

# Package ‘manCTMed’

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Psychological Methods.

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BootPara	<i>Parametric Bootstrap</i>
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## Description

The function generates simulated datasets based on a fitted model and refits the model to each generated dataset using the `dynr` package.

## Usage

```
BootPara(
  fit,
  path,
  prefix,
  taskid,
  B = 1000L,
  ncores = NULL,
  seed = NULL,
  clean = TRUE
)
```

## Arguments

<code>fit</code>	R object. Output of the <code>FitDynr()</code> , <code>FitMx()</code> , <code>IllustrationFitDynr()</code> , or <code>IllustrationFitMx()</code> , functions.
<code>path</code>	Path to a directory to store bootstrap samples and estimates.
<code>prefix</code>	Character string. Prefix used for the file names for the bootstrap samples and estimates.
<code>taskid</code>	Positive integer. Task ID.
<code>B</code>	Positive integer. Number of bootstrap samples.
<code>ncores</code>	Positive integer. Number of cores to use.
<code>seed</code>	Integer. Random seed.
<code>clean</code>	Logical. If <code>clean = TRUE</code> , delete intermediate files generated by the function.

**See Also**

Other Confidence Interval Functions: [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

**Examples**

```
## Not run:
set.seed(42)
library(dynrr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDy whole(data, taskid = 1)
BootPara(
  fit = fit,
  path = getwd(),
  prefix = "pb",
  taskid = 1,
  B = 1000L
)
## End(Not run)
```

BootParaStdXMY

*Parametric Bootstrap Confidence Intervals for X-M-Y (Standardized)***Description**

The function generates parametric bootstrap method confidence intervals for the mediation model  $X \rightarrow M \rightarrow Y$  (Standardized).

**Usage**

```
BootParaStdXMY(boot, theta_hat, delta_t = 1:30, ncores = NULL)
```

**Arguments**

boot	R object. Output of the <a href="#">BootPara()</a> function.
theta_hat	R object. Output of the <a href="#">ThetaHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
ncores	Positive integer. Number of cores to use.

**See Also**

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```

## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
boot <- BootPara(
  fit = fit,
  path = getwd(),
  prefix = "pb",
  taskid = 1,
  B = 1000L
)
theta_hat <- ThetaHat(fit)
ci <- BootParaStdXYM(boot = boot, theta_hat = theta_hat)
plot(ci)
plot(ci, type = "bc")

## End(Not run)

```

BootParaStdXYM

*Parametric Bootstrap Confidence Intervals for X-Y-M (Standardized)*

## Description

The function generates parametric bootstrap method confidence intervals for the mediation model  $X \rightarrow Y \rightarrow M$  (Standardized).

## Usage

```
BootParaStdXYM(boot, theta_hat, delta_t = 1:30, ncores = NULL)
```

## Arguments

boot	R object. Output of the <a href="#">BootPara()</a> function.
theta_hat	R object. Output of the <a href="#">ThetaHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
ncores	Positive integer. Number of cores to use.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```

## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
boot <- BootPara(
  fit = fit,
  path = getwd(),
  prefix = "pb",
  taskid = 1,
  B = 1000L
)
theta_hat <- ThetaHat(fit)
ci <- BootParaStdXYM(boot = boot, theta_hat = theta_hat)
plot(ci)
plot(ci, type = "bc")

## End(Not run)

```

BootParaStdYMX

*Parametric Bootstrap Confidence Intervals for Y-M-X (Standardized)*

## Description

The function generates parametric bootstrap method confidence intervals for the mediation model  $Y \rightarrow M \rightarrow X$  (Standardized).

## Usage

```
BootParaStdYMX(boot, theta_hat, delta_t = 1:30, ncores = NULL)
```

## Arguments

boot	R object. Output of the <a href="#">BootPara()</a> function.
theta_hat	R object. Output of the <a href="#">ThetaHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
ncores	Positive integer. Number of cores to use.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```

## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
boot <- BootPara(
  fit = fit,
  path = getwd(),
  prefix = "pb",
  taskid = 1,
  B = 1000L
)
theta_hat <- ThetaHat(fit)
ci <- BootParaStdYMX(boot = boot, theta_hat = theta_hat)
plot(ci)
plot(ci, type = "bc")

## End(Not run)

```

## BootParaXMY

### *Parametric Bootstrap Confidence Intervals for X-M-Y*

## Description

The function generates parametric bootstrap method confidence intervals for the mediation model  $X \rightarrow M \rightarrow Y$ .

## Usage

```
BootParaXMY(boot, phi_hat, delta_t = 1:30, ncores = NULL)
```

## Arguments

boot	R object. Output of the <a href="#">BootPara()</a> function.
phi_hat	R object. Output of the <a href="#">PhiHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
ncores	Positive integer. Number of cores to use.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```

## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
boot <- BootPara(
  fit = fit,
  path = getwd(),
  prefix = "pb",
  taskid = 1,
  B = 1000L
)
phi_hat <- PhiHat(fit)
ci <- BootParaXYM(boot = boot, phi_hat = phi_hat)
plot(ci)
plot(ci, type = "bc")

## End(Not run)

```

## BootParaXYM

### *Parametric Bootstrap Confidence Intervals for X-Y-M*

## Description

The function generates parametric bootstrap method confidence intervals for the mediation model  $X \rightarrow Y \rightarrow M$ .

## Usage

```
BootParaXYM(boot, phi_hat, delta_t = 1:30, ncores = NULL)
```

## Arguments

boot	R object. Output of the <a href="#">BootPara()</a> function.
phi_hat	R object. Output of the <a href="#">PhiHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
ncores	Positive integer. Number of cores to use.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```

## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
boot <- BootPara(
  fit = fit,
  path = getwd(),
  prefix = "pb",
  taskid = 1,
  B = 1000L
)
phi_hat <- PhiHat(fit)
ci <- BootParaYMX(boot = boot, phi_hat = phi_hat)
plot(ci)
plot(ci, type = "bc")

## End(Not run)

```

## BootParaYMX

### *Parametric Bootstrap Confidence Intervals for Y-M-X*

## Description

The function generates parametric bootstrap method confidence intervals for the mediation model  $Y \rightarrow M \rightarrow X$ .

## Usage

```
BootParaYMX(boot, phi_hat, delta_t = 1:30, ncores = NULL)
```

## Arguments

boot	R object. Output of the <a href="#">BootPara()</a> function.
phi_hat	R object. Output of the <a href="#">PhiHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
ncores	Positive integer. Number of cores to use.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXYM\(\)](#), [BootParaXYM\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
boot <- BootPara(
  fit = fit,
  path = getwd(),
  prefix = "pb",
  taskid = 1,
  B = 1000L
)
phi_hat <- PhiHat(fit)
ci <- BootParaYMX(boot = boot, phi_hat = phi_hat)
plot(ci)
plot(ci, type = "bc")

## End(Not run)
```

Compress

*Compress Replication*

## Description

Compress Replication

## Usage

```
Compress(taskid, repid, output_folder)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

## See Also

Other Compression Functions: [IllustrationCompress\(\)](#)

DeltaStdXYM

*Delta Method Confidence Intervals for X-M-Y (Standardized)*

## Description

The function generates delta method confidence intervals for the mediation model  $X \rightarrow M \rightarrow Y$  (Standardized).

## Usage

```
DeltaStdXYM(theta_hat, delta_t = 1:30)
```

## Arguments

- |           |  |
|-----------|--|
| theta_hat | R object. Output of the <a href="#">ThetaHat()</a> function. |
| delta_t   | Numeric vector. Vector of time intervals.                    |

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDydnr(data, taskid = 1)
theta_hat <- ThetaHat(fit)
ci <- DeltaStdXYM(theta_hat)
plot(ci)

## End(Not run)
```

DeltaStdXYM

*Delta Method Confidence Intervals for X-Y-M (Standardized)*

## Description

The function generates delta method confidence intervals for the mediation model  $X \rightarrow Y \rightarrow M$  (Standardized).

**Usage**

```
DeltaStdXYM(theta_hat, delta_t = 1:30)
```

**Arguments**

- theta\_hat R object. Output of the [ThetaHat\(\)](#) function.  
 delta\_t Numeric vector. Vector of time intervals.

