Package ‘fastmatrix’

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Type Package

Title Fast Computation of some Matrices Useful in Statistics

Version 0.2-3

Description Small set of functions to fast computation of some matrices and operations useful in statistics.

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URL https://faosorios.github.io/fastmatrix/

LazyLoad yes

NeedsCompilation yes

R topics documented:

array.mult *Array multiplication*

Description

Multiplication of 3-dimensional arrays was first introduced by Bates and Watts (1980). More extensions and technical details can be found in Wei (1998).

Usage

```
array.mult(a, b, x)
```

Arguments

a	a numeric matrix.
b	a numeric matrix.
x	a three-dimensional array.

Details

Let $\mathbf{X} = (x_{tij})$ be a 3-dimensional $n \times p \times q$ where indices t, i and j indicate face, row and column, respectively. The product $\mathbf{Y} = \mathbf{AXB}$ is an $n \times r \times s$ array, with $\mathbf{A}$ and $\mathbf{B}$ are $r \times p$ and $q \times s$ matrices respectively. The elements of $\mathbf{Y}$ are defined as:

$$y_{tkl} = \sum_{i=1}^{p} \sum_{j=1}^{q} a_{ki} x_{tij} b_{jl}$$

Value

`array.mult` returns a 3-dimensional array of dimension $n \times r \times s$.

References

Bates, D.M., Watts, D.G. (1980). Relative curvature measures of nonlinearity. *Journal of the Royal Statistical Society, Series B* **42**, 1-25.

Wei, B.C. (1998). *Exponential Family Nonlinear Models*. Springer, New York.

See Also

`array`, `matrix`, `bracket.prod`.

Examples

```
x <- array(0, dim = c(2,3,3)) # 2 x 3 x 3 array
x[,,1] <- c(1,2,2,4,3,6)
x[,,2] <- c(2,4,4,8,6,12)
x[,,3] <- c(3,6,6,12,9,18)

a <- matrix(1, nrow = 2, ncol = 3)
b <- matrix(1, nrow = 3, ncol = 2)

y <- array.mult(a, b, x) # a 2 x 2 x 2 array
y
```

bracket.prod *Bracket product*

Description

Bracket product of a matrix and a 3-dimensional array.

Usage

```
bracket.prod(a, x)
```

Arguments

a	a numeric matrix.
x	a three-dimensional array.

Details

Let $\mathbf{X} = (x_{tij})$ be a 3-dimensional $n \times p \times q$ array and $\mathbf{A}$ an $m \times n$ matrix, then $\mathbf{Y} = [\mathbf{A}][\mathbf{X}]$ is called the bracket product of $\mathbf{A}$ and $\mathbf{X}$, that is an $m \times p \times q$ with elements

$$y_{tij} = \sum_{k=1}^{n} a_{tk} x_{kij}$$

Value

`bracket.prod` returns a 3-dimensional array of dimension $m \times p \times q$.

References

Wei, B.C. (1998). *Exponential Family Nonlinear Models*. Springer, New York.

See Also

`array`, `matrix`, `array.mult`.

Examples

```
x <- array(0, dim = c(2,3,3)) # 2 x 3 x 3 array
x[,,1] <- c(1,2,2,4,3,6)
x[,,2] <- c(2,4,4,8,6,12)
x[,,3] <- c(3,6,6,12,9,18)

a <- matrix(1, nrow = 3, ncol = 2)

y <- bracket.prod(a, x) # a 3 x 3 x 3 array
y
```

dupl.cross *Matrix crossproduct envolving the duplication matrix*

Description

Given the order of two duplication matrices and matrix x, this function performs the operation: $\mathbf{Y} = \mathbf{D}_n^T \mathbf{X} \mathbf{D}_k$, where $\mathbf{D}_n$ and $\mathbf{D}_k$ are duplication matrices of order n and k, respectively.

Usage

```
dupl.cross(n = 1, k = n, x = NULL)
```

Arguments

n	order of the duplication matrix used pre-multiplying x.
k	order of the duplication matrix used post-multiplying x. By default k = n is used.
x	numeric matrix, this argument is required.

Details

This function calls dupl.prod to performs the matrix multiplications required but **without forming** any duplication matrices.

See Also

dupl.prod

Examples

```
D2 <- duplication(n = 2, matrix = TRUE)
D3 <- duplication(n = 3, matrix = TRUE)
x <- matrix(1, nrow = 9, ncol = 4)
y <- t(D3) %*% x %*% D2

z <- dupl.cross(n = 3, k = 2, x) # D2 and D3 are not stored
all(z == y) # matrices y and z are equal!

x <- matrix(1, nrow = 9, ncol = 9)
z <- dupl.cross(n = 3, x = x) # same matrix is used to pre- and post-multiplying x
z # print result
```

`dupl.info` *Compact information to construct the duplication matrix*

Description

This function provides the minimum information required to create the duplication matrix.

