

Package ‘twopDist’

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Description This package fits some asymmetric models and asymmetric distributions by maximum likelihood estimation (MLE) or penalized MLE.

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TPLapD
Two-piece Laplace Distribution

Description

Density, distribution function, quantile function and random generation for the 3-parameter two-piece Laplace distribution (TPLapD). The 2-parameter TPLapD may be used for classical quantile regression.

Usage

```
dtwoplalplace(x, location = 0, scale = 1, skewpar = 0.5, log.arg = FALSE)
ptwoplalplace(q, location = 0, scale = 1, skewpar = 0.5)
qtwoplalplace(p, location = 0, scale = 1, skewpar = 0.5)
rtwoplalplace(n, location = 0, scale = 1, skewpar = 0.5)
```

Arguments

<code>x, q</code>	vector of quantiles.
<code>p</code>	vector of probabilities.
<code>n</code>	number of observations. If <code>length(n) > 1</code> then the length is taken to be the number required.
<code>location</code>	location parameter.
<code>scale</code>	positive scale parameter.
<code>skewpar</code>	skew/asymmetric parameter, between zero and one.
<code>log.arg</code>	Logical. If <code>log.arg = TRUE</code> then the logarithm of the density is returned.

Value

`dtwoplalplace` gives the density, `ptwoplalplace` gives the distribution function, `qtwoplalplace` gives the quantile function, and `rtwoplalplace` generates random deviates.

Author(s)

Arash Ardalan

Examples

```
location = 0; scale = 1.5; skewpar = 0.25; nn = 201
x = seq(-3.5, 6.5, len = nn)
y = dtwoplalplace(x, location, scale, skewpar)

## Not run:
plot(x, y, type = "l", las = 1, ylim = c(0, 1.2), ylab =
  paste("ftwoplalplace(location = ", location, ", scale = ", scale,
    ", skewpar = ", skewpar, ")"),
  col = "blue", cex.main = 0.8,
  main = "Blue is density, red is cumulative distribution function",
  sub = "Purple lines are the 10, 20, ..., 90 percentiles")
lines(x, ptwoplalplace(x, location, scale, skewpar), col = "red")
probs = seq(0.1, 0.9, by = 0.1)
```

```

Q = qtwoplalplace(probs, location, scale, skewpar)
lines(Q, dtwoplalplace(Q, location, scale, skewpar), col = "purple",
      lty = 3, type = "h")
lines(Q, ptwoplalplace(Q, location, scale, skewpar), col = "purple",
      lty = 3, type = "h")
abline(h = probs, col = "purple", lty = 3)
abline(h = 0, col = "gray")
max(abs(ptwoplalplace(Q, location, scale, skewpar) - probs)) # Should be 0

## End(Not run)

```

TPLogisD

Two-piece Logistic Distribution

Description

Density, distribution function, quantile function and random generation for the 3-parameter two-piece logistic distribution (TPLogisD). The 2-parameter TPLogisD may be used for quantile/expectile type regression. This family also is more efficient and more robust in comparison with classical quantile regression.

Usage

```

dtwoplogis(x, location = 0, scale = 1, skewpar = 0.5, log.arg = FALSE)
ptwoplogis(q, location = 0, scale = 1, skewpar = 0.5)
qtwoplogis(p, location = 0, scale = 1, skewpar = 0.5)
rtwoplogis(n, location = 0, scale = 1, skewpar = 0.5)

```

Arguments

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations. If <code>length(n) > 1</code> then the length is taken to be the number required.
location	location parameter.
scale	positive scale parameter.
skewpar	skew/asymmetric parameter, between zero and one.
log.arg	Logical. If <code>log.arg = TRUE</code> then the logarithm of the density is returned.

Details

See [twoplogis3](#) and [twoplogis](#), the **twopDist** family function for estimating the parameters, for the formula of the probability density function and other details.

Value

`dtwoplogis` gives the density, `ptwoplogis` gives the distribution function, `qtwoplogis` gives the quantile function, and `rtwoplogis` generates random deviates.

