

# Package ‘EvalEst’

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**Title** Dynamic Systems Estimation - Extensions

**Description** Provides functions for evaluating (time series) model estimation methods. These facilitate Monte Carlo experiments of repeated simulations and estimations. Also provides methods for looking at the distribution of the results from these experiments, including model roots (which are an equivalence class invariant).

**Depends** R (>= 2.5.0), tfplot, dse (>= 2007.10-1)

**Imports** setRNG, tframe (>= 2007.5-3), stats, graphics

**LazyLoad** yes

**License** GPL-2

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coef.TSmodelEstEval     *Specific Methods for coef*

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## Description

See the generic function description.

## Usage

```
## S3 method for class 'TSmodelEstEval'
coef(object, criterion.args=NULL, ...)
## S3 method for class 'TSEstModelEstEval'
coef(object, criterion.args=NULL, ...)
```

## Arguments

`object`        an object (model) from which to extract coefficients(parameters).  
`criterion.args` arguments to be passed to this method when it is called by `EstEval`.  
`...`            (further arguments, currently disregarded).

## Details

The methods `***.ee` are intended mainly to be called from `EstEval` as criterion for evaluating an estimation method. See `coef`.

## See Also

[EstEval](#) [coef](#)

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distribution	<i>Generate distribution plots of Monte Carlo simulations</i>
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## Description

Generate distribution plots of Monte Carlo simulations.

## Usage

```

distribution(obj, ...)
  ## S3 method for class 'TSdata'
distribution(obj, ..., bandwidth=0.2,
            select.inputs = seq(length= nseriesInput(obj)),
            select.outputs= seq(length=nseriesOutput(obj)))
  ## Default S3 method:
distribution(obj, ..., bandwidth=0.2, series=NULL)

  ## S3 method for class 'MonteCarloSimulations'
distribution(obj,
            series=seq(dim(obj$simulations)[2]),
            x.sections=TRUE, periods=1:3, graphs.per.page=5, ...)

```

## Arguments

obj	The result of MonteCarloSimulations.
bandwidth	passed to density or ksmooth.
series	The series which should be plotted. The default gives all series.
select.inputs	series to be plotted. (passed to selectSeries)
select.outputs	series to be plotted. (passed to selectSeries)
x.sections	If TRUE then kernel density estimates are plotted for periods indicated by periods. If FALSE then a time series plots of the mean and estimates 1 and 2 standard deviations from the mean. Periods is ignored if x.sections is FALSE.
periods	The periods at which the distribution should be calculated and plotted. The default gives the first three.
graphs.per.page	integer indicating number of graphs to place on a page.
...	(further arguments, currently disregarded).
select	integer vector indicating roots to be plotted. If select is not NULL then roots are sorted by magnitude and only the indicated roots are plotted. For example, select=c(1,2) will plot only the two largest roots.

## Details

Kernel estimates of the densities (series by series, not joint densities) are estimated using ksmooth (if available) or density (if available) to produce density plots. Output graphics can be paused between pages by setting par(ask=TRUE).

**Value**

None

**See Also**

[tfplot.MonteCarloSimulations](#)

**Examples**

```
data("eg1.DSE.data.diff", package="dse")
model <- estVARXls(eg1.DSE.data.diff)
z <- MonteCarloSimulations(model)
distribution(z)
```

**distribution.coefEstEval**

*Plot distribution of estimates*

**Description**

Plot distribution of estimates.

**Usage**

```
## S3 method for class 'coefEstEval'
distribution(obj, ..., Sort=FALSE, bandwidth=0.2,
graphs.per.page=5)
## S3 method for class 'rootsEstEval'
distribution(obj, ..., mod=TRUE, invert=FALSE, Sort=FALSE,
bandwidth=0.2, select=NULL)
```

**Arguments**

<b>obj</b>	an object as returned by EstEval.
<b>Sort</b>	if Sort is true then sort is applied. This helps (a bit) with estimation methods like black.box which may not return parameters of the same length or in the same order.
<b>bandwidth</b>	passed to density or ksmooth.
<b>graphs.per.page</b>	integer indicating number of graphs to place on a page.
<b>...</b>	other objects to be plotted (not working for some methods).
<b>invert</b>	logical indicating if the inverse of roots should be plotted
<b>mod</b>	logical indicating if the modulus of roots should be plotted
<b>select</b>	integer vector indicating roots to be plotted. If select is not NULL then roots are sorted by magnitude and only the indicated roots are plotted. For example, select=c(1,2) will plot only the two largest roots.

