

# Package ‘harbinger’

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**Title** An Unified Time Series Event Detection Framework

**Version** 1.0.737

**Description** By analyzing time series, it is possible to observe significant changes in the behavior of observations that frequently characterize events. Events present themselves as anomalies, change points, or motifs. In the literature, there are several methods for detecting events. However, searching for a suitable time series method is a complex task, especially considering that the nature of events is often unknown. This work presents Harbinger, a framework for integrating and analyzing event detection methods. Harbinger contains several state-of-the-art methods described in Salles et al. (2020) <[doi:10.5753/sbbd.2020.13626](https://doi.org/10.5753/sbbd.2020.13626)>.

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<https://cefet-rj-dal.github.io/harbinger/>

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detect	<i>Detect events in time series</i>
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**Description**

Event detection using a fitted Harbinger model

**Usage**

```
detect(obj, ...)
```

**Arguments**

obj	harbinger object
...	optional arguments.

**Value**

a data frame with the index of observations and if they are identified or not as an event, and their type

**Examples**

```
# See ?hanc_ml for an example of anomaly detection using machine learning classification
# See ?hanr_arima for an example of anomaly detection using ARIMA
# See ?hanr_fbiad for an example of anomaly detection using FBIAD
# See ?hanr_garch for an example of anomaly detection using GARCH
# See ?hanr_kmeans for an example of anomaly detection using kmeans clustering
# See ?hanr_ml for an example of anomaly detection using machine learning regression
# See ?hanr_cf_arima for an example of change point detection using ARIMA
# See ?hanr_cf_ets for an example of change point detection using ETS
# See ?hanr_cf_lr for an example of change point detection using linear regression
# See ?hanr_cf_garch for an example of change point detection using GARCH
# See ?hanr_cf_scp for an example of change point detection using the seminal algorithm
# See ?hmo_sax for an example of motif discovery using SAX
# See ?hmu_pca for an example of anomaly detection in multivariate time series using PCA
```

---

hanct_dtw	<i>Anomaly detector using DTW</i>
-----------	-----------------------------------

---

**Description**

Anomaly detection using DTW The DTW is applied to the time series. When seq equals one, observations distant from the closest centroids are labeled as anomalies. When seq is grater than one, sequences distant from the closest centroids are labeled as discords. It wraps the tslust presented in the dtwclust library.

**Usage**

```
hanct_dtw(seq = 1, centers = NA)
```

**Arguments**

```
seq           sequence size
centers       number of centroids
```

**Value**

hanct\_dtw object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 1
dataset <- har_examples$example1
head(dataset)

# setting up time series regression model
model <- hanct_dtw()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hanct\_kmeans

*Anomaly detector using kmeans*

---

**Description**

Anomaly detection using kmeans The kmeans is applied to the time series. When seq equals one, observations distant from the closest centroids are labeled as anomalies. When seq is grater than one, sequences distant from the closest centroids are labeled as discords. It wraps the kmeans presented in the stats library.

**Usage**

```
hanct_kmeans(seq = 1, centers = NA)
```

**Arguments**

seq	sequence size
centers	number of centroids

**Value**

hanct\_kmeans object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 1
dataset <- har_examples$example1
head(dataset)

# setting up time series regression model
model <- hanct_kmeans()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hanc\_ml

*Anomaly detector based on machine learning classification*

---

**Description**

Anomaly detection using daltoolbox classification. A training and test set should be used. The training set must contain labeled events. A set of preconfigured of classification methods are described in <https://cefet-rj-dal.github.io/daltoolbox/>. They include: cla\_majority, cla\_dtree, cla\_knn, cla\_mlp, cla\_nb, cla\_rf, cla\_svm

**Usage**

```
hanc_ml(model)
```

**Arguments**

model	DALToolbox classification model
-------	---------------------------------

**Value**

hanc\_ml object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 17
dataset <- har_examples$example17
dataset$event <- factor(dataset$event, labels=c("FALSE", "TRUE"))
slevels <- levels(dataset$event)

# separating into training and test
train <- dataset[1:80,]
test <- dataset[-(1:80),]

# normalizing the data
norm <- minmax()
norm <- fit(norm, train)
train_n <- transform(norm, train)

# establishing decision tree method
model <- hanc_ml(cla_dtree("event", slevels))

# fitting the model
model <- fit(model, train_n)

# evaluating the detections during testing
test_n <- transform(norm, test)

detection <- detect(model, test_n)
print(detection[(detection$event),])
```

---

hanr\_arima

*Anomaly detector using ARIMA.*

---

**Description**

Anomaly detection using ARIMA The ARIMA model adjusts to the time series. Observations distant from the model are labeled as anomalies. It wraps the ARIMA model presented in the forecast library.

**Usage**

