

Package ‘obshetsurr’

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Type Package

Title Assessing Surrogate Heterogeneity in Observational Data

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Description Provides functions to estimate the proportion of the treatment effect explained (PTE) for a surrogate as a function of multiple baseline covariates in an observational setting, with options for different base learners. More details will be available in the future in: Knowlton, R., Parast, L. (2025) ``Assessing Surrogate Heterogeneity in Real World Data Using Meta-Learners.''

License GPL

Imports stats, matrixStats, mgcv, grf

NeedsCompilation no

Depends R (>= 3.5.0)

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boot.var	<i>Calculate bootstrapped variance estimates.</i>
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Description

Calculates bootstrapped variance estimates of delta, delta.s, and R.s, and optionally calculates p-values for identifying strong surrogates.

Usage

```
boot.var(df.train, df.test, type, numeric_predictors, categorical_predictors,
         threshold, use.actual.control.S)
```

Arguments

`df.train` A dataframe containing training data.

`df.test` A dataframe containing testing data.

`type` Options are "linear", "gam", "trees", or "all"; type of base learners to use.

`numeric_predictors` The column names in the dataframes that represent numeric baseline covariates.

`categorical_predictors` The column names in the dataframes that represent categorical baseline covariates.

`threshold` An optional threshold to test individuals for the null hypothesis that $PTE > threshold$.

`use.actual.control.S` TRUE or FALSE, if user prefers to use the actual observed values for the surrogate in the control group instead of predicting values from the base learners.

Value

A dataframe is returned, which is the `df.test` argument with new columns appended for the estimated variances of δ , $\delta.s$, and $R.s$, as well as p-values if a threshold is provided.

<code>estimate.PTE</code>	<i>Estimate the proportion of the treatment effect explained</i>
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Description

Fits base learners using the specified type of model on the training data, and uses those models to calculate δ , $\delta.s$, and $R.s$ on the testing set.

Usage

```
estimate.PTE(df.train, df.test, type, numeric_predictors, categorical_predictors,
             use.actual.control.S)
```

Arguments

`df.train` A dataframe containing training data.

`df.test` A dataframe containing testing data.

`type` Options are "linear", "gam", "trees", or "all"; type of base learners to use.

`numeric_predictors` The column names in the dataframes that represent numeric baseline covariates.

`categorical_predictors` The column names in the dataframes that represent categorical baseline covariates.

`use.actual.control.S` TRUE or FALSE, if user prefers to use the actual observed values for the surrogate in the control group instead of predicting values from the base learners.

Value

A dataframe is returned, which is the `df.test` argument with new columns appended for the estimates of `delta`, `delta.s`, and `R.s`.

exampledata_test	<i>Example testing data</i>
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Description

Example testing data

Usage

```
data("exampledata_test")
```

Format

A data frame with 200 observations on the following 9 variables.

X1 a numeric baseline covariate of interest

X2 a numeric baseline covariate of interest

X3 a numeric baseline covariate of interest

X4 a numeric baseline covariate of interest

X5 a numeric baseline covariate of interest

X6 a numeric baseline covariate of interest

G the treatment assignment, where 1 indicates treated and 0 indicates control

S the surrogate marker

Y the primary outcome

Examples

```
data(exampledata_test)
names(exampledata_test)
```

exampledata_train	<i>Example training data</i>
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Description

Example training data

Usage

```
data("exampledata_train")
```

Format

A data frame with 1800 observations on the following 9 variables.

X1 a numeric baseline covariate of interest
 X2 a numeric baseline covariate of interest
 X3 a numeric baseline covariate of interest
 X4 a numeric baseline covariate of interest
 X5 a numeric baseline covariate of interest
 X6 a numeric baseline covariate of interest
 G the treatment assignment, where 1 indicates treated and 0 indicates control
 S the surrogate marker
 Y the primary outcome

Examples

```
data(exampledata_train)
names(exampledata_train)
```

obs.het.surr	<i>Estimate the proportion of the treatment effect explained by the surrogate marker as a function of multiple baseline covariates in an observational setting</i>
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Description

Assesses surrogate heterogeneity in real world data by estimating the proportion of the treatment effect explained as a function of baseline covariates. Optionally tests individuals for strong surrogacy based on a threshold.

Usage

```
obs.het.surr(df.train, df.test, type, var.want = FALSE, threshold = NULL,
  use.actual.control.S = FALSE)
```

Arguments

df.train	A dataframe containing training data. Must have columns G (treatment assignment), S (surrogate marker), and Y (primary outcome), in addition to the baseline covariates of interest.
df.test	A dataframe containing testing data. Must contain the same baseline covariate columns as the training data.
type	Options are "linear", "gam", "trees", or "all"; type of base learners to use.
var.want	TRUE or FALSE, if variance estimates are wanted.
threshold	An optional threshold to test individuals for the null hypothesis that PTE > threshold; must have var.want = TRUE to return p-values.
use.actual.control.S	TRUE or FALSE, if user prefers to use the actual observed values for the surrogate in the control group instead of predicting values from the base learners.

Value

A dataframe is returned, which is the `df.test` argument with new columns appended for the estimates and corresponding variances of `delta`, `delta.s`, and `R.s`. If a threshold is specified, returns a p-value for the null hypothesis that $PTE > \text{threshold}$.

Author(s)

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References

Knowlton, R. and Parast, L. (2025) "Assessing Surrogate Heterogeneity in Real World Data Using Meta-Learners." Under Review.

Examples

```
data(exampladata_train)
data(exampladata_test)
obs.het.surr(df.train = exampladata_train, df.test = exampladata_test, type = "all", var.want = FALSE)
```

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