**Details**

`ksmooth` is applied if available to get a smoothed estimate of the distribution of the estimates. If `ksmooth` is not available then `density` is applied if it is available.

**Value**

`None`

**See Also**

[EstEval](#)

**Examples**

```
data("eg1.DSE.data.diff", package="dse")
model <- estVARXls(TSdata(output=outputData(eg1.DSE.data.diff)), max.lag=2)
# now use this as the true model
z <- EstEval(model,
              estimation="estVARXls", estimation.args=list(max.lag=2))
distribution(z)
tfplot(z)
```

[EstEval](#)

*Evaluate an estimation method*

**Description**

Evaluate an estimation method.

**Usage**

```
EstEval(model, replications=100, rng=NULL, quiet=FALSE,
        simulation.args=NULL,
        estimation=NULL, estimation.args=NULL,
        criterion ="coef", criterion.args =NULL)

is.EstEval(obj)
```

**Arguments**

<code>model</code>	A TSmodel.
<code>replications</code>	The number of simulations.
<code>rng</code>	The RNG and starting seed.
<code>quiet</code>	If TRUE then no information is printed during estimation.
<code>simulation.args</code>	A list of any arguments to pass to simulate.
<code>estimation</code>	A character string indicating the estimation routine to use.

```

estimation.args          A list of any arguments to pass to the estimation routine.
criterion               A function to apply to the results of estimation to extract the information which
                        is to be retained.
criterion.args          A list of any arguments to be passed to the criterion function.
obj                     an object.

```

## Details

estimation.args and criterion.args should be NULL if no args are needed. If model is an object of class 'EstEval' or 'simulation' then the model and the seed!!! are extracted so the evaluation will be based on the same generated sample. criterion can be 'coef', 'roots', 'TSmodel', or 'TSEstModel'. With the default (coef) or with TSmodel the other criteria can be reconstructed (when the estimation method finds a known form for the model - which is not always the case, for example with estBlackBox methods). If criterion = 'roots' then criterion.args= list(verbose=FALSE) is advised.

## Value

A list with element result of length replications, each element containing the results of criterion(estimation(simulate(model))). Other elements of the list contain information from the supplied arguments.

## See Also

[simulate](#) [MonteCarloSimulations](#) [distribution](#) [forecast](#) [CovWRTtrue](#)

## Examples

```

data("eg1.DSE.data.diff", package="dse")
model <- estVARXls(TSdata(output=outputData(eg1.DSE.data.diff)))
z <- EstEval(model,
              estimation="estVARXls", estimation.args=list(max.lag=2))
tfplot(z)
zz <- EstEval(model,
              estimation="estVARXls", estimation.args=list(max.lag=2),
              simulation.args=list(sampleT=50, sd=1.5))
is.EstEval(zz)

```

generateSSmodel	<i>Randomly generate a state space model</i>
-----------------	--

## Description

Randomly generate a state space model.

## Usage

```
generateSSmodel(m,n,p, stable=FALSE)
```

**Arguments**

<code>n,m,p</code>	Input, state and output dimensions.
<code>stable</code>	TRUE or FALSE indicating if the model must be stable.

**Details**

Randomly generate a state space model. If stable is true then the largest root will have magnitude less than 1.0.

**Value**

An SS TSmodel.

