

# Package ‘SDRcausal’

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**Title** SDRcausal

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**Description** Provides two semiparametric estimators, imp.ate and ipw.ate.

**Encoding** UTF-8

**Imports** stats, ggplot2

**Suggests** nloptr

**LazyData** true

**License** GPL (>= 2)

**RoxygenNote** 7.1.0

**NeedsCompilation** yes

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aipw.ate

*Combines IPW and IMP estimators to form the augmented IPW, AIPW***Description**

Augmented IPW (AIPW) as in Ghosh, Ma, & De Luna (2020).

**Usage**

```
aipw.ate(y, treated, imp, ipw)
```

**Arguments**

y	Observed response
treated	Binary vector indicating treatment
imp	imp_output object from imp.ate()
ipw	ipw_output object from ipw.ate()

**Value**

Average treatment effect (ATE) for the augmented IPW (AIPW)

**References**

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

**Examples**

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess
alp <- SDRcausal::alpha_guess

# Perform semiparametric imputation
imp <- SDRcausal::imp.ate(x, y, trt, b1, b0,
  explicit_bandwidth = TRUE, bwc_dim_red1 = 1, bwc_impute1 = 1,
  bwc_dim_red0 = 1, bwc_impute0 = 1)

# Perform semiparametric inverse probability weighting
ipw <- SDRcausal::ipw.ate(x, y, trt, alp, bwc_dim_red = 8,
  bwc_prop_score = 8)

# Calculate the Augmented IPW (AIPW)
aipw <- SDRcausal::aipw.ate(y, trt, imp, ipw)
```

---

aipw.var	<i>Estimates Augmented Inverse Probability variance</i>
----------	---

---

### Description

Variance of the Augmented IPW as in Ghosh, Ma, & De Luna (2020).

### Usage

```
aipw.var(
  x,
  y,
  treated,
  imp,
  ipw,
  bandwidth_scale1,
  bandwidth_scale0,
  bandwidth_scale_pr,
  kernel = "EPAN",
  explicit_bandwidth = TRUE,
  gauss_cutoff = 0.001,
  num_deriv_h = 1e-08,
  verbose = FALSE
)
```

### Arguments

x	Covariate matrix
y	Response vector
treated	Binary vetor indicating treatment
imp	imp_output object from imp.ate()
ipw	ipw_output object from ipw.ate()
bandwidth_scale1	Scaling of the calculated bandwidth, m1
bandwidth_scale0	Scaling of the calculated bandwidth, m0
bandwidth_scale_pr	Scaling of the calculated bandwidth, pr
kernel	Specifies which kernel function to be used
explicit_bandwidth	Specifies if bandwidth_scale will be used as the bandwidth or if it will be calculated as $bw = bandwidth\_scale * sd(x * beta) * n^{(1/5)}$ .
gauss_cutoff	Cutoff value for Gaussian kernel
num_deriv_h	Step size of numerical derivative.
verbose	Specifies if the program should print output while running.

### Value

The variance of Augmented IPW

## References

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

## Examples

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess
alp <- SDRcausal::alpha_guess

# Perform semiparametric imputation
imp <- SDRcausal::imp.ate(x, y, trt, b1, b0,
  explicit_bandwidth = TRUE, bwc_dim_red1 = 1, bwc_impute1 = 1,
  bwc_dim_red0 = 1, bwc_impute0 = 1)

# Perform semiparametric inverse probability weighting
ipw <- SDRcausal::ipw.ate(x, y, trt, alp, bwc_dim_red = 10,
  bwc_prop_score = 18)

# Calculate the variance of the Augmented IPW (AIPW)
var <- SDRcausal::aipw.var(x, y, trt, imp, ipw,
  bandwidth_scale1 = imp$bw1, bandwidth_scale0 = imp$bw0,
  bandwidth_scale_pr = ipw$bw_pr)
```

---

aipw2.ate

---

*Improved Augmented IPW (IAIPW)*


---

## Description

Combines IPW and IMP estimators to form the improved augmented IPW, IAIPW as in Ghosh, Ma, & De Luna (2020).