Author(s)

Arash Ardalan

See Also[twoplogis3](#).**Examples**

```

location = 1; scale = 1.2; skewpar = 0.25; nn = 201
# location = 1, scale = 1.2;
x = seq(-1.5, 6.5, len=nn)
y = dtwoplogis(x, location, scale, skewpar)

## Not run:
plot(x, y, type="l", las=1, ylim = c(0, 1.2), ylab =
  paste("ftwoplogis(location=", location, ", scale=", scale,
        ", skewpar=", skewpar, ")"),
  col="blue", cex.main = 0.8,
  main="Blue is density, red is cumulative distribution function",
  sub="Purple lines are the 10, 20, ..., 90 percentiles")
lines(x, ptwoplogis(x, location, scale, skewpar), col="red")
probs = seq(0.1, 0.9, by = 0.1)
Q = qtwoplogis(probs, location, scale, skewpar)
lines(Q, dtwoplogis(Q, location, scale, skewpar), col = "purple",
      lty = 3, type = "h")
lines(Q, ptwoplogis(Q, location, scale, skewpar), col = "purple",
      lty = 3, type = "h")
abline(h = probs, col = "purple", lty = 3)
abline(h = 0,col= "gray")
max(abs(ptwoplogis(Q, location, scale, skewpar) - probs)) # Should be 0

## End(Not run)

```

 TPNormD

Two-piece Normal Distribution

Description

Density, distribution function, quantile function and random generation for the 2 and 3-parameter two-piece normal distribution (TPND). The 2-parameter TPND may be used for quantile/expectile regression.

Usage

```

dtwopnorm(x, location = 0, scale = 1, skewpar = 0.5, log.arg = FALSE)
ptwopnorm(q, location = 0, scale = 1, skewpar = 0.5)
qtwopnorm(p, location = 0, scale = 1, skewpar = 0.5)
rtwopnorm(n, location = 0, scale = 1, skewpar = 0.5)

```

Arguments

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations. If <code>length(n) > 1</code> then the length is taken to be the number required.
location	location parameter.
scale	positive scale parameter.
skewpar	skew/asymmetric parameter, between zero and one.
log.arg	logical. If <code>log.arg = TRUE</code> then the logarithm of the density is returned.

Details

See [twopnorm3](#) and [twopnorm](#) the **twopDist** family function for estimating the parameters, for the formula of the probability density function and other details.

Value

`dtwopnorm` gives the density, `ptwopnorm` gives the distribution function, `qtwopnorm` gives the quantile function, and `rtwopnorm` generates random deviates.

Author(s)

Arash Ardalan

See Also

[twopnorm3](#)

Examples

```
location = 1; scale = 1.2; skewpar = 0.25; nn = 201
x = seq(-1.5 , 6.5, len = nn)
y = dtwopnorm(x, location, scale, skewpar)

## Not run:
plot(x, y, type = "l", las = 1, ylim = c(0, 1.2), ylab =
  paste("twopnorm(location = ", location, ", scale = ", scale,
        ", skewpar = ", skewpar, ")"),
  col = "blue", cex.main = 0.8,
  main = "Blue is density, red is cumulative distribution function",
  sub = "Purple lines are the 10, 20, ..., 90 percentiles")
lines(x, ptwopnorm(x, location, scale, skewpar), col = "red")
probs = seq(0.1, 0.9, by = 0.1)
Q = qtwpnorm(probs, location, scale, skewpar)
lines(Q, dtwopnorm(Q, location, scale, skewpar), col = "purple",
  lty = 3, type = "h")
lines(Q, ptwopnorm(Q, location, scale, skewpar), col = "purple",
  lty = 3, type = "h")
abline(h = probs, col = "purple", lty = 3)
abline(h = 0, col = "gray")
max(abs(ptwopnorm(Q, location, scale, skewpar) - probs)) # Should be 0

## End(Not run)
```

 TPNormLapD

Two-piece Laplace Distribution

Description

Density, distribution function, quantile function and random generation for the 3-parameter two-piece normal Laplace distribution (TPNormLapD). This distribution from one side is short tail(normal) and in the other side(Laplace) is long tail, see Ardalan et. All (2012). The 2-parameter TPNormLapD may be used for quantile type regression model.

Usage

```
dtwopnormlap(x, location = 0, scale = 1, skewpar = 0.5, log.arg = FALSE)
ptwopnormlap(q, location = 0, scale = 1, skewpar = 0.5)
qtwopnormlap(p, location = 0, scale = 1, skewpar = 0.5)
rtwopnormlap(n, location = 0, scale = 1, skewpar = 0.5)
```

Arguments

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations. If <code>length(n) > 1</code> then the length is taken to be the number required.
location	location parameter.
scale	positive scale parameter.
skewpar	skew/asymmetric parameter, between zero and one.
log.arg	Logical. If <code>log.arg = TRUE</code> then the logarithm of the density is returned.

