

# Package: tpr (via r-universe)

September 12, 2024

**Type** Package

**Title** Temporal Process Regression

**Version** 0.3-3

**Author** Jun Yan <jun.yan@uconn.edu>

**Maintainer** Jun Yan <jun.yan@uconn.edu>

**Description** Regression models for temporal process responses with time-varying coefficient.

**Depends** R (>= 4.0), stats, lgtdl

**License** GPL (>= 3)

**NeedsCompilation** yes

**Date/Publication** 2022-10-17 07:10:02 UTC

**Repository** <https://jun-yan.r-universe.dev>

**RemoteUrl** <https://github.com/cran/tpr>

**RemoteRef** HEAD

**RemoteSha** 84461c594e22be6f87a1bca0919fcc76911750c2

## Contents

tpr-package . . . . .	2
ci.plot . . . . .	2
dnase . . . . .	3
tpr . . . . .	5
tpr.pfit . . . . .	7
tpr.test . . . . .	8

<b>Index</b>	<b>10</b>
--------------	-----------

---

 tpr-package

*Temporal Process Regression*


---

### Description

Fit regression models for temporal process responses with time-varying and time-independent coefficients.

### Details

An overview of how to use the package, including the most important functions

### Author(s)

Jun Yan <jun.yan@uconn.edu>

### References

Fine, Yan, and Kosorok (2004): Temporal Process Regression. *Biometrika*.

---

 ci.plot

*Confidence Interval Plot*


---

### Description

Plotting time-varying coefficient with pointwise confidence.

### Usage

```
ci.plot(x, y, se, level = 0.95, ylim = NULL, newplot = TRUE,
        fun = gaussian()$linkinv, dfun = gaussian()$mu.eta, ...)
```

### Arguments

x	the x coordinate
y	the y coordinate
se	the standard error of y
level	confidence level
ylim	the range of y axis
newplot	if TRUE, draw a new plot
fun	a transform function
dfun	the derivative of the transform function
...	arguments to be passed to plot

**Author(s)**

Jun Yan <jun.yan@uconn.edu>

---

dnase

*rhDNase Data*

---

**Description**

Randomized trial of rhDNase for treatment of cystic fibrosis

**Usage**

data(dnase)

**Format**

A data frame with 767 observations on the following 6 variables.

id subject id

rx treatment arm: 0 = placebo, 1 = rhDNase

fev forced expiratory volume, a measure of lung capacity

futime follow time

iv1 IV start time

iv2 IV stop time

**Details**

During an exacerbation, patients received intravenous (IV) antibiotics and were considered unsusceptible until seven exacerbation-free days beyond the end of IV therapy.

A few subjects were infected at the time of enrollment, for instance a subject has a first infection interval of -21 to 7. We do not count this first infection as an "event", and the subject first enters the risk set at day 7.

**Source**

Therneau and Grambsch (2000). Modeling Survival Data: Extending the Cox model. Springer. <http://www.mayo.edu/hsr/people/therneau/book/data/dnase.html>

**References**

Yan and Fine (2008). Analysis of Episodic Data with Application to Recurrent Pulmonary Exacerbations in Cystic Fibrosis Patients. JASA.

## Examples

```

## This example steps through how to set up for the tpr function.
## Three objects are needed:
##   1) response process (an object of "lgtdl")
##   2) data availability process (an object of "lgtdl")
##   3) a time-independent covariate matrix

data(dnase)

## extracting the unique id and subject level information
dat <- unique(dnase[,c("id", "fuptime", "fev", "rx")])

## construct temporal process response for recurrent event
rec <- lapply(split(dnase[,c("id", "iv1", "fuptime")], dnase$id),
  function(x) {
    v <- x$iv1
    maxfu <- max(x$fuptime)
    ## iv1 may be negative!!!
    if (is.na(v[1])) c(0, maxfu + 1.0)
    else if (v[1] < 0) c(v[1] - 1, v[!is.na(v)], maxfu + 1.0)
    else c(0, v[!is.na(v)], maxfu + 1.0)
  })

yrec <- lapply(rec,
  function(x) {
    dat <- data.frame(time=x, cov=1:length(x)-1)
    len <- length(x)
    dat$cov[len] <- dat$cov[len - 1]
    as.lgtdl(dat)
  })

## construct temporal process response for accumulative days exacerbation
do1.acc <- function(x) {
  gap <- x$iv2 - x$iv1 + 1
  if (all(is.na(gap))) yy <- tt <- NULL
  else {
    gap <- na.omit(gap)
    yy <- cumsum(rep(1, sum(gap)))
    tt <- unlist(sapply(1:length(gap), function(i)
      seq(x$iv1[i], x$iv2[i], by=1.0)))
  }
  yy <- c(0, yy, rev(yy)[1])
  if (!is.null(tt[1]) && tt[1] < 0)
    tt <- c(tt[1] - 1, tt, max(x$fuptime) + 1.0)
  else tt <- c(0, tt, max(x$fuptime) + 1.0)
  as.lgtdl(data.frame(time=tt, cov=yy))
}

yacc <- lapply(split(dnase[,c("id", "iv1", "iv2", "fuptime")], dnase$id),
  do1.acc)

## construct data availability (or at risk) indicator process