## Usage

```
aipw2.ate(y, treated, imp, ipw)
```

## Arguments

y	Observed response
treated	Binary vector indicating treatment
imp	imp_output object from imp.ate()
ipw	ipw_output object from ipw.ate()

**Value**

Average treatment effect (ATE) for the improved augmented IPW (IAIPW)

**References**

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

**Examples**

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess
alp <- SDRcausal::alpha_guess

# Perform semiparametric imputation
imp <- SDRcausal::imp.ate(x, y, trt, b1, b0,
  explicit_bandwidth = TRUE, bwc_dim_red1 = 1, bwc_impute1 = 1,
  bwc_dim_red0 = 1, bwc_impute0 = 1)

# Perform semiparametric inverse probability weighting
ipw <- SDRcausal::ipw.ate(x, y, trt, alp, bwc_dim_red = 10,
  bwc_prop_score = 18)

# Calculate the Improved Augmented IPW (AIPW2)
iaipw <- SDRcausal::aipw2.ate(y, trt, imp, ipw)
```

---

b10\_fun

*Calculates B1/0*


---

**Description**

Calculates Eq 2.8 or 2.10 in Ghosh, Ma, & De Luna (2020).

**Usage**

```
b10_fun(x, treated, dm, beta, kernel, bandwidth, gauss_cutoff)
```

**Arguments**

x	Projection of covariate matrix on CMS
treated	Binary vector indicating treatment.
dm	Derivative of imputed values
beta	CMS
kernel	Specifies which kernel function to be used
bandwidth	Specifies if bandwidth_scale will be used as the
gauss_cutoff	Cutoff value for Gaussian kernel

**Value**

B1/0 matrix

---

b_fun	<i>Calculates B1/0</i>
-------	------------------------

---

**Description**

Calculates Eq 2.8 or 2.10 in Ghosh, Ma, & De Luna (2020).

**Usage**

```
b_fun(x, treated, alpha_hat, h, kernel, bandwidth, bandwidth_pr, verbose)
```

**Arguments**

x	Projection of covariate matrix on CMS
treated	Treated
alpha_hat	Derivative of imputed values
h	CMS
kernel	Specifies which kernel function to be used
bandwidth	Kernel bandwidth
bandwidth_pr	Kernel bandwidth for probability
verbose	Specifies if the program should print output while running.

**Value**

B1/0 matrix

---

cms.ps.semi	<i>Estimates the Central Mean Space (CMS)</i>
-------------	---

---

**Description**

Semiparametric estimation of the Central Mean Space (CMS) as in Ghosh, Ma, & De Luna (2020).  
To be used with SDRcausal::ps.semi().

**Usage**

```

cms.ps.semi(
  x,
  treated,
  alpha_initial,
  solver = "optim",
  kernel = "EPAN",
  explicit_bandwidth = FALSE,
  bandwidth_scale = 1,
  gauss_cutoff = 0.001,
  penalty = 10,
  n_before_pen = 5,
  root_tol = 0.001,
  n_threads = 1,
  verbose = FALSE,
  ...
)

```

**Arguments**

x	Covariate matrix
treated	Binary vetor indicating treatment
alpha_initial	Initial guess of CMS
solver	Specifies which solver to be used. Current options optim and cobyla (from nloptr package).
kernel	Specifies which kernel function to be used, current options are: "EPAN", "QUARTIC", and "GAUSSIAN".
explicit_bandwidth	Specifies if bandwidth_scale will be used explicitly as the bandwidth.
bandwidth_scale	Scaling of the calculated bandwidth, or in case of explicit_bandwidth = TRUE the bandwidth.
gauss_cutoff	cutoff value for Gaussian kernel
penalty	Penalty for the optimizer if a probability is outside (0, 1). Added to the function value in optim as: $\text{penalty}^n$ , where n is the number of probabilities outside (0, 1).
n_before_pen	Number of probabilities outside the range (0, 1) to accept during dimension reduction.
root_tol	Tolerance which makes the program warn if optim stops at a value higher than root_tol.
n_threads	Sets number of threads for parallel run. Set to 0 serial. If n_threads exceeds maximum number of threads, sets n_threads to max_threads - 1. To use max_threads, set to n_threads to max_threads of system.
verbose	Specifies if the program should print output while running.
...	Additional parameters passed to solver.

**Value**

A list containing the final alpha, bandwwidth used, and the output of optim

## References

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

## See Also

[stats::optim]

## Examples

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
alp <- SDRcausal::alpha_guess

# Perform semiparametric dimension reduction for treated
cms <- SDRcausal::cms.ps.semi(x, trt, alp,
                             explicit_bandwidth = TRUE, bandwidth_scale = 8)
```

