

Package ‘ggchangepoint’

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Type Package

Title Combines Changepoint Analysis with 'ggplot2'

Version 0.3.0

Description R provides fantastic tools for changepoint analysis, but plots generated by the tools do not have the 'ggplot2' style. This tool, however, combines 'changepoint', 'changepoint.np' and 'ecp' together, and uses 'ggplot2' to visualize changepoints. It provides a unified 'ggcpt' S3 result class, 'broom'-style tidy/glance/augment methods, 'autoplot()', composable geoms ('geom_changepoint()', 'geom_cpt_segment()', 'geom_cpt_ci()', 'stat_changepoint()'), a unified 'cpt_detect()' dispatcher with method introspection via 'cpt_methods()', wrappers for several optional engines (WBS, WBS2, NOT, MOSUM, FPOP, Isolate-Detect, TGUH), a method comparison module, accuracy metrics, data simulation, and canonical test signals.

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Encoding UTF-8

Imports changepoint, changepoint.np, dplyr, ecp, generics, ggplot2 (>= 3.4.0), lifecycle, Rdpack, tibble, utils

RdMacros Rdpack

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Suggests rmarkdown, knitr, testthat (>= 3.0.0), wbs, breakfast, not, mosum, fpop, IDetect, future, future.apply

VignetteBuilder knitr

URL <https://pursuitofdatascience.github.io/ggchangepoint/>

BugReports <https://github.com/PursuitOfDataScience/ggchangepoint/issues>

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Contents

annotate_segments	3
augment.ggcpt	3
autoplot.ggcpt	4
cpt_detect	5
cpt_methods	6
cpt_metrics	6
cpt_metrics_annotated	7
cpt_penalty	8
cpt_simulate	9
cpt_wrapper	10
ecp_wrapper	11
fpop_wrapper	12
geom_changepoint	12
geom_cpt_ci	13
geom_cpt_segment	13
ggcptplot	14
ggcpt_compare	15
ggcpt_compare_table	16
ggcpt_eval	17
ggcpt_methods	17
ggcpcplot	18
glance.ggcpt	19
idetct_wrapper	20
is_ggcpt	20
mosum_wrapper	21
new_ggcpt	21
not_wrapper	22
print.ggcpt	23
signal_blocks	23
signal_fms	24
signal_mix	24
signal_stairs	25
signal_teeth	25
stat_changepoint	26
summary.ggcpt	27
tguh_wrapper	27
theme_ggcpt	28
tidy.ggcpt	28
wbs2_wrapper	29
wbs_wrapper	29

Index

30

annotate_segments	<i>Annotate segments with alternating shading</i>
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Description

Adds alternating shaded rectangles to highlight segments between changepoints.

Usage

```
annotate_segments(cp, n, fill = c("grey90", "white"), alpha = 0.5, ...)
```

Arguments

cp	Changepoint indices (including 0 and n).
n	Length of the series.
fill	Colors for alternating segments. Defaults to c("grey90", "white").
alpha	Alpha for fill. Defaults to 0.5.
...	Additional arguments passed to annotate.

Value

A list of ggplot annotations.

augment.ggcpt	<i>Augment a ggcpt object</i>
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Description

Returns the original data with added columns: seg_id, .fitted, .resid, and is_changepoint.

Usage

```
## S3 method for class 'ggcpt'
augment(x, ...)
```

Arguments

x	A ggcpt object.
...	Additional arguments (ignored).

Value

A tibble with the original data plus augment columns.

autoplot.ggcpt	<i>Autoplot a ggcpt object</i>
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Description

Renders a changepoint detection result as a ggplot. The raw series is drawn as a line (with optional points), changepoints are shown as vertical lines, and (optionally) fitted segment levels are overlaid.

Usage

```
## S3 method for class 'ggcpt'
autoplot(
  object,
  show_segments = FALSE,
  show_ci = FALSE,
  cptline_alpha = 1,
  cptline_color = "blue",
  cptline_type = "solid",
  cptline_linewidth = 0.5,
  show_points = NULL,
  show_line = TRUE,
  ...
)
```

Arguments

<code>object</code>	A ggcpt object.
<code>show_segments</code>	Logical. Whether to draw the fitted segment means. Defaults to FALSE.
<code>show_ci</code>	Logical. Whether to draw confidence intervals for changepoint locations (if available). Defaults to FALSE.
<code>cptline_alpha</code>	Alpha for changepoint lines. Defaults to 1.
<code>cptline_color</code>	Color for changepoint lines. Defaults to "blue".
<code>cptline_type</code>	Linetype for changepoint lines. Defaults to "solid".
<code>cptline_linewidth</code>	Linewidth for changepoint lines. Defaults to 0.5.
<code>show_points</code>	Logical. Whether to draw data points. Auto-off above 500 obs.
<code>show_line</code>	Logical. Whether to draw the line. Defaults to TRUE.
<code>...</code>	Additional arguments passed to ggcptplot_internal.