```
hanr_arima()
```

**Value**

hanr\_arima object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 1
dataset <- har_examples$example1
head(dataset)

# setting up time series regression model
model <- hanr_arima()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hanr\_fbiad

*Anomaly detector using FBIAD*

---

**Description**

Anomaly detector using FBIAD

**Usage**

```
hanr_fbiad(sw_size = 30)
```

**Arguments**

sw\_size            Window size for FBIAD

**Value**

hanr\_fbiad object Forward and Backward Inertial Anomaly Detector (FBIAD) detects anomalies in time series. Anomalies are observations that differ from both forward and backward time series inertia [doi:10.1109/IJCNN55064.2022.9892088](https://doi.org/10.1109/IJCNN55064.2022.9892088).

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 1
dataset <- har_examples$example1
head(dataset)

# setting up time series regression model
model <- hanr_fbiad()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hanr\_garch

*Anomaly detector using GARCH*

---

**Description**

Anomaly detection using GARCH The GARCH model adjusts to the time series. Observations distant from the model are labeled as anomalies. It wraps the ugarch model presented in the rugarch library.

**Usage**

```
hanr_garch()
```

**Value**

hanr\_garch object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 14
dataset <- har_examples$example14
head(dataset)
```



```
# setting up time series regression model
model <- hanr_garch()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hanr_histogram	<i>Anomaly detector using histogram</i>
----------------	---

---

## Description

Anomaly detector using histogram

## Usage

```
hanr_histogram(density_threshold = 0.05)
```

## Arguments

`density_threshold`  
It is the minimum frequency for a bin to not be considered an anomaly. Default value is 5%.

## Value

`hanr_histogram` object histogram based method to detect anomalies in time series. Bins with smaller amount of observations are considered anomalies. Values below first bin or above last bin are also considered anomalies.>.

## Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 1
dataset <- har_examples$example1
head(dataset)

# setting up time series regression model
model <- hanr_histogram()
```

```
# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hanr\_ml

*Anomaly detector based on machine learning regression.*

---

### Description

Anomaly detection using daltoolbox regression The regression model adjusts to the time series. Observations distant from the model are labeled as anomalies. A set of preconfigured regression methods are described in <https://cefet-rj-dal.github.io/daltoolbox/>. They include: ts\_elm, ts\_conv1d, ts\_lstm, ts\_mlp, ts\_rf, ts\_svm

### Usage

```
hanr_ml(model, sw_size = 15)
```

### Arguments

model	DALToolbox regression model
sw_size	sliding window size

### Value

hanr\_ml object

### Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 1
dataset <- har_examples$example1
head(dataset)

# setting up time series regression model
model <- hanr_ml(ts_elm(ts_norm_gminmax(), input_size=4, nhid=3, actfun="purelin"))

# fitting the model
model <- fit(model, dataset$serie)
```

```
detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

han_autoencoder	<i>Anomaly detector using autoencoder</i>
-----------------	---

---

### Description

Anomaly detector using autoencoder

### Usage

```
han_autoencoder(input_size, encode_size)
```

### Arguments

input_size	Establish the input size for the autoencoder anomaly detector. It is the size of the output also.
encode_size	The encode size for the autoencoder.

### Value

han\_autoencoder object histogram based method to detect anomalies in time series. Bins with smaller amount of observations are considered anomalies. Values below first bin or above last bin are also considered anomalies.>.

### Examples

```
# setting up time series regression model
#Use the same example of hanr_fbiad changing the constructor to:
model <- han_autoencoder(3,1)
```

---

harbinger	<i>Harbinger</i>
-----------	------------------

---

### Description

Ancestor class for time series event detection

### Usage

