Changepoint Analysis Using R

Robert Maidstone





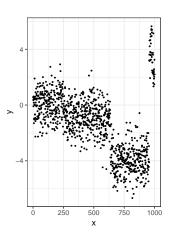
17 October, 2018

What are Changepoints?

change-point *n.* (*a*) a point at which something changes; the point on the scale of a measuring device representing this (now *rare*); (*b*) Statistics the point at which the probability distribution of a sequence of random variables changes (frequently attributive, as **change point analysis**, **change point problem**, etc.).

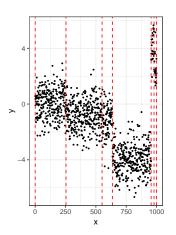
Change in mean

```
set.seed(14)
m<-5
n<-1000
true_cps <- c(0,sort(sample(1:(n-1),m)),n)
means <- rnorm(m+1,0,4)
y<-c()
for(i in 1:(m+1)){
    j <- (true_cps[i]+1):true_cps[i+1]
    y[j]<-rnorm(length(j),means[i],1)
}</pre>
```



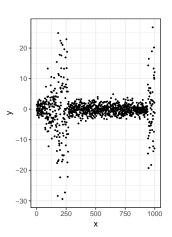
Change in mean

```
set.seed(14)
m<-5
n<-1000
true_cps <- c(0,sort(sample(1:(n-1),m)),n)
means <- rnorm(m+1,0,4)
y<-c()
for(i in 1:(m+1)){
    j <- (true_cps[i]+1):true_cps[i+1]
    y[j]<-rnorm(length(j),means[i],1)
}</pre>
```



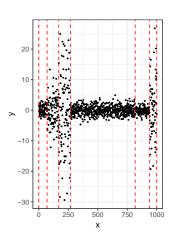
Change in variance

```
set.seed(12)
m<-5
n<-1000
true_cps <- c(0,sort(sample(1:(n-1),m)),n)
sd <- runif(m+1,1,20)
y<-c()
for(i in 1:(m+1)){
    j <- (true_cps[i]+1):true_cps[i+1]
    y[j]<-rnorm(length(j),0,sd[i])
}</pre>
```



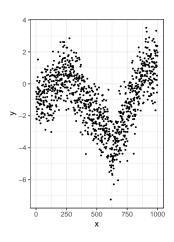
Change in variance

```
set.seed(12)
m<-5
n<-1000
true_cps <- c(0,sort(sample(1:(n-1),m)),n)
sd <- runif(m+1,1,20)
y<-c()
for(i in 1:(m+1)){
   j <- (true_cps[i]+1):true_cps[i+1]
   y[j]<-rnorm(length(j),0,sd[i])
}</pre>
```



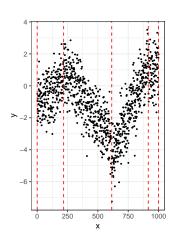
Change in trend

```
set.seed(110)
m < -3
n < -1000
true_cps \leftarrow c(0, sort(sample(1:(n-1), m)), n)
slope \leftarrow rnorm(m+1,0,.01)
intercept <- rnorm(1,0,1)</pre>
y<-c()
for(i in 1:(m+1)){
  j <- (true_cps[i]+1):true_cps[i+1]</pre>
  if(i==1){
  for(jind in j){
    y[jind] <- intercept+(jind-true_cps[i])*
       slope[i] + rnorm(1,0,1)
  }else{
    for(jind in j){
      y[jind]<-y[j[1]-1]+(jind-true_cps[i])*</pre>
         slope[i] + rnorm(1,0,1)
```

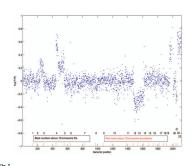


Change in trend

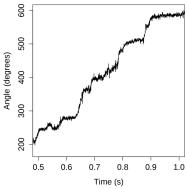
```
set.seed(110)
m < -3
n < -1000
true_cps \leftarrow c(0, sort(sample(1:(n-1), m)), n)
slope \leftarrow rnorm(m+1,0,.01)
intercept <- rnorm(1,0,1)</pre>
y<-c()
for(i in 1:(m+1)){
  j <- (true_cps[i]+1):true_cps[i+1]</pre>
  if(i==1){
  for(jind in j){
    y[jind] <- intercept+(jind-true_cps[i])*
       slope[i] + rnorm(1,0,1)
  }else{
    for(jind in j){
      y[jind]<-y[j[1]-1]+(jind-true_cps[i])*</pre>
         slope[i] + rnorm(1,0,1)
```



Real World Examples



(a) Copy number at genomic positions in a human breast tumor sample (Chen & Wang, 2009).



(b) Rotation of a bacterial flagella motor (Maidstone, 2016).