Value

`dtwopnormlap` gives the density, `ptwopnormlap` gives the distribution function, `qtwopnormlap` gives the quantile function, and `rtwopnormlap` generates random deviates.

Author(s)

Arash Ardalan

References

Arash Ardalan, S. M. Sadooghi-Alvandi and A.N. Nematollahi (2012), The Two-Piece Normal-Laplace Distribution, 35 pages (Comm. in Statist.: Theory & Methods. Accepted).

Examples

```
location = 0; scale = 1.5; skewpar = 0.5; nn = 201
x = seq(-3.5, 6.5, len = nn)
y = dtwopnormlap(x, location, scale, skewpar)
```

```
## Not run:
```

```

plot(x, y, type = "l", las = 1, ylim = c(0, 1.2), ylab =
  paste("ftwopnormlap(location=", location, ", scale=", scale,
    ", skewpar=", skewpar, ")"),
  col = "blue", cex.main = 0.8,
  main = "Blue is density, red is cumulative distribution function",
  sub = "Purple lines are the 10, 20, ..., 90 percentiles")
lines(x, ptwopnormlap(x, location, scale, skewpar), col = "red")
probs = seq(0.1, 0.9, by = 0.1)
Q = qtwopnormlap(probs, location, scale, skewpar)
lines(Q, dtwopnormlap(Q, location, scale, skewpar), col = "purple",
  lty = 3, type = "h")
lines(Q, ptwopnormlap(Q, location, scale, skewpar), col = "purple",
  lty = 3, type = "h")
abline(h = probs, col = "purple", lty = 3)
abline(h = 0, col = "gray")
max(abs(ptwopnormlap(Q, location, scale, skewpar) - probs)) # Should be 0

## End(Not run)

```

TPTD

*Two-piece Student's t-distribution***Description**

Density, distribution function, quantile function and random generation for the 5-parameter two-piece Student's t-distribution (TPTD). The 2-parameter TPTD (with given skewpar, dfL and dfR) may be used for quantile type regression.

Usage

```

dtwopt(x, location = 0, scale = 1, skewpar = 0.5, dfL = 2, dfR = 2,
  log.arg = FALSE)
ptwopt(q, location = 0, scale = 1, skewpar = 0.5, dfL = 2, dfR = 2)
qtwopt(p, location = 0, scale = 1, skewpar = 0.5, dfL = 2, dfR = 2)
rtwopt(n, location = 0, scale = 1, skewpar = 0.5, dfL = 2, dfR = 2)

```

Arguments

x, q	vector of quantiles.
p	vector of probabilities.
n	number of observations. If <code>length(n) > 1</code> then the length is taken to be the number required.
location	location parameter.
scale	positive scale parameter.
skewpar	skew/asymmetric parameter, between zero and one.
log.arg	logical. If <code>log.arg = TRUE</code> then the logarithm of the density is returned.
dfL	the left degree of freedom.
dfR	the right degree of freedom.

Value

dtwopt gives the density, ptwopt gives the distribution function, qtwopt gives the quantile function, and rtwopt generates random deviates.

Author(s)

Arash Ardalan

Examples

```
location = 0; scale = 1.5; skewpar = 0.25; nn = 201
x = seq(-3.5 ,6.5, len = nn)
y = dtwopt(x, location, scale, skewpar)

## Not run:
plot(x, y, type = "l", las = 1, ylim = c(0, 1.2), ylab =
  paste("ftwopt(location = ", location, ", scale = ", scale,
        ", dfL = dfR = ", 2, ", skewpar = ", skewpar, ")"),
  col = "blue", cex.main = 0.8,
  main = "Blue is density, red is cumulative distribution function",
  sub = "Purple lines are the 10, 20, ..., 90 percentiles")
lines(x, ptwopt(x, location, scale, skewpar), col = "red")

probs = seq(0.1, 0.9, by = 0.1)
Q = qtwopt(probs, location, scale, skewpar)
lines(Q, dtwopt(Q, location, scale, skewpar), col = "purple",
  lty = 3, type = "h")
lines(Q, ptwopt(Q, location, scale, skewpar), col = "purple",
  lty = 3, type = "h")
abline(h = probs, col = "purple", lty = 3)
abline(h = 0, col = "gray")
max(abs(ptwopt(Q, location, scale, skewpar) - probs)) # Should be 0

## End(Not run)
```

twoplogistic

Two-piece Logistic Distribution Family Function

Description

Maximum likelihood estimation of the 1, 2 and 3-parameter two-piece logistic distributions (TPLogisD). The 1-parameter TPLogisD may be used for quantile type regression.

