

CAP 4453

Robot Vision

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Administrative details

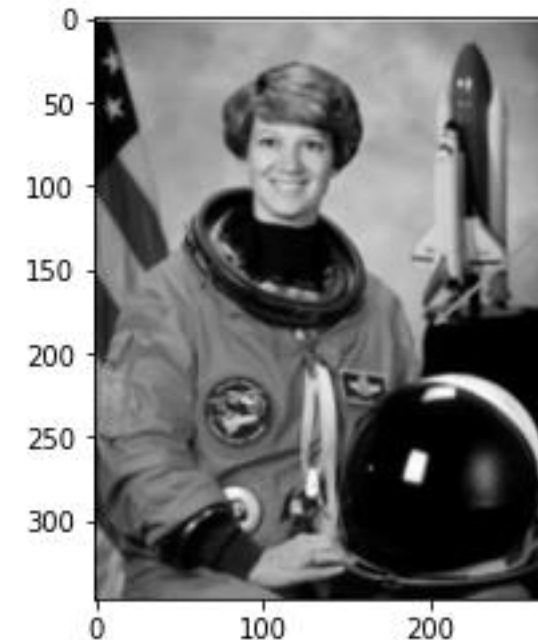
- Issues submitting homework

- #make sure you are the right data type and values. You can use histogram to check it
- # 0-255 if uint8
- # 0-1 if float

```
import matplotlib.pyplot as plt
import cv2

imFile = 'C:\\Users\\gonza\\OneDrive\\Teaching\\CAP4453\\class12\\img.png'

im = cv2.imread(imFile)
img = cv2.cvtColor(im, cv2.COLOR_BGR2GRAY)
plt.imshow(img, cmap='gray'); plt.show()
```



Administrative details

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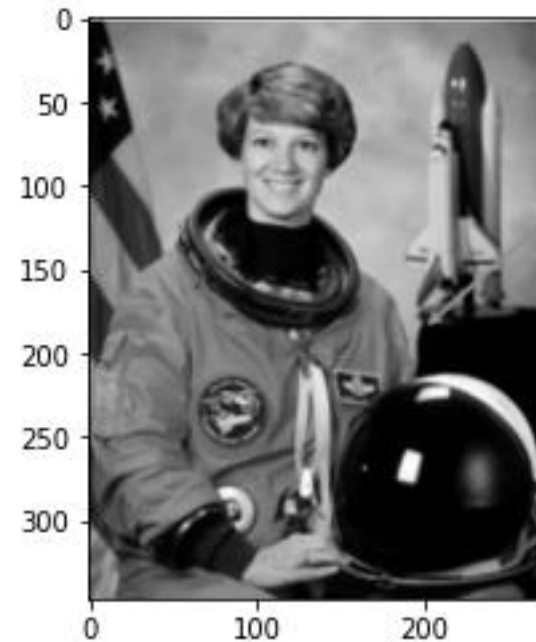
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img = cv2.cvtColor(im, cv2.COLOR_BGR2GRAY)
plt.imshow(img, cmap='gray'); plt.show()
```

```
img2= img+10
plt.imshow(img2, cmap='gray'); plt.show()
```



Administrative details

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- #make sure you are the right data type and values. You can use histogram to check it
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import matplotlib.pyplot as plt
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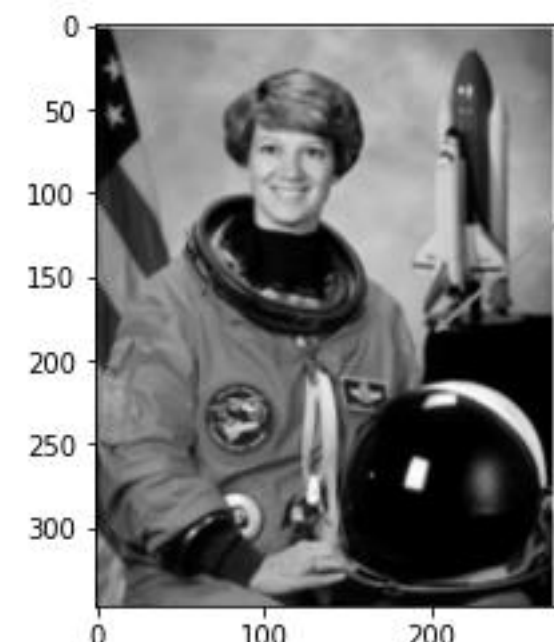
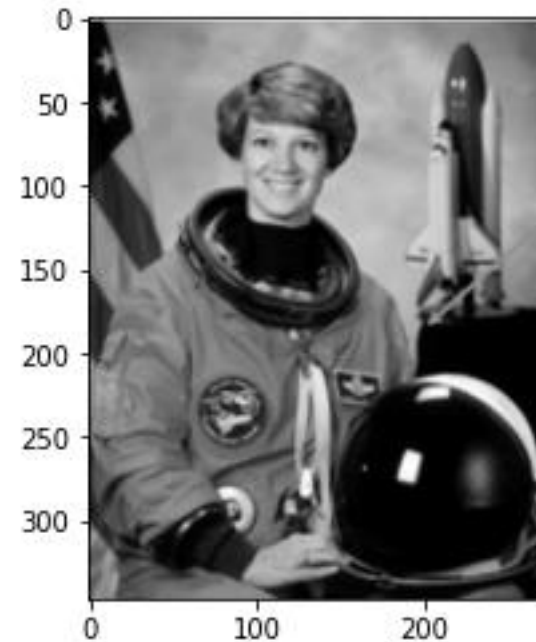
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img = cv2.cvtColor(im, cv2.COLOR_BGR2GRAY)
plt.imshow(img, cmap='gray'); plt.show()
```

```
img2= img+10
```

```
img3 = img.astype(float)
img4 = img3+10
print(img4.max())
plt.imshow(img4, cmap='gray'); plt.show()
```

→ 265.0





Administrative details

- Issues submitting homework
 - `return kernel / np.sum(kernel)` #note that this normalize the filter to 1. is it desirable?
 - Try it to normalize to
 - `return kernel / (2*np.sum(kernel))`
 - `return 2*kernel / np.sum(kernel)`
 - `return kernel / np.sum(kernel)`
 - Check the obtained brightness of the image



Administrative details

- Issues submitting homework

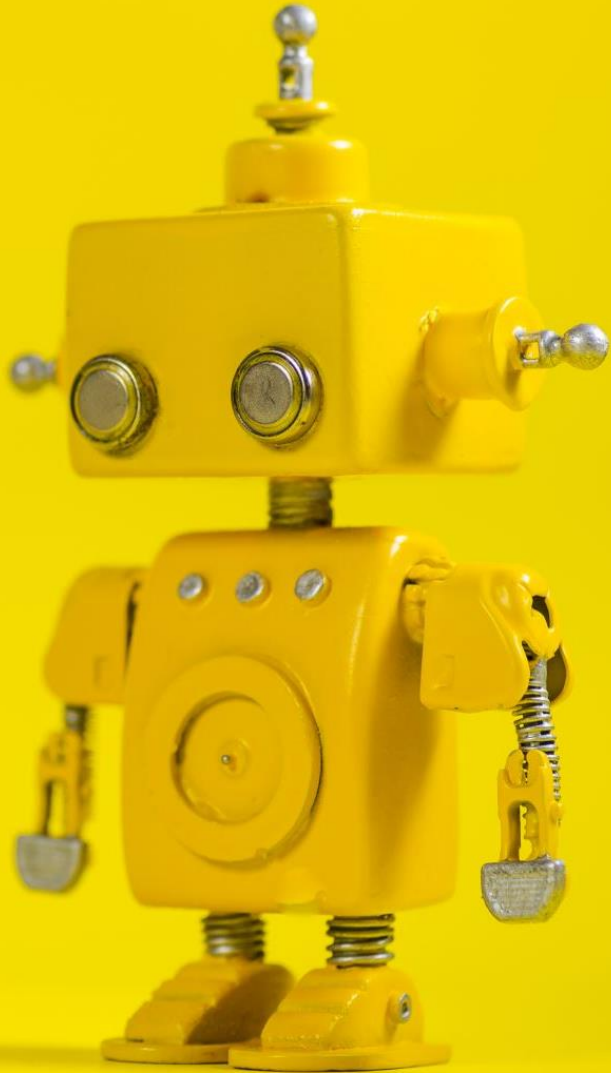
- Plot your partial results
 - Some plotting functions update the range when you are using floats (astronaut image)
 - if you want to be sure on what you are getting:
 - Make sure your image is in the range 0 to 255 (many ways to do it: normalize by max value, `min(255,img)`)
 - Convert the type back to uint8 using `astype(uint8)`
 - Then plot



Credits

- Some slides comes directly from:
 - Yogesh S Rawat (UCF)
 - Noah Snavelly (Cornell)
 - Ioannis (Yannis) Gkioulekas (CMU)
 - Mubarak Shah (UCF)
 - S. Seitz
 - James Tompkin
 - Ulas Bagci
 - L. Lazebnik

Short Review from last class



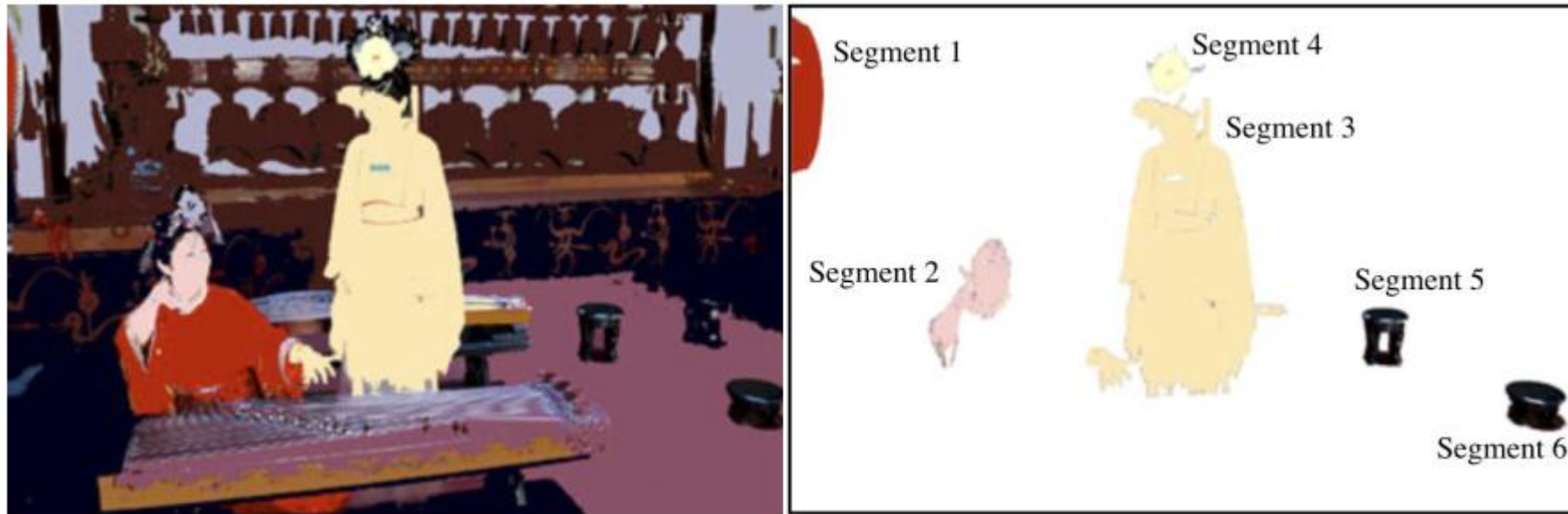


Outline

- ~~Image segmentation basics~~
- ~~Thresholding based~~
 - ~~Binarization~~
 - ~~Otsu~~
- ~~Region based~~
 - ~~Merging~~
 - ~~Splitting~~
- Clustering based
 - K-means (SLIC)

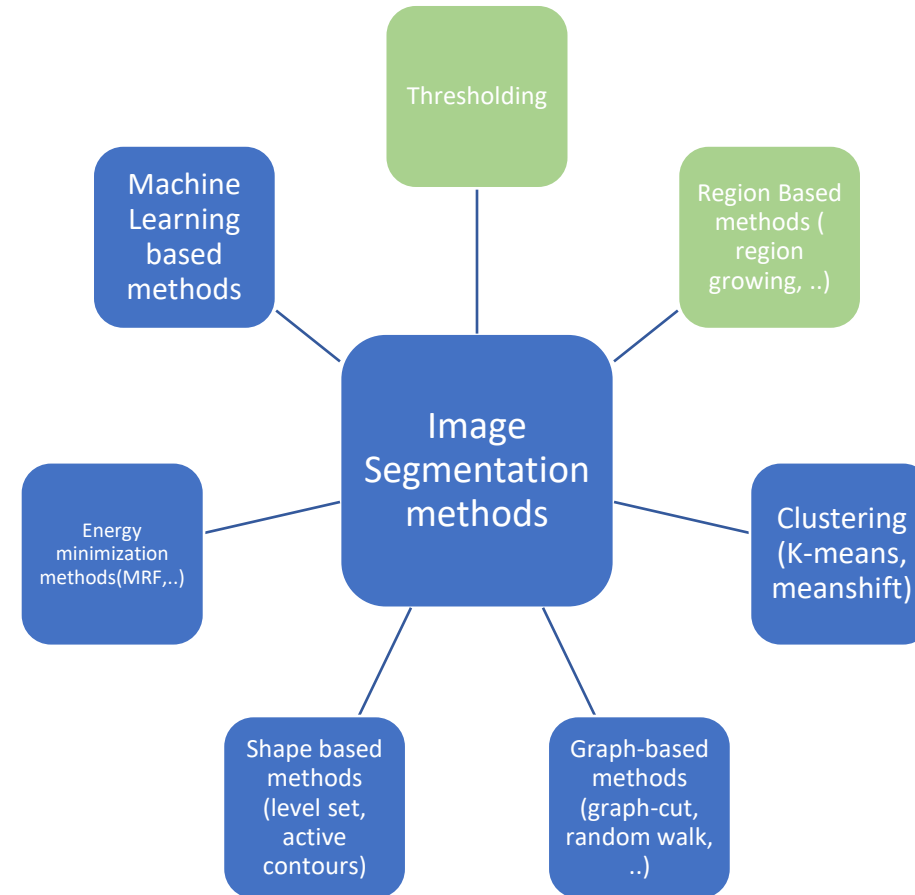
Image segmentation

- Image segmentation partitions an image into regions called segments.