Value

A ggplot object.

cpt_detect

*Unified changepoint detection dispatcher***Description**

Runs one or more changepoint detection methods on a sequence and returns a tidy ggcpt result object. This is the recommended entry point for most users.

Usage

```
cpt_detect(x, method = "pelt", change_in = "mean", penalty = "MBIC", ...)
```

Arguments

x	A numeric vector (the data series).
method	Detection method. See cpt_methods() for the complete table. Methods available in this release: "pelt", "binseg", "segneigh", "amoc", "np", "ecp", "fpop", "wbs", "wbs2", "not", "mosum", "idetect", "tguh". Methods that ship with optional (Suggests) engine packages will prompt you to install them if missing. Planned methods — "smuce", "hsmuce", "kcp", "cpm", "robust", "decafs", "sn", "inspect", "sbs", "bcp", "bocpd", "strucchange", "segmented" — are listed in cpt_methods() with their target release.
change_in	What to detect change in. One of "mean", "var", "meanvar", "slope", "distribution". Defaults to "mean".
penalty	Penalty type or value. Either a character string ("MBIC", "BIC", "AIC", "Hannan-Quinn") or a numeric penalty value. Defaults to "MBIC". Numeric penalties are honoured by the functional-pruning method (fpop); the changepoint-based methods expect one of the character options.
...	Additional arguments passed to the specific wrapper.

Value

A ggcpt object.

Examples

```
set.seed(2022)
x <- c(rnorm(100, 0, 1), rnorm(100, 10, 1))
result <- cpt_detect(x, method = "pelt", change_in = "mean")
result
ggplot2::autoplot(result)
```

cpt_methods

Introspect available changepoint detection methods

Description

Returns a tibble describing every method the package knows about — those that are wired and those that are planned — along with their capabilities and installation status. Useful for discovering what can be run and what needs to be installed.

Usage

```
cpt_methods()
```

Value

A tibble with columns:

method Method name as passed to `cpt_detect()`.

change_in What types of change the method can detect.

engine The upstream R package that implements the method.

status "available" (wired in this release) or "planned" (future).

installed TRUE if the engine package is installed, FALSE if it is a Suggests engine that is missing, NA for planned methods.

target_release The release that plans to wire this method, or NA for currently available methods.

Examples

```
cpt_methods()
```

cpt_metrics

Changepoint accuracy metrics

Description

Computes standard accuracy metrics comparing predicted changepoints to ground truth, including precision/recall/F1 with margin, covering metric, Hausdorff distance, (adjusted) Rand index, annotation error, and MAE/RMSE of matched locations.

Usage

```
cpt_metrics(pred, truth, n, margin = 5)
```

Arguments

pred	Predicted changepoint indices (integer vector).
truth	Ground truth changepoint indices (integer vector).
n	Length of the series.
margin	Tolerance margin for matching (default 5).

Value

A tibble with columns: n, n_pred, n_truth, precision, recall, f1, covering, hausdorff, rand_index, annotation_error, mae_matched, rmse_matched.

Examples

```
cpt_metrics(c(100, 200), c(100, 200), n = 300)
cpt_metrics(c(101, 205), c(100, 200), n = 300, margin = 5)
```

cpt_metrics_annotated *Multi-annotator evaluation*

Description

Computes averaged covering and F1 scores against multiple annotation sets, as used in the Turing Change Point Dataset benchmark.

Usage

```
cpt_metrics_annotated(pred, annotations, n, margin = 5)
```

Arguments

pred	Predicted changepoint indices.
annotations	A list of ground-truth annotation vectors.
n	Length of the series.
margin	Tolerance margin (default 5).

Value

A tibble with averaged metrics.

cpt_penalty	<i>Construct changepoint penalties</i>
-------------	--

Description

Helper to construct standard penalty values for use with changepoint detection methods. Returns a numeric penalty value.

Usage

```
cpt_penalty(type, n = NULL, k = 1, value = NULL)
```

Arguments

type	Penalty type: "None", "BIC" (or "SIC"), "MBIC", "AIC", "Hannan-Quinn", "sSIC", or "Manual".
n	Series length. Required for BIC, MBIC, AIC, Hannan-Quinn, sSIC.
k	Number of parameters per changepoint (typically 2 for mean+variance, 1 for mean-only). Defaults to 1.
value	Numeric value for Manual type.

Value

A numeric penalty value.

