

Lecture 14: Beyond K Means
Clustering

### Recap

- Kmeans algorithm
- Expectation Maximization
- Centroid initialization with kmeans++
- Kmeans decision boundary
- Kmeans limitations with oblong data clusters
- Kmeans limitations with outliers
- Using Kmeans and silhouette analysis for outlier detection



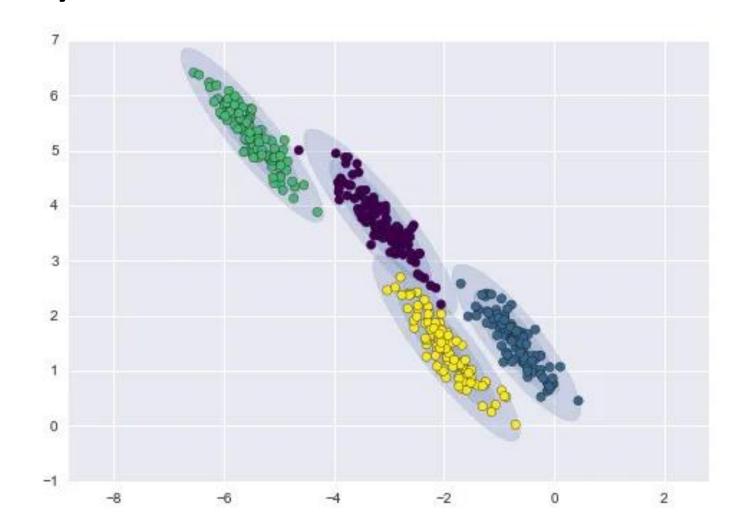
# Mixture Models

### Another way of looking at clusters

- Hard versus soft clustering
- Hard:
  - Clusters don't overlap
  - Elements belong to a cluster or they don't
  - E.g. K-Means
- Soft
  - Clusters may overlap
  - Strength of association between cluster and instances
  - •60% confidence of cluster1 membership, 40% cluster2
  - E.g. Gaussian Mixture Model Clustering

# Mixture Model clustering

- Probabilistic way of doing soft clustering
  - •Soft clustering Cluster membership is not 0/1
  - Probability
- Each cluster is a generative model (probability distribution)
- But parameters (mean and covariance) are unknown
  - Discovered as part of clustering
- Parameters are discovered by EM
  - •EM in GMM is EM in true sense ©

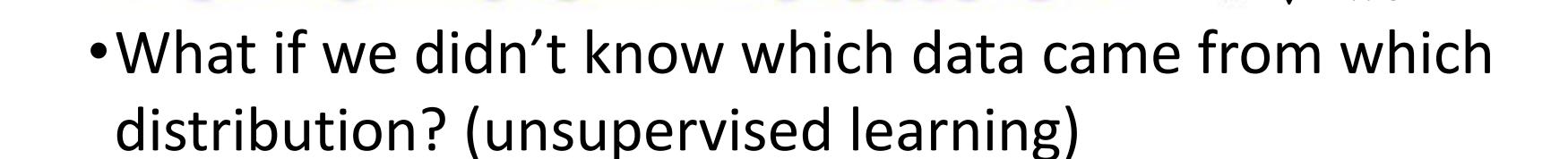


### Recap: Estimate Mean & variance in generative ML

- We are given data x1 .. Xn
  - Mean and variance not known
  - Easy to find from labeled data

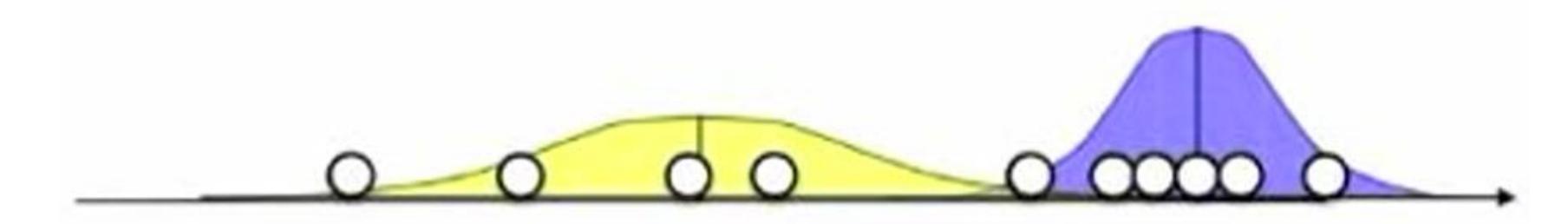
$$\mu_{blue} = \frac{\sigma_{blue}}{n_b}$$
 $\sigma_{blue}^2 = \frac{\sum (x_i - \mu_{blue})^2}{n_{blue}}$ 

 $x_1 + x_2 + \dots + x_{n_b}$ 



### Gaussian Mixture Models

- Chicken & Egg
  - We need labels to determine mean, variance
    - $\circ$   $\circ$   $\circ$   $\circ$   $\circ$
  - We need mean, variance to predict labels



•If we knew mean/var, we could guess which point came from which Gaussian. But we don't know

### Gaussian Mixture Models Intuition

- We are given data points
- They came from k Gaussian distributions
- We don't know which point came from which Gaussian

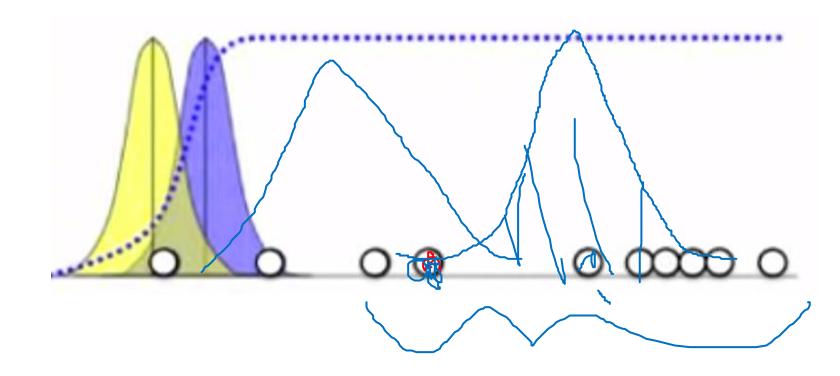


- Solution: EM
- Start with two Gaussian with random mean and variances
  - Just like kmeans centroids



### Gaussian Mixture Models EM

- 2 Gaussian with random mean and variances
- For each point xi, does this look like it came from a or b?



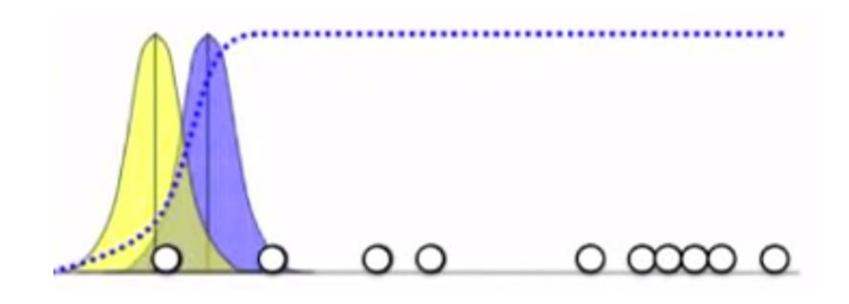
- Soft assignment
- No max P calc

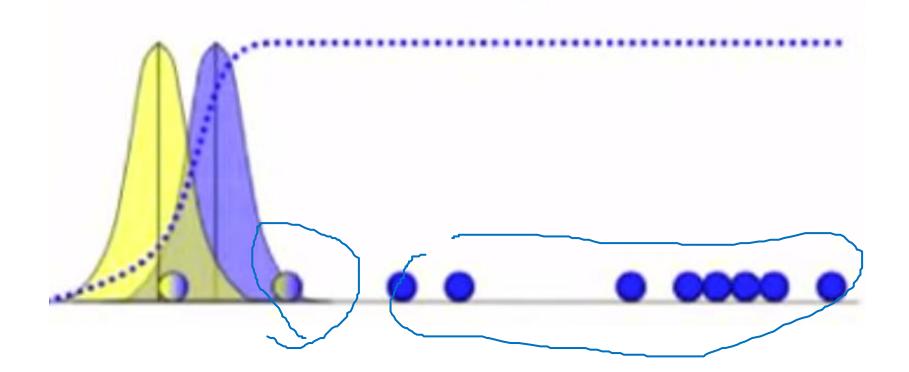
$$P(x^{(i)}|b) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2} \left(\frac{x^{(i)} - \mu_b}{\sigma_b}\right)^2}$$

