

# Package ‘twoexp’

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**Type** Package

**Title** The Two Parameter Exponential Distribution

**Version** 0.1.0

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**Description** Density, distribution function, quantile function, and random generation function, maximum likelihood estimation (MLE), penalized maximum likelihood estimation (PMLE), the quartiles method estimation (QM), and median rank estimation (MEDRANK) for the two-parameter exponential distribution. MLE and PMLE are based on Mengjie Zheng (2013)<[https://scse.d.umn.edu/sites/scse.d.umn.edu/files/mengjie-thesis\\_masters-1.pdf](https://scse.d.umn.edu/sites/scse.d.umn.edu/files/mengjie-thesis_masters-1.pdf)>. QM is based on Entisar Elgmati and Nadia Gregni (2016)<[doi:10.5539/ijsp.v5n5p12](https://doi.org/10.5539/ijsp.v5n5p12)>. MEDRANK is based on Matthew Reid (2022)<[doi:10.5281/ZENODO.3938000](https://doi.org/10.5281/ZENODO.3938000)>.

**License** GPL-3

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cdfplot	<i>Distribution function plot of the two-parameter exponential distribution</i>
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### Description

Distribution function plot of the two-parameter exponential distribution with theta and beta

### Usage

```
cdfplot(x, theta, beta)
```

### Arguments

x	vector of quantile.
theta	location parameter, where $\theta > 0$ .
beta	scale parameter, where $\beta > 0$ .

### Value

a distribution function plot of the two-parameter exponential distribution

### Examples

```
x <- seq(0, 20, by=0.01)
theta <- 6
beta <- 2
cdfplot(x, theta, beta)
```

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medrank	<i>Median rank method to estimate parameters of the two-parameter exponential dist.</i>
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### Description

Median rank method to estimate parameters of the two-parameter exponential dist.

### Usage

```
medrank(x, methods = c("B"))
```

### Arguments

x	vector of quantile (or a data set).
methods	there are some of median rank methods as follows; "B" stand for Benard median rank method (default), "BL" stand for Blom method, "MKM" stand for Hazen (Modified Kaplan Meier) method, "OT" stand for The one-third method, and "C" stand for Cunane method

### Value

the estimate three values for the two-parameter exponential dist. as follows: `theta.hat` gives the estimate location parameter, `beta.hat` gives the estimate scale parameter, and `lamda.hat` gives the estimate the rate.

### Source

Reid, M. (2022). Reliability – a Python library for reliability engineering (Version 0.8.2) [Computer software]. Zenodo. doi: [10.5281/ZENODO.3938000](https://doi.org/10.5281/ZENODO.3938000).

### Examples

```
x1 <- c(25,43,53,65,76,86,95,115,132,150) # test a data set
medrank(x1,"B") # Benard method (default) or medrank(x1)
```

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mle_tpexp	<i>Maximum likelihood estimation for the two-parameter exponential dist.</i>
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### Description

To estimate the location (or shift) and scale parameters for the two-parameter exponential distribution based on maximum likelihood method. See detail in source

**Usage**

```
mle_tpexp(x, theta = 0, beta = 1)
```

**Arguments**

x                    vector of quantile (or a data set).  
 theta                location parameter, where  $\theta > 0$ .  
 beta                 scale parameter, where  $\beta > 0$  and  $rate = 1/\beta$ .

**Value**

the estimate three values for the two-parameter exponential dist. as follows: theta.hat gives the estimate location parameter, beta.hat gives the estimate scale parameter, and lamda.hat gives the estimate the rate.

**Source**

Zheng, M. (2013). *Penalized Maximum Likelihood Estimation of Two-Parameter Exponential Distributions [Master's thesis]*. [https://scse.d.umn.edu/sites/scse.d.umn.edu/files/mengjie-thesis\\_masters-1.pdf](https://scse.d.umn.edu/sites/scse.d.umn.edu/files/mengjie-thesis_masters-1.pdf)

**Examples**

```
x1 <- c(25,43,53,65,76,86,95,115,132,150) # test a data set
mle_tpexp(x1)
x2 <- c(20,15,10,25,35,30,40,70,50,60,90,100,80,5) # test a data set
mle_tpexp(x2)
```

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 pdfplot

*Density plot of the two-parameter exponential distribution*


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**Description**

Density plot of the two-parameter exponential distribution with theta and beta

**Usage**

```
pdfplot(x, theta, beta)
```

**Arguments**

x                    vector of quantile.  
 theta                location parameter, where  $\theta > 0$ .  
 beta                 scale parameter, where  $\beta > 0$ .

**Value**

a density plot of the two-parameter exponential distribution

**Examples**

```
x <- seq(0,20,by=0.01)
theta <- 6
beta <- 2
pdfplot(x,theta,beta)
```

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pmle_tpexp	<i>Penalized maximum likelihood estimation for the two-parameter exponential dist.</i>
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**Description**

To estimate the location (or shift) and scale parameters for the two-parameter exponential distribution based on penalized maximum likelihood method. See detail in [source](#)

**Usage**

```
pmle_tpexp(x, theta = 0, beta = 1)
```

**Arguments**

x	vector of quantile (or a data set).
theta	location parameter, where $\theta > 0$ .
beta	scale parameter, where $\beta > 0$ and $rate = 1/\beta$ .