Penalty semantics across engines

The same penalty name may be interpreted differently by different engines:

- **changepoint-based methods** (PELT, BinSeg, SegNeigh, AMOC): accept character penalties ("MBIC", "BIC", "AIC", "Hannan-Quinn", "None") and pass them to the upstream **changepoint** package. These methods do *not* accept raw numeric penalty values.
- **Functional-pruning methods** (fpop): accept numeric penalties only. When a character penalty is supplied via `cpt_detect()`, it is resolved to a numeric value using `cpt_penalty()` before dispatch.
- **Search-based methods** (WBS, WBS2, NOT, MOSUM, IDetect, TGUH): use internal model-selection criteria (e.g., sSIC, threshold) and generally *ignore* the penalty argument. Specify thresholds via the wrapper's own arguments.
- MBIC in `cpt_penalty()` uses the Zhang-Siegmund (2007) formula $0.5(k+1) \log n + \log \binom{n}{k}$, which differs from the **changepoint** package's MBIC. Use the character "MBIC" with **changepoint**-based methods to get the engine's native MBIC.

Examples

```
cpt_penalty("BIC", n = 100)
cpt_penalty("AIC", n = 100)
cpt_penalty("Manual", value = 5)
```


cpt_simulate

*Generate simulated changepoint data***Description**

Creates a synthetic time series with known changepoints for testing and benchmarking.

Usage

```
cpt_simulate(
  n,
  changepoints = integer(),
  change_in = c("mean", "var", "meanvar", "slope"),
  params = NULL,
  noise = c("gauss", "t", "ar1", "rw"),
  sd = 1,
  df = 3,
  rho = 0,
  seed = NULL
)

rcpt(...)
```

Arguments

n	Length of the series.
changepoints	Integer vector of changepoint locations (last index of each segment before the change).
change_in	What changes: "mean", "var", "meanvar", or "slope".
params	A list of parameters per segment. For mean changes, a vector of segment means. For var changes, a vector of segment sds. For meanvar, a list of lists with mean and sd per segment. For slope, a list with intercept and slope per segment.
noise	Noise type: "gauss" (Gaussian), "t" (Student-t), "ar1" (AR(1)), or "rw" (random walk).
sd	Noise standard deviation (for Gaussian and t). Defaults to 1.
df	Degrees of freedom for t-noise. Defaults to 3.
rho	AR(1) autocorrelation parameter. Defaults to 0.
seed	Optional seed for reproducibility.
...	Passed to <code>cpt_simulate</code> .

Value

A tibble with columns index, value, and seg_id. The true changepoints are stored in the true_changepoints attribute.

Examples

```
dat <- cpt_simulate(200, changepoints = c(100), change_in = "mean",
                  params = c(0, 10), seed = 2022)
attr(dat, "true_changepoints")
```

cpt_wrapper

Changepoint wrapper

Description

This function wraps a number of cpt functions from the changepoint package and the cpt.np() function from the changepoint.np package. It is handy that users can use this function to get the same changepoint results as these functions output individually. Moreover, it returns a tibble that inherits the tidyverse style. Functions from the changepoint package do require data normality assumption by default, yet changepoint.np is a non-parametric way to detect changepoints and let data speak by itself. If user sets change_in as np (or cpt_np), a seed should be set before using the function for the sake of reproducibility. For more details on the changepoint and changepoint.np packages, please refer to their documentation.

Usage

```
cpt_wrapper(data, change_in = "mean_var", cp_method = "PELT", ...)
```

Arguments

data	A numeric vector.
change_in	Choice of mean_var, mean, var, and np (or cpt_np for backward compatibility). Each choice corresponds to cpt.meanvar(), cpt.mean(), cpt.var() and cpt.np() respectively. The default is mean_var.
cp_method	A wide range of choices (i.e., AMOC, PELT, SegNeigh or BinSeg). Please note when change_in is np or cpt_np, PELT is the only option.
...	Extra arguments for each cpt function mentioned in the change_in section.

Value

A tibble including which point(s) is/are the changepoint along with raw changepoint value corresponding to that changepoint. Changepoint locations follow the convention of the changepoint package: the last index of the left segment.

References

Killick R, Eckley I (2014). “changepoint: An R package for changepoint analysis.” *Journal of statistical software*, **58**(3), 1–19.

Examples

```
set.seed(2022)
cpt_wrapper(c(rnorm(100,0,1),rnorm(100,0,10)))
cpt_wrapper(c(rnorm(100,0,1),rnorm(100,10,1)))
```

ecp_wrapper	<i>ecp wrapper</i>
-------------	--------------------

Description

The ecp package provides a non-parametric way to detect changepoints. Unlike the changepoint package, it does not assume raw data to have any formal distribution. This wrapper function wraps two functions from the ecp package, i.e., `e.divisive()` and `e.agglo()`. Users can use either function by switching the `algorithm` argument. Before using the wrapper function, seed should be set for the sake of reproducibility.