$$P(b|x^{(i)}) = \frac{P(x^{(i)}|b)P(b)}{P(x^{(i)}|b)P(b) + P(x^{(i)}|a)P(a)}$$

$$P(a|x^{(i)}) = \frac{P(x^{(i)}|a)P(a)}{P(x^{(i)}|b)P(b) + P(x^{(i)}|a)P(a)}$$

### Gaussian Mixture Models EM





Rinse and Repeat

$$p_{b_i} = P(b|x^{(i)}) = \frac{P(x^{(i)}|b)P(b)}{P(x^{(i)})}$$

$$p_{a_i} = P(a|x^{(i)}) = \frac{P(x^{(i)}|a)P(a)}{P(x^{(i)})}$$

Calculate the new mean/var

$$\mu_b = \frac{p_{b_1}x^{(1)} + p_{b_2}x^{(2)} + ...p_{b_n}x^{(n)}}{p_{b_1} + p_{b_2} + ... + p_{b_n}}$$

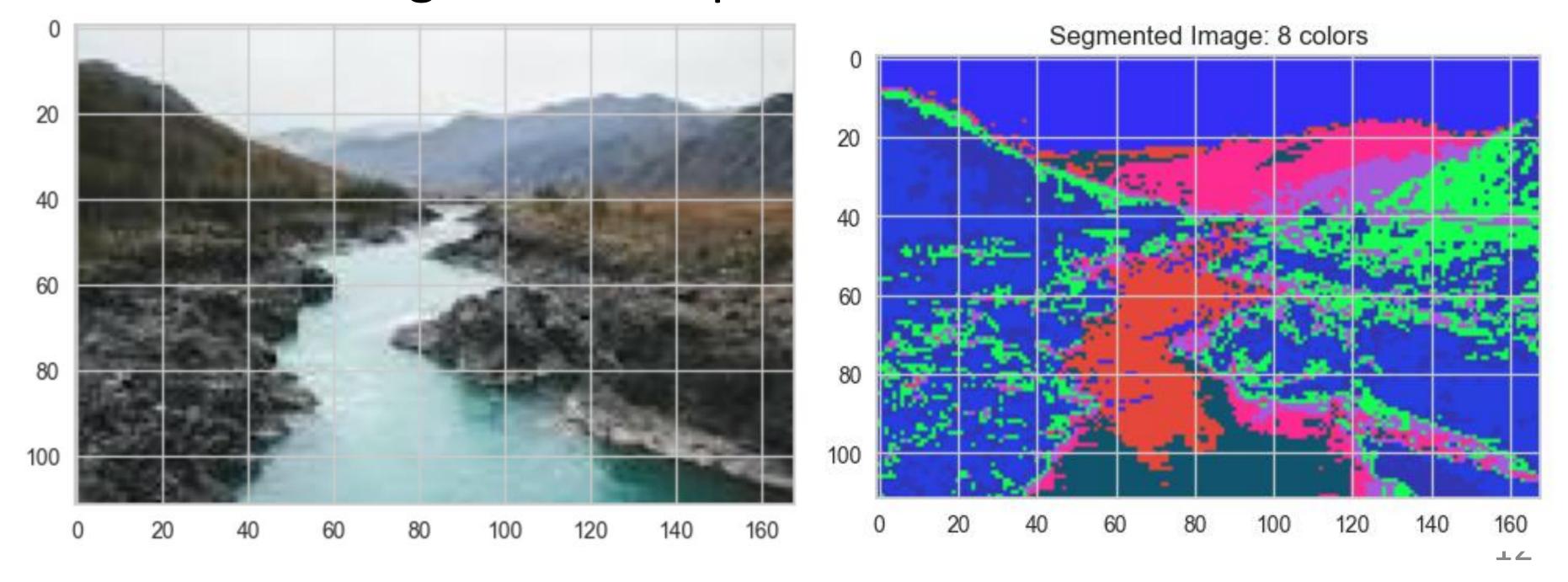
$$\mu_a = \frac{p_{a_1}x^{(1)} + p_{a_2}x^{(2)} + ...p_{a_n}x^{(n)}}{p_{a_1} + p_{a_2} + ... + p_{a_n}}$$

### GMM clustering applications

- Real life scenarios are mixtures, never black and white
- Finance
  - Investment Portfolio construction
  - Identifying stocks for growth (outliers in a good way)

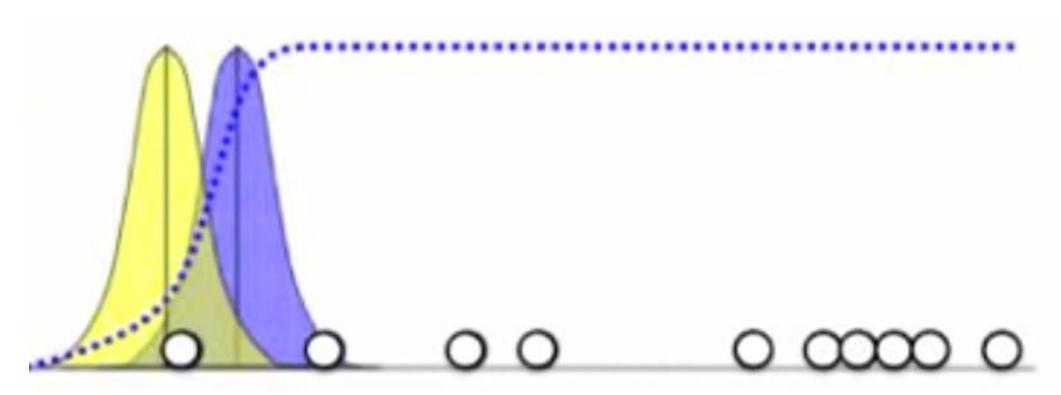
# GMM clustering applications

- Image segmentation (less expensive way)
  - Split into patches & cluster based on image characteristics
  - Medical image locate specific structures



### GMM clustering applications

- One tool of many towards explainable solution
  - Population to 2 wheeler, 4 wheeler ratio
  - Accident stats, Poisson mixtures for pattern identification



