

Package: PartialTL (via r-universe)

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Title Partial Transfer Learning for Causal Estimation

Version 0.1.0

Description Implements partial transfer learning (PTL) for causal effect estimation using source and target data, with bootstrap-based source detection. Provides data generating processes and nuisance functions for simulation.

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URL <https://github.com/JackQuu/PartialTL>

BugReports <https://github.com/JackQuu/PartialTL/issues>

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Imports MASS, DoubleML, ncvreg

Suggests lgr, mlr3, mlr3learners, glmnet

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PartialTL-package *Partial Transfer Learning for Causal Estimation and Source Detection*

Description

Implements partial transfer learning (PTL) and heterogeneous partial transfer learning (HPTL) for causal effect estimation using source and target data. Uses double machine learning for nuisance estimation and cross-fitting. Provides bootstrap-based source detection to identify which sources are transferable to the target. Includes data generating processes and nuisance functions for simulation.

Details

Main functions:

- `fit_PTL`: PTL causal estimate (single source).
- `fit_HPTL`: Heterogeneous PTL with multiple sources and covariate modules.
- `boot_detection`: Bootstrap-based source detection.
- `run_bootstrap_E_hat`: Bootstrap for nuisance function estimation.
- `DGP`: Data generating process for simulations.
- `RMSE`: Root mean squared error for evaluation.
- `f_0`, `g_0`, `f_k`, `g_k`: Nuisance functions for target and source models.

Run `demo(package = "PartialTL")` for demo scripts.

Author(s)

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References

(Add references to the partial transfer learning and DML literature.)

boot_detection	<i>Bootstrap-Based Source Detection</i>
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Description

Identifies which sources are transferable to the target by comparing bootstrap estimates of the nuisance function on the target and each source. A source is detected as transferable if the difference is below a threshold proportional to the combined standard error.

Usage

```
boot_detection(D_t, X_t, Y_t, D_s_all, X_s_all, Y_s_all, source_sizes, B,
              ml_f, ml_g)
```

Arguments

D_t	Target treatment; $n_t \times 1$.
X_t	Target design matrix; $n_t \times p$.
Y_t	Target outcome; $n_t \times 1$.
D_s_all	Source treatments concatenated by row; rows split by source_sizes.
X_s_all	Source design matrices concatenated by row.
Y_s_all	Source outcomes concatenated by row.
source_sizes	Integer vector of length K: sample size of each source.
B	Number of bootstrap replications.
ml_f	Outcome learner for DoubleML.
ml_g	Treatment learner for DoubleML.

Value

A data frame with columns Source (label) and detected.source (list of logical vectors of length B per source).

See Also

[run_bootstrap_E_hat](#), [fit_PTL](#), [fit_HPTL](#)

Description

Generates (X, D, Y) for simulations: multivariate normal X with AR(1)-type covariance, treatment D from a linear confounding model plus noise, and outcome $Y = \rho * D + f(X) + \text{error}$.

Usage

```
DGP(n, q, p, p.nonzero, rho, Beta, Gamma, mu = rep(10, p), sigma = 0.5,
    f_func = f_0, g_func = g_0, seed = NULL)
```

Arguments

<code>n</code>	Sample size.
<code>q</code>	Causal dimension (kept for interface; unused in current body).
<code>p</code>	Nuisance dimension (number of covariates).
<code>p.nonzero</code>	Number of non-zero coefficients for Beta and Gamma.
<code>rho</code>	Causal effect (scalar).
<code>Beta</code>	Nuisance coefficient vector for outcome; $p \times 1$.
<code>Gamma</code>	Nuisance coefficient vector for treatment; $p \times 1$.
<code>mu</code>	Mean vector of X ; length p .
<code>sigma</code>	Base for AR(1)-type covariance: $\Sigma_{i,j} = \sigma^{ i-j }$.
<code>f_func</code>	Function $f(X, \text{Beta})$ for outcome nuisance; default <code>f_0</code> .
<code>g_func</code>	Function $g(X, \text{Gamma})$ for treatment equation; default <code>g_0</code> .
<code>seed</code>	Optional random seed.

Value

A list with components `D` (treatment), `X` (design matrix), `Y` (outcome), and `data` (`cbind(Y, D, X)`).

See Also

[f_0](#), [g_0](#), [f_k](#), [g_k](#), [fit_PTL](#), [RMSE](#)

Examples

```
set.seed(1)
n <- 50
p <- 10
p.nz <- 3
Beta <- matrix(c(rep(0.5, p.nz), rep(0, p - p.nz)))
Gamma <- matrix(c(rep(0.3, p.nz), rep(0, p - p.nz)))
dat <- DGP(n, q = 1, p, p.nz, rho = 0.5, Beta, Gamma, seed = 1)
str(dat)
```

f_0 *Outcome Nuisance Function (Target Model)*

Description

Linear outcome nuisance function for the target model: $f_0(X, \beta) = X\beta$.

Usage

f_0(X, Beta)

Arguments

X Design matrix; $n \times p$.
Beta Coefficient vector (column matrix); $p \times 1$.

Value

Numeric vector of outcome nuisance values; length n.

See Also

[g_0](#), [f_k](#), [g_k](#), [DGP](#)

f_k *Outcome Nuisance Function (Source Model)*

Description

Linear outcome nuisance function for source models: $f_k(X, \beta) = X\beta$. Same form as f_0; used in DGP for source data.

Usage

f_k(X, Beta)

Arguments

X Design matrix; $n \times p$.
Beta Coefficient vector (column matrix); $p \times 1$.