Usage

```
ecp_wrapper(data, algorithm = "divisive", min_size = 2, seed = NULL, ...)
```

Arguments

<code>data</code>	A numeric vector (for univariate) or matrix/data.frame (for multivariate).
<code>algorithm</code>	Either <code>divisive</code> or <code>agglo</code> . <code>divisive</code> is the default.
<code>min_size</code>	Minimum number of observations between change points. By default is 2. This argument is only applied when <code>algorithm = "divisive"</code> .
<code>seed</code>	Optional. A seed for reproducibility of the stochastic permutation test.
<code>...</code>	Extra arguments to pass on either from <code>e.divisive()</code> or <code>e.agglo()</code> .

Value

A tibble includes which point(s) is/are the changepoint along with raw changepoint value corresponding to that changepoint. Changepoint locations follow the ecp package convention: the first index of the right segment. When no changepoint is found, an empty tibble is returned (0 rows).

References

James NA, Matteson DS (2013). “ecp: An R package for nonparametric multiple change point analysis of multivariate data.” *arXiv preprint arXiv:1309.3295*.

Examples

```
set.seed(2022)
ecp_wrapper(c(rnorm(100,0,1),rnorm(100,0,10)))
ecp_wrapper(c(rnorm(100,0,1),rnorm(100,10,1)))
```

fpop_wrapper	<i>FPOP wrapper — Functional Pruning Optimal Partitioning</i>
--------------	---

Description

Wraps the fpop package for optimal changepoint detection via functional pruning.

Usage

```
fpop_wrapper(x, penalty = NULL, ...)
```

Arguments

x	A numeric vector.
penalty	Penalty value. Defaults to $2 * \log(\text{length}(x))$ (BIC).
...	Additional arguments passed to <code>fpop::Fpop()</code> .

Value

A ggcpt object.

geom_changepoint	<i>Changepoint vertical rules geom</i>
------------------	--

Description

Draws vertical lines at changepoint locations. Mimics `geom_vline` but designed to work with the tidy changepoint data frames returned by the package. Can be used as a standalone layer: `geom_changepoint(data = cp_df, aes(xintercept = cp))`.

Usage

```
geom_changepoint(
  mapping = NULL,
  data = NULL,
  ...,
  na.rm = FALSE,
  show.legend = NA
)
```

Arguments

mapping	Set of aesthetic mappings created by <code>ggplot2::aes()</code> . Requires <code>xintercept</code> .
data	A data frame with changepoint information.
...	Other arguments passed to <code>geom_vline</code> .
na.rm	If <code>FALSE</code> , missing values are removed.
show.legend	Whether to show legend.

Value

A ggplot layer.

geom_cpt_ci	<i>Changepoint confidence interval geom</i>
-------------	---

Description

Draws horizontal whiskers for changepoint-location confidence intervals (e.g. from MOSUM, stepR, strucchange, segmented).

Usage

```
geom_cpt_ci(mapping = NULL, data = NULL, ..., na.rm = FALSE, show.legend = NA)
```

Arguments

mapping	Aesthetic mappings. Requires x, xmin, xmax, and y.
data	A data frame with CI information.
...	Other arguments passed to geom_errorbarh.
na.rm	If FALSE, missing values are removed.
show.legend	Whether to show legend.

Value

A ggplot layer.

geom_cpt_segment	<i>Changepoint segment level geom</i>
------------------	---------------------------------------

Description

Draws horizontal segments representing the estimated level of each segment between changepoints. Typically used with data from augment().

Usage

```
geom_cpt_segment(
  mapping = NULL,
  data = NULL,
  ...,
  na.rm = FALSE,
  show.legend = NA
)
```

Arguments

<code>mapping</code>	Aesthetic mappings. Requires <code>x</code> , <code>xend</code> , <code>y</code> , <code>yend</code> .
<code>data</code>	A data frame with segment information.
<code>...</code>	Other arguments passed to <code>geom_segment</code> .
<code>na.rm</code>	If <code>FALSE</code> , missing values are removed.
<code>show.legend</code>	Whether to show legend.

Value

A ggplot layer.

ggcptplot

Plot for the changepoint package

Description

The plot for changepoints detected by the changepoint package is a line plot for the raw data and the vertical lines representing each changepoint. The x-axis is the row number of the raw data in the original data vector. The plot inherits `ggplot2`, meaning users can add `ggplot2` functions on top the changepoint plot for customization.

