPtr value

**Fields**

ptr External pointer to the C++ class

times vector of times

events list of events

times vector of times

events list of events

**Methods**

cancel(predicate) Method to cancel events that satisfy some predicate

clear() Method to clear the event queue

empty() Method to check whether there are no events in the queue

initialize(lower = TRUE) Method to initialize the object. lower argument indicates whether lowest priority or highest priority

pop() Method to remove the head of the event queue and return its value

push(priority, event) Method to push an event with a given priority

cancel(predicate, ...) Method to remove events that satisfy some predicate

clear() Method to clear the event queue

empty() Method to check whether there are no events in the queue

pop() Method to remove the head of the event queue and return its value

push(time, event) Method to insert the event at the given time

final() Method for finalising the simulation

handleMessage(event) Virtual method to handle the messages as they arrive

init() Virtual method to initialise the event queue and attributes

reset(startTime = 0) Method to reset the event queue

run(startTime = 0) Method to run the simulation

scheduleAt(time, event) Method that adds attributes for the event time and the sendingTime, and then insert the event into the event queue



**Examples**

```

pq = pqueue()
pq$push(3,"Clear drains")
pq$push(4, "Feed cat")
pq$push(5, "Make tea")
pq$push(1, "Solve RC tasks")
pq$push(2, "Tax return")
while(!pq$empty())
  print(pq$pop())

pq = new("PQueueRef")
pq$push(3,"Clear drains")
pq$push(4, "Feed cat")
pq$push(5, "Make tea")
pq$push(1, "Solve RC tasks")
pq$push(2, "Tax return")
while(!pq$empty())
  print(pq$pop())

pq = new("EventQueue")
pq$push(3,"Clear drains")
pq$push(4, "Feed cat")
pq$push(5, "Make tea")
pq$push(1, "Solve RC tasks")
pq$push(2, "Tax return")
while(!pq$empty())
  print(pq$pop())

DES = setRefClass("DES",
  contains = "BaseDiscreteEventSimulation",
  methods=list(
    init=function() {
      scheduleAt(3,"Clear drains")
      scheduleAt(4, "Feed cat")
      scheduleAt(5, "Make tea")
      scheduleAt(1, "Solve RC tasks")
      scheduleAt(2, "Tax return")
    },
    handleMessage=function(event) print(event)))

des = new("DES")
des$run()
## Not run:
testRsimulation1 <- function() {
  ## A simple example
  Simulation <-
    setRefClass("Simulation",
      contains = "BaseDiscreteEventSimulation")
  Simulation$methods(
    init = function() {
      scheduleAt(rweibull(1,8,85), "Death due to other causes")
      scheduleAt(rweibull(1,3,90), "Cancer diagnosis")
    }
  )
}

```

```

    },
    handleMessage = function(event) {
      if (event %in% c("Death due to other causes", "Cancer death")) {
        clear()
        print(event)
      }
      else if (event == "Cancer diagnosis") {
        if (runif(1) < 0.5)
          scheduleAt(now() + rweibull(1,2,10), "Cancer death")
        print(event)
      }
    }
  })
  Simulation$new()$run()
}

## An extension with individual life histories
testRsimulation2 <- function(n=100) {
  Simulation <-
    setRefClass("Simulation",
      contains = "BaseDiscreteEventSimulation",
      fields = list(state = "character", report = "data.frame"))
  Simulation$methods(
    init = function() {
      report <-< data.frame()
      state <-< "Healthy"
      scheduleAt(rweibull(1,8,85), "Death due to other causes")
      scheduleAt(rweibull(1,3,90), "Cancer diagnosis")
    },
    handleMessage = function(event) {
      report <-< rbind(report, data.frame(state = state,
        begin = attr(event,"sendingTime"),
        end = currentTime,
        event = event,
        stringsAsFactors = FALSE))
      if (event %in% c("Death due to other causes", "Cancer death")) {
        clear()
      }
      else if (event == "Cancer diagnosis") {
        state <-< "Cancer"
        if (runif(1) < 0.5)
          scheduleAt(now() + rweibull(1,2,10), "Cancer death")
      }
    },
    final = function() report)
  sim <- Simulation$new()
  do.call("rbind", lapply(1:n, function(id) data.frame(id=id,sim$run())))
}

## reversible illness-death model
testRsimulation3 <- function(n=100) {
  Simulation <-
    setRefClass("Simulation",
      contains = "BaseDiscreteEventSimulation",