```

```

tu <- max(dat$futime) + 0.001
rt <- lapply(1:nrow(dat),
  function(i) {
    x <- dat[i, "futime"]
    time <- c(0, x, tu)
    cov <- c(1, 0, 0)
    as.lgtdl(data.frame(time=time, cov=cov))
  })

## time-independent covariate matrix
xmat <- model.matrix(~ rx + fev, data=dat)
## time-window in days
tlim <- c(10, 168)
good <- unlist(lapply(yrec, function(x) x$time[1] == 0))

## fully functional temporal process regression

## for recurrent event
m.rec <- tpr(yrec, rt, xmat[,1:3], list(), xmat[,-(1:3),drop=FALSE], list(),
  tis=10:160, w = rep(1, 151), family = poisson(),
  evstr = list(link = 5, v = 3))
par(mfrow=c(1,3), mgp=c(2,1,0), mar=c(4,2,1,0), oma=c(0,2,0,0))
for(i in 1:3) ci.plot(m.rec$tis, m.rec$alpha[,i], sqrt(m.rec$svalpha[,i]))
## hypothesis test of significance
## integral test, covariate index 2 and 3
sig.test.int.ff(m.rec, idx=2:3, ncut=2)
sig.test.boots.ff(m.rec, idx=2:3, nsim=1000)
## constant fit
cfit <- cst.fit.ff(m.rec, idx=2:3)

## goodness-of-fit test for constant fit
gof.test.int.ff(m.rec, idx=2:3, ncut=2)
gof.test.boots.ff(m.rec, idx=2:3, nsim=1000)

## for cumulative days in exacerbation
m.acc <- tpr(yacc, rt, xmat[,1:3], list(), xmat[,-(1:3),drop=FALSE], list(),
  tis=10:160, w = rep(1, 151), family = gaussian(),
  evstr = list(link = 1, v = 1))
par(mfrow=c(1,3), mgp=c(2,1,0), mar=c(4,2,1,0), oma=c(0,2,0,0))
for(i in 1:3) ci.plot(m.acc$tis, m.acc$alpha[,i], sqrt(m.acc$svalpha[,i]))

```

**Description**

Regression for temporal process responses and time-independent covariate. Some covariates have time-varying coefficients while others have time-independent coefficients.

**Usage**

```
tpr(y, delta, x, xtv=list(), z, ztv=list(), w, tis,
    family = poisson(),
    evstr = list(link = 5, v = 3),
    alpha = NULL, theta = NULL,
    tidx = 1:length(tis),
    kernstr = list(kern=1, poly=1, band=range(tis)/50),
    control = list(maxit=25, tol=0.0001, smooth=0, intsmooth=0))
```

**Arguments**

y	Response, a list of "lgtld" objects.
delta	Data availability indicator, a list of "lgtld" objects.
x	Covariate matrix for time-varying coefficients.
xtv	A list of list of "lgtld" for time-varying covariates with time-varying coefficients.
z	NOT READY YET; Covariate matrix for time-independent coefficients.
ztv	NOT READY YET; A list of list of "lgtld" for time-varying covariates with time-independent coefficients.
w	Weight vector with the same length of tis.
tis	A vector of time points at which the model is to be fitted.
family	Specification of the response distribution; see family for glm; this argument is used in getting initial estimates.
evstr	A list of two named components, link function and variance function. link: 1 = identity, 2 = logit, 3 = probit, 4 = cloglog, 5 = log; v: 1 = gaussian, 2 = binomial, 3 = poisson
alpha	A matrix supplying initial values of alpha.
theta	A numeric vector supplying initial values of theta.
tidx	indices for time points used to get initial values.
kernstr	A list of two names components: kern: 1 = Epanechnikov, 2 = triangular, 0 = uniform; band: bandwidth
control	A list of named components: maxit: maximum number of iterations; tol: tolerance level of iterations. smooth: 1 = smoothing; 0 = no smoothing.