Usage

```
twoplogis(skewpar = 0.5, llocation = "identity", lscale = "loge",
  elocation = list(), escale = list(), iscale = NULL,
  parallelLocation = FALSE, digt = 2, sameScale = TRUE,
  intparloc = FALSE, imethod = 1, zero = -2)

twoplogis3(llocation = "identity", elocation = list(),
```

```
lscale      = "identity",  escale = list(),
lskewpar    = "identity",  eskewpar = list(),
imethod = 1, zero = 2:3)
```

Arguments

<code>skewpar</code>	Numeric vectors with $0 < skewpar < 1$. Most users will only specify <code>skewpar</code> since the estimated location parameter corresponds to the p th regression quantile/expectile, which is easier to understand. See below for details.
<code>llocation, lscale, lskewpar</code>	Character. Parameter link functions for location parameter μ , scale parameter σ , asymmetry parameter <code>skewpar</code> .
<code>elocation, escale, eskewpar</code>	List. Extra argument for each of the links. See <code>earg</code> in Links for general information.
<code>iscale</code>	Optional initial values. If given, it must be numeric and values are recycled to the appropriate length. The default is to choose the value internally.
<code>parallelLocation, intparloc</code>	Logical. Should the quantiles be parallel on the transformed scale (argument <code>llocation</code>)? Assigning this argument to <code>TRUE</code> circumvents the seriously embarrassing quantile crossing problem. The argument <code>intparloc</code> applies to intercept term; the argument <code>parallelLocation</code> applies to other terms.
<code>digit</code>	Passed into Round as the <code>digits</code> argument for the <code>tau</code> values; used cosmetically for labelling.
<code>sameScale</code>	Logical. Should the scale parameters be equal? It is advised to keep <code>sameScale=TRUE</code> unchanged because it does not make sense to have different values for each <code>tau</code> value.
<code>imethod</code>	Initialization method. Either the value 1, 2, 3 or 4.
<code>zero</code>	See CommonVGAMffArguments for more information. Where possible, the default is to model all the σ and <code>skewpar</code> as an intercept-only term.

Details

These **twopDist** family functions implement one variant of two-piece logistic distribution (TPLogisD) suitable for quantile/expectile regression. Its density function is

$$f(y; \mu, \sigma, p) = \frac{1}{4\sigma} \begin{cases} \operatorname{sech}^2\left(\frac{(y-\mu)}{4\sigma p}\right), & y < \mu, \\ \operatorname{sech}^2\left(\frac{(y-\mu)}{4\sigma(1-p)}\right), & y \geq \mu. \end{cases}$$

Here, the ranges are for all real y and μ , positive σ and positive p .

The special case $p = 0.5$ corresponds to the (symmetric) logistic distribution.

The enumeration of the linear/additive predictors used here is to first have all the location parameters, followed by all the scale parameters. Finally, for `twoplogis3()`, the last one is the asymmetry parameter.

It is known that the maximum likelihood estimate of the location parameter μ corresponds to the regression quantile/expectile estimate of the classical quantile regression approach. An important property of the this family is that $P(Y \leq \mu) = p$.

Thus `twoplogis()` may be used as an alternative to `rq` in the **quantreg** package.

In general, the response must be a vector or a 1-column matrix.

Value

An object of class "vglmff" (see [vglmff-class](#)). The object is used by modelling functions such as [vglm](#) and [vgam](#)

In the `extra` slot of the fitted object are some list components which are useful, e.g., the sample proportion of values which are less than the fitted quantile curves.

Note

The **VGAM** family functions use Fisher scoring. Convergence may be slow and half-stepping is usual (although one can use `trace = TRUE` to see which is the best model and then set `maxit` to fit that model).

Author(s)

Arash Ardalan

References

Ardalan, A. and Yee, T. W. (2012), Expectile/quantile regression via two-piece distributions. In preparation.

See Also

[twopnormlms.bcn](#).