Usage

```
ggcptplot(
  data,
  change_in = "mean_var",
  cp_method = "PELT",
  ...,
  cptline_alpha = 1,
  cptline_color = "blue",
  cptline_type = "solid",
  cptline_linewidth = 0.5,
  cptline_size = lifecycle::deprecated(),
  index = NULL,
  show_points = NULL,
  show_line = TRUE
)
```

Arguments

<code>data</code>	A numeric vector.
<code>change_in</code>	Choice of <code>mean_var</code> , <code>mean</code> , <code>var</code> , and <code>np</code> (or <code>cpt_np</code> for backward compatibility). Each choice corresponds to <code>cpt.meanvar()</code> , <code>cpt.mean()</code> , <code>cpt.var()</code> and <code>cpt.np()</code> respectively. The default is <code>mean_var</code> .

cp_method	A wide range of choices (i.e., AMOC, PELT, SegNeigh or BinSeg). Please note when change_in is np or cpt_np, PELT is the only option.
...	Extra arguments for each cpt function mentioned in the change_in section.
cptline_alpha	The value of alpha for the vertical changepoint line(s), default is 1, meaning no transparency.
cptline_color	The color for the vertical changepoint line(s), default is blue.
cptline_type	The linetype for the vertical changepoint line(s), default is solid.
cptline_linewidth	The linewidth for the vertical changepoint line(s), default is 0.5.
cptline_size	Deprecated. Use cptline_linewidth instead.
index	Optional. A vector of x-axis labels (e.g. dates) of the same length as data.
show_points	Logical. Whether to draw data points. Defaults to TRUE when length(data) <= 500, FALSE otherwise.
show_line	Logical. Whether to draw the line. Defaults to TRUE.

Value

A line plot with data points along with the vertical lines representing changepoints.

Examples

```
ggcptplot(c(rnorm(100,0,1),rnorm(100,0,10)))
ggcptplot(c(rnorm(100,0,1),rnorm(100,10,1)))
```

ggcpt_compare

Compare multiple changepoint detection methods

Description

Runs several detectors on the same data and returns a faceted or overlaid ggplot comparison. Respects `future::plan()` for parallel execution if the `future.apply` package is available.

Usage

```
ggcpt_compare(
  x,
  methods = c("pelt", "binseg", "amoc"),
  layout = c("facet", "overlay"),
  change_in = "mean",
  seed = NULL,
  ...
)
```

Arguments

<code>x</code>	A numeric vector (the data series).
<code>methods</code>	Character vector of method names (passed to <code>cpt_detect</code>).
<code>layout</code>	Layout type. "facet" (default) shows one panel per method; "overlay" draws all changepoints in one panel, colour-coded.
<code>change_in</code>	What to detect change in. Passed to each detector.
<code>seed</code>	Optional seed for reproducible parallelism. Passed to <code>future.apply::future_lapply()</code> as <code>future.seed</code> .
<code>...</code>	Additional arguments passed to each detector.

Value

A ggplot object.

Examples

```
set.seed(2022)
x <- c(rnorm(100, 0, 1), rnorm(100, 10, 1))
ggcpt_compare(x, methods = c("pelt", "binseg"))
```

<code>ggcpt_compare_table</code>	<i>Comparison table</i>
----------------------------------	-------------------------

Description

Returns a tidy tibble combining the results of multiple detectors on the same series.

Usage

```
ggcpt_compare_table(
  x,
  methods = c("pelt", "binseg", "amoc"),
  change_in = "mean",
  ...
)
```

Arguments

<code>x</code>	A numeric vector (the data series).
<code>methods</code>	Character vector of method names.
<code>change_in</code>	What to detect change in.
<code>...</code>	Additional arguments passed to each detector.

Value

A tibble with columns `method`, `cp`, `cp_value`.

ggcpt_eval	<i>Evaluation visualization</i>
------------	---------------------------------

Description

Overlays predictions and ground truth on the series with tolerance windows, colouring true positives, false positives, and misses.

Usage

```
ggcpt_eval(pred, truth, data_vec, margin = 5)
```

Arguments

pred	Predicted changepoint indices.
truth	Ground truth changepoint indices.
data_vec	The original data vector (for context).
margin	Tolerance margin (default 5).

Value

A ggplot object.

ggcpt_methods	<i>Coerce, format, and plot ggcpt objects</i>
---------------	---

Description

Convenience S3 methods for working with ggcpt objects: coerce the changepoints to a tibble or data frame, render a one-line summary string, or produce the default plot (a base-graphics fallback that delegates to [autoplot.ggcpt](#)).

Usage

```
## S3 method for class 'ggcpt'
as_tibble(x, ..., .name_repair = NULL)

## S3 method for class 'ggcpt'
as.data.frame(x, row.names = NULL, optional = FALSE, ...)

## S3 method for class 'ggcpt'
format(x, ...)

## S3 method for class 'ggcpt'
plot(x, ...)
```

Arguments

`x` A ggcpt object.
`...` Additional arguments passed to methods.
`.name_repair` Passed to [as_tibble](#).
`row.names, optional`
 Passed to [as.data.frame](#).