**See Also**

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#),  
[BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#),  
[DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#),  
[MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

**Examples**

```
## Not run:  

set.seed(42)  

library(dynr)  

sim <- GenData(taskid = 1)  

data <- RandomMeasurement(sim)  

fit <- FitDy whole(data, taskid = 1)  

theta_hat <- ThetaHat(fit)  

ci <- DeltaStdXYM(theta_hat)  

plot(ci)  

  
## End(Not run)
```

DeltaStdYMX

*Delta Method Confidence Intervals for Y-M-X (Standardized)***Description**

The function generates delta method confidence intervals for the mediation model  $Y \rightarrow M \rightarrow X$  (Standardized).

**Usage**

```
DeltaStdYMX(theta_hat, delta_t = 1:30)
```

**Arguments**

- theta\_hat R object. Output of the [ThetaHat\(\)](#) function.  
 delta\_t Numeric vector. Vector of time intervals.

**See Also**

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#),  
[BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#),  
[DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#),  
[MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

**Examples**

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
theta_hat <- ThetaHat(fit)
ci <- DeltaStdYMX(theta_hat)
plot(ci)

## End(Not run)
```

**Description**

The function generates delta method confidence intervals for the mediation model  $X \rightarrow M \rightarrow Y$ .

**Usage**

```
DeltaXMY(phi_hat, delta_t = 1:30)
```

**Arguments**

- |         |  |
|---------|--|
| phi_hat | R object. Output of the <a href="#">PhiHat()</a> function. |
| delta_t | Numeric vector. Vector of time intervals.                  |

**See Also**

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#),  
[BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#),  
[DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#),  
[MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
phi_hat <- PhiHat(fit)
ci <- DeltaXYM(phi_hat)
plot(ci)

## End(Not run)
```

DeltaXYM

*Delta Method Confidence Intervals for X-Y-M*

## Description

The function generates delta method confidence intervals for the mediation model  $X \rightarrow Y \rightarrow M$ .

## Usage

```
DeltaXYM(phi_hat, delta_t = 1:30)
```

## Arguments

- |         |  |
|---------|--|
| phi_hat | R object. Output of the <a href="#">PhiHat()</a> function. |
| delta_t | Numeric vector. Vector of time intervals.                  |

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
phi_hat <- PhiHat(fit)
ci <- DeltaXYM(phi_hat)
plot(ci)

## End(Not run)
```

DeltaYMX

*Delta Method Confidence Intervals for Y-M-X***Description**

The function generates delta method confidence intervals for the mediation model  $Y \rightarrow M \rightarrow X$ .

**Usage**

```
DeltaYMX(phi_hat, delta_t = 1:30)
```

**Arguments**

- |         |  |
|---------|--|
| phi_hat | R object. Output of the <a href="#">PhiHat()</a> function. |
| delta_t | Numeric vector. Vector of time intervals.                  |

**See Also**

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

**Examples**

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDynr(data, taskid = 1)
phi_hat <- PhiHat(fit)
ci <- DeltaYMX(phi_hat)
plot(ci)

## End(Not run)
```

FigPlotEffects

*Plot Total, Direct, and Indirect Effects***Description**

Effects for the model  $X \rightarrow M \rightarrow Y$ .

**Usage**

```
FigPlotEffects(dynamics = 0, std = FALSE, max_delta_t = 30, xmy = TRUE)
```

### Arguments

dynamics	Integer. dynamics = 0 for original drift matrix, dynamics = -1 for near-neutral dynamics, and dynamics = 1 for stronger damping.
std	Logical. If std = TRUE, standardized total, direct, and indirect effects. If std = FALSE, unstandardized total, direct, and indirect effects.
max_delta_t	Numeric. Maximum time interval.
xmy	Logical. If xmy = TRUE, plot the effects for the $x \rightarrow m \rightarrow y$ mediation model. If xmy = FALSE, plot the effects for the $y \rightarrow m \rightarrow x'$ mediation model.

### Author(s)

Ivan Jacob Agaloos Pesigan

### See Also

Other Figure Functions: [FigScatterPlotCoverage\(\)](#), [FigScatterPlotPower\(\)](#), [FigScatterPlotSeBias\(\)](#), [FigScatterPlotType1\(\)](#), [IllustrationFigPlotEffects\(\)](#), [IllustrationFigScatterPlotCoverage\(\)](#), [IllustrationFigScatterPlotPower\(\)](#), [IllustrationFigScatterPlotSeBias\(\)](#)

### Examples

[FigPlotEffects\(\)](#)

## FigScatterPlotCoverage

*Plot Coverage Probabilities*

### Description

Coverage probabilities for the model  $X \rightarrow M \rightarrow Y$ .

### Usage

```
FigScatterPlotCoverage(results, delta_t = NULL, dynamics = 0, std = FALSE)
```

### Arguments

results	Summary results data frame.
delta_t	Vector of time-interval value. If delta_t = NULL, use all available time-intervals
dynamics	Integer. dynamics = 0 for original drift matrix, dynamics = -1 for near-neutral dynamics, and dynamics = 1 for stronger damping.
std	Logical. If std = TRUE, standardized total, direct, and indirect effects. If std = FALSE, unstandardized total, direct, and indirect effects.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**See Also**

Other Figure Functions: [FigPlotEffects\(\)](#), [FigScatterPlotPower\(\)](#), [FigScatterPlotSeBias\(\)](#), [FigScatterPlotType1\(\)](#), [IllustrationFigPlotEffects\(\)](#), [IllustrationFigScatterPlotCoverage\(\)](#), [IllustrationFigScatterPlotPower\(\)](#), [IllustrationFigScatterPlotSeBias\(\)](#)

**Examples**

```
data(results, package = "manCTMed")
FigScatterPlotCoverage(results)
FigScatterPlotCoverage(results, delta_t = 1:14)
FigScatterPlotCoverage(results, delta_t = 15:30)
```

**FigScatterPlotPower**     *Plot Statistical Power*

**Description**

Statistical power for the model  $X \rightarrow M \rightarrow Y$ .

**Usage**

```
FigScatterPlotPower(results, delta_t = NULL, dynamics = 0, std = FALSE)
```

**Arguments**

results	Summary results data frame.
delta_t	Vector of time-interval value. If <code>delta_t = NULL</code> , use all available time-intervals
dynamics	Integer. <code>dynamics = 0</code> for original drift matrix, <code>dynamics = -1</code> for near-neutral dynamics, and <code>dynamics = 1</code> for stronger damping.
std	Logical. If <code>std = TRUE</code> , standardized total, direct, and indirect effects. If <code>std = FALSE</code> , unstandardized total, direct, and indirect effects.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**See Also**

Other Figure Functions: [FigPlotEffects\(\)](#), [FigScatterPlotCoverage\(\)](#), [FigScatterPlotSeBias\(\)](#), [FigScatterPlotType1\(\)](#), [IllustrationFigPlotEffects\(\)](#), [IllustrationFigScatterPlotCoverage\(\)](#), [IllustrationFigScatterPlotPower\(\)](#), [IllustrationFigScatterPlotSeBias\(\)](#)

## Examples

```
data(results, package = "manCTMed")
FigScatterPlotPower(results)
FigScatterPlotPower(results, delta_t = 1:14)
FigScatterPlotPower(results, delta_t = 15:30)
```

**FigScatterPlotSeBias** *Plot Standard Error Bias*

## Description

Standard Error Bias for the model  $X \rightarrow M \rightarrow Y$ .

## Usage

```
FigScatterPlotSeBias(results, delta_t = NULL, dynamics = 0, std = FALSE)
```

## Arguments

results	Summary results data frame.
delta_t	Vector of time-interval value. If <code>delta_t = NULL</code> , use all available time-intervals
dynamics	Integer. <code>dynamics = 0</code> for original drift matrix, <code>dynamics = -1</code> for near-neutral dynamics, and <code>dynamics = 1</code> for stronger damping.
std	Logical. If <code>std = TRUE</code> , standardized total, direct, and indirect effects. If <code>std = FALSE</code> , unstandardized total, direct, and indirect effects.

