

Package ‘tidytreatment’

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Type Package

Title Tidy Methods for Bayesian Treatment Effect Models

Version 0.2.2

Description Functions for extracting tidy data from Bayesian treatment effect models, in particular BART, but extensions are possible. Functionality includes extracting tidy posterior summaries as in 'tidybayes' <<https://github.com/mjskay/tidybayes>>, estimating (average) treatment effects, common support calculations, and plotting useful summaries of these.

Encoding UTF-8

LazyData true

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URL <https://github.com/bonStats/tidytreatment>

BugReports <https://github.com/bonStats/tidytreatment/issues>

Language en-US

Depends R (>= 3.1.0)

Suggests knitr, rmarkdown, BART, ggplot2, testthat (>= 3.0.0), withr

VignetteBuilder knitr

RoxygenNote 7.1.1

Imports tidybayes, purrr, tidyr, dplyr, readr, rlang

Enhances bartMachine

Config/testthat/edition 3

NeedsCompilation no

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avg_treatment_effects *Get (conditional) average treatment effect draws from posterior*

Description

(C)ATE = (Conditional) Average Treatment Effects newdata specifies the conditions, if unspecified it defaults to the original data. Assumes treated column is either a integer column of 1's (treated) and 0's (nontreated) or logical indicating treatment if TRUE.

Usage

```
avg_treatment_effects(
  model,
  treatment,
  newdata,
  subset = "all",
  common_support_method,
  cutoff,
  ...
)
```

Arguments

model	A supported Bayesian model fit that can provide fits and predictions.
treatment	A character string specifying the name of the treatment variable.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
subset	Either "treated", "nontreated", or "all". Default is "all".
common_support_method	Either "sd", or "chisq". Default is unspecified, and no common support calculation is done.
cutoff	Cutoff for common support (if in use).
...	Arguments to be passed to <code>tidybayes::fitted_draws</code> typically scale for BART models.

Value

A tidy data frame (tibble) with treatment effect values.

bartmodel1

Example model 1

Description

Model fit with simulated data from simulated dataset `suhillsim1`.

Usage

```
bartmodel1
```

Format

Object of type `BART::wbart`

Details

Propensity score estimated and included suhillsim1 for fitting the model.

Source

<https://github.com/bonStats/tidyreatment/tree/master/data-raw>

bartmodel1_modelmatrix

Model matrix used for bartmodel1

Description

Useful for testing tidyreatment package functions.

Usage

bartmodel1_modelmatrix

Format

Object of type BART::wbart

Source

<https://github.com/bonStats/tidyreatment/tree/master/data-raw>

covariate_importance *Counts of variable overall inclusion*

Description

Inclusion metric for bartMachine and BART are scaled differently. bartMachine averaged over number of trees, in addition to number of MCMC draws.

Usage

covariate_importance(model, ...)

Arguments

model Model
 ... Arguments to pass to particular methods.

Value

Tidy data with counts of variable inclusion, when interacting with treatment variable.

covariate_with_treatment_importance
Counts of variable inclusion when interacting with treatment

Description

Counts of variable inclusion when interacting with treatment

Usage

```
covariate_with_treatment_importance(model, treatment, ...)
```

Arguments

model	Model
treatment	A character string specifying the name of the treatment variable.
...	Arguments to pass to particular methods.

Value

Tidy data with counts of variable inclusion, when interacting with treatment variable.

fitted_draws.bartMachine
Get fitted draws from posterior of bartMachine model

Description

Get fitted draws from posterior of bartMachine model

Usage

```
## S3 method for class 'bartMachine'  
fitted_draws(  
  model,  
  newdata,  
  value = ".value",  
  ...,  
  n = NULL,  
  include_newdata = TRUE,  
  include_sigsqs = FALSE  
)
```

Arguments

model	A bartMachine model.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
value	The name of the output column for fitted_draws; default ".value".
...	Not currently in use.
n	Not currently implemented.
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with fitted values.

fitted_draws.lbart *Get fitted draws from posterior of lbart model*

Description

Get fitted draws from posterior of lbart model

Usage

```
## S3 method for class 'lbart'
fitted_draws(
  model,
  newdata,
  value = ".value",
  ...,
  n = NULL,
  include_newdata = TRUE,
  include_sigsqs = FALSE
)
```