Value

`as_tibble()` and `as.data.frame()` return the changepoints table; `format()` returns a length-one character string; `plot()` returns a ggplot object.

Examples

```

set.seed(2022)
res <- cpt_detect(c(rnorm(50), rnorm(50, 5)), method = "pelt")
as_tibble(res)
as.data.frame(res)
format(res)

```

ggecpplot

Plot for the ecp package

Description

The plot for changepoints detected by the ecp package is a line plot for the raw data and the vertical lines representing each changepoint. The x-axis is the row number of the raw data in the original data vector. The plot inherits ggplot2, meaning users can add ggplot2 functions on top the changepoint plot for customization.

Usage

```

ggecpplot(
  data,
  algorithm = "divisive",
  min_size = 2,
  ...,
  cptline_alpha = 1,
  cptline_color = "blue",
  cptline_type = "solid",
  cptline_linewidth = 0.5,
  cptline_size = lifecycle::deprecated(),
  index = NULL,
  show_points = NULL,
  show_line = TRUE
)

```

Arguments

<code>data</code>	A numeric vector (for univariate) or matrix/data.frame (for multivariate).
<code>algorithm</code>	Either divisive or agгло. divisive is the default.
<code>min_size</code>	Minimum number of observations between change points. By default is 2. This argument is only applied when <code>algorithm = "divisive"</code> .
<code>...</code>	Extra arguments to pass on either from <code>e.divisive()</code> or <code>e.agгло()</code> .
<code>cptline_alpha</code>	The value of alpha for the vertical changepoint line(s), default is 1, meaning no transparency.
<code>cptline_color</code>	The color for the vertical changepoint line(s), default is blue.
<code>cptline_type</code>	The linetype for the vertical changepoint line(s), default is solid.
<code>cptline_linewidth</code>	The linewidth for the vertical changepoint line(s), default is 0.5.
<code>cptline_size</code>	Deprecated. Use <code>cptline_linewidth</code> instead.
<code>index</code>	Optional. A vector of x-axis labels (e.g. dates) of the same length as data.
<code>show_points</code>	Logical. Whether to draw data points. Defaults to TRUE when <code>length(data) <= 500</code> , FALSE otherwise.
<code>show_line</code>	Logical. Whether to draw the line. Defaults to TRUE.

Value

A line plot with data points along with the vertical lines representing changepoints.

Examples

```
ggcplot(c(rnorm(100,0,1),rnorm(100,0,10)))
ggcplot(c(rnorm(100,0,1),rnorm(100,10,1)))
```

<code>glance.ggcpt</code>	<i>Glance at a ggcpt object</i>
---------------------------	---------------------------------

Description

Returns a one-row summary of a changepoint detection result.

Usage

```
## S3 method for class 'ggcpt'
glance(x, ...)
```

Arguments

<code>x</code>	A ggcpt object.
<code>...</code>	Additional arguments (ignored).

Value

A one-row tibble with columns: n, n_changepoints, method, change_in, penalty_type, penalty_value, cp_convention, total_cost (if available), runtime (NA).

idetect_wrapper	<i>Isolate-Detect wrapper</i>
-----------------	-------------------------------

Description

Wraps the IDetect package. Requires the IDetect package.

Usage

```
idetect_wrapper(x, seed = NULL, ...)
```

Arguments

x	A numeric vector.
seed	Optional seed for reproducibility.
...	Additional arguments passed to IDetect::ID().

Value

A ggcpt object.

is_ggcpt	<i>Test if an object is a ggcpt object</i>
----------	--

Description

Test if an object is a ggcpt object

Usage

```
is_ggcpt(x)
```

Arguments

x	An object to test.
---	--------------------

Value

TRUE if x inherits from ggcpt.

mosum_wrapper	<i>MOSUM wrapper — Moving Sum</i>
---------------	-----------------------------------

Description

Wraps the mosum package for moving-sum-based changepoint detection.

Usage

```
mosum_wrapper(x, G = NULL, multiscale = FALSE, seed = NULL, ...)
```

Arguments

x	A numeric vector.
G	Bandwidth. If NULL, automatically selected.
multiscale	Logical. Use multiscale MOSUM? Defaults to FALSE.
seed	Optional seed for reproducibility.
...	Additional arguments passed to mosum::mosum().