```

```

        fields = list(state = "character", everCancer = "logical",
                      report = "data.frame"))
Simulation$methods(
  init = function() {
    report <- data.frame()
    state <- "Healthy"
    everCancer <- FALSE
    scheduleAt(rweibull(1,8,85), "Death due to other causes")
    scheduleAt(rweibull(1,3,90), "Cancer diagnosis")
  },
  handleMessage = function(event) {
    report <- rbind(report, data.frame(state = state,
                                      everCancer = everCancer,
                                      begin = attr(event,"sendingTime"),
                                      end = currentTime,
                                      event = event,
                                      stringsAsFactors = FALSE))

    if (event %in% c("Death due to other causes", "Cancer death")) {
      clear()
    }
    else if (event == "Cancer diagnosis") {
      state <- "Cancer"
      everCancer <- TRUE
      if (runif(1) < 0.5)
        scheduleAt(now() + rweibull(1,2,10), "Cancer death")
      scheduleAt(now() + 10, "Recovery")
    }
    else if (event == "Recovery") {
      state <- "Healthy"
      scheduleAt(now() + rexp(1,10), "Cancer diagnosis")
    }
  },
  final = function() report)
sim <- Simulation$new()
do.call("rbind", lapply(1:n, function(id) data.frame(id=id,sim$run())))
}

## cancer screening
testRsimulation4 <- function(n=1) {
  Simulation <-
    setRefClass("Simulation",
                contains = "BaseDiscreteEventSimulation",
                fields = list(state = "character", report = "data.frame"))
Simulation$methods(
  init = function() {
    report <- data.frame()
    state <- "Healthy"
    scheduleAt(rweibull(1,8,85), "Death due to other causes")
    scheduleAt(rweibull(1,3,90), "Cancer onset")
    scheduleAt(50,"Screening")
  },
  handleMessage = function(event) {
    report <- rbind(report, data.frame(state = state,

```

```

begin = attr(event,"sendingTime"),
end = currentTime,
event = event,
stringsAsFactors = FALSE))
if (event %in% c("Death due to other causes", "Cancer death")) {
  clear()
}
else if (event == "Cancer onset") {
  state <- event
  dx <- now() + rweibull(1,2,10)
  scheduleAt(dx, "Clinical cancer diagnosis")
  scheduleAt(dx + rweibull(1,1,10), "Cancer death")
  scheduleAt(now() + rweibull(1,1,10), "Metastatic cancer")
}
else if (event == "Metastatic cancer") {
  state <- event
  cancel(function(event) event %in%
    c("Clinical cancer diagnosis","Cancer death")) # competing events
  scheduleAt(now() + rweibull(1,2,5), "Cancer death")
}
else if (event == "Clinical cancer diagnosis") {
  state <- event
  cancel(function(event) event == "Metastatic cancer")
}
else if (event == "Screening") {
  switch(state,
    "Cancer onset" = {
      state <- "Screen-detected cancer diagnosis"
      cancel(function(event) event %in%
        c("Clinical cancer diagnosis","Metastatic cancer"))
    },
    "Metastatic cancer" = {}, # ignore
    "Clinical cancer diagnosis" = {}, # ignore
    "Healthy" = {
      if (now()<=68) scheduleAt(now()+2, "Screening")
    }
  )
}
else stop(event)
},
final = function() report)
sim <- Simulation$new()
do.call("rbind", lapply(1:n, function(id) data.frame(id=id,sim$run())))
}

## ticking bomb - toy example
testRsimulation5 <- function(n=1) {
  Simulation <-
    setRefClass("Simulation",
      contains = "BaseDiscreteEventSimulation",
      fields = list(report = "data.frame"))
  Simulation$methods(
    init = function() {
      report <- data.frame()

```



**Value**

data-frame  
 No return value, called for side effects  
 No return value, called for side effects  
 No return value, called for side effects  
 No return value, called for side effects  
 No return value, called for side effects  
 No return value, called for side effects

---

RNGStream	<i>S3 class to work with RngStream objects</i>
-----------	--

---

**Description**

S3 class to work with RngStream objects  
 Use RNGStream as an old class  
 With method for RNGStream S3 class

**Usage**

```
RNGStream(nextStream = TRUE, iseed = NULL)

## S3 method for class 'RNGStream'
with(data, expr, ...)
```

**Arguments**

nextStream	whether to move to the next stream (default=TRUE)
iseed	set seed after changing RNG (otherwise keep the current seed)
data	object of type RNGStream
expr	expression using the RNGStream
...	other arguments passed to eval()

**Value**

list of class RNGStream with components:

**resetRNGkind** function to reset to the previous RNG and seed  
**seed** function to return the current seed  
**open** function to use the current seed  
**close** function to make the current seed equal to .Random.seed  
**resetStream** function to move back to start of stream