## Author(s)

Ivan Jacob Agaloos Pesigan

## See Also

Other Figure Functions: [FigPlotEffects\(\)](#), [FigScatterPlotCoverage\(\)](#), [FigScatterPlotPower\(\)](#), [FigScatterPlotType1\(\)](#), [IllustrationFigPlotEffects\(\)](#), [IllustrationFigScatterPlotCoverage\(\)](#), [IllustrationFigScatterPlotPower\(\)](#), [IllustrationFigScatterPlotSeBias\(\)](#)

## Examples

```
data(results, package = "manCTMed")
FigScatterPlotSeBias(results)
```

**FigScatterPlotType1** *Plot Type I Error*


---

### Description

Type I error for the model  $Y \rightarrow M \rightarrow X$ .

### Usage

```
FigScatterPlotType1(results, delta_t = NULL, dynamics = 0, std = FALSE)
```

### Arguments

<code>results</code>	Summary results data frame.
<code>delta_t</code>	Vector of time-interval value. If <code>delta_t</code> = <code>NULL</code> , use all available time-intervals
<code>dynamics</code>	Integer. <code>dynamics</code> = 0 for original drift matrix, <code>dynamics</code> = -1 for near-neutral dynamics, and <code>dynamics</code> = 1 for stronger damping.
<code>std</code>	Logical. If <code>std</code> = <code>TRUE</code> , standardized total, direct, and indirect effects. If <code>std</code> = <code>FALSE</code> , unstandardized total, direct, and indirect effects.

### Author(s)

Ivan Jacob Agaloos Pesigan

### See Also

Other Figure Functions: [FigPlotEffects\(\)](#), [FigScatterPlotCoverage\(\)](#), [FigScatterPlotPower\(\)](#), [FigScatterPlotSeBias\(\)](#), [IllustrationFigPlotEffects\(\)](#), [IllustrationFigScatterPlotCoverage\(\)](#), [IllustrationFigScatterPlotPower\(\)](#), [IllustrationFigScatterPlotSeBias\(\)](#)

### Examples

```
data(results, package = "manCTMed")
FigScatterPlotType1(results)
FigScatterPlotType1(results, delta_t = 1:14)
FigScatterPlotType1(results, delta_t = 15:30)
```

---

**FitDynr***Fit the Model using the dynr Package*

---

## Description

The function fits the model using the [dynr::dynr](#) package.

## Usage

```
FitDynr(data, taskid)
```

## Arguments

- |        |   |
|--------|---|
| data   | R object. Output of the <a href="#">RandomMeasurement()</a> function. |
| taskid | Positive integer. Task ID.  |

## See Also

Other Model Fitting Functions: [FitMx\(\)](#), [IllustrationFitDynr\(\)](#), [IllustrationFitMx\(\)](#), [IllustrationMCPhiSigma\(\)](#), [IllustrationPrepData\(\)](#), [PhiHat\(\)](#), [ThetaHat\(\)](#)

## Examples

```
## Not run:  
set.seed(42)  
library(dynr)  
sim <- GenData(taskid = 1)  
data <- RandomMeasurement(sim)  
fit <- FitDynr(data, taskid = 1)  
summary(fit)  
  
## End(Not run)
```

---

**FitMx***Fit the Model using the OpenMx Package*

---

## Description

The function fits the model using the [OpenMx::OpenMx](#) package.

## Usage

```
FitMx(data, taskid)
```

**Arguments**

- `data` R object. Output of the [RandomMeasurement\(\)](#) function.  
`taskid` Positive integer. Task ID.

**See Also**

Other Model Fitting Functions: [FitDynr\(\)](#), [IllustrationFitDynr\(\)](#), [IllustrationFitMx\(\)](#), [IllustrationMCPhiSigma\(\)](#), [IllustrationPrepData\(\)](#), [PhiHat\(\)](#), [ThetaHat\(\)](#)

**Examples**

```
## Not run:
set.seed(42)
library(OpenMx)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitMx(data, taskid = 1)
summary(fit)

## End(Not run)
```

GenData

*Simulate Data***Description**

The function simulates data using the [simStateSpace::SimSSMOUFixed\(\)](#) function.

**Usage**

```
GenData(taskid)
```

**Arguments**

- `taskid` Positive integer. Task ID.

**See Also**

Other Data Generation Functions: [IllustrationGenData\(\)](#), [RandomMeasurement\(\)](#)

**Examples**

```
## Not run:
set.seed(42)
sim <- GenData(taskid = 1)
plot(sim)

## End(Not run)
```

---

 IllustrationBootPara *Parametric Bootstrap (Illustration)*


---

## Description

The function generates simulated datasets based on a fitted model and refits the model to each generated dataset using the `dynr` package.

## Usage

```
IllustrationBootPara(
  fit,
  path,
  prefix,
  taskid,
  B = 1000L,
  ncores = NULL,
  seed = NULL
)
```

## Arguments

<code>fit</code>	R object. Fitted CT-VAR model.
<code>path</code>	Path to a directory to store bootstrap samples and estimates.
<code>prefix</code>	Character string. Prefix used for the file names for the bootstrap samples and estimates.
<code>taskid</code>	Positive integer. Task ID.
<code>B</code>	Positive integer. Number of bootstrap samples.
<code>ncores</code>	Positive integer. Number of cores to use.
<code>seed</code>	Integer. Random seed.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
library(dynr)
sim <- IllustrationGenData(seed = 42)
data <- IllustrationPrepData(sim)
fit <- IllustrationFitDynr(data)
summary(fit)
```

```

IllustrationBootPara(
  fit = fit,
  path = getwd(),
  prefix = "pb",
  taskid = 1,
  B = 1000L,
  seed = 42
)

## End(Not run)

```

**IllustrationCompress    *Compress Replication (Illustration)***


---

### Description

Compress Replication (Illustration)

### Usage

```
IllustrationCompress(taskid, repid, output_folder)
```

### Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.

### Value

The output is saved as an external file in `output_folder`.

### Author(s)

Ivan Jacob Agaloos Pesigan

### See Also

Other Compression Functions: [Compress\(\)](#)

---

**IllustrationFigPlotEffects**

*Plot Total, Direct, and Indirect Effects (Illustration)*

---

**Description**

Effects for the model  $X \rightarrow M \rightarrow Y$ .

**Usage**

```
IllustrationFigPlotEffects(std = FALSE, max_delta_t = 30)
```

**Arguments**

- std            Logical. If std = TRUE, standardized total, direct, and indirect effects. If std = FALSE, unstandardized total, direct, and indirect effects.
- max\_delta\_t    Numeric. Maximum time interval.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**See Also**

Other Figure Functions: [FigPlotEffects\(\)](#), [FigScatterPlotCoverage\(\)](#), [FigScatterPlotPower\(\)](#), [FigScatterPlotSeBias\(\)](#), [FigScatterPlotType1\(\)](#), [IllustrationFigScatterPlotCoverage\(\)](#), [IllustrationFigScatterPlotPower\(\)](#), [IllustrationFigScatterPlotSeBias\(\)](#)

**Examples**

```
IllustrationFigPlotEffects(std = FALSE)  
IllustrationFigPlotEffects(std = TRUE)
```

---

**IllustrationFigScatterPlotCoverage**

*Illustration Plot Coverage Probabilities*

---

**Description**

Coverage probabilities for the model  $X \rightarrow M \rightarrow Y$ .

**Usage**

```
IllustrationFigScatterPlotCoverage(illustration_results)
```

**Arguments**

`illustration_results`  
 Summary results data frame.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**See Also**

Other Figure Functions: [FigPlotEffects\(\)](#), [FigScatterPlotCoverage\(\)](#), [FigScatterPlotPower\(\)](#),  
[FigScatterPlotSeBias\(\)](#), [FigScatterPlotType1\(\)](#), [IllustrationFigPlotEffects\(\)](#), [IllustrationFigScatterPlot\(\)](#),  
[IllustrationFigScatterPlotSeBias\(\)](#)

**Examples**

```
data(illustration_results, package = "manCTMed")
IllustrationFigScatterPlotCoverage(illustration_results)
```

**IllustrationFigScatterPlotPower**  
*Illustration Plot Statistical Power*

**Description**

Statistical Power for the model  $X \rightarrow M \rightarrow Y$ .

**Usage**

