## **NUS-ISS**Intelligent Sensing and Sense Making





#### Module 3 - Workshop on sensor signal processing

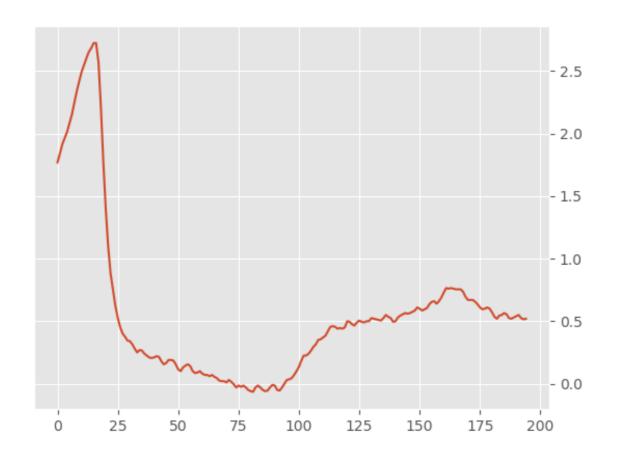
by Nicholas Ho

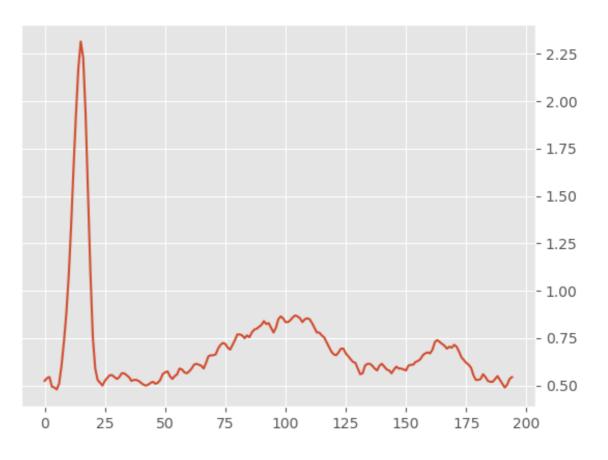
© 2020 National University of Singapore. All Rights Reserved.

#### **Problem**

•How similar are these two signals?

•In which manners are they similar?

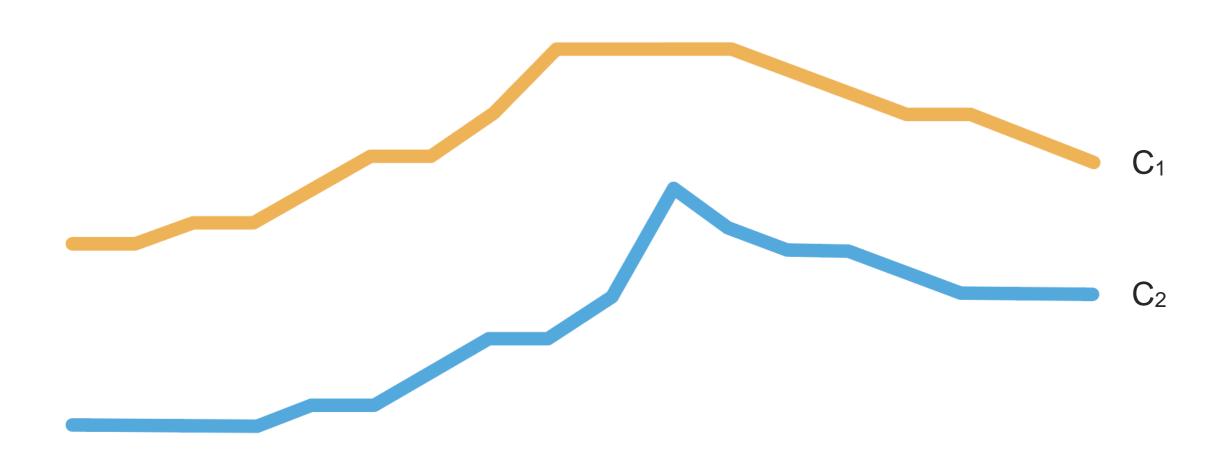




#### **Similarity**

How similar are these two signals

•In what ways are these two signals similar to each other?

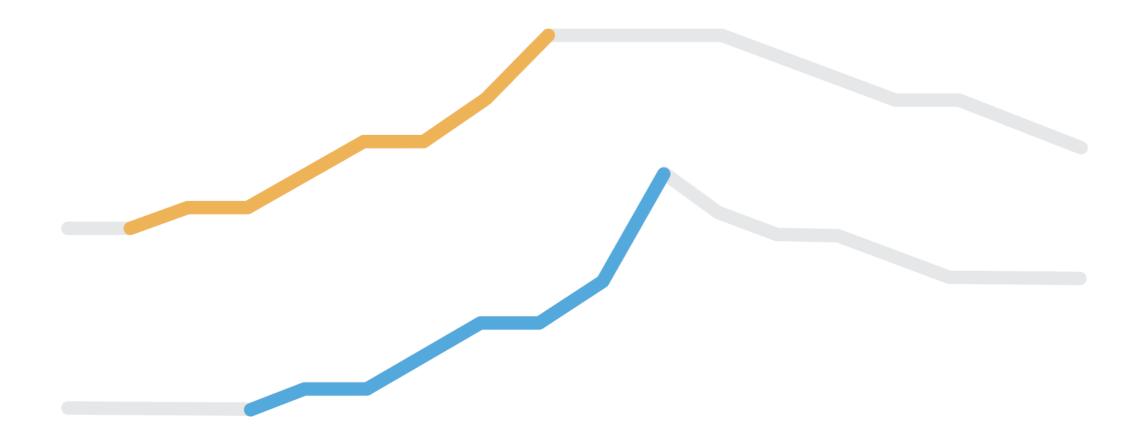


#### **Similarity**

How similar are these two signals

....?

•In what ways are these two signals similar to each other?



#### **Similarity**

How similar are these two signals

•In what ways are these two signals similar to each other?

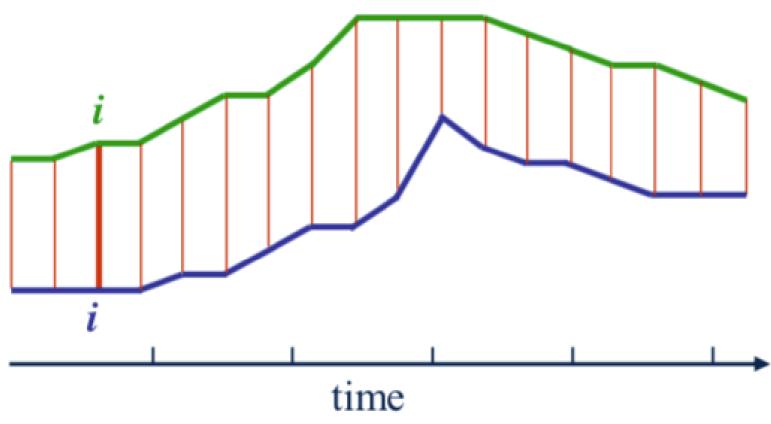


#### Can we try ...

Euclidean, Manhattan .....?

•We can measure the similarity of two signals by calculating the distance between the *i*-th point on one signal and the *i*-th point on another signal

•Simple concept, but could not capture the similarity in shape

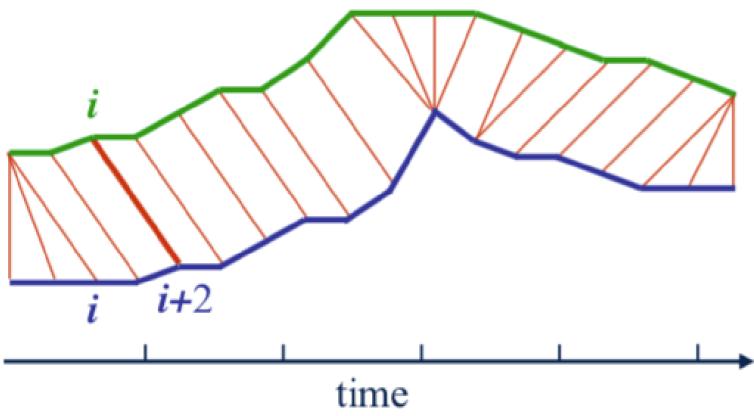


Source: "Dynamic Time Warping Algorithm", by Elena Tsiporkova

#### How about ...

non-linear alignment .....?

