Package 'CBPS'

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Title R Package for Covariate Balancing Propensity Score

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Imports arm

Description CBPS is an R package that implements the covariate balancing propensity score proposed by Imai and Ratkovic (2012). The propensity score is estimated such that it maximizes the resulting covariate balance as well as the prediction of treatment assignment. The method, therefore, avoids an iteration between model fitting and balance checking. Currently, the package can only deal with binary treatment in the cross section setting but in the future it will be extended to other settings including non-binary treatments and causal inference with longitudinal data.

LazyLoad yes

LazyData yes

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R topics documented:

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Description

CBPS and CBPS.fit estimate propensity score such that both covariate balance and prediction of treatment assignment are maximized. The method, therefore, avoids an iterative process between model fitting and balance checking and implements both simultaneously.

Usage

```
CBPS(formula, data, na.action, ATT, method = "over", ...)
CBPS.fit(treat, X, ATT, X.bal = X, method, ...)
```

Arguments

an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment (formula), typically the environment from which CBPS is called. na.action A function which indicates what should happen when the data contain NAs. The default is set by the na.action setting of options, and is na.fail if that is unset. ATT Set to TRUE to find the average treatment effect on the treated, and FALSE to find the average treatment effect. Default is TRUE. treat A vector of treatment assignments. X A covariate matrix. X .bal A matrix to be balanced. Currently not accepted. "over" for fitting an over-identified model that combines the propensity score and covariate balancing conditions; "exact" for fitting a model that only contains the covariate balancing conditions. Other parameters to be passed through to optim()	formula	An object of class formula (or one that can be coerced to that class): a symbolic description of the model to be fitted.
default is set by the na.action setting of options, and is na.fail if that is unset. ATT Set to TRUE to find the average treatment effect on the treated, and FALSE to find the average treatment effect. Default is TRUE. treat A vector of treatment assignments. X A covariate matrix. X.bal A matrix to be balanced. Currently not accepted. method "over" for fitting an over-identified model that combines the propensity score and covariate balancing conditions; "exact" for fitting a model that only contains the covariate balancing conditions.	data	to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment(formula), typically the environment
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X A covariate matrix. X.bal A matrix to be balanced. Currently not accepted. method "over" for fitting an over-identified model that combines the propensity score and covariate balancing conditions; "exact" for fitting a model that only contains the covariate balancing conditions.	ATT	
 X.bal A matrix to be balanced. Currently not accepted. method "over" for fitting an over-identified model that combines the propensity score and covariate balancing conditions; "exact" for fitting a model that only contains the covariate balancing conditions. 	treat	A vector of treatment assignments.
method "over" for fitting an over-identified model that combines the propensity score and covariate balancing conditions; "exact" for fitting a model that only contains the covariate balancing conditions.	Χ	A covariate matrix.
and covariate balancing conditions; "exact" for fitting a model that only contains the covariate balancing conditions.	X.bal	A matrix to be balanced. Currently not accepted.
Other parameters to be passed through to optim()	method	and covariate balancing conditions; "exact" for fitting a model that only contains
····		Other parameters to be passed through to optim()

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Details

Fits covariate balancing propensity scores.

Value

coefficients A named vector of coefficients

residuals The working residuals from the final iteration of CBPS

fitted.values The fitted mean values, obtained by transforming the linear predictors by the

inverse of the binomial function

rank The numeric rank of the fitted model

family The family object used.

deviance Minus twice the log-likelihood of the CBPS fit. Will be lower than the maximum

likelihood deviance.

weights The CBPS balancing weights calculated from the propensity scores.

y The treatment vector used

x The covariate matrix

model The model frame

converged Convergence value. Returned from the call to optim().

call The matched call

formula The formula supplied.
data The data argument.

J The J-statistic at convergence.

df The degrees of freedom.

var The covariance matrix, evaluated numerically from optim().

Author(s)

Marc Ratkovic, Kosuke Imai, and Christian Fong; The CBPS function is based on the code for version 2.15.0 of the glm function implemented in the stats package, originally written by Simon Davies. This documentation is likewise modeled on the documentation for glm and borrows its language where the arguments and values are the same.

