#### PyCon ID 2019

## Introduction to Change Point Analysis

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#### Outline

1

#### **Introduction**

What is change point analysis?

2

#### Methodology

How do we find the "change"?

3

#### **Conclusions**

What have we learned?

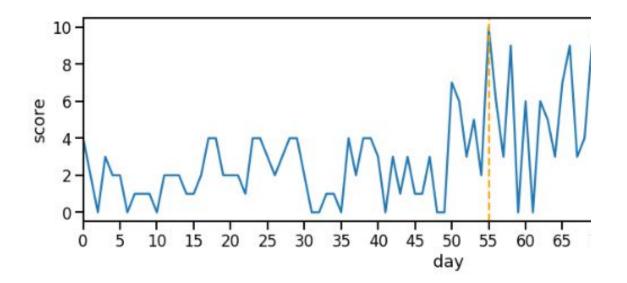
#### 1. Introduction

What is change point analysis?

## Whoa!

At the end of your peaceful Friday, a product manager came and asked a question...



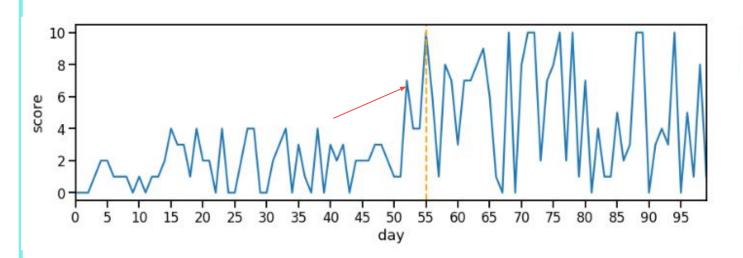




Our reviews are getting better!

It's because of our new feature release, isn't it?

#### Is it *really* getting better?





## What is change point analysis?

Given a series of data, change point analysis involves **detecting the number and** location of change points, **locations in the data where some feature**, for example the mean, **changes**.

#### There are two types of change point analysis ...

#### Offline

- all data are processed in one go
- main goal: accurate detection of changes

#### Online

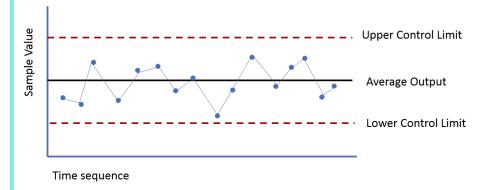
- data must be processed quickly "on the fly" before new data arrives
- main goal: the **quickest**detection of a change after it
  has occured

#### 2. Methodology

How do we find the "change"?

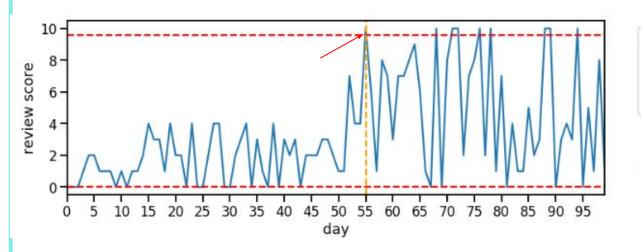
#### A. Control charts

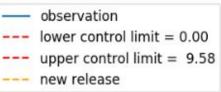
- Common in process control
- Use average, lower & upper control limit
- Focus on **point-wise** error rate
- Lower & upper limit is determined based on standard deviation



Example of control chart (source)

#### Sample control chart





upper control limit =  $\mu + 2\sigma$ lower control limit =  $\mu - 2\sigma$ 

The first observation which lies above upper control limit: day 55.