```
IllustrationFigScatterPlotPower(illustration_results)
```

**Arguments**

`illustration_results`  
 Summary results data frame.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**See Also**

Other Figure Functions: [FigPlotEffects\(\)](#), [FigScatterPlotCoverage\(\)](#), [FigScatterPlotPower\(\)](#),  
[FigScatterPlotSeBias\(\)](#), [FigScatterPlotType1\(\)](#), [IllustrationFigPlotEffects\(\)](#), [IllustrationFigScatterPlot\(\)](#),  
[IllustrationFigScatterPlotSeBias\(\)](#)

## Examples

```
data(illustration_results, package = "manCTMed")
IllustrationFigScatterPlotPower(illustration_results)
```

---

IllustrationFigScatterPlotSeBias  
Illustration Plot Standard Error Bias

---

## Description

Standard Error Bias for the model  $X \rightarrow M \rightarrow Y$ .

## Usage

```
IllustrationFigScatterPlotSeBias(illustration_results)
```

## Arguments

illustration\_results  
Summary results data frame.

## Author(s)

Ivan Jacob Agaloos Pesigan

## See Also

Other Figure Functions: [FigPlotEffects\(\)](#), [FigScatterPlotCoverage\(\)](#), [FigScatterPlotPower\(\)](#),  
[FigScatterPlotSeBias\(\)](#), [FigScatterPlotType1\(\)](#), [IllustrationFigPlotEffects\(\)](#), [IllustrationFigScatterPlot\(\)](#),  
[IllustrationFigScatterPlotPower\(\)](#)

## Examples

```
data(illustration_results, package = "manCTMed")
IllustrationFigScatterPlotSeBias(illustration_results)
```

---

**IllustrationFitDynr** *Fit the Model using the dynr Package (Illustration)*


---

### Description

The function fits the model using the [dynr::dynr](#) package.

### Usage

```
IllustrationFitDynr(data)
```

### Arguments

data	R object. Output of the <a href="#">IllustrationPrepData()</a> function.
------	--

### See Also

Other Model Fitting Functions: [FitDynr\(\)](#), [FitMx\(\)](#), [IllustrationFitMx\(\)](#), [IllustrationMCPhiSigma\(\)](#), [IllustrationPrepData\(\)](#), [PhiHat\(\)](#), [ThetaHat\(\)](#)

### Examples

```
## Not run:
library(dynr)
sim <- IllustrationGenData(seed = 42)
data <- IllustrationPrepData(sim)
fit <- IllustrationFitDynr(data)
summary(fit)

## End(Not run)
```

---

**IllustrationFitMx** *Fit the Model using the OpenMx Package (Illustration)*


---

### Description

The function fits the model using the [OpenMx::OpenMx](#) package.

### Usage

```
IllustrationFitMx(data)
```

### Arguments

data	R object. Output of the <a href="#">IllustrationPrepData()</a> function.
------	--

**See Also**

Other Model Fitting Functions: [FitDynr\(\)](#), [FitMx\(\)](#), [IllustrationFitDynr\(\)](#), [IllustrationMCPhiSigma\(\)](#), [IllustrationPrepData\(\)](#), [PhiHat\(\)](#), [ThetaHat\(\)](#)

**Examples**

```
## Not run:
library(OpenMx)
sim <- IllustrationGenData(seed = 42)
data <- IllustrationPrepData(sim)
fit <- IllustrationFitMx(data)
summary(fit)

## End(Not run)
```

**IllustrationGenData**     *Simulate Data (Illustration)*

**Description**

The function simulates data using the [simStateSpace::SimSSMOUFixed\(\)](#) function.

**Usage**

```
IllustrationGenData(seed = NULL, n = 133, m = 101, delta_t_gen = 0.1)
```

**Arguments**

seed	Integer. Random seed.
n	Positive integer. Sample size.
m	Positive integer. Measurement occasions.
delta_t_gen	Numeric. Time interval used to generate data.

**See Also**

Other Data Generation Functions: [GenData\(\)](#), [RandomMeasurement\(\)](#)

**Examples**

```
## Not run:
sim <- IllustrationGenData(seed = 42)
plot(sim)

## End(Not run)
```

**IllustrationMCPhiSigma**

*Generate a Sampling Distribution of Drift Matrices and Process Noise Covariance Matrices (Illustration)*

**Description**

The function generates a sampling distribution of drift matrices and process noise covariance matrices using the Monte Carlo method.

**Usage**

```
IllustrationMCPhiSigma(fit, R = 20000L, seed = NULL)
```

**Arguments**

fit	R object. Fitted CT-VAR model.
R	Positive integer. Number of Monte Carlo replications.
seed	Integer. Random seed.

**See Also**

Other Model Fitting Functions: [FitDynr\(\)](#), [FitMx\(\)](#), [IllustrationFitDynr\(\)](#), [IllustrationFitMx\(\)](#), [IllustrationPrepData\(\)](#), [PhiHat\(\)](#), [ThetaHat\(\)](#)

**Examples**

```
## Not run:
library(dynr)
sim <- IllustrationGenData(seed = 42)
data <- IllustrationPrepData(sim)
fit <- IllustrationFitDynr(data)
IllustrationMCPhiSigma(fit, seed = 42)

## End(Not run)
```

**IllustrationPrepData**    *Prepare Data Before Model Fitting (Illustration)***Description**

The function converts the output of [IllustrationGenData\(\)](#) into a data frame.

**Usage**

```
IllustrationPrepData(sim)
```

## Arguments

**sim** R object. Output of the [IllustrationGenData\(\)](#) function.

## See Also

Other Model Fitting Functions: [FitDynr\(\)](#), [FitMx\(\)](#), [IllustrationFitDynr\(\)](#), [IllustrationFitMx\(\)](#), [IllustrationMCPhiSigma\(\)](#), [PhiHat\(\)](#), [ThetaHat\(\)](#)

## Examples

```
## Not run:
sim <- IllustrationGenData(seed = 42)
data <- IllustrationPrepData(sim)
head(data)
dim(data)

## End(Not run)
```

illustration\_dist      *Illustration Sampling Distribution*

## Description

Illustration Sampling Distribution

## Usage

```
data(illustration_dist)
```

## Format

A matrix with 1000 rows and 27 columns:

**phi\_xx** Elements of the drift matrix.

**sigma\_xx** Elements of the process noise covariance matrix.

**theta\_xx** Elements of the measurement error covariance matrix.

**mu0\_x** Elements of the initial condition mean vector.

**sigma0\_xx** Elements of the initial condition covariance matrix.

## Author(s)

Ivan Jacob Agaloos Pesigan

**illustration\_dist\_dt** *Illustration Sampling Distribution Discrete Time - Time Interval of 1*

### Description

Illustration Sampling Distribution Discrete Time - Time Interval of 1

### Usage

```
data(illustration_dist_dt)
```

### Format

A matrix with 1000 rows and 15 columns:

**beta\_xx** Elements of the matrix of lagged coefficients.

**psi\_xx** Elements of the process noise covariance matrix.

### Author(s)

Ivan Jacob Agaloos Pesigan

**illustration\_dist\_dt\_mc**

*Illustration Sampling Distribution Discrete Time - Time Interval of 1  
(Monte Carlo Method)*

### Description

Illustration Sampling Distribution Discrete Time - Time Interval of 1 (Monte Carlo Method)

### Usage

```
data(illustration_dist_dt_mc)
```

### Format

A matrix with 20000 rows and 15 columns:

**beta\_xx** Elements of the matrix of lagged coefficients.

**psi\_xx** Elements of the process noise covariance matrix.

### Author(s)

Ivan Jacob Agaloos Pesigan

---

illustration\_dist\_mc *Illustration Sampling Distribution (Monte Carlo Method)*

---

**Description**

Illustration Sampling Distribution (Monte Carlo Method)

**Usage**

```
data(illustration_dist_mc)
```

**Format**

A matrix with 20000 rows and 15 columns:

**phi\_xx** Elements of the drift matrix.

**sigma\_xx** Elements of the process noise covariance matrix.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

illustration\_dist\_med *Illustration Sampling Distribution Total, Direct, and Indirect Effects - Time Interval of 1*

---

**Description**

Illustration Sampling Distribution Total, Direct, and Indirect Effects - Time Interval of 1

**Usage**

