

# Package ‘cbar’

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**Type** Package

**Title** Contextual Bayesian Anomaly Detection in R

**Version** 0.1.3

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**Description** Detect contextual anomalies in time-series data with Bayesian data analysis. It focuses on determining a normal range of target value, and provides simple-to-use functions to abstract the outcome.

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**URL** <https://github.com/zedoul/cbar>

**BugReports** <https://github.com/zedoul/cbar/issues>

**Depends** R (>= 3.3.0)

**Imports** Boom, bsts, dplyr, magrittr, ggplot2, stats

**Suggests** datasets, knitr, testthat

**License** Apache License 2.0 | file LICENSE

**LazyData** true

**RoxygenNote** 6.0.1

**NeedsCompilation** no

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<b>bsts_model</b>	<i>Create bsts model</i>
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## Description

Create bsts model

## Usage

```
bsts_model(.data, spec = NULL, ...)
```

## Arguments

.data	training set
spec	cbar.model.spec object
...	params for bsts_spec_static

## Value

bsts which is a bsts model

## References

Scott, S. L., & Varian, H. R. (2014). Predicting the present with bayesian structural time series. International Journal of Mathematical Modelling and Numerical Optimisation, 5(1-2), 4-23.

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`bsts_spec_static`      *Specify bsts model for static linear regression*

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## Description

Specify bsts model for static linear regression

## Usage

```
bsts_spec_static(.data, sigma_guess = NULL, upper_limit = NULL,
  sd_prior_sample_size = 32, expected_model_size = 3, expected_r2 = 0.8,
  prior_df = 50, niter = 1000, ping = 0, model_options = NULL, ...)
```

## Arguments

<code>.data</code>	time-series data to be trained
<code>sigma_guess</code>	an argument for <code>bsts::bsts</code>
<code>upper_limit</code>	an argument for <code>bsts::bsts</code>
<code>sd_prior_sample_size</code>	an argument for <code>bsts::bsts</code>
<code>expected_model_size</code>	an argument for <code>bsts::bsts</code>
<code>expected_r2</code>	an argument for <code>bsts::bsts</code>
<code>prior_df</code>	an argument for <code>bsts::bsts</code>
<code>niter</code>	an argument for <code>bsts::bsts</code>
<code>ping</code>	an argument for <code>bsts::bsts</code>
<code>model_options</code>	an argument for <code>bsts::bsts</code>
<code>...</code>	params for <code>bsts_model</code>

## Value

`cbar.model.spec` object for model specification

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**cbar***cbar package*

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## Description

**cbar:** Contextual Bayesian Anomaly Detection in R

This function generates **cbar** object to detect contextual anomaly and to abstract analysis output.

## Usage

```
cbar(.data, ref_period, mea_period, apply_standardized = T, interval = 0.95,  
...)
```

## Arguments

.data	data table with datetime, y, and predictors
ref_period	performance reference period
mea_period	performance measurement period
apply_standardized	whether it will standardized data or not
interval	credible interval. 0.95 by default.
...	params for bsts_model

## Details

See the README on [Github](#)

For the input `.data`, note that you should use `datetime` for the first column name. Also, you should use `numeric` type for other columns.

## Examples

```
library(cbar)

.data <- mtcars
rownames(.data) <- NULL
datetime <- seq(from = Sys.time(), length.out = nrow(.data), by = "mins")
.data <- cbind(datetime = datetime, .data)

ref_session <- 1:16
mea_session <- 17:nrow(.data)

obj <- cbar(.data, ref_session, mea_session)
```

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destandardized	<i>Destandardize a vector</i>
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**Description**

Destandardize a vector

**Usage**

```
destandardized(y_hat, y_mu, y_sd)
```

**Arguments**

y_hat	standardized numeric vector
y_mu	a mean value of unstandarized vector
y_sd	a standard deviation of unstandarized vector

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inference	<i>Infer from predictive posetrior prediction of bst model</i>
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**Description**

Infer from predictive posetrior prediction of bst model

**Usage**

```
inference(.model, alpha = 0.05)
```

**Arguments**

.model	bsts model
alpha	percentile for anomaly

**Value**

`data.frame` with observations and predictions

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<code>plot_error</code>	<i>Print estimation error plot</i>
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### Description

Print estimation error plot

### Usage

```
plot_error(.cbar, xlab = "", ylab = "Estimation error", method = "diff",
...)
```

### Arguments

.cbar	cbar object
xlab	a label for x-axis
ylab	a label for y-ayis
method	diff
...	params for boxplot

### Value

boxplot object

---

<code>plot_error_</code>	<i>Print estimation error plot</i>
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### Description

