Package: chopthin (via r-universe)

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Type Package

Title The Chopthin Resampler

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Description Resampling is a standard step in particle filtering and in sequential Monte Carlo. This package implements the chopthin resampler, which keeps a bound on the ratio between the largest and the smallest weights after resampling.

License GPL-3

Imports Rcpp (>= 0.11.2)

LinkingTo Rcpp

Suggests testthat

RoxygenNote 6.0.1

NeedsCompilation yes

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Repository https://agandy.r-universe.dev

RemoteUrl https://github.com/cran/chopthin

RemoteRef HEAD

RemoteSha 9df0149cd0b2dc32727db535b6231b7ac4efd861

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chopthin

Description

A fast implementation of the Chopthin resampler. Can be used as the resampling step in particle filters and in sequential Monte Carlo.

Usage

chopthin(w, N, eta = 5.828427, normalise = TRUE, checks = TRUE)

Arguments

W	a vector of weights
Ν	target number of particles
eta	upper bound on the ratio between the weights. Must be $>=4$. If eta=Inf then only thinnig is performed, requiring the number of weights to be at least N.
normalise	Flag for controlling if the returned weights should be normalised. If TRUE (de- fault) then the sum of the returned weights will sum to N. If FALSE then the returned weights have the same sum as the original weights (within the numeri- cal precision).
checks	Flag controlling if checks on the input and the result should be performed. Default TRUE.

Value

A list with two elements: new weights and indices of the ancestors of the new particles. The weights are normalised to add up to N.

References

A Gandy and F. D-H Lau. The chopthin algorithm for resampling. IEEE Transactions on Signal Processing, 64(16):4273–4281, 2016

Examples

```
chopthin(runif(10),N=10)
chopthin(runif(10),N=20,4)
chopthin(runif(10),N=5)
chopthin(runif(10),N=1)
```

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