- Image segmentation creates segments of connected pixels by analyzing some similarity criteria:
 - intensity, color, texture, histogram, features

Image segmentation methods



Otsu thresholding

- Definition: The method uses grey-value histogram of the given image I as input and aims at providing the best threshold (foreground/background)
- Otsu's algorithm selects a threshold that maximizes the between-class variance σ_b^2 or minimize within-class variance σ_w^2

Option 1: maximum of:

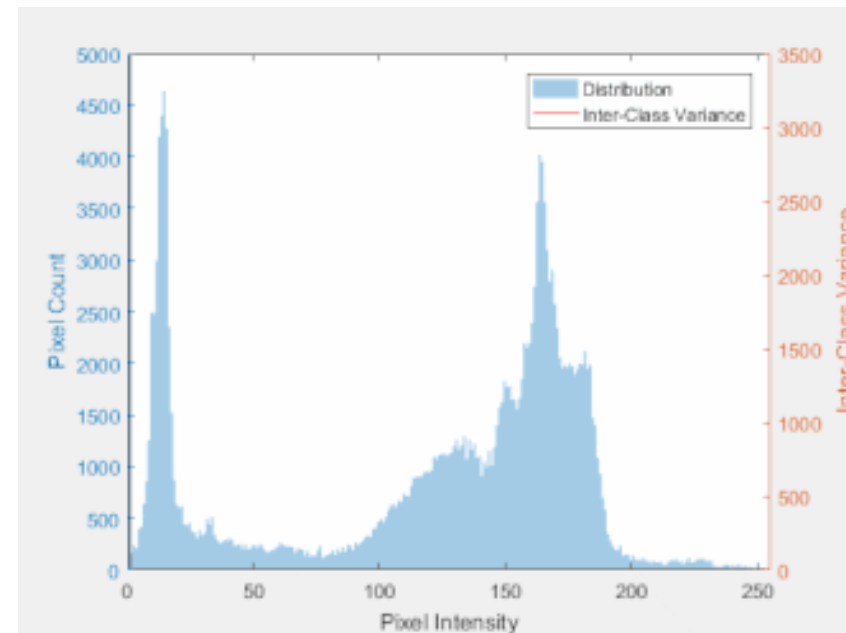
$$\sigma_b^2(t) = w_1(t)w_2(t)[\mu_1(t) - \mu_2(t)]^2$$

$$\mu_1(t) = \sum_{i=1}^t \frac{iP(i)}{w_1(t)}$$

$$w_1(t) = \sum_{i=1}^t P(i)$$

$$\mu_2(t) = \sum_{i=t+1}^I \frac{iP(i)}{w_2(t)}$$

$$w_2(t) = \sum_{i=t+1}^I P(i)$$



Otsu thresholding

- Definition: The method uses grey-value histogram of the given image I as input and aims at providing the best threshold (foreground/background)
- Otsu's algorithm selects a threshold that maximizes the between-class variance σ_b^2 .

Option 2: minimum of:

$$\sigma_w^2(t) = w_1(t)\sigma_1^2(t) + w_2(t)\sigma_2^2(t)$$

$$w_1(t) = \sum_{i=1}^t P(i) \quad P(i) = \frac{n_i}{n}$$

$$w_2(t) = \sum_{i=t+1}^I P(i)$$

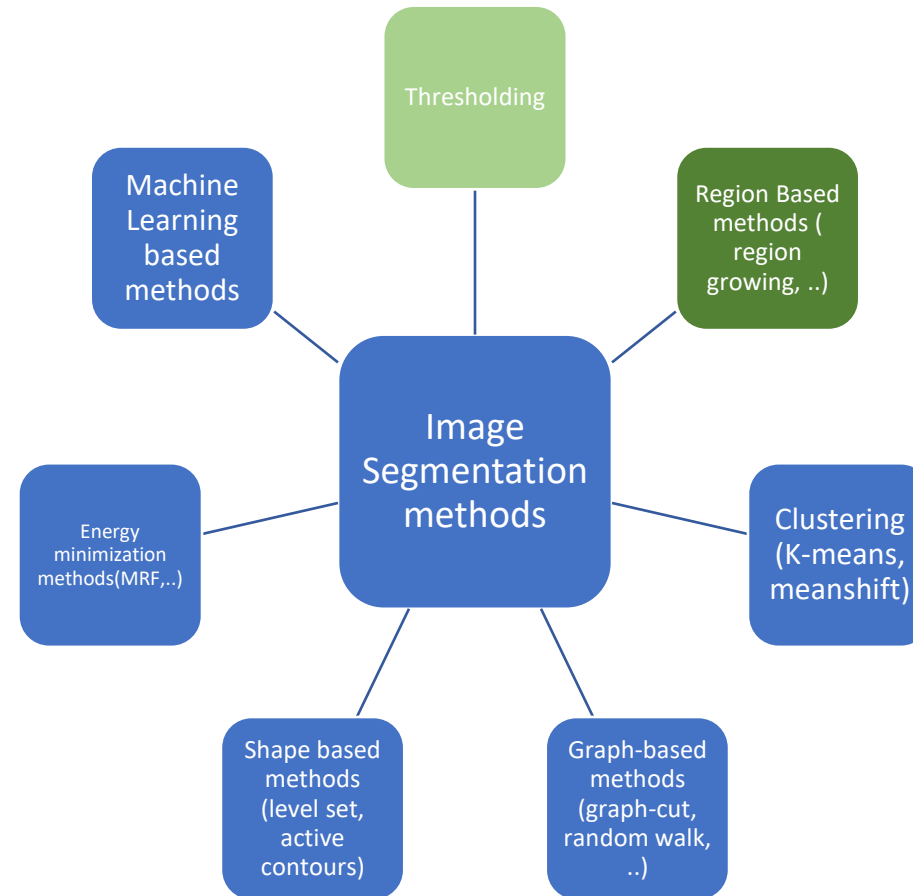
$$\sigma_1^2(t) = \sum_{i=1}^t [i - \mu_1(t)]^2 \frac{P(i)}{w_1(t)}$$

$$\sigma_2^2(t) = \sum_{i=t+1}^I [i - \mu_2(t)]^2 \frac{P(i)}{w_2(t)}$$

$$\sigma_{total}^2 = \underbrace{w_1(t)\sigma_1^2(t) + w_2(t)\sigma_2^2(t)}_{\text{within-class variance}} + \underbrace{w_1(t)w_2(t)(\mu_2^2(t) - \mu_1^2(t))}_{\text{between-class variance}}$$

↓
Minimize
↓
Maximize

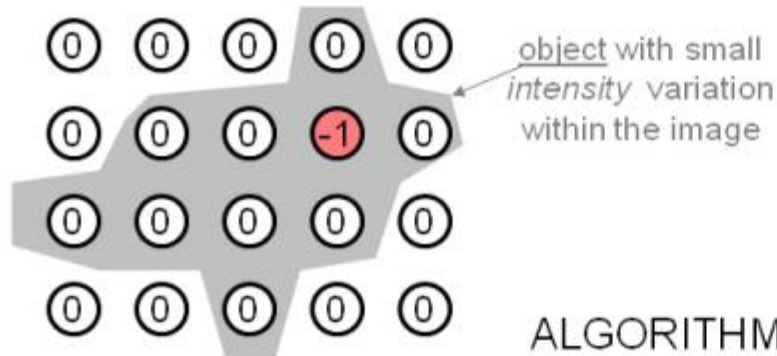
Image segmentation methods



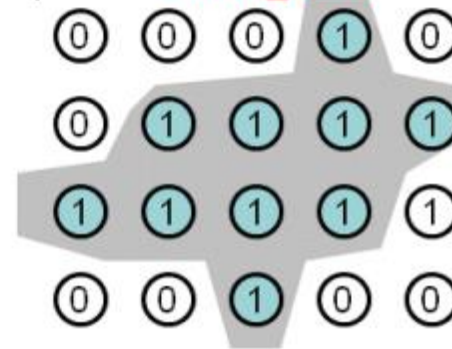
Region Growing Implementation

growRegion: red nodes are the "active_front" (queue or stack)

add seed into **active_front**

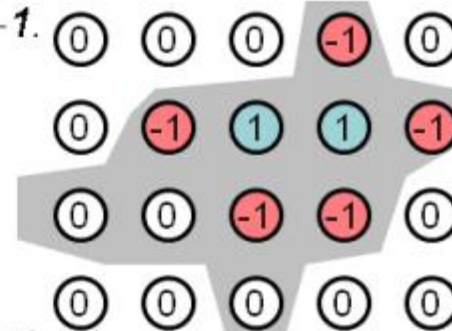
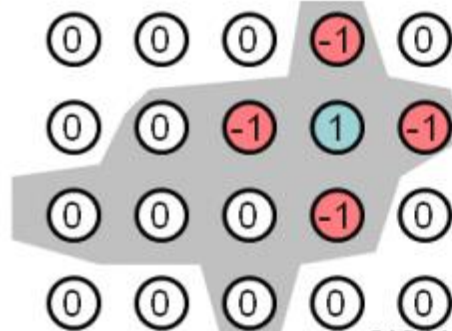


stop when **active_front** is empty



ALGORITHM:

Remove pixel p from **active_front** and mark it as $region[p] = 1$.
 Add all neighbors q such that:
 $region[q] == 0, |I_p - I_q| < T$
 and set $region[q] = -1$.