```
harbinger()
```

**Value**

Harbinger object

**Examples**

```
# See ?hanc_ml for an example of anomaly detection using machine learning classification
# See ?hanr_arima for an example of anomaly detection using ARIMA
# See ?hanr_fbiad for an example of anomaly detection using FBIAD
# See ?hanr_garch for an example of anomaly detection using GARCH
# See ?hanr_kmeans for an example of anomaly detection using kmeans clustering
# See ?hanr_ml for an example of anomaly detection using machine learning regression
# See ?hanr_cf_arima for an example of change point detection using ARIMA
# See ?hanr_cf_ets for an example of change point detection using ETS
# See ?hanr_cf_lr for an example of change point detection using linear regression
# See ?hanr_cf_garch for an example of change point detection using GARCH
# See ?hanr_cf_scp for an example of change point detection using the seminal algorithm
# See ?hmo_sax for an example of motif discovery using SAX
# See ?hmu_pca for an example of anomaly detection in multivariate time series using PCA
```

---

har\_conv1d

*Conv1D*

---

**Description**

Creates a time series prediction object that uses the Conv1D. It wraps the pytorch library.

**Usage**

```
har_conv1d(preprocess = NA, input_size = NA, epochs = 10000L)
```

**Arguments**

preprocess	normalization
input_size	input size for machine learning model
epochs	maximum number of epochs

**Value**

a har\_conv1d object.

**Examples**

```
library(daltoolbox)
data(sin_data)
ts <- ts_data(sin_data$y, 10)
ts_head(ts, 3)
```

```
samp <- ts_sample(ts, test_size = 5)
io_train <- ts_projection(samp$train)
io_test <- ts_projection(samp$test)

model <- har_conv1d(ts_norm_gminmax(), input_size=4, epochs = 10000L)
model <- fit(model, x=io_train$input, y=io_train$output)

prediction <- predict(model, x=io_test$input[1,], steps_ahead=5)
prediction <- as.vector(prediction)
output <- as.vector(io_test$output)

ev_test <- evaluate(model, output, prediction)
ev_test
```

---

har\_eval

*Evaluation of event detection*

---

## Description

Evaluation of event detection (traditional hard evaluation)

## Usage

```
har_eval()
```

## Value

har\_eval object

## Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using the time series 14
dataset <- har_examples$example14
head(dataset)

# setting up time change point using GARCH
model <- hcp_garch()

# fitting the model
model <- fit(model, dataset$serie)

# making detections
detection <- detect(model, dataset$serie)
```

```
# filtering detected events
print(detection[(detection$event),])

# evaluating the detections
evaluation <- evaluate(har_eval(), detection$event, dataset$event)
print(evaluation$confMatrix)

# plotting the results
grf <- har_plot(model, dataset$serie, detection, dataset$event)
plot(grf)
```

---

har\_eval\_soft

*Evaluation of event detection*

---

## Description

Evaluation of event detection using SoftED [doi:10.48550/arXiv.2304.00439](https://doi.org/10.48550/arXiv.2304.00439)

## Usage

```
har_eval_soft(sw_size = 15)
```

## Arguments

sw\_size            tolerance window size

## Value

har\_eval\_soft object

## Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using the time series 14
dataset <- har_examples$example14
head(dataset)

# setting up time change point using GARCH
model <- hcp_garch()

# fitting the model
model <- fit(model, dataset$serie)

# making detections
detection <- detect(model, dataset$serie)
```

```

# filtering detected events
print(detection[(detection$event),])

# evaluating the detections
evaluation <- evaluate(har_eval_soft(), detection$event, dataset$event)
print(evaluation$confMatrix)

# plotting the results
grf <- har_plot(model, dataset$serie, detection, dataset$event)
plot(grf)

```

---

har\_examples

*Synthetic time series for event detection*


---

## Description

A list of univariate time series for event detection

- example1: a single (uncommon value) anomaly example in a stationary time series
- example2: a single (common value) anomaly example in a stationary time series
- example3: a single anomaly example in a trend time series
- example4: a change point in a time series
- example5: a change point with anomaly (uncommon value) in a time series
- example6: a change point with anomaly (common value) in a time series
- example7: a change point with anomaly (uncommon value) in a seasonal time series
- example8: a change point with anomaly (common value) in a seasonal time series
- example9: multi behavior time series (1-200: stationary, 201-400: trend, 401-600: structural break; 601-800: heteroscedasticity, 801:1000 random walk)
- example10: multiple anomalies
- example11: stationary time series
- example12: trend time series
- example13: structural break
- example14: heterocedasticity
- example15: motif
- example16: discord
- example17: repetitive anomaly
- example18: repetitive anomaly

#'

## Usage

```
data(har_examples)
```

**Format**

A list of time series.

**Source**

[Harbinger package](#)

**References**

[Harbinger package](#)

**Examples**

