Test PMLSeg

Ninh

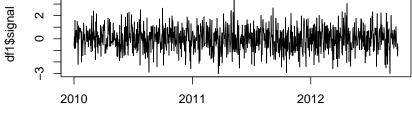
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Introduction

This document presents results from several tests of the PMLSeg package, including:

- Test of the Segmentation function with and without offsets, considering the following examples:
 - ► Ex1 : zero mean + IID noise
 - Ex2: periodic mean + IID noise
 - ► Ex3 : periodic mean + monthly variance
- ► Tests of other functions such as:
 - ▶ PlotSeg to visualize segmentation results
 - Cluster_screening to detect groups of close change-points (usually due to outliers) and determine whether to keep or remove the cluster. The cluster in this document is determined by a group of close change-points within 80 days (this threshold is changeable).
 - Validation to validate the detected change-points with the help of metadata.

Ex1 time series



df1\$date

```
head(df1, 3)

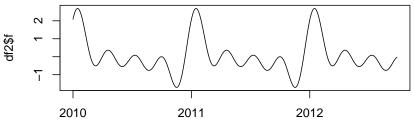
#> date signal

#> 1 2010-01-01 -0.6264538

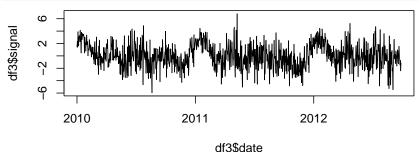
#> 2 2010-01-02 0.1836433

#> 3 2010-01-03 -0.8356286
```

Ex2 time series : Add the functional with 4 Fourier series components, each with a coefficient of $0.5\,$



 $\mathsf{Ex}3$ time series : Apply the same functional as $\mathsf{Ex}2,$ adjusted for monthly variance



Harmonize Formats of 3 Dataframes for Testing

```
df2 <- df2 %>% select(date, signal)
df3 <- df3 %>% select(date, signal)

names(df1)
#> [1] "date" "signal"
names(df2)
#> [1] "date" "signal"
names(df3)
#> [1] "date" "signal"
```

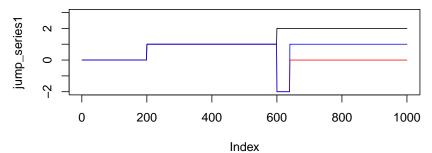
Preliminary Settings

Generate different offset series to add to the original series :

```
# Function to generate jump series
generate_jump_series <- function(jump_indices, jump_amp, length_series) {</pre>
  jump_series <- rep(0, length_series)</pre>
  jump indices <- c(1, jump indices, length series + 1)
  changes <- rep(0, length series)
  changes[jump indices[-length(jump indices)]] <- jump amp</pre>
  jump series <- cumsum(changes)</pre>
  return(jump_series)
# No cluster
jump ind1 \leftarrow c(200, 600)
jump amp1 \leftarrow c(0, 1, 1)
# One cluster formed by the second and third change-points, which need to be
# replaced by the middle point
jump_ind2 \leftarrow c(200, 600, 640)
jump_amp2 \leftarrow c(0, 1, -3, 2)
# One cluster formed by the second and third change-points, which need to be removed
iump amp3 \leftarrow c(0, 1, -3, 3)
```

Premilinary setting

```
# Generate jump series
jump_series1 <- generate_jump_series(jump_ind1, jump_amp1, length_series)
jump_series2 <- generate_jump_series(jump_ind2, jump_amp2, length_series)
jump_series3 <- generate_jump_series(jump_ind2, jump_amp3, length_series)
# Visualize jump series
plot(jump_series1, type = "l", ylim = c(-2,3))
lines(jump_series2, col = "red")
lines(jump_series3, col = "blue")</pre>
```



Premilinary setting

Testing the Segmentation Function

When the series is homogeneous

```
library(PMLseg)
# Ex.1
seg1a = Segmentation(OneSeries = df1, FunctPart = FALSE)
str(seg1a)
#> List of 5
#> $ Tmu : 'data.frame': 1 obs. of 5 variables:
#> ..$ begin: int 1
#> ..$ end : int 1000
#> ..$ mean : num -0.00423
#> ..$ se : num 0.0336
#> ..$ np : num 1000
#> $ FitF : logi FALSE
#> $ CoeffF : logi FALSE
#> $ MonthVar: num [1:12] 1.089 0.887 1.334 1.092 1.21 ...
#> $ SSR : num 933
seg1a$Tmu
#> begin end
                     mean.
#> 1 1 1000 -0.004229094 0.03363045 1000
```

No change-points are detected, as shown in the Tmu dataframe, which lists all segments of the series. The mean is close to 0. Additionally, the results include not only the Tmu dataframe but also other outputs such as the fitted functional part FitF (which is not fitted when FunctPart is set to FALSE), coefficients of functional elements CoeefF, monthly variance MonthVar, and the Sum of Squares of Residuals SSR.