Value

A ggcpt object.

new_ggcpt	<i>Create a ggcpt object</i>
-----------	------------------------------

Description

Create a ggcpt object

Usage

```
new_ggcpt(
  changepoints = tibble::tibble(cp = integer(), cp_value = numeric()),
  segments = tibble::tibble(seg_id = integer(), start = integer(), end = integer(), n =
    integer(), param_estimate = numeric()),
  data = tibble::tibble(index = integer(), value = numeric()),
  method = character(),
  change_in = character(),
  penalty = list(type = NA_character_, value = NA_real_),
  fit = NULL,
  call = NULL,
  cp_convention = "left",
  runtime = NA_real_
)
```

Arguments

changepoints	A tibble with columns cp and cp_value.
segments	A tibble with segment information: seg_id, start, end, n, param_estimate.
data	A tibble with index and value.
method	Character. The detection method used.
change_in	Character. What was detected (e.g. "mean", "var", "meanvar").
penalty	A list with type and value.
fit	The raw upstream object.
call	The matched call.
cp_convention	Character. The convention for reporting changepoint locations: "left" (last index of left segment, used by changepoint) or "right" (first index of right segment, used by ecp). Defaults to "left".
runtime	Numeric. Elapsed detection time in seconds, if measured. Defaults to NA.

Value

An object of class ggcpt.

not_wrapper	<i>NOT wrapper — Narrowest-Over-Threshold</i>
-------------	---

Description

Wraps the not package for changepoint detection via the Narrowest-Over-Threshold method.

Usage

```
not_wrapper(x, contrast = "pcwsConstMean", seed = NULL, ...)
```

Arguments

x	A numeric vector.
contrast	Contrast type. One of "pcwsConstMean", "pcwsLinContMean", "pcwsLinMean", "pcwsConstMeanVar". Defaults to "pcwsConstMean".
seed	Optional seed for reproducibility.
...	Additional arguments passed to not::not().

Value

A ggcpt object.

print.ggcpt	<i>Print a ggcpt object</i>
-------------	-----------------------------

Description

Print a ggcpt object

Usage

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

Arguments

x	A ggcpt object.
...	Additional arguments (ignored).

signal_blocks	<i>Blocks test signal</i>
---------------	---------------------------

Description

The classic Donoho-Johnstone blocks test signal with known changepoints.

Usage

```
signal_blocks(n = 2048, seed = NULL)
```

Arguments

n	Length of the signal. Defaults to 2048.
seed	Optional seed.

Value

A tibble with columns index and value. The true_changepoints attribute contains the known changepoint locations.

References

Donoho, D. L. and Johnstone, I. M. (1994). Ideal spatial adaptation by wavelet shrinkage. *Biometrika*, 81(3), 425-455.

signal_fms	<i>FMS (Four-Metric-Segments) test signal</i>
------------	---

Description

A piecewise-constant test signal from the WBS/NOT literature.

Usage

```
signal_fms(n = 2000, seed = NULL)
```

Arguments

n	Length of the signal. Defaults to 2000.
seed	Optional seed.

Value

A tibble with columns index and value.

signal_mix	<i>Mix test signal</i>
------------	------------------------

Description

A piecewise-constant/linear signal from the literature.

Usage

```
signal_mix(n = 2000, seed = NULL)
```

Arguments

n	Length of the signal. Defaults to 2000.
seed	Optional seed.

Value

A tibble with columns index and value.

signal_stairs	<i>Stairs test signal</i>
---------------	---------------------------

Description

A monotonically stepping signal (staircase).

Usage

```
signal_stairs(n = 2000, seed = NULL)
```

Arguments

n	Length of the signal. Defaults to 2000.
seed	Optional seed.

Value

A tibble with columns index and value.

signal_teeth	<i>Teeth test signal</i>
--------------	--------------------------

Description

A piecewise-constant signal with regularly spaced changepoints.

Usage

```
signal_teeth(n = 2000, seed = NULL)
```

Arguments

n	Length of the signal. Defaults to 2000.
seed	Optional seed.

Value

A tibble with columns index and value.

stat_changepoint	<i>Changepoint detection stat</i>
------------------	-----------------------------------

Description

Runs changepoint detection inside the ggplot pipeline. Useful for quick exploration: `ggplot(df, aes(t, y)) + geom_line() + stat_changepoint(method = "pelt")`. Draws vertical lines at detected changepoint locations.

Usage

```
stat_changepoint(  
  mapping = NULL,  
  data = NULL,  
  geom = "vline",  
  position = "identity",  
  ...,  
  method = "pelt",  
  change_in = "mean",  
  na.rm = FALSE,  
  show.legend = NA  
)
```

Arguments

<code>mapping</code>	Aesthetic mappings.
<code>data</code>	A data frame.
<code>geom</code>	The geometric object to use (default: "vline").
<code>position</code>	Position adjustment.
<code>...</code>	Other arguments passed to the geom.
<code>method</code>	Detection method (passed to <code>cpt_detect</code>).
<code>change_in</code>	What to detect change in (passed to <code>cpt_detect</code>).
<code>na.rm</code>	If FALSE, missing values are removed.
<code>show.legend</code>	Whether to show legend.