- Elastic alignment between points of two signals produces a better, more intuitive similarity measure
- Allow similar shapes to match even if they are out of phase



Source: "Dynamic Time Warping Algorithm", by Elena Tsiporkova

#### **Distance**

Another term to say 'similiarity'

Consider two distinct signals

$$\mathbf{x} = [x_1, x_2, ..., x_i, ... x_m]$$
  
 $\mathbf{y} = [y_1, y_2, ..., y_i, ... y_n]$ 

 The distance between the two signals is defined as

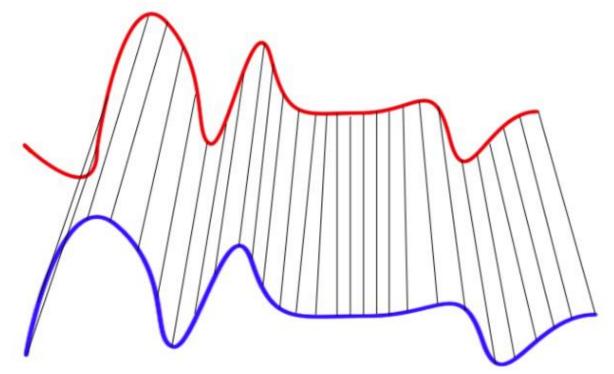
$$d_{s}(\mathbf{x},\mathbf{y})$$

Euclidean distance between two signals:

$$d_s(\mathbf{x}, \mathbf{y}) = \sqrt{\sum_i (x_i - y_i)^2}$$

Problem: two signals must be of same length!

- An algorithm to measure similarity between two temporal sequences (signal), which may vary in speed
- DTW calculates an optimal match between two given sequences
- Sequences are warped along time dimension to determine similarity independent of variations in time
- •DTW produces warping path, which enables alignment between two signals

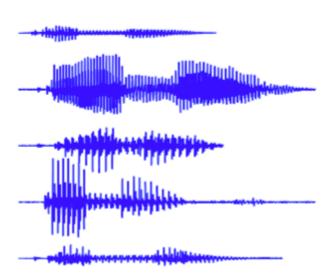


Source: https://th.wikipedia.org/wiki/Dynamic\_time\_warping#/media/Fil e:Euclidean\_vs\_DTW.jpg

Usage

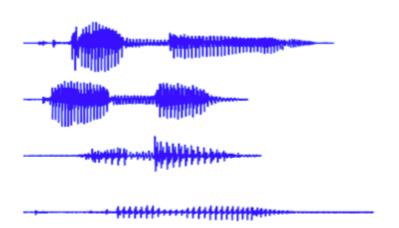
Commonly used in speech recognition





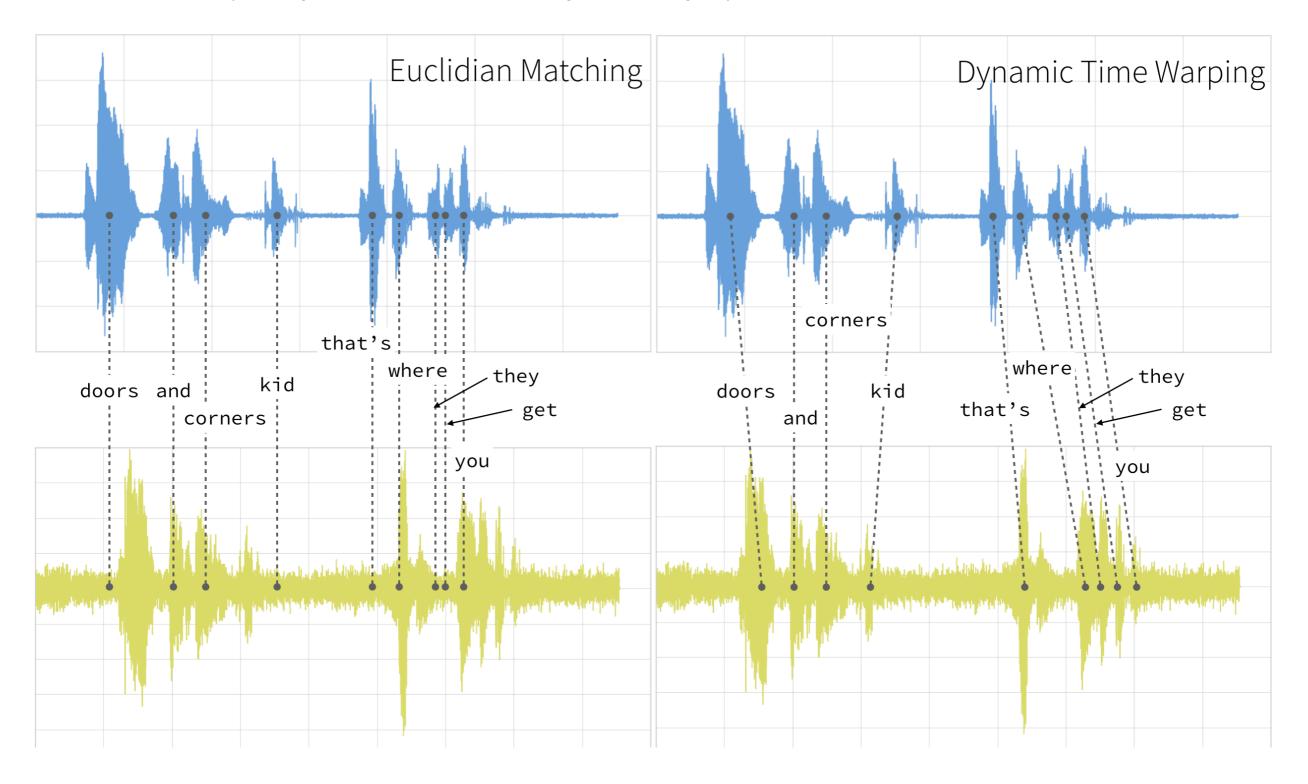
 Individual never pronounces one word twice in exact way

 People never pronounce one word with the same timing

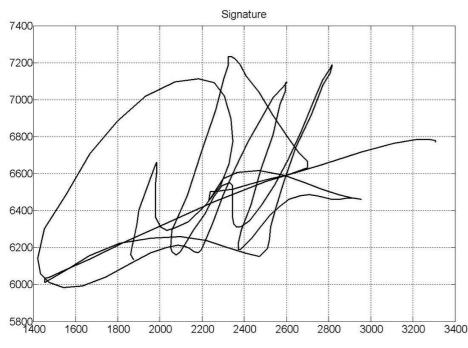


Source: "Speech Recognition - Intro and DTW", by Jan Černocky

Vs Euclidean (Comparison of Audio Clips Example)



Dynamic signature recognition

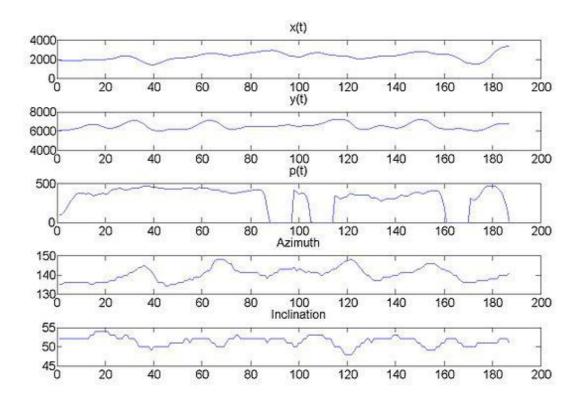


Source: "Speech Recognition - Intro and DTW", by Jan Černocky

### Users sign their signature on digital tablet

- Dynamic information captured:
- x position
- y position
- pressure
- azimuth
- inclination