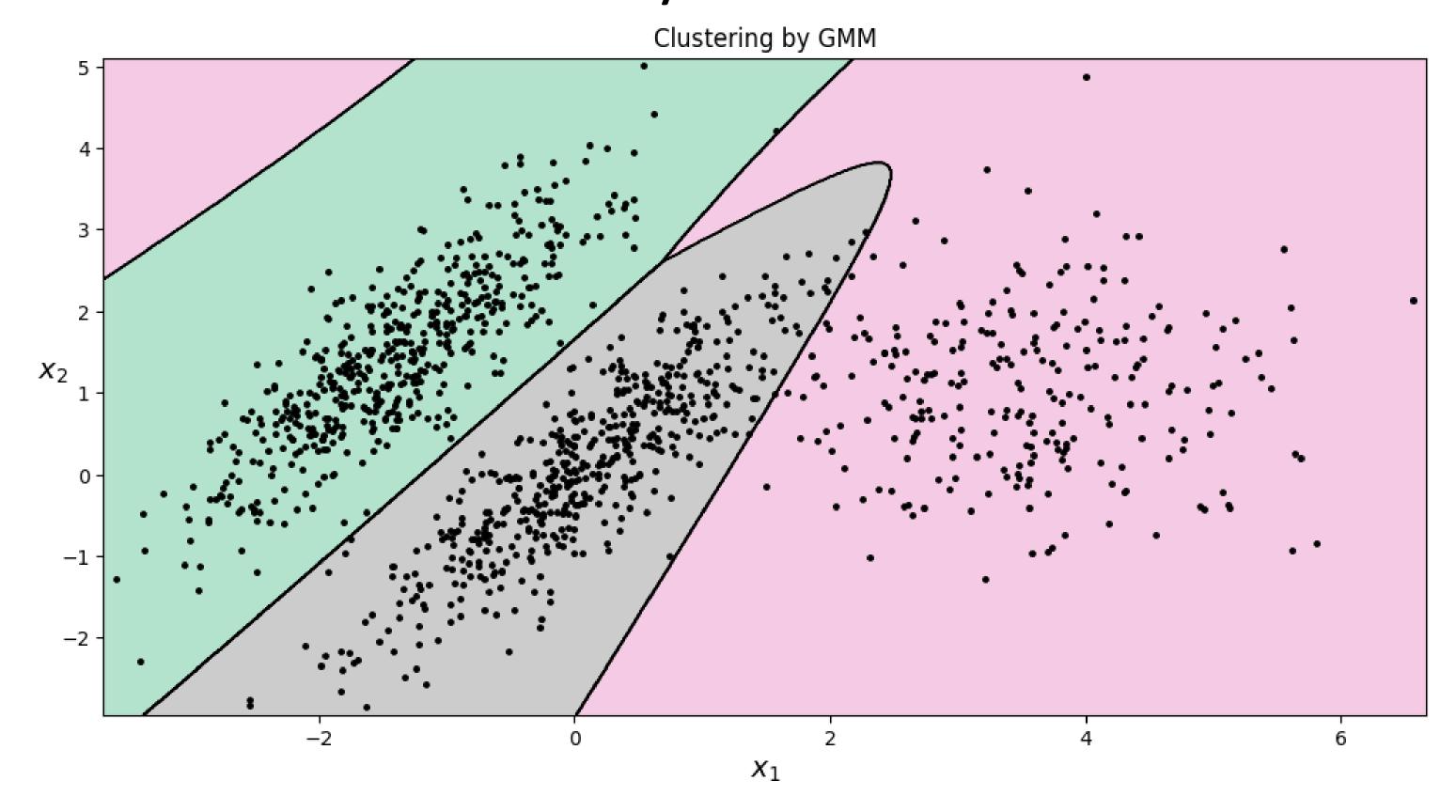
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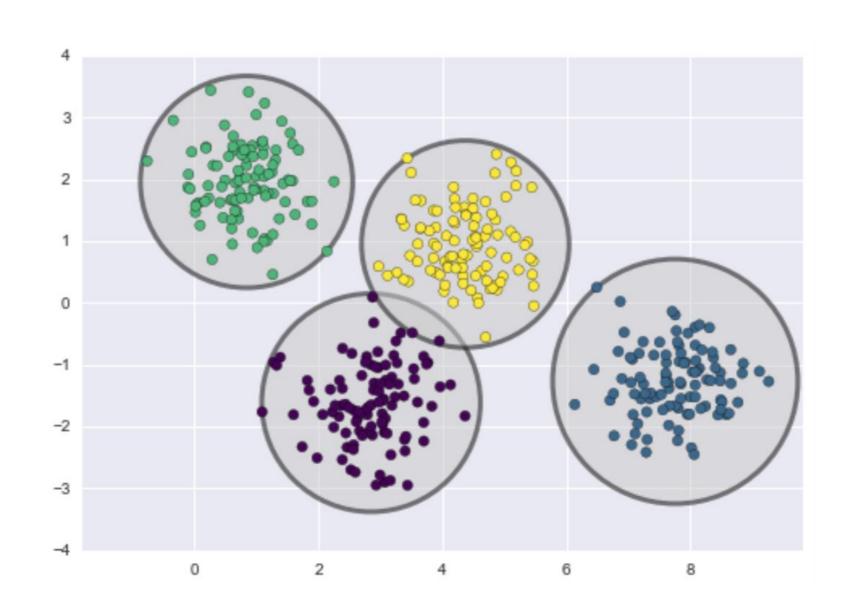
# GMM decision boundary

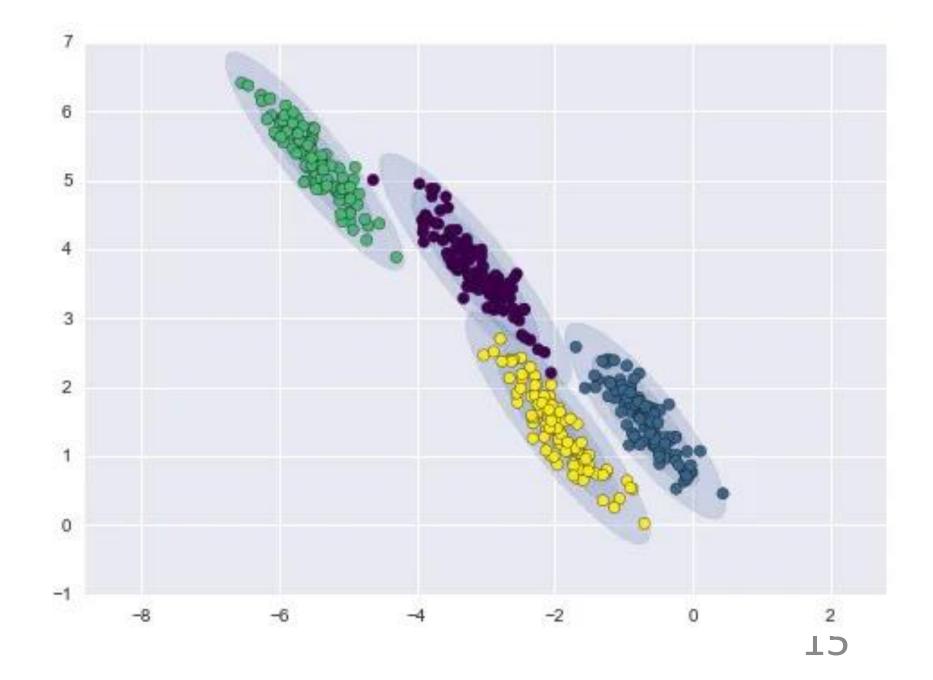
Quadratic decision boundary



### GMM model

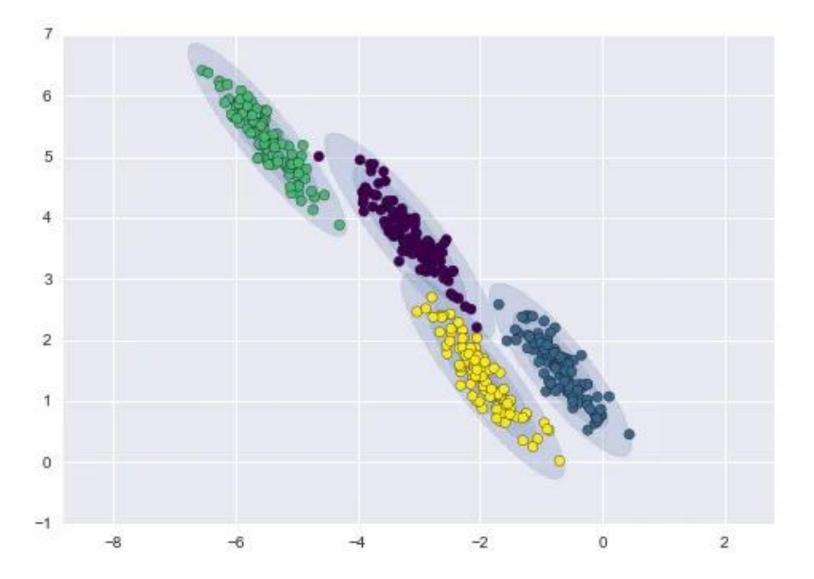
- Clustering every time some new data comes is expensive
- Could use extracted mean, variance, covariance as model
  - Multivariate Gaussian model from scratch!!