Region splitting and Merging Segmentation

0	1	0	0	7	7	7	7
1	0	2	2	7	7	7	7
0	2	2	2	7	7	7	7
4	4	2	2	7	7	7	7
0	0	1	1	3	3	7	7
1	1	2	2	3	7	7	7
2	4	3	0	5	7	7	7
2	3	3	5	5	0	7	7

original image

0	1	0	0	7	7	7	7
1	0	2	2	7	7	7	7
0	2	2	2	7	7	7	7
4	4	2	2	7	7	7	7
0	0	1	1	3	3	7	7
1	1	2	2	3	7	7	7
2	4	3	0	5	7	7	7
2	3	3	5	5	0	7	7

split 1

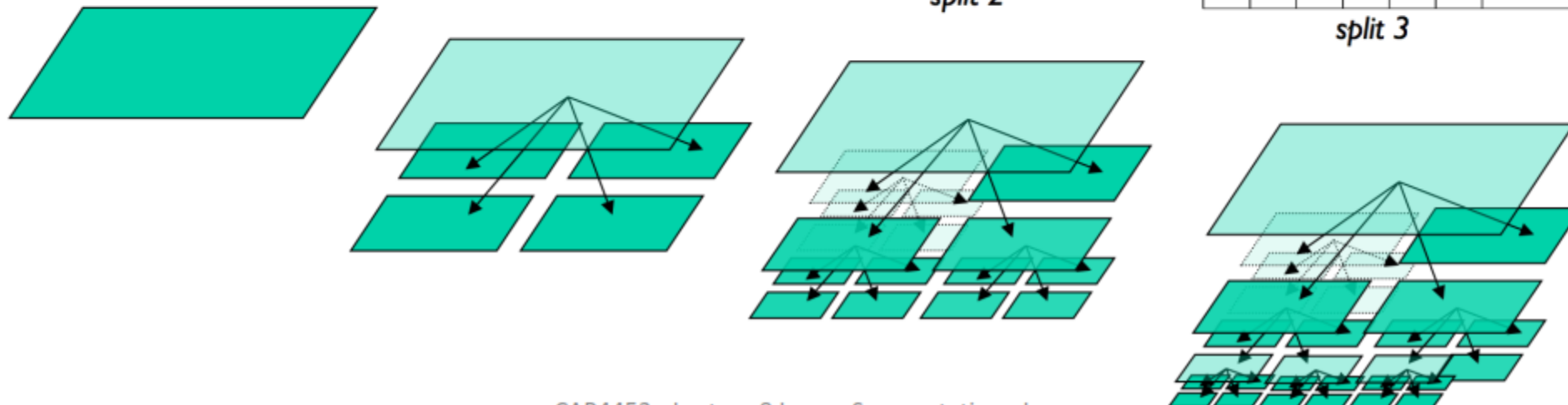
0	1	0	0	7	7	7	7
1	0	2	2	7	7	7	7
0	2	2	2	7	7	7	7
4	4	2	2	7	7	7	7
0	0	1	1	3	3	7	7
1	1	2	2	3	7	7	7
2	4	3	0	5	7	7	7
2	3	3	5	5	0	7	7

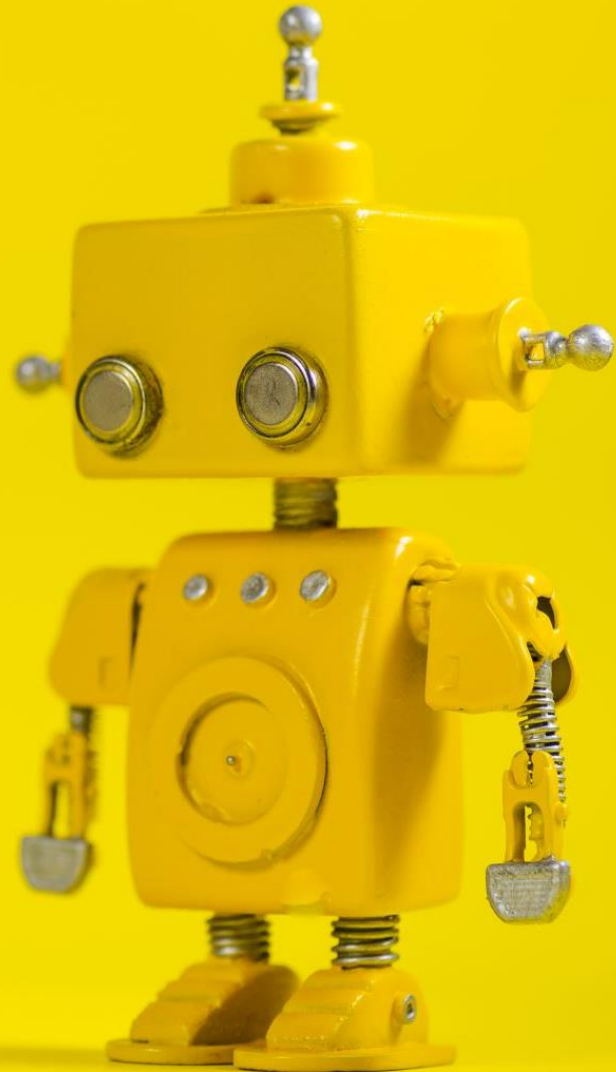
split 2

0	1	0	0	7	7	7	7
1	0	2	2	7	7	7	7
0	2	2	2	7	7	7	7
4	4	2	2	7	7	7	7
0	0	1	1	3	3	7	7
1	1	2	2	3	7	7	7
2	4	3	0	5	7	7	7
2	3	3	5	5	0	7	7

split 3

http://





Robot Vision

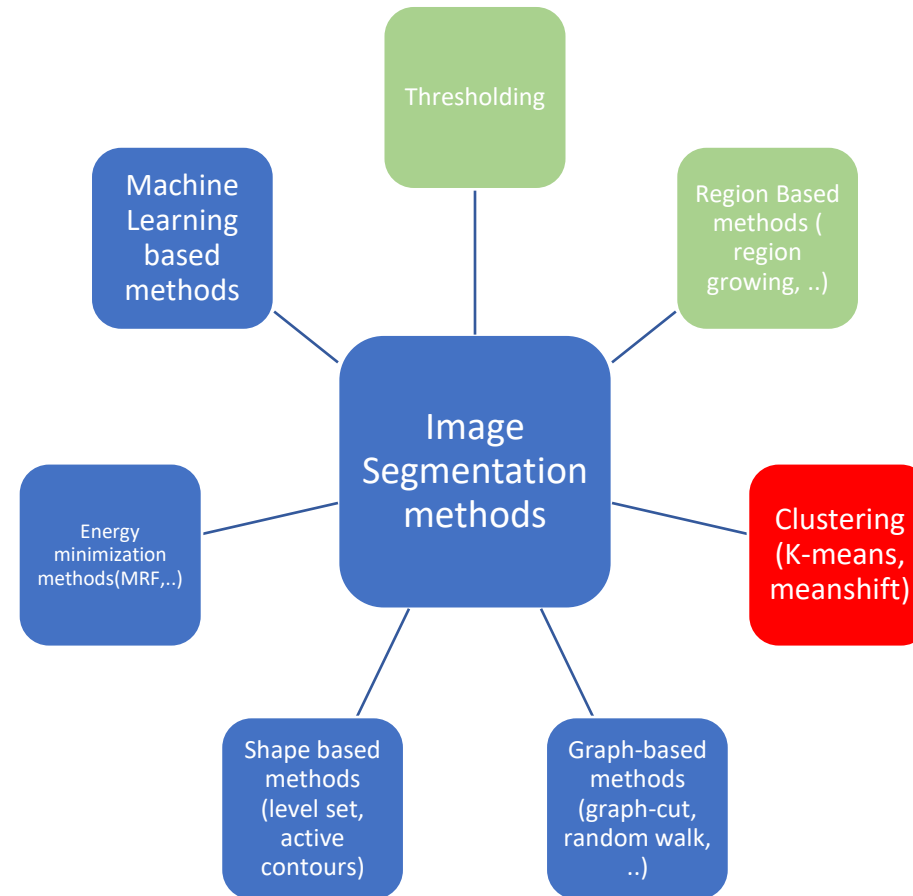
8. Segmentation II



Outline

- ~~Image segmentation basics~~
- ~~Thresholding based~~
 - ~~Binarization~~
 - ~~Otsu~~
- ~~Region based~~
 - ~~Merging~~
 - ~~Splitting~~
- **Clustering based**
 - K-means
 - Superpixels (SLIC)

Image segmentation methods





What is Clustering?

- Organizing data into classes such that:
 - High intra-class similarity
 - Low inter-class similarity
- Finding the class labels and the number of classes directly from the data (as opposed to classification tasks)

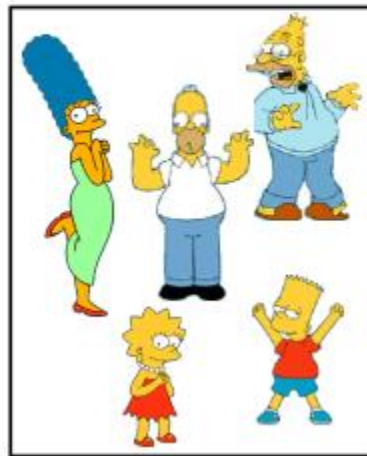
What is a natural grouping ?



What is a natural grouping ?



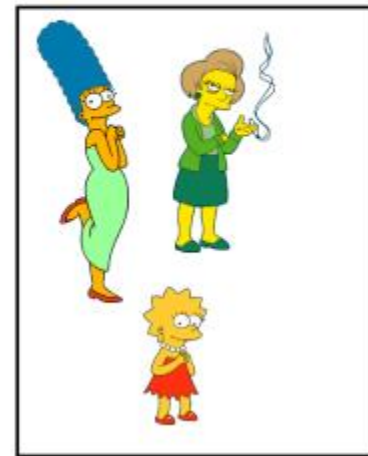
Clustering is subjective



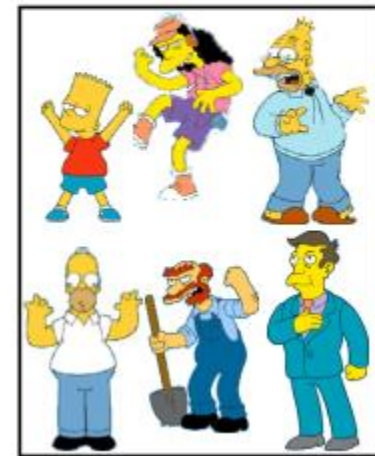
Simpson's family



School employees



Females



Males

What is similarity ?



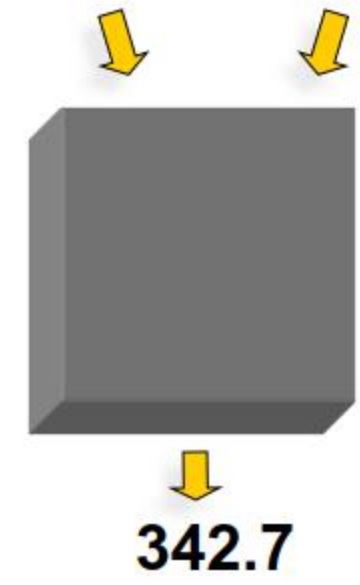
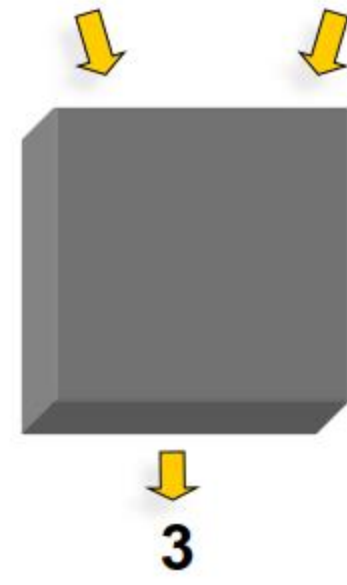
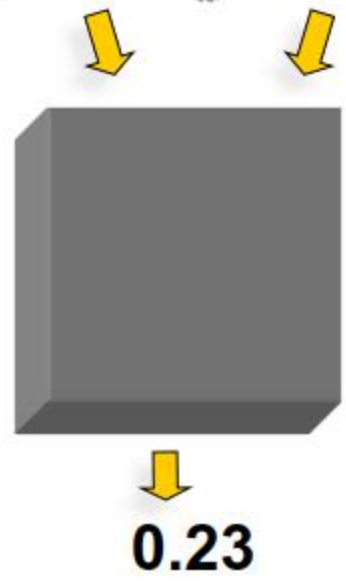
What is similarity ?