Arguments

model	A model from BART package.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
value	The name of the output column for fitted_draws; default ".value".
...	Not currently in use.
n	Not currently implemented.
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with fitted values.

fitted_draws.mbart	<i>Get fitted draws from posterior of mbart model</i>
--------------------	---

Description

Get fitted draws from posterior of mbart model

Usage

```
## S3 method for class 'mbart'
fitted_draws(
  model,
  newdata,
  value = ".value",
  ...,
  n = NULL,
  include_newdata = TRUE,
  include_sigsqs = FALSE
)
```

Arguments

model	A model from BART package.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
value	The name of the output column for fitted_draws; default ".value".
...	Not currently in use.
n	Not currently implemented.
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with fitted values.

fitted_draws.mbart2 *Get fitted draws from posterior of mbart2 model*

Description

Get fitted draws from posterior of mbart2 model

Usage

```
## S3 method for class 'mbart2'
fitted_draws(
  model,
  newdata,
  value = ".value",
  ...,
  n = NULL,
  include_newdata = TRUE,
  include_sigsqs = FALSE
)
```

Arguments

model	A model from BART package.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
value	The name of the output column for fitted_draws; default ".value".
...	Not currently in use.
n	Not currently implemented.
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with fitted values.

fitted_draws.pbart *Get fitted draws from posterior of pbart model*

Description

Get fitted draws from posterior of pbart model

Usage

```
## S3 method for class 'pbart'  
fitted_draws(  
  model,  
  newdata,  
  value = ".value",  
  ...,  
  n = NULL,  
  include_newdata = TRUE,  
  include_sigsqs = FALSE  
)
```

Arguments

model	A model from BART package.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
value	The name of the output column for fitted_draws; default ".value".
...	Not currently in use.
n	Not currently implemented.
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with fitted values.

fitted_draws.wbart *Get fitted draws from posterior of wbart model*

Description

Get fitted draws from posterior of wbart model

Usage

```
## S3 method for class 'wbart'  
fitted_draws(  
  model,  
  newdata,  
  value = ".value",  
  ...,  
  n = NULL,  
  include_newdata = TRUE,  
  include_sigsqs = FALSE  
)
```

Arguments

model	A model from BART package.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
value	The name of the output column for fitted_draws; default ".value".
...	Not currently in use.
n	Not currently implemented.
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with fitted values.

fitted_draws_BART	<i>Get fitted draws from posterior of BART-package models</i>
-------------------	---

Description

Get fitted draws from posterior of BART-package models

Usage

```
fitted_draws_BART(
  model,
  newdata = NULL,
  value = ".value",
  ...,
  include_newdata = TRUE,
  include_sigsqs = FALSE,
  scale = "real"
)
```

Arguments

model	A model from BART package.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
value	The name of the output column for fitted_draws; default ".value".
...	Arguments to pass to predict (e.g. BART::predict.wbart).
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?
scale	Should the fitted values be on the real, probit or logit scale?

Value

A tidy data frame (tibble) with fitted values.

has_common_support	<i>Evaluate if observations have common support.</i>
--------------------	--

Description

The common support identification methods are based on Hill and Su (2013). Loosely speaker, an individuals treatment effect estimate has common support if the counterfactual estimate is not too uncertain. The estimates are uncertain when the prediction is 'far away' from other observations. Removing estimates without common support can be beneficial for treat effect estimates.

Usage

```
has_common_support(model, treatment, method, cutoff, modeldata = NULL)
```

Arguments

model	A supported Bayesian model fit that can provide fits and predictions.
treatment	A character string specifying the name of the treatment variable.
method	Method to use in determining common support. 'chisq', or 'sd'.
cutoff	Cutoff point to use for method.
modeldata	Manually provide model data for some models (e.g. from BART package)

Details

Hill, Jennifer; Su, Yu-Sung. *Ann. Appl. Stat.* 7 (2013), no. 3, 1386–1420. doi:10.1214/13-AOAS630. <https://projecteuclid.org/euclid.aoas/1380804800>

Value

Tibble with a row for each observation and a column indicating whether common support exists.

```
has_tidyreatment_methods
```

Check if a model class has required generic methods for tidyreatment functions.