**Examples**

```
z <- generateSSmodel(2,3,1)
```

genMineData

*Generate Data***Description**

Generate data for Monte Carlo experiments

**Usage**

```
genMineData(umodel, ymodel, uinput=NULL, sampleT=100,
            unoise=NULL, usd=1, ynoise=NULL, ysd=1, rng=NULL)
build.input.models(data, max.lag=NULL)
build.diagonal.model(multi.models)
```

**Arguments**

<code>umodel</code>	Model for input data.
<code>ymodel</code>	Model for output data.
<code>sampleT</code>	Number of periods of data to generate.
<code>unoise</code>	Input noise.
<code>usd</code>	Standard deviationof input noise.
<code>ynoise</code>	Output noise.
<code>ysd</code>	Standard deviation of output noise.
<code>rng</code>	RNG setting.
<code>multi.models</code>	A list of TSEstModels.
<code>data</code>	data from which to build models.
<code>max.lag</code>	number of lags in the estimated models.
<code>uinput</code>	Input data to umodel.

## Details

This function generates test data using specified models. `umodel` is used to generate data corresponding to input data and `ymodel` is used to generate data corresponding to output data. The result of `umodel` is used as input to `ymodel` so the input dimension of `ymodel` should be the output dimension of `umodel`. Typically the `ymodel` would be degenerate in some of the input variables so the effective inputs are a subset. If `umodel` requires input data it should be specified in `uinput`. If `noise` is `NULL` then normal noise will be generated by `simulate`. This will be iid  $N(0, I)$ . The RNG will be set first to `rng` if it is specified. If `unoise` or `ynoise` are specified they should be as expected by `simulate` for the specified `umodel` and `ymodel`.

`genMineData` uses `build.input.models`, which makes a list of univariate `TSeestModels`, one for each series in `inputData(data)` estimated by `estVARXls` with `max.lag` lags. `genMineData` then uses `build.diagonal.model` which builds one diagonal model from a list of models returned by `build.input.models`. It uses the AR part only.

## Value

A `TSdata` object.

## See Also

[simulate](#)

## Examples

```
data("eg1.DSE.data.diff", package="dse")
umodel <- build.diagonal.model(
  build.input.models(eg1.DSE.data.diff, max.lag=2))
z <- TSdata(output=outputData(eg1.DSE.data.diff),
             input = inputData(eg1.DSE.data.diff))
ymodel <- TSmodel(estVARXls(z, max.lag=3))
sim.data <- genMineData(umodel, ymodel)
```

## Description

Run multiple simulations

## Usage

```
is.MonteCarloSimulations(obj)
MonteCarloSimulations(model, simulation.args=NULL,
                      replications=100, rng=NULL, quiet =FALSE, ...)
## Default S3 method:
MonteCarloSimulations(model, simulation.args = NULL,
                      replications = 100, rng = NULL, quiet =FALSE, ...)
```

```

## S3 method for class 'TSmodel'
MonteCarloSimulations(model, simulation.args=NULL,
                      replications=100, rng=NULL, quiet=FALSE, ...)

## S3 method for class 'TSEstModel'
MonteCarloSimulations(model, simulation.args=NULL,
                      replications=100, rng=NULL, quiet=FALSE, ...)

## S3 method for class 'EstEval'
MonteCarloSimulations(model, simulation.args=NULL,
                      replications=100, rng=getRNG(model), quiet=FALSE, ...)

## S3 method for class 'MonteCarloSimulations'
MonteCarloSimulations(model,
                      simulation.args=NULL, replications=100, rng=getRNG(model), quiet=FALSE, ...)

```

## Arguments

<code>model</code>	an object from which a model can be extracted. The model must have an associated <code>simulation</code> method (e.g. a <code>TSmodel</code> ).
<code>simulation.args</code> ,	A list of arguments in addition to <code>model</code> which are passed to <code>simulate</code> .
<code>replications</code>	The number of simulations.
<code>rng</code>	The RNG and starting seed.
<code>quiet</code>	logical indicating if printing and many warning messages should be suppressed.
<code>obj</code>	an object.
<code>...</code>	arguments passed to other methods.

## Details

This function runs many simulations using `simulate`. Often it not be necessary to do this since the seed can be used to reproduce the sample and many functions for testing estimation methods, etc., will produce samples as they proceed. This function is useful for verification and for looking at the stochastic properties of the output of a model. If `model` is an object of class `EstEval` or `simulation` then the model and the seed!!! are extracted so the same sample will be generated. The default method expects the result of `simulate(model)` to be a matrix. There is a `tfplot` method (time series plots of the simulations) and a `distribution` method for the result. The latter plots kernel estimates of the distribution of the simulations at specified periods.