#### B. Change point analysis

- Can detect subtle changes
   frequently missed by control charts
- Can be conducted once all observations are collected, to identify change-wise error rate
- Based on **mean** or **variance**



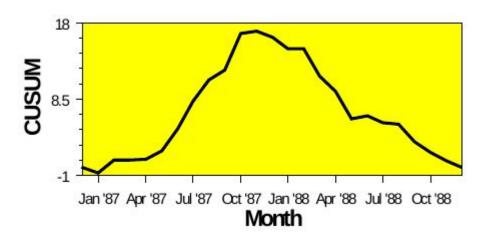
## Single change point analysis

Method 1: Cumulative Sum (CUSUM)

#### Cumulative Sum (CUSUM)

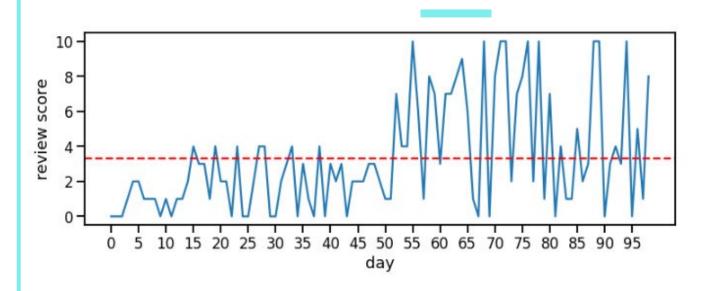
#### Steps:

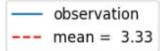
- 1. Calculate **mean** value of **all observations** ( $\bar{y}$ )
- 2. Calculate **residuals**: difference between y<sub>i</sub> and  $\bar{y}$
- 3. Set cumulative sum of residuals at 0: S<sub>0</sub> = 0
   4. Calculate cumulative sum of
- 4. Calculate **cumulative sum of** residuals:  $S_i = S_{i-1} + \varepsilon_i$



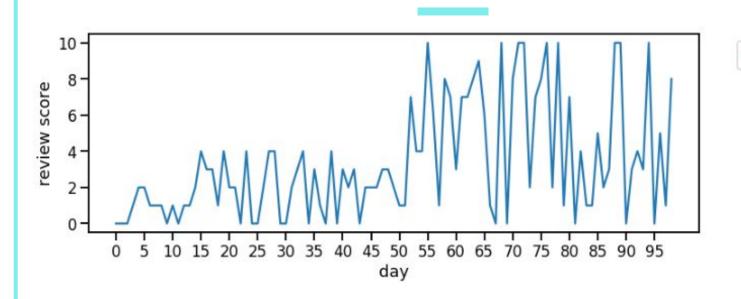
Example of CUSUM plot (source)

#### **CUSUM:** calculate mean





#### **CUSUM:** calculate residuals



residual

#### CUSUM: calculate cumulative sum of residuals



#### CUSUM: how confident are we that the change exists?

- Frequentist method
- **Sampling without replacement**: randomly reorder the observations
- determine number of iteration N
- 2. for each iteration:
  - take random sample without replacement from the observations  $X_1^0, X_2^0, \dots, X_n^0$ , where n = number of observations
  - calculate the sample's cumulative sum of residuals ( $S^0$ ):  $S_1^0, S_2^0, \dots, S_n^{\bar{0}}$
  - · calculate the difference between maximum and minimum residuals in each bootstrap

$$S_{\text{diff}}^0 = S_{max}^0 - S_{min}^0$$

• If  $S_{\rm diff}^0 < S_{\rm diff}$ , it means the result from the sample is consistent with actual observations

confidence level(%) = 
$$100 * \frac{X}{N}$$

where X represents number of iteration which has  $S_{
m diff}^0 < S_{
m diff}$ 

#### CUSUM: how confident are we that the change exists?

```
def calculate_residual_difference(df):
    ## calculate difference between maximum and minimum cumsum residuals
    resid_max = df['residual_cumsum'].max()
    resid_min = df['residual_cumsum'].min()

resid_diff = resid_max - resid_min
    return resid_diff
```

#### CUSUM: how confident are we that the change exists?

```
N = 1000 ## determine number of iteration
X = 0 ## occurrence when sample residual difference < observed residual difference
for i in np.arange(0,N):
    sample = pd.DataFrame(
        np.random.choice(df[0], size = df.shape[0], replace = False)
    sample = calculate cusum residuals( sample)
    sample resid diff = calculate residual difference( sample)
    if sample resid diff < resid diff:</pre>
        X += 1
confidence level = 100 * X / N
print("Confidence level: {:.2f}%".format(confidence level))
```