---

cms.semi

*Estimates the Central Mean Space (CMS)*

---

## Description

Semiparametric estimation of the Central Mean Space (CMS) as in Ghosh, Ma, & De Luna (2020). To be used with `SDRcausal::imp.val()`.

## Usage

```
cms.semi(
  x,
  y,
  treated,
  beta_initial,
  solver = "optim",
  kernel = "EPAN",
  explicit_bandwidth = FALSE,
  bandwidth_scale = 1,
  gauss_cutoff = 0.001,
  penalty = 10,
  n_before_pen = 1,
  root_tol = 0.001,
  n_threads = 1,
  verbose = FALSE,
  ...
)
```



**Arguments**

<code>x</code>	Covariate matrix
<code>y</code>	Response vector
<code>treated</code>	Binary vector indicating treatment
<code>beta_initial</code>	Initial guess of CMS
<code>solver</code>	Specifies which solver to be used. Current options <code>optim</code> and <code>cobyla</code> (from <code>nloptr</code> package).
<code>kernel</code>	Specifies which kernel function to be used, current options are: "EPAN", "QUARTIC", and "GAUSSIAN".
<code>explicit_bandwidth</code>	Specifies if <code>bandwidth_scale</code> will be used as the bandwidth or if it will be calculated as $bw = bandwidth\_scale * sd(x * beta) * n^{1/5}$
<code>bandwidth_scale</code>	Scaling of the bandwidth or the actual bandwidth if explicit bandwidth.
<code>gauss_cutoff</code>	cutoff value for Gaussian kernel
<code>penalty</code>	Penalty for the optimizer if local linear regression fails. Added to the function value in solver as: $penalty^{(n - n\_before\_pen)}$ , where <code>n</code> is the number of llr fails.
<code>n_before_pen</code>	Number of probabilities outside the range (0, 1) to accept during dimension reduction.
<code>root_tol</code>	Tolerance which makes the program warn if <code>optim</code> stops at a value higher than <code>root_tol</code> .
<code>n_threads</code>	Sets number of threads for parallel run. Set to 0 serial. If <code>n_threads</code> exceeds maximum number of threads, sets <code>n_threads</code> to <code>max_threads - 1</code> . To use <code>max_threads</code> , set to <code>n_threads</code> to <code>max_threads</code> of system.
<code>verbose</code>	Specifies if the program should print output while running.
<code>...</code>	Additional parameters passed to <code>optim</code> .

**Value**

A list containing the final beta, the bandwidth used, a warning if `optim` does not converge or converges to a value that is larger than `root_tol`, and the output of `optim`.

**References**

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

**See Also**

[stats::optim]

**Examples**

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
```

```

trt1 <- SDRcausal::treated
trt0 <- rep(1, length(trt1)) - trt1
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess

# Perform semiparametric dimension reduction for treated
cms1 <- SDRcausal::cms.semi(x, y, trt1, b1,
  explicit_bandwidth = TRUE, bandwidth_scale = 1)

# Perform semiparametric dimension reduction for untreated
cms0 <- SDRcausal::cms.semi(x, y, trt0, b0,
  explicit_bandwidth = TRUE, bandwidth_scale = 1)

```

---

example_data	<i>Example data</i>
--------------	---------------------

---

### Description

Data generated as in paper, study 1. Using the betas in betas data. Use beta1/0 for imputation as the initial guess of the central mean space (CMS) and alpha as the initial guess of the CMS for IPW.

### Format

Data used in examples of the SDRcausal package

**covariates** covariate matrix

**outcomes** observed outcome vector

**treated** binary treatment vector

**beta1\_guess** Starting guess for CMS for treated

**beta0\_guess** Starting guess for CMS for untreated

**alpha\_guess** Starting guess for CMS for propensity score

---

imp.ate	<i>Estimates Average Treatment Effect (ATE) by imputation (IMP)</i>
---------	---

---

### Description

Semiparametric estimation of the average treatment effect based on the imputation method described in Ghosh, Ma, & De Luna (2020).

**Usage**