## Handling outliers with GMM

- •GMM is sensitive to outliers
- Can use multivariate Gaussian techniques (Mahalanobis distance) or silhouette analysis
- Can use Minimum Covariance Determinant also



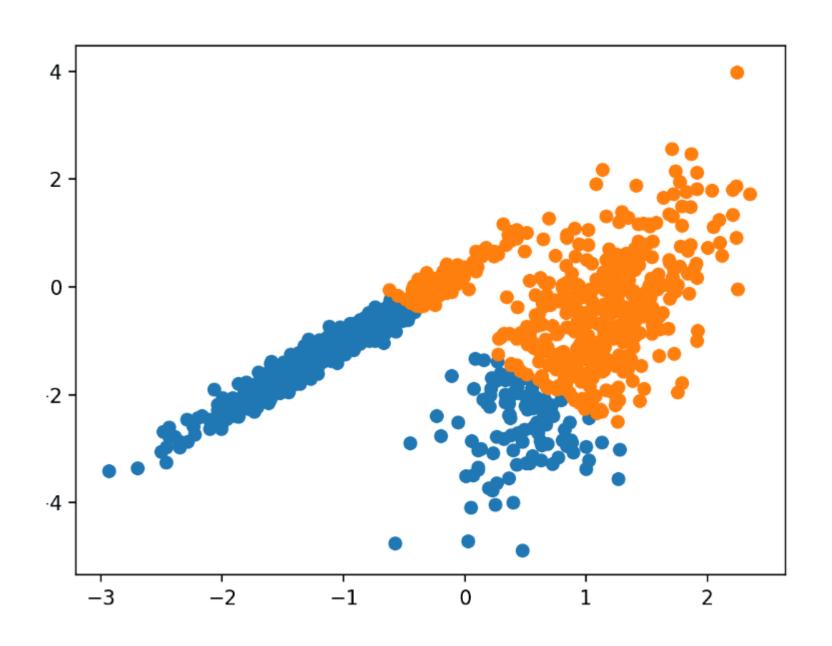


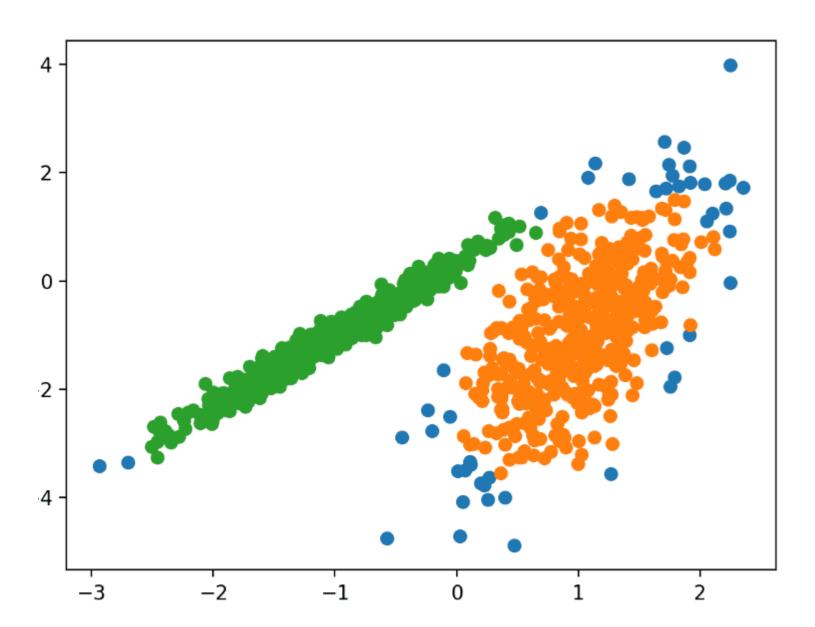
Beyond K-Means & GMM

## Clustering for mixed data types

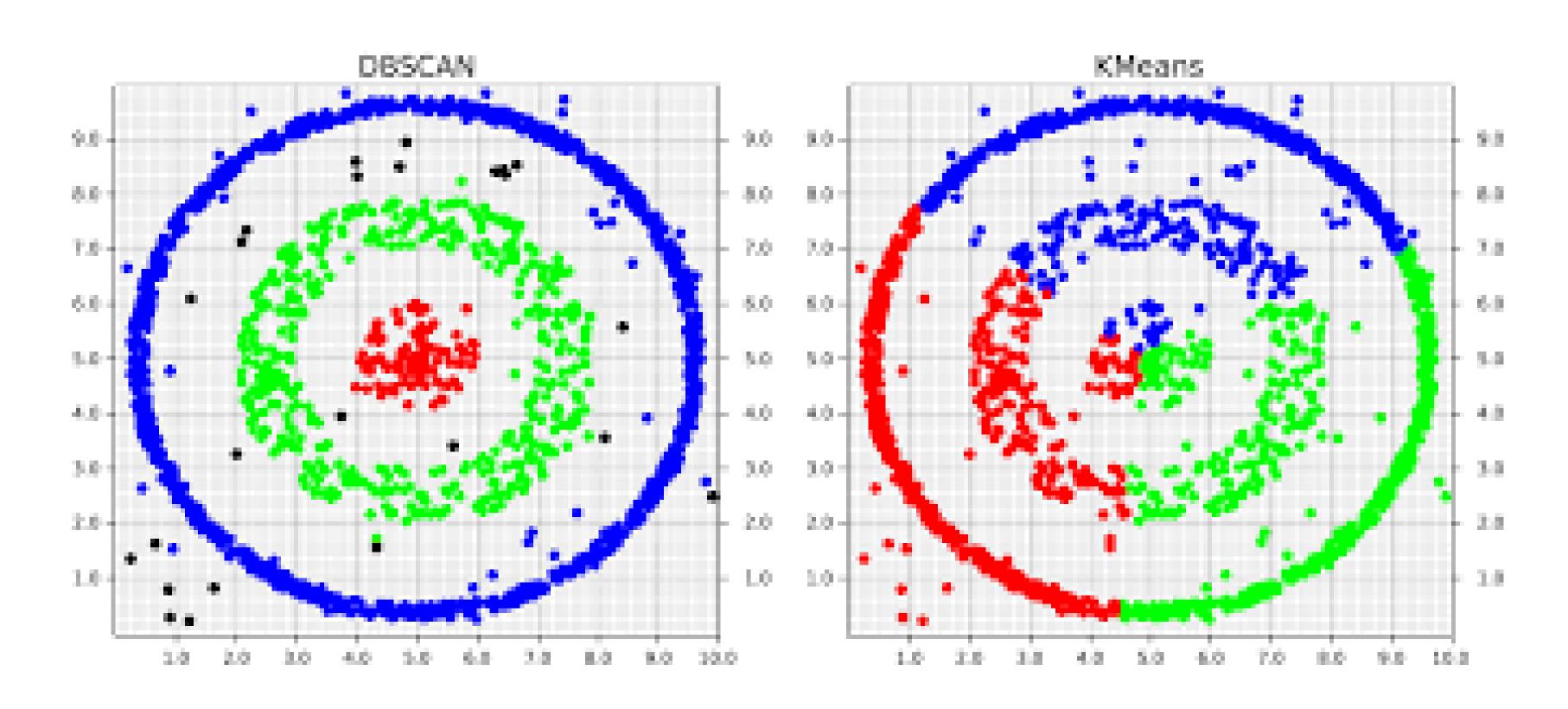
- K-Mode
- K-Prototypes
- K Medoids
  - Gowers distance
    - Manhattan for numeric, Dice for categorical