```
data(illustration_dist_med)
```

**Format**

A matrix with 1000 rows and 27 columns:

**total** Total effect.

**direct** Direct effect.

**indirect** Indirect effect.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**illustration\_dist\_med\_mc**

*Illustration Sampling Distribution Total, Direct, and Indirect Effects - Time Interval of 1 (Monte Carlo Method)*

**Description**

Illustration Sampling Distribution Total, Direct, and Indirect Effects - Time Interval of 1 (Monte Carlo Method)

**Usage**

```
data(illustration_dist_med_mc)
```

**Format**

A matrix with 20000 rows and 27 columns:

**total** Total effect.

**direct** Direct effect.

**indirect** Indirect effect.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**illustration\_dist\_med\_std**

*Illustration Sampling Distribution Standardized Total, Direct, and Indirect Effects - Time Interval of 1*

**Description**

Illustration Sampling Distribution Standardized Total, Direct, and Indirect Effects - Time Interval of 1

**Usage**

```
data(illustration_dist_med_std)
```

**Format**

A matrix with 1000 rows and 27 columns:

**total** Total effect.

**direct** Direct effect.

**indirect** Indirect effect.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

illustration\_dist\_med\_std\_mc

*Illustration Sampling Distribution Standardized Total, Direct, and Indirect Effects - Time Interval of 1 (Monte Carlo Method)*

---

**Description**

Illustration Sampling Distribution Standardized Total, Direct, and Indirect Effects - Time Interval of 1 (Monte Carlo Method)

**Usage**

```
data(illustration_dist_med_std_mc)
```

**Format**

A matrix with 20000 rows and 27 columns:

**total** Total effect.

**direct** Direct effect.

**indirect** Indirect effect.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

illustration\_results    *Illustration Small Scale Simulation Results*

---

**Description**

Illustration Small Scale Simulation Results

**Usage**

```
data(illustration_results)
```

**Format**

A with 22 columns:

**taskid** Task ID.

**replications** Number of replications.

**effect** Total, direct, or indirect effect.

**interval** Time interval.

**parameter** Population parameter.

**method** Method used to generate confidence intervals.

**xmy** Logical. TRUE for x to m to y path.

**std** Logical. TRUE for standardized. FALSE for unstandardized.

**est** Mean parameter estimate.

**se** Mean standard error.

**z** Mean *z* statistic.

**p** Mean *p*-value.

**R** Number of Monte Carlo or bootstrap replications.

**ll** Mean lower limit of the 95% confidence interval.

**ul** Mean upper limit of the 95% confidence interval.

**sig** Proportion of statistically significant results.

**zero\_hit** Proportion of replications where the confidence intervals included zero.

**theta\_hit** Proportion of replications where the confidence intervals included the population parameter.

**sq\_error** Mean squared error.

**se\_bias** Bias in standard error estimate.

**coverage** Coverage probability.

**power** Statistical power.

**Author(s)**

Ivan Jacob Agaloos Pesigan

## Description

The function generates Monte Carlo method confidence intervals for the mediation model  $X \rightarrow M \rightarrow Y$  (Standardized).

## Usage

```
MCStdXMY(theta_hat, delta_t = 1:30, R = 20000L, seed = NULL)
```

## Arguments

theta_hat	R object. Output of the <a href="#">ThetaHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.
seed	Integer. Random seed.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDyN(data, taskid = 1)
theta_hat <- ThetaHat(fit)
ci <- MCStdXMY(theta_hat, seed = 42)
plot(ci)

## End(Not run)
```

---

MCStdXYM*Monte Carlo Method Confidence Intervals for X-Y-M (Standardized)*

---

## Description

The function generates Monte Carlo method confidence intervals for the mediation model  $X \rightarrow Y \rightarrow M$  (Standardized).

## Usage

```
MCStdXYM(theta_hat, delta_t = 1:30, R = 20000L, seed = NULL)
```

## Arguments

theta_hat	R object. Output of the <a href="#">ThetaHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.
seed	Integer. Random seed.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDyN(data, taskid = 1)
theta_hat <- ThetaHat(fit)
ci <- MCStdXYM(theta_hat, seed = 42)
plot(ci)

## End(Not run)
```

## Description

The function generates Monte Carlo method confidence intervals for the mediation model  $Y \rightarrow M \rightarrow X$  (Standardized).

## Usage

```
MCStdYMX(theta_hat, delta_t = 1:30, R = 20000L, seed = NULL)
```

## Arguments

theta_hat	R object. Output of the <a href="#">ThetaHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.
seed	Integer. Random seed.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDyN(data, taskid = 1)
theta_hat <- ThetaHat(fit)
ci <- MCStdYMX(phi_hat, seed = 42)
plot(ci)

## End(Not run)
```

## Description

The function generates Monte Carlo method confidence intervals for the mediation model  $X \rightarrow M \rightarrow Y$ .

## Usage

```
MCXMY(phi_hat, delta_t = 1:30, R = 20000L, seed = NULL)
```

## Arguments

phi_hat	R object. Output of the <a href="#">PhiHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.
seed	Integer. Random seed.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXYM\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDy whole(data, taskid = 1)
phi_hat <- PhiHat(fit)
ci <- MCXMY(phi_hat, seed = 42)
plot(ci)

## End(Not run)
```

## Description

The function generates Monte Carlo method confidence intervals for the mediation model  $X \rightarrow Y \rightarrow M$ .

## Usage

```
MCXYM(phi_hat, delta_t = 1:30, R = 20000L, seed = NULL)
```

## Arguments

phi_hat	R object. Output of the <a href="#">PhiHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.
seed	Integer. Random seed.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCYMX\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDy whole(data, taskid = 1)
phi_hat <- PhiHat(fit)
ci <- MCXYM(phi_hat, seed = 42)
plot(ci)

## End(Not run)
```

## Description

The function generates Monte Carlo method confidence intervals for the mediation model  $Y \rightarrow M \rightarrow X$ .

## Usage

```
MCYMX(phi_hat, delta_t = 1:30, R = 20000L, seed = NULL)
```

## Arguments

phi_hat	R object. Output of the <a href="#">PhiHat()</a> function.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.
seed	Integer. Random seed.

## See Also

Other Confidence Interval Functions: [BootPara\(\)](#), [BootParaStdXMY\(\)](#), [BootParaStdXYM\(\)](#), [BootParaStdYMX\(\)](#), [BootParaXMY\(\)](#), [BootParaXYM\(\)](#), [BootParaYMX\(\)](#), [DeltaStdXMY\(\)](#), [DeltaStdXYM\(\)](#), [DeltaStdYMX\(\)](#), [DeltaXMY\(\)](#), [DeltaXYM\(\)](#), [DeltaYMX\(\)](#), [IllustrationBootPara\(\)](#), [MCStdXMY\(\)](#), [MCStdXYM\(\)](#), [MCStdYMX\(\)](#), [MCXMY\(\)](#), [MCXYM\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(taskid = 1)
data <- RandomMeasurement(sim)
fit <- FitDyN(data, taskid = 1)
phi_hat <- PhiHat(fit)
ci <- MCYMX(phi_hat, seed = 42)
plot(ci)

## End(Not run)
```

---

params	<i>Simulation Parameters</i>
--------	------------------------------

---

### Description

Simulation Parameters

### Usage

```
data(params)
```

### Format

A dataframe with 30 rows and 3 columns:

**taskid** Simulation Task ID.

**n** Sample size.

**dynamics** Dynamics. 0 for original drift matrix, -1 for near-neutral dynamics, and 1 for stronger damping.

### Author(s)

Ivan Jacob Agaloos Pesigan

---

PhiHat	<i>Estimated Drift Matrix</i>
--------	-------------------------------

---

### Description

The function extracts the estimated drift matrix from the fitted model.

### Usage