```

imp.ate(
  x,
  y,
  treated1,
  beta_guess1,
  beta_guess0,
  solver = "optim",
  kernel = "EPAN",
  explicit_bandwidth = FALSE,
  recalc_bandwidth = FALSE,
  bwc_dim_red1 = 1,
  bwc_impute1 = 1,
  bwc_dim_red0 = 1,
  bwc_impute0 = 1,
  gauss_cutoff = 0.001,
  penalty = 10,
  n_before_pen = 5,
  to_extrapolate = TRUE,
  to_truncate = TRUE,
  extrapolation_basis = 5,
  n_threads = 1,
  verbose = TRUE,
  ...
)

```

**Arguments**

x	Covariate matrix
y	Response vector
treated1	Binary vector indicating treatment.
beta_guess1	Initial guess of beta for m1
beta_guess0	Initial guess of beta for m0
solver	Specifies which solver to be used. Current options optim and cobyla (from nloptr package).
kernel	Specifies which kernel function to be used, current options are: "EPAN", "QUARTIC", and "GAUSSIAN".
explicit_bandwidth	Specifies if bandwidth_scale will be used as the bandwidth or if it will be calculated as $bw = \text{bandwidth\_scale} * \text{sd}(x * \text{beta}) * n^{(1/3)}$ .
recalc_bandwidth	Specifies whether the bandwidth should be recalculated after the estimation of alpha (cms.ps.semi).
bwc_dim_red1	Scaling of calculated bandwidth, or if explicit_bandwidth = TRUE used as the bandwidth. For dimension reduction (cms.semi).
bwc_impute1	Scaling of calculated bandwidth, or if explicit_bandwidth = TRUE used as the bandwidth. Recalculated if explicit_bandwidth = FALSE and recalc_bandwidth = TRUE. For imputation.
bwc_dim_red0	See bwc_dim_red1

bwc_impute0	See bwc_impute1
gauss_cutoff	Cutoff value for Gaussian kernel
penalty	Penalty for the optimizer if local linear regression fails. Added to the function value in solver as: $\text{penalty}^{(n - n\_before\_pen)}$ , where $n$ is the number of llr fails.
n_before_pen	Number of probabilities outside the range (0, 1) to accept during dimension reduction.
to_extrapolate	Specifies wheter to extrapolate or not
to_truncate	Specifies wheter to extrapolate or not
extrapolation_basis	Number of data point to base extrapolation on.
n_threads	Sets number of threads for parallel run. Set to 0 serial. If n_threads exceeds maximum number of threads, sets n_threads to max_threads - 1. To use max_threads, set to n_threads to max_threads of system.
verbose	Specifies if the program should print output while running.
...	Additional parameters passed to optim.

### Value

A list containing the average treatment effect of the combination of observed and imputed values (ate), the average treatment effect based on the imputed values only (ate2), the imputed values for treated (m1) and untreated treated (m0), the and the output from optim (op).

### References

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

### See Also

[stats::optim]

### Examples

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess

# Perform semiparametric imputation
imp <- SDRcausal::imp.ate(x, y, trt, b1, b0,
  explicit_bandwidth = TRUE, bwc_dim_red1 = 1, bwc_impute1 = 1,
  bwc_dim_red0 = 1, bwc_impute0 = 1)
```

imp.val

*Estimates imputed values based on CMS***Description**

Performs semiparametric imputation based on the CMS calculated by cms.semi, as in Ghosh, Ma, & De Luna (2020).