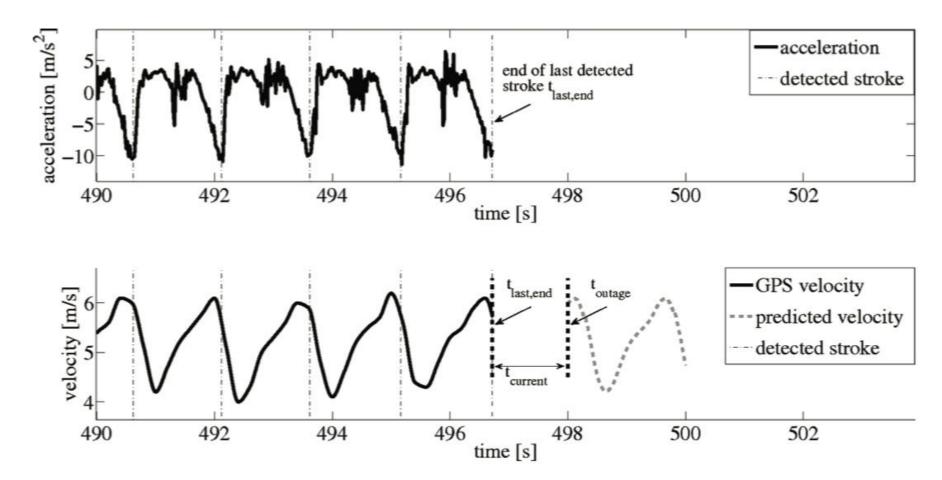
#### Use DTW to check / match signature





Stroke detection (rowing)

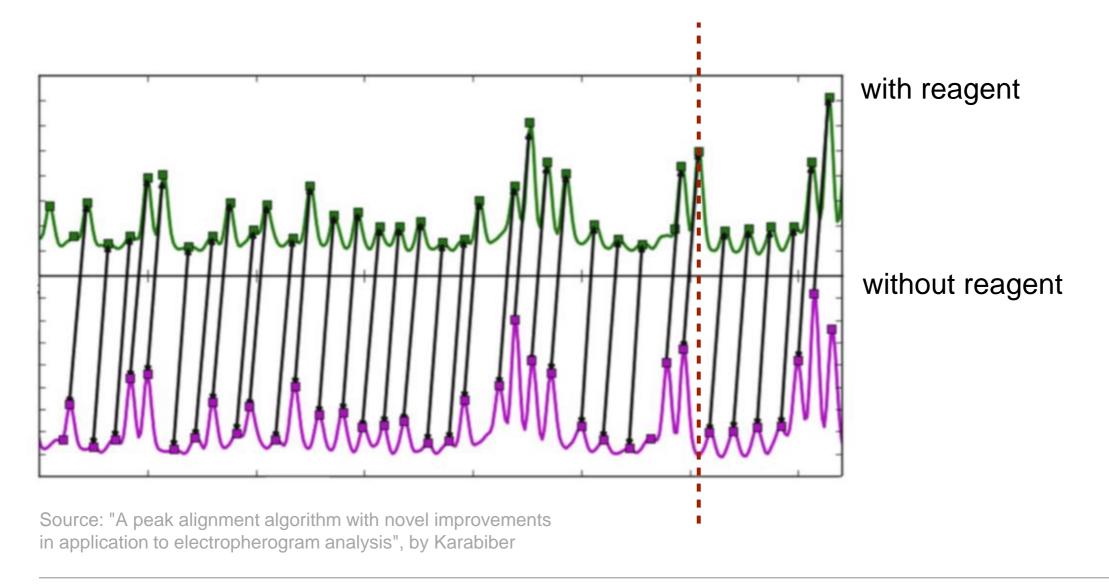
- Use DTW to detect stroke (used in rowing competitions)
- With strokes detected, predict boat's movement and position when sensor transmission lost



Source: "Movement prediction in rowing using a dynamic time warping base stroke detection", by Groh et al.

Peak alignment in DNA sequencing

- Use DTW to align peaks in electropherogram (a plot generated by DNA sequencer)
- Accurate alignment gives better interpretation (e.g. better RNA secondary structure prediction)



Overview of algorithm

### n x m matrix covers the window boundaries

### orange curve = correct wrapping path

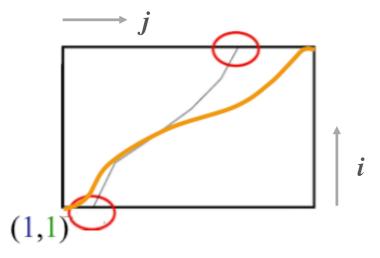
Start by constructing n x m matrix
 D, in which

$$D_{i,j} = d_s(y_i, x_j)$$

where

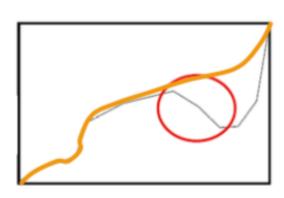
$$d_s(y_i, x_j) = (y_i - x_j)^2$$

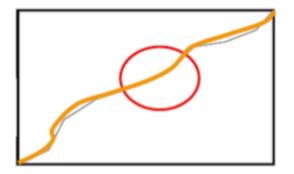
- •Create a warping path w that maps points between x and y, the path w must satisfy the following rules:
  - 1. Boundary conditions
  - 2. Monotonicity
  - 3. Continuity



16 of 39





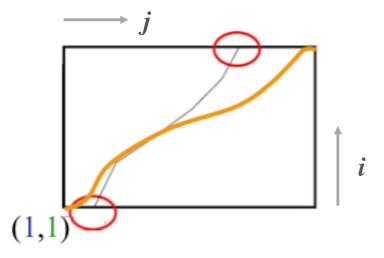


Overview of algorithm

1st Rule: start point from bottom left and end point at top right

**2nd Rule:** Cannot move back in direction

3rd Rule: cannot have a break in between! Must be continuous



17 of 39

Source: "Dynamic time warping algorithm", by Elena Tsiporkova

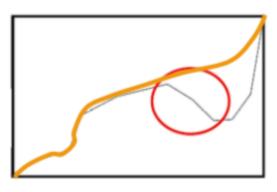
•Start by constructing  $n \times m$  matrix D, in which

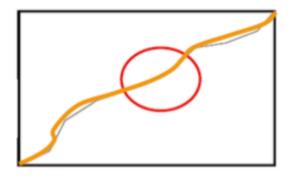
$$D_{i,j} = d_s(y_i, x_j)$$

where

$$d_s(y_i, x_j) = (y_i - x_j)^2$$

- •Create a warping path w that maps points between x and y, the path w must satisfy the following rules:
  - 1. Boundary conditions
  - 2. Monotonicity
  - 3. Continuity





Overview of algorithm

## Codes to import libraries and to perform some setup

- DTW algorithm consists of mainly 3 parts:
  - 1. Compute distance matrix
  - 2. Compute accumulated cost matrix
  - 3. Search the optimal path

> import numpy as np

 To start the code, import the necessary libraries, and setup a bit

```
> import matplotlib.pyplot as plt
> import pandas as pd

> plt.style.use('ggplot')
> plt.rcParams['ytick.right'] = True
> plt.rcParams['ytick.labelright'] = True
> plt.rcParams['ytick.labelright'] = False
```

> plt.rcParams['ytick.labelleft'] = False

#### 1. Compute distance matrix

#### Define two simple short signals:

```
> x = np.array([0.5,0.5,2.0,3.0,2.0,0.0])
> y = np.array([0.0,1.5,1.5,1.5,3.0,2.5,1.0])
```

#### Plot the two signals

1. Compute distance matrix

## applying maths formula to calculate the distance matrix in DTW

 Compute the distance matrix is straightforward, since the matrix is defined as

$$D_{i,j} = d_s(y_i, x_j)$$

and

$$d_s(y_i, x_j) = (y_i - x_j)^2$$

The corresponding code

```
dists
```

```
[[0.25, 0.25, 4. , 9. , 4. , 0. ], [1. , 1. , 0.25, 2.25, 0.25, 2.25], [1. , 1. , 0.25, 2.25, 0.25, 2.25], [1. , 1. , 0.25, 2.25, 0.25, 2.25], [6.25, 6.25, 1. , 0. , 1. , 9. ], [4. , 4. , 0.25, 0.25, 0.25, 6.25], [0.25, 0.25, 0.25, 1. , 4. , 1. , 1. ]]
```

```
> dists = np.zeros((len(y),len(x)))
```

```
> for i in range(len(y)):
    for j in range(len(x)):
        dists[i,j] = (y[i]-x[j])**2
```