```
PhiHat(fit)
```

### Arguments

**fit** R object. Output of the [FitDynr\(\)](#), [FitMx\(\)](#), [IllustrationFitDynr\(\)](#), or [IllustrationFitMx\(\)](#), functions.

### See Also

Other Model Fitting Functions: [FitDynr\(\)](#), [FitMx\(\)](#), [IllustrationFitDynr\(\)](#), [IllustrationFitMx\(\)](#), [IllustrationMCPhiSigma\(\)](#), [IllustrationPrepData\(\)](#), [ThetaHat\(\)](#)

## Examples

```
## Not run:
set.seed(42)
library(dynr)
sim <- GenData(n = 50)
data <- RandomMeasurement(sim)
fit <- FitDynr(data)
PhiHat(fit)

## End(Not run)
```

**RandomMeasurement**      *Simulate Random Measurement*

## Description

The function randomly selects 100 observations from the generated data and replaces the unselected observations with NA.

## Usage

```
RandomMeasurement(sim)
```

## Arguments

**sim**      R object. Output of the [GenData\(\)](#) function.

## See Also

Other Data Generation Functions: [GenData\(\)](#), [IllustrationGenData\(\)](#)

## Examples

```
## Not run:
set.seed(42)
sim <- GenData(taskid = 1)
RandomMeasurement(sim)

## End(Not run)
```

---

**results***Simulation Results*

---

**Description**

Simulation Results

**Usage**

```
data(results)
```

**Format**

A dataframe with 24 columns:

- taskid** Task ID.
- replications** Number of replications.
- effect** Total, direct, or indirect effect.
- interval** Time interval.
- dynamics** Dynamics. 0 for original drift matrix, -1 for near-neutral dynamics, and 1 for stronger damping.
- parameter** Population parameter.
- method** Method used to generate confidence intervals.
- xmy** If TRUE, the mediation model is  $X \rightarrow M \rightarrow Y$ . If FALSE, the mediation model is  $Y \rightarrow M \rightarrow X$ .
- std** If TRUE, standardized total, direct, and indirect effects. If FALSE, unstandardized total, direct, and indirect effects.
- n** Sample size.
- est** Mean parameter estimate.
- se** Mean standard error.
- z** Mean  $z$  statistic.
- p** Mean  $p$ -value.
- R** Number of Monte Carlo replications.
- ll** Mean lower limit of the 95% confidence interval.
- ul** Mean upper limit of the 95% confidence interval.
- sig** Proportion of statistically significant results.
- zero\_hit** Proportion of replications where the confidence intervals contained zero.
- theta\_hit** Proportion of replications where the confidence intervals contained the population parameter.
- sq\_error** Mean squared error.
- se\_bias** Bias in standard error estimate.
- coverage** Coverage probability.
- power** Statistical power.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**Sim**

*Simulation Replication*

**Description**

Simulation Replication

**Usage**

```
Sim(
  taskid,
  repid,
  output_folder,
  overwrite,
  integrity,
  seed,
  ci,
  pb,
  delta_t,
  R,
  B
)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
seed	Integer. Random seed.
ci	Logical. Run simulations for confidence intervals.
pb	Logical. Run simulations for parametric bootstrap confidence intervals.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.
B	Positive integer. Number of bootstrap samples.

**Value**

The output is saved as an external file in output\_folder.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimDynrBootPara

*Simulation Replication - BootPara*

---

**Description**

Simulation Replication - BootPara

**Usage**

```
SimDynrBootPara(  
    taskid,  
    repid,  
    output_folder,  
    seed,  
    suffix,  
    overwrite,  
    integrity,  
    B,  
    ncores = NULL  
)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of manCTMed:::SimSuffix().
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
B	Positive integer. Number of bootstrap samples.
ncores	Positive integer. Number of cores to use.

**Details**

This function is executed via the Sim function.

**Value**

The output is saved as an external file in output\_folder.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

**SimDynrBootParaStdXMY** *Simulation Replication - BootParaStdXMY*

---

**Description**

Simulation Replication - BootParaStdXMY

**Usage**

```
SimDynrBootParaStdXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t
)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
delta_t	Numeric vector. Vector of time intervals.

**Details**

This function is executed via the `Sim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimDynrBootParaStdYMX *Simulation Replication - BootParaStdYMX*

---

**Description**

Simulation Replication - BootParaStdYMX

**Usage**

```
SimDynrBootParaStdYMX(  
  taskid,  
  repid,  
  output_folder,  
  seed,  
  suffix,  
  overwrite,  
  integrity,  
  delta_t  
)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
delta_t	Numeric vector. Vector of time intervals.

**Details**

This function is executed via the `Sim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimDynrBootParaXMY      *Simulation Replication - BootParaXMY*

---

**Description**

Simulation Replication - BootParaXMY

**Usage**

```
SimDynrBootParaXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t
)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>delta_t</code>	Numeric vector. Vector of time intervals.

**Details**

This function is executed via the `Sim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimDynrBootParaYMX      *Simulation Replication - BootParaYMX*

---

## Description

Simulation Replication - BootParaYMX

## Usage

```
SimDynrBootParaYMX(  
    taskid,  
    repid,  
    output_folder,  
    seed,  
    suffix,  
    overwrite,  
    integrity,  
    delta_t  
)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
delta_t	Numeric vector. Vector of time intervals.

## Details

This function is executed via the `Sim` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

**SimDynrDeltaStdXMY**      *Simulation Replication - DynrDeltaStdXMY*

## Description

Simulation Replication - DynrDeltaStdXMY

## Usage

```
SimDynrDeltaStdXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t
)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
delta_t	Numeric vector. Vector of time intervals.

## Details

This function is executed via the `Sim` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

---

SimDynrDeltaStdYMX      *Simulation Replication - DynrDeltaStdYMX*

---

## Description

Simulation Replication - DynrDeltaStdYMX

## Usage

```
SimDynrDeltaStdYMX(  
  taskid,  
  repid,  
  output_folder,  
  seed,  
  suffix,  
  overwrite,  
  integrity,  
  delta_t  
)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of manCTMed:::SimSuffix().
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
delta_t	Numeric vector. Vector of time intervals.

## Details

This function is executed via the Sim function.

## Value

The output is saved as an external file in output\_folder.

## Author(s)

Ivan Jacob Agaloos Pesigan

**SimDynrDeltaXMY**      *Simulation Replication - DynrDeltaXMY*

## Description

Simulation Replication - DynrDeltaXMY

## Usage

```
SimDynrDeltaXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t
)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
delta_t	Numeric vector. Vector of time intervals.

## Details

This function is executed via the `Sim` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

---

SimDynrDeltaYMX      *Simulation Replication - DynrDeltaYMX*

---

## Description

Simulation Replication - DynrDeltaYMX

## Usage

```
SimDynrDeltaYMX(  
    taskid,  
    repid,  
    output_folder,  
    seed,  
    suffix,  
    overwrite,  
    integrity,  
    delta_t  
)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of manCTMed:::SimSuffix().
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
delta_t	Numeric vector. Vector of time intervals.

## Details

This function is executed via the Sim function.

## Value

The output is saved as an external file in output\_folder.

## Author(s)

Ivan Jacob Agaloos Pesigan

SimDynrMCStdXMY

*Simulation Replication - DynrMCStdXMY***Description**

Simulation Replication - DynrMCStdXMY

**Usage**

```
SimDynrMCStdXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t,
  R
)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.

**Details**This function is executed via the `Sim` function.**Value**The output is saved as an external file in `output_folder`.**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimDynrMCStdYMX      *Simulation Replication - DynrMCStdYMX*

---

## Description

Simulation Replication - DynrMCStdYMX

## Usage

```
SimDynrMCStdYMX(  
  taskid,  
  repid,  
  output_folder,  
  seed,  
  suffix,  
  overwrite,  
  integrity,  
  delta_t,  
  R  
)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of manCTMed:::SimSuffix().
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.

## Details

This function is executed via the Sim function.

## Value

The output is saved as an external file in output\_folder.

## Author(s)

Ivan Jacob Agaloos Pesigan

SimDynrMCXMY

*Simulation Replication - DynrMCXMY***Description**

Simulation Replication - DynrMCXMY

**Usage**

```
SimDynrMCXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t,
  R
)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.