Print estimation error plot

### Usage

```
plot_error_(.error, xlab = "", ylab = "Estimation error", method = "diff",
...)
```

### Arguments

.error	error data frame
xlab	a label for x-axis
ylab	a label for y-ayis
method	diff
...	params for boxplot

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**plot\_incprob** *Print inclusion probability plot*

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## Description

Print inclusion probability plot

## Usage

```
plot_incprob(.cbar, threshold = 0.1, horiz = T, cex.names = 0.5,  
            xlab = "Inclusion probability (%)", las = 1, ...)
```

## Arguments

.cbar	cbar object
threshold	a threshold for inclusion probability
horiz	horiz
cex.names	cex.names
xlab	xlab
las	las
...	params for barplot

## Value

boxplot object

---

**plot\_incprob\_** *Print inclusion probability plot*

---

## Description

Print inclusion probability plot

## Usage

```
plot_incprob_(.incprob, threshold = 0.1, horiz = T, cex.names = 0.5,  
              xlab = "Inclusion probability (%)", las = 1, ...)
```

**Arguments**

.incprob	data frame
threshold	a threshold for inclusion probability
horiz	horiz
cex.names	cex.names
xlab	xlab
las	las
...	params for barplot

**Value**

boxplot object

**plot\_ts**

*Print time-series plot*

**Description**

Print time-series plot

**Usage**

```
plot_ts(.cbar, x_label = "", y_label = "", seq_by = NULL)
```

**Arguments**

.cbar	cbar object
x_label	a label for x-axis
y_label	a label for y-axis
seq_by	increment of the sequence, which is NULL by default

**Value**

ggplot object

---

plot_ts_	<i>Print time-series plot</i>
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**Description**

Print time-series plot

**Usage**

```
plot_ts_(target_data, x_label = "", y_label = "", seq_by = NULL)
```

**Arguments**

target_data	data frame
x_label	a label for x-axis
y_label	a label for y-axis
seq_by	increment of the sequence, which is NULL by default

**Value**

ggplot object

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point_prediction	<i>Get point prediction from posterior means and response trajectories</i>
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**Description**

Get point prediction from posterior means and response trajectories

**Usage**

```
point_prediction(y_hat, .posterior_mean, alpha = 0.05)
```

**Arguments**

y_hat	response trajectories
.posterior_mean	posterior mean values
alpha	alpha

**Value**

data.frame for predicted values

posterior\_mean

*Generate posterior mean of the response variable***Description**

This one is used for point prediction based one predictive posterior distribution

**Usage**

```
posterior_mean(.model)
```

**Arguments**

.model	bsts_model
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**Value**

vector that contains poseterior means

print.cbar

*Print cbar object***Description**

Print cbar object

**Usage**

```
## S3 method for class 'cbar'
print(x, ...)
```

**Arguments**

x	cbar object to print
...	further arguments passed to or from other methods

---

response\_trajectory     *Generate trajectories of the response variable*

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### Description

Note that posterior\_state\_samples returns posterior mean, whereas, this one returns posterior mean + noise

### Usage

```
response_trajectory(.model)
```

### Arguments

.model	bsts_model
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### Details

This one is used for lower and upper bounds

### Value

data.frame that contains predicted value y\_hat

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standardized     *Standardize a vector*

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### Description

Standardize a vector

### Usage

```
standardized(y)
```

### Arguments

y	numeric vector
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`summarise_anomaly`      *Summarise anomaly detection result*

### Description

Summarise anomaly detection result

### Usage

```
summarise_anomaly(.cbar, .session = NULL)
```

### Arguments

.cbar	cbar object
.session	names of sessions, which is NULL by default

### Value

`data.frame` that summarises input data with anomaly label

`summarise_incprob`      *Summarise inclusion probability of model*

### Description

Summarise inclusion probability of model

### Usage

```
summarise_incprob(.cbar, threshold = 0.1)
```

### Arguments

.cbar	cbar object
threshold	threshold of inclusion probability, which is .1 by default

### Value

vector that summarises inclusion probabilities for each MCMC samples

---

summarise\_pred\_error *Summarise prediction error of model*

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### Description

This function uses absolute difference and mean absolute percentage error for summarising prediction errors

### Usage

```
summarise_pred_error(.cbar, .session = "measurement")
```

### Arguments

- |          |   |
|----------|---|
| .cbar    | cbar object                                 |
| .session | names of sessions, which is NULL by default |

### Value

`data.frame` with prediction errors

---

summarise\_session *Summarise anomaly in session*

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### Description

Summarise anomaly in session

### Usage

```
summarise_session(.cbar, .session = NULL)
```

### Arguments

- |          |   |
|----------|---|
| .cbar    | cbar object                                 |
| .session | names of sessions, which is NULL by default |

### Value

`data.frame` that summarises outcome for each sessoin

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