1. Compute distance matrix

### **Note:** Output from these codes is inverted

#### i.e. top left cell = D[0,0]; bottom left should be D[0,0] instead

#### dists

issm/m1.3/v1.0

 Compute the distance matrix is straightforward, since the matrix is defined as

$$D_{i,j} = d_s(y_i, x_j)$$

and

$$d_s(y_i, x_j) = (y_i - x_j)^2$$

The corresponding code

```
> dists = np.zeros((len(y),len(x)))
```

```
> for i in range(len(y)):
    for j in range(len(x)):
        dists[i,j] = (y[i]-x[j])**2
```

1. Compute distance matrix

 Create a function to do a plot on the distance matrix

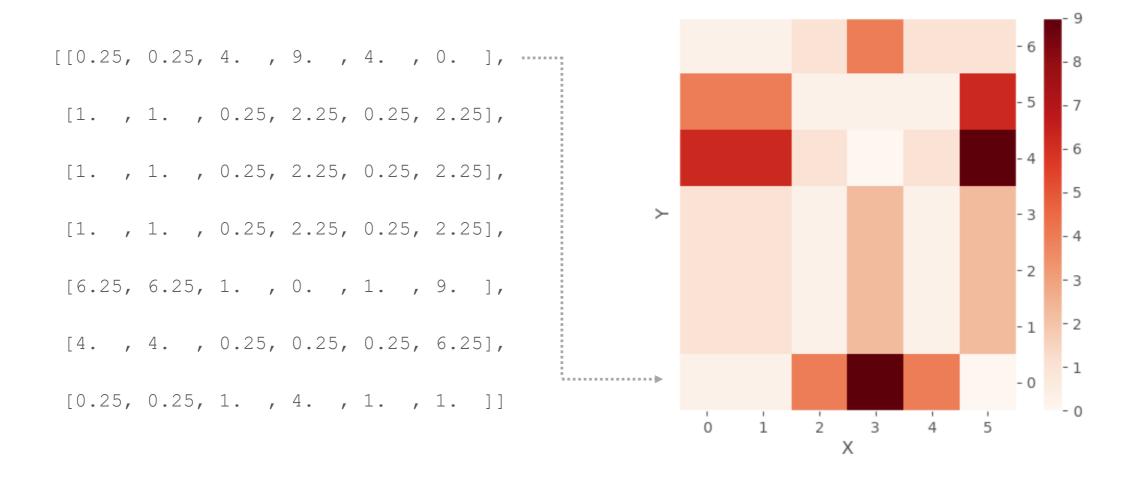
**nverted because** 

```
viridis starts
> def pltDistances(dists,xlab="X",ylab="Y",clrmap="viridis"):
                                                                    from top left
      imgplt = plt.figure()
      plt.imshow(dists,
                 interpolation='nearest',
                 cmap=clrmap)
      plt.gca().invert_yaxis()
      plt.xlabel(xlab)
      plt.ylabel(ylab)
      plt.grid()
      plt.colorbar()
                                                                      · 2
      return imaplt
                                                                        - 2
                                                          3
                                              0
                                                  1
                                                      2
```

> pltDistances(dists,clrmap='Reds')

#### 1. Compute distance matrix

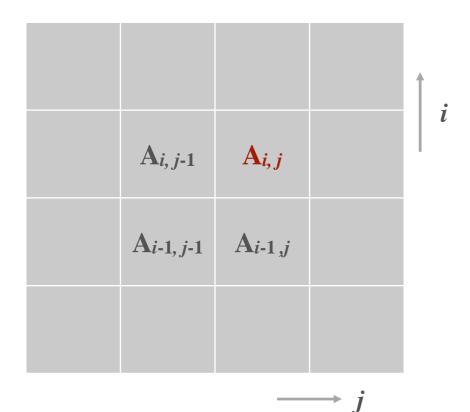
- Do take note, in the plot, the y axis is inverted
- Thus, first row of the matrix corresponds to the last row in the figure



2. Compute accumulated cost matrix

## Cost matrix important to determine wrapping path!

 $A_{i,j}$  equals to  $D_{i,j}$  plus either  $A_{i-1,j-1}$ ,  $A_{i,j-1}$  or  $A_{i-1,j}$ , whichever has the lowest value



The accumulated cost matrix is defined

$$A_{i,j} = D_{i,j} + min(A_{i-1,j}, A_{i,j-1}, A_{i-1,j-1})$$

•When *i* and *j* equals to 0

$$A_{0,0} = D_{0,0}$$

•When *i* equals to 0 (first row)

$$A_{0,j} = D_{0,j} + A_{0,j-1}$$

•When *j* equals to 0 (first column)

$$A_{i,0} = D_{i,0} + A_{i-1,0}$$

2. Compute accumulated cost matrix

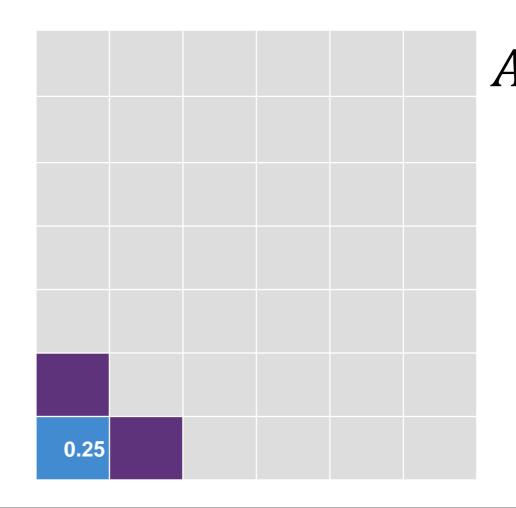
•When *i* equals to 0 (first row)

$$A_{0,j} = D_{0,j} + A_{0,j-1}$$

•When *j* equals to 0 (first column)

$$A_{i,0} = D_{i,0} + A_{i-1,0}$$

0.25	0.25	1	4	1	1	D
4	4	0.25	0.25	0.25	6.25	
6.25	6.25	1	0	1	9	
1	1	0.25	2.25	0.25	2.25	
1	1	0.25	2.25	0.25	2.25	
1	1	0.25	2.25	0.25	2.25	<b>†</b>
0.25	0.25	4	9	4	0	i



2. Compute accumulated cost matrix

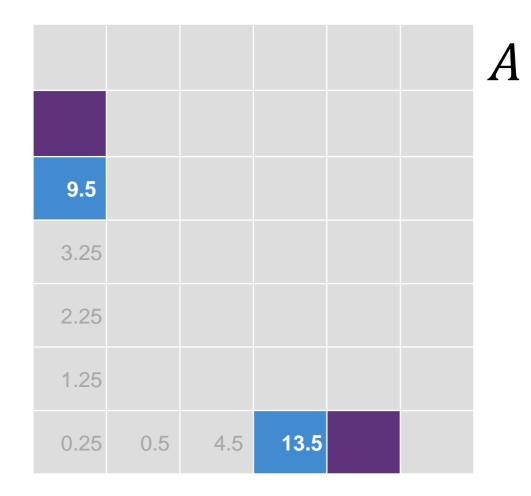
•When *i* equals to 0 (first row)

$$A_{0,j} = D_{0,j} + A_{0,j-1}$$

•When *j* equals to 0 (first column)

$$A_{i,0} = D_{i,0} + A_{i-1,0}$$

	$\boldsymbol{j}$					
0.25	0.25	1	4	1	1	D
4	4	0.25	0.25	0.25	6.25	
6.25	6.25	1	0	1	9	
1	1	0.25	2.25	0.25	2.25	
1	1	0.25	2.25	0.25	2.25	
1	1	0.25	2.25	0.25	2.25	<b>†</b>
0.25	0.25	4	9	4	0	i