## Single change point analysis

Method 2: Structure change model - MSE Estimator

#### Structure Change Model: MSE Estimator

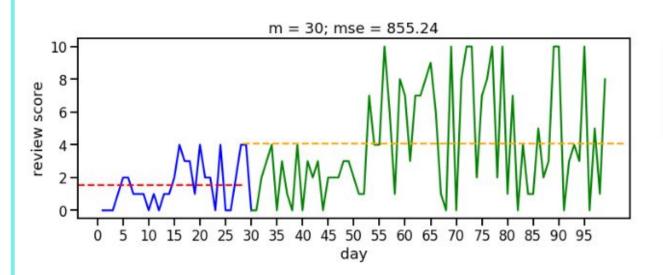
#### Steps:

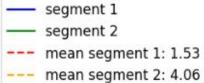
- 1. Split the data into 2 segments
  - segment  $1 = \{1, ..., m\}$
  - segment  $2 = \{m+1, ..., n\}$
- 2. Calculate **average** value of each segment:  $X_1$  and  $X_2$
- 3. Calculate **mean squared error** of observation in **each segment**

$$MSE(m) = \sum_{i=1}^{m} (X_i - \overline{X}_1)^2 + \sum_{i=m+1}^{n} (X_i - \overline{X}_2)^2$$
where  $\overline{X}_1 = \frac{\sum_{i=1}^{m} X_i}{m}$  and  $\overline{X}_2 = \frac{\sum_{i=m+1}^{n} X_i}{n-m}$ 

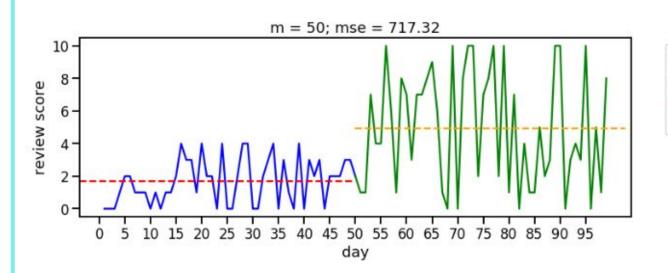
4. Value of **m** which **minimizes the MSE** is the best estimator of the **last point before the change occured** 

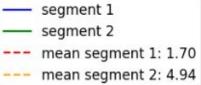
#### MSE estimator: intuition



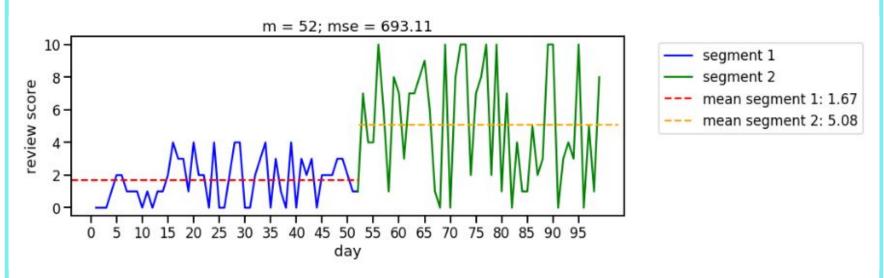


#### MSE estimator: intuition





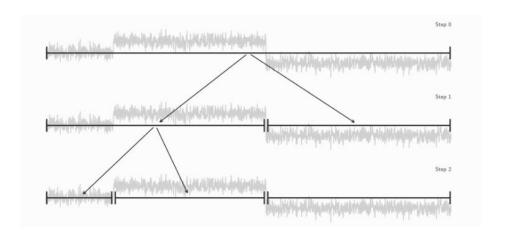
#### MSE estimator: intuition



Value of **m** which **minimizes the MSE** is the best estimator of the **last** point before the change occured → **day 52** 



#### Multiple Change Point: Binary Segmentation



Schematic view of the binary segmentation algorithm (source)

#### Libraries



ruptures bayesloop fbProphet



changepoint bcp strucchange cpm

# Confused? You can apply *Bayesian* approach too!

—Anonymous

#### **Bayesian Approach**

- 1. Set **prior** distribution of  $\mu_1$ ,  $\mu_2$ , and overall  $\sigma$
- 2. The changepoint could occur in  $\tau \in \{1,...,n\}$
- 3. Assign:

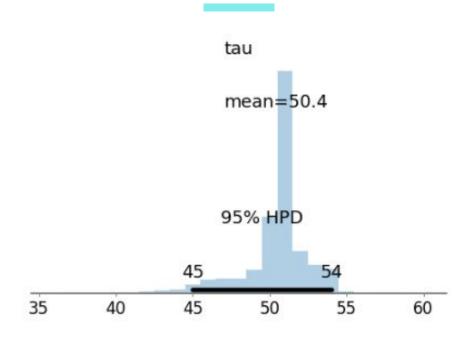
$$\mu = \begin{cases} \mu_1 & \text{if } \tau \ge t \\ \mu_2 & \text{if } \tau < t \end{cases}$$

4. Produce the sample!

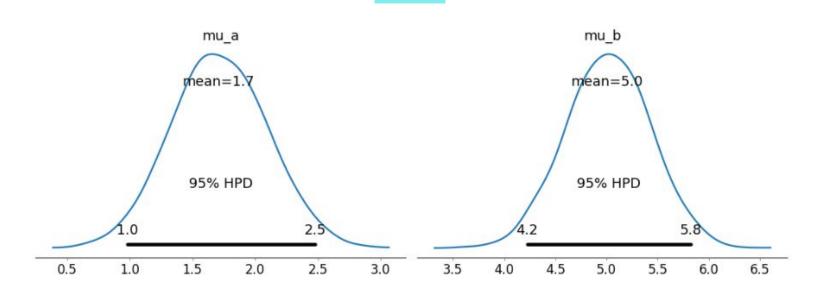
#### Bayesian Approach: PyMC3 - Example

```
import pymc3 as pm
## set number of sample
## set t = time, from 0 to length of observations
samples = 5000 ## number of iteration
t = np.arange(0, len(z)) ## array of observation positions (time)
with pm.Model() as model:
   ## define uniform priors for the mean values
    mu \ a = pm.Uniform('mu \ a', 0, 10)
    mu b = pm.Uniform('mu b', 0, 10)
    sigma = pm.HalfCauchy('sigma', np.std(z))
    tau = pm.DiscreteUniform('tau', t.min(), t.max())
   ## define stochastic variable mu
    mu = pm.math.switch(tau >= t, mu a, mu b)
    observation = pm.Normal('observation', mu, sigma, observed = z)
    trace = pm.sample(samples, step = pm.NUTS())
    burned trace = trace[1000:]
```

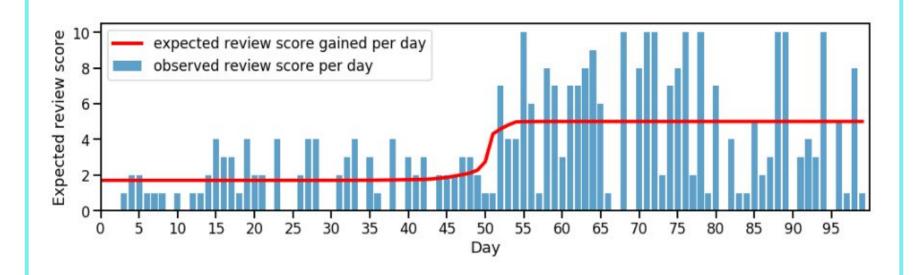
#### Bayesian Approach: PyMC3 - Changepoint distribution



#### Bayesian Approach: PyMC3 - Mean distribution



#### Bayesian Approach: Estimate the change point



#### 3. Conclusions

What have we learned?

### Thanks!



Materials: <a href="https://github.com/elvyna/pycon-id-2019">https://github.com/elvyna/pycon-id-2019</a>

Find me on Twitter: @vexenta

#### Want to learn more?

Killick, R. (2017). Introduction to optimal changepoint detection algorithms. useR! Tutorial 2017

Kass-Hout, T. (2010). Change point analysis. Slideshare.

Bellei, C. (2016). Changepoint Detection. Part I - A Frequentist Approach. [Blog]

Bellei, C. (2017). Changepoint Detection. Part II - A Bayesian Approach. [Blog]

<u>Davidson-Pilon, C. (2015). Chapter 1 - Introduction - PyMC3. Probabilistic Programming and Bayesian Methods for Hackers.</u>

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