Examples

```
# Example 1: Logistic quantile regression
# (1) Preprocessing the data

website = "http://www.stat.auckland.ac.nz/~lee/330/datasets.dir/chd.txt"
chd.df = read.table(website, header = TRUE)

myskew = (1:9) / 10
fit0 <- vglm(chd ~ age,
             fam = twoplogis(llocation = "logit", skewpar = myskew,
                             imethod = 1, parallelLoc = TRUE),
             data = chd.df, trace = TRUE)

fit1 <- vglm(chd ~ bs(age, df = 3),
             fam = twoplogis(llocation = "logit", skewpar = myskew,
                             imethod = 1, parallelLoc = TRUE),
             data = chd.df, trace = TRUE)

# Plot the results
## Not run:
plotObject1 = fit1
plotObject0 = fit0

plot(jitter(chd, 0.2) ~ age, data = chd.df, col = "blue", las = 1)
for (ii in 1:plotObject0@extra$NOS) {
  ooo = with(chd.df, order(age))
  lines(fitted(plotObject0)[ooo, ii] ~ age[ooo], data = chd.df,
```

```

        col = "orange", lwd = 2)
    }

## End(Not run)
# Example 2: quantile regression with B-splines

# (1) Preprocessing the data and sort by wave heights:
require("ismev")
data(wavesurge, package = "ismev")

oo = with(wavesurge, order(wave))

wavesurge = wavesurge[oo, ]

# (2) Scatter plot

## Not run:
plot(surge ~ wave, main = "10:10:90 percentile curves",
     col = "blue", las = 1, pch = 21, cex = 0.8,
     bg = "black", data = wavesurge)

## End(Not run)

# (3) A preliminary fit

myskew = seq(0.10, 0.90, by = 0.1) # A vector!!
fit = vglm(surge ~ bs(wave, degree = 2), data = wavesurge,
          trace = TRUE, fam = twoplogis(myskew))

fit@extra # Gives some info
coef(fit, matrix = TRUE)

# (4) The regression curves
## Not run:
with(wavesurge, matplot(wave, fitted(fit), type = "l",
                       lty = 1:ncol(fitted(fit)), lwd = 3,
                       add = TRUE, col = 1:ncol(fitted(fit))))

text(11, tail(fitted(fit), 1), paste(myskew * 100, "%"))

## End(Not run)

```

twopnorm

Two-piece Normal Distribution Family Function

Description

Maximum likelihood estimation of the 2 and 3-parameter two-piece normal distributions (TPND). The 1-parameter TPND may be used for quantile/expectile regression.

Usage

```
twopnorm(skewpar = 0.5, llocation = "identity", lscale = "loge",
         elocation = list(), escale = list(), iscale = NULL,
         parallelLocation = FALSE, digt = 2, sameScale = TRUE,
         intparloc = FALSE, imethod = 1, zero = -2)
```

```
twopnorm3(llocation = "identity", elocation = list(),
          lscale = "identity", escale = list(),
          lskewpar = "identity", eskewpar = list(),
          imethod = 1, zero = 2:3)
```

Arguments

- skewpar** Numeric vectors with $0 < skewpar < 1$. Most users will only specify *skewpar* since the estimated location parameter corresponds to the *p*th regression quantile/expectile, which is easier to understand. See below for details.
- llocation, lscale, lskewpar** Character. Parameter link functions for location parameter μ , scale parameter σ , asymmetry parameter *skewpar*.
- elocation, escale, eskewpar** List. Extra argument for each of the links. See **earg** in [Links](#) for general information.
- iscale** Optional initial values. If given, it must be numeric and values are recycled to the appropriate length. The default is to choose the value internally.
- parallelLocation, intparloc** Logical. Should the quantiles be parallel on the transformed scale (argument **llocation**)? Assigning this argument to **TRUE** circumvents the seriously embarrassing quantile crossing problem. The argument **intparloc** applies to intercept term; the argument **parallelLocation** applies to other terms.
- digt** Passed into [Round](#) as the **digits** argument for the τ values; used cosmetically for labelling.
- sameScale** Logical. Should the scale parameters be equal? It is advised to keep **sameScale=TRUE** unchanged because it does not make sense to have different values for each τ value.
- imethod** Initialization method. Either the value 1, 2, 3 or 4.
- zero** See [CommonVGAMffArguments](#) for more information. Where possible, the default is to model all the σ and *skewpar* as an intercept-only term.