```
data(har_examples)
serie <- har_examples$example1
```

---

har\_examples\_multi     *Synthetic time series for event detection*

---

**Description**

A list of multivariate time series for event detection

- example1: an single multivariate anomaly

#'

**Usage**

```
data(har_examples_multi)
```

**Format**

A list of multivariate time series.

**Source**

[Harbinger package](#)

**References**

[Harbinger package](#)

**Examples**

```
data(har_examples_multi)
serie <- har_examples_multi$example1
```



---

har_lstm	<i>LSTM</i>
----------	-------------

---

## Description

Creates a time series prediction object that uses the LSTM. It wraps the pytorch library.

## Usage

```
har_lstm(preprocess = NA, input_size = NA, epochs = 10000L)
```

## Arguments

preprocess	normalization
input_size	input size for machine learning model
epochs	maximum number of epochs

## Value

a har\_lstm object.

## Examples

```
library(daltoolbox)
data(sin_data)
ts <- ts_data(sin_data$y, 10)
ts_head(ts, 3)

samp <- ts_sample(ts, test_size = 5)
io_train <- ts_projection(samp$train)
io_test <- ts_projection(samp$test)

model <- har_lstm(ts_norm_gminmax(), input_size=4, epochs = 10000L)
model <- fit(model, x=io_train$input, y=io_train$output)

prediction <- predict(model, x=io_test$input[1,], steps_ahead=5)
prediction <- as.vector(prediction)
output <- as.vector(io_test$output)

ev_test <- evaluate(model, output, prediction)
ev_test
```

---

har_plot	<i>Plot event detection on a time series</i>
----------	--

---

### Description

It accepts as harbinger, a time series, a data.frame of events, a parameter to mark the detected change points, a threshold for the y-axis and an index for the time series

### Usage

```
har_plot(  
  obj,  
  serie,  
  detection,  
  event = NULL,  
  mark.cp = TRUE,  
  ylim = NULL,  
  idx = NULL,  
  pointsize = 0.5,  
  colors = c("green", "blue", "red", "purple")  
)
```

### Arguments

obj	harbinger detector
serie	time series
detection	detection
event	events
mark.cp	show change points
ylim	limits for y-axis
idx	labels for x observations
pointsize	default point size
colors	default colors for event detection: green is TP, blue is FN, red is FP, purple means observations that are part of a sequence.

### Value

A time series plot with marked events

### Examples

```
library(daltoolbox)  
  
#loading the example database  
data(har_examples)
```

```

#Using the time series 1
dataset <- har_examples$example1
head(dataset)

# setting up time change point using GARCH
model <- hanr_arima()

# fitting the model
model <- fit(model, dataset$serie)

# making detections
detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])

# evaluating the detections
evaluation <- evaluate(har_eval_soft(), detection$event, dataset$event)
print(evaluation$confMatrix)

# plotting the results
grf <- har_plot(model, dataset$serie, detection, dataset$event)
plot(grf)

```

---

hcd\_page\_hinkley      *Adapted Page Hinkley method*

---

## Description

Change-point detection method works by computing the observed values and their mean up to the current moment [doi:10.2307/2333009](https://doi.org/10.2307/2333009).

## Usage

```

hcd_page_hinkley(
  min_instances = 30,
  delta = 0.005,
  threshold = 50,
  alpha = 1 - 1e-04
)

```

## Arguments

min_instances	The minimum number of instances before detecting change
delta	The delta factor for the Page Hinkley test
threshold	The change detection threshold (lambda)
alpha	The forgetting factor, used to weight the observed value and the mean

**Value**

hcp\_page\_hinkley object

**Examples**

```
library("daltoolbox")

n <- 100 # size of each segment
serie1 <- c(sin((1:n)/pi), 2*sin((1:n)/pi), 10 + sin((1:n)/pi),
           10-10/n*(1:n)+sin((1:n)/pi)/2, sin((1:n)/pi)/2)
serie2 <- 2*c(sin((1:n)/pi), 2*sin((1:n)/pi), 10 + sin((1:n)/pi),
             10-10/n*(1:n)+sin((1:n)/pi)/2, sin((1:n)/pi)/2)
data <- data.frame(serie1, serie2) #'
event <- rep(FALSE, nrow(data))

model <- fit(hcd_page_hinkley(threshold=3), data)
detection <- detect(model, data)
print(detection[(detection$event),])
```

---

hcp\_amoc

*At most one change (AMOC) method*

---

**Description**

Change-point detection method that focus on identify one change point in mean/variance [doi: 10.1093/biomet/57.1.1](https://doi.org/10.1093/biomet/57.1.1). It wraps the amoc implementation available in the changepoint library.

**Usage**

