

Package ‘pbivnorm’

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Title Vectorized Bivariate Normal CDF

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Author Fortran code by Alan Genz. R code by Brenton Kenkel,
based on Adelchi Azzalini's 'mnormt' package.

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Description Provides a vectorized R function for calculating probabilities from a standard bivariate normal CDF.

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URL <https://github.com/brentonk/pbivnorm>

NeedsCompilation yes

Repository CRAN

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pbivnorm *Standard bivariate normal CDF*

Description

Calculate probabilities from the CDF of a standard bivariate normal distribution.

Usage

```
pbivnorm(x, y, rho = 0, recycle = TRUE)
```

Arguments

<code>x</code>	vector of upper integration limits for the CDF. May also be a two-column matrix, in which case <code>y</code> should not be used.
<code>y</code>	vector of upper integration limits.
<code>rho</code>	correlation parameter.
<code>recycle</code>	whether to automatically recycle the vectors <code>x</code> , <code>y</code> , and <code>rho</code> to conform to whichever is longest. If FALSE, all three must be the same length.

Details

This function returns values identical to those of `biv.nt.prob` in the **mnormt** package, but is vectorized to reduce the number of Fortran calls required for computation of many probabilities.

Value

Numeric vector of probabilities.

Author(s)

Fortran code by Alan Genz (see references). R interface by Brenton Kenkel (<brenton.kenkel@gmail.com>), based on code from Adelchi Azzalini's **mnormt** package.

References

- Genz, A. (1992). Numerical Computation of Multivariate Normal Probabilities. *J. Computational and Graphical Statist.*, **1**, 141–149.
- Genz, A. (1993). Comparison of methods for the computation of multivariate normal probabilities. *Computing Science and Statistics*, **25**, 400–405.
- Genz, A. Fortran code for MVTDSTPACK available at <http://www.math.wsu.edu/math/faculty/genz/software/fort77/mvtdstpack.f> (as of 2011-02-21).

Examples

```

x <- rnorm(10)
y <- rnorm(10)
rho <- runif(10)

pbivnorm(x, y, rho)

X <- cbind(x, y)
pbivnorm(X, rho = rho)

## rho can be a single value, unless recycling is disallowed
rho <- runif(1)
pbivnorm(x, y, rho)

```

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