Test the Segmentation Function

When series is homogeneous

```
# Ex2
seg2a = Segmentation(OneSeries = df2, FunctPart = TRUE)
seg2a$Tmu

#> begin end mean se np

#> 1 1 1000 -0.007048795 0.03366926 1000

# Ex3
seg3a = Segmentation(OneSeries = df3, FunctPart = TRUE)
seg3a$Tmu

#> begin end mean se np

#> 1 1 1000 -0.01249977 0.0488041 1000
```

No change-points are detected in these two examples either.

Test the Segmentation Function

Adding Jump Series to the original Series and running the segmentation

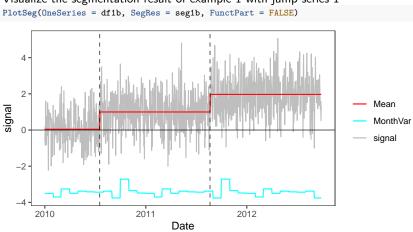
```
# Fx:1
df1b <- df1 %>% mutate(signal = signal + jump_series1)
seg1b = Segmentation(OneSeries = df1b, FunctPart = FALSE)
seg1b$Tmu
#> begin end mean
#> 1 199 0.04231954 0.07503647 199
#> 2 200 598 0.99732352 0.05361806 399
#> 3 599 1000 1.96897996 0.05271989 402
# Ex2
df2b <- df2 %>% mutate(signal = signal + jump series1)
seg2b = Segmentation(OneSeries = df2b, FunctPart = TRUE)
seg2b$Tmu
#> begin end mean se np
#> 1 1 200 0.02694284 0.07473907 200
#> 2 201 598 0.99309421 0.05374394 398
#> 3 599 1000 1.97809762 0.05280951 402
# Ex.3
df3b <- df3 %>% mutate(signal = signal + jump_series1)
seg3b = Segmentation(OneSeries = df3b, FunctPart = TRUE)
seg3b$Tmu
#> begin end mean
#> 2 234 601 1.0163802 0.08082727 368
#> 3 602 1000 1.9701980 0.07739899 399
```

Test the Segmentation Function

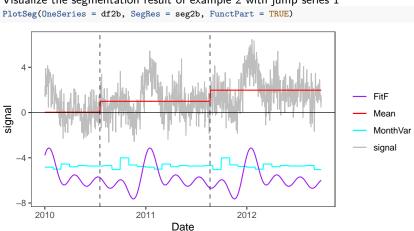
Adding other jump series in the most complicated series (Ex3)

```
df3c <- df3 %>% mutate(signal = signal + jump_series2)
seg3c = Segmentation(OneSeries = df3c, FunctPart = TRUE)
seg3c$Tmu
    beain end
                               se np
                    mean
#> 2 201 599 0.99060960 0.07910378 399
#> 3 600 641 -1.92086925 0.24382647 42
#> 4 642 1000 -0.04575017 0.08162489 359
df3d <- df3 %>% mutate(signal = signal + jump series3)
seg3d = Segmentation(OneSeries = df3d, FunctPart = TRUE)
seg3d$Tmu
#> begin end mean
                               se np
#> 1 1 200 0.04550598 0.10420812 200
#> 2 201 599 0.98930870 0.07913867 399
#> 3 600 639 -1.92932956 0.24728105 40
#> 4 640 1000 0.94993983 0.08153644 361
```

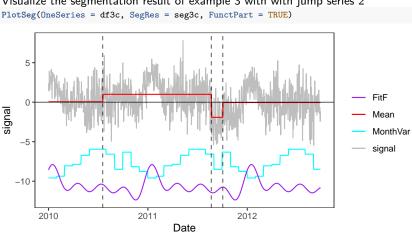
Visualize the segmentation result of example 1 with jump series 1