**Details**This function is executed via the `Sim` function.**Value**The output is saved as an external file in `output_folder`.**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimDynrMCYMX

*Simulation Replication - DynrMCYMX*

---

## Description

Simulation Replication - DynrMCYMX

## Usage

```
SimDynrMCYMX(  
  taskid,  
  repid,  
  output_folder,  
  seed,  
  suffix,  
  overwrite,  
  integrity,  
  delta_t,  
  R  
)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of manCTMed:::SimSuffix().
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
delta_t	Numeric vector. Vector of time intervals.
R	Positive integer. Number of Monte Carlo replications.

## Details

This function is executed via the Sim function.

## Value

The output is saved as an external file in output\_folder.

## Author(s)

Ivan Jacob Agaloos Pesigan

---

**SimFitDynr***Simulation Replication - FitDynr*

---

## Description

Simulation Replication - FitDynr

## Usage

```
SimFitDynr(taskid, repid, output_folder, seed, suffix, overwrite, integrity)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .

## Details

This function is executed via the `Sim` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

---

SimFitMx

*Simulation Replication - FitMx*

---

## Description

Simulation Replication - FitMx

## Usage

```
SimFitMx(taskid, repid, output_folder, seed, suffix, overwrite, integrity)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .

## Details

This function is executed via the `Sim` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

SimFN	<i>Simulation File Name</i>
-------	-----------------------------

**Description**

Simulation File Name

**Usage**

```
SimFN(output_type, output_folder, suffix)
```

**Arguments**

output_type	Character string. Output type.
output_folder	Character string. Output folder.
suffix	Character string. Output of manCTMed:::SimSuffix().

**Value**

Returns a character string file name with the `output_folder` in the OS-specific format.

SimGenData	<i>Simulation Replication - GenData</i>
------------	---

**Description**

Simulation Replication - GenData

**Usage**

```
SimGenData(taskid, repid, output_folder, seed, suffix, overwrite, integrity)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of manCTMed:::SimSuffix().
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Details**

This function is executed via the `Sim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimIllustration

*Simulation Replication (Illustration)*

---

**Description**

Simulation Replication (Illustration)

**Usage**

```
SimIllustration(  
  taskid,  
  repid,  
  output_folder,  
  overwrite,  
  integrity,  
  seed,  
  ci,  
  pb,  
  delta_t,  
  R,  
  B  
)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.
<code>seed</code>	Integer. Random seed.
<code>ci</code>	Logical. Run simulations for confidence intervals.
<code>pb</code>	Logical. Run simulations for parametric bootstrap confidence intervals.

<code>delta_t</code>	Numeric vector. Vector of time intervals.
<code>R</code>	Positive integer. Number of Monte Carlo replications.
<code>B</code>	Positive integer. Number of bootstrap samples.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**SimIllustrationDynrBootPara**  
*Simulation Replication - BootPara*

**Description**

Simulation Replication - BootPara

**Usage**

```
SimIllustrationDynrBootPara(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  B,
  ncores = NULL
)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>B</code>	Positive integer. Number of bootstrap samples.
<code>ncores</code>	Positive integer. Number of cores to use.

**Details**

This function is executed via the `IllustrationSim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimIllustrationDynrBootParaStdXMY  
*Simulation Replication - BootParaStdXMY*

---

**Description**

Simulation Replication - BootParaStdXMY

**Usage**

```
SimIllustrationDynrBootParaStdXMY(  
    taskid,  
    repid,  
    output_folder,  
    seed,  
    suffix,  
    overwrite,  
    integrity,  
    delta_t  
)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>delta_t</code>	Numeric vector. Vector of time intervals.

**Details**

This function is executed via the `IllustrationSim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**SimIllustrationDynrBootParaXMY**  
*Simulation Replication - BootParaXMY*

**Description**

Simulation Replication - BootParaXMY

**Usage**

```
SimIllustrationDynrBootParaXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t
)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>delta_t</code>	Numeric vector. Vector of time intervals.

## Details

This function is executed via the `IllustrationSim` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

`SimIllustrationDynrDeltaStdXMY`

*Simulation Replication - Illustration (DynrDeltaStdXMY)*

## Description

Simulation Replication - Illustration (DynrDeltaStdXMY)

## Usage

```
SimIllustrationDynrDeltaStdXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t
)
```

## Arguments

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>delta_t</code>	Numeric vector. Vector of time intervals.

**Details**

This function is executed via the `IllustrationSim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**SimIllustrationDynrDeltaXMY**

*Simulation Replication - Illustration (DynrDeltaXMY)*

**Description**

Simulation Replication - Illustration (DynrDeltaXMY)

**Usage**

```
SimIllustrationDynrDeltaXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t
)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>delta_t</code>	Numeric vector. Vector of time intervals.

**Details**

This function is executed via the `IllustrationSim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

`SimIllustrationDynrMCPhiSigma`

*Simulation Replication - Illustration (MCPhiSigma)*

**Description**

Simulation Replication - Illustration (MCPhiSigma)

**Usage**

```
SimIllustrationDynrMCPhiSigma(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  R
)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>R</code>	Positive integer. Number of Monte Carlo replications.

## Details

This function is executed via the `IllustrationSim` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

`SimIllustrationDynrMCStdXMY`

*Simulation Replication - Illustration (DynrMCStdXMY)*

## Description

Simulation Replication - Illustration (DynrMCStdXMY)

## Usage

```
SimIllustrationDynrMCStdXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t,
  R
)
```

## Arguments

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>delta_t</code>	Numeric vector. Vector of time intervals.
<code>R</code>	Positive integer. Number of Monte Carlo replications.

## Details

This function is executed via the `IllustrationSim` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

`SimIllustrationDynrMCXMY`

*Simulation Replication - Illustration (DynrMCXMY)*

## Description

Simulation Replication - Illustration (DynrMCXMY)

## Usage

```
SimIllustrationDynrMCXMY(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity,
  delta_t,
  R
)
```

## Arguments

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .
<code>delta_t</code>	Numeric vector. Vector of time intervals.
<code>R</code>	Positive integer. Number of Monte Carlo replications.

### Details

This function is executed via the `IllustrationSim` function.

### Value

The output is saved as an external file in `output_folder`.

### Author(s)

Ivan Jacob Agaloos Pesigan

## **SimIllustrationFitDynr**

*Simulation Replication - IllustrationFitDynr*

### Description

Simulation Replication - `IllustrationFitDynr`

### Usage

```
SimIllustrationFitDynr(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity
)
```

### Arguments

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

### Details

This function is executed via the `IllustrationSim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimIllustrationFitMx    *Simulation Replication - IllustrationFitMx*

---

**Description**

Simulation Replication - IllustrationFitMx

**Usage**

```
SimIllustrationFitMx(  
  taskid,  
  repid,  
  output_folder,  
  seed,  
  suffix,  
  overwrite,  
  integrity  
)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>repid</code>	Positive integer. Replication ID.
<code>output_folder</code>	Character string. Output folder.
<code>seed</code>	Integer. Random seed.
<code>suffix</code>	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .

**Details**

This function is executed via the `IllustrationSim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**SimIllustrationGenData**

*Simulation Replication - IllustrationGenData*

**Description**

Simulation Replication - IllustrationGenData

**Usage**

```
SimIllustrationGenData(
  taskid,
  repid,
  output_folder,
  seed,
  suffix,
  overwrite,
  integrity
)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
seed	Integer. Random seed.
suffix	Character string. Output of <code>manCTMed:::SimSuffix()</code> .
overwrite	Logical. Overwrite existing output in <code>output_folder</code> .
integrity	Logical. If <code>integrity = TRUE</code> , check for the output file integrity when <code>overwrite = FALSE</code> .

**Details**

This function is executed via the `IllustrationSim` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SimIllustrationPara    *Simulation Replication Parametric Bootstrap (Parallel)*

---

## Description

Simulation Replication Parametric Bootstrap (Parallel)

## Usage

```
SimIllustrationPara(  
  taskid,  
  repid,  
  output_folder,  
  overwrite,  
  integrity,  
  seed,  
  B  
)
```

## Arguments

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
seed	Integer. Random seed.
B	Positive integer. Number of bootstrap samples.

## Value

The output is saved as an external file in output\_folder.