## Value

A list of simulations.

## See Also

[simulate](#) [EstEval](#) [distribution](#) [forecast](#) [CovWRTtrue](#)

## Examples

```
data("eg1.DSE.data.diff", package="dse")
model <- estVARXls(eg1.DSE.data.diff)
z <- MonteCarloSimulations(model, simulation.args=list(sampleT=100))
tfplot(z)
distribution(z)
```

**nseries.MonteCarloSimulations**  
*Number of Series*

## Description

Return the number of series.

## Usage

```
## S3 method for class 'MonteCarloSimulations'
nseriesInput(x)
## S3 method for class 'MonteCarloSimulations'
nseriesOutput(x)
```

## Arguments

**x** A featherForecasts object.

## Details

See the generic method.

## Value

An integer.

**print.estimatedModels** *Print Specific Methods*

## Description

See the generic function description.

**Usage**

```
## S3 method for class 'EstEval'
print(x, digits=options()$digits, ...)
## S3 method for class 'MonteCarloSimulations'
print(x, digits=options()$digits, ...)
```

**Arguments**

- x an object to be printed.
- digits a non-null value is used to indicate the number of significant digits. If digits is NULL then the value of digits specified by options is used.
- ... (further arguments, currently disregarded).

**See Also**

[print](#) [summary](#)

`roots.coefEstEval` *Roots Specific Methods*

**Description**

See the generic function description.

**Usage**

```
## S3 method for class 'coefEstEval'
roots(obj, criterion.args=NULL, ...)
## S3 method for class 'rootsEstEval'
roots(obj, ...)
## S3 method for class 'TSestModelEstEval'
roots(obj, criterion.args=NULL, ...)
## S3 method for class 'TSmodelEstEval'
roots(obj, criterion.args=list(randomize = TRUE), ...)
```

**Arguments**

- obj an object from which roots are to be extracted or calculated and printed.
- criterion.args arguments to be passed to this method when it is called by `EstEval`.
- ... arguments to be passed to other methods.

**Details**

The methods `***.ee` are intended mainly to be called from `EstEval` as criterion for evaluating an estimation method.

**See Also**

[roots](#) [stability](#) [EstEval](#)

[seriesNamesInput](#).[MonteCarloSimulations](#)  
*TS Input and Output Specific Methods*

**Description**

See the generic function description.

**Usage**

```
## S3 method for class 'MonteCarloSimulations'
seriesNamesInput(x)
## S3 method for class 'MonteCarloSimulations'
seriesNamesOutput(x)
```

**Arguments**

x an object from which to extract the names of the input or output series.

[summary.EstEval](#) *Summary Specific Methods*

**Description**

See the generic function description.

**Usage**

```
## S3 method for class 'TSEstModelEstEval'
summary(object, ...)
## S3 method for class 'TSmodelEstEval'
summary(object, ...)
## S3 method for class 'EstEval'
summary(object, ...)
## S3 method for class 'MonteCarloSimulations'
summary(object, series=NULL, periods=1:3, ...)
## S3 method for class 'coefEstEval'
summary(object, verbose=TRUE, ...)
## S3 method for class 'rootsEstEval'
summary(object, verbose=TRUE, ...)

## S3 method for class 'summary.TSEstModelEstEval'
```

```

print(x, digits=options()$digits, ...)
  ## S3 method for class 'summary.TSmodelEstEval'
print(x, digits=options()$digits, ...)
  ## S3 method for class 'summary.EstEval'
print(x, digits=options()$digits, ...)
  ## S3 method for class 'summary.MonteCarloSimulations'
print(x, digits=options()$digits, ...)
  ## S3 method for class 'summary.coefEstEval'
print(x, digits=options()$digits, ...)
  ## S3 method for class 'summary.rootsEstEval'
print(x, digits=options()$digits, ...)

```

## Arguments

object	an object for which a summary is to be printed.
x	an object for which a summary is to be printed.
digits	a non-null value is used to indicate the number of significant digits. If digits is NULL then the value of digits specified by options is used.
series	The series which should be plotted. The default NULL gives all series.
periods	optional integer vector indicating periods at which the summary should be calculated.
verbose	logical indicating if a longer summary should be produced.
...	arguments passed to other methods.