Value

A ggplot layer.

summary.ggcpt	<i>Summary of a ggcpt object</i>
---------------	----------------------------------

Description

Provides a human-readable digest of a changepoint detection result, including the segment table with levels and lengths, total cost, penalty, and runtime.

Usage

```
## S3 method for class 'ggcpt'
summary(object, ...)

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

Arguments

object	A ggcpt object.
...	Additional arguments (ignored).
x	A summary.ggcpt object (for print()).

Value

A list with class `summary.ggcpt` containing the summary.

tguh_wrapper	<i>TGUH wrapper</i>
--------------	---------------------

Description

Wraps the breakfast package for Tail-Greedy Unbalanced-Haar detection.

Usage

```
tguh_wrapper(x, ...)
```

Arguments

x	A numeric vector.
...	Additional arguments passed to <code>breakfast::breakfast()</code> .

Value

A ggcpt object.

theme_ggcpt	<i>ggchangeplot theme</i>
-------------	---------------------------

Description

A minimal, publication-ready ggplot2 theme for changepoint plots.

Usage

```
theme_ggcpt(base_size = 11, base_family = "")
```

Arguments

base_size	Base font size. Defaults to 11.
base_family	Base font family. Defaults to "".

Value

A ggplot2 theme object.

Examples

```
library(ggplot2)
ggplot(mtcars, aes(wt, mpg)) + geom_point() + theme_ggcpt()
```

tidy.ggcpt	<i>Tidy a ggcpt object</i>
------------	----------------------------

Description

Returns the changepoints tibble (one row per changepoint).

Usage

```
## S3 method for class 'ggcpt'
tidy(x, ...)
```

Arguments

x	A ggcpt object.
...	Additional arguments (ignored).

Value

A tibble with columns cp, cp_value, and any method-specific columns.

wbs2_wrapper

WBS2 wrapper — Wild Binary Segmentation 2

Description

Wraps the breakfast package. Requires the breakfast package.

Usage

```
wbs2_wrapper(x, ...)
```

Arguments

x	A numeric vector.
...	Additional arguments passed to <code>breakfast::breakfast()</code> .

Value

A ggcpt object.

wbs_wrapper

WBS wrapper — Wild Binary Segmentation

Description

Wraps the wbs package for randomised changepoint detection via Wild Binary Segmentation.

Usage

```
wbs_wrapper(x, n_intervals = 5000, threshold = NULL, seed = NULL, ...)
```

Arguments

x	A numeric vector.
n_intervals	Number of random intervals. Defaults to 5000.
threshold	Manual threshold for detection. If NULL, uses the strengthened Schwarz Information Criterion (sSIC).
seed	Optional seed for reproducibility.
...	Additional arguments passed to <code>wbs::wbs()</code> .

Value

A ggcpt object.

Index

`annotate_segments`, 3
`as.data.frame`, 18
`as.data.frame.ggcpt (ggcpt_methods)`, 17
`as_tibble`, 18
`as_tibble.ggcpt (ggcpt_methods)`, 17
`augment.ggcpt`, 3
`autoplot.ggcpt`, 4, 17

`cpt_detect`, 5
`cpt_methods`, 5, 6
`cpt_metrics`, 6
`cpt_metrics_annotated`, 7
`cpt_penalty`, 8
`cpt_simulate`, 9, 9
`cpt_wrapper`, 10

`ecp_wrapper`, 11

`format.ggcpt (ggcpt_methods)`, 17
`fpop_wrapper`, 12

`geom_changepoint`, 12
`geom_cpt_ci`, 13
`geom_cpt_segment`, 13
`ggcpt_compare`, 15
`ggcpt_compare_table`, 16
`ggcpt_eval`, 17
`ggcpt_methods`, 17
`ggcptplot`, 14
`ggcpcplot`, 18
`glance.ggcpt`, 19

`idetector_wrapper`, 20
`is_ggcpt`, 20

`mosum_wrapper`, 21

`new_ggcpt`, 21
`not_wrapper`, 22

`plot.ggcpt (ggcpt_methods)`, 17

`print.ggcpt`, 23
`print.summary.ggcpt (summary.ggcpt)`, 27

`rcpt (cpt_simulate)`, 9

`signal_blocks`, 23
`signal_fms`, 24
`signal_mix`, 24
`signal_stairs`, 25
`signal_teeth`, 25
`stat_changepoint`, 26
`summary.ggcpt`, 27

`tguh_wrapper`, 27
`theme_ggcpt`, 28
`tidy.ggcpt`, 28

`wbs2_wrapper`, 29
`wbs_wrapper`, 29