

R Package Development by Means of Literate Programming (**noweb**)

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1 Introduction

2 Detecting Peaks/Troughs

2.1 Notation

A uniformly sampled time series $\mathbf{y} = \{y_1, \dots, y_i, \dots, y_T\}$ with T data points is considered. The detection of peak/trough points is achieved by a function $S(i, y_i, T)$ that returns for data point y_i a score value.¹ If this score value surpasses a user-provided threshold value θ , /i.e./, $S(i, y_i, T) \geq \theta$ then the point is considered as a local peak/trough.

Furthermore, in case local peak/trough points appear closely together with respect to time (clustered), then these points can be classified as a burst or bust, respectively.

2.2 Algorithms

In [Palshikar \(2009\)](#) five different score functions S have been suggested. All have in common, that a centred window of size $2*k+1$ around y_i is considered. That is, for a positive integer k the k right neighbours $N^+(i, k, T) = \{y_{i+1}, \dots, y_{i+k}\}$ and the k left neighbours $N^-(i, k, T) = \{y_{i-k}, \dots, y_{i-1}\}$ are employed for assessing y_i as a local peak/trough. The union of $N^-(i, k, T)$ and $N^+(i, k, T)$ is defined as $N(i, k, T) = N^-(i, k, T) \cup N^+(i, k, T)$ and if the centre point is included as $N'(i, k, T) = N^-(i, k, T) \cup y_i \cup N^+(i, k, T)$.

The first function, S_1 , computes the score value as the average of the maximum differences between y_i with its left and right neighbours. The function is defined as:

$$S_1 = \frac{\max(y_i - y_{i-1}, \dots, y_i - y_{i-k}) + \max(y_i - y_{i+1}, \dots, y_i - y_{i+k})}{2} \quad (1)$$

The equation (1) can be casted in R as:

```
1 <score-maxdiff 1>≡  
  scmaxdiff <- function(x, k){
```

¹It suffices to provide a score function for peaks only. Trough points can be detected by using the negative values of the series \mathbf{y} .

```

    cp <- k + 1L
    lmax <- max(x[cp] - head(x, k))
    rmax <- max(x[cp] - tail(x, k))
    (lmax + rmax) / 2.0
}

```

Defines:

`scmaxdiff`, used in chunk 4a.

Instead of using the maximum differences of y_i with its k left and right neighbours as in (1), an alternative is to compute the mean differences and evaluate the average thereof:

$$S_2 = \frac{\frac{(y_i - y_{i-1}, \dots, y_i - y_{i-k})}{k} + \frac{(y_i - y_{i+1}, \dots, y_i - y_{i+k})}{k}}{2} \quad (2)$$

This equation can be casted in R as:

```

2a  <score-diffmean 2a>≡
    scdiffmean <- function(x, k){
      cp <- k + 1L
      ldmean <- x[cp] - mean(head(x, k))
      rdmean <- x[cp] - mean(tail(x, k))
      (ldmean + rdmean) / 2.0
    }

```

Defines:

`scdiffmean`, used in chunk 4a.

Another variation of score computation that has been proposed by [Palshikar \(2009\)](#) is to consider the differences to the mean of the k left and right neighbours, that is:

$$S_3 = \frac{(y_i - \frac{(y_{i-1}, \dots, y_{i-k})}{k}) + (y_i - \frac{(y_{i+1}, \dots, y_{i+k})}{k})}{2} \quad (3)$$

The equation (3) can be casted as R function `scavgdiff` for instance as follows:

```

2b  <score-avgdiff 2b>≡
    scavgdiff <- function(x, k){
      cp <- k + 1L
      lmean <- mean(x[cp] - head(x, k))
      rmean <- mean(x[cp] - tail(x, k))
      (lmean + rmean) / 2.0
    }

```

Defines:

`scavgdiff`, used in chunk 4a.

The fourth proposed score function differs from the previous three in the sense that it does take explicitly the differences between y_i and its neighbours explicitly into account, but tries to capture its information content by means of relative entropy. The entropy of a vector A with elements $A = \{a_1, \dots, a_m\}$ is given as:

$$H_w(A) = \sum_{i=1}^M (-p_w(a_i) \log(p_w(a_i))) \quad (4)$$

where $p_w(a_i)$ is an estimate of the density value at a_i . The score function is now based on computing the entropies of $H(N((k, i, T)))$ and $H(N'(k, i, T))$. Hereby, the densities can be determined by means of a kernel density estimator. The score function is then defined as the difference of the entropies:

$$S_4 = H(N((k, i, T))) - H(N'((k, i, T))) \quad (5)$$

This concept is implemented in the function `scentropy()`. The empirical density is computed by calling `density()`. The ellipsis argument of `scentropy()` is passed down to this function and hereby allowing the user to employ other than the default arguments of `density()`.

3 *⟨score-entropy 3⟩*≡
`scentropy <- function(x, k, ...){`
`cp <- k + 1L`
`dfull <- density(x, ...)$y`
`hfull <- sum(-dfull * log(dfull))`
`dexct <- density(x[-cp], ...)$y`
`hexct <- sum(-dexct * log(dexct))`
`hfull - hexct`
`}`

Defines:

`scentropy`, used in chunk 4a.