26 of 39

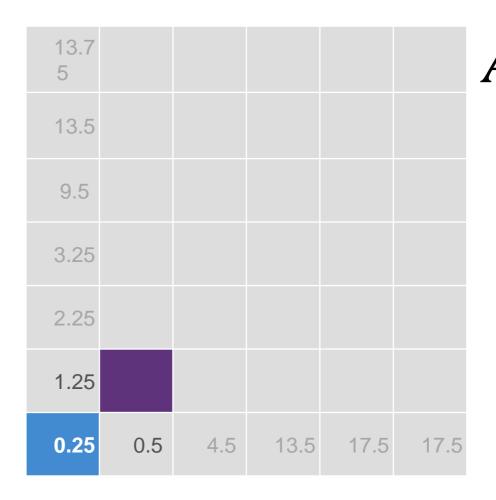
2. Compute accumulated cost matrix

Else

$$A_{i,j} = D_{i,j} + min(A_{i-1,j}, A_{i,j-1}, A_{i-1,j-1})$$

 $\longrightarrow j$ 

0.25	0.25	1	4	1	1	D
4	4	0.25	0.25	0.25	6.25	
6.25	6.25	1	0	1	9	
1	1	0.25	2.25	0.25	2.25	
1	1	0.25	2.25	0.25	2.25	
1	1	0.25	2.25	0.25	2.25	<b>†</b>
0.25	0.25	4	9	4	0	i



2. Compute accumulated cost matrix

Else

$$A_{i,j} = D_{i,j} + min(A_{i-1,j}, A_{i,j-1}, A_{i-1,j-1})$$

----- J

0.25	0.25	1	4	1	1	D
4	4	0.25	0.25	0.25	6.25	
6.25	6.25	1	0	1	9	
1	1	0.25	2.25	0.25	2.25	
1	1	0.25	2.25	0.25	2.25	
1	1	0.25	2.25	0.25	2.25	<b>†</b>
0.25	0.25	4	9	4	0	i

13.7 5 13.5 9.5 2.25 9.5 2.25 1.25 3.25 3.25 1.25 3.25 3.25 5.5 2.25 2.25 3 3.25 5.5 1.25 1.25 0.75 3 3.25 5.5 0.25 0.5 4.5 13.5 17.5 17.5

A

2. Compute accumulated cost matrix

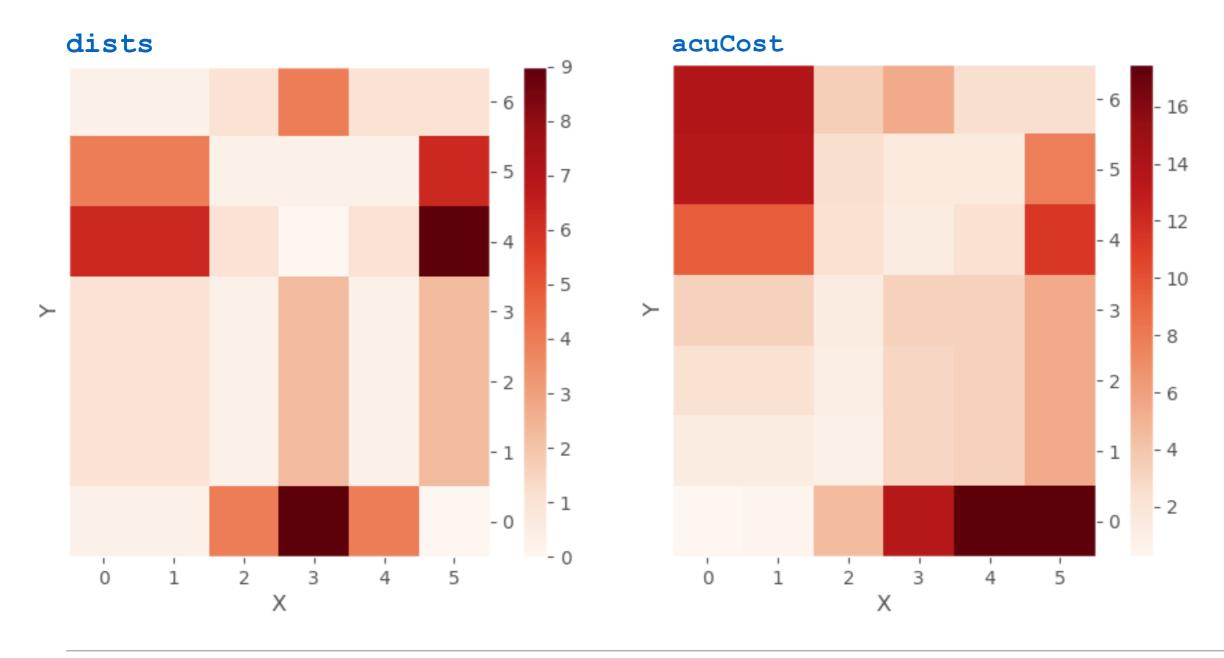
•Name the accumulated cost matrix as acuCost. It has the same shape as dists.

#### Codes to determine the accumulated cost matrix

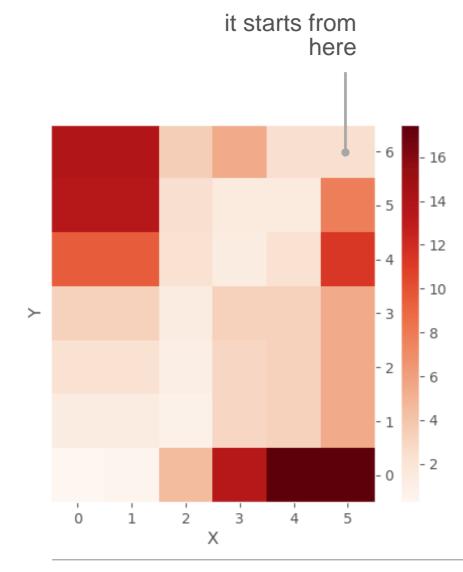
```
= np.zeros(dists.shape)
                        > acuCost
                                                                       starts from 1 because
                        > acuCost[0,0] = dists[0,0]
                                                                    acuCost[0,0] has been filled
 A_{0.0} = D_{0.0}
                                                                           with dists[0,0]
 A_{0,j} = D_{0,j} + A_{0,j-1}
                        > for j in range(1, dists.shape[1]):
                               acuCost[0,j] = dists[0,j] + acuCost[0,j-1]
first row
 A_{i,0} = D_{i,0} + A_{i-1,0}
                        > for i in range(1, dists.shape[0]):
first column
                               acuCost[i,0] = dists[i,0]+acuCost[i-1,0]
                        > for i in range(1, dists.shape[0]):
                               for j in range (1, dists.shape[1]):
                                    acuCost[i,j] = min(acuCost[i-1,j-1],
                                                              acuCost[i-1,j],
 A_{i,j} = D_{i,j} + min(A_{i-1,j}, A_{i,j-1}, A_{i-1,j-1})
                                                              acuCost[i,j-1])+dists[i,j]
                        > pltDistances(acuCost,clrmap='Reds')
```

2. Compute accumulated cost matrix

•Can you see the warping / optimal path?



3. Search the optimal path



Name the warping path as path.