Description

Check if a model class has required generic methods for tidyreatment functions.

Usage

```
has_tidyreatment_methods(model)
```

Arguments

model	Model to be checked.
-------	----------------------

Value

Boolean

highDim_testdataset3 *ACIC2019 High Dimensional Test Dataset*

Description

Dataset from the "Data Challenge" for the Atlantic Causal Inference Conference 2019.

Usage

```
highDim_testdataset3
```

Format

A data frame with 2000 observations, and 187 variables.

Y Outcome variable

A Treatment variable

V1,V2,V3,V4,V5,V6,V7,V8,V9,V10,V11,V12,V13,V14,V15,V16,V17,V18,V19,V20,V21,V22,V23,V24,V25,V26,V27,V28,V29,V30,V31,V32,V33,V34,V35,V36,V37,V38,V39,V40,V41,V42,V43,V44,V45,V46,V47,V48,V49,V50,V51,V52,V53,V54,V55,V56,V57,V58,V59,V60,V61,V62,V63,V64,V65,V66,V67,V68,V69,V70,V71,V72,V73,V74,V75,V76,V77,V78,V79,V80,V81,V82,V83,V84,V85,V86,V87,V88,V89,V90,V91,V92,V93,V94,V95,V96,V97,V98,V99,V100,V101,V102,V103,V104,V105,V106,V107,V108,V109,V110,V111,V112,V113,V114,V115,V116,V117,V118,V119,V120,V121,V122,V123,V124,V125,V126,V127,V128,V129,V130,V131,V132,V133,V134,V135,V136,V137,V138,V139,V140,V141,V142,V143,V144,V145,V146,V147,V148,V149,V150,V151,V152,V153,V154,V155,V156,V157,V158,V159,V160,V161,V162,V163,V164,V165,V166,V167,V168,V169,V170,V171,V172,V173,V174,V175,V176,V177,V178,V179,V180,V181,V182,V183,V184,V185,V186,V187
Other covariates ...

Source

<https://www.mcgill.ca/epi-biostat-occh/seminars-events/atlantic-causal-inference-conference-2019/data-challenge>

posterior_trees_BART *Get posterior tree draws into tibble format from BART model*

Description

Tibble grouped by iteration ('iter') and tree id ('tree_id'). All information calculated by method is included in output.

Usage

```
posterior_trees_BART(model, label_digits = 2)
```

Arguments

model BART model.
label_digits Rounding for labels.

Value

A tibble with columns to

iter Integer describing unique MCMC iteration.

tree_id Integer. Unique tree id with each 'iter'.

node Integer describing node in tree. Unique to each 'tree'-'iter'.

parent Integer describing parent node in tree.

label Label for the node.

tier Position in tree hierarchy.

var Variable for split.

cut Numeric. Value of decision rule for 'var'.

is_leaf Logical. 'TRUE' if leaf, 'FALSE' if stem.

leaf_value

child_left Integer. Left child of node.

child_right Integer. Right child of node.

predicted_draws.bartMachine

Get predict draws from posterior of bartMachine model

Description

Get predict draws from posterior of bartMachine model

Usage

```
## S3 method for class 'bartMachine'  
predicted_draws(  
  object,  
  newdata,  
  value = ".prediction",  
  ...,  
  ndraws = NULL,  
  include_newdata = TRUE,  
  include_fitted = FALSE,  
  include_sigsqs = FALSE  
)
```

Arguments

object	A bartMachine model.
newdata	Data frame to generate predictions from. If omitted, most model types will generate predictions from the data used to fit the model.
value	The name of the output column for predicted_draws; default ".prediction".
...	Not currently in use.
ndraws	Not currently implemented.
include_newdata	Should the newdata be included in the tibble?
include_fitted	Should the posterior fitted values be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with predicted values.

predicted_draws.wbart *Get predict draws from posterior of wbart model*

Description

Get predict draws from posterior of wbart model

Usage

```
## S3 method for class 'wbart'
predicted_draws(
  object,
  newdata,
  value = ".prediction",
  ...,
  ndraws = NULL,
  include_newdata = TRUE,
  include_fitted = FALSE,
  include_sigsqs = FALSE
)
```