**Usage**

```
imp.val(
  x,
  y,
  treated,
  beta_hat,
  kernel = "EPAN",
  explicit_bandwidth = FALSE,
  bandwidth_scale = 1,
  gauss_cutoff = 0.001,
  to_extrapolate = TRUE,
  to_truncate = TRUE,
  extrapolation_basis = as.integer(5),
  verbose = FALSE
)
```

**Arguments**

x	Covariate matrix
y	Response vector
treated	Binary vetor indicating treatment
beta_hat	Locally efficient CMS
kernel	Specifies which kernel function to be used
explicit_bandwidth	Specifies if bandwidth_scale will be used as the bandwidth or if it will be calculated as $bw = bandwidth\_scale * sd(x * beta) * n^{(1/3)}$
bandwidth_scale	Kernel bandwidth
gauss_cutoff	Cutoff value for Gaussian kernel
to_extrapolate	Specifies wheter to extrapolate or not
to_truncate	Specifies wheter to extrapolate or not
extrapolation_basis	Number of data point to base extrapolation on.
verbose	Specifies if the program should print output while running

**Value**

A list containing the reduced space xb, the imputed values and their derivatives.

## References

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

## Examples

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess

# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt1 <- SDRcausal::treated
n <- as.integer(dim(x)[1])
trt0 <- as.integer(rep(1, times = n) - trt1)
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess

# Perform semiparametric dimension reduction for treated
cms1 <- SDRcausal::cms.semi(x, y, trt1, b1,
  explicit_bandwidth = 1, bandwidth_scale = 1)

# Perform semiparametric dimension reduction for untreated
cms0 <- SDRcausal::cms.semi(x, y, trt0, b0,
  explicit_bandwidth = 1, bandwidth_scale = 1)

# Perform semiparametric imputation for treated
m1 <- SDRcausal::imp.val(x, y, trt1, cms1$fb,
  explicit_bandwidth = 1, bandwidth_scale = cms1$bw)

# Perform semiparametric imputation for untreated
m0 <- SDRcausal::imp.val(x, y, trt0, cms0$fb,
  explicit_bandwidth = 1, bandwidth_scale = cms0$bw)
```

---

imp.var

*Estimates IMP variance*

---

## Description

Variance of the IMP as in Ghosh, Ma, & De Luna (2020).

**Usage**

```
imp.var(
  x,
  y,
  treated,
  imp,
  ipw,
  bandwidth_scale1,
  bandwidth_scale0,
  kernel = "EPAN",
  explicit_bandwidth = TRUE,
  gauss_cutoff = 0.001
)
```

**Arguments**

x	Covariate matrix
y	Response vector
treated	Binary vector indicating treatment
imp	imp_output object from imp.ate()
ipw	ipw_output object from ipw.ate()
bandwidth_scale1	Scaling of the calculated bandwidth, or in case explicit_bandwidth the actual bandwidth. For m1 and beta1.
bandwidth_scale0	See bandwidth_scale1. For m0 and beta0.
kernel	Specifies which kernel function to be used
explicit_bandwidth	Specifies if bandwidth_scale will be used as the bandwidth or if it will be calculated as $bw = bandwidth\_scale * sd(x * beta) * n^{(1/3)}$ .
gauss_cutoff	Cutoff value for Gaussian kernel

**Value**

Variance of IMP

**References**

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

**Examples**

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
```

```

b0 <- SDRcausal::beta0_guess
alp <- SDRcausal::alpha_guess

# Perform semiparametric imputation
imp <- SDRcausal::imp.ate(x, y, trt, b1, b0,
  explicit_bandwidth = TRUE, bwc_dim_red1 = 1, bwc_impute1 = 1,
  bwc_dim_red0 = 1, bwc_impute0 = 1)

# Perform semiparametric inverse probability weighting
ipw <- SDRcausal::ipw.ate(x, y, trt, alp, bwc_dim_red = 10,
  bwc_prop_score = 18)

# Calculate the variance of the Augmented IPW (AIPW)
var <- SDRcausal::imp.var(x, y, trt, imp, ipw,
  bandwidth_scale1 = imp$bw1, bandwidth_scale0 = imp$bw0)

```

imp2.var

*Estimates IMP2 variance***Description**

Variance of IMP2 as in Ghosh, Ma, & De Luna (2020).

