

# Package ‘FAdist’

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 FAdist-package

*Distributions that are sometimes used in hydrology*


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### Description

This package contains several distributions that are sometimes useful in hydrology

### Author(s)

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 GAMMA3

*Three-Parameter Gamma Distribution (also known as Pearson type III distribution)*


---

### Description

Density, distribution function, quantile function and random generation for the 3-parameter gamma distribution with shape, scale, and threshold (or shift) parameters equal to shape, scale, and thres, respectively.

### Usage

```

dgamma3(x, shape=1, scale=1, thres=0, log=FALSE)
pgamma3(q, shape=1, scale=1, thres=0, lower.tail=TRUE, log.p=FALSE)
qgamma3(p, shape=1, scale=1, thres=0, lower.tail=TRUE, log.p=FALSE)
rgamma3(n, shape=1, scale=1, thres=0)

```

### Arguments

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations.
shape	shape parameter.
scale	scale parameter.
thres	threshold or shift parameter.
log, log.p	logical; if TRUE, probabilities p are given as log(p).
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

**Details**

If  $Y$  is a random variable distributed according to a gamma distribution (with shape and scale parameters), then  $X = Y+m$  has a 3-parameter gamma distribution with the same shape and scale parameters, and with threshold (or shift) parameter  $m$ .

**Value**

`dgamma3` gives the density, `pgamma3` gives the distribution function, `qgamma3` gives the quantile function, and `rgamma3` generates random deviates.

**References**

Bobee, B. and F. Ashkar (1991). The Gamma Family and Derived Distributions Applied in Hydrology. Water Resources Publications, Littleton, Colo., 217 p.

**See Also**

[dgamma](#), [pgamma](#), [qgamma](#), [rgamma](#)

**Examples**

```
thres <- 10
x <- rgamma3(n=10, shape=2, scale=11, thres=thres)
dgamma3(x, 2, 11, thres)
dgamma(x-thres, 2, 1/11)
```

---

 GenPARETO

*Generalized Pareto Distribution*


---

**Description**

Density, distribution function, quantile function and random generation for the generalized Pareto distribution with shape and scale parameters equal to shape and scale, respectively.

**Usage**

```
dgp(x, shape=1, scale=1, log=FALSE)
pgp(q, shape=1, scale=1, lower.tail=TRUE, log.p=FALSE)
qgp(p, shape=1, scale=1, lower.tail=TRUE, log.p=FALSE)
rgp(n, shape=1, scale=1)
```

**Arguments**

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations.
<code>shape</code>	shape parameter.

scale	scale parameter.
log, log.p	logical; if TRUE, probabilities p are given as log(p).
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

### Details

If  $X$  is a random variable distributed according to a generalized Pareto distribution, it has density  $f(x) = 1/\text{scale} * (1 - \text{shape} * x / \text{scale})^{(1 - \text{shape}) / \text{shape}}$

### Value

dgp gives the density, pgp gives the distribution function, qgp gives the quantile function, and rgp generates random deviates.

### References

Coles, S. (2001) An introduction to statistical modeling of extreme values. Springer

### Examples

```
x <- rgp(1000, -.2, 10)
hist(x, freq=FALSE, col='gray', border='white')
curve(dgp(x, -.2, 10), add=TRUE, col='red4', lwd=2)
```

---

GEV

*Generalized Extreme Value Distribution (for maxima)*

---

### Description

Density, distribution function, quantile function and random generation for the generalized extreme value distribution (for maxima) with shape, scale, and location parameters equal to shape, scale, and location, respectively.

### Usage

```
dgev(x, shape=1, scale=1, location=0, log=FALSE)
pgev(q, shape=1, scale=1, location=0, lower.tail=TRUE, log.p=FALSE)
qgev(p, shape=1, scale=1, location=0, lower.tail=TRUE, log.p=FALSE)
rgev(n, shape=1, scale=1, location=0)
```

### Arguments

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations.
shape	shape parameter.

scale	scale parameter.
location	location parameter.
log, log.p	logical; if TRUE, probabilities p are given as log(p).
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

### Details

If  $X$  is a random variable distributed according to a generalized extreme value distribution, it has density

$$f(x) = 1/\text{scale} * (1 + \text{shape} * ((x - \text{location})/\text{scale}))^{-1/\text{shape} - 1} * \exp(-(1 + \text{shape} * ((x - \text{location})/\text{scale}))^{-1/\text{shape}})$$

### Value

dgev gives the density, pgev gives the distribution function, qgev gives the quantile function, and rgev generates random deviates.