## Author(s)

Ivan Jacob Agaloos Pesigan

SimPara

*Simulation Replication Parametric Bootstrap (Parallel)***Description**

Simulation Replication Parametric Bootstrap (Parallel)

**Usage**

```
SimPara(taskid, repid, output_folder, overwrite, integrity, seed, B)
```

**Arguments**

taskid	Positive integer. Task ID.
repid	Positive integer. Replication ID.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
seed	Integer. Random seed.
B	Positive integer. Number of bootstrap samples.

**Value**

The output is saved as an external file in output\_folder.

**Author(s)**

Ivan Jacob Agaloos Pesigan

SimProj

*Simulation Project Name***Description**

Simulation Project Name

**Usage**

```
SimProj()
```

**Value**

Returns the project name as a character string.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

Sum                    *Summary*

---

**Description**

Summary

**Usage**

`Sum(taskid, reps, output_folder, overwrite, integrity)`

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SumDynrDeltaStdXMY      *Summary (DynrDeltaStdXMY)*

---

**Description**

Summary (DynrDeltaStdXMY)

**Usage**

`SumDynrDeltaStdXMY(taskid, reps, output_folder, overwrite, integrity)`

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Details**

This function is executed via the `Sum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

`SumDynrDeltaStdYMX`      *Summary (DynrDeltaStdYMX)*

**Description**

Summary (DynrDeltaStdYMX)

**Usage**

```
SumDynrDeltaStdYMX(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Details**

This function is executed via the `Sum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SumDynrDeltaXMY

*Summary (DynrDeltaXMY)*

---

**Description**

Summary (DynrDeltaXMY)

**Usage**

```
SumDynrDeltaXMY(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Details**

This function is executed via the `Sum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**SumDynrDeltaYMX**      *Summary (DynrDeltaYMX)*

### Description

Summary (DynrDeltaYMX)

### Usage

```
SumDynrDeltaYMX(taskid, reps, output_folder, overwrite, integrity)
```

### Arguments

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.

### Details

This function is executed via the `Sum` function.

### Value

The output is saved as an external file in `output_folder`.

### Author(s)

Ivan Jacob Agaloos Pesigan

**SumDynrMCStdXMY**      *Summary (DynrMCStdXMY)*

### Description

Summary (DynrMCStdXMY)

### Usage

```
SumDynrMCStdXMY(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.

**Details**

This function is executed via the Sum function.

**Value**

The output is saved as an external file in output\_folder.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SumDynrMCStdYMX      *Summary (DynrMCStdYMX)*

---

**Description**

Summary (DynrMCStdYMX)

**Usage**

SumDynrMCStdYMX(taskid, reps, output\_folder, overwrite, integrity)

**Arguments**

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.

**Details**

This function is executed via the Sum function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

`SumDynrMCXMY`

*Summary (DynrMCXMY)*

**Description**

`Summary (DynrMCXMY)`

**Usage**

```
SumDynrMCXMY(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Details**

This function is executed via the `Sum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SumDynrMCYMX                  *Summary (DynrMCYMX)*

---

**Description**

Summary (DynrMCYMX)

**Usage**

```
SumDynrMCYMX(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.

**Details**

This function is executed via the Sum function.

**Value**

The output is saved as an external file in output\_folder.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

SumFitDynr                  *Summary (FitDynr)*

---

**Description**

Summary (FitDynr)

**Usage**

```
SumFitDynr(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Details**

This function is executed via the `Sum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

`SumIllustration`      *Summary (Illustration)*

**Description**

Summary (Illustration)

**Usage**

```
SumIllustration(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

**SumIllustrationDynrBootParaStdXMY**

*Summary - Illustration (DynrBootParaStdXMY)*

---

**Description**

Summary - Illustration (DynrBootParaStdXMY)

**Usage**

```
SumIllustrationDynrBootParaStdXMY(  
  taskid,  
  reps,  
  output_folder,  
  overwrite,  
  integrity,  
  type = "pc"  
)
```

**Arguments**

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
type	Character string. Confidence interval type.

**Details**

This function is executed via the `IllustrationSum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

**SumIllustrationDynrBootParaXMY**  
*Summary - Illustration (DynrBootParaXMY)*

## Description

Summary - Illustration (DynrBootParaXMY)

## Usage

```
SumIllustrationDynrBootParaXMY(
  taskid,
  reps,
  output_folder,
  overwrite,
  integrity,
  type = "pc"
)
```

## Arguments

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.
type	Character string. Confidence interval type.

## Details

This function is executed via the `IllustrationSum` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

---

SumIllustrationDynrDeltaStdXMY  
Summary - Illustration (DynrDeltaStdXMY)

---

## Description

Summary - Illustration (DynrDeltaStdXMY)

## Usage

```
SumIllustrationDynrDeltaStdXMY(  
  taskid,  
  reps,  
  output_folder,  
  overwrite,  
  integrity  
)
```

## Arguments

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.

## Details

This function is executed via the `IllustrationSum` function.

## Value

The output is saved as an external file in `output_folder`.

## Author(s)

Ivan Jacob Agaloos Pesigan

**SumIllustrationDynrDeltaXMY***Summary - Illustration (DynrDeltaXMY)***Description**

Summary - Illustration (DynrDeltaXMY)

**Usage**`SumIllustrationDynrDeltaXMY(taskid, reps, output_folder, overwrite, integrity)`**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Details**This function is executed via the `IllustrationSum` function.**Value**The output is saved as an external file in `output_folder`.**Author(s)**

Ivan Jacob Agaloos Pesigan

**SumIllustrationDynrMCStdXMY***Summary - Illustration (DynrMCStdXMY)***Description**

Summary - Illustration (DynrMCStdXMY)

**Usage**`SumIllustrationDynrMCStdXMY(taskid, reps, output_folder, overwrite, integrity)`

**Arguments**

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.

**Details**

This function is executed via the `IllustrationSum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

`SumIllustrationDynrMCXMY`

*Summary - Illustration (DynrMCXMY)*

**Description**

Summary - Illustration (DynrMCXMY)

**Usage**

```
SumIllustrationDynrMCXMY(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

taskid	Positive integer. Task ID.
reps	Positive integer. Number of replications.
output_folder	Character string. Output folder.
overwrite	Logical. Overwrite existing output in output_folder.
integrity	Logical. If integrity = TRUE, check for the output file integrity when overwrite = FALSE.

**Details**

This function is executed via the `IllustrationSum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

`SumIllustrationFitDynr`

*Summary - Illustration (FitDynr)*

---

**Description**

Summary - Illustration (FitDynr)

**Usage**

```
SumIllustrationFitDynr(taskid, reps, output_folder, overwrite, integrity)
```

**Arguments**

<code>taskid</code>	Positive integer. Task ID.
<code>reps</code>	Positive integer. Number of replications.
<code>output_folder</code>	Character string. Output folder.
<code>overwrite</code>	Logical. Overwrite existing output in <code>output_folder</code> .
<code>integrity</code>	Logical. If <code>integrity</code> = TRUE, check for the output file integrity when <code>overwrite</code> = FALSE.

**Details**

This function is executed via the `IllustrationSum` function.

**Value**

The output is saved as an external file in `output_folder`.

**Author(s)**

Ivan Jacob Agaloos Pesigan

---

ThetaHat	<i>Estimated Drift Matrix and Process Noise</i>
----------	---

---

## Description

The function extracts the estimated drift matrix and process noise from the fitted model.

## Usage

```
ThetaHat(fit)
```

## Arguments

fit R object. Output of the [FitDynr\(\)](#), [FitMx\(\)](#), [IllustrationFitDynr\(\)](#), or [IllustrationFitMx\(\)](#), functions.

## See Also

Other Model Fitting Functions: [FitDynr\(\)](#), [FitMx\(\)](#), [IllustrationFitDynr\(\)](#), [IllustrationFitMx\(\)](#), [IllustrationMCPhiSigma\(\)](#), [IllustrationPrepData\(\)](#), [PhiHat\(\)](#)

## Examples

```
## Not run:  
set.seed(42)  
library(dynr)  
sim <- GenData(n = 50)  
data <- RandomMeasurement(sim)  
fit <- FitDynr(data)  
ThetaHat(fit)  
  
## End(Not run)
```

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