Details

These **twopDist** family functions implement one variant of two-piece normal distribution (TPND) suitable for quantile/expectile regression. Its density function is

$$f(y; \mu, \sigma, p) = \frac{1}{\sigma\sqrt{2\pi}} \begin{cases} \exp\left(-\frac{(y-\mu)^2}{8\sigma^2 p^2}\right), & y < \mu, \\ \exp\left(-\frac{(y-\mu)^2}{8\sigma^2 (1-p)^2}\right), & y \geq \mu. \end{cases}$$

Here, the ranges are for all real y and μ , positive σ and positive p .

The special case $p = 0.5$ corresponds to the (symmetric) normal distribution.

The enumeration of the linear/additive predictors used here is to first have all the location parameters, followed by all the scale parameters. Finally, for `twopnorm3()`, the last one is the asymmetry parameter.

It is known that the maximum likelihood estimate of the location parameter μ corresponds to the regression quantile/ expectile estimate of the classical quantile regression approach. An important property of the TPND is that $P(Y \leq \mu) = p$.

Thus `twopnorm()` may be used as an alternative to `rq` in the **quantreg** package.

In general, the response must be a vector or a 1-column matrix.

Value

An object of class "vglmff" (see [vglmff-class](#)). The object is used by modelling functions such as [vglm](#) and [vgam](#)

In the `extra` slot of the fitted object are some list components which are useful, e.g., the sample proportion of values which are less than the fitted quantile curves.

Note

The **VGAM** family functions use Fisher scoring. Convergence may be slow and half-stepping is usual (although one can use `trace = TRUE` to see which is the best model and then use `maxit` to fit that model).

Author(s)

Arash Ardalan

References

Ardalan, A. and Yee, T. W. (2012), Expectile/quantile regression via two-piece distribution. In preparation.

See Also

[dtwopnormlms.bcn.](#)

Examples

```
# Example 1: quantile regression with smoothing splines

# (1) Preprocessing the data
data(mmt, package = "VGAM")
melb = data.frame(today = tail(mmt, -1),
                  yesterday = head(mmt, -1))

# Sort by yesterday's temperature:
ooo = with(melb, order(yesterday))
melb = melb[ooo,]
head(melb)

# (2) Scatter plot
## Not run:
plot(today ~ yesterday, data = melb, las = 1,
```

```

      xlab = "Yesterday's Max Temperature",
      ylab = "Today's Max Temperature",
      main = "Melbourne Temperature Data",
      pch = 0, cex = 0.25, col = "blue")
abline(a = 0, b = 1, lty = "dashed")

## End(Not run)
# (3) A preliminary fit
myskew = seq(0.10, 0.90, by = 0.1) # A vector!!
fit = vgam(today ~ s(yesterday, df = 4), data = melb,
           trace = TRUE,
           fam = twopnorm(myskew))

fit@extra # Gives some info
coef(fit, matrix = TRUE)

# (4) The regression curves
## Not run:
with(melb, matplot(yesterday, fitted(fit), type = "l",
                  lty = 1:ncol(fitted(fit)),
                  lwd = 3,
                  add = TRUE, col = 1:ncol(fitted(fit))))

text(42, tail(fit
ted(fit), 1), paste(myskew * 100, "%"))

## End(Not run)

```

twopnormlap

Two-piece Normal Laplace Distribution Family Function

Description

Maximum likelihood estimation of the 1 and 2-parameter two-piece normal Laplace distributions (TPNLD). The 1-parameter TPND may be used for quantile/expectile regression.

Usage

```

twopnormlap(skewpar = 0.5, llocation = "identity", lscale = "loge",
            elocation = list(), escale = list(), iscale = NULL,
            parallelLocation = FALSE, digit = 2, eq.scale = TRUE,
            intparloc = FALSE, imethod = 1, zero = -2)

```

Arguments

skewpar	Numeric vectors with $0 < skewpar < 1$. Most users will only specify <i>skewpar</i> since the estimated location parameter corresponds to the <i>p</i> th regression quantile/expectile, which is easier to understand. See below for details.
llocation, lscale	Character. Parameter link functions for location parameter μ , scale parameter σ .
elocation, escale	List. Extra argument for each of the links. See <code>earg</code> in Links for general information.