References

Imai, Kosuke and Marc Ratkovic. "Covariate Balancing Propensity Score." Unpublished Manuscript, Princeton University. http://imai.princeton.edu/research/CBPS.html

See Also

summary.CBPS

DR DR

Examples

```
## Not run:
###
### Example: propensity score matching
##Load the LaLonde data
data(LaLonde)
## Estimate CBPS via logistic regression
fit <- CBPS(treat ~ age + educ + re75 + re74 + I(re75==0) + I(re74==0), data = LaLonde, ATT = TRUE)
summary(fit)
## matching via MatchIt: one to one nearest neighbor with replacement
library(MatchIt)
m.out <- matchit(treat ~ 1, distance = fitted(fit), method = "nearest", data = LaLonde, replace = TRUE)</pre>
### Example: propensity score weighting
###
## Simulation from Kang and Shafer (2007).
set.seed(123456)
n <- 1000
X \leftarrow mvrnorm(n, mu = rep(0, 4), Sigma = diag(4))
prop <-1 / (1 + exp(X[,1] - 0.5 * X[,2] + 0.25*X[,3] + 0.1 * X[,4]))
treat <- rbinom(n, 1, prop)</pre>
y \leftarrow 210 + 27.4*X[,1] + 13.7*X[,2] + 13.7*X[,3] + 13.7*X[,4] + rnorm(n)
##Estimate CBPS with a misspecificied model
X.mis \leftarrow cbind(exp(X[,1]/2), X[,2]*(1+exp(X[,1]))^{-1}+10, (X[,1]*X[,3]/25+.6)^3, (X[,2]+X[,4]+20)^2)
fit1 <- CBPS(treat ~ X.mis, ATT = TRUE)</pre>
## Horwitz-Thompson estimate
mean(treat*y/fit1$fitted.values)
## Inverse probability weighting
sum(treat*y/fit1$fitted.values)/sum(treat/fit1$fitted.values)
## End(Not run)
```

DR

Doubly Robust Estimator

Description

DR calculates the doubly-robust estimator as as described in Lunceford and Davidian (2004) and originally detailed in Robins, Rotnitzky, and Zhao (1994).

Usage

```
DR(formula, model, data, treat, pscore)
```

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Arguments

formula An object of class formula (or one that can be coerced to that class): a symbolic

description of the model to be fitted.

model Set to "lm" to use a linear model, set to "glm" for the generalized linear model.

data An optional data frame, list or environment (or object coercible by as.data.frame

to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment (formula), typically the environment

from which CBPS is called.

treat A vector of treatment assignments, where 1 indicates assignment to the treat-

ment group and 0 indicates assignment to the control group.

pscore A vector of propensity scores.

Details

Fits covariate balancing propensity scores.

Value

Returns a vector with the point estimate of the doubly-robust estimator as well as its standard error. For use with CBPS, run only with the ATE.

Author(s)

Marc Ratkovic, Kosuke Imai, and Christian Fong.

References

Lunceford, Jared and Marie Davidian. "Stratification and Weighting Via the Propensity Score in Estimation of Causal Treatment Effects: A Comparative Study." Statistics in Medicine 23, 15 October 2004.

See Also

IPW

Examples

```
## Not run:
###
### Example: Doubly robust estimator
###
##Load the LaLonde data
data(LaLonde)
## Estimate CBPS via logistic regression for ATE. Run only with ATE.
fit <- CBPS(treat ~ age + educ + re75 + re74 + I(re75==0) + I(re74==0), data = LaLonde, ATT = FALSE)
## Find doubly robust estimator under GLM
doubly.robust <- DR(re78 ~ age + educ + re75 + re74 + I(re75==0) + I(re74==0), model="glm", data = LaLonde, tr</pre>
```

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```
## End(Not run)
```

IPW

Inverse Probability Weighting Estimators

Description

IPW calculates the inverse probability weighting estimators from Lunceford and Davidian (2004).

Usage