**Details**

This rapper function can be made more user-friendly in the future. For example, evstr can be determined from the family argument.

**Value**

An object of class "tpr":

tis	same as the input argument
alpha	estimate of time-varying coefficients
beta	estimate of time-independent coefficients
valpha	a matrix of variance of alpha at tis
vbeta	a matrix of variance of beta at tis
niter	the number of iterations used
infAlpha	a list of influence functions for alpha
infBeta	a matrix of influence functions for beta

**Author(s)**

Jun Yan <jun.yan@uconn.edu>

**References**

- Fine, Yan, and Kosorok (2004). Temporal Process Regression. *Biometrika*.  
 Yan and Huang (2009). Partly Functional Temporal Process Regression with Semiparametric Profile Estimating Functions. *Biometrics*.

---

tpr.pfit

*Constant fit of coefficients in a TPR model*

---

**Description**

Weighted least square estimate of a constant model for time-varying coefficients in a TPR model.

**Usage**

```
cst.fit.ff(fit, idx)
```

**Arguments**

fit	a fitted object from tpr
idx	the index of the

**Value**

The estimated constant fit, standard error, z-value and p-value.

**Author(s)**

Jun Yan <jun.yan@uconn.edu>

**References**

Fine, Yan, and Kosorok (2004). Temporal Process Regression. *Biometrika*.

**See Also**

[tpr.test](#)

---

tpr.test

*Significance and Goodness-of-fit Test of TPR*

---

**Description**

Two kinds of tests are provided for inference on the coefficients in a fully functional TRP model: integral test and bootstrap test.

**Usage**

```
sig.test.int.ff(fit, chypo = 0, idx, weight = TRUE, ncut = 2)
sig.test.boots.ff(fit, chypo = 0, idx, nsim = 1000, plot = FALSE)
gof.test.int.ff(fit, cfitList = NULL, idx, weight = TRUE, ncut = 2)
gof.test.boots.ff(fit, cfitList = NULL, idx, nsim = 1000, plot = FALSE)
gof.test.boots.pf(fit1, fit2, nsim, p = NULL, q = 1)
```

**Arguments**

fit	a fitted object from tpr
chypo	hypothesized value of coefficients
idx	the index of the coefficients to be tested
weight	whether or not use inverse variation weight
ncut	the number of cuts of the interval of interest in integral test
cfitList	a list of fitted object from <code>cst.fit.ff</code>
nsim	the number of bootstrap samples in bootstrap test
plot	whether or not plot
fit1	fit of H0 model (reduced)
fit2	fit of H1 model (full)
p	the index of the time-varying estimation in fit2
q	the index of the time-independent estimation in fit1

**Value**

Test statistics and their p-values.

**Author(s)**

Jun Yan <jun.yan@uconn.edu>



**References**

Fine, Yan, and Kosorok (2004). Temporal Process Regression. *Biometrika*.

**See Also**

[tpr](#)

**Examples**

## see ?tpr

# Index

- \* **aplot**

- ci.plot, 2

- \* **datasets**

- dnase, 3

- \* **hplot**

- ci.plot, 2

- \* **htest**

- tpr.test, 8

- \* **models**

- tpr.pfit, 7

- \* **multivariate**

- tpr.test, 8

- \* **package**

- tpr-package, 2

- \* **regression**

- tpr, 5

- \* **robust**

- tpr, 5

- tpr.pfit, 7

ci.plot, 2

cst.fit.ff(tpr.pfit), 7

dnase, 3

gof.test.boots.ff(tpr.test), 8

gof.test.boots.pf(tpr.test), 8

gof.test.int.ff(tpr.test), 8

sig.test.boots.ff(tpr.test), 8

sig.test.int.ff(tpr.test), 8

tpr, 5, 9

tpr-package, 2

tpr.pfit, 7

tpr.test, 8, 8