2.3 Combining score methods

4a *<score-wrapper 4a>*≡

```

score <- function(x, k,
                  scoreby = c("vote", "avg", "diff", "max", "ent",
                              "ttype", "hybrid"),
                  tval = 1.0, confby = 3, ...){
  scoreby <- match.arg(scoreby)
  ans <- switch(scoreby,
               vote = scvote(x, k, tval, confby, ...),
               avg = scavdiff(x, k),
               diff = scdiffmean(x, k),
               max = scmaxdiff(x, k),
               ent = scentropy(x, k, ...),
               ttype = scttype(x, k, tval),
               hybrid = schybrid(x, k, tval, ...)
               )
  ans
}

```

Defines:

`score`, never used.

Uses `scavdiff` 2b, `scdiffmean` 2a, `scentropy` 3, and `scmaxdiff` 1.

4b *<score-roxygen 4b>*≡

```

#' Basic scoring methods for local minima and maxima
#'
#' These are basic functions for evaluating the centre
#' point of a time series as local minimum or maximum.
#' Hereby, a score value is computed according to various methods.
#' If the score is positive, the centre point is tentatively
#' classified as a local peak.
#' Incidentally, negative scores indicate a local minima.
#'
#' @param x \code{numeric}, vector of length \code{2 * k + 1}.
#' @param k \code{integer}, the count of left/right neighbours.
#' @param scoreby \code{character}, the scoring method to be used.
#' @param tval \code{numeric}, factor for standard deviation band
#' if \code{scoreby = 'ttype'}.
#' @param confby \code{integer}, count of minimum vote,
#' values in the set \code{3:5}.
#' @param ... ellipsis argument.
#'
#' @name score
#' @family scores
#' @return \code{numeric}, the score value.
NULL

#' @rdname score
#' @export

```

@

The content/structure of the file `score.R` is given as:

```
5a  <score.R 5a>≡
    <score-roxygen 4b>
    <score-wrapper 4a>
    #' @rdname score
    <score-mardiff 1>
    #' @rdname score
    <score-diffmean 2a>
    #' @rdname score
    <score-avgdiff 2b>
    #' @rdname score
    <score-entropy 3>
```

This code is written to file `score.R`.

@

Within this file, all score-related methods and the wrapper-function `score()` is included. The function definitions are interspersed with the roxygen tags, which will be parsed to the Rd-file `score.Rd`.

3 Package structure

3.1 Preliminaries

First, a skeleton of the package

```
5b  <DESCRIPTION.R 5b>≡
    Package: hiker
    Title: Local Peak and Trough of a Time Series
    Version: 0.0.0.9000
    Authors@R: person("Bernhard", "Pfaff", email = "bernhard@pfaffikus.de",
                      role = c("aut", "cre"))
    Description: Methods for detecting local peaks and troughs of a time series.
    Depends: R (>= 3.3.1), zoo, methods
    License: GPL-3
    Encoding: UTF-8
    LazyData: true
```

This code is written to file `DESCRIPTION.R`.

@

3.2 Import directives and S4-classes

```
6 <Allclasses.R 6>≡
  #' @import methods
  NULL
  #' @import zoo
  NULL
  #' @importFrom stats density sd na.omit start end smooth
  NULL
  #' @importFrom utils head tail
  NULL

  # Setting old (aka S3) classes
  setOldClass("zoo")

  #' S4 class \code{HikeR}
  #'
  #' Formal class for classifying local minima and maxima
  #' of a time series.
  #'
  #' @slot ys \code{zoo}, time series with associated scores.
  #' @slot k \code{integer}, count of left/right neighbours around centre point.
  #' @slot scoreby \code{character}, scoring method.
  #' @slot yname \code{character}, name of the series.
  #' @exportClass HikeR
  setClass("HikeR", slots = list(ys = "zoo",
                                k = "integer",
                                scoreby = "character",
                                yname = "character"))

  #' S4 class \code{PTBB}
  #'
  #' Formal class for peaks, troughs, burst, busts and
  #' intermittent phase of a time series.
  #'
  #' @slot pt \code{zoo}, logical: indicating peak/trough points.
  #' @slot type \code{character}, type of point/phase.
  #' @slot h \code{numeric}, the threshold for score evaluation.
  #' @exportClass PTBB
  setClass("PTBB", slots = list(pt = "zoo",
                                type = "character",
                                h = "numeric"))
```

This code is written to file `Allclasses.R`.

@

4 Chunk Index

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5 Identifier Index

scavgdiff: [2b](#), [4a](#)
scdiffmean: [2a](#), [4a](#)
scentropy: [3](#), [4a](#)
scmaxdiff: [1](#), [4a](#)
score: [4a](#)

References

Palshikar, G. (2009). Simple algorithms for peak detection in time-series. In *First Int. Conf. Advanced Data Analysis, Business Analytics and Intelligence*, Ahmedabad, India.