

Package ‘lava.tobit’

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Type Package

Title Latent Variable Models with Censored and Binary Outcomes

Version 0.5

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Description Lava plugin allowing combinations of left and right censored and binary outcomes.

URL <https://github.com/kkholst/lava.tobit>

BugReports <https://github.com/kkholst/lava.tobit/issues>

Depends lava (>= 1.5), mets (>= 1.2.2), mvtnorm, survival

License GPL-3

LazyLoad yes

RoxygenNote 6.0.1

NeedsCompilation no

Repository CRAN

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R topics documented:

binary	2
clprobit	3
lava.tobit	3
lava.tobit.init.hook	4
plotres	5
rbind.Surv	6

Index

7

binary	<i>Defines variables as binary</i>
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Description

Defines variables of a latent variable model (`lvm`-object from the `lava` package) as binary.

Usage

```
## S3 method for class 'lvm'
binary(x, var=NULL, ...)
## S3 replacement method for class 'lvm'
binary(x, ...) <- value
```

Arguments

<code>x</code>	<code>lvm</code> -object
<code>...</code>	Additional arguments parsed to lower-level functions
<code>var</code>	Formula or vector of variable names
<code>value</code>	Formula or vector of variable names

Details

This function sets the status of variables to 'binary'. For use with the `estimate` method a probit-link will be assumed. Used with the `sim` method normal distributed data will be simulated followed by thresholding at 0. To simulate data where the dichotomous variable has a direct effect on the outcome the `distribution` method can be used, e.g. `distribution(m, ~x) <- binomial.lvm("probit")`.

Value

`lvm`-object (or vector of variable names if called without any arguments)

Author(s)

Klaus K. Holst

clprobit	<i>Composite Likelihood for probit latent variable models</i>
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Description

Estimate parameters in a probit latent variable model via a composite likelihood decomposition.

Usage

```
clprobit(x, data, k = 2, type = c("nearest", "all"), pairlist,
         silent = TRUE, ...)
```

Arguments

x	lvm-object
data	data.frame
k	Size of composite groups
type	Determines number of groups. With type="nearest" (default) only neighboring items will be grouped, e.g. for k=2 (y1,y2),(y2,y3),... With type="all" all combinations of size k are included
pairlist	A list of indices specifying the composite groups. Optional argument which overrides k and type but gives complete flexibility in the specification of the composite likelihood
silent	Turn output messages on/off
...	Additional arguments parsed on to lower-level functions

Value

An object of class `clprobit` inheriting methods from `lvm`

Author(s)

Klaus K. Holst

See Also

[estimate](#)

<code>lava.tobit</code>	<i>Estimation and simulation of probit and tobit latent variable models</i>
	<i>Framework for estimating parameters and simulate data from Latent Variable Models with binary and censored observations. Plugin for the lava package</i>

Estimation and simulation of probit and tobit latent variable models
Framework for estimating parameters and simulate data from Latent Variable Models with binary and censored observations. Plugin for the lava package

Description

```
Package:      lava.tobit
Type:        Package
Version:     0.4-5
Date:        2012-03-15
License:     GPL-3
LazyLoad:    yes
```

Author(s)

Klaus K. Holst Maintainer: <kkho@biostat.ku.dk>

Examples

```
library(lava.tobit)
m <- lvm(list(c(y,z) ~ x, y~z))
## Simulate 200 observation from path analysis model
## with all slopes and residual variances set to 1 and intercepts 0:
d <- sim(m,200,seed=1)
## Dichotomize y and introduce censoring on z
d <- transform(d, y=as.factor(y>0), z=Surv(z,z<2))

e <- estimate(m,d,control=list(trace=1),estimator="gaussian")
effects(e,y~x)
```

lava.tobit.init.hook *For internal use*

Description

For internal use

Author(s)

Klaus K. Holst

<code>plotres</code>	<i>Plot distribution of standardized residuals</i>
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Description

Plot empirical (KM) and model-specific cumulative distribution function of standardized residuals

Usage

```
plotres(x, var = lava::endogenous(x),
        ylab = "Cumulative Distribution Function",
        xlab = "Standardized residuals", main, k, ...)
```

Arguments

<code>x</code>	Model, <code>lvmfit</code> object
<code>var</code>	Character vector of (endogenous) variable names
<code>ylab</code>	Label of x-axis
<code>xlab</code>	Label of y-axis
<code>main</code>	Title of plot
<code>k</code>	Optional group number for multiple group analysis
<code>...</code>	Additional argument

Author(s)

Klaus K. Holst

Examples

```
## Not run:

## Simulate data where (y01,y2)
## follows conditional bivariate normal distribution
## given covariate x. Instead of y01 we observe
## right censored version y2
n <- 200
m <- lvm(c(y01,y2) ~ x)
covariance(m) <- y01~y2
set.seed(1)
d <- sim(m,n)
d$cens1 <- rexp(n)
d$status1 <- with(d,y01<cens1)
d$y1 <- with(d, pmin(y01,cens1))

## Estimate model parameters
d$S1 <- with(d, Surv(y1,status1))
```

```
m <- lvm(c(S1,y2)~x); covariance(m) <- S1~y2
e <- estimate(m,d,control=list(trace=1))

## Plot cumulative distribution functions
par(mfrow=c(2,2)); plotres(e); plot(e)

## End(Not run)
```

rbind.Surv*Appending Surv objects***Description**

`rbind` method for `Surv` objects

Usage

```
## S3 method for class 'Surv'
rbind(...)
```

Arguments

... `Surv` objects

Value

`Surv` object

Author(s)

Klaus K. Holst

Examples

```
y <- yl <- yr <- rnorm(10)
yl[1:5] <- NA; yr[6:10] <- NA
S1 <- Surv(yl,yr,type="interval2")
S2 <- Surv(y,y>0,type="right")
S3 <- Surv(y,y<0,type="left")

rbind(S1,S1)
rbind(S2,S2)
rbind(S3,S3)
```

Index

*Topic **models**
 binary, 2
 clprobit, 3
 plotres, 5
*Topic **package**
 lava.tobit, 3
*Topic **regression**
 binary, 2
 clprobit, 3
*Topic **utilities**
 lava.tobit.init.hook, 4
 rbind.Surv, 6

binary, 2
binary<- (binary), 2

clprobit, 3

estimate, 3

lava.tobit, 3
lava.tobit-package (lava.tobit), 3
lava.tobit.color.hook
 (lava.tobit.init.hook), 4
lava.tobit.init.hook, 4
lava.tobit.sim.hook
 (lava.tobit.init.hook), 4

plotres, 5

rbind.Surv, 6

tobit_gradient.lvm
 (lava.tobit.init.hook), 4
tobit_hessian.lvm
 (lava.tobit.init.hook), 4
tobit_logLik.lvm
 (lava.tobit.init.hook), 4
tobit_method.lvm
 (lava.tobit.init.hook), 4

tobit_objective.lvm
 (lava.tobit.init.hook), 4
tobitw_gradient.lvm
 (lava.tobit.init.hook), 4
tobitw_hessian.lvm
 (lava.tobit.init.hook), 4
tobitw_method.lvm
 (lava.tobit.init.hook), 4