**Usage**

```

imp2.var(
  x,
  y,
  treated,
  imp,
  ipw,
  bandwidth_scale1,
  bandwidth_scale0,
  kernel = "EPAN",
  explicit_bandwidth = TRUE,
  gauss_cutoff = 0.001
)

```

**Arguments**

x	Covariate matrix
y	Response vector
treated	Binary vetor indicating treatment
imp	imp_output object from imp.ate()
ipw	ipw_output object from ipw.ate()
bandwidth_scale1	Scaling of the calculated bandwidth, or in case explicit_bandwidth the actual bandwidth. For m1 and beta1.
bandwidth_scale0	See bandwidth_scale1. For m0 and beta0.



kernel                Specifies which kernel function to be used

explicit\_bandwidth        Specifies if bandwidth\_scale will be used as the bandwidth or if it will be calculated as  $bw = \text{bandwidth\_scale} * \text{sd}(x * \text{beta}) * n^{(1/3)}$ .

gauss\_cutoff        Cutoff value for Gaussian kernel

## Value

Variance of IMP

## References

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

## Examples

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess
alp <- SDRcausal::alpha_guess

# Perform semiparametric imputation
imp <- SDRcausal::imp.ate(x, y, trt, b1, b0,
  explicit_bandwidth = TRUE, bwc_dim_red1 = 1, bwc_impute1 = 1,
  bwc_dim_red0 = 1, bwc_impute0 = 1)

# Perform semiparametric inverse probability weighting
ipw <- SDRcausal::ipw.ate(x, y, trt, alp, bwc_dim_red = 10,
  bwc_prop_score = 18)

# Calculate the variance of the Augmented IPW (AIPW)
var <- SDRcausal::imp2.var(x, y, trt, imp, ipw,
  bandwidth_scale1 = imp$bw1, bandwidth_scale0 = imp$bw0)
```

---

ipw.ate

*Estimates average treatment effect through IPW*

---

## Description

Semiparametric estimation of the average treatment effect based on the IPW method described in Ghosh, Ma, & De Luna (2020).

**Usage**

```

ipw.ate(
  x,
  y,
  treated,
  alpha_initial,
  kernel = "EPAN",
  explicit_bandwidth = FALSE,
  recalc_bandwidth = TRUE,
  bwc_dim_red = 1,
  bwc_prop_score = 10,
  gauss_cutoff = 0.001,
  penalty = 10,
  n_before_pen = 1,
  n_threads = 1,
  verbose = TRUE,
  ...
)

```

**Arguments**

<code>x</code>	Covariate matrix
<code>y</code>	Response vector
<code>treated</code>	Binary vector indicating treatment.
<code>alpha_initial</code>	Initial guess of beta for ml
<code>kernel</code>	Specifies which kernel function to be used, current options are: "EPAN", "QUARTIC", and "GAUSSIAN".
<code>explicit_bandwidth</code>	Specifies if <code>bandwidth_scale</code> will be used as the bandwidth or if it will be calculated as $bw = bandwidth\_scale * sd(x * beta) * n^{(1/3)}$ .
<code>recalc_bandwidth</code>	Specifies wheter the bandwidth should be recalculated after the estimation of alpha (cms.ps.semi)
<code>bwc_dim_red</code>	Scaling of calculated bandwidth, or if <code>explicit_bandwidth = TRUE</code> used as the banddwidth. For dimension reduction (cms.ps.semi).
<code>bwc_prop_score</code>	Scaling of calculated bandwidth, or if <code>explicit_bandwidth = TRUE</code> used as the banddwidth. Recalculated if <code>explicit_bandwidth = FALSE</code> and <code>recalc_bandwidth = TRUE</code> . For propensity score.
<code>gauss_cutoff</code>	cutoff value for Gaussian kernel
<code>penalty</code>	Penalty for the optimizer if a probability is outside (0, 1) during dimension reduction. Added to the function value in solver as: $penalty^{(n - n\_before\_pen)}$ , where $n$ is the number of probabilities outside (0, 1).
<code>n_before_pen</code>	Number of probabilities outside the range (0, 1) to accept during dimension reduction.
<code>n_threads</code>	Sets number of threads for parallel run. Set to 0 serial. If <code>n_threads</code> exceeds maximum number of threads, sets <code>n_threads</code> to <code>max_threads - 1</code> . To use <code>max_threads</code> , set to <code>n_threads</code> to <code>max_threads</code> of system.
<code>verbose</code>	Specifies if the program should print output while running.
<code>...</code>	Additional parameters passed to <code>optim</code> .

**Value**

A list containing the average treatment effect (ate), the propensity score (pr), the final alpha (fa), and the output from optim (op).

**References**

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

**See Also**

[stats::optim]

**Examples**

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
alp <- SDRcausal::alpha_guess

# Perform semiparametric inverse probability weighting
ipw <- SDRcausal::ipw.ate(x, y, trt, alp, bwc_dim_red = 8,
                          bwc_prop_score = 8)
```