Arguments

object	A wbart model.
newdata	Data frame to generate predictions from. If omitted, most model types will generate predictions from the data used to fit the model.
value	The name of the output column for predicted_draws; default ".prediction".
...	Use to specify random number generator, default is rng=stats::rnorm.

ndraws Not currently implemented.
 include_newdata Should the newdata be included in the tibble?
 include_fitted Should the posterior fitted values be included in the tibble?
 include_sigsqs Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with predicted values.

predicted_draws_BART *Get predict draws from posterior of BART-package models*

Description

Get predict draws from posterior of BART-package models

Usage

```
predicted_draws_BART(
  object,
  newdata = NULL,
  value = ".prediction",
  ...,
  rng = stats::rnorm,
  include_newdata = TRUE,
  include_fitted = FALSE,
  include_sigsqs = FALSE
)
```

Arguments

object A BART-package model.
 newdata Data frame to generate predictions from. If omitted, most model types will generate predictions from the data used to fit the model.
 value The name of the output column for predicted_draws; default ".prediction".
 ... Arguments to pass to predict (e.g. BART:::predict.wbart).
 rng Random number generator function. Default is rnorm for models with Gaussian errors.
 include_newdata Should the newdata be included in the tibble?
 include_fitted Should the posterior fitted values be included in the tibble?
 include_sigsqs Should the posterior sigma-squared draw be included?

Value

A tidy data frame (tibble) with predicted values.

`residual_draws.bartMachine`*Get residual draw for bartMachine model*

Description

Get residual draw for bartMachine model

Usage

```
## S3 method for class 'bartMachine'
residual_draws(
  object,
  newdata,
  value = ".residual",
  ...,
  ndraws = NULL,
  include_newdata = TRUE,
  include_sigsqs = FALSE
)
```

Arguments

<code>object</code>	bartMachine model.
<code>newdata</code>	Data frame to generate predictions from. If omitted, original data used to fit the model.
<code>value</code>	Name of the output column for <code>residual_draws</code> ; default is <code>.residual</code> .
<code>...</code>	Additional arguments passed to the underlying prediction method for the type of model given.
<code>ndraws</code>	Not currently implemented.
<code>include_newdata</code>	Should the newdata be included in the tibble?
<code>include_sigsqs</code>	Should the posterior sigma-squared draw be included?

Value

Tibble with residuals.

residual_draws.pbart *Get residual draw for pbart model*

Description

The original response variable must be passed as an argument to this function. e.g. ‘response = y’

Usage

```
## S3 method for class 'pbart'  
residual_draws(  
  object,  
  newdata,  
  value = ".residual",  
  ...,  
  ndraws = NULL,  
  include_newdata = TRUE,  
  include_sigsqs = FALSE  
)
```

Arguments

object	wbart model.
newdata	Data frame to generate predictions from. If omitted, original data used to fit the model.
value	Name of the output column for residual_draws; default is .residual.
...	Additional arguments passed to the underlying prediction method for the type of model given.
ndraws	Not currently implemented.
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

Tibble with residuals.

`residual_draws.wbart` *Get residual draw for wbart model*

Description

The original response variable must be passed as an argument to this function. e.g. ‘response = y’

Usage

```
## S3 method for class 'wbart'
residual_draws(
  object,
  newdata,
  value = ".residual",
  ...,
  ndraws = NULL,
  include_newdata = TRUE,
  include_sigsqs = FALSE
)
```

Arguments

<code>object</code>	wbart model.
<code>newdata</code>	Data frame to generate predictions from. If omitted, original data used to fit the model.
<code>value</code>	Name of the output column for <code>residual_draws</code> ; default is <code>.residual</code> .
<code>...</code>	Additional arguments passed to the underlying prediction method for the type of model given.
<code>ndraws</code>	Not currently implemented.
<code>include_newdata</code>	Should the newdata be included in the tibble?
<code>include_sigsqs</code>	Should the posterior sigma-squared draw be included?