Genome of the breast tumor S1514 [39].

Many changepoint methods

Many different changepoint algorithms exist

- Exhaustive Search
- Optimal Partitioning
- PELT
- FPOP (and R-FPOP)
- CROPS
- Segment Neighbourhood Search
- pDPA
- SNIP

- Binary Segmentation
- WBS
- CBS
- SMUCE
- SMOP
- ED-PELT
- E-Divisive
- ECP

Binary Segmentation

Input: A set of data of the form, $(y_1, y_2, ..., y_n)$ where $y_i \in \mathbb{R}$.

A test statistic on the data $\Gamma(\cdot)$,

An estimator of changepoint location $\hat{\tau}(\cdot)$,

A rejection threshold c.

Initialise: Let
$$CP = \emptyset$$
 and $S = \{[1,n]\}$;

Iterate: while $S \neq \emptyset$ do

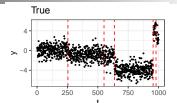
- Choose an element of S; denote this element as [s,t];
- if $\Gamma(\mathbf{y}_{s:t}) < c$ then
 - remove [s,t] from S
- if $\Gamma(\mathbf{y}_{s:t}) \geq c$ then
 - remove [s,t] from S;
 - calculate $r = \hat{\tau}(\mathbf{y}_{s:t}) + s 1$, and add r to \mathcal{CP} ;
 - if $r \neq s$ add [s,r] to S;
 - \blacksquare if $r \neq t-1$ add [r+1,t] to S;

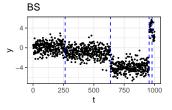
Output: The changepoints recorded in \mathcal{CP} .

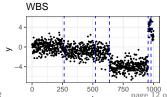
Binary Segmentation using "wbs"

Baranowski and Fryzlewicz (2015)

```
library(wbs)
sbs(y) -> sbs.out
s.cpt <- changepoints(sbs.out,th =4)
s.cpt$cpt.th[[1]] %>% sort
## [1] 264 637 955 979
wbs(v,
    M=5000) -> wbs.out
w.cpt <- changepoints(wbs.out,th=4)</pre>
w.cpt$cpt.th[[1]] %>% sort
## [1] 264 521 637 955 979
```







Cost function representation

Most changepoint detection methods boil down to minimising the sum of some cost function, $\mathcal{C}(\cdot)$, over the segments.

$$\min_{ au_{1:m},m} \left[\sum_{j=0}^{m+1} \mathcal{C}(\mathbf{y}_{ au_j+1: au_{j+1}})
ight]$$

This cost function could be a number of things:

- 1 Negative log-likelihood,
- 2 Negative posterior,
- 3 Minimum Description Length.

Dynamic Programming Methods

Optimal Partitioning: Optimisation based sum of optimal up to last changepoint and the cost between last changepoint and current time (plus a penalty to avoid over fitting).

$$F(\tau^*) = \min_{0 < \tau < \tau^*} [F(\tau) + \mathcal{C}(\mathbf{y}_{(\tau+1):\tau^*}) + \beta].$$

Segment Neighbourhood Search: Optimisation for k segments based on optimal for k-1 segments plus cost for new segment.

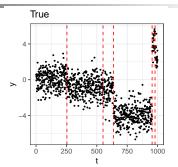
$$q_{1,j}^k = \min_{v \in \{1, \dots, j-1\}} [q_{1,v}^{k-1} + q_{v+1,j}^1].$$

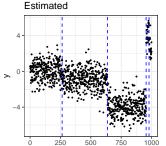
The changepoint package

- Functions for changepoint analysis.
- Can use various changepoint detection methods:
 - Binary Segmentation
 - AMOC
 - Segment Neighbourhood
 - PELT (Optimal Partitioning)
- With various choices of penalty function/value.
- And either a Gaussian or CUSUM test statistic.
- Three headliner functions:
 - cpt.mean
 - cpt.var
 - cpt.meanvar
- Authors: Rebecca Killick, Kaylea Haynes, Idris Eckley, Paul Fearnhead, Jamie Lee.