Distance metrics



Peter Piotr





Outline

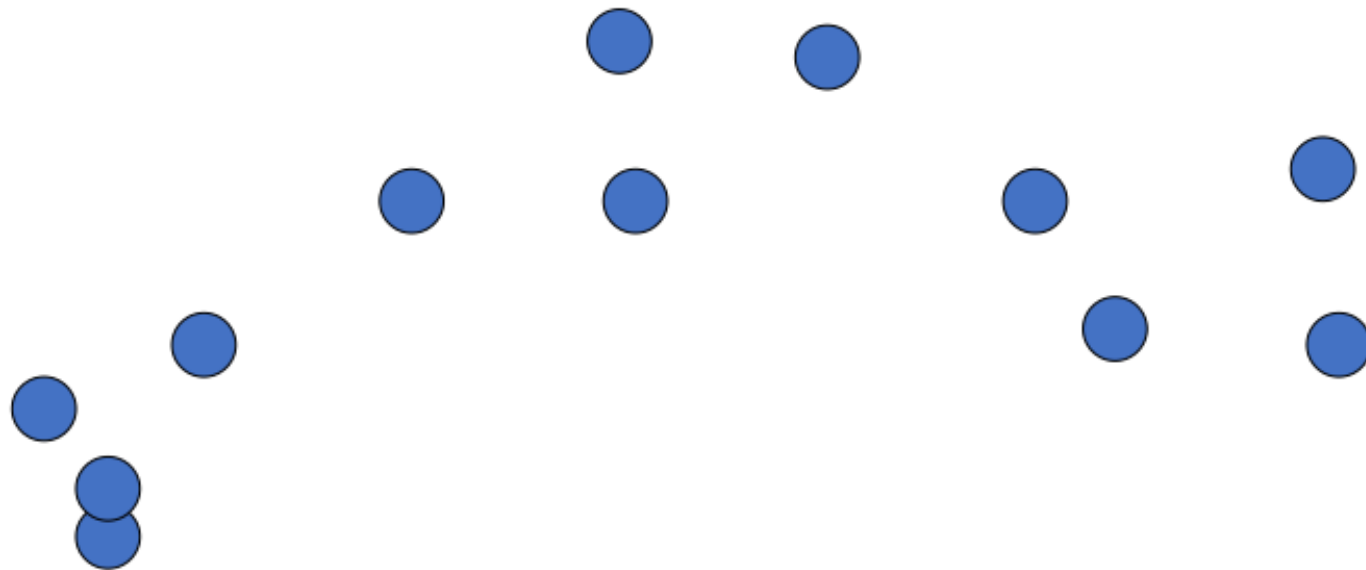
- ~~Image segmentation basics~~
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- ~~Region based~~
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 - ~~Splitting~~
- **Clustering based**
 - **K-means**
 - **Superpixels (SLIC)**



K-means

- Most well-known and popular clustering algorithm:
 - Start with some initial cluster centers
- Iterate:
 - Assign/cluster each example to closest center
 - Recalculate centers as the mean of the points in a cluster

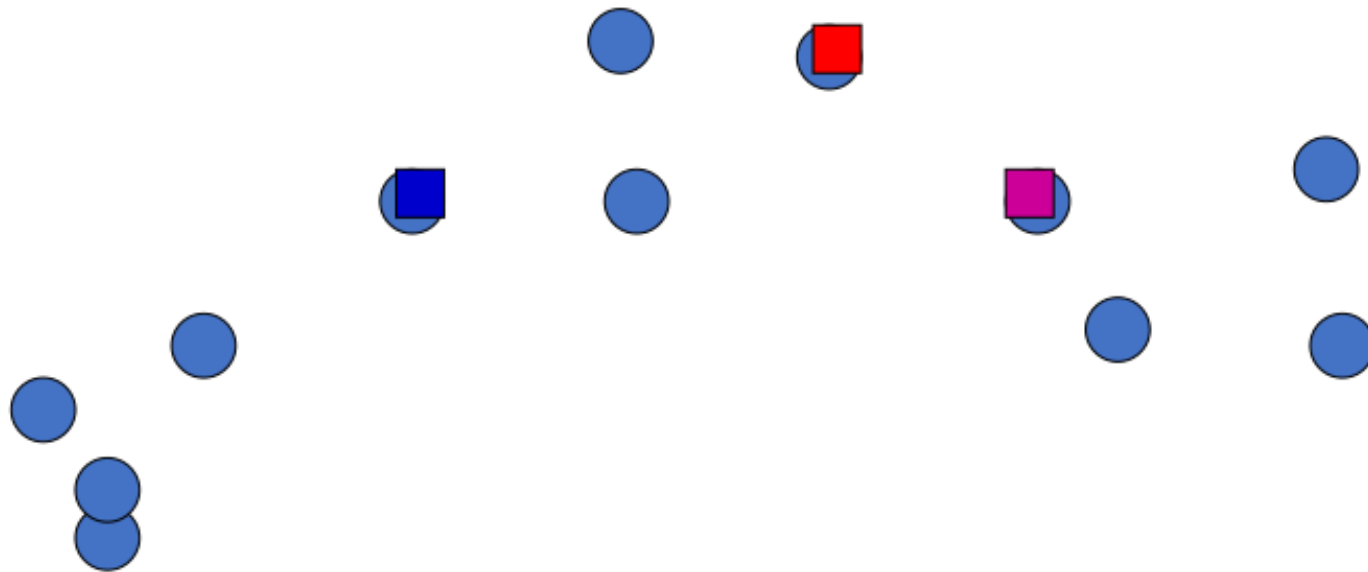
K-means



K-means

Step 0:

- Pick number of classes
- Pick seeds for those classes

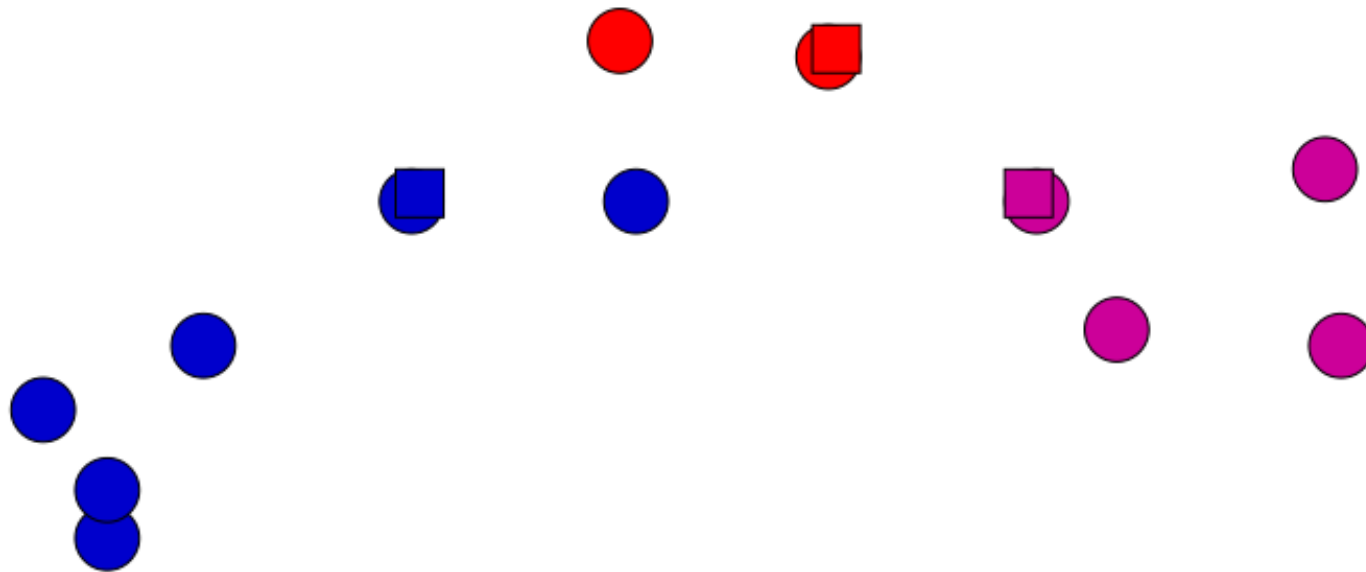


K-means

Iterate:

Assign/cluster each example to closest center

Recalculate centers as the mean of the points in a cluster

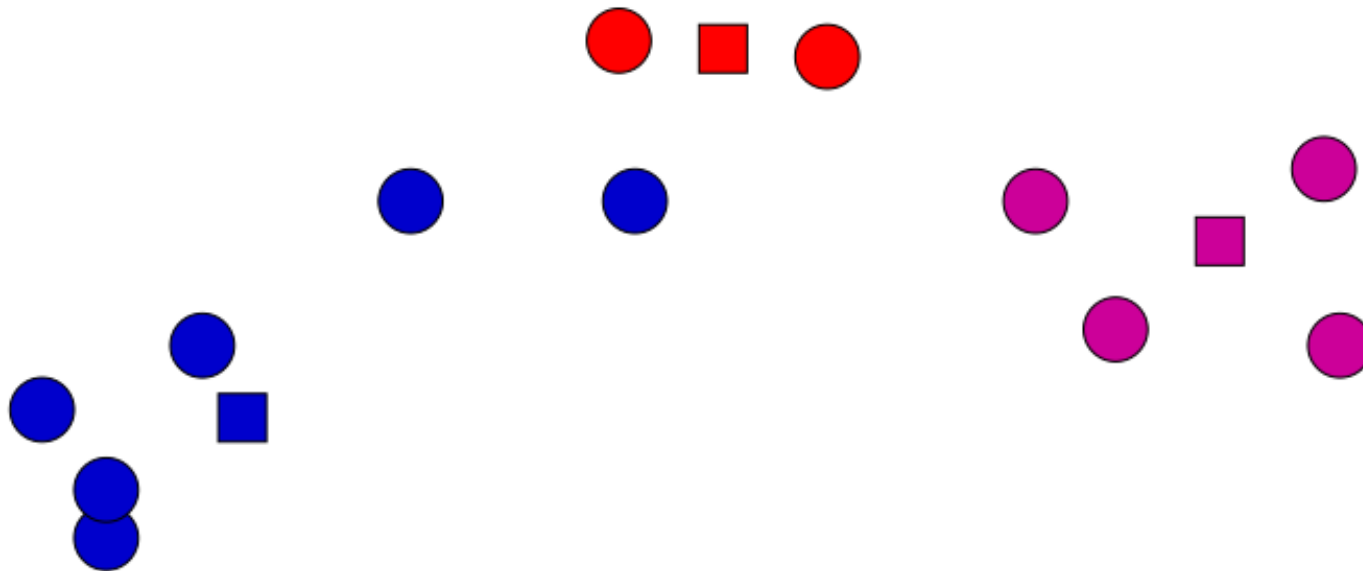


K-means

Iterate:

Assign/cluster each example to closest center

Recalculate centers as the mean of the points in a cluster

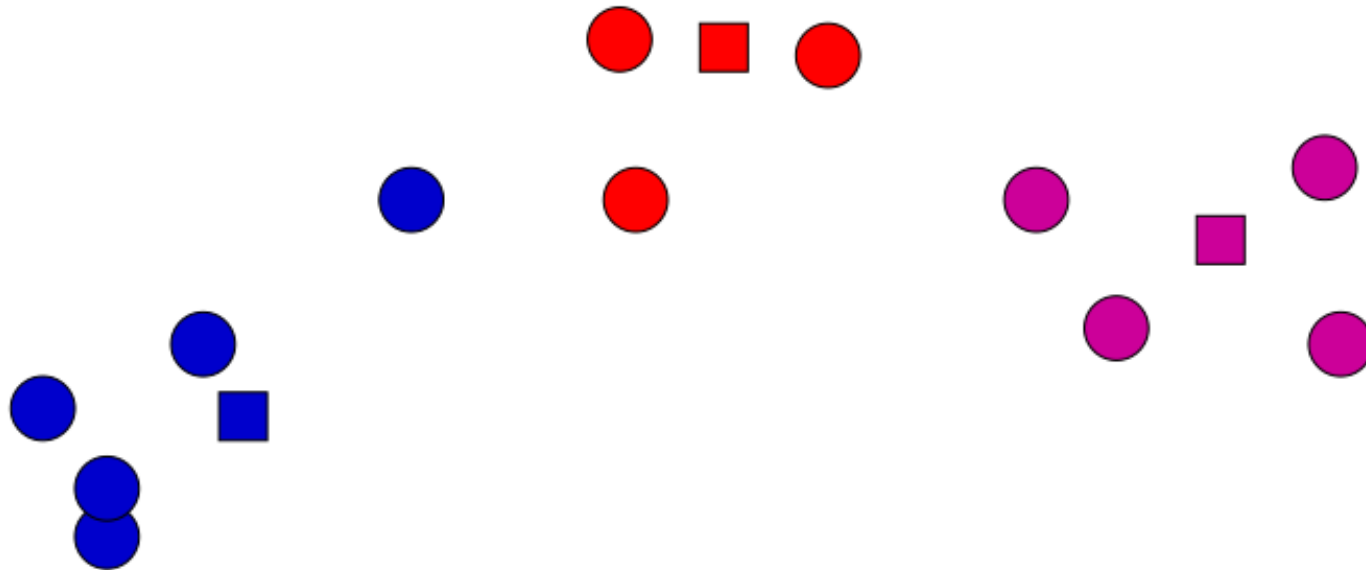


K-means

Iterate:

Assign/cluster each example to closest center

Recalculate centers as the mean of the points in a cluster

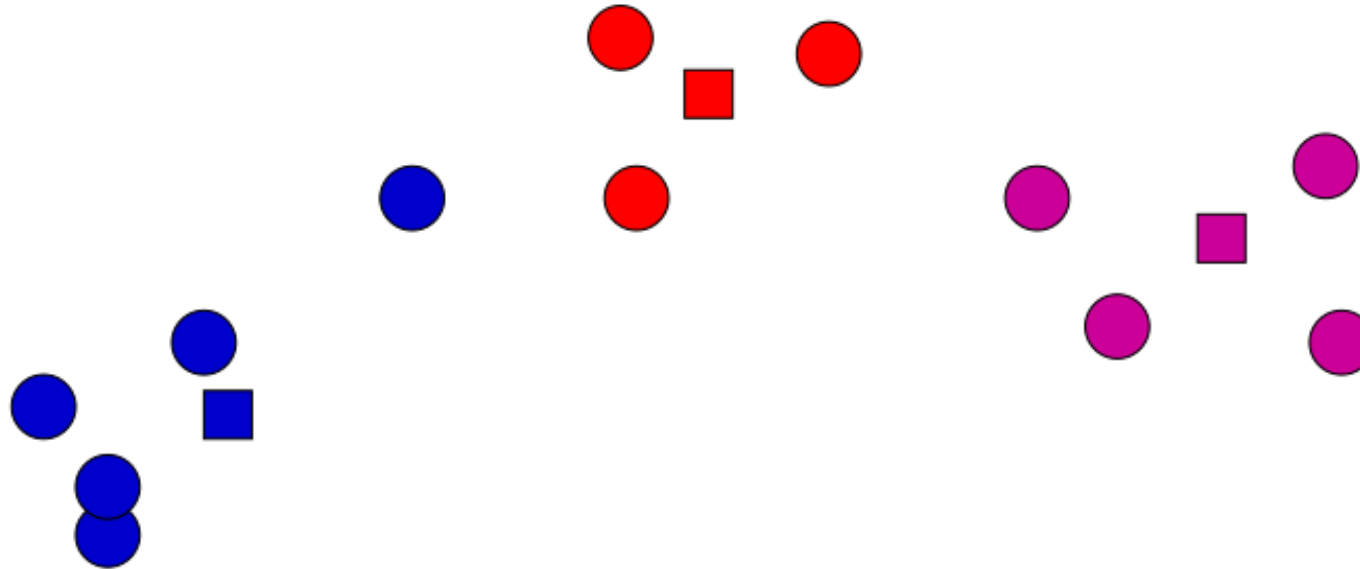


K-means

Iterate:

Assign/cluster each example to closest center

Recalculate centers as the mean of the points in a cluster

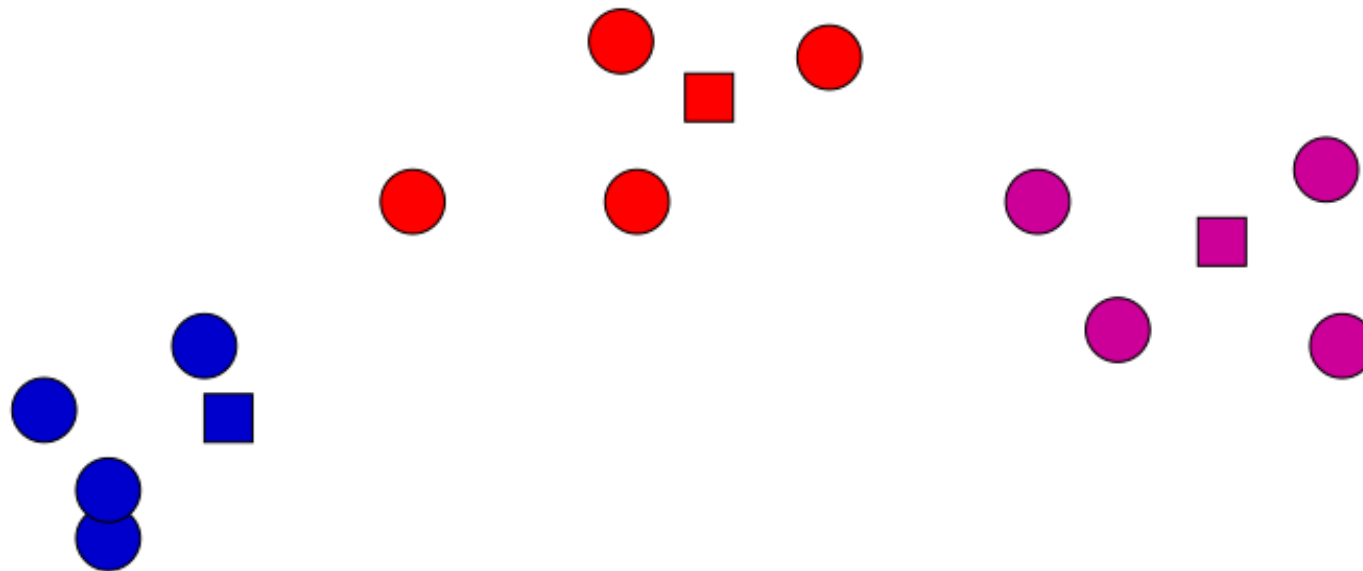


K-means

Iterate:

Assign/cluster each example to closest center

Recalculate centers as the mean of the points in a cluster

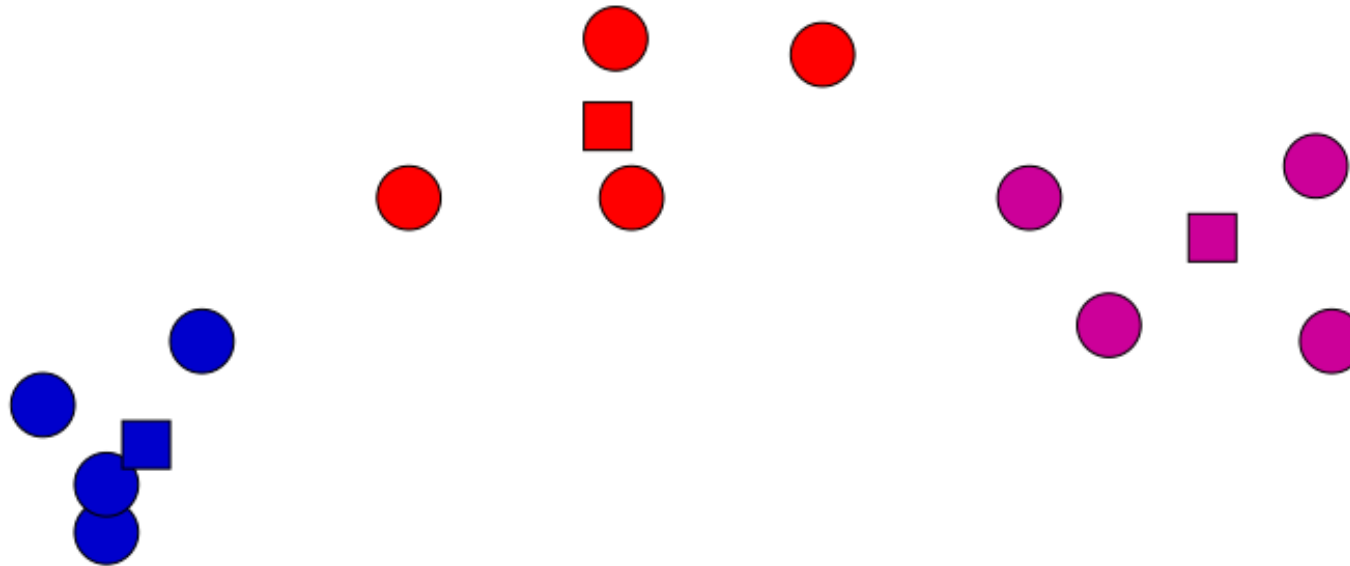


K-means

Iterate:

Assign/cluster each example to closest center

Recalculate centers as the mean of the points in a cluster

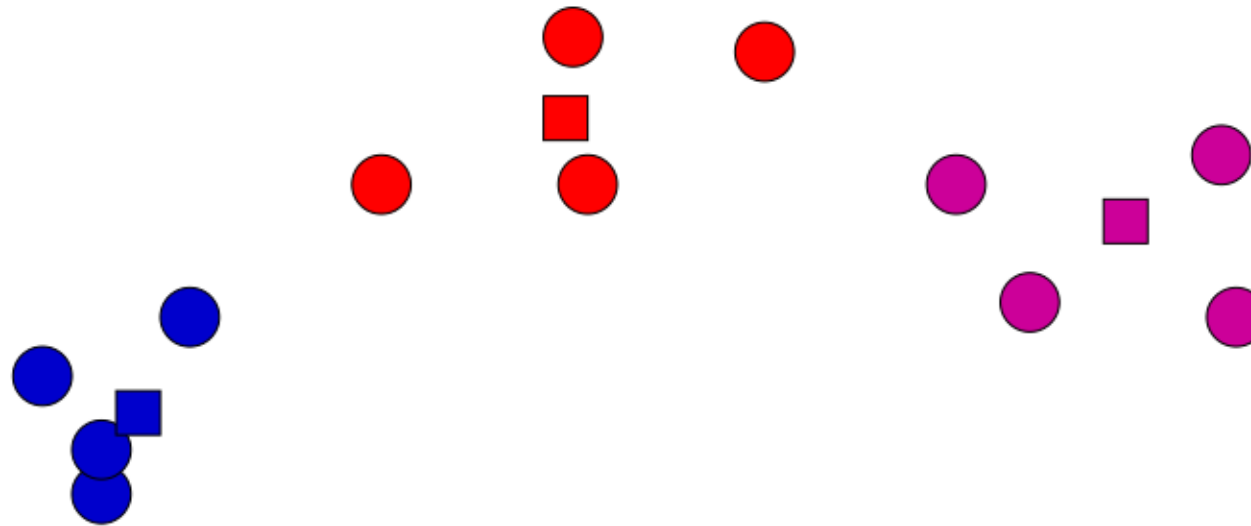


K-means

Iterate:

Assign/cluster each example to closest center

Recalculate centers as the mean of the points in a cluster

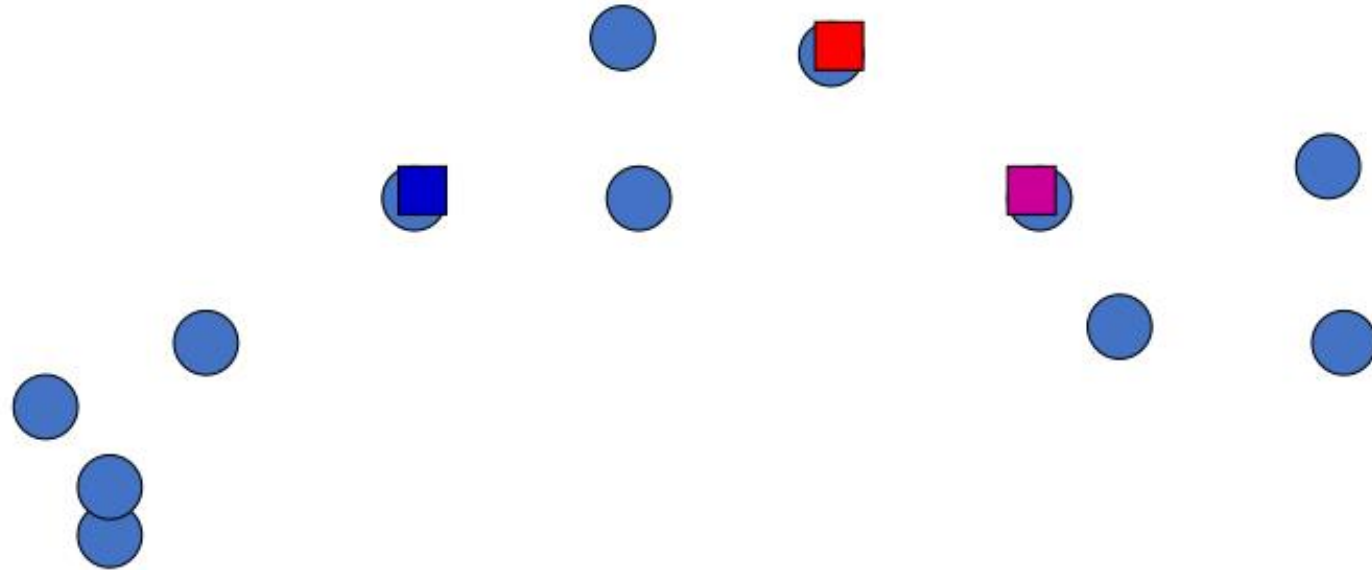


No changes: Done

K-means

Iterate:

- **Assign/cluster each example to closest center**
- Recalculate centers as the mean of the points in a cluster

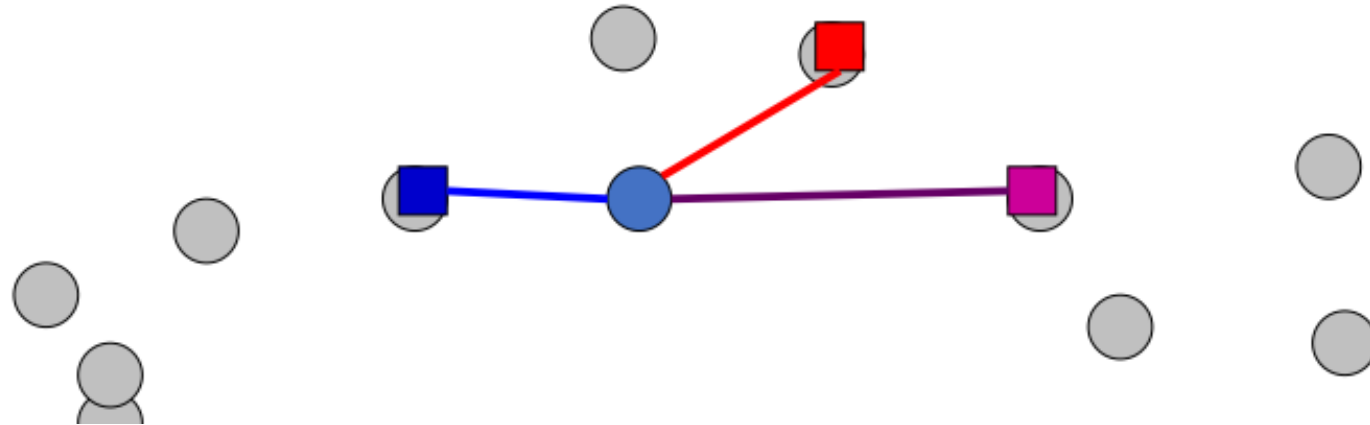


How do we do this?

