Package: pyinit (via r-universe)

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Type Package

Title Pena-Yohai Initial Estimator for Robust S-Regression

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Description Deterministic Pena-Yohai initial estimator for robust S estimators of regression. The procedure is described in detail in Pena, D., & Yohai, V. (1999) <doi:10.2307/2670164>.

Imports robustbase

Suggests testthat

License GPL (>= 2)

URL https://github.com/dakep/pyinit

BugReports https://github.com/dakep/pyinit/issues

NeedsCompilation yes

Biarch true

Copyright See the file COPYRIGHTS for copyright details on some of the functions.

RoxygenNote 7.1.2

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mscale   *Robust M-estimate of Scale*

Description

Compute the M-estimate of scale using the MAD as initial estimate.

Usage

```
mscale(
  x,
  delta = 0.5,
  rho = c("bisquare", "huber", "gauss"),
  cc,
  eps = 1e-08,
  maxit = 200
)
```

Arguments

x	numeric vector.
delta	desired value for the right-hand side of the M-estimation equation.
rho	rho function to use in the M-estimation equation. Valid options are bisquare, huber and gauss.
cc	non-negative constant for the chosen rho function. If missing, it will be chosen such that the expected value of the rho function under the normal model is equal to delta.
eps	threshold for convergence. Defaults to 1e-8.
maxit	maximum number of iterations. Defaults to 200.

Details

This solves the M-estimation equation given by

$$\sum_{i=1}^{n} \rho(x_i/s_n; cc) = ndelta$$

All NA values in x are removed before calculating the scale.

Value

Numeric vector of length one containing the solution s_n to the equation above.

| pyinit | *PY (Pena-Yohai) initial estimates for S-estimates of regression* |

Description

Computes the PY initial estimates for S-estimates of regression.

Usage

```
pyinit(
  x,
  y,
  intercept = TRUE,
  delta = 0.5,
  cc,
  maxit = 10,
  psc_keep,
  resid_keep_method = c("threshold", "proportion"),
  resid_keep_prop,
  resid_keep_thresh,
  eps = 1e-08,
  mscale_maxit = 200,
  mscale_tol = eps,
  mscale_rho_fun = c("bisquare", "huber", "gauss")
)
```

Arguments

x	a matrix with the data, each observation in a row.
y	the response vector.
intercept	logical, should an intercept be included in the model? Defaults to `TRUE`.
delta, cc	parameters for the M-scale estimator equation. If `cc` is missing it will be set to yield consistency under the Normal model for the given `delta` (right-hand side of the M-scale equation).
maxit	the maximum number of iterations to perform.
psc_keep	proportion of observations to keep based on PSCs.
resid_keep_method	
	how to clean the data based on large residuals. If `"threshold"`, all observations with scaled residuals larger than `resid_keep_thresh` will be removed (`resid_keep_thresh` corresponds to the constant C_1 from equation (21) in Pena & Yohai (1999). If `"proportion"`, observations with the largest `resid_keep_prop` residuals will be removed.
resid_keep_prop, resid_keep_thresh	
	see parameter `resid_keep_method` for details.
eps	the relative tolerance for convergence. Defaults to `1e-8`.

mscale_maxit	maximum number of iterations allowed for the M-scale algorithm. Defaults to 200.
mscale_tol	convergence threshold for the m-scale
mscale_rho_fun	A string containing the name of the rho function to use for the M-scale. Valid options are `bisquare`, `huber` and `gauss`.

Value

coefficients	numeric matrix with coefficient vectors in columns. These are regression estimators based on "cleaned" subsets of the data. The M-scales of the corresponding residuals are returned in the entry `objective`. The regression coefficients with smallest estimated residual scale is in the first column, but the others need not be ordered.
objective	vector of values of the M-scale estimate of the residuals associated with each vector of regression coefficients in the columns of `coefficients`.

References

Pena, D., & Yohai, V.. (1999). A Fast Procedure for Outlier Diagnostics in Large Regression Problems. *Journal of the American Statistical Association*, 94(446), 434-445. <doi:10.2307/2670164>

Examples

```
# generate a simple synthetic data set for a linear regression model
# with true regression coefficients all equal to one "(1, 1, 1, 1, 1)"
set.seed(123)
x <- matrix(rnorm(100*4), 100, 4)
y <- rnorm(100) + rowSums(x) + 1
# add masked outliers
a <- svd(var(x))$v[,4]
x <- rbind(x, t(outer(a, rnorm(20, mean=4, sd=1))))
y <- c(y, rnorm(20, mean=-2, sd=.2))

# these outliers are difficult to find
plot(lm(y~x), ask=FALSE)

# use pyinit to obtain estimated regression coefficients
tmp <- pyinit(x=x, y=y, resid_keep_method='proportion', psc_keep = .5, resid_keep_prop=.5)
# the vector of regression coefficients with smallest residuals scale
# is returned in the first column of the "coefficients" element
tmp$coefficients[,1]
# compare that with the LS estimator on the clean data
coef(lm(y~x, subset=1:100))
# compare it with the LS estimator on the full data
coef(lm(y~x))
```

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