Usage

```
dupl.info(n = 1, condensed = TRUE)
```

Arguments

`n`	order of the duplication matrix.
`condensed`	logical. Information should be returned in compact form?

Details

This function returns a list containing two vectors that represent an element of the duplication matrix and is accesed by the indexes in vectors `row` and `col`. This information is used by function `dupl.prod` to do some operations involving the duplication matrix without forming it. This information also can be obtained using function `duplication`

Value

A list containing the following elements:

`row`	vector of indexes, each entry represents the row index of the duplication matrix. Only present if condensed = `FALSE`.
`col`	vector of indexes, each entry represents the column index of the duplication matrix.
`order`	order of the duplication matrix.

See Also

`duplication` `dupl.prod`

Examples

```
z <- dupl.info(n = 3, condensed = FALSE)
z # where are the ones in duplication of order 3?

D3 <- duplication(n = 3, matrix = TRUE)
D3 # only recommended if n is very small
```

`dupl.prod` *Matrix multiplication envolving the duplication matrix*

Description

Given the order of a duplication and matrix `x`, performs one of the matrix-matrix operations:

- $\mathbf{Y} = \mathbf{D}\mathbf{X}$, or
- $\mathbf{Y} = \mathbf{D}^T\mathbf{X}$, or
- $\mathbf{Y} = \mathbf{X}\mathbf{D}$, or
- $\mathbf{Y} = \mathbf{D}\mathbf{X}^T$,

where $\mathbf{D}$ is the duplication matrix of order n. The main aim of `dupl.prod` is to do this matrix multiplication **without forming** the duplication matrix.

Usage

```
dupl.prod(n = 1, x, transposed = FALSE, side = "left")
```

Arguments

`n`	order of the duplication matrix.
`x`	numeric matrix (or vector).
`transposed`	logical. Duplication matrix should be transposed?
`side`	a string selecting if duplication matrix is pre-multiplying `x`, that is `side = "left"` or post-multiplying `x`, by using `side = "right"`.

Details

Underlying `C` code only uses information provided by `dupl.info` to performs the matrix multiplication. The duplication matrix is **never** created.

See Also

`duplication`

Examples

```
D4 <- duplication(n = 4, matrix = TRUE)
x <- matrix(1, nrow = 16, ncol = 2)
y <- crossprod(D4, x)

z <- dupl.prod(n = 4, x, transposed = TRUE) # D4 is not stored
all(z == y) # matrices y and z are equal!
```

duplication — *Duplication matrix*

Description

This function returns the duplication matrix of order `n` which transforms, for a symmetric matrix `x`, `vech(x)` into `vec(x)`.

Usage

```
duplication(n = 1, matrix = FALSE, condensed = FALSE)
```

Arguments

`n`	order of the duplication matrix.
`matrix`	a logical indicating whether the duplication matrix will be returned.
`condensed`	logical. Information should be returned in compact form?.

Details

This function is a wrapper function for the function `dupl.info`. This function provides the minimum information required to create the duplication matrix. If option `matrix = FALSE` the duplication matrix is stored in two vectors containing the coordinate list of indexes for rows and columns. Option `condensed = TRUE` only returns vector of indexes for the columns of duplication matrix.

Warning: `matrix = TRUE` is **not** recommended, unless the order `n` be small. This matrix can require a huge amount of storage.

Value

Returns an n^2 by $n(n+1)/2$ matrix.

References

Magnus, J.R., and Neudecker, H. (1980). The elimination matrix, some lemmas and applications. *SIAM Journal on Algebraic Discrete Methods* **1**, 422-449.

Magnus, J.R., and Neudecker, H. (2007). *Matrix Differential Calculus with Applications in Statistics and Econometrics*, 3rd Edition. Wiley, New York.

See Also

`dupl.info`

Examples

```
z <- duplication(n = 100, condensed = TRUE)
object.size(z) # 40.5 Kb of storage

z <- duplication(n = 100, condensed = FALSE)
object.size(z) # 80.6 Kb of storage

D100 <- duplication(n = 100, matrix = TRUE)
```

```
object.size(D100) # 202 Mb of storage, do not request this matrix!

# a small example
D3 <- duplication(n = 3, matrix = TRUE)
a <- matrix(c( 1, 2, 3,
               2, 3, 4,
               3, 4, 5), nrow = 3)
upper <- vech(a)
v <- D3 %*% upper
all(vec(a) == as.vector(v)) # vectors are equal!
```

---

equilibrate *Column equilibration of a rectangular matrix*

Description

scale is generic function whose default method centers and/or scales the columns of a numeric matrix.

Usage

```
equilibrate(x, scale = TRUE)
```

Arguments

x	a numeric matrix.
scale	a logical value, the columns of x must be scaled to norm unity?.