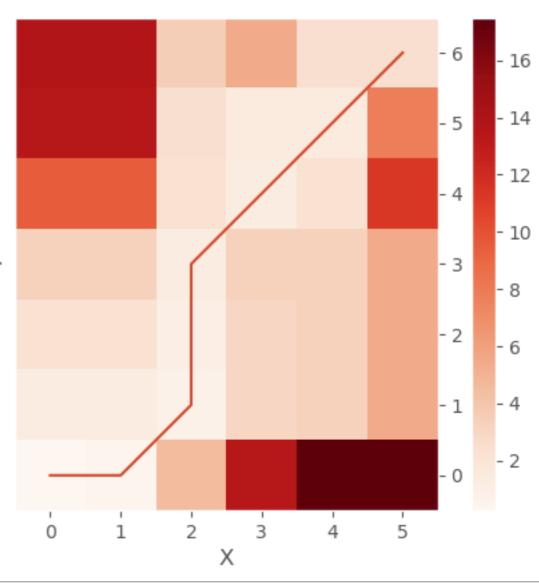
```
a list; j stands for x axis,
> i
          = len(y) - 1
                               i stands for y axis
> j
          = len(x)-1
> path = [[j,i]]
> while (i > 0) and (j > 0):
      if i==0:
                            Wrapping process: Compare 3
          j = j-1
                            neighboring squares and find
      elif j==0:
                              the minimum cost square
          i = i-1
      else:
          if acuCost[i-1,j] == min(acuCost[i-1,j-1],
                                    acuCost[i-1,j],
                                    acuCost[i,j-1]):
              i
                  = i-1
          elif acuCost[i,j-1] == min(acuCost[i-1,j-1],
                                    acuCost[i-1,j],
                                    acuCost[i,j-1]):
                  = j-1
          else:
                  = i-1
                  = \dot{1} - 1
      path.append([j,i])
> path.append([0,0])
```

#### Create a function that plots the path

3. Search the optimal path

```
> def pltCostAndPath(acuCost,path,xlab="X",ylab="Y",clrmap="viridis"):
            = [pt[0] for pt in path]
    Хq
            = [pt[1] for pt in path]
    ру
    imgplt
            = pltDistances (acuCost,
                           xlab=xlab,
                           ylab=ylab,
                            clrmap=clrmap)
    plt.plot(px,py)
    return imaplt
> pltCostAndPath(acuCost,path,clrmap='Reds')
```

## Now we want to plot the path; this code function helps us draw the path



3. Search the optimal path

Rmb your accumulated cost is dependent on the distance:

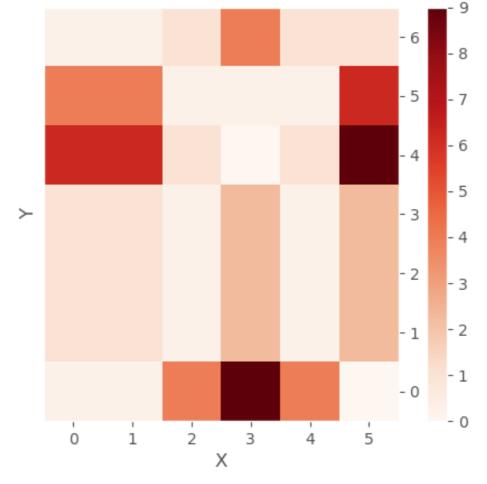
$$A[i,j] = D[i,j] + \dots$$

acuCost

-6
-16
-14
-12
-4
-10
-3
-8
-2
-6
-1
-4
-2

Calculate the cost based on dists,
 which can be considered as a measure for similarity / distance