K-means

Iterate:

- **Assign/cluster each example to closest center**
iterate over each point:
 - get distance to each cluster center
 - assign to closest center (hard cluster)
- Recalculate centers as the mean of the points in a cluster



K-means

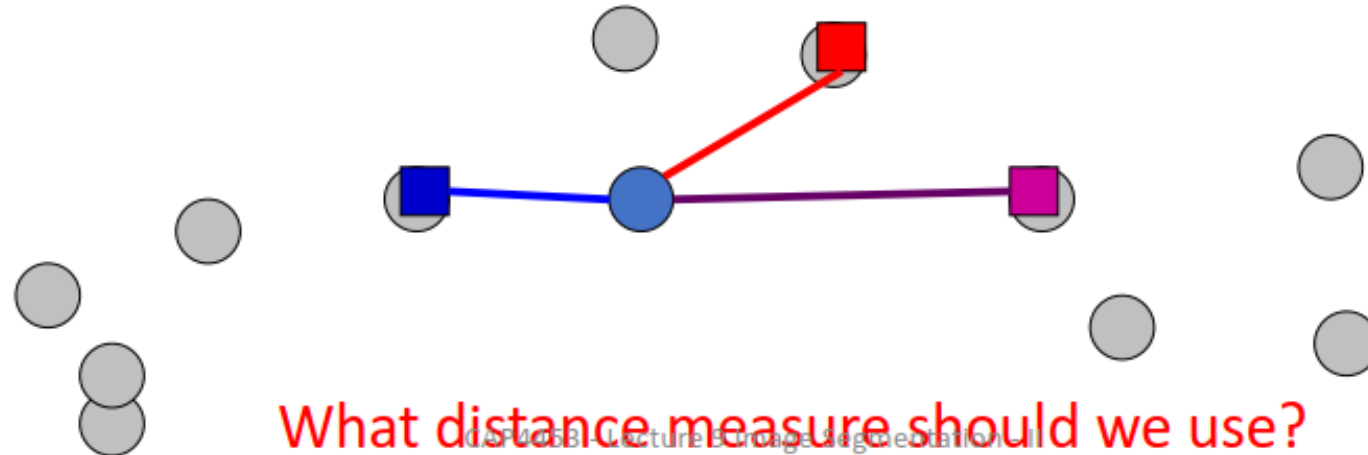
Iterate:

- **Assign/cluster each example to closest center**

iterate over each point:

- get **distance** to each cluster center
- assign to closest center (hard cluster)

- Recalculate centers as the mean of the points in a cluster





Distance measures

$$d = \sqrt{\sum_i (x_i^2 - y_i^2)}$$

- p-norm
- Euclidean norm
- L1-norm

$$\|x\|_p = \left(\sum_i |a_i|^p \right)^{\frac{1}{p}} \quad p \geq 1$$

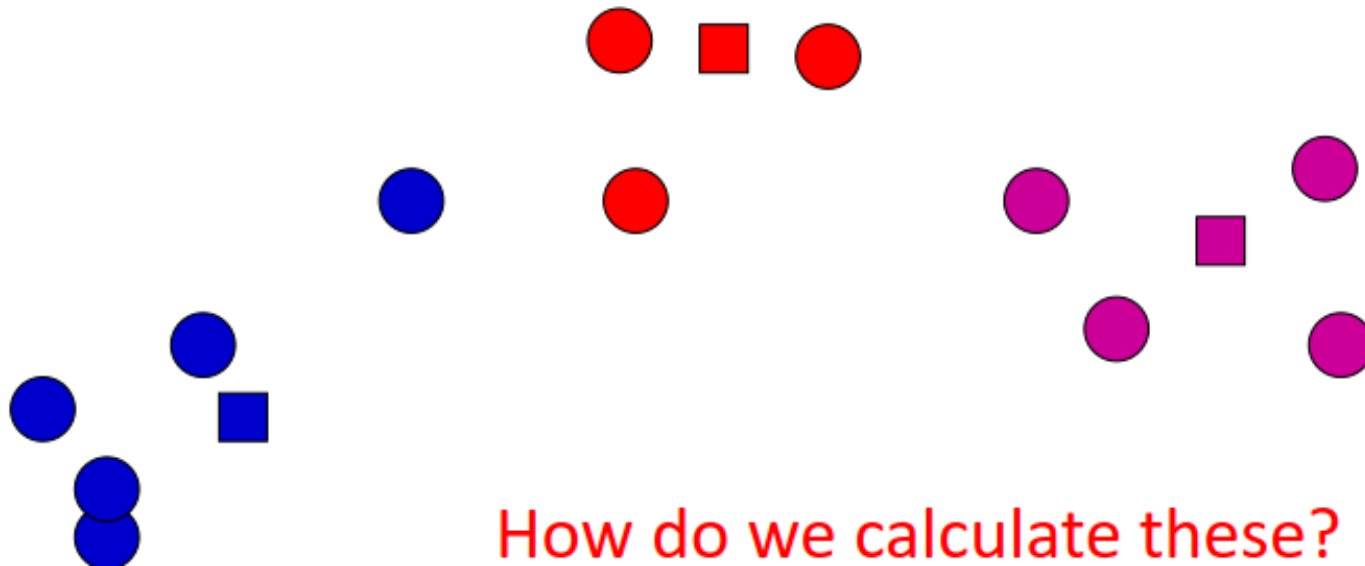
$$\|x\|_2 = \left(\sum_i |a_i|^2 \right)^{1/2}$$

$$\|x\|_1 = \left(\sum_i |a_i| \right)$$

K-means

Iterate:

- Assign/cluster each example to closest center
- Recalculate centers as the mean of the points in a cluster



K-means

Iterate:

- Assign/cluster each example to closest center
- Recalculate centers as the mean of the points in a cluster

$$\mu_j = \frac{\sum_{i:y_i=j} x_i}{\sum_{i:y_i=j} 1}$$



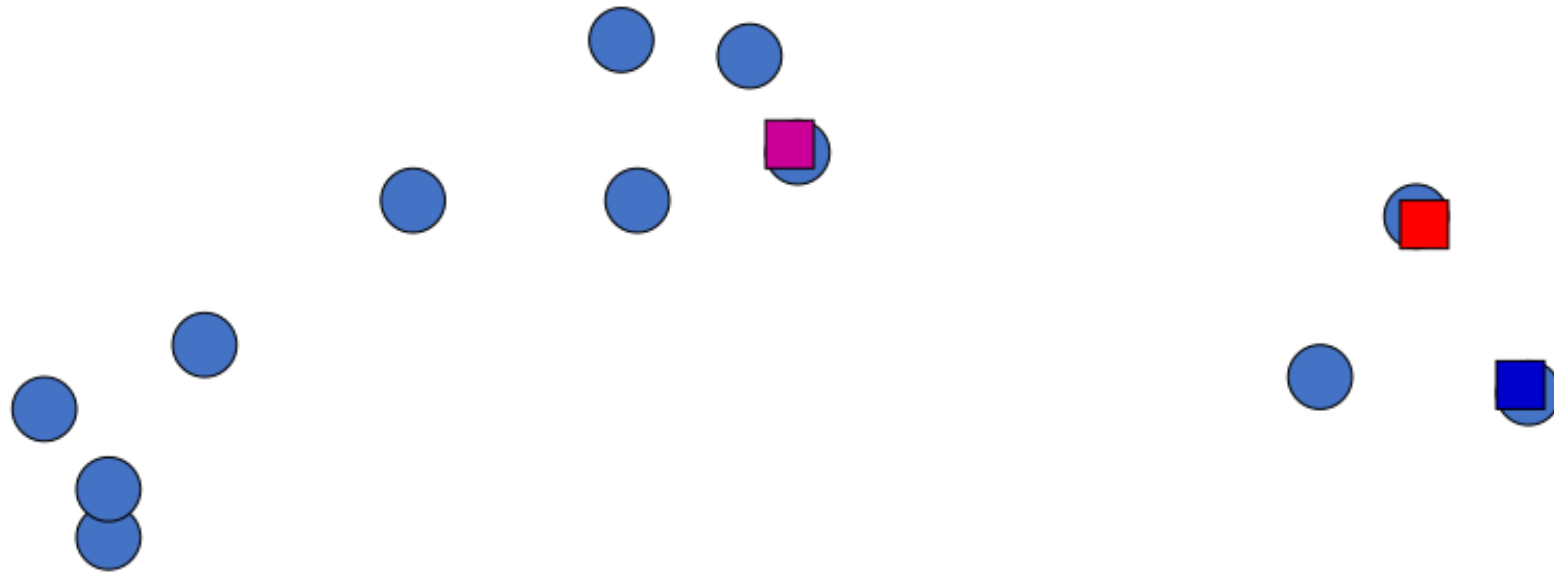
K-means loss function

K-means tries to minimize what is called the “k-means” loss function:

$$loss = \sum_{i=1}^n d^2(x_i, \mu_k), \quad \text{where } \mu_k \text{ is the cluster center for } x_i$$

that is, the sum of the squared distances from each point to the associated cluster center

K-means: initialization



What would happen here?

Seed selection ideas?

CAF4455



Seed choice

Results can vary drastically based on random seed selection

Some seeds can result in poor convergence rate, or convergence to sub-optimal clusterings

Common heuristics

- Random centers in the space
- Randomly pick examples
- Points least similar to any existing center (furthest centers heuristic)
- **Try out multiple starting points**
- Initialize with the results of another clustering method

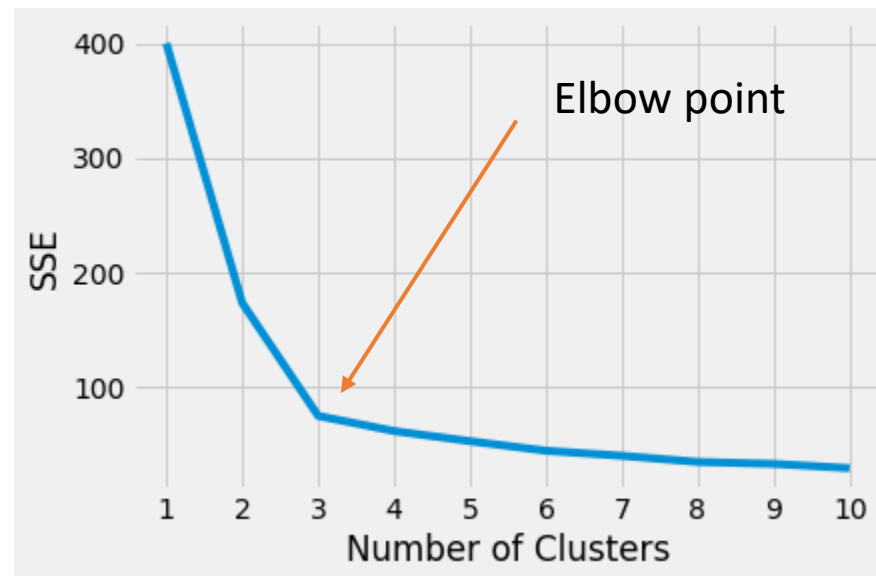


Choosing the Appropriate Number of Clusters

1. The **elbow method**
2. The **silhouette coefficient**

Choosing the Appropriate Number of Clusters

- run several k-means,
- increment k with each iteration
- record the sum of the squared error (SSE)
 - The SSE is defined as the sum of the squared Euclidean distances of each point to its closest centroid

