

# Package ‘bahc’

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**Type** Package

**Title** Filter Covariance and Correlation Matrices with  
Bootstrapped-Averaged Hierarchical Ansatz

**Version** 0.2.0

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**Author** Christian Bongiorno and Damien Challet

**Maintainer** Damien Challet <damien.challet@gmail.com>

**Description** A method to filter correlation and covariance matrices by averaging  
bootstrapped filtered hierarchical clustering. Now with boosting. See Ch. Bon-  
giorno and D. Challet,  
Covariance matrix filtering with bootstrapped hierarchies (2020) <arXiv:2003.05807>.

**License** GPL

**Depends** R (>= 3.5.0), fastcluster, matrixStats

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 7.1.0

**NeedsCompilation** no

**Repository** CRAN

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filterCorrelation      *Compute the BAHC correlation matrix.*

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### Description

Compute the BAHC correlation matrix.

### Usage

```
filterCorrelation(x, Nboot = 100)
```

### Arguments

x                      A matrix:  $x_{i,f}$  is feature  $f$  of object  $i$ .  
 Nboot                  The number of bootstrap copies

### Value

The BAHC-filtered correlation matrix of  $x$ .

### Examples

```
r=matrix(rnorm(1000),nrow=20) # 20 objects, 50 features each
Cor_bahc=filterCorrelation(r)
```

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filterCovariance      *Compute the BAHC covariance matrix.*

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### Description

Compute the BAHC covariance matrix.

### Usage

```
filterCovariance(x, k = 1, Nboot = 100)
```

### Arguments

x                      A matrix:  $x_{i,f}$  is feature  $f$  of object  $i$   
 k                      The order of filtering.  $k=1$  corresponds to BAHC.  
 Nboot                  The number of bootstrap copies

### Value

The BAHC-filtered correlation matrix of  $x$ .

**Examples**

```
r=matrix(rnorm(1000),nrow=20) # 20 objects, 50 features each
sigma=exp(runif(20))
rs=t(sigma %*% r) %*% sigma
Cov_bahc=filterCovariance(rs)
```

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