---

ipw.var

---

*Estimates IPW variance*


---

**Description**

Variance of the IPW as in Ghosh, Ma, & De Luna (2020).

**Usage**

```
ipw.var(
  x,
  y,
  treated,
  imp,
  ipw,
  bandwidth_scale,
  kernel = "EPAN",
  explicit_bandwidth = TRUE,
  gauss_cutoff = 0.001,
  num_deriv_h = 0.001,
  verbose = FALSE
)
```

**Arguments**

x	Covariate matrix
y	Response vector
treated	Binary vector indicating treatment
imp	imp_output object from imp.ate()
ipw	ipw_output object from ipw.ate()
bandwidth_scale	Scaling of the calculated bandwidth, or in case of
kernel	Specifies which kernel function to be used
explicit_bandwidth	Specifies if bandwidth_scale will be used as the bandwidth or if it will be calculated as $bw = bandwidth\_scale * sd(x * beta) * n^{(1/3)}$ .
gauss_cutoff	Cutoff value for Gaussian kernel
num_deriv_h	Step size of numerical derivative.
verbose	Specifies if the program should print output while running.

**Value**

The variance of IPW

**References**

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

**Examples**

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess
alp <- SDRcausal::alpha_guess

# Perform semiparametric imputation
imp <- SDRcausal::imp.ate(x, y, trt, b1, b0,
  explicit_bandwidth = TRUE, bwc_dim_red1 = 1, bwc_impute1 = 1,
  bwc_dim_red0 = 1, bwc_impute0 = 1)

# Perform semiparametric inverse probability weighting
ipw <- SDRcausal::ipw.ate(x, y, trt, alp, bwc_dim_red = 10,
  bwc_prop_score = 18)

# Calculate the variance of the Augmented IPW (AIPW)
var <- SDRcausal::ipw.var(x, y, trt, imp, ipw,
  bandwidth_scale = ipw$bw_pr)
```

---

nw_kernel_regress	<i>The Nadaraya-Watson kernel estimator</i>
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---

## Description

Gives the expected value of  $Y$  given  $X = x$  by kernel regression according to the Nadaraya-Watson kernel estimator to get  $E(Y|X)$ . Note that  $y$  and  $x$  may be vectors or matrices, as long as  $\dim(x)[1] == \dim(y)[1]$ .

## Usage

```
nw_kernel_regress(
  y,
  x,
  bandwidth = 1,
  kernel = "EPAN",
  gauss_cutoff = 0.001,
  verbose = FALSE
)
```

## Arguments

<code>y</code>	$Y$ in $E(Y X)$
<code>x</code>	$X$ in $E(Y X)$
<code>bandwidth</code>	Kernel bandwidth
<code>kernel</code>	Indicates which kernel function to be used
<code>gauss_cutoff</code>	Cutoff value for Gaussian kernel
<code>verbose</code>	Specifies if the program should print output while running.

## Value

Value of kernel regression

## References

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

## Examples

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
x <- SDRcausal::covariates
y <- SDRcausal::outcomes

# Estimating y given x,  $E(y | x)$ 
k <- nw_kernel_regress(y, x, bandwidth = 1)
```

plot.imp

*Plots imputation output***Description**

Plot function for visualisation of imputation output from imp.ate. Note: The function requires ggplot2.