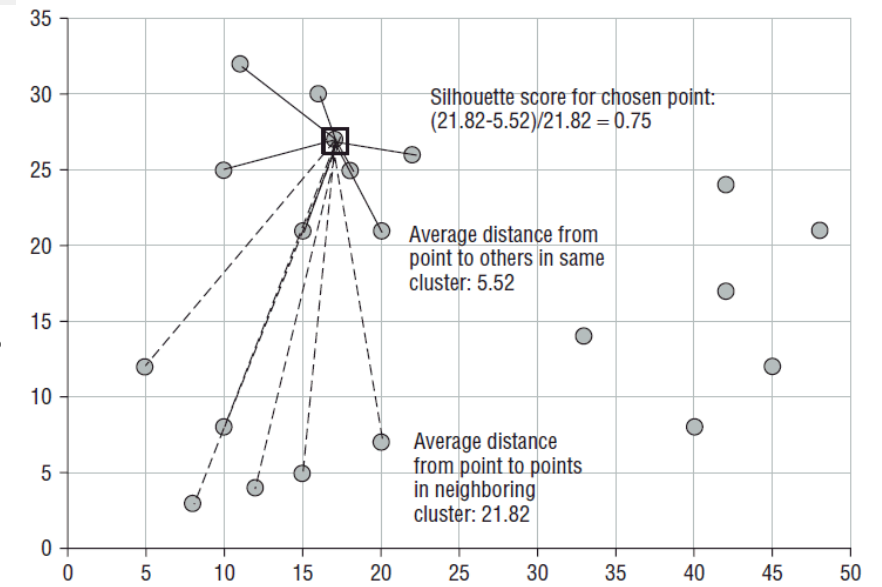


1. The **elbow method**

Choosing the Appropriate Number of Clusters

- run several k-means,
- increment k with each iteration
- Pick max [silhouette coefficient](#)
 1. How close the data point is to other points in the cluster
 2. How far away the data point is from points in other clusters

- $(b - a) / \max(a, b)$. Where,
 - a: intra-cluster distance
 - b: distance between a sample and the nearest cluster that the sample is not a part of.



2. The silhouette coefficient



Segmenting an image with K-means

- Example:
 - Vector: (coordinates i , coordinate j , Color L , Color a , Color b) : 5 dims
 - Distance: Euclidean distance
 - Number of clusters: 10
 - Seeds selected randomly



Kmeans function from scratch

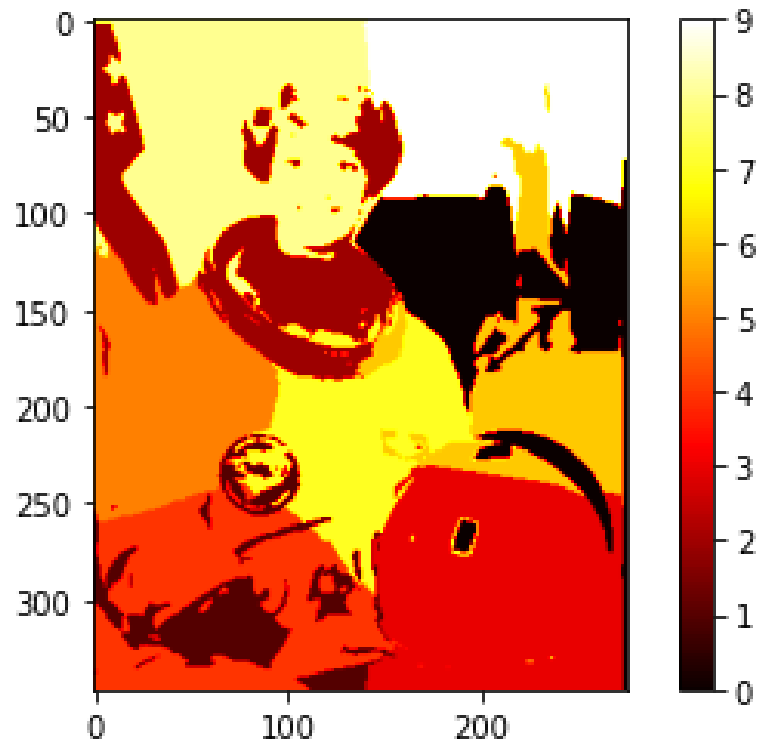
```
11 import numpy as np
12 from scipy.spatial.distance import cdist
13 import matplotlib.pyplot as plt
14 import cv2
15
16 #Defining our function
17 def kmeans(x,k, no_of_iterations):
18     idx = np.random.choice(len(x), k, replace=False)
19     #Randomly choosing Centroids
20     centroids = x[idx, :] #Step 1
21
22     #finding the distance between centroids and all the data points
23     distances = cdist(x, centroids , 'euclidean') #Step 2
24
25     #Centroid with the minimum Distance
26     points = np.array([np.argmin(i) for i in distances]) #Step 3
27
28     #Repeating the above steps for a defined number of iterations
29     #Step 4
30     for _ in range(no_of_iterations):
31         centroids = []
32         for idx in range(k):
33             #Updating Centroids by taking mean of Cluster it belongs to
34             temp_cent = x[points==idx].mean(axis=0)
35             centroids.append(temp_cent)
36
37         centroids = np.vstack(centroids) #Updated Centroids
38
39         distances = cdist(x, centroids , 'euclidean')
40         points = np.array([np.argmin(i) for i in distances])
41
42     return points
43
```



Calling Kmeans

```
45 ●  imFile = 'C:\\Users\\gonza\\OneDrive\\Teaching\\CAP4453\\class12\\img.png'
46
47  im = cv2.imread(imFile)
48  LabImg = cv2.cvtColor(im,cv2.COLOR_BGR2LAB)
49
50
51  i = np.linspace(0, LabImg.shape[0]-1, LabImg.shape[0]).astype(int)
52  j = np.linspace(0, LabImg.shape[1]-1, LabImg.shape[1]).astype(int)
53  xv, yv = np.meshgrid(i, j)
54
55  numpoints = xv.ravel().shape[0]
56
57  L = LabImg[xv.ravel(),yv.ravel(),0].reshape((numpoints,1))
58  a = LabImg[xv.ravel(),yv.ravel(),1].reshape((numpoints,1))
59  b = LabImg[xv.ravel(),yv.ravel(),2].reshape((numpoints,1))
60
61
62  #X=np.concatenate((xv.ravel().reshape((numpoints,1)), yv.ravel().reshape((numpoints,1)),L,a,b))
63  X=np.concatenate((xv.ravel().reshape((numpoints,1)), yv.ravel().reshape((numpoints,1)),L,a,b), axis=1)
64
65
66  points = kmeans(X,10,50)
67
68  newImg = np.zeros((im.shape[0],im.shape[1]))
69  newImg[xv.ravel(),yv.ravel()]=points
70  im1=plt.imshow(newImg, cmap='jet'); plt.colorbar(im1, cmap='jet'); plt.show()
71
```

Results



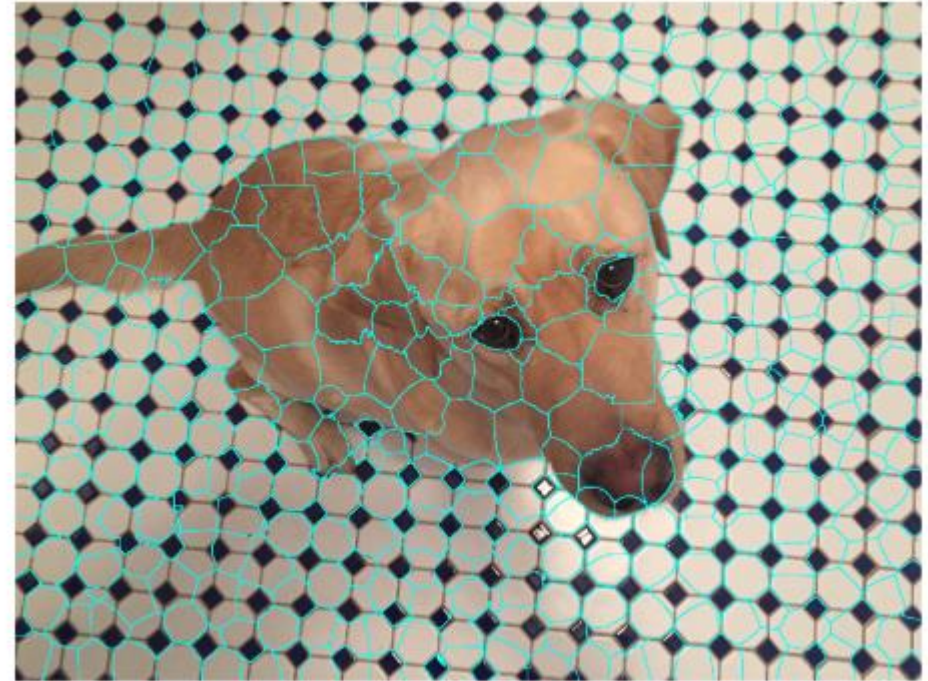


Outline

- ~~Image segmentation basics~~
- ~~Thresholding based~~
 - ~~Binarization~~
 - ~~Otsu~~
- ~~Region based~~
 - ~~Merging~~
 - ~~Splitting~~
- **Clustering based**
 - K-means
 - **Superpixels (SLIC)**

Superpixels

- They carry more information than pixels.
- Superpixels have a perceptual meaning since pixels belonging to a given superpixel share similar visual properties.
- They provide a convenient and compact representation of images that can be very useful for computationally demanding problems.



[Superpixels and SLIC. What is a Superpixel? | by Darshita Jain | Medium](#)



SLIC (Anchanta et. al. TPAMI 2012)

Input:

- a desired number of approximately equally-sized superpixels K

N	Number of pixels in the input image
K	Number of Superpixels used to segment the input image
N/K	Approximate size of each superpixel
$S = \sqrt{N/K}$	For roughly equally sized superpixels there would be a superpixel centre at every grid interval S

Features:

- five-dimensional [labxy] space,
- [lab] is the pixel color vector in [CIELAB](#) color space
 - xy is the pixel position.

Distances:

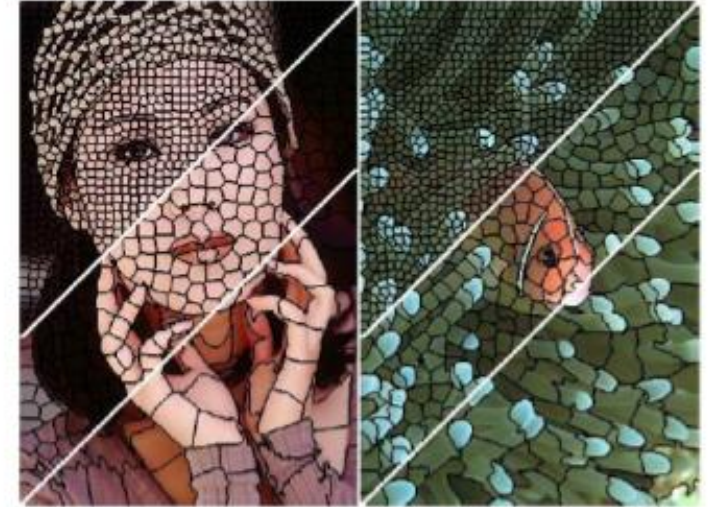
$$d_{lab} = \sqrt{(l_k - l_i)^2 + (a_k - a_i)^2 + (b_k - b_i)^2}$$

$$d_{xy} = \sqrt{(x_k - x_i)^2 + (y_k - y_i)^2}$$

$$D_s = d_{lab} + \frac{m}{S} d_{xy} ,$$

SLIC (Anchanta et. al. TPAMI 2012)