### References

Coles, S. (2001) An introduction to statistical modeling of extreme values. Springer

### Examples

```
x <- rgev(1000, -.1, 3, 100)
hist(x, freq=FALSE, col='gray', border='white')
curve(dgev(x, -.1, 3, 100), add=TRUE, col='red4', lwd=2)
```

---

GUMBEL

*Gumbel Distribution (for maxima)*

---

### Description

Density, distribution function, quantile function and random generation for the Gumbel distribution (for maxima) with scale and location parameters equal to scale and location, respectively.

### Usage

```
dgumbel(x, scale=1, location=0, log=FALSE)
pgumbel(q, scale=1, location=0, lower.tail=TRUE, log.p=FALSE)
qgumbel(p, scale=1, location=0, lower.tail=TRUE, log.p=FALSE)
rgumbel(n, scale=1, location=0)
```

**Arguments**

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations.
<code>scale</code>	scale parameter.
<code>location</code>	location parameter.
<code>log, log.p</code>	logical; if TRUE, probabilities <code>p</code> are given as $\log(p)$ .
<code>lower.tail</code>	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

**Details**

If  $X$  is a random variable distributed according to a Gumbel distribution, it has density  $f(x) = 1/\text{scale} * \exp(-(x-\text{location})/\text{scale} - \exp(-(x-\text{location})/\text{scale}))$

**Value**

`dgumbel` gives the density, `pgumbel` gives the distribution function, `qgumbel` gives the quantile function, and `rgumbel` generates random deviates.

**References**

Coles, S. (2001) An introduction to statistical modeling of extreme values. Springer

**Examples**

```
x <- rgumbel(1000,3,100)
hist(x,freq=FALSE,col='gray',border='white')
curve(dgumbel(x,3,100),add=TRUE,col='red4',lwd=2)
```

---

 KAPPA

*Kappa Distribution*


---

**Description**

Density, distribution function, quantile function and random generation for the kappa distribution with shape and scale parameters equal to shape and scale, respectively.

**Usage**

```
dkappa(x, shape=1, scale=1, log=FALSE)
pkappa(q, shape=1, scale=1, lower.tail=TRUE, log.p=FALSE)
qkappa(p, shape=1, scale=1, lower.tail=TRUE, log.p=FALSE)
rkappa(n, shape=1, scale=1)
```

**Arguments**

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations.
<code>shape</code>	shape parameter.
<code>scale</code>	scale parameter.
<code>log, log.p</code>	logical; if TRUE, probabilities <code>p</code> are given as $\log(p)$ .
<code>lower.tail</code>	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

**Details**

If  $X$  is a random variable distributed according to a kappa distribution, it has density  
 $f(x) = \text{shape}/\text{scale} * (\text{shape} + (x/\text{scale})^{\text{shape}})^{-((\text{shape}+1)/\text{shape})}$

**Value**

`dkappa` gives the density, `pkappa` gives the distribution function, `qkappa` gives the quantile function, and `rkappa` generates random deviates.

**Examples**

```
x <- rkappa(1000,12,10)
hist(x,freq=FALSE,col='gray',border='white')
curve(dkappa(x,12,10),add=TRUE,col='red4',lwd=2)
```

---

KAPPA4

---

*Four-Parameter Kappa Distribution*


---

**Description**

Density, distribution function, quantile function and random generation for the four-parameter kappa distribution with `shape1`, `shape2`, `scale`, and `location` parameters equal to `shape1`, `shape2`, `scale`, and `location`, respectively.

**Usage**

```
dkappa4(x, shape1, shape2, scale=1, location=0, log=FALSE)
pkappa4(q, shape1, shape2, scale=1, location=0, lower.tail=TRUE, log.p=FALSE)
qkappa4(p, shape1, shape2, scale=1, location=0, lower.tail=TRUE, log.p=FALSE)
rkappa4(n, shape1, shape2, scale=1, location=0)
```

**Arguments**

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations.
<code>shape1</code>	shape parameter.
<code>shape2</code>	shape parameter.
<code>scale</code>	scale parameter.
<code>location</code>	location parameter.
<code>log, log.p</code>	logical; if TRUE, probabilities <code>p</code> are given as $\log(p)$ .
<code>lower.tail</code>	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

**Details**

See References

**Value**

`dkappa4` gives the density, `pkappa4` gives the distribution function, `qkappa4` gives the quantile function, and `rkappa4` generates random deviates.