```
hcp_amoc()
```

**Value**

hcp\_amoc object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
head(dataset)

# setting up time series regression model
model <- hcp_amoc()
```

```
# fitting the model
model <- fit(model, dataset$serie)

# execute the detection method
detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hcp\_binseg

*Binary segmentation (BinSeg) method*

---

### Description

Change-point detection method that focus on identify change points in mean/variance [doi:10.2307/2529204](https://doi.org/10.2307/2529204). It wraps the BinSeg implementation available in the changepoint library.

### Usage

```
hcp_binseg(Q = 2)
```

### Arguments

Q                    The maximum number of change-points to search for using the BinSeg method

### Value

hcp\_binseg object

### Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
head(dataset)

# setting up time series regression model
model <- hcp_binseg()

# fitting the model
model <- fit(model, dataset$serie)

# execute the detection method
detection <- detect(model, dataset$serie)
```

```
# filtering detected events
print(detection[(detection$event),])
```

---

hcp\_cf\_arima

*Change Finder using ARIMA*

---

## Description

Change-point detection is related to event/trend change detection. Change Finder ARIMA detects change points based on deviations relative to ARIMA model [doi:10.1109/TKDE.2006.1599387](https://doi.org/10.1109/TKDE.2006.1599387). It wraps the ARIMA model presented in the forecast library.

## Usage

```
hcp_cf_arima(sw_size = NULL)
```

## Arguments

sw\_size            Sliding window size

## Value

hcp\_cf\_arima object

## Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
head(dataset)

# setting up change point method
model <- hcp_cf_arima()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

`hcp_cf_ets`*Change Finder using ETS*

---

**Description**

Change-point detection is related to event/trend change detection. Change Finder ETS detects change points based on deviations relative to trend component (T), a seasonal component (S), and an error term (E) model [doi:10.1109/TKDE.2006.1599387](https://doi.org/10.1109/TKDE.2006.1599387). It wraps the ETS model presented in the forecast library.

**Usage**

```
hcp_cf_ets(sw_size = 7)
```

**Arguments**

```
sw_size      Sliding window size
```

**Value**

```
hcp_cf_ets object
```

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
head(dataset)

# setting up change point method
model <- hcp_cf_ets()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

`hcp_cf_lr`*Change Finder using LR*

---

**Description**

Change-point detection is related to event/trend change detection. Change Finder LR detects change points based on deviations relative to linear regression model [doi:10.1109/TKDE.2006.1599387](https://doi.org/10.1109/TKDE.2006.1599387).

**Usage**

```
hcp_cf_lr(sw_size = 30)
```

**Arguments**

`sw_size`            Sliding window size

**Value**

hcp\_cf\_lr object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
head(dataset)

# setting up change point method
model <- hcp_cf_lr()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```



---

hcp_chow	<i>Chow test method</i>
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**Description**

Change-point detection method that focus on identifying structural changes [doi:10.18637/jss.v007.i02](https://doi.org/10.18637/jss.v007.i02). It wraps the Fstats and breakpoints implementation available in the strucchange library.

**Usage**

```
hcp_chow()
```

**Value**

hcp\_chow object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
head(dataset)

# setting up change point method
model <- hcp_chow()

# fitting the model
model <- fit(model, dataset$serie)

# execute the detection method
detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hcp_garch	<i>Change Finder using GARCH</i>
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**Description**

Change-point detection is related to event/trend change detection. Change Finder GARCH detects change points based on deviations relative to linear regression model [doi:10.1109/TKDE.2006.1599387](https://doi.org/10.1109/TKDE.2006.1599387). It wraps the GARCH model presented in the rugarch library.

**Usage**

```
hcp_garch(sw_size = 5)
```

**Arguments**

```
sw_size          Sliding window size
```

**Value**

```
hcp_garch object
```

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 14
dataset <- har_examples$example14
head(dataset)

# setting up change point method
model <- hcp_garch()

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hcp\_gft

*Generalized Fluctuation Test (GFT)*

---

**Description**

GFT detection method focuses on identifying structural changes [doi:10.18637/jss.v007.i02](https://doi.org/10.18637/jss.v007.i02). It wraps the breakpoints implementation available in the strucchange library.