#### dists

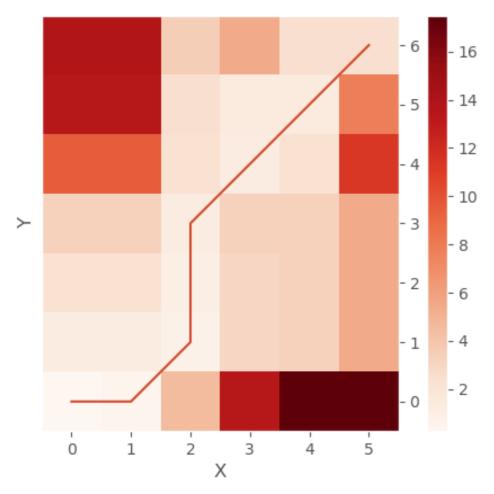


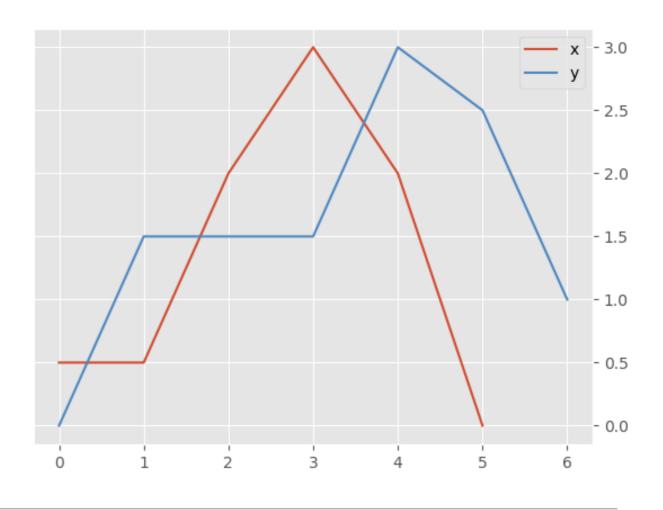
The implication

3. Search the optimal path

## How to illustrate the mapping between the 2 signals (Right) based on the wrapping path in the Cost Matrix (Left)?

#### acuCost





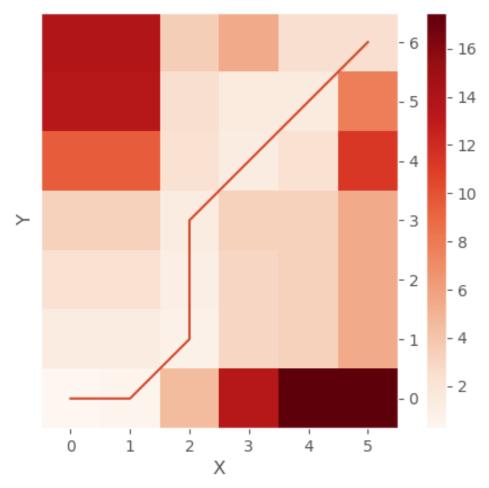
The implication

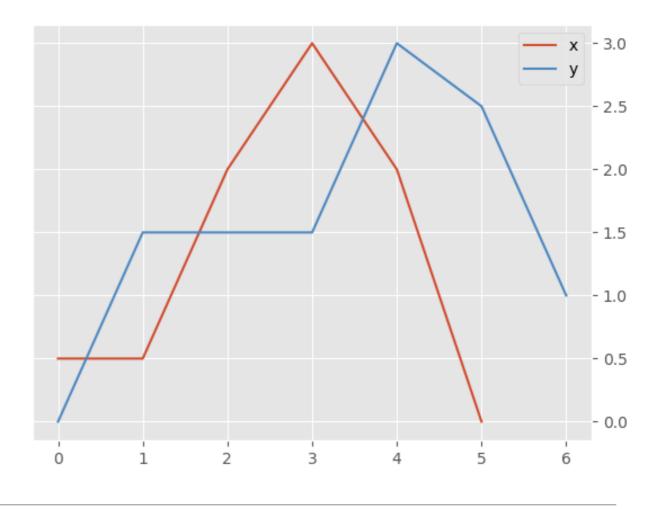
3. Search the optimal path

### How to plot a line that shows the correspondence between point 5 in X and point 6 in Y?

**Ans:** plt.plot([5, 6], [X[5], Y[6]])



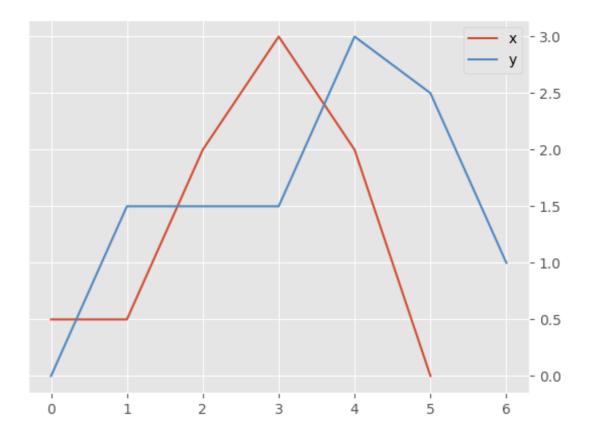


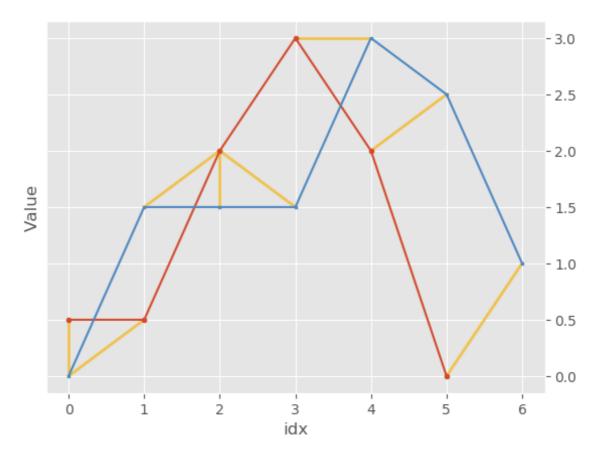


3. Search the optimal path

 Plot the mapping of points between two signals

```
> def pltWarp(s1,s2,path,xlab="idx",ylab="Value"):
    imaplt
                 = plt.figure()
                                  idx1 and idx2 are the interested time parameters
    for [idx1,idx2] in path:
                                                       Plot the connections between
        plt.plot([idx1,idx2],[s1[idx1],s2[idx2]], s1 and s2 (yellow lines)
                  color="C4",
                  linewidth=2)
    plt.plot(s1,
                                                                                            - 3.0
              10-1,
              color="CO",
                                                                                            - 2.5
              markersize=3)
    plt.plot(s2,
                                                                                            - 2.0
              's-',
              color="C1",
                                                                                            - 1.5
              markersize=2)
    plt.xlabel(xlab)
                                                                                            - 1.0
    plt.ylabel(ylab)
                                                                                            - 0.5
    return imaplt
                                                                                             - 0.0
                                                                      3
> pltWarp(x,y,path)
                                                                      idx
```







Another example

## Another example with 2 different signals

#### -1.00 -0.75 -0.50 -0.25 -0.00 -0.25 -0.50 -0.75 -0.75 -1.00

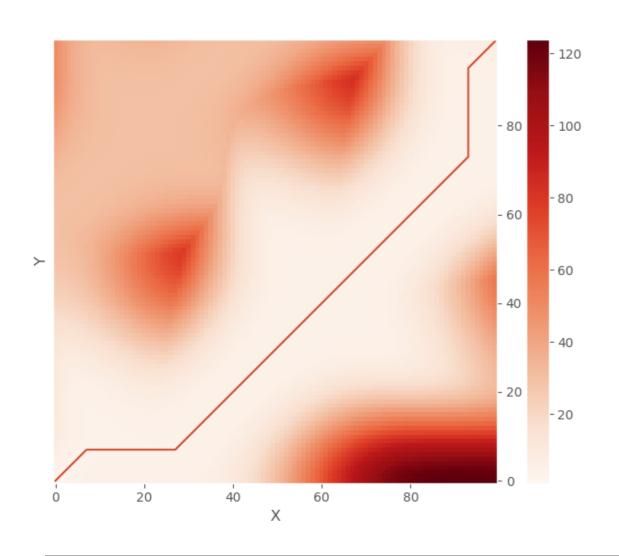
#### Define two signals as:

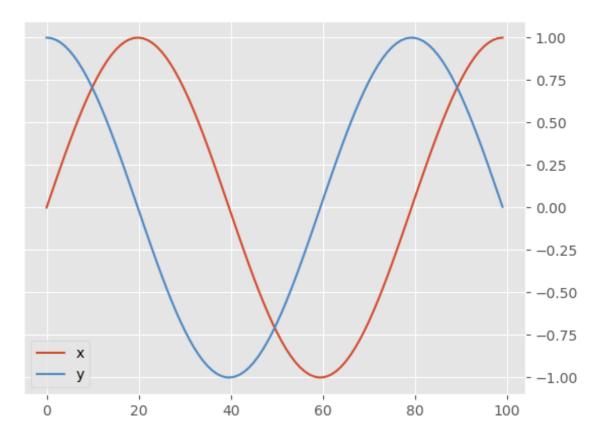
```
> x = np.sin(np.linspace(0,7.85,100))
> y = np.cos(np.linspace(0,7.85,100))
```

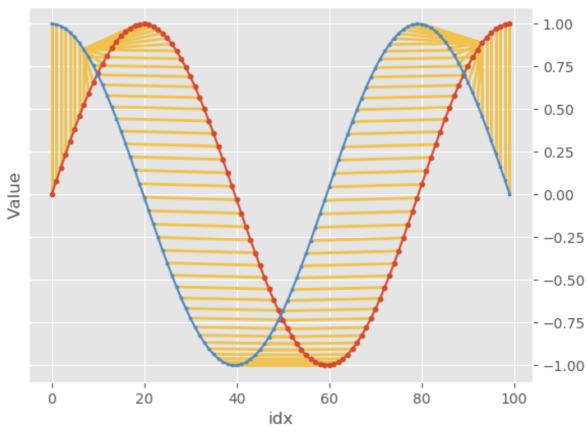
#### Plot the two signals

```
> plt.figure()
> plt.plot(x,
           color="CO",
           label='x')
> plt.plot(y,
           color="C1",
           label='y')
> plt.legend()
```

Another example







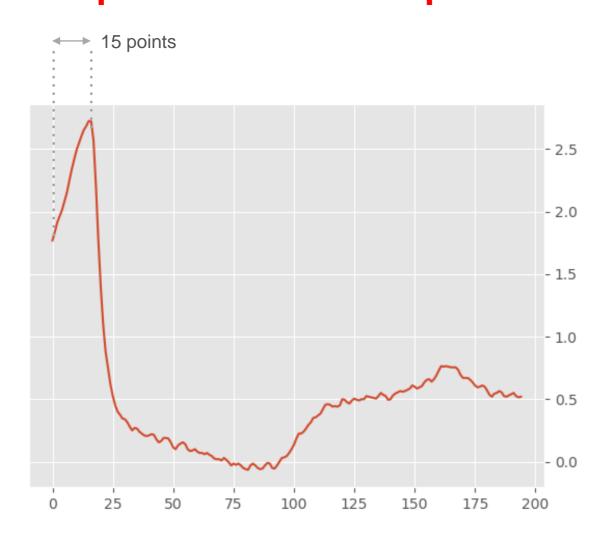
#### Back to the problem

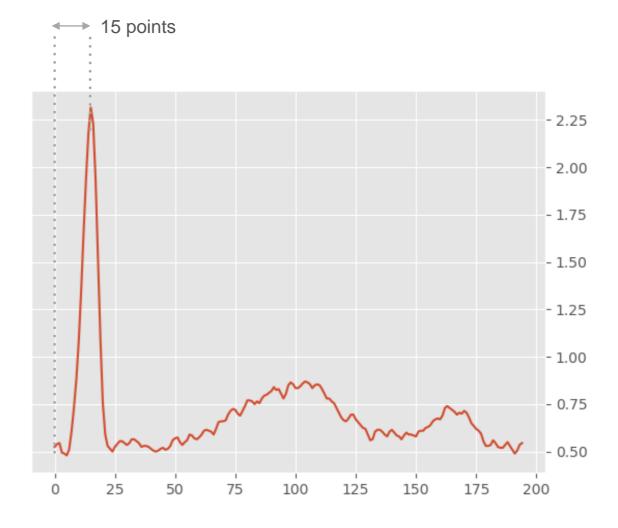
How to start?

Use peak (i.e. findpeak) to estimate start of each ECG heartbeat by shifting 15 points to left from peak

 Before we compute similarity, must segment individual heartbeat signal

 With signal segmented, perform DTW





#### Workshop

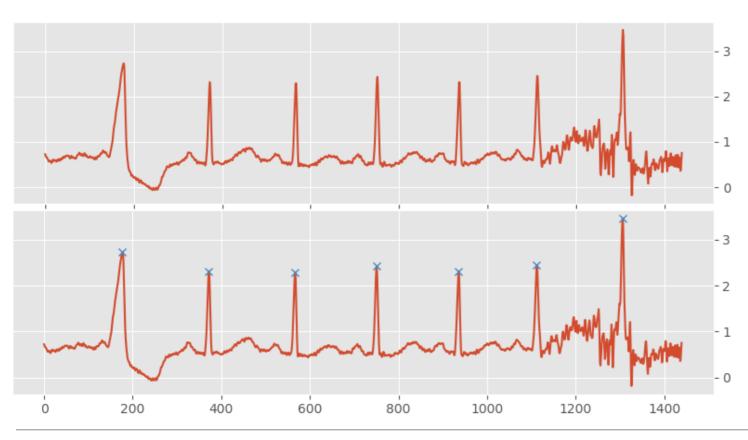
To start

You need to complete a code function to chop (segment) each ECG heartbeat from the complete signal!