<code>iscale</code>	Optional initial values. If given, it must be numeric and values are recycled to the appropriate length. The default is to choose the value internally.
<code>parallelLocation</code> , <code>intparloc</code>	Logical. Should the quantiles be parallel on the transformed scale (argument <code>llocation</code>)? Assigning this argument to <code>TRUE</code> circumvents the seriously embarrassing quantile crossing problem. The argument <code>intparloc</code> applies to intercept term; the argument <code>parallelLocation</code> applies to other terms.
<code>dig</code>	Passed into <code>Round</code> as the <code>digits</code> argument for the <code>tau</code> values; used cosmetically for labelling.
<code>eq.scale</code>	Logical. Should the scale parameters be equal? It is advised to keep <code>eq.scale=TRUE</code> unchanged because it does not make sense to have different values for each <code>skewpar</code> value.
<code>imethod</code>	Initialization method. Either the value 1, 2, 3 or 4.
<code>zero</code>	See Common VGAM Arguments for more information. Where possible, the default is to model all the σ and <code>skewpar</code> as an intercept-only term.

Details

These **twopDist** family functions implement one variant of two-piece normal distribution (TPNDL) suitable for quantile/expectile regression. Its density function is

$$f(y; \mu, \sigma, p) = \frac{1}{\sigma\sqrt{2\pi}} \begin{cases} \exp\left(-\frac{(y-\mu)^2}{8\sigma^2 p^2}\right) & y < \mu \\ \exp\left(-\frac{(y-\mu)^2}{8\sigma(1-p)\sqrt{2\pi}}\right) & y \geq \mu. \end{cases}$$

Here, the ranges are for all real y and μ , positive σ and positive p .

The enumeration of the linear/additive predictors used here is to first have all the location parameters, followed by all the scale parameters.

It is known that the maximum likelihood estimate of the location parameter μ corresponds to the regression quantile/ expectile estimate of the classical quantile regression approach. An important property of the TPNDL is that $P(Y \leq \mu) = p$.

Thus `twopnormlap()` may be used as an alternative to `rq` in the **quantreg** package.

In general, the response must be a vector or a 1-column matrix.

Value

An object of class "vglmff" (see [vglmff-class](#)). The object is used by modelling functions such as `vglm` and `vgam`.

In the `extra` slot of the fitted object are some list components which are useful, e.g., the sample proportion of values which are less than the fitted quantile curves.

Note

The **VGAM** family functions use Fisher scoring. Convergence may be slow and half-stepping is usual (although one can use `trace = TRUE` to see which is the best model and then use `maxit` to fit that model).

Author(s)

Arash Ardalan

References

Arash Ardalan, S. M. Sadooghi-Alvandi and A.N. Nematollahi (2012), The Two-Piece Normal-Laplace Distribution, 35 pages (Comm. in Statist.: Theory & Methods. Accepted).

See Also

[dtwopnormlap lms.bcn.](#)

Examples

```
# Example 1: quantile regression with smoothing splines
```

```
# (1) Preprocessing the data
data(mmt, package = "VGAM")
melb = data.frame(today      = tail(mmt, -1),
                  yesterday = head(mmt, -1))
```

```
# Sort by yesterday's temperature:
ooo = with(melb, order(yesterday))
melb = melb[ooo,]
head(melb)
```

```
# (2) Scatter plot
## Not run:
plot(today ~ yesterday, data = melb, las = 1,
      xlab = "Yesterday's Max Temperature",
      ylab = "Today's Max Temperature",
      main = "Melbourne Temperature Data",
      pch = 0, cex = 0.25, col = "blue", xlim = c(0, 50))
abline(a = 0, b = 1, lty = "dashed")
```

```
## End(Not run)
# (3) A preliminary fit
myskew = seq(0.10, 0.90, by = 0.1) # A vector!!
fit = vgam(today ~ s(yesterday, df = 4), data = melb,
           trace = TRUE,
           fam = twopnormlap(myskew))
```

```
fit@extra # Gives some info
coef(fit, matrix = TRUE)
```

```
# (4) The regression curves
## Not run:
with(melb, matplot(yesterday, fitted(fit), type = "l",
                  lty = 1:ncol(fitted(fit)),
                  lwd = 3,
                  add = TRUE, col = 1:ncol(fitted(fit))))
```

```
text(48, tail(fitted(fit), 1), paste(myskew * 100, "%"))
```

```
## End(Not run)
```

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