**Usage**

```
hcp_gft()
```

**Value**

```
hcp_chow object
```

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
head(dataset)

# setting up change point method
model <- hcp_gft()

# fitting the model
model <- fit(model, dataset$serie)

# execute the detection method
detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hcp\_pelt

*Pruned exact linear time (PELT) method*

---

**Description**

Change-point detection method that focus on identifying multiple exact change points in mean/variance [doi:10.1080/01621459.2012.737745](https://doi.org/10.1080/01621459.2012.737745). It wraps the BinSeg implementation available in the change-point library.

**Usage**

```
hcp_pelt()
```

**Value**

hcp\_pelt object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
```

```
head(dataset)

# setting up change point method
model <- hcp_pelt()

# fitting the model
model <- fit(model, dataset$serie)

# execute the detection method
detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hcp\_scp

*Seminal change point*

---

### Description

Change-point detection is related to event/trend change detection. Seminal change point detects change points based on deviations of linear regression models adjusted with and without a central observation in each sliding window <10.1145/312129.312190>.

### Usage

```
hcp_scp(sw_size = 30)
```

### Arguments

sw\_size            Sliding window size

### Value

hcp\_scp object

### Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 6
dataset <- har_examples$example6
head(dataset)

# setting up change point method
model <- hcp_scp()
```

```
# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hmo\_base36

*Motif discovery using base36*

---

## Description

Motif discovery using base36 [doi:10.1007/s10618-007-0064-z](https://doi.org/10.1007/s10618-007-0064-z)

## Usage

```
hmo_base36(a, w, qtd)
```

## Arguments

a	alphabet size
w	word size
qtd	number of occurrences to be classified as motifs

## Value

hmo\_base36 object

## Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 15
dataset <- har_examples$example15
head(dataset)

# setting up motif discovery method
model <- hmo_base36(37, 3, 3)

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)
```

```
# filtering detected events
print(detection[(detection$event),])
```

---

hmo\_mp

*Motif discovery using Matrix Profile*

---

## Description

Motif discovery using Matrix Profile [doi:10.32614/RJ-2020-021](https://doi.org/10.32614/RJ-2020-021)

## Usage

```
hmo_mp(mode = "stamp", w, qtd)
```

## Arguments

mode	mode of computing distance between sequences. Available options include: "stomp", "stamp", "simple", "mstomp", "scrimp", "valmod", "pmp"
w	word size
qtd	number of occurrences to be classified as motifs

## Value

hmo\_mp object

## Examples

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 15
dataset <- har_examples$example15
head(dataset)

# setting up motif discovery method
model <- hmo_mp("stamp", 4, 3)

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

hmo_sax	<i>Motif discovery using SAX</i>
---------	----------------------------------

---

**Description**

Motif discovery using SAX [doi:10.1007/s10618-007-0064-z](https://doi.org/10.1007/s10618-007-0064-z)

**Usage**

```
hmo_sax(a, w, qtd)
```

**Arguments**

a	alphabet size
w	word size
qtd	number of occurrences to be classified as motifs

**Value**

hmo\_sax object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples)

#Using example 15
dataset <- har_examples$example15
head(dataset)

# setting up motif discovery method
model <- hmo_sax(26, 3, 3)

# fitting the model
model <- fit(model, dataset$serie)

detection <- detect(model, dataset$serie)

# filtering detected events
print(detection[(detection$event),])
```

---

`hmu_pca`*Multivariate anomaly detector using PCA*

---

**Description**

Multivariate anomaly detector using PCA [doi:10.1016/0098-3004\(93\)90090-R](https://doi.org/10.1016/0098-3004(93)90090-R)

**Usage**

```
hmu_pca()
```

**Value**

hmu\_pca object

**Examples**

```
library(daltoolbox)

#loading the example database
data(har_examples_multi)

#Using the time series 9
dataset <- har_examples_multi$example1
head(dataset)

# establishing hmu_pca method
model <- hmu_pca()

# fitting the model using the two columns of the dataset
model <- fit(model, dataset[,1:2])

# making detections using hmu_pca
detection <- detect(model, dataset[,1:2])

# filtering detected events
print(detection[(detection$event),])

# evaluating the detections
evaluation <- evaluate(model, detection$event, dataset$event)
print(evaluation$confMatrix)
```



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