Value

Numeric vector of outcome nuisance values; length n.

See Also

[f_0](#), [g_k](#), [DGP](#)

fit_HPTL

HPTL (Heterogeneous Partial Transfer Learning) Fit

Description

Fits heterogeneous partial transfer learning with multiple sources and covariate modules. Each source has its own module of covariates; PTL is applied per source and results are combined for the target causal estimate.

Usage

```
fit_HPTL(D_t, X_t, Y_t, D_s_all, X_s_all, Y_s_all, source_sizes, module_sizes,
         ml_f, ml_g, fold = 5)
```

Arguments

D_t	Target treatment; $n_t \times q$ matrix.
X_t	Target design matrix; $n_t \times p$ (same covariates as sources).
Y_t	Target outcome; $n_t \times 1$.
D_s_all	Source treatments concatenated by row; dimension is (sum of source_sizes) by q .
X_s_all	Source design matrices concatenated by row; rows split by source_sizes.
Y_s_all	Source outcomes concatenated by row; rows split by source_sizes.
source_sizes	Integer vector of length K : sample size of each source. Must sum to $nrow(Y_s_all)$.
module_sizes	Integer vector of length K : covariate module sizes. The k -th source uses columns $(cumsum(module_sizes)[k-1]+1) : cumsum(module_sizes)[k]$ of X .
ml_f	Outcome learner for DoubleML.
ml_g	Treatment learner for DoubleML.
fold	Number of folds for cross-fitting (default 5).

Value

A list with component `hat_rho_HPTL`: HPTL causal estimate on target.

See Also

[fit_PTL](#), [boot_detection](#)

fit_PTL

*PTL (Partial Transfer Learning) Fit***Description**

Fits partial transfer learning with cross-fitting and outcome reconstruction. Uses double machine learning on the source and SCAD on the outcome nuisance, then transfers to the target via the partial transfer formula.

Usage

```
fit_PTL(D_t, X_t, Y_t, D_s, X_s, Y_s, ml_f, ml_g, fold = 5L)
```

Arguments

D_t	Target treatment; $n_t \times 1$ matrix.
X_t	Target design matrix; $n_t \times p$.
Y_t	Target outcome; $n_t \times 1$.
D_s	Source treatment; $n_s \times 1$.
X_s	Source design matrix; $n_s \times p$.
Y_s	Source outcome; $n_s \times 1$.
ml_f	Outcome learner for DoubleML (e.g. <code>lrm("regr.cv_glmnet")</code>).
ml_g	Treatment learner for DoubleML (same type).
fold	Number of folds for cross-fitting (default 5).

Value

A list with components:

hat_rho_s	Source causal estimate.
beta_hat_s	Source nuisance coefficient estimate.
E_s	Estimated $E[Y - \rho D]$ on source (NA when $q \neq 1$).
hat_rho_PTL	PTL causal estimate on target.

See Also

[fit_HPTL](#), [DGP](#), [RMSE](#)

g_0 *Treatment Nuisance Function (Target Model)*

Description

Linear treatment/confounding function for the target model: $g_0(X, \gamma) = X\gamma$.

Usage

`g_0(X, Gamma)`

Arguments

X Design matrix; $n \times p$.
Gamma Coefficient vector (column matrix); $p \times 1$.

Value

Numeric vector of treatment equation values; length n.

See Also

[f_0](#), [f_k](#), [g_k](#), [DGP](#)

g_k *Treatment Nuisance Function (Source Model)*

Description

Linear treatment/confounding function for source models: $g_k(X, \gamma) = X\gamma$. Same form as `g_0`; used in `DGP` for source data.

Usage

`g_k(X, Gamma)`

Arguments

X Design matrix; $n \times p$.
Gamma Coefficient vector (column matrix); $p \times 1$.

Value

Numeric vector of treatment equation values; length n.

See Also

[g_0](#), [f_k](#), [DGP](#)

RMSE	<i>Root Mean Squared Error</i>
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Description

Computes the root mean squared error of parameter estimates across simulations: for each component, $\sqrt{\text{mean}((\hat{\theta} - \theta)^2)}$.

Usage

```
RMSE(Theta, Theta.hat)
```

Arguments

Theta	True parameter vector; $p \times 1$.
Theta.hat	Matrix of estimates; $p \times n_{sim}$.

Value

Numeric vector of length p: RMSE per parameter component.

See Also

[DGP](#), [fit_PTL](#), [fit_HPTL](#)

Examples

```
th <- c(1, 2)
th_hat <- matrix(c(1.1, 2.2, 0.9, 1.8), nrow = 2)
RMSE(th, th_hat)
```

run_bootstrap_E_hat	<i>Bootstrap for Nuisance Function Estimation</i>
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Description

Runs bootstrap replications to estimate the nuisance quantity $E[Y - D\theta]$ using double machine learning, and returns the bootstrap distribution (mean and variance) of the estimate.

Usage

```
run_bootstrap_E_hat(D, X, Y, B, ml_f, ml_g)
```

Arguments

D	Treatment variable(s); matrix.
X	Design matrix.
Y	Outcome variable.
B	Number of bootstrap replications.
m1_f	Outcome learner for DoubleML.
m1_g	Treatment learner for DoubleML.

Value

A list with components:

E_hat	Numeric vector of length B: bootstrap estimates.
E_hat_mean	Mean of E_hat.
E_hat_var	Variance of E_hat.

See Also

[boot_detection](#)

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