

# Package ‘longitudinalANAL’

October 13, 2023

**Type** Package

**Title** Longitudinal Data Analysis

**Version** 0.2

**Description** Regression analysis of mixed sparse synchronous and asynchronous longitudinal covariates. Please cite the manuscripts corresponding to this package: Sun, Z. et al. (2023) <[arXiv:2305.17715](#)> and Liu, C. et al. (2023) <[arXiv:2305.17662](#)>.

**License** GPL-3

**Encoding** UTF-8

**Depends** dplyr, tibble, MASS, dlm

**RoxygenNote** 7.2.3

**NeedsCompilation** no

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**Repository** CRAN

**Date/Publication** 2023-10-13 12:00:02 UTC

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lda *Longitudinal data analysis*

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### Description

This function provide regression analysis of mixed sparse synchronous and asynchronous longitudinal covariates.

### Usage

```
lda(data_res, data_cov, N, bd, omit, method)
```

### Arguments

data_res	An object of class tibble. The structure of the tibble must be: <code>tibble(id_y=ID, ty=measurement time for response, y=observation for response, x=matrix(observation for synchronous covariates), x_add=matrix(observation for uninterested synchronous covariates))</code> .
data_cov	An object of class tibble. The structure of the tibble must be: <code>tibble(id_z=ID, tz=measurement time for response, z=matrix(observation for asynchronous covariates))</code> .
N	An object of class integer. The sample size.
bd	An object of class vector. If use auto bandwidth selection, the structure of the vector must be: <code>d=c(the maximum bandwidth, the minimum bandwidth, the fold of cross-validation, the number of bandwidth divided)</code> . If use fixed bandwidth, <code>bd=c(the chosen bandwidth)</code> .
omit	An object of class integer indicating the method used to do estimation for synchronous covariates. If use plm method, <code>omit=1</code> ; if use centering method, <code>omit=2</code> ; if use additional covariates information, <code>omit=3</code> .
method	An object of class integer indicating the method used to do estimation for asynchronous covariates. If only deal with omit variable, <code>method=0</code> ; if use two-stage method, <code>method=1</code> ; if use kernel smoothing, <code>method=2</code> .

### Value

a list with the following elements:

est	The estimation for the corresponding parameters.
se	The estimation of standard error for the estimated parameters.

### Examples

```
library(MASS)
library(tibble)
library(dplyr)
```

```

N=100
ty=tz=y=x=z=id_y=id_z=list()
a=b=g=1
ny=rpois(N,5)+1
nz=rpois(N,5)+1
for(i in 1:N){
  ty[[i]]=as.matrix(runif(ny[i]))
  tz[[i]]=as.matrix(runif(nz[i]))
  t.temp=rbind(tz[[i]],ty[[i]])
  n.temp=nz[i]+ny[i]
  corr=exp(-abs(rep(1,n.temp)%*%t(t.temp)-t.temp)%*%t(rep(1,n.temp))))
  corr.e=2^(-abs(rep(1,n.temp)%*%t(t.temp)-t.temp)%*%t(rep(1,n.temp))))
  MX=t.temp^.5
  MZ=rep(0, n.temp)
  x.temp=mvrnorm(1,MX,corr)
  z.temp=mvrnorm(1,MZ, corr)
  z[[i]]=as.matrix(z.temp[1:nz[i]])
  x[[i]]=as.matrix(x.temp[-(1:nz[i])])
  id_z[[i]]=rep(i,nz[i])
  id_y[[i]]=rep(i,ny[i])
  y.temp=a+g*z.temp+x.temp*b+as.matrix(mvrnorm(1,rep(0,n.temp),corr.e))
  y[[i]]=as.matrix(y.temp[-(1:nz[i])])
}
data_cov=tibble(id_z=unlist(id_z),tz=unlist(tz),z=matrix(unlist(z),length(unlist(z))))
data_res=tibble(id_y=unlist(id_y),ty=unlist(ty),x=matrix(unlist(x),length(unlist(x))),y=unlist(y))
bd=0.1
omit=1
method=1
lda(data_res,data_cov,N,bd,omit,method)

```

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ldatv

*Longitudinal data analysis*


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### Description

This function provide regression analysis of mixed sparse synchronous and asynchronous longitudinal covariates with time-varying coefficients.