# K-Means cannot handle asymmetric globular data

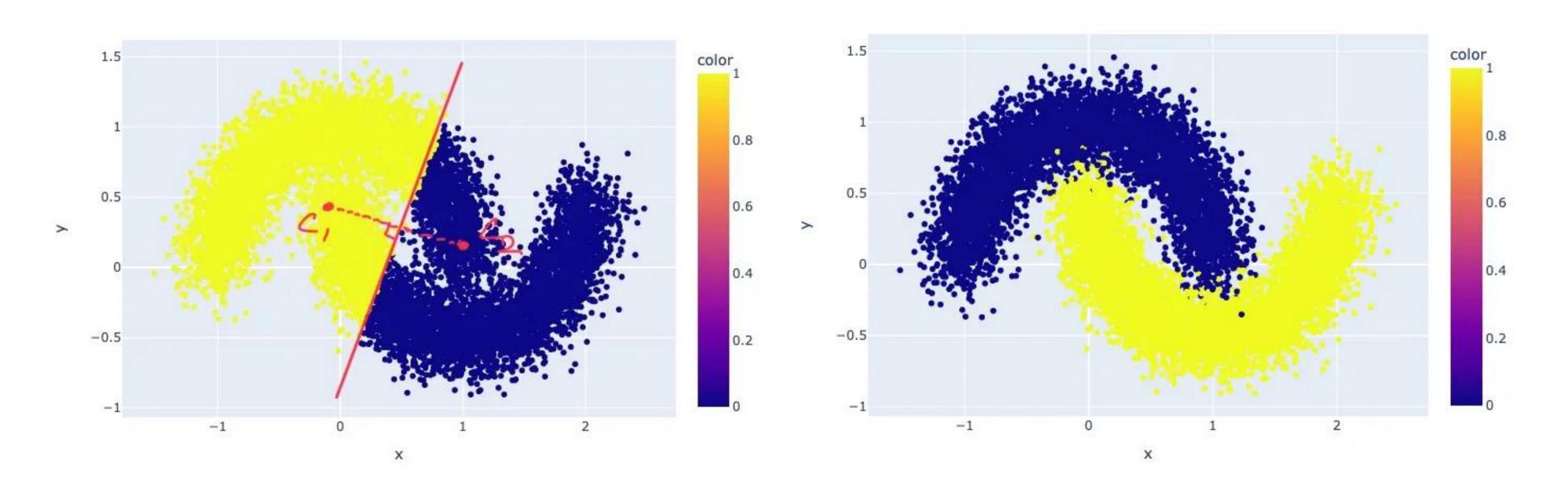




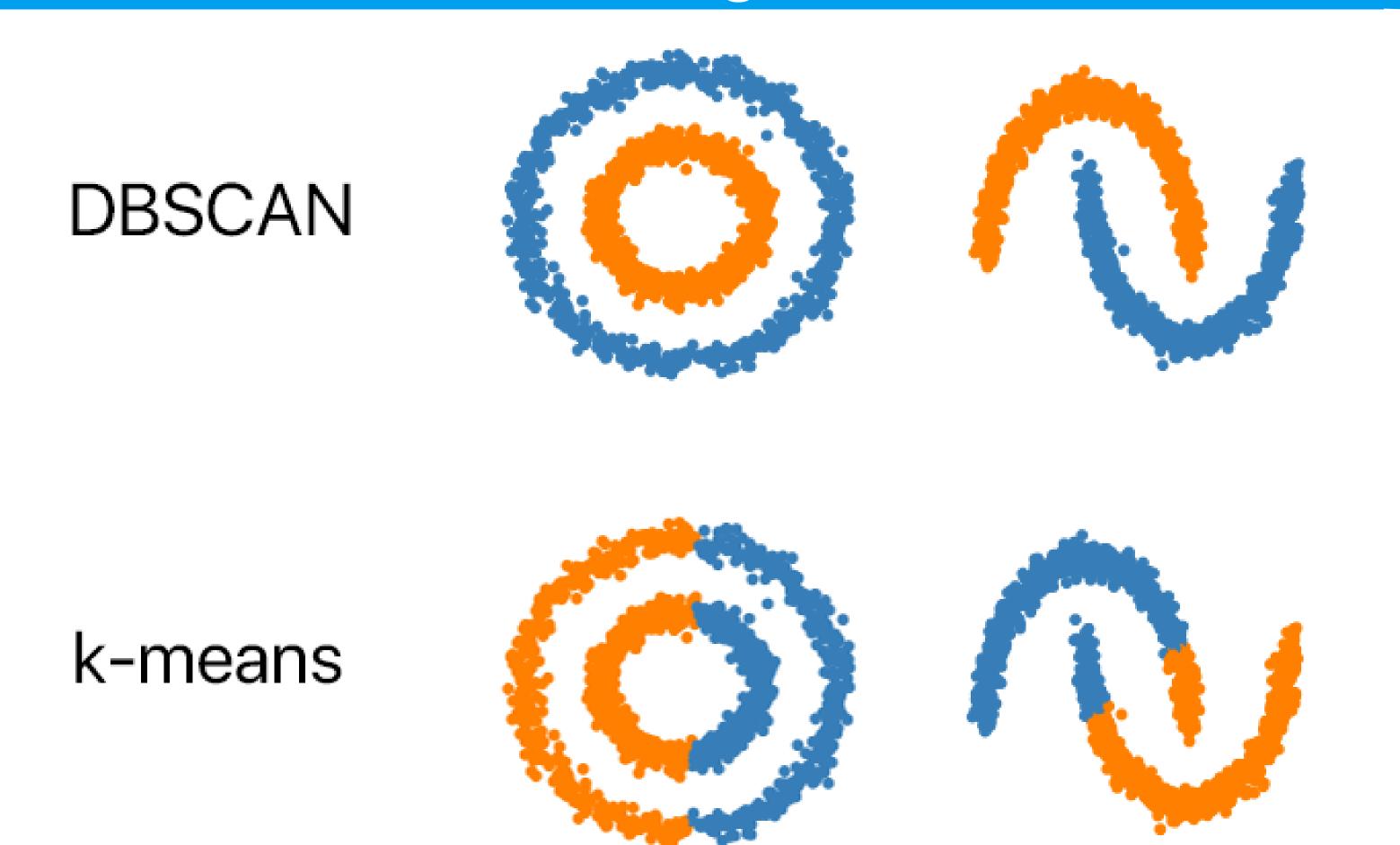
# K-Means cannot handle non globular data



# K-Means cannot handle non globular data



# K-Means cannot handle non globular data



## Different types of clustering (optional)

- DBSCAN
  - Some patterns of non globular data, diff metrics
  - NLP clustering, Cosine distance (1-cos theta)
  - DBSCAN Clearly explained Josh Stammer
    - https://www.youtube.com/watch?v=RDZUdRSDOok
  - Evaluation metric is also different (DBCV etc.)
- BIRCH clustering for large datasets
- Mean shift clustering
  - Finds use in unsupervised image segmentation