Value

Tibble with residuals.

residual_draws_BART *Get residual draw for BART model*

Description

Classes from BART-package models

Usage

```
residual_draws_BART(
  object,
  response,
  newdata = NULL,
  value = ".residual",
  include_newdata = TRUE,
  include_sigsqs = FALSE
)
```

Arguments

object	model from BART package.
response	Original response vector.
newdata	Data frame to generate predictions from. If omitted, original data used to fit the model.
value	Name of the output column for residual_draws; default is .residual.
include_newdata	Should the newdata be included in the tibble?
include_sigsqs	Should the posterior sigma-squared draw be included?

Value

Tibble with residuals.

simulate_su_hill_data *Simulate data with scenarios from Hill and Su (2013)*

Description

Sample n observations with the following scheme:

1. Covariates: $X_j \sim N(0, 1)$.
2. Assignment: $Z \sim \text{Bin}(n, p)$ with $p = \text{logit}^{-1}(a + X\gamma^L + Q\gamma^N)$ where $a = \omega - \text{mean}(X\gamma^L + Q\gamma^N)$.
3. Mean response: $E(Y(0)|X) = X\beta_0^L + Q\beta_0^N$ and $E(Y(1)|X) = X\beta_1^L + Q\beta_1^N$.
4. Observation: $Y \sim N(\mu, \sigma_y^2)$.

Superscript L denotes the linear components, whilst N denotes the non-linear components.

Usage

```
simulate_su_hill_data(
  n,
  treatment_linear = TRUE,
  response_parallel = TRUE,
  response_aligned = TRUE,
  y_sd = 1,
  tau = 4,
  omega = 0,
  add_categorical = FALSE,
  coef_categorical_treatment = NULL,
  coef_categorical_nontreatment = NULL
)
```

Arguments

n	Size of simulated sample.
treatment_linear	Treatment assignment mechanism is linear?
response_parallel	Response surface is parallel?
response_aligned	Response surface is aligned?
y_sd	Observation noise.
tau	Treatment effect for parallel response surfaces. Not applicable if surface is non-parallel.
omega	Offset to control treatment assignment ratios.
add_categorical	Should a categorical variable be added? (Not in Hill and Su)
coef_categorical_treatment	What are the coefficients of the categorical variable under treatment? (Not in Hill and Su)
coef_categorical_nontreatment	What are the coefficients of the categorical variable under nontreatment? (Not in Hill and Su)

Details

Coefficients used are returned in the list this function creates. See Table 1 in Su and Hill (2013) for the table of coefficients. The X_j are in a data.frame named data in the returned list. The formula for the model matrix $[X, Q]$ is named su_hill_formula in the returned list. The coefficients used for the model matrix are contained in coefs. The Su and Hill (2013) simulations did not include categorical variables, but you can add them here using arguments: add_categorical, coef_categorical_treatment, coef_categorical_nontreatment.

Hill, Jennifer; Su, Yu-Sung. Ann. Appl. Stat. 7 (2013), no. 3, 1386–1420. doi:10.1214/13-AOAS630. <https://projecteuclid.org/euclid.aoas/1380804800>

Value

An object of class `suhillsim` that is a list with elements

<code>data</code>	Simulated data in <code>data.frame</code>
<code>mean_y</code>	The mean <code>y</code> values for each individual (row)
<code>args</code>	List of arguments passed to function
<code>formulas</code>	Response formulas used to generate data
<code>coefs</code>	Coefficients for the formulas

<code>suhillsim1</code>	<i>Example simulated dataset 1</i>
-------------------------	------------------------------------

Description

Simulated with `simulate_su_hill_data(...)`, see details. Includes propensity score estimated using BART (`prop_score`), see source.

Usage

```
suhillsim1
```

Format

See `?simulate_su_hill_data` for output format.

Details

```
set.seed(101)
suhillsim1 <- simulate_su_hill_data(n = 100, treatment_linear = FALSE, omega = 0, add_categorical = TRUE,
  coef_categorical_treatment = c(0,0,1),
  coef_categorical_nontreatment = c(-1,0,-1))
```

Source

<https://github.com/bonStats/tidytreatment/tree/master/data-raw>

<code>tidytreatment</code>	<i>tidytreatment: Tidy methods for Bayesian treatment effect models</i>
----------------------------	---

Description

`tidytreatment` provides functions for extracting tidy data from Bayesian treatment effect models, estimating treatment effects, and plotting useful summaries of these.

tidy_ate	<i>Get average treatment effect draws from posterior</i>
----------	--

Description

ATE = Average Treatment Effects Assumes treated column is either a integer column of 1's (treated) and 0's (nontreated) or logical indicating treatment if TRUE.