## See Also

[summary](#) [print](#)

**testEqual.EstEval**      *Specific Methods for Testing Equality*

## Description

See the generic function description.

## Usage

```

## S3 method for class 'EstEval'
testEqual(obj1, obj2, fuzz=0)
## S3 method for class 'MonteCarloSimulations'
testEqual(obj1, obj2, fuzz=1e-16)

```

## Arguments

<code>obj1</code>	an object which is to be compared with the second object.
<code>obj2</code>	an object which is to be compared with the first object.
<code>fuzz</code>	tolerance for numerical comparisons. Values within fuzz will be considered equal.

## See Also

[testEqual](#)

`tfplot.coefEstEval`      *Specific tfplot methods for coefEstEval (EstEval) objects*

## Description

See the generic function description.

## Usage

```
## S3 method for class 'coefEstEval'
tfplot(x, cumulate=TRUE, norm=FALSE, bounds=TRUE,
       invert=FALSE, Sort=FALSE, graphs.per.page = 5, ...)
```

## Arguments

<code>x</code>	an object for which a tfplot is to be produced.
<code>cumulate</code>	logical indicating if the cumulative average of roots should be plotted
<code>invert</code>	logical indicating if the inverse of roots should be plotted
<code>Sort</code>	logical indicating if the roots should be sorted.
<code>graphs.per.page</code>	integer indicating number of graphs to place on a page.
<code>norm</code>	logical indicating if the euclidean norm of roots should be plotted (square root of the sum of squared roots).
<code>bounds</code>	logical indicating if estimated one standard error bounds should be plotted around the lines for the true roots.
<code>...</code>	arguments passed to other methods.

## Details

If `cumulate` is true the cumulative average is plotted. If `norm` is true the norm is used, each parameter is plotted. If `invert` is true the reciprocal is used (before cumulating). If `Sort` is true then sort is applied (before ave). This is not usually recommended but of interest with estimation methods like `black.box` which may not return parameters of the same length or in the same order. Plotting the true lines only makes sense if `truth` is the same length as `result` (and sometimes not even then).

**See Also**

[tfplot](#) [EstEval](#)

**tfplot.MonteCarloSimulations**

*Generate plots of Monte Carlo simulations*

**Description**

Generate plots of Monte Carlo simulations.

**Usage**

```
## S3 method for class 'MonteCarloSimulations'
tfplot(x,
        tf=tframe(x$simulations), start=tfstart(tf), end=tfend(tf),
        series=seq((dim(x$simulations)[2])), select.simulations=seq(dim(x$simulations)[3]),
        graphs.per.page=5, mar=par()$mar, ...)
```

**Arguments**

<code>x</code>	The result of <code>MonteCarloSimulations</code> .
<code>tf</code>	The time frame for plots. see <code>tfplot</code> .
<code>start</code>	The starting period for plots, taken from <code>tf</code> by default.
<code>end</code>	The ending period for plots, taken from <code>tf</code> by default.
<code>series</code>	The series which should be plotted. The default <code>NULL</code> gives all series.
<code>select.simulations</code>	Vector of integers indicating the simulations which should be plotted. The default plots all simulations.
<code>graphs.per.page</code>	The number of graphs to put on a page.
<code>mar</code>	Plot margins (see <code>par</code> ).
<code>...</code>	arguments passed to other methods.

**Details**

This function produces plots of the simulated series. Output graphics can be paused between pages by setting `par(ask=TRUE)`.

**Value**

`None`

**See Also**

[distribution.MonteCarloSimulations](#)

**Examples**

```
data("eg1.DSE.data.diff", package="dse")
model <- estVARXls(eg1.DSE.data.diff)
z <- MonteCarloSimulations(model)
tfplot(z)
```

**tfplot.rootsEstEval**     *Specific tfplot methods for rootsEstEval (EstEval) objects*

**Description**

See the generic function description.