```
IPW(outcome, treat, data, pscore, k)
```

Arguments

8	
outcome	A vector of the outcome variable.
treat	A vector of treatment assignments, where 1 indicates assignment to the treatment group and 0 indicates assignment to the control group.
data	an optional data frame, list or environment (or object coercible by as.data.frame to a data frame) containing the variables in the model. If not found in data, the variables are taken from environment (formula), typically the environment from which CBPS is called.
pscore	A vector of propensity scores.
k	The number of covariates considered in the generation of the propensity scores.

Details

Calculates the inverse probability weighting estimators from Lunceford and Davidian (2004). For use with CBPS, run only with ATE.

Value

Returns a data frame whose columns are the point estimates and standard errors for these inverse probability weighting estimators, and whose rows correspond to IPW1, IPW2, and IPW3.

Author(s)

Marc Ratkovic, Kosuke Imai, and Christian Fong.

References

Lunceford, Jared and Marie Davidian. "Stratification and Weighting Via the Propensity Score in Estimation of Causal Treatment Effects: A Comparative Study." Statistics in Medicine 23, 15 October 2004.

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See Also

DR

Examples

```
## Not run:
###
### Example: propensity score matching
###

##Load the LaLonde data
data(LaLonde)
## Estimate CBPS via logistic regression for ATE (Run with ATE only)
fit <- CBPS(treat ~ age + educ + re75 + re74 + I(re75==0) + I(re74==0), data = LaLonde, ATT = FALSE)
##Calculate the IPW estimators
inverse.prob.wt <- IPW(outcome = re78, treat = treat, data = LaLonde, pscore = fitted(fit), k = length(coef(fit))
## End(Not run)</pre>
```

LaLonde

LaLonde Data for Covariate Balancing Propensity Score

Description

This data set gives the outcomes a well as treatment assignments and covariates for the econometric evaluation of training programs in LaLonde (1986).

Usage

LaLonde

Format

A data frame consisting of 5 columns (including a treatment assignment vector) and 2787 observations.

Source

Data from the National Supported Work Study. A benchmark matching dataset. Columns consist of an indicator for whether the observed unit was in the experimental subset; an indicator for whether the individual received the treatment; age in years; schooling in years; indicators for black and Hispanic; an indicator for marriage status, one of married; an indicator for no high school degree; and reported earnings in 1974, 1975, and 1978. 1974 and 1975 earnings are pre-treatment. 1978 earnings is taken as the outcome variable.

References

LaLonde, R.J. (1986). Evaluating the econometric evaulations of training programs with experimental data. American Economic Review 76, 4, 604-620.

8 summary.CBPS

summary.CBPS

Summarizing Covariate Balancing Propensity Score Estimation

Description

Prints a summary of a fitted CBPS object.

Usage

```
## S3 method for class 'CBPS'
summary(object,...)
```

Arguments

object an object of class "CBPSlogit", usually, a result of a call to CBPS.

... Additional arguments to be passed to summary.

Details

Prints a summmary of a CBPS object, in a format similar to glm. The variance matrix is calculated from the numerical Hessian at convergence of CBPS.

Value

call The matched call.

deviance.residuals

The five number summary and the mean of the deviance residuals.

coefficients A table including the estimate for the each coefficient and the standard error,

z-value, and two-sided p-value for these estimates.

J Hansen's J-Statistic for the fitted model.

Log-Likelihood The log-likelihood of the fitted model.

Author(s)

Marc Ratkovic

See Also

CBPS, summary

vcov.CBPS 9

vcov.CBPS

Calculate Variance-Covariance Matrix for a Fitted CBPS Object

Description

vcov. CBPS Returns the variance-covariance matrix of the main parameters of a fitted CBPS object.

Usage

```
## S3 method for class 'CBPS'
vcov(object,...)
```

Arguments

object An object of class formula (or one that can be coerced to that class): a symbolic

description of the model to be fitted.

... Additional arguments to be passed to vcov.CBPS

Details

This is the CBPS implementation of the generic function vcov().

Value

A matrix of the estimated covariances between the parameter estimates in the linear or non-linear predictor of the model.

Author(s)

Marc Ratkovic, Kosuke Imai, and Christian Fong.

References

This documentation is modeled on the documentation of the generic vcov.

See Also

vcov

Examples

```
###
### Example: Variance-Covariance Matrix
###

##Load the LaLonde data
data(LaLonde)
## Estimate CBPS via logistic regression
fit <- CBPS(treat ~ age + educ + re75 + re74 + I(re75==0) + I(re74==0), data = LaLonde, ATT = TRUE)</pre>
```

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Get the variance-covariance matrix.
vcov(fit)

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