

## cbqr

In this file, we detail the documentation of the functions in `cbqr` implemented in R.

### ❖ `cbqr.gs`

#### Description

`cbqr.gs` implements the methodology censored bent line quantile regression. In the first step, it estimates an informative subset (Tang, Wang, He, & Zhu, 2012). In the second step, it estimates a bent line quantile regression model based on (Li, Wei, Chappell, & He, 2011). The estimation of the change points is based on grid search, and the estimation of regression coefficients is based on `rq` in the package `quantreg`. The standard errors and the interval estimates are provided.

#### Usage

```
cbqr.gs(cy, tt, xx, tau, delta, level, censoring = c('left', 'right'), method = c('ks',  
'nid', 'ker'))
```

#### Arguments

- `cy`: the response (including the censored response).
- `tt`: the covariate that may contain a change point.
- `xx`: the covariates that do not contain a change point.
- `tau`: quantile level
- `delta`: the censoring indicator (0 if censored).
- `level`: significance level. The default is 5%.
- `censoring`: the censoring type. 'right' means censoring from above while 'left' means censoring from below. The default is 'right'.
- `method`: the methods for density estimation involved in asymptotic interval estimation. See Details. The default is 'ks'.

## Details

Similar to [Li et al. \(2011\)](#), we assume the following model for the latent response variable  $Y_{i,j}^*$ ,

$$Y_{i,j}^* = \begin{cases} \alpha_{i,1} + \beta_1 x_{i,j} + \mathbf{z}_{i,j}^\top \boldsymbol{\gamma} + \varepsilon_{i,j} & \text{if } x_{i,j} \leq t \\ \alpha_{i,2} + \beta_2 x_{i,j} + \mathbf{z}_{i,j}^\top \boldsymbol{\gamma} + \varepsilon_{i,j} & \text{if } x_{i,j} > t \end{cases}, \quad i = 1, \dots, n, j = 1, \dots, m_i.$$

Define  $N = \sum_i m_i$ . We only observe  $Y_{i,j} = \max\{Y_{i,j}^*, C\}$ , where  $C$  is a known censoring time point. Following the notations of [Li et al. \(2011\)](#), we are interested in estimating the  $\tau$ th conditional quantile function as

$$Q(\mathbf{w}_{i,j}; \boldsymbol{\theta}_0) = \alpha_i + [\beta_1 I(x_{i,j} \leq t) + \beta_2 I(x_{i,j} > t)](x_{i,j} - t) + \mathbf{z}_{i,j}^\top \boldsymbol{\gamma},$$

where  $\mathbf{w}_{i,j} = (I(i = 1), \dots, I(i = n), x_{i,j}, \mathbf{z}_{i,j}^\top)^\top$  and  $\boldsymbol{\theta}_0 = (\alpha_1, \dots, \alpha_n, \beta_1, \beta_2, \boldsymbol{\gamma}^\top)^\top$ . Therefore, for some small positive constant  $c_N$ , the estimation of  $\boldsymbol{\theta}_0$  is based on

$$\hat{\boldsymbol{\theta}}_N = \underset{\boldsymbol{\theta}}{\operatorname{argmin}} \frac{1}{N} \sum_{i=1}^n \sum_{j=1}^{m_i} \rho_\tau[Y_{i,j} - g(\mathbf{w}_{i,j}; \boldsymbol{\theta})] I[\hat{\pi}(\mathbf{w}_{i,j}) > \tau]$$

where the estimate of the informative subset  $\hat{\pi}(\mathbf{w}_{i,j})$  is obtained by using generalized additive models (GAM) with covariates  $\{\mathbf{w}_{i,j}; i = 1, \dots, n, j = 1, \dots, m_i\}$ .

`method = 'ks'` implements a kernel density estimation, where the residuals are corrected for bias according to the derived Bahadur representation of the estimator. `method = 'ker'` is recommended for larger sample sizes as it may occasionally produce wide intervals when the sample size is small. See details of `method = 'ker'` and `method = 'nid'` in `summary.rq`.

## Value

- `beta.est`: estimates of  $\beta_1, \beta_2$  along with standard error estimates.
- `t.est`: estimates of  $t$  along with standard error estimates.
- `ci.t`: confidence intervals of  $t_0$
- `ci.beta`: confidence intervals of  $\beta_1, \beta_2$
- `gamma.est`: estimates of  $\boldsymbol{\gamma}$  along with standard error estimates.

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## References

- Li, C., Wei, Y., Chappell, R., & He, X. (2011). Bent line quantile regression with application to an allometric study of land mammals' speed and mass. *Biometrics*, 67(1), 242–249.
- Tang, Y., Wang, H. J., He, X., & Zhu, Z. (2012). An informative subset-based estimator for censored quantile regression. *Test*, 21, 635–655.

## Examples

```

subj.m <- seq(from = .5, to = 3, length.out = m)
yy <- c()
tt <- c()
cy <- c()
for (j in 1:m) {
  t <- runif(n, 0, 3)
  t <- t[order(t)]
  tt <- c(tt, t)
  y <- subj.m[j] + .5 * t + 2 * pmax(t - t0, 0) + rnorm(n, sd = .1)

  # censoring
  c <- y
  if (min(which(y > cl)) < Inf) {
    c[min(which(y > cl)):length(c)] <- cl
  }

  yy <- c(yy, y)
  cy <- c(cy, c)
}

# covariate
group <- as.factor(sapply(1:m, function(x) rep(x, n)))
xx <- as.matrix(model.matrix(~ group))
xx <- xx[, - 1]

##### Estimated ISUB

## GAM estimation
delta <- ifelse(yy <= cl, yes = 1, no = 0)

```

```
## Bent QR with estimated ISUB (grid search)
mod.gs <- cbqr.gs(cy = cy, tt = tt, xx = xx, tau = tau, delta = delta)
```