### Usage

```
ldatv(data_res, data_cov, time, N, bd, method, scb)
```

### Arguments

**data\_res** An object of class tibble. The structure of the tibble must be: `tibble(id_y=ID, ty=measurement time for response, y=observation for response, x=matrix(observation for synchronous covariates), x_add=matrix(observation for uninterested synchronous covariates))`.

<code>data_cov</code>	An object of class tibble. The structure of the tibble must be: <code>tibble(id_z=ID, tz=measurement time for response, z=matrix(observation for asynchronous covariates))</code> .
<code>time</code>	An object of class vector. The interest time.
<code>N</code>	An object of class integer. The sample size.
<code>bd</code>	An object of class vector. If use auto bandwidth selection, the structure of the vector must be: <code>bd=c</code> (the maximum bandwidth for h1, the minimum bandwidth for h1, the maximum bandwidth for h2, the minimum bandwidth for h2, the fold of cross-validation, the number of bandwidth divided). If use fixed bandwidth, <code>bd=c</code> (the chosen bandwidth).
<code>method</code>	An object of class integer indicating the method used to do estimation for asynchronous covariates. If use one-stage method, <code>method=1</code> ; if use two-stage method with centering method for the first stage, <code>method=1</code> ; if use two-stage method with time-varying method for the first stage, <code>method=2</code> .
<code>scb</code>	An object of class vector. If need to construct the simultaneous confidence band, the structure of the vector must be: <code>c</code> ( <code>alpha</code> =desirable confidence level, <code>B</code> =repeat times). Otherwise, <code>scb=0</code> .

### Value

a list with the following elements:

<code>est.b</code>	The estimation for the parameter of synchronous covariates.
<code>est.g</code>	The estimation for the parameter of asynchronous covariates.
<code>se.b</code>	The estimation of standard error for the parameter of synchronous covariates.
<code>se.g</code>	The estimation of standard error for the parameter of asynchronous covariates.
<code>c_alpha_x</code>	The empirical percentile used to construct the simultaneous confidence band for the parameter of synchronous covariates.
<code>c_alpha_z</code>	The empirical percentile used to construct the simultaneous confidence band for the parameter of asynchronous covariates.

### Examples

```
library(dplyr)
library(MASS)
library(tibble)
N=400
ty=tz=y=x=x1=z=id_y=id_z=list()
beta<-function(t){
  0.3*(t-0.4)^2
}
gamma<-function(t){
  sin(2*pi*t)
}
ny=rpois(N,5)+1
nz=rpois(N,5)+1
```

```

for(i in 1:N){
  ty[[i]]=as.matrix(runif(ny[i]))
  tz[[i]]=as.matrix(runif(nz[i]))
  t.temp=rbind(tz[[i]],ty[[i]])
  n.temp=nz[i]+ny[i]
  corr=exp(-abs(rep(1,n.temp)%*%t(t.temp)-t.temp%*%t(rep(1,n.temp))))
  corr.e=2^(-abs(rep(1,n.temp)%*%t(t.temp)-t.temp%*%t(rep(1,n.temp))))
  MX=rep(0, n.temp)
  MZ= 2*(t.temp-0.5)^2
  x.temp=mvnorm(1,MX,corr)
  z.temp=mvnorm(1,MZ, corr)
  z[[i]]=as.matrix(z.temp[1:nz[i]])
  x[[i]]=as.matrix(x.temp[-(1:nz[i])])
  id_z[[i]]=rep(i,nz[i])
  id_y[[i]]=rep(i,ny[i])
  y.temp=gamma(t.temp)*z.temp+beta(t.temp)*x.temp+as.matrix(mvnorm(1,rep(0,n.temp),corr.e))
  y[[i]]=as.matrix(y.temp[-(1:nz[i])])
}
data_cov=tibble(id_z=unlist(id_z),tz=unlist(tz),z=matrix(unlist(z),length(unlist(z))))
data_res=tibble(id_y=unlist(id_y),ty=unlist(ty),x=matrix(unlist(x),length(unlist(x))), y=unlist(y))
ldatv(data_res,data_cov,time=0.3,N,bd=c(N^(-0.5),N^(-0.5)),method=1,scb=0)

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