**Value**

the estimate three values for the two-parameter exponential dist. as follows: ptheta.hat gives the estimate location parameter, pbeta.hat gives the estimate scale parameter, and plambda.hat gives the estimate the rate.

**Source**

Zheng, M. (2013). *Penalized Maximum Likelihood Estimation of Two-Parameter Exponential Distributions [Master's thesis]*. [https://scse.d.umn.edu/sites/scse.d.umn.edu/files/mengjie-thesis\\_masters-1.pdf](https://scse.d.umn.edu/sites/scse.d.umn.edu/files/mengjie-thesis_masters-1.pdf)

**Examples**

```
x1 <- c(25,43,53,65,76,86,95,115,132,150) # test a data set
pmle_tpexp(x1)
x2 <- c(20,15,10,25,35,30,40,70,50,60,90,100,80,5) # test a data set
pmle_tpexp(x2)
```

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qm_tpepx	<i>Quartile method estimation of the two-parameter exponential distribution</i>
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### Description

To estimate the location (or shift) and scale parameters for the two-parameter exponential distribution based on quartile method. See detail in source

### Usage

```
qm_tpepx(x, methods = c("Q13"))
```

### Arguments

x	vector of quantile (or a data set).
methods	there are two quartile methods as follows; "Q13" stand for the first and the third quartile method (default), and "Q12" stand for the first and the second quartile (median) method.

### Value

the estimate three values for the two-parameter exponential dist. as follows: qmtheta.hat gives the estimate location parameter, qmbeta.hat gives the estimate scale parameter, and qmlambda.hat gives the estimate the rate.

### Source

Elgmati, E., Gregni, N. (2016). Quartile Method Estimation of Two-Parameter Exponential Distribution Data with Outliers. *International Journal of Statistics and Probability*, 5(5), 12-15. doi: [10.5539/ijsp.v5n5p12](https://doi.org/10.5539/ijsp.v5n5p12)

### Examples

```
x1 <- c(25,43,53,65,76,86,95,115,132,150) # test a data set
qm_tpepx(x1,"Q13") # or qm_tpepx(x1)
qm_tpepx(x1,"Q12")
```

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surplot	<i>Survival function plot of the two-parameter exponential distribution</i>
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**Description**

Survival function plot of the two-parameter exponential distribution with theta and beta

**Usage**

```
surplot(x, theta, beta)
```

**Arguments**

x	vector of quantile.
theta	location parameter, where $\theta > 0$ .
beta	scale parameter, where $\beta > 0$ .

**Value**

a survival function plot of the two-parameter exponential distribution

**Examples**

```
x <- seq(0,20,by=0.01)
theta <- 8
beta <- 1
surplot(x, theta, beta)
```

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tpexp	<i>The two-parameter exponential distribution(tpexp)</i>
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**Description**

Density, distribution function, quantile function, and random generation function for the two-parameter exponential distribution with theta and beta

**Usage**

```
dtpexp(x, theta = 0, beta = 1, log = FALSE)
ptpexp(q, theta = 0, beta = 1, lower.tail = TRUE, log.p = FALSE)
qtpexp(p, theta = 0, beta = 1, lower.tail = TRUE, log.p = FALSE)
rtpexp(n, theta = 0, beta = 1)
```

**Arguments**

<code>x, q</code>	vector of quantile.
<code>theta</code>	location parameter, where $\theta > 0$ .
<code>beta</code>	scale parameter, where $\beta > 0$ and $rate = 1/\beta$ .
<code>log, log.p</code>	logical; (default = FALSE), if TRUE, then probabilities are given as $\log(p)$ .
<code>lower.tail</code>	logical; if TRUE (default), probabilities are $P[X \leq x]$ , otherwise, $P[X > x]$ .
<code>p</code>	vector of probabilities
<code>n</code>	number of observations. If $\text{length}(n) > 1$ , the length is taken to be the number required.

**Value**

`dtpexp` gives the density, `ptpexp` gives the distribution function, `qtpexp` gives the quantile function, and `rtpexp` generates random samples.

**Examples**

```
x <- c(0.1,0.2,0.3,0.4,0.5,0.6,0.7,0.8,0.9,1.0)
dtpexp(x,theta=0,beta=1)
dtpexp(x,theta=0,beta=1,log=TRUE)

q <- c(0.1,0.2,0.3,0.4,0.5,0.6,0.7,0.8,0.9,1.0)
ptpexp(q,theta = 0, beta = 1)
ptpexp(q,theta=0, beta = 1, lower.tail = FALSE)

q <- c(0.1,0.2,0.3,0.4,0.5,0.6,0.7,0.8,0.9,1.0)
p<- ptpexp(q,theta = 0, beta = 1); p
qtpexp(p,theta=0, beta = 1)

rtpexp(5, theta=0, beta=1)
rtpexp(10, theta=1, beta=1.5)
```



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