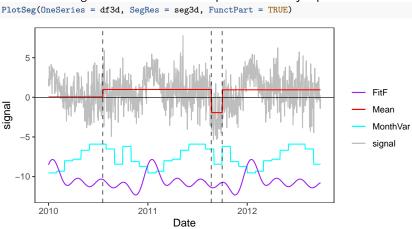
Visualize the segmentation result of example 2 with jump series 1



Visualize the segmentation result of example 3 with with jump series 2



Visualize the segmentation result of example 3 with with jump series 3



Test the Cluster_screening Function

Segmentation sometime detects the cluster like in example 3c, 3d. When it is too close and after cluster, the change in mean is not significant, we will remove such kind of cluster (screening2). If it is significant, we will replace a cluster by the middle point (screening1) and recommend to remove the data within the cluster.

```
screening1 = Cluster screening(Tmu = seg3c$Tmu,
                             MaxDist = 80
str(screening1)
#> List of 3
#> $ UpdatedCP : num [1:2] 200 621
#> $ RemoveData: 'data.frame': 1 obs. of 2 variables:
#> ..$ begin: int 600
#> ..$ end : int 641
#> $ ChangeCP : chr "Yes"
screening2 = Cluster_screening(Tmu = seg3d$Tmu,
                             MaxDist = 80
str(screening2)
#> List of 3
#> $ UpdatedCP : int 200
#> $ RemoveData: 'data.frame': 1 obs. of 2 variables:
#> ..$ beain: int 600
#> ..$ end : int 639
#> $ ChangeCP : chr "Yes"
```

When the screening say it is needed to replace the changepoint, we need to reestimate the mean and other parameters in the Tmu dataframe.

```
seg upd3c = UpdatedParametersForFixedCP(OneSeries = df3c,
                                  ResScreening = screening1)
str(seg upd3c)
#> List of 4
#> $ MonthVar: num [1:12] 1.06 1.28 2.6 2.79 3.92 ...
#> $ Tmu : 'data.frame': 3 obs. of 5 variables:
#> ..$ begin: num [1:3] 1 201 622
#> ..$ end : num [1:3] 200 621 1000
#> ..$ mean : num [1:3] 0.049 0.9937 -0.0451
#> ..$ se : num [1:3] 0.1042 0.0782 0.0807
#> ..$ np : num [1:3] 200 421 379
#> $ FitF : Named num [1:1000] 2.14 2.22 2.3 2.37 2.44 ...
#> ..- attr(*, "names")= chr [1:1000] "1" "2" "3" "4" ...
#> $ CoeffF : Named num [1:11] 0.492 0.511 0.54 0.546 0.446 ...
#> ..- attr(*, "names")= chr [1:11] "cos1" "sin1" "cos2" "sin2" ...
seg upd3c$Tmu
#>
       begin end mean se np
#> mean 2 201 621 0.99366883 0.07818545 421
#> mean 3 622 1000 -0.04511261 0.08072260 379
```

```
seg upd3d = UpdatedParametersForFixedCP(OneSeries = df3d,
                                 ResScreening = screening2)
str(seg upd3d)
#> List of 4
#> $ MonthVar: num [1:12] 1.06 1.28 2.6 2.79 3.92 ...
#> $ Tmu : 'data.frame': 2 obs. of 5 variables:
#> ..$ begin: num [1:2] 1 201
#> ..$ end : int [1:2] 200 1000
#> ..$ mean : num [1:2] 0.0478 0.9727
#> ..$ se : num [1:2] 0.1042 0.0562
#> ..$ np : num [1:2] 200 800
#> $ FitF : Named num [1:1000] 2.13 2.22 2.3 2.37 2.44 ...
#> ..- attr(*, "names")= chr [1:1000] "1" "2" "3" "4" ...
#> $ CoeffF : Named num [1:10] 0.491 0.514 0.545 0.548 0.448 ...
#> ..- attr(*, "names")= chr [1:10] "cos1" "sin1" "cos2" "sin2" ...
seg_upd3d$Tmu
#> begin end mean se np
#> mean 2 201 1000 0.9727184 0.05615832 800
```

Visualize the time series after the updated segmentation

Date

Compare to the result before the cluster screening

PlotSeg(OneSeries = df3d,
SegRes = seg3d)

FitF

Mean

MonthVar

signal

Visualize the time series after the updated segmentation

Date

Compare to the result before the cluster screening

```
PlotSeg(OneSeries = df3c,
SegRes = seg3c)

FitF

Mean

MonthVar

signal
```

Test the Validation function

Validate the detected change-points with respect to the metadata

Visualize results with metadata