Change in mean using "changepoint"

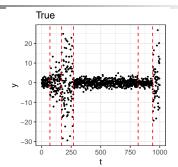
```
changepoint::cpt.mean(y_mean,
                      penalty = "BIC",
                      method = "PELT"
                      ) -> cpt_object #S4
cpt_object %>% changepoint::likelihood()
##
        -2*logLik -2*Loglike+pen
         2839.115
                        2894.377
##
cpt_object %>% changepoint::ncpts()
## [1] 4
cpt_object %>% changepoint::cpts() -> est_cps
qpcR:::rbind.na(tru_cps,est_cps)
##
           [,1] [,2] [,3] [,4] [,5]
## tru_cps 254 551 637 955
                                979
## est cps 264
                 637
                      955
                           979
                                 NA
```

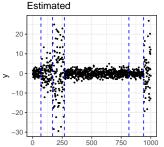




Change in variance using "changepoint"

```
changepoint::cpt.var(y_var,
                     penalty = "BIC".
                     method = "PELT"
                     ) -> cpt_object #S4
cpt_object %>% changepoint::likelihood()
##
        -2*logLik -2*Loglike+pen
                        4464.746
         4395.668
##
cpt_object %>% changepoint::ncpts()
## [1] 5
cpt_object %>% changepoint::cpts() -> est_cps
qpcR:::rbind.na(tru_cps,est_cps)
##
           [,1] [,2] [,3] [,4] [,5]
           70
                 169
                      269 817
                                 940
## tru cps
## est cps
             70
                 169
                      268
                           815
                                940
```



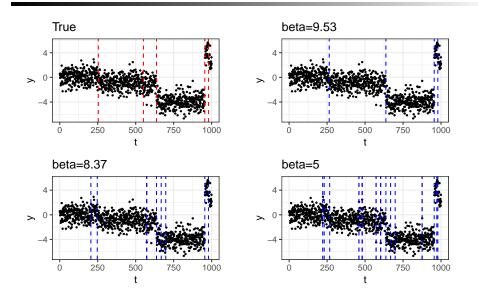


Penalty Choice

Optimal Partitioning: Optimisation based sum of optimal up to last changepoint and the cost between last changepoint and current time (plus a penalty to avoid over fitting).

$$F(\tau^*) = \min_{0 \le \tau < \tau^*} [F(\tau) + \mathcal{C}(\mathbf{y}_{(\tau+1):\tau^*}) + \beta].$$

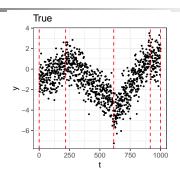
Penalty Choice

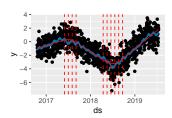


CROPS

```
cpt.mean(y,penalty = "CROPS",
         pen.value=c(5,1000),
         method="PELT",
                                                     1100
                                                   penalised_cost
         class=FALSE) -> crops.out
  [1] "Maximum number of runs of algorithm = 19
                                                     1000
       "Completed runs = 2"
   Г17
      "Completed runs = 3"
   [1]
      "Completed runs = 5"
      "Completed runs = 7"
                                                      900
  [1] "Completed runs = 11"
                                                                  10
                                                                        15
                                                                               20
## [1] "Completed runs = 16"
                                                            numberofchangepoints
  crops.out$cpt.out %>% .[,1:5]
  ##
                                [,1]
                                            [,2]
                                                       [.3]
                                                                   Γ.47
                                                                               [.5]
                             5.0000
                                       5.199941
                                                   5.642361
                                                              6.626451
                                                                          8.256657
  ## beta_interval
  ## numberofchangepoints 20.0000
                                      18.000000 16.000000 12.000000
                                                                         11,000000
  ## penalised_cost
                           890.7315 901.131370 912.416092 938.921896 947.178554
```

Change in trend with Prophet





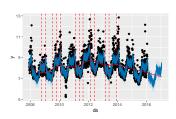
Peyton Manning (Prophet example)

```
df<-read.csv(
   "~/changepointsinR/
   example_wp_log_peyton_manning.csv")
m <- prophet(df)

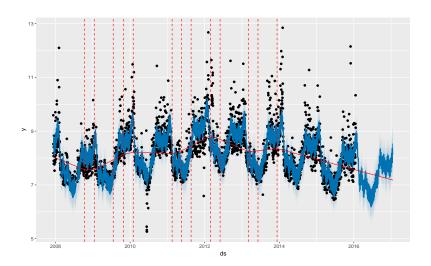
future <- make_future_dataframe(
   m,
   periods = 365)

forecast <- predict(m, future)

plot(m, forecast)+add_changepoints_to_plot(m)</pre>
```

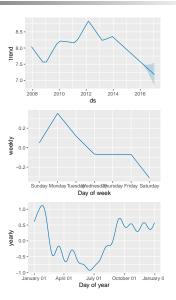


Peyton Manning (Prophet example)



Peyton Manning (Prophet example)

prophet_plot_components(m, forecast)



Discussion

- Many changepoint methods avaliable
 - Not discussed; Multivariate, online changepoint detection, non parametric (changepoint.np,ecp) and more.
- changepoint and wbs packages are a good place to start.
- Computational time can be a major issue in changepoint detection, meaning bespoke solutions can be best.
- Code not always in the best form, but big forcasting packages such as prophet use changepoints too.