**resetSubStream** function to move back to start of sub-stream

**nextSubStream** function to move to next sub-stream

**nextStream** function to move to next stream

the value from the expression

## Examples

```
## set up one stream
s1 <- RNGStream()
s1$open()
rnorm(1)
s1$nextSubStream()
rnorm(1)
## reset the stream
s1$resetStream()
rnorm(2)
s1$nextSubStream()
rnorm(2)

## now do with two streams
s1$resetStream()
s2 <- RNGStream()
with(s1,rnorm(1))
with(s2,rnorm(1))
s1$nextSubStream()
with(s1,rnorm(1))
## now reset the streams and take two samples each time
s1$resetStream()
s2$resetStream()
with(s1,rnorm(2))
with(s2,rnorm(2))
s1$nextSubStream()
with(s1,rnorm(2))
```

---

simulate.survreg

*Simulate event times from a survreg object*

---

## Description

Simulate event times from a survreg object

## Usage

```
## S3 method for class 'survreg'
simulate(object, nsim = 1, seed = NULL, newdata, t0 = NULL, ...)
```

**Arguments**

object	survreg object
nsim	number of simulations per row in newdata
seed	random number seed
newdata	data-frame for defining the covariates for the simulations. Required.
t0	delayed entry time. Defaults to NULL (which assumes that t0=0)
...	other arguments (not currently used)

**Value**

vector of event times with nsim repeats per row in newdata

**Examples**

```
library(survival)
fit <- survreg(Surv(time, status) ~ ph.ecog + age + sex + strata(sex),
              data = lung)
nd = transform(expand.grid(ph.ecog=0:1, sex=1:2), age=60)
simulate(fit, seed=1002, newdata=nd)
simulate(fit, seed=1002, newdata=nd, t0=500)
```

---

summary.SummaryReport *summary method for a SummaryReport object*

---

**Description**

At present, this passes the object to summary and then prints

**Usage**

```
## S3 method for class 'SummaryReport'
summary(object, ...)

## S3 method for class 'summary.SummaryReport'
print(x, ...)

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

## S3 method for class 'SummaryReport'
rbind(...)

## S3 method for class 'SummaryReport'
ascii(
  x,
  include.rownames = FALSE,
```



```

    include.colnames = TRUE,
    header = TRUE,
    digits = c(0, 3, 2, 2, 4, 4),
    ...
)

## S3 method for class 'SummaryReport'
ICER(object1, object2, ...)

## S3 method for class 'ICER.SummaryReport'
ascii(
  x,
  include.rownames = TRUE,
  include.colnames = TRUE,
  header = TRUE,
  digits = c(1, 1, 3, 3, 1, 1, 3, 3, 1),
  rownames = c("Reference", "Treatment"),
  colnames = c("Costs", "(se)", "QALYs", "(se)", "Costs", "(se)", "QALYs", "(se)",
    "ICER"),
  tgroup = c("Total", "Incremental"),
  n.tgroup = c(4, 5),
  ...
)

```

### Arguments

object	SummaryReport object
...	other arguments to pass to ascii
x	an ICER.SummaryReport object
include.rownames	logical for whether to include rownames (default=FALSE)
include.colnames	logical for whether to include colnames (default=TRUE)
header	logical for whether to include the header (default=TRUE)
digits	vector of the number of digits to use for each column
object1	SummaryReport object (reference)
object2	SummaryReport object
rownames	rownames for output
colnames	colnames for output
tgroup	tgroup arg passed to ascii
n.tgroup	arg passed to ascii

### Value

a list of class summary.SummaryReport with components:

**n** Number of simulations  
**indivip** boolean with whether individual values were retained  
**utilityDiscountRate** discount rate for utilities/QALYs  
**costDiscountRate** discount rate for costs  
**QALE** Quality-adjusted life expectancy (discounted)  
**LE** Life expectancy (not discounted)  
**ECosts** Life-time expected costs (discounted)  
**se.QALE** standard error for QALE  
**se.Ecosts** standard error Ecosts

a SummaryReport object  
ascii object  
a list of type ICER.SummaryReport with components:

**n** number of simulations  
**utilityDiscountRate** Discount rate for the utilities/QALE  
**costDiscountRate** Discount rate for the costs  
**s1** summary for object1  
**s2** summary for object2  
**dQALE** QALE for object2 minus QALE for object1  
**dCosts** Costs for object2 minus costs for object1  
**ICER** change of costs divided by change in QALEs  
**se.dQALE** standard error for dQALE  
**se.dCosts** standard error for dCosts

ascii object

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