**Usage**

```
## S3 method for class 'rootsEstEval'
tfplot(x, ...)
## S3 method for class 'rootsEstEval'
plot(x, complex.plane=TRUE, cumulate=TRUE, norm=FALSE,
      bounds=TRUE, transform=NULL, invert=FALSE, Sort=TRUE, ...)
```

**Arguments**

- x                an object for which a tfplot is to be produced.
- complex.plane   logical indicating if the plot should be on the complex plane.
- cumulate        logical indicating if the cumulative average of roots should be plotted
- invert           logical indicating if the inverse of roots should be plotted
- Sort             logical indicating if the roots should be sorted.
- ...              arguments passed to other methods.
- norm             logical indicating if the euclidean norm of roots should be plotted (square root of the sum of squared roots).
- bounds           logical indicating if estimated one standard error bounds should be plotted around the lines for the true roots.
- transform        an optional string indicating the name of a function which should be applied to the roots before plotting.

## Details

If complex.plane is TRUE then all results are plotted on a complex plane and the arguments cumulate and Sort do not apply. If complex.plane is FALSE then a sequential plot of the real and imaginary parts is produced. If cumulate is true the cumulative average is plotted. If mod is true the modulus is used, otherwise real and imaginary are separated. if invert is true the reciprocal is used (before cumulating). if Sort is true then sort is applied (before cumulate but after mod) by the Re part of the root. Some grouping is usually necessary since roots are not in an obvious order but sorting by the real part of the roots could be improved upon.

## See Also

[tfplot](#) [EstEval](#)

[tfplot.TSdata.ee](#)      *Specific Methods for tfplot*

## Description

See the generic function description.

## Usage

```
## S3 method for class 'TSmodelEstEval'
tfplot(x, graph.args=NULL,
        criterion ="coef", criterion.args=NULL, ...)
## S3 method for class 'TSEstModelEstEval'
tfplot(x, graph.args=NULL,
        criterion ="coef", criterion.args=NULL, ...)
## S3 method for class 'EstEval'
tfplot(x, tf=NULL, start=tfstart(tf), end=tfend(tf),
       truth= if(is.TSdata(x$truth)) outputData(x$truth) else x$truth,
       series = seq(length=nseries(truth)),
       Title="Estimated (and true) results",
       ylab = seriesNames(truth), remove.mean = FALSE,
       graphs.per.page=5, mar=par()$mar, reset.screen=TRUE, ...)
```

## Arguments

x	an object for which a tfplot is to be produced.
tf	see <a href="#">tfplot</a> .
start	see <a href="#">tfplot</a> .
end	see <a href="#">tfplot</a> .
truth	true value which will be plotted along with estimates.
Title	string of characters to use for title.
remove.mean	logical indicating if means should be removed before plotting results.

```

ylab           vector of strings for y axis labelling.
graphs.per.page    integer indicating number of graphs to place on a page.
reset.screen   logical indicating if the plot window should be cleared before starting.
series         integer or string indicating the series which should be plotted.
mar            plot margins. See par.
graph.args     list of graphics arguments eventually passed to plot. See par.
criterion      criterion which should be used to extract something from the object which will
then be plotted. See EstEval.
criterion.args arguments to be passed to criterion.
...
arguments passed to other methods.

```

## See Also

[tfplot](#) [EstEval](#)

**TobsMonteCarloSimulations**

*Tframe or Number of Observations*

## Description

Return the number of Observations or the tframe.

## Usage

```

## S3 method for class 'MonteCarloSimulations'
Tobs(x)
## S3 method for class 'MonteCarloSimulations'
tframe(x)

```

## Arguments

x A MonteCarloSimulations object.

## Details

See the generic method.

## Value

An integer or a tframe object.

---

TSdata.coefEstEval      *TS Extractor Specific Methods*

---

### Description

See the generic function description.

### Usage

```
## S3 method for class 'coefEstEval'  
TSeModel(obj)  
## S3 method for class 'coefEstEval'  
TSmodel(obj, ...)
```

### Arguments

obj	an object from which to extract the TSmodel or TSeModel.
...	arguments to be passed to other methods.

### See Also

[TSdata](#) [TSeModel](#) [TSmodel](#)

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