Value

For scale = TRUE, the equilibrated (each column scaled to norm one) matrix. The scalings and an approximation of the reciprocal condition number, are returned as attributes "scales" and "condition".

Examples

```
x <- matrix(c(1, 1, 1,
              1, 2, 1,
              1, 3, 1,
              1, 1,-1,
              1, 2,-1,
              1, 3,-1), ncol = 3, byrow = TRUE)
x <- equilibrate(x)
apply(x, 2, function(x) sum(x^2)) # all 1
```

hadamard — *Hadamard product of two matrices*

Description

This function returns the Hadamard or element-wise product of two matrices x and y, that have the same dimensions.

Usage

```
hadamard(x, y = x)
```

Arguments

x	a numeric matrix or vector.
y	a numeric matrix or vector.

Value

A matrix with the same dimension of x (and y) which corresponds to the element-by-element product of the two matrices.

References

Styan, G.P.H. (1973). Hadamard products and multivariate statistical analysis, *Linear Algebra and Its Applications* **6**, 217-240.

Examples

```
x <- matrix(rep(1:10, times = 5), ncol = 5)
y <- matrix(rep(1:5, each = 10), ncol = 5)
z <- hadamard(x, y)
z
```

matrix.inner — *Compute the inner product between two rectangular matrices*

Description

Computes the inner product between two rectangular matrices calling BLAS.

Usage

```
matrix.inner(x, y = x)
```

Arguments

x	a numeric matrix.
y	a numeric matrix.

Value

a real value, indicating the inner product between two matrices.

Examples

```
x <- matrix(c(1, 1, 1,
              1, 2, 1,
              1, 3, 1,
              1, 1,-1,
              1, 2,-1,
              1, 3,-1), ncol = 3, byrow = TRUE)
y <- matrix(1, nrow = 6, ncol = 3)
matrix.inner(x, y)

# must be equal
matrix.norm(x, type = "Frobenius")^2
matrix.inner(x)
```

`matrix.norm` *Compute the norm of a rectangular matrix*

Description

Computes a matrix norm of `x` using LAPACK. The norm can be the one (`"1"`) norm, the infinity (`"inf"`) norm, the Frobenius norm, the maximum modulus (`"maximum"`) among elements of a matrix, as determined by the value of `type`.

Usage

```
matrix.norm(x, type = "Frobenius")
```

Arguments

`x`	a numeric matrix.
`type`	character string, specifying the *type* of matrix norm to be computed. A character indicating the type of norm desired. `"1"` specifies the **one** norm, (maximum absolute column sum); `"Inf"` specifies the **inf**inity norm (maximum absolute row sum); `"Frobenius"` specifies the **Frobenius** norm (the Euclidean norm of `x` treated as if it were a vector); `"maximum"` specifies the **maximum** modulus of all the elements in `x`.

Details

As function `norm` in package **base**, method of `matrix.norm` calls the LAPACK function `dlange`.

Note that the 1-, Inf- and maximum norm is faster to calculate than the Frobenius one.

Value

The matrix norm, a non-negative number.

Examples

```
# a tiny example
x <- matrix(c(1, 1, 1,
              1, 2, 1,
              1, 3, 1,
              1, 1,-1,
              1, 2,-1,
              1, 3,-1), ncol = 3, byrow = TRUE)
matrix.norm(x, type = "Frobenius")
matrix.norm(x, type = "1")
matrix.norm(x, type = "Inf")

# an example not that small
n <- 1000
x <- .5 * diag(n) + 0.5 * matrix(1, nrow = n, ncol = n)
matrix.norm(x, type = "Frobenius")
matrix.norm(x, type = "1")
matrix.norm(x, type = "Inf")
matrix.norm(x, type = "maximum") # equal to 1
```

`minkowski` *Computes the p-norm of a vector*

Description

Computes a p-norm of vector `x` using BLAS. The norm can be the one (`p = 1`) norm, Euclidean (`p = 2`) norm, the infinity (`p = Inf`) norm. For other values `p >= 1` the underlying Fortran code is based on ideas of BLAS Level 1.

Usage

```
minkowski(x, p = 2)
```

Arguments

`x`	a numeric vector.
`p`	a number, specifying the *type* of norm desired. Possible values include real number greater or equal to 1, or Inf, Default value is `p = 2`.

Details

Method of `minkowski` calls BLAS functions `dasum` (`p = 1`), `dnrm2` (`p = 2`), `idamax` (`p = Inf`). For other values, a Fortran subroutine using unrolled cycles is called.

Value

The vector p-norm, a non-negative number.

Examples

```
# a tiny example
x <- rnorm(1000)
minkowski(x, p = 1)
minkowski(x, p = 1.5)
minkowski(x, p = 2)
minkowski(x, p = Inf)

x <- x / minkowski(x)
minkowski(x, p = 2) # equal to 1
```

`power.method` *Power method to approximate dominant eigenvalue and eigenvector*

Description

The power method seeks to determine the eigenvalue of maximum modulus, and a corresponding eigenvector.