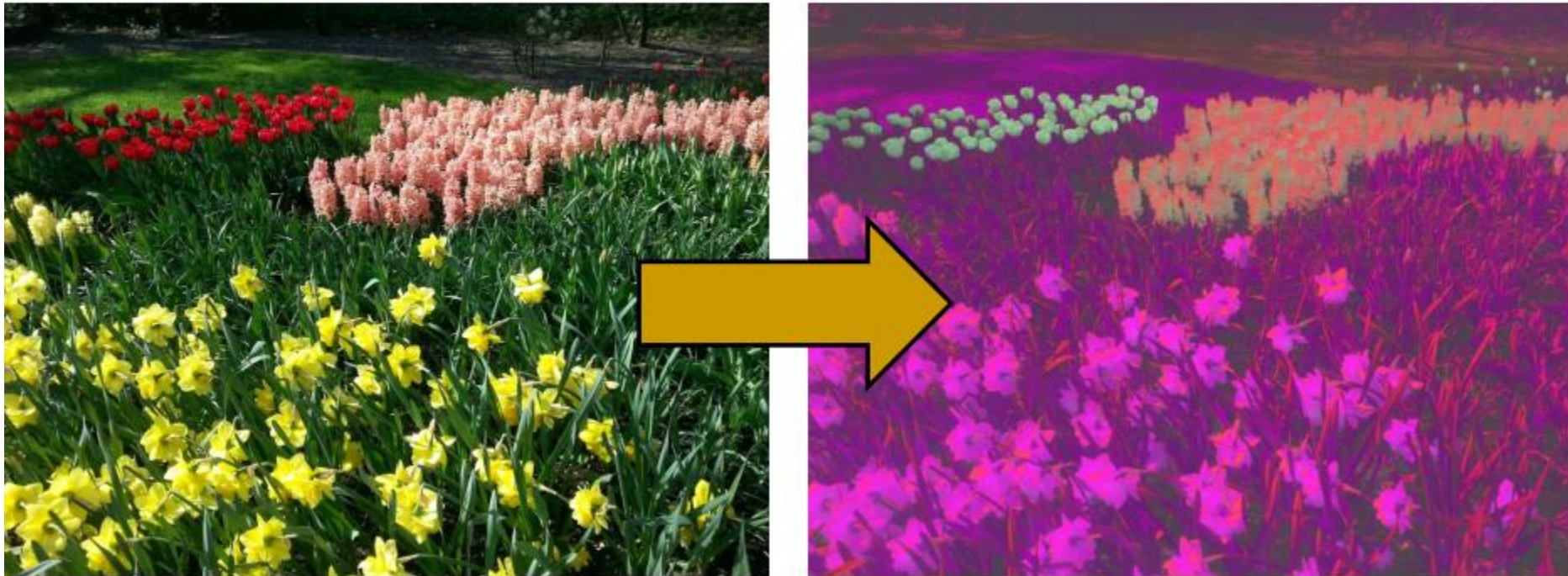
1. Get Features: Lab color, x-y position
2. Initialize cluster centers on pixel grid in steps S
3. Move centers to position in 3×3 window with smallest gradient
4. Compare each pixel to cluster center within $2S$ pixel distance and assign to nearest
5. Recompute cluster centers as mean color/position of pixels belonging to each cluster
6. Stop when residual error is small



- + Fast 0.36s for 320x240
- + Regular superpixels
- + Superpixels fit boundaries
- May miss thin objects
- Large number of superpixels

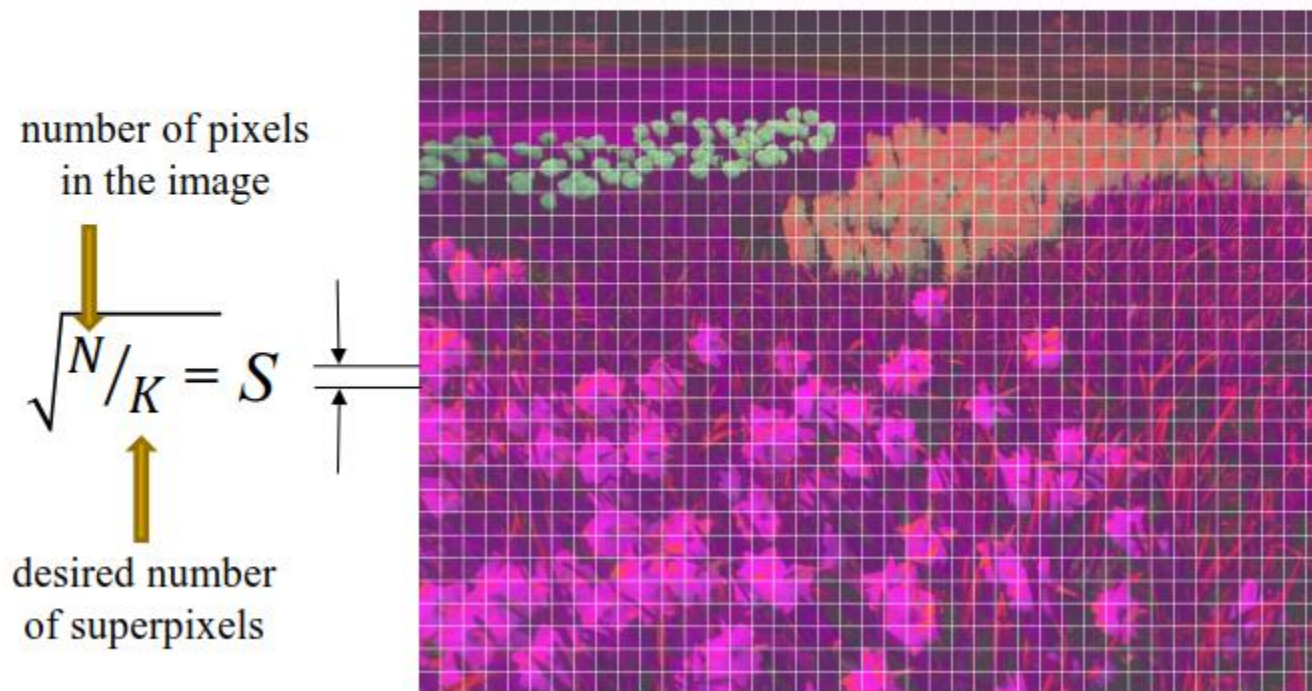
SLIC Example

1. Convert the RGB image to CIELAB color space.



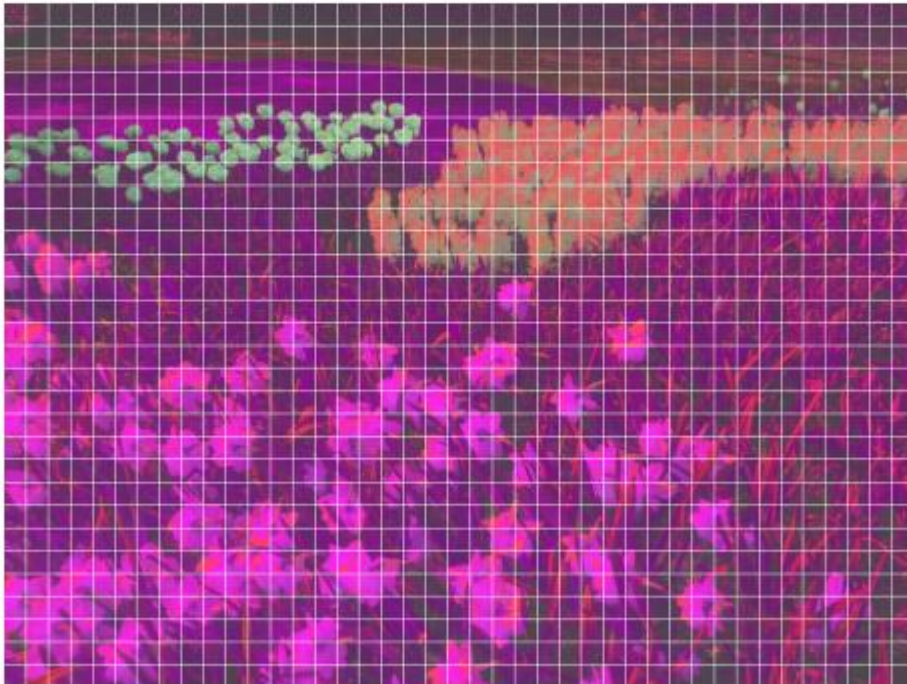
SLIC Example

2. Initialize cluster centers $C_k = [l_k; a_k; b_k; x_k; y_k]^T$ by sampling pixels at regular grid steps S .



SLIC Example

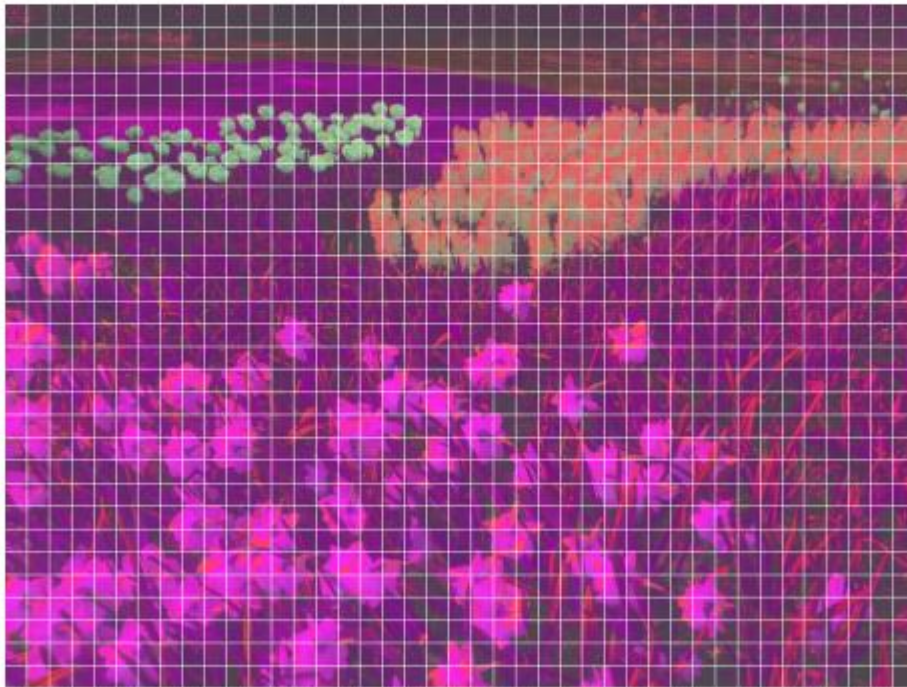
3. Move cluster centers to the lowest gradient position in a 3×3 neighborhood.



This is done to avoid placing them at an edge and to reduce the chances of choosing a noisy pixel

SLIC Example

3. Move cluster centers to the lowest gradient position in a 3×3 neighborhood.



$$G(x,y) = \|I(x+1,y)-I(x-1,y)\|^2 + \|I(x,y+1)-I(x,y-1)\|^2$$

- $I(x,y)$ is the lab vector corresponding to the pixel at position (x,y) ,
- $\|\cdot\|$ is the L2 norm.

SLIC Example

4. A 2D label matrix L as large as the input image will contain the superpixel each pixel belongs to. L is initialized with -1 for all pixels.
(meaning that each pixel belongs to no superpixel in the beginning)

$L =$

-1	-1	-1	-1	-1	-1
-1	-1	-1	-1	-1	
-1	-1	-1			
-1	-1				

SLIC Example

- A 2D distance matrix d as large as the input image will contain the distance of each pixel to the centroid of its superpixel. d is initialized with ∞ for all pixels.

(distance to superpixel centroid in the beginning)

$d =$

∞	∞	∞	∞	∞	∞
∞	∞	∞	∞	∞	
∞	∞	∞			
∞					

$$D = \sqrt{(d_c)^2 + \left(\frac{d_s}{S}\right)^2 m^2}$$

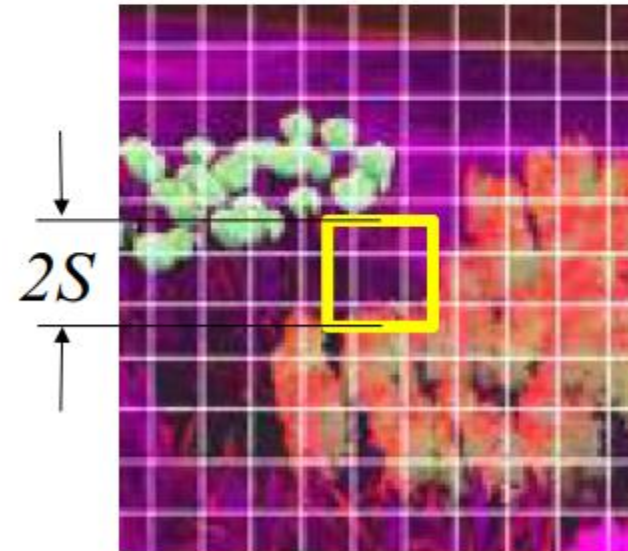
controls the relative importance of shape and color

m large \rightarrow favors more compact (lower area to perimeter ratio) superpixels.

m small \rightarrow favors more adherence to edges.

SLIC Example

```
6. repeat
    for each cluster center  $C_k$  do
        for each pixel  $i$  in a  $2S \times 2S$  region around  $C_k$  do
            Compute the distance  $D$  between  $C_k$  and  $i$ .
            if  $D < d(i)$  then
                set  $d(i) = D$ 
                set  $L(i) = k$ 
            end if
        end for
    end for
    compute new cluster centers.
    compute residual error  $E$ .
until  $E < \text{threshold}$ .
```



Example 1: image size = 735×980 pixels

$K = 1333$ superpixels; $m = 40$



More examples





Questions?