# Package 'diagis'

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<b>Title</b> Diagnostic Plot and Multivariate Summary Statistics of Weighted Samples from Importance Sampling
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Contents
diagis

2 diagis

running_var	•					•				•			•	•	•	•
$running\_weighted\_mean$																
running_weighted_var .																
weighted_mean																
weighted_quantile																
weighted_se																
weighted_var																
weight_plot																

Index

diagis

Auxiliary functions and diagnostic plots for importance sampling

10

#### **Description**

This package contains functions computing weighted (running) summaries and diagonostic plots for importance sampling problems.

#### **Examples**

```
# simple importance sampling example
# true distribution is a standard normal:
p <- function(x) dnorm(x)</pre>
# proposal distribution is normal with sd s
q \leftarrow function(x, s) dnorm(x, 0, s)
# IS weights have finite variance only if s^2 > 1/2
# variance is s/(2-1/s^2)^(3/2)
#optimal case
set.seed(42)
s_opt <- sqrt(2)</pre>
x_{opt} \leftarrow rnorm(1000, sd = s_{opt})
w_{opt} \leftarrow p(x_{opt}) / q(x_{opt}, s_{opt})
weighted_mean(x_opt, w_opt)
weighted_var(x_opt, w_opt)
s_inf <- 0.25
x_{inf} \leftarrow rnorm(1000, sd = s_{inf})
w_{inf} \leftarrow p(x_{inf}) / q(x_{inf}, s_{inf})
weighted_mean(x_inf, w_inf) #!!
weighted_var(x_inf, w_inf) #!!
# diagnostic plots
weight_plot(w_inf)
weight_plot(w_opt)
```

ess 3

ess

Effective sample size

#### Description

Computes the effective sample size (ESS) of importance sampling estimator.

# Usage

```
ess(w, f, x)
```

#### **Arguments**

w A numeric vector of non-negative weights.

f A function used in computing f-specific ESS.

x A numeric vector of samples used to generate w. Used for computing f(x).

#### Value

An effective sample size estimate.

running\_ess

Running effective sample size

#### Description

Computes and returns the running estimate of effective sample size (ESS) of importance sampling estimator.

#### Usage

```
running_ess(w, f, x)
```

#### **Arguments**

w A numeric vector of non-negative weights.

f A function used in computing f-specific ESS.

x A numeric vector of samples used to generate w. Used for computing f(x).

#### Value

An effective sample size estimate.

running\_var

# Description

Computes running mean of a vector or matrix, returning the values from each step.

# Usage

```
running_mean(x, na.rm)
```

# **Arguments**

X	A numeric vector, matrix, three dimensional array, or an mcmc object from the coda package. For matrix, the mean is computed for each column, and for array the sweep is done over the third dimension.
na.rm	If TRUE, NA values in x are omitted from the computation. Default is FALSE.

# Value

A vector containing the recursive mean estimates.

# Description

Computes running variance of a vector, returning the values from each step.

# Usage

```
running_var(x, method = c("moment", "unbiased"), na.rm = FALSE)
```

# Arguments

x	A numeric vector or object that can be coerced to such.
method	Estimator type, either "moment" (default) or "unbiased", which is unbiased only in case of frequency weights.
na.rm	If TRUE, NA values in x are omitted from the computation. Default is FALSE.

# Value

A vector containing the recursive variance estimates.

running\_weighted\_mean Running weighted mean

# Description

Computes running weighted mean of a vector or matrix, returning the values from each step.

# Usage

```
running_weighted_mean(x, w, na.rm)
```

# Arguments

Х	A numeric vector, matrix, three dimensional array, or an mcmc object from the coda package. For matrix, the mean is computed for each column, and for array the sweep is done over the third dimension.
W	A numeric vector of non-negative weights. Will be automatically normalised to sum to one.
na.rm	If TRUE, NA values in x (and corresponding weights in w) are omitted from the computation. Default is FALSE. Only used in vector methods.

#### Value

A vector containing the recursive weighted mean estimates.

running\_weighted\_var Running weighted variance of a vector

# Description

Computes running weighted variance of a vector, returning the values from each step.

#### Usage

```
running_weighted_var(x, w, method = c("moment", "unbiased"), na.rm = FALSE)
```

# Arguments

x	A numeric vector or object that can be coerced to such.
W	A numeric vector of non-negative weights. Will be automatically normalised to sum to one.
method	Estimator type, either "moment" (default) or "unbiased", which is unbiased only in case of frequency weights.
na.rm	If TRUE, NA values in x (and corresponding weights in w) are omitted from the computation. Default is FALSE.