**References**

Hosking, J.R.M. (1994). The four-parameter kappa distribution. IBM Journal of Research and Development, 38(3), 251-258.

**Examples**

```
x <- rkappa4(1000, .1, .2, 12, 110)
hist(x, freq=FALSE, col='gray', border='white')
curve(dkappa4(x, .1, .2, 12, 110), add=TRUE, col='red4', lwd=2)
```

---

 LGAMMA3

*Log-Pearson Type III Distribution*


---

**Description**

Density, distribution function, quantile function and random generation for the log-Pearson type III distribution with `shape1`, `shape2`, and `scale` parameters equal to `shape`, `scale`, and `thres`, respectively.

**Usage**

```
dlgamma3(x, shape=1, scale=1, thres=1, log=FALSE)
plgamma3(q, shape=1, scale=1, thres=1, lower.tail=TRUE, log.p=FALSE)
qlgamma3(p, shape=1, scale=1, thres=1, lower.tail=TRUE, log.p=FALSE)
rlgamma3(n, shape=1, scale=1, thres=1)
```



**Arguments**

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations.
<code>shape</code>	shape1 parameter.
<code>scale</code>	shape2 parameter.
<code>thres</code>	scale parameter.
<code>log, log.p</code>	logical; if TRUE, probabilities <code>p</code> are given as $\log(p)$ .
<code>lower.tail</code>	logical; if TRUE (default), probabilities are $P[X \leq x]$ ; otherwise, $P[X > x]$ .

**Details**

If  $Y$  is a random variable distributed according to a gamma distribution (with shape and scale parameters), then  $X = \exp(Y+m)$  has a log-Pearson type III distribution with shape1 and shape2 parameters corresponding to the shape and  $1/\text{scale}$  parameters of  $Y$ , and with scale parameter  $m$ .

**Value**

`dlgamma3` gives the density, `plgamma3` gives the distribution function, `qlgamma3` gives the quantile function, and `rlgamma3` generates random deviates.

**References**

BOBEE, B. and F. ASHKAR (1991). The Gamma Family and Derived Distributions Applied in Hydrology. Water Resources Publications, Littleton, Colo., 217 p.

**See Also**

[dgamma](#), [pgamma](#), [qgamma](#), [rgamma](#), [dgamma3](#), [pgamma3](#), [qgamma3](#), [rgamma3](#)

**Examples**

```
thres <- 10
x <- rlgamma3(n=10, shape=2, scale=11, thres=thres)
dlgamma3(x, 2, 11, thres)
dgamma3(log(x), 2, 1/11, thres)/x
dgamma(log(x)-thres, 2, 11)/x
```

LLOGIS

*Log-Logistic Distribution***Description**

Density, distribution function, quantile function and random generation for the log-logistic distribution with shape and scale parameters equal to shape and scale, respectively.

**Usage**

```
dllog(x, shape=1, scale=1, log=FALSE)
pllog(q, shape=1, scale=1, lower.tail=TRUE, log.p=FALSE)
qllog(p, shape=1, scale=1, lower.tail=TRUE, log.p=FALSE)
rllog(n, shape=1, scale=1)
```

**Arguments**

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations.
shape	shape parameter.
scale	scale parameter.
log, log.p	logical; if TRUE, probabilities p are given as log(p).
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

**Details**

If  $Y$  is a random variable distributed according to a logistic distribution (with location and scale parameters), then  $X = \exp(Y)$  has a log-logistic distribution with shape and scale parameters corresponding to the scale and location parameters of  $Y$ , respectively.

**Value**

dllog gives the density, pllog gives the distribution function, qllog gives the quantile function, and rllog generates random deviates.

**See Also**

[dlogis](#), [plogis](#), [qlogis](#), [rlogis](#)

**Examples**

```
x <- rllog(10, 1, 0)
dllog(x, 1, 0)
dlogis(log(x), 0, 1)/x
```

LLOGIS3

*Three-Parameter Log-Logistic Distribution***Description**

Density, distribution function, quantile function and random generation for the 3-parameter log-logistic distribution with shape, scale, and threshold (or shift) parameters equal to shape, scale, and thres, respectively.