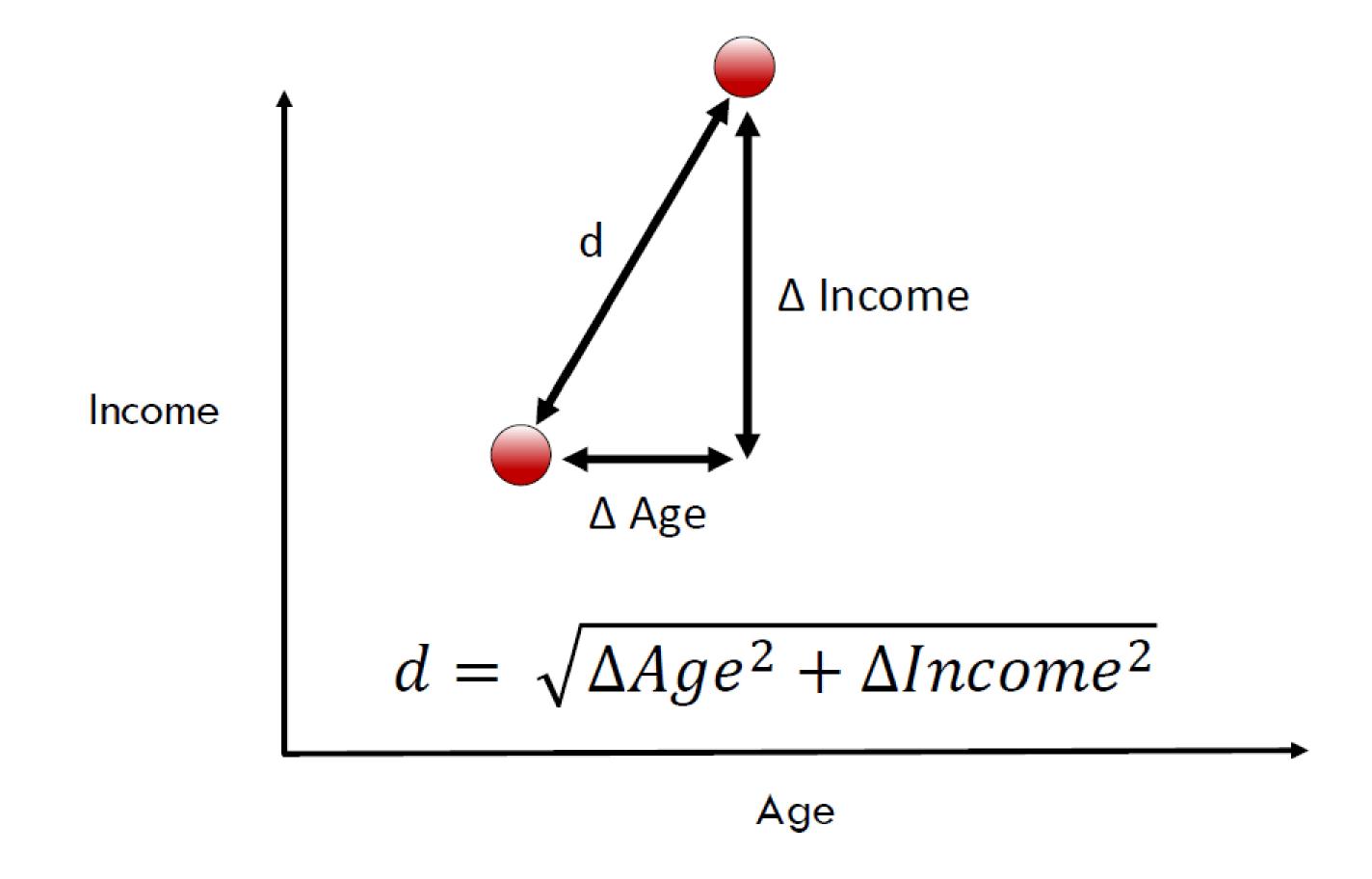
### Case for Distance metrics

- Choice of distance metric is crucial for clustering success
- Also success in any distance based ML method
  - E.g. Nearest Centroid, KNN
- Each metric has strengths and appropriate use cases
  - For e.g. NLP and cosine distance
  - Linear Regression without outliers and Euclidean
- Sometimes metric is chosen by empirical eval (hyperparameter tuning)

