# example\_univar

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This is a simple guide for offline changepoint detection on univariate mean.

There are 3 methods implemented for univariate mean changepoint detection:

- 1. *DP.univar*: perform dynamic programming for univariate mean changepoint detection through l0 penalty.
  - CV.search.DP.univar: perform grid search to select the tuning parameter through Cross-Validation.
- 2. BS.univar: perform standard binary segmentation for univariate mean changepoint detection.
- 3. WBS.univar: perform wild binary segmentation for univariate mean changepoint detection.

In addition, function local refine univar performs local refinement for an initial changepoint estimation.

#### Simulate data

```
library(changepoints)
## Loading required package: gglasso
## Loading required package: glmnet
## Loading required package: Matrix
## Loaded glmnet 4.1-2
## Loading required package: penalized
## Loading required package: survival
##
## Attaching package: 'survival'
## The following object is masked from 'package:gglasso':
##
##
       colon
## Welcome to penalized. For extended examples, see vignette("penalized").
## Loading required package: ks
## Loading required package: MASS
delta = 10 # 2*delta represents the minimum gap between boundaries
sigma2 = 1 # error variance
set.seed(0)
y = c(rep(0, 50), rep(1, 50), rep(0, 50), rep(1, 50)) + rnorm(200, mean = 0, sd = sqrt(sigma2)) # univa
```

```
cpt_true = c(50, 100, 150)
n = length(y) # sample size
```

### Perform dynamic programming

```
gamma.set = c(0.01, 0.5, 1, 5, 10, 50) # a set of tuning parameters for DP
DP_result = CV.search.DP.univar(y, gamma.set, delta = 5) # grid search through cross-validation
##
                         [,2]
                                   [,3]
                                            [,4]
                                                       [,5]
               [,1]
                                                                [,6]
## cpt hat
              numeric,7 numeric,6 numeric,3 numeric,3 numeric,0
## K_hat
              7
                        6
                                  6
                                            3
                                                      3
                                                                0
## test error 97.19456
                        97.13118 97.13118 88.23253
                                                      88.23253
                                                                107.8672
## train_error 81.46384 81.54316 81.54316 87.80528 87.80528 128.2893
min_idx = which.min(DP_result$test_error) # select gamma achieves the minimum validation error
cpt_DP_hat = unlist(DP_result$cpt_hat[[min_idx]]) # estimated changepoints by DP
cpt_DP_hat
## [1] 47 101 147
Hausdorff.dist(cpt_DP_hat, cpt_true)
## [1] 3
cpt_DPlr_hat = local.refine.univar(cpt_DP_hat, y, w = 1/3) # perform local refinement
cpt_DPlr_hat
## [1] 47 101 150
Hausdorff.dist(cpt_DPlr_hat, cpt_true)
## [1] 3
```

## Perform standard binary segmentation

```
tau BS = 3 # threshold parameter for BS
BS_result = threshold.BS(BS.univar(y, 1, n, delta), tau_BS)
BS_result$BS_tree_trimmed # trace BS
## [[1]]
     current parent location
                                 value
## 1
           1
                  1
                         150 5.190086
##
## [[2]]
     current parent location
## 1
           1
                  1
                         101 4.152929
##
## [[3]]
##
     current parent location
                                value
## 1
           1
                          47 5.143882
##
## [[4]]
## [1] current parent
                         location value
## <0 rows> (or 0-length row.names)
##
```

```
## [[5]]
## [1] current parent
                       location value
## <0 rows> (or 0-length row.names)
## [[6]]
## [1] current parent
                        location value
## <0 rows> (or 0-length row.names)
BS result$change points
##
    location
                value level
         150 5.190086
## 2
         101 4.152929
## 3
          47 5.143882
                           3
cpt_BS_hat = sort(BS_result$change_points[,1]) # estimated changepoints by BS
cpt_BS_hat
## [1] 47 101 150
Hausdorff.dist(cpt_BS_hat, cpt_true)
cpt_BSlr_hat = local.refine.univar(cpt_BS_hat, y, w = 1/3) # perform local refinement
cpt_BSlr_hat
## [1] 47 101 150
Hausdorff.dist(cpt_BSlr_hat, cpt_true)
## [1] 3
```

### Perform wild binary segmentation

```
tau_WBS = 3 # threshold parameter for WBS
intervals = WBS.intervals(M = 300, lower = 1, upper = n) # generate random intervals for WBS
WBS_result = threshold.BS(WBS.univar(y, 1, n, intervals$Alpha, intervals$Beta, delta), tau_WBS)
WBS_result$BS_tree_trimmed # trace BS
## [[1]]
   current parent location
## 1
         1
                       150 6.405863
             1
##
## [[2]]
## current parent location
## 1
         1
               1
                     101 6.106824
##
## [[3]]
   current parent location
                              value
## 1
          1
             1
                    47 5.345979
##
## [[4]]
## [1] current parent
                       location value
## <0 rows> (or 0-length row.names)
##
## [[5]]
```

```
## [1] current parent location value
## <0 rows> (or 0-length row.names)
##
## [[6]]
## [1] current parent
                        location value
## <0 rows> (or 0-length row.names)
WBS_result$change_points
##
   location value level
        150 6.405863
## 1
## 2
        101 6.106824
                          2
## 3
         47 5.345979
                          3
cpt_WBS_hat = sort(WBS_result$change_points[,1]) # estimated changepoints by WBS
cpt_WBS_hat
## [1] 47 101 150
Hausdorff.dist(cpt_WBS_hat, cpt_true)
## [1] 3
cpt_WBSlr_hat = local.refine.univar(cpt_WBS_hat, y, w = 1/3) # perform local refinement
cpt_WBSlr_hat
## [1] 47 101 150
Hausdorff.dist(cpt_WBSlr_hat, cpt_true)
## [1] 3
```