**Usage**

```
dllog3(x, shape=1, scale=1, thres=0, log=FALSE)
pllog3(q, shape=1, scale=1, thres=0, lower.tail=TRUE, log.p=FALSE)
qllog3(p, shape=1, scale=1, thres=0, lower.tail=TRUE, log.p=FALSE)
rllog3(n, shape=1, scale=1, thres=0)
```

**Arguments**

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations.
shape	shape parameter.
scale	scale parameter.
thres	threshold (or shift) parameter.
log, log.p	logical; if TRUE, probabilities p are given as log(p).
lower.tail	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

**Details**

If  $Y$  is a random variable distributed according to a logistic distribution (with location and scale parameters), then  $X = \exp(Y) + m$  has a 3-parameter log-logistic distribution with shape and scale parameters corresponding to the scale and location parameters of  $Y$ , respectively; and threshold parameter  $m$ .

**Value**

dllog3 gives the density, pllog3 gives the distribution function, qllog3 gives the quantile function, and rllog3 generates random deviates.

**See Also**

[dlogis](#), [plogis](#), [qlogis](#), [rlogis](#), [dllog](#), [pllog](#), [qllog](#), [rllog](#)

**Examples**

```

m <- 100
x <- rlllog3(10,1,0,m)
dllog3(x,1,0,m)
dllog(x-m,1,0)
dlogis(log(x-m),0,1)/(x-m)

```

LNORM3

*Three-Parameter Lognormal Distribution***Description**

Density, distribution function, quantile function and random generation for the 3-parameter lognormal distribution with shape, scale, and threshold (or shift) parameters equal to shape, scale, and thres, respectively.

**Usage**

```

dlnorm3(x, shape=1, scale=1, thres=0, log=FALSE)
plnorm3(q, shape=1, scale=1, thres=0, lower.tail=TRUE, log.p=FALSE)
qlnorm3(p, shape=1, scale=1, thres=0, lower.tail=TRUE, log.p=FALSE)
rlnorm3(n, shape=1, scale=1, thres=0)

```

**Arguments**

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations.
<code>shape</code>	shape parameter.
<code>scale</code>	scale parameter.
<code>thres</code>	threshold (or shift) parameter.
<code>log, log.p</code>	logical; if TRUE, probabilities <code>p</code> are given as $\log(p)$ .
<code>lower.tail</code>	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

**Details**

If  $Y$  is a random variable distributed according to a normal distribution (with location(mean) and scale(standard deviation) parameters), then  $X = \exp(Y)+m$  has a 3-parameter lognormal distribution with shape and scale parameters corresponding to the scale and location parameters of  $Y$ , respectively; and threshold parameter  $m$ .

**Value**

`dlnorm3` gives the density, `plnorm3` gives the distribution function, `qlnorm3` gives the quantile function, and `rlnorm3` generates random deviates.

**See Also**

[dnorm](#), [pnorm](#), [qnorm](#), [rnorm](#), [dlnorm](#), [plnorm](#), [qlnorm](#), [rlnorm](#)

**Examples**

```
m <- 100
x <- rlnorm3(10,1,0,m)
dlnorm3(x,1,0,m)
dlnorm(x-m,0,1)
dnorm(log(x-m),0,1)/(x-m)
```

WEIBULL3

*Three-Parameter Weibull Distribution***Description**

Density, distribution function, quantile function and random generation for the 3-parameter Weibull distribution with shape, scale, and threshold (or shift) parameters equal to shape, scale, and thres, respectively.

**Usage**

```
dweibull3(x,shape,scale=1,thres=0,log=FALSE)
pweibull3(q,shape,scale=1,thres=0,lower.tail=TRUE,log.p=FALSE)
qweibull3(p,shape,scale=1,thres=0,lower.tail=TRUE,log.p=FALSE)
rweibull3(n,shape,scale=1,thres=0)
```

**Arguments**

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations.
<code>shape</code>	shape parameter.
<code>scale</code>	scale parameter.
<code>thres</code>	threshold (or shift) parameter.
<code>log, log.p</code>	logical; if TRUE, probabilities p are given as log(p).
<code>lower.tail</code>	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .

**Details**

If  $Y$  is a random variable distributed according to a Weibull distribution (with shape and scale parameters), then  $X = Y+m$  has a 3-parameter Weibull distribution with shape and scale parameters corresponding to the shape and scale parameters of  $Y$ , respectively; and threshold parameter  $m$ .

**Value**

dweibull3 gives the density, pweibull3 gives the distribution function, qweibull3 gives the quantile function, and rweibull3 generates random deviates.

**See Also**

[dweibull](#), [pweibull](#), [qweibull](#), [rweibull](#)

**Examples**

```
m <- 100
x <- rweibull3(10,3,1,m)
dweibull3(x,3,1,m)
dweibull(x-m,3,1)
```

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