Usage

```
tidy_ate(model, treatment, common_support_method, cutoff, ...)
```

Arguments

model	A supported Bayesian model fit that can provide fits and predictions.
treatment	A character string specifying the name of the treatment variable.
common_support_method	Either "sd", or "chisq". Default is unspecified, and no common support calculation is done.
cutoff	Cutoff for common support (if in use).
...	Arguments to be passed to tidybayes::fitted_draws typically scale for BART models.

Value

A tidy data frame (tibble) with treatment effect values.

tidy_att	<i>Get average treatment effect on treated draws from posterior</i>
----------	---

Description

ATT = average Treatment Effects on Treated Assumes treated column is either a integer column of 1's (treated) and 0's (nontreated) or logical indicating treatment if TRUE.

Usage

```
tidy_att(model, treatment, common_support_method, cutoff, ...)
```

Arguments

model	A supported Bayesian model fit that can provide fits and predictions.
treatment	A character string specifying the name of the treatment variable.
common_support_method	Either "sd", or "chisq". Default is unspecified, and no common support calculation is done.
cutoff	Cutoff for common support (if in use).
...	Arguments to be passed to <code>tidybayes::fitted_draws</code> typically scale for BART models.

Value

A tidy data frame (tibble) with treatment effect values.

treatment_effects	<i>Get (individual) treatment effect draws from posterior</i>
-------------------	---

Description

CTE = Conditional Treatment Effects (usually used to generate (C)ATE or ATT) `newdata` specifies the conditions, if unspecified it defaults to the original data. Assumes treated column is either a integer column of 1's (treated) and 0's (nontreated) or logical indicating treatment if TRUE.

Usage

```
treatment_effects(
  model,
  treatment,
  newdata,
  subset = "all",
  common_support_method,
  cutoff,
  ...
)
```

Arguments

model	A supported Bayesian model fit that can provide fits and predictions.
treatment	A character string specifying the name of the treatment variable.
newdata	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
subset	Either "treated", "nontreated", or "all". Default is "all".
common_support_method	Either "sd", or "chisq". Default is unspecified, and no common support calculation is done.

cutoff	Cutoff for common support (if in use).
...	Arguments to be passed to <code>tidybayes::fitted_draws</code> typically scale for BART models.

Value

A tidy data frame (tibble) with treatment effect values.

```
treatment_effects.default
```

Get treatment effect draws from posterior

Description

CTE = Conditional Treatment Effects (or CATE, the average effects) `newdata` specifies the conditions, if unspecified it defaults to the original data. Assumes treated column is either a integer column of 1's (treated) and 0's (nontreated) or logical indicating treatment if TRUE.

Usage

```
## Default S3 method:
treatment_effects(
  model,
  treatment,
  newdata,
  subset = "all",
  common_support_method,
  cutoff,
  ...
)
```

Arguments

<code>model</code>	A supported Bayesian model fit that can provide fits and predictions.
<code>treatment</code>	A character string specifying the name of the treatment variable.
<code>newdata</code>	Data frame to generate fitted values from. If omitted, defaults to the data used to fit the model.
<code>subset</code>	Either "treated", "nontreated", or "all". Default is "all".
<code>common_support_method</code>	Either "sd", or "chisq". Default is unspecified, and no common support calculation is done.
<code>cutoff</code>	Cutoff for common support (if in use).
...	Arguments to be passed to <code>tidybayes::fitted_draws</code> typically scale for BART models.

Value

A tidy data frame (tibble) with treatment effect values.

variance_draws	<i>Get variance draws from posterior of BART models</i>
----------------	---

Description

Models from BART-package include warm-up and skipped MCMC draws.

Usage

```
variance_draws(model, value = ".sigma_sq", ...)
```

Arguments

model	A model from a supported package.
value	The name of the output column for variance parameter; default ".sigma_sq".
...	Additional arguments.

Value

A tidy data frame (tibble) with draws of variance parameter

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