6 weighted\_quantile

#### Value

A vector containing the recursive weighted variance estimates.

weighted_	mean
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Weighted mean

# Description

Computes a weighted mean of a vector, matrix, or a three dimensional array.

#### Usage

```
weighted_mean(x, w, na.rm)
```

# Arguments

X	A numeric vector, matrix, three dimensional array, or an mcmc object from the coda package. For matrix, the mean is computed for each column, and for array the sweep is done over the third dimension.
W	A numeric vector of non-negative weights. Will be automatically normalised to sum to one.
na.rm	If TRUE, NA values in x (and corresponding weights in w) are omitted from the computation. Default is FALSE. Only used in vector methods.

#### Value

A weighted mean.

ghted_quantile Weighted quantiles
-----------------------------------

# Description

Computes a weighted quantiles of a vector or matrix. Based on the formula in Wikipedia (see the vignette) which is one of many ways to compute weighted quantiles.

#### Usage

```
weighted_quantile(x, w, probs = probs, na.rm)
```

weighted\_se 7

# Arguments

x	A numeric vector or matrix. For matrix, the quantiles are computed for each column.
W	A numeric vector of non-negative weights. Will be automatically normalised to sum to one.
probs	A numeric vector of probabilities with values between 0 and 1.
na.rm	If TRUE, NA and NaN values in x (and corresponding weights in w) are omitted from the computation. Default is FALSE. Additional missing values in w are not allowed.

#### Value

A weighted variance.

# Note

Compared to some other R functions, here the weights are regarded as probability weights, not frequency weights.

weighted_se	Weighted standard error	

# Description

Computes a weighted standard error of a vector or matrix.

#### Usage

```
weighted_se(x, w, na.rm)
```

# Arguments

Х	A numeric vector or matrix. For matrix, standard errors are computed for each column
W	A numeric vector of non-negative weights. Will be automatically normalised to sum to one.
na.rm	If TRUE, NA values in x (and corresponding weights in w) are omitted from the computation. Default is FALSE.

#### Value

A weighted variance.

# Note

Compared to some other R functions, here the weights are regarded as probability weights, not frequency weights.

8 weight\_plot

weighted_var Weighted covariance
----------------------------------

# Description

Computes a weighted variance/covariance of a vector, matrix or a three dimensional array.

#### Usage

```
weighted_var(x, w, method, na.rm)
```

#### **Arguments**

x	A numeric vector, matrix or three dimensional array. For matrix, covariances are computed between columns. For array, marginal covariances are computed for each column, i.e. for $m \times n \times k$ array function returns $m \times m \times n \times m \times n$ array.
W	A numeric vector of non-negative weights. Will be automatically normalised to sum to one.
method	Estimator type, either "moment" (default) or "unbiased", which is unbiased only in case of frequency weights.
na.rm	If TRUE, NA values in x (and corresponding weights in w) are omitted from the computation. Default is FALSE.

#### Value

A weighted variance.

#### Note

Compared to some other R functions, here the weights are regarded as probability weights, not frequency weights.

weight_plot Diagnostic plot of importance sampling weights	
--	--

# Description

Function weight\_plot plots four figures given the weight vector w: Plot of largest weights, sorted graph of all weights, running variance estimate of weights, and running effective sample size estimate of weights.

# Usage

```
weight_plot(w)
```

weight\_plot 9

#### **Arguments**

w Vector of weights.

#### **Examples**

```
#' importance sampling from too narrow distribution
#' weights have infinite variance
set.seed(1)
x_inf <- rnorm(1000, sd = 0.1)
w_inf <- dnorm(x_inf) / dnorm(x_inf, 0, 0.1)
weight_plot(w_inf)
x_opt <- rnorm(1000, sd = sqrt(2))
w_opt <- dnorm(x_opt) / dnorm(x_opt, 0, sqrt(2))
weight_plot(w_opt)</pre>
```

# **Index**

```
diagis, 2
diagis-package (diagis), 2
ess, 3
running_ess, 3
running_mean, 4
running_var, 4
running_weighted_mean, 5
running_weighted_var, 5
weight_plot, 8
weighted_mean, 6
weighted_quantile, 6
weighted_se, 7
weighted_var, 8
```