Usage

```
power.method(x, only.value = FALSE, maxiter = 100, tol = 1e-8)
```

Arguments

`x`	a symmetric matrix.
`only.value`	if `TRUE`, only the dominant eigenvalue is returned, otherwise both dominant eigenvalue and eigenvector are returned.
`maxiter`	the maximum number of iterations. Defaults to `100`
`tol`	a numeric tolerance.

Value

When `only.value` is not true, as by default, the result is a list with components `"value"` and `"vector"`. Otherwise only the dominan eigenvalue is returned. The performed number of iterations to reach convergence is returned as attribute `"iterations"`.

See Also

`eigen` for eigenvalues and eigenvectors computation.

Examples

```
n <- 1000
x <- .5 * diag(n) + 0.5 * matrix(1, nrow = n, ncol = n)

# dominant eigenvalue must be (n + 1) / 2
z <- power.method(x, only.value = TRUE)
```

`sherman.morrison` *Sherman-Morrison formula*

Description

The Sherman-Morrison formula gives a convenient expression for the inverse of the rank 1 update $(\mathbf{A} + \mathbf{b}\mathbf{d}^T)$ where $\mathbf{A}$ is a $n \times n$ matrix and $\mathbf{b}$, $\mathbf{d}$ are n-dimensional vectors. Thus

$$(\mathbf{A} + \mathbf{b}\mathbf{d}^T)^{-1} = \mathbf{A}^{-1} - \frac{\mathbf{A}^{-1}\mathbf{b}\mathbf{d}^T\mathbf{A}^{-1}}{1 + \mathbf{d}^T\mathbf{A}^{-1}\mathbf{b}}.$$

Usage

```
sherman.morrison(a, b, d = b, inverted = FALSE)
```

Arguments

`a`	a numeric matrix.
`b`	a numeric vector.
`d`	a numeric vector.
`inverted`	logical. If `TRUE`, `a` is supposed to contain its *inverse*.

Details

Method of `sherman.morrison` calls BLAS level 2 subroutines `dgemv` and `dger` for computational efficiency.

Value

a square matrix of the same order as `a`.

Examples

```
n <- 10
ones <- rep(1, n)
a <- 0.5 * diag(n)
z <- sherman.morrison(a, ones, 0.5 * ones)
z
```

`sweep.operator` *Gauss-Jordan sweep operator for symmetric matrices*

Description

Perform the sweep operation (or reverse sweep) on the diagonal elements of a symmetric matrix.

Usage

```
sweep.operator(x, k = 1, reverse = FALSE)
```

Arguments

`x`	a symmetric matrix.
`k`	elements (if `k` is vector) of the diagonal which will be sweeped.
`reverse`	logical. If `reverse = TRUE` the reverse sweep is performed.

Details

The symmetric sweep operator is a powerful tool in computational statistics with uses in stepwise regression, conditional multivariate normal distributions, MANOVA, and more.

Value

a square matrix of the same order as `x`.

References

Goodnight, J.H. (1979). A tutorial on the SWEEP operator. *The American Statistician* **33**, 149-158.

Examples

```
# tiny example of regression, last column contains 'y'
xy <- matrix(c(1, 1, 1, 1,
               1, 2, 1, 3,
               1, 3, 1, 3,
               1, 1,-1, 2,
               1, 2,-1, 2,
               1, 3,-1, 1), ncol = 4, byrow = TRUE)
z <- crossprod(xy)
z <- sweep.operator(z, k = 1:3)
cf <- z[1:3,4] # regression coefficients
RSS <- z[4,4]  # residual sum of squares

# an example not that small
x <- matrix(rnorm(1000 * 100), ncol = 100)
xx <- crossprod(x)
z <- sweep.operator(xx, k = 1)
```

`vec` *vectorization of a matrix*

Description

This function returns a vector obtained by stacking the columns of x

Usage

```
vec(x)
```

Arguments

`x`	a numeric matrix.

Value

Let `x` be a n by m matrix, then `vec(x)` is a nm-dimensional vector.

Examples

```
x <- matrix(rep(1:10, each = 10), ncol = 10)
x
y <- vec(x)
y
```

`vech` *vectorization the lower triangular part of a square matrix*

Description

This function returns a vector obtained by stacking the lower triangular part of a square matrix.

Usage

```
vech(x)
```

Arguments

`x`	a square matrix.

Value

Let `x` be a n by n matrix, then `vech(x)` is a $n(n+1)/2$-dimensional vector.

Examples

```
x <- matrix(rep(1:10, each = 10), ncol = 10)
x
y <- vech(x)
y
```

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