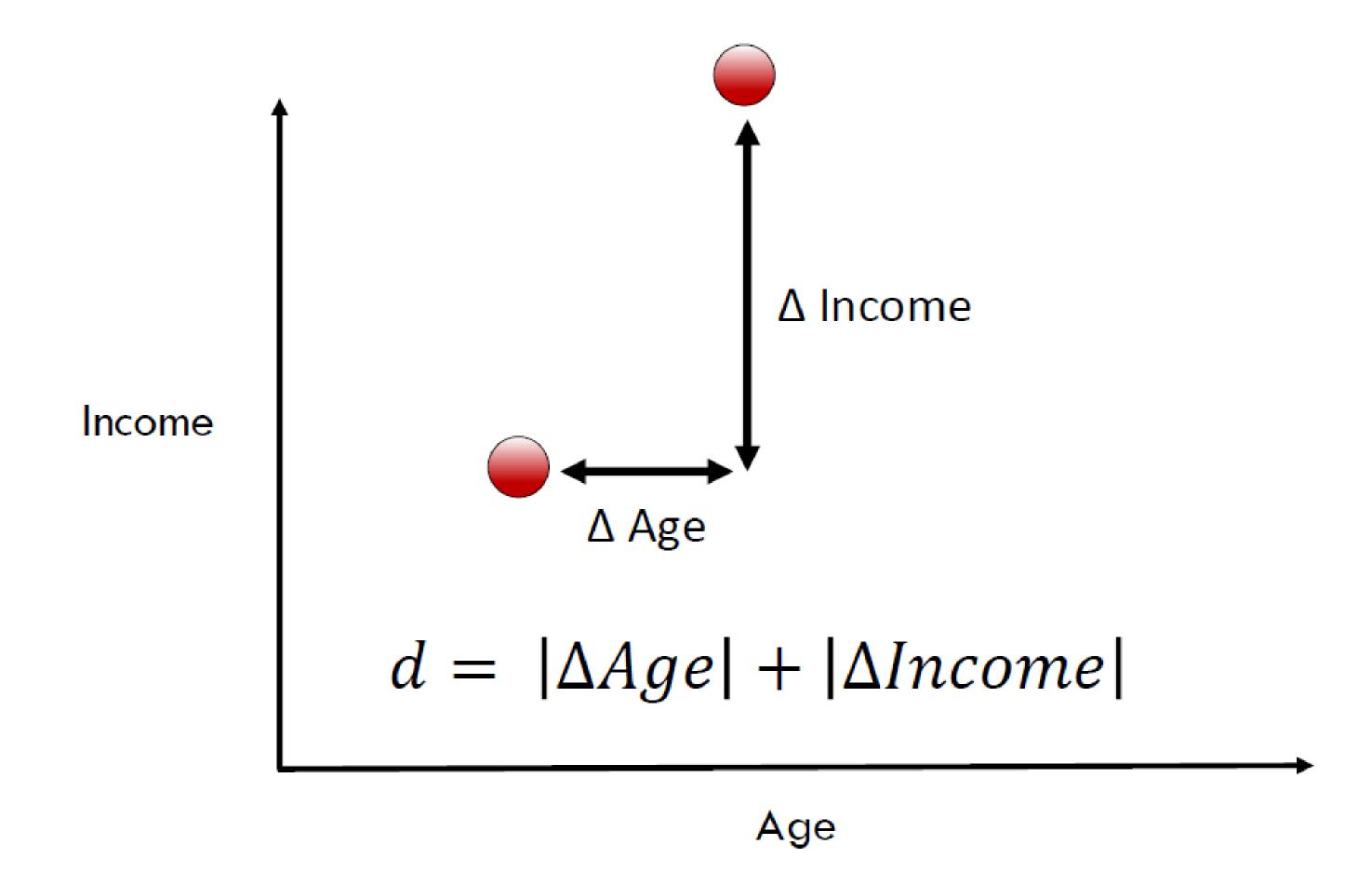


# Distance Metric

### Euclidean distance



### Manhattan distance

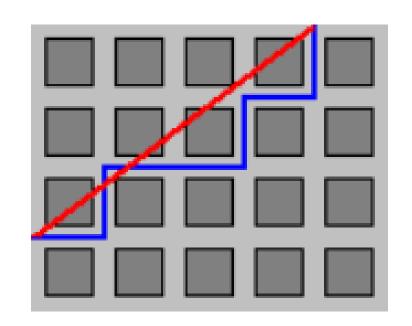


### p norm as generalization of Euclidean

- Numerical attribute: Euclidean distance
  - Pros: Symmetrical, Spherical

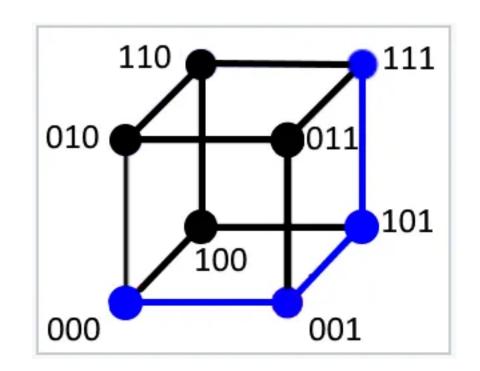
- $\mathbf{E} \mathcal{D}(x, x') = \sqrt{\sum_{i=1}^{a} (x_i x'_i)^2}$
- Cons: Sensitive to extreme value of any single attribute
- Minkowski distance (p-norm)
  - •P=1 Manhattan, p=2 Euclidean

$$\mathcal{D}(x, x') = \sqrt[p]{\sum_{i=1}^{d} |x_i - x_i'|^p}$$

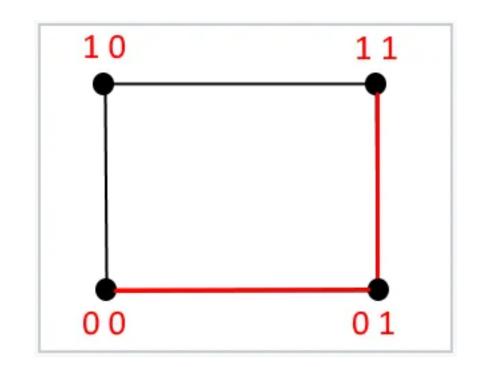


### Default choice of distance

- Categorical attribute: Hamming distance
  - Number of attributes where x, x' differ
  - Best for categorical attributes



$$\mathcal{D}(x, x') = \sum_{i=1}^{d} |x_i - x'_i|$$



$$\mathcal{D}(x, x') = \sum_{i=1}^{a} \mathbb{I}_{x_i = x'_i}$$

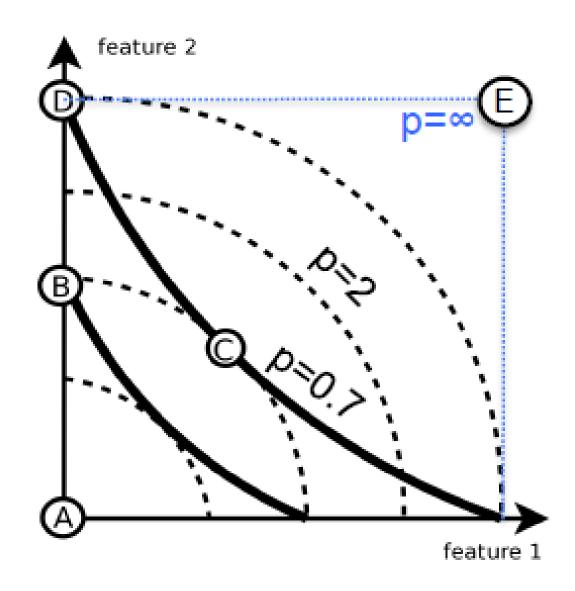
# Special case of One Hot encoded feature

Category	F1	F2	F3
Low	0	0	1
Medium	0	1	0
High	1	0	0

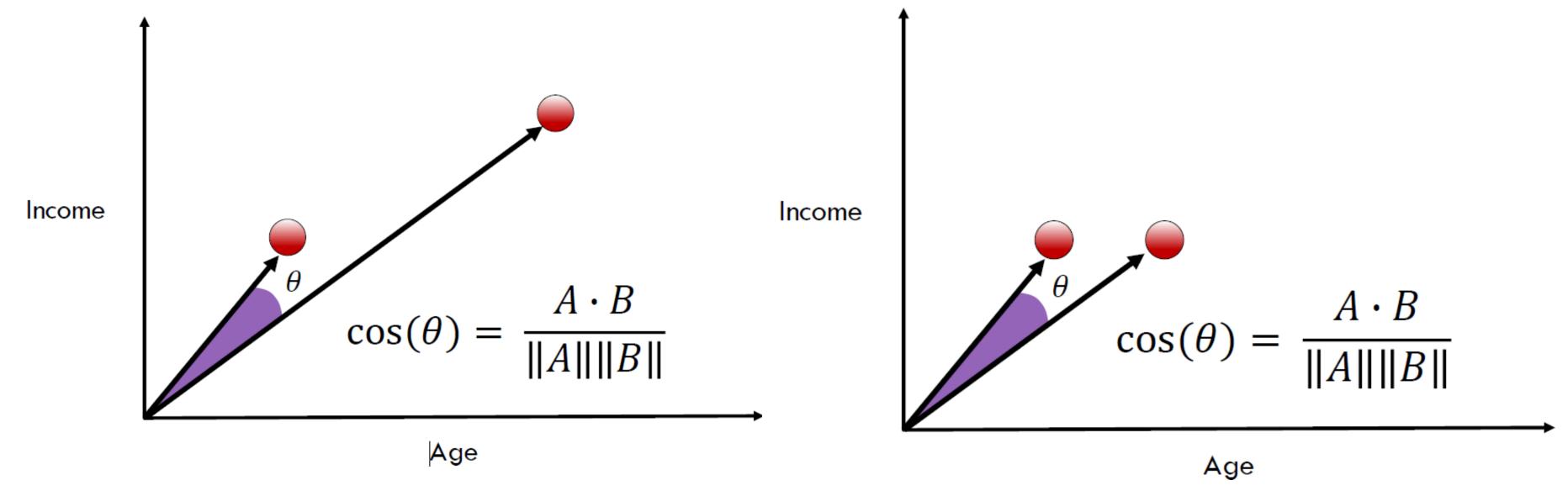
### p norm for different p

- Minkowski distance (p-norm) p=0
  - When x not equal to x'
  - When x equal to x'
  - Behaves like Hamming distance
- P=infinity  $\mathcal{D}(x,x') = max_d|x_i x_i'|$
- Different p

$$\mathcal{D}(x, x') = \sqrt[p]{\sum_{i=1}^{d} |x_i - x_i'|^p}$$



# Cosine similarity & Cosine distance



- Euclidean distance does not matter
- Angular separation matters (e.g. NLP)
- A theta decreases Cosine similarity increases
- •Cosine distance  $=1-cos\theta$

### Jaccard similarity & Jaccard distance

Applies to set similarity, community similarity

Sentence A: "I like chocolate ice cream."

set  $A = \{I, like, chocolate, ice, cream\}$ 

Sentence B: "Do I want chocolate cream or vanilla cream?"

set  $B = \{Do, I, want, chocolate, cream, or, vanilla\}$ 

- •Jaccard similarity  $= \frac{A \cap B}{A \cup B} = \frac{3}{9}$
- •Jaccard distance  $=1-\frac{A\cap B}{A\cup B}=1-\frac{3}{9}=\frac{6}{9}$