Load the data

 Create a function with the below signature. The output is a list consists of all the ECG segments in a ECG signal

def extractECG(ecg,pks,offset=15):



#### **General Procedures:**

- 1. Load data
- 2. Perform peak detection
- 3. Extract segments based on the detected peaks

#### Workshop

Compute the accumulated costs, plot the optimal paths and warp

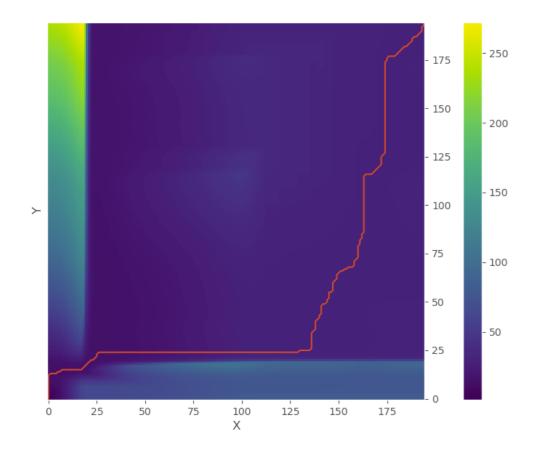
# If you are done early, try on other signals (i.e. other than the 2<sup>nd</sup> column of the ecg2D data)

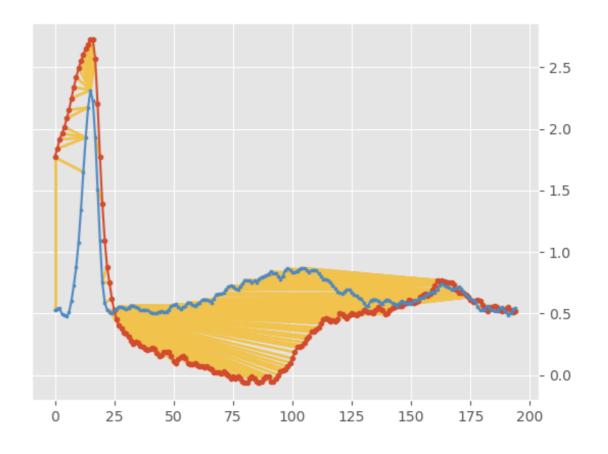
 Make the comparisons between segment

1 and 2

2 and 3

2 and 6



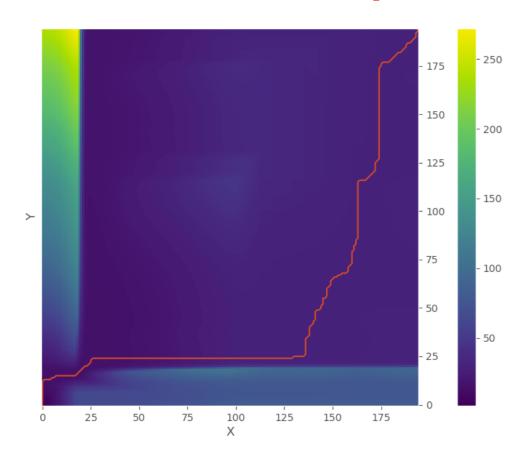


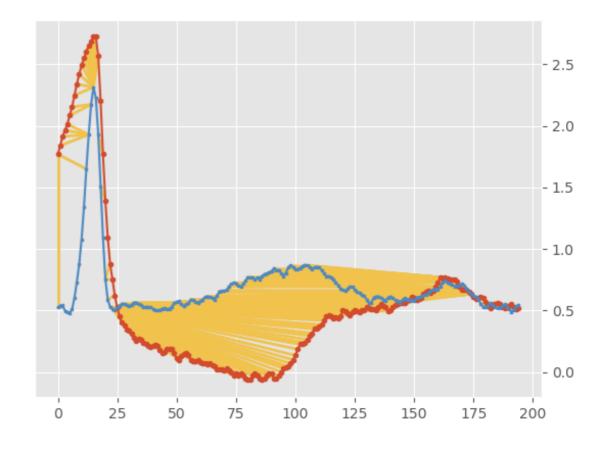
#### Workshop Hints for extractECG function

#### Tips how to chop the ecg:

Assume, you have all the peaks being identified, and 'pks' is the list to hold all the position of peaks. Assume 'ecg' is the ECG signals. Then you need to go through a "for-loop", for each iteration, you do:

#### Another Tip: You need to ignore the last peak





#### Workshop Hints for extractECG function

#### Tips how to chop the ecg:

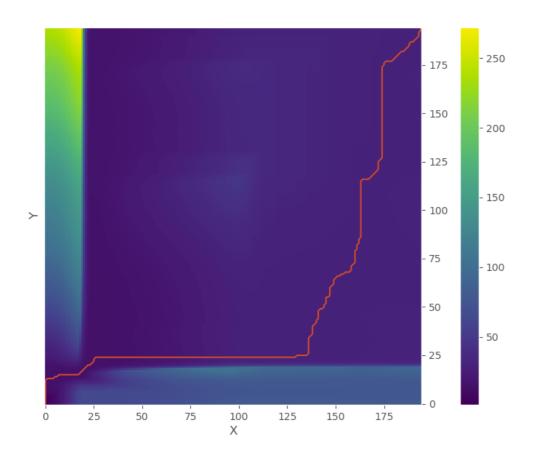
```
def extractECG(ecg,pks,offset=15):
    seqs = []
    if pks[0]-offset < 0:
        start = 1
    else:
        start = 0
    for i in range(start, len(pks)-1):
        seg = ecg[(pks[i]-offset):(pks[i+1]-offset)]
        segs.append(seg)
    return segs
EcgSegs = extractECG(ECGs, Pks)
```

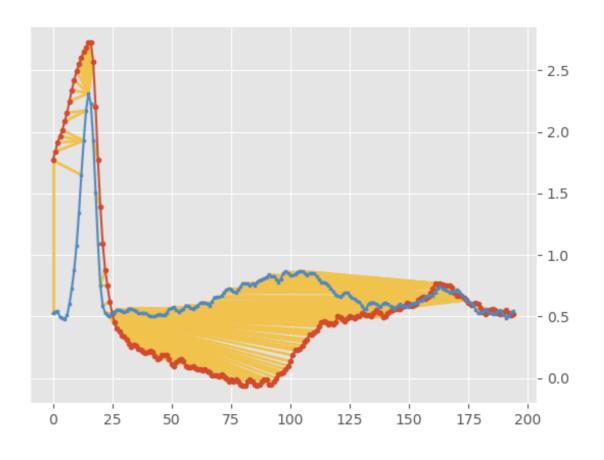
#### **Workshop Hints for Performing DTW on ECG segments**

#### Example for comparing between segments 1 and 2:

```
dist = computeDists(segs[0], segs[1])
acuCost = computeAcuCost(dist)

(path,cost) = doDTW(segs[0], segs[1], dist, acuCost)
pltCostAndPath(acuCost, path)
pltWarp (segs[0], segs[1], path)
```





#### **Appendix: Comparison among Various Methods**

To measure similarity between signals

	Pros	Cons		
Euclidean/Manhattan distance	Easy to implement; straightforward	Only works if the signals to be compared are of same length and preferably similar shape		
DTW	Not necessary for signals to be of same length and shape;	Heavy computational burden (worse than Euclidean/Manhattan distance method);		
	Computes faster than LCSS	Unable to work if one of the signals is a partial type		
Longest common	Not necessary for signals to be of same length and shape;	Heavy computational burden;		
subsequence (LCSS)	Allows for the use of partial and noisy signals	Computes slower than DTW		
Developed Longest	Not necessary for signals to be of same length and shape;			
Common Subsequence	Allows for the use of partial and noisy signals;	Heavy computational burden;		
(DLCSS)	Most accurate among the other methods	Computes slower than DTW and LCSS		