**Usage**

```
## S3 method for class 'imp'
plot(x, ..., covariates, y, treated)
```

**Arguments**

x	imp_output object from imp.ate()
...	Other parameters
covariates	Covariate matrix
y	Response vector
treated	Binary vetor indicating treatment

**Value**

A list of ggplot plots of observed and imputed values (pl\_imp), imputed treated values vs CMS (pl\_m1), and imputed untreated values vs CMS (pl\_m0).

**Examples**

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
covariates <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
b1 <- SDRcausal::beta1_guess
b0 <- SDRcausal::beta0_guess
alp <- SDRcausal::alpha_guess

# Perform semiparametric imputation
imp <- SDRcausal::imp.ate(covariates, y, trt, b1, b0,
  explicit_bandwidth = TRUE, bwc_dim_red1 = 1, bwc_impute1 = 1,
  bwc_dim_red0 = 1, bwc_impute0 = 1)

# Plotting
plots <- plot(imp , covariates = covariates, y=y, treated = trt)
```

---

plot.ipw	<i>Plots IPW output</i>
----------	-------------------------

---

## Description

Plot function for visualisation of IPW output from ipw.ate. Note: The function requires ggplot2.

## Usage

```
## S3 method for class 'ipw'  
plot(x, ..., treated, covariates)
```

## Arguments

x	ipw_output object from ipw.ate()
...	Other parameters
treated	Binary vetor indicating treatment
covariates	Covariate matrix

## Value

ggplot plot of the propensity score vs CMS.

## Examples

```
# Using example data from package SDRcausal  
library(SDRcausal)  
  
# Import example data  
covariates <- SDRcausal::covariates  
y <- SDRcausal::outcomes  
trt <- SDRcausal::treated  
b1 <- SDRcausal::beta1_guess  
b0 <- SDRcausal::beta0_guess  
alp <- SDRcausal::alpha_guess  
  
# Perform semiparametric imputation  
ipw <- SDRcausal::ipw.ate(covariates, y, trt, alp, bwc_dim_red = 8,  
  bwc_prop_score = 8)  
  
# Plotting  
plots <- plot(ipw, treated = trt, covariates = covariates)
```

---

ps.semi	<i>Estimates propensity score</i>
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---

### Description

Semiparametric estimation of the propensity score as in Ghosh, Ma, & De Luna (2020). To be used with `SDRcausal::cms.ps.semi()`.

### Usage

```
ps.semi(
  x,
  treated,
  alpha_hat,
  kernel = "EPAN",
  explicit_bandwidth = FALSE,
  bandwidth_scale = 1,
  verbose = FALSE
)
```

### Arguments

x	Covariate matrix
treated	Binary vetor indicating treatment
alpha_hat	Locally efficient CMS
kernel	Kernel specification
explicit_bandwidth	Specifies if bandwidth_scale will be used as the bandwidth or if it will be calculated as $bw = bandwidth\_scale * sd(x * beta) * n^{(1/3)}$ .
bandwidth_scale	Scaling of calculated bandwidth, or if explicit_bandwidth = TRUE used as the banddwidth.
verbose	Specifies if the program should print output while running.

### Value

A list containing the estimated propensity scores values and their derivatives, and the bandwidth used.

### References

Ghosh, T., Ma, Y., & De Luna, X. (2020). Sufficient dimension reduction for feasible and robust estimation of average causal effect. *Statistica Sinica*, accepted.

### Examples

```
# Using example data from package SDRcausal
library(SDRcausal)

# Import example data
```



```
x <- SDRcausal::covariates
y <- SDRcausal::outcomes
trt <- SDRcausal::treated
alp <- SDRcausal::alpha_guess

# Perform semiparametric dimension reduction
cms <- SDRcausal::cms.ps.semi(x, trt, alp,
  explicit_bandwidth = TRUE, bandwidth_scale = 8)

# Estimate propensity score
pr_score <- SDRcausal::ps.semi(x, trt, cms$fa,
  bandwidth_scale = 8)
```

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