### Custom distance metrics

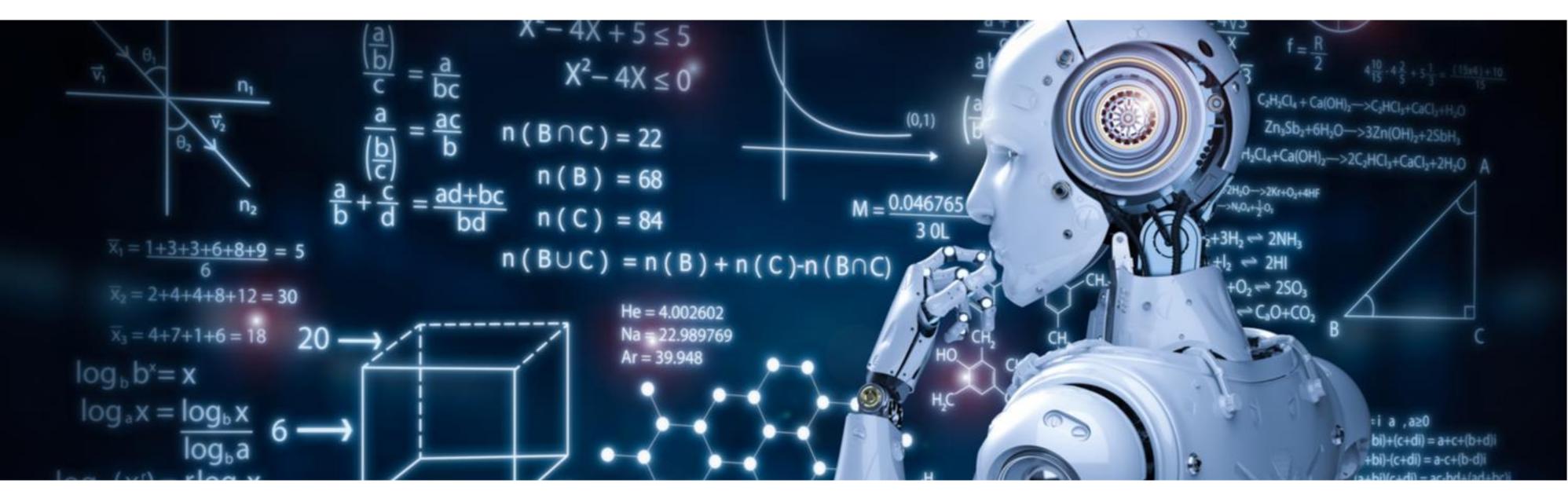
- •Always non negative  $||x_1-x_2||>=0$
- $||x_1 x_2|| = 0 \iff x_1 = x_2$
- $\|x_1 x_2\| = \|x_2 x_1\|$
- •Triangle inequality ||x+y|| <= ||x|| + ||y||

Intuitively true:
Sides of triangle

Should satisfy all of the above (generally speaking)

### Distance metrics in sklearn

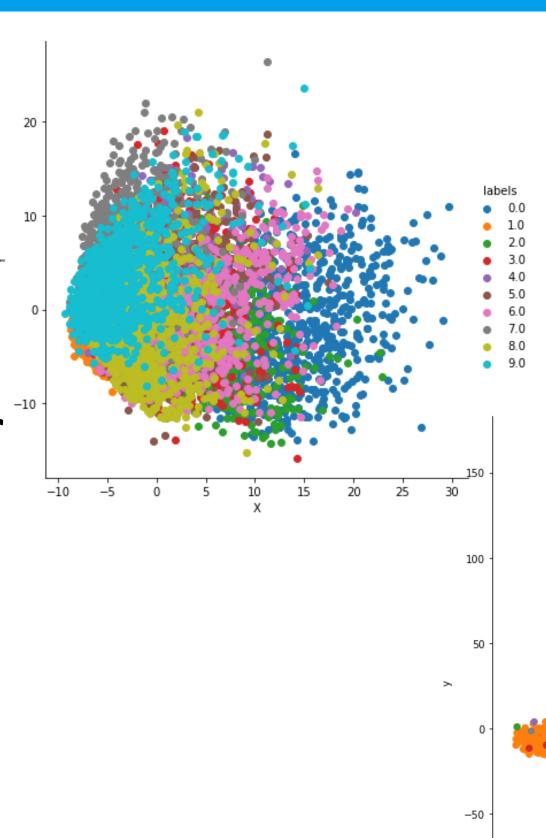
- •from sklearn.metrics import pairwise\_distances
- •dist = pairwise\_distances(X, X', metric="")
- •metric = euclidean, manhattan, cosine, jaccard

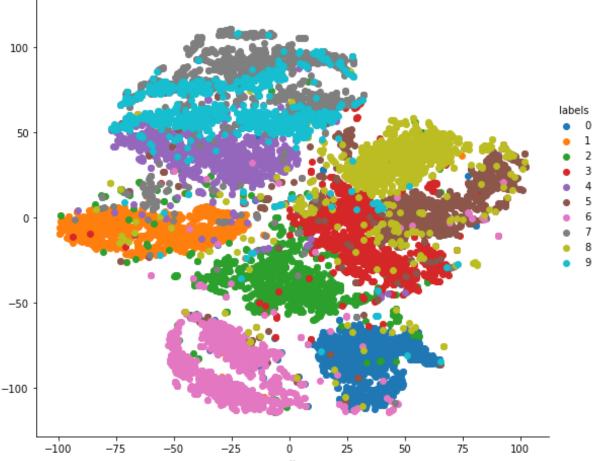


# Dimensionality Reduction + Clustering for visualization (optional)

## Dimensionality reduction

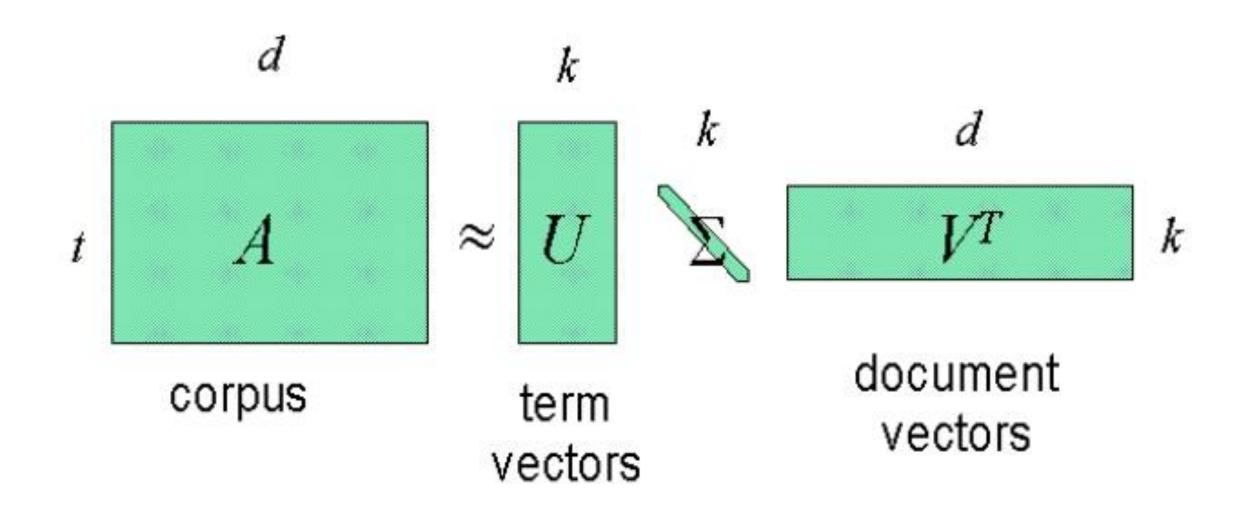
- Apply dimensionality reduction
  - PCA
  - •T-SNE
- •Then clustering in 2-D or 3-D
- E.g. PCA and t-SNE with MNIST for 2D clustering
  - Notice t-SNE gives better cluster
     separation





# Dimensionality reduction

- Truncated SVD
  - Sparse Matrix cases (such as TF-IDF)



# Clustering after dim reduction

- Truncated SVD
  - Sparse Matrix cases (such as TF-IDF)

